

Computational Chemistry

An introduction to the
computational chemistry
package, Gaussian 03

Matt McKenzie
LSU-HPC / LONI

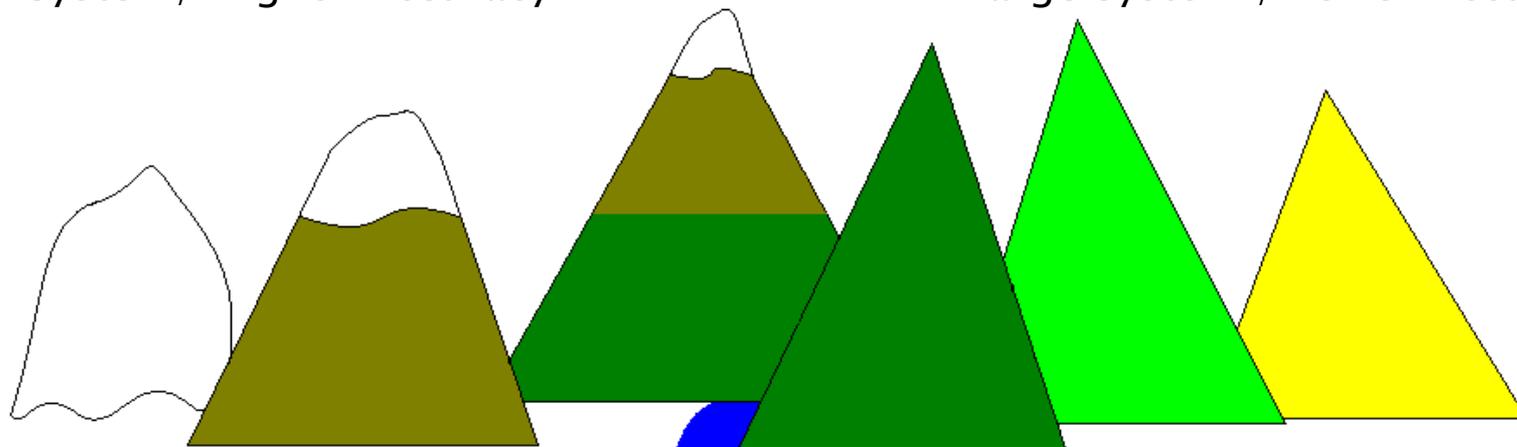
Topics

- ▶ Basic Theory
 - Gaussian specific
 - ▶ Gaussian job creation and submission
 - ▶ Analysis of output
- 

Computational Chemistry Landscape

Small System / Higher Accuracy

Large System / Lower Accuracy



Quantum Mechanics

MO theory, *ab initio*, DFT

Thermodynamics

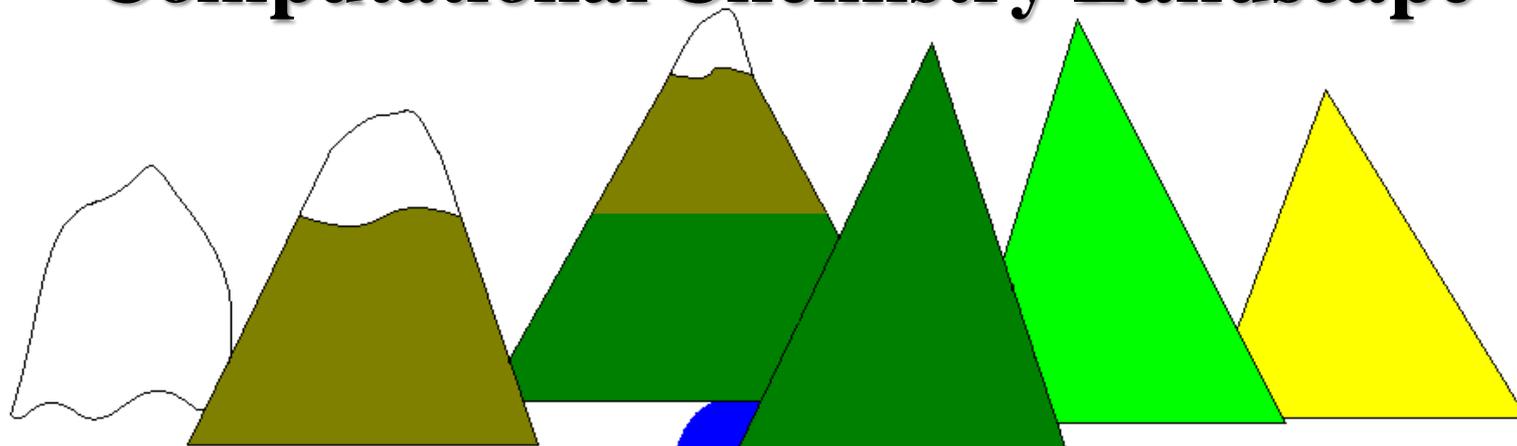
Molecular Mechanics
MD, MC, Virtual Screening

Basis sets

Force fields

Statistical Mechanics

Computational Chemistry Landscape

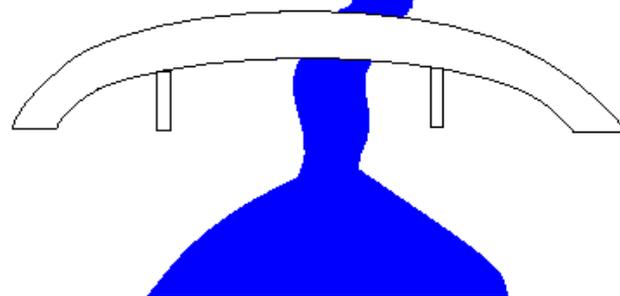


Quantum Mechanics

Thermodynamics

Gaussian 03, Gamess
Qumax

NAMD, LAMMPS,
Gromacs*, Amber*,
AutoDock



PINY, NWChem

Statistical Mechanics

CPMD, QM/MM

Theory Overview

- ▶ Pre-req. = knowledge of quantum mechanics
 - ▶ Goal: Obtain a fundamental knowledge of *ab initio* calculations using Gaussian
 - Cannot cover the entire computational *ab initio* topic
 - ▶ How wave functions are constructed
 - ▶ Levels of theory used
 - ▶ Focus on small molecules, gas phase
- 

Theory

- ▶ Solutions to the Schrödinger equation
- ▶ $(\hat{H}-E)\Psi(r_1, r_2, \dots, r_N)=0$

- ▶ Ways to solve:
- ▶ Numerical: Approx. partial derivatives of Ψ as finite differences of values of Ψ

