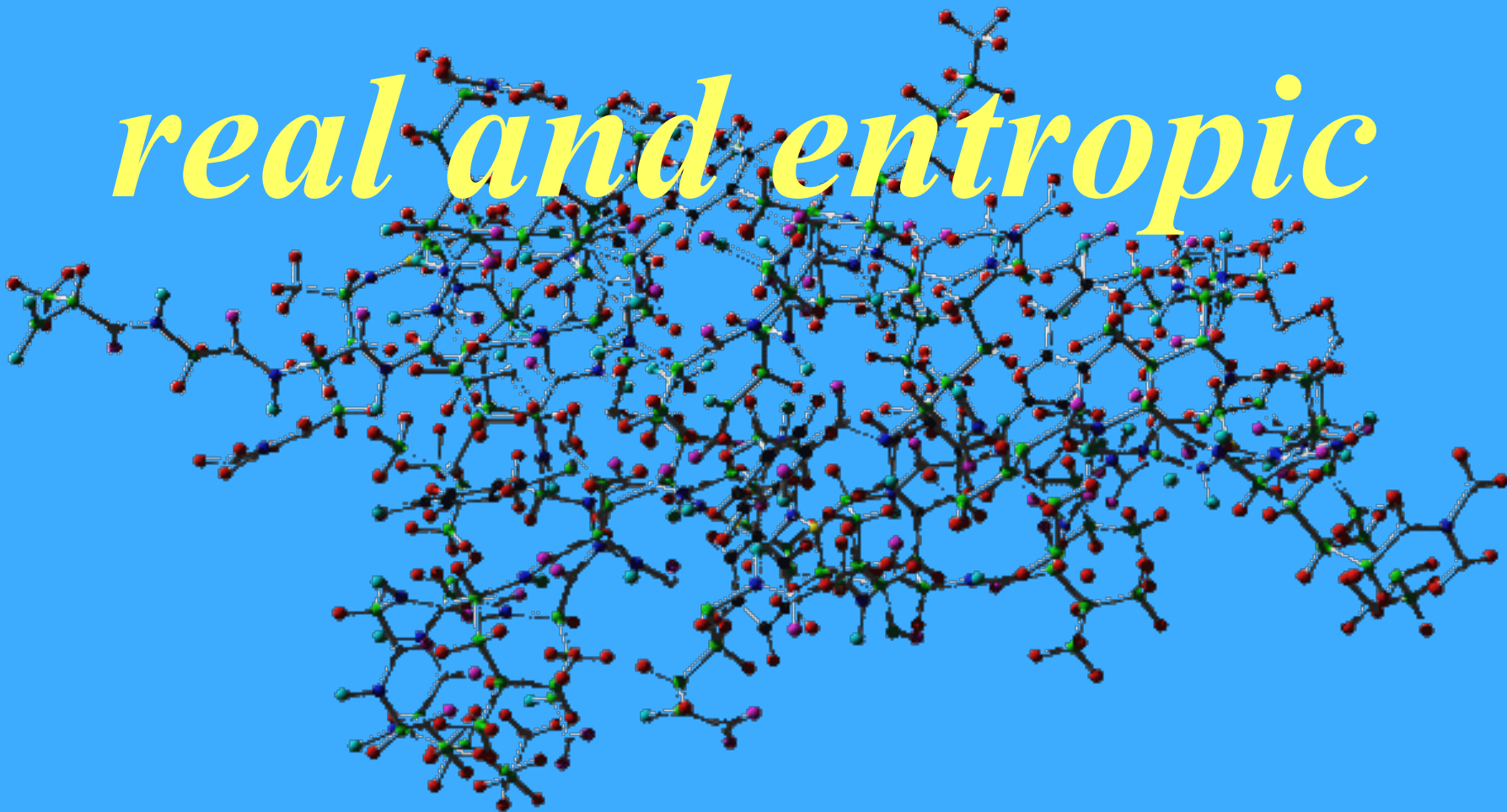


# *Interactions real and entropic*



$$\Delta G = \Delta H - T\Delta S$$

# ***Biological macromolecules are stabilized by physical interactions:***

**Bond energy (interaction energy) is the work required to bring the two atoms (objects) from infinite distance to the length of the bond.**

*Always has a negative value!*

***Strong***

***Weak***

*-200 - -800 kJ/mol (-50 - -200 kcal/mol)*

***Covalent bonds***

*-40 - -400 kJ/mol (-10 - -100 kcal/mol)*

***Ionic interactions***

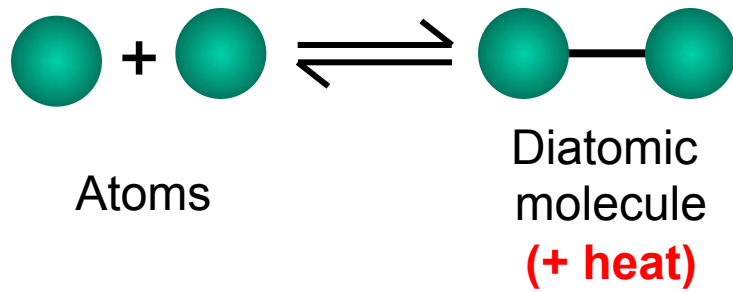
***Ion-dipole interactions***

***Hydrogen bonds***

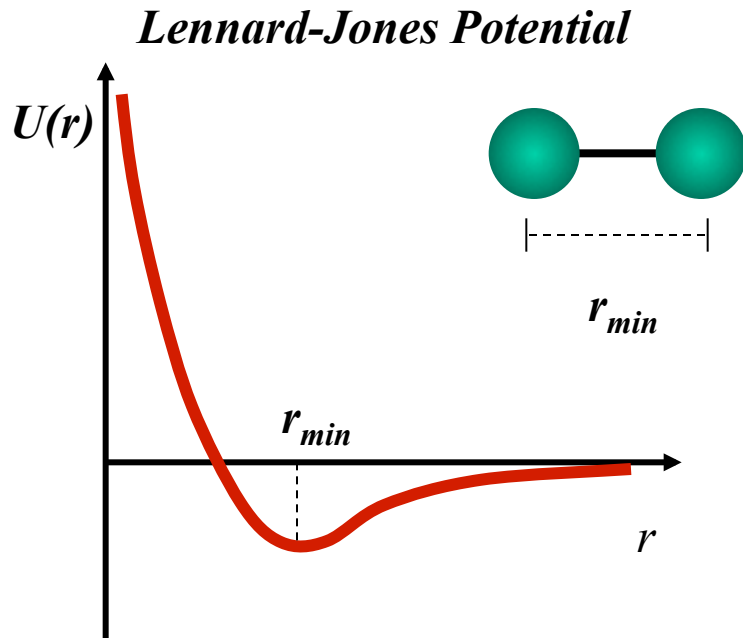
***Van der Waals interactions***

***Hydrophobic effect***

$$\text{C}_{12}\text{H}_{22}\text{O}_{12} + \text{C}_3\text{H}_8\text{O}_3 \rightleftharpoons 5\text{C}_2\text{H}_6\text{O} + 5\text{CO}_2$$



*Bio-polymers are held together by covalent bonds between subunits.*



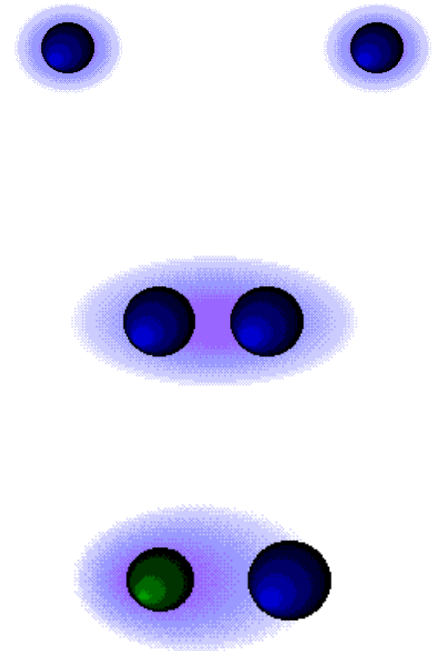
*A covalent bond* is formed between the two non-metals which share a pair of valence electrons so that each obtains a filled valence shell.

