

Intermediate HPC

Introduction to Unix for HPC

What is HPC?

- HPC, or high-performance computing, refers to the application of supercomputers or clusters of computers to computational problems that typically arise through scientific inquiry
- 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

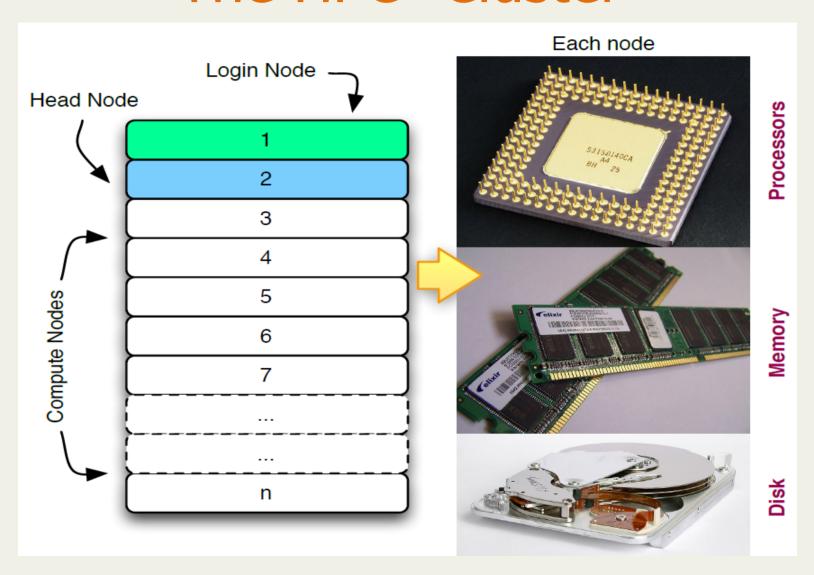
HPC machines

System	Memory Architecture	Cores	Nodes	Memory
Octane (training machine)	Distributed	48	3	48GB
Orange	Distributed	1,600	100	8ТВ
NCI – (Raijin)	Distributed	57,472	3592	158TB

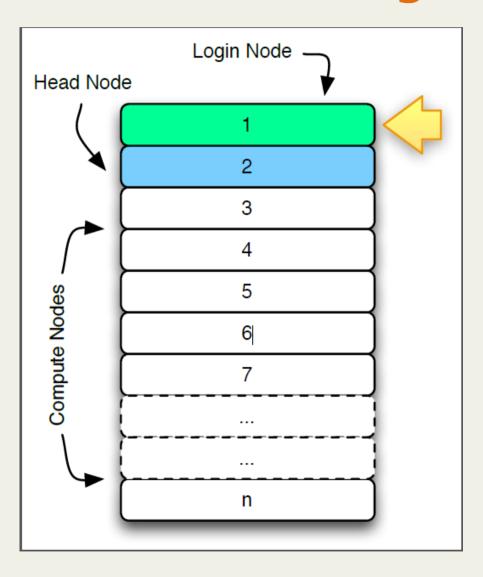
The typical HPC workflow

- In HPC we talk about **jobs**, these are simply commands we wish to run and requests for resources (e.g. compute time, disk space, memory requirements, setup of s/w env's etc.)
- Generally time consuming & resource intensive.
- Jobs are typically run non-interactively,
- Can be run interactively for testing purposes
- We add our jobs to a queue.
- When machines have free resources jobs run
- Once jobs are complete, we can inspect their output.

The HPC "Cluster"



