

# Introduction to Computational Quantum Chemistry

## Lesson 07: Reaction Coordinates and Transition State Calculations



# REVIEW: Potential Energy Surface

- the PES of a molecule is obtained by the total interactions of:
  - nuclear-nuclear repulsion
  - electron-electron interaction
  - electron-nuclear attraction
- important Points on the PES
  - stationary points:

$$\frac{\partial E}{\partial q_i} = 0 \quad (1)$$

- local minimum:

$$\frac{\partial^2 E}{\partial q_i^2} > 0 \text{ for all degrees of freedom} \quad (2)$$

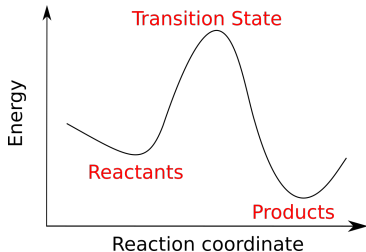
- $n^{\text{th}}$  order saddle point:

$$\frac{\partial^2 E}{\partial q_i^2} < 0 \text{ for } n \text{ degrees of freedom} \quad (3)$$



## REVIEW: Potential Energy Surface (*Cont.*)

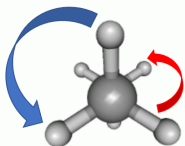
- a two-dimensional cut from a multidimensional PES illustrates one direction view of the reaction coordinates
- the energy differences of  $TS - R$  is the activation energy while the  $P - R$  is the reaction energy



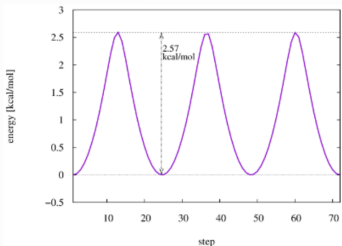
- however, these are only stationary points on a much larger potential energy surface (PES). The actual landscape of this surface can also be explored to see how the various stationary points connect.

# Potential Energy Scan

- it is often useful to scan the potential energy surface (PES), optimizing all other degrees of freedom for each particular value of the scanned variable(s).
- possible to adjust distances as well as atomic and dihedral angles



*Dihedral scan of ethane.*



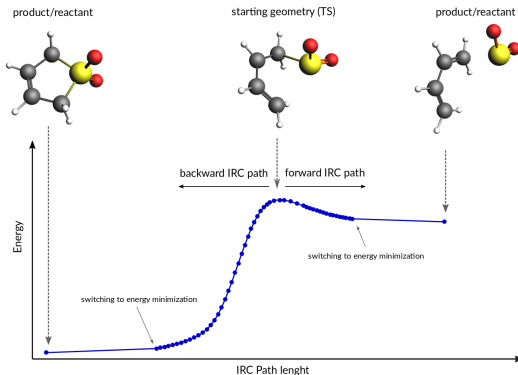
*Energy diagram of the dihedral scan of ethane.*

## Potential Energy Scan, *Cont.*

- PES scan may provide a rough estimate of a pathway between reactants, TS, and products, assuming the **coordinate(s) for the scan has been chosen wisely**
- PES scan is often used in development of classical force fields

# Intrinsic Reaction Coordinate (IRC)

- TS (transition state), the Hessian needs to display the required number of negative eigenvalues
- it is also necessary to **confirm the TS** connections to both sides (reactants and products).



## Intrinsic Reaction Coordinate (IRC), *Cont.*

- the Intrinsic Reaction Coordinate (IRC) is the minimum energy reaction pathway (MERP) in mass-weighted cartesian coordinates between the **TS** to its reactants **R** and products **P**.
- the molecule takes moving down the product and reactant valleys with zero kinetic energy.
- the Gonzalez-Schlegel method for following the coordinate can be used in Gaussian using the **IRC** keyword, latest version using (HPC algorithm).

# Transition State Methods

- mapping reaction coordinates and search of transition state (TS)
  - manual optimizations search for REACTANTS, TS, PRODUCTS (with aid from PES Scan)
  - advance methods such as QST2,QST3, etc



# ACTIVITY 1: PES SCAN

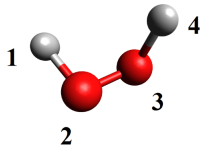
- rotational transition state in hydrogen peroxide (H-O-O-H) using Gaussian
  - (MANUAL) frozen optimization per angle:

```
#P HF/6-31G(d) opt=Z-Matrix

first step at d4=0.0

0 1
H1
02 1 r2
03 2 r3 1 a3
H4 3 r2 2 a3 1 d4

r2=1.0
r3=1.3
a3=110.
d4=0.0 F
```



- the last line of the Z-Matrix describes a value of 0.0 degree for the H/O/O/H dihedral angle **d4**, the tailing character **F** indicating that this variable is frozen and **not to be varied** during the geometry optimization, then change it from 0.0 to 180

