

HPC

Introduction to Unix and HPC

What is HPC?

- HPC, or high-performance computing, refers to the <u>application of supercomputers or clusters of</u> <u>computers to computational problems that</u> <u>typically arise through scientific inquiry.</u>
- HPC is useful when a computational problem:
 - <u>Is too large</u> to solve on a conventional laptop or workstation (because it requires too much memory or disk space) or
 - Would take too long (because the algorithm is complex, the dataset is large, or data access is slow) or
 - Are too many High Throughput Computing

Parallelism on HPC

- HPC systems often derive their computational power by <u>exploiting parallelism</u>
- Programs for HPC systems must be split up into many smaller "sub-programs" which can be executed in parallel on different processors
- HPC systems can offer parallelism at a much larger scale, with 100's or 1000's, or (soon) even millions of tasks running concurrently.
- Writing <u>parallel software can be challenging</u>, and many existing software packages do not already support parallelism & may require development.
- NOTE: Many tasks cannot be parallelised

Reasons to use HPC

- You have a program that can be recompiled or reconfigured to use optimized numerical libraries that are available on HPC systems but not on your own system.
- HPC applications are already installed on the HPC machines which is a non-trivial task
- You have a "parallel" problem, e.g. you have a single application that needs to be rerun many times with different parameters.
- You have an application that has already been designed with parallelism
- To make use of the <u>large memory</u> available
- Our facilities are <u>reliable</u> and regularly backed up

When not to use HPC?

- You have a <u>single threaded job</u> which will only run one job at a time (typical of MatLab users)
- You rely on <u>Databases</u>
- You have a lot of <u>data to transfer</u> between your local machine and the HPC on a continuous basis (e.g. per job)
- You <u>need to have a GUI</u> to interact with your program

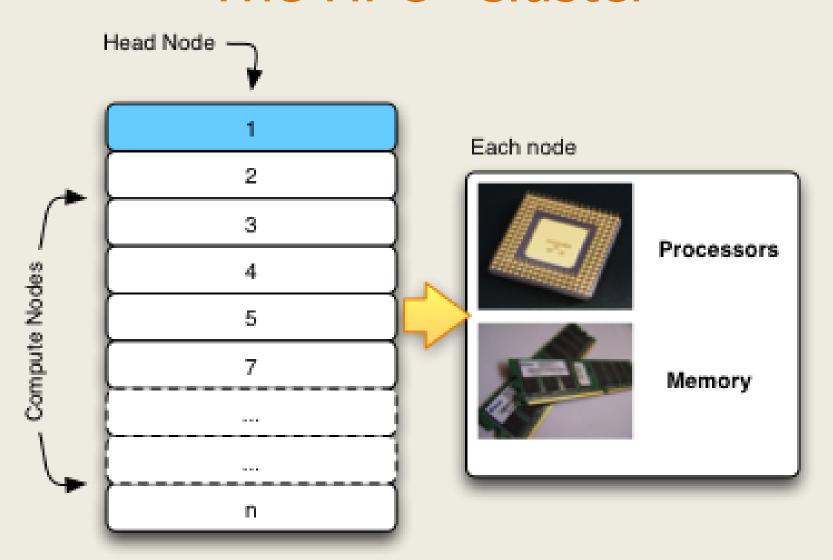
HPC machines

| System | Memory Architecture | Cores | Nodes | Memory |
|---------------------------------------|------------------------|--------|-------|--------|
| Octane (training machine) | Distributed | 48 | 3 | 48GB |
| Orange | Distributed | 1,600 | 100 | 8ТВ |
| NCI – (Vayu) | Distributed | 11,936 | 1492 | 37ТВ |
| NCI – (Name TBA) • available May 2013 | Distributed | 57,472 | 3592 | 158TB |

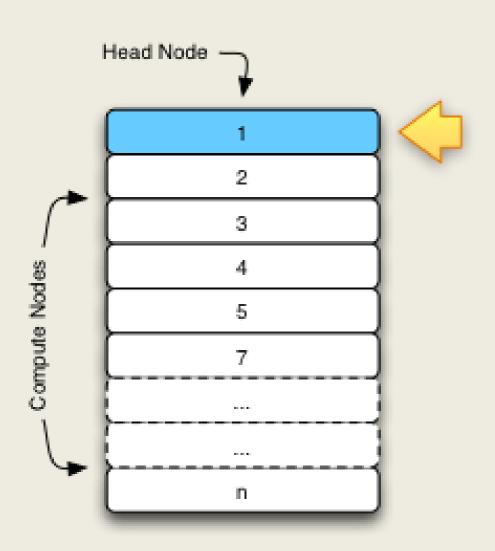
The typical HPC workflow

- In HPC we talk about jobs, these are simply commands we wish to run.
- They are generally time consuming and resource intensive.
- Jobs are typically run non-interactively, but can also be run interactively
- We add our jobs to a queue.
- When the machine has free resources the jobs run.
- Once jobs have completed, we can inspect their output.

