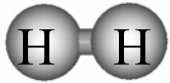


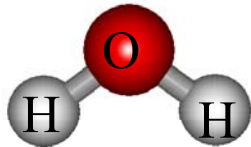
Introduction to Geometry Optimization

**Computational Chemistry lab
2014**

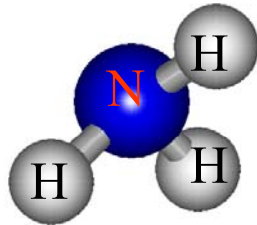
Determination of the molecule configuration



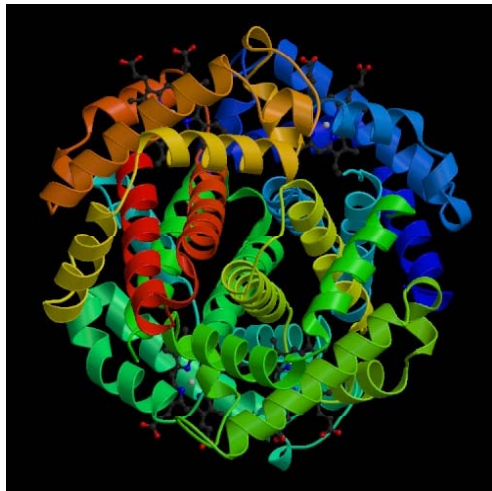
- Diatomic molecule – determine the interatomic distance



- Triatomic molecule – determine distances and the angle



- Four-atomic molecule – determine angles and distances



$3N - 6$ parameters for an N -atomic molecule ($N > 2$)
($3N$ atomic coordinates - 3 c.o.m coordinates - 3 angles)

Optimal configurations correspond to minima of the energy as a function of these parameters

Solution of the Schrödinger equation in the Born-Oppenheimer approximation:

$$H_{tot} = (T_n + V_n) + T_e + V_{ne} + V_e = (H_n) + H_e$$

Electronic SE: $H_e \Psi(r, R) = E_e(R) \Psi(r, R)$

Ab-initio or *semi-empirical* solution

$E_e(R)$ – **electronic energy**

R – **nuclear coordinates**

$V_n(R)$ – **nuclei-nuclei interaction**

$E(R) = V_n(R) + E_e(R) = \text{potential energy surface (PES)}$
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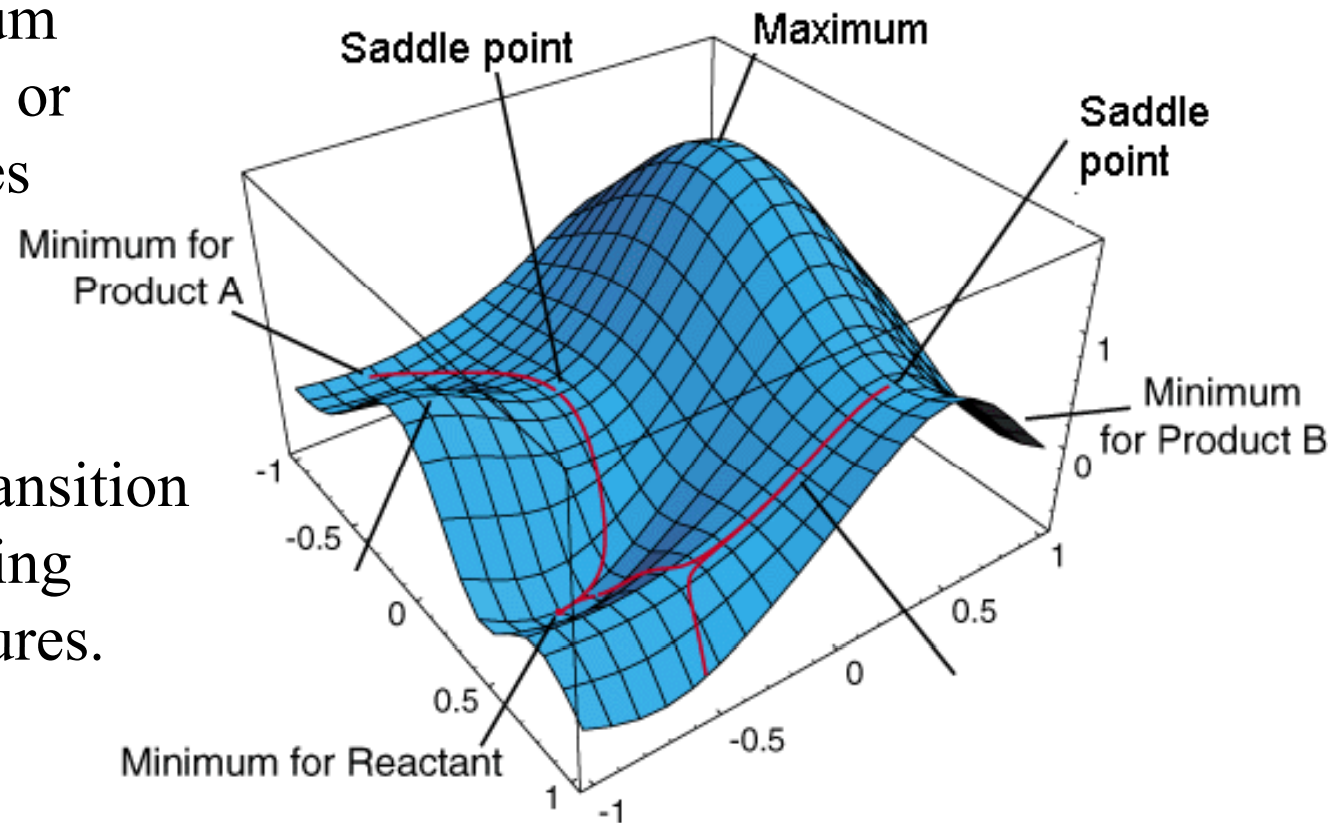
Molecular mechanics: PES ← molecular force field

Potential Energy Surface (PES)

