

Primer to Building Models in RMG

Nth year RMG Users

27th Sept 2013



Outline

We will walk through main sections of a typical RMG input file:

- Databases
- Reactants and reactor conditions
- Pressure dependence parameters
- Model termination
- Guiding RMG Kinetics

Databases

```
//Example for the oxidation mechanism for nBuOH.
```

```
Database: RMG_database
```

```
PrimaryThermoLibrary:
```

```
Name: RMG-minimal
```

```
Location: primaryThermoLibrary
```

```
END
```

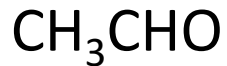
```
PrimaryTransportLibrary:
```

```
Name: GRIMech3.0
```

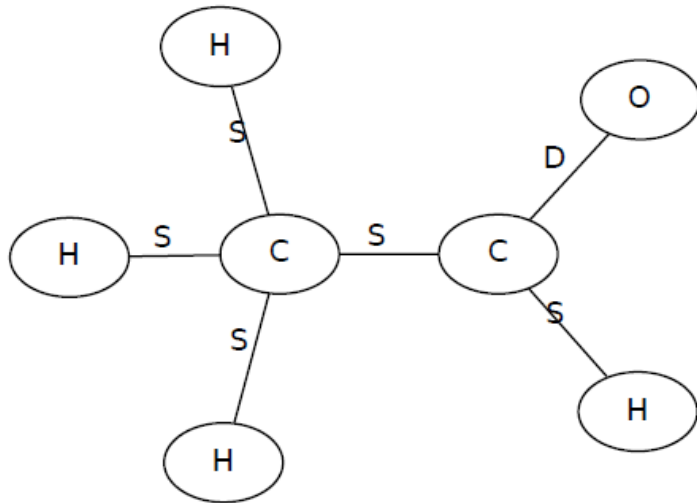
```
Location: GRI-Mech3.0
```

```
END
```

Reactor Conditions



```
1 C 0 {2,S}
2 C 0 {1,S} {3,D}
3 O 0 {2,D}
```



TemperatureModel: Constant (K) 1350

PressureModel: Constant (atm) 1

InitialStatus:

nBuOH (mol/cm³) 0.0338

Reactor conditions

```
1 C 0 {2,S} 2 C 0 {1,S} {3,S}
```

```
3 C 0 {2,S} {4,S}
```

```
4 C 0 {3,S} {5,S}
```

```
5 O 0 {4,S}
```

O₂ (mol/cm³) 0.4056

```
1 O 1 {2,S}
```

```
2 O 1 {1,S}
```

END

InertGas:

