

MANUAL

Polyrate 17-C

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Executive summary:

Polyrate is a computer program for the calculation of chemical reaction rates of polyatomic species (and also atoms and diatoms as special cases) by variational transition state theory (VTST); conventional transition state theory is also supported. Bimolecular and unimolecular reactions and gas-phase, solid-state, and gas-solid-interface reactions are all included. *Polyrate* can perform variational transition state theory (VTST) calculations on reactions with both tight and loose transition states. For tight transition states it uses reaction-path (RP) variational transition state theory of Garret and Truhlar, and for loose transition states it uses variable-reaction-coordinate (VRC) variational transition state theory of Georgievskii and Klippenstein. For RP calculations there are options to use curvilinear coordinates along a minimum energy path (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations;

Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results) or a variationally optimized reaction path and to add multidimensional tunneling contributions by means of a transmission coefficient; the treatment of loose transition states is based on variable-reaction-coordinate VTST with single-faceted and multifaceted dividing surfaces.

The methods used for tight transition states are conventional transition state theory, canonical variational transition state theory (CVT), and microcanonical variational transition state theory (μ VT) with multidimensional semiclassical approximations for tunneling and nonclassical reflection. The tunneling approximations available are zero-curvature tunneling (ZCT), small-curvature tunneling (SSCT), large-curvature-tunneling (LCT), and optimized multidimensional tunneling (OMT). The SCT option is the centrifugal dominant semiclassical adiabatic ground-state tunneling, and the LCT options include both LC3 and LC4 including tunneling into excited states. One may also treat specific vibrational states of selected modes with translational, rotational, and other vibrational modes treated thermally.

Pressure-dependent rate constants for elementary reactions can be computed using system-specific quantum RRK theory (SS-QRRK) with the information obtained from high-pressure-limit VTST calculation as input by using the *SS-QRRK* utility code. The *SS-QRRK* utility program is part of the *POLYRATE* distribution, and its usage is described in a separate manual.

For tight transition states, several options are available for reaction paths, vibrations transverse to the reaction path, and transition state dividing surfaces. Reaction paths may be calculated by the Euler steepest-descent, Euler stabilization, Page-McIver, or variational reaction path algorithms. Vibrations away from the reaction path may be defined by rectilinear, nonredundant curvilinear, or redundant curvilinear coordinates. Generalized-transition-state dividing surfaces may be defined on the basis of gradient directions or by the re-orientation of the dividing surface algorithm. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.) Vibrational frequencies may be scaled or unscaled (it is highly recommended to scale the frequencies along the reaction coordinate by a scale factor that is developed for reproducing accurate ZPE).

For loose transition states, rate constants may be calculated for canonical or microcanonical ensembles or energy and total angular momentum resolved microcanonical ensembles. For loose transition states in barrierless association reactions, single-faceted or multifaceted dividing surfaces based on a variable reaction coordinate are used.

Potential energy surfaces may be analytic functions evaluated by subroutines, or they may be implicit surfaces defined by electronic structure input files containing energies, gradients, and force constants (Hessians) at selected points on a reaction path. The use of electronic structure calculations to calculate rate constants or other dynamical quantities without an analytic potential energy surface is called direct dynamics.

Analytic surfaces may be used for variational transition state theory and any of the types of tunneling calculations, single-level and dual-level calculations based solely on electronic structure input files may be used for variational transition state theory and zero-curvature or small-curvature tunneling, and dual-level calculations based on using an analytic potential energy surface as the lower level and using an electronic structure input file as the higher level may be used for variational transition state theory and any of the types of tunneling calculations.

Polyrate supports six options for direct dynamics, namely (i) straight single-level direct dynamics, (ii) zero-order interpolated variational transition state theory (IVTST-0), (iii) first-order interpolated variational transition state theory (IVTST-1), (iv) interpolated variational transition state theory by mapping (IVTST-M), (v) variational transition state theory with interpolated single-point energies (VTST-ISPE), and (vi) variational transition state theory with interpolated optimized corrections (VTST-IOC). VTST-IOC and VTST-ISPE are examples of dual-level direct dynamics; and VTST-IOC is also called triple-slash dynamics (*///*), where the triple slash denotes higher-order corrections of the geometries at which energetic and Hessian corrections are calculated. Dual-level methods may be applied with electronic structure data for both levels (dual-level direct dynamics), or it may be applied with an analytical potential energy surface for the lower level and electronic structure data for the higher level. When dual-level methods are employed with electronic structure data for both levels, the lower level may be either straight direct dynamics or IVTST-M.

This version of *Polyrate* contains 112 test runs, and 46 of these are for direct dynamics calculations; 85 of the test runs are single-level runs, and 27 are dual-level calculations.

Polyrate is designed to be used in conjunction with interfaces to electronic structure calculations for direct dynamics. Currently the following interfaces are available, although some of them were created with older versions of *Polyrate*:

<u>Electronic structure package</u>	<u>Interface</u>
CHARMM with CHARMMRATE	CRATE
GAMESS (or with GAMESSPLUS)	GAMESSPLUSRATE
GAUSSIAN 16/GAUSSIAN 09	GAUSSRATE
JAGUAR	JAGUARATE
MC-TINKER	MC-TINKERATE
MORATE	MOPAC
MULTILEVEL	MULTILEVELRATE
NWCHEM	NWCHEMRATE

CONTENTS

TITLE PAGE AND EXECUTIVE SUMMARY 1

CONTENTS..... 4

MANUAL..... 1

PART I GENERAL INFORMATION..... 13

1. INTRODUCTION 13

2. REFERENCES 24

 2.A. RECOMMENDED CITATION AND OTHER *Polyrate* REFERENCES 24

 2.B. REFERENCES FOR INCORPORATED CODE 25

3. AVAILABILITY AND LICENSING 26

4. *POLYRATE* SPECIFICS 27

 4.A. *Polyrate* authors 27

 4.B. Hardware platforms 29

 4.C. Compiler versions 32

5. PROGRAM AND MANUAL DESCRIPTION 36

 5.A. Code description, portability, and special files 36

 5.B. Description of the manual 36

 5.B.1. Referencing conventions 38

 5.B.2. Book chapters 38

PART II THEORY BACKGROUND AND CALCULATION DETAILS ON RP-VTST.... 39

6. THEORETICAL BACKGROUND ON RP-VTST 39

 6.A. Basic theory 39

 6.B. Vibrations, Rotations, and Translations 42

 6.C. Partition functions 45

 6.D. Tunneling methods 46

 6.E. Reactant and product properties 48

 6.F. Saddle point properties 50

 6.G. Reaction path 51

 6.G.1. SADDLE option 52

 6.G.2. NOSADDLE option 55

 6.H. Reaction-path curvature 55

6.I. Transmission coefficients.....	57
6.J. Reaction rates including tunneling.....	59
6.K. Imaginary-frequency modes transverse to the reaction path	61
6.L. Equilibrium constant calculation	62
6.M. Omitting the product	62
6.N. Optimized multidimensional tunneling calculations	63
6.N.1. COMT method	63
6.N.2. μ OMT method	63
6.O. Interpolated VTST and mapped interpolation	63
6.O.1. IVTST-0 and IVTST-1	64
6.O.2. Interpolated VTST by mapping (IVTST-M)	65
6.O.3. The mapped interpolation (MI) algorithm with the hooks	67
6.P. Interpolated optimized corrections (VTST-IOC), a form of dual-level VTST	67
6.Q. Interpolated algorithm for large-curvature tunneling calculations	69
7. PROGRAM STRUCTURE	70
7.A. Overview of program flow	70
7.B. Determining the sign of s	76
7.C. Restart options	77
8. CALCULATIONS USING THE HOOKS.....	79
8.A. Methods for interpolating beyond SLM and SLP with the hooks	79
8.B. Zero of energy.....	79
8.C. User-Supplied analytic potential energy surface	80
8.C.1. Interface	80
8.C.2. Utility routines	83
8.D. Calculations using an electronic structure program through hooks	86
9. CALCULATIONS BASED ON AN ELECTRONIC STRUCTURE INPUT FILE	91
9.A. Relation of <i>Polyrate</i> input to electronic structure package output	93
9.B. Orientation of the molecule at generalized transition states	94
9.C. Methods for interpolation of electronic structure data with files fu30 and fu40	95
9.D. Methods for interpolation of electronic structure data with file fu31	97
10. SELECTED DETAILS OF THE CALCULATIONS	98
10.A. Extrapolation of the reaction-path data beyond the range of the original grid	98
10.B. Options for anharmonicity	100
10.B.1. WKB anharmonicity based on a quadratic-quartic fit	101
10.B.2. WKB anharmonicity based on the true potential	102
10.B.3. Hindered rotor treatments	103

10.C. Alternate method for the calculation of numerical second derivatives	108
10.D. Reaction-path following methods.....	109
10.E. Interpolation of small-curvature effective reduced mass	111
10.F. Reaction-path curvature calculation.....	111
10.G. Solid-state and gas-solid interface reactions	112
10.H. VTST-IOC calculation, including ICR, ICL, and ICA schemes	112
10.I. VTST-ISPE calculation.....	116
10.J. Alternative methods for mapped interpolation	119
10.K. Quantized-reactant-state tunneling calculation	120
10.L. Curvilinear generalized normal mode analysis.....	120
10.M. IVTST0FREQ option.....	122
10.N. LCG4 option.....	126
10.O. MS-VTST, MP-VTST, FMS-VTST, and FMR-VSTT calculations by using both the <i>POLYRATE</i> and <i>MSTOR</i> programs.....	128
<u>PART III THEORY BACKGROUND FOR VRC-VTST.....</u>	135
11. THEORETICAL BACKGROUND ON VRC-VTST	135
11.A. Variable reaction coordinate.....	135
11.B. Multifaceted dividing surface	141
11.C. Overview of program flow for VRC-VTST calculation.....	142
11.D. Implementation details of the <i>E</i> , <i>J</i> -resolved microcanonical VRC algorithm.....	142
<u>PART IV DESCRIPTION OF INPUT FILES.....</u>	145
12. DESCRIPTION OF INPUT FOR RP-VTST	145
12.A. General description of file fu5 input.....	145
12.A.1. Section descriptions	147
12.A.2. GENERAL section	148
12.A.3. ENERGETICS section.....	156
12.A.4. SECOND section	159
12.A.5. OPTIMIZATION section	162
12.A.6. REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, AND START sections	168
12.A.7. PATH section.....	188
12.A.8. TUNNEL section	217

12.A.9. RATE section.....	226
12.B. General description of file fu29 input.....	238
12.B.1. E1GEOM Section	239
12.B.2. E1GRAD Section.....	239
12.B.3. E1HESS, P1HESS, P2HESS, R1HESS, R2HESS, and TSHESS sections	239
12.B.4. IVTST Section	240
12.B.5. P1FREQ, P2FREQ, R1FREQ, R2FREQ, and TSFREQ sections	241
12.C. General description of file fu30 input.....	242
12.D. General description of file fu31 input.....	251
12.D.1. Section descriptions	251
12.D.2 GENERAL section	253
12.D.3. REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and SADDLE sections	257
12.E. General description of file fu40 input.....	260
12.E.1 Section Descriptions.....	260
12.E.2 GEN40 section	261
12.E.3 R140, R240, P140, P240, WR40, and WP40 sections	267
12.E.4 SADDLE40 section.....	270
12.E.5 POINT40 section.....	272
12.F. General description of unit fu50 input	277
12.G. General description of unit fu51 input.....	288
12.G.1. Section Descriptions	288
12.G.2 ISPEGEN section.....	289
12.G.3 POINT section	292
13. DESCRIPTION OF INPUT FOR VRC-VTST.....	294
13.A Section descriptions of input file fu5.....	294
13.A.1. GENERAL section	294
13.A.2. ENERGETICS section.....	297
13.A.3. OPTIMIZATION section	298
13.A.4. REACT1, REACT2, PROD1, AND START sections	299
13.A.5. PATH section.....	302
13.A.6. RATE section.....	303
14. GENERAL DESCRIPTION OF LCT FILES	304
<u>PART V MORE GENERAL PROGRAM PACKAGE INFORMATION.....</u>	305

15. INSTALLING AND USING <i>POLYRATE</i>	305
15.A. Instructions for a Unix environment.....	305
15.A.1. Installation	305
15.A.2. Testing	311
15.A.3. Using <i>POLYRATE</i>	313
15.B. Comments for expert users	316
15.C. Running in parallel using MPI.....	317
16. SAMPLE TEST RUNS.....	320
16.A. Chart of the potential energy functions and data files	320
16.B. Descriptions of sample test runs	320
16.C.1. Test run ch5j1tr1	324
16.C.2. Test run ch5j1tr2	326
16.C.3. Test run ch5j1tr3	327
16.C.4. Test run ch5j1tr4	328
16.C.5. Test run ch5j1tr5	329
16.C.6. Test run ch5j1tr6	330
16.C.7. Test run ch5j1tr7	331
16.C.8. Test run ch5j1tr8	332
16.C.9. Test run ch5j1tr9	333
16.C.10. Test run ch5j1tr10	334
16.C.11. Test run ch5j1tr11	335
16.C.12. Test run ch5j2tr1	336
16.C.13. Test run ch5j1tr12	337
16.C.15. Test run ch5j2tr3	339
16.C.16. Test run ch5j2itr1	340
16.C.17. Test run ch5j2itr2	341
16.C.18. Test run ch5fu29tr1	342
16.C.19. Test run ch5fu29tr2	343
16.C.20. Test run ch5fu30tr1	344
16.C.21. Test run ch5fu30tr2	345
16.C.22. Test run ch5fu30tr3	346
16.C.23. Test run ch5fu30tr4	347
16.C.24. Test run ch5fu30tr5	348
16.C.25. Test run ch5fu30tr6	349
16.C.26. Test run ch5fu31tr1	350
16.C.27. Test run ch5fu40tr1	351

16.C.28. Test run ch5fu40tr2	352
16.C.29. Test run ch5fu40tr3	353
16.C.30. Test run ch5fu40tr4	354
16.C.31. Test run ch5fu40tr5	355
16.C.32. Test run ch5fu40tr6	357
16.C.33. Test run ch5fu50tr1	358
16.C.34. Test run ch5fu50tr2	359
16.C.35. Test run ch5fu50tr3	360
16.C.36. Test run ch5fu51tr1	361
16.C.37. Test run ch5icfu30tr1	362
16.C.38. Test run ch5icfu30tr2	363
16.C.39. Test run ch5icfu31tr1	364
16.C.40. Test run ch5icfu40tr1	365
16.C.41. Test run ch5icfu40tr2	366
16.C.42. Test run clhbrtr1	367
16.C.43. Test run clhbrtr2	369
16.C.44. Test run clhbrtr3	370
16.C.45. Test run clhbrtr4	371
16.C.46. Test run clhbr1tr5	372
16.C.47. Test run cmctr1	373
16.C.48. Test run cmctr2	374
16.C.49. Test run cmctr3	375
16.C.50. Test run cwmctr1	376
16.C.51. Test run cwmctr2	377
16.C.52. Test run cwmctr3	378
16.C.53. Test run h3tr1	379
16.C.54. Test run h3tr2	380
16.C.55. Test run h3tr3	381
16.C.56. Test run h3tr4	382
16.C.57. Test run ho2tr1	383
16.C.58. Test run ho2tr2	384
16.C.59. Test run ho2tr3	385
16.C.60. Test run ho2tr4	386
16.C.61. Test run ho2tr5	387
16.C.62. Test run nh3tr1	388
16.C.63. Test run nh3tr2	389

16.C.64. Test run oh3tr1	390
16.C.65. Test run oh3tr2	391
16.C.66. Test run oh3tr3	392
16.C.67. Test run oh3tr4	393
16.C.68. Test run oh3tr5	394
16.C.69. Test run oh3tr6	395
16.C.70. Test run oh3tr7	396
16.C.71. Test run oh3tr8	397
16.C.72. Test run oh3tr9	398
16.C.73. Test run oh3tr10	399
16.C.74. Test run oh3fu30tr1	400
16.C.75. Test run oh3fu30tr2	401
16.C.76. Test run oh3fu30tr3	402
16.C.77. Test run oh3fu30tr4	403
16.C.78. Test run oh3fu30tr5	404
16.C.79. Test run oh3fu30tr6	405
16.C.80. Test run oh3fu31tr1	406
16.C.81. Test run oh3fu31tr2	407
16.C.82. Test run oh3fu40tr1	408
16.C.83. Test run oh3fu40tr2	409
16.C.84. Test run oh3fu40tr3	410
16.C.85. Test run oh3fu40tr4	411
16.C.86. Test run oh3fu40tr5	412
16.C.87. Test run oh3fu40tr6	413
16.C.88. Test run hnitr1	414
16.C.89. Test run h2nitr1	415
16.C.90. Test run ch4ohtr1	416
16.C.91. Test run ch4ohtr2	417
16.C.92. Test run ch4ohtr3	418
16.C.93. Test run ch4ohtr4	419
16.C.94. Test run ch4cltr1	420
16.C.95. Test run ch4cltr2	421
16.C.96. Test run ch4otr1	422
16.C.97. Test run ohcltr1	423
16.C.98. Test run ohcltr2	424
16.C.99. Test run ohcltr3	425

16.C.100. Test run ohcltr4	426
16.C.101. Test run ohcltr5	427
16.C.102. Test run ch3tr1	428
16.C.103. Test run h3o2fu31tr1	429
16.C.104. Test run h3o2fu31tr2.....	430
16.C.105. Test run h3o2fu31tr3.....	431
16.C.106. Test run h3o2fu31tr4.....	432
16.D. Summary of test runs	433
17. DESCRIPTION OF FILES IN RELEASE OF VERSION 17	435
17.A. FORTRAN files and subprograms, MODULE files, and PES subroutines.....	435
17.A.1. <i>Polyrate</i> subprograms	435
17.A.2 User-supplied subprograms and module files.....	475
17.A.3 FORTRAN files	475
17.B. Utility program detmi.f	477
17.C. Utility program hrotor.f	479
17.D. Utility programs findl.f and findb.f	482
17.E. Utility program rotate30.f, rotate31.f, and rotate40.f.....	482
17.F. Utility program <i>SS-QRRK</i>	485
18. UNIX C SHELL SCRIPTS AND PERL SCRIPTS	486
18.A. Installation script	486
18.B. Potential energy surface compiling scripts	487
18.C. Source code compiling script.....	487
18.D. Linking script.....	488
18.E. Test run scripts	488
19. FILE USAGE	490
20. MEMORY REQUIREMENTS	494
21. TEST RUNS AND TEST RUN TIMINGS	495
21.A. Extent of testing	495
21.B. Timings	495
22. CITATIONS INDICATED BY BRACKETS IN THE TEXT	500
23. BIBLIOGRAPHY	504
24. ERRATA	512
25. <u>APPENDIX: POLYRATE REVISION HISTORY</u>	513
25.B. <u>ASCII MANUAL REVISION HISTORY</u>.....	576

25.C.	<u>POSTSCRIPT AND PDF MANUALS</u>	576
25.D.	<u>INCOMPATIBILITIES OF <i>POLYRATE</i>–VERSION 2017 WITH PREVIOUS VERSIONS</u>	576
25.E.	<u>PROGRAM DISTRIBUTION HISTORY</u>	580
25.E.4.	<u>OTHER</u>	582
25.E.5.	<u>CURRENT</u>	582
26.	ACKNOWLEDGMENTS	583
27.	INDEX	584

Part I General Information

1. INTRODUCTION

Polyrate is a program for the analysis of reactants, products, and transition states of chemical reactions and for calculating variational transition state theory (VTST) rate constants and multidimensional semiclassical tunneling probabilities. The current version of the program performs two qualitatively different kinds of VTST calculations:

- RP-VTST: reaction–path VTST for arbitrary kinds of reactions with tight transition states.
- VRC-VTST: variable-reaction-coordinate VTST only for loose transition states of bimolecular association reactions.

and it also can calculate pressure-dependent rate constants by using system-specific quantum Ramsperger–Rice–Kassel theory:

- SS-QRRK: system-specific quantum RRK theory

RP-VTST and VRC-VTST are included in *POLYRATE* proper, and SS-QRRK is performed in a separate utility code that is included as part of the *POLYRATE* package. The *SS-QRRK* utility code is described in a separate manual, which is also included as part of the *Polyrate* package.

The following articles contain summaries of the technical background for the methods in *Polyrate*:

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comp.chem.umn.edu/Truhlar/docs/C58.pdf
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doi.org/10.1201/9781420028027.ch22
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In *Polyrate*, analysis of reactants and products can be performed for up to two reactants and two products for RP-VTST and for two reactants and one product for VRC-VTST. *Polyrate* can also calculate equilibrium constants, so one can calculate the reverse rate constant from the forward rate constant and the equilibrium constant. The RP-VTST method can be applied to gas-phase, solid-phase, and gas-solid-interface reactions. Generalized transition state theory calculations may be carried out using the harmonic approximation for vibrations with either rectilinear or curvilinear coordinates in RP-VTST calculations.

It is highly recommended to use curvilinear coordinates (via the "CURV2" or "CURV3" options) rather than Cartesian coordinates for generalized-normal mode analysis along the MEP in VTST calculations; Cartesian coordinates will often yield

unphysical imaginary frequencies along the reaction coordinate, and these will cause unphysical VTST results. One should always check in the long output that the generalized normal mode frequencies as functions of the reaction coordinate do not contain any imaginary frequencies; imaginary frequencies may lead to unreliable or wrong results.

A limited amount of anharmonicity may also be included. The RP-VTST rate constants can be computed using canonical variational theory (CVT), and microcanonical variational theory (μ VT). Conventional transition state theory (TST) may also be employed. VRC-VTST calculations currently only can be applied to gas-phase reactions using CVT, μ VT, and total energy and total angular momentum-resolved microcanonical variational theory, which for short may be called E,J -resolved microcanonical variational theory (E,J - μ VT).

For thermal or ground-state reactions using RP-VTST, tunneling probabilities can be computed using the zero-curvature tunneling method (ZCT), the small-curvature tunneling (SCT) method, the large-curvature tunneling (LCT) method, or an optimized multidimensional tunneling method (μ OMT or COMT). The SCT calculations are based on the centrifugal-dominant small-curvature semiclassical ground-state (CD-SCSAG) method, and the LCT calculations are based on the large-curvature ground-state method, version 3 or 4 (LCG3 or LCG4). For state-selected excited-state reactions only the ZCT and SCT methods are available in this version. In addition, for ground-state reactions of exoergic or thermoneutral reactions, LCT calculations may be summed over certain excited final states, and for endoergic reactions, LCT calculations may be summed over certain excited states of reactants. (These summations are a required part of the method when these contributions are not negligible). The recommended tunneling procedure when LCT calculations are carried out is the microcanonical optimized multidimensional tunneling method (μ OMT) in which, at each energy, the tunneling probability is approximated as the larger of the values calculated by the SCT and LCT methods. The option of carrying out a tunneling calculation is not available in conjunction with VRC-VTST option.

The *Polyrate* program computes rate constants by either single-level or dual-level approaches for RP-VTST calculations and by single-level approach for VRC-VTST calculations.

The single-level calculations and the lower level of dual-level calculations using RP-VTST may be based on either an analytic potential energy function, which is called the potential energy surface (PES) or potential energy function (PEF), or on various types of interpolations of electronic structure input data at selected points along the minimum energy path. VRC-VTST calculations can only be used for single-level calculations based on a global PES or direct dynamics interfaced with an electronic structure package.

When using a PES, the user must supply a subroutine, which, for a given set of Cartesian coordinates of each of the atoms, returns the potential energy and its first derivatives with respect to the Cartesian coordinates. This subroutine is called by a set of *Polyrate* interface routines called hooks. (In particular, the “hooks” consist of subprograms PREP, PREPJ, EHOOK, GHOOK, HHOOK, and OHOOK plus a few utilities, e.g., YTRANS and YNEWT, that are discussed in Section 8.C.). Note that VRC-VTST calculations don't need gradient information if optimized geometries and vibrational frequencies of reactants and/or product are provided in the input file. When using electronic structure input data without an analytic surface, the user must supply, as an input file to the program, the geometries, energies, and force constants or frequencies for the reactants, products, and saddle point, and optionally these quantities and the gradient at a sequence of points on the minimum energy path. These data are then used for dynamics calculations.

For RP-VTST calculations using dual-level approach, the data along the reaction path may be “corrected” using higher-level data supplied by the user at selected points either for reaction paths based on potential energy surfaces or for those based on electronic structure input files. A generic name for such dual-level RP-VTST methods is VTST-IOC. We currently recognize three variants of the IC methods, namely IOC (interpolated optimized corrections), IOE (interpolated optimized energies), and ISPE (interpolated single point energies). The IOC and IOE methods are also called triple-slash (///) VTST. In the IOC method, the frequencies, potential energies, and moment-of-inertia determinants along the reaction path that are calculated from the potential energy surface are “corrected” using the results of higher-level electronic structure calculations (supplied by the user) at selected points. The O in IOC indicates that any corrections at stationary points are based on geometries optimized at the higher level. If corrections were made at nonstationary points, they would be based on reaction paths calculated at the higher level. The IOE approach is a subcase of IOC in which only the energies and moment-of-inertia determinants are corrected, but the corrected energies are based on geometries optimized

at the higher level. The final dual-level method, called VTST-ISPE, corrects only the energies along the reaction path using higher-level electronic structure calculations (supplied by the user) at the stationary points and selected points along the path. This kind of calculation is called a double-slash (//) calculation. In general the acronyms VTST-IOC and VTST-IOE or the triple slash notation is reserved for cases when the correction involves a higher-level geometry optimization of the transition state. The VTST-ISPE notation refers to using higher-level single-point energies at the lower-level saddle point or at the lower-level saddle point plus selected points along the lower-level reaction path. Thus VTST-IOE is expected to be more accurate than VTST-ISPE, even though one neglects the corrections to the frequencies.

In addition to the straight VTST method and to dual-level VTST-IOC and VTST-ISPE, the code also has single-level IVTST options for RP-VTST calculations. With the IVTST philosophy the user supplies the geometry, energy, and frequencies or Hessian at the stationary points and additionally at as many points along the path as are computationally feasible. *POLYRATE* has the functionality to perform VTST calculations with information at the stationary points and no points along the reaction path, called IVTST-0, with one point along the path on either side of the saddle point, called IVTST-1, or with an arbitrary number of points along the path on either side of the saddle point (as long as there is at least one nonstationary point on either side of the saddle point). The latter option is carried out by the mapped interpolation (MI) algorithm and is called IVTST-M. In the current version of *POLYRATE*, IVTST-M is available with only the default IVTST-M options when electronic structure input data is read from file fu30 or fu40, and it is available with all the IVTST-M options for files fu31 and the hooks. Note that IVTST-M can also be used as the lower level of a dual-level calculation, but in the present version of *POLYRATE* IVTST-0 and IVTST-1 cannot be used that way.

In addition to the use of the MI algorithm as the central element of the IVTST-M method with electronic structure input files, since version 8.0 of *POLYRATE* we have allowed the use of the MI algorithm as an "extrapolation" algorithm when using the hooks for a calculation with a saddle point. If the hooks are used to interface with an electronic structure calculation, this provides an alternative way to perform IVTST-M direct dynamics calculations. If the hooks are used with an analytic potential, this replaces the previously used method for extrapolating the adiabatic potential and effective masses beyond the edge of the originally calculated grid. Finally we note that the energy part of the MI algorithm is used in the VTST-ISPE version of VTST-IC.

There is also an option, called the RESTART option, in which the input consists in part of reaction-path information calculated in a previous run of the program. The RESTART option is not available for dual-level calculations.

Derivative versions of *POLYRATE*, called CHARMMRATE, GAMESSPLUSRATE, GAUSSRATE, JAGUARATE, MC-TINKERATE, MORATE, MULTILEVELRATE, and NWCHEMRATE, which compute rate constants from potentials, gradients, and Hessians calculated by electronic structure methods, are also available.

CHARMMRATE is a module of the CHARMM program; CHARMMRATE can compute enzyme-catalyzed rates by ensemble-averaged VTST. An interface to the CHARMMRATE module of the CHARMM package based on *POLYRATE* version 9.0, called CRATE, is available as a separate distribution.

An interface to the GAMESSPLUS (or GAMESS) electronic structure packages based on *POLYRATE* version 9.3, called GAMESSPLUSRATE, is available as a separate distribution.

An interface to the GAUSSIAN16 (or GAUSSIAN98 or GAUSSIAN03 or GAUSSIAN09) electronic structure packages based on *POLYRATE* version 2017, called GAUSSRATE, is available as a separate distribution.

An interface to the JAGUAR package based on *POLYRATE* version 9.6, called JAGUARATE, is available as a separate distribution.

An interface to the MC-TINKER package based on *POLYRATE* version 9.1, called MC-TINKERATE, is available as a separate distribution.

MORATE uses the semiempirical molecular orbital package MOPAC instead of an analytic potential energy function. The electronic structure input file option and MORATE are both examples of direct dynamics, i.e., rate or dynamics calculations based directly on electronic structure information without the intermediary of an analytic fit to the potential energy surface. The MORATE program is not distributed as part of the *POLYRATE* package, but it is available as a separate distribution from the University of Minnesota. At the time of release of *POLYRATE*—version 2016, the latest version of MORATE is based on *POLYRATE*—version 8.5.1.

An interface to the MULTILEVEL package based on *POLYRATE* version 9.1, called MULTILEVELRATE, is available as a separate distribution.

An interface to the NWCHEM package based on *POLYRATE* version 9.7, called NWCHEMRATE, is available as a separate distribution.

AMSOLRATE is an interface with the AMSOL electronic structure package. AMSOLRATE can also calculate rate constants for reactions in liquid-phase solutions. AMSOLRATE can also be used for gas-phase calculations with semiempirical molecular orbital theory, but we recommend MORATE for that purpose because it produces less noisy frequencies. Therefore we do not mention AMSOLRATE in most lists of available interface programs.

The various direct dynamics methods obtain their input about the system in different ways:

- Single-level direct dynamics calculations may be carried out using file fu29, file fu30, file fu31, file fu40, or file fu50 for the electronic structure information (RP-VTST).
- VTST-IOC dual-level direct dynamics calculations may be carried out using file fu30, fu31, or file fu40 for the lower-level information and file fu50 for the higher-level information (RP-VTST).
- VTST-ISPE dual-level direct dynamics calculations may be carried out using file fu30, fu31, or file fu40 for the lower-level information and file fu51 for the higher-level information (RP-VTST).
- Single-level calculations based on an analytic potential energy surface obtain the potential surface information from subprograms SETUP and SURF, which are called by the hooks (RP-VTST and VRC-VTST).
- VTST-IOC dual-level calculations based on an analytic potential energy surface for the lower level obtain the potential surface information from subprograms SETUP and SURF (called by the hooks) and they obtain the higher-level electronic structure information from file fu50 (RP-VTST).
- VTST-ISPE dual-level calculations based on an analytic potential energy surface for the lower level obtain the potential surface information from subprograms SETUP and SURF (called by the hooks) and they obtain the higher-level electronic structure information from file fu51 (RP-VTST).

We note that files fu30, fu31, and fu40 accomplish essentially the same purpose. There are two differences between fu30 and fu40 is that file fu30 requires formatted data, whereas file fu40 is based on keyword input. The second difference is that there is a WRITEFU30 keyword, but again there is no WRITEFU40 keyword. File fu31 is also based on keyword input, but it differs from file fu40 in that certain IVTST-M options are supported with fu31 but not with fu40. A second difference is that there is a WRITEFU31 keyword, but there is no WRITEFU40 keyword. See also Sect. 11.D.1. At some point in

the future we may stop supporting fu30, but for now we will keep it for the convenience of users of previous versions. New users are encouraged to ignore fu30 and to use fu31 instead.

Throughout this manual it will be useful to keep in mind that there are really several types of single-level VTST calculations, namely

1. Straight VTST
 - 1.A. Straight VTST through the hooks
 - 1.B. Straight VTST with file fu30 or file fu40
2. IVTST-M
 - 2.A. IVTST-M with file fu30 or fu40
 - 2.B. IVTST-M with file fu31
3. IVTST-0 or IVTST-1.

Note that RP-VTST calculations can be performed using any of these options, but VRC-VTST calculations can only be performed using option 1.A. Calculations of type 1.A, 1.B, 2.A, and 2.B have many points of similarity, and in fact once *Polyrate* has all the save grid arrays filled with data, these types of runs become identical. Further the MI algorithm that constitutes the central element of IVTST-M calculations (2.A or 2.B) can also be used in straight VTST (1.A or 1.B), but for straight VTST is considered as an algorithm for extending the save grid to improve numerical convergence rather than being considered as the main ingredient of the method. However runs of type 3 do not use the save grid. In this manual we try to alert the reader whether a statement is true for one or another of these types of calculations, but not for all four types. However, the reader should be aware that sometimes we fail to mention this because it is assumed clear from the context or because the manual needs more work.

Dual-level calculations can be performed with type 1.A, 1.B, 2.A, or 2.B calculations as the lower level.

The main options for RP-VTST calculations are summarized in the following table:

Dynamics approach	hooks	fu29	fu30, fu31, or fu40	fu50
TST	TST	TST	TST	TST
VTST	TST/ZCT CVT CVT/ZCT CVT/SCT CVT/LCT μ VT μ VT/ZCT μ VT/SCT μ VT/LCT		TST/ZCT CVT CVT/ZCT CVT/SCT μ VT μ VT/ZCT μ VT/SCT	
IVTST-0		TST TST/ZCT		TST TST/ZCT
IVTST-1		TST TST/ZCT CVT CVT/ZCT CVT/SCT		
	hooks plus fu50		fu30, fu31, or fu40 plus fu50	
VTST-IOC	TST TST/ZCT CVT CVT/ZCT CVT/SCT CVT/LCT		TST TST/ZCT CVT CVT/ZCT CVT/SCT	

Table continued:

Dynamics approach	hooks plus fu50		fu30, fu31, or fu40 plus fu50	
VTST-IOC	μ VT μ VT/ZCT μ VT/SCT μ VT/LCT		μ VT μ VT/ZCT μ VT/SCT	
	hooks plus fu51		fu30, fu31, or fu40 plus fu51	
VTST-ISPE	μ VT μ VT/ZCT μ VT/SCT		μ VT μ VT/ZCT μ VT/SCT	

Note that whenever both SCT and LCT methods are available, then the μ OMT method is available as well.

Also note: When an analytical potential is used, the hooks call SETUP and SURF. However, the hooks capability is more general, and the hooks can be used to interface *POLYRATE* with electronic structure codes for direct dynamics; examples are the CRATE, GAMESSPLUSRATE, GAUSSRATE, JAGUARATE, MC-TINKERATE, MORATE, MULTILEVELRATE, and NWCHMERATE programs, which are interfaces to CHARMM with the CHARMMRATE module, GAMESSPLUS, GAUSSIAN, JAGUAR, MC-TINKER, MOPAC, MULTILEVEL, and NWCHEM, respectively.

2. REFERENCES

2.A. RECOMMENDED CITATION AND OTHER *Polyrate* REFERENCES

The recommended CITATION for the current version of the *Polyrate* code is

J. Zheng, J. L. Bao, R. Meana-Pañeda, S. Zhang, B. J. Lynch, J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, A. Ratkiewicz, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar, *Polyrate*—version 2017, University of Minnesota, Minneapolis, 2018.

References for methods used in the code are given in Section 23 of this manual.

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2.B. REFERENCES FOR INCORPORATED CODE

1. The code for evaluating the Cartesian **B** matrix and **C** tensor for stretches, nondegenerate bends, and torsions for the general polyatomic case is from J. Cioslowski and M. Challacombe, ANHTRAX, QCPE Program No. 609 (Quantum Chemistry Program Exchange, Bloomington, IN, 1991). (used with permission)
2. The EF code for stationary point optimization is from J. Barker, F. Jensen, H. S. Rzepa, A. Stebbings, LOCATION OF TRANSITION STATES IN AMPAC AND MOPAC USING EIGENVECTOR FOLLOWING (EF), QCPE Program No. 597 (Quantum Chemistry Program Exchange, Bloomington, IN, 1990). (used with permission)

3. AVAILABILITY AND LICENSING

All versions are copyrighted.

Versions 1.6 and later are licensed. Licenses are available from authorized distributors. Currently the only authorized distributors are the University of Minnesota and CPC Program Library; however the CPC Program Library currently has only old versions.

All license fees are waived, but the license must be signed. Copies of the license and instructions for obtaining the code are available at the University of Minnesota World Wide Web site, <http://comp.chem.umn.edu/Polyrate>.

Polyrate distribution at the University of Minnesota is currently handled by

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4. POLYRATE SPECIFICS

4.A. Polyrate authors

Versions 1.0 - 1.1:	A. D. Isaacson, D. G. Truhlar, S. N. Rai, R. Steckler, G. C. Hancock, B. C. Garrett, and M. J. Redmon
Versions 1.5–1.5.2:	A. D. Isaacson, D. G. Truhlar, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. N. Truong, T. Joseph, B. C. Garrett, and R. Steckler
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Versions 1.7–1.7.1:	A. D. Isaacson, D. G. Truhlar, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. N. Truong, T. Joseph, D.-h. Lu, V. S. Melissas, B. C. Garrett, and R. Steckler
Versions 2.0–2.5.2:	D.-h. Lu, T. N. Truong, B. C. Garrett, R. Steckler, A. D. Isaacson, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, V. S. Melissas, and D. G. Truhlar
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Versions 3.0.2–4.5.1:	D.-h. Lu, T. N. Truong, V. S. Melissas, G. C. Lynch, Y.-P. Liu, B. C. Garrett, R. Steckler, A. D. Isaacson, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, and D. G. Truhlar
Versions 5.0–5.1:	Y.-P. Liu, G. C. Lynch, W.-P. Hu, V. S. Melissas, R. Steckler, B. C. Garrett, D.-h. Lu, T. N. Truong, A. D. Isaacson, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, and D. G. Truhlar
Versions 6.0–6.5:	R. Steckler, W.-P. Hu, Y.-P. Liu, G. C. Lynch, B. C. Garrett, A. D. Isaacson, D.-h. Lu, V. S. Melissas, T. N. Truong, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, and D. G. Truhlar
Version 6.6:	R. Steckler, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, B. C. Garrett, A. D. Isaacson, D.-h. Lu, V. S. Melissas, T. N. Truong, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, and D. G. Truhlar
Version 6.6.1:	R. Steckler, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, Y.-Y. Chuang, B. C. Garrett, A. D. Isaacson, D.-h. Lu, V. S. Melissas, T. N. Truong, S. N. Rai, G. C. Hancock, J. G. Lauderdale, T. Joseph, and D. G. Truhlar

- Versions 7.0–7.1: R. Steckler, Y.-Y. Chuang, E. L. Coitiño, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, P. Fast, S. Clayton, V. S. Melissas, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 7.1.1–7.2: R. Steckler, Y.-Y. Chuang, E. L. Coitiño, P. L. Fast, J. C. Corchado, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, S. Clayton, V. S. Melissas, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 7.3–7.4.3: R. Steckler, Y.-Y. Chuang, P. L. Fast, E. L. Coitiño, J. C. Corchado, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, S. Clayton, V. S. Melissas, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 7.8–7.9.1: J. C. Corchado, Y.-Y. Chuang, P. L. Fast, Jordi Villà, E. L. Coitiño, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, S. Clayton, V. S. Melissas, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Version 8.0: Y.-Y. Chuang, J. C. Corchado, P. L. Fast, Jordi Villà, E. L. Coitiño, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, M. Z. Gu, I. Rossi, S. Clayton, V. S. Melissas, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 8.1–8.5.1: Y.-Y. Chuang, J. C. Corchado, P. L. Fast, Jordi Villà, W.-P. Hu, Y.-P. Liu, G. C. Lynch, C. F. Jackels, K. A. Nguyen, M. Z. Gu, I. Rossi, E. L. Coitiño, S. Clayton, V. S. Melissas, B. L. Lynch, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 8.6–8.7.2: J. C. Corchado, Y.-Y. Chuang, P. L. Fast, Jordi Villà, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, V. S. Melissas, B. J. Lynch, I. Rossi, E. L. Coitiño, A. Fernandez-Ramos, J. Pu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 8.7.3–9.3: J. C. Corchado, Y.-Y. Chuang, P. L. Fast, Jordi Villà, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, V. S. Melissas, B. J. Lynch, I. Rossi, E. L. Coitiño, A. Fernandez-Ramos, J. Pu, T. V. Albu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 9.3.1–9.5: J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, B. J. Lynch, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 9.6–9.7: J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, B. J. Lynch, J. Zheng, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar

- Version 2008: The authors of this version are J. Zheng, S. Zhang, B. J. Lynch, J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Version 2010: The authors of this version are J. Zheng, S. Zhang, B. J. Lynch, J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Version 2016: The authors of this version are J. Zheng, J. L. Bao, R. Meana-Pañeda, S. Zhang, B. J. Lynch, J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, A. Ratkiewicz, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar
- Versions 2017 to 2017-C: The authors of these versions are J. Zheng, J. L. Bao, R. Meana-Pañeda, S. Zhang, B. J. Lynch, J. C. Corchado, Y.-Y. Chuang, P. L. Fast, W.-P. Hu, Y.-P. Liu, G. C. Lynch, K. A. Nguyen, C. F. Jackels, A. Fernandez Ramos, B. A. Ellingson, V. S. Melissas, J. Villà, I. Rossi, E. L. Coitiño, J. Pu, T. V. Albu, A. Ratkiewicz, R. Steckler, B. C. Garrett, A. D. Isaacson, and D. G. Truhlar

4.B. Hardware platforms

- Versions 1.0–1.5: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS)
- Versions 1.6–1.7: VAX 11/780 (VMS 4.3), Cray-2 and Cray X-MP (both UNICOS)
- Version 2.0: VAX 11/780 (VMS 4.3), Cray-2 and Cray X-MP (both UNICOS 5.0), Cray Y-MP (UNICOS 5.1.8)
- Version 2.1: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS 5.1.8), Cray X-MP (UNICOS 5.1)
- Version 2.5: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS 5.1.9), Cray X-MP (UNICOS 5.1), Cray Y-MP (UNICOS 5.1.10), IBM 3090 600J (AIX), IBM RS/6000 model 530 (AIX), Silicon Graphics IRIS-4D/320GTX (IRIX System V.3), Sun SparcStation 4/360 (SunOS 4.0.3)
- Version 2.6: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS 5.1.9), Cray X-MP-EA (UNICOS 5.1.9), IBM 3090 600J (AIX), Silicon Graphics IRIS-4D/310GTXB (IRIX System V 3.3.1)

Version 2.7: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS 5.1.9), Cray Y-MP (UNICOS 5.1.9), Cray X-MP-EA (UNICOS 5.1.9), IBM 3090 600J (AIX), Silicon Graphics IRIS-4D/310GTXB (IRIX System V 3.3.1)

Version 3.0: VAX 11/780 (VMS 4.3), Cray-2 (UNICOS 5.1.9), Cray X-MP-EA (UNICOS 5.1.9), IBM 3090 600J (AIX), Silicon Graphics IRIS-4D/310GTXB (IRIX System V 3.3.1), Sun SparcStation 4/360 (SunOS 4.1.1)

Versions 3.0.7, 3.9.1, 3.9.2: Cray-2 (UNICOS 5.1.9), Cray X-MP-EA (UNICOS 5.1.9)

Version 4.0: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), Cray Y-MP (UNICOS 6.1.5a), IBM RS/6000 model 550 (AIX 3.1.5), Silicon Graphics IRIS-4D/310GTXB (IRIX System 4.0.1), Sun SparcStation 4/360 (SunOS 4.0.3), VAX 6000-510 (VMS 5.4), VAX 11/780 (VMS 4.3)

Version 4.0.1: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), Cray Y-MP (UNICOS 6.1.5a), IBM RS/6000 model 550 (AIX 3.1.5), Silicon Graphics IRIS-4D/440VGX, IRIS-4D/310GTXB, IRIS-4D/320VGX, IRIS-4D/210VGX, IRIS-4D/35VGX, IRIS/INDIGO (all IRIS computers use IRIX 4.0.1 operating system), Sun SparcStation 4/360 (SunOS 4.0.3), VAX 6000-510 (VMS 5.4), VAX 11/780 (VMS 4.3)

Version 4.1: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), IBM RS/6000 model 550 (AIX 3.1.5)

Version 4.2: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), Cray Y-MP (UNICOS 6.1.5a), IBM RS/6000 model 550 (AIX 3.1.5), Silicon Graphics IRIS-4D/310GTXB (IRIX 4.0.1), Sun SparcStation 4/360 (SunOS 4.0.3)

Version 4.3: IBM RS/6000 model 550 (AIX 3.1.5)

Version 4.4: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), Cray Y-MP (UNICOS 6.1.5a), IBM RS/6000 model 550 (AIX 3.2), Silicon Graphics IRIS-4D/310GTXB (IRIX 4.0.1), Sun SparcStation 4/360 (SunOS 4.0.3)

Version 4.5: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), IBM RS/6000 model 550 (AIX 3.2)

Version 5.0: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 6.1), Cray Y-MP (UNICOS 6.1.7), Cray M92 (UNICOS 6.1), IBM RS/6000 model 550 (AIX 3.2), Silicon Graphics IRIS-4D/310GTXB (IRIX 4.0.1), Sun SparcStation IPX (SunOS 4.1.2), DEC 3000/500 AXP (OSF/1)

Version 5.0.1: Cray-2 (UNICOS 6.1), Cray X-MP-EA (UNICOS 7.C.2), Cray C90 (UNICOS 7.C.2), IBM RS/6000 model 550 (AIX 3.2), DEC 3000/500 AXP (OSF/1), HP 730 and HP 735 (HP-UX 9.01)

- Version 5.1: Cray-2 (UNICOS 7.0.5), Cray X-MP-EA (UNICOS 7.C.2), Cray Y-MP (UNICOS 7.C.3), Cray C90 (UNICOS 7.C.2), IBM RS/6000 model 550 (AIX 3.2), Silicon Graphics IRIS-4D/310GTXB (IRIX 4.0.5), Sun SparcStation IPX (SunOS 4.1.2), HP 730 and HP 735 (HP-UX 9.01)
- Version 6.0: Cray-2 (UNICOS 7.0.5), Cray X-MP-EA (UNICOS 7.C.3), Cray C90 (UNICOS 7.C.3), IBM RS/6000 model 550 (AIX 3.2.3), Silicon Graphics IRIS-4D/35TG (IRIX 4.0.5), Silicon Graphics IRIS-Indigo R4000 (IRIX 5.2), Sun SparcStation IPX (SunOS 4.1.2), Sun SparcStation LX (SunOS 4.1.2)
- Version 6.2: Cray-2 (UNICOS 7.0.5), Cray X-MP-EA (UNICOS 7.C.3), Cray C90 (UNICOS 7.C.3), IBM RS/6000 model 550 (AIX 3.2.5), Silicon Graphics IRIS-4D/310GTXB (IRIX 4.0.5), Silicon Graphics IRIS-Indigo R4000 (IRIX 5.2), Sun SparcStation IPX (SunOS 4.1.2)
- Version 6.5: Cray-2 (UNICOS 7.0.5), Cray X-MP-EA (UNICOS 7.C.3), Cray C90 (UNICOS 7.C.3), IBM RS/6000 model 550 (AIX 3.2.5), DEC Alpha 3000/300 LX (OSF/1 v2.0), Silicon Graphics IRIS-Indigo R4000 (IRIX 5.2), Sun SparcStation IPX (SunOS 4.1.2)
- Version 6.6: Cray-2 (UNICOS 8.0.3), Cray X-MP-EA (UNICOS 8.0.3), Cray C90 (UNICOS 8.0.3), IBM RS/6000 model 550 (AIX 3.2.5), DEC Alpha 3000/300 LX (OSF/1 v2.0), Silicon Graphics IRIS-Indigo R4000 (IRIX 5.2), Silicon Graphics IRIS-Power Challenge L R8000 (IRIX 6.1), Sun SparcStation IPX (SunOS 4.1.2), Power Macintosh 7100/66 (Mac OS 7.5.2)
- Version 6.6.1: Cray-2 (UNICOS 8.0.3), Cray C90 (UNICOS 8.0.3), IBM RS/6000 model 550, 590 (AIX 4.1), Silicon Graphics IRIS-Indigo R4000 (IRIX 5.3), Silicon Graphics IRIS-Power Challenge L R8000 (IRIX 6.1), Sun SparcStation IPX (SunOS 5.4)
- Version 7.0: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), Silicon Graphics IRIS-Power Challenge L R8000 (IRIX 6.1), Sun SparcStation IPX (SunOS 5.4), DEC Alpha 3000/300 (OSF/1)
- Version 7.1: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), Silicon Graphics IRIS-Power Challenge L R8000 (IRIX 6.1)
- Version 7.1.1: Cray C90 (UNICOS 8.0.3),
- Version 7.2: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), and Silicon Graphics IRIS-Power Challenge L R8000 (IRIX 6.1)
- Versions 7.3–7.3.1: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), and Silicon Graphics Indigo-2 R10000 (IRIX 6.2)
- Versions 7.4–7.4.3: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), Silicon Graphics Indigo-2 R10000 (IRIX 6.2), and Dell XPS Pro200n (Linux, RedHat 4.2)

- Versions 7.8–7.9.2: Cray C90 (UNICOS 8.0.3), IBM RS/6000 models 550 and 590 (AIX 4.1), Silicon Graphics Indigo-2 R10000 (IRIX 6.2), and Micron Millennia Pro (RedHat Linux 4.2, kernel 2.0.30).
- Version 8.0: IBM RS/6000 model 590 (AIX 4.3), Silicon Graphics Indigo-2 R10000 (IRIX 6.2), and Dual Intel Pentium II-333 (RedHat Linux 4.3, kernel 2.0.33).
- Versions 8.1–8.5.1: IBM SP with PowerPC 604e processors (AIX 4.3.1) and Silicon Graphics Power Challenge L with R10000 (IRIX 6.5).
- Versions 8.6–8.7.3: IBM SP with Power3 processors (AIX 4.3.4.0) and SGI Origin 2000 R12000 (IRIX 6.5.12f).
- Versions 8.9–9.1: IBM SP with Power3 processors (AIX 4.3.4.0), SGI Origin 2000 R12000 (IRIX 6.5.12f), Sun Blade 2000 (Solaris 8), and Compaq XP1000 AlphaStation (Tru64 4.0f).
- Versions 8.9–9.1: IBM SP with Power3 processors (AIX 4.3.4.0), SGI Origin 2000 R12000 (IRIX 6.5.12f), Sun Blade 2000 (Solaris 8), and Compaq XP1000 AlphaStation (Tru64 4.0f).
- Versions 9.4-2008: See Tables in Section 4.C.
- Version 2010: SGI Altix with Itanium 2 processors, IBM BladeCenter with Dual-core AMD Opteron processors, SGI Altix XE 1300 Linux Cluster, and Sun Fire X4600 Linux Cluster.
- Version 2016: HP c7000 Linux Cluster (CentOS 6.8) and HP Haswell Linux Cluster (CentOS 6.8).
- Version 2017: HP c7000 Linux Cluster (CentOS 6.8) and HP Haswell Linux Cluster (CentOS 6.8).

4.C. Compiler versions

- Versions 7.4–7.4.3: Cray C90 (cft77 Version 6.0.4.19), IBM RS/6000 models 550 and 590 (XL Fortran Version 3.2), Silicon Graphics Indigo-2 R10000 (Fortran 77 GCS Version 6.2), and Dell XPS Pro200n (f2c-gcc, kernel 2.0.30).
- Versions 7.8–7.9.2: Cray C90 (cft77 Version 6.0.4.19), IBM RS/6000 models 550 and 590 (XL Fortran Version 3.2), Silicon Graphics Indigo-2 R10000 (Fortran 77 GCS Version 6.2), and Micron Millennia Pro (f2c-gcc, kernel 2.0.30).
- Version 8.0: IBM RS/6000 model 590 (XL Fortran Version 3.2), Silicon Graphics Indigo-2 R10000 (Fortran 77 GCS Version 6.2), and Dual Intel Pentium II-333 (f2c-gcc, kernel 2.0.33).

Versions 8.1–8.5.1: IBM SP with PowerPC 604e processors (XL Fortran Version 6.0.1) and Silicon Graphics Power Challenge L with R10000 (MIPSpro 7.2.1.3m).

Versions 8.6–8.7.3: IBM SP with Power3 processors (XL Fortran 6.1.0.3) and SGI Origin 2000 R12000 (MIPSpro 7.3.1.2m).

Versions 8.9–9.1: IBM SP with Power3 processors (XL Fortran 6.1.0.3), SGI Origin 2000 R12000 (MIPSpro 7.3.1.2m), Sun Blade 2000 (Sun WorkShop[tm] 6 update 2 Fortran 77 5.3), and Compaq XP1000 AlphaStation (Compaq Fortran V5.4-1283-46ABA).

Versions 9.4-9.7:

Platform	Operating System	Compilers
IBM Regatta, Power4 processors	AIX 5.3	XL Fortran version 8.1.1
		g77 from gcc version 3.4.1
IBM Netfinity, Dual Pentium III processors	Redhat Enterprise Linux 3	Portland group fortran compiler version 5.2
		g77 from gcc version 3.3.3
SGI Altix, Itanium 2 processors	SuSE Linux	Intel ifort version 9.0
		g77 from gcc version 3.3.3
IBM BladeCenter, Dual-core AMD Opteron processors	SueSE Linux	g77 from gcc version 3.3.3

Version 2008:

Platform	Operating System	Compilers
IBM Regatta, Power4 processors	AIX 5.3	XL Fortran version 8.1.1
		g77 from gcc version 3.4.1
SGI Altix, Itanium 2 processors	SuSE Linux	Intel ifort version 9.0
		g77 from gcc version 3.3.3
IBM BladeCenter, Dual-core AMD Opteron processors	SueSE Linux	g77 from gcc version 3.3.3
SGI Altix XE 1300 Linux Cluster	SuSE Linux	gfortran openmpi/gnu version 1.2.4 or higher

Version 2010:

Platform	Operating System	Compilers
SGI Altix, Itanium 2 processors	SuSE Linux	Intel ifort version 11.0
		gfortran from gcc version 4.3.3
IBM BladeCenter, Dual-core AMD Opteron processors	SuSE Linux	Intel ifort version 11.0
		gfortran from gcc version 4.3.3
SGI Altix XE 1300 Linux Cluster	SuSE Linux	gfortran openmpi/(gnu or intel) version 1.3.2 or higher
Sun Fire X4600 Linux Cluster	SuSE Linux	Intel ifort version 11.0
		gfortran from gcc version 4.3.3 openmpi/(gnu or intel) version 1.3.2 or higher

Version 2016

Platform	Operating System	Compilers
HP c7000 Linux Cluster	CentOS 6.8	Intel ifort version 13.1.3
		gfortran from gcc version 4.7.2
		Openmpi(gnu or intel)
HP Haswell Linux Cluster	CentOS 6.8	Intel ifort version 13.1.3
		gfortran from gcc version 4.7.2
		Openmpi (gnu or intel)

Version 2017

Platform	Operating System	Compilers
HP c7000 Linux Cluster	CentOS 6.8	Intel ifort version 13.1.3
		gfortran from gcc version 4.7.2
		Openmpi(gnu or intel)
HP Haswell Linux Cluster	CentOS 6.8	Intel ifort version 13.1.3
		gfortran from gcc version 4.7.2
		Openmpi (gnu or intel)

5. PROGRAM AND MANUAL DESCRIPTION

5.A. Code description, portability, and special files

The *Polyrate* program is written in FORTRAN 90 with the module and dynamically allocated memory features.

Users should be aware that, beginning with version 2008, *Polyrate* can perform two kinds of calculations. One is RP-VTST, and the other one is VRC-VTST, as explained in Section 1. These two kinds of calculations involve different subroutines to generate different executable programs. RP-VTST has both serial and MPI parallel versions and VRC-VTST has only an MPI parallel version and requires one to link to the SPRNG library. Starting with version 2010, the SPRNG package is included in the *Polyrate* distribution for the user's convenience, and the *Polyrate* installation script includes compiling the SPRNG library before linking *Polyrate* subroutines to this library. Therefore installation of SPRNG library and *Polyrate* program can be done with one script. To use RP-VTST, no pre-installed library or package is required.

The SPRNG package can be also obtained separately from the web page:
<http://sprng.cs.fsu.edu/>

5.B. Description of the manual

This manual is divided into five parts.

Part I has five sections and includes general information about *POLYRATE*.

Part II has five sections and includes theoretical background and computational details for RP-VTST. Section 6 provides an overview of theoretical background for the dynamical calculations. Section 7 gives information related to the program structure. Sections 8 and 9 describe the calculations using a potential energy surface (Section 8) or direct dynamics calculations, which use an electronic structure input file (Section 9). Section 10 provides selected details of the calculations, especially for procedures that are not fully described in publications.

Part III has one section, Section 11, to describe the theoretical background of VRC-VTST.

Part IV has 3 sections for description of input files. Section 12 describes the complete *POLYRATE* input and selected aspects of the output for RP-VTST calculations. Section 13 describes the input file for VRC-VTST. Section 14 describes files that give extra details of LCT calculations.

Part V gives more general information about the *POLYRATE* program package. Section 15 contains instructions on how to install, test, and use the *POLYRATE* program. Section 16 gives sample runs (the *POLYRATE* test suite) that may be useful in understanding the input or testing the program. Section 17 lists the files in the program distribution, and it has a brief explanation of each subprogram. Section 18 contains instructions on using the Unix scripts that are provided. Section 19 describes file usage. Section 20 contains tables of compiling and linking commands and memory requirements on various computers and with various operating systems. The memory requirements were obtained using the "size" command on the various Unix computers. Section 21 contains test run timing information. Section 22 provides other references used as footnotes in the text, and Section 23 is the *POLYRATE* bibliography.

Much of the mathematical formulation underlying the dynamical calculations is presented in a book chapter entitled "Variational Transition State Theory with Multidimensional Tunneling," which is Ref. 70 in the bibliography (Sect. 23), and should be consulted along with this manual. "Generalized Transition State Theory" (GTST), which is Ref. 5 in Sect. 23, is also a useful chapter, especially for an introduction to the older core ideas, although Ref. 70 supersedes it. If these books are unavailable in your library, you may obtain a complimentary copy of the chapters from the senior author (DGT); please see

<http://comp.chem.umn.edu/Polyrate>

A PDF file with a complete compilation of all known errors in Ref. 5 is also given at this Web site.

Section 25 of this manual contains the *POLYRATE* revision history.

5.B.1. Referencing conventions

Reference numbers in this manual refer to references in Sections 23. In all cases the reference number is accompanied by the section number to avoid confusion.

5.B.2. Book chapters

Reference 5 in Section 23 is sometimes referred to simply as "the GTST book chapter". Reference 70 in Section 23 is sometimes referred to as "the VTST/MT book chapter." These two book chapters explain the methods used in *Polyrate*. See

<http://comp.chem.umn.edu/Polyrate>

Part II Theory Background and Calculation Details on RP-VTST

6. THEORETICAL BACKGROUND ON RP-VTST

This section provides an overview of the dynamical theory underlying the calculation procedures. Further details of the methods for gas-phase bimolecular and unimolecular reactions may be found in the "Generalized Transition State Theory" (GTST) book chapter (Ref. 5, Sect. 23). Details of new methods developed after that chapter was written are described in Section 10 of this manual, if not fully published, or in the additional references in Section 21, especially the VTST/MT book chapter (Ref. 70 of Sect. 23).

6.A. Basic theory

As described in the Introduction, the program can be used to calculate rate constants for reactions of one or two reactant molecules to form one or two product molecules, using a variety of generalized transition state theory methods. In order to simplify the discussions in the remainder of this introductory overview section, we consider only the calculation of forward and reverse thermal rates using both conventional transition state theory (TST) and canonical variational theory (CVT). Microcanonical variational theory (μ VT) are described in detail in the GTST book chapter. The applications of VTST methods to gas-phase unimolecular or bimolecular reactions or gas-solid-interface reactions or the calculation of adiabatic and diabatic state-selected rate constants are described in the references included in Section 22.

Note that the improved canonical variational theory (ICVT) option is removed from version 2016 and the much more efficient algorithm, the extended Beyer-Schwinehart (BS) algorithm, is used for calculating number of available states for μ VT. If one sees a need for the more accurate threshold treatment of ICVT, one should instead use μ VT because μ VT with BS algorithm is even more efficient and complete than ICVT and is now coded efficiently.

The conventional transition-state-theory canonical rate constant involves a symmetry number [Ref. 1, Sect. 22] times kT/h times a ratio of a partition function evaluated at the

saddle point to one evaluated at the reactants, and, optionally, times a temperature-dependent transmission coefficient, which in this program, takes into account tunneling and non-classical reflection effects. The partition functions are evaluated as products of electronic, vibrational, and rotational partition functions, and for the reactants of a bimolecular reaction, a relative translational partition function per unit volume. In *Polyrate*, vibrational partition functions are defined such that they tend to a Boltzmann factor of the zero point energy at low temperature (not to unity as is common in statistical mechanics books), and the corresponding local zero of energy is used consistently for the rest of the calculations.

To calculate rate constants by the CVT method various items of information are needed along the reaction path. In *Polyrate*, the reaction path is always taken as the minimum energy path (MEP) through mass-scaled (i.e. isoinertial) coordinates [Refs. 2-5, Sect. 22] except for the treatment of loose transition state using variable-reaction-coordinate (VRC) algorithm with single-faceted or multifaceted (MF) dividing surface (the VRC algorithm and MF dividing surfaces are discussed in detail in Section 11). The descriptions here are all referred to MEP. Following Fukui [Ref. 5, Sect. 22], one may call this the "intrinsic" reaction path. (In *Polyrate*, we always call it the MEP.) The distance along the MEP is called s , or the reaction coordinate. For reactions with a saddle point, the MEP is specifically defined to be the path of steepest descent through mass-scaled Cartesian coordinates from the saddle point ($s = 0$) to the reactants (at which $s = -\infty$ for a bimolecular reaction, or s equals a finite negative number for a unimolecular reaction) and the path from this saddle point to the products (at which $s = +\infty$ for a bimolecular reaction, or s equals a finite positive number for a unimolecular product). We define a sequence of generalized transition states corresponding to the reaction coordinate to be hypersurfaces which are orthogonal to the MEP [Ref. 6, Sect. 22]. All quantities calculated for a generalized transition state that intersects the MEP at a particular value s of the reaction coordinate are labeled by that value of s (as well as the temperature T when they depend on temperature). The generalized TST canonical forward rate constant $k_f^{\text{GT}}(T, s)$ is then evaluated like the conventional one but with saddle point properties replaced by generalized transition-state (GT) ones. Thus, at each generalized transition state along the reaction path the following information is (or may be, depending on the options chosen) needed:

- $V_{\text{MEP}}(s)$ — potential along the minimum energy path,
- $|\mathbf{I}(s)|$ — determinant of the moment of inertia tensor,

- $X(s)$ — geometry of the generalized transition state,
- $\nabla V_{\text{MEP}}(s)$ — the gradient of the generalized transition state,
- $\mathbf{L}_m(s)$ — eigenvector of the generalized normal mode m ,
- $\omega_m(s)$ — generalized normal mode frequencies.

In addition, depending on the anharmonicity option and the method chosen to include tunneling, one may also require some or all of the following additional functions:

- $k_m^{(3)}(s)$ — principal cubic force constants for the generalized normal modes,
- $k_m^{(4)}(s)$ — principal quartic force constants for the generalized normal modes,
- $B_{mF}(s)$ — reaction-path curvature coupling [Ref. 7, Sect. 22] between the reaction coordinate (mode F) and generalized normal mode m . Sometimes we refer to the set of $B_{mF}(s)$ with $m=1, 2, \dots, F-1$, as the curvature vector \mathbf{B}_F .

Note that “principal” force constants are diagonal in the normal mode index; that is why $k_3(s)$ and $k_4(s)$ are short one-dimensional arrays. The functions listed above and the reaction path itself can be obtained through the hooks, which can in turn be interfaced to an analytic potential energy surface or an electronic structure package, or alternatively this data (or information which can be used to calculate this data) can be obtained from electronic structure calculations carried out separately from *Polyrate*. In the latter case the MEP information is input only on a relatively sparse grid of s values, and the program interpolates this information to arbitrary s values. This option is called the electronic structure input file option (POTENTIAL = *unit30*, *unit31*, or *unit40*).

The isoinertial coordinate system used by *Polyrate* is defined by eq. (2) of Ref. 1 of Sect. 23 or, equivalently, by eq. (2) of Ref. 5 of Sect. 23. These coordinate systems have the same reduced mass, μ , for all degrees of freedom. Physical observables and the MEP are independent of μ , but numerical values of s are not. Thus it is essential to specify μ if s values or plots of barriers vs. s are considered. If μ is set equal to 1 amu, then the numerical value of s in mass-scaled coordinates (which have units of length) is the same as the numerical value of the distance along the usual spectroscopists’ mass-weighted coordinates (which have units of mass^{1/2} length) if the mass weighting is amu^{1/2}. Therefore setting μ equal to 1 amu procedure is especially recommended if the goal is to

make the treatment transparent to those who are only used to the spectroscopists' mass weighting scheme.

The reverse rate constant is obtained from the forward one by detailed balance. The canonical variational theory [Ref. 6, Sect. 22] forward rate constant is obtained by minimizing the generalized transition state theory rate constant with respect to s :

$$k_f^{CVT}(T) = \min_s k_f^{GT}(T, s) = k_f^{GT}[T, s_*^{CVT}(T)] \quad (1)$$

Generalized-transition-state-theory rate constants can also be written in terms of the generalized transition-state (GT), standard-state (0), forward (f) free energy of activation, $\Delta G_f^{GT,0}(T, s)$. The location of the variational transition state is then determined by maximizing the free energy of activation with respect to s .

Section 6.C gives a brief explanation of the evaluation of the partition functions and Section 6.D summarizes methods used for transmission coefficients.

6.B. Vibrations, Rotations, and Translations

This section gives a brief explanation of how *Polyrate* uses the SPECIES keyword (see Section 12.A.6) to determine the number of rotations and vibrations used in calculating the partition functions and the number of degrees of freedom which are projected out by the projection operator (when the PROJECT option is on, see Section 12.A.6) for the given reactant, product, transition state, or generalized transition state. The table below gives the valid values of the SPECIES keyword and a description of what the values mean, along with the number of vibrations and rotations in the partition function for each type of species, and the number of modes which are projected out to remove translations, rotations, and the reaction coordinate.

Description	SPECIES keyword value	Partition Function		Projection Operator			
		Vibrations	Rotations	Translations	Rotations	Reaction Coordinate	Mandatory?
atomic reactant or product	<i>atomic</i>	0	0	N.A.
linear reactant or product	<i>linrp</i>	$3N - 5$	2	3	2	0	optional
nonlinear reactant or product	<i>nonlinrp</i>	$3N - 6$	3	3	3	0	optional
solid-state reactant or product	<i>ssrp</i>	$3N$	0	0	0	0	prohibited
linear transition state	<i>lints</i>	$3N - 6$	2	3	2	0	optional
nonlinear transition state	<i>nonlints</i>	$3N - 7$	3	3	3	0	optional
solid-state transition state	<i>ssts</i>	$3N - 1$	0	0	0	0	prohibited
linear GTS	N.A.	$3N - 6$	2	3	2	1	mandatory
nonlinear GTS	N.A.	$3N - 7$	3	3	3	1	mandatory
solid-state GTS	N.A.	$3N - 1$	0	0	0	1	mandatory

N.A., Not Applicable; N , number of atoms

For an atomic reactant or product the value *atomic* should be specified for SPECIES. Since the species is a single atom there are no vibrations or rotations, nor is there any Hessian to consider projecting. Therefore the PROJECT keyword (see Section 12.A.6) cannot be used with an atomic species.

Gas-phase reactants and products that are not atomic and transition states can be either linear or nonlinear, specified by *linrp*, *nonlinrp*, *lints*, and *nonlints* respectively. After removing the 2 rotations and three translations for a linear molecule there are $3N - 5$ vibrations for the linear species, and after removing the 3 rotations and 3 translations for a nonlinear molecule there are $3N - 6$ vibrations for a nonlinear species. A gas-phase transition state has one additional mode removed due to the reaction coordinate and therefore has $3N - 6$ real vibrations for a linear molecule and $3N - 7$ real vibrations for a nonlinear molecule. The number of vibrations for a generalized transition state is calculated in the same manner as for a conventional transition state (which is a saddle point, also called a transition structure). Projecting the Hessian at a transition state is not required, but sometimes it leads to a more stable calculation of vibrational frequencies and eigenvectors, so it is a recommended option to project out translations and rotations at a reactant, product, or transition state. For generalized transition states (points on the reaction path where the gradient does not vanish), projecting out the reaction coordinate is mandatory if the resulting frequencies are to represent motions orthogonal to the reaction coordinate. In such cases it is also mandatory to project out translations and rotations, if they exist (they always exist for gas-phase cases, but not for solid-state cases).

Solid-state species are different than gas-phase species as concerns the number of vibrations. For solid state species there are no rotations or translations which are removed because a rotation or a translation of a species relative to a surface changes the potential energy of the whole system. Therefore there are $3N$ vibrations for a reactant or product species and $3N - 1$ for a transition state or generalized transition state species. The reaction coordinate is removed for a solid-state species in the same manner as it is for a gas-phase species.

6.C. Partition functions

The electronic partition function for a given species is computed quantum mechanically by summing over the Boltzmann factors of the electronic energy levels of the species times their degeneracies. In practice, since the energy spacing between electronic multiplets is generally quite large, we usually include in the summation only those levels comprising the energetically lowest multiplet, e.g., the $J = 0, 1,$ and 2 levels of $O(^3P)$.

The rotational partition function is approximated by the classical limit [Refs. 8 and 9, Sect. 22]. For a linear reactant, product, transition state, or generalized transition state, the partition function is evaluated from the unique moment of inertia. For a nonlinear species, the product of the three principal moments of inertia is required and is conveniently obtained from the determinant of the moment of inertia tensor. All symmetry numbers are left out of all rotational partition functions; the symmetry numbers need only to be included in the overall symmetry number mentioned above [Ref. 1, Sect. 22].

For bimolecular reactions, the relative translational partition function per unit volume is calculated classically.

For vibrational partition functions, a separable approximation is assumed, whereby the overall vibrational partition function is a product of independent normal modes of a reactant, product, or saddle point or the generalized normal modes of a generalized transition state. Vibrational modes are treated classically or quantized, either quantum mechanically or semiclassically. Anharmonic methods are limited to intramode anharmonicity in normal modes expressed as linear combinations of rectilinear coordinates and do not include mode-mode coupling as a supported option at this time. The torsional potential anharmonicity along reaction path can be included using the formula of the multi-structural torsional method but only with one input structure. Classical vibrational partition functions are evaluated using the classical expression for the harmonic oscillator. Quantum mechanical vibrational partition functions are evaluated by the harmonic approximation, by summing Boltzmann factors using quantum or semiclassical energy levels.

Frequencies can be scaled for partition function calculations in CVT calculations when the harmonic approximation is employed. The scaling is not used in the calculation of transmission coefficients (either by CAG or tunneling).

6.D. Tunneling methods

Different semiclassical methods are used to evaluate transmission coefficients depending on the degree of coupling between the MEP and other degrees of freedom through the MEP curvature. There are three general types of transmission coefficients: zero-curvature tunneling (ZCT), small-curvature tunneling (SCT) and large-curvature tunneling (LCT). The methods included in this version for each type of transmission coefficient for thermal reactions are as follows:

- ZCT: minimum-energy-path semiclassical adiabatic ground-state (MEPSAG) method [Refs. 4 and 10, Sect. 22 and Ref. 5, Sect. 23].
- SCT: centrifugal-dominant small-curvature semiclassical adiabatic ground-state (CD-SCSAG) method [Refs. 29 and 31, Sect. 23].
- LCT: large-curvature ground-state version-3 (LCG3) method [Refs. 5, 21, 29, 30, and 34, Sect. 23] and large-curvature ground-state version-4 (LCG4) method [Ref. 54, Sect. 23].

The current theory of these tunneling methods appears in “Variational Transition State Theory with Multidimensional Tunneling,” which is Ref. 59. The final version (CD-SCSAG, which replaced the original SCSAG) of the SCT method is described in Refs. 29 and 31 in Sect. 23, and the final versions of the LCT method are described in Refs. 29, 30, 34 (LCG3 including tunneling into excited states), and Ref. 54 (LCG4 including tunneling into excited states) in Sect. 23. The final versions of these methods were used in all references of Sect. 23 that have reference number 30 or higher. A critical element of the SCT calculation is that the reduced mass, μ (See Section 6.A), is replaced by an effective mass $\mu_{eff}(s)$ in calculating the tunneling probabilities. See Section 6.I of this manual for further discussion.

Notice that the G in MEPSAG, CD-SCSAG, LCG3, and LCG4 refers to the fact that the transmission coefficient for thermal reactions is always calculated for the ground (G) state of the reactants in the exoergic direction of a reaction, as discussed elsewhere (Ref. [13] Sect. 22 and Refs. 29, 30, and 34 of Sect. 23). The transmission coefficient for the endoergic direction of a reaction is then obtained by detailed balance, which requires that

the transmission coefficients for thermal reactions be the same in both directions. The LCG3 and LCG4 methods can include tunneling into excited states of the product in the exoergic direction, and the algorithm in *POLYRATE* is general enough to treat even cases that involve a well on the product side in the exoergic direction. (Wells on the reactant side do not require special considerations in this regard.) The ground-state approximation for the transmission coefficient is equivalent to assuming that the energy-dependent transmission probability curve for each successive quantized level of the transition state has the same functional dependence on energy as that for the ground state but is shifted up in energy by the excitation energy evaluated at the variational transition state [Ref. 47, Sect. 22].

When reaction-path curvature is large, it may be *necessary* to include excited states of the product of the exothermic direction in order to obtain accurate results, and the user of *POLYRATE* is responsible for checking this by checking convergence with respect to the argument of the `EXCITED` keyword in the `TUNNEL` section.

VTST methods are also applicable to state-selected reactants if the quantum number of the selected mode remains adiabatic, to a good approximation, all the way to the transition state. Excited-state tunneling probabilities may be defined using similar methods to those mentioned above, and they are called MEPSA, CD-SCSA, and LC3 or LC4, i.e., we don't include the G when the method is applied to excited states of reactants. These abbreviations and the ZCT, SCT, and LCT abbreviations are also useful when we wish to refer to the semiclassical method without reference to the initial state. The present version of *POLYRATE* allows the MEPSA and CD-SCSA options for excited-state reactions but does not allow LC3 or LC4, which is not coded in such generality.

Regions of large reaction-path curvature can promote vibrational nonadiabaticity. The partial-reaction-path (PRP) adiabaticity approximation [Ref. 13A, Sect 21] assumes adiabaticity in the excited-state vibration up to the first point along the reaction coordinate where an appreciable local maximum in the reaction-path curvature occurs, which we denote s_* . The vibration is assumed to relax nonadiabatically into the ground vibrational state after s_* . The adiabatic potential for tunneling is the excited-state one up to s_* and the ground-state one for $s > s_*$. Excited-state PRP calculations can be implemented using the `STATE` and `STATEOPT` keywords in the `RATE` section. An example of an excited-state PRP calculation is provided in test run `ohcltr4`.

The multidimensional tunneling methods used in *POLYRATE* have been tested extensively against accurate quantal results [Ref. 53, Sect 20].

6.E. Reactant and product properties

The first step in the calculation is to determine the properties of the reactants and products. We will specifically discuss reactant A (the first reactant, which may be an atom or molecule) as an example in this subsection; the methods for the other reactant, in the case of a bimolecular reaction, or for products are analogous.

The calculation begins by determining the equilibrium geometry and energy for reactant A on the potential energy surface. The reactant geometry is optimized by using either the Newton-Raphson (NR) method with Brent line minimization [Ref. 14, Sect. 22], the Broyden-Fletcher-Goldfarb-Shanno (BFGS) method [Ref. 15, Sect. 22], or the Eigenvector Following (EF) method [Refs. 54-58, Sect. 22] to search over a restricted set of atomic Cartesian coordinates. These options are not available for the program packages *MORATE* and *AMSOLRATE*, in which the geometry optimization is only carried out in the electronic structure program (which is specified using option *HOOK*). To do this, the user must fix (somewhat arbitrarily) 6 of the Cartesian coordinates (5 for a linear species) and search over the rest. For example, the user might fix atom 1 at the origin, atom 2 on the x-axis, and the atom 3 in the x-y plane. (In the case of solid-state or gas-solid-interface reactions, there are no translations or rotations because the moving atoms are adsorbed on or embedded in a stationary lattice; thus for such cases, the full set of atomic Cartesians of the movable atoms is used). Starting from an initial guess, an improved estimate for the position of the minimum is obtained by taking a step ΔR_0 , where ΔR_0 is obtained by solving a simultaneous set of equations (given in the GTST book chapter) involving the matrix of second derivatives of the potential energy and the gradient. Once ΔR_0 is obtained by direct calculation of the inverse Hessian matrix (NR method) or estimated using information from the previous geometry optimization steps (BFGS method), a new guess for the position of the minimum is obtained using either the NR step or a variable step size which is determined by the Brent line minimization method [Ref. 16, Sect. 22] (BFGS method). For the EF method, the step is determined by the hessian. If the hessian has the wrong number of negative eigenvalue, then the partitioned rational function (P-RFO) approximation is used. If the hessian has one

negative eigenvalue, then Newton-Raphson step is used. The detailed EF algorithm is given in [Ref. 56, Sect. 22].

For all methods, convergence is reached when the maximum component of the gradient in the space of optimized coordinates is less than a prescribed amount *GCOMP*. In order to increase the stability of the optimization, the maximum component of the step, ΔR_0 , is not allowed to be larger than the prescribed amount *SCALE* for the NR and BFGS methods. However, the *SCALE* keyword has no effect on the EF method. The accurate Hessian is recalculated every *HREC* optimization steps. In the case of the NR method, if *HREC* is not equal to 1, the Hessian is kept frozen until recalculated. The BFGS method is the default. The EF method has a few extra options, namely, *RMAX*, *RMIN*, *OMIN*, *DDMAX*, and *DDMAXTS*, to control the size of the trust region and tune up the optimization procedure.

Next, the origin is moved to the center of mass of reactant A. The moment of inertia or the product of the principal moments of inertia is then determined. Finally, a normal mode analysis is performed on reactant A. Since, in mass-scaled Cartesians, motion in any direction is associated with the same reduced mass (that is why we call these isoinertial coordinates), this is easily accomplished by diagonalizing the force constant matrix. The force constant matrix is calculated by numerical first derivatives of the analytic or input first derivatives of the potential. In the case of gas-phase reactions, we can optionally project out the contamination of the translational and rotational motions in the force constant matrix at the stationary point before diagonalizing it to yield 6 zero eigenvalues (5 for a linear molecule) corresponding to overall translation and rotation. The remaining eigenvalues are all positive, corresponding to bound normal mode frequencies, and their associated (normalized) eigenvectors L_m give the normal mode directions. After the frequencies are calculated, the frequencies and corresponding eigenvectors are placed in “canonical order.” Canonical order is defined as follows:

First define *G*:

Stationary point: $G = R + T =$ 6 for nonlinear molecule,
5 for linear molecule,
0 for gas-solid interface.

Reaction path: $G = R + T + 1$

Then the frequencies are ordered as follows:

1. The *G* frequencies with the smallest absolute values go last, ordered in order of decreasing magnitude.

2. Of the remaining frequencies, first put the real ones, in decreasing order, then put the imaginary ones, in order of increasing magnitude.

By computing or reading the third and fourth derivatives of the potential energy along these normal mode directions, anharmonicity may be included separately in each of the modes, as discussed elsewhere [Ref. 1, Sect. 23]. Depending on the method chosen for treating the anharmonicity, the zero point energy is then computed for reactant A by summing up the zero point energies in each of the normal modes.

The calculations in this section are then repeated for the other reactant (if it exists) and for product(s). From these results, the adiabatic energy of reactants, the classical endoergicity for the reaction, ΔE , and the product adiabatic energy are obtained.

Notice that the normal mode analysis can only be carried in rectilinear coordinates for reactants and products, whereas the saddle point and the nonstationary points can be treated either in rectilinear or in curvilinear coordinates. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.)

6.F. Saddle point properties

The calculation of the saddle point geometry and properties is next. The saddle point position is determined by means of the Newton-Raphson method, the BFGS method, or the EF method as described in Subsection 6.E. The default for transition state searches is the Newton-Raphson method. This optimization provides the saddle point's classical barrier height $V_{\text{MEP}}(s=0)$. Next, the origin is moved to the center of mass of the full N -atom system. The moment of inertia is obtained as discussed for reactants. Next the force constant matrix is calculated in mass-scaled Cartesians, and optionally the contributions of the overall translational and rotational motions are projected out. The normal mode analysis can be carried out in either the rectilinear or curvilinear coordinates for the saddle point. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.) In the case of using rectilinear coordinates, a diagonalization of the $3N \times 3N$ matrix of force constants in mass-scaled Cartesian coordinates is performed. Upon

diagonalization, we again obtain 6 zero eigenvalues (5 for a linear molecule, 0 for solid state), $3N-7$ ($3N-6$ or $3N-1$) positive eigenvalues corresponding to bound normal mode frequencies, and one negative eigenvalue corresponding to the imaginary frequency whose eigenvector provides the reaction coordinate direction at the saddle point. After the frequencies are calculated, the frequencies and corresponding eigenvectors are placed in canonical order (which is defined in Section 6.E). The imaginary frequency is used at a later stage to compute the Wigner tunneling correction (see Subsection 6.J). By computing or reading the third and fourth derivatives along the normal mode directions as discussed above, anharmonicity can be included separately in each of the $3N-7$ ($3N-6$ for a linear molecule, or $3N-1$ for solid state) bound normal modes (see Subsection 10.B of this manual), and the zero point energy (ZPE) for the transition state is computed from the sum of the zero point energies of each of the modes. The ground-state adiabatic energy at the saddle point is then calculated as:

$$V_a^G(s=0) = V_a(n=0, s=0) = V_{\text{MEP}}(s=0) + \text{ZPE}(s=0) \quad (2)$$

The normal mode analysis can also be carried using redundant (or nonredundant) curvilinear coordinates. In this case, Wilson's *FG* matrix method is used to obtain the projected force constant matrix in internal coordinates. See Section 10.L. for further discussion of curvilinear internal coordinates.

6.G. Reaction path

In this section, we discuss how the reaction path (or, more precisely, a set of points along an approximation to the reaction path) and various properties along it are determined. In the present version of *Polyrate*, the reaction path is always taken as the minimum-energy path (i.e. the path of steepest descents) in mass-scaled coordinates; this is called the MEP. Note that the path of steepest descent is the same in any isoinertial coordinate system. [Ref. 49, Sect. 22] There are four cases, and the current version of *POLYRATE* allows three of them:

In START	In PATH	
STATUS = 0	SADDLE	Reaction has a saddle point. Initial guess of its geometry is provided in the START section of the fu5 input file. The geometry of the saddle point is optimized. The first step is followed along the imaginary frequency normal mode (if <i>FIRSTSTEP</i> is <i>nmode</i>) or with the cubic starting algorithm based on

		this mode (if FIRSTSTEP is <i>cubic</i>).
STATUS = 2	SADDLE	This is the same as above except it assumes that the geometry in the START section is already fully optimized, and it skips the optimization step.
STATUS = 0	NOSADDLE	Not supported.
STATUS = 2	NOSADDLE	Reaction does not have a saddle point. It is a strictly downhill association reaction. The reaction path is started by following the negative gradient at the geometry provided in the START section.

Most of the test runs in the current version of *Polyrate* correspond to STATUS = 0/SADDLE. An example of STATUS = 2/SADDLE is test run oh3tr1. All fu29, fu30, fu31, and fu40 test runs use STATUS = 2/SADDLE. An example of STATUS = 2/NOSADDLE is ho2tr2.

6.G.1. SADDLE option

Taking the saddle point as the first point on the reaction path, an initial step of prescribed length is taken according to one of two options. Either it is taken based on a cubic expansion of the potential energy surface [Ref. 17, Sect. 22] using the normalized eigenvector corresponding to the imaginary frequency and the curvature vector, both of which are calculated at the saddle point, or it is taken in the direction given solely by the normalized eigenvector corresponding to the imaginary-frequency normal mode at the saddle point, as was done in the original version of the program [Ref. 1, Sect. 23] described in the GTST book chapter.

In the original straight VTST calculations [Ref. 18, Sect. 22; Ref. 1, Sect. 23] the MEP was calculated by the Euler steepest-descent (ESD) method in which repeated steps of length Δs are then taken in the direction of the negative of the normalized gradient. The whole process is then repeated for the other direction of leaving the saddle point, thus providing the entire reaction path. The distance between successive points along the reaction path (δs) is called the reaction path stride or the gradient step size, and the set of such points is called the fine grid or the gradient grid. Thus, at each point on the fine grid both the potential and gradient are required.

To reduce the numerical error in the above approach due to wandering about the true steepest-descent path, the program allows for a stabilization algorithm to be employed.

This algorithm [Ref. 18, Sect. 23] may be considered to be a modified version of the prescriptions of Ishida et al. [Ref. 19, Sect. 22] and Schmidt et al. [Ref. 20, Sect. 22]. It is called ES1 (Euler stabilization method, version 1), and it is described elsewhere [Ref. 18, Sect. 22]. In the original version of the ES1 algorithm, the stabilization step was omitted whenever the vector difference between successive normalized gradients was less than 10^{-8} , but now this is only a default value and the user is free to choose among other possible values which might be more appropriate for the reacting system under study. This is controlled by the keyword DIFFD.

The ES1 method involves two auxiliary parameters, the stabilization step size DELTA2 and the tolerance in the allowed magnitude DIFFD of the vector difference between successive normalized gradients, and the accuracy achieved for a given amount of computer time can be sensitive to these values. These values have been approximately optimized in a systematic study of methods for calculating minimum energy paths (see Ref. 27 in Sect. 23). In that study, "standard" values were determined for both parameters. These "standard" values are: a) the stabilization step size DELTA2 equal to the gradient step size SSTEP, and b) the tolerance in the allowed magnitude of the vector difference between successive normalized gradients equal to 0.01. When the ES1 method is used with these choices for the stabilization step size and the tolerance in the allowed magnitude of the vector difference between successive normalized gradients, the algorithm is called the optimized ES1 method or ES1*.

The program also allows one to use the Page-McIver algorithm [Ref. 17, Sect. 22], rather than the ESD or Euler stabilization method, to compute the reaction path. In our implementation of the Page-McIver method, the first step away from the saddle point is again taken as described above, and subsequent steps are taken using a local quadratic approximation (LQA) as described in [Ref. 17, Sect. 22] or in the hard-copy manual for version 1.5.

POLYRATE can also use the variational reaction path algorithm based on the Euler steepest-descents method, called VRP(ESD) [Ref. 49, Sect. 23], to construct the reaction path.

Methods for calculating the reaction path are discussed further in Sect. 10.D.

In order to obtain the partition functions needed to evaluate $k^{\text{GT}}(T, s)$, a generalized normal mode analysis is performed at regular intervals along the reaction path. This is done at the points referred to as the save grid. The generalized normal mode analysis can be carried out in either rectilinear (Cartesian) or curvilinear coordinates. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.)

The algorithm used for rectilinear coordinate calculations is discussed by Miller, Handy, and Adams [Ref. 7, Sect. 22]. In particular, for $s \neq 0$ the rectilinear or Cartesian generalized normal modes orthogonal to the reaction path are obtained from the projected force constant matrix

$$\mathbf{F}^P(s) = [1 - \mathbf{P}(s)]\mathbf{F}(s)[1 - \mathbf{P}(s)], \quad (3)$$

where $\mathbf{F}(s)$ is the full ($3N \times 3N$) Cartesian force constant matrix evaluated at a point on the MEP, and $[1 - \mathbf{P}(s)]$ is a projector used to project the direction along the reaction path as well as overall rotations and translations out of the space associated with the non zero eigenvalues. Diagonalization of $\mathbf{F}^P(s)$ gives 7 zero and $3N-7$ positive eigenvalues (6 zero and $3N-6$ positive eigenvalues for linear molecules, or 1 zero and $3N-1$ positive eigenvalues for adsorbed species) corresponding to the bound generalized normal modes orthogonal to the reaction coordinate. After the frequencies are calculated, the frequencies and corresponding eigenvectors are placed in canonical order (which is defined in Section 6.E). If the REORDER keyword is specified, the code will attempt to correlate the modes from point to point in terms of their physical characters rather than their magnitudes. Thus the first mode would be the mode with the highest overlap with the first mode at the previous point, rather than the highest-frequency mode, and so forth. The REORDER option would be useful for studying diabatic correlations of vibrational modes. Once the frequencies have been reordered anharmonicity corrections, the zero point energy, and the ground-state adiabatic energy are computed as they were at the saddle point.

The curvilinear generalized normal mode analysis is carried out by first transforming the Cartesian force constant matrix into a curvilinear coordinate system, then projecting out the gradient components, and finally diagonalizing the projected curvilinear force

constant matrix according to Wilson **GF** method [Refs. 36 and 37 of Sect. 22] as described in Refs. 41, 43, 45, and 46 of Sect. 23. Generalized normal mode frequencies, the zero point energy, and the ground-state adiabatic energy are then computed as they were at the saddle point. Methods for calculating the curvilinear generalized normal mode frequencies are discussed further in Sect. 10.M.

Section 10 and the GTST book chapter give details of the WKB method for computing anharmonic vibrational energies.

Starting with *POLYRATE* version 4.0, vibrational modes corresponding to torsional motion can be treated with the hindered-internal-rotator approximation [Refs. 24, 29, and 34, Sect. 23].

6.G.2. NOSADDLE option

The NOSADDLE option is used only for strictly downhill association reactions. It requires the user to input an initial guess for a point on the MEP such that the variational transition state lies between this point and the product, which is the associated molecule. Let $s = 0$ denote the initial guess, and let s^P denote the value of s at the product. Typically, one will compute the MEP only up to some value s_2 that is less than s^P . Furthermore, the first few points are typically not useful because it takes a few steps for the path generated by the steepest descent algorithm to become sufficiently close to the MEP. If this occurs by $s = s_1$, where $0 \leq s_1 \leq s_2$, then it is the user's responsibility to ensure that the variational transition state lies between s_1 and s_2 .

Starting with *POLYRATE* version 2008, VRC option is recommended instead of NOSADDLE option for barrierless association reactions. The VRC option uses a variable reaction coordinate and single-faceted or multifaceted dividing surfaces. See Sections 11 and 13 for details.

6.H. Reaction-path curvature

The curvature of the reaction path is used in both the SCT and LCT calculations. The curvature of the reaction path is a vector with $F - 1$ components, where F is the number of internal degrees of freedom ($F = 3N - 6$ for a gas-phase reaction with a nonlinear reaction path, $F = 3N - 5$ for a gas-phase reaction with a linear reaction path, and $F = 3N$

for a solid-state reaction employing a fixed lattice, as in the embedded-cluster method). In this manual the components of the curvature vector are denoted B_{mF} , $m = 1, 2, \dots, F-1$, where m labels a generalized-normal-mode direction.

When using an analytic potential energy surface or electronic structure input data from unit fu30 or unit fu40, there are three choices for how to compute B_{mF} . The first two choices are based on eq. (98) of the GTST book chapter [Ref. 5, Sect. 23]. This formula involves a numerical derivative of the negative gradient, and this derivative may be computed by central or one-sided finite differences. The third choice is to use eq. (48) of a paper by Page and McIver [Ref. 17, Sect. 22]. Notice that eq. (98) of the GTST book chapter is the same as eq. (47) of Page and McIver paper. An advantage of eq. (48) is that it does not involve numerical differentiation of the gradient. Both formulas require the generalized-normal-mode eigenvectors so the curvature is calculated on the Hessian grid (as opposed to the gradient grid). The evaluation of the curvature by taking the derivative of a quadratic fit to the gradient at three points (i.e., a central finite difference) is specified by the option *dgrad* of the keyword CURV; the evaluation by taking the derivative of a one-sided two-point fit to the gradient is called *oneside*, and the evaluation by eq. (48) of Page and McIver is called *dhess* because it involves a derivative of the Hessian. Although in principle, *dhess* can be used even when the reaction path is calculated by the ESD method, this can be dangerous because the accuracy of a numerical derivative requires a small step size. When the ESD method is used to calculate the reaction path, the gradient may be known on a much finer grid than the Hessian grid, and numerical differentiation of the Hessian may be inaccurate. If the *dhess* option is used with an ESD integration scheme, one should be especially careful to converge the Hessian with respect to INH.

These methods all break down (become unstable) at or very near to the saddle point. Thus the code does not obtain the curvature vector directly at the saddle point. Instead it obtains the value of B_{mF} at the saddle point by linear interpolation of the absolute values at the closest negative- s save grid point and the closest positive- s save grid point. That is, if the saddle point is s_{NS} , one obtains the curvature at the saddle point as an average of the values at s_{NS-1} and s_{NS+1} if INTMU equals 1, and one obtains the curvature at the saddle point as an average of the values at s_{NS-2} and s_{NS+2} if INTMU equals 3. The curvature at the saddle point is used to calculate $\mu_{eff}(s)$ at s_{NS} . There is an option to compute $\mu_{eff}(s)$ at s_{NS} , s_{NS-1} , and s_{NS+1} by linear interpolation of the values at s_{NS-2}

and s_{NS+2} . This option corresponds to setting INTMU equal to 3 in the PATH section, this is currently the default.

When using the electronic structure data from unit fu31, the method for computing B_{mF} is automatically chosen based on whether or not a gradient is available at the closest point on both sides of the Hessian point. If gradients are available on both sides of the point then the three gradients are fit to a quadratic function and the derivative of this function is calculated and then evaluated at the Hessian grid point. If the gradient is only available on one side then the derivative is evaluated by one-sided difference. For obtaining the curvature vector at the saddle point the procedure described above is also used.

However, if the user selects to perform a direct interpolation of the exponent in eq. 10 of Ref. 48 in Sect. 23, the IVTST-M calculation will obtain μ_{eff} at the saddle point from a spline fitting of the exponent as a function of the mapping variable z . See Section 6.N.2 for more IVTST-M details.

In contrast, when data is obtained from unit fu29, which is used for IVTST-1 calculations, the curvature at the saddle point is computed using an alternative formula presented as equation (49) in the Page and McIver article [Ref. 17 of Sect. 22], and its value at the "extra point" is computed again using eq. (48) of the same reference. The value at the saddle point is actually used in the IVTST-1 calculation, whereas the value at the extra point is computed for informational purposes only.

See Section 10.F for further discussion of the curvature calculations.

6.I. Transmission coefficients

The temperature-dependent ground-state transmission coefficient is the ratio of the Boltzmann average of the quantum energy-dependent ground-state transmission probability to the Boltzmann average of the energy-dependent transmission probability corresponding to classical-reaction-coordinate motion with other degrees of freedom in the ground state. Both transmission probabilities are zero for an energy lower than the maximum of the ground-state energy of the reactants and of the products. For μVT , the threshold energy for the classical transmission probability is the maximum of the adiabatic ground-state energy $V_a^G(s)$, i.e., V_a^{AG} ; it is $V_a^G(s=0)$ for TST and $V_a^G(s=s_*^{CVT})$ for CVT. One can factor the ground-state transmission coefficient into

two factors. The first factor is a ratio of the Boltzmann average of the quantum transmission probability to the Boltzmann average of the classical transmission probability with the threshold energy at the maximum of the adiabatic ground-state energy. The second factor is a ratio of the Boltzmann average of the classical transmission probability with the threshold energy at the maximum of the adiabatic ground-state energy to the Boltzmann average of the classical transmission probability with the threshold at the value that is consistent with the level of variational theory being applied. In the FORTRAN unit fu6 output, the first factor is denoted as κ^{ZCT} , κ^{SCT} , κ^{LCG3} , κ^{LCG4} and the second factor is denoted as $\kappa^{\text{TST/CAG}}$ and $\kappa^{\text{CVT/CAG}}$ for the TST and CVT levels of theory, respectively. Since this CAG factor is exactly 1 in the case of μVT , it is not listed in the output. The final result is denoted as, e.g., $\kappa^{\text{CVT/ZCT}}$ for a transmission coefficient evaluated with the ZCT (or MEPSAG) method at the CVT level.

The CAG factor is *required* for consistency in TST and CVT calculations (but not in μVT calculations), although it is often close to unity. The CAG correction was introduced in Refs. 10 and 44 of Sect. 22, and the reader is referred to those references for further discussion.

The MEPSAG, CD-SCSAG, and LCG3 or LCG4 transmission coefficients are simply denoted ZCT, SCT, and LCT, respectively, in recent papers, as discussed in Section 6.D above. The ZCT method should be considered simply as an approximation to SCT to be used when insufficient information is available to compute the reaction-path curvature. The SCT transmission coefficient is always larger than (or equal to) the ZCT one. The SCT transmission coefficient is expected to be larger than and more accurate than the LCT one when reaction-path curvature is small over the classically forbidden segment of the reaction path; the LCT transmission coefficient is expected to be larger than and more accurate than the SCT one when reaction-path curvature is large in the classically forbidden section of the $V_a^{\text{G}}(s)$ curve. These two situations are discussed in more detail in discussion remarks at Faraday Symposium 29 on Potential Energy Surfaces and Organic Reaction Paths [Ref. 45 in Sect. 22], and they are illustrated (in color) in an earlier review article [Ref. 46 in Sect. 22].

Whenever there is enough information available to calculate LCT transmission coefficients there is also enough data available to calculate SCT ones (although the converse is not true). In such cases the recommended best estimate is obtained by the

microcanonical optimized multidimensional tunneling (μ OMT) approximation that is discussed in Section 6.M.2 below.

The optimized multidimensional tunneling approximations are reviewed below. In particular, Subsection 6.M.1 reviews the canonical optimized multidimensional tunneling (COMT) method, and Subsection 6.M.2 discusses the microcanonical optimized multidimensional tunneling method (μ OMT).

6.J. Reaction rates including tunneling

In this subsection, we describe the calculation of the reaction rate constants as discussed in Subsection 6.A.

For a given temperature T , various transmission coefficients $\kappa(T)$ are computed using methods selected by the user, all of which lead to different approximations to the rates. Next, the s -independent quantities are determined, i.e., the electronic, translational, rotational, and vibrational partition functions for reactant(s) and product(s), as well as the electronic partition function for the generalized transition state and the prefactor for the rate constant.

Next, for each s at which a generalized normal mode analysis was performed the program computes the rotational and vibrational partition functions of the generalized transition state to get the generalized free energy of activation at this value of s . Using the nearest points to the maximum, parabolic and quartic fits are used to obtain the interpolated maximum of $\Delta G_f^{GT,0}(T,s)$. The final location and generalized free energy of activation at this maximum are obtained from the quartic fit (the parabolic fit is just for comparison).

The calculation of the kappa factors is described next. The Wigner tunneling correction for conventional TST is also computed, but only for informational purposes. It is not a recommended method because it is just the first two terms in a slowly convergent or divergent series. For thermal reaction rates, the MEPSAG, CD-SCSAG, and LCG3 or LCG4 transmission probabilities are averaged over a thermal distribution and divided by the thermal average of the classical transmission probabilities. As mentioned previously, CD-SCSAG, LCG3, and LCG4 methods take into account the coupling between the

reaction path curvature and the vibrational modes. In this version of *POLYRATE*, the LCG3, LCG4, and CD-SCSAG methods treat vibrational modes harmonically. (The current version of the program will not perform LCG3 or LCG4 calculations if anharmonic options other than the hindered-rotator method are selected. In the case of CD-SCSAG, zero-point energies and vibrational partition functions can be treated using one of the various anharmonicity methods, and the CD-SCSAG kappa factor will change due to the change in the adiabatic ground state potential. However, the CD-SCSAG method is still based on a harmonic approximation.) The procedures for calculating the transmission probabilities and transmission coefficients for the MEPSAG method are described completely in the GTST book chapter, and the description of the CD-SCSAG method can be found in Refs. 29 and 31 of Sect. 23. For the LCG3 or LCG4 calculation the basic idea is in the GTST book chapter and additional details of the implementation are described in Refs. 21, 29, 31, and 34 of Sect. 23. (Note that the value of s along an LCG3 or LCG4 tunneling path is a continuous function of progress along the path, as described correctly in the GTST book chapter and not as described in Ref. 21 of Sect. 23.)

Early versions of *POLYRATE* used a method called SCSAG. This has now been replaced by CD-SCSAG. The two methods are identical if the reaction-path curvature has only one nonzero component; otherwise they differ, and CD-SCSAG is based on a more correct analysis of the geometry of corner-cutting tunneling.

In the *POLYRATE* program, the Kronrod quadrature for MEPSAG and CD-SCSAG tunneling calculations is performed using the independent variable s and abscissa g_i rather than y and y_i ; see page 99 of the GTST book chapter.

Furthermore, for both the $\theta(E)$ integral over s , eq. (94), and the Boltzmann average, eq. (91) of the GTST book chapter, the range of integration is divided into 1-3 equal segments, and the same order of Kronrod quadrature is used in each. For the $\theta(E)$ integral we use NSEG2 segments with $(2*NQTH+1)$ -point Kronrod quadrature in each, and for the Boltzmann integral we use NSEG segments with $(2*NQE+1)$ -point Kronrod quadrature in each. Kronrod's method provides $NQTH$ -point and NQE -point Gauss-Legendre quadrature results as well. It is important to check the convergence of these integrals by comparing the N -point and $(2N+1)$ -point results.

The CD-SCSAG and LCG3 or LCG4 transmission probabilities are also used to calculate optimized multidimensional tunneling approximations called COMT and μ OMT, as discussed in Sections 6.M.1 and 6.M.2.

Using the various generalized transition state approximations and $\kappa(T)$ factors, various forward and reverse rate constants at temperature T can be obtained, e.g., the combination of an CVT rate constant with an MEPSAG tunneling correction yields an CVT/MEPSAG rate constant.

If the VTST-IOC correction is selected, the program updates all the energy, frequency, and determinant of moment of inertia information to the corrected values and recalculates the tunneling probabilities. A new set of corrected rate constants is then obtained.

If the VTST-ISPE correction is selected, the program spline fits the differences of the lower-level energies and the higher-level energies using the IVTST-M algorithm. The interpolated values are then added on to the lower-level and a new set of corrected rate constants is obtained. See Section 10.J for a detailed description.

6.K. Imaginary-frequency modes transverse to the reaction path

When a generalized normal mode eigenvalue is negative, i.e., the harmonic frequency of a mode transverse to the reaction path is imaginary (recall that the eigenvalues are squares of the frequencies), the program takes the following actions:

1. When an independent-normal mode method is being used, the vibrational partition function of that mode is set equal to 1.0×10^{10} .
2. The zero-point energy of that mode, as required for the calculation of the vibrationally adiabatic ground-state potential curve for tunneling calculations, is set equal to zero.
3. The turning point distance of that mode is set equal to 1.0×10^{10} bohr for the LCG3 or LCG4 calculations.
4. For the CD-SCSAG method, no contribution from this mode is included whether the harmonic or anharmonic approximation is selected.

6.L. Equilibrium constant calculation

In the current version of the code this calculation is the last to be performed. However, the calculation is carried out in two different subroutines, TSOUT and FINOUT. If a TST-only calculation is requested, then all the rate calculations are carried out in the subprogram TSRATE, which calls TSOUT. TSOUT prints the rate constants to the fu6 file and the fu15 (summary) file, and it calculates and prints the equilibrium constants using the TST rate constants.

The subprogram RATE, which calls FINOUT, calculates the rate constants with canonical variational theory (CVT). FINOUT computes and prints final rate constants, activation energies, and equilibrium constants (from the CVT rate constants).

In both cases the equilibrium constants are the last quantities calculated in the respective subprograms.

6.M. Omitting the product

POLYRATE can be run without calculations on the product if reverse rate constants and equilibrium constants are not calculated. This requires specifying the FORWARDK keyword in the RATE section of the fu5 input.

What the code will do then depends on whether there is a WELLP section in the fu5 input. If there is a WELLP section, the interpolation extends to the product-side well as described at Section 3.1 of Ref. 48 of Section 23. If there is not a WELLP section, then the interpolation extends to the last nonstationary point on the product side. Thus $V_{\text{MEP}}(s)$ is cut off (down to the product value) at that point, and if tunneling extends into the region beyond the last nonstationary point on the product side, it will be overestimated.

6.N. Optimized multidimensional tunneling calculations

6.N.1. COMT method

The canonical optimized multidimensional tunneling (COMT) temperature-dependent transmission coefficient is the larger of the thermally averaged temperature-dependent transmission coefficients calculated by the CD-SCSAG method and the LCG3 or LCG4 method at a given temperature. The COMT method was first used in Ref. 28 of Sect. 23 (where it was simply called MT). It was first called COMT in Ref. 34 of Sect. 23.

6.N.2. μ OMT method

The microcanonical optimized multidimensional tunneling (μ OMT) temperature-dependent transmission coefficient is the ratio of the Boltzmann average of the μ OMT quantum transmission probability to the Boltzmann average of the classical transmission probability, where the μ OMT quantum transmission probability is the larger of the energy-dependent transmission probabilities evaluated by the CD-SCSAG method and the LCG3 or LCG4 method at a given energy. This method was introduced in Ref. 34 of Sect. 23. See Also Ref. 45 in Sect. 22.

6.O. Interpolated VTST and mapped interpolation

Calculating $V_{\text{MEP}}(s)$, $|\mathbf{I}(s)|$, and $\omega(s)$ as needed for variational transition-state theory and tunneling calculations from *ab initio* electronic structure theory can be computationally demanding. Thus, it is sometimes desirable to interpolate these quantities based on electronic structure data only at the reactant, product, saddle point, and optionally a small number of additional points (close to the saddle point) along the MEP. Variational transition-state theory based on interpolated data at a few points on the MEP [Ref. 43 in Sect. 22 and Refs. 26, 32, and 48 in Sect. 23] is called interpolated VTST or IVTST.

Three implementations of IVTST are included in this version of the *POLYRATE* program. The first is zero-order IVTST (IVTST-0) and the second is first-order IVTST (IVTST-1) using the global interpolation approach. The third is IVTST-M which is described in Section 6.O.2 and 6.O.3. Permutation of frequencies is not included in any of these methods. TST rates and ZCT transmission coefficients can be calculated with IVTST-0;

in addition CVT rates and SCT transmission coefficients can be calculated with IVTST-1. Moreover, IVTST-M may be used for TST, CVT, μ VT, and CUS rates and ZCT and SCT transmission coefficient calculations.

6.O.1. IVTST-0 and IVTST-1

IVTST-0 and IVTST-1 calculations [Ref. 26, Sect. 23] can be performed for bimolecular reactions by using the IVTST0 or IVTST1 keywords in the GENERAL section. Alternatively, IVTST-0 calculations can be performed using file fu50. Presently IVTST-0 and IVTST-1 methods are only available for bimolecular reactions with two reactants and two products.

The IVTST-0 method assumes that information is only available at three locations on the potential energy surface, namely at the reactant (R), saddle point, and product (P). Details of the interpolation procedures can be found in Ref. 26 of Sect. 23. The IVTST-0 rate constant is the conventional TST rate constant with ZCT tunneling. Note that ZCT tunneling is the only available tunneling method for IVTST-0. ZCT tunneling calculations for zero-order IVTST are carried out using

$$V_a^G(s) = V_{\text{MEP}} + \text{ZPE}(s),$$

where $V_{\text{MEP}}(s)$ is represented by an Eckart function and $\text{ZPE}(s)$ is obtained from interpolated ω values.

The IVTST-1 method requires information for one extra point ($s = s_1$) near the saddle point in addition to the saddle point, R, and P. The $V_{\text{MEP}}(s)$ curve for the IVTST-1 calculation is also approximated by an Eckart function with a constraint that the Eckart function goes through the extra point at $s = s_1$. To determine the location of the variational transition state, the forward standard-state free energy of activation profile, $\Delta G_f^{\text{GT},0}(T,s)$ is calculated using the interpolated $V_{\text{MEP}}(s)$, $|\mathbf{I}(s)|$, and $\omega(s)$. The SCT transmission coefficients can be obtained for first-order IVTST using the interpolated $\mu_{\text{eff}}(s)$.

IVTST-0 calculations can also be performed by using the keyword ICOPT in the GENERAL section and the keyword OIVTST in the unit fu50 input (see Subsection 12.F). In the current version, this method is available only for bimolecular reactions with two products, and therefore the keywords MEPTYPER and MEPTYPEP for the unit fu50 input

should both be set to 2. In addition, the keywords MEPTYPER and MEPTYPEP should not be set to *well*, and the keyword FREQMAT should not be used. The hindered rotor approximation is currently the only anharmonicity option that can be used with IVTST-0 when using file unit fu50 for the IVTST-0 calculation. At this time there are no anharmonicity options for IVTST-0 nor IVTST-1 calculations when using file unit fu29.

For the IVTST-0 calculations, only conventional transition state theory rate constants and ZCT transmission coefficients can be calculated. Variational transition state theory at the CVT level and both ZCT and small-curvature tunneling (SCT) transmission coefficients can be obtained for IVTST-1. Details can be found in Ref. 26 of Sect. 23.

6.O.2. Interpolated VTST by mapping (IVTST-M)

IVTST-M is a method for obtaining the information required for a VTST calculation all along the reaction-path from the information available at a small number of points. The only limitations on using the IVTST-M algorithm are that a saddle point exists and reaction path information is available for least one nonstationary point on each side of the saddle point. In addition, as in all IVTST methods, information is required for 3 – 5 stationary points (reactants, saddle point, products, and optionally one or two wells). The level of IVTST-M is denoted by two numbers, H and G , indicating respectively the number of nonstationary points at which Hessians are available and the number of nonstationary points at which gradients are available. Thus, IVTST-M-6/54 indicates that the interpolation is based in 6 nonstationary Hessian points, at which geometries, energies, gradients, and Hessians are available, and 48 points (which is 54 minus 6) for which only geometries, energies, and gradients are available. The mapping procedure transforms $V_{\text{MEP}}(s)$, $\mathbf{II}(s)$, $\omega(s)$, and $B_{m,F}(s)$ from functions of s into functions of z where z is a new variable which always has a finite value, whereas s can have infinite values if either forward or reverse reaction is bimolecular. The parameter z is determined as

$$z = \frac{2}{\pi} \arctan\left(\frac{s - s_0}{L}\right), \quad (1)$$

with s_0 and L being two parameters obtained from the forward and reverse barrier heights [Ref. 48 in Sect. 23].

The new $V_{\text{MEP}}(z)$, $\mathbf{II}(z)$, $\omega(z)$, and $B_{m,F}(z)$ functions are interpolated by using splines under tension.

The method employs different approaches for the interpolation of $V_{\text{MEP}}(s)$, depending on the number of reactants and products in the reaction. If the reaction is unimolecular in either the forward or reverse direction, a cubic polynomial for estimating 10 energies between the reactant (or product) and the first (or last) point on the reaction path (or between the reactant and the first point and between the product and the last point) is calculated before carrying out the spline under tension fit. The cubic polynomial can be function of either s or z . The user should base the choice of s or z on experience or should test the two choices (using function of s or z) to see which one gives reasonable potential energy curve. The user can also choose not to use the polynomial to generate the 10 extra energies; however, the default is to generate them by using a cubic polynomial function of z . If the forward reaction is bimolecular, the interpolation on the reactant side of the saddle point is based on the energies from the input plus 10 additional energies estimated from an Eckart polynomial with a range parameter which is a function of s . Similar considerations apply on the product side if there are two products.

If for a bimolecular reaction or a reaction with two products, there are wells on the reactant and/or product side, the interpolation can be done using the information on these stationary points with the unimolecular procedures, which at least in principle can make the interpolation more accurate, especially when the well is deep and/or close to the saddle point. But if a well is present, it can also be ignored, and the interpolation can be carried out using reactants and/or products. If the well is ignored, then since for bimolecular reactions (or bimolecular sides of the reaction path) the interpolation is based on an Eckart potential, no point on the reactant side of the reaction path can be used that has an energy lower than the reactants energy, and no point on the product side can be used that is more stable than the products.

The IVTST-M algorithm has been implemented to be used with the hooks and the electronic structure input files, unit fu30, fu31, and fu40. The method is fully operational when using the hooks and the unit fu31 input file. When using units fu30 and fu40, only the IVTST-M default options are available. To use the IVTST-M algorithm with unit fu30, LOPT(2) must be set equal to -1 ; and with fu40, MAXLPTS must be set equal to -1 .

Whenever IVTST-M calls TSPACK for a spline-under-tension fit, it sets the variable IENDL equal to 3. This means that the two extra pieces of data needed for the spline fit are generated by the ENDSLPL subroutine using a local spline under tension.

6.O.3. The mapped interpolation (MI) algorithm with the hooks

As mentioned in Section 1 of this manual, the mapped interpolation (MI) algorithm can be used in two ways by *POLYRATE*. The first way is to perform IVTST-M calculations using electronic structure input file fu30, fu40, or fu31. This method is described fully in Sections 6.N.2, 10.J, and 11.D of this manual. The second way is in conjunction with the hooks; this requires some additional comments, which are given in the rest of this section.

A calculation with the hooks involves two stages. At the end of the first stage, one has stored s , $V_{\text{MEP}}(s)$, and the gradients on the fine grid and s , $V_{\text{MEP}}(s)$, $\omega(s)$, $B_{mF}(s)$, and $|\mathbf{I}(s)|$ are stored on the save grid. (The grids are defined in Sect. 7.A) Both grids extend from $s = \text{SLM}$ on the reactant side to $s = \text{SLP}$ on the product side, and there are $G + 1$ points (including the saddle point but excluding reactants and products) on the fine grid and $H + 1$ points (including the saddle point but excluding reactants and products) on the save grid. Then mapped interpolation is carried out to add $\text{EXNSTEP} * \text{EXSTEP}$ points on each side of the grids. Furthermore SLM is replaced by $\text{SLM} - \text{EXNSTEP} * \text{EXSTEP}$, and SLP is replaced by $\text{SLP} + \text{EXNSTEP} * \text{EXSTEP}$. As should be clear from Section 12.A.7, EXNSTEP and EXSTEP may actually be different on the two sides.

6.P. Interpolated optimized corrections (VTST-IOC), a form of dual-level VTST

In VTST calculations with approximate potential energy surfaces, it is sometimes desirable to correct the classical energies, vibrational frequencies, and moment-of-inertia determinants of some points along the MEP (e.g., the reactant, saddle point, product, van der Waals complex, ion-molecule complex, etc.) with accurate (higher-level) results. This is accomplished by applying correction procedures to the calculated energy, frequencies, and moment-of-inertia determinant along the MEP. The corrections are calibrated such that the corrected results match the accurate values at those selected points, and they correspond to interpolating these corrections at other points. When the corrections involve data from higher-level optimizations of the stationary points, the method is called VTST-IOC, a special case of VTST-IC. The original correction procedure for frequencies is denoted ICA, and two improved procedures are denoted ICR and ICL. (In general we recommend the ICL procedure.) The ICA procedure was introduced in version 5.1 of *POLYRATE*, and the ICL and ICR procedures were introduced in version 7.0.

All three procedures are based on the correction at 3 points, in particular, the saddle point and two stationary points, one on each side of MEP.

There are also two procedures for correcting $V_{\text{MEP}}(s)$ in IOC calculations: the original method, which was called NOECKART in versions 7.2 – 7.9.2 of *POLYRATE* but now is called the single-Eckart (SECKART) method of interpolating V_{MEP} , and the newer method, which was called ECKART in versions 7.2 – 7.9.2 but now is called double-Eckart (DECKART). The ECKART option was introduced in version 7.2. As implied by the names of the correction, is either a single Eckart function or a difference of two Eckart terms. If there is an appreciable well on either side of MEP, it is recommended that one use the molecular complex corresponding to the well as the stationary point on that side of the saddle point; otherwise it is recommended that one use the reactant or the product. However, if the well on the MEP is not very deep or if accurate data for the complex corresponding to the well are not available, the user can neglect the well and choose an appropriate functional form for the correction function as if there were no well at all.

As discussed in Section 1, the VTST-IOC method can be used either with an analytic potential energy surface or with electronic structure input data for the lower-level surface. In either case, the information calculated from the potential energy surface or read from the electronic structure input file will be corrected by the higher-level values.

The functional forms we use for the correction functions are an Eckart function, a hyperbolic tangent, a cut-off Gaussian, or a cut-off hyperbolic tangent, depending on the shape of MEP under consideration and the relative correction values at the 3 stationary points. See Refs. 36 and 44 of Sect. 23 for details. Details for the ICA procedure are available in Subsection 10.H and in Ref. 36 of Sect. 23 and details for the ICR and ICL procedures are available in Subsection 10.H and in Ref. 44 of Sect. 23. Details of the double Eckart procedure are available in Ref. 44 of Sect. 23.

If the higher-level data is obtained at electronic structure level X, and the lower-level data is obtained at level Y, the VTST-IOC calculation may be denoted X///Y. The triple slash notation is similar to the double slash notation that is widely used in electronic structure calculations. In particular, // means “at a stationary point geometry optimized by...” whereas /// is loosely translated as “along a reaction path calculated by...”. More specifically, though, we reserve the IOC and triple slash notations for algorithms where the higher-level information supplied to “correct” the transition state data involves a higher-

level geometry optimization. If only single-point energies are used, the ISPE and double slash notation should be used.

6.Q. Interpolated algorithm for large-curvature tunneling calculations

In LCT calculations, for each energy, a tunneling path is defined as a straight line in isoinertial coordinates between two reaction-coordinate turning points lying on the MEP, on the reactant side and on the product side of the maximum of the vibrationally adiabatic potential. This straight-line is divided into three zones, two of them are in the reactant and product valleys, respectively, and they are vibrationally adiabatic, and another zone describes the middle zone of the straight-line path and is vibrationally nonadiabatic. In the interpolated LCT (or ILCT) algorithm, every time the potential energy is needed along the nonadiabatic segment of the tunneling path, this energy is calculated by means of the spline fit to a number of already calculated points. Details of the methods are available in Ref. 55 of Sect. 23.

7. PROGRAM STRUCTURE

7.A. Overview of program flow

As mentioned in the Introduction, the *POLYRATE* program has several possible ways to obtain the reaction-path information necessary for rate constant calculations. The first option is to calculate this information directly from a set of interface subprograms called hooks. The second is to read this information from an electronic structure input file (fu30, fu31, or fu40) consisting of geometries, gradients, Hessians, etc. at points on the MEP obtained from electronic structure calculations. The third option is to read the information from a restart file which has been saved from a previous *POLYRATE* run. The fourth option is to read in higher-level “corrections” at selected points along the MEP; these corrections are read from file fu50 for VTST-IOC and from file fu51 for VTST-ISPE. Finally, for the IVTST calculations, electronic structure data at selected points may be read from file fu29 (IVTST-0 and IVTST-1) or fu50 (IVTST-0). These options are implemented in the *POLYRATE* program as follows.