*Bond formed due to presence of a stable minimum in the interaction potential energy*

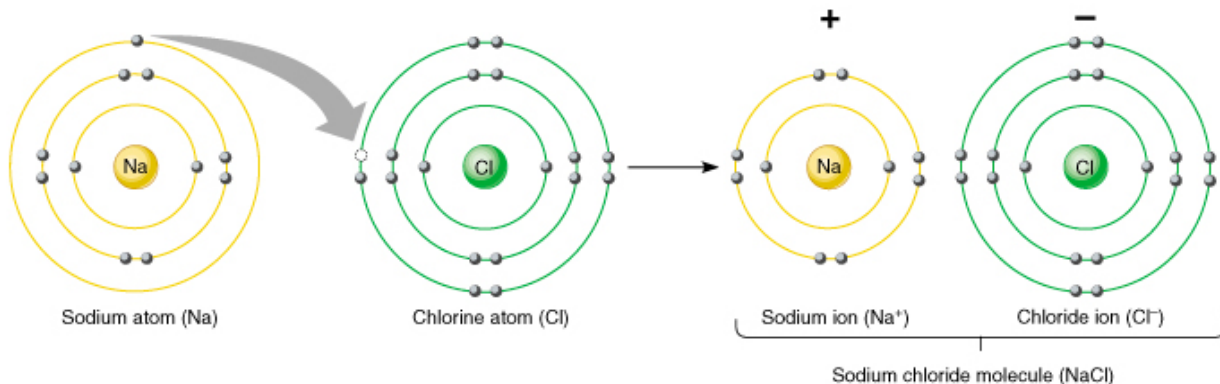
-200 to -800 kJ/mol  
-50 to -200 kcal/mol

*A non-polar covalent bond*

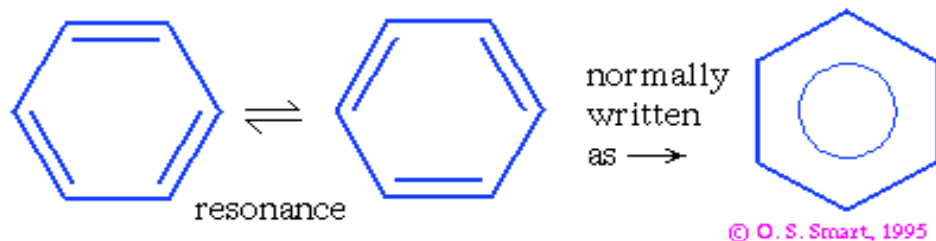
*A polar covalent bond*



# *Ionic bonds*

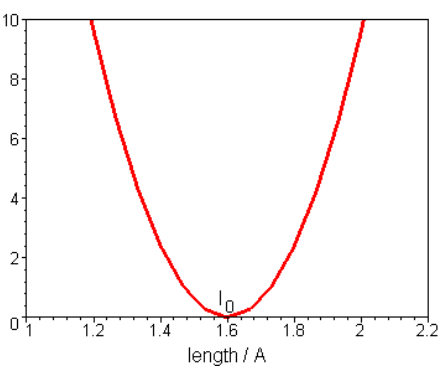


✚ The valence electrons interact and the metal transfers its valence electrons to the non-metal.

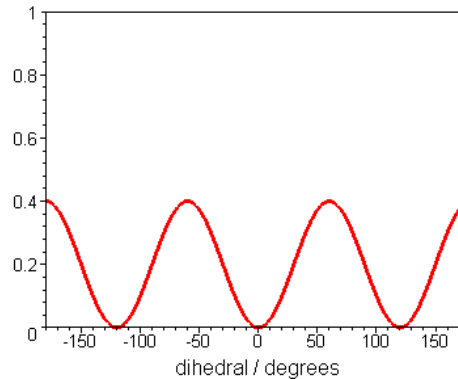
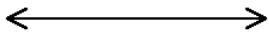


✚ **Delocalized bonding** – a resonance hybrid between alternate structures e.g., benzene

# Bonding potential (covalent bonds)

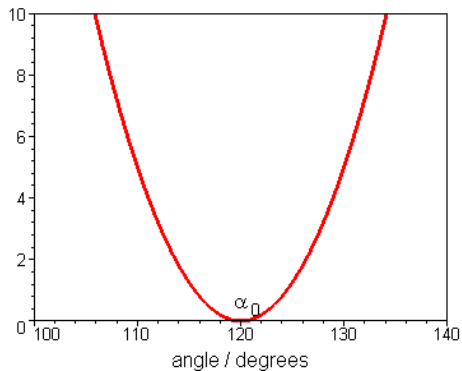
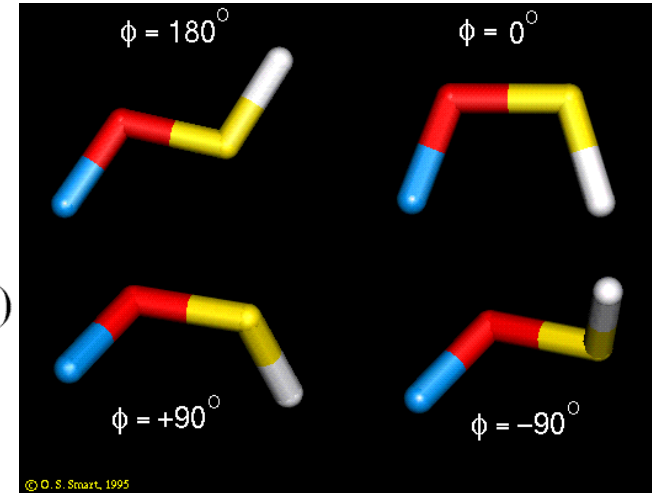
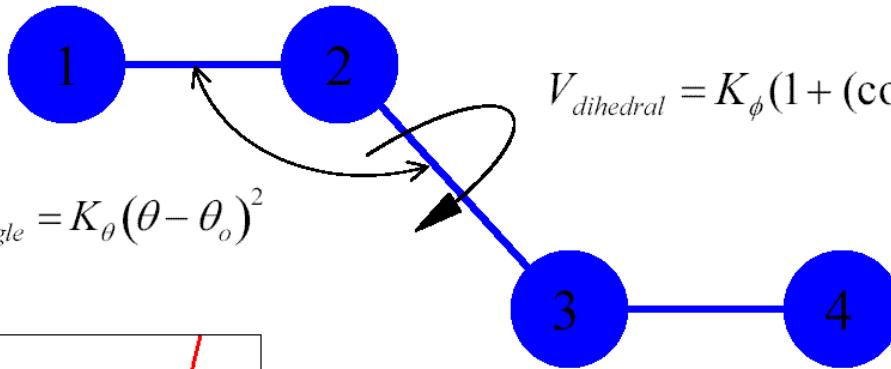


