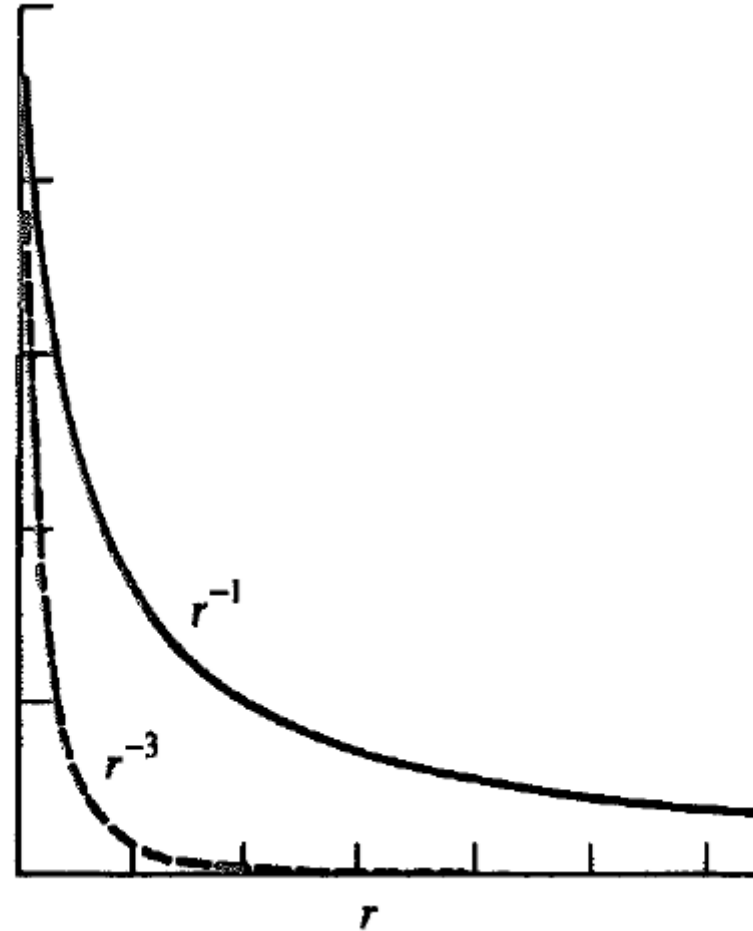


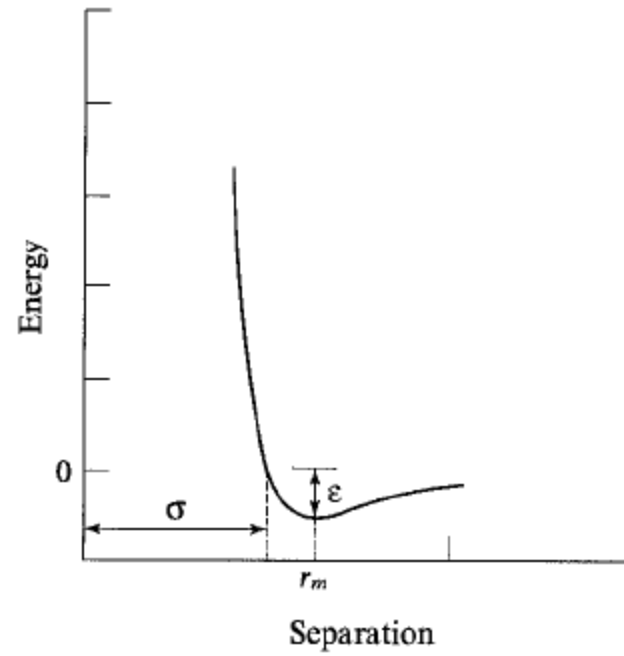
$$\begin{aligned} \mathcal{V}(\mathbf{r}^N) = & \sum_{\text{bonds}} \frac{k_i}{2} (l_i - l_{i,0})^2 + \sum_{\text{angles}} \frac{k_i}{2} (\theta_i - \theta_{i,0})^2 + \sum_{\text{torsions}} \frac{V_n}{2} (1 + \cos(n\omega - \gamma)) \\ & + \sum_{i=1}^N \sum_{j=i+1}^N \left( 4\epsilon_{ij} \left[ \left( \frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left( \frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] + \frac{q_i q_j}{4\pi\epsilon_0 r_{ij}} \right) \end{aligned}$$