First, the MAIN program reads all the keyword input. These keywords are analyzed for possible conflicts (i.e., unsupported combinations of options) in subroutine OPTION. If it is a restart run then the program reads in the restart information in RESTOR, and the rate constant calculation proceeds as described in the last paragraph of this section.

Otherwise, all other input data from unit fu5 are read in and checked by READ5 and REDGEO.

The GENERAL section of keyword input (Section 12.A.2) is always read first. Depending on the options chosen the properties of the reactants, products, and generalized transition states are calculated by calls to the hooks or read in from an electronic structure input file (or both). The user can choose at most one of the two switch keywords IVTST0 and IVTST1, or use the DL keyword. This choice plus the choice made for the POTENTIAL keyword in the ENERGETICS section (Section 12.A.3) determines the basic outline of the rest of the calculations as indicated in the following table. That table assumes that the hooks call SURF and SETUP, which is the standard way to use the hooks when *Polyrate* is used as a stand-alone code. (Subprograms SURF and SETUP are described in Subsection 8.C, and a more complete description of the hooks is provided in the rest of Section 8). In the present version of the code, user can also turn on the VRC option, which is off by default, to calculate rate for barrierless association reactions using a variable reaction coordinate and multifaceted dividing surfaces. When VRC is turned on, potential energies

can only be calculated through hooks since the Monte Carlo method is used for the variable-reaction-coordinate calculations with single-faceted or multifaceted dividing surfaces. Gradient and Hessian are not required for all Monte Carlo samples.

DL	IVTST0	IVTST1	POTENTIAL	Action
<i>None</i>	off	Off	<i>hooks</i>	Single-level calculation based on whole reaction path calculated from an analytic PES.
<i>none</i>	off	Off	<i>unit30, unit31, or unit40</i>	Single-level direct dynamics based on whole reaction path calculated from an electronic structure input file on unit fu30, fu31, or fu40
<i>none</i>	on	Off	<i>unit29 or unit50</i>	Single-level direct dynamics IVTST-0 calculation based on electronic structure input data from unit fu29 or fu50
<i>none</i>	off	On	<i>unit29</i>	Single-level direct dynamics IVTST-1 calculation based on electronic structure input data from unit fu29.
<i>ic</i>	off	Off	<i>hooks</i>	Dual-level direct dynamics with lower-level data from analytic PES and higher-level data from unit fu50
<i>ic</i>	off	Off	<i>unit30, unit31, or unit40</i>	Dual-level direct dynamics with lower-level data from unit fu30, fu31, or fu40 and higher-level data from unit fu50
<i>ispe</i>	off	Off	<i>hooks</i>	Dual-level direct dynamics with lower-level data from analytic PES and higher-level data from unit fu51
<i>ispe</i>	off	Off	<i>unit30, unit31, or unit40</i>	Dual-level direct dynamics with lower-level data from unit fu30, fu31, or fu40 and higher-level data from unit fu51

In the analytic potential energy surface case, the equilibrium geometries and properties of the reactant and product species are optionally obtained in subroutine REACT. For polyatomic species, REACT calls POLYAT, and the user must supply an estimate of the equilibrium geometry. POLYAT calls NEWT to perform a Newton-Raphson search for the location where the first derivatives of V are zero, CENTER to place the origin at this location and evaluate the determinant of the moment of inertia tensor, and NORMOD to do a vibrational analysis. The vibrational analysis involves determining the normal mode eigenvectors and frequencies, and, if desired, one of several procedures for including anharmonicity is invoked in ANHARM. If a diatomic species is treated as a Morse oscillator (DIATOM keyword) REACT calls DIATOM, and the equilibrium geometry, dissociation energy, and range parameter must be supplied by the user.

Next, if using the RP-VTST option without NOSADDLE, the program will determine the saddle point geometry and properties for the tight transition state. This is done in SADDLE and MAIN, which in turn call the same subroutines used by POLYAT. The program will handle 0 or 1 saddle points. For one saddle point the user must supply an approximate geometry for its location. For 0 saddle point, user can choose NOSADDLE option, which requires the user to input an initial guess for a point on the MEP such that the variational transition state lies between this point and the product. However, rather than using NOSADDLE, the VRC option in GENERAL section is recommended for barrierless association reactions, and this option does not require to input any geometry for transition state.

When using the hooks, *POLYRATE* starts either at the saddle point just determined or from an input starting-point geometry, then computes the minimum energy path (MEP) by a choice of several methods (done in the PATH section of file fu5). The distance along the MEP is called s ; the user can make s be zero at the saddle point, negative in the reactant region, and positive in the product region. The initial step can be based on a cubic expansion of the potential energy surface determined from the normalized eigenvector of the imaginary-frequency normal mode and from the curvature vector, both of which are calculated at the saddle point, or the initial step may be taken along the normal coordinate for the unbound motion at the starting point. When the default ESD method is used (RPM keyword set equal to *esd*), the negative of the gradient in isoinertial coordinates is followed using a fixed gradient step size DEL (specified by the SSTEP keyword). At regular intervals (specified by the INH keyword) along the MEP, the program computes

the generalized-transition-state properties necessary to evaluate the generalized free energy of activation and to perform semiclassical tunneling corrections. (We emphasize that both DEL and the regular intervals are measured in the isoinertial coordinates. Thus the units are sometimes called mass-scaled angstroms instead of angstroms. Thus if REDM, specified by the SCALEMASS keyword which is described in Subsection 12.A, is changed, DEL and the regular intervals will have different meanings.) Note that the grid determined by DEL is called the fine grid or the gradient grid. The grid defined by the INH keyword is called the save grid or Hessian grid. This is discussed further in Subsection 10.D. The moment of inertia calculation and Miller-Handy-Adams projections (eq. 31 of Ref. 5) or curvilinear-reaction-coordinate projections (eq. 17 of Ref. 41) are carried out in PROJECT; the vibrational analysis and the calculation of the effective mass for tunneling are done in NORMOD. In addition, if there is a saddle point the mapped interpolation (MI) algorithm can be used to interpolate the energy, the frequencies, the determinant of the moment of the inertia tensor, and the curvature components to extend one or both ends of the MEP for tunneling calculations. (See Sections 6.N.3 and 8.A for more details.) When there is no saddle point a simple exponential extrapolation of the classical and adiabatic energies and the other quantities used for tunneling calculations can be used at one or both ends of the MEP. (See Sect. 10.A for more details.) The EXFIRST and EXSECOND keywords in the PATH section control interpolation (extrapolation) options. Interpolation (extrapolation) is especially useful in situations where a minimum is reached in the classical potential before the adiabatic barrier is low enough for computing converged tunneling probabilities. Notice that extending the grid is sometimes called extrapolation, but it is also called interpolation because one used the reactant or product information that technically it is indeed interpolation.

When the MEP calculation is completed, the reaction-path information can be saved by writing it to unit fu1 for later restart runs. We note that there is an upper limit of NSDM points at which the reaction path information can be saved in a given *POLYRATE* executable file. The parameter NSDM is defined in the file aamod.f90 (see Subsection 5.A.2).

In the electronic structure input file case, the geometries, energies, vibrational frequencies or harmonic force constant matrices, and (optionally) higher-order force constants of the reactant(s), product(s), saddle point, and generalized transition states are read in from unit fu30, fu31, or fu40. (A more detailed statement of the available input options is given in Section 9.) The number of points on the MEP available from electronic structure

calculations is usually small; thus when converging tunneling probabilities it is sometimes necessary to extend the reaction path beyond the last input data point on one or both sides of the MEP. This is done by specifying larger values for the SLM and SLP keywords in the PATH section. The special interpolation procedures are described in Subsections 9.C and D.

The generalized free energy of activation is then computed at each of the points on the save grid (at which the reaction path information was saved), and the CVT rates are calculated for an input set of temperatures (done in RATE). We find the maximum generalized free energy of activation by local interpolation using 5-point Lagrangian interpolation. The result calculated with 3-point Lagrangian interpolation is printed as a check.

Various tunneling corrections can also be included by a call to KAPVA. The final output of forward and reverse reaction rate constants with and without the various tunneling corrections, as well as output of activation energies, analyses of bottleneck properties, equilibrium constants, and the ratio of the CVT to the conventional transition state theory rate constants, are provided by FINOUT.

If the VTST-IOC option is selected (DL keyword set to *ic* or ICOPT keyword specified), the program will then update (done in ZOCUPD) all the MEP information based on corrections calculated from the higher-level values read from the unit fu50 input file and calculate a new set of rate constants based on the corrected values by the same procedures as described above.

If the VTST-ISPE option is selected (DL keyword set to *ispe*), the program will update the energies along the MEP based on a spline fit to the differences of the lower-level energies and the higher-level energies read from the unit fu51 input file. A new set of rate constants is then calculated, based on the corrected energies, by the same procedures described above.

If the IVTST-0 method is selected, after the MAIN program reads all the information provided at reactants, products, and saddle point, subroutine GIVTST is called. Then the process of generating the information required at points along the reaction path by interpolation is started using specific routines for IVTST instead of the ones employed for calculations based on analytic potentials or fu30, fu31, and fu40 input files. If the

IVTST-1 method is selected, the electronic structure information at the stationary points plus the data at an extra point located close to the transition state is read by the MAIN program from file fu29 and then the subroutine GIVTST is called.

7.B. Determining the sign of s

If there is a saddle point, the reaction coordinate s is, by convention, positive on the product side of the saddle point and negative on the reactant side. In the program, however, the default is that s is defined as positive in the direction of the unbound normal mode at the saddle point, and it is defined as negative in the opposite direction. When one is using the hooks, the definition of the sign of s does not affect the calculated rate constants when LCG3 or LCG4 calculations including tunneling into excited states are not carried out, but it does affect the LCT probabilities when contributions from tunneling into excited states are included, since the current coding of that case requires consistency with the convention. When one is using fu30, fu31, or fu40, the choice of SIGN also affects whether SCT calculations or LCT tunneling into the ground state are correct, as explained in the next paragraph. Also, confusion might arise when one specifies the range of the reaction path to be calculated or discusses the location of the optimized variational transition state. We therefore advise users to override the default, if necessary, to follow the established convention of s being positive on the product side. The following procedure is recommended to ensure this. First, make a run without computing the path (do not include the PATH section) and without computing the properties of reactants and products (do not include REACT1, REACT2, PROD1, and/or PROD2 sections), and look at the normal mode eigenvector corresponding to the normal mode with imaginary frequency at the saddle point. If this vector points towards products, set the SIGN variable keyword in the PATH section to *product* in full runs. If not, set it to *reactant* (or omit it, since *reactant* is the default). There is another keyword, IDIRECT, in the PATH section which determines the direction of the first step from the saddle point in calculating the MEP. If SIGN is set correctly and if IDIRECT is set to 1 (which is the default), the MEP toward the product side (which is determined by the keyword SIGN discussed above) will be calculated first and the reactant-side MEP will be calculated after the limit on the product-side MEP is reached. If SIGN is set correctly and if IDIRECT is set to -1, the reactant-side MEP will be calculated first instead. The value of IDIRECT should not affect the calculated results.

In practice, the SIGN variable works differently with the hooks and with files 30, 31, and 40. When one is using the hooks, SIGN actually changes the sign of the value of s that is assigned to given geometry on the reaction path. When one is using the electronic structure input file options, one cannot change the sign of s . However, it is still necessary that SIGN be set correctly. If SIGN is set wrong, then the reaction path curvature at the saddle point will not be computed correctly. The correct setting for SIGN when one is using fu30, fu31, or fu40 is determined by the following rule: when using fu30, fu31, or fu40, if the positive direction of s in the output is not associated with the direction toward products, then change the choice for the SIGN variable in the PATH section of the input and repeat the run.

7.C. Restart options

There are two ways to restart calculations in *POLYRATE*.

The first restart option is controlled by the RESTART keyword in the GENERAL section. This option is designed to allow calculations at additional temperatures without recalculating the information stored on the save grid. This option uses fu1. If the VTST-IOC option is used with RESTART, one can change the lower level but not the higher level, i.e., the fu50 file cannot be changed for the RESTART run. Furthermore, if the original run was not a dual-level run, the restarted run cannot be a dual-level run.

The second restart option allows one to restart calculations without repeating all of the work for the stationary points (reactants, products, and/or saddle point). This restart option is controlled by the STATUS keyword in the REACT1, REACT2, PROD1, PROD2, and START section. Each of the input sections for the stationary points (REACT1, REACT2, PROD1, PROD2, and START) allows one to input optimized geometries, energies, and frequencies from previous jobs. The user also has the option of only calculating reactants, or only calculating products, or only calculating a saddle point to obtain the optimized geometries, energies, and frequencies for the given stationary point before proceeding with the rest of the calculation. Thus it is possible to optimize the stationary points in separate runs, and then read in the data for those stationary points in later runs to continue. To do this, the user can run *POLYRATE* with only the *GENERAL, *ENERGETICS, *SECOND, *OPTIMIZATION, *REACT1, and *REACT2 sections in the fu5 file. If the calculation is a supermolecule calculation then both reactant 1 and reactant 2 must be

specified in the *REACT1 and *REACT2 sections. However, if the calculation is not a supermolecule calculation then it is possible to calculate reactant 1 and reactant 2 separately using only the *REACT1 section and leaving out the *REACT2 section. Note that when calculating only reactant 2, the information for reactant 2 is placed in the *REACT1 section and the *REACT2 section is left out. The same scenario holds for products, i.e., one can include *REACT1 or *REACT1 and *REACT2 but leave out *START. If the user wishes to optimize the saddle point without optimizing the reactants and products then the saddle point information is placed in the *START section and *POLYRATE* is run with only the *GENERAL, *ENERGETICS, *SECOND, *OPTIMIZATION, and *START sections in the fu5 file. An fu61 file, which contains the optimized properties, is created after every *POLYRATE* run. The information on this file can be used to build a complete fu5 file or update an existing fu5 file with optimized information. Thus, effectively, one can optimize each stationary point separately or in any combination of stationary points, add the newly optimized information to an existing or a new fu5 file, and restart the calculation to optimize the remaining stationary points or complete the full calculation. See Section 12.A.6.

8. CALCULATIONS USING THE HOOKS

8.A. Methods for interpolating beyond SLM and SLP with the hooks

This section describes the procedures used when interpolating between the SLM end point on the save grid and the reactants and between the SLP end point on the save grid and the products, respectively. There are two procedures that are used. Note that SLM and SLP keywords are only for MEP, and invalid for variable reaction coordinate.

The first procedure is used when the reaction being studied has a saddle point. (In this case the SADDLE keyword in the PATH section is specified or taken as the default.) Reactions with saddle points either do not involve interpolation, or they use the mapped interpolation (MI) algorithm to extend the reaction path beyond SLM and SLP. The options for this interpolation are controlled by the EXFIRST and EXSECOND keywords in the PATH section. The MI algorithm is described in Sections 6.N.2, 6.N.3, and 10.J and in Ref. 48 of Sect. 23.

The second procedure is used when the reaction being studied does not have a saddle point. For this type of reaction the NOSADDLE keyword must be specified in the path section. Reactions without saddle points always use a two-point extrapolation to extend the reaction path beyond SLM and SLP. The options for this extrapolation are controlled by the EXFIRST and EXSECOND keywords in the path section. The two-point extrapolation is described in Section 10.A.

8.B. Zero of energy

In *POLYRATE* the variable EZERO always denotes the energy that corresponds to reactants in the potential energy subroutine, and it is used to convert the potential energy to the zero of energy that is used in the rest of *POLYRATE*. EZERO refers to the energy of a single reactant (for a unimolecular reaction) or the sum of energies of infinitely separated reactants (for a bimolecular reaction). In the language of potential energy surfaces, EZERO denotes the potential energy of the reactant or reactants at their classical equilibrium geometries. In the language of electronic structure calculations, EZERO denotes the sum of the electronic energy and the nuclear repulsion (for *ab initio* calculations). It is necessary to understand this variable in order to understand the input

options available with the EZERO keyword in the GENERAL section; the rest of this section provides the necessary explanation.

In *POLYRATE* the user must ensure that EZERO is set equal to a value consistent with the potential energy subprogram (when an analytic potential energy subprogram is used) or with the electronic structure data (when the electronic structure input file option is used). This can be accomplished in one of two ways: (i) by setting the EZERO keyword to *calculate* and letting the program calculate the zero of energy, or (iii) by setting the EZERO keyword to *read* and directly inputting a nonzero value for EZERO which is consistent with the potential energy subprogram.

Setting the EZERO keyword to be equal to *calculate* causes the program to calculate the zero of energy after the reactants have been optimized to their equilibrium geometries at "infinite" separation. Because the zero of energy is not computed until the optimization procedure, all energies for the reactant species which are printed to the long output file on unit fu6 before the zero of energy is computed will have an incorrect zero of energy. For a unimolecular system the zero of energy is set equal to the potential energy of the system at its optimized reactant equilibrium geometry. For a bimolecular system the zero of energy is set equal to the sum of the potential energies of the two reactants after both reactants have been optimized to their equilibrium geometries. This can be done with the SUPERMOL option by placing the two reactants at a large separation distance given implicitly by the two geometries specified by the GEOM keyword in the REACT1 and REACT2 sections of the unit fu5 input data file. (Note that, when the SUPERMOL option is used, this sum is given by the energy at the end of the second reactant's optimization).

8.C. User-Supplied analytic potential energy surface

8.C.1. Interface

The protocol of the subprograms for evaluating the potential energy from an analytic function is as follows. The user must supply two subroutines, SETUP and SURF, which accept and return all information from and to *POLYRATE* in atomic units. The subroutine SETUP has one integer type variable, n3tm, in its parameter list:

subroutine setup (N3TM)

In the argument list of the call statement to SETUP the *POLYRATE* code passes the variable N3TM where:

N3TM = $3 \times \text{NATOMS}$
(This variable is used to dimension the arrays for the Cartesian coordinates and the derivatives of the energy with respect to the Cartesian coordinates for a given system.)

NATOMS = maximum number of atoms in the *POLYRATE* code
(It is the actual number of atoms in the system set in the input file.)

SETUP may optionally open data file(s) which contain the potential parameters, read these parameters, set up any necessary data, and close any file(s) which were opened. SETUP is called once and only once in each complete run of the program.

Note that SURF and SETUP are called by the hooks. In particular, SETUP may be called by PREP and/or PREPJ, and SURF may be called by EHOOK, GHOOK, HHOOK, and OHOOK. No other routines call SURF and SETUP.

Users should dimension arrays in SETUP carefully since inconsistency of array dimensioning is the major potential source of error when passing information between subprograms. Also, users should resolve any conflict in the assignment of the input and output I/O units between *POLYRATE* and SETUP. For the I/O unit numbers used by *POLYRATE*, see Section 19. As noted there, unit 4 is reserved for use by the potential routines, i.e., by SETUP. If SETUP requires more than one file, unit 7 is recommended for the second file. Alternatively the FU# parameters in the aamod.f90 file can be modified to avoid conflicts between the FORTRAN units used in *POLYRATE* and those used by SETUP and SURF. In addition, users should not name subroutines with the same name as an existing *POLYRATE* subroutine. A list of subroutine names and a short description of the subroutines used by *POLYRATE* are given in subsection 17.A.1.

The subroutine SURF evaluates the potential and its first derivatives as functions of the Cartesian coordinates, and it has 4 elements in its parameter list. These are V, X, DX, and N3TM, as follows:

```
subroutine surf (V, X, DX, N3TM)
```

where:

V = potential energy in hartrees

X = array of dimension N3TM which contains the Cartesian coordinates in bohrs of all the atoms in the system

DX = array of dimension N3TM which contains the first derivatives (in hartrees per bohr) of the energy with respect to each of the Cartesian coordinates. DX should be determined analytically if possible as all higher-order derivatives needed by *POLYRATE* are calculated numerically from these first derivatives.

N3TM = as defined above

The arrays X and DX must be of dimension N3TM within the subroutine SURF.

The order of the coordinates is (x, y, z) for atom 1, (x, y, z) for atom 2, ..., (x, y, z) for atom N, where the order of the atoms is determined by the order used in the ATOMS list keyword in the GENERAL section. (see Subsection 12.A.2).

Note that whenever the potential is read from FORTRAN unit fu30 file, SURF and SETUP are not needed and may be replaced by their dummy versions, for example, those included in the dumpot.f file.

During the calculation of the MEP, the quantities V, X, and DX are optionally written (using the PRSAVERP keyword in the PATH section) by *POLYRATE* to the file linked to the FORTRAN unit fu6 for each point at which reaction path information is saved (each save grid point). Since the user is often interested in the values of additional quantities calculated by subroutine SURF (such as interatomic distances, curvilinear internal coordinates, or various contributions to the potential energy) at these save points, we also provide the array POTINF(NPOTPT) in rate_const module in the *POLYRATE* program as a means of passing such information from subroutine SURF to the main program, in which the array POTINF is optionally written to the file linked to the FORTRAN unit fu6. (This is controlled by the use of the POTINF keyword in the PATH section). The maximum number of pieces of data from the potential which may be printed at each save grid point is 99. Unlike the parameter lists described above, the use of the common block SURFCM in the

user-supplied SURF subroutine is user-driven. The information stored in the array POTINF is printed exactly as stored. Note that values assigned to the array POTINF have no effect on any values calculated by *POLYRATE*.

8.C.2. Utility routines

The *POLYRATE* program assumes that the user-supplied potential energy function subprogram SURF accepts the values of the atomic Cartesian coordinates and returns the potential energy and the derivatives of the potential energy with respect to the atomic Cartesians. However, not all potential energy functions are written in terms of atomic Cartesians; some potential energy functions are written in terms of bond lengths. With this in mind, four utility subprograms are provided with *POLYRATE*. Two of the utility subprograms are for three-body systems and the other two are for four-body systems.

Two types of utility subprograms are provided. One type converts atomic Cartesians to bond lengths, and the other uses the chain rule to convert the derivatives of the energy with respect to the bond lengths to derivatives with respect to the atomic Cartesians.

The subprograms COORD#, where # = 3 for a three-body system and # = 4 for a four-body system, convert the atomic Cartesians to bond lengths. In the case of COORD3, nine Cartesians are converted to three bond lengths. The order of the Cartesians and the bond lengths are as follows:

X(1) — X(3):	X, Y, Z for atom A
X(4) — X(6):	X, Y, Z for atom B
X(7) — X(9):	X, Y, Z for atom C
R(1):	R(A–B)
R(2):	R(B–C)
R(3):	R(C–A)

The Cartesians are passed to COORD3 through the parameter list (X, N3TM) (X and N3TM are defined in Subsection 8.C.1) and the bond lengths are passed to the potential subprogram through the module POTCM3:

```

module potcm3
  integer :: nsurf, nder, ndum(8)
  real(8) :: r(3),energy, dedr(3)
end module potcm3

```

COORD3 assumes that all the coordinates (Cartesians and bond lengths) are in atomic units. The subprogram also assumes that the potential is written such that the bond specifications are as defined above. For example, if the potential for ClHBr is written such that

```
R(1):          R(Cl-H)
R(2):          R(H-Br)
R(3):          R(Br-Cl)
```

the user must define Cl as the first atom (atom A), H as the second atom (atom B), and Br as the third atom (atom C) in the input data in unit fu5.

The subprogram COORD4 does the same calculation as COORD3 but for a four-body system. In COORD4 the coordinates are defined as follows:

```
X(1) — X(3):   X, Y, Z for atom A
X(4) — X(6):   X, Y, Z for atom B
X(7) — X(9):   X, Y, Z for atom C
X(10) — X(12): X, Y, Z for atom D
R(1):          R(A-B)
R(2):          R(A-C)
R(3):          R(A-D)
R(4):          R(B-C)
R(5):          R(B-D)
R(6):          R(C-D)
```

The subprograms CHAIN#, where # = 3 for a three-body system and # = 4 for a four-body system, convert the derivatives of the energy with respect to the bond lengths to derivatives with respect to the atomic Cartesians. In the case of CHAIN3, three derivatives with respect to the bond lengths are converted to nine derivatives with respect to the atomic Cartesians. The order of the derivatives is as follows:

```
DEDR(1):      DEDR for the bond length A-B
DEDR(2):      DEDR for the bond length B-C
DEDR(3):      DEDR for the bond length C-A
DX(1):        DEDX for atom A
DX(2):        DEDY for atom A
DX(3):        DEDZ for atom A
DX(4):        DEDX for atom B
DX(5):        DEDY for atom B
DX(6):        DEDZ for atom B
DX(7):        DEDX for atom C
DX(8):        DEDY for atom C
DX(9):        DEDZ for atom C
```

The derivatives with respect to the bond lengths are passed through the common block POTCM as defined above, and the Cartesian information is passed through the parameter list (X, DX, N3TM). This subprogram assumes the same definitions as COORD3.

The subprogram CHAIN4 performs the same calculation as CHAIN3, but for a four-body system with the following coordinate definitions:

DEDR(1):	DEDR for the bond length A-B
DEDR(2):	DEDR for the bond length A-C
DEDR(3):	DEDR for the bond length A-D
DEDR(4):	DEDR for the bond length B-C
DEDR(5):	DEDR for the bond length B-D
DEDR(6):	DEDR for the bond length C-D
DX(1):	DEDX for atom A
DX(2):	DEDY for atom A
DX(3):	DEDZ for atom A
DX(4):	DEDX for atom B
DX(5):	DEDY for atom B
DX(6):	DEDZ for atom B
DX(7):	DEDX for atom C
DX(8):	DEDY for atom C
DX(9):	DEDZ for atom C
DX(10):	DEDX for atom D
DX(11):	DEDY for atom D
DX(12):	DEDZ for atom D

These subprograms can be used with any three- or four-body potential, which is written in terms of bond lengths. For example, for a three-body potential made up of two subprograms, PREPOT and POT, where PREPOT does the initializations for the potential, i.e. performs the role of SETUP, and POT does the actual potential and derivative evaluations given the geometry in terms of bond lengths, the user could provide SETUP and SURF routines to link *POLYRATE* with PREPOT and POT.

The subprogram SETUP would simply call PREPOT. For example:

```
subroutine setup (n3tm)
  implicit double precision (a-h, o-z)
  call prepot
  return
end
```

The SURF subprogram would first call COORD3, then call POT, and then call CHAIN3 before returning to the *POLYRATE* program. For example:

```

subroutine surf (v, x, dx, n3tm)
implicit double precision (a-h, o-z)
use potcm3
dimension x(n3tm), dx(n3tm)
call coord3(x, n3tm)
call pot
v = energy
call chain3 (x, dx, n3tm)
return
end

```

8.D. Calculations using an electronic structure program through hooks

A very useful feature of the *Polyrate* program is its ability to link an electronic structure program to obtain reaction path information directly. All energetic information needed by *Polyrate* for single-level runs or for the lower level of a dual-level run is obtained in one of two ways. The first way is via a set of general hook routines. The second way is from an electronic structure file (fu29, fu30, fu31, or fu40). This section of the manual describes the hook routines.

The hook routines are actually quite general. They can be used to call an analytical potential surface, and this special case is described in section 8.C. Alternatively they could be used for direct links to electronic structure programs (as in MORATE, GAUSSRATE, GAMESSPLUSRATE, JAGUARATE, or NWCHEMRATE) or in any way that the user finds useful. In the case of the connection to an electronic structure program, the user is responsible for obtaining or writing the necessary routines that interface the hook routines and the electronic structure program. In some cases, e.g., MORATE, GAUSSRATE, GAMESSPLUSRATE, JAGUARATE, and NWCHEMRATE such interface routines have been prepared by *Polyrate* developers, but users are invited to prepare their own interface routines as well.

We note that historically *POLYRATE* called SURF and SETUP directly. Beginning with version 7.0, the hooks were inserted as an interface between the *Polyrate* program and the source of potential information. The goal is for *Polyrate* to be completely blind as to what happens on the other side of this interface. We refer to this as complete modularity

between the dynamics and the potential, and we believe that this modularity has been fully accomplished since version 7.3.

The hook routines are PREP, PREPJ, EHOOK, GHOOK, HHOOK, and OHOOK. The source code for these routines is located in the src/ subdirectory in hooks.f. This subdirectory also contains six utility subprograms (YTRANS, YSECEN, YSECEP, YDERV2, YDER24, and YNEWT). The role of each of the hooks is described below. Before proceeding with these individual descriptions, we first make a few general comments. The basic concept of the hooks is as follows: Whenever *Polyrate* needs an energy, it calls EHOOK. Whenever *Polyrate* needs a gradient (or an energy and a gradient at the same geometry) it calls GHOOK. Whenever *Polyrate* needs a Hessian (or a Hessian plus an energy and/or gradient, all at the same geometry), it calls HHOOK. *Polyrate* has its own geometry optimization routines, and it may employ them using energy, gradient, and/or Hessian information from calls to EHOOK, GHOOK, and HHOOK. However, in many cases it is more efficient to call external optimization routines. These can be accessed by calls to OHOOK. Finally, we note that PREP and PREPJ are available for initialization tasks, i.e., for passing geometry-independent information to the other side of the interface.

The versions of the hooks distributed with *Polyrate* itself only call SURF and SETUP, except for HHOOK, which calls GHOOK. This does not convey the full usefulness of the hooks. The hooks are better appreciated by studying one of the direct dynamics codes built on *Polyrate*, e.g., MORATE, GAUSSRATE, GAMESSPLUSRATE, JAGUARATE, and NWCHEMRATE. But the *Polyrate* versions of the hooks provide clear examples of their use that can be understood by all users, independent of whichever electronic structure codes they are familiar with.

Since version 7.3, HHOOK computes Hessians by taking numerical derivatives of gradients obtained by calls to GHOOK. The code is identical to the code for this purpose on the *Polyrate* side of the interface. As mentioned in the previous paragraph, this does not convey the full power of this hook. This hook is designed to allow the user to take advantage of analytic Hessians if they are available. If analytical Hessians are available, the user should modify HHOOK to use them. A similar generic version of OHOOK is also provided.

PREP:

PREP is the driver for all preparation routines needed to set up the hooks. This routine is called once after input file fu5 has been read in and before the table of input is written to the output file. This call occurs before any calculations are done. This routine may be used to call electronic structure preparation routines. File fu70 is reserved for use by PREP.

To interface with a new electronic structure program one just needs to add a new block to the 'if block' contained in this subroutine with a call to the appropriate routine. Note that this is an optional hook. Not all electronic structure programs need initialization. When PREP is not needed, one should use a dummy routine with a simple return command inside.

PREPJ:

PREPJ is the species-dependent driver for all preparation routines needed to set up the hooks. Any initializations that are dependent on whether the reactants, products, saddle point, or generalized transition state is computed are done here. PREPJ is called up to six times, namely it is called once each at the starts of the sections for REACT1, REACT2, PROD1, PROD2, START, and PATH. In other words, PREPJ is called before each respective reactant, product, or saddle point is studied and again before the reaction path is computed. A variable, JTYPE, specifies which particular species is being computed; see Section 19 on file usage.

To interface with a new electronic structure program one just needs to add a new block to the 'if block' contained in this subroutine with a call to the appropriate routine. Note that this is an optional hook. Not all electronic structure programs need any species dependent initializations. When the PREPJ is not needed, user should prepare a dummy routine with a simple return command inside.

EHOOK, GHOOK, HHOOK, and OHOOK always communicate with the *Polyrate* side of the interface using mass-scaled Cartesians in bohrs and energies in hartrees. In the versions of the hooks distributed with *POLYRATE*, either mass-scaled Cartesians or unscaled Cartesians may be used for communicating with SURF. In the versions of SURF that are distributed with *POLYRATE*, the energy is computed using the Cartesian coordinates, so the hooks first convert to unscaled Cartesians before calling SURF. Similar transformations

are required for gradients and Hessians; for example, SURF returns gradients in unscaled coordinates, but GHOOK converts them to mass-scaled coordinates.

EHOOK:

EHOOK is the energy interface subprogram of *Polyrate*.

To interface with a new electronic structure program, one needs to add a new block to the 'if block' with a call to an interface routine that accepts Cartesian coordinates (unscaled) and returns the energy. If the *read* option is chosen for the EZERO keyword, then the zero of energy should be consistent with the EZERO value specified in the input file. Remember that the user must also specify the new interface potential by adding another option to the POTENTIAL keyword and implementing the new possible value of the associated IPOT variable in the FORTRAN code.

GHOOK:

GHOOK is the gradient (first derivative) interface subprogram of *Polyrate*.

To interface with a new electronic structure program, one needs to add a new block to the 'if block' with a call to an interface routine that accepts Cartesian coordinates (unscaled) and returns the energy and first derivatives. The developer of a new interface must also specify the new interface potential by adding another option to the POTENTIAL keyword and implementing the new possible value of the associated IPOT variable.

HHOOK:

HHOOK is the Hessian (second derivative) interface subprogram of *Polyrate*.

For the second derivatives, the potential is specified by the HESSCAL keyword of the SECOND section of the fu5 input file and is coded in FORTRAN using the IPOT variable. If the keyword option *ghook* is used in the HESSCAL keyword, numerical difference methods from *POLYRATE* are applied; if the *hhook* option is used, then the second derivatives are taken from HHOOK. The latter option is designed to allow HHOOK to obtain analytic second derivatives from electronic structure routines. In the current version of *POLYRATE*, the *ghook* and *hhook* options are essentially the same. If the *hhook* option is used, the second derivatives are computed with the hooks routines YSECEN and YSECEP, whereas if the *ghook* option is used, the second derivatives are computed with *POLYRATE* subroutines SECCEN and SECCEP. However, the numerical

methods used in YSECEN and YSECEP are the same as those in SECCEN and SECCEP. HHOOK must convert the Hessian to mass-scaled coordinates before returning to the *POLYRATE* side of the interface.

To interface with a new electronic structure program, one needs to remove the subroutine calls in the ISPOT equals to 1 block and add a new block with a call to an interface routine that accepts Cartesian coordinates (unscaled) and returns the energy, first derivatives, and Hessian matrix. Note one must ensure that the gradient and Hessian conform to the same orientation. See subsection 12.A.4 for the description of the keyword options used to calculate second derivatives.

OHOOK:

OHOOK is the geometry optimization interface subprogram of *Polyrate*. This routine accepts an initial guess to the mass-scaled Cartesian coordinates of the full system and returns the optimized geometry in mass-scaled Cartesian coordinates and the energy. A variable, IOP, is used to specify which chemical species is being optimized: for IOP = 1, or 2; the chemical species is reactant 1 or 2, for IOP = 3, or 4; the chemical species is product 1 or 2; for IOP = 5, the chemical species is the saddle point.

The geometry optimization is specified by the OPTMIN and OPTTS keywords and is coded in FORTRAN using the IGPO variable. If the keyword option *bfgs* or *nr* is used for either of these keywords, IGPO is 0, and the optimization is performed by NEWT, which is a *POLYRATE* subroutine. The OHOOK option (IGPO = 1) is designed to allow the OHOOK routine to call an electronic structure package directly. In the generic version of the OHOOK routine, the option calls YNEWT, which is supplied as part of the generic hooks. (Notice that YNEWT is a slightly simplified version of the *POLYRATE* routine NEWT.) To interface with a new electronic structure program, one must replace the call to YNEWT when IGPO equals to 1 with a call to an interface routine that accepts Cartesian coordinates (unscaled) and returns the optimized geometry and energy. Notice that, in the current version of *POLYRATE*, the method used to optimize the minima and saddle point should be the same (i.e. either both from *POLYRATE* or both from electronic structure package). For the description of the keyword options used in geometry optimization, see subsection 12.A.5.

9. CALCULATIONS BASED ON AN ELECTRONIC STRUCTURE INPUT FILE

One major option of the *POLYRATE* program is the ability to use MEP information obtained from electronic structure calculations in the form of an input file instead of using an analytic potential energy surface. This input file will contain MEP information for a finite range of the reaction path, namely s in the interval $[SLM, SLP]$. It is the user's responsibility to ensure that these input geometries do indeed lie on the MEP or a good approximation to it.

When one of the electronic structure files (fu30, fu31, or fu40) is used the *POLYRATE* program is never supposed to make calls to the potential energy surface subprograms SETUP and SURF. However, to satisfy all external references when linking the program the subprograms SETUP and SURF must be provided. A potential energy surface file, dumpot.f, distributed with the *POLYRATE* program has been designed specifically for use with this option of the code. This potential energy surface file contains dummy SETUP and SURF routines which print a warning message and stop if subprogram SURF is called.

When the reaction-path information is calculated from electronic structure calculations and is used as input for dynamical calculations, the necessary information (listed later in this paragraph) is read from a separate data file which is linked to the FORTRAN unit fu30, fu31, or fu40 and is keyed to the normal input in the file which is linked to the FORTRAN unit fu5. Throughout this manual, unit fu30, unit fu31, and unit fu40 are called the electronic structure input files. The input data in unit fu5 remains the same as when the calculation is done with an analytic potential energy surface except that the POTENTIAL variable keyword in the GENERAL section is set to *unit30*, *unit31*, or *unit40*. A line-by-line description of the MEP input for unit fu30 is provided in Section 12.C, a description of the input when using unit fu31 is given in Section 12.D., and a description of the input when using unit fu40 is provided in Section 12.E. Using the information from unit fu30, unit fu31, and unit fu40, the quantities $V(s)$, $|\mathbf{I}(s)|$, $\omega(s)$, $k_3(s)$, $k_4(s)$, and $B_{mF}(s)$ are interpolated to locations on the save grid and are stored in save grid arrays to be used for Lagrangian interpolation as needed for finding variational transition states and as needed for tunneling calculations. The type of information used to obtain the tables of the MEP quantities is as follows:

Potential

$V_{\text{MEP}}(s)$ is always read directly.

Determinant of the moment of inertia tensor

$|\mathbf{I}(s)|$ is always computed from the geometry of the complex. For reactants and products at infinite separation $|\mathbf{I}(s)|$ is infinite, so for large $|s|$ the program interpolates the inverse of $|\mathbf{I}(s)|$ instead of $|\mathbf{I}(s)|$ itself. To obtain $|\mathbf{I}(s)|$ for finite s it is necessary to read the $3N$ Cartesian coordinates $R(s)$ or the mass-scaled ones $x(s)$.

Gradients

The gradient $\nabla V_{\text{MEP}}(s)$ must be read in. The gradient is denoted DX. The code has options to read either the gradient itself or the negative of the gradient, which is the force. In addition, the gradient or force may be in either unscaled or mass-scaled coordinates.

Frequencies

$\omega(s)$ can be read directly or computed from the force constant matrix $\mathbf{F}(s)$. If $\mathbf{F}(s)$ is read, both $\omega(s)$ and the coefficient matrix \mathbf{L} which diagonalizes \mathbf{F} (or the projected \mathbf{F}) are obtained. A total of $3N-7$ (for a nonlinear generalized transition state), $3N-6$ (for a linear generalized transition state), or $3N-1$ (for solid-state or gas-solid interface generalized transition state) frequencies or a $3N \times 3N$ force constant matrix is read.

Anharmonicities

Either the cubic and quartic force constants $k_3(s)$ and $k_4(s)$ are read directly, or they are omitted. If read, a total of $3N-7$ (nonlinear), $3N-6$ (linear), or $3N-1$ (solid-state or interface) of each will be read. Even if $k_3(s)$ and $k_4(s)$ are not read, anharmonicities can still be included by using the Morse I approximation.

Reaction-path curvature

The reaction-path curvature components $B_{mF}(s)$ can either be read in or they can be computed from the gradients. These curvature components are needed for SCT calculations but are not used for CVT or ZCT calculations.

The user should always supply the following information for all points on the Hessian grid: s , V_{MEP} , the geometry, the gradient, and either the frequencies or the Hessian. Depending on the options chosen, one or more of the following should also be supplied for all points on the Hessian grid: the frequencies, the Hessian (i.e., the matrix of second derivatives of the potential), the cubic force constants, the quartic force constants, and the curvature components. Alternatively, the curvature components can be calculated by the program. For this purpose, when using files fu30 or fu40, the program can use only the gradients at the save points or it can use the gradient at up to two additional s values on either side of each Hessian point. However only the latter option is available when using file fu31.

The user must also supply information about the reactants and the products, if it is needed.

9.A. Relation of *Polyrate* input to electronic structure package output

GAUSSIAN94 prints the forces on the standard output file, and it prints the gradients on the formatted check point file; however, on the latter the gradients are incorrectly labeled as forces. We prefer to take the gradients from the formatted checkpoint file because GAUSSIAN's output format on the formatted checkpoint file is more similar to *POLYRATE*'s input format (for fu30, fu31, and fu40), so the user can directly cut and paste.

A different issue arises when one uses the GAMESS electronic structure package. GAMESS writes the gradient in both un-normalized and normalized form. *Polyrate* requires the un-normalized gradient if one uses curvilinear coordinates, so the user should be sure to take the gradient from the correct place in the GAMESS output.

9.B. Orientation of the molecule at generalized transition states

When an electronic structure calculation program is used to determine the optimized structures for the generalized transition states, the electronic structure code sometimes translates and/or rotates the molecule to a “standard”, a “Z-matrix”, or other orientation. However, *POLYRATE* requires precisely the orientation obtained by following the minimum energy path in mass-scaled Cartesians. For example, if the ESD method is used, the points along the path should be related by equations (26) and (27) of the GTST book chapter [Ref. 5 of Section 23]. Using an incorrect orientation can introduce errors in the gradients and curvatures and can result in errors in the tunneling calculations. The user of *POLYRATE* is urged to be very cautious in understanding these issues if he or she uses an electronic structure package that re-orientes the molecule. Special keywords are sometimes required to prevent re-orientation.

For example, for single-point calculations that ask for a gradient or a Hessian or both, when the GAUSSIAN input has the geometry in Cartesian coordinates and the NOSYMM keyword is specified, then GAUSSIAN will not re-orient the molecule, and the gradients and Hessians that it prints out in the formatted checkpoint file (Test.FChk file) are consistent with the orientation of the input geometry. Notice that it is impossible to avoid re-orientation of geometry if the Z-matrix is given in internal coordinates, since GAUSSIAN automatically generates a set of Cartesian “Z-matrix” coordinates in which the first atom is placed at the origin, the second one on the Z axis, and the third one in the X-Z plane. This can be avoided using Cartesian coordinates for defining the input geometry. A second aspect of GAUSSIAN the user should be aware of is that even when using Cartesian coordinates for the input geometry in single-point calculations, if the NOSYMM keyword is not specified, both the gradient and the force constants matrix written to the *formatted checkpoint file* will correspond to a geometry translated and re-oriented according to what GAUSSIAN calls the “standard orientation”. However, notice that the forces and the force constants matrix printed out in the GAUSSIAN *long output file* are still consistent with the initial input geometry orientation; thus they can be used for dynamics without introducing errors in the calculation.

Re-orientation issues are more complicated when one considers reaction paths. It appears to be impossible to generate a reaction path with GAUSSIAN without having the geometries re-oriented if the transition state geometry required to perform a reaction path

calculation is input in internal coordinates. However, if the geometry of the transition state is given in Cartesian coordinates, and if the computed force constants are also read in Cartesian coordinates from a checkpoint file corresponding to a frequency calculation performed including NOSYMM in the keyword list, the reaction path will have the correct orientation to be used in a direct dynamics calculation. Whether the NOSYMM keyword is or is not used in the actual reaction path calculation does not affect the origin or the orientation of the respective points.

An alternative method that we have used in the past to generate the reaction paths is to employ the electronic structure package GAMESS. However, although GAMESS can be used to generate a non-re-oriented reaction path, we could not find a convenient way to prevent GAMESS from re-orienting when it calculates the gradients and Hessians. (We did find a way to avoid re-orientation, but it is inconvenient because it re-orders the atoms.) One solution is to use GAMESS to generate un-re-oriented geometries along the reaction path, and then use GAUSSIAN (with Cartesian coordinates input and NOSYMM) to generate gradients and Hessians in a formatted checkpoint file. Our test of GAMESS were done with the 9-11-96 version.

9.C. Methods for interpolation of electronic structure data with files fu30 and fu40

This section describes the procedures used with file fu30 when LOPT(2) is positive or with file fu40 when MAXLPTS is positive.

For values of s between those on the input grid, the MEP quantities are obtained by n -point Lagrangian interpolation. The maximum number of points used in the interpolation is input to the program; the actual number of interpolation points is the minimum of this input value [LOPT(2) in unit fu30] and the total number of points in the grid. Thus, if only 3 points are used in the grid, only up to second order Lagrangian interpolation can be used.

For unimolecular reactions, association reactions, or reactions with wells, the reactant, product, or well geometry (or both) will occur at finite values of s . In these cases Lagrangian interpolation is also used for values of s approaching the reactant, product, or well s value.

In runs using analytic potential energy functions, there are two grids: (i) the fine grid, also called gradient grid, with step size $\delta s = \text{SSTEP}$ and (ii) the save grid, where frequencies and the Hessian are calculated and stored, with step size $\Delta s = \text{INH} * \text{SSTEP}$. See also sections 7, 10.D, and 12.A.7.

In runs using file fu30 or file fu40, there are three grids which are similar to the two grids described above; however their use is somewhat different; also their step sizes are determined differently. First is the gradient grid. This consists of all the s values at which gradient information is read in records 35, 38, or 41 of unit fu30 or the equivalent records of unit fu40. This grid is not required to be evenly spaced, so δs does not exist, and the SSTEP value read from fu5 is ignored. Second is the basic input grid, which is called the sparse grid in Ref. 14 of Section 23. When using file fu30, this is the same as the gradient grid if $\text{LOPT}(6) \leq 2$; it is a subset of the gradient grid if $\text{LOPT}(6)$ is 3 or ≥ 4 . The basic input grid is the grid of points for which one reads energies in record 31 of unit fu30 and either frequencies (record 46) or Hessians (record 44). Third is the save grid. Input electronic structure information and calculated reaction path curvatures are interpolated from the basic input grid plus saddle point information to the save grid by n -point Lagrangian interpolation. The step size for the save grid is $\Delta s = \text{INH} * \text{SSTEP}$. This interpolation can also be done by the IVTST-M method (For unit fu30, $\text{LOPT}(2) = -1$; for unit fu40, the MAXLPTS keyword in *GEN40 is set to -1) instead of by n -point Lagrangian interpolation. When using the IVTST-M method with files fu30 and fu40, the interpolation can only be calculated with the default IVTST-M options.

A comment on the calculation of $B_{mF}(s)$ may be helpful here. When an analytic potential is used, and the default value (*dgrad*) is accepted for the CURV keyword, $B_{mF}(s)$ is calculated by a quadratic fit of points on the gradient grid; this fit is carried out in subroutine BCALC. In this fit, one uses points with $s \leq 0$ to calculate $B_{mF}(s)$ for $s \leq -\delta s$, and one uses points with $s \geq 0$ to calculate $B_{mF}(s)$ for $s \geq +\delta s$. Then one interpolates between $-\delta s$ and $+\delta s$ to get $B_{mF}(s = 0)$. Notice that none of the quadratic fits involves data with both $s > 0$ and $s < 0$. Thus in eq. (98) [see Section 24.A, item 15], $d\{[\text{sign}(s)]v_{i\gamma}(s)\}/ds$ becomes $[\text{sign}(s)]dv_{i\gamma}(s)/ds$. In contrast, when an electronic structure input file is used, files fu30, fu31, or fu40, one calculates $B_{mF}(s)$ by a quadratic fit to points on the whole gradient grid in subroutine RPHB. Since it is now possible for a given 3-point quadratic fit to involve both negative and positive s , the $\text{sign}(s)$ function must be implemented before the fit is carried out because $[\text{sign}(s)]v_{i\gamma}(s)$ is a continuous function, but $v_{i\gamma}(s)$ is not.

9.D. Methods for interpolation of electronic structure data with file fu31

This section concerns the procedures used with file fu31. The fu31 file is currently incompatible with the IVTST0FREQ option. In version 2010, the ISPE and FREQSCALE options are compatible. The default versions of these procedures may also be used with file fu30 by setting $LOPT(2) = -1$ or with file fu40 by setting $MAXLPTS = -1$.

In runs using file fu31 there are also three grids, similar to those described above for files fu30 and fu40, which store input and interpolated information. The gradient and the basic input grids for file fu31 are the same as described above for files fu30 and fu40. However, when using file fu31, the input grid information is always interpolated onto the save grid using IVTST-M instead of by n -point Lagrangian interpolation. After data is interpolated to the save grid, the rest of the calculation is the same as when using file fu30 or fu40 or when using the hooks.

IVTST-M is described in Sections 6.N.2 and 10.J and in detail in Ref. 48 of Sect. 23.

10. SELECTED DETAILS OF THE CALCULATIONS

Most of the methods used by the program are described in detail, including equations, in the GTST book chapter (Ref. 5 in Sect. 23). All methods not discussed in the GTST book chapter are described in the present section of the manual. See also Refs. 14, 29, 30, and 47 in Sect. 23.

Subsections 10.A-10.L describe in detail various options that may be used with either of the two kinds of calculations—those based on an analytic potential energy surface or those based on an electronic structure input file.

10.A. Extrapolation of the reaction-path data beyond the range of the original grid

In some cases the variational transition state lies in the interval [SLM, SLP], which is the range of the save grid, but the tunneling calculation requires information over a wider range of s to converge.

For reactions with a saddle point this is handled almost automatically by the mapped interpolation (MI) algorithm, which allows one to extend the range of the reaction path used without calculating more energies, gradients, or Hessians. For reactions with a saddle point the MI algorithm is available to interpolate the energy, the determinant of the moment of inertia, the frequencies, and the curvature components between SLM and the reactants or SLP and the products. This may be thought of as an interpolation between SLM and reactants and an interpolation between SLP and products, or it may be thought of as an extrapolation beyond SLM toward reactants and beyond SLP toward products. (That is why we sometimes call it interpolation and sometimes call it extrapolation.) The end result is that all information is interpolated over the entire range from reactants to transition state to products. The keywords EXFIRST and EXSECOND in the PATH section control the options for the interpolation. If the EXFIRST and EXSECOND keywords are not specified then no interpolation is performed. The MI algorithm is explained in Sections 6.N.2, 6.N.3, 10.J, and 12.A.7 and in Ref. 48 of Sect. 23.

For reactions without a saddle point and using RP-VTST *POLYRATE* includes a simple two-point extrapolation to extrapolate the vibrationally adiabatic ground-state potential

and effective reduced mass, μ_{eff} , from the MEP end points to the reactants and/or products. *The asymptotic extrapolation procedures that may be used for reactions without a saddle point are described in this subsection.*

When using units fu30 and fu40 the two-point extrapolation is used by default to extrapolate between the last save grid point and SLM or SLP. However, the IVTST-M algorithm can be specified to interpolate between the last save grid point and SLM or SLP by setting LOPT(2) = -1 for unit fu30 and by setting MAXLPTS = -1 for unit fu40. File fu31 always uses the IVTST-M algorithm to interpolate from the last save grid point to SLM or SLP. See Sections 6.N.2 and 10.J and Ref. 48 of Sect. 23 for more details about the IVTST-M algorithm. In all cases, when using the electronic structure file input options (units fu30, fu31, or fu40), the extrapolation or interpolation is applied using data on the save grid, not the basic input grid (i.e., not the sparse grid). The extrapolation or interpolation is specified by increasing the range of SLM and SLP in the path section.

Precise calculation of the transmission coefficients through the adiabatic barrier for reactions involving two reactant or product species often requires calculation of the ground-state (G) adiabatic energy $V_a^G(s)$ at large absolute values of the reaction coordinate. But calculation of the MEP at large $|s|$ is often impractical, especially in the asymptotic regions near reactants or products, where a very small gradient step size is often required for good accuracy because of the small magnitude of the gradient. To alleviate this problem, the program contains an option to calculate $V_a^G(s)$ and $\mu_{\text{eff}}(s)$ at large $|s|$ by a two-point extrapolation method, controlled by the EXFIRST and EXSECOND keywords in the PATH Section of the fu5 input when the hooks are used and the NOSADDLE keyword is specified in the PATH section. This two-point extrapolation can only be used for reactions without a saddle point or with electronic structure input files fu30 and fu40. See the above discussion for more details. Here we discuss the method in terms of $V_a^G(s)$, but the functional form is the same for $\mu_{\text{eff}}(s)$. [A similar extrapolation is also made for $V_{\text{MEP}}(s)$ but this is for informational purposes only, e.g., it may be useful for plotting.]

The method used for bimolecular reactions when the calculation is based on a potential energy surface involves a simple exponential function which is fit to the values of $V_a^G(s)$ at the positive ($s = \text{SLP}$) or negative ($s = \text{SLM}$) limits of the reaction coordinate for which $V_a^G(s)$ has been accurately determined and to the adiabatic ground-state energies of the products, $V_a^G(s = +\infty)$ or reactants, $V_a^G(s = -\infty)$. Thus

$$V_a^G(s) = V_a^G(s = +\infty) + [V_a^G(\text{SLP}) - V_a^G(s = +\infty)]e^{-\alpha(s-\text{SLP})}$$

for extrapolation towards products ($s > \text{SLP}$), and

$$V_a^G(s) = V_a^G(s = -\infty) + [V_a^G(\text{SLM}) - V_a^G(s = -\infty)]e^{\alpha(s-\text{SLM})}$$

for extrapolation towards reactants ($s < \text{SLM}$). The exponential range parameter α is supplied by the user, or the default (1.0 a_0^{-1}) can be used. Different values of α can be used in the same calculation for extrapolations towards products and reactants.

If the keyword EXFIRST and EXSECOND are not specified, or EXNSTP is zero, no extrapolation is performed. This is equivalent to using $\alpha = \infty$, i.e., this is equivalent to setting $V_a^G(s)$ equal to its value at $s = \text{SLP}$ for all $s > \text{SLP}$ and to its value at $s = \text{SLM}$ for all $s < \text{SLM}$.

As far as inputting a value of α , the user may use any method he or she likes to get α . One suggestion for guesstimating α is a 2-point fit of the exponential on to the end of the Hessian grid data: Let s_x be a point near SLP (s_x could be the second-to-last point, or slightly farther back if precision of $V_a^G(s)$ info is lacking). Then require exponential to fit $V_a^G(s = s_x)$ and $V_a^G(s = \text{SLP})$, which gives

$$\alpha = \frac{\left\{ \ln[V_a^G(s = s_x) - V_a^G(s = +\infty)] / [V_a^G(s = \text{SLP}) - V_a^G(s = +\infty)] \right\}}{(s_x - \text{SLP})}$$

However, if the value chosen for α affects the results, then this represents the basic uncertainty in the values; a wider range for $V_a^G(s)$ will be needed to obtain a better estimation of α .

10.B. Options for anharmonicity

In addition to the standard harmonic option, several anharmonic options are available using the keywords SSTOR, TOR, MORSE, MORSEQQ, MORMODEL, QQWKB, QQSEMI, and WKB. Anharmonicity may be included in reactants, products, or generalized transition states, but only in conjunction with the independent-normal-mode (INM) approximation

in rectilinear coordinates except the SSTOR option. Two of these options, Morse I and Morse I quadratic-quartic (MORSE and MORSEQQ respectively with the MORMODEL variable keyword set to *morse*i**), are described in the GTST book chapter (see pp. 87-92 of Ref. 5 in Sect. 23) and were first used in the calculations reported in Ref. 1 of Sect. 23. The options defined by the QQSEMI and QQWKB keywords are two WKB options that will be discussed in Subsection 10.B.1. The VARIABLE and VRANGE keyword options allow a different anharmonicity method to be used for each mode in each region of the MEP. When any of the above anharmonicity options are used, anharmonicity is computed for all vibrational levels of a given mode by the same method. An additional anharmonic option, specified by the WKB keyword, allows one to use a presumably more accurate, but more time consuming method (the WKB method based on the true potential along each generalized normal mode) for the ground-state energy level of each mode for modes treated thermally, or for the specified energy level for state-selected modes. When this option is chosen, the methods used for higher-energy states of thermal modes are still determined by an additional anharmonicity keyword. If another anharmonicity keyword is not included the higher-energy states will be determined using the harmonic approximation. Details of the implementation of the WKB keyword option will be discussed in Subsection 10.B.2. All WKB anharmonicity options are available only for calculations based on the hooks. Whenever the WKB approximation is used, it is employed in rectilinear coordinates, even if internal coordinates are used for the frequencies and for non-WKB energy levels in the partition functions.

10.B.1. WKB anharmonicity based on a quadratic-quartic fit

When either QQWKB or QQSEMI keywords is specified, the program uses the primitive or uniform WKB method, respectively, to compute the energy levels of a given generalized normal mode vibration from a quadratic-quartic potential fit to the true potential. The quadratic-quartic potential is determined by fitting to the potential computed at two distances, specified by the user in the mass-scaled coordinate system, along the generalized normal mode. The energy levels of this potential can be computed by either a primitive WKB method [Ref. 21, Sect. 22] (QQWKB) or, for double well potentials, by a semiclassical method using parabolic connection formulas [Ref. 22, Sect 20] (QQSEMI). In these methods the phase integrals are calculated analytically. The calculations are described in Ref. 17 of Sect. 23.

10.B.2. WKB anharmonicity based on the true potential

When the WKB keyword is specified, the program uses the WKB method to compute the anharmonic ground- or first-excited-state energy levels of all generalized normal mode vibrations from the true potential along the modes. The method is described in more detail in Ref. 13 of Sect. 23. Note that WKB energy levels are not computed for reactant and product diatomic species, but instead either harmonic or anharmonic energy levels (as specified by the other anharmonicity options) are used.

Because the numerical solution of WKB eigenvalues requires repeated calculation of the potential and may therefore be quite time consuming, this approach is used only for the zero point ($n = 0$) eigenvalues for thermal modes or the specified energy level for state-selected modes. Higher energy vibrational levels, used in the calculation of the partition functions, are obtained by harmonic or anharmonic approximations discussed previously. In particular, when the WKB-with-the-true-potential option is applied to evaluate the ground-state energy levels, the vibrational partition function of a bound motion is calculated as follows:

$$Q_{\text{vib},m} = e^{-\varepsilon_{0,m}^{\text{WKB}}/kT} + \sum_{n=1}^{\infty} e^{-(n+1/2)\hbar\omega_m/kT}$$

$$Q_{\text{vib},m} = Q_{\text{vib},m}^{\text{HO}} - e^{-\hbar\omega_m/2kT} + e^{-\varepsilon_{0,m}^{\text{WKB}}/kT}$$

where $\varepsilon_{0,m}^{\text{WKB}}$ is the WKB ground-state energy of mode m , and $Q_{\text{vib},m}^{\text{HO}}$ is the vibrational partition function in the harmonic oscillator approximation. If the harmonic frequency is imaginary, the partition function is set to 10^{10} . Note that in the WKB method the turning points are determined during the calculation of the vibrational energies, and they can be used directly in the calculation of the effective reduced mass as a function of s .

This method is used in the test runs oh3tr3, ohcltr2, ohcltr3, and ohcltr4.

10.B.3. Hindered rotor treatments

All of the statistical thermochemical methods in which torsions are treated as hindered internal rotations are explained in Refs. 52, 62, 87, and 100. These references explain the methods in detail, and they also give original references.

Beginning with version 8.2 of *POLYRATE*, a torsional mode can be treated in the full generality of Ref. 52. Please note the errata that have been added to Ref. 52. We label the method developed in Ref. 52 as the Chuang-Truhlar (CT) method.

Beginning with version 2010 of *POLYRATE*, several methods in Ref. 62 are also implemented. These additional methods are the reference Pitzer-Gwinn (RPG) method, the segmented- reference Pitzer-Gwinn (SRPG) method, the Ayala-Schlegel (AS) method, and the segmented Ayala-Schlegel (SAS) method.

Beginning with version 2016, *POLYRATE*, can treat the torsional anharmonicity along reaction path (saddle point and nonstationary points) by using coupled torsional potentials in the multistructural method including torsional anharmonicity (MS-T(C)). Note that only a single structure (SS) version is taken into account since only one saddle point is given in a single *POLYRATE* input file. Therefore we call this the SS-T method. Multistructural (MS) effect should be taken into account separately by using the MSTOR program.

There are two kinds of hindered rotor treatments: multiconformer (MC) methods and multistructure (MS) methods. The MC methods involve identifying one or more of the normal modes (or generalized normal modes) as a torsion about of specified bond. When frequencies are required in MC methods they are normal-mode frequencies. The MS methods do not require associating specific torsion with specific normal modes. When torsional frequencies are required in MS methods, they are calculated in internal coordinates based on a coupled model potential. In *POLYRATE* calculations, only one structure of the transition state or one reaction path is taken into account; therefore only single structure torsional potential anharmonicity (SS-T) along the reaction path is taken into account.

10.B.3.a. Multiconformer methods

10.B.3.a.i. CT method

If mode m is treated as a hindered rotor (HR) by the CT method, its partition function tends to the limit of a harmonic oscillator (Q_m^{HO}) at low temperature and goes to the limit of a free rotor (Q_m^{FR}) at high temperature, and the interpolation is designed so that it remains reasonably accurate for the hindered-rotor regime between these limiting cases. The CT method is:

$$Q_m^{\text{HR}} = Q_m^{\text{MC-HO}} \tanh\left(\frac{Q_m^{\text{FR}}}{Q_m^{\text{I}}}\right),$$

where the multiconformer harmonic oscillator (MC-HO) approximation for mode m is $Q_m^{\text{MC-HO}}$, the free rotor approximation for mode m is Q_m^{FR} , and Q_m^{I} is an intermediate result. They are given by

$$Q_{\text{tor}}^{\text{MC-HO}} = \sum_{j=1}^P \frac{e^{-\beta(U_j + \hbar\omega_{\text{tor},j}/2)}}{1 - e^{-\beta\hbar\omega_{\text{tor},j}}}$$

$$Q^{\text{FR}} = \frac{(2\pi I k T)^{1/2}}{\hbar\sigma}$$

$$Q^{\text{I}} = \sum_{j=1}^P \frac{e^{-\beta U_j}}{\hbar\beta\omega_{\text{tor},j}}$$

where P is number of distinguishable minima along the torsion coordinate, β is $(k_{\text{B}}T)^{-1}$, k_{B} is Boltzmann constant, U_j is the energy of the j th well relative to the global minimum, $\omega_{\text{tor},j}$ is the torsional frequency of the j th well along the normal-mode or generalized-normal-mode coordinate, I is the averaged torsional moment of inertia of the P minima, σ is the total symmetry number ($\sigma = M/P$, M is the total number of minima along the torsion coordinate in the range $[0, 2\pi)$). Note that usually $P = 1$ or $P = M$. Therefore, if M/P is not an integer, the program prints an error message and stops. The CT method approximates the one-dimensional hindered-rotator problem with a simple model potential of the form

$$V = \frac{W}{2}(1 - \cos M\phi)$$

where W is the height of the potential barrier, and ϕ is a coordinate that parameterizes the internal rotational motion. For a potential of the above form with a geometry-independent moment of inertia, the barrier height W_j , moment of inertia I_j , and one-dimensional torsional frequency $\omega_{\text{tor},j}$ are interrelated by

$$\omega_{\text{tor},j} = M \left(\frac{W_j}{2I_j} \right)^{1/2}$$

If the frequencies $\omega_{\text{tor},j}$ are approximated as the same for all of the P minima and the energies U_j are approximated as 0 (same as the global minimum energy) for all of the P minima, we label this as single-frequency-and-energy (SFE) approximation. If only the frequency is approximated as the same for all the distinguishable minima, we label it as the single-frequency (SF) approximation. Otherwise, the treatment is labeled FULL.

Note that the single-conformer (SC) approximation in Ref. 52 is another name for the special case of $P=1$. Therefore, starting from version 2010 of *POLYRATE* program, this SC approximation is not used as a stand-alone method.

10.B.3.a.ii RPG method

The torsional Pitzer–Gwinn method based on a reference potential gives the torsional partition function as

$$Q_{\text{tor}}^{\text{RPG}} = \frac{1}{P} \sum_{j=1}^P \frac{Q_{\text{tor}}^{\text{HO}}(\omega_{\text{tor},j})}{Q_{\text{tor}}^{\text{CHO}}(\omega_{\text{tor},j})} Q^{\text{FR}} \exp(-\beta W / 2) I_0(\beta W / 2)$$

where $\omega_{\text{tor},j}$ is the frequency at the well j , W is the averaged barrier height of all the wells to their left and right maxima, and I_0 is a modified Bessel function. If one uses the frequency associated with the minimum of the 1D torsional reference potential to approximate all the $\omega_{\text{tor},j}$, this equation reduces to the Eq. 24 in Ref. 62. The reader is referred to Ref. 62 for all further details.

10.B.3.a.iii. AS method

AS method uses a correction factor to adjust the RPG method

$$Q_{\text{tor}}^{\text{AS}} = \frac{1 + P_2 \exp[-\beta W / 2]}{1 + P_1 \exp[-\beta W / 2]} Q_{\text{tor}}^{\text{RPG}}$$

where P_1 and P_2 are fifth order polynomial functions of $x \equiv 1/Q^{\text{FR}}$ and $y \equiv \beta W / 2$ which is given by Eq. 31 and 32 in Ref. 62.

10.B.3.a.iv. SRPG method

If the hindered-rotator potential does not closely resemble the form of

$$V = \frac{W}{2}(1 - \cos M\phi),$$

it may be convenient to decompose the partition function into a sum of contributions from particular wells and to fit the potential between each peak and minimum using a simple cosine line shape. For each well, the contribution may be divided into the part to the left of the minimum and the part to the right of the minimum, and each part of the classical partition function may be evaluated analytically. This is referred as the segmented-reference classical (SRC) approximation, and it yields

$$Q^{\text{SRC}} = \sum_{j=1}^P Q_j^{\text{SRC}}$$

where

$$Q_j^{\text{SRC}} = \exp(-\beta U_j) Q_j^{\text{FR}} \times \left[\frac{(\phi_j - \phi_j^L)}{2\pi} \exp(-\beta W_j^L / 2) I_0(\beta W_j^L / 2) + \frac{(\phi_j^R - \phi_j)}{2\pi} \exp(-\beta W_j^R / 2) I_0(\beta W_j^R / 2) \right]$$

where U_j is the energy of the local minimum centered at ϕ_j relative to the energy at the global minimum, W_j^L and W_j^R are the left and right barrier heights located at ϕ_j^L and ϕ_j^R , respectively, and

$$Q_j^{\text{FR}} = \frac{\sqrt{2\pi I_j kT}}{\hbar \sigma}$$

Quantum effects may be approximated by the Pitzer-Gwinn approximation to yield the segmented-reference Pitzer-Gwinn method

$$Q_{\text{tor}}^{\text{SRPG}} = \sum_{j=1}^P \frac{Q_j^{\text{HO}}(\omega_{\text{tor},j})}{Q_j^{\text{CHO}}(\omega_{\text{tor},j})} Q_j^{\text{SRC}}$$

10.B.3.a.v. SAS method

The AS method has a segmented approach that is analogous to the SRPG method, and we refer to this analog to the segmented Ayala-Schlegel (SAS) method, in which each occurrence of $\exp(-\beta W_j / 2)$ in the SRPG method is replaced by a term like

$$\exp(-\beta W_j / 2) \frac{1 + P_2(I_j, W_j) \exp(-\beta W_j / 2)}{1 + P_1(I_j, W_j) \exp(-\beta W_j / 2)}$$

where P_1 and P_2 are given by Eq. 31 and 32 in Ref. 62.

The test runs ch4ohtr1, ch4ohtr3, ch4ohtr4, cwmctr2, cwmctr3, and h3o2fu31tr1 are examples of using the hindered rotor options.

10.B.3.b. MS-T method

Starting from version 2016 of *POLYRATE*, the torsional-potential anharmonicity along the MEP can be calculated using the MS-T method with a coupled torsion potential (i.e., MS-T(C)), but only the input single transition structure is considered as generalized transition states in the rate constant calculation; this is the SS-T treatment, i.e. *POLYRATE* used by itself does not compute the Multipath VTST (MP-VTST) rate constant, which requires multiple transition structures in order to compute the variational and transmission coefficients that correspond to their MEPs. Note that reactants and products are still treated by harmonic oscillator approximation. To obtain rate constants calculated by the full MS-T partition functions for reactants, products, and GTS, as well as the MP-VTST rate constant, the user has to use MSTOR program (which is a separate program) to calculate the rate constants at the conventional transition state and then use the variational effects (ratio of CVT rate to TST rate) and transmission coefficients calculated by *POLYRATE* program (see the MSTOR manual for more details).

The conformational-rovibrational partition function using the MS-T method for a species with J conformational structures is

$$Q_{\text{con-rovib}}^{\text{MS-T}} = \sum_{j=1}^J Q_j^{\text{T}}$$

$$Q_j^{\text{T}} = Q_{\text{rot},j} \exp(-\beta U_j) Q_j^{\text{QH}} T_j$$

where Q_j^{T} is the contribution of structure j to the rovibrational partition function with torsional anharmonicity, $Q_{\text{rot},j}$ is the classical rotational partition function for structure j , U_j is its relative potential energy, Q_j^{QH} is its QH vibrational partition function, and T_j is a factor to account torsional-potential anharmonicity for all torsional modes in the structure j . In particular, T_j is

$$T_j = (2\pi\hbar\beta)^{t/2} \frac{\prod_{m=1}^F \omega_{j,m}}{F-t} \frac{\sqrt{\det \mathbf{D}_j}}{t} \prod_{\eta=1}^t \exp(-\beta W_{j,\eta} / 2) I_0(\beta W_{j,\eta} / 2) \prod_{\bar{m}=1}^t \bar{\omega}_{j,\bar{m}} \prod_{\tau=1}^t M_{j,\tau}$$

where t is number of torsional modes, F is number of bound vibrational modes (stretches, bends, and torsions), $\omega_{j,m}$ and $\bar{\omega}_{j,\bar{m}}$ are respectively the normal mode harmonic frequencies and the torsion-projected normal mode frequencies (both sets scaled), \mathbf{D}_j is the moment of inertia matrix of Kilpatrick and Pitzer, $M_{j,\tau}$ is the periodicity of torsion τ in structure j , $W_{j,\eta}$ is the torsional barrier height based on a coupled torsional potential, and I_0 is a modified Bessel function. Note that for nonstationary points along a reaction path j (starting from the saddle point j), the vibrational mode along reaction coordinate is projected out and the torsional-potential anharmonicity T_j is calculated for each nonstationary points where Hessian is calculated or available. Therefore, vibrational partition functions of points along reaction path includes the torsional-potential anharmonicity and then torsional-potential anharmonicity is taken into account for searching variational transition state.

To calculate T_j , one only needs to input the set of $M_{j,\tau}$ values for torsions and projected frequencies have to be calculated using non-redundant internal coordinates.

10.C. Alternate method for the calculation of numerical second derivatives

In order to calculate the Hessian matrix, the second derivatives of the potential with respect to the Cartesian coordinates are computed numerically, using the analytic first derivatives supplied by the analytic potential energy surface subprogram. By default, the method of central differences is used to compute these second derivatives. The formula is given by equation (16) in the GTST book chapter (Ref. 5 of Sect. 23).

Two alternate options, selected by setting the NUMTYP variable keyword to *quadratic* or *fourth*, were added for use in cases where the default method may lead to numerical instability. In the most accurate option, NUMTYP = *fourth*, the second derivatives are computed by fitting the analytic first derivatives to a fourth order polynomial. For NUMTYP = *quadratic*, the second derivatives are computed from a quadratic fit of the potential.

The parameters NUMSTEP and DLX2, which are specified by the variable keywords NUMSTEP and DLX2 respectively, in the input file which is linked to the FORTRAN unit fu5, determine the step size for calculating second derivatives of the potential, and, if needed, for cubic and quartic force constants as well. The second derivatives are needed for two purposes: locating stationary points and computing vibrational potentials.

The DLX2 parameter is used for locating the stationary point, and the NUMSTEP parameter is used for calculating the force constants. When anharmonicity options are chosen that require cubic and quartic force constants, which are third and fourth derivatives of the potential, these are evaluated as numerical second and third derivatives, respectively, of the analytic gradient. The step size used for these numerical derivatives is always NUMSTEP.

10.D. Reaction-path following methods

Our recommendations, as described in Ref. 27 of Sect. 23, for the most appropriate methods for calculating the MEP and the reaction-path information required for variational transition state theory calculations with semiclassical transmission coefficients are as follows: (i) the Page-McIver algorithm, with the Hessian recalculated at every step, and with a cubic starting step, (ii) the ES1* algorithm (the ES1 method with DELTA2 = SSTEP and DIFFD = 0.01) with a Hessian step size three times larger than the gradient step size, and with a quadratic starting step, and (iii) the Euler steepest-descents (ESD) algorithm with a Hessian step size nine times larger than the gradient step size, and with a quadratic starting step. The RPM keyword in the PATH section allows the user to choose between the above possibilities or to choose other methods for computing the MEP. A list of all methods available in this version is as follows:

The RPM keyword is set to *esd* for the ESD method. This method is discussed further in Ref. 18 of Sect. 23.

The RPM keyword is set *es1* for the ES1 method of Ref. 18 of Sect. 23. This is an improved version of the methods proposed by Ishida et al. [Ref. 19, Sect. 22] and by Schmidt, Gordon, and Dupuis [Ref. 20, Sect. 22]. The ES1 method allows for a user-selected criterion for whether or not a stabilization is carried out.

The RPM keyword is set to *pagem* for the Page-McIver method, which is called the local quadratic approximation (LQA) in the original reference [Ref. 17, Sect. 22], for calculating the MEP.

The RPM keyword is set to *vrpe* for the VRP method based on the Euler steepest-descents algorithm for the zero-order step, which is called VRP(ESD). This method is discussed further in Ref 49 of Sect. 23.

The ESD, ES1, and Page-McIver methods can be used with the RODS algorithm [Ref. 47, Sect. 23], to reorient the generalized-transition-state-theory dividing surface. This is done by adding the RODS keyword in the PATH section of file fu5. The RODS algorithm is not recommended for use with the VRP(ESD) algorithm since the generalized-transition-state-theory dividing surface has already been optimized by the VRP(ESD) algorithm.

Each of the above methods can be used with either a quadratic first step (i.e., a first step along the imaginary-frequency normal mode) or a cubic starting algorithm in which the first step off the saddle point is based on a cubic expansion of the potential energy surface using the imaginary-frequency normal mode eigenvector and the curvature vector. This is specified by the FIRSTSTEP variable keyword also in the PATH section. For a quadratic first step set FIRSTSTEP = *nmode* and for a cubic first step set the variable to *cubic*.

A discussion of the above methods and useful hints for choosing the most appropriate one, especially for direct dynamics calculations, is given in Ref. 27 of Sect. 23.

We use the following conventions for the step size of the save grid. The save grid is defined as the set of points where the frequencies are stored for straight VTST runs, these are also the points where the Hessian matrix is calculated and where generalized transition-state normal mode analyses are performed and saved. The INH keyword determines how many gradient steps are taken before a Hessian is calculated or read in. In other words, the step size of the save grid is required to be an integer multiple, INH, of the gradient grid step size. Hence, the save grid step size = INH × SSTEP.

There is another option, which is specified by the SFIRST keyword, in which the first NFSTEP steps are taken with a different gradient step size FSIZE. Then the gradient step size, SSTEP, is used for the remaining steps.

10.E. Interpolation of small-curvature effective reduced mass

For the small-curvature tunneling calculations (SCT), the effective reduced mass is needed. The default method for computing the effective reduced mass from the values stored on the Hessian grid, which includes $s = 0$, is spline interpolation. An alternate method, selected using the LAGRANGE keyword within the SCTOPT list of the TUNNEL section of the input is Lagrangian interpolation of order n . This new method is recommended when using medium-to-large Hessian grid step sizes for which spline fits tend to oscillate. The step size used for either of the interpolation methods is $\text{INH} \times \text{SSTEP}$ specified by the user with the SSTEP and INH keyword in the input data file. The alternate method is used in the sample runs described in Subsections 16.C.3, 16.C.9, and 16.C.29.

10.F. Reaction-path curvature calculation

See Section 6.H for a brief introduction to reaction-path curvature.

When the CURV variable keyword in the PATH section of the input is set to *oneside* or *dgrad*, the calculation of the component $B_{mF}(s)$ of the reaction-path curvature along the direction of mode m at a point s on the reaction path Hessian grid, is carried out by eq. (98) of the GTST book chapter. (This is equivalent to the second form of eq. (47) of [Ref. 17, Sect. 22].) This method requires the computation of the derivative of the gradient of the potential with respect to the reaction coordinate s (see pages 96-98 in the GTST book chapter). In early versions of the polyatomic VTST code this derivative was computed using one-sided forward differences; see eq. (105) of the GTST book chapter. An option, CURV = *dgrad* has been added to use a more accurate method, namely, central differences (which is equivalent to a 3-point quadratic fit). When this new option was added the one-sided difference method was changed to one-sided backward differences to allow for a more consistent set of algorithms. It is recommended, however, that the method of backward differences, CURV = *oneside*, never be used. The step size used for these numerical derivatives is that of the gradient grid, SSTEP.

Another option, CURV = *dhess*, calculates the reaction-path curvature components using eq. (48) of Page and McIver [Ref. 17, Sect. 22]. This method does not require the

derivative of the gradient. A sample run with this option is discussed in Subsection 16.C.4. Preliminary tests indicated somewhat higher accuracy with *dgrad*.

For any of these choices of *CURV*, the code does not compute the curvature at $s = 0$ directly. Instead, the curvature at $s = 0$ is computed linear interpolation as described by the *INTMU* keyword in the *PATH* section.

10.G. Solid-state and gas-solid interface reactions

A solid-state reaction, which includes gas-solid interface reactions as a special case, is a reaction in which *N* primary atoms are considered to move in the presence of a fixed solid. This is called the embedded cluster model; see Ref. 9 of Sect. 23. For solid-state reactions, the *SPECIES* should be *ssrp* or *ssts* as explained in section 6.B.

10.H. VTST-IOC calculation, including ICR, ICL, and ICA schemes

The original VTST-IOC method is the VTST-IOC method that is described in Ref. 36 of Sect. 23, and the modified version that was the default in *POLYRATE* from version 7.2 – 7.9.2 is described in Ref. 44 of Sect. 23. In this manual we use the phrase VTST-IC to refer to any scheme involving interpolated corrections from a higher level and VTST-IOC to refer to any VTST-IC scheme in which the data for corrections involves a higher-level optimization. Thus VTST-IC and VTST-IOC can involve not only the original ICA scheme of Ref. 36 of Sect. 23 for interpolating corrections to frequencies but also to the ICR and ICL options described in Ref. 44 of Sect. 23 and also described toward the end of this section. To use any IOC method, the user has to prepare an input file (unit fu50; see Subsection 12.F) which contains the necessary higher-level information. The *DL* keyword must be set equal to *ioc* and/or the *ICOPT* keyword must also be given in the *GENERAL* section of the unit fu5 input file.

If the MEP is finite (unimolecular reaction with a single product) or semi-infinite (unimolecular reaction with two products or bimolecular reaction with a single product) or has wells (i.e., precursor and/or successor complexes) that the user wishes to use in the interpolation (i.e., if *MEPTYPE* or *MEPTYPEP* is (or both are set to *well*), the user has to

tell the program where the MEP ends and/or some properties of the complexes corresponding to the wells (see Subsection 12.F).

The VTST-IOC method demands an explicit matching of the frequencies, i.e., the user must indicate which mode at the saddle point corresponds to each mode at the other two stationary points involved in the interpolation (or else accept the default as explained later in this paragraph). For a bimolecular reaction with two products (MEPTYPEP and MEPTYPEP are both set to *two*), the recommended method is to let the program match the frequencies in descending order (which is the default in the current version), i.e., the highest frequency at the saddle point is matched to the highest frequency of either of the two asymptotic species, and so on. If the user knows (by symmetry and/or by inspecting the uncorrected MEP output) that the matching should not be done in descending order, then the user can use the FREQMAT keyword to input the matching order manually in unit fu50. When MEPTYPEP and MEPTYPEP are not both set to *two*, one of the normal modes at the reactants and/or products turns into the reaction coordinate motion, and the user has to specify the frequency matching explicitly, so the FREQMAT keyword must be used.

If a well is used for interpolation on the product side of the MEP, and the user is not interested in the reverse rate constant, then the accurate product information is not required. The user can use the NOPRODFREQ keyword to tell the program that no correction for the product frequencies is needed. Otherwise the user has to use the PRODFREQ keyword to input the corrected frequencies for the product in unit fu50.

The original IOC method involved the ICA option, which corrects all the frequencies, and sometimes it will generate a negative correction that changes a small real frequency into a negative frequency (not an imaginary frequency). In order to prevent this unphysical result, one may use IVTST-0 for one or more low frequencies, or, preferably, one may use one of two new interpolated correction options, namely the ICL (Interpolated Correction based on Logarithm of ratios) and the ICR (Interpolated Correction based on Ratios) options.

We note that all three IOC interpolation options for frequencies correct the lower-level frequencies only when they are real. If the lower-level frequencies are imaginary, the program does nothing, so the frequencies remain imaginary. (The recommended way to handle this situation is to use the LOWFREQ option in the fu50 file. This option treats

selected low-frequency vibrations by IVTST-0, while carrying out the rest of the calculation by the VTST-IOC method.)

The differences between the original ICA option and the ICL and ICR options are best illustrated by equations. Let $\omega_m^{\text{LL}}(s)$, $m = 1, 2, \dots, F-1$ (where $F = 3N - 6$ for an N -atom nonlinear generalized transition state) denote the lower-level frequencies of the generalized transition state at s . In the original IC method, which is called “interpolated corrections–additive” (ICA), the dual-level (DL) frequencies are given by

$$\omega_m^{\text{DL}}(s) = \omega_m^{\text{LL}}(s) + f^{\text{ICA}}(s)$$

where

$$f^{\text{ICA}}(s) = \text{interpolant}\left\{\omega_m^{\text{HL}}(s) - \omega_m^{\text{LL}}(s)\right\}_{s=-\infty,0,\infty}.$$

The “interpolated corrections based on the ratio” (ICR) and “interpolated corrections based on the logarithm” (ICL) methods are defined by

$$\omega_m^{\text{DL}}(s) = \omega_m^{\text{LL}}(s) f^{\text{ICR}}(s)$$

where

$$f^{\text{ICR}}(s) = \text{interpolant}\left\{\omega_m^{\text{HL}}(s)/\omega_m^{\text{LL}}(s)\right\}_{s=-\infty,0,\infty}$$

and

$$\omega_m^{\text{DL}}(s) = \omega_m^{\text{LL}}(s) \exp\left[f^{\text{ICL}}(s)\right]$$

where

$$f^{\text{ICL}}(s) = \text{interpolant}\left\{\ln\left[\omega_m^{\text{HL}}(s)/\omega_m^{\text{LL}}(s)\right]\right\}_{s=-\infty,0,\infty}.$$

There are also options for interpolating $V_{\text{MEP}}(s)$. The original option, called SECKART, is described in Ref. 36 of Sect. 23. In the DECKART option, the dual-level $V_{\text{MEP}}(s)$ is defined as:

$$V_{\text{MEP}}^{\text{DL}}(s) = V_{\text{MEP}}^{\text{LL}}(s) + \left[V_{\text{Eck}}^{\text{HL}}(s) - V_{\text{Eck}}^{\text{LL}}(s)\right]$$

where

$$V_{\text{Eck}}(s) = \frac{AY}{1+Y} + \frac{BY}{(1+Y)^2} + C$$

$$Y = \exp\left(\frac{s - S_0}{L}\right)$$

$$A = V_{\text{MEP}}(s = +\infty) - V_{\text{MEP}}(s = -\infty)$$

$$C = V_{\text{MEP}}(s = -\infty)$$

$$B = (2V^\ddagger - A - 2C) \pm \sqrt{(V^\ddagger - C)(V^\ddagger - A - C)};$$

we choose

$$S_0 = -L \ln \left(\frac{A+B}{B-A} \right)$$

so that the maximum of $V_{\text{Eck}}(s)$ occurs at $s = 0$. V^\ddagger is the classical barrier height, and L is the range parameter. In $V_{\text{Eck}}^{\text{HL}}(s)$, L is determined from the imaginary frequency of the higher-level calculation by the equation

$$L^{\text{HL}} = \sqrt{\frac{2V^\ddagger(V^\ddagger - A)}{\mu(\omega^\ddagger)^2 B}}$$

where μ is the scaling mass (normally set to 1.0 amu), and ω^\ddagger is the imaginary frequency at the higher level. In $V_{\text{Eck}}^{\text{IL}}(s)$, L is determined from a four-point fit to $V_{\text{MEP}}(s)$ at reactants, products, saddle point, and one extra point as discussed in Refs. 36 and 44 in Sect. 23.

Versions 7.2 – 7.9.2 used the so-called improved scheme (i.e. *icl* for vibrations and *deckart* for $V_{\text{MEP}}(s)$) as the default VTST-IOC method. Beginning with version 8.0 the defaults are *icl* and *seckart*. To perform the calculation by the original IOC method, a combination of *ica* and *seckart* was required. The Eckart function fit to 3 stationary points and ω^\ddagger at the higher level can only be used for bimolecular reactions.

The following table summarizes the IOC options for interpolating corrections to vibrations and to the V_{MEP} curve:

Option no.	Keywords
1	<i>ica, seckart</i>
2	<i>icl, deckart</i>
3	<i>icl, seckart</i>
4	<i>icr, seckart</i>
5	<i>ica, deckart</i>
6	<i>icr, deckart</i>

Option 1 was the original default, option 2 was the recommended choice and default in versions 7.3 – 7.9.2, and option 3 is the current recommended choice and default.

In order to perform VTST-IOC calculations using any of the *ica*, *icl*, or *icr* choices for interpolating vibrational information, we suggest the following procedure for a bimolecular reaction:

- Step 1. Calculate the rate constants at the lower level. At the end of the reaction path section in the long output file (i.e. before the reaction rate calculation), the range parameter L is calculated automatically from a four-point fit. In the same section, an L value obtained from the imaginary frequency at the lower level is also calculated, but this is only for comparison.
- Step 2. Input the range parameter L calculated from the four-point fit as the value for the RPL keyword in the fu50 input file. Or calculate the range parameter L for the RPL keyword in the fu50 input file with the findl.f program in /util. (See Sect. 15.D.) This utility program asks the user to input the V_{MEP} values at reactants, saddle point, and products (0 , barrier height, endoergicity) and the (s, V_{MEP}) values of an extra point which lies somewhere between $s = -\infty$ and $s = 0$ (between the reactants and saddle point).
- Step 3. Use the RESTART keyword to calculate the rate constants from the interpolated corrections with the higher-level information given in fu50. (Note: The RESTART option is not working properly in this version, you must do the lower-level calculations again if you are dealing with an analytic surface.)

For a unimolecular reaction, the B parameter can be calculated from the findb.f program in /util.

10.I. VTST-ISPE calculation

The VTST-ISPE method for calculating reaction rates is a dual-level approach. ISPE denotes that the higher-level information consists of interpolated single-point energies. The fully converged reaction path is calculated at a lower level of theory Y, e.g. MP2/6-31G, and single-point energies are calculated at a higher level of theory X, e.g. CCSD(T)/cc-pVDZ, where the geometry for the single-point calculation is taken from level Y, denoted X//Y. In order to circumvent the limitation of the VTST-IOC formulas that are designed for calculations in which only corrections at stationary points are used, the VTST-ISPE calculations use the mapped interpolation formulas rather than the VTST-IOC formulas. However, whereas the IVTST-M method requires at least one nonstationary point on both sides of the saddle point, VTST-ISPE can handle the case where corrected energies are provided only at one nonstationary point on either side of the saddle point.

To calculate the reaction rates with the VTST-ISPE algorithm, one should calculate a converged reaction path at the lower level Y. Then a spline under tension is used to interpolate the energy difference of a few single-point level-X energies along the level-Y MEP and the energy from the lower level Y (*i.e.*, $V^{X/Y} - V^Y$). As in our previous interpolated variational transition state theory with mapping (IVTST-M) method, a mapping function is used to provide a systematic interpolating procedure for both bimolecular and the unimolecular reaction. Like IVTST-M, which does not support LCT calculations at this time, VTST-ISPE supports only ZCT and SCT tunneling options at this time.

To map the reaction coordinate s into the z space, the following expression is used:

$$z = \frac{2}{\pi} \arctan\left(\frac{s - s_0}{L}\right) \quad (1)$$

Information from the lower level is used to obtain the parameters s_0 and L for the mapping. First we calculate

$$s_A^0 = -\sqrt{\frac{V_{\text{MEP}}(s=0) - V_{\text{MEP}}(s_X)}{|\omega^\ddagger|^2 \mu}} \quad ; X = \text{R, RW} \quad (2)$$

and

$$s_B^0 = \sqrt{\frac{V_{\text{MEP}}(s=0) - V_{\text{MEP}}(s_X)}{|\omega^\ddagger|^2 \mu}} \quad ; X = \text{P, PW} \quad (3)$$

where μ is the scaling mass, ω^\ddagger is the imaginary frequency of the saddle point at the lower level, and R, RW, P, and PW denote the reactants, the reactant well, the products, and the product well at the lower level. Then we calculate

$$s_A = -\min(|s_A^0|, 2s_B^0) \quad (4)$$

and

$$s_B = -\min(2|s_A^0|, s_B^0) \quad (5)$$

Finally, the parameters s_0 and L are obtained as:

$$s_0 = \frac{s_A + s_B}{2} \quad (6)$$

and

$$L = \frac{-|s_A| + s_B}{2} \quad (7)$$

The parameters s_0 and L obtained from equations (5) and (6) are used in equation (1) to map the reaction coordinate s into z . For a unimolecular reaction with a unimolecular

product ($A \rightarrow B$), where the reaction coordinate s occupies a finite interval, this maps a finite interval in s into a finite interval in z . For a bimolecular reaction with bimolecular products ($A + B \rightarrow C + D$), where the s variable occupies interval $[-\infty, \infty]$, the mapped z values are located in the finite interval $[-1, 1]$. For $A \rightarrow B + C$ or $A + B \rightarrow C$, equation (1) maps a semi-infinite interval of s to a finite interval of z . In all cases, a spline fit to the differences of the classical energies is applied with this finite range of z . Therefore, we can write the interpolated energy along the reaction path at the dual level as:

$$V_{\text{DL}} = V_{\text{LL}} + \text{spline}(\Delta V, z) \quad (8)$$

where DL means dual level (VTST-ISPE), and LL means the lower level. ΔV is the difference of the classical energies between higher-level and lower-level at a particular s value.

In the current version of *POLYRATE*, ISPE calculations may be used with the ZCT and SCT tunneling methods but not with LCT or μ OMT.

To carry out a VTST-ISPE calculation, an external file fu51 is used to input the values of the higher-level energies. Please refer to Section 12.G for a description of fu51. The VTST-ISPE method can be applied with higher-level calculations at only the reactants, products, and saddle point, or it may be applied with more points. In the latter case, extra points are used to correct the energies along the MEP, the POINT sections are required for the extra points.

The interpolation can be carried with or without the wells along the MEP. It is recommended to use the well option if there is a complex along the MEP in order to obtain the best interpolation. For bimolecular reactions, the energies of the saddle point, wells, products, and extra points that are input via fu50 should be with respect to the sum of the energies of the reactants at the higher-level (which is set to zero). There are two test runs using the VTST-ISPE method, the ch5fu51tr1 test run which is a bimolecular reaction with the J1 surface, and the cmctr3 test run, which is a unimolecular reaction with two wells. For both reactions, we use the two turning points at the representative energy of the SCT calculation at 300 K as the extra points, and we evaluate the single-point energies with the MP2/6-31G** method.

The VTST-ISPE method can be applied for a minimum of three high-level points, namely the reactants, saddle point, and products. The calculations are performed using unit fu51 with the stationary points being required in the ISPEGEN section of unit fu51, and the

POINT section being included if and only if additional points along the reaction path are added. A POINT section is needed for each additional high-level point along the reaction path.

See Ref. 51 of Sect. 23 for more discussion.

10.J. Alternative methods for mapped interpolation

The mapped interpolation (MI) methods for IVTST-M calculations and for extrapolation beyond SLM and SLP are described in Ref. 48 of Sec. 21. In addition to the standard methods described in that paper, *POLYRATE* contains a few variants that may be used to test sensitivity to the approximations made in interpolation.

The first variant concerns the interpolation of $V_{\text{MEP}}(s)$ when there is only one reactant or product or when interpolation is based on a well. The default is to use a cubic polynomial in s to fill in 10 points between the end of the grid and the finite-distance stationary point. Variant 1, with the *nopolys* option for the IVTSTMOPT keyword in the PATH section of the fu5 input file, replaces the 10 points that in the standard algorithm are calculated from a local cubic polynomial by a single point located a small distance (δs) from the reactant (or product or well), whose energy is approximated using the lowest frequency at the stationary point. Thus the spline under tension is fit to $14 + m$ or $5 + m$ points rather than $23 + m$ points. (See Ref. 48 for the definitions of m and other details.)

The second variant allowed in MI calculations is an alternative way to calculate the effective reduced mass for CD-SCSAG calculations. The standard way involves interpolating each curvature component. The alternative method, keyword INTMUEFF in the GENERAL section of the fu31 input file, involves interpolating the argument of the exponential function in eq. (16) of Ref. 48 instead of interpolating curvature components. When the INTMUEFF keyword is used in conjunction with a VTST-IOC calculation, no higher-level correction to $\mu^{\text{SC}}(s)$ will be made. The INTMUEFF keyword is only available when using unit fu31.

10.K. Quantized-reactant-state tunneling calculation

For unimolecular reactions at low temperatures, most of the reactant molecules have their reaction-coordinate generalized vibrational modes in the ground state. Thus the semiclassical treatment, which assumes continuous tunneling energies, is inappropriate. Starting in version 6.2, we have included a new method that performs the tunneling calculations only at the energies corresponding to the eigenstates of the reaction-coordinate mode of the reactant for the case of a unimolecular reaction. (Note that all other modes are still assumed to be in their ground states.) The reaction-coordinate eigenstates can be determined using the harmonic or WKB approximation. Details of the theoretical background and an application to the diffusion of hydrogen atoms on a Ni surface can be found in Ref. 37 in Sect. 23. The keywords used to perform this type of tunneling calculation can be specified in the TUNNEL section in the unit fu5 input file. See Section 12.A.8 under the keyword QRST for details.

10.L. Curvilinear generalized normal mode analysis

The generalized normal mode analysis that is performed at each Hessian grid point (see Section 9.C) on the MEP may be carried out either in Cartesian coordinates or in internal coordinates. The COORD variable keyword in the PATH section makes it possible to select the type of coordinates to be used. The default option for the COORD variable keyword is *cart* which specifies Cartesian coordinates. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.) Two options are available to select a generalized normal mode analysis in nonredundant curvilinear coordinates: *curv1* and *curv2*. If either of the curvilinear internal coordinate options is selected for COORD, the Cartesian gradient vector \mathbf{G} and force constant matrix \mathbf{F} at the save point are first transformed to the curvilinear internal gradient vector \mathbf{g} and curvilinear internal force constant matrix \mathbf{f} using equations 30 and 31 in Ref. 37 of Sect. 22 or using equations 10 and 11 in Ref. 46 of Sect. 23.

When the option *curv1* is selected, the method of Ref. 41 of Sect. 23 is used for the generalized normal mode analysis. The option *curv1* is only available for nonredundant internal coordinates for reactions involving a total of three atoms, and it

may be used whether or not the geometry along the reaction path is linear. When the *curv1* option is chosen, the curvilinear internal coordinates—two stretches and a nondegenerate bend for nonlinear triatomics or two stretches and a degenerate bend for linear triatomics—are automatically defined by the program. In this method, the **B** matrix and **C** tensor are constructed using the difference Cartesian coordinate formulas from Pariseau et al. [Ref. 38, Sect. 22] for the bond stretch and nonlinear valence bend coordinates and from Hoy et al. [Ref. 39, Sect. 22] for the degenerate linear bend coordinate. The difference Cartesian **B** matrix and **C** tensor are then transformed to Cartesian coordinates.

When the option *curv2* or *curv3* is selected, the method of Ref. 43 or 46 of Sect. 23 is used for generalized normal mode analysis. The options *curv2* and *curv3* can be used for any number of atoms, but *curv2* is used for nonredundant internal coordinates, and *curv3* is used for redundant ones. Note that stretch, nonlinear or linear bend, and dihedral curvilinear internal coordinates are allowed for this method. (The extension to linear bends was presented in Ref. 45 of Sect. 23.) When the *curv2* option is chosen, the user is required to define the curvilinear internal coordinates used for the curvilinear generalized normal mode analysis via the INTDEF keyword. In this method the Cartesian **B** matrix and **C** tensor are constructed by using explicit FORTRAN statements [Ref. 40, Sect. 22] for the **B** matrix and **C** tensor derived by Challacombe and Cioslowski [Ref. 41, Sect. 22], except for the linear bend case where the derivation is presented in Ref. 45 of Sect. 21. If the curvilinear internal coordinate is a doubly degenerate linear bend, a symbol “=” is used to indicate the connections between atoms within the INTDEF keyword.

Next, at a nonstationary point along the MEP, the curvilinear internal force constant matrix **f** is projected as

$$\mathbf{f}^p = [1 - \mathbf{p}(\mathbf{B}\mathbf{u}\mathbf{B}^T)]\mathbf{f}[1 - (\mathbf{B}\mathbf{u}\mathbf{B}^T)\mathbf{p}],$$

where

$$\mathbf{p} = \frac{\mathbf{g}\mathbf{g}^T}{\mathbf{g}^T(\mathbf{B}\mathbf{u}\mathbf{B}^T)\mathbf{g}},$$

before applying the Wilson **GF** method [Ref. 36, Sect. 22] of vibrational analysis to obtain frequencies and generalized normal modes. The matrix **u** is a $(3N \times 3N)$ matrix with the reciprocal of each inverse mass appearing three times for each atom on the diagonal, and the matrix **B** is the Wilson B matrix.

If `POTENTIAL = unit30` or `unit40` (see Section 12.A.3), normal mode analyses at stationary points are always carried out in Cartesian coordinates. If `POTENTIAL = hooks`, normal mode analyses at minima are carried out in Cartesian coordinates, and the normal mode analysis at the transition state is carried out in curvilinear coordinates. Note that options for anharmonicity (e.g., `TOR` and `MORSE` keywords in Section 12.A.6) have not been implemented for curvilinear internal coordinates. Therefore, keyword options `curv1`, `curv2` and `curv3` should not be used with anharmonicity options.

At stationary points, one should obtain the same frequencies by using either curvilinear or rectilinear coordinates, except for small differences due to round off. However at non-stationary points the generalized normal mode frequencies do depend on the coordinate system — see Refs. 25, 41, 43, 45, and 46 in Section 23. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.)

In the current version, both `curv1` and `curv2` are restricted to use a nonredundant set of internal coordinates. Redundant internal coordinates are always chosen by the option `curv3`. In *POLYRATE*, the number of curvilinear internal coordinates cannot exceed $3N-6$ for the `curv1` and `curv2` options. The option `curv3` allows the number of internal coordinates to be as large as $3N + 6$.

10.M. IVTSTOFREQ option

In some cases, we found that the directly calculated frequencies of the lowest real-frequency modes are not stable or not accurate along the reaction path, and this can cause problems in evaluating the zero point energies, partition functions, and tunneling probabilities. To ameliorate this problem, we have added a new scheme for interpolating frequencies. This scheme is called IVTST-0-for-frequencies; it is a single-level analog of the method used for interpolating differences of frequencies in the VTST-IC method [Ref. 36, Sect. 23]. In the IVTST-0-for-frequencies scheme, the interpolation procedure has one of four possible forms, depending on the molecularity of the end points of the interpolation. Unimolecular reactions always have a molecularity of one at the negative s terminus. Interpolations for bimolecular reactions may have molecularity two at the negative s terminus, or – instead of extending all the way to reactants, the interpolation may extend only to a reactant-side well where the molecularity is one. Similar considerations apply on the product side.

MEP type 1. Bimolecular/bimolecular interpolation

A bimolecular/bimolecular interpolation is for a reaction that has two reactants and two products. Either there are no wells present on either side of the reaction path or (more likely) wells are present but are not used in the interpolation, i.e., they are not recognized. There are two functional forms used for this case,

a. If $(\omega^\ddagger > \omega_R)$ and $(\omega^\ddagger > \omega_P)$ or if $(\omega^\ddagger < \omega_R)$ and $(\omega^\ddagger < \omega_P)$, where $\omega_m^\ddagger = \omega_m(s = 0)$, $\omega_R = \omega_m(s = -\infty)$, and $\omega_P = \omega_m(s = +\infty)$, an Eckart function is used to approximate the frequencies along the reaction path:

$$\omega_m(s) = \frac{A_m Y}{1 + Y} + \frac{B_m Y}{(1 + Y)^2} + C_m \quad (1)$$

where

$$A_m = \omega_m(s = +\infty) - \omega_m(s = -\infty) \quad (2)$$

$$B_m = (2\omega_m^\ddagger - A_m - 2C_m) \pm 2\sqrt{[(\omega_m^\ddagger - C_m)(\omega_m^\ddagger - A_m - C_m)]} \quad (3)$$

$$C_m = \omega_m(s = -\infty) \quad (4)$$

$$Y = \exp\left(\frac{s - S_{0,m}}{L}\right) \quad (5)$$

$$S_{0,m} = -L \ln\left(\frac{A_m + B_m}{B_m - A_m}\right). \quad (6)$$

Unlike the original VTST-IC algorithm, we use a single-level analytical expression for estimating the value of L instead of using a fit involving a lower level. The analytical expression is [Ref. 26, Sect. 23]

$$L^2 = -\frac{2V^\ddagger(V^\ddagger - A)}{\mu(\omega^\ddagger)^2 B} \quad (7)$$

$$A = V_{\text{MEP}}(s = +\infty) - V_{\text{MEP}}(s = -\infty) \quad (8)$$

$$C = V_{\text{MEP}}(s = -\infty) \quad (9)$$

$$B = (2V^\ddagger - A - 2C) \pm 2\sqrt{[(V^\ddagger - C)(V^\ddagger - A - C)]} \quad (10)$$

$$V^\ddagger = V_{\text{MEP}}(s = 0) \quad (11)$$

where V_{MEP} is the potential energy along the reaction path, and ω^\ddagger is the imaginary frequency at the saddle point.

b. If $(\omega_R > \omega^\ddagger > \omega_P)$ or if $(\omega_R < \omega^\ddagger < \omega_P)$, a hyperbolic tangent function is used to approximate the frequencies along the reaction path:

$$\omega_m(s) = \frac{1}{2} A_m \tanh\left(\frac{s - T_{0,m}}{L}\right) + E_m \quad (12)$$

where

$$E_m = \frac{\omega_m(s = +\infty) + \omega_m(s = -\infty)}{2} \quad (13)$$

$$T_{0,m} = -\frac{L}{2} \ln\left(\frac{1 + x_m}{1 - x_m}\right) \quad (14)$$

$$x_m = 2\frac{\omega_m^\ddagger - E_m}{A_m} \quad (15)$$

c. For cases where two consecutive of the three frequencies are the same (i.e., $\omega^\ddagger = \omega_R$ or $\omega^\ddagger = \omega_P$), a constant value or used at one side of the reaction path and a symmetric Eckart function fit is used on the other. For example, if $\omega_R = \omega^\ddagger \neq \omega_P$, then

$$\omega_m(s) = \begin{cases} \omega_R & s \leq 0 \\ \omega_P + B_m Y / (1 + Y)^2 & s > 0 \end{cases} \quad (16)$$

where

$$B_m = (2\omega_m^\ddagger - \omega_P) \pm \sqrt{2}(\omega_m^\ddagger - \omega_P) \quad (17)$$

$$Y = \exp\left(\frac{s}{L}\right) \quad (18)$$

and L is given by eqs. (7)–(11).

MEP type 2. Two reactants to one product (or well)

We introduce the notation P–W which denotes the product *or* the well, whichever is being used as the right-side edge of the interpolation region. In this type of reaction the function used to fit the frequencies depends on the s value, which is in the range of $-\infty < s < s_{\text{P-W}}$.

a. If $s < 0$, a hyperbolic tangent function is used to approximate the frequencies as described in the previous section for base 1b.

b. If $0 \leq s \leq s_{\text{P-W}}$, a cut-off hyperbolic tangent is used to approximate the frequencies:

$$\omega_m(s) = \frac{1}{2} A_m \tanh\left(\frac{s - T_{0,m}}{L} \left(\frac{s_{\text{P-W}}}{s_{\text{P-W}} - s}\right)\right) + E_m \quad (19)$$

where A_m , $T_{0,m}$, and E_m are the same as previous section. L is evaluated using eqs. (7)–(11). This makes $\omega_m(s)$ continuous with one continuous derivative.

If $\omega_{m,\text{R}} = \omega_m^\ddagger$ or $\omega_m^\ddagger = \omega_{m,\text{P-W}}$, then the value of $\omega_{m,\text{R}}$ or $\omega_{m,\text{P-W}}$ is modified by ± 2 cm^{-1} to make the function monotonic.

MEP type 3. One reactant (or well) to two products

This type of reaction is similar to type II but the functional forms are reflected through $s = 0$.

MEP type 4. One reactant (or well) to one product (or well)

We introduce the notation R–W which denotes the reactant or the reactant-side well, whichever is being used as the left-side edge of the interpolation region.

a. If $(\omega^\ddagger > \omega_{\text{R-W}})$ and $(\omega^\ddagger > \omega_{\text{P-W}})$ or if $(\omega^\ddagger < \omega_{\text{R-W}})$ and $(\omega^\ddagger < \omega_{\text{P-W}})$, two cut-off Gaussian functions are used to approximate the frequencies along the reaction path:

$$\omega_m(s) = A_{m,x} \exp\left(-\frac{B_{m,x}}{1 - \left(\frac{s}{s_x}\right)^2}\right) + C_{m,x}; \quad x = \text{R-W,P-W} \quad (20)$$

$$A_{m,x} = (\omega_m^\ddagger - \omega_x) \exp(B_{m,x}); \quad x = \text{R-W,P-W} \quad (21)$$

$$C_{m,x} = \omega_x; \quad x = \text{R-W,P-W}. \quad (22)$$

The parameter $B_{m,x}$ is evaluated using a single-level analytical expression rather than a fit to a lower-level surface. In particular we use

$$B_{m,x} = \frac{s_x^2 \mu (\omega^\ddagger)^2}{2(V^\ddagger - V_x)}; \quad (23)$$

b. If $\omega_{m,R-W} = \omega_m^\ddagger$ or $\omega_{m,P-W} = \omega_m^\ddagger$ then the frequency is set equal to a constant on one side of the saddle and approximated by a symmetrical cut-off Gaussian is used for the other side on the reaction path. For example, if $\omega_{m,R-W} = \omega_m^\ddagger \neq \omega_{m,P-W}$, then the frequency is set to ω_{R-W} when $s \leq 0$ and evaluated using eqns. (20–23) when $s > 0$.

c. If $(\omega_{R-W} > \omega^\ddagger > \omega_{P-W})$ or if $(\omega_{R-W} < \omega^\ddagger < \omega_{P-W})$, a cut-off hyperbolic tangent function is used to approximate the frequencies along the reaction path.

$$\omega_m(s) = \frac{1}{2} A_m \tanh \left[\frac{s - T_{0,m}}{L} \left(\frac{s_{P-W}}{s_{P-W} - s} \right) \left(\frac{|s_{R-W}|}{s - s_{R-W}} \right) \right] + E_m \quad (24)$$

where A_m , $T_{0,m}$, and E_m are determined by eqns. (2,13–16), and L is evaluated using eq. (7).

The keyword option IVTSTOFREQ is used to apply the IVTSTOFREQ method for interpolating the vibrational frequencies along the path. If either side of the reaction path is unimolecular or if one recognizes an intermediate structure (indicated by *WELLR or *WELLP sections), then an arc distance from saddle point has to be provided for the interpolation (by using the SR and SP keyword options).

The equations in this section are not currently published; the only available reference is this manual.

10.N. LCG4 option

Beginning with version 8.6, the options LCT3 and LCT4 are both available, but one cannot use both of them in the same run. These two options replace the previous LCT option.

The LCTOPT keyword includes four new options.

The option *vadavg* can be used with LCT3 or LCT4, and it averages the vibrationally adiabatic potential in a doubly vibrationally adiabatic region. Specifically,

$$V_{\text{eff}}(\xi) = V_{\text{eff}}^{\text{III}}(\tilde{s}_0, \xi_{\text{I}}) + \frac{\xi - \xi_{\text{III}}}{\xi_{\text{I}} - \xi_{\text{III}}} \left[V_{\text{eff}}^{\text{I}}(\tilde{s}_0, \xi_{\text{I}}) - V_{\text{eff}}^{\text{III}}(\tilde{s}_0, \xi_{\text{III}}) \right]$$

where ξ_{III} and ξ_I indicate the beginning and end of the region that overlaps. The default is *novadavg* and the vibrational adiabatic potential in this region is given by

$$V_{\text{eff}}(\xi) = \min[V_{\text{eff}}^{\text{III}}(\tilde{s}_0, \xi_{III}), V_{\text{eff}}^{\text{I}}(\tilde{s}_0, \xi_I)]$$

The option *vefsrch numb* only can be used with the LCT4 option, and it switches on the search for nonadiabatic regions that are completely adiabatic in the LCT3 method. In particular, it becomes operative at the tunneling energy after (higher than) that for which the last nonadiabatic region was found according to the LCG3 algorithm, and the search is carried out for a total of *numb* energies beginning at that point. The search locates the maximum of the vibrationally adiabatic potential at a given tunneling energy and compares the value with the actual potential. If the actual potential is smaller than the adiabatic potential, then the procedure ends, otherwise the nonadiabatic region is extended until the vibrationally adiabatic potential is smaller than the actual potential. The default is *vefsrch 999* because this search is required for correct LCG4 calculations. There is also an option called *novvefsrch* that is equivalent to *vefsrch 0*.

For example, if 80 linear paths are calculated, and the 40 lowest-energy paths have nonadiabatic regions in LCG3, setting *numb* equal to 20 will cause the program to search for nonadiabatic regions along paths 41–60. This avoids the search for nonadiabatic regions along the 20 highest-energy paths where it is likely that nonadiabatic regions may exist only due to round-off; it allows one to avoid extra calculations at these energies very close to the effective barrier top because they have no significant effect on the results.

A restart option has been added, and it is described by two new keywords in LCTOPT. In particular, *lctstr* stores the information in unit fu49 (this is the default), and *lctsrst* restarts a calculation from unit poly.fu48. An old calculation stored in fu49 can be restored directly if fu49 is renamed as fu48. The default is *no_lctsrst*. If *lctsrst* is specified, but the unit poly.fu48 does not exist, the program stops automatically. With *lctsrst* the LCTOPT options have to be the same as in previous calculations, that is, *ngamp* and *ngtheta* cannot be changed.

To incorporate the restart option in *POLYRATE* the modules were modified. The old LCG3 subroutine (renamed in version 8.5 as LCG) reads from unit fu48 and stores the information in the common block LCRST1 which is linked to LCVEF.

This restart option will work in *POLYRATE*, *MORATE*, and *GAUSSRATE*. When the restart option is used with *GAUSSRATE*, the restarting data for the LCT calculation will be read from fu48 instead of esp.fu83.

This restarting procedure is simple and flexible, and it allows the use of an old LCG3 fu48 file to restart an LCG3 calculation or to start a new LCG4 calculation, or vice versa. All the combinations are possible. The restart option is specially useful when we want to calculate tunneling into vibrational accessible excited states. In a first run we can do the calculation of the ground state large curvature transmission factor. This information is saved in unit FU49. We can include tunneling into vibrationally excited final states in a second run if still worth it after looking at the output.

Further details of the LCG4 algorithm are provided in Ref. 54 in Sect. 23.

10.O. MS-VTST, MP-VTST, FMS-VTST, and FMR-VSTT calculations by using both the *POLYRATE* and *MSTOR* programs

For molecules and transition states that have multiple conformations generated by torsions, we recommend MS-VTST [61], MP-VTST [62,63,72], FMS-VTST [68], or FMP-VTST [68] calculations. Such calculations can be carried out by making runs with *POLYRATE* and *MSTOR* programs. The *MSTOR* program [64-66] is a program for applying the internal-coordinate torsional anharmonicity method [70,71,68] for treating torsional anharmonicity (MS-T); it is available as a separate package at

<http://comp.chem.umn.edu/mstor/>.

The internal-coordinate torsional anharmonicity method is a practical thermochemical method for including multiple structures and torsional anharmonicity to calculate conformational–vibrational–rotational partition function for complex systems. This scheme obviates the need of earlier approaches to identify torsional motions with specific normal modes, and it approximately accounts for coupling of torsional modes to one another and to other modes. It includes the conformational free energy associated with the existence of multiple structures (conformations) and the additional effects of torsional

potential anharmonicity. The original paper [70] included methods based on a subset of structures as well as the method using all structures. The method containing all structures was originally call MS-AS, but now we usually use all structures and, as explained in ref. [64], we simply refer to the method employing all structures as MS-T.

In the original papers introducing MS-VTST [61] and MP-VTST [62,63,72], we included multistructural torsional anharmonicity only at stationary points; tunneling transmission coefficients and recrossing transmission coefficients were calculated in the quasiharmonic (QH) approximation. (A quasiharmonic calculation is one that uses the harmonic oscillator formulas for vibrational partition functions but with effective (scaled) frequencies rather than the harmonic frequencies; the effective frequencies are usually chosen to improve the estimate of the zero point energy [69].) In Ref. [68] we introduced the capability to include multistructural anharmonicity all along the reaction path. To be precise, we will label calculations by the original methods as MS-VTST [61] and MP-VTST, and we will label calculations with multistructural anharmonicity all along the reaction path [68] as full MS-VTST (FMS-VTST) and full MP-VTST (FMP-VTST).

In carrying out MS-VTST and MP-VTST calculations, *POLYRATE* is used to carry out single-structure quasiharmonic (SS-QH) calculations for reaction paths. In carrying out FMS-VTST and FMP-VTST calculations, *POLYRATE* is used to carry out single-structure quasiharmonic what we shall call single-structure torsion (SS-T) calculations. Throughout this discussion, we use the word structure as a synonym for conformation. An SS calculation involves a single conformational structure of each reactant and a single conformational structure of the transition state. SS-QH calculations use the QH approximation along the reaction path, and SS-T calculations use the MS-T approximation along the reaction path.

A note about chirality: when one is carrying out a single-structure calculation to compare with experiment, if the structure has a distinguishable mirror image, then one needs to include an extra factor of two in its rate constant, as discussed elsewhere [67]. However, in this section, we literally mean a single structure and the factor of two is not included in such a case.

Some preliminaries about notation: (1) When we say "variational effects" we mean the difference between a variational transition state theory calculation and a conventional transition state theory calculation. In MS-VTST, MP-VTST, FMS-VTST, and FMP-VTST calculations, variational effects are included by multiplying a conventional transition state theory rate constant by a transmission coefficient that is called a recrossing transmission coefficient. (2) The methods explained here may be used with a variety of approximations for the tunneling, for example, ZCT, SCT, LCT, μ OMT,

or LAT. In general we use the abbreviation MT ("multidimensional tunneling") to refer to any of these. (3) In our original paper

This subsection explains how to perform MS-VTST, MP-VTST, FMS-VTST, and FMP-VTST calculations by combining results from *POLYRATE* and MSTOR. In brief, the MSTOR program is used to perform all the calculations for the stationary points (reactants, saddle points, and products), and *POLYRATE* is used to perform SS-QH or SS-MP reaction path calculations to obtain variational effects and transmission coefficients.

In order to carry out either an MS-VTST or an MP-VTST calculation, the first step is to find all the conformational structures of the transition state (that is, all the saddle points leading from reactants to products along parallel reaction paths (as contrasted to reaction paths in series)) and all the structures of the reactant or reactants. Structures of the transition state are called transition structures, and structures of the reactant(s) are called reactant equilibrium structures or just reactant structures. Next one numbers the transition structures so that the lowest-energy one is called $k = 1$, and in another list one orders the reactant equilibrium structures so that the lowest-energy one is called $k = 1$. The energy difference between the lowest-energy transition structure and the lowest-energy reactant equilibrium structure is called the classical barrier height, V^\ddagger .

Then one uses MSTOR to calculate MS-T partition functions for the transition state and the reactant(s). For the reactant of a unimolecular reaction, this partition function is called $Q^{\text{MS-T,R}}$. For the reactants of a bimolecular reaction, the product of the partition functions of the two reactants (Q^{R1} and Q^{R2}) is also called $Q^{\text{MS-T,R}}$, but the quantity we will need to calculate rate constants in the bimolecular case is Φ (the partition function per unit volume) rather than Q (the partition function) so that one has to convert Q to Φ when calculating bimolecular rate constants. Note also that it is possible that MS-T will be used for only one of the two reactants. If the reactant R2 does not have torsion, then one may use a simpler method, for example, the harmonic or quasiharmonic approximation, for Q^{R2} . Then $\Phi^{\text{MS-T,R}}$ is the bimolecular reactants' partition function per unit volume (with its zero of energy at the sum of the potential energies of the lowest-energy reactant structures), and $Q^{\text{MS-T,R}}$ is the total partition function of a unimolecular reactant (with its zero of energy at the potential energy of the lowest-energy reactant structure).

The partition function for the transition state is called $Q^{\text{MS-T},\ddagger}$. Note that this partition function is sum over contributions from distinguishable structures:

$$Q^{\text{MS-T},\ddagger} = \sum_{k=1}^K Q_k^{\text{T},\ddagger} \quad (1)$$

where K is the number of distinguishable conformational structures of the transition state, and $Q_k^{\text{T},\ddagger}$ is the rovibrational partition function of saddle point k with torsional potential anharmonicity (T). Note that the difference in energy between the zero of energy used for the transition state partition functions $Q_k^{\text{T},\ddagger}$ and that used for the reactant partition function is taken here to be the classical barrier height, which is the potential energy of the lowest-energy conformational structure ($k = 1$) of the saddle point minus the potential energy of the lowest-energy equilibrium structure of the reactant.

The next step is to perform one or more SS calculations with *POLYRATE*; this is explained next.

10.O.1. MS-VTST

The MS-VTST rate constants including multidimensional tunneling are given by the following formulas for a bimolecular reaction and a unimolecular reaction respectively:

$$k_{\text{bimol}}^{\text{MS-CVT/MT}} = \kappa_1^{\text{QH}} \Gamma_1^{\text{QH-CVT}} \frac{k_B T}{h} \frac{Q_{\text{elec}}^{\ddagger} \sum_{k=1}^K Q_k^{\text{T},\ddagger}}{\Phi^{\text{MS-T,R}}} \exp(-\beta V^{\ddagger}) \quad (2)$$

$$k_{\text{unimol}}^{\text{MS-CVT/MT}} = \kappa_1^{\text{QH}} \Gamma_1^{\text{QH-CVT}} \frac{k_B T}{h} \frac{Q_{\text{elec}}^{\ddagger} \sum_{k=1}^K Q_k^{\text{T},\ddagger}}{Q^{\text{MS-T,R}}} \exp(-\beta V^{\ddagger}) \quad (3)$$

where κ_1^{QH} is the quasiharmonic tunneling transmission coefficient for the reaction path passing through transition structure 1, β is $1/k_B T$, k_B is Boltzmann's constant, T is temperature, h is Planck's constant, $Q_{\text{elec}}^{\ddagger}$ is the electronic partition function of the conventional transition state (which is denoted as \ddagger); and the CVT recrossing transmission coefficient for structure 1 is $\Gamma_1^{\text{QH-CVT}}$, which is given by the ratio of the quasiharmonic CVT rate constant for the reaction path through transition structure 1 to the quasiharmonic conventional TST one for transition structure 1.