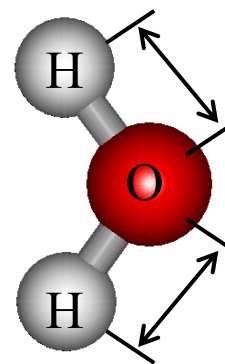
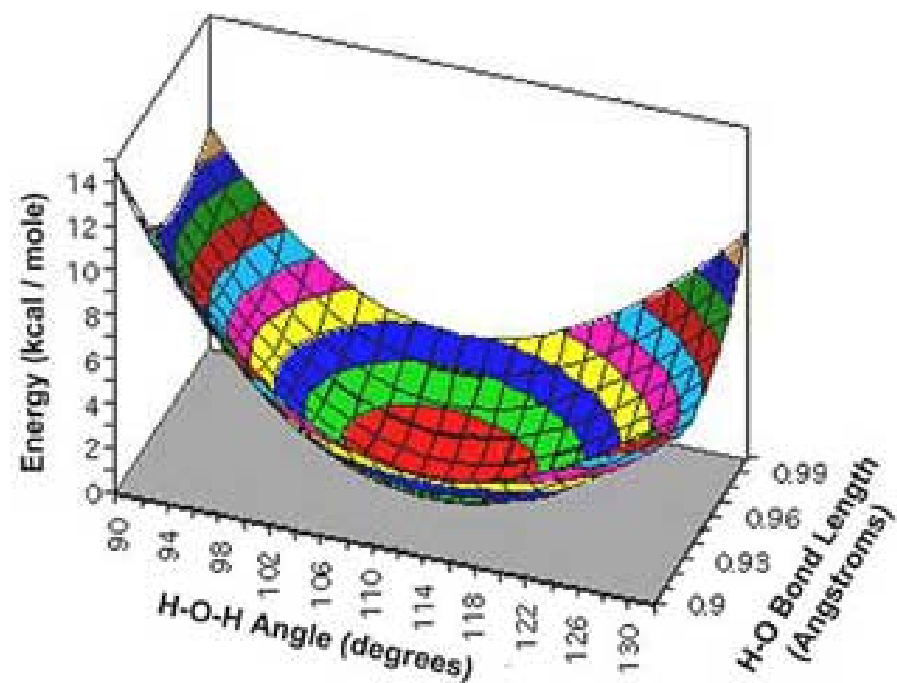
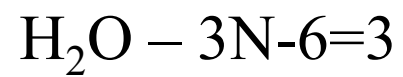
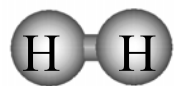
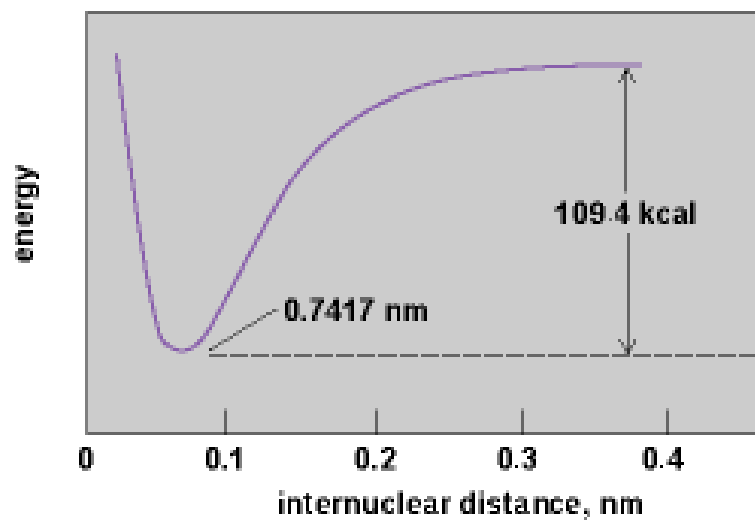
A hypersurface in $3N-6$ -dimensional hyperspace, where N is the number of atoms.

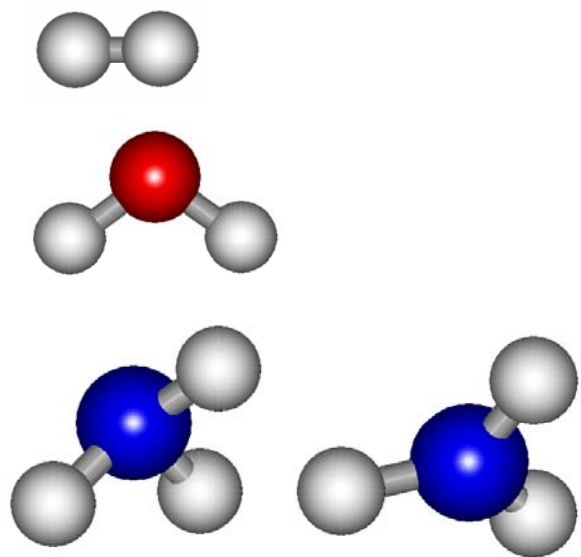
Minima— equilibrium structures of one or several molecules

Saddle points - transition structures connecting equilibrium structures.