The HPC "Cluster"

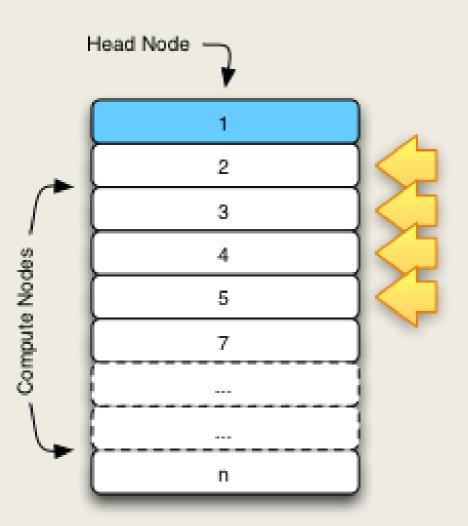


The Head Node



- Interactive programs
- SSH sessions
- Testing
- Compiling
- Queuing jobs

Compute Nodes



- These run your jobs
- Managed by the scheduler
- Typically you will not interact with the nodes directly (some users may need to)

Queuing Systems

- Portable Batch System (PBS) is the name of computer software that performs job scheduling.
 Its primary task is to allocate computational tasks, i.e., batch jobs, among the available computing resources.
- The following versions of PBS are currently available:
 - OpenPBS
 - TORQUE
 - PBS Professional (PBS Pro)
 - ANU PBS
- Guide to PBS: http://hpc.sissa.it/pbs/pbs.html

Queuing Systems cont.

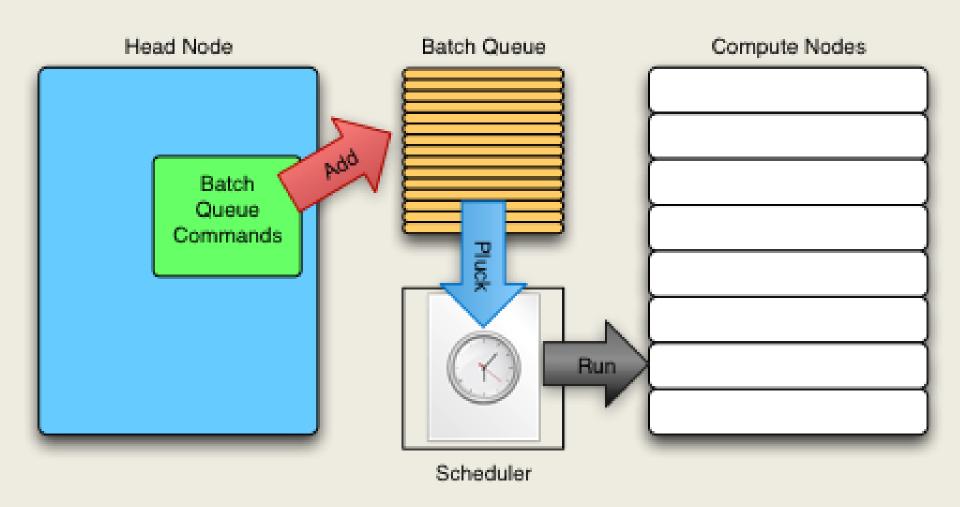
- Another popular batch system is **SLURM** (Simple Linux Utility for Resource Management)
 - Open source, fault-tolerant, and highly scalable cluster management and job scheduling system for large and small Linux clusters.
 - Very useful for use on clusters
 - Platform Tools used by IBM
 - Used by many supercomputers, e.g. TERA 100 at CEA (Europe's most powerful supercomp.)
- Many banks and commercial entities using batch systems

ANU PBS vs PBS Pro

- ANU PBS is a customised version of PBS based on OpenPBS 2.3 maintained by ANU
- Details of ANU PBS modifications are found here: http://anusf.anu.edu.au/~dbs900/PBS/local_modifications.html

| Batch System | ANU PBS | PBS PRO |
|----------------|-------------------------|------------------|
| Machines using | Vayu & NCI New Facility | Orange & Octane |
| Code Base | OpenPBS 2.3 | PBS Professional |
| Licence | ANU Licence | Altair Licence |