In order to obtain κ_1^{QH} and $\Gamma_1^{\text{QH-CVT}}$, one performs an SS-QH calculation with *POLYRATE*. The κ_1^{QH} is printed out in the *POLYRATE* output file and $\Gamma_1^{\text{QH-CVT}}$ is calculated

$$\Gamma_1^{\text{QH-CVT}} = \frac{k^{(1),\text{QH-CVT}}}{k^{(1),\text{QH-TST}}} \quad (4)$$

10.O.2. FMS-VTST

An FMS-VTST calculation is the same as an MS-VTST one except for the transmission coefficients. One replaces κ_1^{QH} and $\Gamma_1^{\text{QH-CVT}}$ by κ_1^{T} and $\Gamma_1^{\text{T-CVT}}$, which are obtained by running SS-T calculations in *POLYRATE* with the keyword *SSTOR*. In particular, the recrossing transmission coefficient $\Gamma_1^{\text{T-CVT}}$ is calculated as

$$\Gamma_1^{\text{T-CVT}} = \frac{k^{(1),\text{T-CVT}}}{k^{(1),\text{T-TST}}} \quad (5)$$

10.O.3. MP-VTST

In MP-VTST, we calculate the transmission coefficient as a properly weighted average over tunneling transmission coefficients and variational transmission coefficients for several reaction paths instead of calculating transmission coefficients only from one path as is done in MS-VTST. This subsection gives the procedure to perform MP-VTST calculations.

The MP-VTST rate constants including multidimensional tunneling are given by the following formulas for a bimolecular reaction and a unimolecular reaction respectively:

$$\begin{aligned} k_{\text{bimol}}^{\text{MP-CVT/MT}} &= \frac{k_B T}{h} \frac{Q_{\text{elec}}^\ddagger}{\Phi^{\text{MS-T,R}}} \exp(-\beta V^\ddagger) \sum_{k=1}^K \kappa_k^{\text{QH}} \Gamma_k^{\text{QH-CVT}} Q_k^{\text{T},\ddagger} \\ &\approx \gamma \frac{k_B T}{h} \frac{Q_{\text{elec}}^\ddagger}{\Phi^{\text{MS-T,R}}} \sum_{k=1}^K Q_k^{\text{T},\ddagger} \exp(-\beta V^\ddagger) \end{aligned} \quad (6)$$

$$k_{\text{unimol}}^{\text{MP-CVT/MT}} \approx \gamma \frac{k_B T}{h} \frac{Q_{\text{elec}}^\ddagger}{Q^{\text{MS-T,R}}} \sum_{k=1}^K Q_k^{\text{T},\ddagger} \exp(-\beta V^\ddagger) \quad (7)$$

where the MP-VTST generalized transmission coefficient γ is approximated by

$$\gamma = \frac{\sum_{k=1}^P \kappa_k^{\text{QH}} \Gamma_k^{\text{QH-CVT}} \mathcal{Q}_k^{\text{T},\ddagger}}{\sum_{k=1}^P \mathcal{Q}_k^{\text{T},\ddagger}} \quad (8)$$

To evaluate the quantities in these formulas, we have to make P runs of *POLYRATE* (where $P \leq K$), one for each of the P user-selected distinguishable structures of the transition state (note that these structures are numbered by $k = 1, \dots, P$). Each of the runs gives a quasiharmonic CVT rate constant $k^{(k),\text{QH-CVT}}$, a quasiharmonic conventional TST rate constant $k^{(k),\text{QH-TST}}$, a quasiharmonic tunneling transmission coefficient κ_k^{QH} , and a quasiharmonic CVT recrossing transmission coefficient $\Gamma_k^{\text{QH-CVT}}$. The first three are printed in the *POLYRATE* output file, and the fourth is given by

$$\Gamma_k^{\text{QH-CVT}} = \frac{k^{(k),\text{QH-CVT}}}{k^{(k),\text{QH-TST}}} \quad (9)$$

The other quantities needed for MP-VTST rate constants can be calculated by using the *MSTOR* program.

Note also that Γ_k^{CVT} is independent of which reactant structure is used, but the choice of reactant structure affects the lower-energy limit of the integration used to calculate κ_k , and thus it could affect the values calculated for the rate constants at low temperature. Therefore all the additional *Polyrate* runs ($k = 2, \dots, P$) needed to upgrade MS-VTST to MP-VTST should use structure k of the transition state and the lowest-energy structure of the reactant(s). Note that one has to calculate a reaction path long enough to make sure the κ_k is converged. In case the saddle point k is connected with a reactant(s) conformation much higher in energy than the lowest-energy structure of the reactant(s), there is possibilities that the κ_k is overestimated because of the program is trying to integrate κ_k to the lowest-energy structure of the reactant(s). This is one aspect if the calculation that needs to be monitored carefully if calculations are made at low temperature.

Note that the wavy equal sign in equation (6) can be replaced by a straight equal sign if P equals K . When $P = 1$, the MP-VTST rate reduces to the MS-VTST rate.

Alternatively one also can calculate the MP-VTST rate constants by the following procedure. Let's define the multistructural torsional anharmonicity factors for reactants (R1 and R2 for bimolecular reactions, R1 = R for unimolecular reactions) and for saddle point 1 (lowest-energy saddle point) as

$$F^{R1} = \frac{Q^{\text{MS-T,R1}}}{Q_1^{\text{QH,R1}}} \quad (10)$$

$$F^{R2} = \frac{Q^{\text{MS-T,R2}}}{Q_1^{\text{QH,R2}}} \quad (11)$$

$$F^{\ddagger,1} = \frac{Q^{\text{MS-T},\ddagger}}{Q_1^{\text{QH},\ddagger}} \quad (12)$$

Then we obtain the MP-VTST rate constants as

$$k_{\text{bimol}}^{\text{MP-CVT/MT}} = \gamma \frac{F^{\ddagger,1}}{F^{R1} F^{R2}} k_{\text{bimol}}^{(1),\text{QH-TST}} \quad (13)$$

$$k_{\text{unimol}}^{\text{MP-CVT/MT}} = \gamma \frac{F^{\ddagger,1}}{F^{R1}} k_{\text{unimol}}^{(1),\text{QH-TST}} \quad (14)$$

10.O.4. FMP-VTST

An FMP-VTST calculation is the same as an MP-VTST one except for the transmission coefficients. One replaces κ_k^{QH} and $\Gamma_k^{\text{QH-CVT}}$ by κ_k^{T} and $\Gamma_k^{\text{T-CVT}}$ in equation (8); these quantities are obtained by running SS-T calculations in *POLYRATE* with the keyword SSTOR. In particular, the recrossing transmission coefficient $\Gamma_k^{\text{T-CVT}}$ is calculated as

$$\Gamma_k^{\text{T-CVT}} = \frac{k^{(k),\text{T-CVT}}}{k^{(k),\text{T-TST}}} \quad (15)$$

Part III Theory Background for VRC-VTST

11. THEORETICAL BACKGROUND ON VRC-VTST

For an association reaction without a barrier or with a negligible barrier, we recommend using the multifaceted (MF) extension of variable-reaction-coordinate variational transition state theory (VRC-VTST). The references for VRC-VTST is Y. Georgievskii and S. J. Klippenstein, *J. Chem. Phys.* **2003**, *118*, 5442, and the reference for the extension of VRC-VTST to use MF dividing surfaces is Y. Georgievskii and S. J. Klippenstein, *J. Phys. Chem. A* **2003**, *107*, 9776.

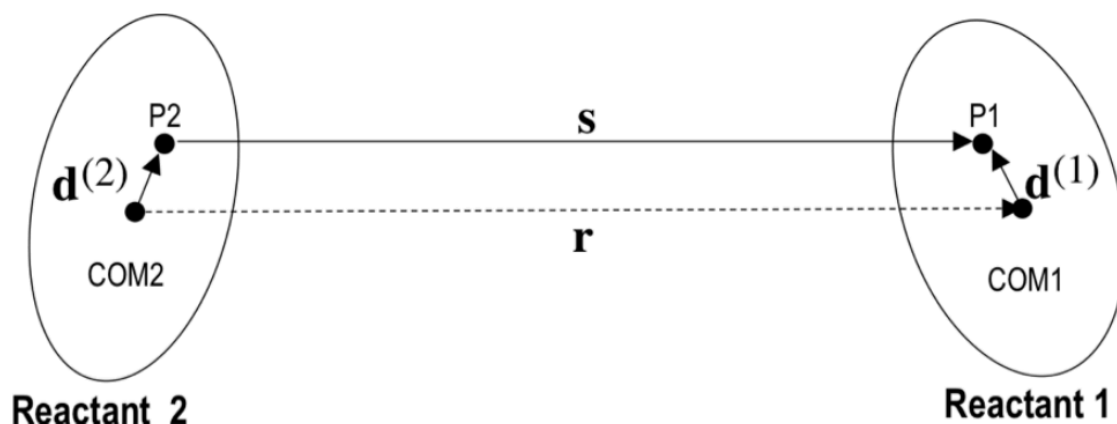
In VRC-VTST, the reaction coordinate and dividing surfaces are defined in a different way than in RP-VTST. In VRC-VTST, the reaction coordinate s is defined by the distance between a pivot point on one reactant and that on another, and the dividing surfaces are defined in terms of pivot points tied to each of the reactants. Choices of the number and location of these pivot points yield different definitions of the reaction coordinates and hence of the generalized transition state theory dividing surfaces, and reactive flux is variationally minimized with respect to both these choices and the distance between pivot points in the two reactants. The original version of VRC-VTST employed one pivot point on each reactant, and the MF extension involves two and more pivot points on one or both reactants. In this manual we consider the use of MF dividing surfaces to a special case of VRC-VTST. The evaluations of the reactive flux involves Monte Carlo integration over the coordinate space degrees of freedom of the classical phase space representation of the generalized transition state partition function for arbitrary orientations of the reagents over a wide range of interfragment separations.

Because there is no saddle point for a barrierless association reaction, the conventional TST rate is not calculated. The START section only needs the electronic degeneracy information for electronic partition function calculations. Association rate constants can be calculated by CVT, μ VT, or $E, J - \mu$ VT. Tunneling calculations cannot be performed for this type of calculation.

11.A. Variable reaction coordinate

When a single-faceted dividing surface is used, the reaction coordinate s for a barrierless association reaction is defined as the distance between two pivot points, one associated

with each of the two reactants. Since s is defined as a distance here, its value is always positive. Thus, it has a different meaning from the s used in RP-VTST. The following scheme shows the definition of the reaction coordinate s for the case of a single-faceted dividing surface.



Here, the two ovals represent the two reactants. COM1 and COM2 are their centers of mass. P1 and P2 are pivot points associated with reactant 1 and reactant 2, respectively. $\mathbf{d}(1)$ and $\mathbf{d}(2)$ are vectors connecting a pivot point with its associated reactant's center of mass, respectively. When the distance s between two pivot points is fixed, the directions of \mathbf{s} , $\mathbf{d}(1)$, and $\mathbf{d}(2)$ are sampled over the whole phase space simultaneously. The lengths of these vectors are fixed during the sampling over their orientations.

In the VRC-VTST approach, the modes of motion are divided into two sets, conserved modes and transitional modes. The conserved modes correspond to the vibrational modes of the two reactants. In the complex of two reactants, these modes usually correspond to the modes with highest vibrational frequencies. VRC-VTST, at least in its present state of development, assumes that changes of conserved modes are negligible during the reaction process. The transitional modes include the other vibrational modes except conserved modes in the complex and the overall rotations. These modes change from free rotation and translation at infinite separation of the reactants to vibrational motion and overall rotation during the reaction process. The new vibrational modes are very floppy and are best thought of as liberations.

The high-pressure limit rate constant for the generalized transition state at location s along the reaction coordinate and at the temperature T can be expressed as

$$k(T,s) = \frac{\hbar^2}{2\pi} g_e \frac{\sigma_1 \sigma_2}{\sigma^\ddagger Q_1 Q_2} \left(\frac{2\pi}{\mu k_B T} \right)^{\frac{3}{2}} \int dE e^{-E/k_B T} dJ N(E,J,s) \quad (1)$$

where s is the value of reaction coordinate, which is defined above as the distance between two pivot points; g_e is the ratio of the electronic partition function of the transition state to the product of the electronic partition functions of reactants; μ is the reduced mass; Q_1 and Q_2 are the rotational partition functions of the reactants calculated without symmetry numbers; J is the unitless total angular momentum quantum number; $N(E,J,s)$ is the number of accessible states of the generalized transition state s for total energy E and angular momentum $J\hbar$; and σ_1 , σ_2 , and σ^\ddagger are the rotational symmetry numbers for the reactants and transition state, respectively.

The keyword SIGMAF in RATE section (see Section 13.A.6) is equal to $\sigma_1 \sigma_2 / \sigma^\ddagger$. For reactions in which the two reactants are the same, SIGMAF is 0.5; for reactions in which the two reactants are different, SIGMAF is 1.0. This is explained in the following paragraph. As background for this explanation, we first note that eq. (137) in the review by Bao and Truhlar¹ is identical to eq. (4) in the *PNAS* paper by Bao et al.,² although in the *PNAS* paper we include the rotational symmetry factors in the reactant partition functions, whereas in eq. (137) of the review we separated them. Note also that σ_1 and σ_2 should have their correct symmetry numbers as usual.

The next question is: What is σ^\ddagger ? Georgievskii and Klippenstein³ (in their eq. 2) define

$$\sigma = \sigma_1 \sigma_2 / \sigma^\ddagger$$

and they make the assumption that – since the transition state is loose – there is free rotation of both species, such that the transition state partition function contains the product of the two rotational partition functions. Then

$$\sigma^\ddagger = \sigma_1 \sigma_2 \sigma_{\text{add}}$$

¹ J. L. Bao and D. G. Truhlar, Variational Transition State Theory: Theoretical Framework and Recent Developments, *Chem. Soc. Rev.*, 2017, **46**, 7548-7596. doi.org/10.1039/C7CS00602K

² J. L. Bao, X. Zhang and D. G. Truhlar, Association of CF₂ and Dissociation of C₂F₄ by Variational Transition State Theory and System-Specific Quantum RRK Theory, *Proc. Natl. Acad. Sci. U. S. A.*, 2016, **113**, 13606. doi.org/10.1073/pnas.1616208113

³ Y. Georgievskii and S. J. Klippenstein, Long-range transition state theory, *J. Chem. Phys.*, 2005, **122**, 194103. doi.org/10.1063/1.1899603

where σ_{add} is the additional symmetry gained in the transition state. Because we sample all possible orientations (Euler angles) of the transition state classically in VRP-TST, we must divide by two when the association reactants are identical because the phase space may be divided into two quantum mechanically indistinguishable spaces when the two reactants are identical. This means that σ_{add} equals 1 if the species are different and equals 2 if they are the same. Therefore $\sigma_1\sigma_2/\sigma^\ddagger$ equals 1 if the species are different and equals 0.5 if they are the same. One could call this the loose-transition-state assumption for the symmetry number of the transition state.

The number of accessible states for the transitional modes with total energy E and angular momentum quantum number J can be written as

$$N(E, J, s) = \langle N_q(E, J, Q, s) \rangle_{\Omega} \quad (2)$$

where $\langle \dots \rangle_{\Omega}$ denotes the average over all possible orientations of the two reactants and of the orientation of the vector s ; Q means a specific configuration of the system. Since this multidimensional integral is carried out by Monte Carlo integration, the configurations of the system are randomly sampled over the whole phase space corresponding to a given s . In our implementation, one specific random configuration is determined by the following procedure:

(1) P1 is always taken as the origin of the system for every random configuration.

(2) The position of P2 is determined by a random solid angle $\Omega_{12} \equiv (\theta_{12}, \phi_{12})$

$$\mathbf{x}_{P2} = \mathbf{A}_{12}^{-1}(\Omega_{12}) \begin{pmatrix} 0 \\ 0 \\ s \end{pmatrix} \quad (3)$$

where \mathbf{x}_{P2} is the Cartesian coordinate of pivot point P2; $\mathbf{A}_{12}^{-1}(\Omega_{12})$ is the inverse rotation matrix,

$$\mathbf{A}_{12}^{-1} = \begin{pmatrix} \cos \theta_{12} \cos \phi_{12} & -\sin \phi_{12} & \sin \theta_{12} \cos \phi_{12} \\ \cos \theta_{12} \sin \phi_{12} & \cos \phi_{12} & \sin \theta_{12} \sin \phi_{12} \\ -\sin \theta_{12} & 0 & \cos \theta_{12} \end{pmatrix} \quad (4)$$

(3) Since the Cartesian coordinates of P1 (\mathbf{x}_{P1}) and P2 (\mathbf{x}_{P2}) are known, the Cartesian coordinates of the center of mass of the k th reactant ($\mathbf{x}_{\text{com}k}$) are given by

$$\mathbf{x}_{\text{com}k} = \mathbf{A}_k^{-1}(\Omega_k) \begin{pmatrix} 0 \\ 0 \\ d^{(k)} \end{pmatrix} + \mathbf{x}_{P_k}, \quad (k = 1, 2) \quad (5)$$

$$\mathbf{A}_k^{-1}(\Omega_k) = \begin{pmatrix} -\sin\psi_k \sin\phi_k + \cos\theta_k \cos\psi_k \cos\phi_k & -\cos\psi_k \sin\phi_k - \cos\theta_k \cos\phi_k \sin\psi_k & \sin\theta_k \cos\phi_k \\ \sin\psi_k \cos\phi_k + \cos\theta_k \cos\psi_k \sin\phi_k & \cos\psi_k \cos\phi_k - \cos\theta_k \sin\phi_k \sin\psi_k & \sin\theta_k \sin\phi_k \\ -\cos\psi_k \sin\theta_k & \sin\psi_k \sin\theta_k & \cos\theta_k \end{pmatrix} \quad (6)$$

where $\mathbf{A}_k^{-1}(\Omega_k)$ is the inverse rotation matrix, and $\Omega_k = (\psi_k, \theta_k, \phi_k)$ is solid angle for the k th reactant; $d^{(k)}$ is the length of vector $\mathbf{d}^{(k)}$.

(4) Since the internal geometries of the reactants are fixed at the input ones, and their center of mass positions are determined in step 3, the coordinates of the generalized transition states are simply obtained by a translation.

The quantity $N_q(E, J, Q, s)$ can be loosely viewed as number of accessible states for a specific configuration Q with total energy E and angular momentum $J\hbar$, which can be expressed as

$$N_q(E, J, Q, s) = \Phi J^2 \mu s^2 \frac{8\pi^{k_a-1} \prod_{k=1}^2 \prod_{i=1}^{v_k} \sqrt{2\pi I_i^{(k)}}}{\Gamma(v/2-1) \prod_{i=1}^3 \sqrt{2\pi I_i(Q)}} \times \left\langle \max \left(E - V(Q) - \sum_{i=1}^3 \frac{J_i^2}{2I_i(Q)}, 0 \right)^{\frac{v}{2}-2} \right\rangle_{\Omega_J}, \quad (7)$$

$$\Phi = \sqrt{1 + \mu \sum_{k=1}^2 \sum_{i=1}^{v_k} \left(\mathbf{n}^{(12)} \times \mathbf{d}^{(k)} \cdot \mathbf{n}_i^{(k)} \right)^2 / I_i^{(k)}}, \quad (8)$$

where $\langle \dots \rangle_{\Omega_J}$ denotes the averaging over all possible orientations of the vector of angular momentum \mathbf{J} while its length is fixed. In equation (7), Φ is a kinematic factor given by Equation (8); v equals $v_1 + v_2 + 3$ where v_k ($k = 1, 2$) is the dimensionality of the orientational space for the k th reactant; $I_i^{(k)}$ denotes the i th moment of inertia of the k th reactant, and I_i is the i th moment of inertia for the system as a whole; $\mathbf{n}^{(12)}$ is the unit vector directed from the second pivot point to the first one; $\mathbf{d}^{(k)}$ is the vector connecting the center of mass of the k th reactant to its pivot point; $\mathbf{n}_i^{(k)}$ is the unit vector directed along the i th principal axis of the k th reactant; k_a is the number of monatomic reactant, which should be equal to 0 or 1.

Although Equation (8) for the kinematic factor Φ is valid for both linear and nonlinear molecules, it is convenient to rewrite the term for linear molecule in a different form:

$$\mu I^{-1} \sum_{i=1}^2 (\mathbf{n}^{(12)} \times \mathbf{d} \cdot \mathbf{n}_i)^2 = \mu I^{-1} [1 - (\mathbf{n}^{(12)} \cdot \mathbf{n}_0)^2], \quad (9)$$

where \mathbf{n}_0 is the unit vector directed along the axis of the molecule; I is the moment of inertia.

There are three ways to perform variational calculations. They correspond to optimizing the transition state dividing surface using three different ensembles. The resulting rate constants are labeled canonical variational theory (CVT), microcanonical variational theory (μ VT), and energy and total angular momentum resolved microcanonical variational theory (E, J - μ VT), respectively. We now discuss these three choices:

(i) Energy and total angular momentum resolved microcanonical variational theory:

For E, J - μ VT theory, $N(E, J)$ is minimized with respect to s for each E and J . Then the integral in Equation (1) is calculated using the optimized $N(E, J)$.

The minimum of the number of accessible states is obtained by interpolation from a quadratic fit to $N(E, J, s)$ at the three values of s nearest to the minimum. We call this method a 3-point fit; 3-point fits are also used in μ VT and CVT. The integrals over E and J for the optimized $N(E, J)$ in Equation (1) are approximated by Gauss-Laguerre and Simpson's rule quadratures, respectively.

(ii) Microcanonical variational theory:

For microcanonical variational theory (μ VT), the integral over J in Equation (1) is carried out first to get $N(E, s)$. By integrating Equation (7) over J and substituting the result into Equation (2), the number of accessible state $N(E, s)$ is obtained as the following expression:

$$N(E, s) = \mu s^2 \frac{2\pi^{ka-2}}{\Gamma((v+1)/2)} \prod_{k=1}^2 \prod_{i=1}^{v_k} \sqrt{2\pi I_i^{(k)}} \times \langle \Phi \max(E - V(Q), 0)^{(v-1)/2} \rangle_{\Omega} \quad (10)$$

Then $N(E, s)$ is minimized with respect to s for each E , and the optimized $N(E)$ is used for the integral over E .

(ii) Canonical variational theory:

For canonical variational theory (CVT), $N(E, J, s)$ is calculated for all E and J at a single s value. Then the rate constants $k(T, s)$ is minimized with respect to s for each T . The analytic result of the integral in Equation (1) is expressed as:

$$Z(T, s) = \int dE e^{-E/kT} dJ N(E, J, s) \\ = 2\pi^{ka-2} \mu s^2 (kT)^{(v+1)/2} \prod_{k=1}^2 \prod_{i=1}^{v_k} \sqrt{2\pi I_i^{(k)}} \left\langle \Phi e^{-V(Q)/kT} \right\rangle_{\Omega} \quad (11)$$

11.B. Multifaceted dividing surface

When one uses single-faceted dividing surfaces, two pivot points (with one on each of the associating reactants) and a fixed distance between them form a spherical dividing surface. But in many cases the reactants of association reactions have multiple binding sites. Multiple binding sites on one or both reactants require more pivot points to increase the flexibility of the shape of the dividing surface. Let's assume that n_1 pivot points are assigned to reactant 1 and n_2 pivot points are assigned to reactant 2, and the reaction coordinate value is s . Therefore there are $n_1 \times n_2$ pairs of pivot points. Each pair of pivot points corresponds a sphere, and the points on each sphere are selected according to the following condition to form a composite dividing surface:

$$\min_{i=1, j=1}^{n_1, n_2} r_{i,j} = s \quad (12)$$

where $r_{i,j}$ is the distance between the i th and j th pivot points associated with the first and the second reactant, respectively; s is the value of reaction coordinate. The meaning of Equation (12) is that at one particular configuration of the complex at any of the spheres formed by the $n_1 \times n_2$ pairs of pivot points, if the minimum of the distances between pivot points associated with different reactants equal to the value of reaction coordinate, this configuration will be accepted in Monte Carlo sampling. Therefore the composite dividing surface has $n_1 \times n_2$ elementary surfaces; each of the elementary surfaces is called a facet, and the composite surface is called a multifaceted (MF) dividing surface.

In variational calculations, various locations of pivot points should be considered to minimize the reaction rate. Users are responsible to try various sets of locations of pivot points; that is, *POLYRATE* has not automated this. The optimal locations of pivot points are the locations that minimize the reaction rate; but the user must be careful that the MF

dividing surface defines by the chosen locations of the pivot points separates reactants from products.

11.C. Overview of program flow for VRC-VTST calculation

When the VRC method is used for loose transition states by turning on the VRC keyword in the GENERAL section, the program will not use the MEP as the reaction path, but will instead use variable reaction coordinates and either single-faceted or multifaceted dividing surfaces. After the MAIN program reads all the information provided at reactants, subroutine VRCTST is called. Note that user gives the information about two reactants in this case, and the START section only need the ELEC keyword for electronic degeneracy information. If only the forward reaction rate constants for association need to be calculated, the product information is not required. In VRC-VTST calculations, the MAIN program and subroutine VRCTST are parallelized using MPI, and they work independently on different processors to read the same fu5 input file and perform Monte Carlo sampling. During the Monte Carlo sampling, only energies are calculated (for various geometries along the reaction path), and gradients and Hessians are not required. When the Monte Carlo sampling is finished on all processors, one processor (rank 0 in MPI) is used to collect the data calculated on all the processors and to perform the rest of the calculations to get reaction rate constants.

11.D. Implementation details of the E, J -resolved microcanonical VRC algorithm

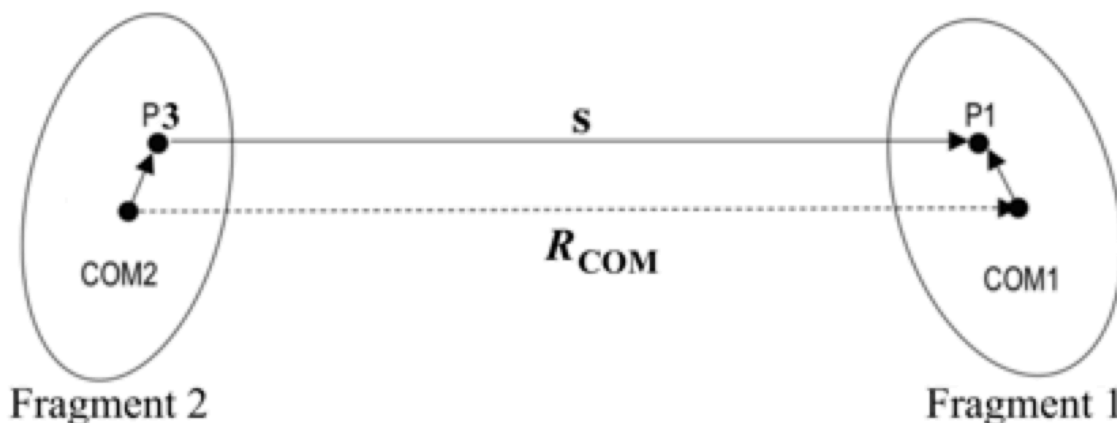
This section contains material from the Supporting Information Appendix of ref. [75] of Sect. 22.

We here describe the implementation details of the E, J -resolved microcanonical VRC algorithm for the case of two pivot points on fragment 1 and two on fragments 2; the generalization to a larger number of pivot points is straightforward. We denote the pivot points on fragment 1 as point 1 and point 2 and the pivot points on fragment 2 as point 3 and point 4. The range of reaction coordinate s searched for a variational transition state is from s_l to s_u (“l” stands for lower edge of the range, and “u” stands for upper edge of the range), and the search for a variational transition state is carried out by varying s with increments of size s_{step} .

We define the vector s as a vector connecting two pivot points, one in each fragment. This is illustrated in Scheme S1, where we are considering a particular pair of pivot points, namely pivot point 1 of fragment 1 and pivot point 3 of fragment 2, so we only show the vector s that connects these two pivot points, and only these two pivot are shown (other pivot points on fragments 1 and 2 are not shown).

In the computer code, we loop over the pairs of pivot points with one on each fragment (1-3, 1-4, 2-3, 2-4), and for each pair of pivot points, the vector s is defined as the vector that connects the pivot point on fragment 1 with the pivot point on fragment 2.

The magnitude of s is denoted as s ; this is the same as the reaction coordinate s mentioned in the first paragraph.



Scheme S1

Step 1: Set the length of the reaction coordinate s to s_1 .

Step 2. Do for m from 1 to 4. For each m , we consider a pair of pivot points with one on each fragment. Index $m = 1$ corresponds to pair 1-3, $m = 2$ represents the pivot points pair 1-4, $m = 3$ represents the pivot points pair 2-3, $m = 4$ represents the pivot points pair 2-4. For our explanation we will use $m = 1$ as an example; in particular, $m = 1$ corresponds to choosing pivot point 1 and pivot point 3. For case $m = 1$ or 2, P1 is pivot point 1, and for $m = 3$ or 4, P1 is pivot point 2. For the case $m = 1$ or 3, P3 is pivot point 3; and for $m = 2$ or 4, P3 is pivot point 4.

Each performance of Step 2C-1 is a Monte Carlo sample. The number of Monte Carlo samples per value of m is denoted as N_{MC} , which is a number specified by the user, and which is the same notation as used in the input and the code. That is, for each m , there are N_{MC} Monte Carlo samples. Because each pair of pivot points determines one facet of the dividing surface, and we have four pairs (i.e., we have a four-faceted dividing surface), the total number of Monte Carlo samples is $4N_{MC}$. In eq 12 of *J. Phys. Chem. A* **2003**, *107*, 9776, which will be used in Step 4, the variable M is equal to $4N_{MC}$.

Step 2A for this pair m : Put fragment 1 in a specific *initial* orientation (this starting orientation is determined randomly) with respect to P1; set the vector s (as shown in Scheme S1) in a specific *initial* direction (this direction is also determined randomly as a start direction) so that P3 position has been decided; and then put the fragment 2 in a specific *initial* orientation (this starting orientation is determined randomly) with respect to P3. Therefore the coordinates of the whole system and other pivot points are determined. Calculate the length of the vector R_{COM} that connects the centers of mass of the fragments.

Step 2B for this pair m : Do for i from $(m-1)N_{MC}+1$ to $(m-1)N_{MC}+N_{MC}$. Starting from the *initial* configuration determined in Step 2A, we now start doing Monte Carlo sampling, i.e., randomly orientating the two fragments, while keeping the length of vector s as specified in Step 1. When orientating the fragments, P1 is *always* taken as the origin of the coordinates system. Then, the position of P3 is determined by a random

solid angle. After reorienting s , also reorient fragment 2 (the solid angles for these reorientations are determined randomly). This now yields configuration i .

Step 2C for this i :

Step 2C-1: Fragment 1 has pivot points 1 and 2, and fragment 2 has pivot points 3 and 4, and the distances between each of the various pairs of pivot points (with one on each fragment) can be computed. Compute the distance r_{ij} (where $i = 1, 2$, and $j = 3, 4$) for each pair of pivot points (1-3, 1-4, 2-3, 2-4) for current Monte Carlo sample configuration. If any of the four r_{ij} is greater than or equal to s (no matter which pair of the pivot points satisfies this condition), then go to Step 2C-2, otherwise abandon this configuration. If this configuration is abandoned, it does not count in M' in eq. 13 of *J. Phys. Chem. A* **2003**, 107, 9776; if this configuration is accepted, then it increases the value of M' by 1.

Step 2C-2: For the specific configuration accepted in Step 2C-1, compute the moment of inertia of each fragment and the unit vector along its principal axis; and calculate the kinematic factor Φ (eq. 2.40 in *J. Chem. Phys.* **2003**, 118, 5442). Next compute number of states $N_q(E, J, s, i)$ for this value of the reaction coordinate s and this specific configuration i , and for each combination of total energy E and angular momentum vector \mathbf{J} (the length of which is denoted as J), averaged over all the possible orientations of vector \mathbf{J} while its length is fixed; this is eq 9 of *J. Phys. Chem. A* **2003**, 107, 9776, and it is computed at using eq 2.39 in *J. Chem. Phys.* **2003**, 118, 5442. Note that we included the correction factor of 2π that is indicated in reference 12 of *J. Phys. Chem. A* **2003**, 107, 9776.

Step 2D: If you have not yet done N_{MC} samples for this m , go back to Step 2B, except start with the present configuration rather than the initial one. If you have completed this m , go to step 2E.

End do loop over index i .

Step 2E: If m is less than 4, increment m by 1 and carry out Steps 2A to 2D for the next pair of pivot points.

End do loop over index m .

Step 3. Now compute the generalized transition state approximation (for a transition state at reaction coordinate value s) of the number of states for the transitional modes at total energy E and total angular momentum J . This is given by

$$N^{GT}(E, J, s) = \frac{1}{M} \sum_{i=1}^{M'} N_q(E, J, s, i) \quad (13)$$

Step 4: If $s < s_u$, increase the length s of vector \mathbf{s} by s_{step} , and go back to Step 1. After all the values of s specified by the user have been considered, go to Step 5.

Step 5: In E, J -resolved microcanonical variational theory, for each E and J , minimize $N(E, J, s)$ with respect to s . This yields

$$N^{\mu VT}(E, J) = N^{GT}(E, J, s_*) \quad (14)$$

where s_* is the value of s that minimizes the number of states. Then plug $N^{\mu VT}(E, J)$ into the rate constant expression and evaluate the integral to obtain the final rate constant $k(T)$.

PART IV DESCRIPTION OF INPUT FILES

12. DESCRIPTION OF INPUT FOR RP-VTST

In the following subsections, a record-by-record description of the data for each input file that is used for RP-VTST calculations is given:

Subsection 12.A: the input data needed from unit fu5;

Subsection 12.B: the input data needed from unit fu29;

Subsection 12.C: the input data needed from unit fu30;

Subsection 12.D: the input data needed from unit fu31;

Subsection 12.E: the input data needed from unit fu40;

Subsection 12.F: the input data needed from unit fu50;

Subsection 12.G: the input data needed from unit fu51.

12.A. General description of file fu5 input

The input for FORTRAN unit fu5 is of the free format keyword style. The fu5 input consists of up to twelve sections. These are defined in sub-section 12.A.1. The input file consists of section headers (section headers are always preceded by a star, *) and associated keywords.

The keywords are of three types: switch, variable, and list. A switch keyword acts as a toggle for an option. A switch keyword, for example SWITCHKEY, usually has an associated form, for example NOSWITCHKEY, that causes the opposite action. A variable keyword always requires an argument. These are usually used for parameters that have to have a value, such as step sizes, convergence criteria, etc. The argument must appear on the same line as the variable keyword. The third type is the list keyword. A list keyword has one or more level-2 keywords associated with it; these keywords may be switch or variable keywords, and the latter kind have arguments. The use of a list keyword takes the form:

```
LEVEL1KEYWORD
  LEVEL2KEYWORDA
  LEVEL2KEYWORDB
  . . .
```

END

where the END line is mandatory.

All switch and variable keywords have default values, and these types of keywords need be given only if the user wishes to override the default. Some, but not all, list keywords have defaults as well. (Additional keywords, called level-2 keywords or subkeywords, are associated with some list keywords — see above. Sometimes, but not always, level-2 keywords have defaults.) List keywords start a subsection that is always terminated with an END line.

Except for the input of the TITLE keyword in the GENERAL section, all input is case insensitive, and anything after a pound sign (#) on a line is assumed to be a comment. Blank lines are ignored. It is highly recommended that the novice user print out one of the test run input files and compare it with the descriptions in the following subsections. A suggested file for this purpose is ch5j2tr3.dat. The rest of subsection 12.A details the keywords for each of the twelve sections. Please note that every keyword must be used in its proper section, i.e., between the occurrence of its starred header and the occurrence of the next starred header (or, for the last section, between the proper starred header and the end of the file).

12.A.1. Section descriptions

The following input sections correspond to the major parts of a RP-VTST rate calculation. Each section begins with a line giving the section name, preceded by an “ * ”. An input file can contain all or some of these sections in any order except for a restart calculation. For a restart calculation, the first section in the file must be the GENERAL section and the REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, START, and PATH sections cannot be used. Keywords for each section are discussed further in the rest of this subsection. For runs using the fu30, fu31, or fu40 electronic structure file input options, one should not include sections SECOND or OPTIMIZATION.

<u>Section Name</u>	<u>Description</u>
*GENERAL	general data such as the title and defining the atoms involved
*ENERGETICS	options for computing potential energy and first derivatives
*SECOND	options for computing second derivatives
*OPTIMIZATION	methods for performing geometry optimizations
*REACT1	definition and properties of the first reactant
*REACT2	definition and properties of the second reactant
*PROD1	definition and properties of the first product
*PROD2	definition and properties of the second product
*WELLR	definition and properties of the reactant side well
*WELLP	definition and properties of the product side well
*START	definition and properties of the saddle pt. or starting geom.
*PATH	options for computing the MEP
*TUNNEL	options for computing the tunneling corrections
*RATE	options for computing the reaction rate

12.A.2. GENERAL section

The GENERAL section is used to give a title that can be used to identify the run, define all the atoms in the reacting system, specify the potential energy surface or electronic structure input option to be used, set up restart criteria (if applicable), and specify whether IVTST-0, IVTST-1, VTST-IOC, or VTST-ISPE is to be used. (IVTST-0 can also be specified on file fu50.) If this is a restart run then the first keyword given in this section must be TITLE. It is good practice to always make this the first keyword so that input files can be easily identified. The table below lists all the valid keywords for the GENERAL section. Indented keywords are level-2 keywords, and the others are level-1 keywords. Level-2 keywords are always associated with a level-1 list keyword and must come after that keyword and before any other level-1 keyword or starred header. In the alphabetical glossary that follows a more complete explanation for each keyword is given. In a standard run this section would have the TITLE and ATOMS keywords. Note that ATOMS is a required keyword.

When using electronic structure input files, IVTST-M is on if and only if one uses file fu31 or if one sets $LOPT(2) = -1$ in file fu30 or if one sets $MAXLPTS = -1$ in file fu40. Otherwise IVTST-M is off, which means that any interpolation *POLYRATE* needs to perform to fill the save grid storage arrays is calculated by Lagrangian interpolation.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
TITLE	list	blank	title with 5 lines maximum
ATOMS	list	required	atoms in system
CHECK	switch	off	check the input data
CLASSVIB	switch	off	classical treatment of vibrations for vibrational partition function
DL	variable	none	interpolated corrections
ICOPT	list	not used	interpolated correction options
BOTH	switch	off	based on corrected and uncorrected data
CORRECT	switch	on	based on corrected data
ENERGY	variable	seckart	Eckart function fit to $V_{MEP}(s)$
FREQ	variable	icl	interpolated correction of frequency
IVTST0	switch	off	zero-order IVTST
IVTST1	switch	off	first-order IVTST
INPUNIT	variable	angstrom	specifies the units in fu5 input file
OUTUNIT	variable	angstrom	specifies the units in fu6 and fu14 output files
MDMOVIE	switch	off	write MDMOVIE input file

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
RESTART	list	not used	restart options
WRITEFU1	switch	off	write restart file on unit 1
READFU1	switch	off	read restart file from unit 1
READFU1&2	switch	off	read restart files from units 1&2
WRITEFU3	switch	off	read from 1&2, write to 3
MERGE	variable	0	an option for merging MEPs
SUPERMOL	switch	on	supermolecule mode for reactants and products
WRITE62	variable	none	write information in the fu62 file
WRITEFU30	switch	off	write an fu30 file
WRITEFU31	switch	off	write an fu31 file

Glossary of GENERAL keywords

ATOMS

ATOMS is a required list keyword that is used to specify all the atoms in the system. For each atom the user must specify n , a sequential number unique to each atom in the molecule, and a label, either the atomic number or symbol of atom n . The mass of the atom in atomic mass units (amu) can also be specified following the label. If it is not included then the mass of the most prominent isotope is used. This program uses the universal atomic mass scale, i.e., the mass of ^{12}C is 12.000000. (For this scale, the correct abbreviation is u, although most people use the old-fashioned abbreviation amu.) The mass is supplied from an internally stored table using the atomic number or symbol as given by the user. Note that the unique number n will be used in other sections to specify the atom. The example below corresponds to monodeuterated water.

Example:

```
ATOMS
  1   O
  2   H   1.0078
  3   H   2.0140
END
```

CHECK

CHECK is a switch keyword that tells the program to stop immediately after checking the input data files. It is especially useful to use this keyword before a long or complicated calculation to make sure that all the data are input correctly. The default is off.

CLASSVIB (NOCLASSVIB)

CLASSVIB is a switch keyword that determines if the vibrational partition functions are to be calculated assuming a classical treatment of the vibrations. The default is that vibrations are treated as quantized. If CLASSVIB is used, all vibrational partition functions are calculated using the classical expression for harmonic oscillators; therefore, no anharmonicity should be specified for any modes.

DL

DL is a variable keyword that determines if the dual-level interpolated corrections are done. (See Subsection 10.H.) The default is *none*. See Section 12.F. This keyword can also be used to perform IVTST-0 calculations with unit fu50; see Section 6.N. The valid options are

<u>Option</u>	<u>Description</u>
<i>ioc</i>	dual-level using fu50 (VTST-IOC)
<i>ispe</i>	dual-level using fu51 (VTST-ISPE)
<i>none</i>	dual-level calculation is not performed (default)

Example:

```
DL   ioc
```

ICOPT

ICOPT is a list keyword that sets options for interpolated optimized corrections. If this keyword is used, then it is assumed that VTST-IOC is desired, and the DL keyword is set to *ioc*. Note that if neither *icr* nor *ica* is specified, the correction defaults to the ICL method, and if *energy deckart* is not specified then the single Eckart function is used. (This differs from versions 7.2 – 7.9.2 and earlier) The ICA, ICL, and ICR schemes, and the two ways to use Eckart functions are discussed in section 10.H. The valid options are given below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>both</i>	separate calculations will be based on the corrected and uncorrected data	off
<i>correct energy</i>	calculation will be based only on the corrected data interpolated optimized corrections for energies: <i>deckart</i> - difference of two Eckart functions fit to the $V_{MEP}(s)$ curve at higher level and Eckart function fit to lower level <i>seckart</i> - do not use Eckart function fit to the $V_{MEP}(s)$ curve at higher level but rather use a single Eckart fit to the difference	on <i>seckart</i>
<i>freq</i>	interpolated optimized corrections for frequencies: <i>ica</i> - interpolated correction based on arithmetic differences <i>icl</i> - interpolated correction based on logarithm of ratios <i>icr</i> - interpolated correction based on ratios <i>none</i> - no corrections	<i>icl</i>

Example:

```
ICOPT
      ENERGY DECKART
      FREQ    ICA
      BOTH
END
```

INPUNIT

INPUNIT is a variable keyword that specifies the units for the geometries and distances given in the fu5 input file. This keyword will have no effect on selecting the units for the geometries and variables for distances in other input files (units fu29, fu30, fu31, fu40, or fu50). The default is *ang*.

<u>Option</u>	<u>Description</u>
<i>ang</i>	angstroms
<i>au</i>	atomic units

Example:

```
INPUNIT    AU
```


IVTST0 IVTST1

IVTST0 and IVTST1 are switch keywords to turn on the zero- and first-order Interpolated VTST (IVTST-0 and IVTST-1) calculations, respectively (See Subsection 6.N.). The default is off.

MDMOVIE

MDMOVIE is a switch keyword that determines if an input file for MDMOVIE should be written out. MDMOVIE is a visualization program that allows animation of motion of the system along the MEP. The default is off.

OUTUNIT

OUTUNIT is a variable keyword that specifies the units for the geometries and distances given in the fu6 and fu14 output file. The default is *ang*.

<u>Option</u>	<u>Description</u>
<i>ang</i>	angstroms
<i>au</i>	atomic units

Example:

```
OUTUNIT    AU
```

RESTART

RESTART is a list keyword that gives the user control over the restart capability of the run or designates the current run as a restart calculation. See Section 7.C. for general comments on this option. If this keyword is not used, a future calculation at a different temperature must start from scratch. Alternatively a user can use the *writeln* option within RESTART and the reaction path will be written to unit fu1. In future runs, the user would just need to specify the *readfu1* option within RESTART for the same reaction path to be read from unit fu1. All the valid options are given below. Note that *merge* requires an argument specifying the increment value. The default is that this keyword is not included, i.e. this is not a restart run and no restart file should be written.

<u>Option</u>	<u>Definition</u>
<i>writeln</i>	write restart information to unit fu1 for future use
<i>readfu1</i>	restart calculation using information on unit fu1
<i>readfu1&2</i>	restart calculation using merged MEP, MEP #1 on unit fu1 is merged with MEP #2 on unit fu2 and the merged file is not saved.
<i>writelnfu3</i>	the merged MEP from <i>readfu1&2</i> is written to unit fu3
<i>merge x</i>	increment in <i>s</i> to be added to the <i>s</i> values of MEP #2 when it is merged with MEP #1. This is required if the READFU1&2 option is used. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.
<i>writelnfu30</i>	write an fu30 file
<i>writelnfu31</i>	write an fu31 file

Example:

```
RESTART
```

```
      READFU1
END
```

NOTE: When writing an fu30 file (*writefu30* option) the user needs to supply an fu30 file with the necessary LOPT(1-40) options.

SUPERMOL (NOSUPERMOL)

SUPERMOL is a switch keyword which indicates that the supermolecule mode is used for reactants and products. The default for SUPERMOL in *POLYRATE* is on. This keyword is used only when the keyword POTENTIAL is set equal to *hooks*, and the choice has no effect if there is only one reactant and one product. The use of supermolecule mode is most easily explained by an example. Consider a bimolecular reaction with reactants A and B. For an analytical potential energy function or a non-size-extensive electronic structure method, one would usually calculate the energy of reactants by placing A at a distance of 20–100 a_0 from B and calculating the energy of the AB supermolecule. However, for size-extensive methods like Hartree-Fock and MP2, one would usually prefer to calculate the energies of A and B separately and add them. The latter requires NOSUPERMOL.

For a bimolecular system, NOSUPERMOL causes *POLYRATE* to calculate the zero of energy as the sum of the potential energies of the two reactants using two separate calculations in which each reactant is treated separately. This is not very useful with an analytical potential energy surface. When NOSUPERMOL is specified, the geometries in REACT1 and REACT2 each contain less than the full number of atoms. In the case of SUPERMOL, the first reactant is optimized and its energy is calculated by having the second reactant frozen at a long distance (which is specified through the definition of the geometries in section REACT2 in file fu5). Then the second reactant is optimized with the first one already at its equilibrium geometry, but far away, and the sum of their energies is obtained as the energy of the supermolecule calculation, i.e., when SUPERMOL is on one does not add the energies from REACT1 and REACT2 at the end of the calculation. The same considerations apply to PROD1 and PROD2 if there are two products.

TITLE

TITLE is a list keyword that allows the user to give a title to the run. The user can specify up to five lines of 80 characters each. Like all list variables, an end must be given at the end of the TITLE subsection. This must be the first keyword list in the GENERAL section for a restart calculation.

Example:

```
      TITLE
      CH3 + H2 -> CH4 + H TEST RUN
      ESD/HARMONIC
END
```

WRITE62

WRITE62 is a variable keyword that determines if a subset of the stationary point information from file fu61 should be printed in fu62 file. The default is *none*. The valid options are

<u>Option</u>	<u>Description</u>
<i>none</i>	no information is printed in fu62
<i>viball</i>	frequencies (in cm^{-1}) are printed for all stationary points
<i>vibsp</i>	frequencies (in cm^{-1}) are printed only for the saddle point

Example:

```
write62  vibsp
```

WRITEFU30

WRITEFU30 is a switch keyword that determines if an output fu30 file should be written out. Notice that the points are not in the required ascending order. Therefore the points in the created fu30 file need to be reordered by the user before using the file. The default is off.

WRITEFU31

WRITEFU31 is a switch keyword that determines if an output fu31 file should be written out. The written file is ready to use with the *unit31* option for the POTENTIAL keyword in the ENERGETICS section of the fu5 input file. The options for writing the fu31 file cannot be selected. It will always be written in atomic units, and the program will always write unscaled gradients and packed Hessians in the fu31 file.

12.A.3. ENERGETICS section

The ENERGETICS section is used to define the method for computing the potential energy of the reacting system and the corresponding first derivatives. The table below lists all the valid keywords for the ENERGETICS section. In the alphabetical glossary that follows a more complete explanation for each keyword is given.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
EZERO	variable	<i>calculate</i>	method for determining zero of E
EZUNIT	variable	<i>a.u.</i>	units of zero of energy
POTENTIAL	variable	<i>hooks</i>	type of PES to be used

Glossary of ENERGETICS keywords

EZERO

EZERO is a variable keyword specifying how the zero of energy should be determined. This keyword is discussed briefly in detail in Section 8.B. *Polyrate* always sets the zero of energy equal to the energy of the reactants, which is defined to be the sum of the energies of both reactants for a bimolecular reaction or to be the energy of the single reactant for a unimolecular reaction. However, the other option is to read in the zero of energy with option *read #*. The default is *calculate*.

<u>Option</u>	<u>Definition</u>
<i>calculate</i>	If <i>calculate</i> is chosen the program will set EZERO to the value that it calculates for the energy of the reactants.
<i>read #</i>	The <i>read #</i> option, where # is the value of the energy of the reactants in units specified by the EZUNIT keyword, causes the program to set EZERO to #.

Example:

```
EZERO read 0.1354
```

Notes:

- (1) If the *read #* option is chosen, the user must ensure that # is the value of the energy of the reactants in units specified by the EZUNIT keyword.
- (2) If the WKB method is used for any of the vibrational modes of reactant 1, then the *read #* option must be used. Otherwise the user can choose *calculate*.
- (3) Another possible use of *read #* would be to allow efficient calculations for cases where the reactant energy is already known and would be expensive to recalculate. However *POLYRATE* always calls at least one of the hook routines for the reactants, so this efficiency must be implemented by the user if desired.

EZUNIT

EZUNIT is a variable keyword specifying the units used for the zero of energy. The default is *a.u.* The other valid option is *kcal*. Note 1 a.u. = 627.5095 kcal/mol. This keyword must be read *before* reading the value of the zero of energy with the *read #* option for the keyword EZERO, if that option is used.

<u>Option</u>	<u>Definition</u>
<i>a.u.</i>	atomic units (hartrees)
<i>kcal</i>	kcal/mol

Example:

```
EZUNIT kcal  
EZERO read 100.1354
```

POTENTIAL

POTENTIAL is a variable keyword that allows the user to specify the source of the potential energy information for a single-level calculation or for the lower level of a dual-level calculation. In the current version of *POLYRATE* there are five valid options: *hooks*, *unit29*, *unit30*, *unit31*, or *unit40*. The option *hooks* means that the user is providing potential information through the hooks. When *Polyrate* is used as a stand-alone package, the hooks call SURF and SETUP; this is how one supplies an analytic potential energy surface (as discussed in Section 8 of this manual). If *unit29* is used the information is read from file fu29 for an IVTST-0 or IVTST-1 calculation. Specification of one of the keywords *unit30*, *unit31*, or *unit40* means that the MEP data is to be read from file fu30, fu31, or fu40, respectively. (as discussed in Sections 9 and 12.C, 12.D, and 12.E of this manual) The fu30 and fu40 options (*unit30* and *unit40*, respectively) are called the electronic structure input file options in older versions of *POLYRATE*.

The default is *hooks*. Derivative versions of *POLYRATE* will also have various options for interfacing with electronic structure packages like ACES II, GAUSSIAN, MOPAC, AMSOL, etc. This may occur through the hooks (modular mode) or in a less general fashion (non-modular mode), but now electronic structure input can be used with any of the five choices of POTENTIAL.

<u>Option</u>	<u>Definition</u>
<i>hooks</i>	use the hooks
<i>unit29</i>	read in from electronic structure file fu29
<i>unit30</i>	read in from electronic structure file fu30
<i>unit31</i>	read in from electronic structure file fu31
<i>unit40</i>	read in from electronic structure file fu40

Example:

```
POTENTIAL unit30
```

12.A.4. SECOND section

The SECOND section is used to define the method for computing the Hessian of the reacting system when the POTENTIAL keyword is set to the *hooks* option; this section should not be included if the option selected for POTENTIAL is *unit30*, *unit31*, or *unit40*. The table below lists all the valid keywords for the SECOND section. In the alphabetical glossary that follows a more complete explanation for each keyword is given.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
HESSCAL	variable	<i>ghook</i>	how to calculate Hessians

Second derivative *POLYRATE* methods:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Other:</i>			
FPRINT	switch	off	print second derivative information
NUMSTEP	variable	1×10^{-4} au	step size for 2nd derivatives of <i>V</i>
NUMTYPE	variable	<i>Central</i>	method for computing 2nd deriv.

Glossary of SECOND keywords

FPRINT (NOFPRINT)

FPRINT is a switch keyword that is used if the normal mode energies, eigenvectors and eigenvalues are to be printed following the normal mode analysis in fu6 output file. Default is NOFPRINT.

HESSCAL

HESSCAL is a variable keyword specifying how to calculate Hessians. The default is *ghook* which means *POLYRATE* will calculate the Hessians by finite differences from the gradients as specified by NUMTYP. Another option is *hhook* which means the Hessians are evaluated by calls to the HHOOK subprogram. See Section 8.C.

<u>Option</u>	<u>Definition</u>
<i>ghook</i>	call GHOOK to get first derivatives, and use central differences to compute the second derivatives of the potential. The step size used is given by the NUMSTEP keyword.
<i>hhook</i>	call HHOOK to get second derivatives.

Example:

```
HESSCAL  ghook
```

NUMSTEP

NUMSTEP is a variable keyword giving the step size in unscaled angstrom (default) or in unscaled bohr (by specifying the keyword INPUNIT AU) for the numerical second derivative of potential energy function with respect to the unscaled atomic coordinates for frequency calculations. The default value is 1.0×10^{-4} bohr.

Example:

```
NUMSTEP  0.00005
```


NUMTYP

NUMTYP is a variable keyword that is used to specify the method for computing the numerical second derivatives of the potential energy with respect to unscaled atomic Cartesians if HESSCAL is *ghook*. See Section 10.C. The default is *central*. The allowed options are:

<u>Option</u>	<u>Definition</u>
<i>central</i>	use central differences of analytic first derivatives to compute the second derivatives of the potential. The step size used is given by the NUMSTEP keyword.
<i>quadratic</i>	use a quadratic fit of the potential to compute the second derivatives of the potential. The step size used is given by the NUMSTEP keyword.
<i>fourth</i>	use a fourth order polynomial fit to the analytic first derivatives to compute the second derivatives of the potential. Step sizes used are NUMSTEP and 2*NUMSTEP.

Example:

NUMTYP *fourth*

12.A.5. OPTIMIZATION section

The OPTIMIZATION section is used to specify the method for carrying out geometry optimizations when the POTENTIAL keyword selection is *hooks*. This section is not needed if POTENTIAL is set to *unit30*, *unit31*, or *unit40*. The table below lists all the valid keywords for the OPTIMIZATION section. In the alphabetical glossary that follows, a more complete explanation for each keyword is given.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
OPTMIN	variable	<i>bfgs</i>	how to optimize minima
OPTTS	variable	<i>nr</i>	how to optimize transition state

Two optimization routines are available in *POLYRATE*. The first is quasi-Newton with BFGS Hessian updates. The alternative is Newton-Raphson with Brent line minimization, which is called the NR method. For minima (equilibrium structures of reactants and products), the default optimizer is BFGS. For saddle points (transition states), the default optimizer is the NR method. See Section 6.E for further details.

POLYRATE optimization methods:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i><u>Common:</u></i>			
PRINT	switch	off	print geometrical parameters and derivatives after each opt step
<i><u>Other:</u></i>			
DLX2	variable	1×10^{-5} au	step size for first derivative of grad in geometry optimization
EFOPT	list	off	indicate the option for EF
GCOMP	variable	1×10^{-5}	maximum allowed component of the gradient for convergence at the minima
TSGCOMP	variable	1×10^{-5}	maximum allowed component of the gradient for convergence at the saddle point
HREC	variable	10	recalculate Hessian every HREC step for convergence at minima
TSHREC	variable	5	recalculate Hessian every TSHREC step for convergence at saddle point
NITER	variable	50	max number of iterations for opt
RETRY	switch	off	switch to full Newton method if BFGS optimization fails
SCALE	variable	0.2	step size scaling factor in Newton search
SDSTART	switch	on (minima opt) off (TS opt)	the first optimization step is steepest-descent
STPTOL	variable	1×10^{-5} au	“optimization failure” parameter

Glossary of OPTIMIZATION keywords

DLX2

DLX2 is a variable keyword used to specify the step size in unscaled angstrom (default) or in unscaled bohr (by specifying the keyword INPUNIT AU) for calculating the first derivatives of the gradients of the potential energy function in geometry optimization. The default value is $1 \times 10^{-5} a_0$.

Example:

```
DLX2                0.000012
```

EFOPT

EFOPT is a list keyword used to specify information for the Eigenvector Following (EF) algorithm. There are five possible variables that may be specified, namely, RMAX, RMIN, OMIN, DDMAX, and DDMAXTS. The RMAX and RMIN variables bracket the ratio of the calculated energy to the predicted energy; the defaults are RMIN = 0 and RMAX = 4. The OMIN variable is used to indicate the minimum of the overlap in the direction which is determined by the dot product between the eigenvector and the previous followed direction; the default is 0.0 for reactants and products and 0.8 for a saddle point. The DDMAX and DDMAXTS variables are used to set the maximum of the trust radius (in angstroms); the DDMAX variable is used for the reactants and products, and the DDMAXTS variable is used for the saddle point. The defaults are DDMAX = 0.5 and DDMAXTS = 0.3.

Example:

```
EFOPT
  RMAX    2.0
  RMIN    1.2
  OMIN    0.2
END
```

GCOMP TSGCOMP

GCOMP and TSGCOMP are variable keywords used to specify the convergence criteria for optimization of minima and transition states, respectively. The optimization is considered converged if the maximum component of the gradient in atomic units is smaller than the value specified. The default value is $1 \times 10^{-5} E_h/a_0$ for both these keywords. Sometimes, if the saddle point region is very flat, 1×10^{-5} is not accurate enough to produce a good reaction path; in such cases we recommend 1×10^{-7} .

Example:

```
GCOMP                0.000001
```

HREC TSHREC

HREC and TSHREC are variable keywords used to specify how often an accurate Hessian is to be computed in a BFGS or Newton-Raphson calculation. See Section 6.E for more

details. Notice that if the number specified is equal to 1 in a Newton-Raphson calculation (*nr* option for keywords OPTMIN or OPTTS), a full Newton method is used. At a step when the Hessian is not recalculated accurately, the Hessian is either estimated using the BFGS formula (*bfgs* option for keywords OPTMIN and OPTTS) or kept frozen (*nr* option for keywords OPTMIN and OPTTS). The defaults are HREC equal to 10 and TSHREC equal to 5.

Example:

HREC 1

NITER

NITER is a variable keyword used to specify the maximum number of iterations to be used in optimizing the geometry. The default value is 50.

Example:

NITER 75

OPTMIN

OPTMIN is a variable keyword that determines how geometries of minima are optimized. The default is *bfgs*.

<u>Option</u>	<u>Definition</u>
<i>bfgs</i>	specifies to use the Broyden-Fletcher-Goldfarb-Shanno Hessian update method. The Hessian is calculated using the option specified in HESSCAL in section SECOND at the initial geometry of the search; at subsequent geometries, the Hessian is estimated using information about the Hessian at the previously geometry and the gradient at the previous and current geometry. (When the search is complete, the Hessian is again calculated by the method specified in HESSCAL, and this Hessian is used for frequencies.)
<i>ef</i>	specifies to use the Eigenvector Following algorithm.
<i>nr</i>	specifies to use the Newton-Raphson algorithm with Brent line minimization.
<i>ohook</i>	optimizes the geometry by calling hooks subprogram OHOOK using options specified in files fu71 through fu74. See Section 8.C.

Example:

OPTMIN *bfgs*

OPTTS

OPTTS is a variable keyword that determines how the geometry of the transition state is optimized. The default is *nr*.

<u>Option</u>	<u>Definition</u>
<i>bfgs</i>	specifies to use the Broyden-Fletcher-Goldfarb-Shanno Hessian update method. The Hessian is calculated using the option specified in HESSCAL in section SECOND at the initial geometry of the search; at subsequent geometries, the Hessian is estimated using information about the Hessian at the previous geometry and the gradient at the previous and current geometry. (When the search is complete, the Hessian is again calculated by the method specified in HESSCAL, and this Hessian is used for frequencies.)
<i>ef</i>	specifies to use the Eigenvector Following algorithm.
<i>nr</i>	specifies to use the Newton-Raphson algorithm with Brent line minimization..
<i>ohook</i>	optimizes TS geometry by calling hook subprogram OHOOK using options specified in file fu75. See Section 8.C.

Example:

```
OPTTS      nr
```

PRINT (NOPRINT)

PRINT is a switch keyword for printing the geometry and the derivatives after each step in the geometry optimization in fu6 output file. Default is off, i.e., NOPRINT.

RETRY (NORETRY)

RETRY is a switch keyword specifying to switch to full Newton search (HREC = 1, no linear search) if the optimization stalls (see STPTOL keyword), instead of quitting. Since this could be quite time consuming, this option has to be used with care. The default is NORETRY.

Note: the difference between full Newton and quasi-Newton is that full Newton calculates a full new Hessian at every step whereas quasi-Newton uses an approximate updated Hessian. In many cases, if the standard search fails it is better to restart the optimization with a different geometry than to use full Newton.

SCALE

SCALE is a variable keyword giving the unitless step size scaling factor in the Newton search for a stationary point. The default value is 0.2.

Example:

```
SCALE      1.1
```

SDSTART (NOSDSTART)

SDSTART is a switch keyword that specifies to start the optimization with a steepest-descent step. This is equivalent to defining the Hessian at the starting point as a unit matrix. This keyword is meaningful only in minima optimizations where BFGS is specified. This option proves useful in speeding up convergence of the optimization when the starting point happens to be outside the quadratic region of the surface around the minimum (the initial guess geometry is bad). The default is SDSTART.

STPTOL

STPTOL is a variable keyword that specify the conditions for optimization failure. The default unit is angstrom and can also be bohr by specifying the keyword INPUNIT AU. If the maximum component of the Newton optimization step is smaller than the value specified by STPTOL, the optimization is considered stalled since the geometry is not changing anymore, and the program quits. This could happen if the (estimated) Hessian used is bad; that in turn probably means that either HREC or SCALE are too large. The default value is $1 \times 10^{-5} a_0$.

Example:

```
STPTOL 0.0001
```

12.A.6. REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, AND START sections

The REACT1, REACT2, PROD1, PROD2, WELLR, WELLP and START sections all have the same set of keywords. A separate section is needed for each reactant and product in the system and for the saddle point or nonstationary starting point. Thus for a system with one reactant, one product, and a saddle point the user needs to include the REACT1, PROD1, and START sections in the input file. For simplicity the keywords have been broken into two categories: basic and anharmonicity. This is only for clarity in the manual. The keywords in these categories can be used in any order, except for the STATUS keyword which must be given as the first option line of a section. Also for simplicity the keywords in each category have been labeled as either required, common, or other. The required keyword (GEOM) must be included. The keywords labeled common are those that might be expected to be found reasonably frequently in input files for typical *POLYRATE* runs. This labeling is only intended as an aid to the novice user. Again an alphabetical glossary of all the keywords follows the two summary tables.

Notes that START refers to the saddle point if there is one and to the starting point if there is no saddle point. Also note that the anharmonicity keywords in the START section control the anharmonicity all along the reaction path since it must be consistent with the START point.

The first restart option (See Section 7.C) does not use any of these sections, instead it is controlled by the RESTART keyword in the GENERAL section. However, the second restart option (See Section 7.C) is controlled by the STATUS keyword of this section. See the glossary for the definition and proper usage of the STATUS keyword.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i><u>Required:</u></i>			
GEOM	list	required	Cartesian coordinates of atoms
<i><u>Common:</u></i>			
CONSTANT	list	not used	coordinates held constant in opt.
ELEC	list	all degen. = 1	electronic degeneracies
ENERGY	variable	none	energy of the stationary points
EIGENVECTOR	list	none	eigenvector of the saddle point
FREQ	switch	on	perform normal-mode analysis
FREQUNIT	variable	waven	units of the frequencies
HESSIAN	list	none	Hessian of the saddle point
INITGEO	variable	geom	location of input data that specifies initial geometries
STATUS	variable	0	status of the stationary point work
SPECIES	variable	nonlinrp	species type (linearity and phase)
VIB	list	none	frequencies of the stationary points
<i><u>Other:</u></i>			
DIATOM	list	do not use Morse function	treat diatomic as a Morse function
RE	variable	none	Morse equilibrium bond length
BETA	variable	none	Morse range parameter
DE	variable	none	Morse dissociation energy
LINAXIS	variable	z-axis	orientation of a linear molecule
PROJECT	switch	on	project out overall trans. and rot.

Anharmonicity:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Common:</i>			
ANTLR	variable	1×10^{-8}	criterion on Morse x_e
DEMIN	variable	0.159	lowest dissociation energy (D)
DQQP	variable	0.0001 au, 0.001 au	two distances for quad.- quar. fit
HARMONIC	switch	on	all modes harmonic
MORSE	switch	off	all modes Morse
MORSEQQ	switch	off	all modes Morse/quadratic-quartic
MORMODEL	variable	<i>morsei</i>	Morse options
PRMODE	switch	off	print XMOL movie clips
QQWKB	switch	off	WKB with a quadratic-quartic fit
QQSEMI	switch	off	semi-classical WKB with a quadratic-quartic fit to potential
TOR	list	not used	information for modes treated as torsions
TOROPT	list	not used	more information for modes treated as torsions
VANHAR	list	not used	each mode is treated individually
VRANGE	list	not used	each mode is treated individually for a given range of s
WKB	list	not used	WKB with true potential
WKBTOL	variable	1×10^{-8}	tolerance for finding eigenvalues
KBQUAD	variable	40	no. of quad. points in phase integrals
KBPRINT	switch	off	print turning points

Glossary of REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and START keywords

ANTLR

ANTLR is a variable keyword whose argument is the criterion on the unitless Morse anharmonicity parameter, x_e , such that the quadratic-quartic model is used if $x_e < \text{ANTLR}$. The *antlr* keyword is only used with the *morseqq* method. The default value is 1×10^{-8} .

Example:

```
ANTLR      0.000013
```

CONSTANT

CONSTANT is a list keyword used to specify which of the Cartesian coordinates should be held constant during the Newton search to find the species' equilibrium geometry. In this list the unique atom number is used along with *x*, *y*, and/or *z* for those coordinates to be held constant. The default is that all coordinates are optimized. In most cases the user needs to specify at least six Cartesian coordinates to be constant for a nonlinear species and five for a linear species. Note that if this keyword is specified, the default is erased, and the user should input all the coordinates which are to be held constant.

Example:

```
CONSTANT
  1      x      y      z
  2      y      z
  3      z
END
```

DEMIN

DEMIN is a variable keyword used to input the lowest reactant or product dissociation energy in a.u., measured from the reactant limit. This is called *D* in published papers. It is needed for any of the Morse anharmonicity options and for μVT calculations. There is a default value of $0.159 E_h$ ($= 100 \text{ kcal/mol}$) for this variable but it is dangerous to use the default. The user is strongly encouraged to understand *D* if Morse anharmonicity is used.

Example:

```
DEMIN      0.1733
```

DIATOM

DIATOM is a list keyword that is used if the particular reactant or product is a diatomic molecule and is to be treated as a Morse oscillator. The default is to treat diatomics by the same methods as any other kind of molecule; thus this keyword would not be used. Note that if the DIATOM option is chosen, then the user must specify all three options, R_e , β , and D_e , which are the Morse parameters for equilibrium geometry, dissociation energy, and range parameter, respectively. β , and D_e are in atomic units and R_e is in angstrom by default and also can be in bohr by specifying the keyword INPUNIT AU. There are no defaults for these options. The example below corresponds to the O_2 molecule.

Example:

```
DIATOM
      RE          2.3158
      BETA        1.4694
      DE          0.1779
END
```

DQQP

DQQP is a variable keyword with two arguments, which are respectively the first and second step sizes used when fitting the quadratic-quartic potential to two points along the normal coordinate. It is in angstrom by default and also can be in bohr by specifying the keyword INPUNIT AU. Note that the first argument (the first step size) must be less than the second. DQQP is to be used only if one of the keywords QQWKB or QQSEMI is specified. Default is $1 \times 10^{-4} a_0$ and $1 \times 10^{-3} a_0$.

Example:

```
DQQP 0.03 1.2
```

EIGENVECTOR

This keyword should be used only in a START section and it should be used if and only if the STATUS option is equal to 6. EIGENVECTOR is a list keyword that is used to specify the eigenvectors of the saddle point in an unscaled Cartesian coordinate system. The eigenvectors of the stationary point should be given in full matrix form with the columns being a particular mode. In addition, the axes must be consistent with the geometry in GEOM.

Example:

```
EIGENVECTOR
  L11  L21  L31  L41  L51  L61
  L12  L22  L32  L42  L52  L62
  L13  L23  L33  L43  L53  L63
  L14  L24  L34  L44  L54  L64
  L15  L25  L35  L45  L55  L65
  L16  L26  L36  L46  L56  L66
END
```

Explanation of example:

L_{ij} is the j th component of the i th eigenvector. Note that $j = 1$ is for the x component of atom 1, $j = 2$ is for the y component of atom 1, $j = 3$ is for the z component of atom 1, $j = 4$ is for the x component of atom 2, etc. Note also that the eigenvalues must be in reverse canonical order (this order is described in Sect. 6.E). Thus, I=1 is for the imaginary-frequency normal mode, then come the eigenvectors corresponding to translations and rotations, and finally come the true vibrations, in order of increasing frequency.

ELEC

ELEC is a list keyword used to specify the degeneracy and energy in atomic units of each electronic state of this species. Up to three states are allowed. The ground state should be given first. The default is for all three states to be singly degenerate. The example given below is for the ³P multiplet of an oxygen atom.

Example:

```
ELEC
  5      0.00000000
  3      0.00072218
  1      0.00103201
END
```

ENERGY

This keyword should be used if and only if STATUS \geq 4. ENERGY is a variable keyword that is used to specify the energies (a.u.) of a reactant, product, or saddle point.

Example:

```
ENERGY    -2.4221970828E-02
```

FREQ (NOFREQ)

The FREQ keyword is used to specify whether normal mode analysis is carried out. The default is FREQ.

FREQUNIT

The FREQUNIT keyword is used to specify the units of the frequencies read by the FREQ keyword. The default is *au*.

<u>Option</u>	<u>Definition</u>
<i>waven</i>	Frequencies are given in cm^{-1} .
<i>au</i>	Frequencies are given in atomic units.

GEOM

GEOM is a list keyword that is required for all calculations except those with the readfu1 or readfu1&2 option selected for RESTART in the GENERAL section. In this list the Cartesian coordinates of all atoms in this particular reactant, product or reaction path species must be given along with each unique atom number. If STATUS (see below) is 0, the user should choose a geometry that is a guess to the species' stationary-point geometry. If status is \geq 2, then the geometry should be the correct geometry, not just a guess. (See the ATOMS keyword in the GENERAL section for the definition of the unique atom number. This number must be consistent for each atom in ATOMS and GEOM.) For a linear species, the user must specify a linear geometry. The example below is for the same deuterated water molecule used in the ATOMS example. It is in angstrom by default and also can be in bohr by specifying the keyword INPUNIT AU (not mass-scaled).

Example:

```
GEOM
  1    0.000000    0.000000  0.000000
  2   -0.241497   -0.945827  0.000000
  3    0.956805    0.000000  0.000000
END
```

HESSIAN

If the option of the STATUS keyword is 4, the Hessian of the saddle point should be read in as a triangular form in unscaled Cartesians. HESSIAN is a list keyword that is used to indicate the Hessian (hartrees per square bohr) of the saddle point. This keyword is only valid for the START section, and can not be used in the REACT1, REACT2, PROD1, and PROD2 sections. In addition, the axes must be consistent with the geometry given in GEOM. The format is free format, with any number of entries per line and entries on the same line separated by one or more spaces. Since the Hessian should be a symmetric matrix we read only the lower triangular part.

Example:

```
HESSIAN
      F11  F21  F22  F31  F32  F33
      F41  F42  F43  F44  F51  F52
      F53  F54  F55  F61  F62  F63
      F64  F65  F66

END
```

Explanation of the example:

The element F_{jj} of the Hessian matrix \mathbf{F} corresponds to the same order of components as explained for j under the EIGENVECTOR keyword.

HARMONIC

HARMONIC is a switch keyword that is used when one wishes to treat all vibrational modes harmonically. It is the default; i.e., if no keyword is given for anharmonicity then the harmonic approximation is used. The default is that harmonic is on, so when it is specified, the purpose is just for clarity (i.e., to make the default explicit).

INITGEO

INITGEO is a variable keyword that is used to indicate the location of the input data that specifies the initial reactant 1, reactant 2, product 1, product 2, and starting geometries. The default is *geom*. The valid options are:

<u>Option</u>	<u>Definition</u>
<i>geom</i>	the initial geometries are taken from the lists following the GEOM keywords in the react1, react2, prod1, prod2, and start sections of the fu5 input file.
<i>hooks</i>	the initial geometries are taken from the external input files fu71-fu78.

Example:

```
INITGEO geom
```

LINAXIS

LINAXIS is a variable keyword that is used to indicate the orientation of a linear reactant, product, or starting molecule. The default is that LINAXIS is off if the initial geometry is nonlinear and that it is on if the initial geometry is linear. If LINAXIS is on, but no option is specified, the default value is *z-axis*. The valid options are:

<u>Option</u>	<u>Definition</u>
<i>x-axis</i>	the linear molecule is aligned with the x-axis
<i>y-axis</i>	the linear molecule is aligned with the y-axis
<i>z-axis</i>	the linear molecule is aligned with the z-axis

MORMODEL

MORMODEL is a variable keyword that is used to specify which Morse model is to be used. This keyword is used only if the MORSE or MORSEQQ switch is on. The default is *morsei*. Note that the value of DEMIN is specified by the user using the keyword DEMIN. All the valid options are given below (see Ref. 1 in Sec. 21 for explanations of these models):

<u>Option</u>	<u>Definition</u>
<i>morsei</i>	Morse I: $V_{\text{MEP}} + \text{local D} = \text{DEMIN}$
<i>morseii</i>	Morse II: local D from V'''
<i>morseiii</i>	Morse III: Morse II if $(V_{\text{MEP}} + \text{local D}) > \text{DEMIN}$; otherwise Morse I
<i>morseia</i>	Morse Ia: like Morse I except use hyperradius to determine side of MEP on which concave-side turning point is to be found if SCT or LCT tunneling correction is done.
<i>morseiia</i>	Morse IIIa: like Morse III except replace Morse I by Morse Ia, i.e., find direction of concave side from hyperradius when using Morse I and find it using 3rd derivative when using Morse II

Example:

```
MORMODEL morseiii
```

MORSE

MORSE is a switch keyword that is used if all vibrational modes are to be treated using the Morse approximation with the same value of MORMODEL. If this option is chosen, then the Morse model is specified by the keyword MORMODEL. Default is off.

MORSEQQ

MORSEQQ is a switch keyword that is used if all vibrational modes are to be treated using Morse/quadratic-quartic approximation with the same value of MORMODEL. This means that the Morse model is used for modes with Morse anharmonicity parameter $x_e \geq \text{ANTLR}$, and the quadratic-quartic option based on second and fourth derivatives of the potential along the generalized normal mode is used for modes with $x_e < \text{ANTLR}$. If this option is chosen, then the Morse model is specified by the keyword MORMODEL. Default is off.

PRMODE

PRMODE is a switch keyword used to print movie clips of the vibrational modes for the XMOL program in XYZ format. The file name of a movie clip consists of two parts. The first part has 4 letters which is used to indicate the type of the species: R1MD for reactant 1, R2MD for reactant 2, P1MD for product 1, P2MD for product 2, RWMD for reactant well, PWMD for product well, or TSMD for the saddle point. And the second part has a three-digit integer which is used to indicate the index of the vibrational mode in decreasing order. Therefore, the maximum number of vibration modes for which this keyword can be used is 999. Currently, each clip has 20 frames.

PROJECT (NOPROJECT)

PROJECT is a switch keyword that tells the program to use projection operators to remove overall translations and rotations from the second derivatives at the stationary points. This projection is not optional along the reaction path, but is optional at stationary points. Even at stationary points, one obtains more stable numerical results if projection is applied. The default is PROJECT.

QQSEMI

QQSEMI is a switch keyword that is used if all vibrational modes are to be treated by using a quadratic-quartic fit to the potential at two distances (specified by the DQQP keyword) along the generalized normal mode with the uniform semiclassical WKB method for computing energy levels. See Section 10.B.1. Default is off.

QQWKB

QQWKB is a switch keyword that is used if all vibrational modes are to be treated using the WKB method with a quadratic-quartic fit to the potential. The fit is done at two distances, specified by the DQQP keyword, along the generalized normal mode with the primitive WKB method for computing energy levels. See Section 10.B.1. Default is off.

SPECIES

SPECIES is a variable keyword that is used to specify the linearity and phase of the reactant, product, or starting species. See Section 6.B for discussion. The default is *nonlinrp*. The valid options are given below.

<u>Option</u>	<u>Definition</u>
<i>atomic</i>	species is a single atom
<i>linrp</i>	species is a linear reactant or product
<i>nonlinrp</i>	species is a nonlinear reactant or product
<i>ssrp</i>	species is a solid-state reactant or product
<i>lints</i>	species is a linear transition state or generalized transition state
<i>nonlints</i>	species is a nonlinear transition state or generalized transition state
<i>ssts</i>	species is a solid-state transition state or generalized transition state

Example:

```
SPECIES      linrp
```

STATUS

STATUS is a variable keyword that is used to indicate the status of the stationary points. When the STATUS keyword is used it must be given as the first option line of the *REACT1, *REACT2, *PROD1, *PROD2, *WELLR, *WELLP, or *START section in which the keyword is used. The options are:

option	*REACT1, *REACT2, *PROD1, *PROD2, *WELLR, or *WELLP	*START
0	calculate geometry, energy, and frequencies	calculate geometry, energy, frequencies, and eigenvectors
2	read in geometry; calculate energy and frequencies	read in geometry; calculate energy, frequencies, and eigenvectors
4	not available	read in geometry, energy and Hessian; calculate frequencies and eigenvectors
6	read in geometry, energy, and frequencies	read in geometry, energy, frequencies, and eigenvectors*

* Note: if *cubic* is chosen as the value of the *FIRSTEP* keyword in the *PATH* section, then option 6 is not valid.

Note that in the *START* section if *STATUS* is 4, then the *ENERGY* and *HESSIAN* lists must be given and if *STATUS* is 6, then the *ENERGY*, *FREQ*, and *EIGENVECTOR* lists must be given. The *GEOM* keyword is always required.

Example:

```
STATUS 2
```

TOR

TOR is a list keyword that is used to specify the information needed for torsional modes that will be treated using the hindered-internal-rotator approximation. Note that the MS-T method is not included in this keyword list, but MS-T can be specified in the *PATH* section using the *SSTOR* keyword. The hindered internal rotator approximation is only used for a subset of the modes, and this keyword is used to give the parameters for each of those modes. The user must also use the *VANHAR* or *VRANGE* (for generalized transition states) keyword to specify which mode or modes are to be treated as hindered internal rotation. There should be one line in the TOR list for every mode treated as a hindered rotor. There are no defaults. If *VANHAR* or *VRANGE* specifies that a mode is to be treated as a hindered rotor, the line for that mode must be present in the TOR list.

The *SRPG* is the recommended method when using the TOR keyword. Other methods may be appropriate for certain systems, but careful analysis should accompany any other choice. See Ref. 52 and 62 of Sect. 23 for a study and comparison of torsion methods.

The input required for each hindered rotator mode is:

MODE P M METHOD N
 where

MODE = mode number m
 P = number P of distinct minima along the torsional coordinate
 M = number M of total minima along the torsional coordinate ($M \geq P$)
 METHOD = one of six methods defined in Section 10.3:
 CT Chuang-Truhlar interpolatory method
 RPG Torsional Pitzer-Gwinn method
 SRPG Segmented Torsional Pitzer-Gwinn method
 AS Ayala-Schlegel method
 SAS Segmented Ayala-Schlegel method
 FR Free Rotor
 N = number N of atoms in first subgroup

The user needs to be cautious with the definition of number of distinct minima along the torsional coordinate. For example, H_2O_2 has two minima that have the same energy and frequencies, but they are mirror images of one another, and one cannot be superimposed to the other. Therefore they are distinguishable. The total symmetry number is the ratio of M/P , which should not be a non-integer number. Here are a few examples for the values of P and M .

<u>Molecule</u>	<u>P</u>	<u>M</u>
$\text{CH}_3\text{-CH}_3$	1	3
$\text{ClCH}_2\text{-CH}_3$	1	3
HO-OH	2	2
$\text{ClCH}_2\text{-CH}_2\text{Cl}$	3	3
$\text{ClCH}_2\text{-CH}_2$	3	6

Among the six methods for torsional treatment, SRPG is recommended. Note each method requires a different set of keywords. For more input information see the TOROPT section below.

Example:

```
TOR
      10 1      3      SRPG 4
END
```

TOROPT

TOROPT is a list keyword that must be given for modes treated as hindered rotors. Each record of TOROPT is a single line of the form:

MODE DATATYPE list of values

An END line signifies the end of all records in the list.

Which quantities may appear in the list of values depends on DATATYPE. The allowed values for DATATYPE and the associated lists are as follows:

DATATYPE list

	ISB	i_1, i_2, \dots, i_N (the indices of the atoms in the first subgroup)
	NBOND	indices of the two atoms that define the rotation axis (used only when the curvilinear option is used to calculate the moment(s) of inertia)
	CTSHEME	OW/RO/CO/RW/CW/RWO/CWO (one of seven schemes defined in Ref. 52 of Section 23 for CT method)
method)	CTLEVEL	FULL/SF/SC (one of three levels for CT method)
	OMEGA	$\omega_2, \dots, \omega_P$ (the frequencies in cm^{-1})
	U	U_2, U_3, \dots, U_P (excitation energies in cm^{-1})
	I	I_2, I_3, \dots, I_P (moments of inertia in m.a.^2)
	READI	I_1 (moment of inertia in m.a.^2)
	W	W_1, W_2, \dots, W_P (barrier heights in cm^{-1})
height from	WL	WL_1, WL_2, \dots, WL_P (WL_j is the barrier height from the minimum j to the maximum on its left in cm^{-1})
height from	WR	WR_1, WR_2, \dots, WR_P (WR_j is the barrier height from the minimum j to the maximum on its right in cm^{-1})
	RATIOL	RL_1, RL_2, \dots, RL_P (RL_j is the torsional angle difference between the minimum j and the maximum on its left in degrees)
angle	RATIOR	RR_1, RR_2, \dots, RR_P (RR_j is the torsional angle difference between the minimum j and the maximum on its right in degree)

The methods specified in the TOR section need different DATATYPE values and the associated lists. Note that ISB and NBOND are always required except when CTSHEME = OW or READI is used. An explanation of DATATYPE requirement for each method is given below

1. The RPG and AS methods require the following DATATYPE values:

ISB, NBOND, I, WL, and WR

In these methods the moment of inertia of the torsional motion I_1 is calculated by the curvilinear model using the input geometry in fu5 file, and the others are given by the user. The user can run jobs to obtain I_2, \dots, I_P using the corresponding geometries. The barrier height used in the RPG and AS methods is taken as the average of the input WL and WR values. The frequency of the lowest minimum is used.

2. The SRPG and SAS methods require the following DATATYPE values:

ISB, NBOND, I, U, WL, WR, RATIOL, and RATIOR

The SPRG and SAS methods need the same information as their corresponding non-segmented counterparts except that U, RATIOI and RATIOOR are also required.

3. The FR method require the following DATATYPE values:

ISB, NBOND, and I

4. The CT method requires the following DATATYPE values:

ISB, NBOND, CTScheme, CTLEVEL, OMEGA, U, and I.

CTScheme list is one of the seven schemes defined in Ref. 52 of Section 23:

OW	Scheme ωW
RO	Scheme $R\omega$
CO	Scheme $C\omega$
RW	Scheme RW
CW	Scheme CW
RWO	Calculate I with the R-scheme, read in W , obtain ω from normal mode or generalized normal-mode analysis
CWO	Calculate I with the C-scheme, read in W , obtain ω from normal mode or generalized normal-mode analysis (default)

CTLEVEL list is one of the three approximation levels used for CT method defined in Ref. 52 of Sect. 23:

FULL	full (default)
SF	single-frequency
SFE	single-frequency and single-energy

Note that the SFE method treats the P minima using the same I , ω , and W as the input geometry in fu5 file and do count the contributions of the P minima to the partition function. The single-conformer (SC) method defined in Ref. 52 of Sect. 23 is a special case of SFE method when P equals 1.

In the seven distinct schemes, R represents using the rectilinear model to calculate the reduced moments of inertia of the torsional motion (see refs. 24, 29, and 30 in Sect. 23 for detailed description of the rectilinear model), and C represents using the curvilinear model to calculate the reduced moments of inertia of the torsional motion (Ref. 59 in Sect. 22 gives a detailed description of the curvilinear model). Compared with the rectilinear model, for which one only needs to indicate which atoms are in each of the two fragments of the hindered rotation, the curvilinear model requires this information and also a choice of internal rotation axis (by using the NBOND keyword) and an assignment (by using the ISB keyword) of the lighter fragment as the rotational top. The first five schemes use the relationship given by Eq. (12) of Ref. 52 in Sect. 23 to calculate either I , W , or ω from the other two. The RWO and CWO schemes require I , W , and ω , and use these variables in the equations of Ref. 52 in Sect. 23 without forcing a relationship between them.

If CTScheme = CO or $C\omega$, then the sum of the masses of the atoms in ISB must be less than or equal to the sum of the masses of the atoms not in ISB. Note that the

values of P and N that appear in these definitions are specified using the TOR keyword. In the current version of *POLYRATE*, the values of OMEGA, U, and W are assumed to be independent of the value of the reaction coordinate s . That is, only values for the stationary points can be read in, and for the non-stationary points along the reaction path, the only hindered rotor parameters that can depend on s are the value of ω_1 , which is obtained directly from generalized normal mode analysis if the OW, RO, or CO scheme is used, and from the moments of inertia of the lowest minimum along the torsion mode, which are computed from the geometry if the CO or CW scheme is used and from the geometry and normal mode eigenvector if the RO or RW schemes is used.

The set of data types expected in the TOROPT list for CT method depend on the choices of CTScheme and CTLEVEL and the number of distinguishable minima P . In particular, the expected records are:

<u>Choices in</u>			<u>DATATYPES expected in TOROPT</u>
<u>P</u>	<u>CTLEVEL</u>	<u>CTSCHEME</u>	Other expected DATATYPES
1	any	OW	W (W_1 only)
1	any	RO	ISB
1	any	CO	ISB, NBOND
1	any	RW	ISB, NBOND, W (W_1 only)
1	any	CW	ISB, NBOND, W (W_1 only)
>1	FULL	OW	U, OMEGA, W
>1	FULL	RO	ISB, OMEGA, U, I,
>1	FULL	CO	ISB, NBOND, OMEGA, U, I,
>1	FULL	RW	ISB, U, I, W
>1	FULL	CW	ISB, NBOND, U, I, W
>1	SF	OW	U, W
>1	SF	RO	ISB, U, I
>1	SF	CO	ISB, NBOND, U, I
>1	SF	RW	ISB, U, I, W (W_1 only)
>1	SF	CW	ISB, NBOND, U, I, W (W_1 only)
>1	SC	OW	W (W_1 only)
>1	SC	RO	ISB
>1	SC	CO	ISB, NBOND
>1	SC	RW	ISB, W (W_1 only)
>1	SC	CW	ISB, NBOND, W (W_1 only)

If values are not expected but are given anyway, then these values will be ignored. The program will not check the correspondence of the TOR and TOROPT keyword options; it is users' responsibility to check if the calculation is meaningful. The defaults are ω_j = frequency obtained from normal mode analysis at point on reaction path ($j = 1, 2, \dots, P$), $W_j = 1000 \text{ cm}^{-1}$ ($j = 1, 2, \dots, P$), $U_j = 0$, ISBN = {1 2 3 4}, and NBOND = {4 5}.

The READI option is used to read in the moment of inertia for the initial configuration.

Example for using CT method with OW and FULL options:

```
TOR
      17 2 2 CT 4
END
TOROPT
      17 CTLEVEL FULL
      17 CTScheme OW
      17 U 1000.
      17 ISB 1 2 3 4
      17 OMEGA 110.
END
```

Example for using CT method with CW and SC options:

```
TOR
      18 1 1 CT 3
END
TOROPT
      18 CTLEVEL SC
      18 CTScheme CW
      18 ISB 3 4 5
      18 NBOND 2 6
      18 W 1450.0
END
```

Example for using SRPG method:

```
TOR
      6 2 2 SRPG 2
END
TOROPT
      6 ISB 1 4
      6 NBOND 1 2
      6 WL 2545.505 376.905
      6 WR 376.905 2545.505
      6 RATIOL 112.504 67.496
      6 RATOR 67.496 112.504
```

VANHAR

VANHAR is a list keyword that is used to specify variable anharmonicity, i.e., that each vibrational mode is to be treated differently. The default are that all modes be treated harmonically. This keyword is used to specify those modes that are to be treated anharmonically. The valid choices are *harmonic*, *morse*, *morseqq*, *qqwkb*, *qqsemi*, and *tor*. (Note that modes for which the harmonic option is wanted need not be included in the list since it is the default.) Note that for each of these anharmonicity options there may be other variables that have to be set, and in each case there is keyword with the same name as the choices just given, i.e., *harmonic*, *morse*, *morseqq*, *qqwkb*,

qqsemi, and tor. In specifying the modes, note that they are always numbered in order of decreasing frequency, regardless of symmetry, and that degenerate modes get two numbers.

The input required is:

```
mode          anharmonicity method
```

Example:

```
VANHAR
      3      tor
      5      qqsemi
END
```

VIB

If the option of the STATUS keyword is 6, frequencies of the stationary point should be read. VIB is a list keyword that is used to specify the frequencies (a.u. or cm^{-1} as determined by FREQUIT) of a reactant, product, or saddle point. The number of frequencies to be read is always F , where F is $3N - 6$ for nonlinear stationary points, $3N - 5$ for linear stationary points, and $3N$ for solid-state stationary points. For reactants and products there should be F real frequencies, and they should be given in decreasing order of their magnitude. For saddle points, the $F - 1$ real frequencies should be given first in decreasing order followed by the imaginary frequency given last as a negative number.

Example:

```
VIB
 1.72881D-02 1.72881D-02 1.62501D-02 8.24925D-03
 8.07930D-03 8.07884D-03 7.06232D-03 7.06187D-03
 6.58527D-03 3.28102D-03 3.28102D-03 -1.24729D-02
END
```

VRANGE

VRANGE is a list keyword similar to VANHAR except that a range of s values can also be specified for each mode. This means that except for the given modes and their respective ranges of s , each mode will be treated harmonically. Again the valid anharmonicity choices are: *harmonic*, *morse*, *morseqq*, *qqwk*, *qqsemi*, and *tor*. Note that for each of these anharmonicity options there may be other variables that have to be set. For more information, please see the keyword glossary entry for each method. This keyword should only be used in the START section. The default unit of s is angstrom and can also be bohr by specifying the keyword INPUNIT AU.

The input required is:

```
smin          smax          mode number          anharmonicity
```

The example below shows the VRANGE input if one wanted to use the Morse I approximation in the regions $0.01 \leq s \leq 0.5$ and $-0.15 \leq s \leq -0.01$ for mode #2 and the harmonic approximation everywhere else.

Example:

```
VRANGE
      -0.15      -0.01      2      morsei
      0.01       0.50      2      morsei
END
```

WKB

WKB is a list keyword that is used if the zero point eigenvalues are to be computed using the WKB method with the true potential. See Sections 10.B and 12.B.2. The WKB keyword applies only to the ground state; another anharmonicity method must be chosen for the excited states, or else the excited states will be calculated harmonically. The options in this list control the WKB calculation. They are defined as follows:

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>wkbtol</i>	tolerance for finding WKB eigenvalues in hartrees	1×10^{-8}
<i>kbquad</i>	number of quadrature points used in calculating WKB phase integrals	40
<i>kbprint</i>	switch for printing values of turning points	off

Example:

```
WKB
      wkbtol      0.0000002
      kbquad      30
      kbprint
END
```

12.A.7. PATH section

The PATH section contains the keywords that control the calculation of the reaction path and the properties along the path. This section is not used in a restart calculation. For simplicity the keywords have been broken into five categories: basic, integrators, step sizes for the MEP, extrapolation, and print. This is only for clarity in the manual. The keywords in any of these five categories can be used in any order. Also for simplicity the keywords in each category have been labeled as either common or other. The keywords labeled common are those that might be expected to be found reasonably frequently in input files for typical *POLYRATE* runs. This labeling is only intended as an aid to the novice user. Again an alphabetical glossary of all the keywords follows the five summary tables.

If only a conventional transition state theory rate constant calculation is desired, one needs to include this section only if the Hessians are input in mass-scaled coordinates and SCALEMASS does not equal 1.0 amu.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i><u>Common:</u></i>			
CARTRP	Switch	on	Whether the normal mode analysis for reactants and products is carried out in Cartesian coordinate
COORD	Variable	<i>cart</i>	type of coordinate system to be used for computing vibrational frequencies and generalized normal modes along the MEP
IDIRECT	Variable	<i>1</i>	initial direction off of saddle pt.
INTDEF	List	not used	define curvilinear internal coordinates
NSTEPS	Variable	<i>99999</i>	max no. of steps along MEP
SADDLE	Switch	yes	whether the reaction has a saddle point
SCALEMASS	Variable	<i>1.0</i>	mass-scaled coordinates scale factor
SIGN	Variable	<i>reactant</i>	direction of unbound eigenvector
SRANGE	List	not used	range of <i>s</i> for MEP
SLP	Variable	<i>+1.0 au</i>	upper limit on <i>s</i> before it is extended by interpolation
SLM	Variable	<i>-1.0 au</i>	lower limit on <i>s</i> before it is extended by interpolation
<i><u>Others:</u></i>			
CURV	Variable	<i>dgrad</i>	method for computing curvature
DLX3	Variable	<i>1 × 10⁻⁴ au</i>	step size for 3 rd derivatives of PES
FCSCALE	List	not used	the force constant scaling factors for specified internal coordinate
FREQINCR	Variable	not used	Increase low frequencies to a certain number that is as the value of this keyword
FIRSTSTEP	Variable	<i>nmode</i>	method for first step

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Others:</i>			
FREQSCALE	Variable	1	Scale factor for frequencies
HESS	Switch	on	Compute the Hessian along MEP
INTMU	Variable	1	Number of points on each side of the saddle point for interpolation of $\mu_{\text{eff}}(s)$
ISWR	Variable	Not used	mode index of the reactant frequency which is to be scaled
ISWP	Variable	not used	mode index of the product frequency which is to be scaled
IVTSTMOPT	List	not used	IVTST-M options
LBEXP	Variable	3	mapping for curvature components
POLYS	switch	on	include 10 extra points for spline fitting
SINCW	variable	0.01 au	step size for 10 extra points
TENSION	variable	-1.0	tension for spline under tension
IVTST0FREQ	list	not used	modifies the frequency of a specified mode along the reaction path
NEXTPT	switch	off	only compute next point on MEP
REORDER	switch	off	order frequencies along MEP by overlaps
SCALERP	switch	off	scales frequencies in VIB section if turned on
SCLPT	switch	off	whether scale the potential if FCSCLAE is used
SPECSTOP	List	not used	special stop for MEP
CURVE	Variable	vag	the curve on which the check for the

				special stop is made
POINT	Variable	<i>savegrid</i>		the grid on which the end point for the special stop is chosen
PERCENTDOWN	Variable	<i>50.</i>		the percentage down from the saddle point for the special stop
SSPECIAL	list	<i>no special s val.</i>		special <i>s</i> values for N.M. analysis
SYMMETRY	switch	<i>off</i>		symmetric reaction path
VSCALE	variable	<i>1</i>		scale factor for MEP

Integrators:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<u>Common:</u>			
ES1OPT	list	not used	Euler with stabilization
DELTA2	variable	5×10^{-5} au	step size δ_2 for stabilization
DIFFD	variable	1×10^{-8}	magnitude of difference vector
RPM	variable	ESD	reaction path following method
<u>Others:</u>			
RODS	switch	off	re-orient the dividing surface

Step sizes for the MEP:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<u>Common:</u>			
SSTEP	variable	5×10^{-4} au	step size for computing MEP
<u>Others:</u>			
INH	variable	9	ratio of gradients to Hessians
INI	variable	1	interval of gradient interpolation step
SFIRST	list	not used	special step size for first n steps
NFSTEP	variable	0	number of initial special steps
FSIZE	variable	1×10^{-5} au	step size for the special steps

Reaction coordinate for the VRC:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Common:</i>			
SVRC	list	not used	Reaction coordinate setting for reaction path in variable reaction coordinate algorithm for loose transition state

Extrapolation:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Others:</i>			
EXFIRST	list	not used	extrapolate first direction from saddle point.
EXNSTEP	variable	0	number of steps
EXREACT	switch	off	extrapolation is towards reactants
EXPROD	switch	on	extrapolation is towards products
EXSTEP	variable	0.03 au	step size
EXALPHA	variable	1.0	extrapolation range parameter
EXSECOND	list	not used	extrapolate second direction from saddle point
EXNSTEP	variable	0	number of steps
EXREACT	switch	on	extrapolation is towards reactants
EXPROD	switch	off	extrapolation is towards products
EXSTEP	variable	0.03 au	step size
EXALPHA	variable	1.0	extrapolation range parameter

Note: The extrapolation is not performed unless either or both the EXFIRST and EXSECOND keywords are specified.

Print:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Others:</i>			
POTINF	variable	0	print n pieces of PES information
PRPATH	list	not used	print reaction-path information to units fu25-fu27
PRINTSTEP	switch	off	print geom. at each save point
PRDISTMX	variable	20	print interatomic distance matrices
PRSAVERP	variable	9999	print geometry, V , and V' at each save point
PRSAVEMODE	variable	9999	print normal mode information at each save point
SDEBG1	variable	not used	s value for starting debug printing of generalized normal mode vibrational analysis along the MEP
SDEBG2	variable	not used	s value for ending debug printing of generalized normal mode vibrational analysis along the MEP
SPRNT	variable	0	extra reaction-path Hamiltonian information printed at large s
SSPECPR	switch	off	print information at special s values

Nonequilibrium Solvation:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
BATH	list	not used	nonequilibrium solvation
BATHTEMP	variable	298.0	temperature of the solvent in K
CUPLCNST	list	not used	coupling constants in mass-scaled atomic units (A)
DIFFUSION	list	not used	diffusion coefficients in $10^{-5} \text{ cm}^2 \text{ s}^{-1}$
FRICITION	variable	0	solvation time in fs

Torsional Anharmonicity

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
SSTOR	list	not used	MS-T torsional anharmonicity along the reaction path by carrying out a single-structure (SS) calculation as described in Section 10.O.
MTOR	variable	<i>no default</i>	local periodicities of torsions

Glossary of PATH keywords

BATH

BATH is a list keyword specifying the parameters used for a nonequilibrium solvation calculation. For detailed description, see Ref. 53 in Sect. 23. There are 4 options:

<u>Keyword</u>	<u>Description</u>
BATHTEMP	temperature of the solvent in K
CUPLCNST	atomic coupling constants of the solvent coordinate to the solute in mass-scaled atomic units
DIFFUSION	effective atomic diffusion constants in $10^{-5} \text{ cm}^2 \text{ s}^{-1}$
FRICITION	solvent friction time in fs (Note: the friction time is also called the solvation time)

Example:

```
BATH
  FRICITION 10.0      # in fs
  DIFFUSION   # in 10-5 cm2s-1
    1  8.08
    2 12.62
    3  1.72
    4  3.04
    5 33.08
    6 192.27
    7  50.52
  END
  BATHTEMP 298.0
END
```

CARTRP (NOCARTRP)

The keyword CARTRP is used to declare explicitly that the reactant and product normal mode analysis is carried out in Cartesian coordinates. The purpose of adding this keyword is to allow the use of curvilinear coordinate normal mode analysis. In versions of *POLYRATE* earlier the 8.6, the normal mode analysis for reactants and products was always carried out in Cartesian coordinates even though the normal mode analysis for transition states can be carried out in curvilinear internal coordinates. The default is CARTRP.

Note: CARTRP should not be used when scaling factors for principal force constants in curvilinear internal coordinates (FCSCALE) are used.

COORD

COORD is a variable keyword specifying the type of coordinate system for calculating the bound-state vibrational frequencies and normal mode eigenvectors along the MEP. It is very important to use curvilinear coordinates rather than Cartesian coordinates for generalized normal mode analysis along MEP in VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results. See Sections 6.G.1 and 10.M. The default is *cart* (Cartesian coordinates). The valid options are:

<u>Option</u>	<u>Definition</u>
<i>curv1</i>	use curvilinear internal coordinates for generalized normal mode analyses using method of Ref. 41 of Sect. 23
<i>curv2</i>	use curvilinear internal coordinates for generalized normal mode analyses using method of Ref. 43 of Sect. 23
<i>curv3</i>	use redundant internal coordinates for generalized normal mode analyses using method of Ref. 46 of Sect. 23
<i>cart</i>	use rectilinear, i.e., Cartesian coordinates for generalized normal mode analyses

Note that if the *curv1*, *curv2*, or *curv3* option is used, the INTDEF keyword discussed below is required.

Example:
 COORD *curv2*

CURV

CURV is a variable keyword used to specify the method for computing the reaction path curvature components. See Sections 6.H and 10.F for more information about this issue. The default is *dgrad*. The valid options are:

<u>Option</u>	<u>Definition</u>
<i>dgrad</i>	use a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate
<i>oneside</i>	use one-sided differences for the derivative of the gradient with respect to the reaction coordinate
<i>dhess</i>	use Hessian, generalized-normal mode eigenvectors, and gradient to calculate the reaction path curvature

Example:
 CURV *dgrad*

DLX3

DLX3 is a variable keyword for specifying the step size used for numerical differentiation in unscaled atomic Cartesians to obtain the 3rd derivatives of *V* from the Hessian at the saddle point. These derivatives are required when the cubic starting algorithm is employed (see FIRSTSTEP). The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default is $1 \times 10^{-4} a_0$.

Example:
 DLX3 0.00005

ES1OPT

ES1OPT is a list keyword for using the Euler with stabilization integrator. The user must give the step size for the finite-difference derivative (DELTA2 keyword) and the tolerance parameter for the magnitude (in a.u.) of the difference vector between successive normalized gradients (DIFFD keyword). See Section 10.D. The default unit of step size DELTA2 is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The defaults are $DELTA2 = 5 \times 10^{-5} a_0$ and $DIFFD = 1 \times 10^{-8}$.

Example:

```
ES1OPT
  DELTA2    0.00001
  DIFFD     0.000001
END
```

EXFIRST

EXFIRST is a list keyword for specifying the extrapolation beyond the limits (SLM and SLP) of the Hessian grid for the first direction from the saddle point when using the hooks. (See Section 7.B for a discussion of which direction is first.) The EXFIRST and EXSECOND keywords cannot be used with units fu30, fu31, or fu40. The extrapolation procedure is different for reactions with and without a saddle point. The particular extrapolation procedure to be used is chosen based on the SADDLE(NOSADDLE) keyword in the PATH section.

For reactions without a saddle point (NOSADDLE specified in the PATH section) the extrapolation procedure is described in Section 10.A; it is designed to extrapolate $V_a G(s)$ and $\mu_{\text{eff}}(s)$ toward bimolecular reactants or products and should not be used to extrapolate toward unimolecular reactants or products. If EXFIRST is omitted, no extrapolation is performed in the first direction, that is, by default no extrapolation is performed. If no extrapolation is performed, V_{MEP} is set equal to the reactant or product value beyond the limits, and μ_{eff} is set to μ .

For reactions with a saddle point (SADDLE is specified in the PATH section; or SADDLE is accepted as the default) the extrapolation procedure is based on the mapped interpolation algorithm, which is described in Sections 6.N.3 and 10.J and in Ref. 48 of Sect. 23. The mapped interpolation algorithm is used to extrapolate V_{MEP} , $|I|$, ω , and B_{mF} . If EXFIRST is omitted, no extrapolation is performed in the first direction, that is by default no extrapolation is performed.

The EXALPHA keyword is only used for reactions without a saddle point.

If the EXFIRST keyword is not specified, then no extrapolation is performed. If the EXFIRST keyword is specified with an empty list then the extrapolation is performed with the default values listed below. The keywords in this list and their respective default values are explained below.

<u>Keyword</u>	<u>Description</u>	<u>Default</u>
EXREACT	extrapolation is towards reactants	off
EXPROD	extrapolation is towards products	on
EXNSTEP	number of steps to be taken in extrapolation region	0
EXSTEP	mass-scaled step size in the extrapolation region for computing the adiabatic potential and other quantities needed for the tunneling calculations. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.	0.03 a ₀
EXALPHA	exponential range parameter α in reciprocal angstroms (only used with NOSADDLE). It can also be in reciprocal bohrs by the keyword INPUNIT AU.	1.0 a ₀ ⁻¹

Example:

```
EXFIRST
  EXPROD
  EXNSTEP    50
  EXSTEP     0.1
  EXALPHA    0.5
END
```

Example (empty list, defaults only):

```
EXFIRST
END
```


EXSECOND

EXSECOND is a list keyword that is very similar to EXFIRST but is used for specifying the extrapolation beyond the limits (SLM and SLP) of the Hessian grid for the second direction from the saddle point when using the hooks. (See Section 7.B for a discussion of which direction is first.) The EXFIRST and EXSECOND keyword can not be used with units fu30, fu31, or fu40. See also the discussion of EXFIRST. The EXALPHA keyword is only used when NOSADDLE is specified in the PATH section. If the EXSECOND keyword is not specified then no extrapolation is performed. If the EXSECOND keyword is specified with an empty list then the extrapolation is performed with the default values listed below. The valid keyword options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
EXREACT	extrapolation is towards reactants	on
EXPROD	extrapolation is towards products	off
EXNSTEP	number of steps to be taken in extrapolation region	0
EXSTEP	mass-scaled step size in the extrapolation region for computing the adiabatic potential and other quantities needed for the tunneling calculations The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.	0.03 a ₀
EXALPHA	exponential range parameter α in reciprocal angstroms. It can also be in reciprocal bohrs by the keyword INPUNIT AU.	1.0 a ₀ ⁻¹

Example:

```
EXSECOND
  EXREACT
  EXNSTEP    50
  EXSTEP     0.1
  EXALPHA    0.5
END
```

Example (empty list, defaults only):

```
EXSECOND
END
```

FCSCALE

FCSCALE is a list keyword used to specify scaling factors for principal force constants of the selected curvilinear internal coordinates defined in the INTDEF keyword. This keyword should be used only when internal coordinates defined by COORD *curv2* or COORD *curv3* options are used, and the switch keyword FREQ is on. To specify the selected curvilinear internal coordinates involving force constant scaling, the connectivities of atoms are required to be defined using the INTDEF keyword as well as the FCSCALE keyword is used. Each item in the list is in the form of an integer sequence number followed by a floating-point scaling factor. The sequence number denotes the ordinal number of the specific coordinate in the list of all the curvilinear internal coordinates defined under the INTDEF keyword. (Note: This is not the ordinal number in internal storage after the code internally re-orders the coordinates.) The corresponding diagonal element in the force constant matrix will be scaled by the scaling factor. All scaling factors are set to 1.0000 as the default. In the following example, the sequence number 8 in FCSCALE represents the stretch between atom 1 and atom 2; thus the diagonal

element in the force constant matrix involving that stretch will be scaled by a scaling factor of 0.9734.

Example:

```
INTDEF
  1=2=3
  2-3-4 2-3-5 2-3-6 4-3-5 4-3-6 5-3-6
  1-2 2-3 3-4 3-5 3-6
  4-3-5-6
END

FCSCALE
  8 0.9734
END
```

When FCSCALE is used, NOCARTRP should be declared explicitly because the reactant and product normal mode analysis should be carried out in curvilinear coordinates.

The actual effect of the force constant scaling is controlled by the keyword SCLPT, in which either the scaling of the potential or the scaling of frequencies is specified. The default is scaling the frequencies only (NOSCLPT).

Note: Another different way to scale frequencies is with FREQSCALE, in which all frequencies are scaled by the same scaling factors.

FIRSTSTEP

FIRSTSTEP is a variable keyword specifying the method for taking the first step off the saddle point or starting geometry. The default is *nmode*. The valid arguments are:

<u>Option</u>	<u>Definition</u>
<i>nmode</i>	take the first step along the imaginary-frequency normal mode
<i>cubic</i>	use the cubic starting algorithm (see STEP 3)
<i>gradi</i>	take the first step along the negative of the gradient direction [use this only if NOSADDLE is selected; see Section 6.G]

Example:

```
FIRSTSTEP  cubic
```

FREQSCALE

FREQSCALE is a variable keyword for specifying the factor by which to scale the harmonic vibrational frequencies for partition function calculations. If FREQSCALE is used, the scaling will be applied to all stationary point frequencies and generalized normal mode frequencies for generalized transition states along the reaction path. Furthermore, the vibrationally adiabatic potential energy curve V_a^G will be calculated by adding the scaled ZPE to V_{MEP} and all tunneling calculations will be based on this new V_a^G . The default is not to scale the frequencies. For kinetics, the scale factor that one should choose is the one for reproducing the correct ZPE, i.e., the ZPE scale factor, which is available on the website: <https://comp.chem.umn.edu/freqscale/version3b2.htm>.

Please notice that, if the vibrational frequencies are directly read in from the .dat input file, the FREQSCALE will *not* be applied on these user-input vibrational frequencies; and therefore, usually, the user-input frequencies in .dat file should have already been scaled by the user with the same frequency scale factor for the reaction path. This feature thus also provides the user with an opportunity to use different scale factors for stationary points and for the saddle point and the non-stationary points on the reaction path.

Example:
FREQSCALE 0.91

FREQINCR

FREQINCR is a variable keyword for specifying the value to which the frequencies that are lower than this value to increase (unit is cm^{-1})

Example:
FREQINCR 200

HESS (NOHESS)

HESS is a switch keyword that determines if the Hessian should be computed along the MEP. This switch is defaulted to on and in most cases one would want to compute the Hessian. Most likely this switch will only be used for a special case when the Hessian is not desired, and thus NOHESS will be used. The default is HESS.

IDIRECT

IDIRECT is a variable keyword used to specify the direction of the initial step from the starting point. The allowed arguments for this variable are ± 1 : +1 if the initial direction is the direction toward products and -1 if it is towards reactants. See Section 7.B. The default is 1.

Example:
IDIRECT 1

INH

INH is a variable keyword used to specify the ratio of the number of gradients to the number of Hessians calculated along the MEP. INH must be an integer value. The default is 9.

Example:
INH 4

INI

INI is a variable keyword used to specify the interval of the gradient interpolation steps in the IVTST-M calculation. INI must be an integer value less than or equal to INH. The default is 1.

Example:

```
INI 2
```

INTDEF

INTDEF is a list keyword used to specify the curvilinear internal coordinates used for the vibrational frequencies and calculations of generalized normal modes. (It is very important to use curvilinear coordinates rather than Cartesian coordinates for VTST calculations; Cartesian coordinate will very likely yield imaginary frequencies along the reaction coordinate, which will cause unphysical VTST results.) The allowed types are

- stretches ($i-j$),
- bends ($i-j-k$),
- dihedrals ($i-j-k-l$), and
- doubly degenerate linear bends ($i=j=k$)

where i, j, k , and l are the atom-number-labels defined in the GENERAL section.

Note: when the bond angle is close to 180 degrees (i.e., greater than about 175 degrees), using the doubly degenerate linear bending mode approximation will prevent dividing by zero or a number close to zero during computations.

Example:

```
INTDEF
      1-2 1-3 3-4 2-1-3 1-3-4 2-1-3-4
END
```

Note: The pound sign (#) cannot be used as a comment card within the INTDEF keyword list.

INTMU

INTMU is a variable keyword that indicates the number of points on each side of the saddle point used for the interpolation of the effective reduced mass at the saddle point. See Section 6.H. The default value is 1. The valid options are:

<u>Option</u>	<u>Definition</u>
1	1 point on either side of the saddle point; thus interpolation is based on 2 points
3	3 points on either side of the saddle point; thus interpolation is based on 6 points

Example:

```
INTMU 1
```

ISWP

ISWP is a variable keyword for specifying the index of the product frequency that will be scaled by a factor of $(VSCALE)^{1/2}$. This keyword is only needed if the VSCALE keyword is specified and there is only one product species.

Example:

```
ISWP          9
```

ISWR

ISWR is a variable keyword for specifying the index of the reactant frequency that will be scaled by a factor of $(VSCALE)^{1/2}$. This keyword is only needed if the VSCALE keyword is specified and there is only one reactant species.

Example:

```
ISWR          9
```

IVTSTMOPT

IVTSTMOPT is a list keyword for specifying the IVTST-M options for the curvature components and unimolecular reactions.

The mapping procedure for the curvature components is slightly different from the mapping for the rest of the parameters. The exponent in equation 19 of Ref. 48 in Sect. 23 has been optimized in order to make the changes of the interpolated curvature components along the reaction path as close as possible to the changes observed in converged calculations. Therefore, the user is strongly advised against changing this parameter without a good reason. *If* one does have a good reason, though, the exponent can be set by means of the LBEXP variable option. This keyword can only take positive odd integer values.

There are two methods for performing IVTST-M calculations for the unimolecular side (or sides) of reactions with one reactant or/and one product, or wells in one of the channels: (i) to carry out a spline-under-tension fit (as a function of z) based only on the input data, or (ii) to carry out a spline-under-tension fit based on ten extra energy points. The ten extra points are calculated by means of a cubic polynomial fitted to the unimolecular reactant, unimolecular product, or well, a point calculated from the lowest real frequency at that minimum, and the two points on the reaction path with the largest $|s|$ values on the unimolecular side. In the second option generating the extra point from the frequency requires the choice of a small step-size, whose value is recommended to be the same as the gradient step-size used in the calculation. The *polys* option specifies the treatment to be given to the unimolecular side (or sides), and, the *sincw* option specifies the value of the step-size. The default is to use a cubic polynomial for estimating 10 extra energies, with a step-size along the lowest frequency mode of $0.01 a_0$ in s , which is mass scaled.

<u>Option</u>	<u>Definition</u>
<i>lbexp</i>	value of the exponent for interpolating the curvature components. The default is 3.
<i>polys</i>	use a cubic polynomial for estimating 10 energy points before spline-fitting $V_{\text{MEP}}(s)$.
<i>polyz</i>	use a cubic polynomial for estimating 10 energy points before spline-fitting $V_{\text{MEP}}(z)$ (default).
<i>nopolys</i>	do not include 10 extra points, and base the spline fit on input data only plus one extra computed as described in Section 10.J.
<i>sincw</i>	variable keyword to indicate the step-size along the lowest-frequency mode to fit the cubic polynomial. The default unit is mass-scaled angstroms and it can also be in mass-scaled bohr by specifying the keyword INPUNIT AU. The default value is $0.01 a_0$. this value is used only if <i>polys</i> or <i>polyz</i> is selected.
<i>fixmuef</i>	keyword used to apply a global interpolation to estimate the effective reduced mass μ_{eff} of the SCT algorithm at the saddle point when the IVTST-M algorithm is used with hooks. The default option is to use local Lagrangian interpolation for the effective reduced mass at the saddle point with either one point or three points (see INTMU keyword option) one each side of the saddle point. The global interpolation can be used to obtain a smoother value of μ_{eff} at the saddle point; it should be used if and only if the points close to the saddle point suffer from numerical instability.

tension keyword used to specifically give a tension factor in interpolation using spline under tension. If this keyword not used, the program will automatically determine the tension factors. The value for this keyword must be between 0 an 85. We recommend to use this keyword only in case very large tension factor is needed.

Example:

```
IVTSTMOPT
    lbexp    5
    polys
    sincw    0.005
END
```

IVTST0FREQ

IVTST0FREQ is a list keyword that modifies the frequency of a particular mode along the reaction path. A section for each mode begins with the list option keyword MODE which contains lines with the following keyword options:

<u>Option</u>	<u>Definition</u>
<i>index</i>	index of the vibrational mode to be interpolated (this is determined from the frequencies of the saddle point arranged in decreasing order).
<i>reactant x</i>	used to indicate the frequency (in cm^{-1}) at the reactant side ($s < 0$) for the interpolation with value x . When the reactant is unimolecular or a reactant well present, the s value has to be provided with option SR. If a reactant well exists for a bimolecular reaction, the interpolation uses the frequency at the well instead of the reactants.
<i>start x</i>	used to indicate the frequency at $s = 0$ for the interpolation.
<i>product x</i>	used to indicate the frequency at the product side ($s > 0$) for the interpolation with value x . When the product is unimolecular or a product well present, the s value has to be provided with option SP. If product well exists for a bimolecular reaction, the interpolation uses the frequency at the well in stead of the products.
<i>sr x</i>	used to indicate the s value at the limit of reactant side ($s < 0$). The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default value is -1.0 bohr.
<i>sp x</i>	used to indicate the s value at the limit of product side ($s < 0$). The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default value is $+1.0$ bohr.

The IVTST0FREQ algorithm is similar to the frequency interpolation algorithm of the VTST-IC method. See Section 10.M for a detailed description.

Notice that IVTST0FREQ can be used to interpolate a given mode, say i , (where modes are numbered in order of decreasing frequency) from the saddle point to reactants and products. It cannot interpolate between mode i at the saddle point and mode $i + 1$ at reactants (which might sometimes be useful for unimolecular reactions). It cannot interpolate between mode i at the saddle point and mode i at an intermediate location between the saddle point and reactants.

Note that IVTST0FREQ is currently incompatible with using fu31.