Global Minimum - that point that is the lowest value in the PES

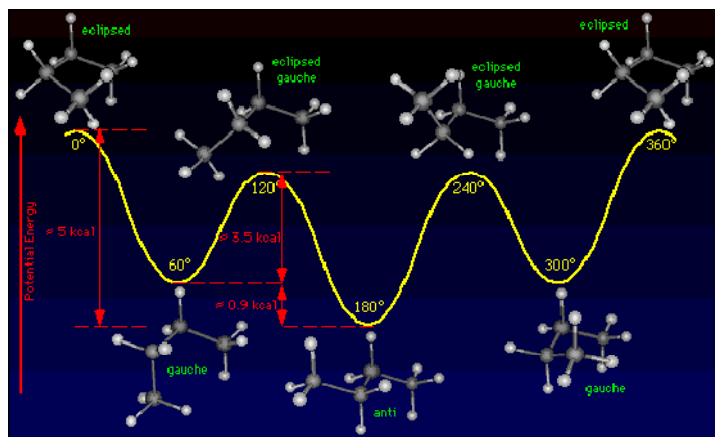




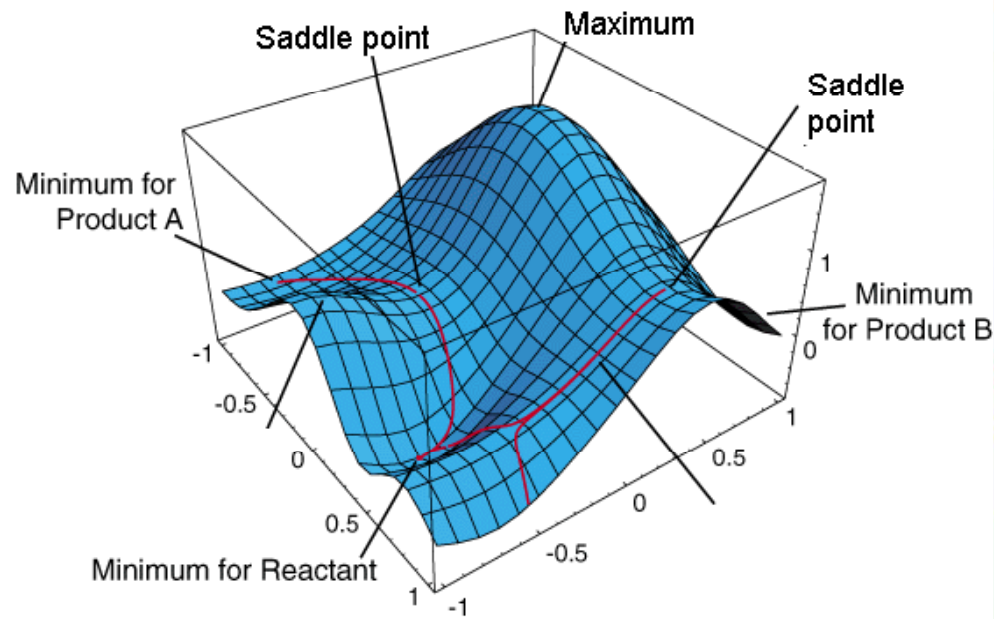
- One configuration is known for H₂
- One configuration is known for H₂O
- The planar and pyramidal configurations are known for NH₃

The number of local minima typically goes exponentially with the number of variables (degrees of freedom). **Combinatorial Explosion Problem**

Possible Conformations (3^n) for linear alkanes CH₃(CH₂)_{n+1}CH₃



$n = 1$	3
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$n = 10$	59,049
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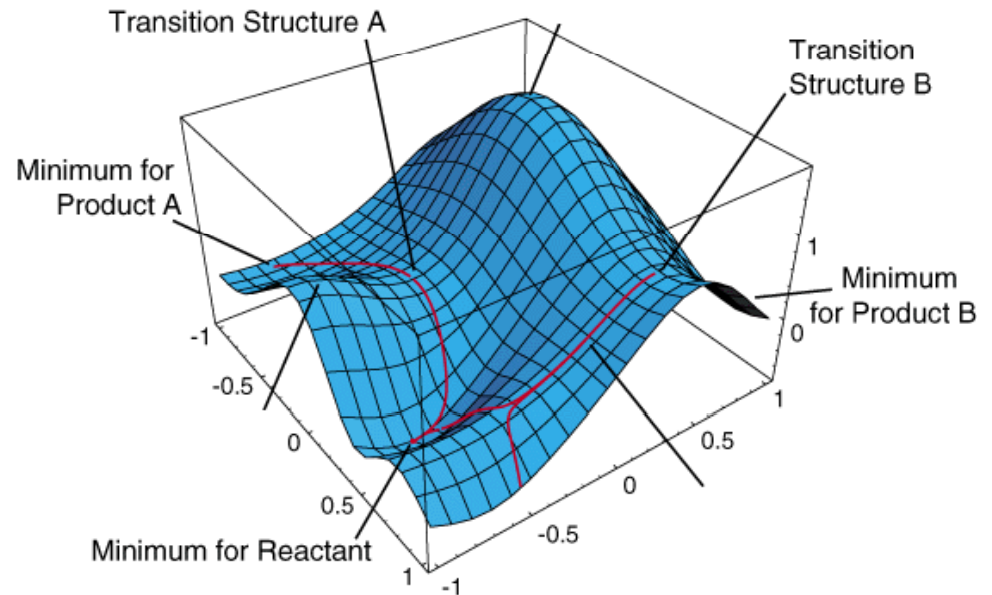


Global minimum
- Mariana Trench
 in the Pacific Ocean

Optimization/Goals

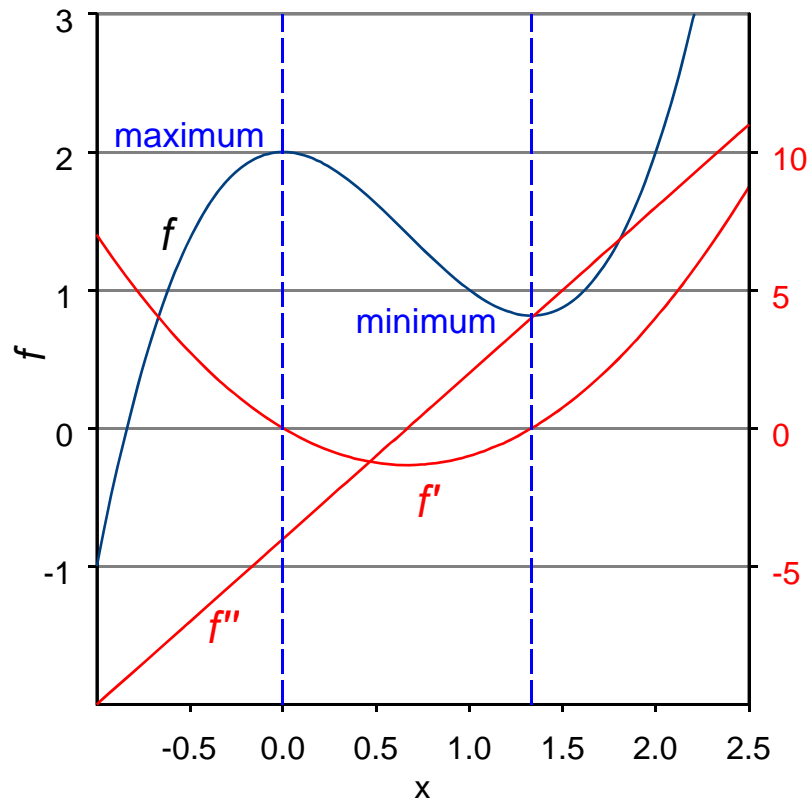
- Find the **Local Minimum** Structure
- Find the **Global Minimum** Structure
- Find the **Transition State** Structure

We would like to Find the **Reaction Pathway** Connecting Two Minima and Passing through the Transition State Structure