The Batch Queuing System



Batch Queuing component

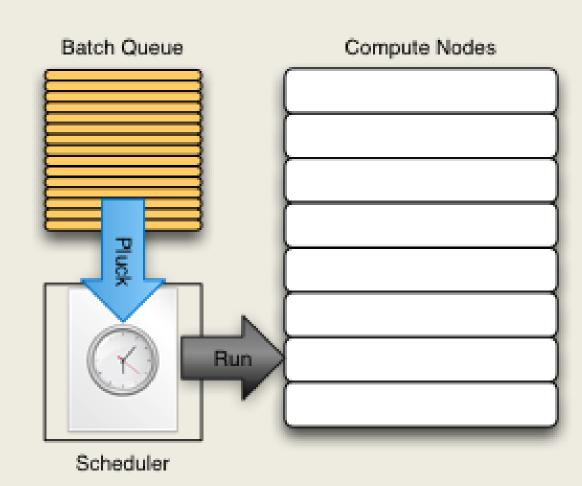
Head Node

Batch Queue Commands

- The batch system is a normal program
- Lets you add and remove jobs from the queue and monitor the queue
- Script/command line driven

The Scheduler component

- Allocates jobs to compute nodes
- Optimizes usage of resources
- "Optimize" can mean many things
- Non-trivial
- Never interact with directly



PBSPro Commands

In order to use the batch system productively, we need to know how to perform three actions:

- Add a job to the queue
- Remove a job from the queue
- See where our job is in the queue

| Command | Description |
|----------------------------------|-----------------------------------------------------------------|
| qsub <job-script></job-script> | Submit a job (add to queue) Returns a <job-number></job-number> |
| qdel <job-number></job-number> | Delete job (remove from the queue) |
| qstat <job-number></job-number> | Monitor jobs |
| qalter <job-number></job-number> | Modifies the attributes of the job or jobs |

Module Package

In order to use the available modules more productively, we need to know how to perform four actions:

- Add a job to the queue
- Load or unload modules for use
- List all available modules

| Command | Description |
|--------------------------|----------------------------------------------------------------|
| module avail | Will list all available module files in the current MODULEPATH |
| module load/unload | Will load/unload a modulefile into the shell environment |
| module list | List all loaded modules |
| module show [modulefile] | Will show information about the modulefile |

Monitoring the queue

| Command | Description |
|------------------------------------|---------------------------------------|
| qstat -a | List all jobs in the queue |
| qstat -u <username></username> | List all jobs of a particular user |
| qstat -f <job-number></job-number> | Show detailed information about a job |

Exercise 1: Monitoring the queue with qstat

Add a job to the queue

- To add a job to the queue, we write a job script.
- The job script is a simply a script.
- It has some special comments that pass information to PBSPro.
- When we want to queue the job, we pass its filename as a parameter to qsub, e.g.
 - qsub <job-script>
- The batch queuing system will return a number that uniquely identifies the job.

A sample PBS job script

NOTE: This script will not NCI facilities

```
#!/bin/bash
# Request resources
# * 10 minutes wall time to run
#PBS -1 walltime=00:10:00
# * 1 node, 1 processor
#PBS -1 nodes=1:ppn=1
# * 100 megabytes physical memory allocated to job
\#PBS -1 mem=100mb
# Specify a project code (for accounting)
#PBS -P a40
cd $PBS O WORKDIR
# Specify the job to be done
date
sleep 10
date
```

You've got mail!

```
# Set email address
#PBS -M fred@intersect.org.au
# Send an email when jobs
# begins (b), gets aborted (a)
# and ends (e)
#PBS -m abe
```

Exercise 2: Submitting a sample job

Useful Environment Variables

These are available in the context of your job script.

| Command | Description |
|---------------|-------------------------------------------------|
| PBS_O_WORKDIR | The directory the job was submitted from |
| PBS_JOBID | The job number given when the job was submitted |

Job limits on Orange

- 200 hours of walltime
- 64GB of memory per standard node.
 e.g. 128GB for 2 nodes etc.
- 256GB of **memory** per large memory nodes
- NOTE: If you grab a node with 64GB, you can effectively use about 60GB as the OS uses memory

Priorities of Jobs

In order of importance, jobs are prioritised in this order:

- 1. Resources available to the project
- 2. Walltime
- 3. Number of jobs (fair share)

Best strategy

- Submit jobs constantly/daily
- Have about 10-20 jobs in the machine
- Be realistic with walltime
- Don't ask for resources you don't need!

NCI Facilities

| Disk Type | Disk Usage |
|----------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| VAYU | Sun Constellation Cluster with 1492 nodes, each containing 2 quad core Nehalem processors summing up to 11,936 cores. 37TB RAM and 800 TB disk space. Commissioned in 2010. The unit of <u>shared memory parallelism is the node</u>, which comprises dual 8-core processors, i.e., 16 cores. |
| NCI Upcoming System (available in May) | 57,472 cores in the compute nodes; Approximately 160 TBytes of main memory; Infiniband FDR interconnect; and Approximately 10 PBytes of usable fast file system (for short-term scratch space). Will be commissioned in its entirety in early 2013. |