$$V_{bond} = K_b (b - b_o)^2$$



$$V_{dihedral} = K_\phi (1 + (\cos n\phi - \delta))$$

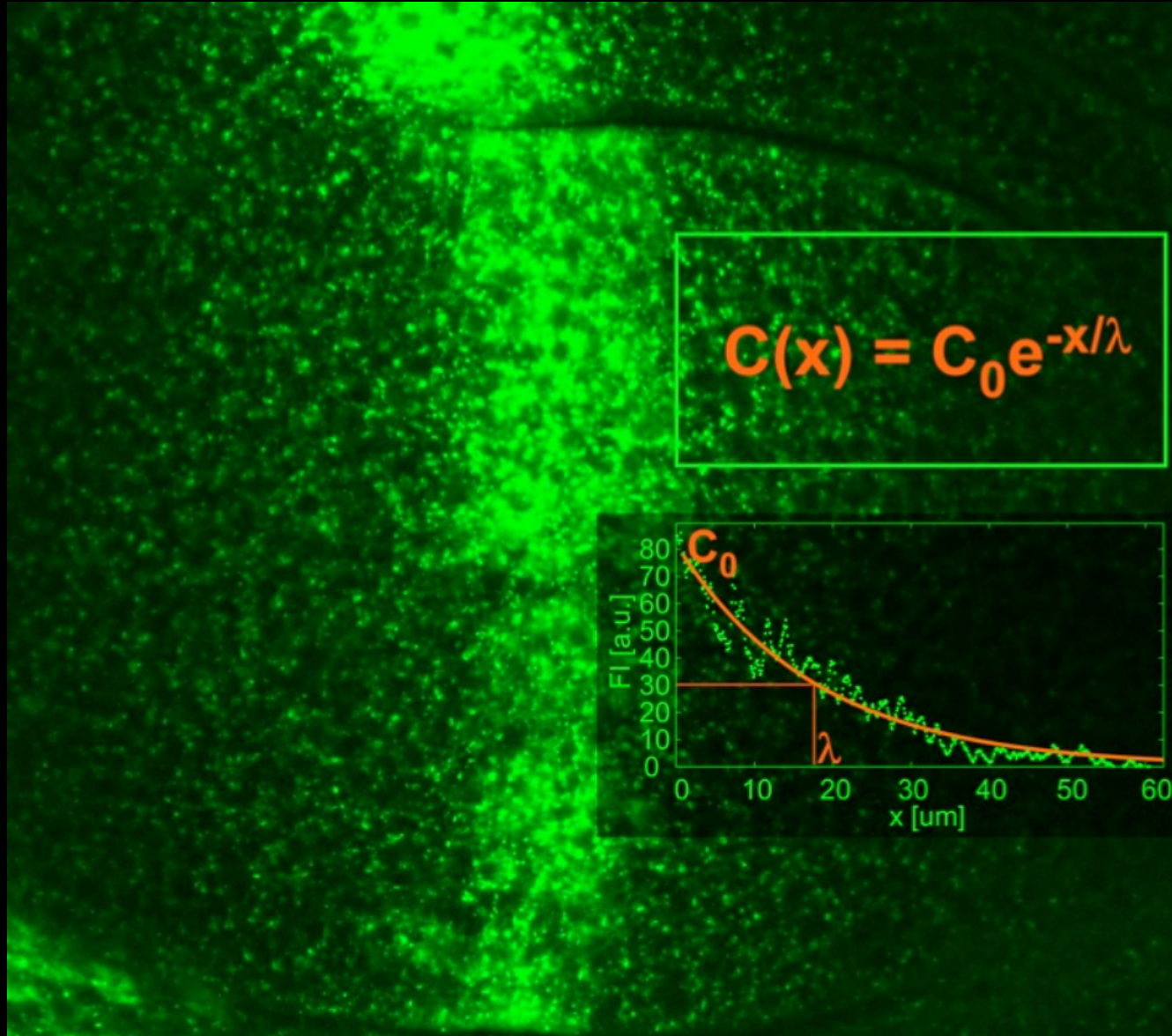
$$V_{angle} = K_\theta (\theta - \theta_o)^2$$



*$V_{bonding}$  is big, but  $\Delta V_{bonding}$  between conformations is not so big  $\rightarrow V_{nonbonding}$  is what counts most in biological system*



# *Biophysics*



# *Intra- and Inter-molecular interactions is what biology is all about.*

***Weak Interactions are additive***

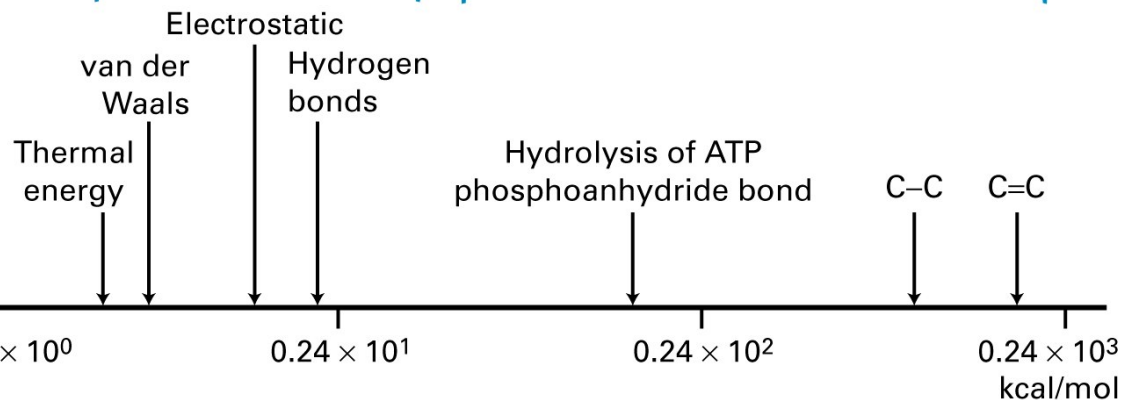
***Weak interactions are dynamic***

*(interactions form, break, re-form constantly)*

Interaction type	Range	Directional	Energy (kJ/mol)	Energy ( $k_B T$ )
Covalent	very short	yes	100 - 900	40 - 360
Hydrogen bond	very short	yes	10 - 40	4 - 16
Charge-charge	$\propto r^{-1}$	no	503	200
Charge-dipole	$\propto r^{-2}$	yes	97	39
Dipole-dipole	$\propto r^{-3}$	yes	19	7.5
van der Waals	$\propto r^{-6}$	no	0.5 - 5	0.2 - 2