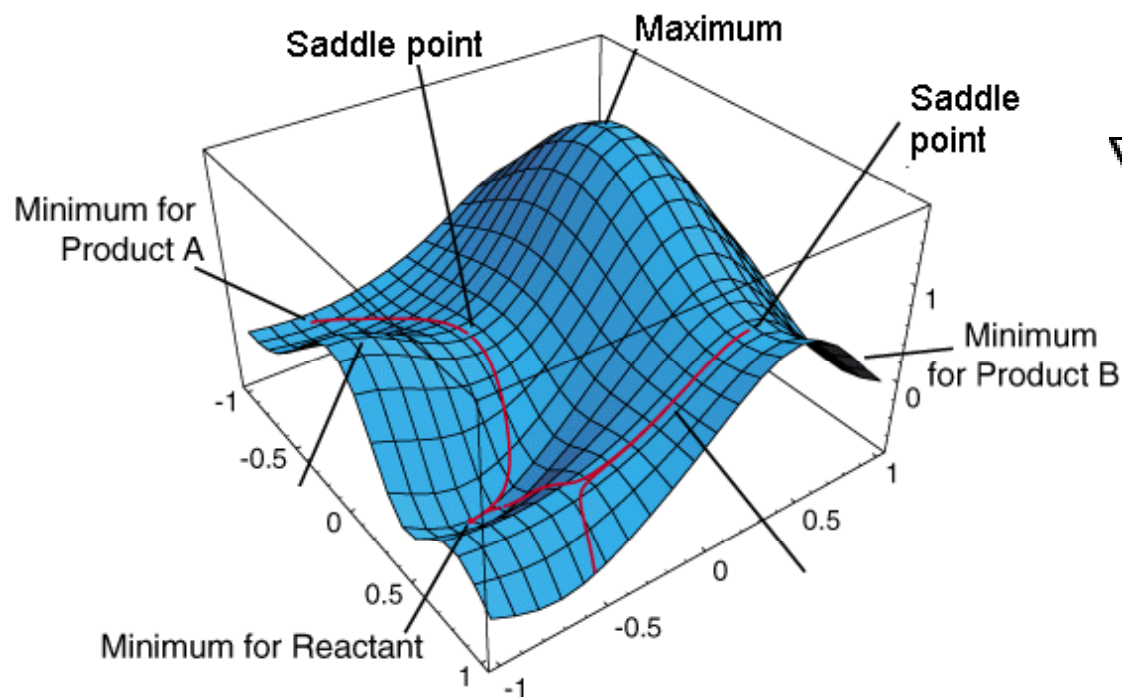
Geometry optimization is the name for the procedure that attempts to find the configuration of minimum energy of the molecule.

One-dimensional optimization



- $f'(x)=0$ – stationary point
- $f''(x)<0$ – maximum
- $f''(x)>0$ – minimum

Multidimensional Optimization



$$\nabla f = \left(\frac{\partial f}{\partial x_1}, \dots, \frac{\partial f}{\partial x_n} \right)$$

- gradient

The coordinates can be Cartesian (x,y,z..), or internal, such as bondlength and angle displacements)

Force: $F = -\nabla E$ (Potential energy)

$\nabla f = 0$ - Stationary Point (minimum, maximum, or saddle point)

Hessian matrix

A matrix of second-order derivatives of the energy with respect to atomic coordinates (e.g., Cartesian or internal coordinates)

Sometimes called force matrix – matrix size of $(3N-6) \times (3N-6)$

$$H_{ij}(f) = \frac{\partial^2 f}{\partial x_i \partial x_j}$$

Quadratic approximation

Approximate the complex energy landscape by harmonic potentials

around a stationary point $(x_1^{(st)}, \dots, x_n^{(st)})$ $[\nabla E(x_1^{(st)}, \dots, x_n^{(st)}) = 0]$

$$E(x_1, \dots, x_n) = E(x_1^{(st)}, \dots, x_n^{(st)}) + \frac{1}{2} \sum_{i=1}^n \sum_{j=1}^n H_{ij}(E)(x_i - x_i^{(st)})(x_j - x_j^{(st)})$$

$$H(f) = \begin{bmatrix} \frac{\partial^2 f}{\partial x_1^2} & \frac{\partial^2 f}{\partial x_1 \partial x_2} & \cdots & \frac{\partial^2 f}{\partial x_1 \partial x_n} \\ \frac{\partial^2 f}{\partial x_2 \partial x_1} & \frac{\partial^2 f}{\partial x_2^2} & \cdots & \frac{\partial^2 f}{\partial x_2 \partial x_n} \\ \vdots & \vdots & \ddots & \vdots \\ \frac{\partial^2 f}{\partial x_n \partial x_1} & \frac{\partial^2 f}{\partial x_n \partial x_2} & \cdots & \frac{\partial^2 f}{\partial x_n^2} \end{bmatrix}$$

$$E(x_1, \dots, x_n) = E(x_1^{(st)}, \dots, x_n^{(st)}) + \frac{1}{2} \sum_{i=1}^n \sum_{j=1}^n H_{ij}(E)(x_i - x_i^{(st)})(x_j - x_j^{(st)})$$

Hessian matrix diagonalization (eigenproblem) $\varepsilon_k l_j^{(k)} = \sum_{i=1}^n H_{ij} l_i^{(k)}$

Eigenvalues $\varepsilon_k = m\omega_k^2$ ω_k - vibrational frequencies

Eigenvectors $l_j^{(k)}$ give **normal coordinates** $q_k = \sum_{i=1}^n l_i^{(k)}(x_i - x_i^{(st)})$

$$E(x_1, \dots, x_n) = E(x_1^{(st)}, \dots, x_n^{(st)}) + \frac{1}{2} \sum_{k=1}^n \varepsilon_k q_k^2$$