N₂ (mol/cm³) 0.5606

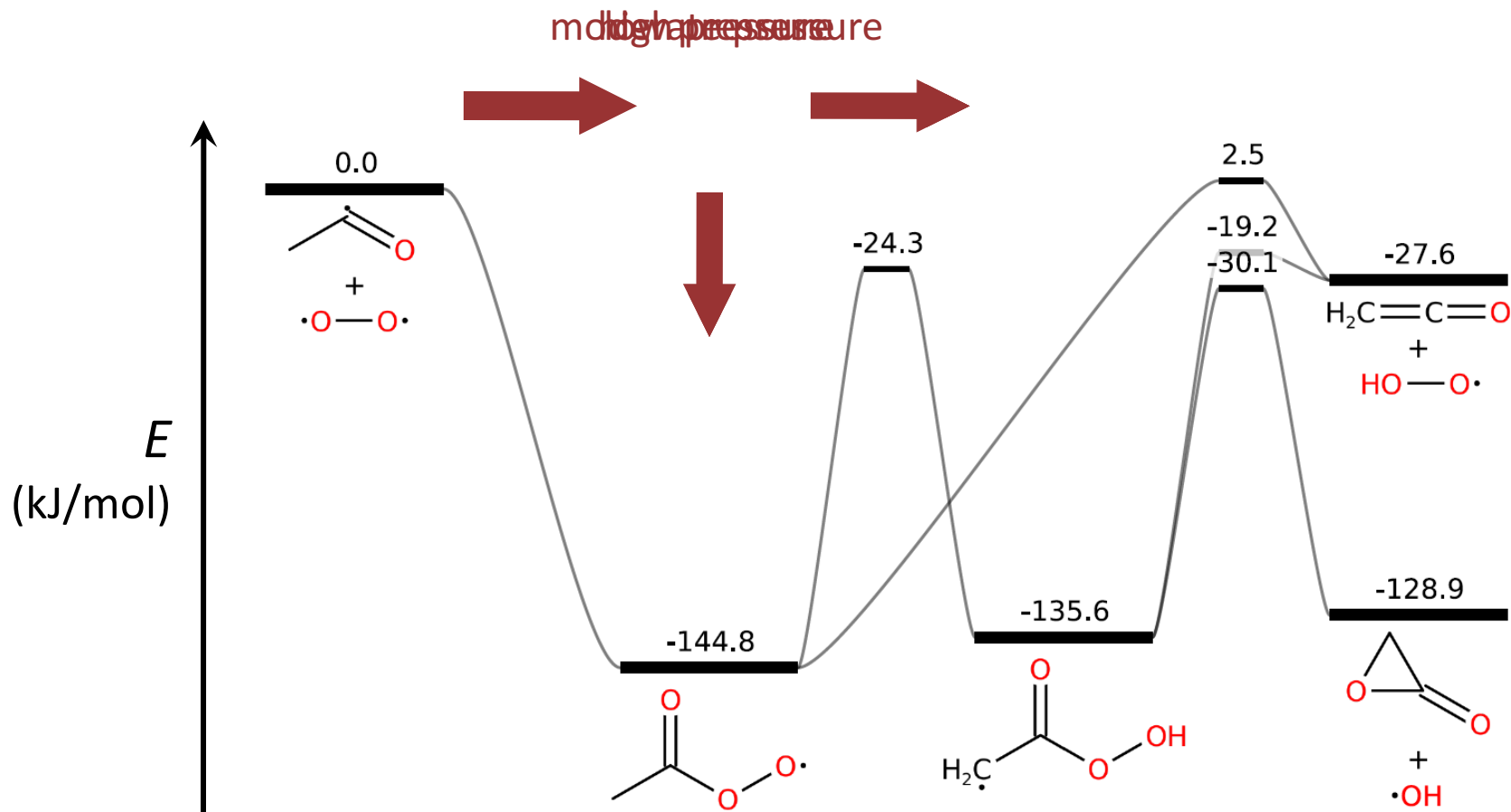
Ar (mol/cm³) 0.0

END

Pressure Dependence

```
//Pressure dependence specific parameters  
SpectroscopicDataEstimator: FrequencyGroups  
PressureDependence: ModifiedStrongCollision  
  
MaxAtomsForPressureDependence: 20  
// Size of adduct above which pdep will not be performed  
  
PDepKineticsModel: Chebyshev 6 4  
TRange: (K) 290.0 3000.0 8  
PRange: (bar) 0.01 100.0 5
```

Changing pressure can dramatically change product branching ratios in multi-well networks.