## Noncovalent interactions

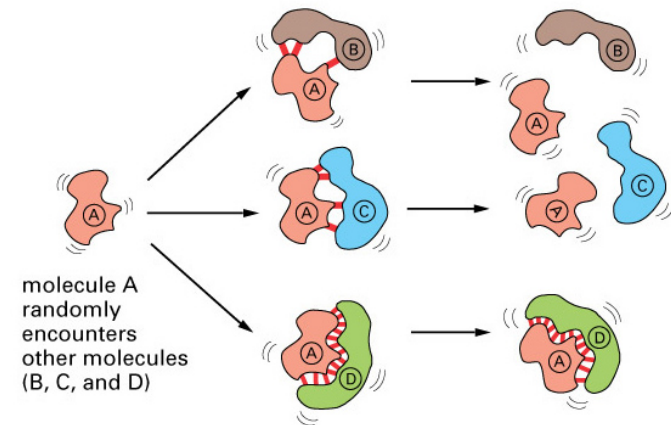
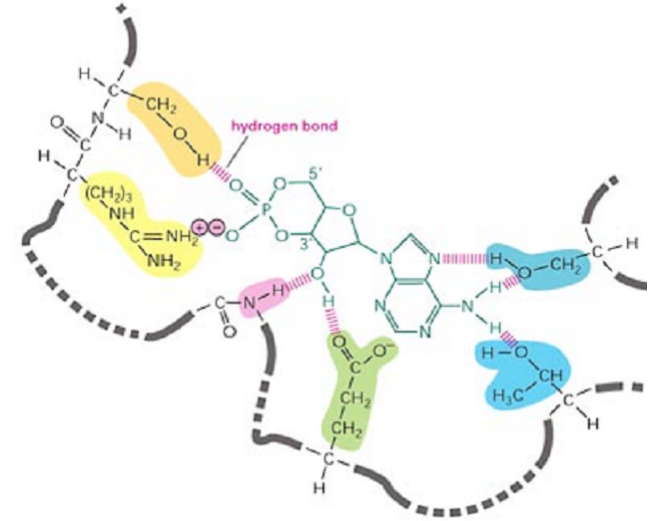
## Covalent bonds





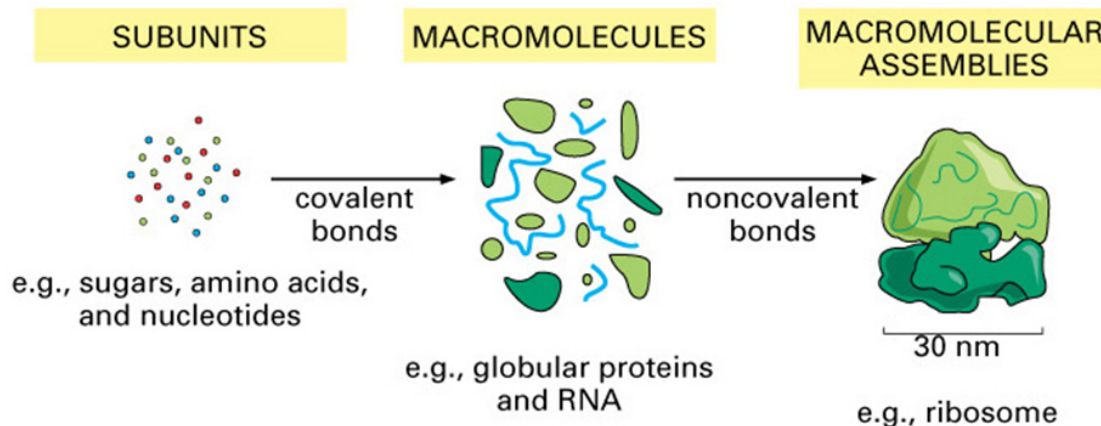
# Weak interactions

- They determine the shape of macromolecules
- They produce reversible self-assembly of presynthesized subunits into specific structures
- They determine the specificity of most molecular interactions
- Molecules or supramolecular aggregates changes upon environmental changes which affect the strengths of weak bonds



the surfaces of molecules A and B, and A and C, are a poor match and are capable of forming only a few weak bonds; thermal motion rapidly breaks them apart

the surfaces of molecules A and D match well and therefore can form enough weak bonds to withstand thermal motion; they therefore stay bound to each other

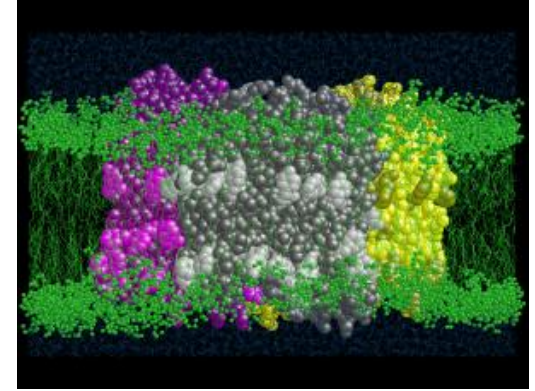


# Energy of macromolecules and aggregates

There are two types of interactions

$$V = \sum_{i < j} \frac{q_i q_j}{4\pi\epsilon_0 r_{i,j}} + \sum_{i < j} \left( \frac{A_{i,j}}{r_{i,j}^{12}} - \frac{B_{i,j}}{r_{i,j}^6} \right) +$$

 noncovalent



$$+ \sum_{bonds} \frac{1}{2} k_{i,j}^b (r_{i,j} - b_{i,j}^0)^2$$

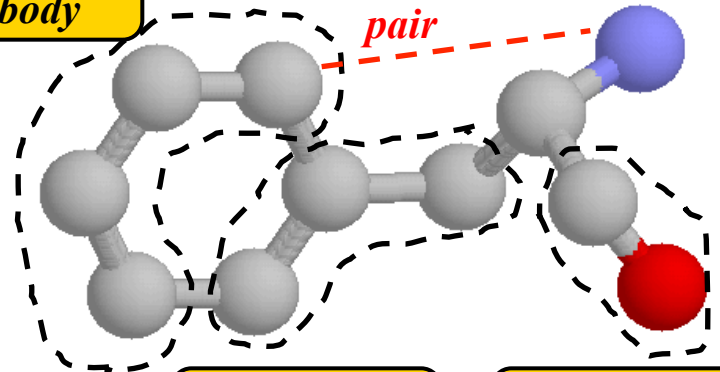
$$+ \sum_{angels} \frac{1}{2} k_{i,j,k}^\theta (\theta_{i,j,k} - \theta_{i,j,k}^0)^2 +$$