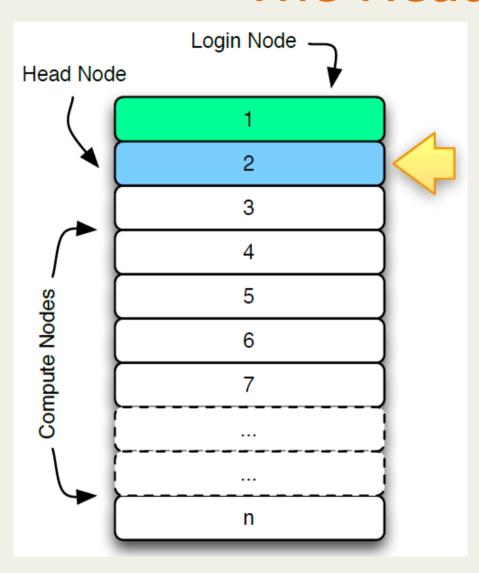
The Login Node



Login Node

- Interactive programs
- SSH sessions
- Testing
- Compiling

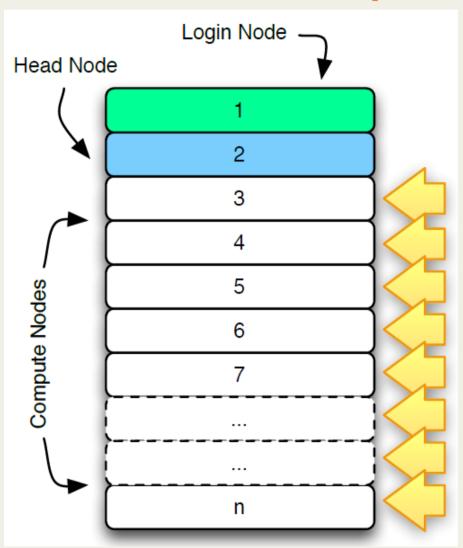
The Head Node



Head Node

Queuing jobs

Compute Nodes



- These nodes run your jobs
- Managed by the scheduler
- Typically you won't 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)
- 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

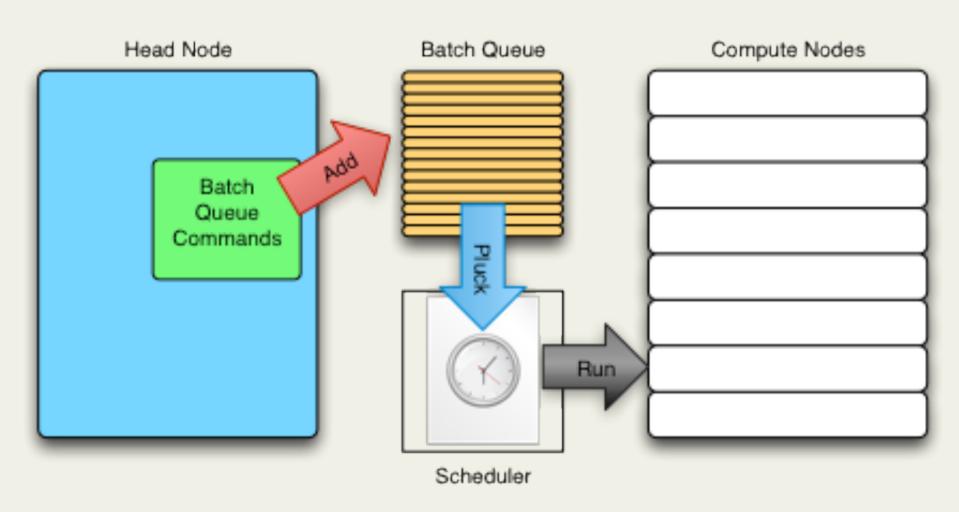
PBS Pro

PBS Pro is the batch system used on all three systems available through your Intersect membership:

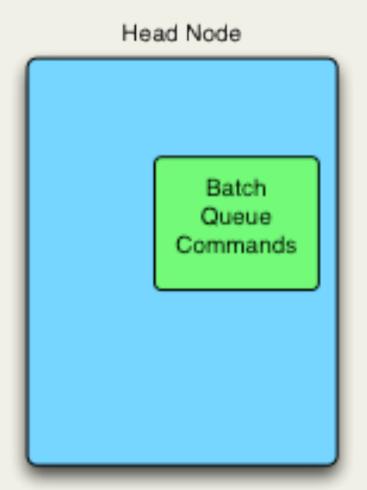
- Octane (training machine)
- Orange
- Raijin(NCI)

Batch System	PBS PRO
Machines using	Orange, Octane, Raijin
Code Base	PBS Professional
Licence	Altair Licence

The Batch Queuing System



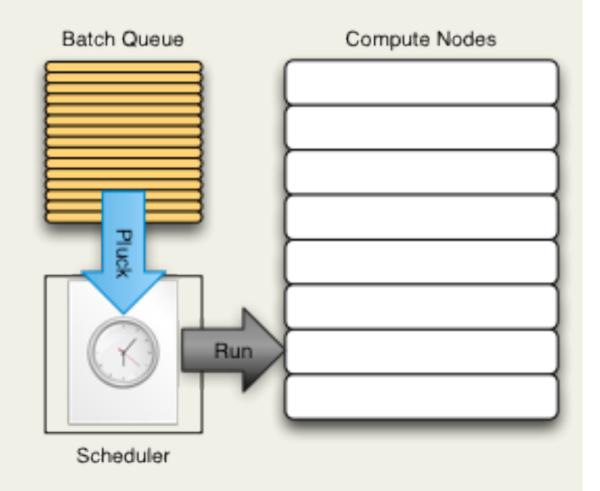
Batch Queuing component



- 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



PBS Pro 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

Exercise 1

Monitoring the queue with qstat

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



All about Modules

What do Modules do?

- Set up the environment for a software package
- Adds paths to executables to \$PATH
- May change other shell variables, and/or load other modules.
- Allow you to have different versions of the same software package, e.g. load the Intel compilers v23 with module load intel/12, or if you require an older version, you can load it with module load intel/9

All about Modules (cont.)