Methods for estimating pressure dependence kinetics in RMG

PressureDependence: ModifiedStrongCollision

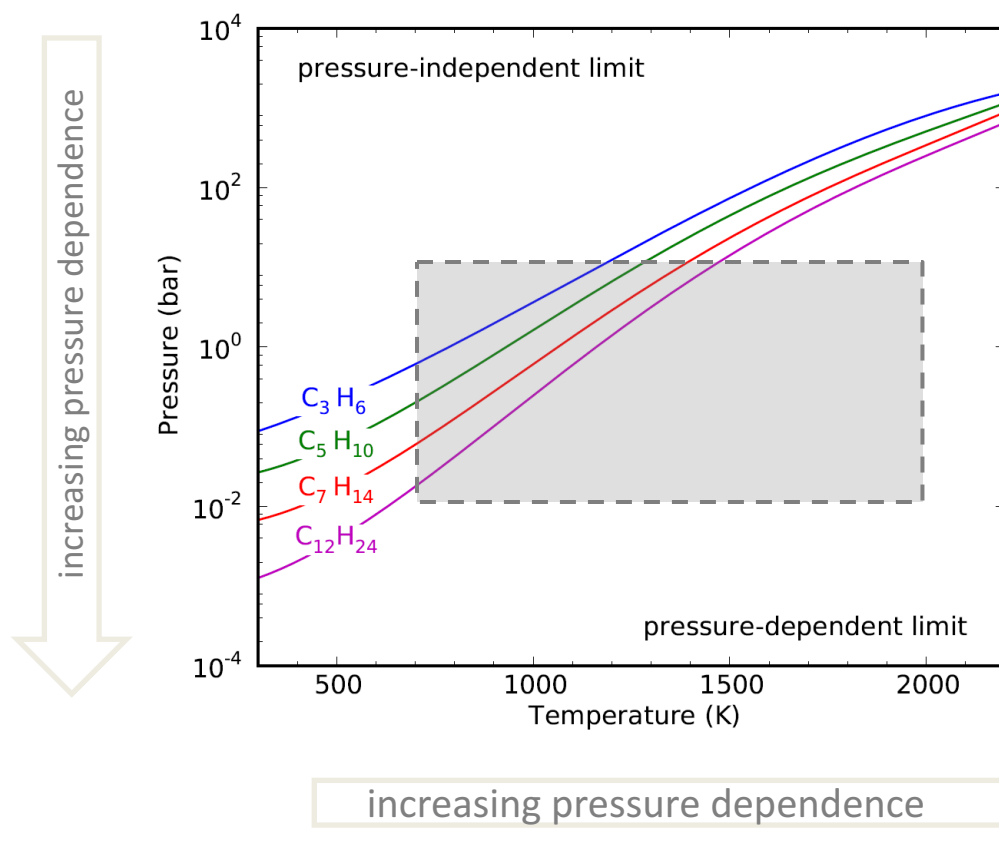
PressureDependence: ReservoirState

	MSC	RS	CSE
accuracy	okay	good (low T) bad (high T)	very accurate
speed	very fast	fast	very slow
robustness	very robust	robust	not robust



Maximum size for pressure dependent networks

MaxAtomsForPressureDependence: 20



switchover pressure – indicating the onset of pressure dependence as a function of temperature and molecular size

Interpolation model for pressure dependent kinetics

PDepKineticsModel: Chebyshev 6 4

PDepKineticsModel: PDepArrhenius

- **Option #1: Chebyshev**
 - Output is in CHEB parameters no physical significance
 - 6 points in Temperature and 4 points in Pressure has been found to be quite accurate and avoids overfitting
- **Option #2: PLOG**
 - Easier to understand the output
 - Usually the fit is not very accurate

Termination Criteria

FinishController:

(1) Goal Conversion: nBuOH 0.9

//(1) Goal ReactionTime: 0.1 (sec)

(2) Error Tolerance: 0.1

DynamicSimulator: DASSL

Conversions: AUTO

Atol: 1e-18

Rtol: 1e-8

Guiding RMG Kinetics

```
PrimaryKineticLibrary:  
END
```

```
ReactionLibrary:  
Name: nBuOH  
Location: nBuOH  
END
```

```
SeedMechanism:  
Name: GRIMech3.0  
Location: GRI-Mech3.0  
GenerateReactions: yes  
END
```