Example:

For a bimolecular/bimolecular case

```
IVTST0FREQ
  MODE
    index      11
    reactant   0
    start      300
    product    0
```



```

        END
    END
    For a unimolecular/bimolecular case
    IVTST0FREQ
        MODE
            index      11
            reactant   50
            start      200
            product    0
            sr         -2.0
        END
    END

```

NEXTPT (NONEXTPT)

NEXTPT is a switch keyword that only works if POTENTIAL = *unit30* or *unit 40*. When turned on it will cause *POLYRATE* to compute only the next point on the MEP by taking a single step from the last point in the electronic structure input file, and then stop. The method used for the step will be ESD (default), or Page-McIver (if PAGEM is specified). The step size used in this procedure is specified using the SSTEP keyword. The direction in which one is following the path (towards reactants or products) is specified using the IDIRECT keyword in conjunction with NEXTPT (IDIRECT = 1 to calculate the next point to the right; IDIRECT = -1 to calculate the next point to the left). For example, using the ESD method, turning on NEXTPT causes the following calculation to be performed

$$\mathbf{x}_{\text{next-point}} = \mathbf{x}_{\text{current}} - \delta s \hat{g}_{\text{current}}$$

$$s_{\text{next-point}} = s_{\text{current}} + (\text{IDIRECT}) \delta s$$

where $\mathbf{x}_{\text{current}}$ is the mass-scaled geometry corresponding to the last point input on fu30 or fu40, and \hat{g}_{current} is a unit vector in the direction of the mass-scaled gradient at that point. Thus, to go left, the last point read in should have negative s , and to go right, it should have positive s .

This option is useful when using *POLYRATE* as a driver to determine geometries for ab initio calculations. The only print option in unit fu30 compatible with the use of NEXTPT is LOPT(1) = 0. The default is NONEXTPT.

NSTEPS

NSTEPS is a variable keyword for specifying the maximum number of steps (after the first step) along the negative of the gradient of V to be taken in each direction from the saddle point or starting full-system geometry. The minimum number of steps is the number required to reach the limits specified by SRANGE. The default is 99999.

Example:

```

    NSTEPS      5000

```

POTINF

POTINF is a variable keyword that specifies how many pieces of data from the potential energy subroutine are written along the MEP to FORTRAN unit fu6. The default is 0. See Subsection 8.C.1 for details.

Example:

```
POTINF      4
```

PRDISTMX

PRDISTMX is a variable keyword that indicates the interval along the Hessian grid for the printing of interatomic distance matrices. The i - j element of the interatomic distance matrix is the distance (in mass-scaled bohr) between atoms i and j . The default is set to print the interatomic distance matrix at every 20th point along the Hessian grid.

Example:

```
PRDISTMX    20
```

PRINTSTEP (NOPRINTSTEP)

PRINTSTEP is a switch keyword that specifies that the geometry at each step along the MEP be printed in fu6 output file. The default is NOPRINTSTEP. See the SPRNT keyword below.

PRPATH

PRPATH is a list keyword used to set the options for printing reaction-path information to units fu25-fu27. This keyword is especially useful for analyzing reaction-path data. The unit fu25 and fu26 output can easily be imported to a spreadsheet, and unit fu27 output is designed to be read into the XMOL program to visualize the geometry changes along the reaction path. (The XMOL program is a X-windows-based molecule viewer and format converter developed by Wasikowski and Klemm at the Minnesota Supercomputing Center, Inc.) When this keyword is specified, the program will automatically write the energetic information (V_{MEP} and V_a^{G}) to unit fu25. The valid options are defined below.

<u>Option</u>		<u>Description</u>	<u>Default</u>
<i>coord</i>	a_i	up to 4 atoms, from the full system, can be specified for which the bond distances(2, 3, or 4 atoms specified), the bond angles(3 or 4 atoms specified), and the dihedral angle(4 atoms specified) are calculated and written to unit fu28.	off
<i>interval</i>	n	write the reaction-path information every n Hessian grid points.	1
<i>freq</i>	m_i	GTS frequency mode numbers to be printed to unit fu26; a total of up to ten modes can be specified and printed.	not used
<i>xmol</i>		write XMOL XYZ input file to unit fu27.	off

Example:

```
PRPATH
  COORD      1  2  3  6
  INTERVAL   5
```

```

          1  2  3
        FREQ
        XMOL
      END

```

PRSAVEMODE

PRSAVEMODE is a variable keyword that specifies that (in addition to the items listed for PRSAVERP) the normal mode eigenvectors, energies, curvatures, and frequencies be printed at a given multiple of the save point. The default is 9999.

PRSAVERP

PRSAVERP is a variable keyword that specifies that the geometry, potential (V), curvatures, and gradient (V') at a given multiple of the save point along the MEP be printed. The default is 9999.

REORDER (NOREORDER)

REORDER is a switch keyword that specifies that the frequencies be ordered according to overlap calculations along the MEP. The default is NOREORDER, which means the frequencies are in canonical order. See Sections 6.E, 6.F, and 6.G for further discussion.

RODS (NORODS)

RODS is a switch keyword which specifies that the RODS algorithm is to be used to re-orient the generalized transition state theory dividing surface. The RODS algorithm can be used with any of the integrators, ESD, ES1, or PAGEM. See Sect. 10.D. The default is NORODS, which turns RODS off. When RODS is off, the generalized-transition-state-theory dividing surface is normal to the gradient.

RPM

RPM is a variable keyword that indicates which integrator is to be used for following the reaction path. The default is *esd*. The valid options are defined below.

<u>Option</u>	<u>Description</u>
<i>esd</i>	use the Euler steepest-descents integrator to follow the reaction path; this is the name we now use for the Euler algorithm, which is often called the Euler single-step algorithm in textbooks
<i>es1</i>	use the Euler-with-stabilization integrator to follow the reaction path
<i>pagem</i>	use the Page-McIver integrator to follow the reaction path
<i>vrpe</i>	use the VRP(ESD) algorithm to follow the reaction path

Example:

```
RPM    pagem
```

SADDLE (NOSADDLE)

SADDLE is a switch keyword to indicate whether the reaction has a saddle point or not. See Sect. 6.G. The default value is SADDLE.

SCALEMASS

SCALEMASS is a variable keyword for the reduced mass μ in amu used in defining the mass-scaled coordinates. This reduced mass is called the scaling mass. See Refs. 1 and 5 of Sect. 23 for an explanation of the scaling mass. Notice that no physical observable depends on μ , but a single value must be used consistently in the whole calculation. Note also that certain nonobservable intermediate quantities, most notably the numerical value of s at any point along the MEP, *do* depend on μ . The default is 1.0 amu which means that the mass-scaled coordinates in bohrs have the same numerical values as mass-weighted coordinates in amu^{1/2} bohrs. (These are the mass-weighted coordinates made popular by Wilson, Decius, and Cross in Ref. [36] of Sect. 22.) The variable μ is discussed in Section 6.A of this manual.

Example:

```
SCALEMASS 0.195
```

SCALERP

SCALERP is a switch keyword for scaling previously calculated frequencies that appear in the VIB section of input. This keyword is off by default. Frequencies that have already been calculated with the FREQSCALE option are scaled, and therefore this option would not be required. Frequencies that have been calculated without scaling, and then used with the FREQSCALE option would require this keyword in order for the program to scale them correctly.

SCLPT (NOSCLPT)

SCLPT is a switch keyword to specify the behavior of the force constant scaling option. (Note: Details of the force constant scaling option can be found in the description of the keyword FCSCALE).

If the keyword SCLPT is turned on along with the keyword FCSCALE, the potential surface is actually scaled. The scaled force constant matrix will be used to obtain both the frequencies and the normal mode eigenvectors in the normal mode analysis. Thus the MEP is altered when FCSCALE is used with those force-constant involved reaction-path following algorithms, such as VRP(ESD), Page-McIver, etc. Also, the first step on the reaction path will be changed as well if the normal mode eigenvector with the imaginary frequency is used to determine the direction of the first step.

However, if this switch is turned off using NOSCLPT, the force constant scaling is actually just a way to change some of the frequencies. After frequencies are obtained from the scaled force constant matrix, the un-scaled force constants are recovered. The normal mode eigenvectors all come from the un-scaled force constants. Users should be aware that the MEP is not altered using this strategy. The only quantity changed is the frequencies. Those scaled frequencies will be used in the calculations of ZPE, vibrational partition functions, and tunneling probabilities.

SDEBG1 SDEBG2

SDEBG1 and SDEBG2 are variable keywords used to specify the s limits of the interval on the MEP at which to have debug information of the vibrational frequencies and generalized normal-mode calculations output to unit fu6. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. (Note: the value of SDEBG1 has to be smaller than SDEBG2).

Example:

```
SDEBG1 0.499
SDEBG2 0.501
```

SFIRST

SFIRST is a list keyword that allows a special step size for the first n steps. The user must give the number of initial steps to be taken in each direction from the initial geometry, including the initial step (NFSTEP) and the step size for each of these steps (FSIZE). The default unit of FSIZE is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The defaults are NFSTEP = 0 and FSIZE = 1×10^{-5} a₀.

Example:

```
SFIRST
      NFSTEP      10
      FSIZE       0.0001
END
```

SIGN

SIGN is a variable keyword used to ensure the conventional definition of the sign of s , $s < 0$ for the reactant side and $s > 0$ for the product side, is followed. The allowed arguments for this variable are *reactant* and *product*: *product* if the unbound eigenvector at the saddle point points toward the product side and *reactant* if the eigenvector points toward the reactant side. The SIGN keyword is discussed further in Section 7.B. The default is *reactant*.

Example:

```
SIGN product
```

SPECSTOP

SPECSTOP is a list keyword that may be used to specify the range over which the MEP is to be computed. This keyword can function in conjunction with the SRANGE keyword that is presented below. The SPECSTOP keyword may be used to limit the range over which the MEP is calculated to a smaller range than that specified by the SRANGE keyword. The MEP calculation will be stopped when the value of the potential energy (V_{MEP}) or the value of the vibrationally adiabatic ground-state potential energy (V_{a}^{G}) becomes smaller than a critical value. This critical value is defined based on the values of potential energy or zero-point-inclusive potential energy at the saddle point and reactants or products. For example, if the V_{a}^{G} curve is chosen, the MEP will be determined until the $V_{\text{a}}^{\text{G}}(s)$ becomes smaller than the value calculated as: $[(1-x) V_{\text{a}}^{\text{G}}(s=0) + x V_{\text{+ZPE}}(\text{R or P})]$ where $V_{\text{+ZPE}}(\text{R or P})$ is the higher of the zero-point-inclusive potential energy for the reactants and products, and x is equal to $X/100$ (X is the argument of PERCENTDOWN subkeyword). Note that $V_{\text{+ZPE}}(\text{R or P})$ is equivalent to $V_{\text{a}}^{\text{G}}(\text{R or P})$ for a bimolecular reaction but not for an unimolecular reaction for which $V_{\text{a}}^{\text{G}}(\text{R or P})$ does not include the zero-point-energy contribution of the reaction coordinate motion. Similarly, if the V_{MEP} curve is chosen, the MEP will be calculated until the $V_{\text{MEP}}(s)$ becomes smaller than the value calculated as: $[(1-x) V_{\text{MEP}}(s=0) + x V_{\text{MEP}}(\text{R or P})]$ where $V_{\text{MEP}}(\text{R or P})$ is the higher of the zero-point-exclusive potential energy for the reactants and products, and x is again equal to $X/100$ (X is the argument of PERCENTDOWN subkeyword). Finally, the MEP limits can be chosen either on the fine grid (also called gradient grid) or the save grid. In the current version of the code, this keyword cannot be used if the calculation requires the use of a restart file (fu1, fu2, or fu3) or the use of an electronic structure input file (fu29, fu30, fu31, or fu40). The valid keyword options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default*</u>
CURVE	the curve on which comparison is made (the options are <i>vmep</i> and <i>vag</i>)	<i>vag</i>
POINT	the grid on which the end point is chosen (the options are <i>savegrid</i> and <i>finegrid</i>)	<i>savegrid</i>
PERCENTDOWN	percent down from the saddle point	50.

Example:

```
SPECSTOP
      CURVE          vmep
      POINT          finegrid
      PERCENTDOWN    63.4
END
```

Example (empty list, defaults only):

```
SPECSTOP
END
```

* These are the defaults if SPECSTOP is specified. If it is not specified, then PERCENTDOWN is effectively infinity, i.e., only SRANGE is used to stop the reaction path calculation.

SPRNT

SPRNT is a variable keyword that, if PRINTSTEP is on, requires that $|s| > \text{SPRNT}$ for any reaction-path information to be printed. Default is 0. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.

Example:

```
SPRNT 0.1
```

SRANGE

SRANGE is a list keyword needed to specify the limits on the reaction coordinate, s for which the MEP is to be computed. These limits, SLM and SLP, are the ranges over which the MEP is computed, i.e., $\text{SLM} \leq s \leq \text{SLP}$. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default values are $\text{SLM} = -1.0 a_0$ and $\text{SLP} = +1.0 a_0$.

Example:

```
SRANGE
  SLP 1.50
  SLM -1.75
END
```

SSPECIAL

SSPECIAL is a list keyword used to give all the special s values at which normal mode analyses are to be performed. These values of s will be included in the Hessian grid. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default is to not include any special s values. See keyword SSPECPR below.

Example:

```
SSPECIAL
  0.14
  0.34
 -0.76
END
```

SSPECPR

SSPECPR is a switch keyword that works in conjunction with SSPECIAL. It turns on the printing of generalized normal mode vibrational energies, frequencies, and eigenvectors for the special s values regardless of whatever other print keywords are specified. The default is off.

SSTOR

SSTOR is a list keyword that specifies the parameters used for the torsional-potential anharmonicity calculations along reaction path by the SS-T method. Note that when only a single structure among all conformational structures is considered, the MS-T method is reduced to the SS-T method as described in Section 10.O. The SSTOR keyword also turns on the SS-T calculations along reaction path. The valid keyword options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default*</u>
MTOR	local periodicities of torsions, i.e. $M_{j,\tau}$ values	<i>no default</i>

Example:

```
SSTOR
      MTOR      3 3 3 3
END
```

SSTEP

SSTEP is a variable keyword that specifies the step size along the mass-scaled MEP for following the gradient of V , except that the first n steps can be different (see SFIRST). The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The default is $5 \times 10^{-4} a_0$.

Example:

```
SSTEP 0.0001
```

Note that SSTEP is ignored if POTENTIAL = *unit30*, *unit31*, or *unit40* is specified in the GENERAL Section. See Section 9.C.

SYMMETRY (NOSYMMETRY)

SYMMETRY is a switch keyword that specifies that the reaction path is symmetric. When this is turned on only one direction of the MEP from the saddle point is calculated, and the MEP information from the other direction is obtained by reflection. Default is NOSYMMETRY.

VSCALE

VSCALE is a variable keyword for specifying the factor by which to scale the potential energy along the MEP. If VSCALE is 0 or 1, no scaling is done. The default is 1.

Example:

```
VSCALE 1
```


12.A.8. TUNNEL section

The TUNNEL section contains the keywords that control the tunneling calculations. This section is not needed if tunneling calculations are not desired. If this section is not included then all defaults are ignored, and the default SCT, ZCT, and Wigner calculations are not performed. For simplicity the keywords have been broken into two categories: basic and large-curvature. This is only for clarity in the manual. The keywords in any of these categories can be used in any order. Also for simplicity the keywords in each category have been labeled as either *common* or *other*. The keywords labeled *common* are those that might be expected to be found reasonably frequently in input files for typical *POLYRATE* runs. This labeling is only intended as an aid to the novice user. Again an alphabetical glossary of all the keywords follows the two summary tables.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Common:</i>			
SCT	switch	on	small-curvature tunneling
ZCT	switch	on	zero-curvature tunneling
<i>Others:</i>			
EMIN	variable	0	cutoff E for computing percent tunneling
QRST	list	not used	quantized-reactant-state tunneling
HARMONIC	switch	on	harmonic approximation
HARMWKB	switch	off	harmonic-WKB approximation
SYMWKB	switch	off	WKB and symmetric V_a^G
MODE	variable	required	number of the selected vibrational reaction-path mode
SRW	variable	required	s value of the reactant
STATES	variable	all	number of states included in calculation
QUAD	list	no changes	change quadrature parameters
NSEGBOLTZ	variable	1	segments for Boltzmann integral
NSEGTHETA	variable	1	segments for theta integral
NQE	variable	40	quadrature points for Boltzmann integral
NQTH	variable	40	quadrature points for theta integral
SCTOPT	list	<i>spline</i>	small-curvature tunneling options
LAGRANGE	variable	4	Lagrange interpolation for μ_{eff}
SPLINE	switch	on	spline fit for interpolating μ_{eff}
WIGNER	switch	on	Wigner tunneling

Large-curvature tunneling:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Common:</i>			
ALLEXCIT	switch	on	tunneling into all excited states
EXCITED	variable	0.0	tunneling into n th excited state
LCT	switch	off	large-curvature tunneling
LCTDETAIL	list	not used	print extra details in LCT calculations
<i>Others</i>			
LCTOPT	list	not used	large-curvature tunneling options
ILCT	switch	off	interpolated LCG calculation
INTERPOLATE	variable	4	order of interpolation (see glossary)
LCGMETHOD	variable	4	method for LCT calculations
LCTRST	switch	off	restart a previous LCT calculation
LCTSTR	switch	on	store information from a LCT calculation for a possible restart
NGAMP	variable	120	quadrature points for tunneling amplitude
NGTHETA	variable	120	quad. pts. for theta integral
NILCGPT	variable	9	number of interpolated points at each tunneling energy
PRFREQ	switch	off	print information (see glossary)
PRPROB	switch	off	print uniform probabilities
RTPJAC	switch	off	print information in pseudo-Jacobian coordinates
VADAVG	switch	off	average the overlapping of two adiabatic potentials
VEFSRCH	variable	999	search for possible nonadiabatic LCG4 regions

Glossary of TUNNEL keywords

ALLEXCIT (NOALLEXCIT)

ALLEXCIT is a switch keyword that specifies that the large-curvature tunneling calculations (see LCT) are to be performed with tunneling into all accessible excited states in the LCT exit arrangement. Default is on. The opposite switch, NOALLEXCIT, specifies that tunneling calculations be performed with tunneling into only the ground state.

Note that the maximum number of excited states is set by the MAXPS parameter in the aamod.f90 file (see Subsection 5.A.2).

EMIN

EMIN is a variable keyword whose argument specifies the energy (in kcal/mol) that is the cutoff point in computing the percent tunneling that comes from below E_{\min} . The default is 0.0 .

Example:

```
EMIN 48.35
```

EXCITED

EXCITED is a variable keyword whose argument is the state n . The LCT tunneling calculation is performed with tunneling into up to the state with quantum number n , where $n = 0, 1, \dots$. Default is 0. The example below is for the case where tunneling will be performed for the first excited state in the LCT exit arrangement (if accessible) as well as for the ground state. Note that this keyword shouldn't be used together with the ALLEXCIT or NOEXCIT keyword.

Example:

```
EXCITED 1
```

LCT3 or LCT4

One of these two keywords is compulsory if large curvature calculations are required. The default is off. If LCT3 (the old LCT) or LCT4 (see Section 10.4. for details) are on, then the COMP and μ OMT results (described in Section 6.I) are always calculated. Of the two options (LCT3 and LCT4), LCT4 is recommended; LCT4, being the final version, is usually just denoted LCT.

LCTDETAIL

LCTDETAIL is a list keyword that prints extra details of the LCT calculations to Fortran unit fu41 to fu47 files. The valid subkeyword options are defined below. See Section 14.

<u>Option</u>	<u>Description</u>
<i>state n</i>	<i>n</i> is the quantum number of the vibrational state (0 is the ground state) for which information is to be printed.
<i>interval m</i>	<i>m</i> is the number of intervals of the energy grid the user wants to print out for this particular quantum state; LB and UB are the indices of the lower and upper bounds of the energy grid interval, respectively.
<i>lb₁ ub₁</i>	
<i>lb₂ ub₂</i>	
⋮	
<i>lb_m ub_m</i>	

Example:

```
LCTDETAIL
  STATE      0
  INTERVAL    2
    20      22
    30      32

  STATE      1
  INTERVAL    1
    30      31
END
```

LCTGRID

This is a list keyword that specifies the grid to carry out the **ILCT1D** or **ILCT2D** calculations. The valid options are defined below

<u>Option</u>	<u>Description</u>
<i>state n</i>	<i>n</i> is the quantum number of the vibrational state (0 is the ground state) for which information is to be printed.
<i>Npoints</i>	are the single point energy calculations carried out at each tunneling energy and fitted to a spline. <i>Npoints</i> must be specified with ILCT1D . The default is <i>Npoints</i> equal to 9.
<i>NumberE NumberX</i>	The 2D grid to perform the ILCT2D calculation can be specified here. <i>NumberE</i> are the number of tunneling energies and <i>NumberX</i> is the number of points within each tunneling energy.

LCTOPT

LCTOPT is a list keyword that is used to set options for the large-curvature tunneling calculations. If this keyword is used, then it is assumed that the large-curvature tunneling calculations are desired and LCT is automatically switched on. The valid options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>ILCT1D</i>	This switch keyword is used to calculate LCT3 or LCT4 transmission factors. At each tunneling energy performs a 1D spline under tension fit of the nonadiabatic region. The number of energies calculated along a given tunneling path is specified in LCTGRID.	off
<i>ILCT2D</i>	This switch keyword can be used either to calculate LCT3 or LCT4 transmission factors, although we recommend its use to evaluate only the latter. This is a 2D spline under tension fit based on a grid of tunneling energies and points along the tunneling path for each tunneling energy. The grid is specified in LCTGRID.	off
<i>interpolate</i>	order of interpolation used for the LCG3 theta integrals, the excited state potential, the vibrational period, and the sine of the angle between the MEP and the tunneling path.	4
<i>lcgmethod</i>	Select the method for LCT calculations. Two values are possible: 3 and 4, for LCG3 and LCG4 methods respectively.	4
<i>lctrst</i>	restart a previous large curvature calculation from unit fu48. This is useful to calculate LCT4 transmission factors from a previous LCT3 run.	off
<i>lctstr</i>	store information in unit fu49 for a possible restart.	on
<i>ngamp</i>	number of points for Gauss-Legendre quadrature and trapezoidal rule in computing the total tunneling amplitude along the incoming trajectory in the LCG3 and LCG4 methods.	120
<i>ngtheta</i>	number of points for Gauss-Legendre quadrature in computing the theta integral over the straight-line tunneling path in the LCG3 and LCG4 methods	120
<i>prfreq</i>	print information in fu22 file about the frequency of the mode $p(s)$, the classical energy $V_{\text{MEP}}(s)$, and the vibrationally adiabatic potential energy as a function of s and $n_p(V_p(n_p, s))$, and all the uniformized probabilities at each n_p .	off
<i>prprob</i>	print all uniformized probabilities at each n_p to fu22	off
<i>rtpjac</i>	print the MEP and representative tunneling path turning points in mass-scaled pseudo-Jacobian coordinates to fu24	off
<i>vadavg</i>	average the overlapping of two adiabatic potentials instead of choosing the minimum of them.	off
<i>vefsrch numb</i>	search for possible nonadiabatic LCG4 regions in totally adiabatic LCG3 regions. The search starts from that tunneling energy until <i>numb</i>	0

Example:

```
LCTOPT
  lctrst
  ngamp      120
  ngtheta    120
END
```

QRST

QRST is a list keyword that is used to specify the options for the quantized-reactant-state tunneling calculations. This option should only be used with unimolecular reactions. It is described in Refs. 8 and 37 of Sect. 23. Be sure that MXWKB is large enough if you use this (See Section 5.A.2. MXWKB must be greater than or equal to the total number of energy levels.) STATES subkeyword determines the number of quantized reactant states to be included in the tunneling calculations. The options are *all* (all possible states are included in the calculation), *0* (only the vibrational ground state will be considered), *1* (only the ground state and the first vibrational states will be considered), etc. STQVIB subkeyword determines the number of quantized reactant states to be included in calculation of the reactant vibrational partition function. The options are *all* (all possible states are included in the calculation), and *same* (the number of states used for calculating the reactant vibrational partition function is the same as the number of states used for tunneling contributions). The valid options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>harmonic</i>	use the harmonic approximation to determine the energy levels of the reaction-path mode	on
<i>harmwkb</i>	between the reactant and saddle point use the WKB approximation to find the energy levels but use the harmonic potential as the effective potential to the left of the reactant	off
<i>symwkb</i>	use the WKB approximation to find the energy levels and assume that the effective potential is symmetric with respect to the reactant	off
<i>mode</i>	the number of the selected vibrational reaction-path mode	required
<i>srw</i>	the value of the reaction coordinate at the position of the reactant. The default unit is angstrom and it can also be in bohr by specifying the keyword INPUNIT AU.	required
<i>states</i>	the number of quantized reactant states to be included in the tunneling calculation	<i>all</i>
<i>stqvib</i>	the number of quantized reactant states to be included in the vibrational partition function calculation	<i>all</i>

Example:

```
QRST
  SYMWKB
  STATES      5
  MODE        3
  SRW         -2.42
  STQVIB     all
END
```

QUAD

QUAD is a list keyword that is used to specify the Gauss-Legendre quadrature points in each segment used in computing the Boltzmann average (NQE) and theta integrals (NQTH) in the tunneling calculations. The valid options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>nqe</i>	number of Gauss–Legendre quadrature points in each segment to be used for computing the Boltzmann average in the tunneling calculations; allowed values are 2-7, 9, 10, 12, 15, 20, 30, and 40	40
<i>nqth</i>	number of Gauss–Legendre quadrature points in each segment to be used for computing the theta integrals in the tunneling calculations; allowed values are 2-7, 9, 10, 12, 15, 20, 30, and 40	40
<i>nsegboltz</i>	number of equal segments for integrals that average the energy–dependent tunneling probabilities over a Boltzmann distribution of energies	1
<i>nsegtheta</i>	number of equal segments for the theta integrals	1

Example:

```
QUAD
  nqe      30
  nqth    30
  nsegboltz 2
  nsegtheta 2
END
```

SCT (NOSCT)

SCT is a switch keyword that specifies that small–curvature tunneling calculations be performed. Default is SCT.

SCTOPT

SCTOPT is a list keyword that is used to specify the method for interpolating the effective mass for the small–curvature tunneling calculations. See Section 10.E. The default is SPLINE. The valid options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>lagrange n</i>	Lagrangian interpolation of order n is used to interpolate $\mu_{\text{eff}}(s)$	4
<i>spline</i>	spline fit is used to interpolate $\mu_{\text{eff}}(s)$	on

Example:

```
SCTOPT
  lagrange 5
END
```


WIGNER

WIGNER is a switch keyword that specifies that the Wigner transmission coefficient be computed. Default is on because it is inexpensive, but it is not a reliable method. In the current version of *POLYRATE*, this option cannot be turned off.

ZCT (NOZCT)

ZCT is a switch keyword that specifies that zero-curvature tunneling calculations be performed. Default is ZCT.

12.A.9. RATE section

The RATE section contains the keywords that control the rate constant calculations. This section is not needed if rate constants are not desired. If this section is not included then all defaults are ignored, and the default CVT and TST calculations are not performed. For simplicity the keywords have been broken into three categories: basic, methods, and print. This is only for clarity in the manual. The keywords in any of these three categories can be used in any order. Also for simplicity the keywords in each category have been labeled as either *common* or *other*. The keywords labeled *common* are those that are found in most input files for a standard *POLYRATE* run. This labeling is only intended as an aid to the novice user. Again an alphabetical glossary of all the keywords follows the three summary tables.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Common:</i>			
BOTHK	switch	on	forward and reverse rate constants
EACT	list	not used	activation energy calculation
REVKEXP	variable	0	scale factor for reverse rates
SIGMAF	variable	1	sym. factor for forward reaction
SIGMAR	variable	1	sym. factor for reverse reaction
TEMP	list	see below	temperatures for rate calculations
<i>Others:</i>			
ANALYSIS	list	not used	detailed analysis of TST/CVT ratio
EDGEOK	switch	on	continue calculation even if ΔG_{\max} is near end of grid
FORWARDK	switch	off	forward rate constant only
GSPEC	list	not used	special grid for ΔG calculation
SMAX	variable	1.0 a _o	max. s for special grid
SMIN	variable	-1.0 a _o	min. s for special grid
GTEMP	list	not used	special grid for ΔG calc. for given T
GTLOG	switch	off	free energy of activation by logarithms
STATE	variable	<i>therm</i>	thermal or state-selected rate constant
STATEOPT	list	<i>thermal</i>	options for STATE keyword
SPTOPT	list		options for simple pert. theory
CORIOLIS	switch	off	include Coriolis term
PRINT	switch	off	extra data printed out
VPFEXP	variable	0	scale factor for vibrational partition function

Methods:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<u>Common:</u>			
CVT	switch	on	canonical VTST
<u>Others:</u>			
TST	switch	on	conventional transition state theory
MUVT	switch	off	microcanonical VTST
MUVTOPT	list		options for μ VT
FIT	variable	<i>gas53</i>	primary and secondary fits
NITER	variable	<i>30</i>	Gauss-Laguerre quadrature
PREENERGY	variable	<i>0</i>	# of energies on the grid to print
SLOWER	variable	<i>0.0</i>	lower limit of MEP for searching
SUPPER	variable	<i>0.0</i>	upper limit of MEP for searching
CUS	variable	off	canonical unified statistical model
US	switch	off	unified statistical calculation

Print:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Others:</i>			
PRDELG	switch	off	print free energy curve
PRPART	variable	not used	print reactant, product, and saddle point partition functions
PRVIB	list	not used	region to print extra vib mode info
SX	variable	0.0	s at beginning of desired region
SY	variable	0.0	s at the end of desired region

Glossary of RATE keywords

ANALYSIS

ANALYSIS is a list keyword used to specify the temperatures (in Kelvin) at which a detailed analysis of the conventional TST/CVT ratio is to be performed. The default is not to do a detailed analysis at any temperature. (Note: the rate constant must be computed at all temperatures on this list; see the TEMP keyword.)

Example:

```
ANALYSIS
    298.
    400.
    800.
END
```

BOTHK (NOBOTHK)

BOTHK is a switch keyword that specifies that both the forward and reverse rate constants and the equilibrium constant are to be calculated. If NOBOTHK is specified no rate constants are computed; use FORWARDK if only the forward rate constant is desired. The default is BOTHK. (See REVKEXP for scaling the output for the reverse rate constant.)

CUS (NOCUS)

CUS is a variable keyword that specifies that rate constants should be calculated with the canonical unified statistical model. See Ref. 51 in Sect. 22. The input argument of this keyword indicates the maximum number of local maxima in the free energy of activation vs. reaction coordinate profiles that should be included at each temperature. If the number of local maxima that are found is less than this argument, then all maxima are used. The CUS calculations will also include the lowest local minimum between any pair of included local maxim. When the argument of CUS equals 1, the CUS result is exactly the same as CVT. The default is NOCUS.

Example:

```
CUS      2
```

CVT (NOCVT)

CVT is a switch keyword that specifies that generalized transition state theory and canonical variational theory rate constants be calculated. The default is CVT. If NOCVT is specified and NOTST is not specified, rate constants will be calculated but only at the conventional-TST level. In this case, the calculations are carried out in subprogram TSRATE (as opposed to RATE or GIVTST). TSRATE does not include anharmonicity.

EACT

EACT is a list keyword used to specify the pairs of temperatures (in Kelvin) for which to compute Arrhenius activation energies (maximum of 10 pairs) from the calculated rate constants. Each line should first give the lower temperature and then the upper temperature for the pair. The default is not to compute any activation energies. (Note: the rate constant must be computed at all temperatures in this list; see the TEMP keyword.)

Example:

```
EACT
    298.  400.
    400.  600.
END
```

EDGEOK (NOEDGEOK)

EDGEOK is a switch keyword that specifies that the calculation be continued even if V_a or ΔG^{ICVT} is near an end point, i.e., even if the grid does not appear wide enough to find variational transition states. The default is EDGEOK.

FORWARDK (NOFORWARDK)

FORWARDK is a switch keyword that specifies that only the forward rate constant be calculated. If FORWARDK is specified, equilibrium constants will not be computed. The default is NOFORWARDK.

GSPEC

GSPEC is a list keyword that is used to specify that the generalized free energy of activation profile only be computed for the range $\text{SMIN} \leq s \leq \text{SMAX}$ for all temperatures. The default is to compute the free energy curve over the entire range of s . The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The defaults for the options are $\text{SMAX} = 1 \text{ a}_0$ and $\text{SMIN} = -1 \text{ a}_0$.

Example:

```
GSPEC
    SMAX  1.50
    SMIN -0.05
END
```

GTEMP

GTEMP is a list keyword that is used to specify that the generalized free energy of activation profile be computed over a different range of s for each temperature. Each entry in the list should contain the temperature in Kelvin and then SMAX and SMIN (in bohr) for that temperature. The default is to compute the free energy curve over the entire range of s . The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU. The defaults for the options are $\text{SMAX} = 1 \text{ a}_0$ and $\text{SMIN} = -1 \text{ a}_0$.

Example:

```
GTEMP
    298.  1.50  -0.10
```

```
400. 1.00 -0.25  
END
```

GTLOG (NOGTLOG)

GTLOG is a switch keyword that specifies that the generalized free energy of activation should be calculated by using the logarithms of the vibrational partition functions. This type of calculation is preferred when the vibrational partition function has an extremely small value (as may be the case for very low temperatures and/or for very big systems because we put the zero of energy for the partition coefficient at the lowest value of the potential rather than at the zero point energy). The default is NOGTLOG.

MUVT (NOMUVT)

MUVT is a switch keyword that specifies that μ VT rate constants should be calculated. The default is NOMUVT if US and MUVTOPT are off, and the default is MUVT if US is on or MUVTOPT is given.

MUVTOPT (NOMUVTOPT)

MUVTOPT is a list keyword that is used to set options for the μ VT calculation. If MUVTOPT is given then it is assumed that the μ VT calculation is desired and the default for MUVT becomes on. The valid options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>																																								
<i>niter</i>	number of Gauss-Laguerre quad. pts. for computing μ VT rate constants	30																																								
<i>preenergy</i>	number of energies on the grid to be printed out; PREENERGY must be $\leq \min[6, niter]$	0																																								
<i>slower</i>	lower limit, respectively, of the region of the MEP over which the GT minimum of $N_{\mu VT}$ will be searched for. (This is only specified if one wishes to limit the search. Leave set to the default of zero to search over the entire s range.) The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.	0.0																																								
<i>supper</i>	upper limit, of the region of the MEP over which the GT minimum of N will be searched for. (This is only specified if one wishes to limit the search. Leave set to the default of zero to search the entire s range.) The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.	0.0																																								
<i>fit</i>	variable that specifies the degree of the primary and secondary fits. The allowed values are given below. For a gas-phase reaction, <i>gas53</i> is the recommended value, and for a reaction at a gas-solid interface, <i>gsolid13</i> is recommended. Sometimes there is a numerical problem with the 5-point fit; in such a case it is recommended to use <i>gas31</i> rather than <i>gas53</i> .	<i>gas53</i>																																								
	<table><thead><tr><th><u>value</u></th><th><u>primary</u></th><th><u>secondary</u></th><th><u>phase</u></th></tr></thead><tbody><tr><td><i>gas53</i></td><td>5</td><td>3</td><td>gas</td></tr><tr><td><i>gas15</i></td><td>1</td><td>5</td><td>gas</td></tr><tr><td><i>gas35</i></td><td>3</td><td>5</td><td>gas</td></tr><tr><td><i>gas31</i></td><td>3</td><td>1</td><td>gas</td></tr><tr><td><i>gas13</i></td><td>1</td><td>3</td><td>gas</td></tr><tr><td><i>gsolid13</i></td><td>1</td><td>3</td><td>gas-solid interface</td></tr><tr><td><i>gsolid15</i></td><td>1</td><td>5</td><td>gas-solid interface</td></tr><tr><td><i>gsolid31</i></td><td>3</td><td>1</td><td>gas-solid interface</td></tr><tr><td><i>gsolid53</i></td><td>5</td><td>3</td><td>gas-solid interface</td></tr></tbody></table>	<u>value</u>	<u>primary</u>	<u>secondary</u>	<u>phase</u>	<i>gas53</i>	5	3	gas	<i>gas15</i>	1	5	gas	<i>gas35</i>	3	5	gas	<i>gas31</i>	3	1	gas	<i>gas13</i>	1	3	gas	<i>gsolid13</i>	1	3	gas-solid interface	<i>gsolid15</i>	1	5	gas-solid interface	<i>gsolid31</i>	3	1	gas-solid interface	<i>gsolid53</i>	5	3	gas-solid interface	
<u>value</u>	<u>primary</u>	<u>secondary</u>	<u>phase</u>																																							
<i>gas53</i>	5	3	gas																																							
<i>gas15</i>	1	5	gas																																							
<i>gas35</i>	3	5	gas																																							
<i>gas31</i>	3	1	gas																																							
<i>gas13</i>	1	3	gas																																							
<i>gsolid13</i>	1	3	gas-solid interface																																							
<i>gsolid15</i>	1	5	gas-solid interface																																							
<i>gsolid31</i>	3	1	gas-solid interface																																							
<i>gsolid53</i>	5	3	gas-solid interface																																							

Example:

```
MUVTOPT
  niter 40
END
```

PRDELG (NOPRDELG)

PRDELG is a switch keyword that specifies that $\Delta G(T,s)$ be printed to fu6. Default is NOPRDELG.

PRPART (NOPRPART)

PRPART is a variable keyword. Its argument specifies that the reactant and product (*rp*) partition functions, the saddle point (*t*) partition functions, or the reactant, product, and saddle point (*rp**t*) partition functions should be printed to fu6. When the reactant partition function is printed for a gas-phase unimolecular reaction the internal free energy of reactant is also printed. The default is not to print any partition functions or free energies at the stationary points.

Example:

```
PRPART    rp
```

PRVIB

PRVIB is a list keyword that is used to specify the initial and final *s* values, with keywords *sx* and *sy* respectively, that delineate a region for which extra vibrational mode and Q_{vib} information be printed to fu6. The default is no extra printing. The default unit is angstroms and it can also be in bohr by specifying the keyword INPUNIT AU.

Example:

```
PRVIB
      SX      0.0
      SY      0.6
END
```

REVKEXP

REVKEXP is a variable keyword whose argument specifies the value *n* in the scale factor 10^{-n} for the reverse rate constant. The final rate will be calculated in units of $10^{-n} \text{ cm}^3 \text{ molecule}^{-1} \text{ s}^{-1}$ if there are two products or in units of 10^{-n} s^{-1} if there is one product. Default is $n = 0$, i.e., a scale factor of 1.

Example:

```
REVKEXP    10
```

SIGMAF

SIGMAF is a variable keyword whose argument specifies the number of equivalent ways for forming the GTS for the forward reaction. The default value is 1.

Example:

```
SIMGAF      1
```

SIGMAR

SIGMAR is a variable keyword whose argument specifies the number of equivalent ways for forming the GTS for the reverse reaction. The default value is 1.

Example:

```
SIMGAR      1
```

STATE

STATE is a variable keyword whose argument specifies if this is to be a thermal or state-selected rate constant calculation. The default is *therm*. The valid arguments are:

<u>Option</u>	<u>Description</u>
<i>therm</i>	thermal run
<i>adiab</i>	adiabatic state-selected; STATEOPT is required
<i>diab</i> <i>x</i>	diabatic state-selected; <i>x</i> is the value of the reaction coordinate at which the adiabatic/diabatic switch is made; STATEOPT is required

Example:

```
STATE adiab
```

STATEOPT

STATEOPT is a list keyword that is used to specify how each mode in each reactant, product, and transition state is to be treated. By default each mode is treated thermally. The user needs to specify each mode explicitly that is to be restricted to the ground state or first excited state as shown below. Each input line in the list requires:

reatyp *nmode* *modtyp*

The allowed values of the variable *reatyp* are:

Value	Description
<i>react1</i>	reactant #1
<i>react2</i>	reactant #2
<i>prod1</i>	product #1
<i>prod2</i>	product #2
<i>ts</i>	transition state and along the reaction path
<i>switts</i>	reaction path after the adiabatic/diabatic switch (see STATE)

nmode is an integer specifying the vibrational mode number

The allowed values of the variable *modtyp* are:

Value	Description
<i>thermal</i>	mode is thermal (default)
<i>ground</i>	mode is restricted to the ground state
<i>first</i>	mode is restricted to the first excited state

Example:

```
STATEOPT
  react1  1  ground
  react2  1  first
  prod1   1  thermal
  prod1   2  ground
  prod1   3  first
  ts      4  ground
  ts      5  first
  switts  4  first
  switts  5  ground
END
```

TEMP

TEMP is a list keyword used to specify the temperatures (in Kelvin) at which rate constants be computed. Unless NOBOTHK or FORWARDK is selected the equilibrium constants will also be calculated at these temperatures. The default is to compute the rate constants at 7 temperatures: 200, 300, 400, 600, 1000, 1500, and 2400.

Example:

```
TEMP
  298.
  400.
  600.
  800.
END
```

TST (NOTST)

TST is a switch keyword that specifies that the conventional transition state theory rate constants be calculated. The default of TST is on. If any other rate constants are calculated, i.e., CVT, ICVT, US, or μ VT, then the conventional rate constants will automatically be calculated (NOTST will have no effect) except when VRC is turned on

US (NOUS)

US is a switch keyword that specifies that unified statistical rate constants be calculated. If US is on, then MUVT is automatically turned on. The default of US is off, which is the same as specifying NOUS.

VPFEXP

VPFEXP is a variable keyword that that specifies that the total vibrational partition functions be scaled by 10^n where n is the argument. The default is no scaling.

Example:
VPFEXP 2

12.B. General description of file fu29 input

File fu29 is only needed if IVTST0 or IVTST1 is specified in the GENERAL section of the unit fu5 input. This unit contains input data for IVTST rate calculations. The first section of this input file must be *IVTST. All input sections must start with the section name preceded by an "*", and must end with an END line (a line containing "end"). Keywords and input data for each section are discussed further in the rest of this subsection. Sample inputs can be found in several test runs described in Section 16.

<u>Section Name</u>	<u>Description</u>
*E1GEOM	Cartesian coordinates of the extra point on the MEP
*E1GRAD	Cartesian gradient vector of the extra point on the MEP
*E1HESS	Cartesian Hessian of the extra point on the MEP
*IVTST	general data for IVTST calculations (must come first)
*P1FREQ	frequencies of the 1st product
*P2FREQ	frequencies of the 2nd product
*P1HESS	Cartesian Hessian of the 1st product
*P2HESS	Cartesian Hessian of the 2nd product
*R1FREQ	frequencies of the 1st reactant
*R2FREQ	frequencies of the 2nd reactant
*R1HESS	Cartesian Hessian of the 1st reactant
*R2HESS	Cartesian Hessian of the 2nd reactant
*TSH Hess	Cartesian Hessian of the saddle point
*TSFREQ	frequencies of the saddle point

When IVTST0 is specified, the user can choose between giving the Hessian or the frequencies for the stationary points (reactants, saddle point and products) according to the kind of information available for the system. When running an IVTST1 calculation

the use of Hessians for the stationary and extra points is required, since this information is needed in order to calculate the SCT transmission factors.

12.B.1. E1GEOM Section

The E1GEOM section of the file fu29 input is used to give the Cartesian coordinates of the extra point on the MEP. This section is only needed for first-order IVTST calculations.

Example:

```
*E1GEOM
0.000000  0.000000  0.000000
0.456364  1.749560  0.000000
1.808100  0.000000  0.000000
END
```

12.B.2. E1GRAD Section

The E1GRAD section of the file fu29 input is used to give the Cartesian gradient vector of the extra point on the MEP. This section is only needed for first-order IVTST calculations.

Example:

```
*E1GRAD
0.000000  0.000000  0.000000
0.056364  0.009560  0.000000
0.008100  0.000000  0.000000
END
```

12.B.3. E1HESS, P1HESS, P2HESS, R1HESS, R2HESS, and TSHESS sections

The E1HESS, P1HESS, P2HESS, R1HESS, R2HESS, and TSHESS sections of file fu29 input are used to give the Cartesian Hessian matrix of the extra point on the MEP, products, reactants, and the transition state in packed form, respectively. The E1HESS section is only needed for first-order IVTST calculations.

Example:

```
*E1HESS
0.000000
0.056364  0.009560
0.008100  0.000000  0.000000
0.008100  0.000000  0.000000  0.080100
0.008100  0.000000  0.000000  0.080100  0.080100
0.008100  0.000000  0.000000  0.080100  0.080100
0.080100
END
```

12.B.4. IVTST Section

The IVTST section of the file fu29 input is used to give general data for IVTST calculations. The table below lists all valid subkeywords for the IVTST section. A detailed explanation of each keyword is given alphabetically in the glossary below.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
BARRI	variable	required	classical barrier height in kcal/mol
DELTAE	variable	required	classical energy of the products in kcal/mol
DELEX1	variable	not used	the classical energy of the extra point in kcal/mol (only needed for the IVTST1)
PRINT	switch	0	output control
SEX1	variable	not used	the reaction coordinate at the extra point in atomic units (only needed IVTST1)
UNITS	switch	0	units for geometry, gradient, and Hessian matrix

Glossary of IVTST keywords

BARRI

BARRI is a keyword that specifies the classical barrier height in kcal/mol. In case a conversion of units is required for obtaining this value in kcal/mol, the conversion factor used should be the same as that internally employed by *POLYRATE* as specified in the percon.inc file.

Example:

```
BARRI      5.0
```

DELTAE

DELTAE is a keyword that specifies the classical energy of the products relative to the classical energy of the reactants in kcal/mol. This is a required keyword. In case a conversion of units is required for obtaining this value in kcal/mol, the conversion factor used should be the same as that internally employed by *POLYRATE* as specified in the percon.inc file.

Example:

```
DELTAE    -5.0
```


DELEX1

DELEX1 is a keyword that specifies the classical energy of the extra point relative to the classical energy of the reactants in kcal/mol. This keyword is only needed for first-order IVTST calculations.

Example:

```
DELEX1    -4.4
```

PRINT

PRINT is a keyword for output control: 0—normal (default), 1—debug.

Example:

```
PRINT     1
```

SEX1

SEX1 is a keyword that specifies the reaction coordinate at the extra point in bohr. This keyword is only needed for first-order IVTST calculations.

Example:

```
SEX1      0.1
```

UNITS

UNITS is a keyword that specifies the units for the geometry, gradients and Hessian matrix: UNITS = 0: The geometry will be in bohrs, and both the gradients, and the Hessian matrix will be in mass-scaled atomic units (default). UNITS = 1: The geometry will be in mass-scaled Å, and both the gradient vectors and the Hessian matrix will be in non-mass-scaled atomic units.

Example:

```
UNITS     0
```

12.B.5. P1FREQ, P2FREQ, R1FREQ, R2FREQ, and TSFREQ sections

The P1FREQ, P2FREQ, R1FREQ, R2FREQ, and TSFREQ sections of the file fu29 input are used to give the frequencies (in cm^{-1}) of the product and reactant molecules and the saddle point, respectively. The imaginary frequency of the saddle point must be the last input value in the TSFREQ section.

Example:

```
*R1FREQ
    2000.0    1000.0
END
```

12.C. General description of file fu30 input

File fu30 contains the information typically obtained from electronic structure calculations which is required for performing a direct dynamics calculation (see Section 9 for a detailed description of this capability). This option is selected by setting the keyword POTENTIAL in the ENERGETICS section equal to *unit30*.

File fu30 is organized in the following way: 1) a one-line title; 2) 2 lines with numerical options for routing and output; and 3) several blocks of data containing the specific information provided for each point on the reaction path according to the options selected (s, V, frequency matching data, Cartesian coordinates, gradient, frequencies or Hessian matrix, anharmonicity data, curvature component data, etc.).

The user should be aware that gradients provided in the fu30 file must correspond to structures that have not been arbitrarily reoriented along the reaction path in order to obtain meaningful results in the calculation of the curvature components B_{mF} . See Section 9 for a fuller discussion of this issue.

A record-by-record description (where a record is one or more lines) of the structure of the FORTRAN unit fu30 follows. If the keyword POTENTIAL = *unit30*, POLYRATE can be used with the dummy potential energy surface file provided, dumpot.f, since the potential routines SURF and SETUP will not be called.

<u>Record</u>		<u>Format</u>
	(Records 1 and 2 are read in subprogram RPHSET)	
1	One-line title	(80A1)
2	Routing and output options (LOPT(I), I = 1, 40)	(*)
	LOPT(1) LEVEL OF PRINT	
	≤ 0, print out only titles and options	
	= 1, also print out summary	
	= 2, also echo input as it is read	
	= 3, also print out intermediate information about fitting exponential functions and print interpolated results each time	
	out RPHINT is called	
	≥ 4, also print out debugging information in exponential fitting	

LOPT(2) MAXIMUM NUMBER OF POINTS USED IN LAGRANGE INTERPOLATION IN THE PROCEDURE OF SUBSECTION 9.C.

= -1, use IVTST-M for interpolating. See Sect. 9.C.

Note: When using file fu30, the user cannot select any options for the IVTST-M calculation; it will be performed according to the default options, and the number of gradient and Hessian points have to be the same and equal to NPTS (see below). Therefore, only IVTST-M-NPTS/NPTS calculations are allowed. The curvature components will be interpolated using a value of 3 for the exponent in the mapping function. For unimolecular reactions or reactions with wells, a cubic polynomial fitting to V is done previous to its spline fitting, taking a step of $0.01 a_0$ along the lowest frequency mode for performing this fitting. The only way to change any of these options or to use a different number of gradient and Hessian points is by using the fu31 input file, which is specifically designed for IVTST-M calculations.

LOPT(3) OPTIONS FOR COORDINATES (X) AND GRADIENTS (DX)

≤ -2,	X in mass-scaled Cartesians, DX in unscaled Cartesians
= -1,	X in unscaled Cartesians, DX in mass-scaled Cartesians
= 0,	X and DX in unscaled Cartesians
= 1,	X and DX in mass-scaled Cartesians
= 99,	X in unscaled Cartesians, -DX in mass-scaled Cartesians
= 100,	X in unscaled Cartesians, -DX in unscaled Cartesians

Note: DX denotes the gradient, and -DX denotes the negative of the gradient which is the force. For dynamics calculations in Cartesian coordinates, the normalized gradients (or forces) can be used without any effects on the results. However, unnormalized gradients have to be used for the curvilinear option. Notice the ‘Cartesian Forces’ from GAUSSIAN formatted check point output file are actually the Cartesian gradient components.

LOPT(4) OPTIONS FOR FREQUENCY OR HESSIAN MATRIX INPUT

There are three issues:

- (1) Are the frequencies read in from unit fu30, or are they obtained by diagonalizing the Hessian matrix \mathbf{F} or the projected Hessian \mathbf{F}^P ?
- (2) If the frequencies are calculated from the Hessian, how is \mathbf{F} read in, in packed (triangular) form, in packed form with full lines, or in unpacked (full matrix) form?
- (3) What coordinate system was used in computing \mathbf{F} ?

LOPT(4)	(1)	(2)	(3)
≤ -2,	diagonalization	packed	mass-scaled coordinates
= -1,	diagonalization	packed	unscaled coordinates

= 0, frequencies are read
 = 1, diagonalization full matrix unscaled coordinates
 = 2, diagonalization full matrix mass-scaled coordinates

- Notes: (1) If LOPT(4) \neq 0, the choice of coordinates (mass-scaled or not) must be consistent with LOPT(3).
 (2) If the Hessian is read, the code assumes it is the non-projected Hessian.
 (3) Packed full lines means 5 elements per line; whereas packed triangular means F(1,1) on first line, F(2,1) and F(2,2) on second line, etc.

LOPT(5) ANHARMONICITY INPUT

\leq 0, no anharmonicity data read
 = 1, read the third derivatives k_3 with respect to the normal coordinates at each grid point
 \geq 2, also read the fourth derivatives k_4 with respect to the normal coordinates at each grid point

LOPT(6) OPTIONS FOR CURVATURE COMPONENT B_{mF} INPUT

\leq 0, B_{mF} are set to zero everywhere
 = 1, B_{mF} are read directly
 = 2, B_{mF} are computed. Derivative of the gradient with respect to s is obtained from a quadratic fit of the gradient at three points on the basic input grid (defined in Section 9.C), with the middle point being the current point on the Hessian grid.
 = 3, B_{mF} are computed. Derivative of the gradient is computed by one-sided finite difference, using the gradient on the basic input grid and a gradient input at one extra point near the basic input grid point.
 \geq 4, B_{mF} are computed. Derivative of the gradient is computed from a quadratic fit obtained using the gradient on the basic input grid and gradients input at an extra point on each side of the basic input grid point.

Note: For LOPT(6) $>$ 1, B_{mF} are computed and require the coefficient matrix from the diagonalization of the Hessian matrix. Therefore, LOPT(4) must be nonzero if LOPT(6) $>$ 1.

LOPT(7)-LOPT(40): Not used in the present version of *POLYRATE*.

Loop over records 3 through 14 for each reactant species and then for each product species. Records 3-25 are read in subprogram RPHRD2.

3 One-line title (80A1)

4 S, V (*)

S = value of s in bohrs at reactant (or product). This only has meaning for a unimolecular and/or association reaction, for which s is finite at the reactant and/or product; otherwise S is ignored.

V = potential (in atomic units) at reactants (or products).

For an atomic reactant or product, skip records 5 through 14, **add a blank line after record 4**, and then continue with the next species.

5 (IFREQ(I), I = 1, NFREQ) (*)

IFREQ = indices for reordering frequencies, anharmonicities, and curvature components in the interpolation table for the reactant or product species, in order to make the ordering consistent with that at the saddle point and/or along the MEP.

Note: Index 1 corresponds to the largest frequency at the saddle point, index 2 to the second largest frequency, and so on. The imaginary frequency is not included in this ordering.

Note that this matching is not necessary if you are just calculating conventional TST reaction rates.

NFREQ is the number of bound vibrational modes for the species.

For a Morse diatom reactant or product, read record 6; otherwise go to record 7.

6 SGN

SGN = sign used to multiply the third derivative of the diatomic potential. This can be used to make the sign of the third derivative of this mode in the asymptotic reactant or product region match up with the correlating mode in the interaction region. This ensures a smooth extrapolation into the asymptotic region.

For a Morse diatom reactant or product, skip records 7 through 14. If LOPT(4) = 0 (frequencies directly input), skip records 7 and 8.

7 One-line title (80A1)

8 N records of (F(I,J), J = 1, JF) N records of (*)

\mathbf{F} = Hessian matrix (in atomic units) for the reactant or product species. Second derivatives are with respect to Cartesian coordinates; LOPT(4) determines whether the coordinates are mass-scaled or not. $JF = N$ for LOPT(4) > 0 and $JF = I$ for LOPT(4) < 0, where $N = 3 * NRATOM$, NRATOM = number of atoms in the species, and I indexes the N records in this set ($I = 1, \dots, N$).

If LOPT(4) \neq 0, skip records 9 and 10.

9 One-line title (80A1)

10 (FREQ(I), I = 1, NFREQ) (*)

FREQ = vibrational frequencies (**in atomic units**) for the reactant or product species.

If LOPT(5) < 1 (no anharmonicity), skip records 11 through 14.

11 One-line title (80A1)

12 (XK3(I), I = 1, NFREQ) (*)

XK3 = third derivatives of V with respect to motion along the normal coordinates for the reactant or product species.

If LOPT(5) < 2 (just third derivatives), skip records 13 and 14.

13 One-line title (80A1)

14 (XK4(I), I = 1, NFREQ) (*)

XK4 = fourth derivatives of V with respect to motion along the normal coordinates for the reactant or product species.

After each reactant or product species except for the last product, go back to record 3.
After the last product, go to the next record.

If there is no saddle point, skip records 15 through 27.

15 One-line title (80A1)

16 S, V (*)

S = a dummy quantity and is ignored.
 V = potential (in atomic units) at the saddle point.

17 (IFREQ(I), I = 1, NFREQ) (*)

These quantities are the same as in record 5, but now they refer to the bound vibrational modes of the saddle point. NFREQ is now the number of bound vibrational modes of the saddle point.

If LOPT(4) = 0, skip records 18 and 19.

18 One-line title (80A1)

19 $N3$ records of (F(I,J), J = 1, JF) $N3$ records of (*)

These quantities are the same as in record 8, but now they refer to the saddle point. $JF = N3$ for LOPT(4) > 0 and $JF = I$ for LOPT(4) <

0, where I indexes the $N3$ records in this set ($I = 1, \dots, N3$), and where $N3 = 3 * \text{NATOM}$, NATOM = number of atoms in the system.

If $\text{LOPT}(4) \neq 0$, skip records 20 and 21.

20 One-line title (80A1)

21 (FREQ(I), $I = 1, \text{NFREQ}$) (*)

These quantities are the same as in record 10, but now they refer to the bound vibrational modes of the saddle point.

If $\text{LOPT}(5) < 1$ (no anharmonicity), skip records 22 through 25.

22 One-line title (80A1)

23 (XK3(I), $I = 1, \text{NFREQ}$) (*)

These quantities are the same as in record 12, but now they refer to the bound vibrational modes of the saddle point.

If $\text{LOPT}(5) < 2$ (just third derivatives), skip records 24 and 25.

24 One-line title (80A1)

25 (XK4(I), $I = 1, \text{NFREQ}$) (*)

These quantities are the same as in record 14, but now they refer to the bound vibrational modes of the saddle point.

If $\text{LOPT}(4) \neq 0$, skip records 26 and 27.
Records 26-31 are read in subprogram RPHRD1.

26 One-line title (80A1)

27 WSTAR (*)

WSTAR = absolute value of the imaginary frequency at the saddle point.

If the reaction path is calculated, read records 28 to 63.

28 One-line title (80A1)

29 NPTS (*)

NPTS = number of s points along the MEP (not including the saddle point) at which reaction-path Hamiltonian data is to be read (maximum of 200). Unless only TST reaction rates are calculated, file fu30 must contain a minimum of one point on each side of the saddle point. Unless NEXTPT is on, these points must be arranged in order of increasing S .

Loop over records 30 through 52 NPTS times, **from the lowest value of s to the highest value of s** . Records 30-50 are read in subprogram RPHRD2.

30 One-line title (80A1)

31 S, V (*)

S = value of s (in bohr) at the grid location along the MEP.

V = potential (in hartrees) at s on the MEP.

32 One-line title (80A1)

33 $(X(I), I = 1, N3)$ (*)

X = Cartesian coordinates of the GTS at s on the MEP.

$N3 = 3 * (\text{number of atoms in the total system})$. Ordering is the same as specified in the ATOM keyword of the FORTRAN unit fu5 data.

Note: LOPT(3) determines whether the coordinates are mass-scaled or not.

34 One-line title (80A1)

35 $(DX(I), I = 1, N3)$ (*)

DX = gradient vector (in atomic units) at s along the MEP. DX can be the vector of derivatives of the potential with respect to the Cartesian coordinates or it can be normalized. LOPT(3) determines whether the coordinates are mass-scaled or not. Ordering is the same as for $X(I)$ in record 33.

Note: Some electronic structure codes print the force rather than the gradient. Recall that the force is the negative of the gradient.

If LOPT(6) ≤ 2 , skip records 36 through 41.

36 One-line title (80A1)

37 SB(1) (*)

SB(1) = value of s (in bohr) at a location along the MEP close to the current grid point.

38 $(DX(I), I = 1, N3)$ (*)

These quantities are the same as in record 35, but now they refer to the location SB(1) on the MEP.

If LOPT(6) ≤ 3 , skip records 30 through 41.

39 One-line title (80A1)

40 SB(2) (*)

SB(2) = value of s (in bohr) at another location along the MEP close to the current grid point.

41 (DX(I), I = 1, N3) (*)

These quantities are the same as in record 35, but now they refer to the location SB(2) on the MEP.

42 (IFREQ(I), I = 1, NFREQ) (*)

These quantities are the same as in record 17, but now they refer to the bound generalized normal modes of the current grid point along the MEP.

If LOPT(4) = 0, skip records 43 and 44.

43 One-line title (80A1)

44 N3 records of (F(I,J), J = 1, JF) (*)

These quantities are the same as in record 19, but now they refer to the current grid point along the MEP.

If LOPT(4) \neq 0, skip records 45 and 46.

45 One-line title (80A1)

46 (FREQ(I), I = 1, NFREQ) (*)

These quantities are the same as in record 21, but now they refer to the current grid point along the MEP.

If LOPT(5) < 1 (no anharmonicity), skip records 47 through 50.

47 One-line title (80A1)

48 (XK3(I), I = 1, NFREQ) (*)

These quantities are the same as in record 23, but now they refer to the current grid point along the MEP.

If LOPT(5) < 2 (just third derivatives), skip records 51 and 52.

49 One-line title (80A1)

50 (XK4(I), I = 1, NFREQ) (*)

These quantities are the same as in record 25, but now they refer to the current grid point along the MEP.

If LOPT(6) = 1 (B_{mF} are read), read records 51 and 52.
Records 51-63 are read in RPHRD1.

51 One-line title (80A1)

52 (BFS(I), I = 1, NFREQ) (*)

BFS = reaction-path curvature components (in atomic units) at the
current grid point along the MEP.

Return to record 30 until records 30 through 52 have been read NPTS times.

12.D. General description of file fu31 input

This file is only needed if the POTENTIAL keyword of the ENERGETICS section of file fu5 is equal to *unit31*. The input is in free format keyword style and is composed of several sections. When *unit31* is used, the code always uses the IVTST-M algorithm.

12.D.1. Section descriptions

<u>Section Name</u>	<u>Description</u>
*GENERAL	general data such as options for input and IVTST-M
*REACT1	information for the first reactant
*REACT2	information for the second reactant (if any)
*PROD1	information for the first product
*PROD2	information for the second product (if any)
*WELLR	information for the well on the reactant side (if any)
*WELLP	information for the well on the product side (if any)
*SADDLE	information for the saddle point
*POINT	information for a point along the MEP

Each section must appear once and only once except POINT, which has to appear at least twice (since IVTST-M is not defined for less than one point on each side of the reaction path). The POINT section should be repeated once for each of the nonstationary points along the MEP. A saddle point is also required for carrying out an IVTST-M calculation. Therefore, the fu31 input file cannot be used for reactions without a saddle point.

The program is insensitive to the order in which the points are entered. This is one of the main differences between the fu31 input file and the fu30 and fu40 files. This allows us to use an fu31 file written by means of the *writefu31* option for the RESTART keyword in the GENERAL section of the fu5 input file without any change. Another difference is that Hessians are not required at each point; since the calculation of the reaction path always gives a value of the reaction coordinate, *s*, the energy, geometry, and

gradients, this information is required, but the Hessian information is not. This way, the user can take advantage of the information calculated for points between two Hessian points.

Since the IVTST-M calculation is always based on frequencies ordered in canonical order, no frequency indexes are required. Moreover, since anharmonicity is not supported for IVTST-M calculations, in the current version of *POLYRATE*, no anharmonic data is allowed. From all other points of view, the fu31 input file is very similar to a fu30 or fu40 input file.

12.D.2 GENERAL section

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
GEOM	list	no changes	change defaults for reading geometries
GRADS	list	no changes	change defaults for reading gradients
HESSIAN	list	no changes	change defaults for reading Hessians
INTMUEFF	switch	off	interpolation of the exponent in the effective tunneling mass
KIES	switch	off	recalculation of the <i>s</i> values

Glossary of GENERAL keywords

GEOM

GEOM is a list keyword for selecting the units of the geometries input in the fu31 file. The units are completely defined by a combination of two keywords; thus, if we want to input the mass-scaled geometries in angstroms we will have to select both *ms* and *ang*. If one of the options is not given, the default value is assumed.

<u>Option</u>	<u>Definition</u>
---------------	-------------------

<i>ang</i>	Geometries are given in Å.
<i>au</i>	Geometries are given in atomic units (default).
<i>ms</i>	Geometries are given in mass-scaled Cartesian coordinates.
<i>uns</i>	Geometries are given in unscaled Cartesian coordinates (default).

Example:

```
GEOM
      ang
      ms
END
```

GRADS

GRADS is a list keyword for selecting the units of the gradients (or forces) input in the fu31 file. The units are completely defined by a combination of three keywords.

<u>Option</u>	<u>Definition</u>
---------------	-------------------

<i>ang</i>	Gradients are given in Å.
<i>au</i>	Gradients are given in atomic units (default).
<i>ms</i>	Gradients are given in mass-scaled Cartesian coordinates.
<i>uns</i>	Gradients are given in unscaled Cartesian coordinates
(default).	
<i>forces</i>	Forces are given; the gradients are obtained from them.
<i>grads</i>	Gradients are given (default).

Example:

```
GRADS
      ang
      uns
      forces
END
```

HESSIAN

HESSIAN is a list keyword for selecting the units of the Hessian matrices input in the fu31 file. The units are completely defined by a combination of three keywords.

Option Definition

<i>ang</i>	Gradients are given in Å.
<i>au</i>	Gradients are given in atomic units (default).
<i>ms</i>	Gradients are given in mass-scaled Cartesian coordinates.
<i>uns</i>	Gradients are given in unscaled Cartesian coordinates (default).
<i>full</i>	Full Hessian matrices are given.
<i>packed</i>	Packed Hessian matrices are given (default).

Example:

```
HESSIAN
      full
END
```

If *packed* is selected, the lower triangular elements of **F** should be listed in order such that $F_{ij} = P(\frac{1}{2}i(i-1) + j)$, where $P(k)$ is element k of the input list and F_{ij} is the element of **F** in row i and column j . The number of elements listed per line is arbitrary.

Examples of both input forms for a 6 x 6 matrix is shown below:

Full matrix - example 1

```
F11  F12  F13  F14  F15  F16
F21  F22  F23  F24  F25  F26
F31  F32  F33  F34  F35  F36
F41  F42  F43  F44  F45  F46
F51  F52  F53  F54  F55  F56
F61  F62  F63  F64  F65  F66
```

Full matrix - example 2

```
F11  F12  F13  F14  F15  F16  F21  F22  F23
F24  F25  F26  F31  F32  F33  F34  F35  F36
F41  F42  F43  F44  F45  F46  F51  F52  F53
F54  F55  F56  F61  F62  F63  F64  F65  F66
```

Packed matrix - example 1

```
F11
F21  F22
F31  F32  F33
F41  F42  F43  F44
F51  F52  F53  F54  F55
F61  F62  F63  F64  F65  F66
```

Packed matrix - example 2

```
F11  F21  F22  F31  F32  F33
F41  F42  F43  F44  F51  F52
F53  F54  F55  F61  F62  F63
```

F_{64} F_{65} F_{66}

The order of the coordinates is explained in Section 12.E.1.

Note that GAUSSIAN94 writes the packed Hessian matrix in a slightly different way in the standard output file, but the Hessian in the formatted checkpoint file (requested by their FormCheck keyword) is in precisely the packed form accepted by POLYRATE and so is convenient for copying and pasting into POLYRATE fu40 files.

Note that the exact \mathbf{F} matrix is symmetric, but due to finite precision in computations, the actual matrix one has to input is usually nonsymmetric. If one inputs a nonsymmetric full matrix, the code uses the unsymmetric matrix. If a user wants the code to use a symmetric matrix, we recommend taking an arithmetic average of F_{ij} and F_{ji} for the resulting symmetrized elements in packed form.

INTMUEFF (NOINTMUEFF)

The interpolation of the effective mass for CD-SCSAG tunneling can be done in two ways in an IVTST-M calculation: by means of the interpolation of the curvature components and evaluation of the effective mass, or by means of the direct interpolation of the effective mass, specifically, of the term inside the exponential of the equation 16 in Ref. IVTST-M. The latter option can be selected by means of the INTMUEFF switch keyword, but the former option (NOINTMUEFF) is recommended and it is set as the default option.

KIES (NOKIES)

In a calculation of kinetic isotope effects the values of the reaction coordinate, s , calculated for one isotopomer are not valid for another isotopomer, since the values of s are defined in a mass-scaled space. The KIES keyword allows the calculation of the kinetic-isotope-effects-specific values of s . These values are calculated as the straight line between two consecutive points in the isotope-specific mass-scaled space. Therefore, if the KIES keyword is on, the values of s included in the fu31 input file will only be used for ordering the points along the reaction path according to the standard procedure: first the points with negative s , then the saddle point, and then the points with positive s , ordered in increasing value of s . The geometries in file fu31 must be given in unscaled coordinates (GEOM = *uns*). Nevertheless, if the calculation of new values of s gives place to a permutation in the position of two points (which will not occur as a consequence of just a change in an atomic mass) the program will not detect it, and the interpolation will probably fail. The default is off (NOKIES) and its use is only recommended for a KIE's calculation with RODS using a the reaction path calculated with a different isotope.

12.D.3. REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and SADDLE sections

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
VMEP	variable	(see Glossary)	value of $V_{MEP}(s)$
SMEP	variable	(see Glossary)	value of s
GEOM	list	(see Glossary)	geometries
GRADS	list	(see Glossary)	gradients
HESSIAN	list	(see Glossary)	Hessians
ENERXN	variable	(see Glossary)	reaction energy
ENERSAD	variable	(see Glossary)	barrier height
ENEWELLR	variable	(see Glossary)	energy of the well on the reactant side of the reaction-path
ENEWELLP	variable	(see Glossary)	energy of the well on the product side of the reaction-path

Glossary of REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and SADDLE keywords

ENERSAD

ENERSAD is a variable keyword that indicates the value of the barrier height, in atomic units (measured by taking the zero of energy at reactants). Its only acceptable position is in the SADDLE section, where it is required.

ENERWELLR

ENERWELLR is a variable keyword that indicates the energy of the well on the reactant side of the reaction path, in atomic units (measured by taking the zero of energy at reactants). Its only acceptable position is in the WELLR section.

ENERWELLP

ENERWELLP is a variable keyword that indicates the energy of the well on the product side of the reaction path, in atomic units (measured by taking the zero of energy at reactants). Its only acceptable position is in the WELLP section.

ENERXN

ENERXN is a variable keyword that indicates the reaction endoergicity, in atomic units (measured by taking the zero of energy at reactants). Its only acceptable positions are the PROD2 section for reactions with two products, and the PROD1 section for reactions with one product.

GEOM

GEOM is a list keyword that specifies the Cartesian coordinates at the current grid point. Its only acceptable position is the POINT sections, where it is required; the geometries for the stationary points are read from the fu5 input file.

GRADS

GRADS is a list keyword that specifies the Cartesian gradients (or forces) at the current grid point. Its only acceptable position is the POINT sections, where it is required, since the gradients for the stationary points are assumed to be zero.

HESS

HESS is a list keyword that specifies the Cartesian Hessian matrices. It is required in all the REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and SADDLE sections (if they exist and they are not atomic species). It is not required in all the POINT sections, but there must be at least one POINT section with Hessian information on each side of the reaction path.

SMEP

SMEP is a variable keyword that specifies the value of the reaction coordinate s at the current grid point, in mass-scaled atomic units. This keyword is required and only accepted in the POINT sections; for reactants, products, and wells the value of s is calculated as described in Ref. 48 in Sect. 23. If the KIES keyword is on, the values in SMEP are only used for ordering the points, but they could take any value that allow the program to sort the points in its right order.

VMEP

VMEP is a variable keyword that specifies the value of $V_{\text{MEP}}(s)$ at the current grid point, in atomic units and setting the zero of energy at reactants. This keyword is required and only accepted in the POINT sections.

12.E. General description of file fu40 input

This file is needed only if the POTENTIAL keyword of the ENERGETICS section of file fu5 is equal to *unit40*. The input from FORTRAN unit fu40 is in free format keyword style. All input sections must start with the section name preceded by an “*”.

12.E.1 Section Descriptions

<u>Section Name</u>	<u>Description</u>
*GEN40	general data such as options for input and extrapolation
*EXTRAP40	extrapolation information for the reactants and products
*R140	information for the first reactant
*R240	information for the second reactant
*P140	information for the first product
*P240	information for the second product
*WR40	information for reactant side well
*WP40	information for product side well
*SADDLE40	information for the saddle point
*POINT40	information for a point along the MEP

The POINT40 section should be repeated once for each of the points along the MEP excluding the reactant limit, product limit, points used for extrapolation, and saddle point; i.e., POINT40 should be repeated once for each of the points $s_1, s_2, s_3, \dots, s_N$ (in the notation of pages 96-99 of Ref. 14 of section 23) except the saddle point, and if there is a saddle point, that point should not be specified in a POINT40 section. Each section appears at most once except POINT40, which appears NPTS times. The sections can be in any order except that the POINT40 sections must be in order of increasing s . The saddle point, if any, is specified in the SADDLE40 section. In reading in gradients and Hessians, the coordinates are always in this order: x, y, z of atom 1, x, y, z of atom 2, etc.

12.E.2 GEN40 section

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
TITLE	list	blank	job title or comment
PRINT	variable	<i>noextra</i>	print options
MAXLPTS	variable	3	maximum number of points used in Lagrange interpolation
GEOMUNIT	variable	<i>bohr</i>	unit for coordinate input
GRADUNIT	variable	<i>bohr</i>	unit for gradient input
FREQSOURCE	variable	<i>read</i>	option for obtaining frequencies
FREQUNIT	variable	<i>waven</i>	option for frequency input
HESSFORM	variable	<i>full</i>	form of Hessian matrix
HESSUNIT	variable	<i>bohr</i>	option for Hessian matrix input
ANHARM	variable	<i>none</i>	option for anharmonicity input
CURVATURE	variable	<i>setzero</i>	curvature calculation options
GRADDER	variable	<i>noextra</i>	option for computing the derivative of the gradient
NPTS	variable	2	number of points along the MEP at which data is to be read

Glossary of GEN40 keywords

ANHARM

ANHARM is a variable keyword for selecting anharmonicity input.

<u>Option</u>	<u>Definition</u>
<i>none</i>	No anharmonic data is read (default).
<i>cubic</i>	Read the third derivatives <i>k3</i> with respect to the normal coordinates at each grid point.
<i>quartic</i>	Also read in fourth derivatives <i>k4</i> with respect to the
normal	coordinates at each grid point.

Example:

```
ANHARM cubic
```

CURVATURE

CURVATURE is a variable keyword used to select the way in which information about the curvature components B_{mF} is input. For the *compute* option to be used, *freesource* must be set to *hessian* (and the Hessians must be provided in SADDLE40 and POINT40).

<u>Option</u>	<u>Definition</u>
<i>setzero</i>	The B_{mF} are set to zero everywhere (default).
<i>read</i>	The B_{mF} are read directly.
<i>compute</i>	The B_{mF} are computed from the gradient and Hessian as discussed in Sections 6.H and 10.F.

Example:

```
CURVATURE read
```

FREQSOURCE

FREQSOURCE is a variable keyword for selecting how the frequencies should be determined.

<u>Option</u>	<u>Definition</u>
<i>read</i>	Frequencies are read directly (default).
<i>hessian</i>	Frequencies are obtained by diagonalizing the Hessian or projected Hessian matrix.

Example:

```
FREQSOURCE hessian
```

FREQUNIT

FREQUNIT is a variable keyword that specifies the units of the input frequencies.

<u>Option</u>	<u>Definition</u>
<i>waven</i>	Frequencies are given in cm^{-1} (default)
<i>au</i>	Frequencies are given in atomic units.

Example:

```
FREQUNIT  au
```

Note: all frequencies must be input in the same units (all wavenumbers or all atomic units), including the imaginary frequency, WSTAR.

GEOMUNIT

GEOMUNIT is a variable keyword that specifies the units of the Cartesian or mass-scaled Cartesian coordinates. If one of the mass-scaled options is chosen, then the scaling mass μ must be specified in unit fu5 by the variable SCALEMASS in section PATH. If $\mu = 1$ amu is used for scaling, then SCALEMASS does not need to be specified since 1 amu is the default.

<u>Option</u>	<u>Definition</u>
	coordinates are in:
<i>bohr</i>	unscaled bohrs (default)
<i>msbohr</i>	mass-scaled bohrs
<i>ang</i>	unscaled angstroms
<i>msang</i>	mass-scaled angstroms
<i>si</i>	unscaled meters

Example:

```
GEOMUNIT  ang
```

GRADDER

GRADDER is a variable keyword used to select how the derivative of the gradient is obtained for use in calculation of the curvature components B_{mF} . This keyword is utilized only if CURVATURE is set to *compute*.

<u>Option</u>	<u>Definition</u>
<i>noextra</i>	Derivative of the gradient with respect to s is obtained from a quadratic fit of the gradient at three points on the basic input grid, with the middle point being the current point on the Hessian grid (default). In the notation of Section 9.C, <i>noextra</i> means that the gradient grid is the basic input grid.
<i>oneextra</i>	Derivative of the gradient is computed by one-sided finite difference using the gradient on the basic input grid and a gradient input at one extra point near the basic input grid point. <i>oneextra</i> means that the gradient grid has twice as many points as the basic input grid.
<i>twoextra</i>	Derivative of the gradient is computed from a quadratic fit obtained using the gradient on the basic input grid and gradients input at an extra point near and on each side of the basic input grid point.

Example:

```
GRADDER  oneextra
```

GRADUNIT

GRADUNIT is a variable keyword that specifies the units of the gradients (or, if forces are used, the unit of the force).

<u>Option</u>	<u>Definition</u>
	gradients are in:
<i>bohr</i>	hartrees per unscaled bohr (default)
<i>msbohr</i>	hartrees per mass-scaled bohr
<i>ang</i>	hartrees per unscaled Å
<i>msang</i>	hartrees per mass-scaled Å
<i>si</i>	joules per unscaled meter

Example:

```
GRADUNIT  msbohr
```

HESSFORM

HESSFORM is a variable keyword that, if *hessian* is specified in FREQSOURCE, specifies the form of the Hessian matrix.

<u>Option</u>	<u>Definition</u>
<i>full</i>	F is input as the full matrix (default).
<i>packed</i>	F is input in packed form.

Example:

```
HESSFORM  packed
```

If *packed* is selected, the lower triangular elements of **F** should be listed in order such that $F_{ij} = P(\frac{1}{2}i(i-1) + j)$, where $P(k)$ is element k of the input list and F_{ij} is the element of **F** in row i and column j . The number of elements listed per line is arbitrary. Examples of both input forms for a 6 x 6 matrix is shown below:

Full matrix - example 1

```
F11 F12 F13 F14 F15 F16
F21 F22 F23 F24 F25 F26
F31 F32 F33 F34 F35 F36
F41 F42 F43 F44 F45 F46
F51 F52 F53 F54 F55 F56
F61 F62 F63 F64 F65 F66
```

Full matrix - example 2

```
F11 F12 F13 F14 F15 F16 F21 F22 F23
F24 F25 F26 F31 F32 F33 F34 F35 F36
F41 F42 F43 F44 F45 F46 F51 F52 F53
F54 F55 F56 F61 F62 F63 F64 F65 F66
```


Packed matrix - example 1

```
F11
F21 F22
F31 F32 F33
F41 F42 F43 F44
F51 F52 F53 F54 F55
F61 F62 F63 F64 F65 F66
```

Packed matrix - example 2

```
F11 F21 F22 F31 F32 F33
F41 F42 F43 F44 F51 F52
F53 F54 F55 F61 F62 F63
F64 F65 F66
```

The order of the coordinates is explained in Section 12.E.1.

Note that GAUSSIAN94 writes the packed Hessian matrix in a slightly different way in the standard output file, but the Hessian in the formatted checkpoint file (requested by their FormCheck keyword) is in precisely the packed form accepted by POLYRATE and so is convenient for copying and pasting into POLYRATE fu40 files.

Note that the exact \mathbf{F} matrix is symmetric, but due to finite precision in computations, the actual matrix one has to input is usually nonsymmetric. If one inputs a nonsymmetric full matrix, the code uses the unsymmetric matrix. If a user wants the code to use a symmetric matrix, we recommend taking an arithmetic average of F_{ij} and F_{ji} for the resulting symmetrized elements in packed form.

HESSUNIT

HESSUNIT is a variable keyword that, if *hessian* is specified in FREQSOURCE, specifies the units of the Hessian matrix \mathbf{F} . The choice of coordinates (mass-scaled or not) must be consistent with GEOMUNIT.

<u>Option</u>	<u>Definition</u>
	Hessians are in:
<i>bohr</i>	hartrees per unscaled bohr ² (default)
<i>msbohr</i>	hartrees per mass-scaled bohr ²
<i>ang</i>	hartrees per unscaled Å ²
<i>msang</i>	hartrees per mass-scaled Å ²
<i>si</i>	joules per unscaled meter ²

Example:

```
HESSUNIT msbohr
```

Users calculating Hessians by an electronic structure package should read the manual for that package to see what units the Hessian is in. For example, GAUSSIAN94 prints the Hessian in hartrees per unscaled bohr².

MAXLPTS

MAXLPTS is a variable keyword with an integer argument that specifies the maximum number of points used in Lagrange interpolation in the procedure of Subsection 9.C. The default value for this keyword is 3. If this keyword is set to -1 , the program will perform an IVTST-M calculation using the information read from the fu40 input file. When using file fu40, the user cannot select any options for the IVTST-M calculation; it will be performed according to the default options, and the number of gradient and Hessian points have to be the same and equal to NPTS (see below). Therefore, only IVTST-M-NPTS/NPTS calculations are allowed. The curvature components will be interpolated using a value of 3 for the exponent in the mapping function. For unimolecular reactions or reactions with wells, a cubic polynomial fitting to V is done previous to its spline fitting, taking a step of $0.01 a_0$ along the lowest frequency mode for performing this fitting. The only way to change any of these options or to use a different number of gradient and Hessian points is by using the fu31 input file, which is specifically designed for IVTST-M calculations.

NPTS

NPTS is a variable keyword with an integer argument that specifies the number of points along the MEP (excluding the reactant limit, product limit, points used for extrapolation, and saddle point) at which data is to be read. Unless only TST reaction rates are calculated, file fu40 must contain a minimum of one point on each side of the saddle point. Unless NEXTPT is on, these points must be arranged in order of increasing s . The section POINT40 should occur NPTS times in the file 40 input. The default value for NPTS is 2.

PRINT

PRINT is a variable keyword for selecting print options.

Option	Definition
<i>noextra</i>	Print out titles and options (default).
<i>summary</i>	Also print out summary.
<i>input</i>	Also echo input as it is read.
<i>intermediate</i>	Also print out intermediate information about fitting exponential functions and print out interpolated results each time RPHINT is called.
<i>debugging</i>	Also print out debugging information in exponential fitting.

Example:

```
PRINT summary
```

TITLE

TITLE is a list keyword that allows the user to give a one-line title to the electronic structure file.

12.E.3 R140, R240, P140, P240, WR40, and WP40 sections

The R140, R240, P140, P240, WR40, and WP40 sections all have the same set of keywords. A separate section is needed for each reactant, each product, and each well.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
SVALUE	variable	<i>dgr</i> (or <i>dgp</i>)	value of <i>s</i> (in a.u.) at the reactant (or product)
VVALUE	variable	(required)	potential (in a.u.) at reactant (or product)
HESSIAN	list	(see Glossary)	Hessian matrix for the reactant or product species
IFREQ	list	<i>1 2 3 ...</i>	indices for reordering frequencies, anharmonicities, and curvature components
FREQ	list	(see Glossary)	vibrational frequencies for the reactant or product species
3RDDRIV	list	(see Glossary)	third derivatives of <i>V</i> with respect to motion along the normal coordinates for the reactant or product
4THDERIV	list	(see Glossary)	fourth derivatives of <i>V</i> with respect to motion along the normal coordinates for the reactant or product

Glossary of R140, R240, P140, P240, WR40, and WP40 keywords

3RDDERIV

3RDDERIV is a list keyword that is used to specify the third derivatives of V along the normal coordinates at the reactant, product, or well. If ANHARM is set to *cubic* or *quartic* in the GEN40 section, this keyword is required unless the reactant or product under consideration is monatomic.

4THDERIV

4THDERIV is a list keyword that is used to specify the fourth derivatives of V along the normal coordinates at the reactant, product, or well. If ANHARM is set to *quartic* in the GEN40 section, this keyword is required unless the reactant or product under consideration is monatomic.

FREQ

FREQ is a list keyword that is used to specify the frequencies for each normal mode at the reactant, product, or well species. This keyword is required if FREQSOURCE is set to *read* (the default value) in the GEN40 section and is not used otherwise. The units of the frequencies are determined by the FREQUNIT keyword in the GEN40 section.

HESSIAN

HESSIAN is a list keyword that is used to specify the Hessian matrix at the reactant, product, or well species. This keyword is required if FREQSOURCE is set to *hessian* in the GEN40 section and is not used otherwise. The form and units of the Hessian matrix are determined by the HESSFORM and HESSUNIT keywords in the GEN40 section, respectively.

IFREQ

IFREQ is a required list keyword that is used to specify indices for reordering frequencies, anharmonicities, and curvature components in the interpolation table in order to make the ordering consistent with that at the saddle point and/or along the MEP. The imaginary frequency at the saddle point is not included in this ordering. Therefore, index 1 corresponds to the largest frequency at the saddle point, index 2 corresponds to the second largest frequency, and so on.

SVALUE

SVALUE is a variable keyword that specifies the value of s at the reactant if the reaction is unimolecular or at the well on the reactant side or the value of s at the product if there is only one product or at the well on the product side. The argument of SVALUE can be either a numerical value (in bohr), *dgr* (default guess for reactant or well), or *dgp* (default guess for product or well). SVALUE for reactants has meaning only for a unimolecular reaction, and SVALUE for products has meaning only for a unimolecular rearrangement or an association reaction. SVALUE should be omitted for the reactants if there are two or more reactants, and SVALUE should be omitted for the products if there are two or more products. In the following definitions, s^R is the value of s at the reactant or well on the reactant side, and s^P is the value of s at the product or well on the product side.

<u>Option</u>	<u>Definition</u>
(numerical value)	value of s^R (R140 or WR40 section) or s^P (P140 or WP40 section)
<i>dgr</i>	$s = \max \begin{cases} -2.0a_0 \\ s_1 - (s_2 - s_1) \end{cases}$
<i>dgp</i>	$s = \max \begin{cases} 2.0a_0 \\ s_N + (s_N - s_{N-1}) \end{cases}$

VVALUE

VVALUE is a required variable keyword that specifies the value of the potential (in a.u.) at the reactant or product.

12.E.4 SADDLE40 section

The SADDLE40 section is used to provide information at the saddle point, if it exists. This section should be omitted if there is no saddle point.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
VVALUE	variable	(required)	potential at saddle point
IFREQ	list	1 2 3 ...	indices for reordering frequencies, anharmonicities, and curvature components
HESSIAN	list	(see Glossary)	Hessian matrix for the saddle point
FREQ	list	(see Glossary)	vibrational frequencies for the saddle point
3RDDRIV	list	(see Glossary)	third derivatives of V with respect to motion along the normal coordinates for the saddle point
4THDERIV	list	(see Glossary)	fourth derivatives of V with respect to motion along the normal coordinates for the saddle point
WSTAR	list	(see Glossary)	absolute value of the imaginary frequency at the saddle point

Glossary of SADDLE40 keywords

3RDDERIV

3RDDERIV is a list keyword that is used to specify the third derivatives of V along the bound normal coordinates at the saddle point. This keyword is required if ANHARM is set to *cubic* or *quartic* in the GEN40 section.

4THDERIV

4THDERIV is a list keyword that is used to specify the fourth derivatives of V along the bound normal coordinates at the saddle point. This keyword is required if ANHARM is set to *quartic* in the GEN40 section.

FREQ

FREQ is a list keyword that is used to specify the frequencies for each bound normal mode at the saddle point. This keyword is required if FREQSOURCE is set to *read* (the default value) in the GEN40 section and is not used otherwise. The units of the frequencies are determined by the FREQUIT keyword in the GEN40 section.

HESSIAN

HESSIAN is a list keyword that is used to specify the Hessian matrix at the saddle point. This keyword is required if FREQSOURCE is set to *hessian* in the GEN40 section and is not used otherwise. The form and units of the Hessian are determined by the HESSFORM and HESSUNIT keywords in the GEN40 section, respectively.

IFREQ

IFREQ is a required list keyword that is used to specify indices for reordering frequencies, anharmonicities, and curvature components in the interpolation table for the saddle point.

VVALUE

VVALUE is a required variable keyword that specifies the value of the potential (in a.u.) at the saddle point.

WSTAR

WSTAR is a variable keyword that is used to specify the value of the imaginary frequency at the saddle point. This keyword is required if FREQSOURCE is set to *read* (the default value) in the GEN40 section and is not used otherwise. The default unit for the imaginary frequency is cm^{-1} , but can be changed to atomic units by setting FREQUIT to *au*.

12.E.5 POINT40 section

This section should be repeated for each point along the MEP except for the reactant limit, product limit, points used for extrapolation, and saddle point. The points must be arranged starting from the one with the most negative s value and continuing in the direction of increasing s .

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
SMEP	variable	(required)	value of s at the current grid location along the MEP
VMEP	variable	(required)	potential at the current grid location on the MEP
GEOM	list	(required)	Cartesian coordinates at the current grid point
GRADS	list	(required)	gradient vector at s along MEP
FORCES	list	(required)	force vector at s along MEP
SB1	variable	(see Glossary)	value of s at a location along the MEP near the current grid point
GRADSB1	list	(see Glossary)	gradient vector at the extra point SB1 for computing the curvature B
FORCESB1	list	(see Glossary)	force vector at the extra point SB1 for computing the curvature B
SB2	variable	(see Glossary)	value of s at another location along the MEP near the current grid point
GRADSB2	list	(see Glossary)	gradient vector at the extra point SB2 for computing the curvature B
FORCESB2	list	(see Glossary)	force vector at the extra point SB2 for computing the curvature B
IFREQ	list	1 2 3 . . .	indices for reordering frequencies, anharmonicities, and curvature components

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
HESSIAN	list	(see Glossary)	Hessian matrix at the current grid point
FREQ	list	(see Glossary)	vibrational frequencies at the current grid point
3RDDERIV	list	(see Glossary)	third derivatives of V with respect to motion along the normal coordinates for the current grid point
4THDERIV	list	(see Glossary)	fourth derivatives of V with respect to motion along the normal coordinates for the current grid point
BFS	list	(see Glossary)	reaction path curvature components at the current grid point along the MEP

Glossary of POINT40 keywords

3RDDERIV

3RDDERIV is a list keyword that is used to specify the third derivatives of V along the bound normal coordinates at the current grid point. This keyword is required if ANHARM is set to *cubic* or *quartic* in the GEN40 section.

4THDERIV

4THDERIV is a list keyword that is used to specify the fourth derivatives of V along the bound normal coordinates at the current grid point. This keyword is required if ANHARM is set to *quartic* in the GEN40 section.

BFS

BFS is a list keyword that is used to specify the reaction path curvature components at the current grid point along the MEP. This keyword is required if CURVATURE is set to *read* in the GEN40 section and is not used otherwise.

FORCES

FORCES is a list keyword that is used to specify the forces at the current grid point. This keyword is required if GRADS is not used. The units of the force are determined by the GRADUNIT keyword in the GEN40 section.

FORCESB1

FORCESB1 is a list keyword that is used to specify the force at $s = SB1$. This keyword is required if GRADDER is set to *oneextra* or *twoextra* in the GEN40 section and GRADSB1 is not used; otherwise, it is not used.

FORCESB2

FORCESB2 is a list keyword that is used to specify the force at $s = SB2$. This keyword is required if GRADDER is set to *twoextra* in the GEN40 section and GRADSB2 is not used; otherwise, it is not used.

FREQ

FREQ is a list keyword that is used to specify the frequencies for each bound generalized normal mode at the current grid point. This keyword is required if FREQSOURCE is set to *read* (the default value) in the GEN40 section and is not used otherwise. The units of the frequencies are determined by the FREQUNIT keyword in the GEN40 section.

GEOM

GEOM is a list keyword that specifies the Cartesian coordinates at the current grid location, in units determined by the GEOMUNIT keyword in the GEN40 section.

GRADS

GRADS is a list keyword that is used to specify the gradient at the current grid point. This keyword is required if FORCES is not used. The units of the gradient are determined by the GRADUNIT keyword in the GEN40 section.

GRADSB1

GRADSB1 is a list keyword that is used to specify the gradient at $s = SB1$. This keyword is required if GRADDER is set to *oneextra* or *twoextra* in the GEN40 section and FORCESB1 is not used; otherwise, it is not used.

GRADSB2

GRADSB2 is a list keyword that is used to specify the gradient at $s = SB2$. This keyword is required if GRADDER is set to *twoextra* in the GEN40 section and FORCESB2 is not used; otherwise, it is not used.

HESSIAN

HESSIAN is a list keyword that is used to specify the Hessian matrix at the current grid point. This keyword is required if FREQSOURCE is set to *hessian* in the GEN40 section and is not used otherwise. The form and units of the Hessian are determined by the HESSFORM and HESSUNIT keywords in the GEN40 section, respectively.

IFREQ

IFREQ is a list keyword that is used to specify indices for reordering frequencies, anharmonicities, and curvature components in the interpolation table for the current grid point.

SMEP

SMEP is a required variable keyword that specifies the value of s (in bohr) at the current grid location along the MEP.

SB1

SB1 is a variable keyword that specifies the value of s (in bohr) at a grid location along the MEP close to SMEP. This keyword is required if GRADDER is set to *oneextra* or *twoextra* in the GEN40 section and is not used otherwise.

SB2

SB2 is a variable keyword that specifies the value of s (in bohr) at a grid location along the MEP close to SMEP, but on the opposite side of SMEP from SB1. This keyword is required if GRADDER is set to *twoextra* in the GEN40 section and is not used otherwise.

VMEP

VMEP is a required variable keyword that specifies the value of the potential at the current grid location.

12.F. General description of unit fu50 input

This file is only needed if the DL keyword is set to *ioc* in the GENERAL section of the unit fu5 input. This allows one to calculate zero-order interpolated VTST or interpolated optimized corrections. The input from FORTRAN unit fu50 is in free format keyword style. In the current version of *POLYRATE* there is only one section, VTSTIC, in the unit fu50 input file, and the first keyword in this file must be *VTSTIC. A sample unit fu50 input file is cmcic.dat for the cmctr2 test run. A description of the input to the *POLYRATE* code from FORTRAN unit fu50 follows.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
OIVTST	switch	off	zero-order IVTST calculation
(NO)DEMI	list	no correction	corrected det I
ENERXN	variable	required	corrected exoergicity
ENESAD	variable	required	corrected classical barrier height
(NO)FREQMAT	list	no matching	frequency matching information
(NO)HRMI	variable	99	correct the moment of inertia
FREQIMAG	variable	required	imaginary freq. at saddle point
LOWFREQ	list	no correction	IVTST treatment of low freq.
MEPTYPEP	variable	two	type of reactant-side MEP
MEPTYPEP	variable	two	type of product-side MEP
MIRP	list	not used	corrected moments of inertia for the reactant and product
MITS	list	not used	corrected moments of inertia for the transition state
LCTCOR	variable	no	correction of <i>V</i> in the nonadiabatic region in LCT calculations

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
PCFREQA	list	not used	corrected frequencies of the product-side stationary point
PCFREQS	list	not used	lower-level frequencies of the product-side stationary point
PCINFO	list	not used	product-side stationary point info.
SPC	variable	not used	<i>s</i> value at this point
ENEPCS	variable	not used	lower-level <i>V</i> at this point
ENEPCA	variable	not used	corrected <i>V</i> at this point
(NO)PRODFREQ	list	not used	corrected product frequencies
RCFREQA	list	not used	corrected frequencies of the reactant-side stationary point
RCFREQS	list	not used	lower-level frequencies of the reactant-side stationary point
RCINFO	list	not used	reactant-side stationary point info.
SRC	variable	not used	<i>s</i> value at this point
ENERCS	variable	not used	lower-level <i>V</i> at this point
ENERCA	variable	not used	corrected <i>V</i> at this point
RANGEIC	list	not used	range parameters
RPL	variable	1.0	range parameter L for an Eckart function
BR	variable	5.0	range parameter for a cut-off Gaussian on the reactant side
BP	variable	5.0	range parameter for a cut-off Gaussian on the product side
REACFREQ	list	not used	corrected reactant frequencies
SADFREQ	list	not used	corrected saddle point frequencies
UNITMI	variable	au	units used for det I input

Glossary for file fu50 keywords

OIVTST

OIVTST is a switch keyword that specifies that the zero-order IVTST calculation will be performed. The default is off.

DETM I (NODETM I)

DETM I is a list keyword used to input the corrected determinant of the moment of inertia tensor. Five values of det **I** (which is the determinant of the inertia tensor, also equal to the product of the three principal moments of inertia) have to be input. They are, in the order of input, for the first reactant species, the second reactant species, the saddle point, the first product species, and the second product species. The units used are specified by the UNITM I keyword, which is discussed latter in this section. For an atomic species, the user should input a value of zero for det **I**. The default is NODETM I, i.e., no correction for det **I**.

Example:

```
DETM I
  5.7220E16
  4.9906E13
  9.8871E17
  5.9964E16
  2.3769E13
END
```

ENERXN

ENERXN is a variable keyword used to input the corrected energy of the reaction in kcal/mol, excluding zero point contributions. This is a required keyword. It should be positive if the reaction is endoergic and negative if it is exoergic.

Example:

```
ENERXN      -5.0
```

ENESAD

ENESAD is a variable keyword used to input the corrected classical barrier height in kcal/mol. This is a required keyword.

Example:

```
ENESAD      6.0
```

FREQMAT (NOFREQMAT)

FREQMAT is a list keyword used to specify the frequency matching information. The input consists of n lines where n is the number of real saddle point frequencies. Each line needs three numbers; the first one is the mode number of a mode at the saddle point, and the second and the third ones are the mode numbers of the modes at the reactant-side and product-side stationary points which match that saddle point mode. The default is NOFREQMAT, i.e., the vibrational frequencies are automatically matched in decreasing order. However, if the keywords MEPTYPEP and MEPTYPEP, which are discussed later in this section, are not both equal to *two*, this keyword is required.

Example:

```
FREQMAT
  1      2      1
  2      3      3
  3      4      2
  4      1      4
  5      5      5
  6      6      6
  7      8      7
  8      7      8
  9      9      9
END
```

HRMI (NOHRMI)

HRMI is a variable keyword that specifies the correction method for the moment of inertia of hindered-rotor modes. The default is NOHRMI. The valid arguments for the HRMI keyword are:

<u>Option</u>	<u>Description</u>
<i>all</i>	correct the moments of inertia of all the hindered-rotor modes of the reactants, products, and saddle point.
$-n (n=1, 2, \dots)$	same as above, and set the moments of inertia of the lowest n GTS hindered-rotor modes for all values of s to their values at the saddle point.
$n (n=1, 2, \dots)$	no correction to the moment of inertia at stationary points, but set the moments of inertia of the lowest n GTS hindered-rotor modes for all values of s to their lower-level values at the saddle point.

Example:

```
HRMI -1
```

FREQIMAG

FREQIMAG is a variable keyword used to input the absolute value of the corrected imaginary frequency at saddle point in cm^{-1} . This is a required keyword.

Example:

```
FREQIMAG 900.0
```


LCTCOR

LCTCOR is a variable keyword that specifies the method to correct the potential energy in the nonadiabatic region in LCT calculations. This keyword has an effect only when LCT calculations are performed. The default is *quadlin*, which provides the standard version of the VTST-IC method as explained in Section 3.1 of Ref. 36 of Sect. 23. The valid arguments for this keyword are:

<u>Option</u>	<u>Description</u>
<i>quadlin</i>	linear or quadratic correction depending on the magnitude of the corrections in different regions
<i>linear</i>	linear correction only
<i>nocor</i>	no correction

Example:

```
LCTCOR      nocor
```

Note: The user should take special care to make sure that this correction yields physical results.

LOWFREQ

LOWFREQ is a list keyword used to specify that certain GTS mode frequencies are to be interpolated using the IVTST-0 formula based on their values at three points. The functional forms used to interpolate the specified frequencies are the same as the IVTST-0 functions as explained in Section 4.1 of Ref. 36 of Sect. 23. Notice that this is similar to the IVTST0FREQ keyword of the PATH section of fu5. IVTST0FREQ is used for single-level calculations and lower levels of dual-level calculations, whereas LOWFREQ is used for the higher level of dual-level runs. If, for a given mode, IVTST0FREQ and LOWFREQ are both specified, then LOWFREQ overrides IVTST0FREQ.

Example:

```
LOWFREQ
  11
  12
  13
  14
END
```

MEPTYPER

MEPTYPER is a variable keyword that specifies the type of reactant-side MEP that will be considered in the VTST-IOC calculation. The default value is *two*. The valid arguments for this keyword are:

<u>Option</u>	<u>Description</u>
<i>one</i>	There is only one reactant species and interpolation is carried out between this species and the saddle point.
<i>two</i>	There are two reactant species and interpolation is carried out between these species (using the merged set of frequencies) and the saddle point.
<i>well</i>	The correction on the reactant side is based on a well.

Example:

```
MEPTYPER one
```

MEPTYPEP

MEPTYPEP is a variable keyword that specifies the type of product-side MEP that will be considered in the VTST-IOC calculation. The default is *two*. The valid arguments for this keyword are:

<u>Option</u>	<u>Description</u>
<i>one</i>	There is only one product species and interpolation is carried out between this species and the saddle point.
<i>two</i>	There are two product species and interpolation is carried out between these species (using the merged set of frequencies) and the saddle point.
<i>well</i>	The correction on the product side is based on a well.

Example:

```
MEPTYPEP well
```

MIRP

MIRP is a list keyword used to input the corrected moments of inertia in atomic units of the reactant and product hindered-rotor modes. This moment of inertia is for the internal rotation, which is not the same as the moment of inertia of the overall system. This keyword is needed if HRMI is set to *all* or is less than zero. The input order is reactant 1, reactant 2 (if any), product 1, and product 2 (if any); and within each reactant and product species the modes should be ordered from the highest frequency hindered-rotor mode to the lowest frequency hindered-rotor mode. This keyword can be omitted if there are no hindered-rotor modes in reactants and products.

Example:

```
MIRP
      960.50
END
```

MITS

MITS is a list keyword used to input the corrected moments of inertia in atomic units of the saddle point hindered-rotor modes. The moment of inertia is for the internal rotation, and it is not the same as the moment of inertia of the overall system. This keyword is needed if HRMI is set to *all* or is less than zero. The input should be ordered from the highest frequency hindered-rotor mode to the lowest frequency hindered-rotor mode. This keyword can be omitted if there are no hindered-rotor modes at the saddle point. In the example below, which is for the CH₄ + OH reaction, the symmetry number for the internal rotation of the CH₃ group around the C–H axis in the TS is 3:

Example:

```
MITS
      7690
END
```

PCFREQA

PCFREQA is a list keyword used to input the corrected frequencies (in cm⁻¹) of the product-side stationary point used for the IOC correction. If the number of real frequencies is less than that of the saddle point (i.e., when MEPTYPEP is *two*), add zeros after the last non-zero frequency to input a total of *n* frequencies, where *n* is the number of real frequencies at the saddle point. There should be *n* + 1 input frequencies if MEPTYPEP is not *two*. The order of the frequencies in the input must match the order of the lower-level frequencies, i.e., the first frequency must correspond physically to the highest lower-level frequency, and the second frequency must correspond physically to the second highest lower-level frequency, and so on.

Example:

```
PCFREQA
  3124.0    2282.0    2282.0    2279.0    1489.0
  1489.0    1287.0    1287.0    1111.0    1072.0
  1033.0    1033.0     565.0     415.0     411.0
    0.0         0.0         0.0         0.0         0.0
END
```

(In this example MEPTYPEP is *two* and there are nine atoms in the system. The first product has four atoms and the second product has five atoms. The saddle point has twenty real frequencies but the products have only fifteen real frequencies, so we add five zeros at the end of input.)

PCFREQS

PCFREQS is a list keyword used to input the lower-level frequencies (in cm^{-1}) of the product-side stationary point. This keyword is only needed when MEPTYPEP is *well*. Otherwise the frequencies are just the product frequencies, and the program will have this information when it calculates the product properties. There should be $n+1$ input frequencies, where n is the number of the real frequencies at the saddle point. The frequencies in the input must be in decreasing order.

Example:

```
PCFREQS
 3124.0    2282.0    2282.0    2279.0    1489.0
 1489.0    1287.0    1287.0    1111.0    1072.0
 1033.0    1033.0     565.0     415.0     411.0
END
```

PCINFO

PCINFO is a list keyword used to specify the product-side stationary point information. This keyword is needed only if MEPTYPEP is *one* or *well*. The valid subkeyword options are defined below. There are no defaults for these subkeywords, and the user must supply all the values, except as noted.

<u>Option</u>	<u>Description</u>
<i>spc</i>	s value in bohrs at this point
<i>enepcs</i>	lower-level potential energy in kcal/mol at this point (required only if MEPTYPEP is <i>well</i>)
<i>enepca</i>	corrected potential energy in kcal/mol at this point (required only if MEPTYPEP is <i>well</i>)
<i>nofreqcorr</i>	no frequency correction at the product complex; when this keyword is used the keywords PCFREQA and PCFREQS are not needed

Example:

```
PCINFO
  SPC          3.0
  ENEPCS      -10.0
  ENEPCA       -9.0
END
```

PRODFREQ (NOPRODFREQ)

PRODFREQ is a list keyword used to input the corrected product frequencies in cm^{-1} . It is only needed when MEPTYPEP is *well*; otherwise the corrected product frequencies are input in the PCFREQA keyword. NOPRODFREQ specifies that no correction is needed for the product frequencies, but this is allowed only if MEPTYPEP is *well*. The default is to correct the product frequencies when MEPTYPEP is *well*, so if NOPRODFREQ is not specified, the PRODFREQ keyword is required. See keyword PCFREQA for the required input order for the frequencies.

Example:

```
PRODFREQ
  3092.0    3092.0    2981.0    1453.0    1453.0
  1377.0    1017.0    1017.0     741.0
END
```

RCFREQA

RCFREQA is a list keyword used to input the corrected frequencies (in cm^{-1}) of the reactant-side stationary point that the IOC correction is based on. If the number of real frequencies is less than that of the saddle point (i.e., when MEPTYPEP is *two*), add zeros after the last non-zero frequency to input a total of n frequencies, where n is the number of real frequencies at the saddle point. There should be $n+1$ input frequencies if MEPTYPEP is not *two*. The order of the frequencies in the input must match the order of the lower-level frequencies, i.e., the first frequency must correspond physically to the highest lower-level frequency, and the second frequency must correspond physically to the second highest lower-level frequency, and so on.

Example (see the example for the PCFREQA keyword):

```
RCFREQA
  3124.0    2282.0    2282.0    2279.0    1489.0
  1489.0    1287.0    1287.0    1111.0    1072.0
  1033.0    1033.0     565.0     415.0     411.0
  0.0        0.0        0.0        0.0        0.0
END
```

RCFREQS

RCFREQS is a list keyword used to input the flower-level frequencies (in cm^{-1}) of the reactant-side stationary point. This keyword is only needed when MEPTYPEP is *well*. Otherwise the frequencies are just the reactant frequencies, and the program will have this information when it calculates the reactant properties. There should be $n + 1$ input frequencies, where n is the number of the real frequencies at the saddle point. The frequencies in the input must be in decreasing order.

Example:

```
RCFREQS
  3124.0    2282.0    2282.0    2279.0    1489.0
  1489.0    1287.0    1287.0    1111.0    1072.0
  1033.0    1033.0     565.0     415.0     411.0
END
```

RCINFO

RCINFO is a list keyword used to specify the reactant-side stationary point information. This keyword is needed only if MEPTYPE is *one* or *well*. The valid subkeyword options are defined below. There are no defaults for these subkeywords, and the user must supply all the values, except as noted.

<u>Option</u>	<u>Description</u>
<i>src</i>	<i>s</i> value in bohrs at this point
<i>enercs</i>	lower-level potential energy in kcal/mol at this point (required only if MEPTYPE is <i>well</i>)
<i>enerca</i>	corrected potential energy in kcal/mol at this point (required only if MEPTYPE is <i>well</i>)
<i>nofreqcorr</i>	no frequency correction at the reactant complex; when this keyword is used the keywords RCFREQA and RCFREQS are not needed

Example:

```
RCINFO
  SRC      -3.0
  ENERCS   -5.0
  ENERCA   -4.0
END
```

RANGEIC

RANGEIC is a list keyword used to input the range parameters for the correction functions in the VTST-IOC calculations. The valid subkeyword options are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>rpl</i>	range parameter L in bohrs for an Eckart function	1.0
<i>br</i>	range parameter for a cut-off Gaussian function on the reactant side	5.0
<i>bp</i>	range parameter for a cut-off Gaussian function on the product side	5.0

Example:

```
RANGEIC
  RPL      0.3
  BR      10.0
  BP      8.0
END
```

REACFREQ

REACFREQ is a list keyword used to input the corrected reactant frequencies in cm^{-1} . It is required if MEPTYPE is *well*. Otherwise the reactant frequencies are input in RCFREQA keyword. See keyword RCFREQA for the requirement on the input order of the frequencies.

Example:

```
REACFREQ
  3092.0    3092.0    2981.0    1453.0    1453.0
  1377.0    1017.0    1017.0     741.0
END
```

SADFREQ

SADFREQ is a list keyword used to input the corrected real saddle point frequencies in cm^{-1} . It is a required keyword. See keyword RCFREQA for the required input order for the frequencies.

Example:

```
SADFREQ
  3748.0    3130.0    3259.0    3255.0    1414.0
  1521.0    1494.0    1298.0    1206.0     933.0
  769.0     357.0     316.0     40.0
END
```

UNITMI

UNITMI is a variable keyword that specifies the units used for the DETMI keyword input. The default is *au*. The valid arguments are:

<u>Option</u>	<u>Description</u>
<i>au</i>	atomic units
<i>si</i>	SI units, i.e., kg m^2 or $(\text{kg m}^2)^3$
<i>cgs</i>	c.g.s. units, i.e., g cm^2 or $(\text{g cm}^2)^3$
<i>scale</i>	scaled units, i.e., 10^{-40}g cm^2 or $(10^{-40} \text{g cm}^2)^3$
<i>amu</i>	atomic mass unit, i.e., a.m.u. bohr ² or (a.m.u. bohr ²) ³ .

Example:

```
UNITMI    AMU
```

12.G. General description of unit fu51 input

This file is only needed if the *ispe* option is specified with the DL keyword in the GENERAL section of unit fu5. The single-point energy correction (VTST-ISPE) is used to correct the energies along the reaction path. The input from FORTRAN unit fu51 is in free format keyword style. A description of the input to the *POLYRATE* code from FORTRAN unit fu51 follows.

12.G.1. Section Descriptions

<u>Section Name</u>	<u>Description</u>
*ISPEGEN	general section for stationary point information
*POINT	section for the extra points along the reaction path

The *POINT section should be repeated once for each of the points along the reaction path excluding the reactants, products, saddle point, and wells.

12.G.2 ISPEGEN section

The *ISPEGEN section is used to provide information at the stationary points.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
EXERXN	variable	required	corrected exoergicity
ENESAD	variable	required	corrected classical barrier height
MEPTYPEP	variable	<i>two</i>	type of reactant-side MEP
MEPTYPEP	variable	<i>two</i>	type of product-side MEP
PCINFO	list	not used	product-side stationary point data
SPC	variable	not used	value of s at this point
ENEPCS	variable	not used	lower-level V at this point
ENERCA	variable	not used	corrected V at this point
RCINFO	list	not used	reactant-side stationary point data
SRC	variable	not used	value of s at this point
ENERCS	variable	not used	lower-level V at this point
ENERCA	variable	not used	corrected V at this point

Glossary of ISPEGEN keywords

EXERXN

EXERXN is a variable keyword used to input the corrected classical energy change of the reaction in kcal/mol, excluding zero-point contributions. This is a required keyword. It should be positive if the reaction is endoergic and negative if it is exoergic. In other words, this is the corrected product energy relative to the reactant energy.

Example:

```
EXERXN          -5.0
```

ENESAD

ENESAD is a variable keyword used to input the corrected classical barrier height in kcal/mol. This is a required keyword.

Example:

```
ENESAD          6.0
```

MEPTYPEP

MEPTYPEP is a variable keyword that specifies the type of product-side MEP that will be considered in the VTST-ISPE calculation. The default is *two*. The valid arguments for this keyword are:

<u>Option</u>	<u>Description</u>
<i>one</i>	There is one product and wells, if any, are not recognized.
<i>two</i>	There are two products and wells, if any, are not recognized.
<i>well</i>	A product well is recognized.

Example:

```
MEPTYPEP  one
```

MEPTYPER

MEPTYPER is a variable keyword that specifies the type of reactant-side MEP that will be considered in the VTST-ISPE calculation. The default is *two*. The valid arguments for this keyword are:

<u>Option</u>	<u>Description</u>
<i>one</i>	There is one reactant and wells, if any, are not recognized.
<i>two</i>	There are two reactants and wells, if any, are not recognized.
<i>well</i>	A reactant well is recognized.

Example:

```
MEPTYPER  one
```

PCINFO

PCINFO is a list keyword used to specify the product-side stationary point information. This keyword is needed only if MEPTYPEP is set to *one* or *well*. There are no defaults for these options, the user must supply all values except as otherwise noted. The valid options for this keyword are:

<u>Option</u>	<u>Description</u>
<i>spc</i>	value of <i>s</i> , in bohrs, at this point
<i>enepcs</i>	lower-level potential energy, in kcal/mol, at this point with respect to the reactant (required if and only if MEPTYPEP is set to <i>well</i>)
<i>enepca</i>	corrected potential energy, in kcal/mol, at this point with respect to the reactant (required if and only if MEPTYPEP is set to <i>well</i>)

Example:

```
PCINFO
      spc          3.0
      enepcs     -10.0
      enepca     -9.0
```

RCINFO

RCINFO is a list keyword used to specify the reactant-side stationary point information. This keyword is needed only if MEPTYPEP is set to *one* or *well*. There are no defaults for these options, the user must supply all values except as otherwise noted. The valid options for this keyword are:

<u>Option</u>	<u>Description</u>
<i>src</i>	value of <i>s</i> , in bohrs, at this point
<i>enercs</i>	lower-level potential energy, in kcal/mol, at this point with respect to the reactant (required if and only if MEPTYPEP is set to <i>well</i>)
<i>enerca</i>	corrected potential energy, in kcal/mol, at this point with respect to the reactant (required if and only if MEPTYPEP is set to <i>well</i>)

Example:

```
PCINFO
      src          -3.0
      enercs     -5.0
      enerca     -4.0
```

12.G.3 POINT section

The POINT section should be repeated for each point along the MEP, except for the reactants, products, saddle point, and wells. Each new POINT section must begin with *POINT. The points can be arranged in any order.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
DVMEP	variable	variable	difference in the corrected energy at point
SMEP	variable	required	value of s at point
VMEP	variable	variable	corrected energy at point

Notice that in the POINT section, the keyword SMEP is required and either VMEP or DVMEP is needed for indicating the energy at a higher-level or difference of the energy at a higher-level to the lower-level one.

Glossary of POINT keywords

DVMEP

DVMEP is a variable keyword that can be used to specify the energy difference between the higher-level along the MEP from the lower-level ones. Using the VMEP keyword to indicate the higher-level potential energies along the lower-level MEP has a limitation, namely that the point chosen has to be a save-grid point of the lower-level path. However, using the DVMEP keyword, the point chosen does not have to be a save-grid point of the lower-level path because the difference in potential energies from the higher-level to the lower-level is read instead of calculated.

Example:

```
VMEP      5.1608
```

SMEP

SMEP is a variable keyword that specifies the value of s , in bohrs, at the current point along the MEP.

Example:

```
SMEP      -0.8
```

VMEP

VMEP is a variable keyword that specifies the energy at the current point along the MEP.

Example:

```
VMEP      10.5
```

13. DESCRIPTION OF INPUT FOR VRC-VTST

In VRC-VTST rate calculations, input file fu5 is the only one required; the other input files used in RP-VTST calculations are not needed.

13.A Section descriptions of input file fu5

The following input sections correspond to the major parts of a VRC-VTST rate calculation. Each section begins with a line giving the section name, preceded by an “*”. Keywords for each section are discussed further in the rest of this subsection. In each section, we list the keywords used in VRC-VTST rate calculations. The keywords that are only used for VRC-VTST are discussed in this section; the keywords that can be used for both RP-VTST and VRC-VTST are not discussed again and users can find the details in Section 12.

<u>Section Name</u>	<u>Description</u>
* GENERAL	general data such as the title and defining the atoms involved
* ENERGETICS	options for computing potential energy
* REACT1	definition and properties of the first reactant
* REACT2	definition and properties of the second reactant
* START	Only used to define the electronic degeneracy for generalized transition state
* PROD1	definition and properties of the first product
* PATH	definition of reaction coordinate range and step size
* RATE	options for computing the reaction rate

13.A.1. GENERAL section

The table below lists all the valid keywords for the GENERAL section. Indented keywords are level-2 keywords, and the others are level-1 keywords. Level-2 keywords are always associated with a level-1 list keyword and must come after that keyword and before any

other level-1 keyword or starred header. In the alphabetical glossary that follows the box that are only used for VRC-VTST rate calculations. In any run for VRC-VTST this section would have the TITLE, ATOMS, and VRC keywords.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
TITLE	list	blank	title with 5 lines maximum
ATOMS	list	required	atoms in system
CHECK	switch	off	check the input data
INPUNIT	variable	angstrom	specifies the units in fu5 input file
OUTUNIT	variable	angstrom	specifies the units in fu6 and fu14 output files
SUPERMOL	switch	on	supermolecule mode for reactants and products
VRC	switch	Off	use variable reaction coordinate to treat association reaction with loose transition state
VRCOPT	list	not used	variable reaction coordinate algorithm options
JMAX	variable	300	maximum angular momentum quantum number J in $E, J - \mu VT$
JSTEP	variable	5	step size in the integral over J in $E, J - \mu VT$
GTYPE	variable	4	type of random number generator in SPRNG
NITER	variable	4	Gauss-Laguerre quadrature
NMC	variable	1000	number of Monte Carlo samples per facet in multifaceted dividing surface

Glossary of GENERAL keywords only used for VRC-VTST

VRC

VRC is a switch keyword that allows the user to employ the variable reaction coordinate algorithm to treat loose transition states. When this keyword is turned on, the PROD1, PROD2, WELLR, WELLP, and START sections are not used in fu5 input file. Only forward reaction rates will be calculated in this case. If each reactant has only one pivot point, a spherical (i.e. single-faceted) dividing surface will be used. If more than one pivot point is assigned to one or two of the reactants, a multifaceted dividing surface will be used. Any keywords related the MEP algorithm are not used when VRC is turned on.

VRCOPT

VRCOPT is a list keyword that allows the user control over the parameters used in variable reaction coordinate algorithm. If VRCOPT is given then it is assumed that the VRC calculation is desired and the default for VRC becomes on. The valid *options* are defined below.

