

PCFC QM Subgroup: An introduction to MESS

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1 Introduction

1.1 What is MESS?

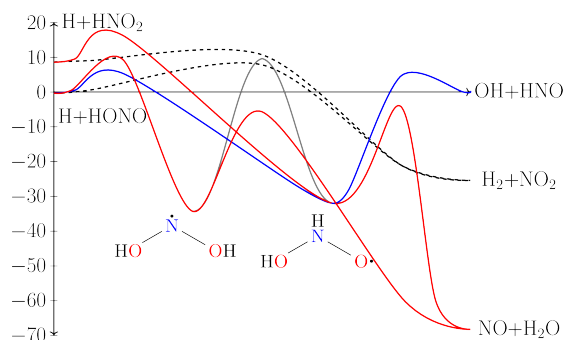
- MESS is the **Master Equation System Solver** (Publication[1]), a component of the PAPR suite of chemical kinetics codes
 - Developers: Y. Georgievskii, C. F. Goldsmith, J. A. Miller, M. P. Burke, and S. J. Klippenstein
 - PAPR: <https://tcg.cse.anl.gov/papr/codes/mess.html>
- Conformers on a potential energy surface are connected via transition states
- Ab initio calculations provide necessary input information to calculate partition functions and determine rates via transition state theory (TST)

1.2 $\text{H}_2\text{N}_1\text{O}_2$ Potential Energy Surface

Consider simple PES as example system (Publication[2])

- System consists of three kinds of nodes (conformers):
 - BIMOLECULAR: a pair of fragments which are lumped together, *e.g.* $\text{H}+\text{HNO}_2$
 - WELL: a unimolecular intermediate, *e.g.* HONOH
 - BARRIER: a transition state (TS) between two wells/bimoleculars
- Each conformer is described by its geometry and frequencies and (relative) energy

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2 MESS Input File

2.1 Input file overview

- Global section (unnamed header) specifies parameters for solution of the master equation
- Specify temperatures, pressures, calculation parameters
- Two major choices for calculation method
 - DIRECT: fuller, more expensive calculations (not so good at low temperature)
 - LOW-EIGENVALUE: assume which eigenvalues are chemical and which are relaxational, numerically cheap, not accurate at high temperatures
- MODEL definition follows and begins with collision parameters
- All conformers are then described to define system

2.1.1 Header Information

```

TemperatureList [K]          400.0 450.0 500.0 550.0 600.0 650.0 700.0 750.0
PressureList [atm]           0.01 0.1 0.316 1.0 3.16 10.0 31.6 100.0
EnergyStepOverTemperature    .2
ExcessEnergyOverTemperature  30
ModelEnergyLimit [kcal/mol]  400
CalculationMethod            direct #low-eigenvalue #direct
WellCutoff                   10
ChemicalEigenvalueMax        0.2
GlobalCutoff [kcal/mol]      -45.0
Model
  EnergyRelaxation
    Exponential
      Factor [1/cm]          200
      Power                   .85
      ExponentCutoff          15
  End
  
```

```

CollisionFrequency
LennardJones
  Epsilons [1/cm]          6.95  285.2
  Sigmas [angstrom]        2.55  5.182
  Masses [amu]             4.0  59
End

```

2.1.2 Bimolecular Input

```

!
!Bimolecular R1:  Reactants H + HNO2
!

```

```

Bimolecular          hno2      # H+HNO2
Fragment      H
! Current data taken from:
! / gpfs / data / cfgoldsm / cfgoldsm / qchem / gas_phase / h / h / dft / h_b2plypd3_ccpvtz . l
Atom
  Mass [amu]      1
  ElectronicLevels [1/cm]      1
    0      2
End
Fragment      HNO2
! Current data taken from:
! / gpfs / data / cfgoldsm / mfuller / qchem / noh / n1o2 / h1n1o2 / hno2 / hno2_b2plypd3_ccpvtz . l
RRHO
  Geometry [angstrom]      4
    N      0.000000      0.000000      0.000000
    O      0.000000      0.000000      1.219352
    O      0.955111      0.000000      -0.758013
    H      -0.933369      0.000000      -0.450837

  Core      RigidRotor
  SymmetryFactor      2
End

  Frequencies [1/cm]      6
    784.8  1047.3  1365.3
    1513.7  1643.6  3225.1

  ZeroEnergy [kcal / mol]      0.0
  ElectronicLevels [1/cm]      1
    0      1
End
GroundEnergy [kcal / mol]      8.56
End