Reaction Libraries & Seed Mechanisms

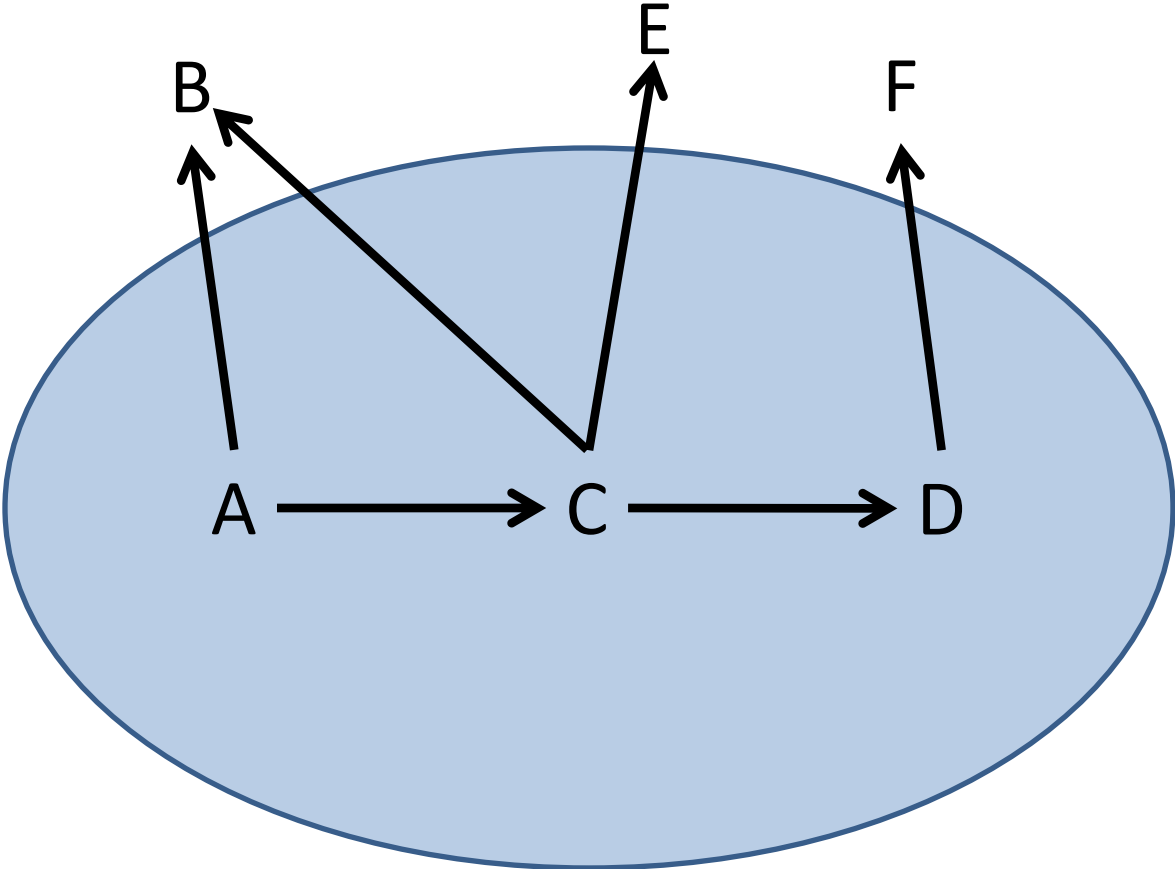
Many are already available in `RMG_database/kinetics_libraries`

Contain trusted kinetic parameters calculated via experiments or quantum chemistry

Include mechanisms for small and large(-ish) molecules in combustion and pyrolysis

