# Single Point Energies And Geometry Optimizations

## **Single Point Energy Calculations**

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 Solution of Schrödinger Equation for Molecule with Specified Geometric Structure

- ► *E* and geometry
- v and thermodynamic properties
- ► NMR
- ► Electronic distributions
- Other static properties
- Validity
  - ► Reasonable structure
- Choice of level
- Choice of basis set

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#### Purpose

- Only "affordable" calculation
- Starting point for optimization
- Estimate time of optimization
- Calculate specific molecular properties after optimization
- Calculate accurate values of *E* and related properties at a higher level of theory following lower level optimization

## **Potential Energy Surfaces (PES)**

E and other properties are function of geometry

- Mathematical relationship of E as a function of structure
- Surface has as many dimensions as number of internal degrees of freedom in molecule
  - Diatomic molecule AB
  - $-r_{AB}$
  - Nonlinear triatomic molecule ABC
  - $-r_{AB}$  and  $r_{BC}$  (unless identical) - A-B-C bond angle
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## **PES Features**

Minima

- Bottom of "Valley" on PES
  Changing any geometric parameter increases E
  Equilibrium structures of molecule
- Different conformers
- Structural isomers
- Reactant and product



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#### **Geometry Optimization**

- Locating Extremum of PES
  - Minimization gives equilibrium structure(s) Saddle point corresponds to transition structure
- Calculus at Extremum for Multidimensional Function

  - Gradient = g<sub>i</sub> = ∂E/∂v<sub>i</sub> = 0
     Force = -(gradient) = 0
  - "stationary point" is location on PES where forces = 0
  - Force constant =  $\partial^2 E / \partial \mathbf{v}_i^2$
  - Specifies "curvature" of surface at point
  - All > 0 at stationary point
  - All > 0 except one < 0 at (1st order) saddle point</li>
    "Hessian" is matrix of second derivatives



#### Analysis

- Convergence criteria
- Forces = 0 or less than chosen cut-off value
   Next structure change is very small or below chosen cut-
- off value – Often additional criteria are used
- Optimization methods for structure change based on decrease in E
- Use gradient and force constant values
- More later...

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#### Optimization Results

- ▶ Was a minimum determined?
- Check by doing a single point calculation of vibrational frequencies (all positive)
- Was a global minimum determined?
- Check PES map
- More later...
- Was a saddle point determined?
- Check by doing a single point calculation of vibrational frequencies (all positive except one is negative/imaginary)

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## **Optimization Methods**

"Simple" Method

- Procedure
  - Minimize E with respect to only one variable holding all others constant
  - Change variable to optimum value
  - Repeat minimization with respect to a second variable holding all others constant
  - Change second variable to optimum value
  - Continue for each variable

#### Problems

- Variables are not independent
   Linear molecules like acetylene are often problems
- Several cycles are needed
   "Evpapaive" for large males
- "Expensive" for large molecules
- "Real" methods
  - Use all variables
  - ► Three general approaches: SD, CG, NR

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## **Optimization Methods**

Steepest Descent (SD)

- Change coordinates in direction opposite the maximum of gradient using a unit vector s<sub>i</sub> = -g<sub>i</sub>/|g<sub>i</sub>|
- Line search method for distance
  - Find 3 points such that inner point is lower in E than outer points
- Thus at least one minimum lies between outer points
- Iterate decreasing distance
- Recalculate g<sub>i+1</sub> and repeat



- Advantages of SD Always locate (local) minimum
- Problems of SD
  - ► **g**<sub>i+1</sub> is perpendicular to **g**<sub>i</sub> Lose any further *E* lowering possible by **g**,
    Path oscillates around minimum path
- Rate of convergence decreases near minimum
- Undesireable behavior in long, narrow valley

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#### Modifications of SD

- ▶ Fit 3 points to function
- Differentiate to find minimum
- Arbitrary step size
- Start with predetermined value or value proportional to gi
- Calculate E at new position
- If E is less
  - Increase step size
- Calculate E at new position
- Repeat until E increases
- If E is greater or increases
- Decrease step size
- Calculate E at new position
- Iterate

