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 <u>ace3p@slac.stanford.edu</u> : First and Last names, nationality, phone, address, and email)
- General overview at http://www.nersc.gov/nusers/systems/franklin/
- Or go directly to
 <u>http://www.nersc.gov/nusers/systems/franklin/running_jobs/</u>
- 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 AC3P executables for Franklin in the directory ~candel/.community/

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/run.pbs.

```
#PBS -N JOBNAME
#PBS -q regular <- which queue to use (regular is fine)
#PBS -1 mppwidth=128 <- requested number of CPUs (multiple of 4)
#PBS -1 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</pre>
```

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

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

<u>Notes on the parameters of aprun, the "parallel job starter"</u> -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 -1 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 gsub run.pbs
- Look at the status of the batch jobs, and get the Job ID gstat -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 -1 walltime=HH:MM <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/