- ▶ Expansion Method: model Ψ ,
- ▶ $\Psi(r_1, r_2, \dots, r_N) \approx \sum_i c_i \eta_i(r_1, r_2, \dots, r_N)$
- ▶ The set of η_i forms a basis representation of Ψ

Theory – Basis Sets

- ▶ Expansion Method
- ▶ $(\hat{H}-E)\Psi(r_1, r_2, \dots, r_N)=0$
- ▶ $\Psi(r_1, r_2, \dots, r_N) \approx \sum_i c_i \eta_i(r_1, r_2, \dots, r_N)$
- ▶ The set of η_i forms a basis representation of Ψ

From dictionary.com,

- ▶ Basis
- ▶ 1. the bottom or base of anything; the part on which something *stands or rests*.
- ▶ 2. *anything upon which something is based*; fundamental principle; groundwork.
- ▶ 3. the principal constituent; *fundamental ingredient*.
- ▶ 4. a *basic fact, amount, standard, etc., used in making computations*, reaching conclusions, or the like: The nurse is paid on an hourly basis. He was chosen on the basis of his college grades.
- ▶ 5. Mathematics. *a set of linearly independent elements of a given vector space having the property that every element of the space can be written as a linear combination of the elements of the set.*
 - $\Psi(r_1, r_2, \dots, r_N) \approx \sum_i c_i \eta_i(r_1, r_2, \dots, r_N)$
- ▶ Latin Origin: step, place one stands on, pedestal

Theory – Basis sets

- ▶ $\Psi(r_1, r_2, \dots, r_N) \approx \sum_i c_i \eta_i(r_1, r_2, \dots, r_N)$
- ▶ Instead of the whole system, let's focus on a molecular orbital

$$\varphi_i = \sum_{n=1}^N c_{ni} \eta_n$$

Molecular orbital coefficient
Basis function

- ▶ **Increase η = higher quality** of the wave function
- ▶ **Increase η = higher computational cost**

Basis Functions– STO's & GTO's

Slater-Type Orbitals

$$\eta^{STO}(\mathbf{r}) = x^l y^m z^n e^{-\zeta r}$$

Similar to angular momentum

$$L = l + m + n$$

Gaussian Type Orbital

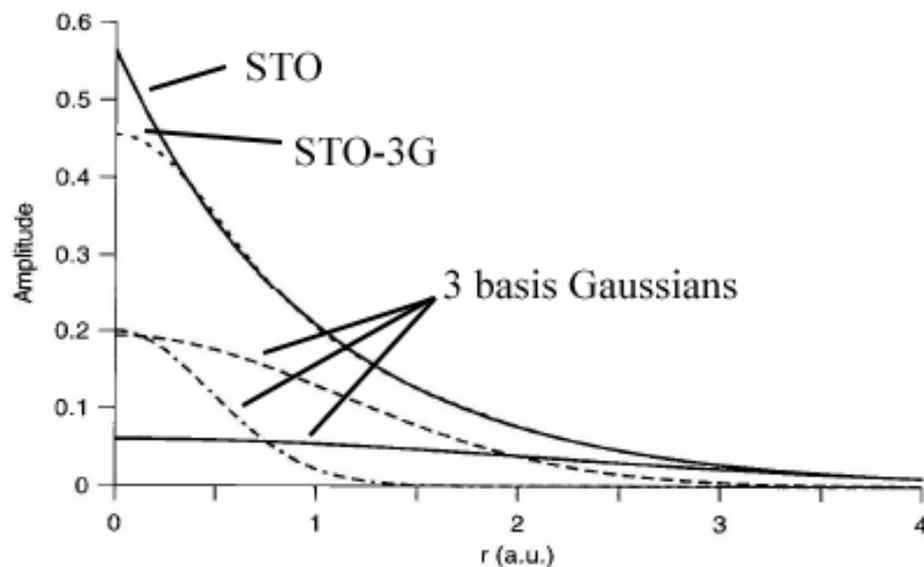
$$\eta^{GTO}(\mathbf{r}) = x^l y^m z^n e^{-\alpha r^2}$$

ζ & α are the **orbital exponents**.

The radial 'size'

A **large** α means the electron is held **tight** to the nucleus.

A **small** α means the electron can **roam far** from the nucleus.