NCI Facilities

| VAYU | New NCI Facilities |
|-----------------------------------|---------------------------------|
| 66% of Nodes have 32Gb (2Gb/core) | 96.5% of Nodes have 24Gb/node |
| 33% of Nodes have 64Gb (4Gb/core) | 3.2% of Nodes have 48Gb/node |
| 2% of Nodes have 128Gb (8Gb/core) | 0.3% of Nodes have 96Gb/node |

Software on NCI

| Area | Software |
|--------------------------|---------------------------------------------------|
| Computational Chemistry | ABINIT, Amber, CPMD*, GULP*, NAMD*, Molpro etc. |
| Bioinformatics | AbySS, BEAST, BIOPERL, Cufflinks, MAW, etc. |
| Math Libraries | ARPACK, BLACS, Boost, FFTW, GSL, MKL, Tao |
| Statistics & Maths Env's | Maple*, Mathematica*, MatLab*, Octave*, R, Stata* |

- Asterisked items indicates that discussion with NCI facility staff is required before use (Licensing issues)
- http://nf.nci.org.au/facilities/software/index.php

Orange Physical Disks -

Which of the 3 disks to use and when?

| Disk Type | Disk Type | Disk Usage |
|------------------|--------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Panasas | 59Tb | Parallel global file system All nodes see the Panasas disk Very fast for large files Very slow for small files System director blade creates metadata for each file |
| SGI | 50Tb | An NFS mounted file system Uses old technology, therefore very robust Scales nicely for clusters up to 100 Nodes (very good for Orange) |
| Local Scratch | 200Tb | Exist in each node No network is necessary making these the fastest disk If you have a lot of I/O, you should copy your data to here and work here |

Disk Partitions - Orange

| /home | |
|----------------|-------------------------------|
| Mounted under: | /home/username |
| Disk Type | SGI Disks |
| Size: | 60GB default |
| Backed up: | Yes |
| Speed: | Intermediate disk (SGI Disks) |
| Life time: | Permanent |

Disk Partitions - Orange

| /projects/project-name | | |
|------------------------|--------------------------------------------------------|--|
| Mounted under: | /projects/project-name | |
| Disk Type | Panasas Disk | |
| Size: | no default size | |
| Backed up: | Yes | |
| Speed: | High speed | |
| Life time: | Till end of the running year - merit allocation period | |

 There will also be some "repository space" for large datasets, such as bioinformatics databases

Disk Partitions - Orange

| /data2 | |
|----------------|---------------------|
| Mounted under: | /data2 on each node |
| Disk Type | xx |
| Size: | Limit of disk - 2TB |
| Backed up: | No |
| Speed: | Fastest |
| Life time: | Job duration |

Warning: This partition is shared among users, so can be "filled up" (with other jobs) while your job is running!

Disk Partitions

You can find out more about the partitions on the HPC machine using the **df** command.

| Command | Description |
|---------|------------------------------------------------------------------|
| df -h | Show disk free space for all partitions in human readable format |

You can find out more about current disk usage, using the **du** command.

| Command | Description |
|----------|---------------------------------------------------------------|
| du -hs . | Show disk usage of current directory in human readable format |

Quotas

- There is no quota on scratch disks for performance reasons.
- The quota on /projects/project-name depends on your allocation.
- 60 GB soft limit for /home.
- 80 GB hard limit for /home (30 days).

More Info on NCI & Orange

- Read more about Orange and NCI Facilities
 - http://www.intersect.org.au/hpc-news
 - http://www.intersect.org.au/orange
 - http://www.intersect.org.au/nci_next
 - http://www.intersect.org.au/orangehandbook

Resource Allocation Round

- Merit-based system by which Intersect members can gain access to our HPC facilities
- Applications reviewed by HPC staff (for <u>technical</u> <u>complexity</u> and track record) and the Intersect Resource Allocation Committee (for <u>research merit</u>)
- Applications to Intersect's HPC systems will be made through NCI's forms in October each year
- Applications must be <u>made by Academic Staff</u> at an Intersect member institutions. PhD students can make use of the facilities, the lead CI must be an academic staff member.
- Questions to: hpc_support@intersect.org.au

Register with NCI (step 1)

Register **a new Id with NCI**:

http://nf.nci.org.au/accounts/forms/user_registration.php

This will provide your details to NCI

Register with NCI (step 2)

Apply for **a project from NCI**:

https://nf.nci.org.au/accounts/projects_new/ APP_form.php

This will provide link your Id to your Project

Project Registration Form

- Pick <u>INTERSECT under partner/scheme</u> on the first page of the project registration form or else you won't get access to Orange
- If you're unsure about which machine to get access to, email hpc_support@intersect.org.au who can advise you
- You can add accounts to an existing project also!

Conclusion

- In this course we have covered the basics of the Unix command line, transferring data, and the specifics of our HPC machine
- As different machines have different PBS systems, scripts that work on Octane may not work on NCI facilities
- Please complete our survey!
- Any questions?