$$+ \sum_{dihedrals} k^\phi (1 + \cos(n(\phi - \phi^0)))$$

 covalent

*Torsion angles  
Are 4-body*

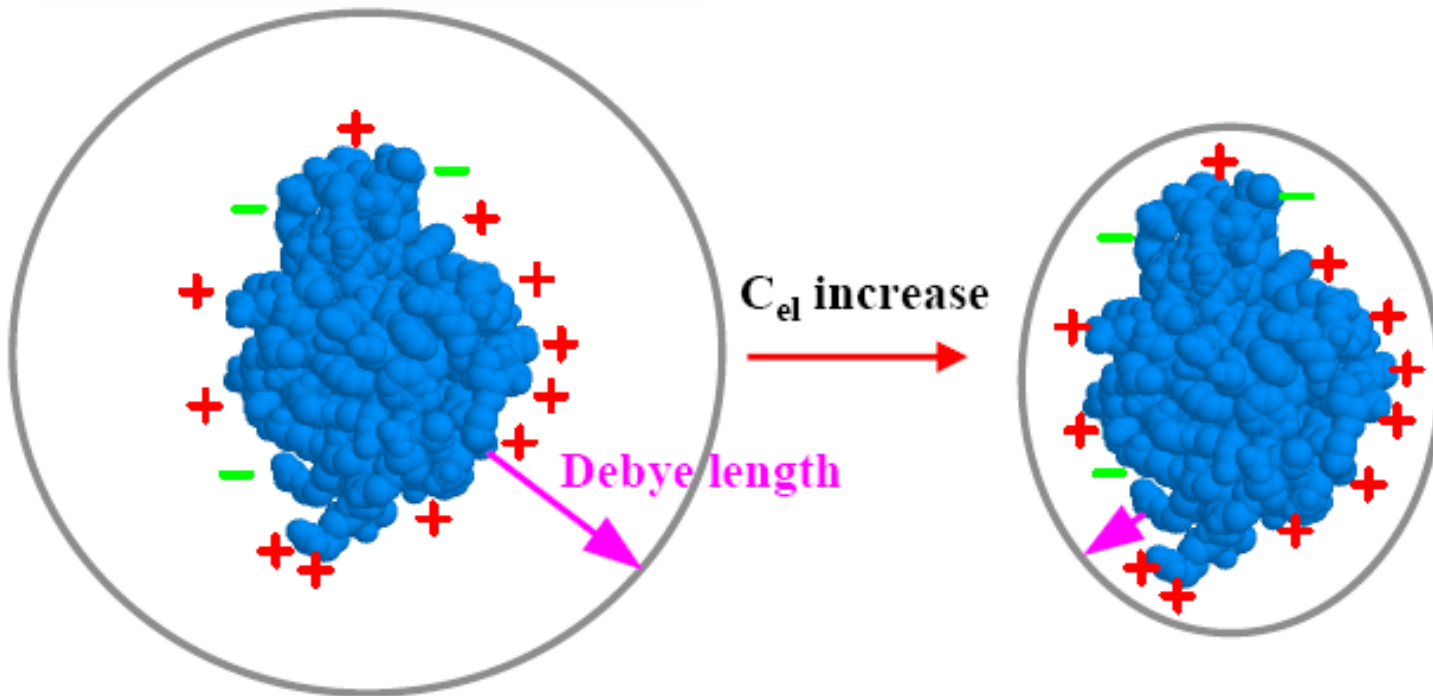
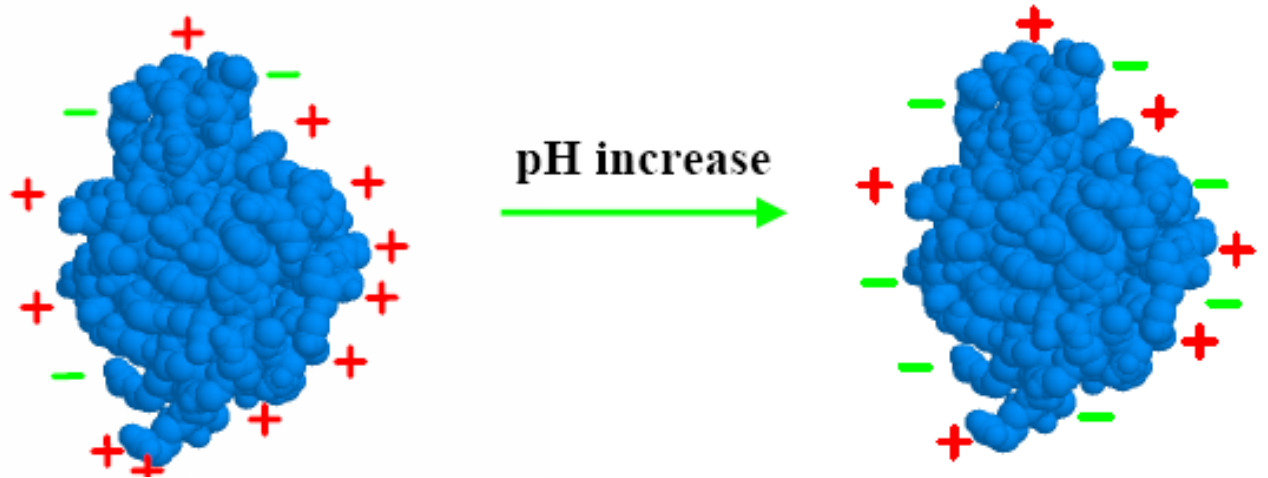
*Non-bonded  
pair*



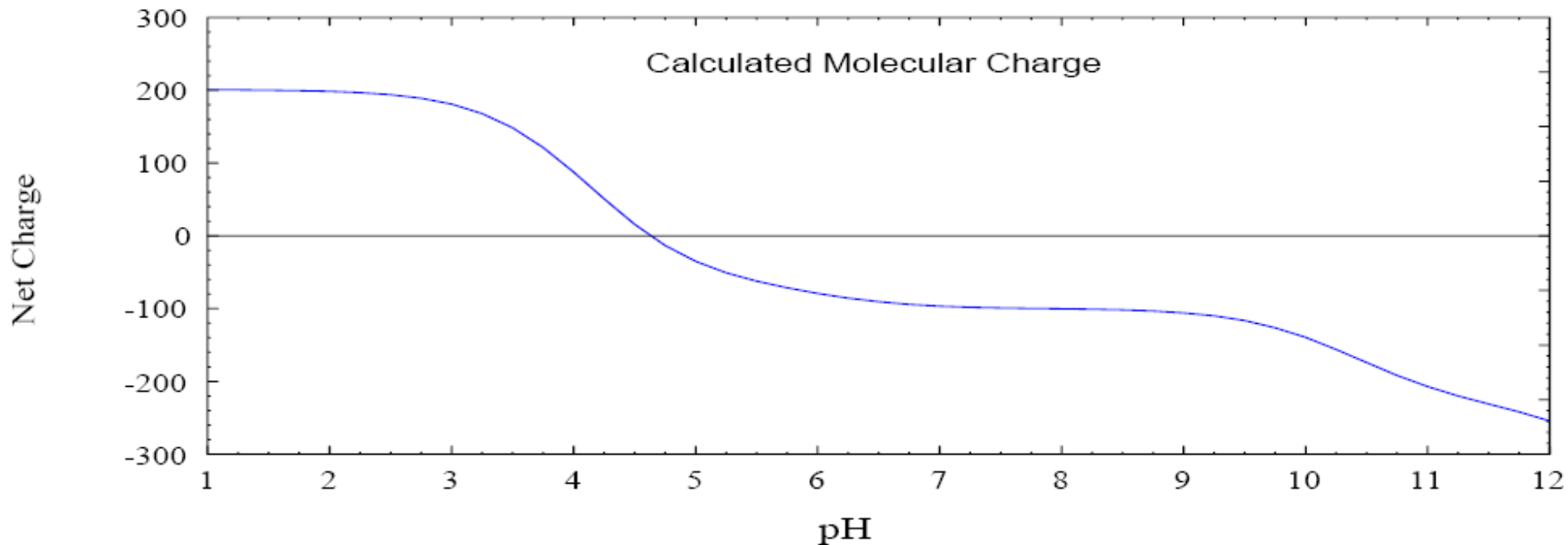
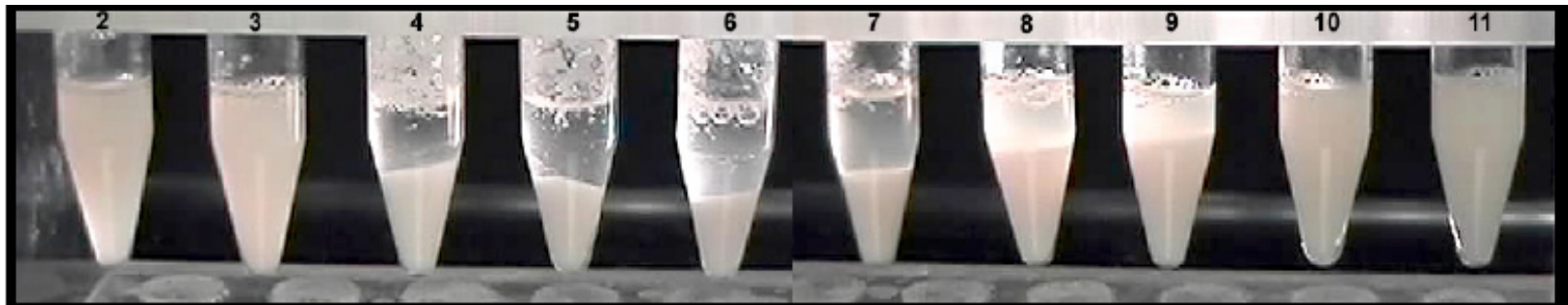
*Angles  
Are 3-body*