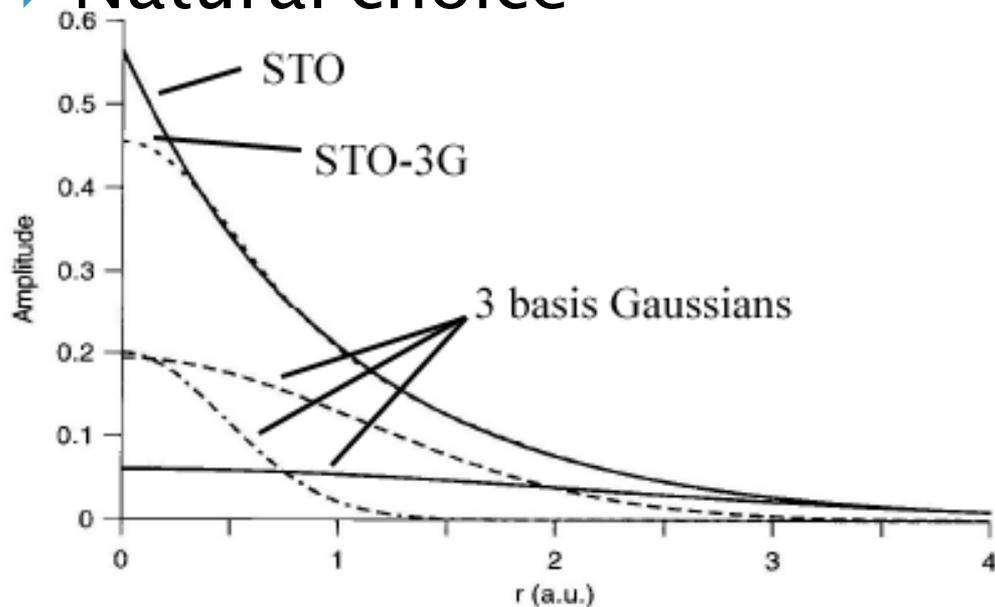


STO vs. GTO

STO

- ▶ Correct cusp at $r \rightarrow 0$
- ▶ Desired decay $r \rightarrow \infty$

- ▶ Correctly mimics H orbitals
- ▶ Natural choice



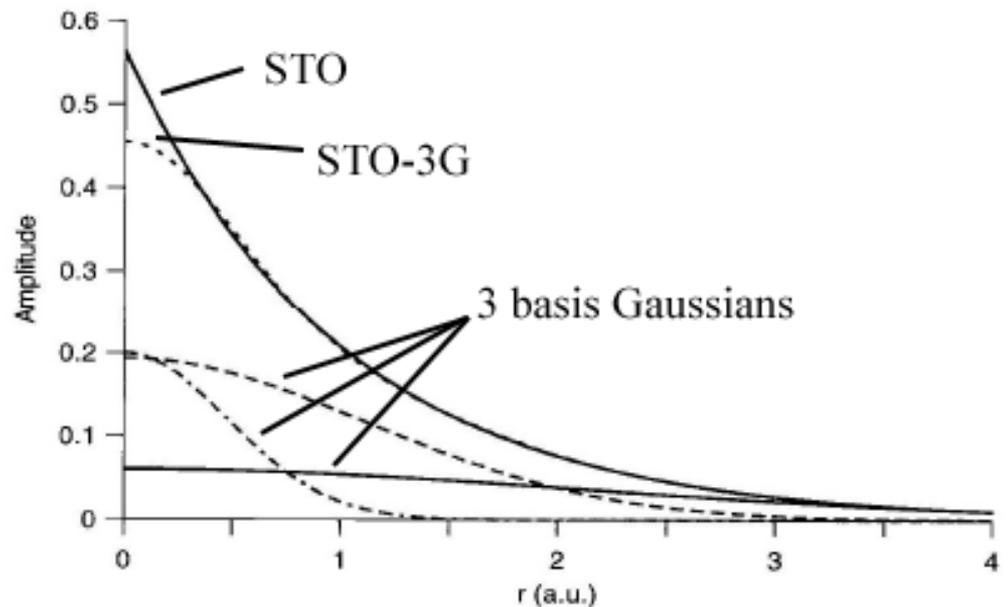
GTO

- ▶ Wrong
- ▶ Wrong
- ▶ Gaussian x gaussian = gaussian
- ▶ Mathematically & computationally easier

Contracted Gaussian Functions (CFG)

- ▶ Product of 2 gaussians is another gaussian
- ▶ Integration is easy (versus STO's)
- ▶ Loss of accuracy BUT increase the number of gaussians used
- ▶ STO-nG, where n = # GTOs used

$$\eta_{\tau}^{CGF} = \sum_1^n d_{n\tau} \eta^{GTO}$$



-C			
S	3	1.00	
	0.1722560000D+03	0.6176690000D-01	
	0.2591090000D+02	0.3587940000D+00	3-21G
	0.5533350000D+01	0.7007130000D+00	
SP	2	1.00	
	0.3664980000D+01	-0.3958970000D+00	0.2364600000D+00
	0.7705450000D+00	0.1215840000D+01	0.8606190000D+00
SP	1	1.00	
	0.1958570000D+00	0.1000000000D+01	0.1000000000D+01
	α	d_s	d_p

$$\varphi_i = \sum_{n=1}^N d_{ni} \eta_n \quad \eta^{GTO}(\mathbf{r}) = x^l y^m z^n e^{-\alpha r^2} = \varphi^{GF}$$

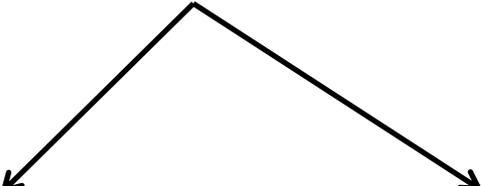
$$\varphi_{2s}(r) = \sum_{i=1}^3 d_{2s} \varphi_{2s}^{GF}(r, \alpha_{2s}) + d'_{2s} \varphi_{2s}^{GF}(r, \alpha'_{2s})$$

Sum of x G's
w/ c_i's

$$\varphi_{2s}(r) = -0.395 \varphi_{2s}^{GF}(r, 3.664) + 1.215 \varphi_{2s}^{GF}(r, 0.771) + 1.00 \varphi_{2s}^{GF}(r, 0.195)$$

Extended Basis Set: Split Valence

- ▶ Minimal basis set: STO-3G
 - A single CGF for each atomic orbital
- ▶ Double-Zeta (DZ) Basis Sets
 - Multiple basis functions corresponding to each atomic orbital = greater flexibility
 - Inert core orbitals: single CGF
 - Valence orbitals: double set of CGF's


$$\varphi_{2s}(\mathbf{r}) = d\varphi_{2s}^{STO}(r, \zeta_1) + d\varphi_{2s}^{STO}(r, \zeta_2)$$

Extended Basis Sets (cont.)

3-21G

- ▶ Core = 1s orbital, a sum of 3 gaussians
- ▶ Each valence orbital is split into 2 parts
- ▶ Valence = 2s & 2p orbitals:
 - Inner part a sum of 2 gaussians
 - Outer part is 1 gaussian

▶ Triple-Zeta (TZ)

- Inner = 1 CGF
- Valence = 3 CGF's
- 6-311G

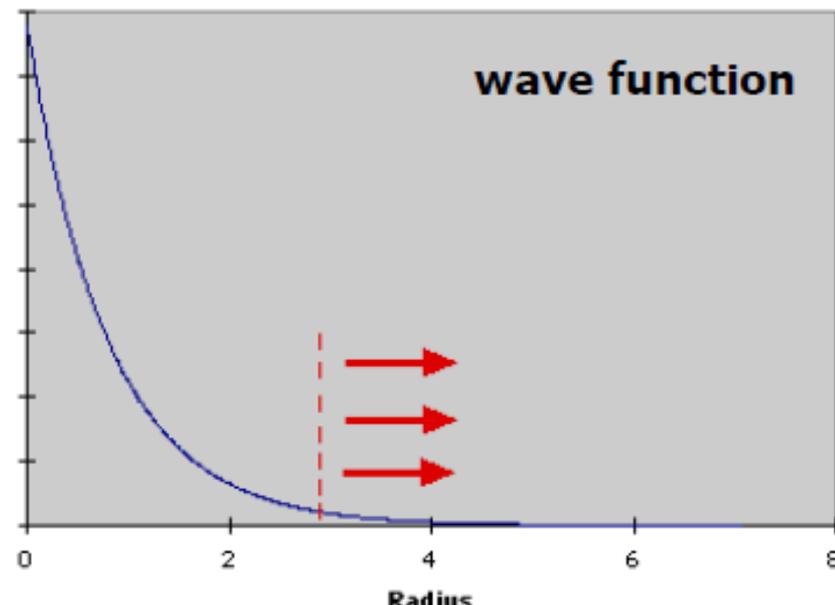
Additional Basis Functions

- ▶ Polarization Functions:
 - Distortion of molecular orbitals due to the molecular environment
 - Functions with a higher angular momentum than any occupied atomic orbital
 - p-functions for H-He **
 - d-functions for Li-Ca *
 - f-functions for transition metals
 - 6-31G(d,p) = 6-31G** = polarization functions added to elements H-Ca

