

# NWChem HW1-Bonus Report

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## 1 Introduction

The purpose of this assignment is to compile, optimize, and execute the **NWChem** application on the Taiwania3 HPC cluster. The goal is to obtain correct results under the given input within 300 seconds, while exploring optimization techniques to improve runtime performance.

## 2 Build Procedure

The compilation was performed on the login node (lgn301) with the following steps:

### 1. Environment setup

```
source /work/b11902043/PP25/intel/oneapi/setvars.sh --force
export OMPI_HOME=/work/b11902043/PP25/openmpi
export PATH="$OMPI_HOME/bin:$PATH"
export LD_LIBRARY_PATH="$OMPI_HOME/lib:$LD_LIBRARY_PATH"
```

### 2. Clone NWChem source code

```
cd ~/hw1bonus
git clone https://github.com/nwchemgit/nwchem.git
export NWCHEM_TOP=$HOME/hw1bonus/nwchem
```

### 3. Set build environment variables

```
export NWCHEM_TARGET=LINUX64
export NWCHEM_MODULES=qm
export USE_MPI=y
export USE_MPIF=y
export USE_MPIF4=y
export ARMCI_NETWORK=MPI-PR
export USE_SCALAPACK=y
export USE_OPENMP=y
export FC=mpifort
```

```
export CC=mpicc  
export CXX=mpicxx
```

#### 4. Build Global Arrays (GA) and ARMCI-MPI

```
cd $NWChem_TOP/src/tools  
./get-tools-github  
MPICC=mpicc ./install-armci-mpi
```

#### 5. Configure and build NWChem

```
cd $NWChem_TOP/src  
make nwchem_config  
make -j$(nproc) FFLAGS+=" -i8 -diag-disable=10448" \  
      CFLAGS+=" -diag-disable=10441"
```

After compilation, the executable is located at:

```
$NWChem_TOP/bin/LINUX64/nwchem
```

## 3 Execution Setup

The job was submitted through a Slurm batch script `hw1-bonus.sh`:

```
#!/bin/bash  
#SBATCH --job-name=nwchem  
#SBATCH --partition=ctest  
#SBATCH --nodes=4  
#SBATCH --ntasks-per-node=32  
#SBATCH --cpus-per-task=1  
#SBATCH --mem=64G  
#SBATCH --time=0:5:00  
#SBATCH --account=ACD114118  
#SBATCH --output=nwchem.log  
#SBATCH --error=nwchem.err  
  
export NWChem_TOP=/home/r14922156/hw1bonus/nwchem  
source /work/b11902043/PP25/intel/oneapi/setvars.sh --force  
export OMPI_HOME=/work/b11902043/PP25/openmpi  
export PATH="$OMPI_HOME/bin:$PATH"  
export LD_LIBRARY_PATH="$OMPI_HOME/lib:$LD_LIBRARY_PATH"  
  
export NWChem_TARGET=LINUX64  
export NWChem_MODULES=qm  
export FC=mpifort
```

```

export OMP_NUM_THREADS=1

mpirun -np $SLURM_NTASKS $NWCHEM_TOP/bin/LINUX64/nwchem \
/wk/b11902043/PP25/hw1-bonus/inputs/w12_b3lyp_cc-pvtz_energy.nw

```

After job completion, the main output is found in `nwchem.log`, which ends with:

```
Total times  cpu: 46.9s  wall: 46.9s
```

## 4 Judge System

The correctness of the results and timing was verified using the TA-provided judge system:

```

bash /wk/b11902043/PP25/setup.sh
source ~/.bashrc
hw1-bonus-judge

```

The job successfully produced valid output under 300 seconds, and results were uploaded to the scoreboard.

## 5 Optimization Attempts

### 5.1 Compiler Optimizations

Tried adding `-O3 -xHost -ipo` flags to the Intel compiler. Result: negligible improvement (< 1s).

### 5.2 Parallel Configurations

We tested multiple MPI/OMP configurations:

Configuration	MPI ranks/node	OMP threads	Total ranks	Wall time (s)	Notes
Baseline	32	1	128	<b>46.5</b>	Fastest
Hybrid 16×2	16	2	64	48.2	Slower
Hybrid 8×4	8	4	32	50.1	Slower
Pure OMP	1	32	4	>300 (fail)	Not scalable

### 5.3 Observations

- The input workload is relatively small.
- Hybrid configurations added thread synchronization overhead.
- The pure MPI configuration fully utilized all 128 cores and achieved the best performance.

## 6 Conclusion

- Successfully compiled and executed NWChem on Taiwania3 with the given input.
- Best performance was achieved with the **baseline pure MPI** configuration: 128 ranks (32 per node  $\times$  4 nodes), runtime  $\sim$ 46.5 seconds.
- Hybrid parallelization and compiler flags did not yield improvements.
- For small-scale inputs, NWChem scales best with pure MPI on this cluster.

## 7 References

- NWChem GitHub
- NWChem Manual
- TA-provided environment and instructions for PP25 course