<u>Option</u>	<u>Description</u>	<u>Default</u>
<i>jmax</i>	maximum angular momentum quantum number J calculated for the number of accessible state $N(E, J)$ in $E, J - \mu VT$	300
<i>jstep</i>	step size of angular momentum quantum number J in the integral over J to calculate the contribution to reactive flux from transitional modes	5
<i>gtype</i>	the type of random number generator in SPRNG package. Type 4 is Multiplicative Lagged Fibonacci Generator in SPRNG version 2.0. It is not recommended to change the default value. User needs to refer SPRNG manual to change the value to the desired type.	4
<i>niter</i>	number of Gauss-Laguerre quadrature points for total energy E .	4
<i>nmc</i>	number of Monte Carlo sampling for each facet in a multifaceted dividing surface. In parallel calculation, the actual number used in the calculation may be changed according to the number of processors used. If the remainder when NMC is divided by number of processors is not zero or NMC is smaller than number of processors, the number of sampling for each processor will be $\text{int}(\text{NMC}/\text{number of processors}) + 1$. User need to be careful to balance the nmc and number of processors used in parallel calculation.	1000

Example:

```
VRCOPT
  NMC      600
  JMAX     500
  JSTEP    10
END
```


13.A.2. ENERGETICS section

The ENERGETICS section is used to define the method for computing the potential energy of the reacting system. The table below lists all the valid keywords for the ENERGETICS section. In the alphabetical glossary a more complete explanation for each keyword is given.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
EZERO	variable	<i>calculate</i>	method for determining zero of E
EZUNIT	variable	<i>a.u.</i>	units of zero of energy
POTENTIAL	variable	<i>hooks</i>	type of PES to be used

Glossary of ENERGETICS keywords

EZERO

Refer to Section 12.A.3

EZUNIT

Refer to Section 12.A.3

POTENTIAL

POTENTIAL is a variable keyword that allows the user to specify the source of the potential energy information for a single-level calculation. There are only one valid option for a VRC-VTST calculation, namely *hooks*. The option *hooks* means that the user is providing potential information through the hooks. When *Polyrate* is used as a stand-alone package, the hooks call SURF and SETUP; this is how one supplies an analytic potential energy surface. When *Polyrate* is used in conjunction with an interface program, e.g. GAUSSRATE, the hooks call the corresponding electronic structure program to perform electronic structure calculations to obtain potential energy.

<u>Option</u>	<u>Definition</u>
<i>hooks</i>	use the hooks

Example:

POTENTIAL *hooks*

13.A.3. OPTIMIZATION section

The OPTIMIZATION section is used to specify the method for carrying out geometry optimizations when the POTENTIAL keyword selection is *hooks*. In principle, the keywords used for RP-VTST can also be used for VRC-VTST to optimize reactants. But these keywords have not been tested for VRC-VTST calculations. Therefore it is strongly recommended to input the optimized geometries instead of using *POLYRATE* to optimize the reactant geometries because a semi-global analytical potential energy surface when using *POLYRATE* as a stand-alone package is required in VRC-VTST calculation and it usually doesn't have analytical derivatives used for geometry optimization.

13.A.4. REACT1, REACT2, PROD1, AND START sections

The summary table below gives the keywords that can be used for a VRC-VTST calculations in REACT1, REACT2, PROD1, and START sections. Note that in VRC-VTST calculations an additional keyword is introduced for setting the Cartesian coordinates of pivot points used for the variable reaction coordinate. These pivot points are only required in the REACT1 and REACT2 sections.

In START section, only the ELEC keyword is used. This gives electronic degeneracy of the generalized transition state. No other keyword is required in this section when *POLYRATE* runs as a stand-alone program. When running in direct dynamics with other rate program, e.g. GAUSSRATE, the STATUS keyword should be set to 6 to avoid any electronic structure calculation for START section.

Basic:

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
<i>Required:</i>			
GEOM	list	required	Cartesian coordinates of atoms
PIVOT	list	Required only for reactants	Cartesian coordinates of pivot points for reactants
<i>Common:</i>			
ELEC	list	all degen. = 1	electronic degeneracies
ENERGY	variable	none	energy of the stationary point
FREQ	switch	on	perform normal-mode analysis
FREQUNIT	variable	waven	units of the frequencies
INITGEO	variable	geom	location of input data that specify initial geometries
STATUS	variable	0	status of the stationary point work
SPECIES	variable	nonlinrp	species type (linearity)
VIB	list	none	frequencies of the stationary points

Glossary of REACT1 and REACT2 keywords

PIVOT

PIVOT is a list keyword that is used to define the locations of pivot points. The Cartesian coordinates of pivot points must be given along with their identification number. The identification number can be the same for pivot points on the different reactants. The origin of the Cartesian coordinates must be the same as the origin of the corresponding reactant Cartesian coordinates, so that they can give the positions relative to the corresponding reactant.

Example:

```
PIVOT
  1   0.000000   0.000000   0.200000
  2   0.000000   0.000000  -0.200000
END
```

13.A.5. PATH section

The PATH section contains the keywords that control the calculation of the reaction path. In VRC-VTST rate calculations, only one keyword, SVRC, is allowed in this section.

SVRC

SVRC is a list keyword that specifies the reaction coordinate s range and step size used in variable reaction coordinate and single-faceted or multifaceted dividing surface algorithm for loose transition state. If VRC keyword in GENERAL section is turned on, user needs to specify this keyword only in the PATH section, and the other keywords in the PATH section are not required.

Example:

```
SVRC  
  2.0  4.0  0.1  
END
```

This example tells the program that the s range is 2.0 – 4.0 Å (note that default unit is Å) and that the step size is 0.1 Å.

13.A.6. RATE section

The RATE section contains the keywords that control the rate constant calculations. Note that a conventional TST calculation is not a valid option for VRC-VTST rate since there is no barrier on potential energy surface. Only forward rate constants (bimolecular association rate) are calculated for VRC-VTS and the reverse unimolecular dissociation rate calculation is not implemented yet. The VRC-VTST rate constants are calculated by CVT, μ VT, and $E, J - \mu$ VT. An alphabetical glossary of all the keywords follows the summary table.

<u>Keyword</u>	<u>Type</u>	<u>Default</u>	<u>Description</u>
SIGMAF	variable	1.0	sym. factor for forward reaction
SIGMAR	Variable	1.0	sym. Factor for reversed reaction
TEMP	list	see below	temperatures for rate calculations

SIGMAF

SIGMAF is a variable keyword whose argument specifies the number of equivalent ways for forming the GTS for the forward reaction. Note that only two values are used in VRC-VTST calculations. The value should be 1 for the case where the two reactants are different, and the value should be 0.5 for the case two reactants are identical.

SIGMAR

SIGMAF is a variable keyword whose argument specifies the number of equivalent ways for forming the GTS for the reversed reaction.

14. GENERAL DESCRIPTION OF LCT FILES

In the calculation of LCT probabilities, relevant quantities such as theta integrals (i.e., imaginary action integrals) are evaluated on the grid defined by the input parameters NQTH and NGTHETA. The stored information is then used for interpolation. When extra output of the LCT calculations is chosen (LCTDETAIL keyword), the details of the calculation of these quantities at certain grid points, which are specified by user, are given in files poly.fu41 through poly.fu47. The contents of each output file are listed in the table below. In the table, 'each tunneling path' refers to a tunneling path corresponding to an energy quadrature point, IE, that is within the range specified in the keyword LCTDETAIL in the TUNNEL section of unit fu5 input.

File name	Fortran Unit	Contents
poly.fu41	fu41	the theta integrand and the weighting factor at each point on each tunneling path, within its adiabatic region.
poly.fu42	fu42	the theta integrand, the effective potential, the classical potential, and the potential correction terms, for each point on each tunneling path, in its non-adiabatic region
poly.fu43	fu43	the theta integrals and the boundaries of the nonadiabatic region of each tunneling path
poly.fu44	fu44	the length of the tunneling path
poly.fu45	fu45	the corresponding s value on the MEP for each point on each tunneling path
poly.fu46	fu46	the adiabatic potential and the cosine of the angle between the tunneling path and the gradient vector of the MEP at the s value to which the point on the tunneling path corresponds for each point on each tunneling path
poly.fu47	fu47	the vibrational turning points, the projection onto each normal mode coordinate, and the curvature components for each point on each tunneling path

Part V More General Program Package Information

15. INSTALLING AND USING *POLYRATE*

A step-by-step procedure for installing, testing, and using *POLYRATE* is given in this section. The file structure and the Perl and C Shell scripts should make *POLYRATE* reasonably easy to install, test, and use.

Subsection 15.A discusses running *POLYRATE* in a Unix environment, including Linux. Subsection 15.B contains information for users who do not wish to use the provided Unix scripts or who wish to run *POLYRATE* in a non-Unix environment.

15.A. Instructions for a Unix environment

If the *POLYRATE* program has been received in tar.gz format, the recommended directory structure will be created automatically when this file is uncompressed and untarred, and the files will be put in their canonical places as listed in Subsection 15.A.1.

15.A.1. Installation

Before you start, make sure all the file names are in lower case. For the purpose of illustration, we use upper case for sample directory names, e.g., PDIR. Here PDIR is the directory above the *Polyrate17* directory, i.e., it is the parent directory where you are going to install *POLYRATE*.

Polyrate may be downloaded as a single tar.gz file with a structure that is illustrated in STEP 1. To uncompress and untar the *Polyrate* tar.gz file in the canonical place the user has to execute:

```
user@machine [~/Polyrate17] % tar -xzvf Polyrate17.tar.gz
```

STEP 1 Verify that the following directory structure was created properly when the file was uncompressed and untarred. User may use the 'ls -R' command.

```
user@machine [~/Polyrate17] % ls -R
```

```
PC/          doc/          obj/          script/       src/          sprng2.0/     testrun/
configure*   exe/          poten/        set_machine.jc*  testo/        util/
./PC:
check_test.pl  poly_ch4cl/  poly_ch5/     poly_cwmc/     poly_hni/     poly_oh3/
clean_test.bat poly_ch4o/   poly_clhbr/   poly_h2ni/     poly_ho2/     poly_ohcl/
clean_update.bat poly_ch4oh/  poly_cmc/     poly_h3/       poly_nh3/     run_all.bat

./PC/poly_ch4cl:
poly_ch4cl.mak poly_ch4cl.mdp run_ch4cl.bat

./PC/poly_ch4o:
poly_ch4o.mak poly_ch4o.mdp run_ch4o.bat

./PC/poly_ch4oh:
run_ch4oh.bat

./PC/poly_ch5:
poly_ch5.mak poly_ch5.mdp run_ch5.bat run_lola.bat run_lola.bat.bak

./PC/poly_clhbr:
poly_clhbr.mak poly_clhbr.mdp run_clhbr.bat

./PC/poly_cmc:
poly_cmc.mak poly_cmc.mdp run_cmc.bat

./PC/poly_cwmc:
poly_cwmc.mak poly_cwmc.mdp run_cwmc.bat

./PC/poly_h2ni:
poly_h2ni.mak poly_h2ni.mdp run_h2ni.bat

./PC/poly_h3:
poly_h3.mak poly_h3.mdp run_h3.bat

./PC/poly_hni:
hnitrl.dat hnitrl.fu15 hnitrl.fu6 poly_hni.mak pothni.dat
hnitrl.fu14 hnitrl.fu5 hnitrl.fu61 poly_hni.mdp run_hni.bat

./PC/poly_ho2:
poly_ho2.mak poly_ho2.mdp run_ho2.bat

./PC/poly_nh3:
poly_nh3.mak poly_nh3.mdp run_nh3.bat

./PC/poly_oh3:
poly_oh3.mak poly_oh3.mdp run_oh3.bat

./PC/poly_ohcl:
poly_ohcl.mak poly_ohcl.mdp run_ohcl.bat
```

```

./doc:

./exe:

./obj:

./poten:
c2h7pot5.f ch4o_module.f90 chain4.F coord3.F dumpot.F h3.F ho2.F ohcl.F
potch5j2.dat potcwmc3b.dat poth2nilat.dat setup4.F ch3.F ch5.F clhbr.F coord4.F h2ni.F
h3_module.F nh3.F potch4o.dat potcmc1.dat potcwmcvib.dat pothni.dat surf3.F
ch4o.F chain3.F cmc.F cwmc.F h2o2.F hni.F oh3.F potch5j1.dat potcmc2.dat
poth2niads.dat setup3.F surf4.F

./script:
comp_pot.jc* comp_src.jc* exe_chk.jc* link_exe.jc* potcomp.jc*

./src:
aamod.f90 dattim.F ef.F fromlapack.F hooks.F intbsv2.F.bak ivtstm.F Makefile
polyag.F polymq.F rtpjac.F util.f90 acespoly.F dummy_mpi.F energetics.F givtst.F
intbsv1.F intbsv3.F main.F poly31.F polyh1.F polyrr.F sprng_f.h vrcstst_mpi.F
alloc.f90 dumpot.F fromblas.F headr.F intbsv2.F interface.F main_vrcmpi.F poly40.F
poly_mpi.F polysz.F torsion.f90

./src/acesrate:
acespoly.F

./src/morate:
mopoly.F

./testo:
ch3tr1.fu15 ch5fu40tr2.fu15 ch5j1tr2.fu15 cmctr1.fu15 nh3tr1.fu6 oh3tr2.fu15
ch4cltr1.fu15 ch5fu40tr3.fu15 ch5j1tr3.fu15 cmctr2.fu15 nh3tr2.fu15 oh3tr3.fu15
ch4cltr2.fu15 ch5fu40tr4.fu15 ch5j1tr4.fu15 cmctr3.fu15 oh3fu30tr1.fu15 oh3tr4.fu15
ch4ohtr1.fu15 ch5fu40tr5.fu15 ch5j1tr5.fu15 cwmctr1.fu15 oh3fu30tr2.fu15 oh3tr5.fu15
ch4ohtr2.fu15 ch5fu40tr6.fu15 ch5j1tr6.fu15 cwmctr2.fu15 oh3fu30tr3.fu15 oh3tr6.fu15
ch4ohtr3.fu15 ch5fu50tr1.fu15 ch5j1tr7.fu15 cwmctr3.fu15 oh3fu30tr4.fu6 oh3tr7.fu15
ch4ohtr4.fu15 ch5fu50tr2.fu15 ch5j1tr8.fu15 h2nitr1.fu15 oh3fu30tr5.fu6 oh3tr8.fu15
ch4otr1.fu15 ch5fu50tr3.fu15 ch5j1tr9.fu15 h3o2fu31tr1.fu15 oh3fu30tr6.fu15 oh3tr9.fu15
ch5fu29tr1.fu15 ch5fu51tr1.fu15 ch5j2itr1.fu15 h3tr1.fu15 oh3fu31tr1.fu15 ohcltr1.fu15
ch5fu29tr2.fu15 ch5icfu30tr1.fu15 ch5j2itr2.fu15 h3tr2.fu15 oh3fu31tr2.fu15 ohcltr2.fu15
ch5fu30tr1.fu15 ch5icfu30tr2.fu15 ch5j2tr1.fu15 h3tr3.fu15 oh3fu40tr1.fu15 ohcltr3.fu15
ch5fu30tr2.fu15 ch5icfu31tr1.fu15 ch5j2tr2.fu6 h3tr4.fu15 oh3fu40tr2.fu15 ohcltr4.fu15
ch5fu30tr3.fu15 ch5icfu40tr1.fu15 ch5j2tr3.fu15 hnitr1.fu15 oh3fu40tr3.fu15 ohcltr5.fu15
ch5fu30tr4.fu15 ch5icfu40tr2.fu15 clhbrtr1.fu15 ho2tr1.fu15 oh3fu40tr4.fu6
ch5fu30tr5.fu15 ch5j1tr10.fu15 clhbrtr2.fu15 ho2tr2.fu15 oh3fu40tr5.fu6
ch5fu30tr6.fu15 ch5j1tr11.fu15 clhbrtr3.fu15 ho2tr3.fu15 oh3fu40tr6.fu15
ch5fu31tr1.fu15 ch5j1tr12.fu15 clhbrtr4.fu15 ho2tr4.fu15 oh3tr10.fu15
ch5fu40tr1.fu15 ch5j1tr1.fu15 clhbrtr5.fu15 ho2tr5.fu15 oh3tr1.fu15

./testrun:
ch3 ch5 clhbr nh3 runch3.jc runch5.jc runh2ni.jc runho2.jc
ch4cl check_all.jc cmc h3 h3o2 oh3 runch4cl.jc runcelhbr.jc runh3o2.jc runnh3.jc
ch4o check_test.jc cwmc hni ohcl runch4oh.jc runcmc.jc runh3.jc runoh3.jc

```

ch4oh clean_up.jc h2ni ho2 run_all.jc runch4o.jc runewmc.jc runhni.jc runohel.jc

./testrun/ch3:
ch3tr1.dat ch3tr1.jc

./testrun/ch4cl:
ch4cltr1.dat ch4cltr1.fu29 ch4cltr1.jc* ch4cltr2.dat ch4cltr2.fu29 ch4cltr2.jc*

./testrun/ch4o:
ch4otr1.dat ch4otr1.jc*

./testrun/ch4oh:
ch4ohic.dat ch4ohiv.dat ch4ohiv2.dat ch4ohtr1.dat ch4ohtr1.jc* ch4ohtr2.dat ch4ohtr2.fu29
ch4ohtr2.jc*

./testrun/ch5:
ch5fu29tr1.dat ch5fu30tr5.fu30 ch5fu40tr4.jc* ch5icfu30tr1.dat ch5icfu40tr2.jc* ch5j1tr7.dat
ch5fu29tr1.fu29 ch5fu30tr5.jc* ch5fu40tr5.dat ch5icfu30tr1.fu30 ch5j1tr1.dat ch5j1tr7.jc*
ch5fu29tr1.jc* ch5fu30tr6.dat ch5fu40tr5.fu40 ch5icfu30tr1.fu50 ch5j1tr1.jc* ch5j1tr8.dat
ch5fu29tr2.dat ch5fu30tr6.fu30 ch5fu40tr5.jc* ch5icfu30tr1.jc* ch5j1tr10.dat ch5j1tr8.jc*
ch5fu29tr2.fu29 ch5fu30tr6.jc* ch5fu40tr6.dat ch5icfu30tr2.dat ch5j1tr10.jc* ch5j1tr9.dat
ch5fu29tr2.jc* ch5fu31tr1.dat ch5fu40tr6.fu40 ch5icfu30tr2.fu30 ch5j1tr11.dat ch5j1tr9.jc*
ch5fu30tr1.dat ch5fu31tr1.fu31 ch5fu40tr6.jc* ch5icfu30tr2.fu50 ch5j1tr11.jc* ch5j2itr1.dat
ch5fu30tr1.fu30 ch5fu31tr1.jc* ch5fu50tr1.dat ch5icfu30tr2.jc* ch5j1tr12.dat ch5j2itr1.fu29
ch5fu30tr1.jc* ch5fu40tr1.dat ch5fu50tr1.fu50 ch5icfu31tr1.dat ch5j1tr12.jc* ch5j2itr1.jc*
ch5fu30tr2.dat ch5fu40tr1.fu40 ch5fu50tr1.jc* ch5icfu31tr1.fu31 ch5j1tr2.dat ch5j2itr2.dat
ch5fu30tr2.fu30 ch5fu40tr1.jc* ch5fu50tr2.dat ch5icfu31tr1.fu50 ch5j1tr2.jc* ch5j2itr2.fu29
ch5fu30tr2.jc* ch5fu40tr2.dat ch5fu50tr2.fu50 ch5icfu31tr1.jc* ch5j1tr3.dat ch5j2itr2.jc*
ch5fu30tr3.dat ch5fu40tr2.fu40 ch5fu50tr2.jc* ch5icfu40tr1.dat ch5j1tr3.jc* ch5j2itr1.dat
ch5fu30tr3.fu30 ch5fu40tr2.jc* ch5fu50tr3.dat ch5icfu40tr1.fu40 ch5j1tr4.dat ch5j2itr1.jc*
ch5fu30tr3.jc* ch5fu40tr3.dat ch5fu50tr3.fu50 ch5icfu40tr1.fu50 ch5j1tr4.jc* ch5j2itr2.dat
ch5fu30tr4.dat ch5fu40tr3.fu40 ch5fu50tr3.jc* ch5icfu40tr1.jc* ch5j1tr5.dat ch5j2itr2.jc*
ch5fu30tr4.fu30 ch5fu40tr3.jc* ch5fu51tr1.dat ch5icfu40tr2.dat ch5j1tr5.jc* ch5j2itr3.dat
ch5fu30tr4.jc* ch5fu40tr4.dat ch5fu51tr1.fu51 ch5icfu40tr2.fu40 ch5j1tr6.dat ch5j2itr3.jc*
ch5fu30tr5.dat ch5fu40tr4.fu40 ch5fu51tr1.jc* ch5icfu40tr2.fu50 ch5j1tr6.jc*

./testrun/clhbr:
clhbrtr1.dat clhbrtr2.dat clhbrtr3.dat clhbrtr4.dat clhbrtr5.dat
clhbrtr1.jc* clhbrtr2.jc* clhbrtr3.jc* clhbrtr4.jc* clhbrtr5.jc*

./testrun/cmc:
cmcic.dat cmctr1.dat cmctr1.jc* cmctr2.dat cmctr2.jc* cmctr3.dat cmctr3.fu51 cmctr3.jc*

./testrun/cwmc:
cwmctr1.dat cwmctr1.jc* cwmctr2.dat cwmctr2.jc* cwmctr3.dat cwmctr3.jc*

./testrun/h2ni:
h2nitr1.dat h2nitr1.jc*

./testrun/h3o2:
h3o2fu31tr1.dat h3o2fu31tr1.jc h3o2fu31tr1.fu31

./testrun/h3:
h3tr1.dat h3tr1.jc* h3tr2.dat h3tr2.jc* h3tr3.dat h3tr3.jc* h3tr4.dat h3tr4.jc*

./testrun/hni:
hnitrl.dat hnitrl.jc*

./testrun/ho2:
ho2ic.dat ho2tr1.jc* ho2tr2.jc* ho2tr4.dat ho2tr5.dat ho2tr1.dat
ho2tr2.dat ho2tr3.dat ho2tr3.jc* ho2tr4.jc* ho2tr5.jc*

./testrun/nh3:
nh3ic.dat nh3tr1.dat nh3tr1.jc* nh3tr2.dat nh3tr2.jc*

./testrun/oh3:
oh3fu30tr1.dat oh3fu30tr4.dat oh3fu30tr7.dat oh3fu40tr1.dat oh3fu40tr4.dat oh3tr1.dat
oh3tr3.jc* oh3tr8.dat
oh3fu30tr1.fu30 oh3fu30tr4.fu30 oh3fu30tr7.fu30 oh3fu40tr1.fu40 oh3fu40tr4.fu40 oh3tr1.jc*
oh3tr4.dat oh3tr8.jc*
oh3fu30tr1.jc* oh3fu30tr4.jc* oh3fu30tr7.jc* oh3fu40tr1.jc* oh3fu40tr4.jc* oh3tr10.dat
oh3tr4.jc* oh3tr9.dat
oh3fu30tr2.dat oh3fu30tr5.dat oh3fu31tr1.dat oh3fu40tr2.dat oh3fu40tr5.dat oh3tr10.jc*
oh3tr5.dat oh3tr9.jc*
oh3fu30tr2.fu30 oh3fu30tr5.fu30 oh3fu31tr1.fu31 oh3fu40tr2.fu40 oh3fu40tr5.fu40 oh3tr11.dat
oh3tr5.jc* tr7old.fu30
oh3fu30tr2.jc* oh3fu30tr5.jc* oh3fu31tr1.jc* oh3fu40tr2.jc* oh3fu40tr5.jc* oh3tr11.jc*
oh3tr6.dat
oh3fu30tr3.dat oh3fu30tr6.dat oh3fu31tr2.dat oh3fu40tr3.dat oh3fu40tr6.dat oh3tr2.dat
oh3tr6.jc*
oh3fu30tr3.fu30 oh3fu30tr6.fu30 oh3fu31tr2.fu31 oh3fu40tr3.fu40 oh3fu40tr6.fu40 oh3tr2.jc*
oh3tr7.dat
oh3fu30tr3.jc* oh3fu30tr6.jc* oh3fu31tr2.jc* oh3fu40tr3.jc* oh3fu40tr6.jc* oh3tr3.dat
oh3tr7.jc*

./testrun/ohcl:
ohcltr1.dat ohcltr1.jc* ohcltr2.dat ohcltr2.jc* ohcltr3.dat ohcltr3.jc* ohcltr4.dat ohcltr4.jc*

./util:
ch4.mi dattim.altix dattim.f dattim.ibm detmi.doc findl.f reord30.f
ch4oh.mi dattim.compaq dattim.f90 dattim.iris detmi.f hrotor.doc rotate/
cwmc.rmi dattim.cray dattim.gnu dattim.linux detmi.jc* hrotor.f
dateclock.c dattim.dec dattim.hp dattim.sun findb.f oh.mi

./util/rotate:
rotate.doc rotate30/ rotate31/ rotate40/

./util/rotate/rotate30:
rotate30.f rotate30.jc* test30.dat test30.fu30

./util/rotate/rotate31:
rotate31.f rotate31.jc* test31.dat test31.fu31

./util/rotate/rotate40:
rotate40.f rotate40.jc* test40.dat test40.fu40

STEP 2 Change your working directory to the *Polyrate17* directory and run the script *configure* by typing *./configure* <Return>. You will be asked if you want this version to be your default version of *POLYRATE*. If you select yes, then the hidden files *.poly_path* and *.poly_ver* will point to the location and the version of *POLYRATE*, respectively that you are installing. Then the script will ask which kind of executable (RP-VTST or VRC-VTST) the Makefile to generate. You should input your choice, i.e., “RP” or “VRC”. Next, the script will automatically detect available C and FORTRAN compilers and ask if you want to use the detected options. Usually, you should answer “yes” so that you can use the automatically detected compilers. If you want to use a different compiler, you should do the followings: (step 1) answer “yes” and proceed with the current configuring; (step 2) go to *src/* directory, and open the Makefile; (step 3) At the beginning of the Makefile, you will see:

```
F77C = /usr/bin/gfortran -O -fno-automatic
CC = /usr/bin/gcc
```

Please change the definition of these two variables (F77C and CC) to the location of your fortran and C compilers.

Note that no executable file is generated in this configuration step and the generated Makefile is for either RP-VTST or VRC-VTST.

STEP 3 Check the Makefile to ensure that the correct compiler options appear. *POLYRATE17* supports dynamic memory but previous versions do not, so compiler options that force static memory are typically required for previous versions. For example, the *g77* compiler requires the flag ‘-fno-automatic’, and the Portland Group compiler requires the flag ‘-Msave’. Due to the various combinations of OS and compilers, the configure script does not always print these flags automatically.

STEP 4 You may proceed directly to testing. The Makefile will compile any files that are needed to execute the test runs. If you want to use your own potential energy surface, you have to add it to the Makefile. The name of the new executable, the file name of the potential energy surface and its object have to be added. Comments in the Makefile will help you make these additions.

STEP 5 The following two compiling problems are common enough to merit special mention. If you get an error relating to module `libc.a(shr.o)`, it is probably because other modules are interfering with the compilation. Unload any extra modules that are running and recompile. If during the compilation of the RP-VTST version, the installation stops because the file `mpif.h` cannot be found, the user should generate a dummy file called `mpif.h` in the `src` directory (for instance, by typing `touch mpif.h` <Return> in the *Polyrate* `src` directory).

15.A.2. Testing

The directory structure and files should be set up as instructed in Subsection 15.A.1. Note that some of the processes described in this section may be time consuming. In a Unix system, you can always add an `&` character at the end of a command line to put the process in the background. In the following instructions, whenever you see a command followed by an `&` character, it means that we suggest that the user run the command in the background.

It is recommended that the user run all the test runs in the distribution package before using *POLYRATE*. We have included many useful scripts to run the test runs, and you can choose how to run the test runs from among several different approaches.

(1) To run all the test runs:

The test runs are grouped into 15 different sets, in particular, `ch3`, `ch4cl`, `ch4oh`, `ch5`, `clhbr`, `cmc`, `cwmc`, `h2ni`, `h3`, `h3o2`, `hni`, `ho2`, `nh3`, `oh3`, and `ohcl` test runs. (See Subsection 16.C.) Each set has its own subdirectory in the *Polyrate17*/`testrun` directory. (See Subsection 15.A.1.) The subdirectories contain the input data files and the job control files for the test runs. The only thing you need to run all the test runs for either RP-VTST or VRC-VTST is to change the working directory to the *Polyrate17*/`testrun` directory and run the script `run_allRPVTST.jc` or `run_allVRCVTST.jc`.

```
./run_allRPVTST.jc & <Return>
```

(or submit it as a batch job)

Note `run_allRPVTST.jc` and `run_allVRCVTST.jc` need different Makefiles. The user has to take the steps described in Section 15.A.1 to generate a Makefile either for VRC-VTST calculations or for RP-VTST calculations.

(2) To run a subset of the test runs

1. To run a set of test runs, use the scripts available in the `testrun/` directory. For example, to run all the h3 test runs, change the working directory to the *Polyrate17/testrun* directory and use the script `runh3.jc`.

(change to the `testrun/` directory)

```
./runh3.jc & <Return>
```

(or any of the other `run{name}.jc` scripts, where `{name}` can be `ch5`, `oh3`, `ho2`, `clhbr`, `cmc`, `cwmc`, `nh3`, `hni`, `h2ni`, `ch4cl`, or `ch4oh`)

3. To run an individual test run, say `ch5j1tr1`, change the directory to `testrun/ch5/` and use the script `ch5j1tr1.jc`.

```
./ch5j1tr1.jc & <Return>
```

(or any of the `{test_run_name}.jc` scripts in that directory)

Note: Do not run more than one test run in the same directory simultaneously. (See Subsection 15.B, item VI.)

(3) To check the test run results:

When you finish running the test runs, you should check if they produced the correct results by comparing the test run short output files (`fu15` files) with the ones in the `testo/` directory. There is a useful script, `check_test.jc`, in the `testrun/` directory to do this after you finish a whole set of test runs. For example, if you have finished all the `oh3` test runs and you want to compare the results with the correct ones, change the working directory to `testrun/` and type `./check_test.jc oh3 <Return>`. The script will use the Unix command `diff` to compare your results with the ones in the directory `testo/`, and generate an output file called `oh3.chk`. You can then look at the contents of that file and see if there are any differences. Since the reaction rates are not calculated in test runs `ch5j2tr2` and `nh3tr1`, the short output files for these two test runs are not created. Thus if you type `check_test.jc ch5 <Return>`, there will be no check for `ch5j2tr2`. You can still use the `diff` command to compare the long output files (`fu6` files) of these test runs with the ones provided in the *Polyrate17/testo* directory. There is a similar script, `check_all.jc`, that you can use to test the full suite of test runs.

If your test runs reproduce the results in *Polyrate17/testo*, the only differences will be the date and time. Otherwise, the script will point out where the results differ from each other. (Of course, you have to know a little bit about the diff command to understand its output. Type *man diff* <Return> if you need help.) In running the test runs on different machines, we found that the differences in the rate constants were never greater than 1%. If your rate constant results do not differ from the ones in the *Polyrate17/testo* directory by more than this percentage, they are probably correct; otherwise, something unexpected might be happening.

(4) Cleaning up:

When you finish checking the testrun results, you may want to clean up the testrun directory to save disk space. The script, *clean_up.jc*, in the *Polyrate17/testrun* directory can be used to remove all the files *POLYRATE* generated when running the test suite. These files include the following: *.fu1, *.fu6, *.fu14, *.fu15, *.fu25, *.fu26, *.fu27, and *.fu61.

15.A.3. Using *POLYRATE*

Many users will use *POLYRATE* mainly as part of a direct dynamics calculations carried out with *GAMESSPLUSRATE*, *GAUSSRATE*, *JAGUARRATE*, *MC-TINKERATE*, *MORATE*, *MULTILEVELRATE*, or *NWCHEMRATE*. However the rest of this section is devoted to using *POLYRATE* as a stand-alone program.

After installation you should have the default directory structure and the correct machine-dependent files. If you are planning to use *POLYRATE* with a direct dynamics code, such as *GAUSSRATE* or *MULTILEVELRATE*, read the instructions in the direct dynamics manual. If you are planning to use *POLYRATE* as a stand-alone code, follow the instructions in this section:

1. *Unless you are performing an electronic structure input file calculation, prepare your own potential energy surface subroutine to do the calculations.* See Section 8 for a discussion of user-supplied potential energy surfaces. Supposing that you have made your own potential energy surface subroutine, you need to compile it before you link it with the *POLYRATE* source code. This can be accomplished by adding the

surface into the Makefile located in the src directory. Comments in the Makefile will show you where to add lines. For example, suppose you have made a surface subroutine for a system that consists of 2 hydrogen atoms and 1 oxygen atom, and you have put it in the *Polyrate17/poten* directory and named *oh2.f*. You have to add the following line where the executables are defined

```
OH2S = ../exe/Polyrate$(VERSION).oh2.serial.exe
```

Then add *../poten/oh2.f* to the POT: list, using a \ in order to continue a line. Also, add the following line to compile the surface, beginning with a tab.

```
$(F77C) -c ../poten/oh2.f
```

Finally, add the lines that create the executable. Use a tab to start the second line.

```
OH2S:$(OBJ) dummy_mpi.o oh2.o
```

```
$(F77C) -o $(OH2) $(OBJ) dummy_mpi.o oh2.o
```

For three- and four-atom systems, the user can take advantage of some pre-written subroutines, namely: *setup*, *surf*, *coord*, and *chain*. If these subroutines were not explicitly included in the *oh2.f* file, the lines to create the executable would look like this (without a break between *oh2.o* and *setup3.o*):

```
OH2S:$(OBJ) dummy_mpi.o oh2.o
$(F77C) -o $(OH2) $(OBJ) dummy_mpi.o oh2.o
setup3.o surf3.o coord3.o chain3.o
```

2. *Prepare the input data files.* Study Section 12 carefully to make the input data files correctly. Because some of the input data files are quite complicated, it may be easier to use some of the test run input data files as templates when preparing your own input data files.
3. *Run the dynamics calculations with POLYRATE.* After you finish steps 1 and 2, you are ready to use *POLYRATE* to do your calculations. Before you submit your job to the computer, it is recommended that you do the following:

- A. Make a directory for each specific reaction. For example, you can make a directory called reaction in the *Polyrate17* directory or somewhere else, and when you are planning to do a calculation on a specific reaction, say $O + H_2$, you can create a directory called oh2 in the *Polyrate17/reaction* directory. You can then use this directory as your working directory for this reaction and put all the input data files and output files in this directory.
- B. Use a job control script (.jc) to handle all the processes involved in running *POLYRATE*. *POLYRATE* only recognizes a certain set of input file names and will generate a set of output files with certain names (see Sections 12, 15.B and 16.C), therefore, the user always needs to copy (or link) the input data files to the default names and rename the output files to more logical names. It is more convenient to use a script to control these tedious processes. Using a script also makes it easier to run the job in background or to submit the *POLYRATE* run as a batch job. In fact, we have included job control files for all test runs in the distribution package. You probably used them when you ran the test runs, and you can easily modify them for your own needs. They all reside in the subdirectories of the *Polyrate17/testrun* directory. In your .jc script, be sure to change the 'set name' and 'gmake' lines to match your job. In the OH_2 example, if you decided the name your files oh2run1, the lines would look like this:

```
set name = oh2run1
```

```
gmake OH2S
```

- C. With gmake in the job control script, the executable will be compiled the first time that it is used, and also anytime there is a change in one of the files that it is dependent on. If a change is made to the source code, the next time that the job control script is run, gmake will recognize that a change has been made and compile the changed code and create a new executable. If the user does not desire this feature, the gmake line may be deleted from the .jc job control script once the executable is compiled for the first time. Executables are put in the exe directory by default.

15.B. Comments for expert users

Of course, users need not use the provided Unix scripts, files, or recommended directory structure. All that is needed is the compilation of the source code, linking the *POLYRATE* object files with appropriate potential energy surface object file(s), and running the executable file(s) with input data file(s). However, there are a few practical issues to be aware of, as follows.

Compilation of POLYRATE

For the RP version of the code, compile

```
dattim.F  ef.F      fromlapack.F hooks.F  intbsv2.F.bak ivtstm.F,  
polyag.F  polymq.F  rtpjac.F   util.f90  acespoly.F dummy_mpi.F  
energetics.F givtst.F, intbsv1.F intbsv3.F  main.F    poly31.F  
polyhl.F  polyrr.F  sprng_f.h  vrcstst_mpi.F, alloc.f90  dumpot.F  
fromblas.F headr.F   intbsv2.F interface.F  main_vrcmpi.F  
poly40.F , poly_mpi.F polysz.F torsion.f90
```

For the RP version of the code compile

```
energetics.F, givtst.F, hooks.F, intbsv1.F, intbsv2.F intbsv3.F,  
interface.F, main.F, Polyrate.F, polyag.F, polyhl.F, polymq.F, polyrr.F,  
polysz.F, rtpjac.F,
```

The *POLYRATE17* program should be compiled with 64-bit precision and dynamic memory allocation.

Compilation of the potential energy surface files

The potential energy surface subroutines provided with the *POLYRATE* program should be compiled with the same precision and memory specifications used for the *POLYRATE* source code. Any user-supplied potential energy surface subroutine must also be compiled with 64-bit precision, but specification for static memory allocation will depend on the way in which the subroutine is written.

Creation of an executable file

Link the *POLYRATE* object file with potential object file(s) to create specific *POLYRATE*-potential executable file(s).

Running POLYRATE

The *POLYRATE* program opens all the input and output files needed for a calculation with the logical file names poly.fu#, where # is an integer. The files poly.fu5 and poly.fu6 are always opened (they correspond to input and output files respectively). The other poly.fu# files are opened depending on which options are set. Therefore, all the necessary input data files must be copied or linked to files with the proper logical names. If the user wants the output files to have different names, the renaming of the poly.fu# output files must be performed after the calculation has completed.

Note: Because the *POLYRATE* program opens and closes its files with the file name specification poly.fu#, these files should be renamed at the end of each calculation, or there will be conflicts when two calculations are run in the same directory.

In addition to the *POLYRATE* input and output files, the user should ensure that the potential data files, if any, which are opened by the potential energy surface subroutine(s), have the correct file names as specified in the OPEN statement(s) in the potential subroutine source code. For the test runs, these names are listed after "FILE NAME =" in Subsection 16.C. Also, these data files need to be in the working directory of the calculation.

15.C. Running in parallel using MPI

The *POLYRATE* program has been expanded to work with multiple processors using MPI. MPI can be very machine dependent, therefore this section will highlight the areas that need to be correct for the parallel feature to work correctly. There are two MPI versions. One is for calculations with MEP and the other one is for calculations with variable reaction coordinate (VRC). These two versions need to be compiled by including different subroutines since they use different parallel schemes.

A. MPI version for calculations with MEP

For samples on how to make MPI work using MEP, see the CH5P sections of the Makefile, and look at the job control scripts in the ch5 testruns. The MPI sections of the Makefile are only printed if an MPI compiler is found by the configure script.

- I. In the Makefile, MPIF77C needs to be set to the correct MPI compiler. The poly_mpi.f source file must be compiled using this compiler, not the serial compiler. Finally, the linking section must use the MPI compiler when creating the executable.
- II. In the job control script (*.jc file), set \$nproc equal to the number of processors you wish to use. Make sure the exe variable is set to the proper executable. Most importantly, the platform on which you are running has a certain way that MPI jobs must be run. By default, the line looks like this:


```
(time mpirun -v -np $nproc $exe) >>& $name.time
```

 where “mpirun -v -np \$nproc \$exe” is the part that actually tells the program how to run. This line will likely need to be edited to work with each platform.
- III. For an example, the testrun ch5j1tr1 has edited this line in the job control script to work on a particular IBM regatta. The line now reads:

```
(time poe $exe -rmpool 1 -nodes 1 -procs $nproc)
>>& $name.time
```

B MPI version for calculations with VRC

When installing *POLYRATE* program, user will be asked whether the MPI executable for variable reaction coordinate need to be compiled by a Perl installation script under *Polyrate17* directory, named *configure*. This script will automatically search MPI compiler for user. But user has to be responsible to make sure the compiler and its options are correct in Makefile since MPI can be very machine dependent.

Since this MPI version for variable reaction coordinate use SPRNG (SCALABLE PARALLEL RANDOM NUMBER GENERATOR LIBRARY) package in Monte Carlo integrations, user needs to install SPRNG before *POLYRATE* installation. Starting with version 2010, SPRGN 2.0 package is included in *POLYRATE* program. The *Polyrate* installation script will compile the SPRNG 2.0 library before generate Makefile for VRC-VTST run. The SPRNG package can also be downloaded from its website at the Florida State University (<http://sprng.cs.fsu.edu/>) and installed separately by users. When linking object files, user

should give the correct path for SPRNG library by changing the \$SPRNLIB variable in Makefile if user desires use separately installed SPRNG library.

16. SAMPLE TEST RUNS

16.A. Chart of the potential energy functions and data files

The following chart gives the name of the potential energy function and data file(s) that are used for each test run. They are files located in the *Polyrate17*/poten directory.

When two data files are listed for the same test run, both data files are required. The file listed first should be assigned to unit 4, and the file listed second should be assigned to unit 7.

TEST RUN(S)	POTENTIAL SOURCE CODE	POTENTIAL DATA FILE(S)
ch3tr1	ch3.F	(none)
ch5j1tr1-ch5j1tr11	ch5.F	potch5j1.dat
ch5j2tr1-ch5j2tr3	ch5.F	potch5j2.dat
ch5fu51.tr1	ch5.F	potch5j1.dat
clhbrtr1-clhbrtr4	clhbr.F	(none)
cmctr1, cmctr2	cmc.F	potcmc1.dat potcmc2.dat
cwmctr1-cwmctr3	cwmc.F	potcwmc3b.dat potcwmcvib.dat

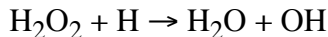
TEST RUN(S)	POTENTIAL SOURCE CODE	POTENTIAL DATA FILE(S)
ch4otr1	ch4o.F	potch4o.dat potch4o.inc
h3tr1-h3tr4	h3.F	(none)
nh3tr1-nh3tr2	nh3.F	(none)
ho2tr1-ho2tr5	ho2.F	(none)
oh3tr1-oh3tr10	oh3.F	(none)
hnitr1	hni.F	pothni.dat
h2nitr1	h2ni.F	poth2niads.dat poth2nilat.dat
ohcltr1-ohcltr4	ohcl.F	(none)

16.B. Descriptions of sample test runs

This section contains descriptions of the sample runs which make up the *POLYRATE* test suite. The input data files and the summary output files from unit fu15 for the test runs are part of the distribution package. For convenience, we have named all the test runs according to the potential energy surface used for the test run.

The following is a list of reactions involved in the test suite and references to published calculations of their reaction rates using *POLYRATE*.

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_3$	
$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$	see Refs. 17 and 20 in Sect. 23
$\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$	see Ref. 21 in Sect. 23
$\text{Cl} + \text{HBr} \rightarrow \text{ClH} + \text{Br}$	see Ref. 10 in Sect. 23
$\text{Cl}^- + \text{CH}_3\text{Cl}^+ \rightarrow \text{ClCH}_3 + \text{Cl}^-$	see Ref. 23 in Sect. 22
$\text{Cl}^-(\text{H}_2\text{O}) + \text{CH}_3\text{Cl}^+ \rightarrow \text{ClCH}_3 + \text{Cl}^-(\text{H}_2\text{O})$	see Ref. 24 in Sect. 22
$\text{H}_2 + \text{H}^+ \rightarrow \text{HH}^+ + \text{H}$	see Ref. 12 in Sect. 23
$\text{OH} + \text{O} \rightarrow \text{HO}_2$	see Ref. 4 in Sect. 23
$\text{O}_2 + \text{H} \rightarrow \text{HO}_2$	see Ref. 4 in Sect. 23
$\text{NH}_3 \rightarrow \text{NH}_3$ (inversion)	see Ref. 7 in Sect. 23
$\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$	see Refs. 1, 3, and 6 in Sect. 23
$\text{H}(\text{a})/\text{Ni}(100) \rightarrow \text{H}(\text{a})/\text{Ni}(100)$	see Ref. 33 in Sect. 22
$\text{H}_2(\text{a})/\text{Ni}(100) \rightarrow 2\text{H}(\text{a})/\text{Ni}(100)$	see Ref. 19 in Sect. 23
$\text{OH} + \text{CH}_4 \rightarrow \text{H}_2\text{O} + \text{CH}_3$	see Refs. 26 and 32 in Sect. 23
$\text{Cl} + \text{CH}_4 \rightarrow \text{HCl} + \text{CH}_3$	see Refs. 22 and 26 in Sect. 23
$\text{CH}_4 + \text{O} \rightarrow \text{CH}_3 + \text{OH}$	see Ref. 52 in Sect. 22
$\text{O} + \text{HCl} \rightarrow \text{OH} + \text{Cl}$	
$\text{O} + \text{HCl}(v=1) \rightarrow \text{OH} + \text{Cl}$	see Refs. 22 and 26 in Sect. 23

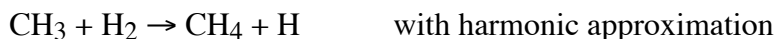


Various options were used for each of these systems to give a thorough test of *POLYRATE*. Some of the most critical options used are discussed in each description; the keywords used for these options are specified in square brackets: e. g. [KEYWORD]. The potential energy surfaces used for the sample runs for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ are surfaces J1 and J2 of Joseph et al. [Ref. 17, Sect. 23]. Surface J1 is also used for the calculations for the reaction $\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$, where Ph denotes a single atom with the mass of a phenyl group (see Ref. 21 in Sect. 23). The calculations for the reaction $\text{Cl} + \text{HBr} \rightarrow \text{ClH} + \text{Br}$ use the extended LEPS surface of Babamov, Lopez, and Marcus [Ref. 26, Sect. 22]. The calculations for the $\text{Cl}^-(\text{H}_2\text{O}_m) + \text{CH}_3\text{Cl} \rightarrow \text{ClCH}_3 + \text{Cl}^-(\text{H}_2\text{O}_m)$ reactions, $m = 0,1$, use the potential energy surface of Tucker and Truhlar [Ref. 23, Sect. 22] and Zhao et al. [Refs. 24 and 27, Sect. 22], while the DMBE surface of Varandas et al. [Ref. 28, Sect. 22] is used for the reaction $\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$. The calculations for the reaction $\text{OH} + \text{O} \rightarrow \text{HO}_2$ and $\text{O} + \text{H}_2 \rightarrow \text{HO}_2$ use the ab initio surface of Melius and Blint [Ref. 29, Sect. 22], and all the runs for the reaction $\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$ use the Schatz-Elgersma fit to the Walch-Dunning ab initio potential energy surface [Refs. 30 and 31, Sect. 22]. The embedded diatomics-in-molecules (EDIM) potential with parameter set 4 of Truong et al. [Refs. 25 and 32, Sect. 22] is used for the H/Ni(100) surface diffusion test run, and the Lee and DePristo [Ref. 33, Sect. 22] potential energy surface, as modified by Truong et al. [Ref. 19, Sect. 23], is used for the $\text{H}_2/\text{Ni}(100)$ system. Furthermore, both Ni surface test runs are carried out assuming that the entire metallic subsystem on which the diatoms are absorbed is rigid. The calculations for the reaction $\text{Cl}^- + \text{CH}_3\text{Cl} \rightarrow \text{ClCH}_3 + \text{Cl}^-$ use the semiempirical surface of Tucker and Truhlar [Ref. 23, Sect. 22], and the calculations for the NH_3 inversion use potential energy surface II of Brown, Tucker, and Truhlar [Ref. 7, Sect. 23]. For the $\text{OH} + \text{CH}_4 \rightarrow \text{H}_2\text{O} + \text{CH}_3$ [Ref. 32, Sect. 23] and $\text{Cl} + \text{CH}_4 \rightarrow \text{HCl} + \text{CH}_3$ [Ref. 22, Sect. 23] reactions, rate calculations are carried out using the IVTST method [Ref. 26, Sect. 23] with *ab initio* data.

It should be pointed out that the results for the ClHBr and the H_3 test runs will differ from the published results. This difference occurs because the published results for those systems were determined using another program, ABCRATE, which calculates anharmonicity corrections differently than *POLYRATE*. In particular, ABCRATE uses curvilinear internal coordinates, and these produce quite different harmonic frequencies, as discussed elsewhere [Ref. 25 and 41, Sect. 23].

Note that the test runs are designed to illustrate the use of the *POLYRATE* program. No implication is made that these runs are converged with respect to all step sizes and grid ranges. One can use these runs to test that the code as received gives the same results on various new platforms by comparing the output for these test runs to the distributed output, even though the results are not necessarily converged with respect to numerical parameters.

16.C.1. Test run ch5j1tr1



This sample run is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with the J1 potential energy surface. All normal modes and generalized normal modes are treated using the default harmonic approximation. The rate constants are computed using CVT and ICVT [CVT, ICVT]. Tunneling corrections are computed using the MEPSAG, CD-SCSAG, and LCG3 methods [ZCT, SCT, SCTOPT, LCTOPT] Note that only the ground-state in the LCG3 exit arrangement is considered for the LCG3 calculation in this run [NOALLEXCIT] this option is chosen to simplify the calculation because we expect on physical grounds that only ground-state-to-ground-state tunneling is important in this case. The LCT keyword was not explicitly given since the default is on when LCTOPT is used. The rate constants are computed for ten temperatures [TEMP], and the activation energies are computed for six pairs of temperatures [EACT]. The MEP is followed using the Page-McIver method [RPM = *pagem*] with a step size of 0.001 bohr [SSTEP]. The Hessian matrix is recalculated and a mode analysis is performed every 0.01 bohr [INH = 100]. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. A restart file containing reaction-path information is written to unit fu1 [RESTART = *writefu1*] for use in test run ch5j1tr2. For both reactants and the first product STATUS = 6 and the geometries, energies, and frequencies are read from fu5, for the second product STATUS = 2 and the geometry is read from fu5, and for the saddle point STATUS = 4 and the geometry, energy, and Hessian are read from fu5. The reactants are input in fu5 [ENERGY, FREQ] to reduce the number of calculations. The results may be compared to those in Table VII of Ref. 17 in Sect. 23.

POLYRATE I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr1.dat	poly.fu5	input data file
ch5j1tr1.fu6	poly.fu6	long output file
ch5j1tr1.fu1	poly.fu1	reaction path info.
ch5j1tr1.fu14	poly.fu14	summary output file
ch5j1tr1.fu15	poly.fu15	summary output file
ch5j1tr1.fu61	esp.fu61	2nd restart option file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.2. Test run ch5j1tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with harmonic approximation

This sample run is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with the J1 potential energy surface. All the parameters are the same as for test run ch5j1tr1 except that the reaction-path information is read from a restart file on unit fu1 [RESTART: readfu1]. In order to perform this test run, the restart file must first be created by the user by running test run ch5j1tr1.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr2.dat	poly.fu5	input data file
ch5j1tr1.fu1	poly.fu1	reaction path info. (input data for a restart calculation)
ch5j1tr2.fu6	poly.fu6	long output file
ch5j1tr2.fu14	poly.fu14	summary output file
ch5j1tr2.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat	UNIT = 4
	FILE NAME = potch5.dat

16.C.3. Test run ch5j1tr3

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with mass-weighted reaction coordinate

This sample run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ is similar to test run ch5j1tr1, except that the reduced mass [SCALEMASS] which defines the mass-scaled Cartesian coordinates, is set equal to the default, 1.0 amu. Thus, these coordinates have the same numerical values as the conventional mass-weighted coordinates in infrared spectroscopy. The vibrational partition functions are multiplied by a factor of 1×10^{10} to avoid underflow [VPFEXP]; note that these changes alone would not change the final rate constants. In addition, though, in this run the effective mass for the CD-SCSAG tunneling calculation is obtained by sixth-order Lagrangian interpolation [SCTOPT: lagrange] instead of a cubic spline fit. Also, there is no LCG3 calculation in this test run.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr3.dat	poly.fu5	input data file
ch5j1tr3.fu6	poly.fu6	long output file
ch5j1tr3.fu14	poly.fu14	summary output file
ch5j1tr3.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat	UNIT = 4
	FILE NAME = potch5.dat

16.C.4. Test run ch5j1tr4

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with harmonic approximation in redundant internal coordinates and the Page-McIver method to calculate reaction-path curvature

This sample run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ demonstrates the calculation of reaction-path curvature components from the Hessian, generalized normal mode eigenvectors, and gradient [$\text{CURV} = \text{dhess}$] using eq. (48) of Page and McIver [Ref. 17, Sect. 22]. The MEP is followed using the Euler steepest-descents (ESD) method [$\text{RPM} = \text{esd}$] with a step size of 0.0001 bohr [SSTEP]. A generalized normal mode analysis is performed every 0.01 bohr along the MEP [$\text{INH} = 100$], and all the vibrations are treated harmonically in redundant internal coordinates [$\text{COORD} = \text{curv3}$]. Both CVT [CVT] and ICVT [ICVT] rate constants are calculated, and tunneling corrections are computed using both MEPSAG and CD-SCSAG methods [$\text{ZCT}, \text{SCTOPT}$].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr4.dat	poly.fu5	input data file
ch5j1tr4.fu6	poly.fu6	long output file
ch5j1tr4.fu14	poly.fu14	summary output file
ch5j1tr4.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat	UNIT = 4
	FILE NAME = potch5.dat

16.C.5. Test run ch5j1tr5

$\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ with LCG3 calculation for the ground state only

This sample run is for the $\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ reaction, where Ph denotes a single particle with the mass of a phenyl group. The CH_5 analytical potential energy surface, J1, is used for this six-atom model problem; see Ref. 21 in Sect. 23 for further discussion. All normal modes and generalized normal modes are treated using the default harmonic approximation. TST, CVT, and ICVT rate constants are computed [TST, CVT, ICVT]. The tunneling calculations for ground-state transmission coefficients are computed using the MEPSAG, CD-SCSAG, and LCG3 methods [ZCT, SCTOPT, LCTOPT]. The program writes a restart file containing the reaction-path information to unit fu1 for use in test run ch5j1tr6 [RESTART = *writeln fu1*]. The MEP is followed using the ESD integrator without stabilization (ESD method) [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.01 bohr [INH = 100]. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. The results may be compared to those in Table 5 in Ref. 21 in Sect. 23, but it should be noted that there is a difference of a few percent between the LCG3 results in Ref. 21 and those calculated with this data file due to improved convergence.

Polyrate I/O FILES

FILE NAME

ch5j1tr5.dat
ch5j1tr5.fu6
ch5j1tr5.fu1
ch5j1tr5.fu14
ch5j1tr5.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu1 reaction path info.
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat

UNIT = 4
FILE NAME = potch5.dat

16.C.6. Test run ch5j1tr6

$\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ with LCG3 calculation for all energetically accessible excited states from a restart file in which the LCG3 calculation was carried out for the ground state only; rate constants are also calculated at more temperatures. (Restart file for unit fu1 is from test run ch5j1tr5)

This sample run, like the preceding test run, is for the $\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ reaction, where Ph denotes a single particle with the mass of a phenyl group. The potential energy surface and all the parameters are the same as for test run ch5j1tr5 except for the tunneling calculations and the temperatures for which the rate constants are calculated. In this run the tunneling calculations for ground-state transmission coefficients are performed with the MEPSAG, CD-SCSAG, and LCG3 methods [ZCT, SCTOPT, LCTOPT]. For the LCG3 calculation, the program determines the highest excited state of the LCG3 exit arrangement which is accessible, and those excited states are included in the LCG3 calculation [ALLEXCIT]. The results may be compared to those from test run ch5j1tr5, in which the LCG3 calculations were done for the ground state only; the present LCG3 transmission coefficients are up to 4% larger.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr6.dat	poly.fu5	input data file
ch5j1tr5.fu1	poly.fu1	reaction path info. (input data for a restart calculation)
ch5j1tr6.fu6	poly.fu6	long output file
ch5j1tr6.fu14	poly.fu14	summary output file
ch5j1tr6.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.7. Test run ch5j1tr7

CH₃ + H₂ → CH₄ + H with Page-McIver mass-weighted reaction path and
IVTST0FREQ

This sample run for the reaction CH₃ + H₂ → CH₄ + H is similar to test run ch5j1tr3, except that the reaction-path curvature components are calculated from the Hessian, generalized normal mode eigenvectors, and gradient [CURV = *dhess*] using eq. (48) of Page and McIver [Ref. 17, Sect. 22]. The MEP is followed using the Page-McIver method [RPM = *pagem*] with a step size of 0.001 bohr [SSTEP]. The Hessian matrix is recalculated and a mode analysis is performed every 0.01 bohr [INH = 100]. The vibrational partition functions are multiplied by a factor of 1×10^{10} to avoid underflow [VPFEXP]; note that this alone would not change the final rate constants. In addition, the effective mass for the CD-SCSAG tunneling calculation is obtained by sixth-order Lagrangian interpolation [SCTOPT: lagrange] instead of the default cubic spline fit. Also, there is no LCG3 calculation in this test run. The IVST0FREQ method [IVTST0FREQ] is used to modify the 10th and 11th vibrational modes along the reaction path.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr7.dat	poly.fu5	input data file
ch5j1tr7.fu6	poly.fu6	long output file
ch5j1tr7.fu14	poly.fu14	summary output file
ch5j1tr7.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
FILE NAME = potch5.dat

16.C.8. Test run ch5j1tr8

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with MEPSAG and CD-SCSAG calculations at two temperatures [one of the test cases in Ref. 27 in Sect. 23]

This sample run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ is one of the test cases in Ref. 27 in Sect. 23. (However, Ref. 27 used SCSAG whereas the current code uses CD-SCSAG). The MEP is calculated from $s = -1.3$ bohr to $s = 0.8$ bohr [SRANGE] using the Euler stabilization method [RPM = *es1*]. The gradient and save step sizes are all set to 0.047 bohr [SSTEP, INH = 1]. The finite difference step size and the tolerance parameter in the ES1 method are set to 0.072 ($1.532 * \text{SSTEP}$) [ES1OPT: delta2] and 0.082 [ES1OPT: DIFFD], respectively. The CVT/MEPSAG and CVT/CD-SCSAG rate constants are calculated [CVT, ZCT, SCT] at 200K and 300K [TEMP]. All the rate constants are within 15% of the converged values obtained in the test run ch5j1tr1.

Polyrate I/O FILES

FILE NAME

ch5j1tr8.dat
ch5j1tr8.fu6
ch5j1tr8.fu14
ch5j1tr8.fu15

Polyrate FILE NAME

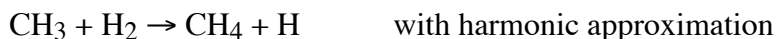
poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat

UNIT = 4
FILE NAME = potch5.dat

16.C.9. Test run ch5j1tr9



This sample run is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with the J1 potential energy surface. All normal modes and generalized normal modes are treated using the default harmonic approximation. The rate constants are computed using CVT and ICVT [CVT, ICVT]. Tunneling corrections are computed using the MEPSAG, CD-SCSAG, and LCG4 methods [ZCT, SCT, SCTOPT, LCTOPT] Note that only the ground-state in the LCG4 exit arrangement is considered for the LCG4 calculation in this run [NOALLEXCIT] this option is chosen to simplify the calculation because we expect on physical grounds that only ground-state-to-ground-state tunneling is important in this case. The LCT keyword was not explicitly given since the default is on when LCTOPT is used. The rate constants are computed for ten temperatures [TEMP], and the activation energies are computed for six pairs of temperatures [EACT]. The MEP is followed using the Page-McIver method [RPM = *pagem*] with a step size of 0.001 bohr [SSTEP]. The Hessian matrix is recalculated and a mode analysis is performed every 0.01 bohr [INH = 100]. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. For both reactants and the first product STATUS = 6 and the geometries, energies, and frequencies are read from fu5, for the second product STATUS = 2 and the geometry is read from fu5, and for the saddle point STATUS = 4 and the geometry, energy, and Hessian are read from fu5. The reactants are input in fu5 [ENERGY, FREQ] to reduce the number of calculations. The results may be compared to those in Table VII of Ref. 17 in Sect. 23.

POLYRATE I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr9.dat	poly.fu5	input data file
ch5j1tr9.fu6	poly.fu6	long output file
ch5j1tr9.fu1	poly.fu1	reaction path info.
ch5j1tr9.fu14	poly.fu14	summary output file
ch5j1tr9.fu15	poly.fu15	summary output file
ch5j1tr9.fu61	esp.fu61	2nd restart option file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.10. Test run ch5j1tr10

$\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ with LCG4 calculation for all energetically accessible excited states from a restart file in which the LCG4 calculation was carried out for the ground state only; rate constants are also calculated at more temperatures. (Restart file for unit fu1 is from test run ch5j1tr5)

This sample run, like the preceding test run, is for the $\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ reaction, where Ph denotes a single particle with the mass of a phenyl group. The potential energy surface and all the parameters are the same as for test run ch5j1tr5 except for the tunneling calculations and the temperatures for which the rate constants are calculated. In this run the tunneling calculations for ground-state transmission coefficients are performed with the MEPSAG, CD-SCSAG, and LCG4 methods [ZCT, SCTOPT, LCTOPT]. For the LCG4 calculation, the program determines the highest excited state of the LCG4 exit arrangement which is accessible, and those excited states are included in the LCG4 calculation [ALLEXCIT].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr10.dat	poly.fu5	input data file
ch5j1tr5.fu1	poly.fu1	reaction path info. (input data for a restart calculation)
ch5j1tr10.fu6	poly.fu6	long output file
ch5j1tr10.fu14	poly.fu14	summary output file
ch5j1tr10.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.11. Test run ch5j1tr11

$\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ with interpolated LCG4 calculation with all energetically accessible excited states.

This sample run for the $\text{CH}_3 + \text{HPh} \rightarrow \text{CH}_4 + \text{Ph}$ is similar than the previous run where Ph denotes a single particle with the mass of a phenyl group. The only difference with the previous test is that the large curvature tunneling energies obtained by the LCG4 method are interpolated to 9 points in the nonadiabatic zone [ILCT1D with LCTGRD equal to 9 for all excited states]. The ILCG4 calculation includes all the final excited states that are accessible [ALLEXCIT]. It is interesting to compare the LCG4 transmission factors with the ones calculated in the previous run in which NGTHETA = 180.

Polyrate I/O FILES

FILE NAME

ch5j1tr11.dat
ch5j1tr11.fu6
ch5j1tr11.fu14
ch5j1tr11.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat

UNIT = 4
FILE NAME = potch5.dat

16.C.12. Test run ch5j2tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with harmonic approximation and ES1 method

This sample run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with the J2 potential energy surface is similar to the one described in Subsection 16.C.1. In addition to employing a different potential energy surface, a major difference between this test run and test run ch5j1tr1 is the method of calculating the MEP. Also, the LCG3 calculation is omitted in this run. To calculate the MEP, a step size of 0.001 bohr is used for the first five steps from the saddle point [SFIRST] and then a larger step size (0.01 bohr) [SSTEP] is used for the remaining steps in combination with the Euler stabilization (ES1) method [RPM = *es1*]. In addition, sixth-order Lagrangian interpolation is used to obtain the values of the effective mass for the CD-SCSAG tunneling calculation [SCTOPT : *lagrange*].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j2tr1.dat	poly.fu5	input data file
ch5j2tr1.fu6	poly.fu6	long output file
ch5j2tr1.fu14	poly.fu14	summary output file
ch5j2tr1.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j2.dat UNIT = 4
FILE NAME = potch5.dat

16.C.13. Test run ch5j1tr12

CH₃ + HPh → CH₄ + Ph with 2D interpolated LCG4 with all energetically accessible excited states.

In this run the CH₃ + HPh → CH₄ + Ph thermal rate constants are calculated with the LCT4 method with all energetically accessible vibrational excited states. As in the previous run Ph denotes a single particle with the mass of the phenyl group. The only difference with previous test is that both the tunneling energies and the points along each tunneling energies are obtained from a grid specified by LCTGRD and fitted to a 2D spline under tension [option ILCT2D in LCTOPT]. The rectangular grid was built from 7 tunneling energies and 11 points along each tunneling energy. It is interesting to compare this run with the previous one, which uses a 1D interpolation and with test ch5j1tr10, which corresponds to the full LCT4 calculation.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5j1tr12.dat	poly.fu5	input data file
ch5j1tr12.fu6	poly.fu6	long output file
ch5j1tr12.fu14	poly.fu14	summary output file
ch5j1tr12.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1	UNIT = 4
	FILE NAME = potch5.dat

16.C.14. Test run ch5j2tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with MEP only

This sample run is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. The RATE and TUNNEL sections are not given so the rate constants are not calculated. The MEP is calculated using the ESD method [RPM = *esd*]. The MEP information is printed [PRINTSTEP] at grid points separated by 0.01 bohr [SSTEP]. The overall translations and rotations are projected out at stationary points [PROJECT].

Polyrate I/O FILES

FILE NAME

Polyrate FILE NAME

ch5j2tr2.dat
ch5j2tr2.fu6

poly.fu5 input data file
poly.fu6 long output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j2.dat

UNIT = 4
FILE NAME = potch5.dat

16.C.15. Test run ch5j2tr3

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with WKB approximation for two modes

This sample run is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using a combination of methods for treating the various normal modes [VANHAR]. All reactants and products are treated harmonically. Along the reaction path the two lowest frequency modes are treated using the uniform WKB method with a quadratic-quartic fit to the potential at two input distances along the generalized normal modes [QQSEMI] and the rest of the modes are treated using the harmonic approximation. The rate constants are calculated at five temperatures [TEMP], and four temperature pairs are used to compute activation energies [EACT]. Both CVT [CVT] and ICVT [ICVT] rate constants are calculated, using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT]. The ESD integrator [RPM = *esd*] is used to calculate the MEP, using a step size of 0.001 bohr [SSTEP]. The limits for the MEP are determined by using two keywords: SRANGE and SPECSTOP. A generalized normal mode analysis is performed every 0.01 bohr [INH = 10].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME
ch5j2tr3.dat	poly.fu5
ch5j2tr3.fu6	poly.fu6
ch5j2tr3.fu14	poly.fu14
ch5j2tr3.fu15	poly.fu15

input data file
long output file
summary output file
summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j2.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.16. Test run ch5j2itr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with zero-order IVTST approximation

This zero-order IVTST test run [IVTST0] is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using the energies, geometries, and vibrational frequencies of the reactants, saddle point, and products from the J2 potential of Joseph et al. [Ref. 17, Sect. 23].* All reactants and products are treated harmonically. The rate constants are calculated at four temperatures [TEMP]. This test run reproduces the results from one of the test cases in Ref. 26 in Sect. 23.

Polyrate I/O FILES

FILE NAME

ch5j2itr1.dat
ch5j2itr1.fu29
ch5j2itr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 long output file

* The data are input by means of the unit fu29 file.

16.C.17. Test run ch5j2itr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ with first-order IVTST approximation

This first-order IVTST test run [IVTST1] is for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using the energies, geometries, and vibrational frequencies of the reactants, saddle point, products, and a point on the MEP at $s = 0.01$ bohr from the J2 potential of Joseph et al. [Ref. 17, Sect. 23].* All reactants and products are treated harmonically. The rate constants are calculated at four temperatures [TEMP]. This test run reproduces the results from one of the test cases in Ref. 26 in Sect. 23.

Polyrate I/O FILES

FILE NAME

ch5j2itr2.dat
ch5j2itr2.fu29
ch5j2itr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 long output file

* The data are input by means of the unit fu29 file.

16.C.18. Test run ch5fu29tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ zero-order IVTST reaction rate with electronic structure data read from file fu29 and frequencies input directly.

This is a zero-order IVTST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [IVTST0]. Electronic structure information calculated at the UHF/STO-3G level (energies and frequencies) is read from unit fu29. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically and the frequencies are in cm^{-1} . The TST rate constants [TST, NOCVT] are calculated at four temperatures [TEMP] with zero-curvature transmission coefficients calculation. [ZCT] The zero-order IVTST calculation can also be done with the fu50 option. A similar test run (ch5fu50tr1) can be used to check the result.

Polyrate I/O FILES

FILE NAME

ch5fu29tr1.dat
ch5fu29tr1.fu29
ch5fu29tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 long output file

16.C.19. Test run ch5fu29tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ zero-order IVTST reaction rate with information read from fu29 and Hessian matrix input in packed form

This is a zero-order IVTST test run [TST,CVT] for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [IVTST0]. Electronic structure information calculated at the UHF/STO-3G level (Hessians of reactants, saddle point, and products in packed format) is read from file fu29. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically. Hessians and geometries are input in unscaled Cartesian coordinates in bohrs. The rate constants are calculated at four temperatures [TEMP]. A similar calculation has been done in the ch5tr29tr1 and ch5fu50tr1 test runs.

Polyrate I/O FILES

FILE NAME

ch5fu29tr2.dat
ch5fu29tr2.fu29
ch5fu29tr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 long output file

16.C.20. Test run ch5fu30tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with frequencies read from fu30 directly

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies and frequencies of reactants, saddle point, and products) is read from file fu30 [POTENTIAL = *unit30*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Harmonic frequencies of reactants and products are input in atomic units [LOPT(4) = 0 in unit fu30]. The rate constants are calculated at four temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu30rt1.dat
ch5fu30tr1.fu30
ch5fu30tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.21. Test run ch5fu30tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu30 and full Hessian matrices input

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies and full matrix format Hessians of reactants, saddle point, and products) is read from file fu30 [POTENTIAL = *unit30*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically. Hessians and geometries are input in unscaled Cartesian coordinates in atomic units [LOPT(4) = 1 and LOPT(3) = 0, respectively, in unit fu30]. The rate constants are calculated at four temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu30tr2.dat
ch5fu30tr2.fu30
ch5fi30tr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.22. Test run ch5fu30tr3

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu30 and Hessian matrix input in packed form

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies and Hessians of reactants, saddle point, and products in packed format) is read from file fu30 [POTENTIAL = *unit30*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically (some anharmonicity options are not supported for conventional TST calculations in *POLYRATE*-2017). Hessians are input in unscaled Cartesian coordinates in atomic units [LOPT(4) = -1 in unit fu30]. The rate constants are calculated at four temperatures [TEMP].

The electronic structure information for this run was generated with the GAMESS electronic structure package (Aug. 11, 1994 – IBM version), which yields an H_2 stretching frequency of 5479.11 cm^{-1} . *POLYRATE*, with this input, yields 5479.18 cm^{-1} . The GAUSSIAN94 electronic structure package yields a slightly different Hessian and a frequency of 5478.64 cm^{-1} .

Notice that the Hessian for H_2 corresponds to the molecule being on the z -axis, whereas the geometry corresponds to the molecule being on the y -axis. For calculating reactant frequencies this does not matter, but such an inconsistency would matter along the reaction path (see Section 9.B). It is worthwhile to point out that commonly-available electronic structure packages may, by default, re-orient a molecule before computing its Hessian.

Polyrate I/O FILES

FILE NAME

ch5fu30tr3.dat
ch5fu30tr3.fu30
ch5fu30tr3.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.23. Test run ch5fu30tr4

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu30 and frequencies input directly

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information (energies and frequencies of the reactants, the products, and the saddle point) at the QCISD/6-311G(d, p) level is read from file fu30 [POTENTIAL = *unit30*]. (The $V_{\text{MEP}}(s)$ has been scaled by a factor of 0.86 to be consistent with Ref. 48 in Section 22. The scaling was done by hand; however, the VSCALE keyword could have been used instead of hand scaling) At this level of electronic structure theory, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF=4]. Harmonic frequencies of reactants and products are input in atomic units and geometries are input in unscaled Cartesian coordinates in bohrs. The rate constants are calculated at twelve temperatures [TEMP].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5fu30tr4.dat	poly.fu5	input data file
ch5fu30tr4.fu30	poly.fu30	input data file
ch5fu30tr4.fu6	poly.fu6	long output file

16.C.24. Test run ch5fu30tr5

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST and CVT reaction rates with information along the reaction path read from fu30

This is a sample run that illustrates the use of electronic structure input file data from unit fu30 [POTENTIAL = *unit30*] in the calculation of rate constants for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. Electronic structure information read from file fu30 (energies, geometries, and Hessians of reactants, saddle point, products and a series of 60 points along the reaction path) was calculated at the UHF/STO-3G level using GAMESS for generating the reaction path (ES1 integrator, mass-weighted Cartesian coordinates, step size = 0.005 amu^{1/2} a₀) and GAUSSIAN94 for calculating the unscaled forces and Hessians (run with geometry input in Cartesian coordinates and the NOSYMM keyword on) at the generalized TST structures every 0.05 amu^{1/2} bohr along the reaction path. The strategy for producing the electronic structure data was carefully selected in order to avoid the reorientation of geometries along the reaction path though the different steps of the process. At this level of electronic structure theory, CH₃ is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Geometries and forces are input in unscaled Cartesian coordinates in atomic units [LOPT(3) = 100 in unit fu30]. Hessians are also input in unscaled Cartesian coordinates in bohr using the packed format as obtained from the GAUSSIAN94 formatted checkpoint file [LOPT(4) = 99]. All normal modes are treated harmonically and TST, CVT, and ICVT rate constants [CVT and ICVT] are calculated at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling transmission factors [ZCT, SCT]. Curvature components are calculated using the derivative of the gradient with respect to *s* obtained from a quadratic fit of the gradient at three points on the basic input grid [LOPT(6) = 2] which is also called the sparse input grid. Note that the results obtained with this test run differ from those previously reported in Ref. 20 in Section 23. Several errors were detected in Ref. 20 (see Section 24.B in this manual for the corresponding errata).

Polyrate I/O FILES

FILE NAME

ch5fu30tr5.dat
ch5fu30tr5.fu30
ch5fu30tr5.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.25. Test run ch5fu30tr6

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST and CVT reaction rates with information along the reaction path read from fu30. Frequencies evaluated in redundant internal coordinates.

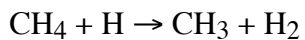
This is a sample run that illustrates the use of electronic structure input file data from unit fu30 [POTENTIAL = *unit30*] in the calculation of rate constants for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. Electronic structure information calculated at the UHF/STO-3G level following the procedure explained in test run ch5fu30tr5 (energies, geometries, and Hessians of reactants, products, saddle point, and a series of 50 points along the reaction path) is read from file fu30. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Geometries and forces are input in unscaled Cartesian coordinates in atomic units [LOPT(3) = 100 in unit fu30]. Hessians are also input in unscaled Cartesian coordinates in atomic units using the packed format as obtained from the GAUSSIAN94 formatted checkpoint file [LOPT(4) = 99]. All normal modes are treated harmonically in redundant internal coordinates [COORD = *curv3*], and TST, CVT, and ICVT rate constants [CVT and ICVT] are calculated at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling transmission factors [ZCT, SCT].

The ‘Cartesian Forces’ from the GAUSSIAN94 formatted checkpoint file are actually Cartesian Gradients; use of them as forces will introduce errors in the transformation from rectilinear to curvilinear internal coordinates. Also, the gradients (or forces) must be un-normalized in order to carry out the transformation correctly.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5fu30tr6.dat	poly.fu5	input data file
ch5fu30tr6.fu30	poly.fu30	input data file
ch5fu30tr6.fu6	poly.fu6	long output file

16.C.26. Test run ch5fu31tr1



IVTST-M-56/6 calculation using fu31

This sample run performs a IVTST-M-56/6 calculation using an fu31 input file. The curvature components are interpolated, the vibrational analysis is carried out in redundant internal coordinates, and ICVT rate constants are calculated. The information in the fu31 input file was obtained from the J1 potential energy surface by means of the WRITEFU31 keyword. Only the forward rate constant were calculated.

Polyrate I/O FILES

FILE NAME

ch5fu31tr1.dat
ch5fu31tr1.fu31
ch5fu31tr1.fu6
ch5fu31tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
 poly.fu31 input data file
poly.fu6 long output file
 poly.fu15 summary output file

16.C.27. Test run ch5fu40tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with frequencies read from fu40 directly

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies, and frequencies of reactants, saddle point, and products) is read from file fu40 [POTENTIAL = *unit40*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Harmonic frequencies of reactants, products, and saddle point are input in atomic units [FREQSOURCE = *read* and FREQUIT = *au* in unit fu40] and geometries are input in unscaled Cartesian coordinates in bohrs [GEOMUNIT = *bohr* in unit fu40]. The rate constants are calculated at four temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu40rt1.dat
ch5fu40tr1.fu40
ch5fu40tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.28. Test run ch5fu40tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu40 and full Hessian matrix for frequency input

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies and full matrix format Hessians [HESSFORM = *full*] of reactants, saddle point, and products) is read from file fu40 [POTENTIAL = *unit40*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically. Hessians and geometries are input in unscaled Cartesian coordinates in atomic units [HESSUNIT = *bohr* and GEOMUNIT = *bohr*, respectively, in unit fu40]. The rate constants are calculated at four temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu40tr2.dat
ch5fu40tr2.fu40
ch5fi40tr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.29. Test run ch5fu40tr3

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu40 and Hessian matrix input in packed form

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level (energies and Hessians of reactants, saddle point, and products in packed format [HESSFORM = *packed*]) is read from file fu40 [POTENTIAL = *unit40*]. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. All reactants and products are treated harmonically. Hessians and geometries are input in unscaled Cartesian coordinates in bohrs [HESSUNIT = *bohr* and GEOMUNIT = *bohr*, respectively, in unit fu40]. The rate constants are calculated at four temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu40tr3.dat
ch5fu40tr3.fu40
ch5fu40tr3.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.30. Test run ch5fu40tr4

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST reaction rate with information read from fu40 and frequencies input directly.

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information (energies and frequencies of reactants, saddle point, and products) at the QCISD/6-311G(d, p) level is read from file fu40 [POTENTIAL = *unit40*]. (The $V_{\text{MEP}}(s)$ has been scaled by a factor of 0.86 to be consistent with Ref. 48 in Section 22. The scaling was done by hand; however, the VSCALE keyword could have been used instead of hand scaling) At this level of electronic structure theory, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF=4]. Harmonic frequencies of reactants and products are input in wavenumbers [FREQSOURCE = *read*, FREQUNIT = *waven*]. The rate constants are calculated at twelve temperatures [TEMP].

Polyrate I/O FILES

FILE NAME

ch5fu40tr4.dat
ch5fu40tr4.fu40
ch5fu40tr4.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.31. Test run ch5fu40tr5

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST and CVT reaction rates with information along the reaction path read from fu40

This is a sample run that illustrates the use of electronic structure input file data from unit fu40 [POTENTIAL = *unit40*] in the calculation of rate constants for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. Electronic structure information read from file fu40 (energies, geometries, and Hessians of reactants, products, saddle point, and a series of 60 points along the reaction path) was calculated at the UHF/STO-3G level using GAMESS for generating the reaction path (ES1 integrator, mass-weighted Cartesian coordinates, step size = 0.005 amu^{1/2} a₀) and GAUSSIAN94 for calculating the unscaled forces and Hessians (run with geometry input in Cartesian coordinates and the NOSYMM keyword on) at the generalized TST structure every 0.05 amu^{1/2} bohr along the reaction path. The strategy for producing the electronic structure data was carefully selected in order to avoid the reorientation of geometries along the reaction path through the different steps of the process. At this level of electronic structure theory, CH₃ is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Geometries and forces are input in unscaled Cartesian coordinates in atomic units [GEOMUNIT = *bohr* and GRADUNIT = *bohr*, respectively, in unit fu40]. Hessians are also input in unscaled Cartesian coordinates in atomic units using the packed format as obtained from the GAUSSIAN94 formatted checkpoint file [HESSUNIT = *bohr* and HESSFORM = *packed*]. All normal modes are treated harmonically and TST, CVT, and ICVT rate constants [CVT and ICVT] are calculated at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling transmission factors [ZCT, SCT]. Curvature components are calculated using the derivative of the gradient with respect to *s* obtained from a quadratic fit of the gradient at three points on the basic input grid [CURVATURE = *compute*, GRADDER = *noextra*], which is also called the sparse input grid. Note that the results obtained with this test run differ from those previously reported in Ref. 20 in Section 23. Several errors were detected in Ref. 20 (see Section 24.B in this manual for the corresponding errata).

Polyrate I/O FILES

FILE NAME

ch5fu40tr5.dat
ch5fu40tr5.fu40
ch5fu40tr5.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.32. Test run ch5fu40tr6

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ TST and CVT reaction rates with information along the reaction path read from fu40. Frequencies are evaluated with redundant internal coordinates.

This is a sample run that illustrates the use of electronic structure input file data from unit fu40 [POTENTIAL = *unit40*] in the calculation of rate constants for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. Electronic structure information calculated at the UHF/STO-3G level following the procedure explained in test run ch5fu40tr5 (energies, geometries, and Hessians of reactants, saddle point, products and a series of 50 points along the reaction path) is read from file fu40. At this level of electronic structure theory, CH_3 is nonplanar; therefore the symmetry number of the forward reaction is 2 [SIGMAF=2]. Geometries and forces are input in unscaled Cartesian coordinates in atomic units [GEOMUNIT = *bohr* and GRADUNIT = *bohr*, respectively, in unit fu40]. Hessians are also input in unscaled Cartesian coordinates in atomic units using the packed format as obtained from the GAUSSIAN94 formatted checkpoint file [HESSUNIT = *bohr* and HESSFORM = *packed*]. All normal modes are treated harmonically in redundant internal coordinates [*curv3*] and TST, CVT and ICVT rate constants [CVT and ICVT] are calculated at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling transmission factors [ZCT, SCT].

The quantities labeled as “Cartesian forces” in the GAUSSIAN94 formatted checkpoint file are actually Cartesian gradients, and using them as forces will introduce errors in the transformation from rectilinear to curvilinear internal coordinates. Also, the gradients (or forces) must be un-normalized in order to carry out the transformation correctly.

Polyrate I/O FILES

FILE NAME

ch5fu40tr6.dat
ch5fu40tr6.fu40
ch5fu40tr6.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.33. Test run ch5fu50tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ zero-order IVTST reaction rate with information read from fu50 and frequencies input directly.

This is a conventional TST test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ [NOCVT and TST]. Electronic structure information calculated at the UHF/STO-3G level is read from file fu50 [IC in unit fu5 and 0IVTST in unit fu50]. All reactants and products are treated harmonically and the frequencies are input in cm^{-1} [RCFREQA, PCFREQA, SADFREQA, and FREQIMAG]. A rate constants are calculated at four temperatures [TEMP]. The similar calculation can be performed with the fu29 option. Both the ch5fu29tr1 and ch5fu29tr2 test runs can be used for checking the output from different input methods.

Polyrate I/O FILES

FILE NAME

ch5fu50tr1.dat
ch5fu50tr1.fu50
ch5fu50tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu50 input data file
poly.fu6 long output file

16.C.34. Test run ch5fu50tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-IOC dual-level reaction rate calculation with information for the higher level read from fu50 and J1 analytic surface as the lower level.

This is a VTST-IOC test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using an analytic surface as the lower level [IC and POTENTIAL = *hooks*]. The potential energy is not corrected in the nonadiabatic region in the LCT calculation [LTCOR = *nocor*]. Electronic structure information calculated at the UHF/STO-3G level is read from file fu50 and the lower-level information is obtained from the J1 surface. All the parameters are the same as for test run ch5j1tr1.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5fu50tr2.dat	poly.fu5	input data file
ch5fu50tr2.fu50	poly.fu50	input data file
ch5fu50tr2.fu6	poly.fu6	long output file
ch5fu50tr2.fu14	poly.fu14	summary output file
ch5fu50tr2.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat UNIT = 4
 FILE NAME = potch5.dat

16.C.35. Test run ch5fu50tr3

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-IOC dual-level reaction rate calculation with information for the higher level read from fu50 and J1 analytic surface as the lower level.

This is a VTST-IOC test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using an analytic surface as the lower level [IC and POTENTIAL = *hooks*]. The potential energy is corrected in the nonadiabatic region in the LCT calculation based on a quadratic correction [LCTCOR = *quadlin*]. Electronic structure information calculated at the UHF/STO-3G level is read from file fu50 and the lower-level information is obtained from the J1 surface. All the parameters are the same as for test run ch5j1tr1.

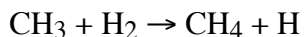
Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5fu50tr3.dat	poly.fu5	input data file
ch5fu50tr3.fu50	poly.fu50	input data file
ch5fu50tr3.fu6	poly.fu6	long output file
ch5fu50tr3.fu14	poly.fu14	summary output file
ch5fu50tr3.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat	UNIT = 4
	FILE NAME = potch5.dat

16.C.36. Test run ch5fu51tr1



VTST-ISPE calculation with MP2/6-31G** and J1

This is a VTST-ISPE test run for the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ using an analytic surface as the lower-level [dl and potential = hooks]. Electronic structure information calculated at the MP2/6-31G** level is read in from file fu51 and the lower-level information is obtained from the J1 surface. Two extra points plus the stationary points used for the VTST-ISPE calculation were obtained from single-point energy calculations at the MP2/6-31G** level with the configurations obtained from the lower-level surface. All the parameters for the lower-level are the same as for the test run ch5j1tr1.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5fu51tr1.dat	poly.fu5	input data file
ch5fu51tr1.fu51	poly.fu51	VTST-
ISPE input file		
ch5fu51tr1.fu6	poly.fu6	long output
file		
ch5fu51tr1.fu14	poly.fu14	
summary output file		
ch5fu51tr1.fu15	poly.fu15	
summary output file		

POTENTIAL ENERGY SURFACE DATA FILE

potch5j1.dat

UNIT=4

FILE NAME=potch5.dat

16.C.37. Test run ch5icfu30tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-ICL-DECKART calculation using file fu30 for the lower level information and rectilinear coordinates

This sample run uses the *unit30* [POTENTIAL = *unit30*] and the IOC [ICOPT: *icl*, *deckart*] options to invoke a dual-level direct dynamics calculation (variational transition state theory with interpolated optimized corrections) on the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. The lower-level reaction path file is constructed at the UHF/STO-3G level, and the higher-level interpolated optimized corrections are calculated from the QCISD(T)/6-311G(d, p) level results. (The $V_{\text{MEP}}(s)$ has been scaled by a factor of 0.86 to be consistent with Ref. 48 in Section 22. The scaling was done by hand; however, the VSCALE keyword could have been used instead of hand scaling) All the modes are treated harmonically. Cartesian coordinates are used [COORD = *cart*]. Both small-curvature and zero-curvature tunneling calculations are carried out [ZCT, SCT]. The interpolated optimized correction is based on the logarithm of ratios [ICOPT: *icl*] and an Eckart fit is used to the energies at 3 stationary points (i.e. reactants, products, and saddle point) [ICOPT: *deckart*]. At the higher level of electronic structure theory employed, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF=4].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ch5icfu30tr1.dat	poly.fu5	input data file
ch5icfu30tr1.fu30	poly.fu30	input data file
ch5icfu30tr1.fu50	poly.fu50	input data file
ch5icfu30tr1.fu6	poly.fu6	output file

16.C.38. Test run ch5icfu30tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-IOC calculation using file fu30 for the lower level information and curvilinear coordinates

This sample run uses the *unit30* option [POTENTIAL = *unit30*] and the IOC [ICOPT: *ica, seckart*] options to invoke a dual-level direct dynamics calculation with interpolated optimized corrections on the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. The lower-level reaction path file is constructed at the UHF/STO-3G level, and the higher-level interpolated optimized corrections are calculated from the QCISD(T)/6-311G(d, p) level results. (The $V_{\text{MEP}}(s)$ has been scaled by a factor of 0.86 to be consistent with Ref. 48 in Section 22. The scaling was done by hand; however, the VSCALE keyword could have been used instead of hand scaling) The original ICA interpolated correction method is used for the IC method. At the higher level of electronic structure theory employed, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF = 4].

Polyrate I/O FILES

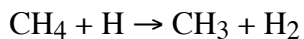
FILE NAME

ch5icfu30tr2.dat
ch5icfu30tr2.fu30
ch5icfu30tr2.fu50
ch5icfu30tr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu50 input data file
poly.fu6 output file

16.C.39. Test run ch5icfu31tr1



VTST-IOC calculation using IVTST-M-56/6 data for the lower level of calculation and using the fu31 input file

This sample test run uses the *unit31* and the IOC options to perform a dual level direct dynamics calculation with interpolated optimized corrections. The lower level is an IVTST-M-6/56 calculation based on the information from a fu31 input file. The higher-level interpolated optimized corrections are calculated at the QCISD(T)/6-311G(d, p) level with $V_{\text{MEP}}(s)$ scaled by a factor of 0.86, as taken from Ref. 48 in Section 22. The ICA method is used for the correction of the frequencies, and the SECKART option is used for the energy correction.

Polyrate I/O FILES

FILE NAME

ch5fu31tr1.dat
ch5fu31tr1.fu31
ch5fu31tr1.fu50
ch5fu31tr1.fu6
ch5fu31tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
 poly.fu31 input data file
 poly.fu50 input data file
poly.fu6 long output file
 poly.fu15 summary output file

16.C.40. Test run ch5icfu40tr1

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-ICL-DECKART calculation using file fu40 for the lower level information and rectilinear coordinates

This sample run uses the *unit40* [POTENTIAL = *unit40*] and the IC [ICOPT: *ic1*, *deckart*] options to invoke a dual-level direct dynamics calculation (variational transition state theory with interpolated optimized corrections) on the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. The lower-level reaction path file is constructed at the UHF/STO-3G level, and the higher-level interpolated optimized corrections are calculated from the QCISD(T)/6-311G(d, p) level results with $V_{\text{MEP}}(s)$ scaled by a factor of 0.86 taken from Ref. 48 in Section 22. All the modes are treated harmonically. Cartesian coordinates are used [COORD = *cart*]. Both small-curvature and zero-curvature tunneling calculations are carried out [ZCT, SCT]. The interpolated correction is based on the logarithm of ratios [ICOPT: *ic1*]. At the higher level of electronic structure theory employed, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF = 4].

Polyrate I/O FILES

FILE NAME

ch5icfu40tr1.dat
ch5icfu40tr1.fu40
ch5icfu40tr1.fu50
ch5icfu40tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu50 input data file
poly.fu6 output file

16.C.41. Test run ch5icfu40tr2

$\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ VTST-IOC calculation using file fu40 for the lower level information

This sample run uses the *unit40* [POTENTIAL = *unit40*] and the IOC [ICOPT: *ica*, *seckart*] options to invoke a dual-level direct dynamics calculation with interpolated optimized corrections on the reaction $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$. The lower-level reaction path file is constructed at the UHF/STO-3G level, and the higher-level interpolated optimized corrections are calculated from the QCISD(T)/6-311G(d, p) level results with $V_{\text{MEP}}(s)$ scaled by a factor of 0.86 taken from Ref. 48 in Section 22. The original ICA interpolated correction procedure is used for the IOC method. At the higher level of electronic structure theory employed, CH_3 is planar; therefore the symmetry number of the forward reaction is 4 [SIGMAF = 4].

Polyrate I/O FILES

FILE NAME

ch5icfu40tr2.dat
ch5icfu40tr2.fu40
ch5icfu40tr2.fu50
ch5icfu40tr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu50 input data file
poly.fu6 output file

16.C.42. Test run `clhbrtr1`

$\text{Cl} + \text{HBr} \rightarrow \text{HCl} + \text{Br}$ with LCG3 calculations for all energetically accessible excited states in the LCG3 exit arrangement, and with the zero of energy of the reactants determined by the program

This sample run is for the reaction $\text{Cl} + \text{HBr} \rightarrow \text{HCl} + \text{Br}$ with an extended LEPS surface. All modes are treated using the default harmonic approximation. TST and CVT rate constants are computed [TST, CVT]. The tunneling calculations of ground-state transmission coefficients are performed using the MEPSAG, CD-SCSAG, and LCG3 methods [ZCT, SCT, LCTOPT]. The MEP is followed with the ESD integrator [RPM = *esd*] with a step size of 0.00001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.01 bohr [INH = 1000]. For the LCG3 calculation, the highest energetically open vibrational state in the LCG3 exit arrangement (HCl + Br) is determined and then all open vibrational states are included in the tunneling calculations [ALLEXCIT]. (As always, for LCG3 calculations only the ground state is considered for the LCG3 entrance arrangement, which is the reactant arrangement of the exoergic direction.) Also, the theta integrals, the vibrationally adiabatic potential, the vibrational period, and the sine of the angle between the MEP and the tunneling path at NGS0 points between each pair of adjacent energy quadrature points are obtained by 2-point interpolation [LCTOPT: *interpolate*]. This is done because the adiabatic potential of the first vibrational excited state is not monotonic on the reactant side and, therefore, unphysical results are obtained using interpolation with an order higher than 2. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. The zero of energy of the reactants is calculated from the energies of the individual molecular reactant species; in other words, the default value of EZERO = *calculate* is used. Note that the degeneracies and energies of the $^2\text{P}_{3/2}$ and $^2\text{P}_{1/2}$ states of Cl and Br are used in the calculation of the reactant and product electronic partition functions. However, for the transition state only the ground electronic state needs to be considered, because all excited electronic states are too high in energy to have a significant effect.

The results differ from those in Ref. 10 in Sect. 23 because Ref. 10 included anharmonicity and treated the bend in curvilinear internal coordinates, whereas the present run is harmonic and treats all vibrations in mass-scaled Cartesians. The treatment in Ref. 10 in Sect. 23 is more reliable.

Polyrate I/O FILES

FILE NAME

clhbrtr1.dat
clhbrtr1.fu6
clhbrtr1.fu14
clhbrtr1.fu15

Polyrate FILE NAME

poly.fu5	input data file
poly.fu6	long output file
poly.fu14	summary output file
poly.fu15	summary output file

16.C.43. Test run clhbrtr2

Cl + HBr \rightarrow HCl + Br with LCG3 calculations in which $n = 3$ is the highest excited state that is included in the LCG3 exit arrangement, and with the zero of energy of the reactants specified in the input

This sample run is similar to the clhbrtr1 test run. All vibrational modes are treated using the default harmonic approximation. TST and CVT rate constants are computed [TST, CVT]. The tunneling calculations of ground-state transmission coefficients are performed using the MEPSAG, CD-SCSAG, and LCG3 methods [ZCT, SCT, LCTOPT]. For the LCG3 calculation, the $n = 0, 1, 2,$ and 3 vibrational states in the LCG3 exit arrangement (HCl + Br) are included in the tunneling calculation [EXCITED = 3]. The highest energetically open vibrational state N_p^{\max} is determined [Ref. 30 in Sect. 23], and the LCG3 tunneling calculation includes the $n = 0$ to $\min[3, N_p^{\max}]$ vibrational states in the LCG3 exit arrangement. (For LCG3 calculations, only the ground state is considered for the LCG3 entrance arrangement, which is the reactant arrangement of the exoergic direction.) The MEP is followed by the ESD method [RPM = *esd*] with a step size of 0.00001 bohr [SSTEP] and a generalized normal mode analysis is performed every 0.01 bohr [INH = 1000]. Just as for test run clhbrtr1, we use 2-point interpolation [LCTOPT: *interpolate*] for the various quantities needed in the LCG3 calculations. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. The energy of the reactants, which is taken as the overall zero of energy, is read from unit fu5 [EZERO = *read*]. The results should be the same as those obtained in the clhbrtr1 test run.

Polyrate I/O FILES

FILE NAME

clhbrtr2.dat
clhbrtr2.fu6
clhbrtr2.fu14
clhbrtr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.44. Test run clhbrtr3

Cl + HBr \rightarrow HCl + Br with LCG4 calculation for all the energetically accessible excited states in the LCG4 exit rearrangement, and with the zero of energy determined by the program

This sample is similar to the clhbrtr1 run. All vibrational modes are treated within the harmonic approximation. The only difference with test run clhbrtr1 is that large curvature transmission coefficients are obtained using the LCG4 method. At each tunneling energy a number of points along the straight path is NGTHETA = 140. The LCG4 tunneling probabilities are evaluated at NGAMP = 40 tunneling paths.

Polyrate I/O FILES

FILE NAME

clhbrtr3.dat
clhbrtr3.fu6
clhbrtr3.fu14
clhbrtr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.45. Test run clhbrtr4

Cl + HBr \rightarrow HCl + Br with interpolated LCG4 calculation for all the energetically accessible excited states in the LCG4 exit rearrangement, and with the zero of energy determined by the program

This sample is similar to the previous clhbrtr3 test run. All vibrational modes are treated within the harmonic approximation. The difference with the previous run is that an interpolation is carried out, at each tunneling energy, by a splines-under-tension procedure. In this case only 3 points were considered at each tunneling energy [keywords ILCT and NILCGIT = 3]. Two of the three points are located at the beginning of the nonadiabatic region and the third point is located at half the distance between the two limiting points. This procedure still makes use of the keyword NGTHETA = 140 but the needed values are produced by the spline interpolation procedure. As in the previous run the calculation takes into account all the possible open vibrational states of the exit channel.

Polyrate I/O FILES

FILE NAME

clhbrtr4.dat
clhbrtr4.fu6
clhbrtr4.fu14
clhbrtr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.46. Test run clhbr1tr5

Cl + HBr \rightarrow HCl + Br with interpolated LCG4 calculation with all the energetically accessible excited states in the LCG4 exit rearrangement. Restart from test run clhbrtr3.

This sample is the similar to the previous test run. The only differences are that this test run makes use of the ILCG2D interpolation option and that a restart file is used to build the 2D rectangular grid. In this case a quite rough grid of 5 5 is specified by LCTGRD.

Polyrate I/O FILES

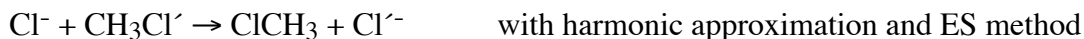
FILE NAME

chbrtr5.dat
chbrtr5.fu6
chbrtr5.fu14
chbrtr5.fu15
chbrtr5.fu48

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file
poly.fu48 LCT restart file

16.C.47. Test run cmctr1



This sample run for the reaction $\text{Cl}^- + \text{CH}_3\text{Cl}^* \rightarrow \text{ClCH}_3 + \text{Cl}^-$ uses the default harmonic approximation for all vibrational modes. The MEP is calculated by the ESD integrator [RPM = *esd*] with a step size of 0.005 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.025 bohr [INH = 5]. Both CD-SCSAG and MEPSAG tunneling probabilities are calculated [ZCT, SCT].

This test run and the next one (cmctr2) form a pair. This one must be run before the next one to find the *s* values of the wells that correspond to complexes on the reactant and product sides of the MEP. The interpolation in test run cmctr2 on both sides of the MEP is based on these wells, and thus we must first find the *s* values at these wells. A relatively long range of the MEP ($s = -7.1$ bohr to $s = 7.1$ bohr) is calculated [SRANGE] and we find that the *s* values at the wells are approximately 7.0 bohr and -7.0 bohr. We also learn from this run that in order for the tunneling calculation to converge, the MEP should be calculated at least from $s = -1.5$ bohr to $s = 1.5$ bohr. The saddle point frequencies match the frequencies of the complexes in descending order (except for the one corresponding to the reaction coordinate). We then use the information above together with some ab initio data to make the input data file for the VTST-IOC calculation in the cmctr2 test run.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME
cmctr1.dat	poly.fu5
cmctr1.fu6	poly.fu6
cmctr1.fu14	poly.fu14
cmctr1.fu15	poly.fu15
	input data file
	long output file
	summary output file
	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potcmc1.dat	UNIT = 4
	FILE NAME = potcmc1.dat
potcmc2.dat	UNIT = 7
	FILE NAME = potcmc2.dat

16.C.48. Test run cmctr2

$\text{Cl}^- + \text{CH}_3\text{Cl} \rightarrow \text{ClCH}_3 + \text{Cl}^-$ harmonic approximation and VTST-IOC calculation

This sample run is similar to the cmctr1 test run except we perform a VTST-IOC calculation in addition to the VTST calculation [ICOPT] .

This reaction has two reactants and two products, but the interpolation on both sides of the saddle point is based on wells that correspond to complexes. We set the complex corresponding to the well on the product side to be the product so we can get its properties, and thus we set MEPTYPEP to *well* and MEPTYPEP to *one*. The correct energy, frequency, and moment-of-inertia-tensor determinant information at reactants, products, saddle point, and the ion-molecule complexes are obtained from the calculations of Tucker and Truhlar [Ref. 34, Sect. 22]. The FREQMAT keyword is used, and the frequency matching information and the *s* value for the ion-molecule complex are obtained from the cmctr1 test run. The range parameters for the unit fu50 input are obtained by the utility programs findl.f and findb.f (see Subsection 17.D) by using additional points on the MEP that have an energy halfway between the energy of the saddle point and the energy of reactant- or product-side stationary point.

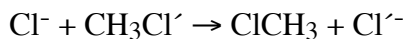
Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
cmctr2.dat	poly.fu5	input data file
cmcic.dat	poly.fu50	input data file
cmctr2.fu6	poly.fu6	long output file
cmctr2.fu14	poly.fu14	summary output file
cmctr2.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potcmc1.dat	UNIT = 4 FILE NAME = potcmc1.dat
potcmc2.dat	UNIT = 7 FILE NAME = potcmc2.dat

16.C.49. Test run cmctr3



VTST-ISPE correction with MP2/6-31G**

This sample test run for the reaction $\text{Cl}^- + \text{CH}_3\text{Cl} \rightarrow \text{ClCH}_3 + \text{Cl}^-$ uses the default harmonic approximation for all vibrational modes. The MEP is followed by the Euler steepest-descent integrator [ESD] with a step size of 0.005 bohr [SSTEP], and a generalized normal mode analysis is performed every 5 gradient steps [INH]. Both CD-SCSAG and MEPSAG tunneling probabilities are calculated [ZCT, SCT].

This test run corrects the energies along the MEP by the VTST-ISPE method. The energies of two extra points plus the stationary points were evaluated at the MP2/6-31G** level with the configuration taken from the potential energy surface. The MEP is calculated from $s = -1.5$ bohr to $s = 1.5$ bohr.

Polyrate I/O FILES

FILE NAME

cmctr3.dat
cmctr3.fu31
cmctr3.fu6
cmctr3.fu14
cmctr3.fu15

Polyrate FILE NAME

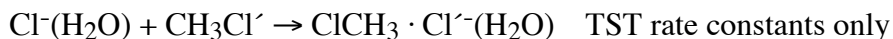
poly.fu5 input data file
poly.fu51 VTST-ISPE input file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potcmc1.dat
potcmc2.dat

UNIT = 4
FILE NAME = potcmc1.dat
UNIT = 7
FILE NAME = potcmc2.dat

16.C.50. Test run cwmctr1



In this sample run the reactant, product, and saddle point properties for the reaction $\text{Cl}^-(\text{H}_2\text{O}) + \text{CH}_3\text{Cl} \rightarrow \text{ClCH}_3 \cdot \text{Cl}^-(\text{H}_2\text{O})$, where $\text{Cl}^-(\text{H}_2\text{O})$ represents a chloride ion in the presence of a water molecule, are calculated, and only forward TST rate constants [TST, FORWARDK, NOCVT] are determined. No transmission coefficients are calculated nor is the minimum energy path determined (RATE and TUNNEL sections are not used). All normal modes and generalized normal modes are treated using the default harmonic approximation.

Polyrate I/O FILES

FILE NAME

cwmctr1.dat
cwmctr1.fu6
cwmctr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu15 summary output file

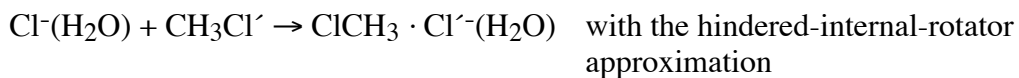
POTENTIAL ENERGY SURFACE DATA FILES

potcwmc3b.dat
potcwmcvib.dat
potcwmcvib.dat

UNIT = 4
UNIT = 7

FILE NAME = potcwmc3b.dat
FILE NAME =

16.C.51. Test run cwmctr2



In this sample run for the same reaction as in the cwmctr1 test run, the torsional mode of the generalized transition state is treated with the hindered-internal-rotator approximation [VANHAR, TOR]; all other modes are treated harmonically (default). The ESD integrator [RPM = *esd*] with a step size of 0.001 bohr [SSTEP] is used to follow the MEP, and a generalized normal mode analysis is performed every 0.01 bohr [INH = 10]. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. A restart file containing reaction-path information is written to unit fu1 [RESTART: WRITEFU1]. Rate constants are computed using CVT only [CVT]. The tunneling corrections are computed using the MEPSAG and CD-SCSAG methods [ZCT, SCT]. The rate constants are calculated for the same six temperatures as in the cwmctr1 test run [TEMP].

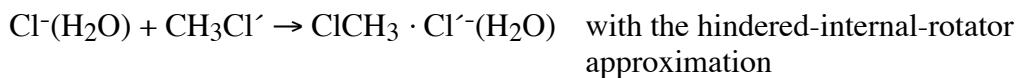
Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
cwmctr2.dat	poly.fu5	input data file
cwmctr2.fu6	poly.fu6	long output file
cwmctr2.fu1	poly.fu1	reaction path info.
cwmctr2.fu14	poly.fu14	summary output file
cwmctr2.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILES

potcwmc3b.dat	UNIT = 4	FILE NAME = potcwmc3b.dat
potcwmcvib.dat	UNIT = 7	FILE NAME = potcwmcvib.dat

16.C.52. Test run cwmctr3



In this sample run for the same reaction as in the cwmctr1 test run, the torsional mode of the generalized transition state is treated with the hindered-internal-rotator approximation with Pitzer's algorithm for the moment of inertia [VANHAR, TOR] and with $P = 3$ and $\sigma_1 = 3$; all other modes are treated harmonically (which is the default). The ESD integrator [RPM = *esd*] with a step size of 0.001 bohr [SSTEP] is used to follow the MEP, and a generalized normal mode analysis is performed every 0.01 bohr [INH = 10]. The curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. A restart file containing reaction-path information is written to unit fu1 [RESTART: WRITEFU1]. Rate constants are computed using CVT only [CVT]. The tunneling corrections are computed using the MEPSAG and CD-SCSAG methods [ZCT, SCT]. The rate constants are calculated for the same six temperatures as in the cwmctr1 test run [TEMP].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
cwmctr3.dat	poly.fu5	input data file
cwmctr3.fu6	poly.fu6	long output file
cwmctr3.fu1	poly.fu1	reaction path info.
cwmctr3.fu14	poly.fu14	summary output file
cwmctr3.fu15	poly.fu15	summary output file

POTENTIAL ENERGY SURFACE DATA FILES

potcwmc3b.dat	UNIT = 4	FILE NAME = potcwmc3b.dat
potcwmcvib.dat	UNIT = 7	FILE NAME = potcwmcvib.dat

16.C.53. Test run h3tr1

$\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with WKB and Morse IIIa approximations and Page-McIver integrator

This sample run uses the Page-McIver method [RPM = *pagem*] for calculating points along the MEP of the reaction $\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with a step size of 0.01 bohr [SSTEP], evaluates the Hessian matrix and performs a generalized normal mode analysis at every calculated point [INH = 1]. Note that the generalized transition state is restricted to linear symmetry [SPECIES = *lints*]. Two different methods are used to treat the vibrational modes [VANHAR]; vibrational modes of the reactant and product molecules and the highest frequency mode of the generalized transition state are treated by the Morse IIIa approximation [MORSE, MORMODEL = *morseiia*], while the two lowest vibrational modes of the generalized transition state are treated by the uniform WKB method applied to a quadratic-quartic fit to the potential along the normal mode coordinate [QQSEMI]. Extrapolation to both reactant and product limits is performed [EXFIRST, EXSECOND]. Both CVT [CVT] and ICVT [ICVT] rate constants are calculated only in the forward direction [FORWARDK] using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT], which are computed with two equal segments in the Boltzmann and theta integrals [NSEGBOLTZ = 2, NSEGTETA = 2].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
h3tr1.dat	poly.fu5	input data file
h3tr1.fu6	poly.fu6	long output file
h3tr1.fu14	poly.fu14	summary output file
h3tr1.fu15	poly.fu15	summary output file

16.C.54. Test run h3tr2

$\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with harmonic approximation and Page-McIver integrator

This sample run uses the Page-McIver method [RPM = *pagem*] for calculating points along the MEP of the reaction $\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with a step size of 0.005 bohr [SSTEP], evaluates the Hessian matrix and performs a generalized normal mode analysis at every 0.01 bohr [INH = 2]. The generalized transition states are restricted to linear symmetry [SPECIES = *lints*]. All vibrational modes are treated by the default harmonic approximation. Both CVT [CVT] and ICVT [ICVT] rate constants are calculated only in the forward direction [FORWARDK], using MEPSAG, CD-SCSAG, and LCG3 transmission coefficients [ZCT, SCT, LCTOPT], in which only the ground state is included in the LCG3 exit arrangement [NOALLEXCIT]. (We expect on physical grounds that only the ground state needs to be considered in this case, so this option is chosen to simplify the calculation.) The overall translations and rotations are projected out at stationary points; this is not necessary at stationary points, but it does improve the stability of the calculations.

Polyrate I/O FILES

FILE NAME

h3tr2.dat
h3tr2.fu6
h3tr2.fu14
h3tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.55. Test run h3tr3

$\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with harmonic approximation, Page-McIver integrator, and symmetric reaction path option

This sample run is the same as test run h3tr2 except that the symmetric reaction-path option [SYMMETRY] is used, and LCG3 transmission coefficients are not calculated, i.e., only MEPSAG and CD-SCSAG transmission coefficients are calculated [ZCT, SCT].

Polyrate I/O FILES

FILE NAME

h3tr3.dat
h3tr3.fu6
h3tr3.fu14
h3tr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.56. Test run h3tr4

$\text{H}_2 + \text{H}' \rightarrow \text{HH}' + \text{H}$ with WKB and Morse IIIa approximations, Page-McIver integrator, and symmetric reaction path option

This sample run is the same as test run h3tr1 except that the symmetric reaction-path option is used [SYMMETRY].

Polyrate I/O FILES

FILE NAME

h3tr4.dat
h3tr4.fu6
h3tr4.fu14
h3tr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.57. Test run ho2tr1

OH + O → HO₂ with Morse I approximation

In this sample run, CVT, μ VT, and US rate constants [CVT, MUVT, MUVTOPT, US] are calculated at two temperatures [TEMP] for the reaction OH + O → HO₂. (The values of the conventional transition state theory rate constants, not computed in this sample calculation because there is no saddle point ([STATUS = 2] in START section and NOSADDLE in the PATH section), are set to 1.0 in the unit fu15 output.) The first step off the starting point is followed along the negative of the gradient direction [FIRSTSTEP = *grad*]. All vibrational modes are treated by the Morse I approximation [MORSE, MORMODEL = *morsei*] The MEP is followed from the input initial geometry of the reactants by the ESD method [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.031 bohr [INH = 310]. Both CD-SCSAG and MEPSAG tunneling corrections are calculated [ZCT, SCT]. Note that the reverse rate constants are scaled by a factor of 1×10^{15} [REVKEXP = 15]. Thirty quadrature points were used in calculating the μ VT rate constants. Note that the degeneracies and energies of the ²Π_{3/2} and ²Π_{1/2} states of OH and ³P₂, ³P₁, and ³P₀ states of O are used in the calculation of the electronic partition functions. The results may be compared to those in Table V of Ref. 4 in Sect. 23.

Polyrate I/O FILES

FILE NAME

ho2tr1.dat
ho2tr1.fu6
ho2tr1.fu14
ho2tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.58. Test run ho2tr2

OH + O → HO₂ with Morse I approximation using curvilinear coordinates

In this sample run, CVT, μ VT, and US rate constants [CVT, MUVT, MUVTOPT, US] are calculated at two temperatures [TEMP] for the reaction OH + O → HO₂ using curvilinear coordinates (O-H and O-O stretches and an H-O-O bend). (The values of the conventional transition state theory rate constants, not computed in this sample calculation because there is no saddle point [STATUS = 2 in START section and NOSADDLE in the PATH section], are set to 1.0 in the unit fu15 output.) The MEP is followed from the input initial geometry of the reactants by the ESD method [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.031 bohr [INH = 310]. Both CD-SCSAG and MEPSAG tunneling corrections are calculated [ZCT, SCT]. Note that the reverse rate constants are scaled by a factor of 1×10^{15} [REVKEXP = 15].

Polyrate I/O FILES

FILE NAME

ho2tr2.dat
ho2tr2.fu6
ho2tr2.fu14
ho2tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.59. Test run ho2tr3

OH + O → HO₂ with WKB approximation for one mode

This sample run for the reaction OH + O → HO₂ uses a combination of methods for the treatment of the vibrational modes [VANHAR]. All modes of the reactant and product species are treated by the default harmonic approximation. The lowest frequency mode of the generalized transition state is also treated harmonically, while the highest frequency mode is treated by the primitive WKB method applied to a quadratic-quartic fit of the potential along the normal mode coordinate [QQWKB]. (The values of the conventional transition state theory rate constants, not computed in this sample calculation because there is no saddle point [STATUS = 2 in START section and NOSADDLE in the PATH section], are set to 1.0 in the unit fu15 output). Points along the MEP are calculated with the ESD integrator [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP]. A generalized normal mode analysis, together with a Hessian matrix evaluation, is performed every 0.155 bohr [INH = 1550]. A generalized normal mode analysis is also performed at eight special save points [SSPECIAL]. CVT [CVT] and ICVT [ICVT] rate constants are calculated at one temperature [TEMP], using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT]. Sixth-order Lagrangian interpolation is used to obtain the values of the effective mass for CD-SCSAG tunneling [SCTOPT: *lagrange*]. The reverse rate constants are scaled by a factor of 1×10^{15} [REVEXP = 15]. Also, restart information is written to unit fu1 [RESTART: WRITEFU1].

Polyrate I/O FILES

FILE NAME

ho2tr3.dat
ho2tr3.fu6
ho2tr3.fu1
ho2tr3.fu14
ho2tr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu1 reaction path info.
poly.fu14 summary output file
poly.fu15 summary output file

16.C.60. Test run ho2tr4

O₂ + H → HO₂ with Morse I approximation

This sample run for the reaction O₂ + H → HO₂ uses the Morse I approximation for all vibrational modes [MORSE, MORMODEL = *morsei*]. The MEP is calculated by the ESD integrator [RPM = *esd*] with a step size of 0.0002 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.01 bohr [INH = 50]. Both CD-SCSAG and MEPSAG tunneling probabilities are calculated [ZCT, SCT].

This test run forms a set with the ho2tr5 test run. Since the MEP is finite on the product side of the saddle point and the next test run involves a VTST-IOC calculation, before running it we must find the *s* value at the product. Thus, in this run we calculate a relatively long range of the MEP (*s* = -2.5 bohr to *s* = 2.5 bohr) [SRANGE] and find that the *s* value at the product is approximately 2.20 bohr. We also learn from this test run that in order for the tunneling calculation to converge, the MEP should be calculated at least from *s* = -1.8 bohr to *s* = 0.8 bohr. Finally we learn that the highest saddle point frequency becomes the reactant frequency at *s* = -∞, and the second and third highest saddle point frequencies become the product frequencies at *s* = 2.20 bohr. The lowest product frequency corresponds to the reaction coordinate. We then use the information above together with some ab initio data to make the input data file for the VTST-IOC calculation in test run ho2tr5.

Polyrate I/O FILES

FILE NAME

ho2tr4.dat
ho2tr4.fu6
ho2tr4.fu14
ho2tr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.61. Test run ho2tr5

O₂ + H → HO₂ with Morse I approximation and VTST-IOC calculation

This sample run is similar to the ho2tr4 test run except we perform a VTST-IOC calculation in addition to the VTST calculation [ICOPT].

This reaction has two reactants and one product and a shallow well on the reactant side of MEP. In this test run we neglect the well and set the MEPTYPER to *two* and MEPTYPEP to *one*. The accurate energy and frequency information and the geometry for calculating the determinant of the moment of inertia tensor at the reactants and product are obtained from *J. Chem. Phys.* **74**, 3482 (1981). The corrected barrier height is set to 1.0 kcal/mol and the corrected saddle point frequencies are obtained by interpolating the correction at reactants and product. Frequency matching information is input by the FREQMAT keyword. We use the DETMI keyword to input the corrected determinant of the moment of inertia tensor, and we use UNITMI to select atomic units input. The frequency-matching information and the *s* value for the product are obtained from the test run ho2tr4. The range parameters in the unit fu50 input are determined by the utility programs findl.f and findb.f (see Subsection 17.D) by using additional points on the MEP that have energies halfway between the energy of the saddle point and the energies of the reactant- and product-side stationary points.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ho2tr5.dat	poly.fu5	input data file
ho2ic.dat	poly.fu50	input data file
ho2tr5.fu6	poly.fu6	long output file
ho2tr5.fu14	poly.fu14	summary output file
ho2tr5.fu15	poly.fu15	summary output file

16.C.62. Test run nh3tr1

NH₃ → NH₃ (inversion) with harmonic approximation and VTST-IOC calculation

This sample run is for the unimolecular reaction corresponding to the unimolecular inversion of ammonia between two pyramidal configurations, using the default harmonic approximation for all vibrational modes. The MEP is calculated by the ESD method [RPM = *esd*] between $s = 1.18$ bohr and $s = -1.18$ bohr [SRANGE] with a step size of 0.0001 bohr [SSTEP]. A generalized normal mode analysis is performed every 0.01 bohr [INH = 100]. MEPSAG and CD-SCSAG tunneling probabilities are calculated [ZCT, SCT], and they can be used to study the tunneling splitting in this reaction. Rate constants are not calculated [NOTST, NOCVT].

The VTST-IOC option is selected [ICOPT] and the accurate energy and frequency information and the geometry for calculating the determinant of the moment of inertia tensor are obtained from *J. Chem. Phys.* **93**, 6570 (1990). The range parameters in the unit fu50 input are determined by the utility programs findl.f and findb.f (see Subsection 17.D) by using additional points on the MEP that have an energy halfway between the energy of the saddle point and the reactant/product energy.

Polyrate I/O FILES

FILE NAME

nh3tr1.dat
nh3ic.dat
nh3tr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu50 input data file
poly.fu6 long output file

16.C.63. Test run nh3tr2

NH₃ → NH₃ (inversion) with harmonic approximation VTST calculation

This sample run is for the unimolecular reaction corresponding to the unimolecular inversion of ammonia between two pyramidal configurations, using the default harmonic approximation for all vibrational modes. The MEP is calculated by the ESD method [RPM = *esd*] between $s = 1.18$ bohr and $s = -1.18$ bohr [SRANGE] with a step size of 0.0001 bohr [SSTEP]. A generalized normal mode analysis is performed every 0.01 bohr [INH = 100]. The TST and CVT rate constants are calculated. The partition functions and the internal free energy of reactant are printed out with PRPART = *rpt*. The GTLOG keyword is used so the generalized free energy of activation is calculated using logarithms of the individual partition functions.