Additional Basis Functions

▶ Diffuse Functions:

- Electrons are bound looser in anions, neutral molecules w/ unshared pairs, zwitterions
- Wave function tail is important
- Remember,
 - $\phi = \text{Constants} \times \exp[-\alpha r^2]$
 - For longer tails, this means
 - $0.1 > \alpha > 0.01$
 - Large radial extent
- 6-31 ' 'G (*'s here for polar.)
- + diffuse functions on first row
- ++ diffuse function on H



Other Basis Sets

These are “*about*” equivalent

Pople split valence	Pople valence triple zeta	Dunning correlation consistent	
6-31G	6-311G	cc-pvdz	aug-cc-pvdz
6-31+G*	6-311+G*	cc-pvtz	aug-cc-pvtz
6-31++G**	6-311++G**	cc-pvqz	aug-cc-pvqz

Diffuse functions
Long-distance interactions
anions

Polarization functions
Angular charge distribution
flexibility

Levels of theory: Hartree–Fock

▶ The variational principle.

- If Ψ is the true wavefunction, then for any model antisymmetric wavefunction Φ

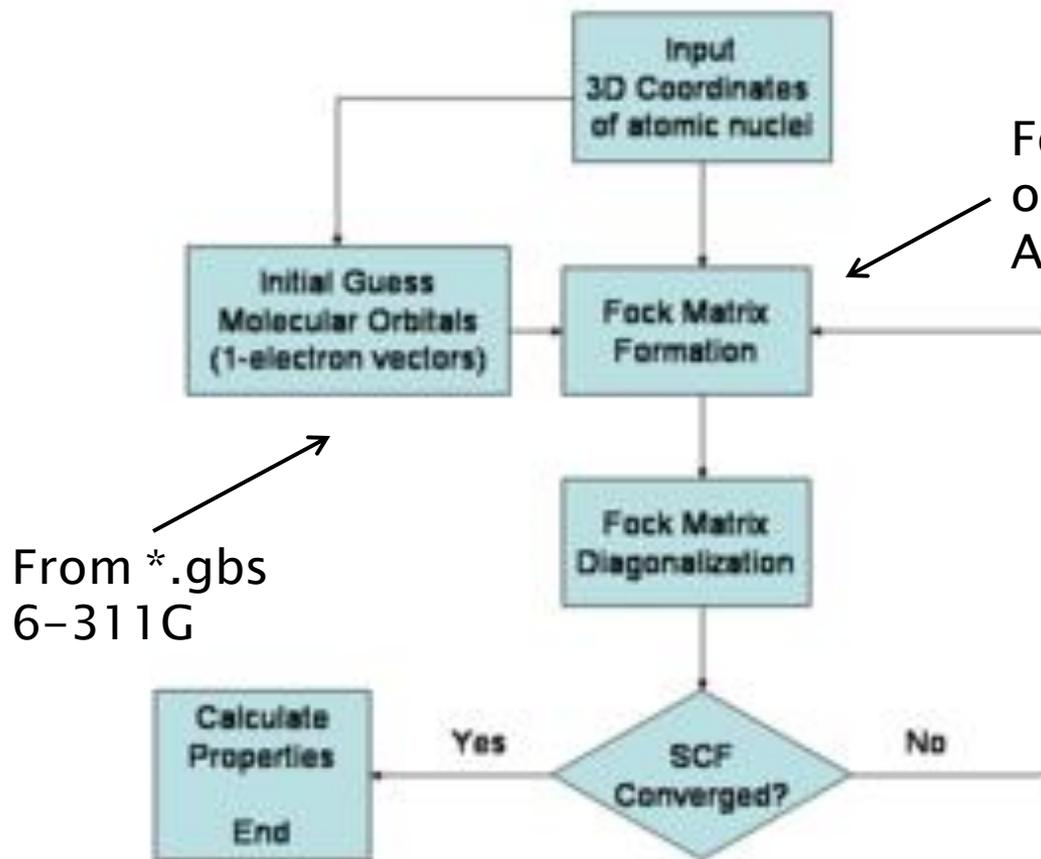
$$E(\Phi^{\text{model}}) > \text{true } E(\Psi)$$

- To find the HF w.f.:
 - Minimize the energy w.r.t. changes in the orbitals
 - $\phi_i \rightarrow \phi_i + \delta\phi_i$
 - Adjustable parameters, the c_{ni} 's, the orbital coeff's
- ▶ E_{HF} forms an upper bound, $E_{\text{exact}} - E_{\text{HF}} = E_{\text{correlation}}$
- HF neglects electron correlation

Simplistic & typical routine Self-consistent field method

Next section

Self-consistent =
Iterative, step by step



Formulation
orbital combo's
Adjust -311

From *.gbs
6-311G

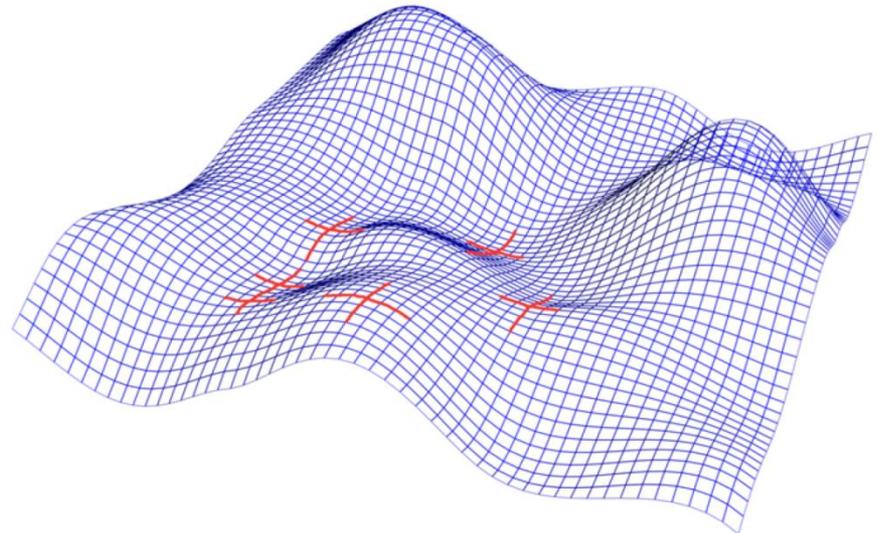
Field (a.k.a. mean field)
where each particle is
subjected to a mean field
(average) created by all
the other particles