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## **Optimization Methods**

Conjugate Gradient Method (CG)

- Mixture of current gradient and previous search direction
- Procedure
- ► First step: SD
- ► Subsequent steps:  $\mathbf{v}_{i+1} = -\mathbf{g}_{i+1} + \gamma_{i+1} \mathbf{v}_i$
- Fletcher-Reeves:  $\gamma_{i+1} = (\mathbf{g}_{i+1} \cdot \mathbf{g}_{i+1})/(\mathbf{g}_i \cdot \mathbf{g}_i)$
- $\text{Polak-Ribiere (common): } \gamma_{i+1} = [(\mathbf{g}_{i+1} \mathbf{g}_i) \bullet \mathbf{g}_{i+1}]/(\mathbf{g}_i \bullet \mathbf{g}_i) \\ \text{Hestenes-Stifel: } \gamma_{i+1} = [(\mathbf{g}_{i+1} \mathbf{g}_i) \bullet \mathbf{g}_{i+1}]/[\mathbf{v}_i \bullet (\mathbf{g}_{i+1} \mathbf{g}_i)]$



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Advantages of CG

- Better convergence
- Always locate (local) minimum
- Problems of CG
  - Require additional computer resources to store g<sub>i</sub> and g<sub>i+1</sub>
  - Expense scales as N<sup>2</sup>

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## **Optimization Methods**

Newton-Raphson (NR)

- Includes Hessian
- Advantages
  - Locates minima and saddle points
  - Convergence is second order near stationary point
- Problems
  - Control of step size
  - Calculation of Hessian required
- Several variations of NR

## **Global Minimum**

- Optimization Results
  - Above methods locate the "nearest" minimum (local?)
  - Conformers and transition structures are related to local minima and saddle points
  - Molecular properties are related to global minimum--the most stable structure
  - Several approaches to finding the global minimum – NO Guarantees!!!

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#### **Global Minimum Methods**

Grid Search

- Simple
  - Calculate all possible energies
  - Prepare energy map or table
  - Choose best structure
- Problems
  - Number of minima increases exponentially with the number of variables
- Not practical for large or biomolecules
   Build structures from optimized fragments
- Optimize portions of molecule



#### Example: $CH_3(CH_2)_{n+1}CH_3$

<i>n</i> = 1	$N = 3^1 = 3$	<i>t</i> = 3 s
n = 2	$N = 3^2 = 9$	<i>t</i> = 9 s
n = 5	<i>N</i> = 3 <sup>5</sup> = 243	<i>t</i> = ~4 min
<i>n</i> = 10	<i>N</i> = 3 <sup>10</sup> = 59 k	<i>t</i> = ~16 h
<i>n</i> = 15	<i>N</i> = 3 <sup>15</sup> = 14 M	<i>t</i> = ~160 d

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#### **Global Minimum Methods**

Other

- Monte Carlo
  - Usually start at a minimum
  - Randomly change one or several angles or bond lengths to generate new geometry
  - ► Calculate E
  - If *E* is lower, accept new geometry
  - If E is greater, retain or reject based on a Boltzmann distribution related to the energy change and T
  - Many variations

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#### Molecular Dynamics

- Based on Newton's equations of motion for atoms
- Molecule overcomes barrier between minima if kinetic energy is great enough
- Similar to climbing out a valley over a mountain
  Kinetic energy is proportional to T
- Use elevated temperatures (600 1200 K)
- High temperatures reduce chance of trapping molecule in a local minimum
- Essentially searches PES in the region of the starting minimum

- Simulated Annealing
  - Molecular dynamics variation
  - Start with high T (2000 3000 K)
  - ▶ Reduce T
  - Molecule trapped in a minimum
     Similar to playing roulette
     Very slow cooling might give global minimum
     Repeat to get many results
  - Choose best
- Genetic Algorithm
- Use several low energy structures as "parents" to generate "off spring" that might be better
- Distance Geometry
- Based on establishing minimum and maximum distances between all pairs of atoms