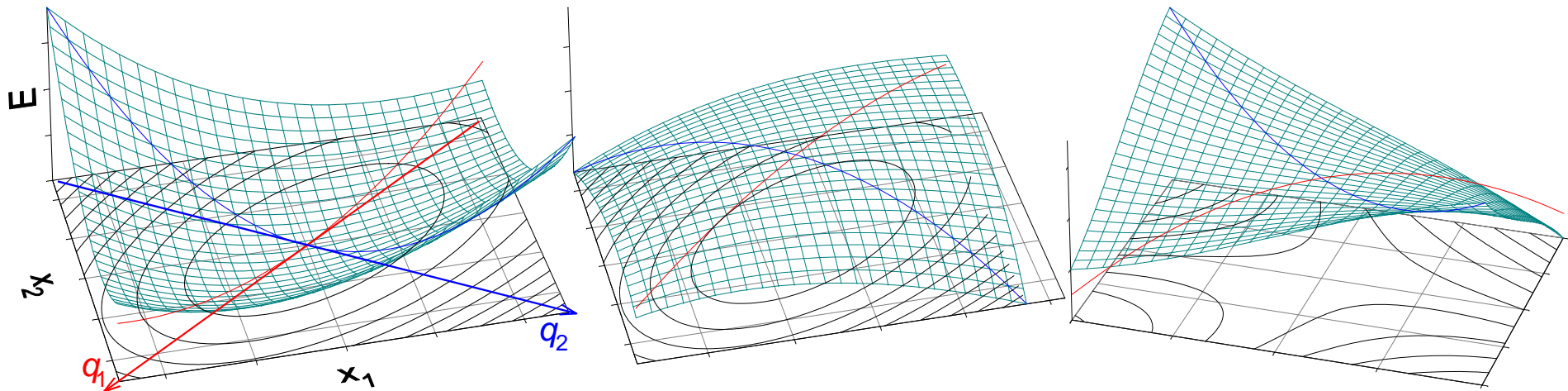
All $\varepsilon_k > 0$ (ω_k - real)

minimum

All $\varepsilon_k < 0$ (ω_k - imaginary) $\varepsilon_k \leftrightarrow 0$ (ω_k - imaginary/real)

maximum

saddle point



Steepest Descent

Steepest descent direction $\mathbf{g} = -\nabla E$

$\mathbf{r} = (x_1, \dots, x_n)$, \mathbf{g} - $(3N-6)$ -dimensional vectors

$$\mathbf{r}_{i+1} = \mathbf{r}_i + \lambda_i \mathbf{g}_i / |\mathbf{g}_i|$$

Advantage:

- local minimization is guaranteed
- fast minimization far from the minimum

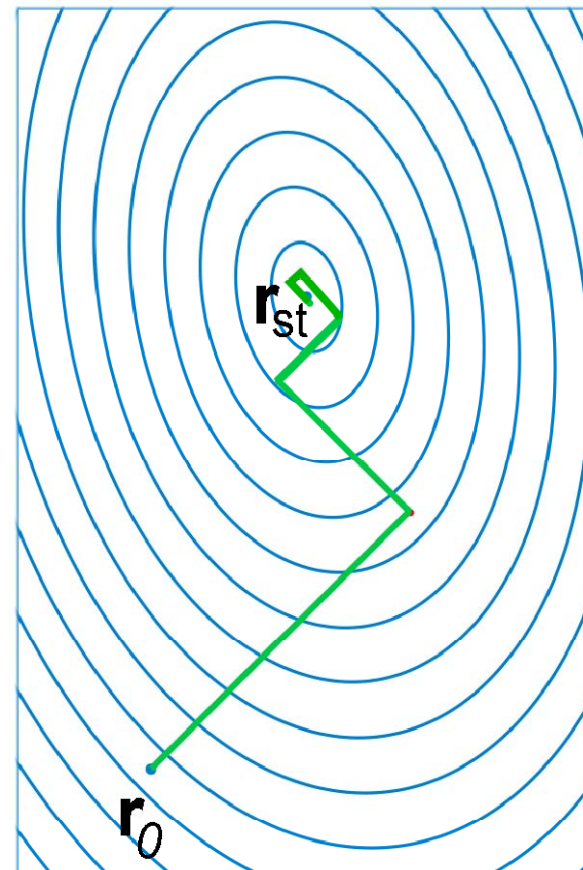
Disadvantage:

- slow descent along narrow valleys
- slow convergence near minima

In HYPERCHEM

$$\text{if } E(\mathbf{r}_{i+1}) < E(\mathbf{r}_i) \text{ then } \lambda_{i+1} = 1.2\lambda_i \text{ else } \lambda_{i+1} = 0.5\lambda_i$$

History gradient information is not kept



Conjugate Gradient Methods

Search direction is chosen using history gradient information

Initial direction – the steepest descent direction, $\mathbf{h}_0 = \mathbf{g}_0 = -\nabla E$

$$\mathbf{r}_{i+1} = \mathbf{r}_i + \xi \mathbf{h}_i$$

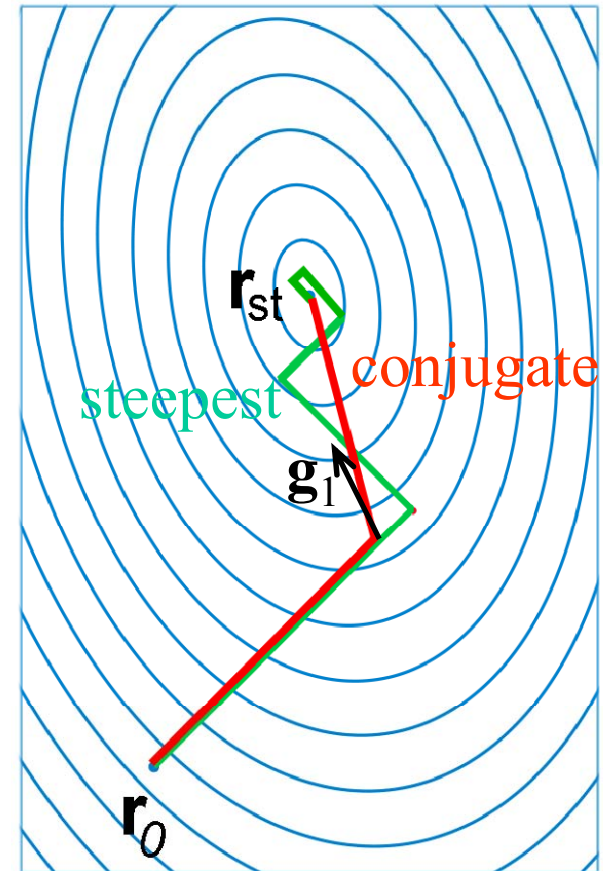
ξ is defined by a one-dimensional minimization along the search direction.

$$\mathbf{h}_{i+1} = \mathbf{g}_{i+1} + \gamma_{i+1} \mathbf{h}_i$$

Fletcher-Reeves method:
$$\gamma_{i+1} = \frac{|\mathbf{g}_{i+1}|^2}{|\mathbf{g}_i|^2}$$

Polak-Ribiere method:
$$\gamma_{i+1} = \frac{(\mathbf{g}_{i+1} - \mathbf{g}_i) \cdot \mathbf{g}_{i+1}}{|\mathbf{g}_i|^2}$$

Polak-Ribiere may be superior for non-quadratic functions.