Molecular Geometry Optimization

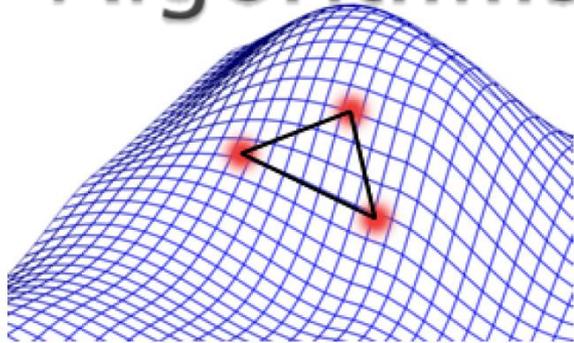
- ▶ Hessian matrix is the square matrix of second-order partial derivatives of a function; that is, it describes the local curvature of a function of many variables.

$$H_{ij} = \frac{\partial^2 E}{\partial x_i \partial x_j}$$

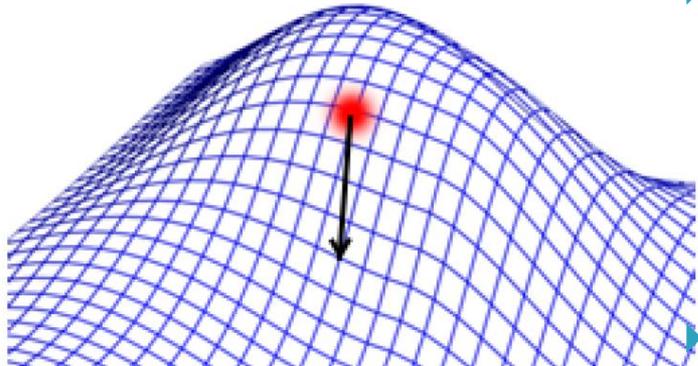
All H eigenvalues positive = Minimum
All H eigenvalues negative = Maximum
Mix of pos. & neg. eigenvalues = Saddle



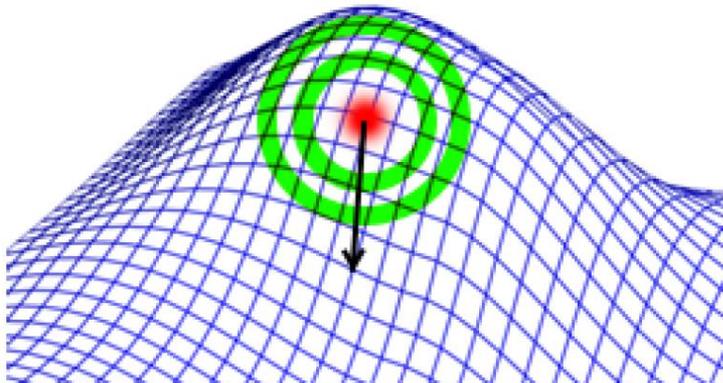
Molecular Geometry Optimization Algorithms



- ▶ Energy only
 - Simplest
 - Might not correspond to the true minimum



- ▶ Gradient following (1st order)
 - Line search, backtracking, gradient descent (steepest descent)
 - Less robust, gradient evaluation at every iteration step



- ▶ Higher order
 - Newton–Raphson, Berny, RFO, GDIIS, ODIIS
 - Hessians & higher order derivations at every step
 - Powerful, more time consuming, VERY dependent on initial configuration /guess

Higher orders of theory

Configuration Interaction

- HF theory is limited by its neglect of electron–electron correlation.
 - Artificially, electrons interact with a SCF, not individual e 's.
 - Most higher ordered levels of theory begin with HF formulation of the w.f.
- “Full CI” includes the Hartree–Fock ground–state determinant and all possible variations.

$$\Psi = a_0 \Psi_0 + \sum_{s=1}^{\infty} a_s \Psi_s$$

- The wavefunction becomes where s includes all combinations of substituting electrons into HF *virtual orbitals*.
 - CISD(T) = Config Inter. of singlets, doublets; triplets combinations
 - The a_s 's are optimized; not so practical if s goes to infinity!
- Partial CI's are useful, and takes less time... ~4 months?

Levels of Theory

Semi-empirical methods
MNDO,AM1,PM3



Hartree-Fock
HF-SCF

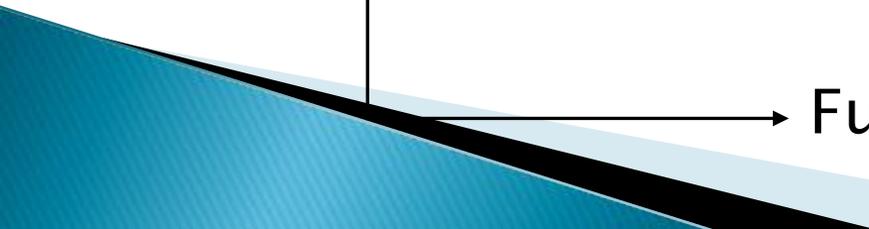
Excitation Hierarchy
CIS,CISD,CISDT
CCS,CCSD,CCSDT