*Bonds  
Are 2-body*

# *The two basic parameters affecting macromolecule interactions*



# *Correspondence between charge and precipitation equilibria - soy protein.*

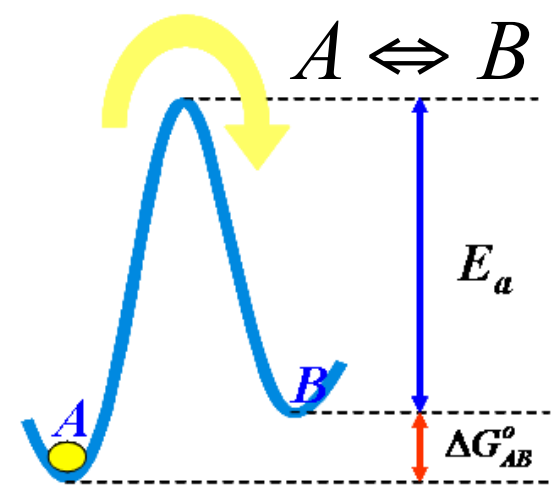


# *Biology works within $kT$*

$1kT = 0.62 \text{ kcal/mole}$  (at  $T=300\text{K}$ )

*At equilibrium*

$$K_{A \leftrightarrow B} = \frac{[B]}{[A]} = e^{-\frac{\Delta G_{A \leftrightarrow B}^0}{k_B T}}$$

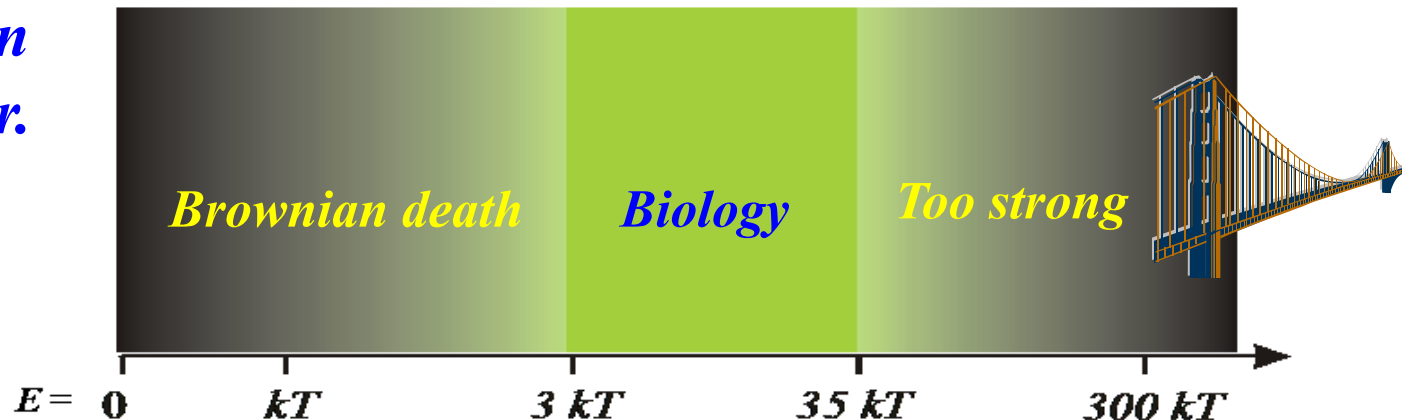


$10^{67}$  years

seconds

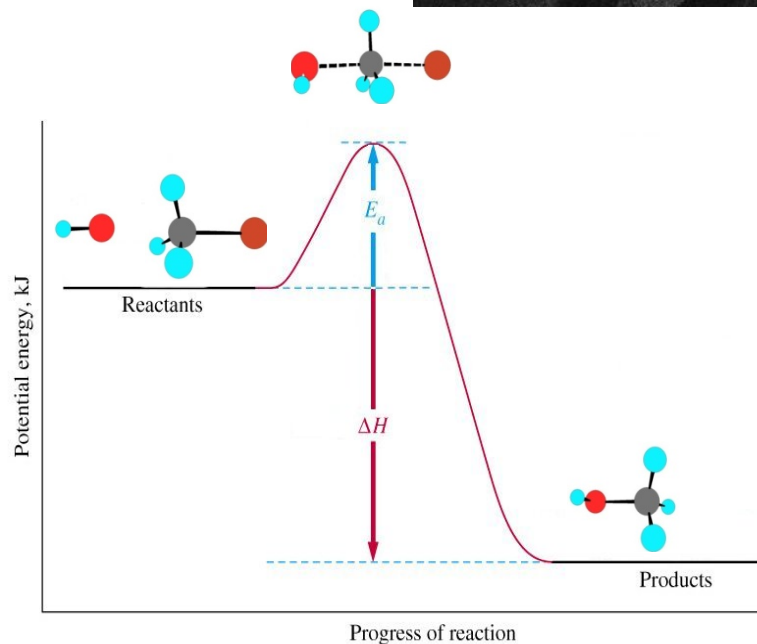
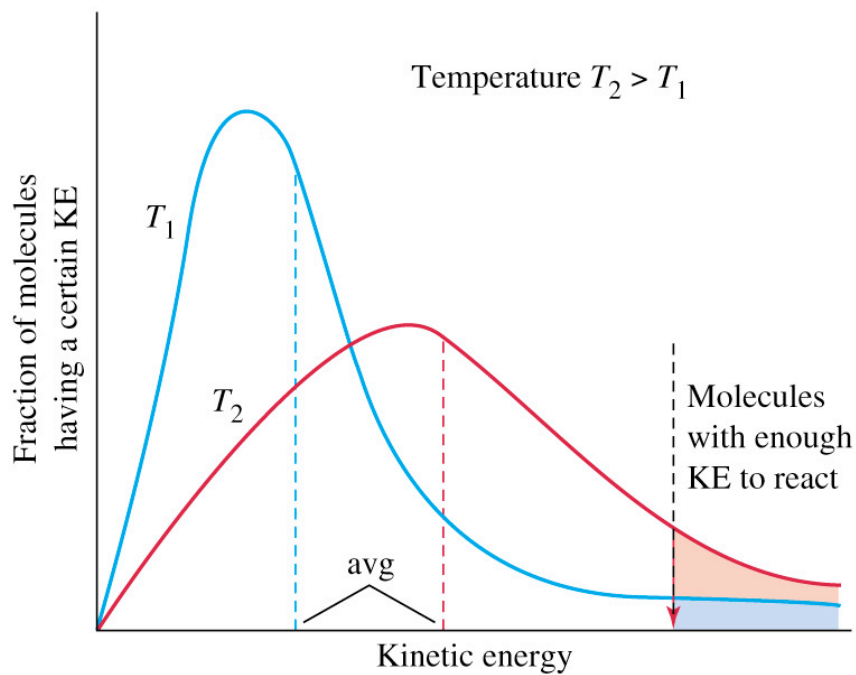
pico-seconds

