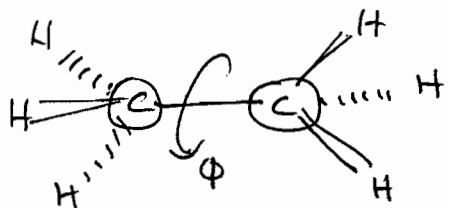


Potential Energy Surface

Once we have chosen a force field, we can analyse the conformations and dynamics of molecules of interest.

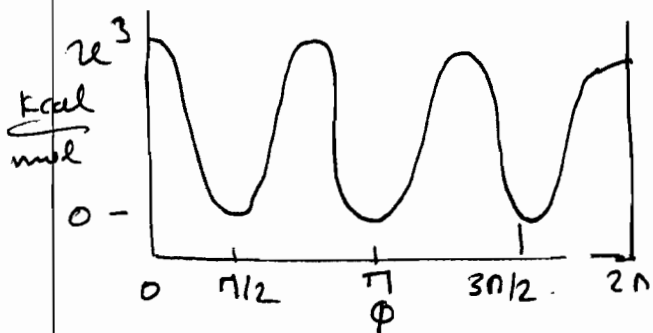
In one dimension, things are simple



e.g. rotation of $\text{CH}_3\text{-CH}_3$
Single bond in ethane

minima at $60^\circ, 180^\circ, 240^\circ$

maxima " $0^\circ, 120^\circ, 240^\circ$



In general

$f(x)$ has extremum if

$$\frac{df}{dx} = 0 \quad \left\{ \begin{array}{l} \text{to check} \\ \text{max or min,} \end{array} \right. \rightarrow \left. \begin{array}{l} \frac{d^2f}{dx^2} > 0 \text{ min;} \\ \frac{d^2f}{dx^2} < 0 \text{ max;} \end{array} \right\}$$

For functions of many variables, things are not so simple; for N atoms, we have in principle $3N$ degrees of freedom (variables), but 6 are trivial (position + rotation of molecule as a whole). There are $3N-6$ internal coordinates

$$U = U(\vec{q}) \quad \text{where} \quad \vec{q} = \begin{pmatrix} q_1 \\ q_2 \\ \vdots \\ q_p \end{pmatrix} \quad \begin{array}{l} p = 3N-6 \\ (3N-5 \text{ for} \\ \text{linear}) \end{array}$$

A stationary point of the function u is one for which

$$\vec{g} = \nabla u = \begin{pmatrix} (\partial u / \partial q_1)_{q_2, \dots, q_p} \\ (\partial u / \partial q_2)_{q_1, q_3, \dots, q_p} \\ \vdots \\ (\partial u / \partial q_p)_{q_1, q_2, \dots, q_{p-1}} \end{pmatrix} = 0$$

Of course, we do not know if a stationary point is a maximum or a minimum until we check the eigenvalues of the Hessian matrix:

$$H = \begin{pmatrix} \frac{\partial^2 u}{\partial q_1^2} & \dots & \frac{\partial^2 u}{\partial q_1 \partial q_p} \\ \vdots & & \vdots \\ \frac{\partial^2 u}{\partial q_p \partial q_1} & \dots & \frac{\partial^2 u}{\partial q_p^2} \end{pmatrix} \quad \begin{array}{l} \text{if all eigenvalues} \\ > 0 \Rightarrow \text{min} \\ \\ \text{if all eigenvalues} \\ < 0 \Rightarrow \text{min} \end{array}$$

if some are < 0 and some $> 0 \Rightarrow$ "saddle point"

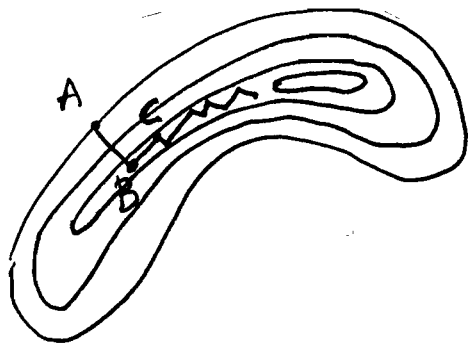
Finding Minima

For functions of many variables, it is impractical to try to evaluate them on a grid. Instead, one uses sophisticated numerical methods developed over many years by applied mathematics researchers.

A few common methods:

- (a) First-order methods: only use $\vec{g} = \nabla u$
the gradient of u

Steepest descent: calculate $\nabla U = \vec{g}$, move along $-\vec{g}$ to a local (linear) minimum; repeat.



$A \rightarrow B \rightarrow C \rightarrow \dots$

Can take a long time to locate minimum; fairly robust

Conjugate Gradient (Fletcher-Reeves, 1964)

iteration (k) is at $\vec{q}^{(k)}$; move along $\vec{V}^{(k)} = -\vec{g}^{(k)} + \gamma^{(k)} \vec{V}^{(k-1)}$ where $\gamma^{(k)} = \frac{(\vec{g}^{(k)})^T \vec{g}^{(k)}}{(\vec{g}^{(k-1)})^T \vec{g}^{(k-1)}}$

Method is faster than steepest descent - does not move just at right angles

1) Second-order Methods

These require calculation of the Hessian, analytically or numerically; they are faster, but may fail to converge if the initial guess is poor.

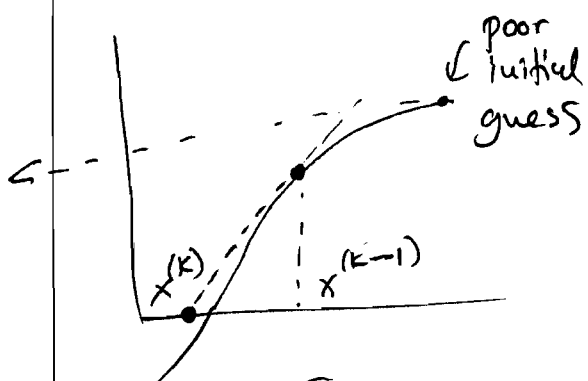
Newton-Raphson

Solves $f(x) = 0$ by setting

$$x^{(k)} = x^{(k-1)} - \frac{f(x)}{f'(x)}$$

← derivative in 1D

For minimization, need $f'(x) = 0$



$$\vec{x}^{(k)} = \vec{x}^{(k-1)} - \left(\underline{H}^{(k-1)} \right)^{-1} \cdot \nabla f$$

(Gives quadratic (fast) convergence near solution)

All energy minimization methods require a numerical criterion for ending the calculation -

- e.g. * maximum tolerance for final gradient
- * maximum number of iterations

Intro to Hyperchem

(from outline notes)

- creating a molecule
- adding H
- selecting a force field, looking at parameters
- energy minimizing the structure
- measuring molecular parameters (bond lengths, angles)
- 1D potential
- Simple "grid" properties: molecular volume + surface area