Gradient history is equivalent to implicit use of the Hessian matrix.

Newton-Raphson Methods

Explicit use of Hessian matrix.

Quadratic approximation :
$$E(\mathbf{r}) = E(\mathbf{r}_0) - \sum_{k=1}^n (F_k q_k - \frac{1}{2} \varepsilon_k q_k^2)$$

q_k - normal coordinates,
$$\mathbf{r} - \mathbf{r}_0 = \sum_{k=1}^n q_k \mathbf{l}_k \quad \nabla E(\mathbf{r}_0) = - \sum_{k=1}^n F_k \mathbf{l}_k$$

ε_k and \mathbf{l}_k - eigenvalues and eigenvectors of the Hessian matrix at \mathbf{r}_0

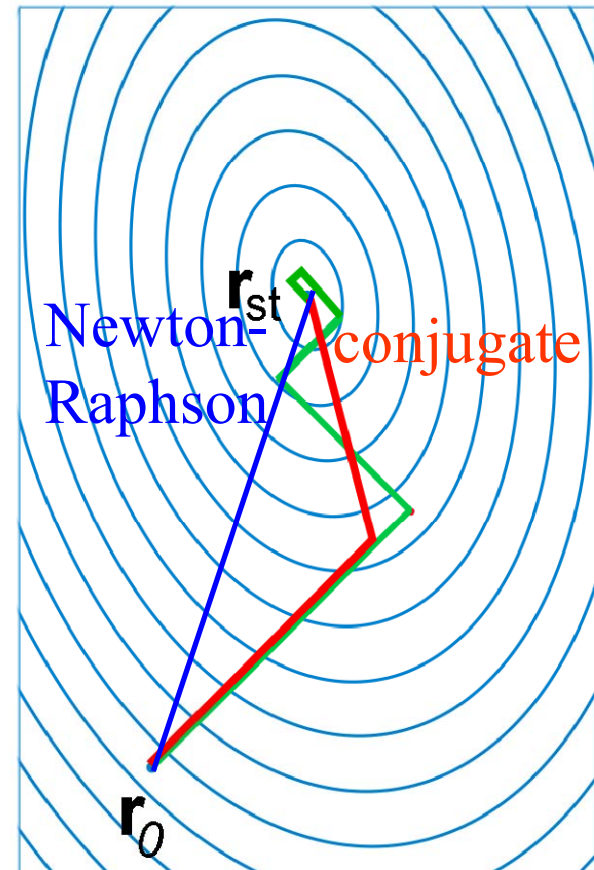
One-step optimization of quadratic functions $q_k = \frac{F_k}{\varepsilon_k} \implies \nabla E(\mathbf{r}) = 0$

For arbitrary functions

$$\mathbf{r}_{i+1} = \mathbf{r}_i + \sum_{k=1}^n \mathbf{l}_k(\mathbf{r}_i) F_k(\mathbf{r}_i) / \varepsilon_k(\mathbf{r}_i)$$

Descent direction for $\varepsilon_k(\mathbf{r}_i) > 0$

Finds the closest stationary point (either minimum, maximum, or saddle point).



Obtaining, storing, and diagonalizing the Hessian

Numerical calculation – $(3N)^2$ calculations of energy

Diagonalization - $(3N)^3$ operations

Memory - $(3N)^2$

Extremely expensive for large molecules.

Block Diagonal Newton-Raphson

$$\left(\begin{array}{ccc|ccc} \frac{\partial^2 E}{\partial x_1^2} & \frac{\partial^2 E}{\partial x_1 \partial y_1} & \frac{\partial^2 E}{\partial x_1 \partial z_1} & 0 & 0 & 0 \\ \frac{\partial^2 E}{\partial x_1 \partial y_1} & \frac{\partial^2 E}{\partial y_1^2} & \frac{\partial^2 E}{\partial y_1 \partial z_1} & 0 & 0 & 0 \\ \frac{\partial^2 E}{\partial x_1 \partial z_1} & \frac{\partial^2 E}{\partial y_1 \partial z_1} & \frac{\partial^2 E}{\partial z_1^2} & 0 & 0 & 0 \\ \hline 0 & 0 & 0 & \frac{\partial^2 E}{\partial x_2^2} & \frac{\partial^2 E}{\partial x_2 \partial y_2} & \frac{\partial^2 E}{\partial x_2 \partial z_2} \\ 0 & 0 & 0 & \frac{\partial^2 E}{\partial x_2 \partial y_2} & \frac{\partial^2 E}{\partial y_2^2} & \frac{\partial^2 E}{\partial y_2 \partial z_2} \\ 0 & 0 & 0 & \frac{\partial^2 E}{\partial x_2 \partial z_2} & \frac{\partial^2 E}{\partial y_2 \partial z_2} & \frac{\partial^2 E}{\partial z_2^2} \end{array} \right)$$

Calculation - $9N$

Diagonalization - $27N$

Memory – $9N$

Gaussian

The Hessian matrix is calculated and processed on the first step only
It is updated using the computed energies and gradients

