

NWChem 101: a tutorial for experienced quantum chemists

Jeff R. Hammond
Leadership Computing Facility
Argonne National Laboratory
Argonne, IL 60439
`jhammond@mcs.anl.gov`

Last edited: December 21, 2009

1 Introduction

The document (101) introduces the molecular quantum chemistry capability of NWChem in pedagogical manner. It should not replace the *NWChem User Manual* (UM), but rather supplement it. The UM is organized by functionality and documents all options while 101 is organized from the most basic features to the most advanced.

1.1 Input File Structure

Figure 1 is a basis NWChem input file which introduces a number of key elements. The important input elements are described briefly in Table 1.1.

The `echo` should be considered essentially since it is the best way to prevent data rot. It is common to modify input files throughout a project, thus the only way to preserve the exact settings used for a given calculation is to retain the input file with the corresponding output. It is tedious to extract the original input settings from the output file, and in some circumstances, these may not be available at all (e.g. when `print low` or `print none` are used).

Figure 1: A very simple input file.

```
1  echo
2
3  start water
4
5  title "a simple calculation of water"
6
7  # COMMENT: these options are system specific
8  memory total 2000 mb
9  permanent_dir /home/jeff/scratch/nwchem
10 scratch_dir /home/jeff/scratch/nwchem
11
12 geometry angstroms # angstroms are the default units
13   O   0.00000000  0.00000000  0.11726921
14   H   0.75698224  0.00000000 -0.46907685
15   H  -0.75698224  0.00000000 -0.46907685
16 end
17
18 basis
19 * library 6-31G*
20 end
21
22 task scf energy
```

The **start** directive is also essential since it controls the names of files that NWChem will create. It is critical that the RTDB file be unique, as running two jobs simultaneously with the same prefix will frequently result in an error or incorrect behavior. Using **title** is optional and less useful than **echo** with a commented input file.

Comments can be introduced to input files using the **#** symbol. These comments will be echoed along with the rest of the input file. One can comment after a regular input line as demonstrated in the example above. Suggested comments include: (1) a description of where the input geometry came from (e.g. "this is the B3LYP/6-31G optimized geometry"), (2) the citation output from the EMSL Basis Set Exchange (bse.pnl.gov), when this is the source of the basis set input, and (3) annotation of undocumented options users, system-specific parameters and **set** directives (described in Section 3.5).

1.2 Geometry Input

Geometry input may be specified in both cartesian and zmatrix format. Point-group symmetry will be detected automatically but in some cases it must be set manually. First, not all modules implement non-degenerate point-group symmetry, in which case the symmetry must be set to D_{2h} or one of its subgroups (see Figure 2). Some modules detect non-degenerate point-groups and disable symmetry altogether so it is recommended that one set the appropriate symmetry manually for TCE jobs. In other cases, it is desirable to use symmetry to generate a molecule from a fragment. An example is given in Figure 3. Finally, an example of a non-trivial zmatrix geometry input is given in Figure 4.

The UM is an excellent resource for the geometry input block so we will not elaborate further on this topic.

Table 1: See text for details.		
directive	line	purpose
<code>echo</code>	1	prints the input file to output stream
<code>start</code>	3	initializes the RTDB and other files using given prefix
<code>title</code>	5	this title will be printed with each module's output
<code>#</code>	7, 12	stops parsing at the <code>#</code> symbol
<code>memory</code>	8	configures memory usage
<code>permanent_dir</code>	9	the directory where persistent files will be stored
<code>scratch_dir</code>	10	the directory where scratch files will be stored
<code>geometry</code>	12	input block for the molecular geometry
<code>basis</code>	18	input block for the basis set
<code>task</code>	22	NWChem will execute the directive given here

Figure 2: The geometry input for an atom with a forced reduction in symmetry.

```

1  geometry
2      symmetry d2h
3      Ne 0.0 0.0 0.0
4  end

```

Figure 3: Geometry input for C₆O using symmetry completion.

```
1 geometry print xyz units bohr
2 symmetry d2h
3 C      0.0000000000      6.5776597000      1.3147448000■
4 C      2.2251790000      5.7277170000      2.6899811000■
5 C      1.3752363000      4.3524807000      4.9151601000■
6 C      4.3524807000      4.9151601000      1.3752363000■
7 C      2.6899811000      2.2251790000      5.7277170000■
8 C      4.9151601000      1.3752363000      4.3524807000■
9 C      5.7277170000      2.6899811000      2.2251790000■
10 C     6.5776597000      1.3147448000      0.0000000000■
11 C     1.3147448000      0.0000000000      6.5776597000■
12 end
```

1.3 Basis Set Input

1.4 Task Input

2 Basis Input Files

2.1 Hartree-Fock (SCF)

2.2 Density-Function-Theory (DFT)

2.3 Perturbation Theory (MP2)

2.4 Coupled-Cluster Theory (CCSD)

2.5 Multi-Configuration SCF (MCSCF)

3 Advanced Input Files

3.1 Multiple Tasks

3.2 Converging SCF/DFT

3.3 Custom Density Functionals

3.4 Undocumented Options⁵

3.5 Set Directives

4 TCE Input Files

Figure 4: A very simple input file.

```
1  geometry units bohr
2      zmatrix
3      X
4      C  1 RXC
5      C  1 RXC  2 A60
6      C  1 RXC  3 A60 2 D180
7      C  1 RXC  4 A60 3 D180
8      C  1 RXC  5 A60 4 D180
9      C  1 RXC  6 A60 5 D180
10     F  1 RXF  2 A60 7 D180
11     F  1 RXF  3 A60 2 D180
12     F  1 RXF  4 A60 3 D180
13     F  1 RXF  5 A60 4 D180
14     F  1 RXF  6 A60 5 D180
15     F  1 RXF  7 A60 6 D180
16     constants
17         RXC      2.63428
18         RXF      5.14195
19         A60      60.00
20         D180    180.00
21     end
22 end
```