

ACE3P CW10 Supercomputing Cheat Sheet, Sep 22 2010

Franklin is a supercomputer at NERSC (National Energy Research Center) in Berkeley, CA. It has a scheduler to run computational jobs based on a queue. The user has to use a job submission script to run a job. Once a job is submitted, the scheduler will decide when and where the job will be executed.

- You first need to get a NERSC Franklin account
(send your information to ace3p@slac.stanford.edu :
First and Last names, nationality, phone, address, and email)
- General overview at <http://www.nersc.gov/nusers/systems/franklin/>
- Or go directly to
http://www.nersc.gov/nusers/systems/franklin/running_jobs/
- Transfer mesh and input file(s) from your desktop to your directory on Franklin's high-performance file system:

```
scp -pr <problem_dir>  
franklin.nersc.gov:/scratch/scratchdirs/<user>/
```

(Note, that since we want access to the high-performance file system, we need to run our ACE3P simulations from within the /scratch directory)

- Login on Franklin:
`ssh franklin.nersc.gov`
- Go to the above directory:
`cd /scratch/scratchdirs/<user>/<problem_dir>/`
- Note that you can find the ACE3P executables for Franklin in the directory
`~candel/.community/franklin/`

How to run a Job on Franklin:

http://www.nersc.gov/nusers/systems/franklin/running_jobs/

- The following is a sample job submission script. You can find it at
~candel/.community/franklin/run.pbs.

```
#PBS -N JOBNAME
#PBS -q regular          <- which queue to use (regular is fine)
#PBS -l mppwidth=128    <- requested number of CPUs (multiple of 4)
#PBS -l walltime=00:30:00 <- requested time (don't need to use it all)
#PBS -j eo              <- combine standard output + error
cd $PBS_O_WORKDIR      <- go to the right place on compute nodes

### Omega3P (with growing amount of memory per process)
#aprun -n 128 -N 4 ~candel/.community/franklin/omega3p input.o3p
#aprun -n 64 -N 2 ~candel/.community/franklin/omega3p input.o3p
#aprun -n 32 -N 1 ~candel/.community/franklin/omega3p input.o3p

### T3P with VolumeMonitor -> Mode conversion (also parallel)
aprun -n 128 -N 4 ~candel/.community/franklin/t3p input.t3p
aprun -n 64 -N 2 ~candel/.community/franklin/acdtool postprocess
volmontomode in.t3p
```

Notes on the parameters of aprun, the “parallel job starter”

- n specifies how many processes overall (distributed computing with MPI)
- N specifies how many processes per compute node are started

Franklin has 4 CPUs and 8 GB of memory per compute node.

#PBS -l mppwidth=<NCPUS> says how many processors (+memory!) your reserve

Depending on problem size, you can fine-tune the available memory per process:

aprun -n <NCPUS> -N 4 starts 4 processes per node, each has around 2 GB

aprun -n <NCPUS>/2 -N 2 starts 2 processes per node, each has around 4 GB

aprun -n <NCPUS>/4 -N 1 starts 1 process per node, it has all the 8 GB for itself!

For Omega3P/S3P and large problems, it may be better to start <NCPUS>/4 processes with 8 GB each than to use <NCPUS> processes with 2 GB each, as memory scalability is hard and the O/S and support libraries also occupy memory. For all other codes, it is usually OK to use all CPUs (-N 4).

Remember to use Solver: CG for T3P, to reduce memory requirements.

- Submit the job

```
qsub run.pbs
```

- Look at the status of the batch jobs, and get the Job ID

```
qstat -u <username>
```

- Watch the job's printout as it runs

```
tail -f <Job ID>.nid0000X.ER  
(e.g., tail -f 6733359.nid0003.ER)
```

- Cancel the job

```
qdel <Job ID> (e.g., qdel 6733359)
```

- Change the request time (before the job starts)

```
qalter -l walltime=HH:MM:SS <Job ID>
```

- Copy the results back to your computer (from your computer)

```
scp -pr franklin.nersc.gov:<directory> .
```

- Clean up on Franklin:

```
1) Move the results back into your project directory  
mv <directory> /project/projectdirs/gc2/<user>/
```

or, even better

```
2) upload it into HPSS (high performance storage system) for archival
```

please consult <http://www.nersc.gov/nusers/systems/hpss/>