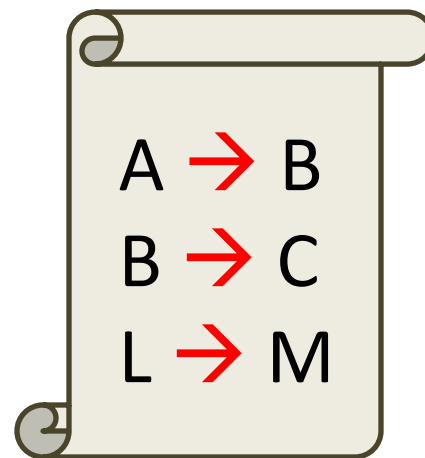
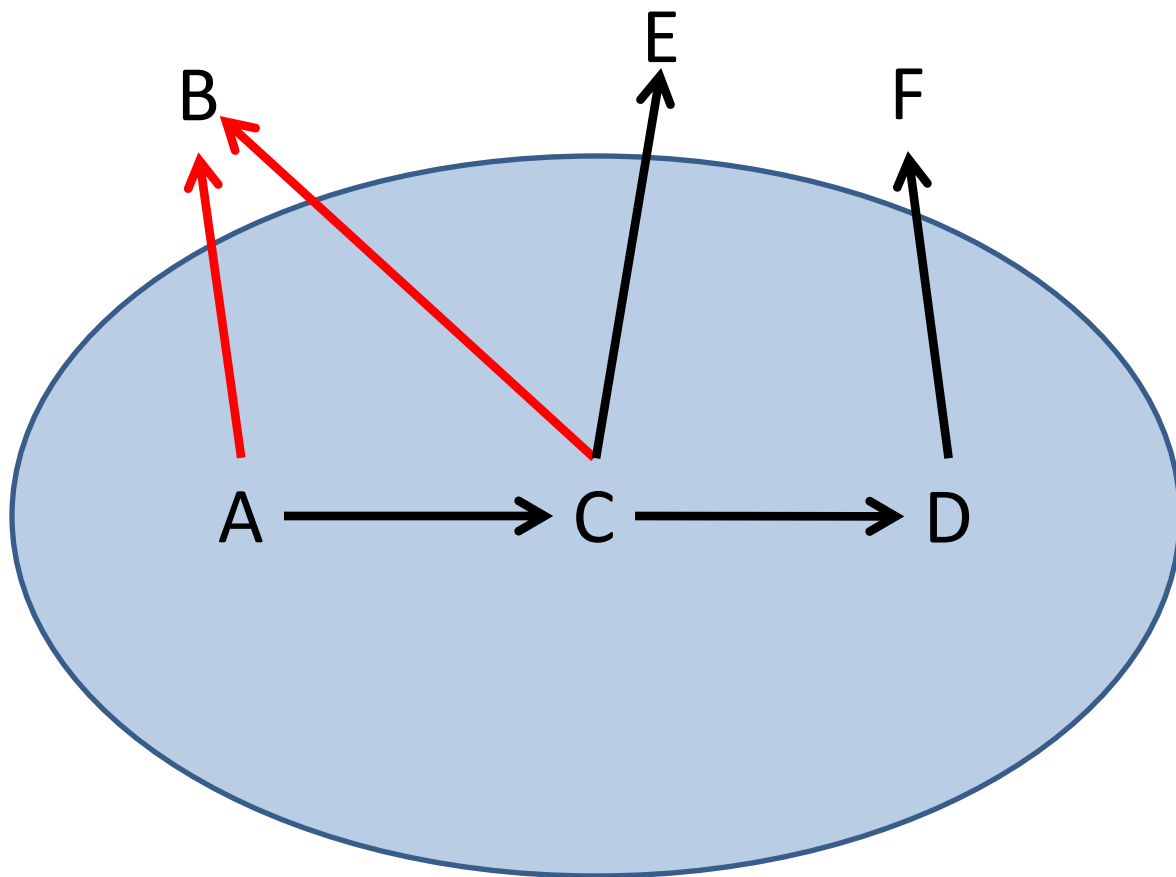
What's the difference?

