Computational Methods Summary

Types of Energy Models (QM or MM)

1. Molecular mechanics (MM)

- Useful for initial calculations on small organics, can do any size up through proteins.
- Uses classical ball-and-spring laws where each force constant, ideal bond length, angle and dihedral angle are empirical parameters. Through-space electrostatic and VDW interactions are also added.
- *Empirical parameters* mean that method is only useful if you have parameters (a **force field**) closely matching every type of linkage in your system.
- Programs available here: Macromodel, AMBER, NAMD, Desmond, Gromacs, more.....

2. Hartree-Fock (QM)

- Mainly used as a starting point for high-level *ab initio* calculations or mixed with DFT (see 3).
- Approximate solution of the Schrödinger equation neglecting instantaneous electron-electron interactions (correlation) unless corrections are applied.
- Energy and orbital coefficients solved for by an iterative self consistent field (SCF) method.
- High-level calculations (configuration interaction, coupled cluster, G3, etc.) can correct for correlation and achieve chemical accuracy but only possible for small numbers of atoms, very time consuming.
- Programs available here: Gaussian, Jaguar, QChem, Gamess, NWChem, more

3. Density functional theory (QM)

- Appropriate for systems up to ca. 100 atoms (sometimes larger), best choice for most organometallics
- Based on an equation analogous to the Schrödinger equation where *energy is a functional of electron density*, not the true wavefunction. Iterative SCF procedure is similar to that of HF.
- Implicitly includes electron correlation, so much **more accurate than HF**, but behavior is sometimes **unpredictable** because the electronic density functional cannot be derived *ab initio*.
- Myriad functionals designed (parameterized) based on certain principles. The most *popular* one is B3-LYP which includes some HF exchange (hybrid DFT). B3LYP has very good performance for organic molecules, but may not suitable for metal complexes and structures with non-covalent interactions. Some recently developed functionals are ωB97X-D, M06, HSE06, BMK, TPSSh, etc...
- Programs available here: Gaussian, Jaguar, QChem, Gamess, NWChem, more.....

Basis Sets (for QM calculations only)

- The set of atomic orbitals whose coefficients are optimized in SCF.
- Each orbital usually represented by a linear combination of Gaussian functions.
- The number of Slate-type orbital (STO) determines the size of a basis set. The bigger the basis set, the better the results, but for DFT not much gain beyond triple-zeta level (three STO).
- **6-31G** is a *popular* double-zeta level split-valance basis set. It uses 6 Gaussian functions to describe inner shell orbitals, while 3 and 1 Gaussian functions for the 1st and the 2nd valance STO, respectively.
- **Polarization functions** (** or (d,p) after G) add d functions to p shells (carbon, phosphorus, etc.) and p functions to hydrogen. *Must use at least d functions on p shells (one *) for acceptable results!*
- **Diffuse functions** (++ before G) add next-principal-quantum-number shell. In DFT, normally only useful for anions, excited states or sometimes late transition metals.
- Effective core potentials (ECP) replace core electrons with a potential felt by valence electrons, simplifies calculation of heavy atoms. The relativistic effect of core electrons can be included in the ECP. Normally needed for atoms heavier than 1st-row transition metals.
- If you have **metals/heavy atoms**, start with **LANL2DZ** but then see our page for suggestions about better basis sets: <u>http://glab.cchem.berkeley.edu/glab/faqs/gauss_custombasis.html</u>

QM Output:

- Can include orbital energies/pictures, excitation energies, partial charges, dipole and multipole moments, polarizabilities, NMR shielding and coupling constants, etc.

Common Types of Geometry Algorithms

- 1. Molecular energy (calculates energy of input geometry no geometry change)
- Can be used on a previously optimized geometry
 - **MM**: Energy is relative to a hypothetical unstrained system.
 - QM: SCF energy, relative to a separated collection of electrons and nuclei.
- Add up energies of species on each side of a **balanced** chemical equation and calculate reaction energy (*Make sure the total number of basis functions are the same in the whole reaction*).

2. Geometry optimization (also called minimization)

- Finds local minimum: the bottom of whichever potential energy well in which a molecule is drawn.
- Iterative process, with the molecular energy calculated between *each* optimization cycle.
- Can optimize all coordinates or constrain one or more coordinates.
- Default convergence criteria are arbitrarily set, you might allow looser or need tighter criteria.

3. Coordinate Scan

- Can be used to manually adjust a distance, angle or torsion systematically. Commonly used for searching torsions (dihedral drive) or as a manual method to locate transition states (TS).

4. Conformational searching

- Best methods randomly sample potential energy surface to find global minimum-energy structure.
- Nearly always confined to MM methods due to number of structures generated/minimized.
- For flexible organic species, perform conformational search before QM geometry optimization.

5. Transition state searching

- Needed to model chemical **kinetics**, locates an energy *maximum* along a reaction coordinate while *minimizing* energy w.r.t. all other coordinates for a reaction mechanism study.
- Normally used with **QM** methods, locating correct TS in the reaction pathway is much more difficult to be successfully achieved than optimization for intermediates.
- A converged TS must have exactly one imaginary vibrational frequency.
- Perform intrinsic reaction coordinate (IRC) calculation for a TS and confirm reactant and product.

6. Vibrational frequencies

- Normally used with **QM** methods, calculates the molecular vibrational frequencies and sometimes IR and/or Raman intensities.
- Also calculates **zero-point energies** (ZPE), thermal enthalpies and thermal Gibbs free energies which should be added to SCF energies when studying reaction kinetics or thermodynamics.
- A true minimum has all real-valued frequencies, a transition state has exactly one imaginary frequency.

7. Molecular dynamics

- Applicable to MM (up to macromolecules) and sometimes QM methods (very small systems).
- Models molecular motion over time, response to forces acting on molecule.
- Will eventually sample *all* accessible conformations at a given temperature, if the simulation is run long enough (this can be very long and in some cases impractical).
- For large systems (proteins, DNA, etc.) more useful than conformational searching.
- Takes into account that reaction trajectories do not always follow the exact minimum-energy path.