## *Electrostatics*

$$\mathcal{V} = \sum_{i=1}^{N_A} \sum_{j=1}^{N_B} \frac{q_i q_j}{4\pi\epsilon_0 r_{ij}}$$

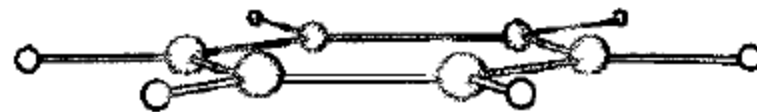
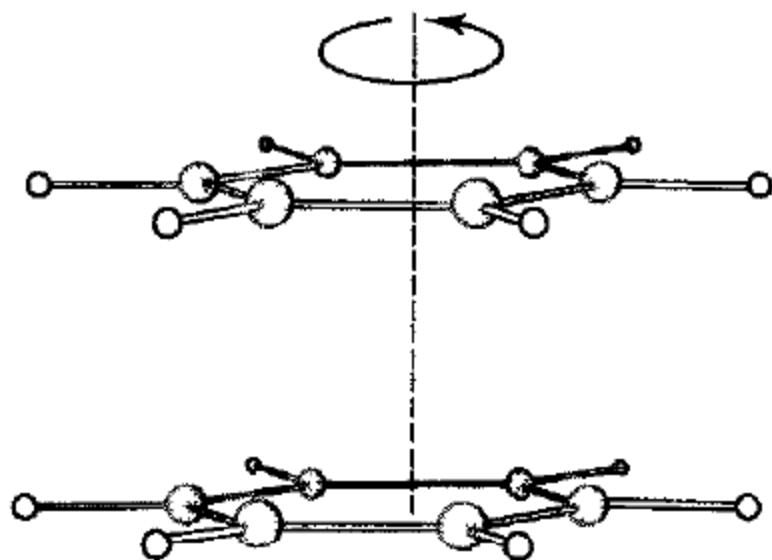


*van der Waals*



$$+ \sum_{i=1}^N \sum_{j=i+1}^N \left( 4\epsilon_{ij} \left[ \left( \frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left( \frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] \right)$$

## *Aromatic Pi-Pi Interactions*

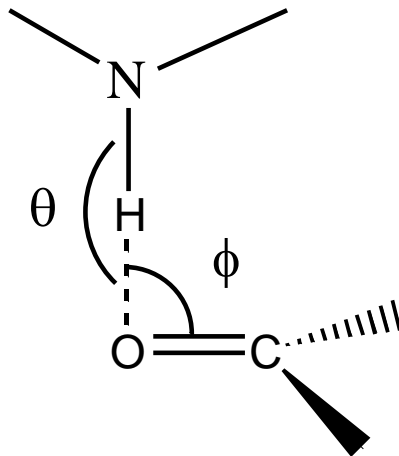


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*Face-to-face (left) and T-shaped (right) orientations of the benzene dimer*

# Hydrogen bonding

$$E_{HB} = D_0 \left[ 5 \left( \frac{R_0}{R} \right)^{12} - 6 \left( \frac{R_0}{R} \right)^{10} \right] F(\theta)$$



$$sp^3 \text{ donor} - sp^3 \text{ acceptor} \quad F = \cos^2 \theta \cos^2 (\phi - 109.5)$$

$$\theta > 90^\circ, \phi - 109.5^\circ < 90^\circ \quad (3)$$

$$sp^3 \text{ donor} - sp^2 \text{ acceptor} \quad F = \cos^2 \theta \cos^2 \phi \quad (4)$$

$$\phi > 90^\circ$$

$$sp^2 \text{ donor} - sp^3 \text{ acceptor} \quad F = \cos^4 \theta \quad (5)$$

$$sp^2 \text{ donor} - sp^2 \text{ acceptor} \quad F = \cos^2 \theta \cos^2 (\max[\phi, \varphi]) \quad (6)$$