General RMG Algorithm



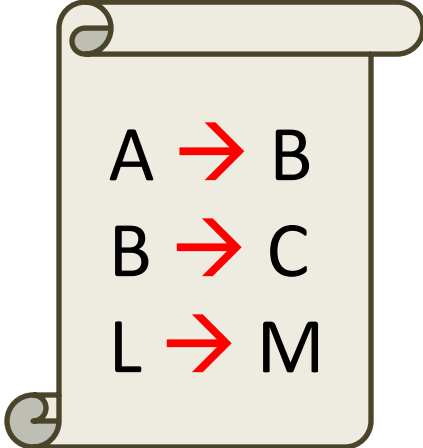
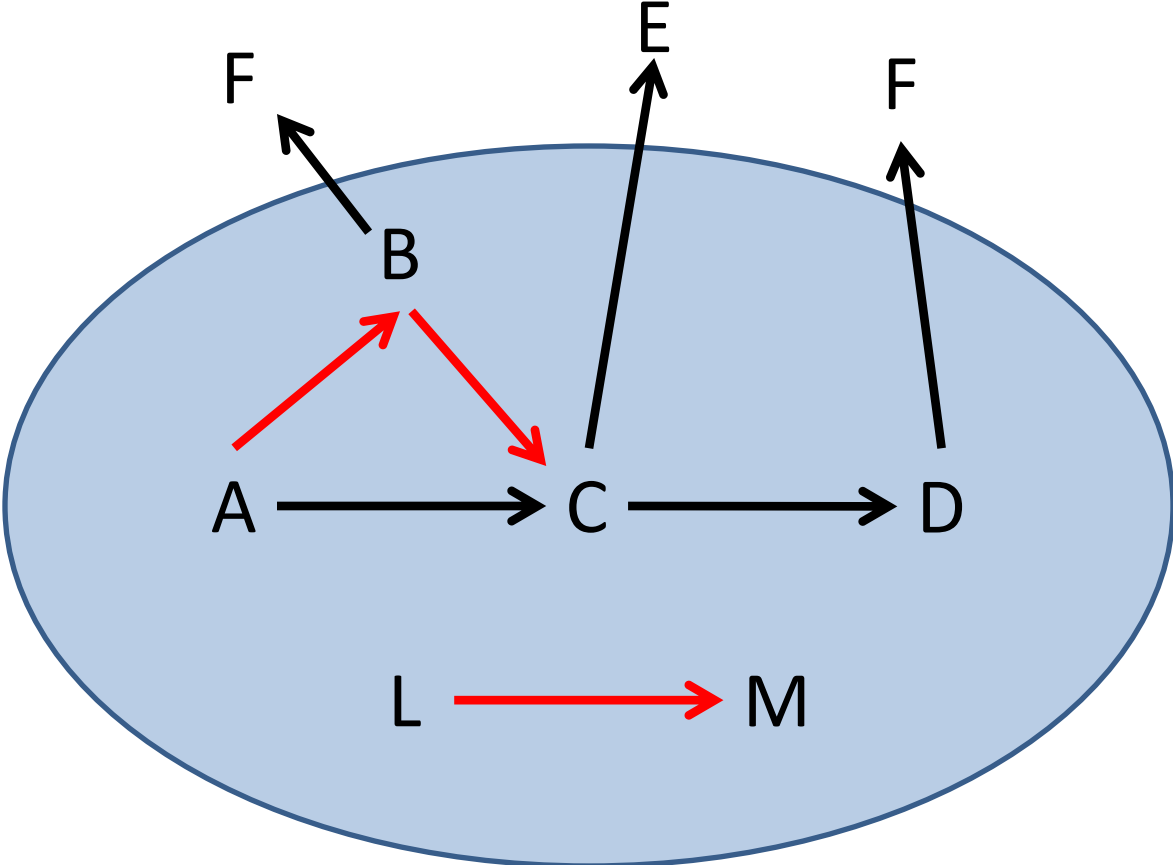
What's the difference?

Reaction Library



What's the difference?

Seed Mechanism



Reaction Libraries & Seed Mechanisms

In a folder within `RMG_database/kinetics_libraries/`

Must contain three files

- `reactions.txt`
- `pdepreactions.txt`
- `species.txt`

Sometime in the future: advanced tips for efficient model generation

- Imposing constraints on:
 - Molecular structure of products
 - Allowed chemical reactions
- Guiding RMG model generation:
 - Seed mechanisms
 - Kinetic/thermo libraries
 - Adding species to input file
- Playing around with termination criteria