*The time it will take  
for Brownian motion  
to overcome an  
energy barrier.*



# *Energetics of transition state theory*

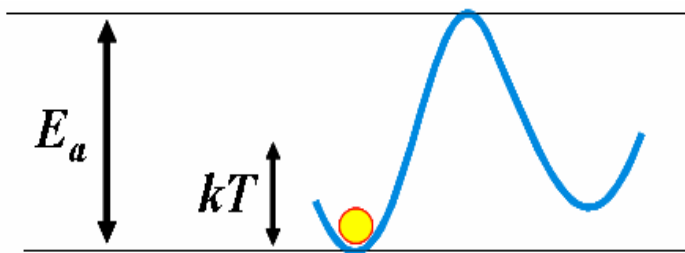
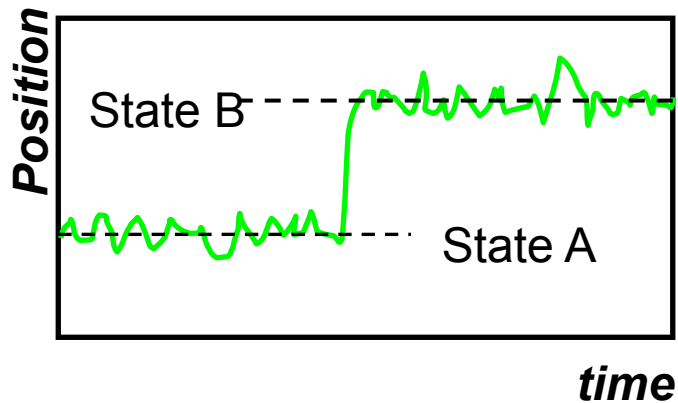
Svante Arrhenius (Nobel Prize, 1903).





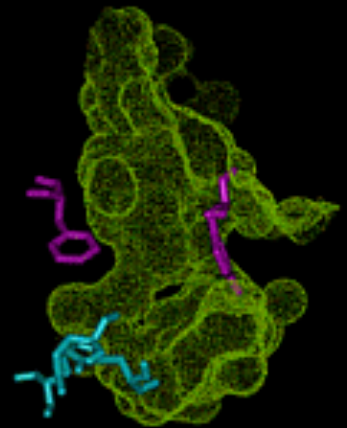
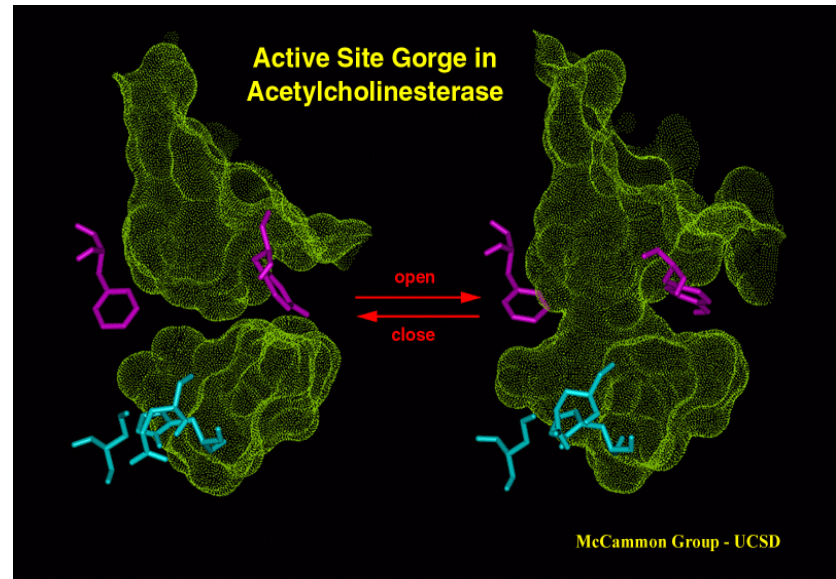
# Thermal motion

- biomolecules are stable enough to make things work,
- allow the systems to play around in order to allow the evolution