```

2.1.3 Well Input

```

!
!Well 2: honho

```

```

!
Well      honho      # HONHO
Species
! Current data taken from:
! /gpfs/data/cfgoldsm/mfuller/qchem/noh/n1o2/h2n1o2/honho/honho_b2plypd3_
RRHO
  Geometry[angstrom]      5
  N      0.000000      0.000000      0.000000
  O      0.000000      0.000000      1.242823
  O      1.246970      0.000000      -0.684257
  H      -0.604321      0.645919      -0.510082
  H      1.844105      -0.334291      0.001465

  Core      RigidRotor
  SymmetryFactor      1
End
  Rotor      Hindered
  Group      5      # atoms in rotating group exclu
  Axis      1 3      # rotational axis
  Symmetry      1      # 360/angle of rotation to reco
  Potential[kcal/mol]      36      # number of equidistant point c
0.00 0.01 0.08 0.20 0.34 0.49 0.65 0.84 1.06 1.33
1.65 2.03 2.48 2.98 3.54 4.16 4.79 5.41 5.93 6.27
6.36 6.16 5.69 5.05 4.34 3.63 3.00 2.44 1.98 1.58
1.24 0.94 0.66 0.41 0.20 0.06
End

  Frequencies[1/cm]      8
  544.7 785.8
  956.3 1298.7 1415.2
  1519.8 3406.8 3749.3

!Torsional frequencies: 279.4

ZeroEnergy[kcal/mol]      -31.94
ElectronicLevels[1/cm]      1
  0 2
End

```

2.1.4 Barrier Input

```

!
!TS abstraction: h+hno2 <--> honho
!
Barrier      B4      hno2      honho      #

! Current data taken from:
! /gpfs/data/cfgoldsm/mfuller/qchem/noh/n1o2/h2n1o2/ts/ts_h_hno2_to_honho/

```

```

RRHO
  Geometry [ angstrom ]      5
  N      0.000000      0.000000      0.000000
  O      0.000000      0.000000      1.211134
  O      0.972148      0.000000      -0.753057
  H     -0.921040      0.065743      -0.468641
  H      1.786284      1.415784      -0.743167

  Core      RigidRotor
  SymmetryFactor      1
End

Frequencies [ 1/cm ]      8
  220.2   518.6   793.0
  1038.6  1355.2  1501.6
  1619.2  3235.5
!Imaginary mode:  -1352.8

ZeroEnergy [ kcal/mol ]    16.16
ElectronicLevels [ 1/cm ]  1
  0      2
Tunneling      Eckart
  ImaginaryFrequency [ 1/cm ]  1352.8
  WellDepth [ kcal/mol ]      7.60
  WellDepth [ kcal/mol ]      48.10
End
End

```

2.2 Generating MESS Input Blocks

- While you could, it is tedious and error-prone to write input files manually
- The utility program WRITEMESS reads a GAUSSIAN output (log) file and writes a conformer template
- Optionally, hindered rotors (GAUSSIAN log files) and energy from a single-point calculation (ORCA, currently) may also be added and written concurrently

3 Getting Results

3.1 Job Submission and Output

- Assuming everything is installed correctly, job submission on the HPC with slurm is simple: we call MESS on the input file
 - If we have an input file for a network of bimoleculars, wells, and barriers, then we call MESS on the input file
 - For two bimoleculars connected by a single transition state, we call ABSTRACTION on the input file

- A log file and output file will be generated by MESS
- Output file can be split into three sections:
 1. System network at the top
 2. Rate tables at each temperature and pressure condition
 3. Rate tables for each species pair
- The output file requires conversion to a recognized format for inclusion in chemical kinetic mechanisms

3.2 Fitting the Output

- A function to fit the output data to pressure-dependent, modified Arrhenius expressions is posted on the MESS webpage, http://tcg.cse.anl.gov/papr/codes/mess/mess_aux/mod_arr.tgz
- You will likely need to make some small changes for your specific system, but this will output CHEMKIN-format PLOG fits for all of the conformer pairs

4 Additional Resources

- The MESS manual: http://tcg.cse.anl.gov/papr/codes/mess/messmanual_3_23_16.pdf
- More examples to run and test: http://tcg.cse.anl.gov/papr/codes/mess/MESS_Examples.tgz

References

- [1] Y. Georgievskii, J. A. Miller, M. P. Burke, S. J. Klippenstein, Reformulation and solution of the master equation for multiple-well chemical reactions, *The Journal of Physical Chemistry A* 117 (2013) 12146–12154. doi:10.1021/jp4060704.
- [2] M. E. Fuller, C. F. Goldsmith, On the relative importance of HONO and HNO₂ in low-temperature combustion, *Proceedings of the Combustion Institute* 37 (1) (2019) 695–702. doi:10.1016/j.proci.2018.06.208.