

Day 2

- Brief Introduction
- Bad News
- Good News
- Software Environment
- PBS Directives
- Client Commands
- Job Priority
- MPI

Brief Introduction

- CRI
- CRI Infrastructure
- My Team
- Me
- Yesterday's session

Bad News

- Users not understanding the software environment (later!)
- Scheduling slow for clinical work
- Resource manager problems
- Scheduler crash
- Scratch space issues
- Job Array Problems
- Job Geometry Changing

Software Environment Confusion

- Problem:
 - Users seem to be confused with how the software environment (Lmod) works
- Solution:
 - Cover it again in today's session
 - User Guide

Clinical Scheduling

- Problem
 - Some clinical work that influences patient care has been slow to run
 - Had purchased exclusive nodes on Tarbell
 - Feel users can take over large memory and mid-tier nodes
 - Giving them exclusive nodes on Gardner would let them
- Solution
 - Partitions
 - Reservations

Proposed Limits

- Overall
 - Processor Cores: 500
 - Memory: 2 TB
- Standard Nodes
 - Processor Cores: 500
- Mid-Tier Nodes
 - Processor Cores: 100 cores
- High-Tier Nodes
 - Processor Cores: 14 cores

Torque Issues

- Problem:
 - Error message: “server is shutting down”
 - Problem with the init script used to start/stop Torque
- Solution:
 - Wrote new init script
 - Fixed in the latest version of Torque (released 3/31/17)

Torque Issues

- Problem:
 - PBS Server crashed (three times!!!)
 - This is an issue with validating PBS directives
 - #PBS -t 1-
 - #PBS walltime=1;00:00
- Solution:
 - Hotfix (fixed some problems)
 - Fixed in the latest Torque Release

Moab Crash

- Problem:
 - Moab crashed hard and was down for close to 8 hours
 - This was due to a corrupt job that needed to be tracked down
- Solution:
 - Hotfix (included in next release as well)
 - Still have yet to determine what caused the corrupt job

Scratch Space Issues

- Problem:
 - Two OS Hard Drives were lost on our Scratch Servers
 - Replacements causing problems too
- Solution:
 - Servers have been removed for maintenance
 - Storage at half capacity

Scratch Space Issues

- Problem:
 - Scratch Space can be slow at times
 - Only have two scratch servers instead of four
 - Negative cache
- Solution:
 - Regain full strength in the scratch environment
 - Lengthen time for cache
 - Make sure cache is always up-to-date

Job Array Problems

- Problem:
 - Job Arrays tend to get stuck
 - Will show up in Torque as running but not Moab
- Solution:
 - Restart pbs_server
 - Fixed in the next release (maybe)

Job Geometry Changes

- Problem:
 - Jobs changed from 4 cores to 1 core
 - Only happened to two users
- Solution:
 - Can be fixed on the sysadmin level
 - Working with Adaptive to handle the issue

Good News

- 113 users are now on Gardner
- 74% of CPU hours for March were on Gardner
- Amount of FLOPS completed in 3 months on Gardner is equivalent to 8 months on Tarbell
- Half of Tarbell is decommissioned
- Most of the positive feedback on Gardner has been computational speed
- New contract with Adaptive Computing (3 years)
- Tony

Tarbell Plans

- Half the cluster decommissioned on 3/31/17
- Who should be using Tarbell
 - Have a Gardner account and a qualifying analysis between the two systems
 - Those who have applied and are waiting for a Gardner account
 - Those who have purchased exclusive nodes on Tarbell
 - Graham School biomedical informatics class
- June 2017 – Rest of Tarbell will be decommissioned

Software Environment

- Tarbell -> Environment Modules
 - Flat module system
 - Modules written in TCL
 - Last Update: December 2012

- Gardner -> Lmod
 - Hierarchical module system
 - Modules written in Lua
 - Last Update: August 2016

Lmod Basics

- See which modules are available to be loaded
 - `module avail`
- Load packages
 - `module load <package1> <package2>`
- See which packages are loaded
 - `module list`
- Unload a package
 - `module unload <package>`

Lmod Basics

- **Swap compilers**
 - `module swap gcc/5.4.0 gcc/6.1.0`
- **Find a module by keyword**
 - `module keyword alignment`
- **List all possible versions of a module**
 - `module spider bwa`
- **Print detailed information for a specific module**
 - `module spider bwa/0.7.5`

Lmod Basics

- Save your loaded modules as the default
 - `module save`
- Restore your default modules
 - `module restore`

Lmod Basics

- Clean up environment
 - `module purge`
- Save a named collection
 - `module save <collection>`
- Restore a named collection
 - `module restore <collection>`
- List all the modules in a collection
 - `module describe <collection>`

Lmod Basics

- **Print help message**
 - `module help <package>`
- **Print description**
 - `module whatis <package>`
- **Get help on Lmod**
 - `module help`

Safety Features of Lmod

- Users can only load one version of a module at a time
 - For example, only one BWA module can be loaded
- Can only load one module from a family at a time
 - Compilers
 - MPI
- Conflict
- Prereq
- Prereq_any

How do I find?

- Perl Modules
 - instmodsh
- Python packages
 - pip freeze
- R packages
 - installed.packages()

Elog Example

- Where do we need help?

PBS Directives

- procs vs. ppn
 - Use procs only for MPI
- mem vs. pmem
- Accounting - #PBS -A
- gpus
- features
- mail - #PBS -M, #PBS -m
- Priority - #PBS -p
- Environment variables - #PBS -V, #PBS -v

Client Commands

- Job monitoring (all jobs)
 - qstat
 - showq
- Job monitoring (individual jobs)
 - qstat -f <jobid>
 - checkjob <jobid>
- Partition monitoring (immediate available resources)
 - showbf
- When will my job start?
 - showstart <jobid>
 - Evaluates historical data, reservations, priority backlogs

mjobctl

- Cancel Job: -c <jobid>
- Hold Job: -h user <jobid>
- Rerun Job: -e <jobid>

Reservations

- When to request them?
 - Deadline
 - Large batch of jobs
- showres
 - Shows all reservations that apply to you

Job Arrays

- `qsub -t <range>%<limit>`
- `$PBS_ARRAYID`
- How to track your job arrays?
 - `showq`
 - `checkjob -v`
 - `qstat -t`

Job Dependencies

- #PBS -W depend=type:jobid[:jobid[:jobid...]]

after	This job may be scheduled after jobs jobid have started
afterok	This job may be scheduled after jobs jobid have completed with no errors
afternotok	This job may be scheduled after jobs jobid have completed with errors
afterany	This job may be scheduled after jobs jobid have completed with or without errors
before	After this job begins, jobs jobid may be scheduled
beforeok	After this job completes without errors, jobs jobid may be scheduled
beforenotok	After this job completes with errors, jobs jobid may be scheduled
beforeany	After this job completes with or without errors, jobs jobid may be scheduled

Quality of Service

- #PBS -l qos=<qos_name>
- Standard = 1000
- Premium = 2000
- Biocore = 3000
- Deadline = 4000
- VIP = 5000

Job Priority

- How does that work.
- Starts at 1000
- Add one every scheduler iteration

MPI Example

- Hello World
- Calculating the volume of a molecule

Upcoming Work

- Trickle – qsub replacement
- Dbuilder
- Data staging
- Software statistics (Tony)
- Viewpoint
- Remote Visualization (?)
- User Guide
- Elog
- Future training (?)