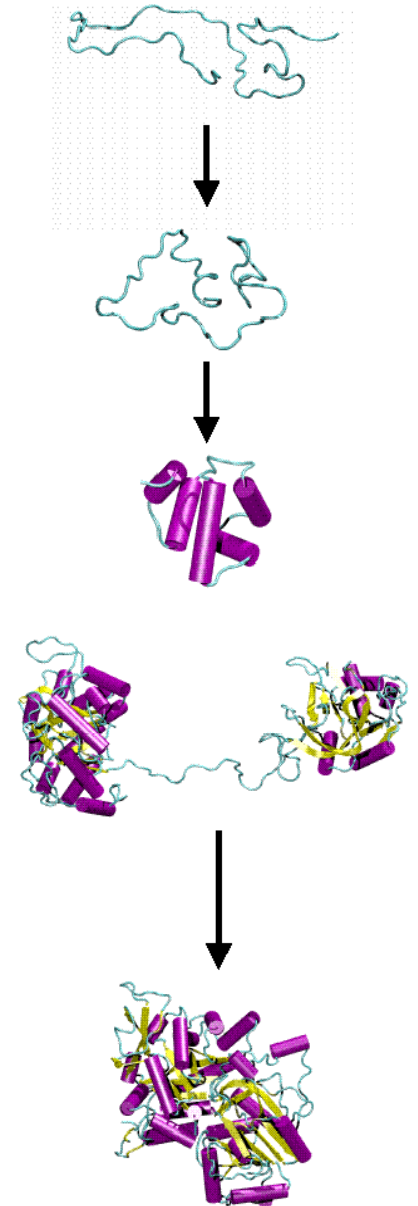
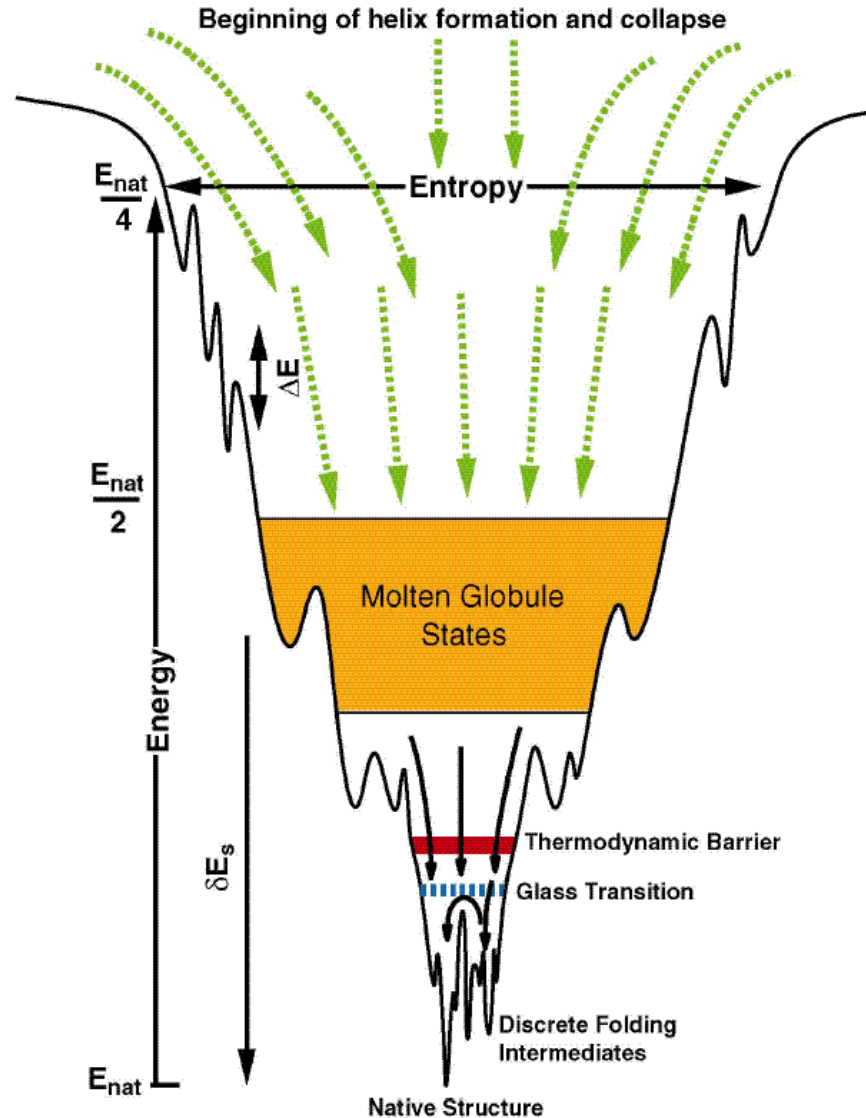
The average waiting time

$$\tau_{A \rightarrow B} = Ce^{\frac{E_a}{kT}}$$



**Conformation** - surface **outline** or **contour** or **3-D orientation** of chemical groups that are free to assume different positions in space without breaking any bonds.

*The folding of a macromolecule*

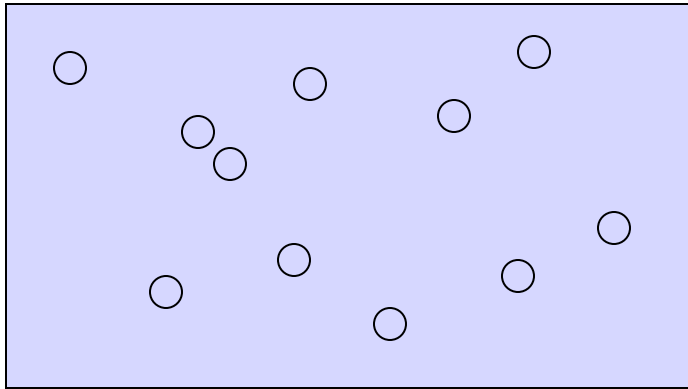


# *Entropic „Forces”*

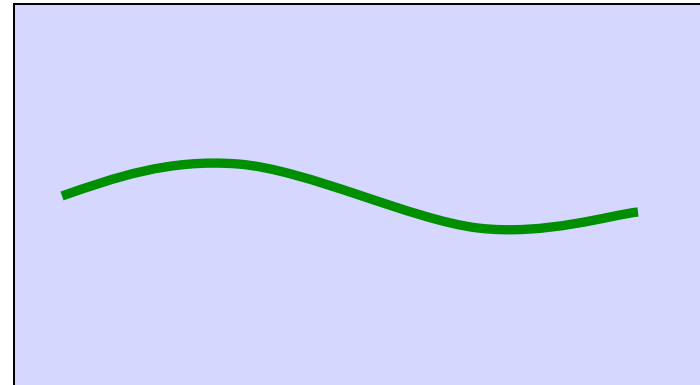
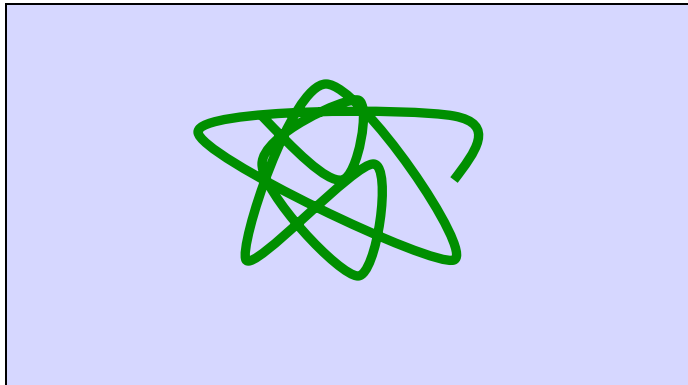
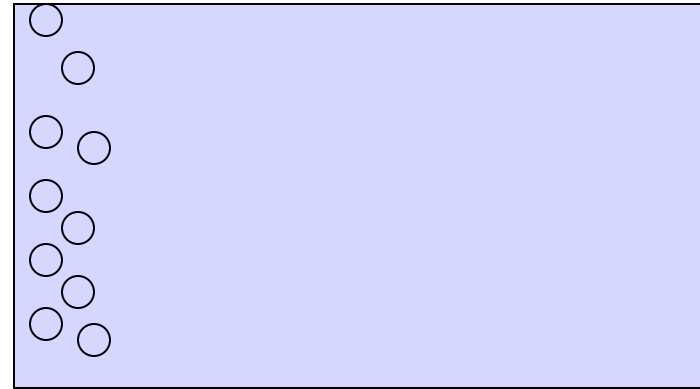
*Disorder is Favorable*



**Probable**



**Improbable**



*The entropic forces can create a situation where two molecules will interact strongly, although there is not a direct “force” between them.*

*To create order ‘work’ must be done*