Perturbation Hierarchy
MP2,MP3,MP4

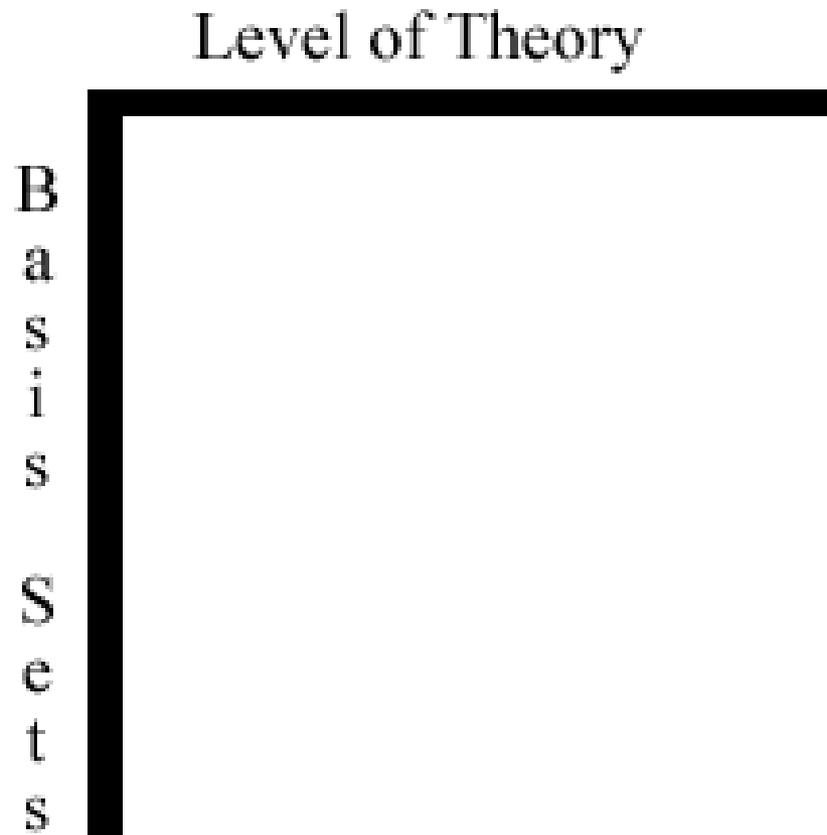
Multiconfigurational HF
MCSCF,CASSCF

Multireference Perturbation
CASPT2, CASPT3

Full CI

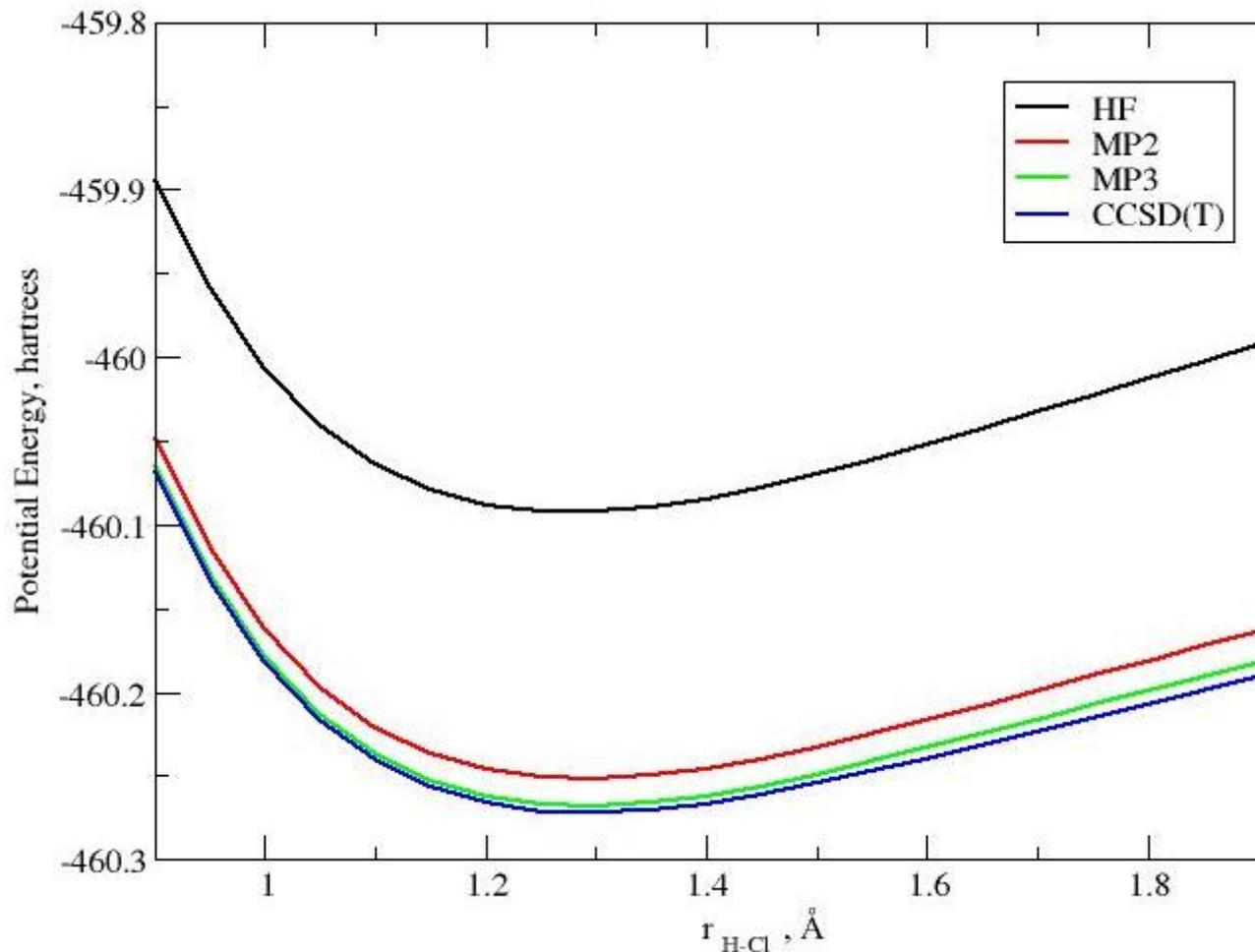


Pople's Diagram



HCl Potential Energy Scan, comparison of levels of theory

Basis Set: aug-cc-pvdz



$\Delta E = \text{CCSD(T)} - \text{xx}$
(Kcal/mol)

HF: 112.931
No é corr.!!

MP2: 12.881

MP3: 2.672

1 a.u. =
1 hartree =
627.5095
Kcal/mol

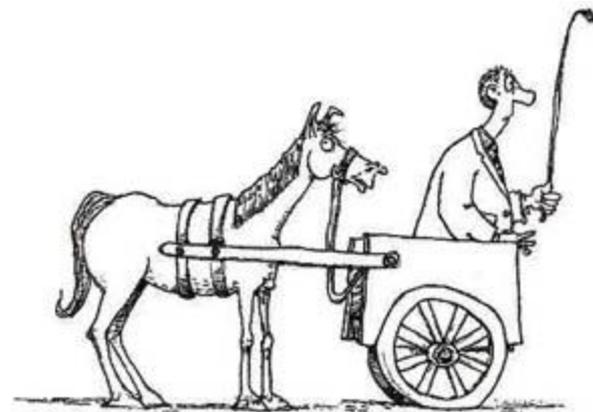
Gaussian runtime

- ▶ Understand how the input file is constructed
 - ▶ Specifying atomic coordinates
 - ▶ How to submit your job to a HPC
- 