This is a somewhat artificial example (because it is a degenerate rearrangement and because ordinary pressures are too low to stabilize the system in on or another well), it is used for testing the code.

Polyrate I/O FILES

FILE NAME

nh3tr2.dat
nh3tr2.fu6
nh3tr2.fu14
nh3tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 input data file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.64. Test run oh3tr1

OH + H₂ → H₂O + H with harmonic approximation and using the VRP(ESD) algorithm

This sample run is for the OH + H₂ → H₂O + H reaction. The zero of energy of the reactants is calculated from the energies of the individual molecular reactant species; in other words, the default value of EZERO = *calculate* is used. All normal modes and generalized normal modes are treated using the default harmonic approximation. The VRP(ESD) algorithm [RPM = *vrpe*] with a step size of 0.001 bohr [SSTEP] is used to follow the MEP, while the initial step from the saddle point is based on a cubic expansion of the potential energy surface [FIRSTSTEP = *cubic*]. The first five steps from the saddle point geometry in each direction are of size 0.0001 bohr [SFIRST]. The Hessian matrix is recalculated and a generalized normal mode analysis is performed every 0.01 bohr [INH = 10].

MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT] are computed, and both CVT [CVT] and ICVT [ICVT] rate constants are calculated at eight temperatures [TEMP]. The derivative of the gradient with respect to the reaction coordinate is found by a quadratic fit [CURV = *dgrad*]. In addition, in this test run as well as in test runs oh3tr2–oh3tr6, extrapolation is used for the first direction from the saddle point (i.e., toward reactants) [EXFIRST].

Polyrate I/O FILES

FILE NAME

oh3tr1.dat
oh3tr1.fu6
oh3tr1.fu14
oh3tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.65. Test run oh3tr2

OH + H₂ → H₂O + H with Morse I-quadratic-quartic approximation

This sample run for the OH + H₂ → H₂O + H reaction uses the Morse I and quadratic-quartic approximations [MORSEQQ, MORMODEL = *morsei*] for the normal and generalized normal modes, i.e., modes with nonzero cubic anharmonicity are treated by the Morse I model, while modes with zero cubic anharmonicity are approximated as quadratic-quartic oscillators. Both the CVT [CVT] and ICVT [ICVT] methods are used. The rate constants are calculated at twelve temperatures [TEMP] and the activation energies are computed for three temperature pairs [EACT]. The ESD integrator [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP] is used to follow the MEP, and a generalized normal mode analysis is performed every 0.01 bohr [INH = 100]. Both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT] are computed. Note that the derivative of the gradient with respect to the reaction coordinate is computed by the one-sided difference method [CURV = *oneside*] and the free energy of activation profile is calculated over a different range of reaction coordinate for each temperature [GTEMP]. The results can be compared to those in Table IV of Ref. 1 in Sect. 23.

Polyrate I/O FILES

FILE NAME

oh3tr2.dat
oh3tr2.fu6
oh3tr2.fu14
oh3tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.66. Test run oh3tr3

OH ($\nu = 0$) + H₂ → H₂O + H with WKB and harmonic approximations

This sample run for the OH ($\nu = 0$) + H₂ → H₂O + H reaction uses the WKB approximation for the ground-state energy of all normal modes [WKB], and the default harmonic approximation for all excited-state energies. The overall translations and rotations are not projected out at stationary points [NOPROJECT]. The MEP is followed with the Euler stabilization (ES1) integrator [RPM = *es1*], with a step size of 0.001 bohr [SSTEP], and the initial step from the saddle point is based on a cubic expansion of the energy surface [FIRSTSTEP = *cubic*]. A value different from the default value for the stabilization criterion (magnitude of the vector difference between successive gradients) is used; it is set equal to 0.001 [ES1OPT: *diffd*]. A generalized normal mode analysis is performed every 0.02 bohr [INH = 20]. CVT [CVT] and ICVT [ICVT] rate constants are calculated at eight temperatures [TEMP]. Both MEPSAG and CD-SCSAG tunneling corrections are calculated [ZCT, SCTOPT], and the curvature components are found by using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. Sixth-order Lagrangian interpolation is used to obtain the values of the effective mass for the CD-SCSAG tunneling calculation [SCTOPT: LAGRANGE].

Polyrate I/O FILES

FILE NAME

oh3tr3.dat
oh3tr3.fu6
oh3tr3.fu14
oh3tr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.67. Test run oh3tr4

Adiabatic OH ($v = 0$) + H₂($v = 1$) → H₂O + H with Morse I quadratic-quartic approximation

This sample run is an example of the calculation of adiabatic state-selected rate constants [STATE = *adiab*] for the reaction OH ($v = 0$) + H₂($v = 1$) → H₂O + H. The highest-frequency normal mode of the generalized transition state is restricted to the first excited state, while the second highest frequency mode is restricted to the ground state. All other normal modes of the generalized transition state are treated thermally. For the reactants, the stretching mode of OH is restricted to the ground state, and the stretching mode of H₂ is restricted to the first excited state. Finally, for the product H₂O, the highest normal mode is restricted to the excited state, the second highest mode is restricted to the ground state, and the lowest mode is treated thermally. The Page-McIver local-quadratic approximation integrator [RPM = *pagem*] with a step size of 0.001 bohr [SSTEP] is used to follow the MEP, the Hessian matrix is recalculated and a generalized normal mode analysis is performed every 0.01 bohr [INH = 10]. All other options and input are the same as for test run oh3tr2 except that only the forward rate constants are calculated [FORWARDK].

Polyrate I/O FILES

FILE NAME

oh3tr4.dat
oh3tr4.fu6
oh3tr4.fu14
oh3tr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.68. Test run oh3tr5

Diabatic OH + H₂($\nu = 1$) → H₂O + H with Morse I quadratic-quartic approximation

This sample run is an example of the calculation of diabatic state-selected rate constants [STATE = *diab*] for the reaction OH + H₂($\nu = 1$) → H₂O + H.

The adiabatic/diabatic switch occurs at $s = -0.31$ bohr. For $s < -0.31$ bohr, the highest and second-highest frequency normal modes of the generalized transition state are restricted to the first excited and ground state, respectively. For $s > -0.31$ bohr, the highest-frequency normal mode is restricted to the ground state, while the second-highest normal mode is restricted to the first excited state. The lowest three normal modes of the GTS are all treated thermally for all values of s . For the reactants, the OH stretch vibration is restricted to the ground state, while the H₂ vibration is restricted to the first excited state. The highest and second-highest frequency normal modes of the product H₂O are restricted to the ground state and first excited state, respectively, while the lowest normal mode is treated thermally. Other options are the same as those for test run oh3tr2, except that only forward rate constants are calculated [FORWARDK] and the free energies are calculated over the same range of s for all temperatures [GSPEC]. The results may be compared to those in Table I of Ref. 3 in Sect. 23.

Polyrate I/O FILES

FILE NAME

oh3tr5.dat
oh3tr5.fu6
oh3tr5.fu14
oh3tr5.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.69. Test run oh3tr6

OH + H₂ → H₂O + H with harmonic approximation and the ES1 integrator

This sample run is for the OH + H₂ → H₂O + H reaction. All normal modes and generalized normal modes are treated using the default harmonic approximation. The ES1* integrator [RPM = *es1*, ES1OPT: DELTA2 = SSTEP and DIFFD = 0.01] with a step size of 0.01 bohr [SSTEP] is used to follow the MEP, while the initial step from the saddle point is based on a cubic expansion of the potential energy surface [FIRSTSTEP = *cubic*]. The Hessian matrix is recalculated and a generalized normal mode analysis is performed every 0.01 bohr [INH = 1].

MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT] are computed, and both CVT [CVT] and ICVT [ICVT] rate constants are calculated at eight temperatures [TEMP]. The derivative of the gradient with respect to the reaction coordinate is found by a quadratic fit [CURV = *dgrad*].

Polyrate I/O FILES

FILE NAME

oh3tr6.dat
oh3tr6.fu6
oh3tr6.fu14
oh3tr6.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.70. Test run oh3tr7

OH + H₂ → H₂O + H with MEPSAG and CD-SCSAG calculations at two temperatures (one of the test cases in Ref. 27 in Sect. 23), WRITEFU30 option

This sample run for the OH + H₂ → H₂O + H reaction is taken from one of the test cases in Ref. 27 in Sect. 23. (However, Ref. 27 used the obsolete SCSAG method whereas the current code uses CD-SCSAG.) The MEP is calculated from $s = -1.0$ bohr to $s = 0.3$ bohr [SRANGE] using the Page-McIver method [RPM = *pagem*] with the cubic starting algorithm [FIRSTSTEP = *cubic*]. The gradient and save step sizes are both set to 0.062 bohr [SSTEP, INH = 1]. The CVT/MEPSAG and CVT/CD-SCSAG rate constants are calculated [CVT, ZCT, SCT] at 200K and 300K [TEMP]. All the rate constants are within 15% of the converged values obtained in the test run oh3tr1. Also, a file is written out in unit fu30 format as an electronic structure input file with the WRITEFU30 option. The points are not in the required ascending order. Therefore, this generated fu30 file needs to be reordered by the user before using the file.

Polyrate I/O FILES

FILE NAME

oh3tr7.dat
oh3tr7.fu6
oh3tr7.fu14
oh3tr7.fu15
tr7old.fu30
oh3tr7.fu30

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file
poly.fu30 old fu30 file
poly.fu30 reordered fu30 file

16.C.71. Test run oh3tr8

OH + H₂ → H₂O + H with MEPSAG and CD-SCSAG calculations at two temperatures, one of the test cases in Ref. 27 in Sect. 23, using curvilinear internal coordinates

This sample run for the OH + H₂ → H₂O + H reaction is taken from one of the test cases in Ref. 27 in Sect. 23. (However, Ref. 27 used the obsolete SCSAG method, whereas the current code uses the CD-SCSAG method that has replaced it. Other changes are that here curvilinear internal coordinates [COORD = *curv2*] are used here to calculate bound-state vibrational frequencies and generalized normal modes, and that the run uses SPECSTOP keyword. The curvilinear internal coordinates used for the O-H + H'-H" → H-O-H' + H" hydrogen atom transfer reaction include the O-H, H'-H", and O-H' stretches, the H-O-H' and H"-H'-O bends, and the H"-H'-O-H dihedral [INTDEF]. The MEP is calculated from $s = -1.0$ bohr to $s = 1.0$ bohr [SRANGE] or until the V_a^G curve goes below 95% from its value at the saddle point to its value at the reactants [SPECSTOP]. The MEP is determined using the Page-McIver method [RPM = *pagem*] with the cubic starting algorithm [FIRSTSTEP = *cubic*]. The gradient and save step sizes are both set to 0.062 bohr [SSTEP, INH = 1]. The CVT/MEPSAG and CVT/CD-SCSAG rate constants are calculated [CVT, ZCT, SCT] at 200K and 300K [TEMP].

Polyrate I/O FILES

FILE NAME

oh3tr8.dat
oh3tr8.fu6
oh3tr8.fu14
oh3tr8.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.72. Test run oh3tr9

$\text{OH} + \text{H}_2 \rightarrow \text{OH}_2 + \text{H}$ VTST calculation including a well on the reactant side and writing a fu31 file

This sample test run writes the fu31 input file from the information calculated using an analytical potential energy surface. The run does not require any fu31 input file and the resulting fu31 file is ready to be used without any change. In addition, a well that is 1.28 kcal/mol more stable than the reactants is optimized, and its frequencies are calculated. The data is written to the fu31 file.

The vibrational analysis is carried out in curvilinear coordinates, and debugging information on this analysis is printed out to the fu6 output file.

Polyrate I/O FILES

FILE NAME

ch5fu31tr1.dat
ch5fu31tr1.fu31
ch5fu31tr1.fu50
ch5fu31tr1.fu6
ch5fu31tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu50 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.C.73. Test run oh3tr10

OH + H₂ → H₂O + H with harmonic approximation and an ESD/RODS reaction path

This sample run uses the WRITEFU31 keyword to generate an fu31 file, and it uses the RODS keyword to make up for deficiencies in the path. All normal modes and generalized normal modes are treated using the default harmonic approximation. The ESD integrator [RPM = *esd*] with a step size of 0.05 bohr [SSTEP] is used to follow the MEP, and the RODS algorithm [RODS] is used to correct the energies and frequencies along the path. The Hessian matrix is recalculated and a generalized normal mode analysis is performed every 0.05 bohr [INH = 1].

MEPSAG tunneling corrections [ZCT] are computed, and CVT [CVT] rate constants are calculated at eight temperatures [TEMP]. The derivative of the gradient with respect to the reaction coordinate is found by a quadratic fit [CURV = *dgrad*].

Polyrate I/O FILES

FILE NAME

oh3tr10.dat
oh3tr10.fu6
oh3tr10.fu14
oh3tr10.fu15
oh3tr10.fu31

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file
poly.fu31 input data file

16.C.74. Test run oh3fu30tr1

$\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$ using electronic structure input file data from file fu30 and curvilinear internal coordinates

This sample run demonstrates the use of electronic structure input file data in the calculation of conventional TST rate constants [$\text{POTENTIAL} = \text{unit30}$] for the reaction $\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$. Hessians are read in packed format in mass-scaled coordinates [$\text{LOPT}(4) = -2$ in unit fu30 input]. Note that even though the reaction path is not required for a conventional TST rate constant calculation, the `PATH` section must be included in the unit fu5 input in order to pass to the program the value of the factor used in scaling the coordinates [`SCALEMASS` in the unit fu5 input], and the data for at least two points along the reaction path have to be input from unit fu30. The curvilinear internal coordinates [$\text{COORD} = \text{curv2}$] used for the $\text{O-H} + \text{H}'\text{-H}'' \rightarrow \text{H-O-H}' + \text{H}''$ hydrogen atom transfer reaction include the O-H, H'-H'', and O-H' stretches, the H-O-H' and H''-H'-O bends, and the H''-H'-O-H dihedral [`INTDEF`]. All normal modes are treated harmonically [`HARMONIC` in the unit fu5 input].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
oh3fu30tr1.dat	poly.fu5	input data file
oh3fu30tr1.fu30	poly.fu30	input data file
oh3fu30tr1.fu6	poly.fu6	long output file
oh3fu30tr1.fu14	poly.fu14	summary output file
oh3fu30tr1.fu15	poly.fu15	summary output file

16.C.75. Test run oh3fu30tr2

OH + H₂ → H₂O + H using electronic structure input file data from file fu30 and curvilinear internal coordinates

This sample run demonstrates the use of electronic structure input file data in the calculation of rate constants [POTENTIAL = *unit30*] for the reaction OH + H₂ → H₂O + H. This type of calculation requires the unit fu30 input file. The normal mode and generalized normal mode frequencies are obtained at each save point on the MEP by first transforming the Cartesian Hessian matrix into curvilinear internal coordinate space, projecting out the gradient direction, and then diagonalizing the projected Hessian using the Wilson **GF** method. The curvilinear internal coordinates [COORD = *curv2*] used for the O-H + H'-H'' → H-O-H' + H'' hydrogen atom transfer reaction include the O-H, H'-H'', and O-H' stretches, the H-O-H' and H''-H'-O bends, and the H''-H'-O-H dihedral [INTDEF]. The curvature components are computed using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [LOPT(6) = 4]. All normal modes and generalized normal modes are treated harmonically [HARMONIC in the unit fu5 input], and only TST and CVT rate constants are calculated [CVT] at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
oh3fu30tr2.dat	poly.fu5	input data file
oh3fu30tr2.fu30	poly.fu30	electronic structure input file data
oh3fu30tr2.fu6	poly.fu6	long output file
oh3fu30tr2.fu14	poly.fu14	summary output file
oh3fu30tr2.fu15	poly.fu15	summary output file

16.C.76. Test run oh3fu30tr3

OH + H₂ → H₂O + H direct dynamics calculation using the electronic structure data read from file fu30, rectilinear coordinates, and IVTST-M-4/4 interpolation.

This sample run demonstrates the use of electronic structure input file data read from unit fu30 in the calculation of rate constants [POTENTIAL = *unit30*] for the reaction OH + H₂ → H₂O + H. This type of calculation requires the unit fu30 input file. The interpolation is done using IVTST-M-4/4 [LOPT(2) = 500]. Note that the normal mode and generalized normal mode frequencies are obtained by diagonalization of the matrix of the second derivatives of the potential with respect to the mass-scaled Cartesian coordinates, which is read in packed form from unit fu30 [LOPT(4) = -2 in the unit fu30 input], and no anharmonic data are included [LOPT(5) = 0]. All normal modes are treated harmonically [HARMONIC in the unit fu5 input], and only TST and CVT rate constants are calculated [CVT] at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
oh3fu30tr3.dat	poly.fu5	input data file
oh3fu30tr3.fu30	poly.fu30	input data file
oh3fu30tr3.fu6	poly.fu6	long output file
oh3fu30tr3.fu14	poly.fu14	summary output file
oh3fu30tr3.fu15	poly.fu15	summary output file

16.C.77. Test run oh3fu30tr4

OH + H₂ → H₂O + H using electronic structure input file data from file fu30 for generating the next point along the reaction path (reactants valley).

This sample run demonstrates the use of the NEXTPT keyword for generating the next point along the reaction path in conjunction with the use of electronic structure input from file fu30 [POTENTIAL = *u n i t 3 0*] for the reaction OH + H₂ → H₂O + H. Note that the print option LOPT(1) and IFITV are set equal to 0 in file fu30 because nonzero values for these variables are incompatible with NEXTPT. Electronic structure information is included for two points on the reaction path (one on each side of the saddle point), with the first one having a positive *s* and the second one having a negative *s* (thus the points are not in increasing order of *s*, but this is permitted when NEXTPT is on). Since the last point has a negative *s*, IDIRECT is set equal to -1. The next point is generated on the reactants side using a step size of 0.0005 amu^{1/2} bohr [SSTEP].

Polyrate I/O FILES

FILE NAME

oh3fu30tr4.dat
oh3fu30tr4.fu30
oh3fu30tr4.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.78. Test run oh3fu30tr5

OH + H₂ → H₂O + H using electronic structure input file data from file fu30 for generating the next point along the reaction path (products valley).

This sample run demonstrates the use of the NEXTPT keyword for generating the next point along the reaction path in conjunction with the use of electronic structure input from file fu30 [POTENTIAL = *unit30*] for the reaction OH + H₂ → H₂O + H. Note that the print option LOPT(1) and IFITV are set equal to 0 in file fu30 because nonzero values for these variables are incompatible with NEXTPT. Electronic structure information is included for two points on the reaction path (one on each side of the saddle point), with the last one being in the products valley. Since the last point has a positive s, IDIRECT is +1. The next point is generated on the products side using a step size of 0.0005 amu^{1/2} bohr [SSTEP].

Polyrate I/O FILES

FILE NAME

oh3fu30tr5.dat
oh3fu30tr5.fu30
oh3fu30tr5.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu30 input data file
poly.fu6 long output file

16.C.79. Test run oh3fu30tr6

$\text{OH} + \text{H}_2 \rightarrow \text{OH}_2 + \text{H}$ fu30 calculation including a well on the reactant side

The oh3fu30tr6 test run is a sample run that uses file fu30 and includes a well on the reactant side in the calculation. The reaction path is interpolated by means of third order Lagrangian interpolation using the points along the reaction path including the reactant well. The file fu30 was created with the writefu30 option, with only two changes being necessary: to input of the value of the reaction coordinate, s , at the reactants well, and to reorder of the points.

Polyrate I/O FILES FILE NAME

oh3fu30tr6.dat
oh3fu30tr6.fu30
oh3fu30tr6.fu6
oh3fu30tr6.fu14
oh3fu30tr6.fu15

Polyrate FILE NAME

poly.fu5	input data file
poly.fu30	input data file
poly.fu6	long output file
poly.fu14	summary output file
poly.fu15	summary output file

16.C.80. Test run oh3fu31tr1

OH + H₂ → OH₂ + H IVTST-M calculation including a well on the reactant side

This sample test run uses the *unit31* option to perform an IVTST-M calculation including a well on the reactant side. This is an IVTST-M-100/2002 calculation based on an fu31 input file identical to the one written in the oh3tr9 test run. The value for the step-size SINCW is set to 0.001 bohr, which is the step-size used in the generation of the fu31 file.

Note that this run cannot be done without interpolating to the well, since the energies given in the fu31 input file for some of the points on the reactant side are below the energy of the infinitely separated reactants. Since the interpolation to two separated products is based on a Eckart potential, which does not allow the presence of wells, the algorithm would fail. Thus, the well is required in the present IVTST-M calculation. If the information given on the reactant side of the reaction path were to be reduced to the point where no energy lower than the reactant energy appears in the fu31 file, the interpolation could be carried out using the reactant information.

The calculated reaction path is extended up to $s = -1.70 a_0$ (using a scaling mass of 1.802023 a.u.). Since the well is estimated to be located at $s = -1.73 a_0$, extending the range of values of s further toward reactants could cause numerical errors, since the reaction path has not been interpolated beyond the well. In general, whenever a finite range of s is used in a IVTST-M calculation (i.e., reactions with only one reactant or only one product, or reactions with wells), the user should check that the range of s given in the fu5 file does not go beyond the values of s estimated for the stationary point (or points).

Polyrate I/O FILES

FILE NAME

oh3fu31tr1.dat
oh3fu31tr1.fu31
oh3fu31tr1.fu6
oh3fu31tr1.fu15

Polyrate FILE NAME

poly.fu5	input data file
poly.fu31	input data file
poly.fu6	long output file
poly.fu15	summary output file

16.C.81. Test run oh3fu31tr2

OH + DH \rightarrow HOD + H direct dynamics calculation of an isotopically substituted reaction using electronic structure data for the isotopically unsubstituted reaction read from file fu31, using ESD(H)/RODS, and using IVTST-M.

This sample run demonstrates how to use the OH + H₂ \rightarrow H₂O + H reaction path along with the ESD(H)/RODS [RODS] and IVTST-M algorithms and file fu31 [POTENTIAL = *unit31*] to obtain the conventional TST, CVT, CVT/ZCT, and CVT/SCT rate constants for the isotopically substituted reaction OH + DH \rightarrow HOD + H. The ESD(H)/RODS method is described in Ref. 50 in Sect. 23. The fu31 input file used in this run was written using the WRITEFU31 option in a run on OH + H₂ (not included in the test suite). The KIES option must be included in the GENERAL section of file fu31 so that the correct value of *s* is calculated for the isotopically substituted reaction.

Polyrate I/O FILES

FILE NAME

oh3fu31tr2.dat
oh3fu31tr2.fu31
oh3fu31tr2.fu6
oh3fu31tr2.fu14
oh3fu31tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.82. Test run oh3fu40tr1

$\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$ using electronic structure input file data from file fu40 and curvilinear internal coordinates

This sample run demonstrates the use of electronic structure input file data in the calculation of conventional TST rate constants [$\text{POTENTIAL} = \text{unit40}$] for the reaction $\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$. Hessians are read in mass-scaled coordinates [$\text{HESSFORM} = \text{msbohr}$, $\text{HESSUNIT} = \text{msbohr}$ in file fu40]. Note that even though the reaction path is not required for a conventional TST rate constant calculation, the PATH section must be included in the unit fu5 input in order to pass to the program the value of the factor used in scaling the coordinates [SCALEMASS in the unit fu5 input] and at least two points along the reaction path have to be input from unit fu40. The curvilinear internal coordinates [$\text{COORD} = \text{curv2}$] used for the $\text{O-H} + \text{H}'\text{-H}'' \rightarrow \text{H-O-H}' + \text{H}''$ hydrogen atom transfer reaction include the O-H, H'-H'', and O-H' stretches, the H-O-H' and H''-H'-O bends, and the H''-H'-O-H dihedral [INTDEF]. All normal modes are treated harmonically [HARMONIC in the unit fu5 input].

Polyrate I/O FILES

FILE NAME

oh3fu40tr1.dat
oh3fu40tr1.fu40
oh3fu40tr1.fu6
oh3fu40tr1.fu14
oh3fu40tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.83. Test run oh3fu40tr2

OH + H₂ → H₂O + H using electronic structure input file data from file fu40 and curvilinear internal coordinates

This sample run demonstrates the use of electronic structure input file data in the calculation of rate constants [POTENTIAL = *unit40*] for the reaction OH + H₂ → H₂O + H. This type of calculation requires the unit fu40 input file. The normal mode and generalized normal mode frequencies are obtained at each save point on the MEP by first transforming the Cartesian Hessian matrix into curvilinear internal coordinate space, projecting out the gradient direction, and then diagonalizing the projected Hessian using the Wilson **GF** method. The curvilinear internal coordinates [COORD = *curv2*] used for the O-H + H'-H'' → H-O-H' + H'' hydrogen atom transfer reaction include the O-H, H'-H'', and O-H' stretches, the H-O-H' and H''-H'-O bends, and the H''-H'-O-H dihedral [INTDEF]. The curvature components are computed using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURVATURE = *compute* and GRADDER = *noextra* in the unit fu40 input]. All normal modes and generalized normal modes are treated harmonically [HARMONIC in the unit fu5 input], and only TST and CVT rate constants are calculated [CVT] at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
oh3fu40tr2.dat	poly.fu5	input data file
oh3fu40tr2.fu40	poly.fu40	input data file
oh3fu40tr2.fu6	poly.fu6	long output file
oh3fu40tr2.fu14	poly.fu14	summary output file
oh3fu40tr2.fu15	poly.fu15	summary output file

16.C.84. Test run oh3fu40tr3

OH + H₂ → H₂O + H direct dynamics calculation using the electronic structure data read from file fu40, using rectilinear coordinates, and IVTST-M-4/4.

This sample run demonstrates the use of electronic structure input file data read from unit fu40 in the calculation of rate constants [POTENTIAL = *unit40*] for the reaction OH + H₂ → H₂O + H. This type of calculation requires the unit fu40 input file. The interpolation is calculated using IVTST-M-4/4 [MAXLPTS = 500]. Note that the normal mode and generalized normal mode frequencies are obtained by diagonalization of the matrix of the second derivatives of the potential with respect to the mass-scaled Cartesian coordinates, which is read in packed form from unit fu40 [HESSFORM = *packed*, HESSUNIT = *msbohr*], and no anharmonic data are included [ANHARM = *none*]. The curvature components are computed using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURVATURE = *compute*, GRADDER = *noextra*]. All normal modes are treated harmonically [HARMONIC in the unit fu5 input], and only TST and CVT rate constants are calculated [CVT] at four temperatures [TEMP], using both MEPSAG and CD-SCSAG tunneling corrections [ZCT, SCT].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
oh3fu40tr3.dat	poly.fu5	input data file
oh3fu40tr3.fu40	poly.fu40	input data file
oh3fu40tr3.fu6	poly.fu6	long output file
oh3fu40tr3.fu14	poly.fu14	summary output file
oh3fu40tr3.fu15	poly.fu15	summary output file

16.C.85. Test run oh3fu40tr4

OH + H₂ → H₂O + H using electronic structure input file data from file fu40 for generating the next point along the reaction path (reactants valley).

This sample run demonstrates the use of the NEXTPT keyword for generating the next point along the reaction path in conjunction with the use of electronic structure input from file fu40 [POTENTIAL = *u n i t 4 0*] for the reaction OH + H₂ → H₂O + H. Electronic structure information is included for two points on the reaction path (one on each side of the saddle point), with the first one having a positive *s* and the second one having a negative *s* (thus the points are not in increasing order of *s*, but this is permitted when NEXTPT is on). Since the last point has a negative *s*, IDIRECT is set equal to -1. The next point is generated on the reactants side using a step size of 0.0005 amu^{1/2} bohr [SSTEP].

Polyrate I/O FILES

FILE NAME

oh3fu40tr4.dat
oh3fu40tr4.fu40
oh3fu40tr4.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.86. Test run oh3fu40tr5

OH + H₂ → H₂O + H using electronic structure input file data from file fu40 for generating the next point along the reaction path (products valley)

This sample run demonstrates the use of the NEXTPT keyword for generating the next point along the reaction path in conjunction with the use of electronic structure input from file fu40 [POTENTIAL = *u n i t 4 0*] for the reaction OH + H₂ → H₂O + H. Electronic structure information is included for two points on the reaction path (one on each side of the saddle point), with the last one being in the products valley. Since the last point has a positive *s*, IDIRECT is +1. The next point is generated on the products side using a step size of 0.0005 amu^{1/2} bohr [SSTEP].

Polyrate I/O FILES

FILE NAME

oh3fu40tr5.dat
oh3fu40tr5.fu40
oh3fu40tr5.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu40 input data file
poly.fu6 long output file

16.C.87. Test run oh3fu40tr6

$\text{OH} + \text{H}_2 \rightarrow \text{OH}_2 + \text{H}$ fu40 calculation including a well on the reactant side

The oh3fu40tr6 test run is a sample run that uses file fu40 and includes as the left-hand edge of the interpolation interval a well on the reactant side in the calculation. The reaction path is interpolated by means of third order Lagrangian interpolation using the points along the reaction path including the reactants well.

Polyrate I/O FILES

FILE NAME

oh3fu40tr6.dat
oh3fu40tr6.fu40
oh3fu40tr6.fu6
oh3fu40tr6.fu14
oh3fu40tr6.fu15

Polyrate FILE NAME

poly.fu5	input data file
poly.fu40	input data file
poly.fu6	long output file
poly.fu14	summary output file
poly.fu15	summary output file

16.C.88. Test run hnitrl

H(a)/Ni(100) \rightarrow H(a)/Ni(100) with harmonic approximation and ESD integrator

This sample run is for the surface diffusion reaction H(a)/Ni(100) \rightarrow H(a)/Ni(100), where both reactant and product are adsorbed species [PHASE = *solid*] on a fixed Ni lattice. In this particular calculation none of the Ni atoms are allowed to move. All vibrational modes are treated by the default harmonic approximation. Points along the MEP are calculated by the Euler steepest-descents (ESD) integrator [RPM = *esd*] with a step size of 0.0001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.02 bohr [INH = 200]. CVT rate constants [CVT] are calculated. In addition, MEPSAG and CD-SCSAG transmission coefficients are included [ZCT, SCT]. The reaction path curvature components are calculated using a quadratic fit to obtain the derivative of the gradient with respect to the reaction coordinate [CURV = *dgrad*]. Note that the electronic partition functions of the transition state and of the reactant are assumed to cancel.

Polyrate I/O FILES

FILE NAME

hnitrl.dat
hnitrl.fu6
hnitrl.fu14
hnitrl.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

pothni.dat

UNIT = 4
FILE NAME = pothni.dat

16.C.89. Test run h2nitr1

$\text{H}_2(\text{a})/\text{Ni}(100) \rightarrow 2\text{H}(\text{a})/\text{Ni}(100)$ with harmonic approximation and ESD integrator

This sample run is for the unimolecular gas-solid-interface reaction $\text{H}_2(\text{a})/\text{Ni}(100) \rightarrow 2\text{H}(\text{a})/\text{Ni}(100)$ where both reactant and products are adsorbed species [PHASE = *solid*] on a fixed Ni lattice consisting of 817 Ni atoms. All vibrational modes are treated by the default harmonic approximation. Points along the MEP are calculated by the ESD integrator [RPM = *esd*] with a step size of 0.001 bohr [SSTEP], and a generalized normal mode analysis is performed every 0.01 bohr [INH = 10]. CVT rate constants are calculated [CVT] at 8 temperatures [TEMP]. In addition, MEPSAG, CD-SCSAG, and LCG3 transmission coefficients are included [ZCT, SCT, LCTOPT], and only the ground state is involved in the LCG3 exit arrangement for the LCG3 tunneling calculation [NOALLEXCIT].

Polyrate I/O FILES

FILE NAME

h2nitr1.dat
h2nitr1.fu6
h2nitr1.fu14
h2nitr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILES

poth2niads.dat
poth2nilat.dat

UNIT = 4
UNIT = 7

FILE NAME = poth2niads.dat
FILE NAME = poth2nilat.dat

16.C.90. Test run ch4ohtr1

OH + CH₄ → H₂O + CH₃ zero-order IVTST calculation

This sample run uses the ICOPT keyword [ICOPT: *zero*] for a zero-order IVTST (*i.e.*, IVTST-0) calculation of the reaction OH + CH₄ → H₂O + CH₃ at 19 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -2.0$ bohr to $s = 2.0$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 20]. The ab initio data taken from *J. Chem. Phys.* **99**, 1013 (1993) are input by means of the unit fu50 file. The ZCT keyword is used since MEPSAG is the only semiclassical tunneling method available for the zero-order IVTST calculation. The lowest-frequency vibrational mode of the generalized transition state is treated as a hindered rotor [VANHAR, TOR], while all of the other modes are treated by the default harmonic approximation.

Polyrate I/O FILES

FILE NAME

ch4ohtr1.dat
ch4ohic.dat
ch4ohtr1.fu6
ch4ohtr1.fu14
ch4ohtr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu50 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.91. Test run ch4ohtr2

OH + CH₄ → H₂O + CH₃ zero-order IVTST calculation

This sample run uses the IVTST0 keyword for a zero-order IVTST (*i.e.*, IVTST-0) calculation of the reaction OH + CH₄ → H₂O + CH₃ at 19 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -2.0$ bohr to $s = 2.0$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 20]. The ab initio data taken from *J. Chem. Phys.* **99**, 1013 (1993) are input by means of the unit fu29 file. The MEPSAG method [ZCT] is the only semiclassical tunneling method available for the zero-order IVTST calculation. All vibrational modes are treated by the default harmonic approximation.

Polyrate I/O FILES

FILE NAME

ch4ohtr2.dat
ch4ohtr2.fu29
ch4ohtr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 output file

16.C.92. Test run ch4ohtr3

OH + CH₄ → H₂O + CH₃ zero-order IVTST calculation

This sample run uses the ICOPT keyword [ICOPT: *zero*] for a zero-order IVTST (*i.e.*, IVTST-0) calculation of the reaction OH + CH₄ → H₂O + CH₃ at 19 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -2.0$ bohr to $s = 2.0$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 20]. The ab initio data taken from *J. Chem. Phys.* **99**, 1013 (1993) are input by means of the unit fu50 file. The ZCT keyword is used since MEPSAG is the only semiclassical tunneling method available for the zero-order IVTST calculation. This sample run is the same as the ch4ohtr1 test run, except that the lowest-frequency vibrational mode of the generalized transition state is treated as a hindered rotor using the CO torsion option.

Polyrate I/O FILES

FILE NAME

ch4ohtr3.dat
ch4ohic.dat
ch4ohtr3.fu6
ch4ohtr3.fu14
ch4ohtr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu50 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.93. Test run ch4ohtr4

OH + CH₄ → H₂O + CH₃ zero-order IVTST calculation

This sample run uses the ICOPT keyword [ICOPT: *zero*] for a zero-order IVTST (*i.e.*, IVTST-0) calculation of the reaction OH + CH₄ → H₂O + CH₃ at 19 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -2.0$ bohr to $s = 2.0$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 20]. The ab initio data taken from *J. Chem. Phys.* **99**, 1013 (1993) are input by means of the unit fu50 file. The ZCT keyword is used since MEPSAG is the only semiclassical tunneling method available for the zero-order IVTST calculation. This sample run is the same as the ch4ohtr1 test run, except that the lowest-frequency vibrational mode of the generalized transition state is treated as a hindered rotor using the CW torsion option.

Polyrate I/O FILES

FILE NAME

ch4ohtr4.dat
ch4ohic.dat
ch4ohtr4.fu6
ch4ohtr4.fu14
ch4ohtr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu50 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.94. Test run ch4cltr1

$\text{CH}_4 + \text{Cl} \rightarrow \text{HCl} + \text{CH}_3$ zero-order IVTST calculation

This sample run uses the IVTST0 keyword for a zero-order IVTST (*i.e.*, IVTST-0) calculation of the reaction $\text{CH}_4 + \text{Cl} \rightarrow \text{HCl} + \text{CH}_3$ at 4 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -1.0$ bohr to $s = 1.5$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 4]. The *ab initio* data are input by means of the unit fu29 file. The MEPSAG method is the only semiclassical tunneling method available for the zero-order IVTST calculation. All vibrational modes are treated by the default harmonic approximation. The $^2\text{P}_{3/2}$ and $^3\text{P}_{1/2}$ states of Cl are used in the calculation of the electronic partition functions [ELEC]. [This test run reproduces the results of one of the test cases in Ref. 26 in Sect. 23.]

Polyrate I/O FILES

FILE NAME

ch4cltr1.dat
ch4cltr1.fu29
ch4cltr1.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 output file

16.C.95. Test run ch4cltr2

$\text{CH}_4 + \text{Cl} \rightarrow \text{HCl} + \text{CH}_3$ first-order IVTST calculation

This sample run uses the IVTST1 keyword for a first-order IVTST (*i.e.*, IVTST-1) calculation of the reaction $\text{CH}_4 + \text{Cl} \rightarrow \text{HCl} + \text{CH}_3$ at 4 temperatures [TEMP]. The classical potential energy curve and the adiabatic ground-state energy curve are approximated by Eckart functions from $s = -1.0$ bohr to $s = 1.5$ bohr [SRANGE]. The Hessian grid spacing is set to 0.02 bohr [INH = 4]. The *ab initio* data are input by means of the unit fu29 file. Both MEPSAG and SCSAG semiclassical tunneling methods [ZCT, SCT] are used, since they are available for the first-order IVTST calculation. All vibrational modes are treated by the default harmonic approximation. The $^2\text{P}_{3/2}$ and $^3\text{P}_{1/2}$ states of Cl are used in the calculation of the electronic partition functions[ELEC]. [This test run reproduces the results of one of the test cases in Ref. 26 in Sect. 23.]

Polyrate I/O FILES

FILE NAME

ch4cltr2.dat
ch4cltr2.fu29
ch4cltr2.fu6

Polyrate FILE NAME

poly.fu5 input data file
poly.fu29 input data file
poly.fu6 output file

16.C.96. Test run ch4otr1

$\text{CH}_4 + \text{O} \rightarrow \text{CH}_3 + \text{OH}$ CUS calculation using harmonic redundant curvilinear coordinates

This sample run calculates canonical unified statistical model (CUS) rate constants including MEPSAG, CD-SCSAG, LCG3, COMT, and μOMT tunneling, based on the information from an analytical potential energy surface. The reaction has two different electronic states, but since they are very close in energy, they are assumed to be represented by the same surface. Thus, the total rate constant should be multiplied by a factor of 2 in order to take into account this electronic degeneracy. This is done by using a factor of 6 for the saddle point electronic degeneracy instead of its value of 3.

The $V_a^G(s)$ curve has two maxima near the saddle point. In order to estimate the recrossing effects due to these maxima a CUS calculation is performed.

Polyrate I/O FILES

FILE NAME

ch4otr1.dat
ch4otr1.fu6
ch4otr1.fu14
ch4otr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

POTENTIAL ENERGY SURFACE DATA FILE

potch4o.dat

UNIT = 4

FILE NAME = potch4o.dat

16.C.97. Test run ohcltr1

O + HCl → OH + Cl ICVT calculation using Morse 1 anharmonicity and internal coordinates

This sample run calculates improved canonical variational theory (ICVT) rate constants including MEPSAG and CD-SCSAG tunneling, based on the information from an analytical potential energy surface. The saddle point and generalized transition states are nonlinear [`SPECIES = nonlints`]. Internal coordinates are defined [`INTDEF` keyword] as the OH (1-2) and HCl (2-3) bond lengths and OHCl (1-2-3) bond angle. Vibrational modes are treated by the Morse I approximation [`MORSE, MORMODEL = morsei`], with a dissociation energy [`DEMIN`] of 0.16968172 E_h.

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME
ohcltr1.dat	poly.fu5
ohcltr1.fu6	poly.fu6
ohcltr1.fu14	poly.fu14
ohcltr1.fu15	poly.fu15

input data file
long output file
summary output file
summary output file

16.C.98. Test run ohcltr2

O + HCl \rightarrow OH + Cl ICVT calculation using WKB-on-the-true-potential for ground-state energies and Morse 1 anharmonicity for excited state energies.

This sample run is the same as ohcltr1, except that the ground-state vibrational energy levels are calculated using the WKB method on the true potential. All excited state energy levels are computed using the Morse I approximation [MORSE, MORMODEL = *morsei*], with a dissociation energy [DEMIN] of 0.16968172 E_h .

Polyrate I/O FILES

FILE NAME

ohcltr2.dat
ohcltr2.fu6
ohcltr2.fu14
ohcltr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.99. Test run ohcltr3

O + HCl(v=1) → OH + Cl Adiabatic excited-state calculation using WKB-on-the-true-potential for the v=1 stretch energy and for the ground-state bend energy and Morse 1 anharmonicity for excited statebend energies.

This sample run is the same as ohcltr2, except that the reaction is for excited-state stretch vibration. The stretch vibration is treated as adiabatic along the full reaction path [STATE adiab] with quantum number v=1 specified using the STATEOPT keyword. The excited-state energy level for the stretch vibration and the ground-state vibrational energy level for the bend vibration are calculated using the WKB method on the true potential. Bend excited state energy levels are computed using the Morse I approximation [MORSE, MORMODEL = *morsei*], with a dissociation energy [DEMIN] of 0.16968172 E_h. The ICVT calculation is limited to $-1.5 a_0 < s < 1.0 a_0$.

Polyrate I/O FILES

FILE NAME

ohcltr3.dat
ohcltr3.fu6
ohcltr3.fu14
ohcltr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.100. Test run ohcltr4

O + HCl(v=1) → OH + Cl Partial-reaction-path (PRP) adiabatic excited-state calculation using WKB-on-the-true-potential for stretch energy and for the ground-state bend energy and Morse 1 anharmonicity for excited state bend energies.

This sample run is the same as ohcltr3, except that stretch vibration is treated as adiabatic in the v=1 state along the reaction path only up to $s = -0.22 a_0$ and then in the ground state for $s > -0.22 a_0$. This is specified using the STATE diab keyword and specifying transition state mode 2 in the first excited state before the diabatic switch and in the ground excited state after the diabatic switch. Thermal treatment of the bend vibration before and after the switch is explicitly indicated in the input. The energy level for the stretch vibration and the ground-state vibrational energy level for the bend vibration are calculated using the WKB method on the true potential. Bend excited state energy levels are computed using the Morse I approximation [MORSE, MORMODEL = *morsei*], with a dissociation energy [DEMIN] of $0.16968172 E_h$. The ICVT calculation is limited to $-1.5 a_0 < s < 1.0 a_0$.

Polyrate I/O FILES

FILE NAME

ohcltr4.dat
ohcltr4.fu6
ohcltr4.fu14
ohcltr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu14 summary output file
poly.fu15 summary output file

16.C.101. Test run ohcltr5

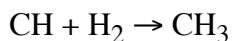
O + HCl → OH + Cl ICVT and MUCVT calculation using Morse 1
anharmonicity and internal coordinates

This sample run is the same as ohcltr1, except that microcanonical variational theory (MUCVT) rate constants are also calculated. The primary and secondary fits use 1-point and 3-point fitting [*MUVTOPT*, *FIT= GAS13*].

Polyrate I/O FILES

FILE NAME	<i>Polyrate</i> FILE NAME	
ohcltr5.dat	poly.fu5	input data file
ohcltr5.fu6	poly.fu6	long output file
ohcltr5.fu14	poly.fu14	summary output file
ohcltr5.fu15	poly.fu15	summary output file

16.C.102. Test run ch3tr1



VRC-VTST calculation

This sample run calculates the variable reaction coordinate variational transition state theory rate constants by canonical variational theory, microcanonical variational theory, and total energy and angular momentum resolved microcanonical variational theory. One pivot point is assigned to each reactant. Pivot point for CH is assigned as the C atom, and pivot point for H₂ is the center of mass of H₂. The range of reaction coordinate s is [1.3, 6.5] Å with a step size 0.2 Å. The number of Monte Carlo samples (keyword *NMC*) is set to 100 for each dividing surface. Note that this number should be set to around 1000 in a production run in order to have a much lower uncertainty of the Monte Carlo integration.

Polyrate I/O FILES

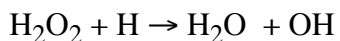
FILE NAME

ch3tr1.dat
ch3tr1.fu6
ch3tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.C.103. Test run h3o2fu31tr1



TST with torsional treatment for H_2O_2 using the segmented-reference Pitzer-Gwinn (SRPG) method

This sample run calculates rate constants by conventional TST. This run is used to show the usage of the SRPG method for treating a torsion. The potential energy surface is provided by the h3o2fu31tr1.fu31 file.

Polyrate I/O FILES

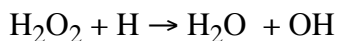
FILE NAME

h3o2fu31tr1.dat
h3o2fu31tr1.fu31
h3o2fu31tr1.fu6
h3o2fu31tr1.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.C.104. Test run h3o2fu31tr2



TST with torsional treatment for H₂O₂ using the segmented-reference Alyala-Schlegel (SAS) method

This sample run calculates rate constants by conventional TST. This run is used to show the usage of the SAS method for treating a torsion. The potential energy surface is provided by the h3o2fu31tr2.fu31 file.

Polyrate I/O FILES

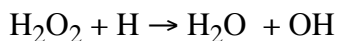
FILE NAME

h3o2fu31tr2.dat
h3o2fu31tr2.fu31
h3o2fu31tr2.fu6
h3o2fu31tr2.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.C.105. Test run h3o2fu31tr3



TST with torsional treatment for H_2O_2 using the reference Pitzer-Gwinn (RPG) method

This sample run calculates rate constants by conventional TST. This run is used to show the usage of the RPG method for treating a torsion. The potential energy surface is provided by the h3o2fu31tr3.fu31 file.

Polyrate I/O FILES

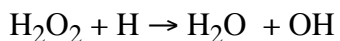
FILE NAME

h3o2fu31tr3.dat
h3o2fu31tr3.fu31
h3o2fu31tr3.fu6
h3o2fu31tr3.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.C.106. Test run h3o2fu31tr4



TST with torsional treatment for H_2O_2 using the
Alyala-Schlegel (AS) method

This sample run calculates rate constants by conventional TST. This run is used to show the usage of the AS method for treating a torsion. The potential energy surface is provided by the h3o2fu31tr4.fu31 file.

Polyrate I/O FILES

FILE NAME

h3o2fu31tr4.dat
h3o2fu31tr4.fu31
h3o2fu31tr4.fu6
h3o2fu31tr4.fu15

Polyrate FILE NAME

poly.fu5 input data file
poly.fu31 input data file
poly.fu6 long output file
poly.fu15 summary output file

16.D. Summary of test runs

This section contains a summary of the test runs which make up the *POLYRATE* test suite.

Analytic potential

ch5j1tr1	ch5j1tr2	ch5j1tr3	ch5j1tr4	ch5j1tr5	ch5j1tr6
ch5j1tr7	ch5j1tr8	ch5j1tr9	ch5j1tr10	ch5j1tr11	ch5j1tr12
ch5j2tr1	ch5j2tr2	ch5j2tr3	clhbrtr1	clhbrtr2	clhbrtr3
clhbrtr4	clhbrtr5	cmctr1	cwmctr1	cwmctr2	cwmctr3
h3tr1	h3tr2	h3tr3	h3tr4	ho2tr1	ho2tr2
ho2tr3	ho2tr4	oh3tr1	oh3tr2	oh3tr3	oh3tr4
oh3tr5	oh3tr6	oh3tr7	oh3tr8	oh3tr9	oh3tr10
hnitr1	h2nitr1	ch4otr1	cmctr3	ch5fu51tr1	nh3tr2
ohcltr1	ohcltr2	ohcltr3	ohcltr4	ohcltr5	ch3tr1

fu30

ch5fu30tr1	ch5fu30tr2	ch5fu30tr3	ch5fu30tr4	ch5fu30tr5	ch5fu30tr6
oh3fu30tr1	oh3fu30tr2	oh3fu30tr3	oh3fu30tr4	oh3fu30tr5	oh3fu30tr6

fu31

ch5fu31tr1	oh3fu31tr2	ohfu31tr1	h3o2fu31tr1	h3o2fu31tr2	h3o2fu31tr3
h3o2fu31tr4					

fu40

ch5fu40tr1	ch5fu40tr2	ch5fu40tr3	ch5fu40tr4	ch5fu40tr5	ch5fu40tr6
oh3fu40tr1	oh3fu40tr2	oh3fu40tr3	oh3fu40tr4	oh3fu40tr5	oh3fu40tr6

IVTST-0 with fu29

ch5j2itr1	ch5fu29tr1	ch5fu29tr2	ch4ohtr2	ch4cltr1
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IVTST-0 with fu50

ch5fu50tr1	ch4ohtr1
------------	----------

IVTST-1 with fu29

ch5j2itr2	ch4cltr2
-----------	----------

VTST-IOC with analytic potential and fu50

ch5fu50tr2 ch5fu50tr3 cmctr2 ho2tr5 nh3tr1

VTST-IOC with fu30, fu31, or fu40 and fu50

ch5icfu30tr1 ch5icfu30tr2 ch5icfu31tr1 ch5icfu40tr1 ch5icfu40tr2

VTST-ISPE with fu30, fu31, or fu40 and fu51

ch5fu51tr1 cmctr3

17. DESCRIPTION OF FILES IN RELEASE OF VERSION 17

The source code is partitioned into 33 FORTRAN files. Note that a full *POLYRATE* source consists of everything in the 33 FORTRAN files that including modules and parameters. A description of each *POLYRATE* subprogram is given in Subsection 17.A.1. Subsection 17.A.2 briefly describes the subprograms *SETUP* and *SURF*, which must be supplied by the user for runs that do not use any of the potentials provided for the test run suite. (Dummy versions that can be used for runs with *POTENTIAL = unit30, unit31, or unit40* are in *dumpot.f.*) The input and output files for the sample runs are explained in Subsection 16.C. The Unix C shell scripts are explained in Subsection 15.A and Section 18. A list of all the files in the distribution package can be found in Subsection 15.A.

The files *polymq.F* and *polysz.F* contain subprogram *MXLNEQ* for solving and inverting sets of simultaneous linear equations and *EISPACK* subroutines for determining matrix eigenvalues and eigenvectors. In the distributed version the machine precision variable *MACHEP* is set equal to 1.0×10^{38} in the *MXLNEQ* subprogram and to 2^{-46} in the *EISPACK* subprograms. We found these settings to be adequate for all machines on which we tested the code.

If you received the *POLYRATE* distribution in a tar format, then the files will automatically be distributed into the default *POLYRATE* directories as described in Section 15, except that this manual may be in a separate file (not tarred with everything else).

17.A. FORTRAN files and subprograms, MODULE files, and PES subroutines

Subsection 17.A.1 contains an alphabetical list and brief descriptions of all the subprograms that make up the *POLYRATE* program, Subsection 17.A.2 describes the *MODULE* files and the user-supplied potential energy subprograms, and Subsection 17.A.3 describes the FORTRAN files.

17.A.1. *Polyrate* subprograms

In many cases a more detailed description and some relevant references are given as comments at the beginning of the FORTRAN code for each subprogram. Note that subroutines may be divided into two groups, *POLYRATE*-proper routines and hooks

routines. Routines that start with a Y or have HOOK in the name are always part of the hooks.

ABSORD

Order the lowest eigenvalues and eigenvectors according to absolute values.

ACALC

Dummy routine in place of the corresponding ACESRATE routine.

ACESET

Dummy routine in place of the corresponding ACESRATE routine.

AITKEN

Performs a Lagrangian interpolation of the effective reduced mass from the reaction-path Hessian grid.

AITKF2

Checks the Aitken interpolation for "bad" values, i.e., interpolants that do not lie within the original range of the data.

AITKNF

Performs Aitken interpolation.

ALFCT

Computes the logarithmic values of factorials of even integers used in the calculation of the Bernoulli numbers.

ANCOEF

Computes the anharmonicity coefficients x_{ij} and the constant term (E_0) from dimensionless normal coordinate force constants evaluated previously by subprogram GRADDR, and then calculates the vibrational ground-state energy and fundamentals from the anharmonicity coefficients and the constant term.

ANGL

Function that calculates the angle i-j-k.

ANGLCS

Function that calculates the sine and cosine of the valence angle between two difference Cartesian vectors.

ANGLV

Calculates the angle between two vectors.

ANHARM

Computes anharmonicities for bound normal modes. The anharmonicity option is selected by the anharmonicity keywords.

ANHDER

Computes the 3rd and 4th derivatives along the generalized normal modes.

ARMUEF

Calculates the argument of the exponential in equation 19 of Ref. 48 in Section 23 for the interpolation of the effective mass for CD-SCSAG tunneling.

ARMUEF1
Calculates μ^{CDSC} for the extrapolated points in IVTST-M with the hooks.

ARRSM
Calculates the moment of inertia for the points along the reaction path given in fu31 and reorders them according to the value of s .

ARRSV
Reorders the value of $V_{\text{MEP}}(s)$ given as input in fu31 according to s .

ASYMALP
Calculates the direction cosines for an asymmetric gyrator.

ASYMCAL
Prints the moment of inertia of a group of asymmetric gyrators.

ASYMENT
Calculates the moment of inertia of an asymmetric gyrator.

ASYMNZD
Sets up the calculation of moment of inertia of an asymmetric gyrator.

ASYMRMI
Calculates the moment of inertia of an asymmetric gyrator.

AVERJ
Calculate an integral over all possible orientation of the angular momentum vector \mathbf{J} with its length fixed in the evaluation of the number of available states.

BANGLE1
Calculates the Wilson B-matrix ($\partial R/\partial X$) for bending.

BCALC
Calculates the curvature components and the total curvature for the small-curvature semiclassical adiabatic tunneling calculations.

BCALCP
Calculates Page-McIver curvature factors B_{mF} .

BCALC0
Calculates the small-curvature effective mass at the saddle point by linear interpolation.

BTORSN
Calculates the Wilson B-matrix ($\partial R/\partial X$) for a torsion.

- BMAT**
Calculates the Wilson B-matrix, $\partial R/\partial X$, where R are the curvilinear internal coordinate vectors and X are the Cartesian coordinate vectors.
- BMAT1**
Forms the Wilson B-matrix row corresponding to a linear stretch coordinate between atom j and atom i . The input is in the form of difference Cartesians.
- BMAT2**
Forms the rows of the Wilson B-matrix corresponding to a bent triatomic.
- BMAT5**
Forms the Wilson B-matrix rows corresponding to degenerate linear bends. The input is in the form of two difference Cartesians defining the linear triatomic. The definitions of the bends are those of Hoy et al. (*Mol. Phys.* **24**, 1265 (1972)).
- BMAT5X**
Forms the Wilson B-matrix rows corresponding to degenerate linear bends. The input is in the form of two difference Cartesians defining the linear triatomic by first determining a transformation matrix to rotate the coordinate system such that the molecule lies along the Z axis.
- BIMAT**
Calculates the generalized inverse of the Wilson B-matrix.
- BTEN1**
Forms the Wilson B-matrix 2nd derivative elements for a bond-length curvilinear internal coordinate in terms of its first derivatives and magnitudes of difference Cartesians.
- BTEN2**
Computes the bent triatomic 2nd derivative Wilson B-matrix elements.
- BTEN5**
Defines the 2nd derivative Wilson B-matrix elements for the degenerate linear bend in terms of the difference Cartesian vectors that define its bonds.
- BTEN5X**
Defines the 2nd derivative Wilson B-matrix elements for the degenerate linear bend in terms of the difference Cartesian vectors that define its bonds so that the molecular frame has the linear bend along the Z axis.
- BTENR**
Transforms the Wilson B-matrix elements from difference Cartesians to Cartesians.
- BTENS**
Calculates the tensor $(X/R \cdot R^2/X^2 \cdot \partial X/\partial R)$.
- BOLTZ**
Performs the Boltzmann average of the tunneling probability by a numerical integration with respect to E .

BRENT
Implements Brent's line minimization method. See Ref. 14 in Sect. 22.

BRNULI
Computes Bernoulli numbers used in the calculation of the WKB phase integral for a double-well potential.

CALCS
Computes the minimum distance between two geometries along the reaction path measured as a straight line between them.

CALCUB
Calculates the coefficients for the cubic polynomial used in the IVTST-M interpolation of $V_{MEP}(s)$ for unimolecular reactions.

CASE
Function that takes a string of 80 characters and converts the upper case letters in the string to lower case letters.

CENTER
Translates the center of mass of the system to the origin. Also calculates the determinant of the moment of inertia tensor, which is equal to the product of the three principal moments of inertia, since they are the eigenvalues of this matrix.

CENTR1
Determines the relative index assignments for a linear triatomic system (i-j-k). Atom i is placed in the center and the j-k assignments are made arbitrarily.

CENTRAL
Determines which atom is the central atom for a bent triatomic by using bond lengths, and calculates the bond angle.

CFLOAT
Function that takes a character string that contains a floating point number and converts it to a floating point number as defined by standard FORTRAN conventions.

CHKFRE
Checks if any of the vibrational frequencies are zero or imaginary.

CHKUP
Retrieves 3rd-order dimensionless normal coordinate force constants needed for calculating the vibrational ground-state energy and fundamentals for simple perturbation theory calculations.

CLASS
Finds the recursion relation for various classes of orthogonal polynomials used to find the nodes in Gaussian-type quadratures.

CMS
Calculates the center of mass of a subset of the system.

COG
Defines a cut-off Gaussian correction function.

COHBT1
Defines a cut-off hyperbolic tangent correction function.

COHBT2
Defines a cut-off hyperbolic tangent correction function.

COLSHF
Performs shifting of Hessian grid data in the storage arrays.

CONOUT
Prints out the curvilinear internal coordinates.

CONV40
Converts the geometries, gradients, and Hessians to atomic units.

CORTRM
Calculates Coriolis constants.

COSINE
Modified cosine function.

CROSSPROD
Calculate cross product of two vectors.

CRVVEC
Computes the curvature vector when unit40 input is used.

CUBST
Evaluates quantities necessary for the cubic-starting algorithm.

CUBIC
Finds the real root of $ax^3 + bx^2 + cx + d = 0$ closest to a given x value.

CUBIC2
Finds all roots (real and complex) of $ax^3 + bx^2 + cx + d = 0$.

CUSSPL
Spline fits the $\Delta G^{GT,0}(s)$ curve and finds the maxima and minima along the curve for a CUS calculation.

CVCOOR
Driver for the curvilinear internal coordinate bound-state analysis of non-linear polyatomics.

DAXPY
BLAS routine calculates $cx + y$.

DATTIM
Calls system-dependent subroutines DATE and TIME and writes the current date and time.

DEFAULT
Sets the defaults for all the parameters and variables used in *POLYRATE*. The defaults for each different section of input is set in a separate subprogram for clarity.

DEFENG
Sets the defaults for keywords used in the ENERGETICS section.

DEFGEN
Sets the defaults for keywords used in the GENERAL section.

DEFLT40
Determines if the needed fu40 keywords were read for the IOPth species. Default values are assigned when necessary.

DEFOPT
Sets the defaults for keywords used in the OPTIMIZATION section.

DEFPAT
Sets the defaults for keywords used in the PATH section.

DEFRAT
Sets the defaults for keywords used in the RATE section.

DEFSEC
Sets the defaults for keywords used in the SECOND section.

DEFSTA
Sets the defaults for keywords used in the all the stationary point sections.

DEFTUN
Sets the defaults for keywords used in the TUNNEL section.

DERIV2
Calculates the Hessian matrix by using a quadratic fit to the potential energy surface. Only the potential is used, not its first derivatives.

DERV
Calculates the curvature vector related to third derivatives for IVTST calculations.

DERV24
Computes the second derivatives of the potential by fitting the analytic first derivatives to a fourth-order polynomial around the desired location and then taking the derivative of this polynomial.

DERVAG

Evaluates the first derivative of an n -dimensional point by the two-point central difference method.

DFGN31

Set defaults for a fu31 file.

DGAMMA

Evaluates the gamma function of order $n \leq 3$.

DGEDI

Computes the determinant and inverse of a matrix using the factors computed by DGEFA.

DGEFA

Factors a double precision matrix by Gaussian elimination.

DIATOM

Computes the equilibrium properties for a diatomic reactant or product treated as a Morse oscillator.

DIGISTR

Function that transform a three digits integer into a string with 3 characters.

DIST

Function that calculates an interatomic distance.

DPMPAR

A service routine for HYBRD1 that computes machine precision.

DOGLEG

A service routine for HYBRD1 that determines the convex combination x of the Gauss-Newton and scaled gradient directions that minimizes $(ax - b)$ in the least-squares sense, subject to the restriction that the Euclidean norm of dx be at most delta.

DOPNM

Calculates the steepest-descent path and the generalized normal modes along the MEP.

DOREPR

Calculates the reactant and product properties.

DOREST

Prepares a restart calculation (i.e., reads data from unit fu1, etc.).

DORODS

Subroutine used to setup a RODS or VRP(ESD) calculation.

DORPH

Prepares the potential energy routines and sets up the MEP interpolation information for an electronic structure input file calculation.

DOSAFR
Calculates the saddle point frequencies.

DOSAGE
Calculates the saddle point geometry.

DDOT
BLAS routine computes the dot product of two vectors.

DOTPRD
Computes the dot product between two vectors.

DPR123
Computes the dot product between two vectors. First vector points from atom 2 to atom 1; second vector points from atom 2 to atom 3.

DPSWAP
Swaps two double precision numbers.

DSCAL
BLAS routine scales a vector by a constant.

DSWAP
BLAS routine interchanges two vectors.

EBND
Computes the energy of a level of a quadratic-quartic potential by the perturbation-variation method of Truhlar.

ECKARS
Interpolates the classical potential curve (IOP = 1) or the ground-state vibrationally adiabatic energy curve (IOP = 2).

ECKART
Defines an Eckart correction function.

ECKRT
Interpolates frequencies with an Eckart or a hyperbolic tangent function.

EHOOK
The driver for the energy routines in the modular version of *POLYRATE*.

EFFBATH
Computes the solvent contribution for the nonequilibrium solvation approximation.

EFMAIN
The driver for the eigenvector following algorithm.

EFPHES
Prints the Hessian matrix in eigenvector following algorithm.

EFORMD
Forms geometry step by eigenvector following algorithm.

EFOVLP
Determines the overlap of the geometry steps in the eigenvector following algorithm.

ELLIP
Computes elliptic integrals of the first and second kinds.

ELRPH
Evaluates generalized normal mode frequencies along the MEP when electronic-structure input is used [POTENTIAL = *unit30* or *unit40*].

ENDRODS
Subroutine used to finish a RODS or VRP(ESD) calculation.

ENDSLP
TSPACK subroutine that estimates the first derivative at the edges of the spline-fit interval to be used as the two extra conditions of the global spline under tension.

ENORM
Calculates the norm of a vector \mathbf{x} .

ENROUT
Writes reaction energetics information to FORTRAN unit fu6.

EPARTI
Function that computes the electronic partition function for IVTST calculations.

EPART
Computes the electronic partition function.

ERF
Calculates the error function.

EVIB
Calculates vibrational energy levels.

EWKB
Uses a Chebyshev quadrature to compute the primitive WKB approximation to the energy of a vibrational level of a quadratic-quartic potential.

EXPND
Expands triangular matrix \mathbf{A} to square matrix \mathbf{B} .

EXTRAP
Provides simple exponential extrapolations to the asymptotic values of quantities used in tunneling calculations.

EXTRP40
Sets the values for extrapolation when using unit fu40 input.

EXTRP
Fits a straight line to two points and carries out interpolations or extrapolations.

F1DI
Evaluates the function VALVAG for LINMNR.

F1DIM
Computes the one-dimensional energy for use in line minimization subroutines.

FACORD
Orders the force constants scaling factors according to the actual sequence of the curvilinear internal coordinate when FCSCALE is used.

FCINPT
Closes all the input files used in READ5.

FCMEP
Closes all the input and output files used in the MEP calculation.

FCN
Calculates an Eckart or a hyperbolic tangent function to be passed into the non-linear equation solver.

FCRATE
Closes all the input and output files used in the rate constant and the tunneling calculations.

FCSCL
Scales diagonal elements of the force constant matrix in curvilinear internal coordinates by corresponding scaling factors when FCSCALE is used.

FDIAG
Diagonalizes the force constant matrix and, optionally, prints out generalized normal mode information.

FDIAG2
Diagonalizes the Cartesian force constant matrix for Hessians from unit 29 input and, optionally, prints out generalized normal mode information.

FDJAC1
A service routine for HYBRD1 that computes a forward-difference approximation to the $n \times n$ Jacobian matrix associated with a specified problem of n functions in n variables. If the Jacobian has a banded form, then function evaluations are saved by only approximating the nonzero terms.

FICLSE
Closes a file given the status with which to close the file and the FORTRAN unit number to which the file is linked.

FINDL
Calculates the value for the RPL keyword in a VTST-IC input file (fu50) from a lower-level calculation.

FINDLV

Calculates the range parameter for an Eckart function.

FINOUT

Computes and prints out the final set of rate constants. Also obtains activation energies and bottleneck properties and analyzes the ratio of transition state theory results to those of canonical variational transition state theory.

FIOPEN

Opens all files needed for a given calculation except for the main input and the main output files (i.e., the files linked to FORTRAN units fu5 and fu6).

FITMAX

Finds the maximum in the adiabatic potential curve and the generalized free energy of activation curve using 3- and 5-point fits.

FITMX

Finds the maximum of function f with a 5-point fit.

FIVFT

Computes a quartic fit through five points for IVTST.

FIVPT

Computes a quartic fit of $F = ax^4 + bx^3 + cx^2 + dx + e$ to five points.

FORMA

Forms the generalized inverse of the B-matrix by the Pulay prescription.

FORMF

Forms the product $\mathbf{A}^T \mathbf{F} \mathbf{A}$, where \mathbf{A} is the A-matrix and \mathbf{F} is the Cartesian force constant matrix.

FORMF2

Forms the second term of curvilinear internal force constant matrix.

FORMG

Forms the G matrix.

FORMBT

Forms one xyz-band of the transformation matrix from Cartesian coordinates to difference Cartesians.

FRPRMN

Conjugate gradients minimization subroutine.

FRSORT

Sorts a set of frequencies into ascending or descending order.

GAUSQD

Calculates the weights and abscissa for N -point Gauss-Legendre quadrature.

GAUSSQ

Computes the nodes and weights for Gaussian-type quadrature.

GBSLVE

For a symmetric tridiagonal $n \times n$ matrix \mathbf{A} , this subroutine finds the component n of the n -dimensional vector \mathbf{v} such that $\mathbf{A}\mathbf{v} = \mathbf{c}$, where \mathbf{c} is a vector of all zeros except for 1 in the n -th position.

GBTQL2

Finds the eigenvalues and the first components of the eigenvectors of an asymmetric tridiagonal matrix by the implicit QL method.

GEMM

Performs fast matrix multiplication. From the BLAS library.

GENSEC40

Reads and processes the GEN40 section of the unit fu40 input.

GETFRQ

Obtains vibrational frequencies in cm^{-1} and generalized eigenvectors.

GFDIAG

This routine effectively diagonalizes the unsymmetric \mathbf{GF} -matrix, formed by product of \mathbf{G} and \mathbf{F} . The method is given by Macintosh and Michaelian (*Can. J. Spectrosc.* **24**, 65 (1979)). Only symmetric matrices are diagonalized here. the \mathbf{GF} product is never explicitly formed.

GFDIA2

This routine effectively diagonalizes the unsymmetric, rectangular \mathbf{GF} matrix by generalized inverse \mathbf{G} matrix method. The method is given in P. Pulay,; G. Fogarasi, *J. Chem. Phys.* **1992**, 96, 2856.

GHOOK

The driver for the first derivative routines in the modular version of *POLYRATE*.

GIVTST

Driver for zero- and first-order interpolated VTST (the global approach) as described in *J. Chem. Phys.* **95**, 8875 (1991).

GRAD

Evaluates derivatives along normal mode (or generalized normal mode) directions.

GRADDR

Calculates 3rd- and 4th-order dimensionless normal coordinate force constants.

GSETA

Dummy routine in place of the corresponding ACESRATE routine.

GSETM

Geometry optimization calculations needed when *POLYRATE* is used with MOPAC.

GSETP

Geometry optimization calculations needed when *POLYRATE* is used with an analytic surface.

HBT
Calculates the hyperbolic tangent at s_{MEP} .

HEADR
Prints out the program header.

HHOOK
The driver for the second derivative routines in the modular version of *POLYRATE*.

HINDRT
Calculates the moment of inertia for internal rotation.

HINDRT1
Calculates the moment of inertia for internal rotation of an asymmetric gyrotor.

HQSC
Semiclassical solution of a harmonic-quartic potential.

HRPART
Calculates the partition functions using the hindered-internal-rotator approximation.

HRSET
Setup calculation of hinder rotor partition function.

HVAL
TSPACK subroutine that evaluates a Hermite interpolatory tension spline at a point.

HYBRD
A service routine for HYBRD1.

HYBRD1
Finds a zero of a system of n nonlinear functions in n variables by a modification of the Powell hybrid method.

ICFDRP
Driver of the curvilinear internal coordinate bound-state analysis for reactants and products.

ICFDIAG
Driver for the curvilinear internal coordinate bound-state analysis of triatomics.

ICINT
Function that takes a character string which is really an integer and converts it to an integer. Error checking that is not possible using *ichar* is done.

ICODAL
Function that translates a set of alphanumeric characters into an integer code.

ICOUTRP
Prints out the curvilinear internal coordinates for reactants and products.

ICSAVE
Copies frequencies and eigenvectors for bound modes from curvilinear internal coordinate work spaces to the locations used by Cartesian coordinate procedures.

ICSVRP
Copies the bound-state analysis results from the curvilinear internal coordinate work spaces to the locations used by Cartesian coordinate procedures.

IDAMAX
BLAS routine finds the index of element having maximum absolute value.

IMOM40
Computes the moment of inertia.

INTCOR
Defines the curvilinear internal coordinates for reactants and products by splitting the global connectivities into subsets.

INTERSC
Finds out the intersection of two integer arrays.

INTF31
Subroutine that reorders the information on stationary points taken from fu31 and puts it on the correspondent arrays.

INTRVL
TSPACK function that locates the interval in the input data for a spline fit that contains a certain point.

INVRT
Inverts a one-dimensional array of size N .

INVRT2
Inverts a two-dimensional array of size N_1, N_2 .

INVRT3
Inverts a three-dimensional array of size N_1, N_2, N_3 .

INERTA
Calculates the moment of inertia for a linear molecule or a diatomic, or the product of the moments of inertia for a non-linear molecule in au.

INITVRC
Initializes the array IATSV(I,J) for the generalized transition state in variable reaction coordinate calculations because START section is not give in the fu5 file.

INITZE
Initializes some of the variables and arrays.

INTAB
Writes out a table of all the input parameters.

INTABS
Writes out a table of all the input parameters associated with the stationary point sections.

INTEGR
Integration driver for the PATH subroutine.

INTEUL
Performs Euler steepest-descents integration.

INTFNC
Calculates and returns the negative of the normalized gradient for the path integration.

INTPM
Page-McIver reaction path algorithm..

INTPM2
Computes $\partial s / \partial t$ for the Page-McIver method.

INTRPL
Calls POLFIT to do polynomial interpolation for a given quantity.

IRCX
Intrinsic-reaction-coordinate pathfinder, using the method of Schmidt, Gordon, and Dupuis.

IRFTG0
Computes the information necessary for dealing with the frequencies, Hessians, and curvature vectors for unit fu40 input.

IRPHZ
Does interpolations for the zero-order IVTST calculations.

IRPH
Does interpolations for the first-order IVTST calculations.

ISWAP
Swaps two integers.

IVTMA
Interpolates the exponent for calculating μ^{CDSC} by IVTST-M through the hooks.

IVTMD
Calculates the extrapolated points with IVTST-M through the hooks.

IVTMH
Sets up the parameters required for IVTST-M through the hooks.

IVTST0
Calculates the classical (i.e., Born-Oppenheimer) potential energy curve and the adiabatic ground-state potential energy curve along the reaction path in a zero-order VTST-IOC calculation.

JACSCF
Computes all eigenvalues and eigenvectors by the Jacobi method.

KAPA
Calculates transmission coefficients for IVTST calculations.

KAPPAS
Calculates the transmission coefficients using the uniform semiclassical WKB method (MEPSAG) and the small curvature method (SCSAG).

KAPVA
Calculates the tunneling corrections for a set of temperatures.

KG1
Nodes and weights for Kronrod quadrature.

LCG34
Calculates the tunneling probabilities $P^{LCG34}(E)$ by the LCG34 method.

LCGIT
Builds the grid for the 1D spline under tension.

LCGNAD
Defines the limit between adiabatic and nonadiabatic regions for the LCT3 and LCT4 methods.

LCGRD
Builds a 2D rectangular grid from selected points of the PES.

LCGSRT
Sorts the uniformized LCG3 or LCG4 probabilities.

LCGTHE
Calculates the primitive tunneling amplitude $T_{tun}(\tilde{s}_0, n_p)$ by integration over the straight-line LCG3 or LCG4 tunneling path using both Gauss-Legendre quadrature and the trapezoidal rule.

LCGTR
Transforms a point of the PES to a point of the 2D rectangular grid.

LCNORM
Calculates the generalized normal mode coordinates corresponding to the scaled displacements along the LCG3 or LCG4 tunneling path. These coordinates are needed for eq. (14) of Ref. 20 in Sect. 23.

LCNWTN
Performs a Newton root search to determine the s value corresponding to a point along the LCG3 or LCG4 tunneling path, using eq. (22) of Ref. 20 in Sect. 23.

LCPATH
Calculates the mass-scaled Cartesian coordinates of the LCG3 or LCG4 tunneling path as given in eq. (8) of Ref. 20 in Sect. 23.

- LCPROB**
Calculates the uniformized LCG3 probabilities from the primitive values $P_{prim}^{LCG3}(E)$ using eq. (30) of Ref. 20 in Sect. 23.
- LCPROJ**
Calculates the projections of the generalized normal mode coordinates onto the LCG3 path using eq. (14) of Ref. 20 in Sect. 23.
- LCSET**
Sets up the LCG3 tunneling path and calculates the angle between the MEP and the tunneling path [eq. (12) of Ref. 20 in Sect. 23] along with the vibrational period of eq. (17) of Ref. 20 in Sect. 23.
- LCSINT**
Performs the numerical integration over tunneling termini along the reaction path as given by eq. (9) of Ref. 20 in Sect. 23.
- LCSTX**
Sets up the tunneling path for excited final states and determines the highest excited state involved in LCG3 calculations.
- LCTP**
Determines the ending terminus of the tunneling path for a given excited state of the LCG3 exit arrangement and a given initiating terminus \tilde{s}_0 using the resonance condition.
- LCTV4**
Searches for a LCT4 nonadiabatic region within a LCG3 adiabatic region.
- LCVCOR**
Calculates the effective potential for the nonadiabatic region in an LCG3 calculation as given in eq. (28) of Ref. 20 in Sect. 23.
- LCVIB**
Calculates the vibrational zero-point energies of eq. (5) of Ref. 20 in Sect. 23.
- LCVMOD**
Calculates the zero-point energy and potential for a given normal mode in an LCG3 calculation.
- LCXINT**
Calculates the integrand in the phase integral along the adiabatic segments of an LCG3 tunneling path.
- LCXTAU**
Calculates the vibrational period, τ , for LCG3 calculations as given by eqs. (17) and (18) of Ref. 20 of Sect. 23.
- LCZCRT**
Determines which region (adiabatic or nonadiabatic) a given point on an LCG3 tunneling path is in by using eqs. (23) and (24) of Ref. 20 in Sect. 23.

LIN
Solves a set of simultaneous linear equations.

LIN2
Solves two simultaneous linear equations in the matrix form $\mathbf{Ax} = \mathbf{b}$.

LININT
Does the linear interpolation.

LINMNR
Finds the minimum along a given direction.

LINMN
One-dimensional minimization search.

LOCATE
Determines the indices on the grid of reaction path coordinates corresponding to a point on the LCG3 tunneling path determined by LCNWTN.

LOCS
Called by AITKF2 to locate the s value about which to center the interpolation grid. The grid is centered about the closest point on the Hessian grid.

LOPT40
Translates the GEN40 input keywords to LOPT values.

LSAME
Service routine for GEMM.

LUDCMP
Performs an LU decomposition using an algorithm from W. H. Press, B. P. Flannery, S. A. Teukolsky, and W. T. Vetterling, *Numerical Recipes* (Cambridge University Press, Cambridge, 1986), p. 31.

LUBKSB
Service routine for LUDCMP.

MAIN
Driver for *POLYRATE*. The main input and output files (i.e., those files linked to FORTRAN units fu5 and fu6) are opened in this subprogram.

MATINV
Performs matrix inversion using LU decomposition routines.

MATX
A general-purpose matrix multiplication routine.

MEPINV
Stores the information saved as functions of the reaction coordinate in order of increasing s by inverting the order of storage.

MEPOUT
Prints MEP information to the long output file, i.e., to unit fu6.

MEPSRT
Does a bubble sort of the MEP information in order of increasing s .

MNBRAK
Finds an interval which contains a minimum.

MOPEG
Dummy routine to disable the keyword MOPAC in *POLYRATE*.

MOPHES
Dummy routine to disable the keyword MOPAC in *POLYRATE*.

MOPOPT
Dummy routine to disable the keyword MOPAC in *POLYRATE*.

MOPSET
Dummy routine to disable the keyword MOPAC in *POLYRATE*.

MTINV2
Performs matrix inversion using singular value decomposition routines

MUCDSC
Computes the centrifugal-dominant small-curvature effective mass.

MUSC
Computes the SCSAG effective mass.

MUSCCD
Computes the CD-SCSAG effective mass.

MXLNEQ
Solves a set of simultaneous linear equations in the matrix form $\mathbf{Ax} = \mathbf{b}$, and evaluates the determinant of \mathbf{A} .

NEARWC
computes the nearest reactant or product endpoint used in frequency interpolation.

NEWT
Does a Newton-Raphson search for a stationary point.

NEXTPT
Computes the next point on the MEP.

NORMOD
Computes generalized normal mode frequencies and directions.

NOROUT
Prints out generalized normal mode information.

NOST31
Processes the information about non-stationary points along the reaction path.

NSTATE

Calculates the total number of accessible states in LCG3 calculations that include contributions from tunneling into excited states.

NUMFRQ

Calculates the number of internal vibrations.

OBTSRT

Compares two elements and stores the bigger one.

OHOOK

The driver for the geometry optimization routines in the modular version of *POLYRATE*.

OPENFI

Opens a file given the unit number, the status, and the file name.

OPT50

Checks for any conflicts in the options for the VTST-IOC calculations.

OPT50C

Checks for consistency between the VTST-IOC options chosen in the unit fu50 input.

OPTION

Checks for consistency between the options chosen in the unit fu5 input and writes out information about the options selected.

ORDS31

Reorders the information stored on the array save31 containing all the information given as an input in fu31.

OVRLP

Computes overlaps between old and new normal mode eigenvectors and warns if a crossing has occurred.

PATH

Calculates the reaction path.

PHID

Computes the phi function needed for uniform semiclassical quantization of a double-well potential.

PHSINT

Computes the phase integral for the WKB method.

PMOI

Calculate principal moment of inertia and their unit vector along the ith principal axis for each reactant or complex of reactants

POLINT

Uses Neville's algorithm to interpolate a given quantity.

POLYAT
Calculates the equilibrium properties of a linear or a nonlinear polyatomic reactant or product.

PRCORD
Prints the bond lengths, bond angles, and dihedral angle along the reaction path.

PREGPT
Spline fits s , V , and I^{-1} for IVTST-M calculations carried out through the hooks.

PREP
The driver for all the preparation routines in the modular version of *POLYRATE*.

PREPJ
The driver for all the preparation routines needed by REACT1, REACT2, PROD1, PROD2, START, and PATH. See Sect. 8.C.

PRMEP
Prints extra MEP information to units fu25-fu27.

PRNTBT
Prints the B-matrix stored in lower triangular form in clusters of 5 rows.

PRNTFC
Prints a lower triangular matrix (in clusters of 5 rows).

PROJTI
Forms the projection operator and calculates the projected Cartesian force constant matrix for IVTST calculations.

PROJCT
Calculates the projected force constant matrix and normalized gradient.

PROJF
Forms the projection operator and projects out the gradient from the curvilinear internal force constant matrix.

PRSQ
Prints out a square matrix.

PSAG
Locates points (translational turning points) where the energy is equal to the adiabatic potential and then computes the imaginary-action integral.

PSATP
Finds turning points on the adiabatic barrier.

PSATP2
Finds turning points on the adiabatic barrier.

PSATX
Finds turning points on the adiabatic barrier.

PSATX2

Finds turning points on the adiabatic barrier.

PTORS

Function that calculates the torsion angle i-j-k-l.

PTQVIB

Calculates vibrational partition functions by simple perturbation theory using the ground-state energy and fundamentals. [See Ref. 23 in Sect. 23 for the equations without Coriolis terms in E_0 and see Ref. 35 in Sect. 22 for the Coriolis contributions to E_0 .].

PTSEC40

Reads and processes the POINT40 sections in the unit fu40 input.

PYC1

TSPACK subroutine that employs a three-point quadratic interpolation method to compute local derivative estimates associated with a set of data points. The derivatives are chosen so that the Hermite interpolatory tension spline fulfils the user selected end conditions.

PYC1P

TSPACK subroutine that employs a three-point quadratic interpolation method to compute local derivative estimates associated with a set of data points. The derivatives are chosen so that the Hermite interpolatory tension spline fulfils the defaulted end conditions.

PYC2

TSPACK subroutine that solves a linear system for a set of first derivatives associated with a Hermite interpolatory tension spline. The derivatives are chosen so that the Hermite interpolatory tension spline fulfils the user selected end conditions.

PYC2P

TSPACK subroutine that solves a linear system for a set of first derivatives associated with a Hermite interpolatory tension spline. The derivatives are chosen so that the Hermite interpolatory tension spline fulfils the defaulted end conditions.

PYCOEF

TSPACK subroutine that computes the coefficients of the derivatives in the symmetric diagonally dominant tridiagonal system associated with the C-2 derivative estimation procedure for a Hermite interpolatory tension spline.

QFORM
Computes the **QR** factorization.

QQPOT
Fits the quadratic-quartic potential to two points along the generalized normal coordinate using two given distances.

QRFAC
This is a service routine for HYBRD1 that uses Householder transformations with column pivoting (optional) to compute a QR factorization of the $m \times n$ matrix **A**.

QRSENE
Calculates the quantized reactant eigenstates for tunneling.

QSLVE
Computes a quadratic fit to the function values at two points and the derivative at one point, and uses the fit to perform an interpolation.

QTQVIB
Calculates the quantized reactant-state partition functions.

QUADFT
Computes a quadratic fit to three points.

QUADR
Solves a quadratic equation.

QUADTW
For a function $y(x) = A + B(x-C)^2$, finds B and C from the given information.

RBASIS
Dummy routine in place of the corresponding ACESRATE routine.

RBATH
Reads the keyword options for the BATH keyword in PATH section.

R1UPDT
A service routine for HYBRD1.

R1MPYQ
A service routine for HYBRD1.

RACES
Reads the ACES namelist options.

RALL31
Reads all the data from fu31 and store it in the save31 array.

RATE
Computes the generalized free energy of activation curves and rate constants.

RAN1
Random number generator from Numerical Recipes.

RANALY
Reads the temperatures at which a detailed analysis is desired.

RATOMS
Reads the atomic symbols and optionally the masses.

RCNST
Reads the coordinates that will be held constant in any optimization of a stationary point geometry.

RD40EXTR
Reads the EXTRAP40 section of the file fu40 input.

RD40GEN
Reads the GEN40 section of the file fu40 input.

RD40RP
Reads the reactants and products sections of the file fu40 input.

RDETIL
Reads the LCG3 output details information.

RDIATM
Reads the diatomic options for stationary points.

RDIFFU
Reads the diffusion coefficients or coupling constants for the nonequilibrium solvation calculation.

RDHESS
Reads a Hessian matrix from file fu40.

RDLN
Reads a line from an input file without conversion.

REACT
Computes the geometry and properties of a reactant or product.

REACTCOM
Calculate center of mass for reactants

READ5
Reads input from the file linked to FORTRAN unit fu5.

READIC
Reads the VTST-IOC correction data from the file linked to FORTRAN unit fu50.

READINT
Reads INTDEF keyword information to get the connectivities.

READLN
Reads a line from an input file and converts all characters to uppercase.

REARG1
Rearranges a set of frequencies to an order that is input by the user.

RECS31
Recalculates all the values of s given in fu31 as distances along straight lines between two consecutive geometries.

REDGEO
Determines the geometry of the reactant, product, or saddle point, i.e., linear or nonlinear, and the number of degrees of freedom.

REDMU
Evaluates the effective mass for a given value of s for IVTST as described in *J. Chem. Phys.* **95**, 8875(1991).

REFOPT
Reads the options for eigenvector following algorithm.

RELEC
Reads the electronic degeneracies for stationary points.

RENERG
Reads the ENERGETICS section of the file fu5 input.

REPFL
Replaces the low-frequency mode along the path when the IVTST0FREQ keyword is used.

RES1
Reads the variables for the Euler-with-stabilization (ES1) integrator.

RESTOR
Reads information from the file linked to FORTRAN unit fu1 for a restart rate constant calculation.

RESTRT
Writes information to the file linked to FORTRAN unit fu1 for a restart rate constant calculation.

RETRP
Reads the EXTRAP section of the file fu5 input.

REZERO
Reads the input for the zero of energy.

REZERU
Reads the unit for the zero of energy.

RFCFAC
Reads FCSCALE keyword information to get the force constant scaling factors when FCSCALE is used.

RFREQ
Reads frequencies from unit fu29 for IVTST calculations.

RGEM31
Reads the options for geometry input in fu31.

RGEN
Reads general information for IVTST calculations from unit fu29.

RGEN31
Reads the options in the general section of an fu31 input file.

RGENER
Reads the GENERAL section of the file fu5 input.

RGEOM
Reads the initial geometry for reactants or products.

RGRD
Read gradient from unit fu29 for IVTST calculations.

RGRD31
Reads the options for gradients input in fu31.

RGSAD
Reads the initial geometry for the saddle point or starting geometry.

RGSPEC
Reads the range of s for which the generalized free energy curve is to be computed.

RGSZM
Reads the initial geometry for the saddle point or starting geometry using z-matrix form.

RGTEMP
Reads the range of s values for each temperature for which the generalized free energy curve is to be computed.

RGZMAT
Reads the initial geometry for the saddle point or starting geometry using z-matrix form. This is used only when the geometry optimization is done using ACES.

RHESS
Reads Hessians from unit fu29 for IVTST calculations.

RHES31
Reads a Hessian matrix from fu31.

RHESSC
Reads HESSCAL options from unit fu5.

RHFSEC
Checks whether frequencies or Hessians are to be read.

RHSS31
Reads the options for Hessian input in fu31.

RICOPT
Reads the VTST-IOC options.

RLCT
Reads the information for large-curvature tunneling calculations.

RLINE
Utility program for reading a line of input and parsing off extra spaces and comments.

RLOWFR
Reads the information given by the LOWFREQ keyword.

RMUVT
Reads μ VT options.

ROPT
Reads the OPTIMIZATION section of the file fu5 input.

ROPT31
Reads the general section of an incomplete fu31 file and sets the options for writing it.

ROPTOP
Reads the options for the OPTIMIN and OPTTS keywords.

RPART
Computes the classical rotational partition function.

RPARTI
Computes the classical rotational partition function for IVTST calculations.

RPATH
Reads the PATH section of the file fu5 input.

RPH31
Sets up the defaults and routes for reading an fu31 file.

RPH40
Sets up the electronic structure information from file fu40.

RPHAIT
Performs NINT-point Lagrangian interpolation of an asymptotic function used in the electronic structure input file option (i.e., the RPH formulation).

RPHB
Calculates the curvature component of mode k .

RPHDXN
Converts from unweighted to mass-scaled gradient and normalizes the gradient vector.

RPHEXP
Evaluates asymptotic exponential functional forms.

RPHFIT
Evaluates parameters of the asymptotic exponential fits.

RPHFT2
Computes 2-point fits to 3-parameter asymptotic exponential functions with the range parameter fixed.

RPHFT3
Computes 3-point fits to 3-parameter asymptotic exponential functions.

RPHG1
Function used in the root search for the range parameter in an asymptotic functional form.

RPHG2
Function used in the root search for the range parameter in an asymptotic functional form.

RPHG5
Function used in the root search for the range parameter in an asymptotic functional form.

RPHINT
Interpolates RPH information.

RPHMEF
Computes the effective tunneling mass for CD-SCSAG from the interpolated value of the exponent in equation 19 in Ref. 48 in Section 23.

RPIVOT
Reads the Cartesian coordinates of pivot points

RPNT31
Reads the information from each point in the fu31 file.

RPHNWT
Performs a Newton search for the zero of one of the functions $G(C)$ defined in Subsection 9.D.1.

RPHRD1
Reads electronic structure input information and sets up tables used for interpolation.

RPHRD2
Reads electronic structure input information.

RPHSET
Sets up electronic structure input interpolation information.

RPHTRF
Transforms the matrix of second derivatives with respect to unweighted coordinates to second derivatives with respect to mass-scaled coordinates.

RPHTRX
Converts mass-scaled to Cartesian coordinates or Cartesian to mass-scaled coordinates.

RPHWRT
Writes reaction path Hamiltonian interpolation information.

RPOTET
Reads the potential type.

RPOTGE
Reads the potential type to be used for optimizing the geometry.

RPRFL
Replaces the low-frequency mode of the stationary points when the IVTST0FREQ keyword is used.

RPRMEP
Reads the parameters for printing the extra MEP information.

RPRVIB
Reads the range of s between which extra vibrational mode information and Q_{vib} are printed.

RPSEC40
Reads and processes a reactant or product section of the file fu40 input.

RQCOM
Calculates the distance between the centers of mass of two reacting species.

RQRST
Reads the information for the quantized-reactant-state tunneling calculation.

RQUAD
Reads the quadrature information for the tunneling calculations.

RRANGE
Reads the range of s values for which the reaction path is to be computed.

RRATE
Reads the RATE section of the file fu5 input.

RRPM
Reads the value of the RPM keyword of the file fu5.

RRSTRT
Reads the restart options.

RSCT
Reads the options for small curvature tunneling.

RSECND
Reads SECOND section of the file fu5 input.

RSFRST
Reads the variables for a special step size for the first n steps.

RSP
EISPACK diagonalization routine. Finds the eigenvalues and eigenvectors of a real symmetric packed matrix.

RSPDRV
Accepts unpacked arrays the way EISPACK subroutine RS does, packs them for RSPP, and calls RSPP to compute eigenvalues and eigenvectors.

RSPECL
Reads special values of s at which a normal mode analysis is to be done.

RSPP
Diagonalizes a real symmetric matrix in packed form.

RSSTOP
Reads the options of the SPECSTOP keyword.

RST
Diagonalizes a real symmetric matrix using RSPP.

RSTAT
Reads all the stationary point sections of the file fu5 input.

RSTATE
Reads the options used with a state-selected rate constant calculation.

RSVRC
Reads the range and step size of variable reaction coordinate

RTEMP
Reads the temperatures at which the rate constant is to be calculated.

RTITLE
Reads the title.

RTOR
Reads the TOR keyword options.

RTOROPT
Reads the TOROPT keyword options.

RTPJAC
Prints the MEP and large curvature representative tunneling termini in mass-scaled Jacobian coordinates.

RTUNNL
Reads the tunneling parameters.

RUNM31
Reads the options for the IVTST-M interpolation on a unimolecular side (or side with a well) on the reaction path.

RVAR6
Reads keyword values in a restart calculation.

RVARY
Reads the variable anharmonicity options desired for reactants, products and or the saddle point.

RVPERT
Reads the options for using simple perturbation theory for the vibrational partition functions.

RVRANG
Reads the anharmonicity options desired for the reaction path for specific regions of the reaction coordinate s .

RVRCOPT
Reads the options for variable reaction coordinate calculations.

RWKB
Reads the WKB anharmonicity options.

RWORD
Parses a character string to find the next word.

RZVAR
Reads the z-matrix variables for the geometry optimization of the reactants, products and the saddle point.

SADDLE
Computes the saddle point geometry (with subroutine NEWT) and other saddle point properties.

SADENG
Computes the saddle point energetics.

SADSEC40
Reads and processes the SADDLE40 section of the file fu40 input.

SAVEXRP
Calculate Cartesian coordinate of each reactant and the other pivot points relative to the first pivot point of reactant 1.

SECCEN
Calculates second derivatives by the method of central differences.

SECCEP
Calculates second derivatives by the method of central differences; only computes part of the Hessian matrix.

SENOUT
Writes saddle point energetics information to unit fu6.

SETANH
Sets variables associated with the variable anharmonicity option.

SETLGS
Converts the variables and flags set by evaluating the keywords to the corresponding LGS flags.

SETUP
Called once to initialize the potential, as discussed in Sect. 8.C.1. Ordinarily, this is supplied by the user. The distribution, however, contains a dummy version for use as described in Section 8.C, as well as sample versions for the test runs.

SETVAR
Converts the variables and flags set by evaluating the keywords to the corresponding *POLYRATE* variables.

SGND3V
Stores the sign of the third derivative in an array for use in computing the Morse turning point.

SHFTSAD
Shifts saddle point information when using unit fu40 input.

SIGS
TSPACK subroutine that determines the value of the tension factors for a spline under tension fit.

SIMPSN
Calculates the phase integral by using Simpson's rule in the quantized-reactant-state tunneling calculation.

SINE
Modified sine function.

SNHCSH
TSPACK subroutine that computes approximate values for modified hyperbolic functions.

SORTG
Sorts frequencies in descending order.

SORT31
Sorts an nx2 array according to the values on the first column.

SPL1B1
Converts the spline fit data supplied by SPL1D1 to four arrays containing the coefficients for local cubic fits.

SPL1B2
Computes the spline interpolation using the cubic polynomial coefficients supplied by SPL1B1.

SPL1D1
Computes the second derivative array for spline interpolations.

SPL2D
Calls the 2D spline under tension.

SPL31
Subroutine that interfaces *Polyrate* and TSPACK. This subroutine is used when the information is given by means of fu31, which allows for a different number of data for each property.

SPLNB
Calculates the value of the curvature components at any point along the reaction path by using spline under tension fitting.

SPLNF
Calculates the value of the frequencies at any point along the reaction path by using spline under tension fitting.

SPLNM
Calculates the value of the moment of inertia at any point along the reaction path by using spline under tension fitting.

SPLNMF
Calculates the value of the effective mass for CD-SCSAG tunneling at any point along the reaction path by using spline under tension fitting.

SPLNV
Calculates the value of the energy along the reaction path, $V_{MEP}(s)$ at any point along the reaction path by using spline under tension fitting.

SPLM31
Calculates the value of the moment of inertia at any point along the reaction path by using spline under tension fitting. This subroutine is used when the input is given in fu31.

SPLRMI
Calculates the value of the moment of inertia at any point along the reaction path by using spline under tension fitting.

SPLSPV
Spline fits ΔV in the ISPE method. This subroutine is used with fu51.

SPLV31
Calculates the value of the energy along the reaction path, $V_{\text{MEP}}(s)$, at any point along the reaction path by using spline under tension fitting. This subroutine is used when the input is given in fu31.

SSAVE
Saves the reaction path results calculated in subroutine PATH.

SSX
Computes the distance between two Cartesian geometries as a straight line in mass-scaled coordinates.

STAT31
Calculates stationary point properties from the information read in fu5 and fu31.

STAUV
Calculates the correction term used in computing the improved generalized transition-state partition function from the generalized transition-state partition function.

STLGSJ
Converts the stationary-point-dependent variables and flags set by evaluating the keywords to the corresponding LGS flags.

STOR
TSPACK subroutine that determines floating point number characteristics.

STVARJ
Converts the stationary-point-dependent variables and flags set by evaluating the keywords to the corresponding *POLYRATE* variables.

SURF
Evaluates the potential, as discussed in Subsection 8.C.1. Ordinarily this is supplied by the user, but the *POLYRATE* distribution provides a dummy version for use as discussed in Section 8.C, as well as sample versions for the test runs.

SVBKSJ
This routine, taken from Numerical Recipes, is for the backward substitution of SV decomposition.

SVBCMP
This routine, taken from Numerical Recipes, is for the singular value decomposition procedure.

TABLE
Writes a summary of selected forward rate constants to the file linked to FORTRAN unit fu15.

TAUV
Calculates the cutoff partition function for improved canonical variational theory.

THETA2
Computes the phase integral for the semiclassical eigenvalues of a single-well potential.

THETA3
Computes the phase integral for the semiclassical eigenvalues of a double-well potential.

TITLE
Reads and writes the title of the job.

TP
Locates the turning points at a given energy for a normal mode vibration.

TPCDSC
Calculates the centrifugal-dominant small-curvature turning point.

TQL2
Computes the eigenvalues and eigenvectors of a symmetric tridiagonal matrix by the QL method [H. Bowdler, R. S. Martin, C. H. Reinsch, and J. H. Wilkinson, *Num. Math* **11** (1968) 293].

TQLRAT
Finds the eigenvalues of a symmetric tridiagonal matrix by the rational QL method [C. H. Reinsch, *Comm. ACM* **16** (1973) 689].

TRANLF
Converts Cartesian to mass-weighted coordinates or mass-weighted to Cartesian coordinates store in lower triangular form.

TRANS
Converts Cartesian to mass-scaled coordinates or mass-scaled to Cartesian coordinates. Also converts derivatives of the potential energy function between these coordinate systems.

TRANS2
Converts Cartesian to mass-scaled coordinates (IOP = 1) or mass-scaled to Cartesian coordinates (IOP = 2) used in IVTST calculations.

TRANG
Transforms gradient in Cartesians to curvilinear internal coordinates.

TRANFC
Transforms Hessian in Cartesians to curvilinear internal coordinates.

TRANSF
 Converts Cartesian to mass-weighted coordinates (IOP = 1) or mass-weighted to Cartesian coordinates (IOP = 2).

TRBAK3
 Forms the eigenvectors of a real symmetric matrix by back transformation of the eigenvectors of the similar symmetric tridiagonal matrix.

TRED3
 Reduces a real symmetric matrix to a symmetric tridiagonal matrix.

TREPT
 Computes a parabolic fit, $F = ax^2 + bx + c$, to three points.

TRLFRP
 Stores the force constant matrix in lower triangle form.

TSC
 Locates points where $E = V_a^{AG}$ and calculates the theta integral.

TSOUT
 Computes and writes the final set of conventional transition state theory rate constants. Also obtains activation energies and equilibrium constants.

TSPSI
 TSPACK subroutine that computes a set of parameter values which define a Hermite interpolatory tension spline.

TSRATE
 Calculates the free energy of activation at the saddle point, individual reactant and product partition functions, and conventional transition state theory rate constants in the harmonic approximation. This routine is only used when the NOCVT keyword is used in the RATE section. Some further discussion is provided in Section 6.L of this manual.

TURNPT
 Computes the concave-side turning point for the zero-point level in the calculation of the curvature component.

UPCSE
 Converts a line of characters to uppercase.

UPCAS
 Converts lower case characters to upper case in the given string.

UPDHR
 Updates the arrays FMIHR and FMIHTS in the VTST-IOC calculations.

UPDMI
Updates the arrays FMOM and FMITS in the VTST-IOC calculations.

VALVAG
Calculates the value of V_a^G for a given projected direction, given V, X, DX, and F.

VECCON
Normalizes the normal mode vectors and converts them to mass-scaled Cartesians.

VECMAG
Obtains the magnitudes of a set of vectors.

VICLCG
Calculates the classical energy corrections in the non-adiabatic region for the LCT method in the VTST-IOC calculations.

VIVT
Computes the Eckart potential and its derivatives.

VPART
Computes the vibrational partition function.

VPARTI
Computes the vibrational partition function using the harmonic approximation.

VRCNEJ
Calculate number of available states for transitional modes using variable reaction coordinate algorithm at canonical, microcanonical, and E, J -resolved microcanonical levels.

VRCTST
Main subroutine to calculate reaction rate for loose transition state using variable reaction coordinate and single-faceted or multifaceted dividing surface.

VSPLIN
Generates spline fits to the vibrationally adiabatic potential curve and the effective reduced mass.

VSPLI2
Generates spline fits to the vibrationally adiabatic potential curve and the effective reduced mass for IVTST calculations.

VTMUSN
Calculates the microcanonical variational sum of states in the calculation of the microcanonical variational rate constant.

WTITLE
Writes the title of the run.

WKBENE
Calculates the energy eigenstates with the WKB approximation.

WKBPOT
Computes the energy levels of a quadratic-quartic function using a semiclassical method with parabolic connection formulae.

WKBVIB
Driver for the calculation of WKB vibrational energies based on the true (unfitted) potential for a given mode.

WRTHOK
Writes stationary point information to file fu61 and, if necessary, to file fu62.

WRTWEL
Writes the information about the wells to file fu6.

WVAR6
Writes the keyword values for a restart calculation.

XDOTP
Calculates the dot product of two vectors.

XERBLA
Error handler for the BLAS routines.

XPROD
Forms the cross product of vectors x and y, placing the result in vector z. All vectors are of dimension three.

YDER24
Computes the second derivatives of the potential by fitting the analytic first derivatives to a fourth-order polynomial around the desired location and then taking the derivative of this polynomial. This is the hooks version of DERV24.

YDERV2
Calculates the Hessian matrix by using a quadratic fit to the potential energy surface. Only the potential is used, not its first derivatives. This is the hooks version of DERIV2.

YNEWT
Does a Newton-Raphson search for a stationary point. This is the hooks version of NEWT.

YSECEN
Calculates second derivatives by the method of central differences. This is the hooks version of SECCEN.

YSECEP
Calculates second derivatives by the method of central differences; only computes part of the Hessian matrix. This is the hooks version of SECCEP.

YTRANS

Converts Cartesian coordinates to mass-scaled coordinates or converts mass-scaled coordinates to Cartesian coordinates. Also converts derivatives of the potential energy function between these coordinate systems. This is the hooks version of TRANS.

ZEROPT

Computes the zero-point energy.

ZERP

Function that computes the zero-point energy.

ZIGAMA

Calculates the incomplete Gamma function of order 3/2.

ZOC3P

Prepares the correction information at 3 points for the VTST-IOC calculations.

ZOCAST

Checks the accuracy of the interpolation in the VTST-IOC calculations.

ZOCFRE

Calculates the corrections for the frequencies for VTST-IOC calculations.

ZOCPAR

Calculates the parameters for the VTST-IOC calculations.

ZOCPRN

Prints some of the information involved in VTST-IOC calculations.

ZOCUPD

Updates the arrays WER, FMOM, CDSCMU, VCLAS, WETS, and VADIB and the variables EPRD and WSTAR with zero-order corrections for VTST-IOC calculations.

ZOCVCL

Calculates the correction for the energy VMEP in VTST-IOC calculations.

ZRIDDR

Solves a nonlinear equation for calculating the range parameter of an Eckart potential.

ZROUT

Explicitly initializes the work space for matrix with zeros to overcome the FORTRAN save effect.

ZRPTE

Evaluates the zero-point energies.

ZSPMEP

Sets up information for ISPE. This subroutine is used with unit fu51.

17.A.2 User-supplied subprograms and module files

MODULE files:

aamod.f90

This FORTRAN 90 file contains all the modules used for passing information between the various subprograms. It also contains all the permanent constants for the *POLYRATE* code. These parameters should not be changed.

alloc.f90

This FORTRAN 90 file dynamically allocates memory for arrays according to the user input file.

User-supplied subprograms:

SETUP

Subprogram supplied by the user that computes any necessary constants for the potential (see Section 8 and examples in the test suite).

SURF

Subprogram supplied by the user that computes the potential and gradient for a set of Cartesian coordinates (see Section 8 and examples in the test suite).

17.A.3 FORTRAN files

acespoly.F

This file contains dummy subroutines used with ACESRATE.

energetics.F

This file contains subprograms for the calculations of properties of the PES. These include subprograms that handle optimizations and numerical second derivatives.

ef.F

This file contains the Eigenvector-Following algorithm for optimizing geometries of the stationary points.

fromblas.F

This file contains all the subroutines from the BLAS math library.

givtst.F

This file contains subprograms for the global version of IVTST calculations.

headr.F

This file contains subprograms that print out the program header.

hooks.F
This file contains subprograms for interfacing electronic structure calculations with *POLYRATE*.

intbsv1.F
This file contains subprograms for the transformations of the Cartesian Hessians into a curvilinear coordinate system at non-stationary points along the MEP and for the curvilinear coordinate vibrational analysis for linear and nonlinear triatomics.

intbsv2.F
This file contains subprograms for the transformations of the Cartesian Hessians into a curvilinear coordinate system at non-stationary points along the MEP and for the curvilinear coordinate vibrational analysis for nonlinear polyatomics. The coordinate transformation of the Hessian is based on the transformation ANHTRAX program written by Matt Challacombe and Jerzy Cioslowski.

intbsv3.F
This file contains subprograms for the transformations of Cartesian Hessians into curvilinear coordinate system at reactants and products and for the curvilinear coordinate vibrational analysis for nonlinear polyatomic.

interface.F
This file contains subprograms for parsing *POLYRATE* keyword input.

ivtstm.F
This file contains subprograms for performing IVTST-M interpolation.

main.F
This file contains the driver for polyatomic VTST calculations.

main_vrcmpi.F
This file contains the driver for polyatomic VRC-TST calculations. It is a substitution of main.F file.

mopoly.F
This file contains dummy subroutines used with MORATE.

polyag.F
This file contains subprograms with names beginning with letters A to G.

polyhl.F
This file contains subprograms with names beginning with letters H to L.

polymq.F
This file contains subprograms with names beginning with letters M to Q.

polyrr.F
This file contains subprograms with names beginning with the letter R.

- polysz.F This file contains subprograms with names beginning with letters S to Z.
- poly31.F This file contains subprograms that are used when `POTENTIAL = unit31`.
- poly40.F This file contains subprograms that are used when `POTENTIAL = unit40`.
- rtpjac.F This file contains subprograms that are used to plot the MEP and large curvature representative tunneling termini in mass-scaled Jacobian coordinates.
- vrctst_mpi.F This file contains subprograms that are used to calculate reaction rate for loose transition states using variable reaction coordinates and single-faceted or multifaceted dividing surfaces. These subprograms are parallelized using MPI.

17.B. Utility program detmi.f

In this version of *POLYRATE*, one of the VTST-IOC options [`DETMI` keyword] allows the user to correct the calculated determinants of moment of inertia tensors with accurate data. However, the values of determinants of moment of inertia tensors are not always available in the literature or in the electronic structure calculation output of standard codes. So, in this version of the *POLYRATE* distribution we provide a small utility program, `detmi.f`, in the `util` subdirectory of the *POLYRATE17* directory. This program reads a molecular geometry in Cartesians (in bohrs or in Angstroms) and the atomic weights (in a.m.u.) of the constituent atoms of the molecule, and calculates the determinant of the moment of inertia tensor of the molecule in all the units supported by the *POLYRATE* unit `fu50` input options [`UNITMI` keyword]. (See subsection 12.F) The input formats are very simple, and a detailed explanation can be found in the file `detmi.doc` in the same subdirectory. A job control file `detmi.jc` and three sample input files, `ch4.mi`, `ch4oh.mi`, and `oh.mi` are also provided.

One can use the compiling and linking commands listed in the Section 15 of this manual to make an executable file `detmi.exe` for any machine. This program is easy to use; however, one has to remember to do the following:

1. Make a data file. The program will read the data from a file called detmi.dat. If your data file has a different name, copy, move, or link your file to this name.
2. When finished, the program will generate an output file called detmi.out. It is recommended that you move (or copy) this file to another file name before doing any other calculations. Otherwise the new output will overwrite the old detmi.out file, and the old information in that file will be lost.
3. There is a simple script detmi.jc in the same sub directory. It can do the above file copying steps for you if you want. It takes the input file name as the only argument.

For example, if your data file is called ch4.mi, type

```
detmi.jc ch4.mi <Return>
```

to do the calculation. The output will be written to the file detmi.out and copied to the file ch4.out.

How to make an input data file:

1. In the first line you have to input three numbers in free format.

The first number is the number of atoms in the molecule.

The second number specifies if the geometry of the molecule is linear or nonlinear. Set the number to 1 if it is linear. Set the number to 0 if the geometry is not linear.

The third number specifies the input units for the Cartesians. Set it to 1 if the units you use are Angstroms. Set the number to 0 if you use bohrs.

For example, if the target molecule is methane and your geometry data in Cartesians are in Angstroms, the first line should look like the following:

```
5 0 1
```

2. The second and the third lines are the calculation titles. You can write whatever you want here (less than 80 characters in one line).

For example:

```
Methane: CH4
QCISD/adj-cc-pVTZ geometry
```

3. The next n lines, where n is the first number you entered in the first line, contains the atomic masses and Cartesian coordinate information. There should be 4 numbers on each line. The first number is the atomic mass in a.m.u. of the atom; the second, third, and fourth numbers are the x , y , and z coordinates of the atom.

For example:

1.007825	0.0000	0.0000	0.0000
12.00000	1.0848	0.0000	0.0000
1.007825	1.4464	1.0228	0.0000
1.007825	1.4464	-0.5114	0.8857
1.007825	1.4464	-0.5114	-0.8857

17.C. Utility program hrotor.f

When some of the vibrational modes of the reactants, products, or GTS are treated as hindered rotations, *POLYRATE* allows the user to input more accurate values of the moments of inertia of these modes in a VTST-IOC calculation [IC and HRMI keywords]. However, usually the user can only get the geometry of a molecule and the vibrational eigenvectors of these modes from an electronic structure calculation output.

The utility program hrotor.f is designed to take the atomic masses and geometry of the molecule and the vibrational eigenvector of a hindered-rotor mode of the molecule and then to calculate the moment of inertia using rectilinear model. It can also calculate the harmonic, free rotor, and hindered rotor partition functions at different temperatures. Detailed explanations can be found in the file hrotor.doc in the *Polyrate17/util* directory and in Refs. 24, 29, and 30 in Sect. 23. A sample input file, cwmc.rmi, is also provided in the *Polyrate17/util* directory.

You can use the compiling and linking commands listed in the Section 15 of this *POLYRATE* manual to make an executable file hrotor.exe for your machine. The program is designed to read data from Unix standard input and write data to Unix standard output. To use the program, type

```
hrotor.exe < input_file_name > output_file_name <Return>
```

For example:

```
hrotor.exe < cwmc.rmi > cwmc.rmi.out <Return>
```

To make a data file follow the steps below:
(Note: All the input data are in free format.)

1. In the first line you have to input 4 numbers.

The first number is the number of atoms in the molecule.

The second number specifies the units of the Cartesian coordinates you are going to input. Set it to 0 if you use bohrs, to 1 if you use Angstroms.

The third number specifies the coordinate system of the vibrational eigenvector you are going to input. Set it to 0 if you use a mass-scaled coordinate system; set it to 1 if you use a non-mass-scaled Cartesian coordinate system.

The fourth number specifies the number of temperatures at which you want to calculate the partition function.

For example, if the target molecule is the saddle point of the reaction in the *POLYRATE* cwmctrl test run, your geometry data in Cartesians are in bohrs, the eigenvector is in mass-scaled coordinates, and you want to calculate the partition function at two temperatures, the first line should look like the following:

```
9 0 0 2
```

2. The second and the third lines are the calculation titles. You can write whatever you want here (less than 80 characters in one line).

For example:

```
Saddle point of the reaction in cwmc test run  
The moment of inertia
```

3. If you set the third number in the first line to 1, skip this step. The next line is the scaling mass in a.m.u. of the mass-scaled coordinate system which the input eigenvector is based on.

For example:

```
1.0000
```

4. The next n lines, where n is the first number you entered in the first line, contain the atomic masses of the constituent atoms in a.m.u.

For example:

```
12.00000  
34.96885  
1.007825  
1.007825  
1.007825  
34.96885  
1.007825  
1.007825  
15.99491
```


5. The next n lines, where n is the first number you entered in the first line, contain the geometry data of the molecule in (unscaled) Cartesian coordinates. There should be 3 numbers on each line, namely the x , y , and z coordinates of the atom.

For example:

```

2.0187  -2.3695  0.1399
1.7442  -2.3695  4.4447
4.0466  -2.3695  0.1399
1.0334  -4.1335  0.1399
1.0334  -0.6054  0.1399
1.7442  -2.3695  -4.1650
-2.3564 -2.3695  1.5703
-2.3564 -2.3695  -1.2906
-3.4890 -2.3695  0.1399

```

6. The next n lines, where n is the first number you entered in the first line, contain the vibrational eigenvector of this hindered-rotor mode. There should be 3 numbers on each line, the x , y , and z components of the eigenvector for the atom corresponding to this line.

For example:

```

0.000  0.567  0.000
0.000  -0.260  0.000
0.000  0.533  0.000
0.308  -0.008  0.000
-0.308 -0.008  0.000
0.000  -0.260  0.000
0.000  -0.116  0.000
0.000  -0.116  0.000
0.000  0.207  0.000

```

7. The next line contains the symmetry and subgroup information. The first number is the symmetry factor of this hindered rotation. The second number is the number of atoms in one of the subgroups. The next m numbers, where m is the second number in this line, are the atom indices of those atoms in that subgroup.

For example:

```

3 4 1 3 4 5

```

8. If you decide to calculate the partition function (i.e., if the fourth number in the first line is greater than 0) you need to input the harmonic frequency in cm^{-1} in a new line and the temperatures in Kelvin in the last line.

For example:

```

163.4794
600.00 1000.00

```

17.D. Utility programs findl.f and findb.f

In VTST-IOC calculations one has to input some range parameters for the correction functions, in particular, the range parameter L for the Eckart function and the parameter B for the cut-off Gaussian function on the reactant and/or product side. These parameters are usually obtained by fitting a lower-level potential energy curve to the corresponding functions. The utility programs findl.f and findb.f, which are located in the *Polyrate17/util* subdirectory, are designed to do the fitting and return the optimized values of L and B . The user can use the compiling and linking commands in Section 15 of this *POLYRATE* manual to make executable files (e.g., findl.exe and findb.exe). To use these programs, just type the name of the executable file, and you will be prompted to input the energies of 3 stationary points for findl.exe, or the energies of 2 stationary points and the s value for one of the stationary points for findb.exe. Then, for both programs, you will be prompted to input the energy and s value for another point along the lower-level reaction path. This point is usually chosen to be the point that has an energy halfway between the saddle point and the stationary point on the endothermic side of the MEP for findl.exe or the reactant/product side stationary point for findb.exe. The programs will then print the optimized value of L or B for the Eckart or cut-off Gaussian function, respectively. The user can then use these values for the unit fu50 input [RANGEIC keyword].

17.E. Utility program rotate30.f, rotate31.f, and rotate40.f

ROTATE x ($x = 30, 31, \text{ or } 40$) is a FORTRAN program that uses as an input file a *POLYRATE* poly.fu30, poly.fu31, or poly.fu40 input file and orients the geometries on that file in a consistent way along the reaction path. This program is based on the Chen orientation algorithm Ref. 60 of Section 22.

The poly.fu30, poly.fu31, and poly.fu40 input files for *POLYRATE* are usually constructed from electronic structure calculations carried out by one of the several available packages. These packages usually re-orient the input geometry in a way that allows the program to take advantage of the symmetry of the system, or just to put the molecule in the orientation preferred by the program. As a consequence, the input geometry is oriented in a different way than the output geometry, gradients, and/or Hessians. This results errors in: 1) reaction path following algorithms and 2) computation of reaction

path curvature. If we do not calculate the reaction path or the curvature of the reaction path, the properties that *POLYRATE* uses for the dynamical calculations are rotationally invariant, and the results will not be affected by any rotation of any geometry. However, if we want to calculate the reaction path curvature, we need to calculate the derivatives of the Cartesian components of the gradient with respect to the reaction coordinate s . For doing this, we require that all the points along the reaction path be oriented in the same way. Otherwise, the numerical differentiation will lead to erroneous results.

The *ROTATEX* program uses a geometry that is input by the user (this is assumed to be the saddle point geometry), and it reorients the first point along each side of the reaction path according to this point. The following points along the reaction path are reoriented using as reference geometry the previous point. This reorientation is done by means of the Chen algorithm, as described in the original paper. However, this algorithm only works for small rotations, so before applying Chen's method the program performs a series of trial rotations in order to obtain the closest orientation to the reference geometry, and next it applies Chen's method on the most similar orientation.

Chen's method is based on the minimization of the distance in mass-scaled Cartesian coordinates of the two geometries. In some cases, this minimization does not converge. When that happens, there is no way to solve the reorientation problem with this program.

The program can also recalculate the values of the reaction coordinate, s , as a straight line between each pair of two consecutive points. This recalculation can be useful, e.g., for calculating kinetic isotope effects.

There are three versions of the program *ROTATEX*: *ROTATE30*, *ROTATE31*, and *ROTATE40*, and they are used for reorienting the geometries in *poly.fu30*, *poly.fu31*, and *poly.fu40* input files, respectively. In order to perform this rotation it is necessary to give the geometry of each point. This is sometimes not given on the *poly.fu30* and *poly.fu40* input files. Thus, when we supply additional information on the gradients of one or two points on each side of the Hessian points in *poly.fu30* or *poly.fu40*, the geometry of these one or two additional points per Hessian point is not given. This is the reason why only *poly.fu30* files with $LOPT(6) = 2$ will work with *ROTATE30*, and only *poly.fu40* input files with $GRADER = NOEXTRA$ will work with *ROTATE 40*.

An additional limitation of the program is that it only works with poly.fu30 and poly.fu40 input files that do not include any anharmonic information.

Along with the source code, a test run is included for each program, as well as a sample script for compiling and running the code.

The program requires two input files: (1) one of the *POLYRATE* input files poly.fu30, poly.fu31, or poly.fu40, and (2) a input file for the ROTATE program. The latter input file is described next.

The first line of the input file consists of seven integers that indicate the number of atoms in each system. The first integer is NATOM, the number of atoms in the saddle point structure (which is assumed to be the same as the number of atoms for the points along the reaction path) followed by six integers indicating the number of atoms in the first reactant species, the second reactant, the first product, the second product, the reactants well, and the products well, respectively. If one or more of these species don't exist, a zero has to be input in its place.

The second line is a list of NATOM + 1 real numbers. The first one indicates the mass used for scaling the Cartesian coordinates (which has to be the same as SCALEMASS in the PATH section of the *POLYRATE* poly.fu5 input file. The following NATOM real numbers are the masses of each of the atoms present in the points along the reaction path, in the same order as in in these points. All the masses have to be input in amu.

The third line is an integer that indicates if the values of *s* read from the poly.fu30, poly.fu31, or poly.fu40 input files have to be kept unchanged or if the user wants to put in their places the values computed by the program. If this integer is set to one, the values of *s* that are read are discarded, and the calculated values are printed in the output file. Any other value leads the program to preserve the *s* values that are input in the poly.fu30, poly.fu31, or poly.fu40 input file.