Things to know about modules

- No <u>default module loaded when you login</u>. You can change this by adding lines to your .bashrc file (BASH users) or similar for other Shells
- Modules exist for many packages but <u>not for all!</u>
- Some modules exclude each other, e.g.
 - You can't load Intel compilers v8 & v9. You can only load one.
 - You can only load 1 MPI (e.g. MPT or Intel or OpenMPI)

Module Commands

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 e.g. module load mpt/2.06
module list	List all loaded modules
module show [modulefile]	Will show information about the modulefile

Add a job to the queue

- To add a job to the queue, we write a job script. A job script is a simply a script.
- The # symbol signifies a comment for the Shell
- It has some special comments that pass info to PBS Pro.
- #PBS is a keyword for PBS and specifies that this line is for PBS. The Shell will ignore it
- 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.

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

A sample PBS job script for Octane

Note the "-1" below is a lowercase letter "L" not a one "1"!

```
#!/bin/bash
# Request resources
# * 10 minutes wall time to run
#PBS -1 walltime=00:10:00
# * 1 node, 1 processor
#PBS -1 select=1:ncpus=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 nobody@intersect.org.au
# Send an email when jobs
# begins (b), gets aborted (a)
# and ends (e)
#PBS -m abe
```

A sample PBS job script for Orange

A small change is required to the job script in order to make it work with the way resources are accounted for on Orange:

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

Exercise 2

Create a script and submit a very simple job

Command	Description
#PBS	#PBS is a keyword for PBS and specifies that this line is for PBS. The Shell will ignore it
#	Signifies a comment for the Shell, e.g. # Next line will create a job
qsub	Submit a job to PBS
qstat	List jobs in the queue
<pre>cat <filename></filename></pre>	Print a file to the terminal (catenate)
less <filename></filename>	Like cat, but less at a time
nano <filename></filename>	Will open file <filename> in a text file editor</filename>
Sample PBS Script	http://www.intersect.org.au/orange-handbook



Exercise 3

Create another script and submit a more realistic sample job

Command	Description
#PBS	#PBS is a keyword for PBS and specifies that this line is for PBS. The Shell will ignore it
#	Signifies a comment for the Shell, e.g. # Next line will create a job
qsub	Submit a job to PBS
qstat	List jobs in the queue
nano <filename></filename>	Will open file <filename> in a text file editor</filename>
<pre>module load <module_name></module_name></pre>	Loads a software module on the HPC machine



Job limits on Orange

- 250 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

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

NCI Facilities

Disk Type	Disk Usage
NCI Raijin	 2 sockets with 8-core CPU's = 16 cores 57,472 cores in the compute nodes Approximately 160 TBytes of main memory; Infiniband FDR (fourteen data rate) interconnect Approximately 42 PBytes of usable fast file system (for short-term scratch space).

NCI Facilities

Raijin

96.5% of Nodes have 24GB/node

3.2% of Nodes have 48GB/node

0.3% of Nodes have 96GB/node

plus a small number of nodes with 1TB/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

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Orange Physical Disks

Which of the 3 disks to use and when?

Туре	Disk Type	Disk Type	Disk Usage
1	Panasas	59Tb	 Parallel global file system All nodes see the Panasas disks directly attached to each node Very fast for large files Can be slow if you copy MANY small files System director blade creates metadata for each file
2	SGI	50Tb	 An NFS mounted file system All nodes see the SGI disks can only be seen via NFS backbone Uses old technology, therefore very robust Scales nicely for clusters up to 100 Nodes (very good for Orange)

Orange Disks (cont.)

Which of the 3 disks to use and when?

Туре	Disk Type	Disk Type	Disk Usage
3	Local	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 One 2Tb disk in each compute node Is only accessible within that node as it's not attached to any network, therefore cannot be accessed by another node NOTE: The lifetime of files on this disk is only for the duration of the runtime of the job – The user must copy back results. If not, the next job will erase any files

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	Scratch Disks
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
- Sample PBS Script & Info on Orange
 - 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

To Apply

Please follow the instructions at:

http://intersect.org.au/time/merit

Briefly this involves:

- Registering for a user account
- Completing personal and career profiles
- Registering a new project
- Completing an application for resources for the project

Applications continued

- 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
 - the specifics of our HPC machine
- As different machines have different PBS systems, scripts that work on Orange may need to be adapted for Raijin (NCI) and vice-versa
- Sample answers to all exercises in this course <u>are available on GitHub</u>

Thanks for attending!

- Please complete our <u>course survey</u> at:
 - http://svy.mk/18c8dHa

Any **further questions**, contact us at

- training@intersect.org.au
- Find out about <u>upcoming courses</u> by signing up to our mailing list
 - http://bit.ly/1aZvRqw

