

# NANO266 Lab 1

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## 1 Introduction

We will begin our lab sessions with a gentle introduction into quantum mechanical modeling of molecules. For this purpose, we will be using computational chemistry techniques to study reaction energies, geometries, vibrational frequencies, etc. We will be using NWChem, an open source quantum chemistry code. Note that all calculations in this lab are fairly simple and on very small molecules. So you can be run them on a modern desktop or laptop in serial mode, i.e., you do not actually need access to a supercomputing cluster to perform these calculations.

For this lab, we will be studying one of the most important reactions in the world - the formation of ammonia from nitrogen and hydrogen. Through the Haber-Bosch process, this is the main industrial reaction for the production of ammonia, which is then used in the production of fertilizers, etc. It is estimated that one-third of the Earth's population is sustained by fertilizer generated from ammonia produced by the Haber process.

In this lab, almost all analyses will be done using Unix-based command line tools as it is important for you to learn how to find data efficiently using these tools. Subsequent labs will provide you with the opportunity to use Python to automate and analyze data more effectively. You are of course welcome to write your own scripts if you have the know-how.

## 2 Initial setup

It is assumed that you have already followed the instructions in the README.md in the root labs folder and have access to nwchem, either on XSEDE or on your own computer or virtual machine. Do a git pull so that you are up to date with the repo. Also, read through the README.md file in the main labs folder and make sure that you have NWChem setup properly. Try typing `nwchem` in your terminal to make sure that everything is working. You will get an error message because there is no input file, but that's not a big deal.

Once you are done with the above, make sure you are in the lab1 folder by doing:

```
cd <path/to/repo>/labs/lab1
```

### 3 Q1 (10 points): Geometry optimization and energy of H2

We will start with one of the simplest molecules, diatomic hydrogen. In the directory, we have provided a sample H2.nw input file. First, open up the input file and understand its structure. Here's a replica of the input file with comments added.

```
memory total 1000 mb      # This specifies the memory for the job.

geometry units angstroms  # This section provides a summary of the input
  H 0 0 0                 # geometry of the molecule. You usually get this
  H 0 0 0.7414            # from an experimental source or some chemistry
end                       # rules

# This nwchem job comprises three steps - geometry optimization, a frequency
# calculation, and a final energy calculation at a larger basis set.

title "H2 dft optimize"   # This is just the title
charge 0                  # We are doing calculations on neutral H2.
basis                    # Specifies the basis set for each atomic species
  H library "6-31G"
end
dft
  mult 1                  # Spin multiplicity of 1, i.e., singlet state
  xc b3lyp                # Exchange functional used is B3LYP.
end
task dft optimize         # Specify that we want to do a geometry optimization

title "H2 dft freq"
charge 0
basis
  H library "6-31G"      # The same basis set must be used for the frequency calculation.
end
dft
  mult 1
  xc B3LYP               # The same functional must be used for the frequency calculation
end
task dft freq            # Specify that we want to do a frequency calculation

title "H2 dft energy"
charge 0
basis
  H library "6-311G"     # A larger basis set is used to get better energies   end
dft
  mult 1
  xc b3lyp
end
task dft energy          # Specify that we want to do an energy calculation
```

Let us first create a separate directory to run the calculation. This makes it easier for us to cleanup after we are done.

```
mkdir scratch
cd scratch
```

Now, we copy the input files we want over, and run `nwchem`:

```
cp ../H2.nw .
nwchem H2.nw > H2.nwout
```

After a very short while, the calculation should complete and the results are in the `H2.nwout` file. To find the final coordinates, search for the final occurrence of:

```
grep -A 8 "geometry" -> H2.nwout
```

This command finds all instances that `"geometry" ->` occurs in `H2.nwout` and prints out 8 lines after each occurrence.

To get the final total energy, we can use `grep` from the command line:

```
grep "Total_DFT_energy" H2.nwout
```

The last energy line gives the total energy in **Hartree**.

For this question, record down the final bond length of H2 in angstroms and the final total energy in eV. Keep all output files until the end of the lab.

#### 4 Q2 (20 points): Geometry optimization and energy of N2

Repeat Q1, but this time with N2. For this question, copy `H2.nw` to `N2.nw` and then modify the file accordingly. The experimental N2 bond length is around 1.1 angstroms.

Again, record down the final bond length of H2 in angstroms and the final total energy in eV.

#### 5 Q3 (20 points): Geometry optimization and energy of NH3

The geometry of ammonia is somewhat more complex, so we have provided an `NH3.nw` file. Perform the same calculation as in Q1.

For NH3, record down the final N-H bond lengths, and also, calculate the angle between the bonds. Compare the calculated values with the experimental ones. Cite the source of your experimental data (e.g., by providing a journal citation or a weblink.)

Also record down the final energy of the NH3 molecule in eV.

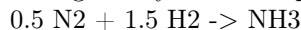
#### 6 Q4 (20 points): Formation enthalpy of NH3.

Calculate the formation enthalpy (per molecule) of NH3 in kJ/mol. To do this, you need not only the energies from Q1-Q3, you also need to extract the thermal correction to the enthalpy. For example,

```
grep "Thermal_correction_to_Enthalpy" H2.nwout
```

Note the units stipulated by NWChem in the output.

The enthalpy  $H$  is then given by the energy + the correction. The formation energy of NH<sub>3</sub> is given by the enthalpy change of the following:



Compare your calculated formation enthalpy with experimental values. The NIST Chemistry Webbook (<http://webbook.nist.gov/chemistry/>) is a good source of data for many common molecules.

## 7 Q5 (30 points)

In this question, you will investigate the effect that functional choice and basis set choice has on the formation enthalpy of NH<sub>3</sub>. Repeat Q1-Q4, but now modify your input files to experiment with the HF, PBE and B3LYP functionals. You can also modify the basis set used between 6-31g and 6-311g. Note that you can either modify the functional or basis set for the geometry optimization and frequency step, or the final energy calculation step, or both.

Discuss the effect of the functional and basis set on the geometries and final energies. What would be a general recommended strategy for other similar calculations if you want to perform the calculations as efficiently as possible while maintaining relatively good accuracy?

Note that this is an open-ended question, and there is no real right or wrong answer. It is more important for you to understand the tradeoffs and come up with a good justification for your recommendation.

## 8 Bonus (10 points)

The formation enthalpy of ammonia is negative, but generally the reaction does not take place (or does so very slowly) under normal conditions. That is why the Haber process is performed under high pressures and temperatures with a catalyst. The main reason is that N<sub>2</sub> is very unreactive. Can you give a rough estimate of the reaction barrier using a few simple calculations, assuming that one has to break the N<sub>2</sub> triple bond for the reaction?