Defining Atomic Coordinates

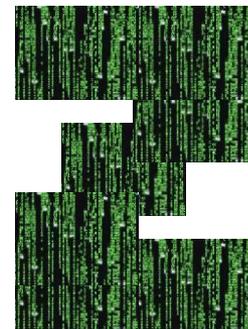
▶ Cartesian Coordinates

- René Descartes
- *Atom x y z*

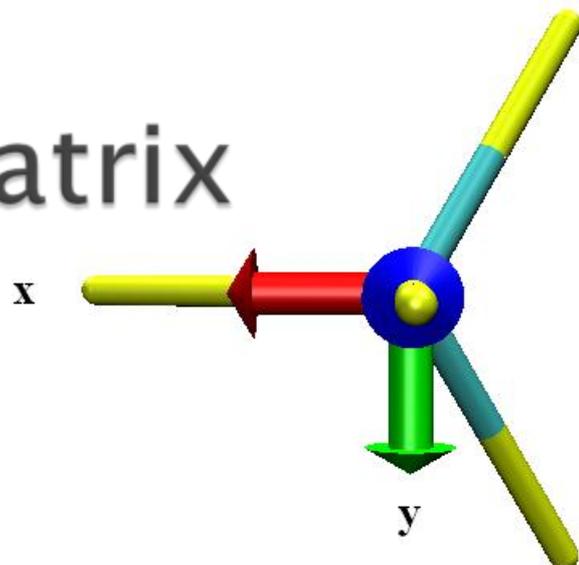
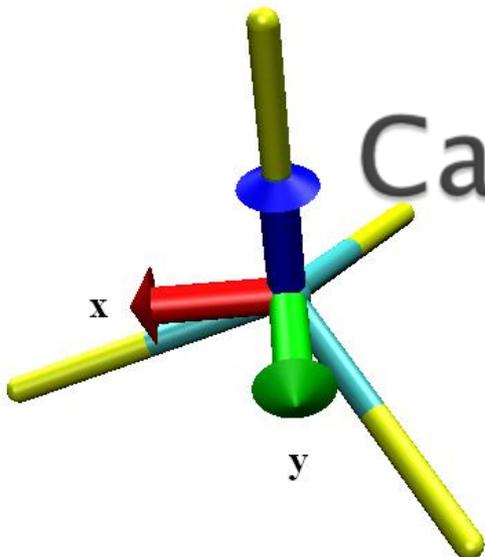


▶ Z matrix

- Internal Coordinate System
- *newatom*
newatom atom1 bond-length
newatom atom1 bond-length atom2 bondangle
newatom atom1 bond-length atom2 bondangle atom3 dihedral-angle



Cartesian vs. Z-Matrix



5

```
C 0.000 0.000 0.000
H 0.000 0.000 1.089
H 1.027 0.000 -0.363
H -0.513 -0.889 -0.363
H -0.513 0.889 -0.363
```

C

```
H 1 1.089
H 1 1.089 2 109.47
H 1 1.089 2 109.47 3 120.000
H 1 1.089 2 109.47 3 -120.000
```

This **atom** is connected to **atom1** having a **bond-length** *and* with **atom2** has a **bondangle** *and* with **atom3** has a **dihedral angle**

Molden

Select Point:

First
Next
Prev

Miscellaneous:

Dens. Mode
Read File
ZMAT Editor
PostScript

Draw Mode:

Solid
 StickColor
 Shade
 Perspect.
 Label
 BackBone

Render Forces:

Forces
Dist. Scale
Dist. Scale
Urscale

Calculate :

Distance
Angle
Dihedral

Convergence:

SF conv.
Geom. conv.

Zoom:

In
Out

Status line:

No coordinates found !

BondLength BondAngle DiHedral Close

C							
C	1	1.400000					
C	2	1.400000	1	120.000000			
C	3	1.400000	2	120.000000	1	0.000000	0
C	4	1.400000	3	120.000000	2	0.000000	0
C	5	1.400000	4	120.000000	3	0.000000	0
H	2	1.089000	1	120.000000	3	180.000000	0
C	3	1.450000	2	120.000000	4	180.000000	0

Apply Changes to current Z-Mat Clear Status All Variables

Cancel Non-Applied Changes

Delete Line Add Line

Substitute atom by Fragment

New Z-mat MapXYZ/Optimise

New Z-mat from screen coordinates:

Reorder Z-matrix

Select by cursor DeSelect

Apply Selection

Linked Variable

Write Z-Matrix Submit Job

File name ?

Format: Gamess Gaussian Mopac Cartesian

Gaussian Job Details

%chk=water.chk

%NProcLinda=2

%NProcShared=4

mp2/6-31g* opt=tight freq

<blank line>

Title Line

<blank line>

O 1

O 0.0 0.0 0.0

H 0.0 1.0 0.0

H 0.0 0.0 1.1

<blank line>

Checkpoint file

nodes

#ppn

Job description

Title

Charge Multi.

Coordinates

...

...

Blank line !

%chk=water.chk

%NProcLinda=2

%NProcShared=4

mp2/6-31g* opt=tight freq

<blank line>

Title Line

<blank line>

O 1

O

H 1 1.0

H 1 1.0 2 104.5

<blank line>

Gaussian Job Details

```
%chk=water.chk  
%NProcLinda=2  
%NProcShared=4  
# mp2/6-31g* opt=tight freq  
<blank line>  
Title Line  
<blank line>  
0 1  
O  
H 1 1.0  
H 1 1.0 2 104.5  
<blank line>
```

Variables section
Defining used variables

```
%chk=water.chk  
%NProcLinda=2  
%NProcShared=4  
# mp2/6-31g* opt=tight freq  
<blank line>  
Title Line  
<blank line>  
0 1  
O  
H 1 roh1  
H 1 roh1 2 ahoh  
<blank line>  
roh1 1.0  
ahoh 104.5
```

If defined by calculation type: bond
stretches, isotopes, continue (serial
job processing)

Gaussian on LSU HPC / LONI

- ▶ Add “+gaussian-03” to your .soft file
- ▶ Resoft

- ▶ On IBM P5 systems
- ▶ Command is
 - runG03 inputfile nprocs queue
 - runG03 input 8 MP5L
 - (LL) ‘llclass’ for queue types

