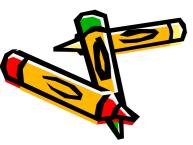


A Brief Introduction

ALE

The range of applicability

- Molecules containing thousands of atoms.
- Organics, oligonucleotides, peptides, and saccharides (metallo-organics and inorganics in some cases).
- Vacuum, implicit, or explicit solvent environments.
- Ground state molecular structure and energy only.
- Thermodynamic and kinetic (via molecular dynamics) properties.



Molecular Modeling

- Nuclei and electrons are lumped into atom-like particles.
- Atom-like particles are spherical (radii obtained from measurements or theory) and have a net charge (obtained from theory).
- Interactions are based on springs and classical potentials "Newtonian" physics laws.
- Interactions must be preassigned to specific sets of atoms.
- Interactions determine the **spatial distribution** of atom-like particles and their **energies**.





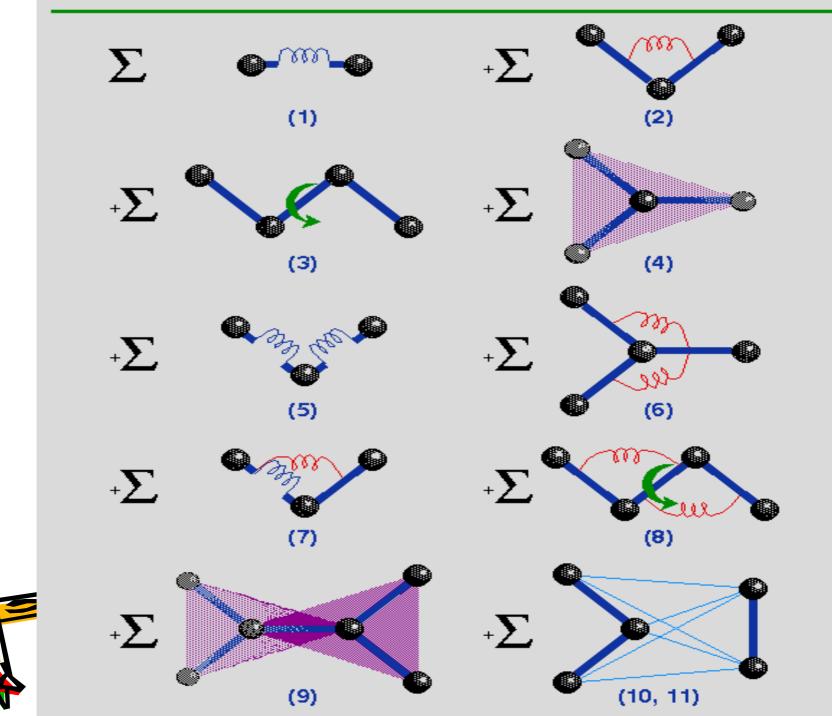
"Steric energy" based on energy increments due to deviation from some "ideal" geometry

$$E_{steric} = E_{stretch} + E_{bend} + E_{torsion}$$

$$+ E_{stretch-bend} + (E_{other-cross-term})$$

$$+ E_{H-bonding} + E_{vdW}$$

$$+ E_{electrostatic} + E_{dipole-dipole}$$



Description of the Classical (Newtonian) Forcefield:

$$E_{\text{pot}} = \sum_{b} D_{b} \left[1 - e^{-\alpha (b - b_{0})} \right] + \sum_{\theta} H_{\theta} (\theta - \theta_{0})^{2} + \sum_{\phi} H_{\phi} \left[1 + s \cos (n\phi) \right]$$
(1)
(2)
(3)

$$+\sum_{\chi} H_{\chi\chi}^{2} + \sum_{b} \sum_{b'} F_{bb'}(b - b_{0}) (b' - b'_{0}) + \sum_{\theta} \sum_{\theta'} F_{\theta\theta'}(\theta - \theta_{0}) (\theta' - \theta'_{0})$$
(4)
(5)
(6)

+
$$\sum_{b} \sum_{\theta} F_{b\theta}(b-b_{0}) (\theta-\theta_{0}) + \sum_{\phi} F_{\phi\theta\theta'} \cos\phi(\theta-\theta_{0}) (\theta'-\theta'_{0}) + \sum_{\chi} \sum_{\chi'} F_{\chi\chi'} \chi\chi'$$
(7)
(8)
(9)

+ $\sum \varepsilon \left[(r^{*}/r)^{12} - 2(r^{*}/r)^{6} \right]$ + $\sum q_{i}q_{i}/\varepsilon r_{ij}$ (10) (11)

Use of Cut-offs

- Van der Waals forces, hydrogen bonding, electrostatic forces, and dipole-dipole forces have dramatic distance dependencies; beyond a certain distance, the force is negligible, yet it still "costs" the computer to calculate it.
- To economize, "cut-offs" are often employed for these forces, typically somewhere between 10 and 15Å.



Forcefield Parameters

- A parameter should be adjusted so that the simulated system reproduces properties of the real system. It does NOT have to equal a microscopic descriptor.
- As yet, there is no universal forcefield. Parameters for particular forcefields can only be used to study particular molecules.
- Forcefield parameters must be cohesive--modelers can't just "plug & play" new parameters without testing
- Forcefield parameters must be referenced



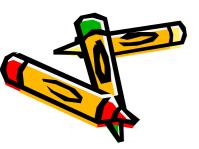
Types of Forcefields

- Diagonal force field: no cross terms (AMBER, CHARMM)
- Matrix force field: with cross terms (MM4, CFF)
- Class II force field: parameters derived from quantum mechanics (ESFF)

• "Fast" force fields: only torsion & hon-bond terms (SCULPT, YETI)

Hyperchem Force Fields

- MM2 / MM3 (Allinger) <u>best</u>; general purpose
- MMX (Gilbert) added TS's, other elements; good
- MM+ (Ostlund) general; good
- OPLS (Jorgenson) proteins and nucleic acids
- AMBER (Kollman) proteins and nucleic acids +
- **BIO+** (Karplus) CHARMm; nucleic acids



Properties Calculated

- Optimized geometry (minimum energy conformation)
- Equilibrium bond lengths, bond angles, and dihedral (torsional) angles
- Dipole moment (vector sum of bond dipoles)
- Enthalpy of Formation.

Steps in Performing MM Calculations

- Construct graphical representation of molecule to be modeled ("front end")
- Select forcefield method and termination condition (gradient, # cycles, or time)
- Perform geometry optimization
- Examine output geometry... is it reasonable?

Search for <u>global</u> minimum.

Energy Minimization

- Local minimum vs <u>global</u> minimum
- Many local minima; only ONE global minimum
- Methods: Newton-Raphson (block diagonal), steepest descent, conjugate gradient, others.