The last information to be input in the ROTATE input file is the geometry in Cartesian coordinates of the saddle point in atomic units. The three Cartesian coordinates for each atom have to be preceded by an integer, which acts as a label. The ordering of the atoms has to be the same as in the points along the reaction path, and the labels have to start with the integer 1 for the first atom and finish with the integer NATOM for the last atom.

The output of the program, poly.fu60, is a file with the format same as poly.fu30, poly.fu31, or poly.fu40, which can be used in a *POLYRATE* calculation directly without further modification.

17.F. Utility program *SS-QRRK*

In the utility program directory *SS-QRRK*, it contains one .exe file, one source code file (.f90) and one directory for test runs and one directory for outputs of test runs. *SS-QRRK* code is a FORTRAN90 code for carrying out system-specific quantum RRK calculations. Please refer to a separate manual for the use of *SS-QRRK* code.

18. UNIX C SHELL SCRIPTS AND PERL SCRIPTS

POLYRATE includes many useful scripts for users. The following subsections discuss the functions of these scripts and explain how to use them.

18.A. Installation script

This script, *configure*, resides in the *Polyrate17* directory. It is used to install the *POLYRATE* program on your computer. Note that this installation is only for generating the Makefile, and no executable is generated in this process. Usually, the first thing you do after uncompressing and untarring the distribution tar file is to run this script. (See Subsection 15.A.) Actually, this script does four things:

1. Makes a hidden file `.poly_path` in your home directory.

The hidden file contains the path to the *Polyrate17* directory in the file system you are using. This lets other scripts in *POLYRATE* easily locate the necessary files. If you lose the hidden file for some reason, or if the file system changes after you install *POLYRATE*, you need to run this script again to generate a new `.poly_path` file.

2. Asks the user to choose which kind of executable the user wants the Makefile to generate: RP-VTST or VRC-VTST. (If you want both, you must run the script twice.)

3. Detects compilers, operating system, and processor type

The script will automatically search for the FORTRAN and C compilers that are correct for the configuration being used.

4. Copies machine-dependent files and puts them into the proper directories with the default file names.

It copies the appropriate date-and-time routine from the *Polyrate17/util* directory to the *Polyrate17/src* directory and names it `dattim.f`.

If VRC-VTST installation is selected by the user, the script will install the SPRNG 2.0 library. If RP-VTST installation is chosen, the script will skip the SPRNG installation step.

5. Creates the Makefile

The Makefile is used to compile and link all of the code needed for a job. If any changes are made to the files used to create the executable, the Makefile will automatically update the executable.

See section 15.A.1 for a description of the directory structure following a successful installation.

18.B. Potential energy surface compiling scripts

This script is distributed in the current version, but it is no longer needed when using Makefile.

There are two potential energy surface subroutine compiling scripts in the *Polyrate17/script* directory. The script `comp_pot.jc` is used to compile all the potential energy surface subroutines for the test runs. (See Subsection 15.A.2.)

The script `potcomp.jc` is designed to compile a single file of potential energy surface subroutines. It is general, and you can use it to compile your own potential subroutines. This script takes two arguments; the first one is the potential file name without extension, and the second one is the number of atoms involved in the potential. For example, suppose you have a set of FORTRAN potential energy subroutines which defines the potential energy surface of a system that consists of one oxygen atom and 3 hydrogen atoms, and the set of subroutines has the file name `myoh3.f`. Put this file in the *Polyrate17/poten* directory. You then change to the *Polyrate17/script* directory and type `potcomp.jc myoh3 4 <Return>`; the script will compile the file `myoh3.f` for you and put the object module `myoh3.o` in the *Polyrate17/obj* directory. You can, if you wish, copy the script into the *Polyrate17/poten* directory so you can use it more conveniently. Remember that you must put your potential source code in the *Polyrate17/poten* directory. It should be noted that since the `nh3` potential energy surface subroutine provided in the distribution package has already been written the way that is required by *POLYRATE*, when compiling this potential energy subroutine with the script `potcomp.jc`, you should use a number that is not 3 or 4 as the second argument. This is because the `nh3.f` subprogram already contains the necessary `SETUP`, `SURF`, `COORD`, and `CHAIN` subroutines.

18.C. Source code compiling script

This script is distributed in current, but it is no longer needed when using Makefile.

The script `comp_src.jc` in the *script/* directory is used for compiling the *POLYRATE* source code. It takes one argument, the parameter `INCLUDE` file number. You probably used this script when you did the testing. (See Subsection 15.A.2.) Usually, after you compile the

source code with a certain parameter INCLUDE file and keep the library file that the script generated, you don't need to recompile the source code unless you need to use a different parameter INCLUDE file. For example, suppose you write your own INCLUDE file named param5.inc, and you want to compile the source code with this INCLUDE file. Put the param5.inc file in the src/ directory, change to the script/ directory, and type comp_src.jc 5 & <Return>. The script will compile the source code with the parameter INCLUDE file param5.inc and generate a library file named libpoly5.a in the obj/ directory. You can then use the linking script described in the next subsection to link this library file with your potential object module to make an executable file. Remember that you must put the parameter INCLUDE file in the src/ directory and always name a new parameter file param#.inc where # can be any integer greater than 4.

18.D. Linking script

This script is distributed in the current version, but it is no longer needed when using Makefile.

The script link_exe.jc in the *Polyrate17/script* directory is used for linking the potential object file with the source code library file to make an executable file. It takes 2 arguments. The first one is the library file number, which is just the parameter INCLUDE file number you used when making the source code library file. (See the Subsection above.) The second argument is the potential file name (without the extension) you want to use. For example, if you want to use the source code library compiled with the parameter file param5.inc and the potential object module myoh3.o to make an executable file, change your working directory to the script/ directory and type link_exe.jc 5 myoh3 <Return>. The script will link the source code library and potential object files together and generate the executable file poly_myoh3.exe and a short summary file poly_myoh3.doc about this new executable file in the exe/ directory.

18.E. Test run scripts

The test run scripts have been discussed in Subsection 15.A.2. These scripts are designed such that you don't need to know much about *POLYRATE* in order to run the test runs. Some of the test runs might take a long time to run (see Section 21) so you should consider running them in background or submitting them as batch jobs. Some test runs

generate very big output files that you may want to delete together with the test run executable files when you need to free up some disk space.

19. FILE USAGE

The *POLYRATE* program uses several files for input data, for storing restart information (optional), and for output. The program explicitly opens and closes all files required for a given run, and the files are assumed to have the file names *poly.fu#*, where # is an integer. The file *poly.fu#* connects to a FORTRAN I/O unit *fu#* which is defined in the file *aamod.f90* (see Subsection 5.A). This means that at the beginning of any run, all the *POLYRATE* input data files which have logical file names differing from *poly.fu#* must be linked or copied to the corresponding *poly.fu#* file. If the user wishes to assign different logical names to the output files, which are also *poly.fu#* files, these reassignments must be carried out after the calculation has completed. Any data file(s) needed by the potential energy surface subroutine is (are) not opened by the *POLYRATE* program and therefore, do not automatically follow the *poly.fu#* format.

One would ordinarily expect FORTRAN unit 5 to be assigned to *fu5*, FORTRAN unit 6 to *fu6*, and so forth. The *fu#* system, however, allows one to avoid conflicts when hooking *POLYRATE* up to other programs, e.g., MOPAC or analytic potentials.

<u>Unit</u>	<u>Usage</u>
fu1	Reaction-path information for restarting the program
fu2	Additional reaction-path information to be merged with that from unit fu1
fu3	Output for merged reaction path information
fu4	Reserved for potential energy surface data (Note: A potential subroutine may read data from more than one file, if desired. In the test suite we always use unit 4 if data is read from only one file, and we use units 4 and 7 for the cases where data is read from two files)
fu5	General input data. This data has several sections—see section 12.A of this manual for the details.
fu6	Output (This is the full output file. Selected subsets of this output are also written to files <i>poly.fu14</i> and <i>poly.fu15</i>)
fu7	Usually reserved for additional potential energy surface data (See Subsections 8.C and 16.B for further discussion)
fu14	Output table of dynamical bottleneck properties
fu15	Output table of selected forward rate constants

fu16	Reserved for output from electronic structure packages
fu22	Detailed output for the LCT calculations [LCTOPT: PRFREQ, PRPROB]
fu24	Output file of MEP and large curvature representative tunneling termini in mass-scaled Jacobian coordinates.
fu25	Output file of the potential energy information along the MEP
fu26	Output file of the generalized transition state frequency information along the MEP
fu27	Generalized transition state geometry information along the MEP in the XYZ input format of the XMOL program
fu28	Bond angles, bond distances, and dihedral angles along the MEP
fu29	Electronic structure input file for IVTST-0 and IVTST-1 calculations
fu30	Formatted electronic structure input file used if POTENTIAL = <i>unit 30</i> in the ENERGETICS Section of file fu5
fu31	Keyword-based electronic structure input file used if POTENTIAL = <i>unit 31</i> in the ENERGETICS section of file fu5.
fu40	Keyword-based electronic structure input file used if POTENTIAL = <i>unit 40</i> in the ENERGETICS section of file fu5
fu41–fu47	Output tables of LCT calculations [LCTDETAIL]
fu48, fu49	Output files for LCT restart calculations
fu50	Input file for higher-level data for VTST-IOC calculations [DL keyword set to <i>ioc</i> or ICOPT]
fu51	Input file for higher-level data for VTST-ISPE calculations [DL keyword set to <i>ispe</i>]
fu60	Reserved for the direct dynamics intermediate file
fu61	Reserved for information about stationary points that can be used to restart a calculation by the second restart method of Section 7. C.
fu62	Reserved for a subset of information that are presented in fu61 if the argument of the WRITE62 keyword is not set to <i>none</i>
fu65	Generalized transition state geometry information along the MEP in the XYZ input format of the MDMOVIE program

fu70	Reserved for input data needed by PREP
fu71	Reserved for input data needed by PREPJ when JTYPE = 1 (REACT1)
fu72	Reserved for input data needed by PREPJ when JTYPE = 2 (REACT2)
fu73	Reserved for input data needed by PREPJ when JTYPE = 3 (PROD1)
fu74	Reserved for input data needed by PREPJ when JTYPE = 4 (PROD2)
fu75	Reserved for input data needed by PREPJ when JTYPE = 5 (START)
fu76	Reserved for input data needed by PREPJ when JTYPE = 6 (PATH)
fu77	Reserved for input data needed by PREPJ when JTYPE = 7 (WELLR)
fu78	Reserved for input data needed by PREPJ when JTYPE = 8 (WELLP)
fu80-fu89	Reserved for electronic structure codes and interfaces

20. MEMORY REQUIREMENTS

This section is no longer valid starting from version 2016 because version 2016 dynamically allocate memory and the size of memory is really determined by the system size in the job. But we keep this section here so that users can know roughly how much memory is required for the test runs in some old machines. In general, memory requirement is not a concern for a modern computer.

Memory (in Megabytes) required for the *POLYRATE* test suite executable files on all the machines on which the code was tested. The parameter INCLUDE file used is param4.inc for the ch5, cwmc, and ch4o test runs, param3.inc for the dumpot test runs, and param2.inc for all other test runs. The memory requirements were determined using version 2008 and the Unix SIZE command.

Test run	dumpot*	ch4o	ch5	clhbr	cmc	cwmc
IBM Power4	28.4	42.9	42.9	20.8	20.9	43.0
SGI Altix	30.0	44.5	44.5	22.3	22.4	44.6

Test run	h2ni	h3	hni	ho2	nh3	oh3
IBM Power4	20.9	20.8	27.9	20.8	20.8	20.8
SGI Altix	22.5	22.4	29.5	22.4	22.4	22.4

Test run	ch3
SGI Altix	369.7

* The dummy potential is used for the fu30, fu31, and fu40 test runs.

Note in version 2010, the number of atoms is set to 30 ((NATOMS = 30) in param.inc and param4.inc so that users don't need change this parameter for most of cases. For this setting, all the test run executables require 206.4 MB memory except that hni needs 213 MB memory and ch3 needs 677.5 MB memory.

21. TEST RUNS AND TEST RUN TIMINGS

21.A. Extent of testing

The full test suite of version 2017 was run on the HP Linux clusters with two quad-core 2.8 GHz Intel Xeon X5560 and two quad-core 2.5 GHz Intel Xeon E5-2680 processors (CentOS Linux).

21.B. Timings

The following timing trials for *POLYRATE* may be of interest. The running time of *POLYRATE* varies quite a bit, depending mainly on the number of atoms, the complexity of the potential, the range of reaction coordinate examined, step sizes, and options for anharmonicity and tunneling. The rest of this section gives some examples of timings. The times (system time + user time) are in CPU seconds.

1) ch4cl test run

	JN FIRE	SGI Altix
ch4cltr1	0.0	0.0
ch4cltr2	0.0	0.0
Total	0.0	0.0

2) ch4o testrun

	JN FIRE	SGI Altix
ch4otr1	2.4	1.3

3) ch4oh test runs

	JN FIRE	SGI Altix
ch4ohtr1)	0.0
ch4ohtr2)	0.0
ch4ohtr3)	0.0
ch4ohtr4	l	0.0
Total	l	0.0

4) ch5 test runs

	JN FIRE	SGI Altix
ch5fu29tr1	0.0	0.0
ch5fu29tr2	0.0	0.0
ch5fu30tr1	0.0	0.0
ch5fu30tr2	0.0	0.0
ch5fu30tr3	0.0	0.0
ch5fu30tr4	0.0	0.0
ch5fu30tr5	0.7	0.7
ch5fu30tr6	0.4	0.3
ch5fu31tr1	1.0	0.9
ch5fu40tr1	0.0	0.0
ch5fu40tr2	0.0	0.0
ch5fu40tr3	0.0	0.0
ch5fu40tr4	0.0	0.0
ch5fu40tr5	0.7	0.7
ch5fu40tr6	0.3	0.2
ch5fu50tr1	0.0	0.0
ch5fu50tr2	0.7	0.6
ch5fu50tr3	0.6	0.6
ch5fu51tr1	0.3	0.3
ch5icfu30tr1	1.5	1.5
ch5icfu30tr2	1.5	1.5
ch5icfu31tr1	2.0	1.6
ch5fcfu40tr1	1.7	1.5
ch5icfu40tr2	1.6	1.5
ch5j1tr1	0.6	0.6
ch5j1tr2	0.3	0.3
ch5j1tr3	0.7	0.8
ch5j1tr4	0.6	0.5
ch5j1tr5	0.6	0.5
ch5j1tr6	0.7	0.6
ch5j1tr7	0.8	0.8
ch5j1tr8	0.1	0.0
ch5j1tr9	0.5	0.4
ch5j1tr10	0.7	0.6
ch5j1tr11	0.4	0.3
ch5j1tr12	0.6	0.5
ch5j2itr1	0.0	0.0
ch5j2itr2	0.0	0.0
ch5j2tr1	0.2	0.1
ch5j2tr2	0.2	0.1
ch5j2tr3	0.2	0.1
Total	20.2	18.1

5) clhbr test runs

	JN FIRE	SGI Altix
clhbrtr1	0.6	0.6
clhbrtr2	0.6	0.6
clhbrtr3	0.6	0.6
clhbrtr4	0.6	0.6
clhbrtr5	0.6	0.6
Total	3.0	3.0

6) cmc test runs

	JN FIRE	SGI Altix
cmctr1	1.4	0.7
cmctr2	0.9	0.4
cmctr3	1.5	0.7
Total	3.8	1.8

7) cwmc test runs

	JN FIRE	SGI Altix
cwmctr1	0.0	0.0
cwmctr2	3.5	2.3
cwmctr3	3.4	2.2
Total	6.9	4.5

8) h2ni test run

	JN FIRE	SGI Altix
h2nitr1	1.9	1.9

9) h3 test runs

	JN FIRE	SGI Altix
h3tr1	0.6	0.2
h3tr2	0.3	0.2
h3tr3	0.2	0.1
h3tr4	0.3	0.2
Total	1.4	0.7

10) hni test run

	SUN FIRE	SGI Altix
hnitr1	4.8	4.3

11) ho2 test runs

	SUN FIRE	SGI Altix
ho2tr1	0.4	0.3
ho2tr2	0.3	0.3
ho2tr3	1.5	1.0
ho2tr4	1.4	0.6
ho2tr5	0.9	0.4
Total	3.5	2.6

12) h3o2 test runs

	JN FIRE	SGI Altix
h3o2fu31tr1	0.0	0.0
h3o2fu31tr1	0.0	0.0
h3o2fu31tr1	0.0	0.0
h3o2fu31tr1	0.0	0.0
Total	0.0	0.0

13) nh3 test run

	JN FIRE	SGI Altix
nh3tr1	0.1	0.1
nh3tr2	0.1	0.1
Total	0.2	0.2

14) oh3 test runs

	JN FIRE	SGI Altix
oh3fu30tr1	0.5	0.5
oh3fu30tr2	0.1	0.0
oh3fu30tr3	0.1	0.0
oh3fu30tr4	0.0	0.0
oh3fu30tr5	0.0	0.0
oh3fu30tr6	0.1	0.1
oh3fu31tr1	0.5	0.3
oh3fu31tr2	1.4	1.1
oh3fu40tr1	0.9	0.5
oh3fu40tr2	0.0	0.0

oh3fu40tr3	0.0	0.0
oh3fu40tr4	0.0	0.0
oh3fu40tr5	0.0	0.0
oh3fu40tr6	0.2	0.1
oh3tr1	1.8	1.8
oh3tr2	0.1	0.1
oh3tr3	0.2	0.2
oh3tr4	0.2	0.2
oh3tr5	0.1	0.1
oh3tr6	0.0	0.0
oh3tr7	0.0	0.0
oh3tr8	0.1	0.1
oh3tr9	0.2	0.2
oh3tr10	0.0	0.0
<hr/>		
Total	6.5	5.3

15) ohcl test runs

	JN FIRE	SGI Altix
ohcltr1	0.3	0.2
ohcltr2	0.9	0.8
ohcltr3	0.9	0.8
ohcltr4	0.9	0.8
ohcltr5	1.9	2.1
<hr/>		
Total	4.9	4.7

16) ch3 test run (MPI run using 4 cores)

	JN FIRE	SGI Altix
ch3tr1	66.5	61.3
<hr/>		

22. CITATIONS INDICATED BY BRACKETS IN THE TEXT

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Note : See Section 22 for citations indicated by brackets indicated in the text.

The following is a list of some publications reporting calculations carried out with *POLYRATE* or with its original version, which was entitled ICVTST. This includes all articles in which new procedures incorporated in *POLYRATE* are described, but it does not include all applications.

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24. ERRATA

All known errata to the GTST book chapter (Ref. 5 in Sect. 23) have been assembled into a PDF file. This file is available at the *Polyrate* Web site.

25. APPENDIX: *Polyrate* REVISION HISTORY

25.A. Computer program revision history

This section contains the revision history subsequent to version 1.1 (August 1987), which was the first distributed version. First and second level revisions (e.g., 2.0 or 2.1) denote enhancements; third level revisions (e.g., 1.5.1) denote bug fixes.

Version 1.5 (September 1988)

1. Projection operators suitable for reactions on solid surfaces are included as well as those for gas-phase nonlinear and linear complexes. An option for solid-state and gas-solid-interface reactions is added employing these projection operators. This can also be used for embedded cluster calculations.
2. Variable dimensions are used, removing the previous restriction to 10 or fewer atoms.
3. An extra option, $LGS(13) = 2$, for unit 5 input is added.
4. An option, $LGS(3) = 3$, is added to allow calculation of the MEP without calculating frequencies of generalized transition states.
5. The less useful anharmonicity options are removed. All methods except harmonic, Morse I/II/III, quadratic-quartic, and WKB are removed.
6. The Newton-Raphson method is improved to use a variable search step determined by the Brent method as discussed in the section "Reactant and product properties" in this ASCII manual.
7. The methods for calculating the MEP are updated. All previous methods except Euler and ES1 are removed. The Page-McIver method is added.
8. The output format is improved.
9. A list of all known errata and typographical errors in the GTST book chapter is included in the manual.

Version 1.5.1 (December 1988) (Manual fix)

1. Test runs W and X were rerun with smaller DELSV to get converged results.

Version 1.5.2 (February 1989) (Bug fixes)

1. In subprogram NOROUT:

```
if (... pmode.eq.false)
```


is non-standard FORTRAN and was changed to

if (... .not.pmode).

2. In subprogram PATH:
When the option of scaling V_{MEP} is chosen, the vibrational frequencies of the reactant and product which correspond to the reaction coordinate are also scaled appropriately by a factor of \sqrt{VFAC} . If VFAC is not zero or one, the indices of the reactant and product vibrational frequencies being scaled are read on unit 5.
3. In subprogram RATE:
For gas-phase reactions which involve only one reactant or product, the relative translational partition function for the forward or reverse rate, respectively, is not calculated unless an external field is present. This affects the reverse rate constants of test runs P-R.
4. In subprogram REACT:
IF(LGS(4) .EQ. 4) is corrected to IF(LGS(6) .EQ. 4). This affects the free format reading of the variables ICODE and IATOM in the case of a unimolecular reaction.
5. Test run W is removed because unstable results are caused by the nature of the potential energy surface.
6. In subprogram OVERLP:
Frequencies and normal mode eigenvectors are also switched when two modes switch. (Note this will introduce a small difference in the SCSAG transmission probability compared to previous results for a given set of numerical parameters, but both algorithms should converge to the same result when the step size is decreased.)
7. In function DGAMMA:
An error that affected the operation of GAUSSQ when it was called for generalized Gauss-Laguerre quadrature with $\langle > 2$ or for Jacobi quadrature with $\langle > 2$, $\textcircled{R} > 2$, or $\langle +\textcircled{R} > 1$ has been corrected, even though *Polyrate* does not require these cases.
8. In subprogram RESTOR:
Errors in reading information about MEP #2 and calculating the reverse rate constants when LGS(8) = 3 have been fixed.

Version 1.6 (July 1989)

1. Two subprograms, FIOPEN and FICLOSE, were added to open and close all files. These were added to make the program more portable, in particular to run under the CTSS operating system. (Note: these statements are commented out in distribution versions from 1.6 to 3.9.3, and they are reorganized and supported in versions 4.0 and later)
2. A 'call link' statement has been added to MAIN to make the program usable on a Cray computer running CTSS. Users of different machines will need to add a stub subprogram called link that just returns when called.
3. The Newton-Raphson search convergence criterion has been modified in subprogram NEWT to search until the norm of the gradient is small. In previous versions the search was terminated when changes in the energy from iteration to iteration became small.

4. The total vibrational partition functions Q_v are calculated in terms of $\ln(Q_v)$ to avoid underflow in medium-size systems, an option has been introduced to allow Q_v functions to be multiplied by a factor of $10^{\text{LGS}(37)}$ to avoid underflow in large systems.
5. Option $\text{LGS}(36)$ has been added to compute the geometry of the next point on the MEP from the electronic structure input data.
6. The reduced mass for mass-scaled coordinates is read in record 5, and the program now calculates relative translational reduced masses for bimolecular reactions internally.

Version 1.6.1 (November 1989) (Bug fixes)

1. In subprogram CENTER:
For a polyatomic molecule with a nonlinear geometry, the new version correctly prints out FMOM, which is the product of the three principal moments of inertia, whereas versions 1.1-1.6 incorrectly printed out the determinant of the moment of inertia tensor without the factor of μ^3 in eq.(47) of the GTST book chapter. This printout error did not affect the rate calculations.
2. In subprogram NORMOD:
For the Page-McIver reaction-path method, unnormalized derivatives of the potential are stored in the array DXP as for the Euler method. This corrected a printout error in versions 1.1-1.6.
3. In subprogram OPTION:
The print statement reflecting the action carried out when $\text{LGS}(17) \neq 0$ has been changed to say that the reverse rate constant will be multiplied by $10^{\text{LGS}(17)}$. This required deleting a minus sign in FORMAT 1340. This also corrected a printout error in versions 1.1-1.6.
4. The manual has been corrected to include subprogram NEXTPT which was added in version 1.6. Furthermore, this subprogram has been corrected so that the geometry of the next point on the MEP can be calculated correctly from RPH input data. In addition, a bug for option $\text{LGS}(36) \neq 0$ is fixed in subprogram RPHRD2, where the generalized normal mode analysis is now performed only if $\text{LGS}(36) = 0$.
5. In the evaluation of turning points for a symmetric adiabatic barrier, the tunneling energies should be selected on a grid from the threshold energy up to the maximum of the adiabatic barrier. The adiabatic barrier maximum obtained by spline fit is always greater than or equal to the largest value of the adiabatic potential on the Hessian grid. Thus the barrier maximum for determining the grid of tunneling energies should be obtained from the fit, and not just set to the largest value on the save grid. Versions 1.1-1.6 used the maximum value from the save grid, but version 1.6.1 uses the value from the fit. This required changes in four subprograms: KAPVA, PSAG, PSATP, and PSATP2.
6. In subprogram SPL1D1:
A non-standard FORTRAN statement involving a transfer into a DO loop is corrected by rearranging the subprogram.

7. The *Polyrate* manual has been corrected and improved. Especially important corrections occur on pages 16, 21, 24, and 25. In addition, descriptions of subprograms INTFNC, PITZ, NEXTPT, and STABLE have been added to the section "Description of FORTRAN subprograms".

Version 1.6.2 (February 1990) (Bug fix)

1. In subprogram OVERLP:
An error introduced in version 1.5.2 in the way that frequencies and eigenvectors are switched at crossings and/or avoided crossings of vibrationally adiabatic potential curves has been corrected so that the effective mass for an SCSAG tunneling calculation is evaluated correctly.

Version 1.6.3 (March 1990) (Bug fixes)

1. In subprogram RATE:
The size of array QVSV, which is only used in the subprogram RATE for vibrational partition functions of reactants and products, is increased from $N3TM = 3 \cdot NATOM$ to $N6TM = 6 \cdot NATOM$, since $QVSV(N3TM)$ may not be sufficiently large for some reactions.
2. In subprogram PATH:
The format of the input data for the variable NST0, which is the number of steps of size DEL1 to be taken in following the reaction path away from the saddle point, was incorrect from versions 1.5 to 1.6.2, and is correct in version 1.6.3. (Note: the format was correct in version 1.1.)

Version 1.7 (March 1990)

1. All input information read from unit 5, except the routing and output options, is now read in and checked by two new subprograms READ5 and REDGEO at the beginning of the calculation. Previously the input was read in many different subprograms and at various stages of the calculation. A new input option, LGS2(1) = 1, is used to request that the program read the input from unit 5, write it on unit 6, and then stop. This is useful for checking input data interactively before submitting a job to a batch queue. This consolidation of input involved changes in subprograms MAIN, DIATOM, FINOUT, PATH, POLYAT, RATE, RESTOR, and SADDLE.
2. Another new option, LGS(29) = 2, is now added for calculating the reaction-path curvature by the method of Page and McIver from the gradient and the Hessian and its eigenvectors. This option involved changes in subprograms BCALC, INTPM, and OPTION. Another test run Z is added to show the use of this option.
3. The normalization factor of the gradient along the reaction path is computed and stored in variable DXNORM of the IRCCM common block which has been rearranged in subprograms INTFNC, INTPM, IRCX, NEXTPT, and NORMOD.

Version 1.7.1 (April-July 1990) (Bug fixes)

1. In subprogram PATH:
The conversion factor for energy, CKCAL, has been corrected so that it is consistent with the value used elsewhere in the code.

2. In versions 1.5-1.7, the frequencies written on the short output file for the reaction-bottleneck properties were truncated to integers rather than rounded in the subprogram FINOUT, and the option, LGS(9) = 0, for calculation of the rate constants without tunneling calculations, did not work because of the undefined array BETA. Both bugs are corrected in version 2.0.

Version 2.0 (July 1990)

1. A new type of semiclassical tunneling calculation is added, in particular the large curvature ground state method, version 3 (LCG3), which is especially appropriate for reactions with large curvature of the reaction path. The implementation of the LCG3 method involved adding 22 new subprograms and functions as well as modifications in 17 other subprograms of POLY1 and 6 subprograms of POLY2. Notice that the LCG3 calculation is available only for the harmonic approximation and for ground-state products as well as ground-state reactants in version 2.0.
2. A new option, LGS2(2) = 1, is now added for explicitly projecting out the translational and rotational motions at stationary points. In some cases this results in more accurate values for nonzero frequencies. This option involved changes in the subprograms: OPTION, CENTER, NORMOD, POLYAT, and SADDLE.
3. An algorithm is added for finding turning points when the tunneling energy is greater than the largest value of the adiabatic potential on the save grid, but less than the maximum of the spline fit. This is accomplished by interpolation between the maximum of the spline fit and the largest values on the grids using the cubic polynomial corresponding to that section of the spline fit. Previous versions did not handle this situation correctly, although the quadrature grid was seldom dense enough for the problem to occur. Installing the new algorithm involved changes in the subprograms: PSAG, PSATP, PSATP2, BOLTZ, FINOUT, FITMAX, VTMUSN, and FIVPT.
4. A new type of input has been introduced for the step sizes of the Hessian and save grids. This enhancement involved changes in the subprograms: INTPM, PATH, READ5, RESTOR, and NEXTPT. See the section on "Reaction-path following methods" in this manual.
5. For version 2.0, the two subprograms INTERP and LINMIN have been renamed INTRPL and LINMN, respectively, so that the interface with MOPAC can be made without subprogram name conflicts.
6. Two new test runs have been added, and all the previous test runs have been modified to test the new enhancements. A new, more convenient convention was introduced for naming test runs. The two new test runs, ch5tr2 and h3tr2, involve LCG3 calculations for two different reactions; the former one uses the restart input file option, and the latter computes the reaction path information from an analytic potential energy surface. The LCG3 option has also been turned on for test runs ch5tr1 and h2nitr1, and test runs oh3tr3 and ch5tr4 have the new stationary-point projection option turned on. In the majority of the test runs, the Hessian and save grid step sizes are now calculated from input values of INH and INM, which specify them as integer multiples of the gradient grid step size. The gradient step and Hessian step size for most test runs have been increased to be closer to optimum choices for 1% accuracy.

7. For version 2.0, all test runs are verified on four different computers, and the system-dependent subprograms, in particular, DATTIM.VMX or DATTIM.UNIX, have been moved to separate files from the rest of the source code. The code is now distributed with sample command files for both VMS and UNICOS.
8. Various changes are made to allow the convenient creation of MORATE from *Polyrate*.

Version 2.0.1 (September 1990) (Bug fix)

1. In subprogram REDGEO:
An error introduced in version 1.7 for reading input for a unimolecular reaction has been corrected so that the information is stored in the correct variable.

Version 2.1 (October 1990)

1. In order to make the code more portable, all statements containing non-ANSI FORTRAN 77 that are flagged by the cft77 compiler on the Cray-2 have been modified to conform to the FORTRAN 77 standard. Also, all common blocks have been made to have the same length every time they occur, and subprogram FICLOSE has been renamed FCLOSE. The code is written to execute in double precision so it runs with REAL*8 floating point variables on VAX, Sun, Silicon Graphics, and IBM computers. On the Cray, REAL*8 is single precision so the code should be compiled with the disable precision (-dp) option.
2. All common blocks and statements used only in MORATE have been commented out with C*M in columns 1-3 so that the memory and execution time of *Polyrate* itself are as small as possible.
3. A symmetric reaction path option, LGS(3) < 0, has been added, and a related test run, h3tr3, has been provided for this option. For the current version, this option is available for calculations with all transmission coefficients except LCG3.
4. A warning was added to the manual about using LCG3 with only ground-state final states. Restricting the final state of the tunneling process to the ground state is not always appropriate.
5. A large number of typos were corrected in the manual, and various clarifications were added.

Version 2.1.1 (November 1990) (Bug fix)

1. In previous versions, the frequencies written on unit fu6 were truncated to integers rather than rounded in subprograms PATH and RESTOR. This bug is fixed in version 2.1.1.

Version 2.5 (November 1990)

1. The main changes in version 2.5 are all related to portability. A single version of *Polyrate* now runs correctly on all eight computers mentioned in the section labeled "computers and operating systems on which the code has been checked". The full 20-run test suite executes successfully on all eight computers. All non-ANSI usage that was uncovered in the course of these tests has now been converted

to standard FORTRAN 77 and all non-portable code has been made portable. The modifications include the declarations of external functions DIGAMMA and ERF in subprograms CLASS and ZIGAMA respectively, a correction in subprogram PSATP2 to avoid possible floating point errors, and the correction of all statements containing non-ANSI FORTRAN 77 in the analytic potential energy surfaces used for the 20 test runs. For example, the subprogram TRIPLET in CH5POT.FOR, which is used in the seven test runs ch5tr1 to ch5tr7, has been renamed TRIPLE, and the variables NCOUNTA and NCOUNTB in H2NIPOT.FOR which is used in test run h2nitr1 have been renamed NCOUNA and NCOUNB, respectively.

2. A large number of precision changes were made to many subprograms to make sure that the code is executed completely in 64-bit precision. The five non-Cray machines on which the code has been tested have 32-bit words, and explicit use of double precision typing ensures REAL*8 precision on these machines. On the Cray computers, the word length is 64 bits, so single precision corresponds to REAL*8. Thus on these machines we use the disable precision (-dp) compiler option so the code again runs with REAL*8 precision. But the non-standard REAL*4 and REAL*8 type declarations do not actually appear in the program.
3. The Boltzmann constant has been replaced with the more accurate value $3.166830 \cdot 10^{-6}$ a.u. in subprogram RATE. Thus, for some test runs the rate constants might be slightly different from the ones in previous versions. The differences should be less than 0.2%.
4. The code is now distributed with sample command files suitable for its execution on all eight computers on which it has been tested. Actually only seven sets of command files are used since the same set of command files can be used for the Cray-2 and Cray X-MP.

Version 2.5.1 (February 1991) (Bug fix)

1. A correction has been made in the manual only. The correction concerns allowed values of other parameters that may be used with the electronic structure input option, LGS(30). The combination of LGS(30) = 1 and LGS(6) = 0 is allowed; however LGS(30) = 1 must be used when LGS(36) = 1. The statement just below LGS(30) = 1 on page 59 in the manual of version 2.5 is corrected.

Version 2.5.2 (March 1991) (Bug fixes)

1. In subprogram BCALC:
A correction has been made for the storage of the turning points of generalized normal modes with imaginary frequencies for LCG3 calculations. In versions 2.0-2.5 these were set to zero. Now they are set more appropriately to infinity, i.e., a very large value (10^{10} bohr). Note that this modification only affects LCG3 calculations, not SCSAG calculations. (For SCSAG calculations the turning points of modes with imaginary frequencies are set equal to zero.)
2. In subprogram IRCX:
The Euler stabilization algorithm now performs correctly, even when the stabilization step is omitted. This correction supplies a previously missing statement, calculating the reaction coordinate s , when the stabilization procedure is skipped. This error existed in versions 1.0-2.5.

3. In subprogram NEXTPT:
An extra READ statement, which was not deleted when all READ statements were moved in version 1.7 and which caused a fatal error by reading beyond the end-of-file mark whenever the option for the calculation of the geometry of the next point on the MEP from electronic structure input data was selected [LGS(36) = 1], was removed.
4. In subprogram OPTION:
An error allowing the user to select options LGS(30) ≠ 1 and LGS(36) = 1 simultaneously was removed. (If LGS(36) = 1, then the only allowed value of LGS(30) is 1.) This error existed in versions 1.6-2.5.
5. In subprogram REDGEO:
An error in the printout of the initial geometry for the optimization of the saddle point has been corrected. This error existed in versions 1.7 - 2.5.
6. A correction has been made to an error introduced in the manual at the time version 1.7 was prepared. The manual stated that the normalization factor of the gradient along the reaction path is computed and stored in variable DXNORM of the IRCCM common block, but actually it is correctly stored in DXMAG.
7. A correction has been made in the manual concerning the input format for variable NST0, the number of steps of size DEL1 to be taken in following the reaction path away from the saddle point, when LGS(13) = 1. Before, it was given in the manual as (.../,20X,I5), but the program actually reads it as (.../,30X,I5). This error existed in the manual since version 1.5.
8. Two subprograms, PITZ and STABLE, that are no longer used, have been removed, and common block MINCOM has been adjusted to have the same length in all subprograms.
9. In function VPART:
The treatment of imaginary frequency generalized normal modes is made consistent with the procedures described in the section "Imaginary frequency modes" in this manual.

Version 2.6 (March 1991)

1. A new method for calculating the initial step from the saddle point with analytical potential energy surfaces is now included. This method bases the first step on a cubic expansion of the potential energy surface, instead of a quadratic one. The new method can be selected through option LGS(31) and is responsible for changes in subprograms INTEUL, INTPM, OPTION, PATH, and READ5.
2. The Page-McIver integration method is now able to use NST0 ↑ 0 in the unit 5 input, which allows it to take smaller steps than the pre-selected gradient step size in the vicinity of the saddle point. This enhancement involved changes in subprograms INTPM and PATH.
3. The user can now select the stabilization criterion for the ES1 integration method; this is equal to the vector difference between successive normalized gradients. The old value, 10⁻⁸, remains as the default. This improvement involved changes in subprograms IRCX, PATH, and READ5.

4. The calculation of the reaction-path curvature by the method of Page and McIver from the gradient and the Hessian and its eigenvectors—option LGS(29) = 2—has been optimized to avoid unnecessary storage. This optimization involved changes in subprograms BCALC, INTFNC, INTPM, IRCX, NORMOD, and PATH.
5. In subprogram OPTION:
Three STOP statements were replaced with LEXIT = .TRUE. statements, to allow the code to flag all incorrectly selected options before stopping. The STOP statements were introduced in version 2.0.
6. The test runs oh3tr1 and oh3tr3 were modified to incorporate the new enhancements. In particular, test run oh3tr1 now uses the Page-McIver method with the cubic starting algorithm and performs five steps of size DEL1 in the vicinity of the saddle point before taking steps of size DEL. Furthermore, test run oh3tr3 now also uses the cubic starting algorithm, and it uses the Euler stabilization method along the reaction path with a bigger stabilization criterion, equal to 10^{-3} . Differences in rate constant results between the old and new test runs in both cases do not exceed 1%.

Version 2.6.1 (June 1991) (Bug fixes)

1. In subprogram EXTRAP:
Minor modifications have been made in order to incorporate changes in subprogram READ5 related to the extrapolation procedure. This corrects wrong statements which existed since version 1.0.
2. In program MAIN:
In versions 1.0-2.6, the energy and gradient were not computed correctly at the starting point when no saddle point calculation was performed. This is now corrected.
3. Also in program MAIN:
Another change has been made in MAIN to correct an error in restart runs when *Polyrate* is used in conjunction with MOPAC as MORATE. This error existed since version 1.6.
4. In subprograms LCNWTN and LCG3:
A bug in the root search to determine the unique s value corresponding to a point along the LCG3 tunneling path has been eliminated. As explained in pages 105-106 of the GTST book chapter, the value of s should be a continuous function of the progress along the tunneling path. In versions 2.0-2.6 of *Polyrate*, the root search was coded incorrectly to find the smallest (i.e., most negative or least positive) root on the reactant side and the greatest root on the product side. This is corrected in the new subprograms so that the iterations start with the value of s for the previous point on the tunneling path. Thus the iterations should converge to the continuous root.
5. In subprogram OPTION:
In versions 1.0-2.6, the statements printed about whether the first step away from the saddle point towards the positive s -direction (products) is along the positive or negative direction of the imaginary frequency eigenvector were incorrect; in particular, FORMAT statements 1110 and 1111 were interchanged. This is now corrected.

6. Also in subprogram OPTION:
A second correction prevents the user from selecting an ICVT, α VT, and/or unified statistical (US) rate constant calculation while also choosing no MEP calculation (LGS(3) = 0) or choosing only an MEP calculation (LGS(3) = 3). This change corrects an error which existed since version 1.0.
7. In subprogram PATH:
Extra statements which reversed the sign of SOB and SOE (the initial and final s values in bohr between which extra vibrational mode and Q_{vib} information is printed) when ISEN is positive (initial step from the saddle point is in the direction of products) have been eliminated. This corrects a bug which existed since version 1.5.
8. Also in subprogram PATH:
An extra statement was added in order to save the imaginary frequency eigenvector when subprogram SSAVE is called for the first time. This corrects a bug which existed since version 1.5.
9. In subprogram READ5:
More statements were introduced to prevent the user from selecting incompatible input for LGS(12), ISEN, and IWHAT for a particular run. This error existed since version 1.0.
10. Also in subprogram READ5:
The output values of SLMG and SLPG for NGSPEC = -1 were wrong in version 1.0-2.6 but are correct now.
11. In subprogram REDGEO:
Minor modifications were introduced to allow the user to choose anharmonicity option LGS(5) = 21 at the saddle point in MORATE runs (in which *Polyrate* is interfaced to MOPAC). This corrects a bug which existed since version 1.6.

Version 2.7 (June 1991)

The user now has the option of calculating conventional transition state theory (TST) rate constants without performing an MEP calculation and without being required to carry out canonical variational transition state (CVT) rate constants. This enhancement can be selected by setting LGS(7) = -1, and it required changes in program MAIN and subprograms OPTION, READ5, and RATE, as well as two new subroutines, TSOUT and TSRATE.

1. In subprogram OPTION:
Seven STOP statements were replaced with LEXIT = .TRUE. statements, to allow the code to flag all incorrectly selected options before stopping. The STOP statements were introduced in version 2.0.
2. The user will now receive a warning whenever inappropriate values are selected for NTEMP and NPAIR. This required modifications in subprogram READ5.

Version 3.0 (July 1991)

1. The large-curvature tunneling method for semiclassical ground-state transmission coefficients, version 3 (LCG3) is now fully implemented for analytic potential energy surfaces in the harmonic approximation. In particular the contributions of all accessible states populated by the tunneling process in the LCG3 exit arrangement (see "Polyrate 4.0: ...", *Comp. Phys. Comm.* **71** (1992) 235) can be evaluated, i.e., tunneling into excited states may now be calculated. The new options are specified by $LGS(9) = 3$ or $LGS(9) = -n$ with $n \geq 2$; $LGS(9) = 3$ causes the program to include all accessible final states, whereas the $LGS(9) = -2, -3, \dots$ allows the user to specify the highest excited state to be included in terms of a maximum quantum number for a quasideiabatic mode. Five subprograms for LCG3 calculations, PROB, SINTGL, XINTGL, XTAU, and VMODE, have been renamed LCPROB, LCSINT, LCXINT, LCXTAU, and LCVMOD, respectively. Three new subprograms, LCG3TH, LCSTX, and LCTP, have been added, and more than twelve subprograms have been modified in conjunction with this enhancement of the LCG3 calculation.
2. A new option, $LGS2(3) = 1$, is now added that allows a user to omit the contributions of low-frequency modes to μ_{eff} when SCSAG tunneling calculations are performed. For this option, the user can provide a frequency ω_0 in units of cm^{-1} , and all modes with frequencies less than ω_0 will be neglected during the calculation of μ_{eff} .
3. Three new test runs have been added. Two new test runs, ch5tr8 and ch5tr9, were introduced to show LCG3 calculations with tunneling into all accessible final states [$LGS(9) = 3$] and to show the omission of curvature from low-frequency modes in SCSAG calculations [$LGS2(3) = 1$]. The third test run, involving Cl + HBr, shows an LCG3 calculation with the option, $LGS(9) = -3$, of a user-specified maximum quantum number for the quasideiabatic mode.
4. For version 3.0, unit 16 is reserved for use by MOPAC so that output previously written to unit 16 now has been assigned to unit 14. This change involves only subprograms FINOUT and FCLOSE.
5. The maximum number of points in the save grid along the MEP has been increased to 1500 in this distributed version.

Version 3.0.1 (August 1991) (Bug fixes)

1. In the subprogram READ5:
A non-standard statement introduced in version 2.7 "IF (LGS(7).LT.0) THEN NGSPEC = 0" is changed to "IF (LGS(7) .LT. 0) NGSPEC = 0".
2. In the subprogram REDGEO:
When $LGS(13) = 2$ is selected for the input format, the index for the coordinates is not stored correctly. It is fixed by changing "INDX(IOP,I) = I" to "INDX(I,IOP) = I". This error exists in versions 2.0-2.7.

Version 3.0.7 (September 1991) (Bug fixes)

1. Not all the FORTRAN statements involving unit 16 were changed to unit 14 in version 3.0. This inconsistency has been fixed and it involved changes in the

subroutine FIOPEN.

2. Subprogram LCG3TH is modified. When only one quadratic point on a tunneling path is in the nonadiabatic region, the effective adiabatic potential for that point is now taken as the average value of the adiabatic potential at the ending grid points of the adiabatic region on the reactant and the product sides.
3. In the subprogram LCPROJ, a ratio of "0/0" could occur when the large-curvature tunneling path is almost parallel to the reaction path, and causes an infinite looping in the subprogram LCSTX. This bug is now fixed.
4. Subprogram PSATP2 was modified so that at energies at or above the maximum of the adiabatic potential on the save grid but less than the maximum of the spline fit to the vibrational adiabatic potential curve, the turning points (both left and right) are set to the value of s at the maximum. This correction only affects LCG3 calculations involving tunneling into excited states.
5. The adiabatic potential and the turning points along each generalized normal mode, which are needed for calculating the contribution of tunneling into excited states in the large-curvature tunneling approximation, were stored only at save points greater than STP0. They are now saved at all the save points along the minimum energy path. This correction involves subprogram LCSTX.
6. The criterion for determining that a point is within the globally multi-valued region is modified in the subprogram LCG3TH.
7. A bug is fixed in subprogram LCG3, where a Kronecker delta needed for calculating the primitive LCG3 tunneling probability was missing.
8. Redundant statements are removed from MAIN.
9. For reactions without a saddle point, a generalized normal mode analysis is now done on the projected force constant matrix for the first point on the reaction path. The first point was previously treated incorrectly. Note that the output files for test runs hootr1, hootr2, and hootr3 are also modified due to this change.
10. For calculations using the cubic initial step option, the stored classical energy at the saddle point was incorrect. This involves corrections in the subprogram PATH. This modification corrects the calculated conventional transition state rate constants of test runs oh3tr1 and oh3tr3.
11. The variable IMAX was used both as an index for the maximum of the adiabatic potential, VS(IMAX), and as an index for the spline segment where VMAX occurs. This error had caused turning points to be set incorrectly to zero in some cases. This involved changes in the subprograms KAPVA, PSAG, PSATP, PSATP2 and VSPLIN.
12. An unnecessary tolerance check in the subprogram FDIAG which could, under some circumstances, lead to an infinite loop has been removed. Also, one of the looping structures in this subprogram has been modified to avoid unnecessary checking.
13. The tolerance criterion in the subprograms VPART, WKBPOT, and HQSC has been improved. This change does not affect the rate constants within the level of accuracy reported, but it does make the iterative calculation more robust.
14. The number of excited states included in an LCG3 calculation:

The maximum excited state into which the p mode can be excited is limited by the parameter MAXPS defined in the file aamod.f90. The program is modified such that if the number of excited states to be included is not specified (i.e., LGS(9) = 3) then the number of states to be included is at most MAXPS even though there may be more accessible states. If the number of states specified by the user is larger than MAXPS and the number of accessible states are also larger than MAXPS, a warning message will be printed and the calculation is carried out with MAXPS states.

15. A bug in the LCG3 calculations including tunneling into excited states was corrected. In versions 3.0 and 3.0.1, the uniformization is carried out only for $n_{\max} = n_p^{\text{Max}}$ rather than for all values from 0 to n_p^{Max} or to the largest value allowed by the dimensions (MAXPS).

Version 3.9 (September 1991)

1. A new small-curvature tunneling approximation, called the centrifugal-dominant small-curvature adiabatic ground-state tunneling approximation, the CD-SCSAG method, is implemented. Two new subprograms are added, and 15 others are modified. The section in the output files with the final results has been restructured accordingly. This involved changes in the subprogram FINOUT. We now distribute short output files of the sample runs using the CD-SCSAG method instead of the original SCSAG method, since the CD-SCSAG method is the one we recommend for the small-curvature cases. The default is now CD-SCSAG.

Version 3.9.1 (September 1991) (Bug fixes)

1. In subprogram VSPLIN, a missing RETURN is now added.
2. Subprogram ANHARM is modified to avoid losing accuracy due to multiple transformations between mass-scaled Cartesian and Cartesian coordinates.
3. The option of optimizing certain Cartesian components to search for a stationary point was not available when LGS2(2) = 1. This problem caused failures in searching for stationary points of certain systems. It is now fixed by modifying subprogram POLYAT.

Version 3.9.2 (November 1991) (Bug fix)

1. Subprogram FDIAG is modified to always give eigenvectors of $3N$ components, whether the species being analyzed is a reactant, a product, or a generalized transition state, where N is the number of atoms in the reaction. Due to this change, subprograms ANHDER, NOROUT, SADDLE, and SGND3V are also modified.

Version 3.9.5 (March 1992) (Bug fixes)

1. Subroutine READ5 is modified to always read in DEMIN when Morse anharmonicity or a \propto VT calculation is requested.
2. The dimensions of certain arrays in subprograms AITKEN, AITKF2, AITKNF, KAPVA, LOCS, SL1B1, SL1B2, and VSPLIN are modified to match the sizes in the corresponding subprograms which pass the arrays through call statements to these

subprograms. Among the machines on which the *Polyrate* program has been tested, a discrepancy due to this bug occurs only on the Sun SparcStation 4/360 (SunOs 4.0.3).

3. A gradient point was previously determined to be a save point if it is less than a gradient step away from an end point specified by the user. This criterion is now changed to be less than half a gradient step to avoid possible discrepancies caused by machine precision. The correction is made in `PATH`.
4. In subprogram `OVERLP`, the array `INDEX`, which stores the indices of the generalized normal mode eigenvectors, was initialized to zero. To avoid the error of trying to retrieve the zeroth array element, this array is now initialized as `INDEX(I) = I`.
5. In subprogram `READ5`:
For `LGS(35) ≠ 0` (i.e., a `MORATE` calculation), `LGS(25)` and `LGS(33)` are set equal to zero, as these options are not currently supported for a calculation that is not based on an analytic potential energy surface.
6. In subprogram `RESTOR`:
A typographical error introduced in version 1.6 has been corrected. This entailed changing the statement "`READ(2, *) (XER(2), I = 1, NTOT2)`" to "`READ(2, *) (XER2(I), I = 1, NTOT2)`". This bug fix will affect any calculations which used the option to merge reaction paths from units 1 and 2 and also involved anharmonicity for the generalized normal modes.
7. In subprogram `LCSET`:
A typographical error in the call statements for the subroutine `INTRPL` has been corrected. This involved modifying the statement

```
CALL INTRPL (XETS, SSUBI, ANHARM, . . . )
```

to

```
CALL INTRPL (XETS, SSUBI, ANHRM, . . . )
```

There were two occurrences of this problem in this subprogram. This bug fix will not affect any currently supported options because it relates to the anharmonicity parameters and it occurs in an `LCG3` subprogram, but anharmonicity is not yet supported with `LCG3` tunneling.
8. In subprogram `LCG3TH`:
The call statement for the subprogram `INTRPL` has been corrected:

```
CALL INTRPL (XETS, SSUBI, ANHARM, . . . )
```

has been changed to

```
CALL INTRPL (XETS, SSUBI, ANHRM, . . . )
```

The `CALL` statements for the subroutine `LCVOR` incorrectly used the variable `XE` in the argument list instead of the array `ANHRM`; this has also been corrected. These changes have no effect on any currently supported options because they affect the anharmonicity parameters in a large-curvature tunneling subprogram, and anharmonicity is not yet supported with `LCG3` tunneling.
9. `FORMAT` statements that were incompatible with `NATOMS` larger than 6 and which would cause output conversion errors for the `cwmctr1` and `cwmctr2` test runs were fixed. This involved changes in the following subroutines: `CENTER`, `NEWT`, `POLYAT`, `SADDLE`, and `REACT`. Also, `FORMAT` statements 2300 and 2330 in `RESTOR` have been corrected; these errors existed since version 3.0.7.
10. Spelling and grammatical errors were corrected in `FORMAT` statements in the `POLYRATE` subroutines `HEADER` and `OPTION`, and in the potential routine `cwmc.f`.

Version 4.0 (March 1992)

1. All the common blocks are byte aligned. All the `PARAMETER` statements and common blocks that are used throughout the code are now placed in three `INCLUDE` files, `param.inc` for the adjustable parameters, `percon.inc` for the fixed parameters, and `common.inc` for the common blocks. This modification makes redimensioning arrays much more convenient. Also, to maintain the code's portability, non-ANSI standard FORTRAN 77 statements were corrected in the following subroutines: `NEWT`, `NEXTPT`, `RATE`, `LCG3TH`, and `LCVMOD`.
2. The dimensions of the arrays for storing the information along the minimum energy path are now controlled by two parameters, `NSDM` and `NSDML`. This modification allows users to reduce the required memory when `LCG3` calculations are not requested.
3. A new method for approximating the partition functions of selected low-frequency vibrational modes corresponding to hindered internal rotation is implemented. This is called the hindered-internal-rotator approximation.
4. A new option, `LCG2(4)`, has been implemented. This flag allows the user to specify a nonzero value of `EZERO`, which is defined such that $V - EZERO = 0$ at the reactants. If a restart file is written to unit `fu1` or read from unit `fu1` then a value for `EZERO` is also written/read. Because of this enhancement old restart data files will not work with this version of the code.
5. All `STOP` statements in all subprograms other than `MAIN` have been changed from `STOP` to `STOP 'subprogram name #'` where `#` is 1,2, etc. sequentially for each subprogram.
6. All `STOP` statements in the subprograms `READ5`, `REDGEO`, and `OPTION` which flagged incorrect input were changed to `LEXIT = .TRUE.` so that the code would check the entire input deck before quitting.
7. All input and output I/O units for the *Polyrate* code are now adjustable. They can easily be changed by modifying the variables `fu#` (e.g., `fu6` for unit 6) defined in the `INCLUDE` file `param.inc`.
8. All the input and output files used by *Polyrate*, except for the files linked to FORTRAN units `fu5` and `fu6`, are explicitly opened and closed in the subprograms `FIOPEN` and `FCLOSE`, respectively. The files for units `fu5` and `fu6` are opened in the subprogram `MAIN`. The default file name is `poly.fu#`, which is linked to FORTRAN I/O unit `fu#` defined in the `INCLUDE` file `param.inc`. For example, if `fu15` is set to unit 15 in `param.inc`, the `poly.fu15` file will be opened and linked to unit 15.
9. The information for restart calculations is now written with formatted statements rather than with free formatting for certain long records. This modification enhances the portability of the *Polyrate* code.
10. The treatment of diatomics has been made more flexible and now allows them to be treated as general linear molecules; because of this, the input data files of all the test runs that involve diatomic molecules are modified such that diatomic molecules are treated in the same manner as linear polyatomic molecules. Before, diatomic molecules were treated as Morse oscillators. This Morse oscillator option is still available and can give accurate results with less effort if the diatomic molecules are indeed Morse oscillators. Due to this change, the `fu30` file, `oh3rph.dat`, for the test

run oh3tr6 is also modified.

11. The test runs are reorganized according to the potential functions. Four sample runs are added, and one is modified: cwmctr1 is an example of calculating the conventional transition state rate constants only; cwmctr2 uses the hindered-internal-rotator approximation; clhbrtr1 is modified to use the LGS2(4) = 1 option, i.e., using *Polyrate* to calculate EZER0; clhbrtr2 is added to demonstrate the option of reading EZER0 in the fu5 input; and, finally, ch5j1tr6 is added to show that LCG3 calculations that include tunneling into excited states can be carried out using the restart information from a run that includes only LCG3 tunneling into the ground state.
12. The manual no longer includes the fu5 input and the fu14 and fu15 short output files for the sample runs. These input and output files are instead included in the *Polyrate* 4.0 distribution package. The advantage of this is that users can make input data files by modifying the sample input files, and the comparison to the sample run outputs can easily be done by using the "difference" or "diff" command.
13. Since the number of accessible states as a function of reaction coordinate, $N^{\mu VT}(s,E)$, is not a continuous function of s , the 3-point and 5-point interpolations to find the maximum, as used in previous versions, may be unstable. Thus we now also print out the maximum value on the grid. This may be used for a "one-point" interpolation, which may be more reliable.
14. A new subprogram, INITZE, has been added to the code. This subprogram initializes variables and arrays used by the program. This subprogram currently does not initialize all the arrays and the variables used by the program.
15. The variable IOT has been changed. In version 4.0 it is a flag used by the LCG3 subprograms to indicate whether or not state-labeled primitive tunneling probabilities should be printed to unit fu22. Note: in versions 3.0.7-3.9.5, the printing of this extra information was controlled by LGS2(5).
16. A new option, LGS2(5), which allows the user to print extra information from the potential energy surface routine to the *Polyrate* long output, unit fu6, has been added. The number of pieces of information which can be printed is controlled by the values of the input variable LGS2(5) and the variable NPOTPT which is set in the INCLUDE file param.inc.

Version 4.0.1 (March 1992) (Bug fixes)

1. In subprogram TSRATE, a line was added to allow for the correct calculation of anharmonicity when $LGS(5) \geq 21$ and $LGS(7) < 0$. Versions 2.7-4.0 gave correct rate constants and equilibrium constants only for the first temperature for this combination of LGS options.
2. FORMAT statements in various subprograms affecting the *Polyrate* output were corrected. This does not affect any results of the calculation.
3. Minor modifications were made to allow the user to control the printing of the bond orders and charges calculated in MORATE. This modification affected the following subprograms: MAIN, PATH, POLYAT, RESTOR, and SADDLE.

4. In subprogram MAIN:
The statement
`IF (LGS(1).EQ.0 . . .)`
has been changed to
`IF (LGS(30).LE.0.AND.LGS(35) = 0)`
This modification removes the redundant check for $LGS(1)=0$. This cleanup of the code is not a bug fix and does not affect any of the test runs.

Version 4.1 (April 1992)

1. In subprogram OVERLP:
Prior to version 4.1, when an LCT calculation was requested, the program re-ordered the generalized normal modes and related quantities, e.g., eigenvectors and eigenvalues, at the current save grid point so that the eigenvector of each mode at the current save grid point had the highest overlap with the eigenvector of the same mode at the previous point. If two modes at the current point each had their highest overlap with the same mode from the previous point, the program would print a message on the unit fu6 file and stop. This was modified in version 4.1. In version 4.1, once a mode at the previous point has been matched to a mode at the current point, it is no longer tested against further modes at the current point. Thus the conflict never occurs.

Version 4.1.1 (April 1992) (Bug fixes)

1. In subprogram REACT:
A bug in the computation of the zero of energy for a bimolecular reaction has been corrected. This bug was introduced in version 4.0.
2. In subprogram VTMUSN:
A call to subprogram TREPT, which computes a parabolic fit of free energy curves, contained one extra argument. This bug fix affects μ VT rate constant calculations using a 3-point primary fit, which is selected by setting IFTMV1 to 3 in the input file on unit fu21. This bug was introduced in version 4.0. It does not affect the results of the test runs.
3. Certain non-ANSI statements in files dattim.sun, dattim.ibm, and dateclock.c have been corrected to ANSI-standard. Non-ANSI statements in subprogram MAIN and potential surface function files hni.f, h2ni.f, and cwmc.f are also corrected.
4. A bug has been corrected in OVERLP. In previous versions, the switching of the eigenvalues and eigenvectors was sometimes incorrect, especially for very low frequency vibrations. The new version corrects the error, and in addition it should be more robust because the matching is now carried out in order of decreasing frequency of the vibrational frequencies.

Version 4.2 (May 1992)

1. Subprogram FDIAG has been restructured. In this version of *Polyrate* the generalized normal modes are now ordered according to the absolute values of their eigenvalues. In addition, for a save point along the minimum energy path the matching of the generalized normal modes with those at the previous save point is carried out between the vibrational normal modes only. That is, the modes corresponding to the translational, rotational, and reaction coordinate degrees of

freedom, which are assumed to be the modes with the smallest absolute eigenvalues, are not used in the matching process. This also involved changes in OVERLP.

2. In subprogram OVERLP:
The IF statement 'IF (NN.EQ.0) GO TO 200' has been modified to be 'IF (NN.LE.1) GO TO 200'.
3. FORMAT statements in several subroutines have been modified to make the output more clear. In particular, the output for the reaction energetics has been improved. The output has also been modified so that most of the information written to unit fu6 is lower case.
4. In subprogram INITZE:
A check for the value of LGS(4) when LGS(35) = 0 has been coded; this was necessary because in *Polyrate* LGS(4) is allowed to have values of 0, 1, and 2, but in *MORATE* LGS(4) can also have a value of 3.
5. Test run oh3tr1 is modified to test the option LGS2(4) = 1. This option allows users to calculate the zero of energy by *Polyrate*.

Version 4.2.1 (June 1992)

1. In the INCLUDE file common.inc:
The size of array LRP, which is only used in the subprogram RATE for calculating vibrational partition functions of reactants and products, is increased from N3TM = 3*NATOM to N6TM = 6*NATOM, since LRP(N3TM) may not be sufficiently large for some reactions.
2. In subprogram INITZE:
The array LRP, which is used to indicate whether or not a particular vibrational mode is thermal or not, is initialized. All modes are initialized to be thermal, as this is the case unless LGS(23) \uparrow 0.

Version 4.3 (June 1992)

1. In subprogram OPTION:
A check for the value of LGS(3) when $|LGS(9)| > 1$ has been coded. This check was added because the LCT calculation is not supported when LGS(3) < 0.
2. In subprogram PATH:
The calculation of the MEP and properties at points on the MEP will now be stopped if the number of save points reaches parameter NSDM, which defines the size of arrays for storing MEP information.

Version 4.3.1 (July 1992) (Bug fixes)

1. In subprogram LCPROB:
In versions 3.0-4.3, the LCT primitive probabilities were uniformized with respect to the primitive probability at the highest energy grid point whereas the correct procedure is to use the value at the maximum energy of the adiabatic ground state potential. This is now corrected.
2. In subprogram LCG3:

In versions 3.0-4.3, the total number of accessible states in the LCT calculation was set equal to the largest value of n_p that satisfies eq. (11) in *Comp. Phys. Comm.* **71** (1992) 235, at the highest energy grid point. The calculation of the maximum value of n_p is now handled correctly, as discussed in *Comp. Phys. Comm.* **75** (1993) 143.

3. In subprograms LCG3TH, LCSTX, LCVCOR, and LCXTAU:
In versions 3.0-4.3, when quantities derived from the vibrational normal mode frequencies were calculated, the contribution from imaginary normal modes were included incorrectly. This is now corrected.
4. In subprogram LCG3TH:
For a given tunneling path, the program determines the boundary of the adiabatic region, which consists of two points, $\{a$ (or x_{ia}) and $\{b$ (or x_{ib}). When $\{a$ (or x_{ia}) is greater than $\{b$ (or x_{ib}), the integrand at $\{b$ (or x_{ib}) was included twice in the calculation of the theta integral. This is now corrected.
5. In subprogram LCG3:
In versions 3.0-4.3, the numerical integration of $T_1^{LCG3}(E, n_p)$ [eq. (35) in *Comp. Phys. Comm.* **71** (1992) 235] was incorrect when $n_p > 0$ and the backward reaction is exoergic. The upper bound of the range of integration should be the reaction coordinate turning point calculated from the adiabatic excited-state potential. However this information was not stored and the turning point calculated from the adiabatic ground-state potential was used. This is now corrected.
6. In subprogram READ5:
An IF statement for checking the conflicts between the option for tunneling and the option for anharmonicity was not written properly. Due to this problem, users were not able to start a calculation if a mode was to be treated as a harmonic oscillator when $LGS(5) \geq 21$ and an LCG3 calculation was requested in the same run. This bug existed in versions 4.0-4.3.
7. In subprogram RESTOR:
Several WRITE statements were mistakenly coded as READ statements. This caused I/O conflicts when information from two restart files was to be merged into a new restart file. This bug existed in versions 2.0-4.3.

Version 4.4 (August 1992)

1. One of the criteria for determining whether a point on an LCG3 tunneling path is within the adiabatic region is to check if that point is within the turning points of each of the generalized normal modes. In subprogram LCZCRT, the generalized normal mode vibrational coordinate for each mode was compared to both turning points of that mode. The checking procedure is now made more efficient by using eqs. (30a) and (30b) of *Comp. Phys. Comm.* **71** (1992) 235, in which the absolute value of the turning point of each generalized normal mode is used in the comparison. Also the algorithm for calculating the vibrational turning points is improved in subprograms LCG3TH and LCSTX. Prior to this version, vibrational turning points were calculated from the frequencies at each save grid point and then the vibrational turning points at various s values were obtained by interpolation from this grid. Starting with version 4.4, only the vibrational frequencies are interpolated, and vibrational turning points are always calculated from the

interpolated frequencies. The two algorithms yield very similar results when the generalized normal mode frequencies are much greater than zero, but the new algorithm is more stable when any frequencies are small (near zero) and noisy. In addition, subprogram LCG3TH has been restructured. This modification reduces unnecessary calculation.

2. The LCT method is now generalized to cases involving a well on the product side of the MEP in the exoergic direction. The program first checks if the adiabatic ground-state potential at the last save grid point on the product side of the MEP (s_+) is less than the value of the product ground-state energy. If so, a new point, called the well boundary or s_{WB} , is identified as the farthest out s value for which $V_a^G(s)$ is greater than or equal to the product ground-state energy. Then the maximum number of accessible states is determined as discussed in *Comp. Phys. Comm.* **75** (1993) 143, using s_{WB} instead of s_+ . The contribution from tunneling paths with termini beyond the well boundary is not included.
3. In the calculation of the LCG3 tunneling amplitude, the theta integral, the sine of the angle between the tunneling path and the MEP, the vibrational period, and the excited state potential are obtained by interpolation at NGS0 points between each pair of adjacent energy quadrature points. Prior to version 4.4, 4-point interpolation was used to calculate these quantities. A new option that allows users to specify the order of interpolation is added in version 4.4 because sometimes a more stable, lower-order interpolation is needed for good accuracy. The default is 4-point interpolation and the desired order is set equal to LGS2(9). The input data files of test runs clhbrtr1 and clhbrtr2 are modified to use 2-point interpolation. Details of these two test runs can be found in the section on sample test runs. This involves modifications in subprogram LCSINT.
4. An option for extra output containing details of LCT calculations has been added. In particular, the extra output includes the theta integrand and the corresponding s value on the MEP at each point along the tunneling path, the boundary of each adiabatic region on the tunneling path, the length of the tunneling path, and the classical potential and the correction terms to the potential for points in the non-adiabatic region. This involves modifications in subprograms FIOPEN, FCLOSE, INITZE, LCG3, LCG3TH, LCZCRT, READ5. The C-shell scripts poly.incr and poly.exec are also modified to accommodate the change in the file usage. The command files for the VMS operating system remain the same since none of the test runs uses this option.
5. In the input data files ch5j1tr1.dat, ch5j1tr2.dat, ch5j1tr5.dat, ch5j1tr6.dat, clhbrtr1.dat, clhbrtr2.dat, h2nitr1.dat, and h3tr2.dat:
The numbers of quadrature points used in the calculation of tunneling probabilities, as specified by NQE, NSEG, NQTH, NSEG2, NG, and NGS0, are decreased to reduce the computation time while still converging the transmission coefficients within 1%. This is possible because the current algorithm is more stable due to modifications 1 and 3.
6. A new subprogram NSTATE has been added. This subprogram is used to calculate the total number of accessible states and to determine if an excited state is open for a given tunneling energy. This modification reduces redundant calculations.

7. In subprogram LCNWTN:
The call to subprogram INTRPL to interpolate the gradient vector is modified. The parameter N3TM is used in the argument of the call instead of the variable NDIMX. This is to clean up the program (not a bug fix).
8. In INCLUDE file percon.inc and subprograms BOLTZ, KAPVA, and LCG3:
The dimension of arrays related to the calculation of tunneling probabilities was declared as 243 in previous versions. It is now set to the parameter NSV, and NSV is set equal to 245 in the INCLUDE file percon.inc. The arrays that were modified are PE, PEC, PED, ESV, PLCG3, TRNPT1, and TRNPT2.
9. In versions 4.0-4.3.1, variables N3 and LSAVE were passed to subprograms LCSET and LCSTX as arguments in the CALL statements while they were also passed through common blocks. The program is now cleaned up such that these variables are now only passed through common blocks. Subprogram LCG3 is also modified due to this change.
10. In subprograms ANHARM, FITMAX, LCG3, LCVIB, MUBAR, PSAG, PSATP2, QQPOT:
Prior to version 4.4, certain array elements in these subprograms were passed to other subprograms as arguments in the CALL statements. However, they were not declared as one-dimensional arrays in the subprograms that are called. In version 4.4, dummy variables are set equal to these array elements, and then the dummy variables are passed to other subprograms as arguments in the CALL statements.
11. In subprogram READ8:
In versions 4.0-4.3.1, the variable NATOM and the array elements LGS(13) and LGS(23) were passed to READ8 as arguments in the CALL statements while they were also passed through common blocks. In addition, the array elements were not declared as one-dimensional arrays in READ8. In version 4.4, NATOM, LGS(13), and LGS(23) are no longer passed in the CALL statements. Due to this change, subprogram READ5 and MAIN are also modified.
12. In subprogram SGND3V:
The variable NEND that was passed to SGND3V as an argument in the CALL statement was not used in SGND3V. It is now removed. Subprogram ANHARM is also modified due to this change.
13. In subprogram TURNPT:
The variable NMODE that was passed to TURNPT as an argument in the CALL statement was not used in TURNPT. It is now removed. Subprogram MUBAR is also modified due to this change.
14. Subprogram FIOPEN has been restructured. With this modification, the names of any files that cause errors as they are opened or attempted to be opened will be given in the poly.fu6 file.
15. In subprogram READ5:
A check for the values read from unit fu5 for the temperatures has been coded. This check is only carried out if NTEMP is greater than 0, and it stops the calculations if any of the temperature values are not greater than zero.

Version 4.5 (September 1992)

1. In subprograms `PATH` and `READ5`:
In previous versions, when the cubic starting algorithm was requested, the third derivative of the potential with respect to the coordinates was evaluated using a step size equal to `DLX`, which is the step size for numerical second derivatives. In version 4.5, this step size is set to `D3LX` specified in the `poly.fu5` input file. The `INCLUDE` file `common.inc` is also modified due to this change.
2. In `INCLUDE` files `poly9995.inc`, `poly9911.inc`, `poly9111.inc`, `poly6775.inc`, `poly6711.inc`, and `common.inc`:
The array `SSPEC`, which contains the special save grid points (see records 18 and 19 for `fu5` input) is now dimensioned to `NMSPEC`, which is specified in the `INCLUDE` file `param.inc`. The subprogram `READ5` is also modified so that a warning message will be written to the `fu6` output if the user-input value of `NSPEC` is greater than `NMSPEC`.
3. In subprogram `MAIN`:
The statement “`STOP 'MORATE MAIN 1'`” is modified to “`STOP 'MAIN 1'`”. Also several commented statements for opening files `morate.fu5` and `morate.fu6` are removed. This is to facilitate the procedure of interfacing *Polyrate* with electronic structure packages.
4. In subprograms `MAIN`, `COLSHF`, `MEPSRT`, `PATH`, `POLYAT`, `REACT`, `RESTOR`, `RESTRT`, `SADDLE`, and `SSAVE`:
The statement “`include 'morate.inc'`” is modified to “`include 'esp.inc'`”. This is to facilitate the procedure of interfacing *Polyrate* with a variety of electronic structure packages.
5. In subprograms `FIOPEN`, `FCLOSE`, `MAIN`, `POLYAT`, `REACT`, and `SADDLE`:
The statements to open the files `esp.fu31—fu35` are added in `FIOPEN`. These files correspond to FORTRAN units `fu31—fu35` which are specified in the `INCLUDE` file `param.inc`. The sample `INCLUDE` files for `param.inc`, i.e., `poly9995.inc`, `poly9911.inc`, `poly9111.inc`, `poly6775.inc`, and `poly6711.inc`, and the C shell script `poly.inccr` are also modified due to this change. This is to facilitate the procedure of interfacing *Polyrate* with a variety of electronic structure packages.
6. Subprogram `HEADER` is renamed as `HEADR`. This is to facilitate the procedure of interfacing *Polyrate* with a variety of electronic structure packages.

Version 4.5.1 (September 1992) (Bug fix)

1. In subprogram `REACT`:
A bug in the computation of the zero of energy for a unimolecular reaction when `LGS(35) = 0` has been corrected. This bug was introduced in version 4.1.1.

Version 4.5.2 (March 1993) (Bug fixes)

1. In subprogram `READ5`:
A bug in the following print statement

```
WRITE (FU6,7200)
```

has been corrected. The statement has been changed to

```
WRITE (FU6,7200) NTEMP
```

This bug did not affect any of the calculations; only the output was affected.

2. In subprogram READ5:
For $LGS(35) \uparrow 0$ (for MORATE calculations), a bug in the read statement for DLX and DERSTP has been corrected. This statement read DLX and DERSTP but it should have read DLX, DERSTP, and D3LX. The statement for setting D3LX equal to DLX if D3LX = 0 and $LGS(35) \uparrow 0$ has also been added to this subprogram.
3. In subprogram RPHRD2:
For $LGS(30) = 1$ and $LOPT(4) = 0$, the FREQ array which is used to store the input frequencies was not initialized before the subprogram read in the frequencies of a new species. This bug did not affect any of the calculations, but it caused some wrong output. This has been fixed by adding appropriate initialization statements.
4. In subprogram HINDRT:
The variable NATOM was reinitialized in the subprogram HINDRT as a local variable. This could cause problems when the vibrational modes of the reactants or products are treated as hindered internal rotators. This bug, which was introduced in version 4.0, has been corrected. None of the test runs in the distribution package are affected by this bug.

Version 5.0 (March 1993)

1. An algorithm to allow frequencies, potential energies, and moment-of-inertia-tensor determinants to be corrected by an interpolation scheme has been added. This method is called VTST-IC (i.e., VTST with interpolated corrections). In the present version, we have included zero-order interpolation, i.e., VTST-IC-0. At the zero-order level, corrections are interpolated by using the accurate values for the reactant or a pre-saddle minimum, the saddle point, and the product or a post-saddle minimum. Various types of functions are used for interpolation depending on the shape of MEP and on the relative magnitudes of the corrections at those points mentioned above. Details of this zero-order interpolation scheme are given in the subsection on IVTST in the sections on theoretical background for dynamics calculations and selected details of the calculations. The various options of this interpolation scheme can be selected by setting LGSIC options in the FORTRAN unit fu50 input. In addition, users should supply accurate values of quantities needed for interpolation on FORTRAN unit fu50. For this option, subprograms COG, COHBT, ECKART, HBT, READ50, UPDMI, VICLCG, ZOC3P, ZOCAST, ZOCFRE, ZOCMI, ZOCPAR, ZOCPRN, ZOCVCL, and ZOCUPD have been added. Subprograms LCG3TH, MUBAR, MUCDSC, PATH, READ5, RESTOR, RESTRT, ZEROPT, and MAIN, and the INCLUDE files common.inc and param.inc have been modified.
2. In version 5.0, if $LGS(30) < 1$, normal mode frequencies and generalized normal mode frequencies are ordered in descending order for all output and input. Related quantities, e.g., normal mode eigenvectors, zero point energies, individual vibrational partition functions, etc., are also ordered accordingly in all output. Subprograms FINOUT, REACT, READ5, and REDGEO have been modified for this change.
3. The canonical optimized multidimensional tunneling (COMT) approximation and the microcanonical optimized multidimensional tunneling (μ OMT) approximation are now implemented. These methods are explained in detail in the subsection on optimized multidimensional tunneling calculations. These methods will be used in evaluating transmission coefficients whenever SCT and LCT calculations are both requested. (In version 5.0, SCT calculations are carried out whenever LCT calculations are carried out, so the critical distinction is whether LCT calculations

are requested.) The new subprogram PBTSRT has been added. Subprograms KAPVA, FINOUT, and TABLE and the INCLUDE file common.inc have been modified.

4. A new option for calculating the vibrational partition functions using simple perturbation theory [Truhlar and Isaacson, *J. Chem. Phys.* **94** (1991) 357] for the case of no degenerate vibrational modes was implemented. (Corrections for Fermi or other resonance are not included yet.) In this method, vibrational partition functions are evaluated using the ground-state energy and the fundamentals, which are calculated using perturbation theory up to first order for the quartic terms and the Coriolis interaction, and 2nd order for the cubic terms. Subprograms ANCOEF, CHKFRE, CHKUP, CORTRM, GRAD, GRADDER, and PTQVIB have been added. The INCLUDE file common.inc and subprograms CENTER, COLSHF, INITZE, MEPSRT, NORMOD, PROJCT, RATE, RESTOR, RESTRT, SSAVE and TSRATE have been modified.
5. In subprogram FINOUT:
In version 5.0, the output of calculated transmission coefficients in the file linked to unit fu6 and rate constants in the files linked to units fu6 and fu15 has been cleaned up, and only the quantities that are requested to be calculated will be printed.
6. Various changes have been made to make the subprograms more modular. They are summarized as follows:
 - 6A. In subprogram TRANS:
The position and the gradient vectors are now passed as arguments in call statements rather than being passed through the common block POTCM.
Subprograms ANHARM, ANHDER, INTFNC, INTPM, LCG3TH, NORMOD, PATH, PHSINT, QQPOT, REACT, TP, and WKBVIB have been modified.
 - 6B. In subprograms RESTRT and RESTOR:
Subprogram RESTRT now writes MEP information to unit IOUT which is passed from the calling subprograms as an argument in the call statements. Subprogram RESTOR has been restructured, and the writing of the information corresponding to the merged MEP to FORTRAN unit fu3 has been removed since this function is now carried out by subprogram RESTRT.
 - 6C. The calculations of quantities needed for the cubic-starting algorithm have been removed from subprogram PATH, and they are now carried out by a new subprogram CUBST.
 - 6D. The calculations related to the electronic structure input file option have been removed from subprogram PATH, and these are now carried out by a new subprogram ELRPH.
 - 6E. The writing of reaction energetics has been removed from subprogram NOROUT, and these are now written by a new subprogram ENROUT.
 - 6F. The writing of saddle point energetics has been removed from subprogram MAIN, and these are now written by a new subprogram SENOUT.
 - 6G. The part of the code that prints MEP information, including V_{MEP} , V_a^G , μ_{CD-SC} , and generalized transition state frequencies has been taken out of the subprogram

PATH and placed in a new subprogram MEPOUT. This subprogram is always called by PATH if the MEP is calculated.

- 6H. The contents of the subprogram MAIN have been grouped into 6 new subprograms, DOREST, DORPH, DOREPR, DOSAGE, and DOPNM. These subprograms are called by MAIN to control the VTST calculations.
- 6I. In subprogram RESTOR:
For a restart calculation, the variable EZERO is echoed to FORTRAN unit fu6. The write statements for writing the MEP information to FORTRAN unit fu6 have been replaced by a call to the subprogram MEPOUT. All remaining FORMAT statements in this subprogram have been modified to be case sensitive and to fit in 80 columns.
- 6J. In subprogram MAIN:
The open statements for the files linked to FORTRAN units fu5 and fu6 have been replaced by calls to the new subprogram FIOOPEN.
- 6K. For $LGS(35) = 1$, i.e., an electronic structure input file calculation, the calculation, printing, and storage of the bond orders and charges have been removed from the code. This involved changes in the following subprograms: COLSHF, MAIN, MEPSRT, PATH, POLYAT, RESTOR, RESTRT, SADDLE, and SSAVE. This was done to make the link between *Polyrate* and any electronic structure package more modular and more transparent.
- 7. Various changes have been made to clean up the program. They are summarized as follows:
 - 7A. The matching of normal modes to those at previous save grid points has been changed. Prior to version 5.0, the usage of arrays was different when LCG3 calculations were requested from when they were not. In this version, the program is restructured so that the matching of normal modes is always performed in the same fashion regardless of which tunneling method is used. Subprograms FDIAG, INITZE, and OVERLP have been modified to accommodate this change.
 - 7B. In subprogram ZEROPT:
The number of degrees of freedom corresponding to rotation, translation, and the reaction path (if GTST), and the number of degrees of freedom of a species are now evaluated in subprogram ZEROPT rather than being passed to ZEROPT. Subprograms RPHINT and RPHRD2 are also modified.
 - 7C. The error-checking on the compatibility of ICODE(IOP) and LGS(34) has been moved to subprogram REDGEO.
 - 7D. Extra blank lines in the long output (FORTRAN unit fu6) have been eliminated, and the layout and the clarity have also been improved. Most of the subprograms that echo information to the long output have been modified.
 - 7E. All the unit conversion and variable initialization that were in the subprogram MAIN have been moved to the subprograms INITZE and READ5.
 - 7F. The number of vibrational modes for each reactant and product species and for the generalized transition state is now only calculated in the subprogram REDGEO and is stored in the array NF. After that, these numbers are referenced through the arrayelements whenever needed. The calculation of these numbers in other parts of the program has been removed.

- 7G. In subprogram BCALC:
Two lines of FORTRAN which were commented out have been removed. These lines were used for debugging during the preparation of version 4.0.
- 7H. The option LGS(26) = 1, which allowed the user to force the points on the MEP to have a particular symmetry, has been removed in this version of the code. This involved changes in the following subprograms: ELRPH, OPTION, PATH, and READ5, and the removal of the dummy subprogram SYMCHK.
- 7I. As a result of item 6K, the option LGS(4) = 3 has been removed; this option controlled the printing of the bond orders and the charges which are no longer calculated in this version of *Polyrate*. This involved changes in the following subprograms: INITZE, MAIN, PATH, POLYAT, SADDLE, and RESTOR.
- 7J. All C*M lines, except those for the statement "include 'esp.inc'", have been removed. This involved changes in the following subprograms: MAIN, POLYAT, REACT, READ5, and SADDLE. This will cause the program to perform a couple more 'if' checks, but it should not slow the code down appreciably.
8. In subprogram NORMOD:
The position vector of a GTS in the mass-scaled Cartesians is now saved in a temporary array before it is transformed to unscaled Cartesians to evaluate the second derivatives. The mass-scaled position vector is restored afterward by using the values in the temporary array. This procedure avoids the round-off error when the position vector is transformed back and forth between the scaled and unscaled Cartesians.
9. The SCSAG tunneling method has been removed because it is obsolete, being replaced by CD-SCSAG. As a consequence, in the unit fu5 input file, the options LGS2(3) and LGS2(6) on record 1A and the variable M0 on record 20 are no longer used. Subprograms BCALC0, ELRPH, FINOUT, KAPVA, MUCDSC, OPTION, PATH, and PSAG have been modified.
10. The call to subprogram MEPSRT by subprogram PATH has been replaced by a call to the new subprogram MEPINV for better performance. The subprogram MEPSRT is now only called by the subprogram RESTOR.
11. The common block POTCM in the INCLUDE file common.inc has been renamed PEFCOM. This has been done to avoid conflicts between this common block and the one used in the three-body analytic potential energy functions that are in the test suite.
12. The code has been modified so that all the files used in a calculation are no longer closed at the end of the calculation by a call to FCLOSE. Instead, the files in a calculation are closed after they are used; in fact, there are three points in the calculation when files are closed. First, after the call to READ5, all the input data files that have been used up to that point are closed by a call to the new subprogram FCINPT. Second, after the MEP calculation, all input and output files used during the MEP calculation are closed by a call to the new subprogram FCMEP. Finally, at the end of the calculation, all files, input and output, which have not been closed are closed by a call to the new subprogram FCRATE. The subprogram FCLOSE no longer exists. The three new subprograms, FCINPT, FCMEP, and FCRATE, call another new subprogram FICLSE; this subprogram closes a file linked to a FORTRAN unit NUNIT with the status FSTAT. This change was made to safeguard against loss of information in the event of a system or code crash.

13. Some changes have been made in the *Polyrate* test runs. They are summarized as follows:
 - 13A. Five new test runs have been added to this version of *Polyrate* to test the VTST-IC options. They are cmctr1, cmctr2, ho2tr4, ho2tr5, and nh3tr1.
 - 13B. The forward symmetry factors SIGMAF in the unit fu5 input data files of the ch5 test runs have been changed to the correct values. The forward rate constants are a factor of 4 smaller for the test runs ch5j1tr5 and ch5j1tr6, and a factor of 2 smaller for other ch5 test runs than they were in the previous versions.
 - 13C. Copies of the short output files linked to FORTRAN unit fu14 have been removed from the *Polyrate* distribution.
 - 13D. For the clhbr, ho2 and oh3 test runs, the data for the potential energy surfaces have been placed inside the respective potential energy surface codes, so that no potential energy surface data need now be read.
14. The INCLUDE files poly9995.inc, poly9911.inc, poly6775.inc, and poly6711.inc have been renamed param4.inc, param3.inc, param2.inc, and param1.inc, respectively, and the include file poly9111.inc has been removed from the *Polyrate* distribution.
15. The subroutine RST has been renamed RSPDRV to avoid the name conflict with the subroutine RST in Cray's science library.

Version 5.0.1 (May 1993) (Bug fixes)

1. For LGS(30) = 1 and LOPT(4) = 0, the imaginary frequency of the saddle point was not printed out correctly in the unit fu6 output file. This was because the program printed the frequencies before the imaginary frequency was read. This has been fixed by printing the frequencies after the imaginary frequency is read from the unit fu30 input file. The subprograms RPHRD1 and RPHRD2 have been modified. This bug was present since version 1.1.
2. For LGS2(1) = 1, and when the hindered-internal-rotator approximation was used, the program printed incorrect data in the unit fu6 output file because of a missing argument in one of the WRITE statements in the subprogram READ5. This has been fixed by adding the appropriate argument. This bug was present since version 4.0.
3. In subprogram LCVMOD:
The variables EVIBM and VMD were not initialized in the subprogram LCVMOD. Due to this error, a non-zero value calculated by the subprogram LCVMOD was used when there should have been no contribution from a generalized normal mode to the correction term of the effective potential for LCG3 tunneling. This has been corrected by adding appropriate initialization statements. This bug was present since version 4.0.

4. The modification in the subprogram MEPSRT due to the simple perturbation theory treatment option introduced in version 5.0 was not put into the new subprogram MEPINV. (See version 5.0 revision history, items 4 and 10.) This has been fixed by adding appropriate statements in the subprogram MEPINV.
5. The forward symmetry factor SIGMAF in the unit fu5 input data files of the ch5 test runs has been corrected. This factor is now 2 in the test runs ch5j1tr5 and ch5j1tr6 and is 4 in the other ch5 test runs. As a result, the forward rate constants are a factor of 2 larger for the test runs ch5j1tr5 and ch5j1tr6, and a factor of 4 larger for other ch5 test runs than they were in version 5.0. Except for the test runs ch5j1tr5 and ch5j1tr6, this factor was correct in the unit fu5 input data files of the ch5 test runs before version 5.0.
6. This version and the previous version (5.0) of *Polyrate* contain unpublished developers' versions of the VTST-IC algorithm which are not in its final form and may have bugs. So the VTST-IC feature should not be used in versions prior to 5.1.

Version 5.1 (December 1993)

1. The VTST-IC method, which was present in undocumented development version in version 5.0, has been included in the form documented in *Faraday Trans. Chem. Soc.* **90** (1994) 1715. It is now the recommended method if more accurate data are available for the reaction path information than the potential routine can provide. This is especially useful when an implicit semiempirical potential energy surface is used, e.g., in a MORATE calculation with version 5.1 or later of MORATE. Numerous changes have been made in several subroutines to implement the final version of VTST-IC, and the unit fu50 input formats have also been changed.
2. The input files for unit 50 for test runs cmctr2, nh3tr1, and ho2tr5 have been modified because of the changes mentioned in item 1. The calculated results are a little bit different from the previous versions.
3. The zero-order IVTST method described in *J. Chem. Phys.* **95** (1991) 8875 has been added to this version for bimolecular reactions. A new test run, ch4ohtr1, has been added in the distribution package to test this method. This option uses file fu50.
4. A new utility program, hrotor.f, which calculates the reduced moment of inertia of a hindered-rotor vibrational mode and the harmonic, free rotor, and hindered-rotor partition functions of this mode, has been added to the distribution package.
5. Two other new utility programs, findb.f and findl.f, which calculate range parameters for VTST-IC calculations, are also included in the distribution package.
6. Two other new test runs, ch5j1tr8 and oh3tr8, which are two of the test cases in *J. Chem. Phys.* **96** (1992) 5758, have been added to the distribution package.

Version 5.1.1 (March 1994) (Bug fixes)

1. In subprogram EXTRAP:
A bug in extrapolating the classical potential energy curve along the MEP on the product side has been corrected. This bug was introduced in or before version 1.5.
2. In subprogram DOREST:

A bug in printing the saddle point energetics for a restart calculation has been corrected. This doesn't effect any reaction rate constant calculations. The bug was introduced in version 5.0.1.

3. In subprogram ANHARM:
A bug in transforming between mass-scaled and unscaled Cartesian coordinates has been corrected. This only effects the calculations using an electronic structure input file (unit fu30), i.e., when LGS(30) = 1. This bug was introduced in version 5.0.
4. The printing of curvature components, which was reversed in the previous version, has been fixed.

Version 6.0 (June 1994)

1. *Polyrate* 6.0 involved a complete change in the structure of the user input files previously numbered fu5, fu21, fu40, and fu50. The user-supplied data needed by the program is now all in FORTRAN units fu5 and fu50. The only other input file is unit fu30, which contains electronic structure information in lieu of a global analytic potential energy surface. The new unit fu5 and fu50 input files are in the keyword style.
For the unit fu5 input, the user options have been broken into nine sections: GENERAL, REACT1, REACT2, PROD1, PROD2, START, PATH, TUNNEL, and RATE. In each of these sections, one uses keywords to specify the parameters, calculation type, and variables for the calculation. (See the keyword section of the *Polyrate* manual for complete details.) This restructuring of the input has added new capabilities to the program. One example is that it is now possible to converge the geometry in each of the stationary point calculations to a different tolerance and to use a different step size for each stationary point. This flexibility comes from changing the previously-used LGS options, which applied to the entire calculation, to keywords that are specified for each reactant, product, and saddle point in the system. This new type of input makes the program much easier to use. Most of the keywords have reasonable defaults. No input files, except those for unit fu30, from any previous version of *Polyrate* will work with version 6.0 or later. The user should find that the inconvenience of converting existing input files to the new format will be offset by the ease of generation of future input files. All the test run input files, for both unit fu5 and unit fu50, in the distribution package have been converted to the new format.
2. Numerous new subprograms have been added in version 6.0 because of the new keyword input user interface. These new subroutines reside in the interface.f file in the source code directory. Subprograms MAIN, OPTION, OPT50, READ5, RESTOR, RESTRT, and ZOCPAR have been modified for the keyword input interface, and subprograms READ8, READ40 and READ50 have been removed.
3. The section of the unit fu6 output file that echoes the selected options has been completely rewritten. It now contains more details than earlier versions, and it is restricted to 80 columns. The part of unit fu6 output that echoes the VTST-IC calculation information has also been refined. This involves changes in the subprogram ZOCPRN.
4. The option [LGS(24)] for including the energy of product diatomics in endoergicity calculations has been removed. In addition, the program now has some built-in intelligence, and the user is no longer required to input quantities that *Polyrate* can determine. For example, the number of frequencies, the mass of the individual

atoms, the number of atoms in the system, the linearity of a species, the number of temperatures, and the number of activation energies are now all determined by the *Polyrate* program.

5. The options LGS(18) and LGS(19) for special treatments for V_{MEP} and vibrational partition functions using M. Quack and J. Troe's method [*Ber. Bunsenges. Phys. Chem.* **81** (1977) 329] have been removed. Because of this, test run ho2tr2 and related files have also been removed.
6. New default values for the range parameters of the correction functions used in the VTST-IC calculation have been included in version 6.0. (See the subsection on the general description of unit fu50 input.) The previous method for calculating the default values is obsolete and is no longer supported. However, it is still strongly recommended that one use the utility programs findl.f and findb.f included in the distribution package to find appropriate values for these range parameters.
7. The normal mode analysis section in the unit fu6 output file for a diatomic Morse oscillator (ICODE = 2 in earlier versions or with the keyword DIATOM in this version) has been modified. The stretching vibration is now labeled as mode 1 instead of mode 6, which is consistent with the output for polyatomic systems.
8. The manual has been converted from an ASCII text format to Postscript format and fully updated to reflect the major changes in the input.
9. Machine-dependent scripts have been added for IRIS Indigo machines.

Version 6.2 (August 1994)

1. The *Polyrate* long output file (unit fu6) has been significantly improved, and all the output lines have been restricted to a maximum of 80 columns. Numerous changes have been made in subroutines that contain output statements.
2. A new quantized-reactant-state tunneling method for unimolecular reactions has been introduced in this version of *Polyrate*. This method can be used together with all the existing tunneling methods. The subroutines BOLTZ, INTAB, KAPVA, RATE, RTUNNL, and SETLGS have been modified, and the new subroutines QRSENE, QTQVIB, RQRST, SIMPSN, and WKBENE have been added into the code. The INCLUDE files common.inc and param[1-4].inc have also been modified. The new keywords for this method can be specified in the TUNNEL section of the unit fu5 input file.
3. Three new output files have been added in this version; they are units fu25, fu26, and fu27. Unit fu25 contains the V_{MEP} and V_{a}^{G} information, and unit fu26 contains generalized transition state vibrational frequency information; these new output files are especially useful for analyzing reaction-path data since they can easily be imported (with minimal editing) to a spreadsheet program for analysis or plotting. Unit fu27 contains the Cartesian coordinates of the generalized transition states in the XYZ input format of the XMOL program (which is *not* supplied with *Polyrate*). The user can thus visualize the geometry changes along the reaction path by using the XMOL program, if available. The subroutines MAIN, FIOPEN, FCRATE, INTAB, RRATE, and SETLGS have been modified for this enhancement, and the new subroutines PRMEP and RPRMEP have been added to the code. The INCLUDE files common.inc and param[1-4].inc have also been modified for this enhancement. The new keywords for generating these new output files can be

specified in the RATE section of the unit fu5 input file.

Version 6.5 (January 1995)

1. The *Polyrate* input file (unit fu5) has been significantly altered in preparation to facilitate a more convenient incorporation of direct dynamics options. New sections, i.e., ENERGY, SECOND, and OPTIMIZATION, have been created. Numerous changes have been made throughout the program to localize all calls to the potential energy subroutine and calculations of first and second derivatives.

Version 6.5.1 (October 1995) (Bug fixes)

1. A bug in calculating the anharmonicity constant for the Morse model has been corrected. The local dissociation energy was calculated incorrectly when not using an analytic potential energy surface. This required modifications in the subroutine ANHARM. Also, when the dual-level dynamics method is used, the anharmonicity constant was not corrected using the higher-level data. This has been fixed by modifying in the subprogram ZOCUPD.
2. A subroutine call in subprogram RATE to the subprogram HRPART had an incorrect number of arguments. Also, one call to FITMAX in subprogram RATE had an incorrect number of arguments. These bugs have been fixed in this version.
3. A bug in subroutine DIATOM has been fixed so that the subprogram works correctly for unit fu30 input.
4. A bug in the extrapolation options in subroutine RPHINT has been fixed.

Version 6.6 (October 1995)

1. An option for performing generalized normal mode analyze in curvilinear coordinates has been added via the COORD keyword in the PATH section. For a linear or bent triatomic i-j-k, the curvilinear internal coordinate definitions are chosen automatically by the program; this triatomic option includes the i-j and j-k stretches and the i-j-k bends. For systems with more than 3 atoms, the INTDEF keyword can be used to define the curvilinear internal coordinates. Test run ho2tr2 was added for the $\text{OH} + \text{O} \rightarrow \text{HO}_2$ reaction. Test runs oh3tr9 and oh3tr10 were added for the $\text{OH} + \text{H}_2 \rightarrow \text{H}_2\text{O} + \text{H}$ reaction. Details for these reactions are given in the section on sample test runs.
2. The zero- and first-order (the global approach) IVTST methods have been added, using fu29. (Recall that IVTST-0 has been available using fu50 since version 5.1, so now these are two ways to do IVTST-0.) Several test runs were added to the distributed test suite. The test runs ch5j2itr1 and ch5j2itr2 were added for the $\text{CH}_3 + \text{H}_2 \rightarrow \text{CH}_4 + \text{H}$ reaction, the test run ch4ohtr2 was added for the $\text{OH} + \text{CH}_4 \rightarrow \text{H}_2\text{O} + \text{CH}_3$ reaction, and the test runs ch4cltr1 and ch4cltr2 were added for the $\text{CH}_4 + \text{Cl} \rightarrow \text{HCl} + \text{CH}_3$ reaction. Details for these reactions are given in the section on sample test runs.
3. The subroutine NEWT has been rewritten in order to support the quasi-Newton BFGS optimization method. This method can be used with both analytic surfaces and direct dynamics, and it is now the default for reactant and

product optimization. Some changes related to this implementation have also been made in subroutines

LINMN, SECCEN, and SECCEP. The keyword CONVERGENCE in the OPTIMIZATION section is no longer valid, and it has been replaced by the GCOMP and TSGCOMP keywords to emphasize that the convergence criterion for optimizations has changed.

4. A new sub-keyword, *nofreqcorr*, has been added to the unit fu50 data file for the keywords RCINFO and PCINFO. This sub-keyword can be convenient when higher-level data are unavailable for precursor or successor complexes. The subprograms READIC, OPT50, ZOCPAR, and ZOCUPD have been modified for this enhancement.
5. The code now runs on the Power Macintosh 7100/66 (Mac OS 7.5.1) computer using the LS FORTRAN compiler (version 1.1b1) available from Language Systems Corporation.

Version 6.6.1 (June-August 1996) (Bug fixes)

1. In file givtst.f:
Several errors are corrected in the IVTST calculations. In particular: the forward relative reduced mass is now calculated correctly for the evaluation of the relative translational partition function when unit fu29 is used for input. Also, the output of the zero- and first-order IVTST methods has been made clearer. The input energy of reaction is now converted to atomic units first, in order to prevent mistakes in the interpolation. The imaginary frequency of the transition state can be input as the last element of the TSFREQ section. The variables VAP and VAR are now used correctly in determining the lower limit of integration over energy in the transmission coefficient calculation. Most of the constants were moved to the INCLUDE file percon.inc for consistency. Finally, the CAG correction has been added. Information about evaluated curvatures has been added to the output.
2. In subroutine ENERG:
Version 6.6 had a bug in runs using the unit fu30 input file when one of the reactants or products is an atom. In such cases the subroutine SURF was called, but SURF should not be called in runs that use the fu30 input file. This has been fixed. Now, one can use the dumpot.f source code file as intended when using the unit fu30 electronic structure file input for a case when one of the reactants or products is an atom.
3. In subroutine RPHRD2:
We have corrected the order of fitting the gradients that are read from the unit fu30 input file. In the corrected version, the points along reaction path are always left-shifted even after the saddle point information has been inserted in the middle of the path. This bug had caused curvature calculations in the vicinity of the saddle point to be wrong.
4. In subroutine RFREQ:
This subroutine has been modified to allow the user to input the imaginary frequency for the ZCT calculation. In this case, the user must input the imaginary frequency as the last frequency of the saddle point.
5. In file main.f:

Anharmonicity is no longer allowed in TST-only calculations.

6. In test run ch4cl:
The data for the electronic partition function of the Cl atom has been changed to the correct values. The degeneracy of the ground state has been changed, and the first excited state has been included with correct atomic spectral data.
7. In test runs ch4ohtr1 and ch4ohtr2:
A comment is added for the reduced moment of inertia in the unit fu50 input file in the ch4ohtr1 test run. A more detailed description of keywords RMITS and RMIRP has been added to the manual with an example of how to include the symmetry number for the hindered-rotor approximation. The symmetry number SIGMAF in ch4ohtr2 has been changed to 12. (Notice: the hindered-rotor approximation is not supported when using the unit fu29 input file.)
8. In subroutine TPCDSC:
This subroutine was fixed to prevent the occurrence of a zero-divided-by-zero case.
9. In subroutine DEFSTA:
MODER(i,j) was changed to MODER(j,i) for correct dimensioning. This change will initialize the array as intended. At the beginning of the subroutine, the elements of this array are set to zero.
10. In test runs oh3tr6 and oh3tr10:
Unused data sections were removed from the input file.
11. In files dattim.ibm and dattim.hp:
These files have been modified to remove the NULL characters which generated unsightly hardcopy.
12. In subroutine ZOCPRN:
The title line for the frequency output of the lower-level and higher-level methods for the VTST-IC method has been clarified. The columns were labeled 'Point 1' and 'Point 2' even though MEPTYPEP or MEPTYPEP was set to *two* or *one*; however, the code was printing a sorted list of the combined frequencies. Now the title has been changed to 'Sorted Reactant' and 'Sorted Product' when there are no wells.
13. In subroutine NOROUT:
The variable SPRNT has been replaced with s in order to make sure the keyword SPRNT works properly.
14. In subroutines TSRATE and RATE:
The unused variable LGS(19) inside these routines has been removed since LGS(19) is not used in the current version of *Polyrate*.
15. In file common.inc:
The common block AMASCM has been added for consistency with other blocks.

Version 7.0 (September 1996)

1. The keywords ICR and ICL were added for new interpolated correction methods. The following changes were made: (1) A new variable IVICM has been added in common.inc. (2) The subroutine MUCDSC in polymq.f, subroutines ZOCAST, ZOCPAR, and ZOCUPD in polysz.f, and subroutines INTAB, RICOPT, SURF, RVAR6, SETLGS, WVAR6 in interface.f were modified for the new methods. Also, LGS2(10) is

used internally to store the choice between the ICA, ICL, and ICR options.

2. A new option for electronic structure input data has been added (`POTENTIAL = unit40`). This option reads the data from unit fu40 in keyword form rather than from unit fu30 in formatted records.
3. In subroutine RPHSET:
In version 7.0, the saddle point geometry is no longer read from both unit fu5 and unit fu30; it is only read from unit fu5. All the unit fu30 input files distributed with the present version of the code have no saddle point geometry. A warning message is printed when the NEXTPT option is used with an option code different from 0, and a stop message is printed after the next point is found.
4. In file interface.f:
Options *quad* and *page* for the keyword CURV have been changed to *dgrad* and *dhess* respectively. The PAGE keyword for activating the Page-McIver integrator has been changed to PAGEM. All these changes require that the user modify the input files prepared since version 6.0. All the examples included in the test run suite whose unit fu5 input files make use of these options have been modified accordingly.
5. In the testrun/ch5 directory:
New test runs are added (ch5fu30tr1, ch5fu30tr2, ch5fu30tr3, ch5fu30tr4, ch5fu30tr5, ch5fu30tr6, ch5fu29tr1, ch5fu29tr2, ch5fu40tr1, ch5fu40tr2, ch5fu40tr3, ch5fu40tr4, ch5fu40tr5, ch5fu40tr6, ch5icfu30tr1, ch5icfu30tr2, ch5icfu40tr1, ch5icfu40tr2, ch5fu50tr1 and ch5fu50tr2) to demonstrate the different input methods allowed in the current version of *Polyrate* for electronic structure data.
6. In the testrun/oh3 directory:
The names of some of the existing test runs have been changed (oh3tr6 and oh3tr10 have become oh3fu30tr3 and oh3fu30tr2, respectively; oh3tr7, oh3tr8 and oh3tr9 have become oh3tr6, oh3tr7 and oh3tr8, respectively). New test runs to illustrate the use of direct dynamics calculations with unit fu30 and unit fu40 input files have been added (oh3fu30tr1, oh3fu30tr4, oh3fu30tr5, oh3fu40tr1, oh3fu40tr2, oh3fu40tr3, oh3fu40tr4, oh3fu40tr5).
7. The manual has been improved with the following major changes: (1) A new section describing the reaction-path curvature calculations has been added. (2) The section describing the reaction rates including tunneling calculations has been improved. (3) A better description of the VTST-IC method has been included in the sections on theoretical background for dynamical calculations and selected details of the calculations. (4) The description of extrapolation methods has been improved. (5) The description of the unit fu30 input file has been improved. (6) A new section has been added to explain the use of the unit fu40 input file for direct dynamics calculations. (7) A summary has been added to the sample test run section. (8) The section summarizing subprograms has been updated. (9) New errata sections have been added for published papers referenced in the footnotes to the text and bibliography sections. (10) More consistent font choices have been used to distinguish subprogram names, variables, keywords, and keyword values.
8. The MDMOVIE keyword has been added in the GENERAL section to generate a file (as unit fu65) for the MDMOVIE program. This change involved changes in the following subroutines: DEFGEN, INTAB, RGENER, RVAR6, WVAR6, and DEFPAT. A new variable, IMDMOV, was added in the file common.inc.

9. In file `set_machine.jc`:
This job control file has been modified for more choices of machines. The compiler options for most of the platforms are for the O3 optimization level, and a new set of scripts for the IBM RS/6000 Model 590 machine is available.
10. The output of *Polyrate* has been cleaned up so that no line exceeds 80 columns, the respective units are included for each value printed, clearer headings are printed together with the numerical information, timing routine prints times to 0.01 seconds instead of 0.0001 seconds and with a more clear format, and the debug information in the VTST-IC section is removed. To print the debug information, remove the comment from the `CALL ZOCAST` line in the `ZOCUPD` routine.
11. In file `givtst.f`:
Improved output is provided for several parts of the IVTST calculations. A table containing the reaction energetics is printed. The parameters for the interpolating functions match the notation used in *J. Chem. Phys.* **95** (1991) 8875. The transmission probabilities are better labeled and the reaction-path curvature results are printed in first-order IVTST calculations. Formulas used in subroutines `BCALP` and `MUSCCD` have been modified to follow the notation used in the references. The `DELTAH` keyword has been removed from the unit `fu29` input file.
12. In subroutine `BANGLE`:
This subroutine has been renamed `BANGLE1` to avoid using the same name as is used in the MOPAC program.
13. In subroutines `RPHRD2` and `RPHDXN` :
The option choices `LOPT(3) = 99`, `LOPT(3) = 100`, and `LOPT(4) = 99` have been added for the unit `fu30` input file.
14. In subroutine `KAPVA`:
A new table is printed in the long output file to display the values of the transmission coefficients obtained with all the possible combinations of the number of quadrature points used in the evaluation of the theta and Boltzmann integrals.

Version 7.0.1 (February 1997) (Bug fixes)

1. In file `poten/ch5.f`:
The data parameter `DIJ` has been changed from 6.27396 to 3.1190 to correctly correspond to *J. Chem. Phys.* **87** (1987) 7024.
2. In subroutine `REACT`:
All occurrences of `LGS(24)` are changed to `LGS2(4)`. This was an error.

Version 7.1 (February 1997)

1. In files `interface.f` and `polyag.f`:
A keyword option `REORDER` has been added to the `PATH` section. The default is set to `NOREORDER`, which forces all calculated frequencies to be in canonical order along the MEP. Previous versions of *Polyrate* have the default set as `REORDER`, where the frequencies are ordered according to overlap calculations. The `DEFPAT`, `INTAB`, `RPATH`, `RVAR6`, `WVAR6`, `NOROUT`, `RFREQ`, `GETFREQ`, `RPHRD2`, `IRFGT0`, and `FDIAG` subroutines were modified, and a new variable, `ireord`, was defined in file `common.f`. Note: All calculated frequencies in the `FREQ` array are stored in

- canonical order (if NOREORDER is used) except at the saddle point, where we store the imaginary frequency in position 1.
2. In file poly40.f:
The default unit for frequencies read from unit fu40 has been changed from atomic units to wavenumbers. The keyword `FREQUNIT` controls the units of the frequencies; values are *waven* for wavenumbers (default value) and *au* for atomic units.
 3. The file `dattim.ibm` has been revised to obtain the date and time without the need for `dateclock.ibm`.
 4. In file `interface.f`:
In the subroutine `RMUVT`, the variables `FIT1` and `FIT2` were changed to `IFIT1` and `IFIT2`.
 5. A change was made to subroutine `SADDLE` to correct the zero of energy in calculations where an electronic structure program is being used in place of an analytic surface or a file of data.
 6. Other changes were made to the `NORMOD`, `POLYAT`, `REACT`, `DORPH`, `FCMEP`, `CENTER`, `DOREPR`, `RATE`, `SADENG`, and `HEADR` subroutines. In general, these changes were to make the linking of *Polyrate* with various electronic structure programs more seamless or to clean up old code.
 7. In subroutine `FINOUT`:
Lines containing `LGS(13)` are removed because this option is no longer used.
 8. In addition, several changes were made that will be supported by later versions of *Polyrate* or in `ACESRATE` or both. These changes should not affect users of *Polyrate*-version 7.1 who do not have the `ACESRATE` interface, but they are listed here for completeness.
 - 8.1. Specialized `ACESRATE` hooks were added to subroutines `ENERG`, `FIRST`, `GEOMOP`, `PREP`, and `SECDER` and to files `common.inc` and `param*.inc`, and the following subroutines were modified to read and process `ACESRATE` keywords: `DEFENG`, `RACES`, `READ5`, `RENERG`, `RGEOM`, `RGSAD`, `RGZMAT`, `RSECND`, `RZVAR`, `DEFOPT`, `DEFPAT`, `DEFSTA`, `INTAB`, `INTABS`, `RGSZM`, `RICOPT`, `ROPT`, `RSTAT`, `RVAR6`, `SETLGS`, `WVAR6`, `FIOPEN`, and `DOREST`.
 - 8.2. The restart option was enhanced. In earlier versions of *Polyrate* for a two-level (VTST-IC) calculation any change in the higher-level potential energy surface required the recomputation of the entire lower-level minimum energy path. When this restart option is supported, it will be possible to read the lower-level path from a unit `ful` restart file. `MAIN` was also restructured to make restart and direct dynamics calculations easier and more flexible.
 - 8.3. The `CALCMUFF` keyword option was added. When this is supported the reduced mass will be calculated when the reaction path is computed. In previous versions of *Polyrate*, the effective reduced mass was only computed if tunneling was also selected. This means that in a restart calculation, if the original run did not include a tunneling calculation, tunneling could not be used in any later calculation without recomputing the entire reaction path.

Version 7.1.0.5 (March 1997)

1. A fourth switch keyword, GAUSSIAN, is added to the OPTIMIZATION and SECOND sections of the unit fu5 input file. The four switch keywords *Polyrate*, MOPAC, ACES, and GAUSSIAN are used by GEOMOP and SECDER. The ROPT and RSECND subroutines are modified for this update. This is a temporary measure so the first version of GAUSSRATE can be implemented with minimal changes to the structure of version 7.1.

Version 7.1.1 (March 1997) (Bug Fixes)

1. In files energetics.f, common.inc, and interface.f:
The keyword SADDLE is introduced to indicate whether the reaction has a saddle point or not. The following subroutines are modified: SADDLE, SADENG, DEFPAT, INTAB, OPTION, RPATH, and SETLGS. To perform the calculations with NOSADDLE correctly, a new *gradi* option of the keyword FIRSTSTEP has to be used to obtain the first point by following the gradient direction. Input files of the test runs ho2tr1, ho2tr2, and ho2tr3 were changed due to this update. In versions 6.0-7.1.0.5, the NOOPT keyword was correct only for reactions without a saddle point. Thus the NOOPT keyword was equivalent to NOOPT, NOSADDLE and FIRSTSTEP = *gradi*.
2. In subroutine SECDER:
The subroutine calls for evaluating the second derivatives of the potential with *quadratic* and *fourth* are corrected for the right number of frequencies. This corrects a mistake that was introduced in version 6.5.

Version 7.1.2 (March 1997) (Bug Fixes)

1. In subroutines RPHDXN, RPHRD2, CVCOOR, and file poly40.f:
The transformation from rectilinear to curvilinear internal coordinates requires the unnormalized gradients. If normalized gradients are given in the unit fu30 and unit fu40 input files, there is no way to perform the transformation from rectilinear to curvilinear internal coordinates. Incorrect statements in the subroutine CVCOOR were removed.
2. In files ch5fu30tr*.fu30, ch5fu40tr*.fu40, ch5icfu30*.fu30, and ch5icfu40*.fu40:
The electronic structure program GAUSSIAN94 gives the Cartesian gradients instead of Cartesian forces in the formatted checkpoint file (even though the output says Cartesian forces). Test runs prepared based on the forces taken from GAUSSIAN94 are corrected. Also, the SIGN of unbound eigenvector at the saddle point is set to *product* instead of *reactant*.
3. In files percon.inc and givtst.f:
The constant CALAU is removed from *Polyrate* to prevent confusion with the parameter of the same name in MOPAC.
4. In the ch4cl* test runs:
The geometries of the stationary points in the input file have been changed. The values have been wrong since version 6.6.

5. In subroutine NEWT:
The output statements for the optimization results are corrected.

Version 7.2 (April 1997)

1. In subroutines REACT and SETLGS:
A portion of the program that calculates the zero of energy has been rewritten. The option *reactants* no longer exists in the current version of program. The zero of energy is determined automatically after the reactants are evaluated unless the *read* option is chosen for the EZERO keyword. Changes were made in test runs ch5j1tr4.dat, clhbrtr1.dat, cwmctr2.dat, cmctr1.dat, cmctr2.dat, ho2tr4.dat, ho3tr5.dat, nh3tr1.dat, oh3tr1.dat, oh3tr7.dat, oh3tr8.dat, and oh3tr3.dat.
2. In file intbsv2.f:
The transformation from rectilinear coordinates to curvilinear internal coordinates of doubly degenerate linear bends has been added. Changes were made in subroutines CVCOOR, CONOUT, ICODAL, BMAT, BIMAT, and BTENS, and in the common.inc file. To use this option, the '=' symbol is used to denote the linear bends in the INTDEF keyword option. The input files of the ch5j1tr4, ch5fu30tr6, and ch5fu40tr6 test runs are modified to illustrate this new curvilinear internal coordinate option.
3. In files interface.f and polysz.f:
New interpolated corrections options have been implemented. First, a difference of two Eckart function fits is used to interpolate the correction to $V_{\text{MEP}}(s)$. The new keyword option is called ECKART. The following subroutines have been modified: DEFPAT, INTAB, RICOPT, ZOCPAR, and ZOCVCL. Second, two improved options are available for interpolating corrections to the frequencies. The original IC method is denoted as ICA, and the two new options are called ICL and ICR. The combination of the ICL option with the ECKART option is taken as the new default. To perform calculations with the original IC method, the NOECKART and ICA keywords are required. The ch5icfu30tr1 and ch5icfu40tr1 test runs are used as examples of the new dual-level ICL/ECKART option. The input files for the ch4ohtr1, cmctr2, ho2tr5, ch5fu50tr2, and nh3tr1 test runs are changed to accommodate the new default setting.
4. In files polysz.f and comp_src.*:
The BLAS library routines are moved into a separate file, fromblas.f, from the main *Polyrate* program. If their machine has a pre-installed BLAS library, users can modify the script comp_src.jc to include the machine library which yields the best performance.
5. In subroutine FINDL:
The subroutine FINDL, previously distributed only as a stand-alone utility, is now also added to the polyag.f file as a subroutine. This subroutine will automatically determine the L parameter for VTST-IC calculations (the value of the keyword RPL in the unit fu50 input file) in any single-level reaction path calculation with a positive barrier.
6. In files interface.f, hooks.f, and polyag.f:
The keyword WRITEFU30 is added to the GENERAL section of the unit fu5 input file to create a *.fu30 file from a calculation using the hooks. (This option is mainly useful to developers for creating test runs.) Test run oh3tr7 is modified to demonstrate this option. The OPENFI and RPHWRT subroutines have been modified.

7. In file givtst.f:
The RHESS, RGRAD, and GIVTST subroutines have been changed to allow free formatted geometry, gradients, and Hessian input from the unit fu29 file.
8. In file interface.f:
A new keyword, EZUNIT, is added to allow the zero of energy to be input in kcal/mol. A new subroutine, REZERU, and a variable, IEZERU, are added, and the input file for clhbrtr2 is changed for this new feature.
9. The keywords SIGMAF and SIGMAR are no longer saved to file ful for restarts. This allows the user to change their values in the RATE section of the restart run.
10. Subroutines ENERG, FIRST, SECDER, and GEOMOP have been renamed EHOOK, GHOOK, HHOOK, and OHOOK, respectively. The options for the potential keyword were changed and the following subroutines were modified: DERIV2, REACT, DIATOM, DOSAGE, F1DIM, LCG3TH, PHSINT, QQPOT, WKBVIB, DERV24, INTPM, NEWT, NORMOD, SECCEN, SECCEP, ANHARM, ANHER, GRAD, TP, INTFNC, CUBST, INTPM, NEWT, NORMOD, POLYAT, REACT, SADDLE, and DOREST. A new keyword, HESSCAL, was added to provide a choice between calculating second derivatives with the GHOOK or HHOOK subprograms. In the SECOND section, HESSCAL = *ghook* replaces the *Polyrate* keyword, and HESSCAL = *hhook* replaces the MOPAC and GAUSSIAN keywords. The keywords *Polyrate*, MOPAC, GAUSSIAN, BFGS, and TSBFGS in the OPTIMIZATION section were replaced by OPTMIN and OPTTS. Subroutines RHESSC and ROPTOP were added to parse the options.
11. The option *analytic* is replaced by *hooks* for the keyword POTENTIAL in the ENERGETICS section of the unit fu5 input file.
12. In subroutine RPHINT:
The interpolation of the curvature components from the unit fu30 or the unit fu40 basic input grid to the save grid (see the section on methods for interpolation and extrapolation of electronic structure data for a description of these grids) is now based on the absolute value instead of signed values in order to avoid problems related to the continuity of the eigenvector sign convention.
13. In subroutine GFDIAG:
When using curvilinear coordinates, a check is added to set the eigenvalue to zero if a value corresponding to a vibration is smaller than 10^{-18} .

Version 7.3 (July 1997)

1. Three new keywords are added to increase modularity, and as a consequence three new variables are added to the file common.inc. The variable ISUP is used to toggle the keyword SUPERMOL (in the GENERAL section), the variable INITG is used to indicate the option chosen by the INITGEO keyword (in the REACT1, REACT2, PROD1, PROD2, and START sections), and the variable INTMU (for the INTMU keyword in the PATH section) is used to specify the number of points used for the interpolation of the reduced mass at the saddle point. All three variables replace the LGS(35) value of the previous version. For ISUP, the following subroutines are modified: DEFGEN, INTAB, RGENER, SETLGS, NORMOD, POLYAT, REACT, SADENG, and CENTER. For INTMU: DEFPAT, INTAB, RPATH, BCALC0, ELRPH, and PATH are modified. For INITG: REACT, DEFOPT, INTAB, RGEOM, RGSAD, ROPT, DOREST, FCMEP, and FIOPEN are modified. Also, the IPOT values are modified as follows:

value	old meaning	new meaning
0	<i>hooks</i>	<i>hooks</i>
1	<i>mopac</i>	<i>unit30/unit40</i>
2	<i>aces</i>	<i>aces</i>
3	<i>gamess</i>	<i>unit29</i>

A new switch keyword, SUPERMOL (and its negation, NOSUPERMOL), has been added to the GENERAL section to indicate that the supermolecule mode is used for reactants and products. The default is on (SUPERMOL). The choice has no effect if there is only one reactant and one product. The following subroutines were modified: CENTER, DEFGEN, INTAB, NORMOD, POLYAT, REACT, REDGEO, RGENER, and SETLGS.