Berny algorithm

**Rational function
optimization (RFO) step**

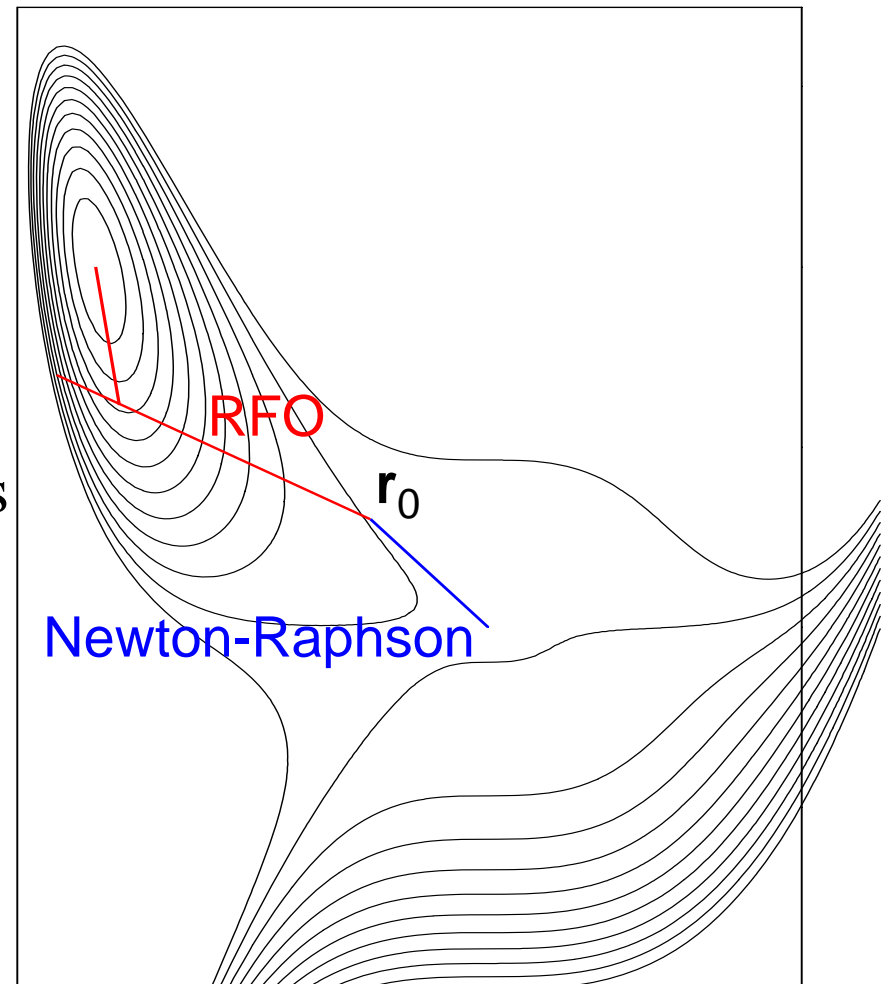
$$\mathbf{r}_{i+1} = \mathbf{r}_i + \sum_{k=1}^n \mathbf{l}_k(\mathbf{r}_i) \frac{F_k(\mathbf{r}_i)}{\varepsilon_k(\mathbf{r}_i) - \lambda}$$

ε_k and \mathbf{l}_k - eigenvalues and eigenvectors
of the Hessian matrix at \mathbf{r}_i

$\lambda < \min(\varepsilon_k)$ ($\lambda < 0$ for $\varepsilon_k < 0$)

- step always toward to a minimum.

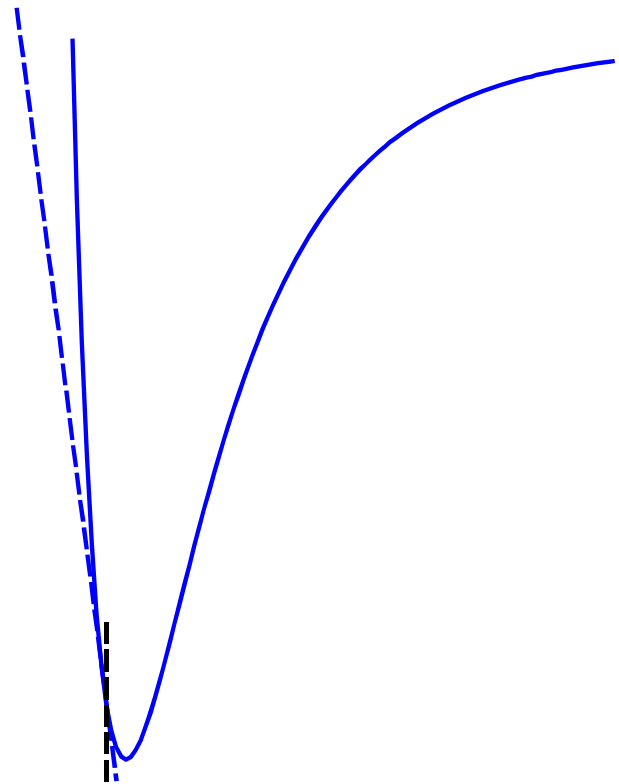
A **linear search** between the latest
point and the best previous point.



Convergence criteria

Gradient or force $\mathbf{F} = -\mathbf{g} = -\nabla E$

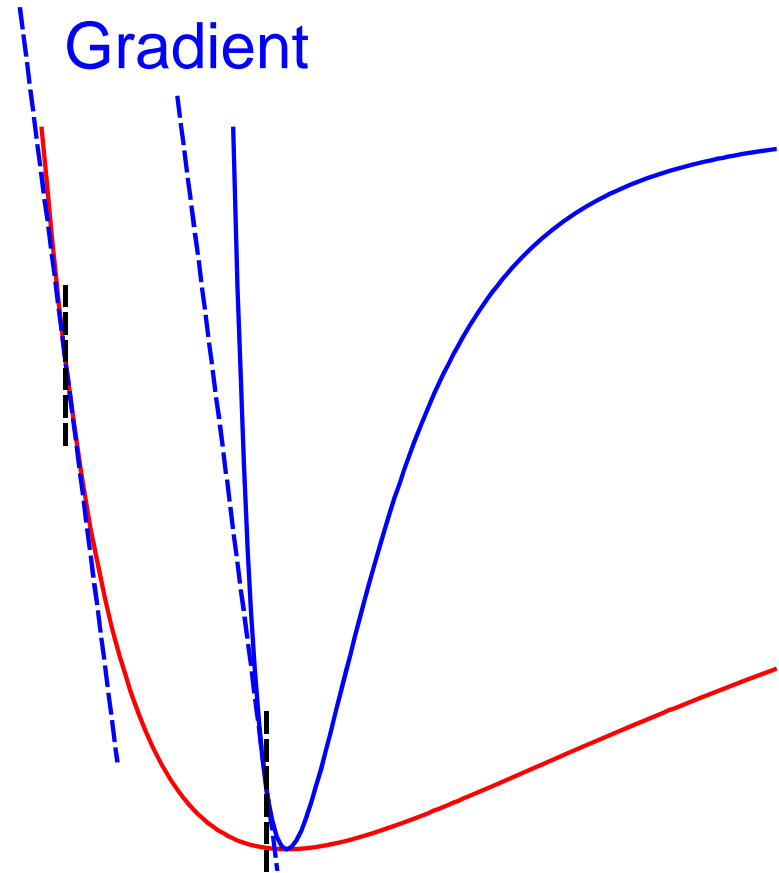
Gradient



Convergence criteria

Gradient or force $\mathbf{F} = -\mathbf{g} = -\nabla E$

Gradient



Convergence criteria

Gradient or force $\mathbf{F} = -\mathbf{g} = -\nabla E$

Predicted displacement $\mathbf{r}_{i+1} - \mathbf{r}_i$

Maximum force:

$$\max(\mathbf{F}) = \max(F_1, \dots, F_n) < \text{threshold}$$

Root-mean-square (RMS) force:

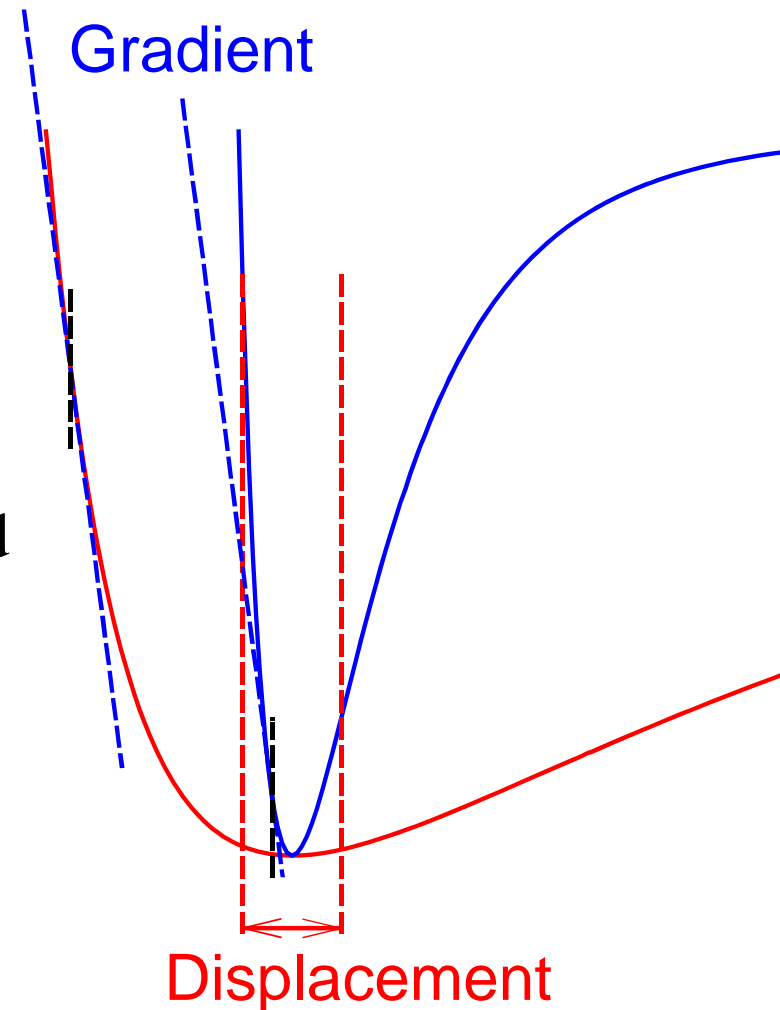
$$|\mathbf{F}| = \sqrt{(F_1^2 + \dots + F_n^2) / n} < \text{threshold}$$

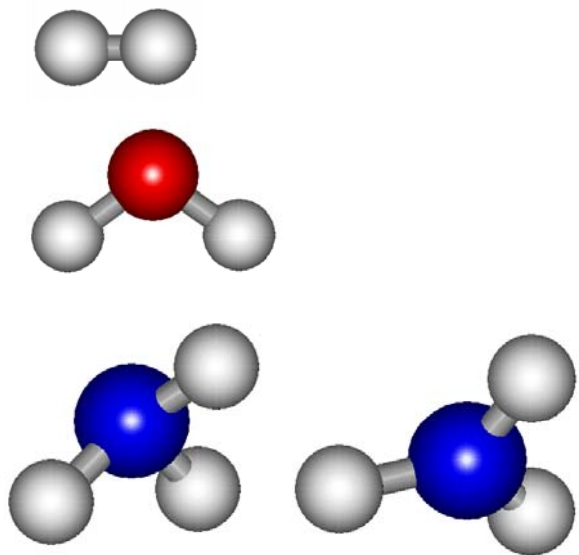
Maximum displacement:

$$\max(\mathbf{r}_{i+1} - \mathbf{r}_i) < \text{threshold}$$

RMS displacement:

$$|\mathbf{r}_{i+1} - \mathbf{r}_i| / \sqrt{n} < \text{threshold}$$

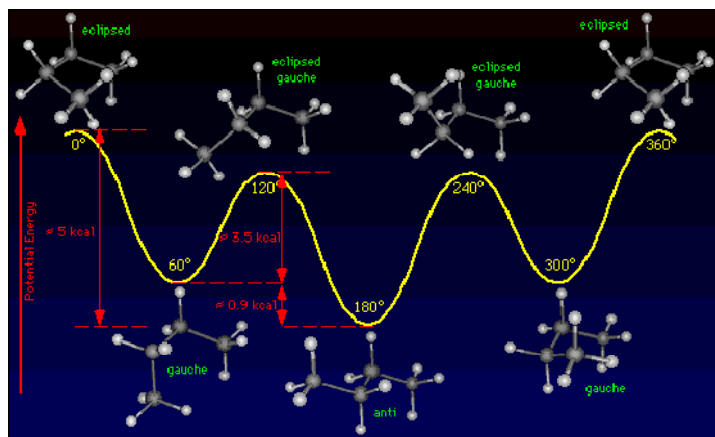




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Global minimization

- **"building up" the structure:** combining a large molecule from preoptimized fragments (protein from preoptimized aminoacids)
- **conformational sampling:** take various starting points for local minimization (simulation of nature)
- **None of these are guaranteed to find the global minimum!**

Global Minimum- conformational sampling

- Molecular Dynamics
- Monte Carlo
- Simulated Annealing
- Tabu search
- Genetic Algorithms
- Ant Colony Optimizations
- Diffusion Methods
- Distance Geometry Methods

