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# **NWCHEM**

# A. Description

NWChem is open source highly scalable high-performance electronic structure computational package for computer simulation of molecular systems, biomolecules, nanostructures, and solid-state systems. It incorporates classical molecular dynamics, Hartree-Fock, Density Functional Theory methods, and a wide range of post-Hartree-Fock methods including Moeller-Plesset and Coupled Cluster techniques for the treatment of electron correlation. See NWChem home page www.nwchem-sw.org for additional information.

### B. How to obtain NWCHEM

NWCHEM is distributed under the terms of the Educational Community License (ECL 2.0). See instructions for obtaining the code.

### C. How to build NWCHEM

Blue Waters does not provide precompiled NWCHEM binaries. The users are expected to obtain their own copy of the code and install it for the personal use.

# C.1 Building CPU version under GNU environment

cd \$HOME

tar zxvf Nwchem-6.6.revision27746-src.2015-10-20.tar.gz cd \$HOME/nwchem-6.6

module swap PrgEnv-cray PrgEnv-gnu module add craype-hugepages8M

export NWCHEM\_TOP=`pwd`

# specify your favourite modules

export NWCHEM\_MODULES="nwdft nwpw driver stepper mp2\_grad rimp2 ccsd

property hessian vib tce"

export FC=ftn
export CC=cc

export MSG\_COMMS=MPI
export TARGET=LINUX64

export NWCHEM\_TARGET=LINUX64
export ARMCI\_NETWORK=MPI-PR

export HAS\_BLAS=yes
export BLAS\_OPT=''
export LIBMPI=''
export USE\_MPI=y
export USE\_MPIF=y
export USE\_MPIF4=y
export USE\_64T032=y

export MA\_USE\_ARMCI\_MEM=y

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```
export BLAS_SIZE=4
export LAPACK_SIZE=4
export SCALAPACK_SIZE=4
export SCALAPACK=-lsci_gnu
export BLASOPT=-lsci_gnu
export GA_DIR=ga-5-4
export USE_NOFSCHECK=TRUE
export USE_N0I0=TRUE
cd $NWCHEM_TOP/src
# perform precision conversion
# apply it only once
# repeated application will break the code
# comment out when recompiling already converted code
make 64_to_32
# compile the code
make realclean
make nwchem_config
make FC=ftn GA_DIR=ga-5-4
```

Successful compilation will create static executable: \$HOME/nwchem-6.6/bin/LINUX64/nwchem

# D. Sample test

Create file input.nw:

```
start guanine
memory total 2000 MB global 1500 MB
title "guanine molecule"
geometry
H 2.5399 1.8161 -0.0360
N 1.5349 1.7071 -0.0144
C 0.5833 2.7012 -0.0107
N -0.6537 2.2306
                  0.0124
C -0.4879 0.8641
                  0.0321
C -1.4774 -0.1880 -0.0008
N -0.8129 -1.4562 0.0346
C 0.5421 -1.6761
                 0.0161
N 1.4396 -0.7242
                 0.0169
C 0.8632 0.5129 -0.0009
H 0.8526 3.7486 -0.0223
0 -2.6927 -0.1362 -0.0450
N 0.9378 -3.0020
                  -0.0675
  1.9335 -3.1015
                  0.0844
H 0.3973 -3.6490
                   0.4919
H -1.4365 -2.2461 -0.0956
end
basis spherical
 * library 6-311++G**
end
scf
noprint "final vectors analysis"
end
```

create run.pbs file having the following content:

#!/bin/bash

maxiter 50 thresh 1000.0

task ccsd energy

freeze core atomic

ccsd

end

### Blue Waters User Portal | NWCHEM

#PBS -l nodes=10:ppn=32:xe

#PBS -1 walltime=00:30:00

#PBS -q debug

#PBS -N ztest

source /opt/modules/default/init/bash

module load craype-hugepages8M

module list

cd \$PBS\_O\_WORKDIR

aprun -n160 -N16 -d 2  $\theta$ -N17 -d 2  $\theta$ -N17 -d 2  $\theta$ -N17 -d 2  $\theta$ -N17 -d 2  $\theta$ -N16 -d 2  $\theta$ -N17 -d 2  $\theta$ -N17 -d 2  $\theta$ -N17 -d 2  $\theta$ -N18 -d 2  $\theta$ -N18

iob.out

Submit the job

qsub run.pbs

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