A new variable keyword, INTMU, has been added to the PATH section to indicate the number of points on each side of the saddle point used for the interpolation of the effective reduced mass at the saddle point. The available options are 1 and 3. Thus, values of 1 and 3 indicate that interpolation is based on 2 and 6 points, respectively. The default is 1. The following subroutines were modified: BCALC0, DEFPAT, ELRPH, INTAB, PATH, and RPATH.

A new variable keyword, INITGEO, has been added to the REACT1, REACT2, PROD1, PROD2, and START sections to indicate the location of the initial geometry. The available options are *geom* and *hooks*, the default being *geom*. The following subroutines were modified: DEFOPT, DOREST, FCMEP, FIOPEN, INTABS, POLYAT, REACT, RGEOM, RGSAD, and RSTAT.

- The switch keywords NONLINEAR and LINEAR and the variable keyword PHASE have been incorporated into the single variable keyword SPECIES. The default has been set to *nonlinrp*. The following subroutines were modified: DEFSTA, INTABS, REDGEO, RSTAT, RVAR6, and WVAR6.
- Subroutine REACT:
If the option *read* of the EZERO keyword in the potential section is used, only the geometries of the reactants are obtained. In the case of INITGEO = *hooks*, optimizations are required because the initial geometries are obtained through the OHOOK subroutine. Since MORATE-version 7.3 requires using INITGEO = *hooks*, the NOOPT option is not implemented with MORATE.
- A new keyword is added to the REACT1, REACT2, PROD1, PROD2, and START sections. The LINAXIS keyword is used to indicate that a molecule is linear and to specify its orientation. There are three options: *x-axis*, *y-axis*, and *z-axis*. The default option is *z-axis*. A new array, IOLIN, is added to the file common.inc. This array is used in subroutines DEFSTA, RSTAT, CENTER. RVAR6, INTABS and WVAR6.
- An option to print the distance matrix was added, and a new variable keyword, PRDISTMX, has been added to the PATH section to indicate the interval along the save grid for the printing of interatomic distance matrices. The default is to print the interatomic distance matrix at every 20th point along the save grid. The following subroutines were modified: DEFPAT, INTAB, NOROUT, and RPATH.
- In subroutine ZOCPAR:
The default value of the keyword RPL was calculated from the lower-level surface inversion 7.2; from version 7.3 on, the default is set to the value calculated from the higher-level calculations.
- An option to calculate the vibrational partition functions with vibrations treated

classically has been added, and a new switch keyword, CLASSVIB, and its negation, NOCLASSVIB, have been added to the GENERAL section to allow the user to choose between classical and quantal vibrations. The default is to use quantized vibrations. The following subroutines were modified: DEFGEN, INTAB, OPTION, and RGENER.

8. In subroutines ICFDIAG and PROFJ:
An unnecessary calculation of the **G** matrix has been removed in order to obtain better performance for the transformation from rectilinear to curvilinear internal coordinates.
9. In subroutines NORMOD and HHOOK:
Two GHOOK subroutine calls with an HHOOK call between them have been removed to eliminate unnecessary work. The HHOOK subprogram is modified to return the gradient at the same time as the Hessian, which is now the expected way for all HHOOK subprograms to be written.
10. The FORTRAN file units fu31-fu36 have been moved to fu71-fu76, and the FORTRAN file units fu80-fu89 have been reserved for use with interface codes.

Version 7.3.1 (July 1997) (Bug fixes)

1. The line “if (icode(jtype).lt.0) lgs(34) = 1” was missing from the interface.f subprogram in version 7.3. This only affects solid-state runs.
2. In the oh3fu30tr3 testrun the value of the SPECIES keyword for the saddle point in version 7.3 is *lints*, and this has been corrected to *nonlints*.

Version 7.4 (August 1997)

1. The option *curv3* is added by making modifications to the file common.inc, and to the OPTION, SETLGS, VECCON, CVCOOR, and READINT subroutines, plus adding the new subroutines BIMAT2, MTINV2, GFDIA2, SVBKS, and SVDCMP. This option is designed for the transformation from Cartesian coordinates to redundant internal coordinates for frequency evaluations at generalized transition states. The maximum number of internal coordinates is $3N + 6$. This new option has been used in the test runs ch5j1tr4, ch5fu30tr6, ch5fu40tr6, and ch5j1tr4. Furthermore the following test runs have been changed to use the *curv1* option: clhbrtr1, ho2tr2, and h3tr3.
2. The unit fu61 file has been created to save the stationary point information so it can be used in a subsequent run, if desired. This effectively creates a new restart capability, as discussed in the section on restart options. After the calculations are done on a given reactant or product, the geometry, energy, and frequencies of the reactant or product are written to the unit fu61. Similarly, for a saddle point, the geometry, energy, frequencies, Hessian, and eigenvectors are written. A new variable keyword, STATUS, is added in the *REACT1, *REACT2, *PROD1, *PROD2, and *START sections with several options.
Other keywords added to support this new option are ENERGY, FREQ, and EIGENVECTOR. The subroutines modified are DEFSTA, INTABG, INTABS, OPTION, RSTAT, RVAR6, STVARJ, WVAR6, DOREPR, FCRATE, FDIAG, FIOPEN, DOSAFR, NORMOD, POLYAT, REACT, SADDLE, and SADENG. A new subroutine WRTHOK is added. The keyword option OPT is removed, and FREQ is changed to a list keyword.

The subroutines SETLGS, DORPER, and MAIN have been modified to allow *Polyrate* to stop correctly after calculating one or more stationary points.

3. The keywords ICA, ICL, and ICR are replaced by options for the new keyword option FREQ in the ICOPT list. A new option, *none*, is added for no frequency corrections in the VTST-IC method. Modifications were made in the ZOCAST, ZOCPAR, ZOC3P, INTAB, and RICOPT subroutines.
4. In the file hooks.f:
Duplicate copies of the subroutines DERIV2, DERV24, NEWT, SECCEP, SECCEN, and TRANS were renamed YDERV2, YDER24, YNEWT, YSECEP, YSECCEN, and YTRANS respectively, and they were added to the hooks.f file to increase modularity.
5. The default for the FREQMAT keyword in the unit fu50 file has been modified for all cases. From now on, the default is to match the frequencies of a unimolecular reaction with wells in descending order. Modifications were made in the OPT50 and ZOC3P subroutines.
6. The switch keywords PRSAVEMODE and PRSAVERP have been changed into variable keywords with default values set to 9999. This means that after every 9999 save grid points, the information on the save grid will be printed instead of after every save grid point. Modifications were made in the file common.inc and in the NORMOD, RPATH, INTAB, DEFPAT, and CVCOOR subroutines.
7. The two extra gradients given in the unit fu30 files for computing the curvature were placed before the Hessian matrix in previous versions of *Polyrate*, but have been moved after the Hessian matrix in version 7.4. Modifications were made to the RPHRD2 subroutine.

Also, in previous versions of *Polyrate* the two extra gradients were always printed out in mass-scaled coordinates. From now on, the two extra gradients will be printed out in the same coordinates as the gradient. The subroutines PATH, RPHDXN, RPHWRT, and RPHRD2 have been modified to allow for this change.

8. The subroutine FIOPEN has been modified to allow *Polyrate* to write the unit fu25, fu26, fu27, and fu30 files at the same time. This modification allows the PRPATH and WRITEFU30 options to be used together.
9. When PRPATH is used, the numerical values of the CD-SCSAG effective reduced mass and of kappa and the determinant of the moment of inertia tensor are written to the unit fu25 file. The following subroutines have been modified: MEPINV, MEPSRT, PATH, SSAVE, BCALC, COLSHF, EXTRAP, and TSRATE.
10. The SGI Power Challenge option in the set_machine.jc script has been divided into two options, one for the SGI R8000 chip set and one for the SGI R10000 chip set.
11. New scripts have been added to support the Linux operating system.

Version 7.4.1 (September 1997) (Bug fix)

1. In files polyrr.f, poly40.f, and common.inc:
When using input units fu30 or fu40 with hindered rotor anharmonicity, the reduced moment of inertia for hindered rotations along the reaction path was set to a fixed value for all the points along the path. Now, the value is interpolated using

Lagrangian interpolation and extrapolated using exponential functions in the same fashion as the moment of inertia. The following subroutines have been modified: GENSEC40, IRFGT0, RPHFIT, RPHINTM, RPHRD2, RPHSET, and SHFTSAD.

Version 7.4.2 (October 1997) (Bug fix)

1. Subroutine ZOPAR and ZOCPRN:
An error in the ICL algorithm has been corrected. The error involved using the energy ratio of frequencies (0 instead of 1) for the case where both the high- and lower-level frequencies go to zero at reactants or products.

Version 7.4.3 (October 1997) (Bug fix)

1. Subroutine ROPTOP:
The dummy argument ISTRT in the subroutine's formal-argument-list has been misspelled, ISTRTI, since version 7.2.

Version 7.4.4 (December 1997) (Bug fixes)

When the geometries in files fu30 and fu40 were given in mass-scaled coordinates, the saddle point geometry, given in file fu5 in unscaled coordinates, was not mass-scaled when it was used with units fu30 and fu40 and therefore, the saddle point geometry was incorrectly used as mass-scaled. This leads to the wrong moment of inertia and the wrong TST rates. The subroutines RPHRD2 and SADSEC40 have been modified to fix this bug.

1. When using the WRITEFU30 option with LOPT(6) = 3 or 4, the extra gradient before the Hessian point was being printed out incorrectly. The extra gradient was un-normalized using the wrong norm. This has been corrected by modifying the PATH subroutine.

Version 7.8 (December 1997)

1. The IVTST-M algorithm has been implemented. For this purpose, the module ivtstm.f has been added, the variables N3S31 and NPT31 have been added into the percon.inc INCLUDE file, and several variables or arrays have been added to the common.inc INCLUDE file. The subprograms added or modified are SPL31, SPLM31, SPLV31, MUCDSC, PSAG, IMOM40, SADSEC40, RPHINT, and RPHRD2.
2. A new input unit, named fu31, has been added. The keyword WRITEFU31 in the GENERAL section and the option *unit31* for the POTENTIAL keyword in the ENERGETICS section of the fu5 input file have been added. The module poly31.f has been added for this change, and the INCLUDE files param.inc, percon.inc and common.inc and the PATH, DOPNM, DOSAFR, FCMEP, FIOPEN, POLYAT, REACT, DEFGEN, INTAB, RLINE, RGENER, RPOTET, SETLGS, RPHINT, and RPHWRT subprograms have been modified.

3. The original extrapolation algorithm for the information in the input units fu30 and fu40 has been removed. The subprograms EXTRP40, RD40EXTR, RD40GEN, RPH40, and RPHRD1 have been modified to reflect these changes.
4. The code has been modified in order to calculate wells in the reactant and/or product channels. The sections WELLR and WELLP have been added in the unit fu5 and WR40 and WP40 in unit fu40. Several arrays in the INCLUDE file common.inc and the variable NARRS in percon.inc have been modified, as well as the subprograms WRTHOK, PATH, DOREPR, DORPH, REACT, DEFT40, EXTRP40, IMOM40, IRFGT0, PTSEC40, RD40EXTR, RD40RP, RPH40, RPSEC40, SADSEC40, DEFSTA, INTAB, INTABG, INTABS, OPTION, READ5, REDGEO, RGEOM, RSTAT, RVAR6, RVARY, SETANH, WVAR6, RPHINT, RPHRD1, RPHRD2, RPHSET, and RPHWRT. The subprogram WRTWEL has been added.
5. The switch keyword RODS has been added to the PATH section of file fu5 to determine if the RODS algorithm is used. The DEFPAT, DOSAFR, INTAB, NORMOD, NOST31, PATH, RPATH, RPHRD2, RVAR6, and WVAR6 subroutines have been modified, the IRODS, VTT, and XNOU variables have been added to common.inc, and a new subroutine RRPM has been added to interface.f.
6. The keywords INM and SSAVE have been removed from the PATH section of file fu5. This effectively makes the Hessian grid and the save grid the same. Therefore, when *Polyrate* calculates the reaction path, the Hessian is saved whenever it is calculated. The Hessian grid step size is now controlled by the INH keyword in the path section. The DEFPAT, ELRPH, INTAB, IVTST0, NEXTPT, OPTION, PATH, QRSENE, RESTOR, RESTRT, and RPHWRT subroutines have been modified accordingly.
7. The CALCS keyword has been added to the path section of file fu5. This new keyword determines how the value of s is calculated when using the RODS algorithm. The INTAB, RPATH, RVAR6, and WVAR6 subroutines have been modified, and the ICALCS variable has been added to common.inc.
8. A new keyword CUS has been added into the RATE section for evaluating the reaction rate constants with the canonical unified statistical model. The file COMMON.INC and the following subroutines were changed: RATE, FINOUT, DEFRAT, INTAB, and RRATE. In addition, three new subroutines were added: CUSSPL, ISWAP, and DPSWAP.
9. The keyword GEOMUNIT has been introduced in the GENERAL section of the fu5 input file to specify the units for the geometries of the stationary points read from the fu5 input file. The DEFGEN, INTAB, RGENER, RGEOM, and RGSAD subroutines have been modified and the RRGU5 subprogram has been added. The IGEOU5 variable has been added to the common.inc file.
10. The subroutine TABLE has been modified in order to print the ZCT results in the fu15 output file.
11. The subroutine RPHWRT has been modified in order to write a cleaner fu30 input file.
12. Subroutine CENTER:
This subroutine has been modified to clarify the output and reduce the amount of the output. The POLYAT, SADENG, IMOM40, and RPHRD2 subroutines have also been modified for this purpose.

13. The frequency index for files fu30 and fu40 has been changed so that index 1 corresponds to the largest frequency at the saddle point, index 2 to the second largest frequency, and so on. The subroutines DEFLT40, RD40RP, and RPHRD2 have been modified accordingly. In addition, the subroutine RPHWRT has been modified so that when the writefu30 keyword is specified the generated unit fu30 file is consistent with this frequency numbering.
14. The following test runs have been added: ch5fu31tr1, ch5icfu31tr1, oh3tr9, oh3tr10, oh3fu30tr6, oh3fu31tr1, oh3fu40tr6, and ch4otr1.
15. A large number of minor corrections and clarifications have been made in the manual.

Version 7.8.1 (February 1998) (Bug fixes)

1. When using the INTMUEFF option with IVTST-M, the values of α_{eff} at the first point on the reactant side were incorrectly stored as the values of α_{eff} at the first point on the product side and vice versa. This has been corrected in subroutine ARMUEF.
2. The moment of inertia and the reduced moment of inertia were both incorrectly interpolated asymptotically to zero instead of asymptotically to infinity when using IVTST-M. The subroutines SPLRMI, SPLNM, and SPLM31 have been modified to reflect this correction.
3. The reduced moment of inertia of the saddle point was incorrectly stored as the product well when using unit fu30. This has been corrected in subroutine RPHRD2.
4. In runs which WRITEFU30 or WRITEFU31 is selected, along with the option for printing the Hessian in unscaled coordinates, the unscaled Hessian was correctly written out but then it was incorrectly substituted for the mass-scaled Hessian in the rest of the run. Subroutine RPHWRT has been corrected to eliminate this error.
5. In the subroutine MEPINV the variable SBKAP was incorrectly specified as SBAP. This typo caused the wrong curvature to be printed to the fu26 file.
6. The check_test.jc script has been modified to avoid checking both ch4o and ch4oh test run files when the ch4o test run files are to be checked.
7. The compiling script comp_src.linux has been corrected to compile the dateclock.c subroutine.
8. In subroutine RPHRD1, a change was made in the calculation of the reaction-path curvature at the points adjacent to the saddle point. This was later found to be incorrect and was removed in version 7.9.3.

Version 7.8.2 (March 1998) (Bug fixes)

1. The subroutine NORMOD has been modified to fix a run-time error which occurs on SUN workstations.
2. The hindered rotor option has been enabled for the reactants, products, and wells. The subroutines modified are SETANH, NORMOD, REACT, HINDRT, and ANHARM and new arrays were added to common.inc.

3. When the hindered rotor approximation was used with unit fu31 the transformation from Cartesian coordinates to mass-scaled coordinates was incorrect. As a consequence, the reduced moment of inertia of the saddle point was miscalculated. The subroutines RPNT31, STAT31, and NOST31 have been modified to account for this correction.
4. The subroutines IRFGT0, LOPT40, NOST31, PATH, and RPHRD2 have been modified to allow SCT tunneling approximations to be calculated when using the RODS algorithm with units fu30, fu31, and fu40, and the hooks.

Version 7.9 (March 1998)

1. A new keyword LOWFREQ has been added to the PATH section. The subroutines RLOWFR and REPFL have been added and the subroutine NORMOD has been modified to account for this new option.
2. To remove conflicts with AMSOL, the subroutines INVRT1 and OVERLP have been changed to INVRT and OVRLP, respectively.
3. The CALCS keyword has been removed from the PATH section because that option is not recommended. The subroutines DORODS, DEFPAT, INTAB, RPATH, RVAR6, and WVAR6 have been modified accordingly.

Version 7.9.1 (April 1998) (Bug fix)

1. Some extraneous print statements have been removed from the PATH subroutine.

Version 7.9.2 (June 1998) (Bug fixes)

1. A typo in the subroutine ROPTOP has been fixed. This typo caused an error in the OPTMIN and OPTTS keywords in the OPTIMIZATION section.
2. Subroutines ZOCPAR and ZOCPRN: Modifications were made to correct errors in the frequency interpolation with the *icl* option.
3. The bug introduced in item 8 of Version 7.8.1 was removed. Now the curvature at the saddle point is correctly calculated by eq. 98 of the GTST book chapter. Notice that even though the curvature at the saddle point itself is obtained by interpolation (see discussion of the INTMU keyword), when INTMU = 1 the imaginary frequency eigenvector at the saddle point is used to obtain the curvature at the points on the save grid or either side of the saddle point. Furthermore the code has been modified so that this calculation is carried out correctly when fu31 is used (which was not true before) as well as when the hooks or other electronic structure input file options are used. Subroutines STAT31, NOST31, RPHRD1, and SHTSAD were modified.

Version 8.0 (August 1998)

1. The mapped interpolation (MI) algorithm, which was added in version 7.8 for use with files fu30, fu31, and fu40, now also works with the hooks. The mapped interpolation option has been used to replace the two-point exponential extrapolation for the $V_a^G(s)$ and $\mu_{eff}(s)$ functions for reactions with a saddle point. For reactions without a saddle point the original two-point extrapolation is still used. Modifications were made to the subroutines ZOCUPD, BCALC, ELRPH, MEPSRT, and PATH. New subroutines IVTMH, IVTMA, IVTMD, and ARMUEF1 have been added. The keywords EXFIRST and EXSECOND in the PATH section are no longer used with the electronic structure input files fu30, fu31, and fu40, and their meaning is changed when they are used with the hooks. Keywords UNIMOL and LBEXP were moved from file fu31 to the PATH section of file unit fu5.
2. The keyword LBEXP has been moved from file fu31 and is now an option for the keyword IVTSTMOPT in the PATH section of file fu5. In addition, the UNIMOL keyword has been removed from file fu31, and its options are now options for the keyword IVTSTMOPT in the PATH section of file fu5. These changes were made to use mapped interpolation with the hooks.
3. In subroutine REACT:
A new array WEW is added to store the frequencies of the wells for interpolation. Energies of the reactant well and product well are stored in the variables EWR and EWP, respectively. The geometries of the stationary points are stored in the corresponding XR arrays. These changes were made to use MI with the hooks. In subroutine DEFPAT:
A new variable keyword INI has been added to indicate the interval of the gradient interpolation steps in the MI algorithm. A check has been added to the subroutine OPTION to make sure that INI " INH. These changes were made to use MI with the hooks.
4. The option *vrpe* has been added to the RPM keyword to allow the user to use the variational reaction path (VRP) algorithm based on the Euler steepest-descents (ESD) method to follow the reaction path. This algorithm is called VRP(ESD) or just VRP. The subroutines DEFPAT, DORODS, ENDRODS, INTAB, NORMOD, NOST31, PATH, RPHRD2, RRPM, RVAR6, and WVAR6 have been modified.
5. The IC keyword has been changed to an option of the DL keyword and is renamed *ioc*. The ICOPT keyword has been renamed ICOPT and the options *eckart* and *noeckart* have been changed to *deckart* and *seckart*, respectively. The default for interpolating $V_{MEP}(s)$ is changed from *deckart* to *seckart*. The *zero* option has been removed from the ICOPT keyword.
6. A new option for correcting energies along the reaction path by the dual-level single-point energies (ISPE) method has been added. The new option *ispe* is available under the DL keyword. The file unit fu51 has been reserved for use with the new *ispe* option. Note that the ISPE option uses the mapped interpolation algorithm and so the formulas differ from an IOC calculation in which only the energy is corrected. Furthermore an IOC calculation involves corrections based on an optimization at a higher level. The IC keyword has been changed to an option of

the DL keyword and is renamed *ioc*.

7. The option *coord* has been added to the PRPATH keyword in the PATH section to print the bond lengths and bond angles along the reaction path. A new subroutine PRCORD has been added and the subroutines INTAB, RPRMEP, NORMOD, FCRATE, FIOOPEN, and RPHRD2 have been modified.
8. The keywords SPT and SPTOPT have been removed, and the associated has been commented out.
9. The NONADIB keyword has been renamed LCTCOR, the option previously called *quadratic* has been renamed *quadlin*, and the default has been changed from *no* to *quadlin*.
10. The LOWFREQ keyword in the PATH section of the fu5 input file has been renamed IVTST0FREQ.
11. The default value for the INTMU keyword of the PATH section of the fu5 input file was changed from 1 to 3.
12. The parameter NSDIM has been moved from percon.inc to the param*.inc files.
13. The RPM keyword option *euler* has been renamed *esd*; subroutines DEFPAT, INTAB, and RRPM have been modified.
14. In subroutine MAIN:
Modifications were made to allow anharmonicity options to be used with TST-only calculations.
15. In function VPART:
The vibrational partition function is set to 10^{10} when using the WKB method with an imaginary frequency normal mode. This value was previously set to 1.
16. In subroutine NOROUT:
Modifications were made to print the ground-state vibrational energies for the saddle point to file unit fu6.
17. Modifications have been made to print 'VTST-IOC' and 'VTST-ISPE' to the output files instead of 'VTST/IC'.
18. In subroutines REACT and NOROUT:
Modifications were made to print 'Harmonic Frequencies' to the output files instead of 'Frequencies'.
19. The common block OUTPUT was renamed OUTCOM in all the subroutines contained in the givtst.f subprogram.
20. The subprograms SORT and UPCASE were renamed SORTG and UPCSE, respectively. In addition the function TORS was renamed PTORS.
21. The IBM PowerPC 604e option has been added to the set_machine.jc script and the necessary script files, *.ibm604, have been added to the script directory.
22. The test runs oh3fu31tr2, ch5fu51tr1, ch5fu50tr3, and cmtr3 have been added to the test suite.

23. The WRITEFU31 keyword has been added to the oh3tr10 test run.
24. The oh3tr1 test run has been modified to use the *vrpe* option of the RPM keyword.

Version 8.1 (July 1999)

1. A new geometry optimization option using the eigenvector following (EF) algorithm was added for the reactants, products, reactant well, product well, and saddle point. The FORTRAN source code is taken from Frank Jensen's program in the MOPAC5.07MN software package (this EF code is originally from Ref. 2 in Sect. 22) and stored in the file ef.f. The subroutines OHOOK, DEFOPT, ROPT, ROPTOP, and INTAB were modified, a new subroutine REFOPT was added, and the compilation scripts comp_src.* are modified to accomplish these changes.
2. A new option for calculating the reduced moment of inertia of an asymmetric rotator using Pitzer's curvilinear algorithm was added. The subroutines RTOR and RTOROPT were added to replace the RHINRT and RHINRTS subroutines in the interface.f file. Subroutines DEFSTAI, INTAB, INTABS, RSTAT, SETANH, ANHARM, CENTER, HRPART, RATE, RPHINT, and TSRATE were modified for this new option. Three new subroutines, HRSET, ASYMRMI and ASYMENT, were added. One can now estimate the effective vibrational frequency based on the rotational barrier height. A new testrun cwmctr3 is added to illustrate the curvilinear option. Also, the hindered rotor treatment has been generalized so that it no longer implicitly assumes that $P = 1$. The old list keyword HINROT was removed and is replaced by new list keywords TOR and TOROPT.
3. The subroutines ICFDIAG, ICSAVE, and CVCOOR were modified to carry out normal mode analysis using curvilinear internal coordinates (with or without redundancy) at the saddle point for the CURV2 and CURV3 options.
4. The PRMODE keyword was added to the REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and START sections for printing movie clips of the vibrational modes for the XMOL program. The subroutines NOROUT, INTABS, DEFSTA, and RSTAT were modified, and a new function DIGISTR was added.
5. The option DVMEP is added to file fu51 for use with the IVTST-ISPE option. By using this option, the point at the higher level doesn't need to be a save-grid point at the lower level. The subroutines READIC and ZSPMEP were modified.
6. The subroutine WRTHOK has been modified for printing out the vibrational frequencies and Hessians for calculations with the SOLID option.
7. In the previous versions of *POLYRATE*, the FREQ list keyword is used when a list of harmonic frequencies is read in as a restart option when the STATUS keyword has the value 6, and the NOFREQ option is used to switch off the normal mode analysis. In this version, the functions of the old keyword FREQ in the REACT1, REACT2, PROD1, PROD2, WELLR, WELLP, and START sections have been separated, and they are now associated with the keywords VIB and FREQ in order to clarify these two distinct functions. The VIB list keyword is used to indicate a list of harmonic

frequencies of the corresponding species when the `STATUS` keyword has the value of 6. The `FREQ` keyword is used as a switch keyword to determine whether the normal mode analysis is carried out in the calculation.

8. The subroutines `IVTMD`, `IVTMH`, and `RPHINT` were modified to support the mapped interpolation VTST algorithm for cases without information about either the product or a product well. The method used in such calls is fully explained in Section 6.M. of this manual.
9. The keyword `FIXMUEF` has been added to the `IVTSTMOPT` list keyword in the `PATH` section. This keyword is used to apply a global interpolation to estimate the effective reduced mass (μ_{eff}) of the SCT algorithm at the saddle point when the IVTST-M algorithm is used with hooks. The default option is to use local Lagrangian interpolation for the effective reduced mass at the saddle point with either one point or three points (see `INTMU` keyword option) on each side of the saddle point. The global interpolation can be used to obtain a smoother value of μ_{eff} at the saddle point; it should be used if and only if the points close to the saddle point suffer from numerical instability.
10. The keywords `RMITS`, `RMIRP`, and `HRRMI` were renamed to `MITS`, `MIRP`, and `HRMI` for input of the moment of inertia of the internal rotation mode in file `fu50` instead of the reduced moment of inertia as the previous versions. The subroutine `ROPT50` has been modified.
11. A check has been implemented in the subroutine `REMOVEC` to ensure that the total number of internal coordinates is greater or equal to the total number of vibrational modes.
12. The subroutine `WRTHOK` has been modified in order to print out correct energies for the reactants, products, wells, and saddle point when using the restart option controlled by the `STATUS` keyword.
13. The subroutine `SSAVE` has been modified to ensure that the array `BKAP` will contain the correct value of the curvature along the reaction path.
14. The `IVTST0FREQ` keyword option has been extended the case of unimolecular reactants and/or products. The subroutine `REPFL` has been modified and a new subroutine `rprfl` has been added to `polytr.f`. The test run `ch5j1tr7` is used as an example for this keyword option.
15. The compilation and linking scripts for IBM-SP with PowerPC 604e processor is upgraded to O3 with 64-bit computation.

Version 8.1.1 (July 1999) (Bug fixes)

1. The BLAS subroutines in the `ef.f` file were moved to the file `fromblas.f` in order to keep all the routines from the mathematical libraries in the same file.
2. The default values of the `SR` and `SP` options in the `IVTST0FREQ` keyword have been

set to -1.0 bohr and $+1.0$ bohr.

3. The keyword option PROJECT in the *START section has been corrected. In previous versions, the translational and rotational modes are always projected. We recommend to project out the translational and rotational modes in order to obtain better vibrational frequencies; however, this is not necessary. The subroutine CENTER has been modified.

Version 8.1.2 (August 1999) (Bug fix)

1. The *.jc scripts in the testrun directory were updated to work with the current version of *Polyrate*.

Version 8.2 (August 1999)

1. New keyword option BATH and its options FRICTION, BATHTEMP, DIFFUSION, and CUPLCNST are added to the PATH section to perform nonequilibrium solvation calculations. There are two limitations for using this new option, 1) the system must be molecule, not solid state (because the redundant internal coordinate system is used) and 2) the equilibrium free energy of solvation is included in the potential function. In this version, there are no test cases for carrying out nonequilibrium solvation calculations using *Polyrate* directly, however, new test cases are added to AMSOLRATE and GAMESOLRATE where the free energy of solvation is approximated using a continuum solvation model. Three new subroutines are added to *Polyrate*: RBATH, RDIFFU, and EFFBATH. The NEWT, SADENG, EHOOK, GHOOK, HHOOK, CVCOOR, IVTMH, DOSAFR, INITZE, NOROUT, PATH, PROJCT, VPART, WRTHOK, and ZOC3P subroutines are modified. Since BATHTEMP is a variable keyword, the nonequilibrium solvation calculation can only be carried out at one temperature.

Version 8.2.1 (November 1999) (Bug fix)

1. The function subprogram UPCASE has been renamed UPCSE. This should have been done in version 8.0 (see item 21 of the version 8.0 revision history).

Version 8.4 (December 1999)

1. The subroutines RATE and TSRATE have been modified to print out the internal free energy of reactant.
2. The nh3tr2 testrun has been created.

Version 8.4.1 (March 2000) (Bug fixes)

1. The file /testo/cwmctr3.fu15 has been replaced with the correct version for this testrun.

2. FORMAT statements were changed to prevent compiler warnings.

Version 8.4.2 (October 2000) (Bug fix)

1. The calculation of the partition function was incorrect when calculated off the saddle point when using curvilinear internal coordinates. The bug was due to the PROJCT subroutine being commented out in versions 8.1-8.4.1. This affects all runs using the options CURV2 and CURV3 in versions 8.1-8.4.1. CURV1 and rectilinear-vibration runs were unaffected by this bug.

Version 8.5 (October 2000)

1. The LCG4 option has been added. The implementation of the LCG4 method in *Polyrate* involved the modification of the following subroutines:
 - a. LCXINT: function that calculates the imaginary action integral at every quadrature point of the adiabatic region.
 - b. LCGTH (old LCG3TH): this subroutine calculates the imaginary action integral between the outer turning points of the vibrational adiabatic potential. Part of the code that was previously in this subroutine has been incorporated into two new subroutines called LCNAD and LCVEF.
 - c. LCNAD: returns the value of the imaginary action integral at every quadrature point of the nonadiabatic region.
 - d. LCVEF: function that calculates the effective potential of the nonadiabatic region.

The output of units fu41–fu44 was modified in the following ways (see page 250 of the *Polyrate*–version 8.4.1 manual) and two new units (fu48 and fu49) were added:

- a. poly.fu41: contains the theta integral and the weighting factor of each point on each tunneling path.
- b. poly.fu42: contains the length of the tunneling path, the classical energy, the potential correction terms (zero-point energy and anharmonic), the interpolated correction if any, and the effective potential for each point in the nonadiabatic region of the tunneling path.
- c. poly.fu43: contains the theta integrals and the boundaries of the nonadiabatic region for each tunneling path.
- d. poly.fu44: contains the length of the tunneling path, the maximum of the effective potential, and its location on the path for each tunneling energy.
- e. poly.fu45-47: same as before.
- f. The unit poly.fu48 is used to restart a LCG calculation. It contains the grid number of a given tunneling energy, the grid number of a point along the

linear path and the value of the calculated potential in the nonadiabatic region (the expensive part of the LCG calculations).

- g. The unit poly.fu49 stores the information that can be read directly from unit poly.fu48 afterwards.
2. We added an option to scale the frequencies for CVT calculations with harmonic vibrations. This implementation involved changes in the main program, and the subroutines REACT, SADENG, ENROUT, SENOUT, NOROUT, RATE, TSRATE, TABLE, TSOUT, VPART, WRTWEL, and several subroutines in the interface.f block as well as changes in common.inc.
3. We added an option to compute energy levels by WKB-on-the-true-potential for excited-state calculations. This implementation involved changes in subroutines RATE, VPART, STAUUV, VTMUSN, and EVIB.
4. We added 6 new test runs, ohcltr1-4, ch5jltr9, and ch5jltr10.

Version 8.5.1 (November 2000) (Bug fix)

1. An error was made in the hindered rotor subroutine, HRSET, in version 8.5. This bug only affects version 8.5.

Version 8.6 (February 2001)

1. The curvilinear internal coordinate bound-mod analysis for reactants and products is added.
2. The scaling force constant keyword FCSCALE is added into PATH section.
3. The keyword SCLPT and CARTRP are added. SCLPT is used to specify the effect of force constant scaling. CARTRP is used to declare explicitly that the normal mode analysis for reactants and products is carried out in Cartesian coordinates.
4. The switch subkeyword RTPJAC is added to the list keyword LCTOPT to specify the MEP and large curvature representative tunneling termini in mass-scaled Jacobian coordinates are printed in fu24.

Version 8.6.1 (August 2001, Bug fixes)

1. The behavior of the keyword FREQSCALE is changed:

Previously the keyword FREQSCALE was used to scale all harmonic vibrational frequencies uniformly but only for partition function calculations. However, the scaled frequencies were not used in calculation of zero point energies or transmission coefficients. The usage of FREQSCALE has been corrected to remove this error. If FREQSCALE is used, the scaling will be applied to all stationary point frequencies and generalized normal mode frequencies for generalized transition states along the reaction path. Furthermore, the vibrationally adiabatic potential

energy curve V_a^G will be obtained by adding the scaled vibrational ZPE to V_{MEP} , and all tunneling calculations will be based on this new V_a^G .

The above changes involving `FREQSCALE` are implemented in subroutine `NORMOD` (in `energetics.f`) by scaling all frequencies right after they are obtained from force constant matrix diagonalizing routines `FDIAG`, `ICFDIAG`, or `ICFDRP`.

2. The bug is fixed for curvature calculations when `fu31` is supplied as electronic structure input file, and `RODS` keyword is used.

When `RODS` is specified, the reaction path curvature vector calculation should be based on the vector normal to the reoriented dividing surface optimized by `RODS` algorithm. It involves a derivative calculation of this vector normal to the dividing surface with respect to the reaction coordinate s . Before the numerical derivative routine is called, the above vectors should be converted to mass-scaled and should be normalized. However, the code in old version did not handle this correctly. We fixed this bug in the current version in the subroutine `NOST31` (`poly31.f`).

Version 8.7 (September, 2001)

1. The normal mode analysis of reactants and products is set to Cartesian coordinates as the default, i.e., `CARTRP` is set as default in `PATH` section. One can specify that the generalized normal mode analysis for reactants and products is applied in curvilinear coordinates by turning `CARTRP` off (`NOCARTRP`). Note that `NOCARTRP` must be used when the force constant scaling keyword `FCSCALE` is specified, and if incompatible options are chosen, a message is printed and the program stops.
2. The default for `EZUNIT` keyword in the `ENERGETICS` section had not been set in previous versions. This bug is fixed, and the default for `EZUNIT` is set to be `au` in the subroutine `DEFENG`.
3. The default setting of `INITGEO` keyword for reactants, products, and saddle points is moved from the subroutine `DEFOPT` to the appropriate subroutine `DEFSTA` because the keyword `INITGEO` is defined under `REACT1`, `REACT`, `PROD1`, `PROD2`, and `START` sections instead of `OPTIMIZATION` section.
4. In subroutine `INTAB`, the writing of `FPRINT` input option in `SECOND` section is taken from the `ghook` "if" statement, so that status of the `FPRINT` switch will always be printed out whether or not `HESSCAL` is set to `ghook` or `hhook`.

Version 8.7.1 (January 2002, Bug fix)

1. Array declarations in `rtpjac.f` and `intbsv3.f` are modified to be compatible with the '-static' compiling option in compiling scripts.

Version 8.7.2 (February 2002, Bug fixes)

1. A bug in subroutine `HRSET` was fixed that affected hindered rotor calculations. This bug only affects version later than 8.5.
2. A bug was fixed in subroutine `REACT`. The geometries for each species should be copied to a local array no matter the initial geometries were obtained from `fu5` or `hooks`. See comments in the source code for details.

3. A bug was fixed in subroutine DEFSTA. The initialization of array INITG(I) should be done from I=1 to 8 not only from I=1 to 5. See comments in the source code for details.

Version 8.7.3 (May 2002, Bug fixes)

1. The initialization of logical variable LUPD in subroutine EFMAIN that controls the use of an updated Hessian for an optimization was not done. This logical variable is now initialized to .TRUE.
2. The initialization of the variable IPRNT that controls some printing options in subroutine EFMAIN was not done. This variable is now passed through a common block.
3. A bug in subroutine KAPVA was causing the wrong header to be printed for LCG3 or LCG4 calculations when using the quantized-reactant-state tunneling method was fixed. This has now been fixed.
4. A bug was fixed in subroutine HRSET that calculates information needed for evaluating the hindered rotor partition function. A wrong index for a vibrational frequency mode had been used.
5. A compilation error with Linux has been fixed. This bug fix involved modifications in the TABLE, FINOUT, and KAPVA subroutines.

Version 8.9 (May 2002)

1. The ILCG option has been added. This is done though two new keywords, ILCT and NILCTPT, that have been added to the list keyword LCTOPT. The implementation of the ILCG method in *Polyrate* involved modifications of the following subroutines:
 - a. RLCT: two new variables, ILCGIT and NSPLIC, have been added. These two variables are also included in common.inc.
 - b. INTAB: the two new keywords, ILCT and NILCTPT, have been included and are now tabulated.
 - c. LCG34 and LCGTH: the modifications have been made to incorporate the interpolated LCG algorithm.
 - d. LCGIT: this is a new subroutine that chooses the points at which the ILCG interpolation is made.
 - e. LCVEF: the modifications allow performing the spline-under-tension interpolation.
 - f. SPL: the dimensionalities of arrays IPAS and SAVARR have been increased.
2. An option for determining the MEP limits based on energetics criteria has been added. This is done though the new keyword SPECSTOP. The implementation of this new feature required addition of a new subroutine, RSSTOP, and modifications to the following subroutines: DEFPAT, ELRPH, INTAB, OPTION, PATH, and

RPATH.

3. The generalized free energy of activation can now be calculated in subroutines RATE and TSRATE either through direct use of the vibrational partition functions or through the use of the logarithms of the vibrational partition functions. Using logarithms enables the calculation of the free energy of activation when the vibrational partition functions (or their product) are susceptible to underflow. The procedure used for the calculating generalized free energy of activation is determined by the GTLOG switch in the RATE section. The implementation of this feature required modifications to the following subroutines: DEFRAT, INTAB, RATE, RRATE, SETLGC, and TSRATE.
4. The tunneling contributions obtained the using quantized tunneling energy method are now calculated, in the BOLTZ subroutine, by using an exponential of an energy difference rather than a product of exponentials. In this way, the results are precisely the same as before, and the calculation is more stable.
5. The STATES subkeyword is introduced under the QRST keyword to limit the number of quantized reactant states that will be included in the tunneling calculation. The implementation of this feature required modifications to the following subroutines: DEFRAT, QRSENE, RQRST, SETLGS, and WKBENE.
6. The format of the output files is changed so that the results obtained with ZCT (MEPSAG) method are labeled as ZCT instead of MEPSAG, and the results obtained with SCT (CD-SCSAG) method are labeled as SCT instead of CD-SCSAG. The modified subroutines are BOLTZ, GIVTST, KAPA, KAPPAS, KAPVA, and TABLE.
7. The set_machine.jc script has been upgraded to include options for SGI Origin R12000 processors, for Sun UltraSparc workstations, and for Compaq AlphaStations.
8. New scripts have been added for SGI Origin R12000 processors, for Sun UltraSparc workstations, and for Compaq AlphaStations.
9. The script comp_src.linux was modified to allow compilation with the fort77 compiler in Mandrake Linux.
10. Two new timing routines have been added. One of them, dattim.compaq, is used for timings on Compaq workstations, and the other, dattim.f90, uses the DATE_AND_TIME routine that is the standard timing routine in Fortran 90.
11. Three new test runs, ch5j1tr11, clhbrtr3, and clhbrtr4, have been added, and two test runs, ch5j2tr and oh3tr8, have been slightly modified to include the SPECSTOP keyword. The running, checking, and cleaning scripts have been updated.
12. A few old variables that were not used anymore have been removed from the code.

Version 8.9.1 (May 2002, Bug fixes)

1. The default setting of STATES subkeyword in QRST keyword is moved from the subroutine DEFRAT to the appropriate subroutine DEFTUN because the keyword QRST is defined under the TUNNEL section instead of the RATE section.

2. The interpolation for determining the tunneling probability at certain energies was done wrongly when using quantized tunneling energy method with a low frequency mode for the reaction coordinate. This bug was fixed in subroutine BOLTZ.
3. A bug was fixed in subroutine QRSENE that was not allowing using quantized tunneling energy method with a given number of quantized states.

Version 9.0 (May 2002)

1. The keyword WRITE62 has been added to the GENERAL section of the unit fu5 input file to create an fu62 file that contains a subset of the stationary point information presented in fu61. The implementation of this feature required modifications to the following subroutines: DEFGEN, INTAB, RGENER, FIOPEN, FCRATE, and WRTHOK.
2. The STQVIB subkeyword is introduced under the QRST keyword to limit the number of quantized reactant vibrational states that will be included in the reactant vibrational partition function. The implementation of this feature required modifications to the following subroutines: DEFTUN, QRSENE, RQRST, KAPVA, and WKBENE.
3. The format of the long output file, fu6, has been slightly modified when using quantized tunneling energy method. Similarly, the format of the output file fu15 has been slightly modified for IVTST calculations.
4. Test run nh3tr2 has been slightly modified to include the GTLOG keyword.
5. A few common blocks have been renamed and are now ended in either COM or CM.
6. A few old variables that were not used anymore have been removed from the code.

Version 9.1 (July 2002)

1. The long output file, fu6, has been modified so it contains the rate constants with non-tunneling contributions (i.e., contributions from energies above the top of the effective barrier). The implementation of this feature required modifications to the following subroutines: BOLTZ, FINOUT, and KAPVA.
2. The default of the CARTRP switch keyword has been set to on. The option of changing the default based on the values of other keywords has been removed. The implementation of this feature required modifications to the following subroutines: DEFPAT, INTAB, and SETLGS.

Version 9.1.1 (not released)

1. One bug affecting the evaluation of the correction potential, V_{corr} , when tunneling occurs into final vibrational excited states was corrected.
2. The restart option of the LCT method, which under some conditions, gave wrong results for tunneling into vibrationally excited states was corrected.

Version 9.2 (July 2003)

1. Modification of subroutine LCG34.
2. The old LCGTH subroutine is now subroutine LCGNAD. It evaluates the LCT3 and LCT4 nonadiabatic regions and stores all the needed information to evaluate the tunneling amplitude.
3. The new subroutine LCGTHE evaluates the theta integral.
4. Old subroutine LCGNAD is now subroutine LCG4V. It searches for a LCT4 nonadiabatic region within a LCG3 adiabatic region.
5. Two common blocks GRDDX and GRDDX2 were added to store the information from the LCGNAD subroutine.
6. Several subroutines called by SPL2D, which incorporate the 2D spline under tension, were added. They are based in the code developed by R. Renka.
7. The rectangular grid needed by the 2D spline under tension is built in LCGRD and printed in unit fu42.
8. The LCGTR subroutine was added.
9. The LCVEF subroutine was slightly modified.
10. Keywords **ILCT1D**, **ILCT2D** and **LCTGRID** were added and the two keywords ILCT and NILCTPT were removed.
11. Subroutine RLCTGRD was created in order to activate the new keywords.
12. Two new test runs, clhbrtr5 and ch5j1tr12, were added.
13. The Linux version of *Polyrate* must be compiled with Portland Fortran 77.

Version 9.3 (November 2003)

1. Version 9.3 runs on Windows 95/98/Me/2000/XP.
2. Version 9.3 can be compiled with the efc compiler and run on the SGI Altix.

Version 9.3.1 (June 2005, Bug fixes)

1. Previous versions had errors in the calculation of the reduced moment of inertia for a torsion using the curvilinear scheme of Pitzer and Gwinn. Version 9.3.1 corrects the errors in the alpha matrix, and uses all right-handed coordinate systems. The CO and CW torsion schemes now work correctly.
2. Version 9.3.1 no longer prints out the table of torsion information before it is calculated, and the value labeled “Total Moment of inertia” is now correct.

3. The ISPE and FREQSCALE options are now compatible with fu31 files. The IVTST0FREQ option remains a known incompatibility.

Version 9.3.2 (not released)

1. Program will now compile and work properly with the gnu and ifort compilers. Since the gnu compiler is available in Linux and for most other operating systems, this greatly increases the portability.

Version 9.4 (May 2006)

1. The 'configure' script is added to automatically detect compilers, and a Makefile is made for easy compilation of Version 9.4.
2. Version 9.4 allows parallel processing using MPI. The code has been rearranged to accommodate this, where the file poly_mpi is used for multiple processors, and the file dummy_mpi is required for serial jobs. This is done in the Makefile.

Version 9.4.1 (June 2006)

1. The 'configure' script has been modified to only print the parallel options if an MPI compiler is found.

Version 9.4.2 (October 2006)

1. A frequency scaling bug has been corrected. Prior to Version 9.3.1, the FREQSCALE option was not compatible with the fu30, fu31, and fu40 potential files. In Versions 9.3.1-9.4.1, the frequencies were inadvertently scaled twice for calculations that did not use a potential file. In Version 9.4.2, the FREQSCALE option works for all PES methods.
2. The torsion options have been made fully compatible for the reactant and products for a chemical reaction. The READI keyword has been created to read in the internal moment of inertia which is needed for calculations that do not calculate the vibrations of the species in the traditional manner (i.e. a restart calculation.)
3. The SSRP species is now compatible with the fu40 file.
4. A bug in the IVTST spline fitting procedure has been corrected.

Version 9.4.3 (November 2006)

1. The file dattim.gnu, which has inadvertently been removed from the 9.4.2 release, has been included in the 9.4.3 release.

Version 9.4.4 (not released)

1. Previous versions did not correctly evaluate torsions where multiple wells required updating as the MEP was calculated. This has been corrected.

Version 9.5 (January 2007)

1. The keyword SCALERP has been added, which causes the frequency scaling factor to be applied to frequencies found in the VIB section of the input file.

Version 9.5.1 (February 2007)

1. Frequencies calculated with internal coordinates are now scaled properly by the factor given by the FREQSCALE option. This was incorrectly implemented in previous versions.

Version 9.6 (February 2007)

1. Options have been added to the TOR and TOROPT sections, allowing for several interpolation methods to be used when calculating a hindered rotor method.
2. The keyword MUVTOPT has two new options “*fit gas13*” and “*fit gas31*” to do 1-point and 3-point fits instead of 3-point and 5-point fits.

Version 9.7 (June 2007)

1. Two new keywords has been added in the GENERAL section, INPUNIT and OUTUNIT. They control the units of geometries and distances in the fu5 input file and in the fu6 output file, respectively. In this version, the default unit is angstrom. The old keyword GEOMUNIT in the GENERAL section of the fu5 file is removed.

Version 2008 (May 2009)

1. The variable reaction coordinates method with either single-faceted or multifaceted dividing surfaces have been implemented in this version for barrierless association reaction. This implementation is parallelized using MPI.
2. The keywords, SIGMAF and SIGMAR, can accept floating point numbers as their values, while previous versions only allow integer values.
3. A new keyword, TENSION, is added to allow the user to give a tension factor for spline-under-tension interpolation in IVTST-M calculations.
4. The default value of the keyword VEFRCH is changed to 999 from 0.
5. In file polyhl.F, subroutine LCGNAD, the following line is changed to remove

some confusion output when LCG3 is used.

Change:

```
WRITE(FU6,134) IE,ENER*CKCAL
```

to

```
IF(ILCT.GT.1) WRITE(FU6,134) IE,ENER*CKCAL
```

6. A bug is fixed for using VADAVG keyword. In file polyhl.F, the following line is changed in subroutine lcgnd.
DO 89 IZ = IZ1, IZ0
TERM1 = (IZ-IZ1)/(IZ0-IZ1)
(wrong)
TERM1 = (QDX(IZ) - QDX(IZ1))/(QDX(IZ0)-QDX(IZ1)) (correct)
7. A bug is fixed for FREQSCALE keyword when doing IVTSTM with fu31 file. In file poly31.F, the following line is added to subroutine nost31
if (ifqfac.eq.1) ws(is,i)=ws(is,i)*freqfac
8. In polymq.F file, a shifting constant is used to solve the overflow and underflow problem in the case that the imaginary action integral λ is too large.

Version 2008-A (not released as a separate version, but fully incorporated in version 2010)

1. A bug is fixed for Monte Carlo error estimation in VRC-VTST calculations.
2. The VRC-VTST output format is reorganized in a more user-friendly way.
3. A bug is fixed so that the WELLR and WELLP energies are written to the fu31 file using relative energies. The old code wrote their absolute electronic energies to the fu31 file.
4. A bug is fixed so that a torsion treatment can be performed in conventional TST rate calculation.
5. A bug is fixed in poly31.F file for initializing the arrays WS, BFS, and FMIRS in subroutine rph31.
6. A bug was fixed for calculating the reduced momentum of inertia using C scheme in subroutine asyment.

Version 2010 (June 2010) (These revisions were made by J. Zheng and D. G. Truhlar)

1. One new keyword "FREQINCR" is added to raise low frequencies to a certain number (e.g., 50 or 100 cm^{-1}) to approximately account for anharmonicity.
2. In rate calculations for barrierless association reactions by VRC-VTST, the reverse reaction (dissociation) rate is also calculated by the equilibrium constant and the association rate. Previously only the association rate was calculated.
3. The SPRNG 2.0 package is included in the *Polyrate* distribution. The *configure*

script will compile the SPRNG 2.0 library if the VRC-VTST installation is performed.

4. In VRC-VTST calculations, in calculating the total number of available states, the integration over the projection on the principal axis (K) of the total angular momentum \mathbf{J} is calculated analytically.
5. Several methods for treating torsions, in particular the RPG, AS, SRPG, and SAS methods, are implemented. The input format and options for the keyword “TOR” and “TOROPT” are reorganized.
6. One test run directory h3o2 is added to show the usage of the SPRG, RPG, AS, and SAS methods for torsion treatment.

Version 2010-A (June 2010) (These revisions were made by J. Zheng and D. G. Truhlar)

1. A bug is fixed for SRPG and SAS method for torsional treatment. This bug only affects the cases in which the total symmetry number is not 1.

Version 2015 (January 2015) (These revisions were made by J. Zheng and D. G. Truhlar)

Important note: This version introduced bugs in CURV2 and CURV3. These bugs have been fixed in version 2016. Users with this version should upgrade to version 2016 or later versions. Users with versions prior to 2015 are not affected by this bug.

1. All common blocks are replaced by Fortran 90 modules. Memory for arrays is dynamically allocated. Therefore, compiling the code with a predefined system size (e.g., number of atoms) is no longer required.
2. The number of states calculation that is required for μ VT is now programmed much more efficiently by the extended Beyer-Schwinehart method. Now there is no significant difference between CVT and μ VT as far as computational cost.
3. The ICVT option is removed. Whenever one needs a more accurate treatment of the threshold, one can simply use μ VT without significantly increasing the computational cost, so there is no longer any motivation for using ICVT.
4. The *single-structure* MS-T method is added with coupled torsional potentials. This capability is added both for saddle point and for including torsional anharmonicity along a reaction path.
5. The standard-state free energy of activation profile is calculated along a reaction path using a standard state of 1 bar rather than an unconventional pressure (that is 1 atm).
6. Various minor bugs have been fixed.

Version 2016 (September 2016) (These revisions were made by J. Zheng, R. Meana-Pañeda, A. Ratkiewicz, and D. G. Truhlar)

1. All common blocks are replaced by Fortran 90 modules. Memory for arrays is dynamically allocated. Therefore, compiling the code with a predefined system size (e.g., number of atoms) is no longer required.

2. The number of states calculation that is required for μ VT is now programmed much more efficiently by the extended Beyer-Schwinehart method. Now there is no significant difference between CVT and μ VT as far as computational cost.
3. The ICVT option is removed. Whenever one needs a more accurate treatment of the threshold, one can simply use μ VT without significantly increasing the computational cost, so there is no longer any motivation for using ICVT.
4. The *single-structure* MS-T method is added with coupled torsional potentials. This capability is added both for stationary points and for including torsional anharmonicity along a reaction path.
5. The standard-state free energy of activation profile is calculated along a reaction path using a standard state of 1 bar rather than an unconventional pressure (that is 1 atm).
6. Various minor bug fixes have been made.

Version 2016-2A (Nov. 2016) (Revisions were made by J. L. Bao and D. G. Truhlar)

1. The *SS-QRRK* code for pressure-dependent rate constants has been included as a utility code.

Version 2017 (Aug. 2017) (Revisions were made by J. L. Bao and D. G. Truhlar)

1. A bug in the CUS option has been fixed.
2. A bug in the QRST option has been fixed. This bug causes a change (especially at high temperatures) in the TST and CVT rate constants when QRST option is turned on.
3. The configure script has been modified in order to fix a problem that affected the installation of VRC version of the *Polyrate*.

Version 2017-C (Jun. 2018) (Revisions were made by J. L. Bao)

1. A bug in the dimensionality in some arrays in *alloc.f90* has been fixed. This bug affects calculations with using more than $3N-6$ redundant internal coordinates with CURV3 option turned on.

25.B. ASCII manual revision history

From version 1.6 through version 5.1, the manual was distributed as an ASCII text file.

25.C. PostScript and PDF manuals

The manuals for versions 6.0 to 7.2 were produced with Microsoft Word 5.0 or 5.1a for the Macintosh and distributed as a PostScript file. The manuals for versions 7.3 to 8.4.1 were produced with Microsoft Word 7.0 for Windows and distributed in PostScript and Portable Document Format (PDF) formats. Beginning with version 8.5 the manual is distributed only in Portable Document Format (PDF) format.

25.D. Incompatibilities of *Polyrate*—version 2017 with previous versions

1. In version 3.9, the default SCT tunneling method was changed from SCSAG to CD-SCSAG. Thus, $LGS(9) \geq 1$ in all previous versions [and $LGS(9) = -2$ in versions 3.0-3.07] requested the SCSAG method, but from version 3.9 to 5.1.1, these options request CD-SCSAG. In versions 6.0 and later, where this kind of choice is controlled by keywords, the CD-SCSAG method remains the default. In fact the original SCSAG method was removed in version 5.0.
2. In the FORTRAN unit fu5 input data file, the input order for the normal modes and generalized normal modes in records 10A, 15, and 15B was changed in passing from version 4.5.1 to version 5.0. In versions 5.0 and later, the modes should be ordered from the highest to the lowest frequency, which is consistent with the universal convention in the spectroscopic literature.
3. The Quack-Troe method for interpolating energies along the reaction path was removed for versions 6.0 and later.
4. No FORTRAN unit fu5 input data files from versions earlier than 6.0 will work in *Polyrate* version 6.0 or later.
5. The input keyword CONVERGENCE in the OPTIMIZATION section was removed as of version 6.6.
6. In version 7.0 and later, variables *quad* and *page* for the keyword CURVE have been changed to *dgrad* and *dhess*, respectively. Furthermore, the PAGE keyword for activating the Page-McIver integrator has been changed to PAGEM.
7. In versions 7.0 and later, the saddle point geometry is no longer read from both the unit fu5 file and the unit fu30 file; it is only read from the unit fu5 file. Additionally, the DELTAH keyword has been removed from the unit fu29 input file.
8. In versions 7.0 and later, for the unit fu30 file, the meaning of $LOPT(3) > 1$ and $LOPT(4) > 2$ have been changed.

9. In versions 7.0 and later, the frequencies along the MEP are ordered in canonical order by default, whereas previous versions ordered the frequencies according to overlap calculations.
 10. In versions 7.1 and later, when using the unit fu40 file, the default unit for frequencies is cm^{-1} , whereas in version 7.0, these frequencies were in atomic units. Note: For the unit fu30 file, frequencies are still in atomic units, but the unit fu40 file has the full functionality of the unit fu30 file (except for LOPT(7) and LOPT(8) not equal to zero) and is more convenient, so we recommend using the unit fu40 file instead of the unit fu30 file.
 11. In versions 6.0-7.1, one could carry out calculations on a reaction without a saddle point simply by using the NOOPT keyword. Starting with version 7.1.1 one must use NOOPT, NOSADDLE, and FIRSTSTEP = *gradi*. The reason we changed this is so that the combination of NOOPT and SADDLE could be allowed.
 12. In version 7.2, the default values for the Interpolated Correction option (ICOPT) are set to ICL and ECKART, whereas previous versions used ICA and NOECKART. See also item 32 where the default is changed again.
 13. In version 7.2, the allowed values for the keyword POTENTIAL were changed; the option *analytic* is renamed *hooks*.
 14. In version 7.2, the keyword *Polyrate* in the SECOND section was changed to HESSCAL = *ghook*.
 15. Starting with version 7.2, the keywords *Polyrate*, MOPAC, GAUSSIAN, BFGS, and TSBFGS in the OPTIMIZATION section of the unit fu5 input file have been replaced by two keywords OPTMIN and OPTTS. GAUSSIAN and MOPAC are now selected using OPTMIN = *ohook* and OPTTS = *ohook*. The *Polyrate* and BFGS combination of keywords is now selected by setting OPTMIN = *bfgs*. *Polyrate* and NOBFGS are selected by setting OPTMIN = *nr*
 16. Starting with version 7.2, the values of the SIGMAF and SIGMAR keywords are no longer read from a restart file. The correct values must be read from the unit fu5 input file; otherwise a default value of 1.0 will be used.
- 9.D.D. Starting with version 7.2, the option *reactants* of the EZERO keyword of the ENERGETICS section of the unit fu5 input file was removed. *Polyrate* will automatically determine the zero of energy (denoted by the option *calculate*) unless the option *read* of the EZERO keyword is used.
- 9.D.D. In version 7.3, the meanings of the values of the variable IPOT have been modified as follows:

value	old meaning	new meaning
0	<i>hooks</i>	<i>hooks</i>
1	<i>mopac</i>	<i>unit30/unit40</i>
2	<i>aces</i>	<i>aces</i>
3	<i>gamess</i>	<i>unit29</i>

19. In subroutine ZOCPAR:
The default value of the keyword RPL was calculated from the lower-level surface in version 7.2; from version 7.3 on, the default is set to the value calculated from the higher-level calculations.

20. In version 7.3, units fu31-36 were renamed fu71-76.
21. In version 7.3, the switch keywords `NONLINEAR` and `LINEAR` and the variable keyword `PHASE` were incorporated into the single variable keyword `SPECIES`. The default has been set to *nonlinrp*.
22. In version 7.4, the keywords `ICA`, `ICL`, and `ICR` are replaced by options for the new keyword option `FREQ` in the `ICOPT` list.
 - 9.D.D. In version 7.4, the default of the `FREQMAT` keyword on unit fu50 has been modified for all cases. From now on, the frequencies of a unimolecular reaction or reactions with wells are matched in descending order by default.
 - 9.D.D. In version 7.4, the keywords `PRSAVERP` and `PRSAVEMODE` were changed from switch keywords to variable keywords with default values of 9999.
25. The two extra gradients given in the unit fu30 files for computing the curvature were placed before the Hessian matrix in previous versions of *Polyrate*, but have been moved after the Hessian matrix in version 7.4.
26. In version 7.8, the `INM` and `SSAVE` keywords have been removed. The save grid information is now saved every `INH`th step. Therefore, the Hessian grid and the save grid are now the same for all calculations.
27. In version 7.8, the `EULER`, `ES1`, and `PAGEM` keywords have been combined into options for the new `RPM` keyword.
28. In version 7.8, the default for the `INH` keyword has been changed from 1 in previous versions to 9.
29. In version 7.9, the `CALCS` keyword has been removed from the `PATH` section.
30. In version 8.0, the mapped interpolation (MI) algorithm, which was added in version 7.8 for use with fu30, fu31, and fu40, now also works with the hooks. The MI option has been used to replace the two-point exponential extrapolation for the $V_a^G(s)$ and $\mu^{\text{eff}}(s)$ functions for reactions with a saddle points. For reactions without a saddle point, the original two-point extrapolation is still used. The keywords `EXFIRST` and `EXSECOND` in the `PATH` section are no longer used with the electronic structure files fu30, fu31, and fu40. See also item 31.
31. In version 8.0, the keyword `LBEXP` has been moved from file fu31 and is now an option for the keyword `IVTSTMOPT` in the `PATH` section of file fu5. In addition, the `UNIMOL` keyword has been removed from file fu31, and its options are now options for the keyword `IVTSTMOPT` in the `PATH` section of file fu5. These changes were made to use MI with the hooks.

32. In version 8.0, the IC keyword has been changed to an option of the DL keyword and is renamed *ioc*. The ICOPT keyword has been renamed ICOPT and the options *eckart* and *noeckart* have been changed to *deckart* and *seckart*, respectively. The default for interpolating $V_{\text{MEP}}(s)$ is changed from *deckart* to *seckart*. The *zero* option has been removed from the ICOPT keyword.
33. In version 8.0, the keywords SPT and SPTOPT have been removed.
34. In version 8.0, the RPM keyword option *euler* has been renamed *esd*.
35. In version 8.0, the NONADIB keyword has been renamed LCTCOR, the *quadratic* option has been renamed *quadlin*, and the default has been changed from *no* to *quadlin*.
36. In version 8.0, the LOWFREQ keyword in the PATH section of the fu5 input file has been renamed IVTST0FREQ.
37. In version 8.0, the default value for the INTMU keyword of the PATH section of the fu5 input file has been changed from 1 to 3.
38. In version 8.0, the vibrational partition function has been set to 10^{10} when using the WKB method with an imaginary frequency normal mode. This value was previously set to 1.
39. In version 8.1, the keyword FREQ has been separated into two keywords, VIB and FREQ. The keyword VIB is used to specify a list of harmonic vibrational frequencies when STATUS is 6 for the reactants, products, wells, and saddle point. The keyword FREQ is used as a switch keyword to indicate whether normal mode analysis is to be carried out.
40. In version 8.1, the old hindered-rotor list keyword HINROT is removed; it is replaced by new list keywords TOR and TOROPT.
41. In version 8.1, the keywords RMITS, RMIRP, and HRRMI were renamed to MITS, MIRP, and HRMI to input the moment of inertia of the internal rotation mode instead of the reduced moment of inertia in a fu50 file. Notice that the difference between the reduced moment of inertia and the moment of inertia is that the reduced moment of inertia is divided by the square root of the symmetry number.
42. In version 8.1, the input format for the keyword IVTST0FREQ has been changed to include new options INDEX, SR, and SP. And the algorithm used for interpolating the vibrational mode also differs from the previous version. See section 10.M for a detailed explanation for the algorithm.
44. In version 8.1.1, the NOPROJECT keyword option in the START section has been fixed in order to work in the way that is described in the manual. In previous versions, the translational and rotational modes of the saddle point were always projected.
45. In version 9.7, the GEOMUNIT keyword in the GENERAL section of the fu5 is removed. Its function is replaced by new keyword INPUNIT, which is also in the GENERAL section of file fu5.

46. In version 2010, the input format and options are reorganized for the TOR and TOROPT keywords.
47. Beginning with versions 2015 and 2016, the ICVT option is removed. Note that version 2015 was removed from circulation due to a bug.

25.E. Program distribution history

25.E.1. Program libraries

Version number sent to	Date sent to library	Library or libraries	Program number
1.1	1987	CPC	ABBD
2.5	Dec. 1990	QCPE	601-version 2.5
4.0	Mar. 1992	QCPE	601-version 4.0
4.0.1	Mar. 1992	CPC	ACJC
4.0.1	Apr. 1992	QCPE	601-version 4.0.1
5.0	Apr. 1993	QCPE	601-version 5.0
5.0.1	May 1993	QCPE	601-version 5.0.1
6.5	Apr. 1995	QCPE	601-version 6.5
6.5	Apr. 1995	CPC	ADBN
6.6	Nov. 1995	QCPE	601-version 6.6
7.3.1	July 1997	QCPE	601-version 7.3.1

25.E.2. Anonymous ftp

Version number	Ftp site	Date available
6.5	SDSC	Mar. 1995
6.6	SDSC	Nov. 1995
7.0	SDSC	September 1996

25.E.3. World Wide Web

Version number	Web sites	Date available
7.0	U of M and SDSC	September 1996
7.1	U of M	February 1997
7.1.1	U of M	March 1997
7.2	U of M	April 1997
7.3	U of M	July 1997
7.3.1	U of M	July 1997
7.4	U of M	August 1997
7.4.1	U of M	September 1997
7.4.2	U of M	October 1997
7.4.3	U of M	October 1997
7.8	U of M	December 1997
7.8.1	U of M	February 1998
7.9	U of M	March 1998
7.9.1	U of M	April 1998
8.0	U of M	July 1998
8.1	U of M	July 1999
8.1.1	U of M	July 1999
8.1.2	U of M	August 1999
8.2	U of M	August 1999
8.2.1	U of M	November 1999
8.4	U of M	December 1999
8.4.1	U of M	March 2000
8.5	U of M	October 2000
8.5.1	U of M	November 2000
8.6	U of M	February 2001
8.6.1	U of M	August 2001
8.7	U of M	September 2001
8.7.1	U of M	January 2002
8.7.2	U of M	February 2002
8.7.3	U of M	May 2002
8.9	U of M	May 2002
8.9.1	U of M	May 2002
9.0	U of M	May 2002
9.1	U of M	July 2002
9.3	U of M	November 2003
9.3.1	U of M	June 2005
9.4	U of M	May 2006
9.4.1	U of M	June 2006
9.4.2	U of M	October 2006
9.4.3	U of M	November 2006
9.5	U of M	January 2007
9.6	U of M	February 2007
9.7	U of M	June 2007
2008	U of M	October 2008
2010	U of M	May 2010
2010-A	U of M	June 2010
2016	U of M	August 2016
2017	U of M	August 2017

The URL for the U of M website is <http://comp.chem.umn.edu/Polyrate>

25.E.4. Other

A few other versions have been distributed on a limited basis.

25.E.5. Current

Current availability is delineated in Section 3 on availability and licensing.

26. ACKNOWLEDGMENTS

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27. INDEX

0IVTST KEYWORD		
DEFINITION	279	
ABSORD SUBPROGRAM	436	
ACALC SUBPROGRAM	436	
ACESET SUBPROGRAM	436	
AITKEN SUBPROGRAM	436	
AITKF2 SUBPROGRAM	436	
AITKNF SUBPROGRAM	436	
ALFCT SUBPROGRAM	436	
ALLEXCIT KEYWORD		
DEFINITION	220	
TESTRUN EXAMPLE	330, 334, 367	
ANALYSIS KEYWORD		
DEFINITION	230	
ANCOEF SUBPROGRAM	436	
ANGL SUBPROGRAM	436	
ANGLCS SUBPROGRAM	436	
ANGLV SUBPROGRAM	436	
ANHARM SUBPROGRAM	73, 436	
Anharmonicity	50, 51, 100, 101, 244	
Anharmonicity keywords	170	
ANHDER SUBPROGRAM	436	
ANTLR KEYWORD		
DEFINITION	171	
ARMUEF SUBPROGRAM	436	
ARMUEF1 SUBPROGRAM	437	
ARRSM SUBPROGRAM	437	
ARRSV SUBPROGRAM	437	
ASYMALP SUBPROGRAM	437	
ASYMCAL SUBPROGRAM	437	
ASYMENT SUBPROGRAM	437	
ASYMNZD SUBPROGRAM	437	
ASYMRMI SUBPROGRAM	437	
ATOMS KEYWORD	82, 148, 295	
DEFINITION	151	
AVERJN SUBPROGRAM	437	
BANGLE1 SUBPROGRAM	437	
BARRI KEYWORD		
DEFINITION	240	
BATH KEYWORD		
DEFINITION	197	
BCALC SUBPROGRAM	437	
BCALCO SUBPROGRAM	437	
BCALCP SUBPROGRAM	437	
BIMAT SUBPROGRAM	438	
BMAT SUBPROGRAM	438	
BMAT1 SUBPROGRAM	438	
BMAT2 SUBPROGRAM	438	
BMAT5 SUBPROGRAM	438	
BMAT5X SUBPROGRAM	438	
BOLTZ SUBPROGRAM	438	
BOTHK KEYWORD		
DEFINITION	230	
BRENT SUBPROGRAM	439	
BRNULI SUBPROGRAM	439	
BTEN1 SUBPROGRAM	438	
BTEN2 SUBPROGRAM	438	
BTEN5 SUBPROGRAM	438	
BTEN5X SUBPROGRAM	438	
BTENS SUBPROGRAM	438	
BTENR SUBPROGRAM	438	
BTORSN SUBPROGRAM	437	
CALCS SUBPROGRAM	439	
CALCUB SUBPROGRAM	439	
Canonical order	49	
Canonical Variational Theory	16	
CARTRP KEYWORD		
DEFINITION	197	
CASE SUBPROGRAM	439	
CD-SCSAG	59	
CENTER SUBPROGRAM	73, 439	
CENTR1 SUBPROGRAM	439	
CENTRAL SUBPROGRAM	439	
CFLOAT SUBPROGRAM	439	
ch3tr1 test run	428	
ch4cltr1 test run	420	
ch4cltr2 test run	421	
ch4ohtr1 test run	416, 418, 419	
ch4ohtr2 test run	417	
ch4otr1 test run	422	
ch5fu29tr1 test run	342	
ch5fu29tr2 test run	343	
ch5fu30tr1 test run	344	
ch5fu30tr2 test run	345	
ch5fu30tr3 test run	346	
ch5fu30tr4 test run	347	
ch5fu30tr5 test run	348	
ch5fu30tr6 test run	349	
ch5fu31tr1 test run	350	
ch5fu40tr1 test run	351	
ch5fu40tr2 test run	352	
ch5fu40tr3 test run	353	
ch5fu40tr4 test run	354	
ch5fu40tr5 test run	355	
ch5fu40tr6 test run	357	
ch5fu50tr1 test run	358	
ch5fu50tr2 test run	359	
ch5fu50tr3 test run	360	
ch5fu51tr1 test run	361	
ch5icfu30tr1 test run	362	
ch5icfu30tr2 test run	363	
ch5icfu31tr1 test run	364	

ch5icfu40tr1 test run.....	365	Rectilinear.....	54
ch5icfu40tr2 test run.....	366	CORTRM SUBPROGRAM.....	440
ch5j1tr1 test run.....	324	COSINE SUBPROGRAM.....	440
ch5j1tr10 test run.....	334	CROSSPROD SUBPROGRAM.....	440
ch5j1tr11 test run.....	335	CRVVEC SUBPROGRAM.....	440
ch5j1tr2 test run.....	326	CUBIC SUBPROGRAM.....	440
ch5j1tr3 test run.....	327	CUBIC2 SUBPROGRAM.....	440
ch5j1tr4 test run.....	328	CUBST SUBPROGRAM.....	440
ch5j1tr5 test run.....	329	CURV KEYWORD	
ch5j1tr6 test run.....	330	DEFINITION.....	198
ch5j1tr7 test run.....	331	DGRAD OPTION.....	111
ch5j1tr8 test run.....	332	TESTRUN EXAMPLE.....	324, 329, 333, 367, 369,
ch5j1tr9 test run.....	333	377, 378	
ch5j2itr1 test run.....	340	DHESS OPTION.....	111
ch5j2itr2 test run.....	341	TESTRUN EXAMPLE.....	331
ch5j2tr1 test run.....	336	TESTRUN EXAMPLE.....	328
ch5j2tr2 test run.....	338	ONESIDE OPTION.....	111
ch5j2tr3 test run.....	339	TESTRUN EXAMPLE.....	391
CHAIN UTILITY SUBPROGRAMS.....	84	CUS KEYWORD	
CHARMMRATE.....	19	DEFINITION.....	230
CHECK KEYWORD		CUSSPL SUBPROGRAM.....	440
DEFINITION.....	151	CVCOOR SUBPROGRAM.....	440
CHKFRE SUBPROGRAM.....	439	CVT.....	39
CHKUP SUBPROGRAM.....	439	TESTRUN EXAMPLE.....	329
CLASS SUBPROGRAM.....	439	CVT KEYWORD	
CLASSVIB KEYWORD		DEFINITION.....	230
DEFINITION.....	151	TEST RUN EXAMPLE.....	324, 333
clhbrtr1 test run.....	367	TESTRUN EXAMPLE.....	328, 332
clhbrtr2 test run.....	369	CVT/MEPSAG.....	61
clhbrtr3 test run.....	370	cwmctr1 test run.....	376
clhbrtr4 test run.....	371	cwmctr2 test run.....	377
cmctr1 test run.....	373	cwmctr3 test run.....	378
cmctr2 test run.....	374	D parameter.....	171
cmctr3 test run.....	375	DATTIM SUBPROGRAM.....	441
CMS SUBPROGRAM.....	439	DAXPY SUBPROGRAM.....	441
COG SUBPROGRAM.....	440	DDOT SUBPROGRAM.....	443
COHBT1 SUBPROGRAM.....	440	DEFAULT SUBPROGRAM.....	441
COHBT2 SUBPROGRAM.....	440	DEFENG SUBPROGRAM.....	441
COLSHF SUBPROGRAM.....	440	DEFGEN SUBPROGRAM.....	441
comp_pot.jc.....	487	DEFLT40 SUBPROGRAM.....	441
comp_src.jc.....	487	DEFOPT SUBPROGRAM.....	441
COMT.....	61, 63	DEFPAT SUBPROGRAM.....	441
configure.....	486	DEFRAT SUBPROGRAM.....	441
CONOUT SUBPROGRAM.....	440	DEFSEC SUBPROGRAM.....	441
CONSTANT KEYWORD		DEFSTA SUBPROGRAM.....	441
DEFINITION.....	171	DEFTUN SUBPROGRAM.....	441
CONV40 SUBPROGRAM.....	440	DELEX1 KEYWORD	
Conventional Transition State Theory.....	16	DEFINITION.....	241
COORD KEYWORD		DELTA2 KEYWORD.....	53, 109
DEFINITION.....	197	DELTA2 PARAMETER.....	198
COORD UTILITY SUBPROGRAMS.....	83	DELTAE KEYWORD	
Coordinates		DEFINITION.....	240
Cartesian.....	54	DEMIN KEYWORD	
Curvilinear.....	54	DEFINITION.....	171

DERIV2 SUBPROGRAM	441	ECKART SUBPROGRAM.....	443
DERV SUBPROGRAM	441	ECKRT SUBPROGRAM	443
DERV24 SUBPROGRAM	441	EDGEOK KEYWORD	
DERVAG SUBPROGRAM.....	441	DEFINITION	231
DETM KEYWORD.....	477	EFFBATH SUBPROGRAM	443
DEFINITION	279	EFMAIN SUBPROGRAM	443
TESTRUN EXAMPLE.....	387	EFOPT KEYWORD	
detmi utility program.....	477	DEFINITION	164
DFGN31 SUBPROGRAM	442	EFORMD SUBPROGRAM	443
DGAMMA SUBPROGRAM	442	EFOVLP SUBPROGRAM	444
DGEDI SUBPROGRAM	442	EFPHES SUBPROGRAM.....	443
DGEFA SUBPROGRAM	442	EHOOK SUBPROGRAM	89, 443
DIATOM KEYWORD	73	EIGENVECTOR KEYWORD	
DEFINITION	172	DEFINITION	173
DIATOM SUBPROGRAM	73, 442	ELEC KEYWORD	
DIFFD KEYWORD.....	53, 109	DEFINITION	173
DIFFD PARAMETER	198	electronic structure input	17, 74, 402, 410
DIGISTR SUBPROGRAM	442	Electronic structure input file	
DIST SUBPROGRAM	442	anharmonicities	92
DL KEYWORD	112	data interpolation	95
DEFINITION	151	Dummy potential energy surface	82, 91
DLX2 KEYWORD		frequencies	92
DEFINITION	164	moment of inertia.....	92
DLX3 KEYWORD		unit fu30	91, 242
DEFINITION	198	unit fu40	91
DOGLEG SUBPROGRAM	442	user-supplied information	93
DOPNM SUBPROGRAM.....	442	ELLIP SUBPROGRAM.....	444
DOREPR SUBPROGRAM	442	ELRPH SUBPROGRAM	444
DOREST SUBPROGRAM.....	442	EMIN KEYWORD	
DORODS SUBPROGRAM	442	DEFINITION	220
DORPH SUBPROGRAM	442	ENDRODS SUBPROGRAM	444
DOSAFR SUBPROGRAM	443	ENDSLP SUBPROGRAM	444
DOSAGE SUBPROGRAM	443	ENERGETICS SECTION	156, 297
DOTPRD SUBPROGRAM	443	ENERGY KEYWORD	
DPMPAR SUBPROGRAM	442	DEFINITION	174
DPR123 SUBPROGRAM.....	443	ENERXN KEYWORD	
DPSWAP SUBPROGRAM	443	DEFINITION	279
DQQP KEYWORD		ENESAD KEYWORD	
DEFINITION	172	DEFINITION	279
RELATED KEYWORDS.....	178	ENORM SUBPROGRAM.....	444
DSCAL SUBPROGRAM.....	443	ENROUT SUBPROGRAM	444
DSWAP SUBPROGRAM	443	EPART SUBPROGRAM	444
dumpot.f	82, 91	EPARTI SUBPROGRAM	444
E1GEOM SECTION		Equilibrium constant.....	62
DEFINITION	239	ERF SUBPROGRAM.....	444
E1GRAD SECTION		ES1 KEYWORD	
DEFINITION	239	DIFFD PARAMETER	
E1HESS SECTION		TESTRUN EXAMPLE	332
DEFINITION	239	ES1 KEYWORD	
EACT KEYWORD		DELTA2 PARAMETER	
DEFINITION	231	TESTRUN EXAMPLE	332
TESTRUN EXAMPLE.....	324, 333	METHOD DESCRIPTION	109
EBND SUBPROGRAM	443	METHOD DESCRIPTION	53
ECKARS SUBPROGRAM	443	TESTRUN EXAMPLE	332, 336, 392

ESLOPT KEYWORD		FIRSTSTEP KEYWORD	110
DEFINITION	198	CUBIC OPTION	
ESD KEYWORD	73, 109	TESTRUN EXAMPLE	390, 392, 395
TESTRUN EXAMPLE	328, 339, 385	DEFINITION	202
Euler stabilization method	53	RELATED KEYWORD	198
Euler steepest-descents (ESD)method	52	FITMAX SUBPROGRAM	446
EVIB SUBPROGRAM	444	FITMX SUBPROGRAM	446
EWKB SUBPROGRAM	444	FIVFT SUBPROGRAM	446
EXCITED KEYWORD		FIVPT SUBPROGRAM	446
DEFINITION	220	FORMA SUBPROGRAM	446
EXFIRST KEYWORD	99	FORMBT SUBPROGRAM	446
DEFINITION	199	FORMF SUBPROGRAM	446
TESTRUN EXAMPLE	379, 390	FORMF2 SUBPROGRAM	446
Expert users	316, 317	FORMG SUBPROGRAM	446
EXPND SUBPROGRAM	444	FORWARDK KEYWORD	
EXSECOND KEYWORD	99	DEFINITION	231
DEFINITION	201	TESTRUN EXAMPLE	379, 380
TESTRUN EXAMPLE	379	FPRINT KEYWORD	
EXTRAP SUBPROGRAM	444	DEFINITION	160
EXTRP SUBPROGRAM	445	FREQ KEYWORD	
EXTRP40 SUBPROGRAM	444	DEFINITION	174
EZERO KEYWORD	80	FREQIMAG KEYWORD	
CALCULATE OPTION		DEFINITION	280
TESTRUN EXAMPLE	367, 390	FREQMAT KEYWORD	113
DEFINITION	157	DEFINITION	280
READ OPTION		TESTRUN EXAMPLE	374, 387
TESTRUN EXAMPLE	369	FREQSCALE KEYWORD	
EZUNIT KEYWORD		DEFINITION	202, 203
DEFINITION	157	FREQUNIT KEYWORD	
F1DI SUBPROGRAM	445	DEFINITION	174
F1DIM SUBPROGRAM	445	FRPRMN SUBPROGRAM	446
FACORD SUBPROGRAM	445	FRSORT SUBPROGRAM	446
FCINPT SUBPROGRAM	445	GAMESSPLUSRATE	19
FCMEP SUBPROGRAM	445	GAUSQD SUBPROGRAM	446
FCN SUBPROGRAM	445	GAUSSQ SUBPROGRAM	446
FCRATE SUBPROGRAM	445	GAUSSRATE	19
FCSCALE KEYWORD		GBSLVE SUBPROGRAM	447
DEFINITION	201	GBTQL2 SUBPROGRAM	447
FCSCL SUBPROGRAM	445	<i>GCOMP KEYWORD</i>	
FDIAG SUBPROGRAM	445	DEFINITION	164
FDIAG2 SUBPROGRAM	445	GEMM SUBPROGRAM	447
FDJAC1 SUBPROGRAM	445	GENERAL SECTION	148, 294
FICLSE SUBPROGRAM	445	Generalized transition state theory	39
File fu30 input		Canonical variational theory (CVT)	39, 42
lopt options	242	Microcanonical variational theory (μ VT)	39
File usage	490	GENSEC40 SUBPROGRAM	447
findb utility program	482	GEOM KEYWORD	80
testrun example	374	DEFINITION	174
FINDL SUBPROGRAM	445	Geometry optimization keywords	
findl utility program	482	<i>Polyrate</i>	163
testrun example	374	GETFRQ SUBPROGRAM	447
FINDLV SUBPROGRAM	445	GFDIAG SUBPROGRAM	447
FINOUT SUBPROGRAM	62, 75, 446	GHOOK SUBPROGRAM	89, 447
FIOPEN SUBPROGRAM	446	GIVTST SUBPROGRAM	447

GRAD SUBPROGRAM	447	ICINT SUBPROGRAM	448
GRADDR SUBPROGRAM	447	ICL KEYWORD	113
GSETA SUBPROGRAM	447	ICODAL SUBPROGRAM	448
GSETM SUBPROGRAM	447	ICOPOT KEYWORD	112
GSETP SUBPROGRAM	447	ICOUTRP SUBPROGRAM	448
GSPEC KEYWORD		ICR KEYWORD	113
DEFINITION	231	ICSAVE SUBPROGRAM	449
TESTRUN EXAMPLE	394	ICSVRP SUBPROGRAM	449
GTEMP KEYWORD		ICVT	
DEFINITION	231	TESTRUN EXAMPLE	328, 329, 339
TESTRUN EXAMPLE	391	ICVT KEYWORD	
GTLOG KEYWORD		TEST RUN EXAMPLE	324, 333
DEFINITION	232	IDAMAX SUBPROGRAM	449
GTST Book Chapter	37	IDIRECT KEYWORD	
h2nitr1 test run	415	DEFINITION	203
h3o2fu31tr1 test run	429, 430, 431, 432	ILCT	69
h3tr1 test run	379	IMODULE FILES	
h3tr2 test run	380	AAMOD.F90	475
h3tr3 test run	381	IMOM40 SUBPROGRAM	449
h3tr4 test run	382	INCLUDE FILES	435
HARMONIC KEYWORD		PARAM.INC	490
DEFINITION	175	INERTA SUBPROGRAM	449
HBT SUBPROGRAM	448	INH KEYWORD	73
HEADR SUBPROGRAM	448	DEFINITION	203
HESS KEYWORD		TESTRUN EXAMPLE	332, 379
DEFINITION	203	INI KEYWORD	
HESSCAL KEYWORD		DEFINITION	204
DEFINITION	160	INITGEO KEYWORD	
HESSIAN KEYWORD		DEFINITION	175
DEFINITION	175	INITZE SUBPROGRAM	449
HHOOK SUBPROGRAM	89, 448	INPUNIT KEYWORD	
HINDERED ROTOR ...	103, 106, 107, 179, 180, 183,	DEFINITION	152
280, 283, 377, 378, 419, 448, 479		Input	
HINDRT SUBPROGRAM	448	file fu29	238
HINDRT1 SUBPROGRAM	448	file fu30	91, 242
hnitr1 test run	414	file fu31	251
ho2tr1 test run	383	file fu40	260
ho2tr2 test run	384	file fu5	145
ho2tr3 test run	385	section descriptions	147, 294
ho2tr4 test run	386	file fu50	277
ho2tr5 test run	387	file fu51	288
HQSC SUBPROGRAM	448	INTAB SUBPROGRAM	449
HREC KEYWORD		INTABS SUBPROGRAM	450
DEFINITION	164	INTCOR SUBPROGRAM	449
HRMI KEYWORD	479	INTDEF KEYWORD	
DEFINITION	280	DEFINITION	204
hrotor utility program	479	INTEGR SUBPROGRAM	450
HRPART SUBPROGRAM	448	INTERPOLATED OPTIMIZED CORRECTIONS	
HRSET SUBPROGRAM	448	(VTST-IOC)	67, 112
HVAL SUBPROGRAM	448	INTERPOLATED VTST	63
HYBRD SUBPROGRAM	448	INTERSC SUBPROGRAM	449
HYBRD1 SUBPROGRAM	448	INTEUL SUBPROGRAM	450
ICFDIAG SUBPROGRAM	448	INTF31 SUBPROGRAM	449
ICFDRP SUBPROGRAM	448	INTFNC SUBPROGRAM	450

INTMU KEYWORD		
DEFINITION	204	
INTPM SUBPROGRAM	450	
INTPM2 SUBPROGRAM	450	
INTRPL SUBPROGRAM	450	
INTRVL SUBPROGRAM	449	
INVRT1 SUBPROGRAM	449	
INVRT2 SUBPROGRAM	449	
INVRT3 SUBPROGRAM	449	
IOCOPT KEYWORD		
DEFINITION	152	
TESTRUN EXAMPLE	374	
ZERO OPTION		
TESTRUN EXAMPLE	416, 418, 419	
IRCX SUBPROGRAM	450	
IRFTG0 SUBPROGRAM	450	
IRPH SUBPROGRAM	450	
IRPHZ SUBPROGRAM	450	
Isoinertial coordinates	49, 73	
ISWAP SUBPROGRAM	450	
ISWP KEYWORD		
DEFINITION	205	
ISWR KEYWORD		
DEFINITION	205	
IVTMA SUBPROGRAM	450	
IVTMD SUBPROGRAM	450	
IVTMH SUBPROGRAM	450	
IVTST SECTION		
DEFINITION	240	
IVTST0 KEYWORD	64	
DEFINITION	153	
IVTST0 SUBPROGRAM	450	
IVTST0FREQ KEYWORD		
DEFINITION	208	
INCOMPATABILITY	208	
IVTST1 KEYWORD	64	
DEFINITION	153	
IVTST-M	65, 119	
IVTSTMOPT KEYWORD		
DEFINITION	206	
JACSCF SUBPROGRAM	451	
JAGUARATE	19	
KAPA SUBPROGRAM	451	
KAPPAS SUBPROGRAM	451	
KAPVA SUBPROGRAM	75, 451	
KG1 SUBPROGRAM	451	
Large-curvature Tunneling	16, 46	
LCG3	46, 59	
LCG3 SUBPROGRAM	451	
LCG4	46, 59	
LCGIT SUBPROGRAM	451	
LCGSRT SUBPROGRAM	451	
LCGTH SUBPROGRAM	451	
LCNORM SUBPROGRAM	451	
LCNWTN SUBPROGRAM	451	
LCPATH SUBPROGRAM	451	
LCPROB SUBPROGRAM	452	
LCPROJ SUBPROGRAM	452	
LCSET SUBPROGRAM	452	
LCSINT SUBPROGRAM	452	
LCSTX SUBPROGRAM	452	
LCT3 KEYWORD		
DEFINITION	220	
LCT4	126	
LCT4 KEYWORD		
DEFINITION	220	
LCTCOR KEYWORD		
DEFINITION	281	
LCTDETAIL KEYWORD		
DEFINITION	221	
OUTPUT DETAILS	304	
LCTOPT KEYWORD		
DEFINITION	222	
INTERPOLATE OPTION		
TESTRUN EXAMPLE	369	
TESTRUN EXAMPLE	324, 329, 330, 333, 334	
LCTP SUBPROGRAM	452	
LCVCOR SUBPROGRAM	452	
LCVIB SUBPROGRAM	452	
LCVMOD SUBPROGRAM	452	
LCXINT SUBPROGRAM	452	
LCXTAU SUBPROGRAM	452	
LCZCRT SUBPROGRAM	452	
LIN SUBPROGRAM	453	
LIN2 SUBPROGRAM	453	
LINAXIS KEYWORD		
DEFINITION	176	
LININT SUBPROGRAM	453	
link_exe.jc	488	
LINMN SUBPROGRAM	453	
LINMR SUBPROGRAM	453	
LOCATE SUBPROGRAM	453	
LOCS SUBPROGRAM	453	
LOPT OPTIONS	242	
LOPT40 SUBPROGRAM	453	
LOWFREQ KEYWORD		
DEFINITION	281	
LSAME SUBPROGRAM	453	
LUBKSB SUBPROGRAM	453	
LUDCMP SUBPROGRAM	453	
MAIN SUBPROGRAM	453	
Manuals	36	
MATINV SUBPROGRAM	453, 454	
MATX SUBPROGRAM	453	
MC-TINKERATE	19	
MDMOVIE KEYWORD		
DEFINITION	153	
Memory usage	494	

MEPINV SUBPROGRAM.....	453	TESTRUN EXAMPLE	383, 384
MEPOUT SUBPROGRAM	453	MXLNEQ SUBPROGRAM.....	454
MEPSAG	46, 59	NATOMS VARIABLE	81
MEPSRT SUBPROGRAM.....	454	NEARW SUBPROGRAM.....	454
MEPTYEP KEYWORD	112	NEWT SUBPROGRAM	73, 454
DEFINITION	282	NEXTPT KEYWORD	
TESTRUN EXAMPLE.....	374, 387	DEFINITION	209
MEPTYPER KEYWORD	112	NEXTPT SUBPROGRAM	454
DEFINITION	282	nh3tr1 test run	388
TESTRUN EXAMPLE.....	374, 387	nh3tr2 test run	389
MF DIVIDING SURFACE	135, 141	NITER KEYWORD	
microcanonical optimized multidimensional		DEFINITION	165
tunneling	16	NOALLEXCIT KEYWORD	
MICROCANONICAL Variational Theory	16	TESTRUN EXAMPLE	380
MINIMUM ENERGY PATH (MEP)		NOCVT KEYWORD	
CALCULATING	52	TESTRUN EXAMPLE	376
MIRP KEYWORD		NONLINEAR KEYWORD	
DEFINITION	283	TESTRUN EXAMPLE	379, 423
MITS KEYWORD		NOOPT KEYWORD	
DEFINITION	283	TESTRUN EXAMPLE	383, 384, 385
MNBRK SUBPROGRAM	454	NOPROJECT KEYWORD	
MODULE FILES	475	TESTRUN EXAMPLE	392
ALLOC.F90.....	475	NORMOD SUBPROGRAM	73, 74, 454
MOPAC program	19	NOROUT SUBPROGRAM	454
MOPEG SUBPROGRAM	454	NOSADDLE KEYWORD	55
MOPHES SUBPROGRAM	454	NOST31 SUBPROGRAM	454
MOPOPT SUBPROGRAM	454	NQE PARAMETER	224
MOPSET SUBPROGRAM	454	NQTH	60
MORATE	19	NQTH PARAMETER	224
MORMODEL KEYWORD	100	NSEG2	60
DEFINITION	176	NSEGBOLTZ KEYWORD	
MORSEI OPTION		TESTRUN EXAMPLE	379
TEST RUN EXAMPLE	423, 424, 425, 426	NSEGHETA KEYWORD	
TESTRUN EXAMPLE	383, 384, 386, 391	TESTRUN EXAMPLE	379
MORSEIIIA OPTION		NSTATE SUBPROGRAM	454
TEST RUN EXAMPLE	379	NSTEPS KEYWORD	
RELATED KEYWORDS.....	176, 177	DEFINITION	209
MORSE KEYWORD	100	NUMFRQ SUBPROGRAM.....	455
DEFINITION	176	NUMSTEP KEYWORD	109
RELATED KEYWORDS.....	176	DEFINITION	160
TESTRUN EXAMPLE....	379, 383, 384, 386, 423,	RELATED KEYWORDS	161
424, 425, 426		NUMTYP KEYWORD.....	108
MORSEQQ KEYWORD	100	DEFINITION	161
DEFINITION	177	NWCHEMRATE	19
RELATED KEYWORDS.....	171, 176	OBTSRT SUBPROGRAM	455
TESTRUN EXAMPLE.....	391	oh3fu30tr1 testrun.....	400
MUCDSC SUBPROGRAM.....	454	oh3fu30tr2 testrun.....	401
MULTILEVELRATE	19	oh3fu30tr3 testrun.....	402
MUSC SUBPROGRAM	454	oh3fu30tr4 testrun.....	403
MUSCCD SUBPROGRAM.....	454	oh3fu30tr5 testrun.....	404
MUVT KEYWORD		oh3fu30tr6 test run.....	405
DEFINITION	232	oh3fu31tr1 test run.....	406
MUVTOPT KEYWORD		oh3fu31tr2 test run.....	407
DEFINITION	233	oh3fu40tr1 testrun.....	408

oh3fu40tr2 testrun	409	PATH SECTION.....	188, 302
oh3fu40tr3 testrun	410	EXTRAPOLATION KEYWORDS	194
oh3fu40tr4 testrun	411	INTEGRATOR KEYWORDS	192
oh3fu40tr5 testrun	412	NES KEYWORDS.....	196
oh3fu40tr6 test run	413	PRINT KEYWORDS	195
oh3tr1 testrun.....	390	SSTOR KEYWORDS.....	196
oh3tr10 testrun.....	399	STEP SIZE KEYWORDS.....	192
oh3tr2 testrun.....	391	SVRC KEYWORDS	193
oh3tr3 test run.....	102	PATH SUBPROGRAM	455
oh3tr3 testrun.....	392	PCFREQA KEYWORD	
oh3tr4 testrun.....	393	DEFINITION	283
oh3tr5 testrun.....	394	PCFREQS KEYWORD	
oh3tr6 testrun.....	395	DEFINITION	284
oh3tr7 testrun.....	396	PCINFO KEYWORD	
oh3tr8 testrun.....	397	DEFINITION	284
oh3tr9 test run.....	398	PHASE KEYWORD	
ohcltr1 test run.....	423	SOLID OPTION	
ohcltr2 test run.....	424	TESTRUN EXAMPLE	414, 415
ohcltr3 test run.....	425	PHID SUBPROGRAM	455
ohcltr4 test run.....	426	PHSINT SUBPROGRAM	455
ohcltr5 test run.....	427	PIVOT KEYWORD	
OHOOK SUBPROGRAM.....	90, 455	DEFINITION	301
OPENFI SUBPROGRAM	455	PMOI SUBPROGRAM	455
OPT50 SUBPROGRAM	455	POLINT SUBPROGRAM	455
OPT50C SUBPROGRAM	455	POLYAT SUBPROGRAM.....	73, 456
OPTIMIZATION SECTION.....	162, 298	<i>POLYRATE</i>	
OPTION SUBPROGRAM	70, 455	DESCRIPTION OF SUBPROGRAMS	435
OPTMIN KEYWORD		DIRECTORY STRUCTURE.....	305
OPTIMIZATION		HIDDEN FILE.....	486
DEFINITION	165	INSTALLATION PROCEDURES.....	305
OPTTS KEYWORD		INSTRUCTIONS FOR GENERAL USE.....	313
OPTIMIZATION		LICENSING.....	26
DEFINITION	166	MEMORY REQUIREMENTS.....	494
ORD31 SUBPROGRAM	455	MODULE FILES	475
Orientation.....	94	PROGRAM STRUCTURE	70
Output		RUNNING TEST RUNS	312
files fu41-fu47	304	TESTING PROCEDURES.....	311
OUTUNIT KEYWORD		potcomp.jc	487
DEFINITION	153	Potential energy surface.....	17, 435
OVLRLP SUBPROGRAM.....	455	POTENTIAL KEYWORD.....	91
P1FREQ SECTION		DEFINITION	158, 297
DEFINITION	241	UNIT30 OPTION	
P1HESS SECTION		TESTRUN EXAMPLE.....	400, 401, 402, 403, 404, 408, 409
DEFINITION	239	UNIT40 OPTION	
P2FREQ SECTION		TESTRUN EXAMPLE	410, 411, 412
DEFINITION	241	POTINF KEYWORD	82
P2HESS SECTION		DEFINITION	210
DEFINITION	239	PRCORD SUBPROGRAM.....	456
PAGEM KEYWORD.....	110	PRDELG KEYWORD	
TESTRUN EXAMPLE.....	324, 331, 333, 379, 393	DEFINITION	233
Partition function		PRDISTMX KEYWORD	
electronic	45	DEFINITION	210
rotational.....	40, 45	PREGPT SUBPROGRAM.....	456
vibrational.....	40, 45, 59		

PREP SUBPROGRAM	88, 456	DEFINITION	234
PREPJ SUBPROGRAM	88, 456	PRPATH KEYWORD	
PRINT KEYWORD		DEFINITION	210
DEFINITION	166, 241	PRSAVEMODE KEYWORD	
PRINTSTEP KEYWORD		DEFINITION	211
DEFINITION	210	PRSAVERP KEYWORD	82
PRMEP SUBPROGRAM.....	456	DEFINITION	211
PRMODE KEYWORD		PRSQ SUBPROGRAM	456
DEFINITION	177	PRVIB KEYWORD	
PRNTBT SUBPROGRAM.....	456	DEFINITION	234
PRNTFC SUBPROGRAM.....	456	PSAG SUBPROGRAM	456
PROD1 SECTION	168	PSATP SUBPROGRAM.....	456
PROD2 SECTION	168	PSATP2 SUBPROGRAM.....	456
PRODFREQ KEYWORD	113	PSATX SUBPROGRAM	456
DEFINITION	285	PSATX2 SUBPROGRAM	457
Product		PTORS SUBPROGRAM.....	457
force constant matrix	50	PTQVIB SUBPROGRAM.....	457
geometry optimization.....	48	PTSEC40 SUBPROGRAM.....	457
zero point energy	51	PYC1 SUBPROGRAM	457
PROGRAM FILES		PYC1P SUBPROGRAM	457
ACESPOLY.....	475	PYC2 SUBPROGRAM	457
EF 475		PYC2P SUBPROGRAM	457
ENERGETICS.....	475	PYCOEF SUBPROGRAM	457
FROMBLAS.....	475	QFORM SUBPROGRAM	458
GIVTST.....	475	QQPOT SUBPROGRAM.....	458
HEADR	475	QQSEMI KEYWORD.....	100, 101
HOOKS	476	DEFINITION	178
INTBSV1.....	476	RELATED KEYWORDS	172
INTBSV2.....	476	TESTRUN EXAMPLE	339, 379
INTBSV3.....	476	QQWKB KEYWORD	100, 101
INTERFACE.....	476	DEFINITION	178
IVTSTM	476	RELATED KEYWORDS	172
MAIN.....	476	TESTRUN EXAMPLE	385
MAIN_VRCMPI	476	QRFAC SUBPROGRAM.....	458
MOPLY	476	QRSENE SUBPROGRAM.....	458
POLY31	477	QRST	
POLY40.....	477	KEYWORD DEFINITION	223
POLYAG	476	QSLVE SUBPROGRAM	458
POLYHL	476	QTQVIB SUBPROGRAM	458
POLYMQ.....	476	QUAD KEYWORD	
POLYRR	476	DEFINITION	224
POLYSZ	477	QUADFT SUBPROGRAM	458
RTPJAC.....	477	QUADR SUBPROGRAM	458
VRCTST_MPI	477	QUADTW SUBPROGRAM	458
PROGRAM SUBPROGRAMS		R1FREQ SECTION	
SETUP	475	DEFINITION	241
SURF	475	R1HESS SECTION	
PROJECT SUBPROGRAM.....	74, 456	DEFINITION	239
PROJECT KEYWORD		R1MPYQ SUBPROGRAM	458
DEFINITION	177	R1UPDT SUBPROGRAM.....	458
TESTRUN EXAMPLE.....	380	R2FREQ SECTION	
PROJF SUBPROGRAM.....	456	DEFINITION	241
PROJTI SUBPROGRAM.....	456	R2HESS SECTION	
PRPART KEYWORD		DEFINITION	239

RACES SUBPROGRAM.....	458	READIC SUBPROGRAM.....	459
RALL31 SUBPROGRAM.....	458	READINT SUBPROGRAM	459
RAN1 SUBPROGRAM	459	READLN SUBPROGRAM	460
RANALY SUBPROGRAM.....	459	REARG1 SUBPROGRAM.....	460
RANGEIC KEYWORD		RECS31 SUBPROGRAM	460
DEFINITION	286	REDGEO SUBPROGRAM	70, 460
RATE SECTION	226, 303	REDMU SUBPROGRAM.....	460
print keywords	229	References	
RATE SUBPROGRAM.....	62, 75, 458	bibliography	504
RATOMS SUBPROGRAM.....	459	convention explained	38
RBASIS SUBPROGRAM.....	458	footnotes to text.....	500
RBATH SUBPROGRAM	458	REFOPT SUBPROGRAM.....	460
RCFREQA KEYWORD		RELEC SUBPROGRAM	460
DEFINITION	285	RENERG SUBPROGRAM.....	460
RCFREQS KEYWORD		REORDER KEYWORD	
DEFINITION	285	DEFINITION	211
RCINFO KEYWORD		REPFL SUBPROGRAM.....	460, 464
DEFINITION	286	RES1 SUBPROGRAM.....	460
RCNST SUBPROGRAM.....	459	RESTART KEYWORD	77, 116
RD40EXTR SUBPROGRAM	459	DEFINITION	153
RD40GEN SUBPROGRAM	459	TESTRUN EXAMPLE	324, 326, 329, 377, 378
RD40RP SUBPROGRAM.....	459	RESTOR SUBPROGRAM	70, 460
RDETIL SUBPROGRAM.....	459	RESTRT SUBPROGRAM.....	460
RDHESS SUBPROGRAM.....	459	RETRY KEYWORD	
RDIATM SUBPROGRAM	459	DEFINITION	166
RDIFFU SUBPROGRAM.....	459	Reverse rate constant.....	42, 61, 113
RDLN SUBPROGRAM	459	REVKEXP KEYWORD	
REACFREQ KEYWORD		DEFINITION	234
DEFINITION	287	TESTRUN EXAMPLE	383, 384, 385
REACT SUBPROGRAM.....	73, 459	REXTRP SUBPROGRAM	460
REACT1 SECTION	168	REZERO SUBPROGRAM.....	460
REACT2 SECTION	168	REZERU SUBPROGRAM.....	461
Reactant		RFCFAC SUBPROGRAM	461
force constant matrix	50	RFREQ SUBPROGRAM	461
geometry optimization.....	48	RGEM31 SUBPROGRAM	461
properties	48	RGEN SUBPROGRAM.....	461
zero point energy	51	RGEN31 SUBPROGRAM.....	461
REACTCOM SUBPROGRAM	459	RGENER SUBPROGRAM.....	461
Reaction coordinate.....	40, 51	RGEOM SUBPROGRAM.....	461
determining the sign of.....	76	RGRD SUBPROGRAM.....	461
Reaction path.....	40, 51, 52, 61	RGRD31 SUBPROGRAM.....	461
Euler stabilization method.....	53	RGSAD SUBPROGRAM.....	461
negative generalized normal mode eigenvalue		RGSPEC SUBPROGRAM	461
.....	61	RGSZM SUBPROGRAM	461
projected force constant matrix	54	RGTEMP SUBPROGRAM	461
save grid.....	54, 74	RGZMAT SUBPROGRAM	461
step sizes.....	110	RHES31 SUBPROGRAM	462
Reaction rate methods keywords.....	228	RHESS SUBPROGRAM.....	461
Reaction rates	59	RHESSC SUBPROGRAM	462
generalized free energy of activation	59	RHFSEC SUBPROGRAM	462
transmission coefficients	59	RHSS31 SUBPROGRAM	462
transmission probabilities.....	59	RICOPT SUBPROGRAM	462
Boltzmann average	58	RLCT SUBPROGRAM	462
READ5 SUBPROGRAM	70, 459	RLINE SUBPROGRAM	462

RLOWFR SUBPROGRAM.....	462	RSPDRV SUBPROGRAM.....	465
RMUVT SUBPROGRAM.....	462	RSPECL SUBPROGRAM.....	465
RODS KEYWORD		RSPP SUBPROGRAM.....	465
DEFINITION.....	211	RSSTOP SUBPROGRAM.....	465
ROPT SUBPROGRAM.....	462	RST SUBPROGRAM.....	465
ROPT31 SUBPROGRAM.....	462	RSTAT SUBPROGRAM.....	465
ROPTOP SUBPROGRAM.....	462	RSTATE SUBPROGRAM.....	465
rotate utility program.....	482	RSVRC SUBPROGRAM.....	465
RPART SUBPROGRAM.....	462	RTEMP SUBPROGRAM.....	465
RPART1 SUBPROGRAM.....	462	RTITLE SUBPROGRAM.....	466
RPATH SUBPROGRAM.....	462	RTOR SUBPROGRAM.....	466
RPH31 SUBPROGRAM.....	462	RTOROPT SUBPROGRAM.....	466
RPH40 SUBPROGRAM.....	462	RTPJAC SUBPROGRAM.....	466
RPHAIT SUBPROGRAM.....	463	RTUNNL SUBPROGRAM.....	466
RPHB SUBPROGRAM.....	463	RUNM31 SUBPROGRAM.....	466
RPHDXN SUBPROGRAM.....	463	RVAR6 SUBPROGRAM.....	466
RPHEXP SUBPROGRAM.....	463	RVARY SUBPROGRAM.....	466
RPHFIT SUBPROGRAM.....	463	RVPERT SUBPROGRAM.....	466
RPHFT2 SUBPROGRAM.....	463	RVRANG SUBPROGRAM.....	466
RPHFT3 SUBPROGRAM.....	463	RVRCOPT SUBPROGRAM.....	466
RPHG1 SUBPROGRAM.....	463	RWKB SUBPROGRAM.....	466
RPHG2 SUBPROGRAM.....	463	RWORD SUBPROGRAM.....	466
RPHG5 SUBPROGRAM.....	463	RZVAR SUBPROGRAM.....	466
RPHINT SUBPROGRAM.....	463	SADDLE KEYWORD.....	52
RPHMEF SUBPROGRAM.....	463	DEFINITION.....	211
RPHNWT SUBPROGRAM.....	463	SADDLE SUBPROGRAM.....	52, 466
RPHRD1 SUBPROGRAM.....	464	SADENG SUBPROGRAM.....	467
RPHRD2 SUBPROGRAM.....	464	SADFREQ KEYWORD	
RPHSET SUBPROGRAM.....	464	DEFINITION.....	287
RPHTRF SUBPROGRAM.....	464	SADSEC40 SUBPROGRAM.....	467
RPHTRX SUBPROGRAM.....	464	Sample test runs.....	321
RPHWRT SUBPROGRAM.....	464	Save grid.....	54, 74
RPIVOT SUBPROGRAM.....	463	SAVEXRP SUBPROGRAM.....	467
RPL KEYWORD.....	116	SCALE KEYWORD	
<i>RPM KEYWORD</i>		DEFINITION.....	166
DEFINITION.....	211	SCALEMASS KEYWORD.....	74
RPNT31 SUBPROGRAM.....	463	DEFINITION.....	212
RPOTET SUBPROGRAM.....	464	TESTRUN EXAMPLE.....	327
RPOTGE SUBPROGRAM.....	464	SCLPT KEYWORD	
RPRMEP SUBPROGRAM.....	464	DEFINITION.....	212
RPRVIB SUBPROGRAM.....	464	Scripts	
RPSEC40 SUBPROGRAM.....	464	comp_pot.jc.....	487
RP-VTST rate constants.....	16	comp_src.jc.....	487
RQCOM SUBPROGRAM.....	464	link_exe.jc.....	488
RQRST SUBPROGRAM.....	464	potcomp.jc.....	487
RQUAD SUBPROGRAM.....	464	SCT KEYWORD	
RRANGE SUBPROGRAM.....	464	DEFINITION.....	224
RRATE SUBPROGRAM.....	465	TESTRUN EXAMPLE.....	324, 333
RRPM SUBPROGRAM.....	465	SCTOPT KEYWORD	
RRSTRT SUBPROGRAM.....	465	DEFINITION.....	224
RSCT SUBPROGRAM.....	465	LAGRANGE OPTION.....	111
RSECND SUBPROGRAM.....	465	TESTRUN EXAMPLE.....	331, 336
RSFRST SUBPROGRAM.....	465	TESTRUN EXAMPLE.....	327
RSP SUBPROGRAM.....	465	TESTRUN EXAMPLE.....	

SPLINE OPTION	328	SPLNMF SUBPROGRAM	468
<i>SDEB1</i> KEYWORD		SPLNV SUBPROGRAM	468
DEFINITION	213	SPLRMI SUBPROGRAM	469
<i>SDEB2</i> KEYWORD		SPLSPV SUBPROGRAM	469
DEFINITION	213	SPLV31 SUBPROGRAM	469
SDSTART KEYWORD		SPRNT KEYWORD	
DEFINITION	167	DEFINITION	215
SECCEN SUBPROGRAM	467	SRANGE KEYWORD	
SECCEP SUBPROGRAM	467	DEFINITION	215
Second derivatives		TESTRUN EXAMPLE	386, 388, 389
central differences	111	SSAVE SUBPROGRAM	469
numerical	108	SSPECIAL KEYWORD	
numerical fourth order polynomial	108	DEFINITION	215
quadratic fit	108	TESTRUN EXAMPLE	385
SECOND SECTION	159	SSPECPR KEYWORD	
SECOND SECTION KEYWORDS		DEFINITION	215
<i>POLYRATE</i>	159	SS-QRRK	2, 13, 485, 575
SENOUT SUBPROGRAM	467	SSTEP KEYWORD	53, 73, 109
SETANH SUBPROGRAM	467	DEFINITION	216
SETLGS SUBPROGRAM	467	TESTRUN EXAMPLE	329, 332, 379
SETUP SUBPROGRAM	80, 85, 467	SSTOR	100, 179, 196, 216
SETVAR SUBPROGRAM	467	SSX SUBPROGRAM	469
SEX1 KEYWORD		START SECTION	168
DEFINITION	241	STAT31 SUBPROGRAM	469
SFIRST KEYWORD	110	STATE KEYWORD	
DEFINITION	213	ADIAB OPTION	
TESTRUN EXAMPLE	336, 390	TESTRUN EXAMPLE	393
SGND3V SUBPROGRAM	467	DEFINITION	235
SHFTSAD SUBPROGRAM	467	DIAB OPTION	
SIGMAF KEYWORD		TESTRUN EXAMPLE	394
DEFINITION	234, 303	STATEOPT KEYWORD	
SIGMAR KEYWORD		DEFINITION	236
DEFINITION	234	STATUS KEYWORD	
SIGN KEYWORD	76	DEFINITION	178
DEFINITION	213	STAU SUBPROGRAM	469
SIGS SUBPROGRAM	467	STLGSJ SUBPROGRAM	469
SIMPSN SUBPROGRAM	467	STOR SUBPROGRAM	469
SINE SUBPROGRAM	467	<i>STPTOL</i> KEYWORD	
Small-curvature Tunneling	16, 46	DEFINITION	167
SNHCSH SUBPROGRAM	468	STVARJ SUBPROGRAM	469
SORT31 SUBPROGRAM	468	SUPERMOL KEYWORD	
SORTG SUBPROGRAM	468	DEFINITION	154
SPECIES KEYWORD		SURF SUBPROGRAM	80, 81, 83, 85, 469
DEFINITION	178	SVBCMP SUBPROGRAM	469
SPECSTOP KEYWORD		SVBKSJ SUBPROGRAM	469
DEFINITION	214	SVRC KEYWORD	
SPL1B1 SUBPROGRAM	468	DEFINITION	302
SPL1B2 SUBPROGRAM	468	SYMMETRY KEYWORD	
SPL1D1 SUBPROGRAM	468	DEFINITION	216
SPL31 SUBPROGRAM	468	TESTRUN EXAMPLE	381, 382
SPLM31 SUBPROGRAM	468	TABLE SUBPROGRAM	470
SPLNB SUBPROGRAM	468	TAUV SUBPROGRAM	470
SPLNF SUBPROGRAM	468	TEMP KEYWORD	
SPLNM SUBPROGRAM	468	DEFINITION	236

RELATED KEYWORDS.....	230, 231	UNITS KEYWORD	
TESTRUN EXAMPLE.....	324, 333, 383, 384	DEFINITION	241
THETA2 SUBPROGRAM.....	470	UPCAS SUBPROGRAM	471
THETA3 SUBPROGRAM.....	470	UPCSE SUBPROGRAM.....	471
TITLE KEYWORD		UPDHR SUBPROGRAM.....	471
DEFINITION	154	UPDMI SUBPROGRAM	472
TITLE SUBPROGRAM	470	US KEYWORD	
TOR KEYWORD		DEFINITION	237
DEFINITION	179	TESTRUN EXAMPLE	383, 384
TESTRUN EXAMPLE.....	377, 378, 416	Utility routines	
TOROPT KEYWORD		chain	84
DEFINITION	180	coord	83
TP SUBPROGRAM	470	VALVAG SUBPROGRAM	472
TPCDSC SUBPROGRAM.....	470	VANHAR KEYWORD	
TQL2 SUBPROGRAM.....	470	DEFINITION	185
TQLRAT SUBPROGRAM	470	TESTRUN EXAMPLE	339, 377, 378, 379, 385
TRANFC SUBPROGRAM	470	VARIABLE KEYWORD	101
TRANG SUBPROGRAM	470	Variable Reaction Coordinate and Multifaceted	
TRANLF SUBPROGRAM.....	470	Dividing Surface algorithm.....	135, 141
TRANS SUBPROGRAM.....	470	VECCON SUBPROGRAM	472
TRANS2 SUBPROGRAM.....	470	VECMAG SUBPROGRAM.....	472
TRANSF SUBPROGRAM.....	471	VIB KEYWORD	
Transmission coefficients.....	42, 46, 47, 59, 60	DEFINITION	186
TRBAK3 SUBPROGRAM	471	VICLCG SUBPROGRAM.....	472
TRED3 SUBPROGRAM.....	471	VIVT SUBPROGRAM.....	472
TREPT SUBPROGRAM	471	VPART SUBPROGRAM	472
TRLFRP SUBPROGRAM	471	VPARTI SUBPROGRAM.....	472
TSC SUBPROGRAM	471	VPFEXP KEYWORD	
TSFREQ SECTION		DEFINITION	237
DEFINITION	241	TESTRUN EXAMPLE	327, 331
TSGCOMP KEYWORD		VRANGE KEYWORD.....	101
DEFINITION	164	DEFINITION	186
TSHESS SECTION		VRC.....	135
DEFINITION	239	VRC KEYWORD.....	55, 70, 295
TSHREC KEYWORD		DEFINITION	296
DEFINITION	164	VRCNEJ SUBPROGRAM	472
TSOUT SUBPROGRAM.....	62, 471	VRCOPT KEYWORD	
TSPSI SUBPROGRAM.....	471	DEFINITION	296
TSRATE SUBPROGRAM.....	62, 471	VRCTST SUBPROGRAM	472
TST		VSCALE KEYWORD	
TESTRUN EXAMPLE.....	329	DEFINITION	216
TST KEYWORD		VSPLI2 SUBPROGRAM.....	472
DEFINITION	237	VSPLIN SUBPROGRAM	472
TESTRUN EXAMPLE.....	376	VTMUSN SUBPROGRAM	472
TUNNEL SECTION.....	217	VTST-IOC	67
Tunneling		VTST-ISPE.....	116
canonical optimized multidimensional.....	63	WELLP SECTION	168
interpolation of effective reduced mass.....	111	WELLR SECTION	168
microcanonical optimized multidimensional		WIGNER KEYWORD	
.....	63	DEFINITION	225
TURNPT SUBPROGRAM.....	471	Wigner tunneling	51, 59
UNITMI KEYWORD		WKB KEYWORD.....	100, 101, 102
DEFINITION	287	DEFINITION	187
TESTRUN EXAMPLE.....	387	TESTRUN EXAMPLE	392

WKB METHOD		YSECEP SUBPROGRAM.....	473
based on true potential.....	102	YTRANS SUBPROGRAM.....	474
PRIMITIVE.....	101	ZCT KEYWORD	
UNIFORM	101	DEFINITION	225
WKBENE SUBPROGRAM	473	TESTRUN EXAMPLE	324, 328, 329, 330, 333,
WKBPOT SUBPROGRAM.....	473	334, 416, 418, 419	
WKBVIB SUBPROGRAM	473	Zero of energy.....	79
WRITE62 KEYWORD		Zero-curvature Tunneling.....	16, 46
DEFINITION	155	ZEROPT SUBPROGRAM	474
WRITEFU30 KEYWORD		ZERP SUBPROGRAM.....	474
DEFINITION	155	ZIGAMA SUBPROGRAM.....	474
WRITEFU31 KEYWORD		ZOC3P SUBPROGRAM	474
DEFINITION	155	ZOCAST SUBPROGRAM	474
WRTHOK SUBPROGRAM	473	ZOCFRE SUBPROGRAM	474
WRTWEL SUBPROGRAM	473	ZOCPAR SUBPROGRAM.....	474
WTITLE SUBPROGRAM	472	ZOCPRN SUBPROGRAM.....	474
WVAR6 SUBPROGRAM	473	ZOCUPD SUBPROGRAM.....	474
XDOTP SUBPROGRAM	473	ZOCVCL SUBPROGRAM.....	474
XERBLA SUBPROGRAM	473	ZRIDDR SUBPROGRAM.....	474
XPROD SUBPROGRAM	473	ZROUT SUBPROGRAM.....	474
YDER24 SUBPROGRAM	473	ZRPTE SUBPROGRAM.....	474
YDERV2 SUBPROGRAM	473	ZSPMEP SUBPROGRAM	474
YNEWT SUBPROGRAM.....	473	μOMT.....	63
YSECEN SUBPROGRAM.....	473	μVT.....	39