Gaussian on Dell x86 maxchines

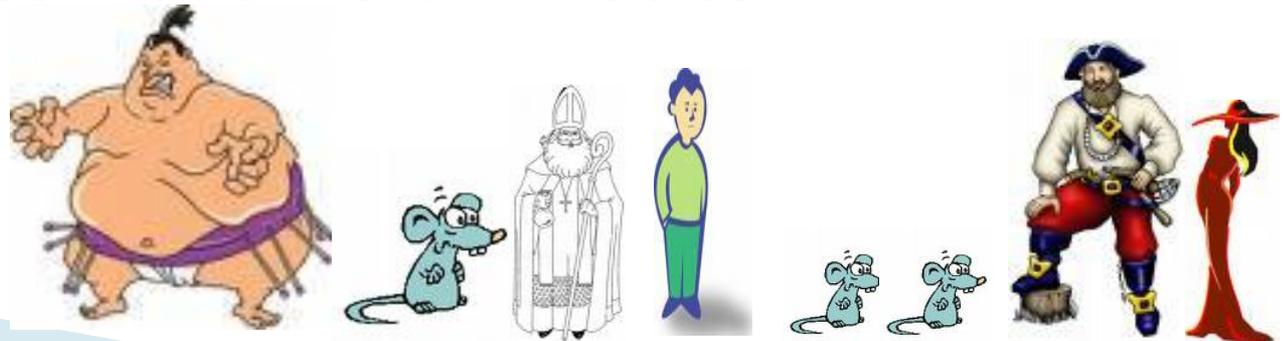
Add the softenv key & resoft

Sample submission script, 'qsub subscript'

```
#!/bin/tcsh
#PBS -A your_allocation      # specify the allocation. Change it to your allocation
#PBS -q checkpt              # the queue to be used.
#PBS -2 nodes=1:ppn=4       # Number of nodes and processors
#PBS -l walltime=1:00:00    # requested Wall-clock time.
#PBS -o g03_output          # name of the standard out file to be "output-file".
#PBS -j oe                   # standard error output merge to the standard output file.
#PBS -N g03test              # name of the job (that will appear on executing the qstat command).
set NPROCS=`wc -l $PBS_NODEFILE |gawk '/{print $1}`
setenv GAUSS_SCRDIR /scratch/$USER
set NODELIST = ( -vv -nodelist "" `cat $PBS_NODEFILE` "" -mp 4)
setenv GAUSS_LFLAGS " $NODELIST "
# cd to the directory with Your input file
cd ~USERNAME/g03test
# Change this line to reflect your input file and output file
g03 < test.inp > g03job.out
```

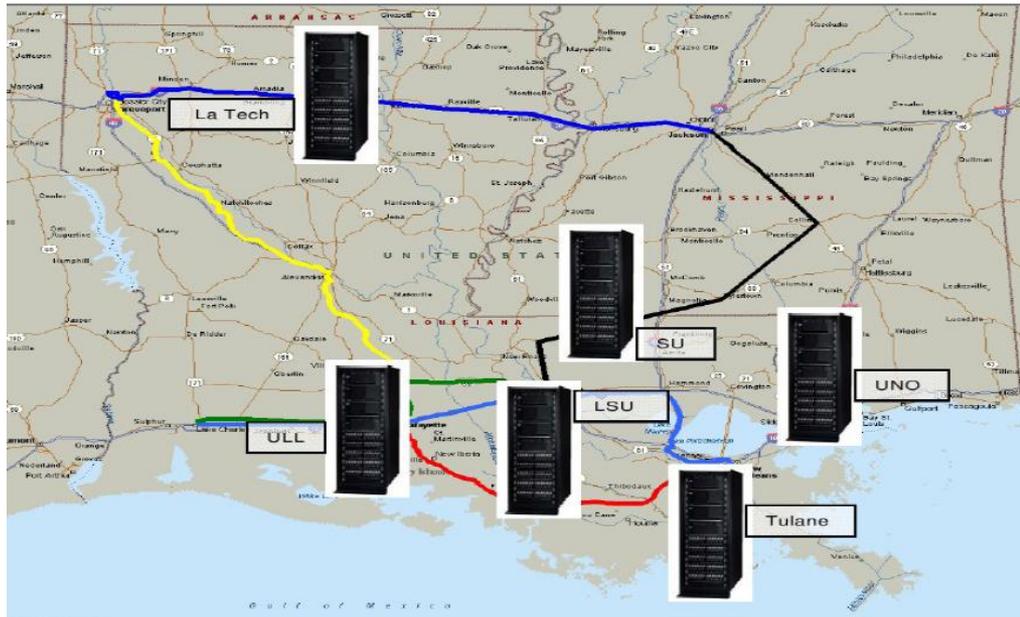
Fighting the Queue

- ▶ **Not all simulations are equal!**
- ▶ **Benchmark your simulation**
 - Walltime vs. # processors
 - Walltime vs. # nodes (w/ half procs idle)
 - Find the optimum configuration
 - Decrease walltime to 1 hour
- ▶ Walltime slightly greater than time needed
- ▶ **THEN:** Tailor your submission script to reflect **YOUR** needed walltime & # nodes



Fighting the Queue

- ▶ Check & monitor the load on other HPCs
- ▶ NOTE: Gaussian license is site specific



- ▶ Researchers tend to mainly work with a few clusters and submit different job types/sizes

Analysis– some pointers

Theory	Basis Set	Ground State Bond Length		
		(hartrees)	(Angstroms)	% Error
HF	3-21G*	-457.981	1.2671	-0.58%
HF	6-31G*	-460.06	1.2662	-0.65%
HF	6-311G*	-460.087	1.2713	-0.26%
HF	cc-pvDZ	-460.09	1.2776	0.24%
HF	cc-pvTZ	-460.107	1.2672	-0.58%
HF	aug-cc-pvDZ	-460.093	1.2767	0.17%
HF	aug-cc-pvTZ	-460.108	1.2675	-0.55%
B3LYP	3-21G*	-458.709	1.2888	1.12%
B3LYP	6-31G*	-460.796	1.2895	1.18%
mp2	6-311G**	-460.244	1.2734	0.09%
CCISD	3-21G*	-458.145	1.2841	0.75%
CCISD	6-31G*	-460.218	1.2854	0.85%
CCISD	cc-pVTZ	-460.371	1.2749	0.03%
Experiment			1.2746	

Geometry optimization of HCl

Is CCISD/cc-pVTZ the best?

Is bond length a good judgment of accuracy of the model?

References

- ▶ <http://www.gaussian.com>
 - ▶ http://www.gaussian.com/g_ur/keywords.htm
 - Gaussview

 - ▶ <http://www.cmbi.ru.nl/molden/molden.html>

 - ▶ <http://www.cscs.ch/molekel/>
- 