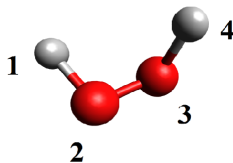
# ACTIVITY 1: PES SCAN, *Cont.*

- (FULL SCAN) can be achieved in a single job:

```
#P HF/6-31G(d) opt=Z-Matrix nosymm
H2O2 rotational potential

0 1
H1
02 1 r2
03 2 r3 1 a3
H4 3 r2 2 a3 1 d4

r2=1.0
r3=1.3
a3=110.
d4=0.0 S 18 +10.0
```



- initial value of **0.0** degree for the H/O/O/H dihedral angle **d4** but also specifies a **scan of 18 steps**, is varied by **+10.0 degrees**.
- in order to avoid problems caused through changes in the point group along the pathway, the **nosymm** keyword is added.

## ACTIVITY 1: PES SCAN, *Cont.*

- examine the output file and look for the keyword "**Summary of Optimized Potential Surface Scan**", you can see the energy eigenvalues as **d4** changes
- add module **gaussview**
- open the file using **gv {filename}**
- go to menu bar **Results > scan**, **Energy** and **RMS** values are presented
- you can visualize the molecular movements by clicking the animate play button

## ACTIVITY 2: Reaction Coordinate and TS Search

- we will study a classic  $\text{SN}_2$  reactions (known to have double well potentials), using Gaussian.
- $\text{SN}_2$  reaction:  $\text{Br}^- + \text{CH}_3\text{Cl} \rightarrow \text{Cl}^- + \text{CH}_3\text{Br}$ 
  - use B3LYP functional, the 6-31+G(d) basis set for all atom
  - first perform a geometry **Optimization** and **Freq** for all Reactants and Products
  - for an initial guess of transition state structure (TS), obtain it from PES scan by varying the both distances of **Br** and **Cl**, refer to this manual: <https://gaussian.com/scan/>
  - NOTE: to save time, [use the coordinates provided in the next page](#), this structure is a good starting point for a PES SCAN
  - refer to this manual: <https://gaussian.com/scan/>

## ACTIVITY 2: Reaction Coordinate and TS Search

- perform your SCAN from this structure (focus on B1 and B2) bond lengths

C

Cl 1 B1

Br 1 B2 2 A2

H 1 B3 2 A3 3 D3

H 1 B4 2 A4 3 D4

H 1 B5 2 A5 3 D5

variables:

B1 2.0 9 0.1

B2 2.0 9 0.1

A2 179.13318

B3 1.06210

A3 89.25624

D3 135.20467

B4 1.06224

A4 90.30745

D4 352.54852

B5 1.08241

A5 91.90132

D5 243.91238



## ACTIVITY 2: Reaction Coordinate and TS Search, *Cont.*

- once you found a good TS candidate, be sure it has strongest imaginary frequency among others that **refers to the bond breaking/forming** of the **Br ... C ... Cl**, that's why it's necessary to perform a preliminary FREQ calculation
- if it's viable, proceed to Transition State Optimization
- for Transition State (TS) minimization use `#P B3LYP/6-31G(d) opt(TS,ModRedundant,noeigentest) nosymm freq`
- the option above only calculate freq once on the initial structure (`calcfc`), freq calculation for every step is also available (`CalcAll`) but is not typically necessary for this simple system



## ACTIVITY 2: Reaction Coordinate and TS Search, *Cont.*

- after a successful TS Optimization, open the structure and examine the frequencies using Gabedit or Avogadro, if the frequencies are okay, you now got a **Transition State Structure**
- use the optimize TS structure and proceed with IRC calculation for the confirmation  
`#P B3LYP/6-31G(d) scf=(tight,direct) int=finegrid  
IRC(calcfc,maxpoints=500,maxcyc=500,stepsize=10)`
- by default IRC run examines every direction, you can also chose specific directions as **Forward** and **Reverse** options
- refer to this manual: <https://gaussian.com/irc/>

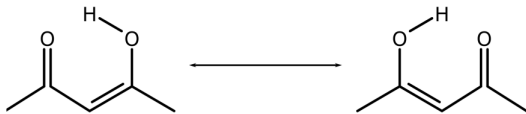
## ACTIVITY 2: Reaction Coordinate and TS Search, *Cont.*

- once you have successful IRC calculation, open **Gaussview** for visualization
  - open the file using **gv** {filename}
  - go to menu bar **Results** > **IRC**, it allows you to view **Total Energy** and **RMS Gradient** along IRC values are presented
  - you can visualize the molecular movements by clicking the animate play button



# ASSIGNMENT

- This process concerns the proton transfer in malonaldehyde, find the transition state geometry:



- use DFT methods PBE, B3LYP, and then try MP2. Be sure to use cc-pVDZ basis set. Verify your obtained transition state geometry.

**END**