---

## Energy functions for protein design

D Benjamin Gordon\*, Shannon A Marshall\* and Stephen L Mayo†

Current Opinion in Structural Biology 1999, 9:509–513

## *Water Models*

### *Simple Point Charge*

	SPC	SPC/E	TIP3P	BF	TIP4P	ST2
$r(\text{OH}), \text{\AA}$	1.0	1.0	0.9572	0.96	0.9572	1.0
HOH, deg	109.47	109.47	104.52	105.7	104.52	109.47
$A \times 10^{-3}, \text{kcal } \text{\AA}^{12}/\text{mol}$	629.4	629.4	582.0	560.4	600.0	238.7
$C, \text{kcal } \text{\AA}^6/\text{mol}$	625.5	625.5	595.0	837.0	610.0	268.9
$q(\text{O})$	-0.82	-0.8472	-0.834	0.0	0.0	0.0
$q(\text{H})$	0.41	0.4238	0.417	0.49	0.52	0.2375
$q(\text{M})$	0.0	0.0	0.0	-0.98	-1.04	-0.2375
$r(\text{OM}), \text{\AA}$	0.0	0.0	0.0	0.15	0.15	0.8

An ensemble is a collection of all possible systems which have different microscopic states but have an identical macroscopic or thermodynamic state.

- Microcanonical ensemble (NVE) : The thermodynamic state characterized by a fixed number of atoms,  $N$ , a fixed volume,  $V$ , and a fixed energy,  $E$ . This corresponds to an isolated system.
- Canonical Ensemble (NVT): This is a collection of all systems whose thermodynamic state is characterized by a fixed number of atoms,  $N$ , a fixed volume,  $V$ , and a fixed temperature,  $T$ .
- Isobaric-Isothermal Ensemble (NPT): This ensemble is characterized by a fixed number of atoms,  $N$ , a fixed pressure,  $P$ , and a fixed temperature,  $T$ .
- Grand canonical Ensemble ( $\mu VT$ ): The thermodynamic state for this ensemble is characterized by a fixed chemical potential,  $\mu$ , a fixed volume,  $V$ , and a fixed temperature,  $T$ .

# Temperature and Pressure Coupling

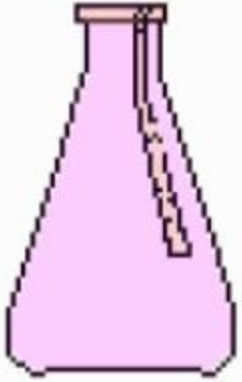
## Berendsen Thermostat and Barostat

$$\frac{dT}{dt} = \frac{T_0 - T}{\tau}$$

$$\left. \frac{dP}{dt} \right|_{\text{bath}} = \frac{P_0 - P}{\tau_P}$$

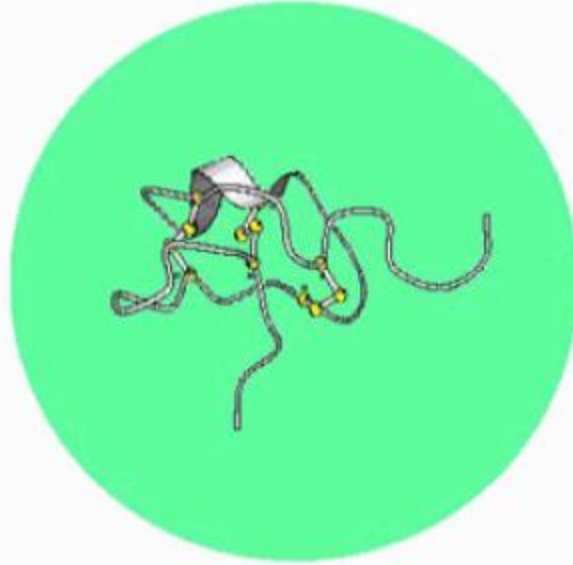


Experiment



Macroscopic

Molecular Simulation



Microscopic

The **Ergodic hypothesis** states

$$\langle A \rangle_{ensemble} = \langle A \rangle_{time}$$

Ensemble average = Time average