Introduction to Milton

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Research Computing

Goals for today

- Part 1: Learning about supercomputers and research computing at WEHI.
- Part 2: Logging on and exploring the Linux Environment.
- Part 3: Learning about software modules and the batch job submission system.
- Part 4: Submitting test jobs.
- Part 5: PBS Command Summaries
- Part 6: WEHI Filesystems

Slide Respository

- These slides are based on material developed by Lev Lafayette from Research Platform Services at the University of Melbourne.
- A copy of these slides and sample code is available at:
 https://github.com/WEHI-ResearchComputing/MiltonIntro
- Send me an email: thomas.e@wehi.edu.au

Research Computing at WEHI

- Part of ITS which is part of professional services
- Operate our high performance computer (HPC) Milton
- Provide support and guidance to researchers
- Data management

Who	Role
Andrew Isaac	Head of Research Computing
Jakub Szarlat	eResearch
Miffy Edwards	Program delivery manager, data management
Stephenie Gibbs	eResearch
Miguel Esteva	eResearch
Junyao Deng	eResearch
Evan Thomas	Research Computing Scientist

Getting help I

- More documentation is available on Catalyst: Catalyst ->
 Research -> Research Computing
 - Documentation website coming soon
- Help is available at: helpdesk@wehi.edu.au.
- Other training is also conducted by:
 - Research Platforms:
 https://research.unimelb.edu.au/infrastructure/research-platform-services
 - Melbourne Bioinformatics:
 https://www.melbournebioinformatics.org.au/

Getting help II

- Join the slack team: https://milton-wehi.slack.com/signup
 - This is user to user support. ITS do not (officially) monitor it.
- Join the MUG mailing list. Send a request to

research.computing@wehi.edu.au

- i ditti. i icipacsk
 - Read the Message of the Day when you login!
 - If you have
 - problems with submitting a job
 - need a new application or extension to an existing application installed
 - submissions are generating unexpected errors etc.,
 - Just have a question
 - Email helpdesk: helpdesk@wehi.edu.au!
 - Do not email individuals; we need consolidated records. Please be informative about the error or issue.
 - The system(s) you were using, e.g. unix500, batch system, etc.
 - Provide details such as modules loaded, etc

Copy and paste errors messages and commands when

Part 1: Supercomputers and HPC

- "Supercomputer" means any single computer system that has exceptional processing power for its time.
- One popular metric (LINPACK) is the number of floating point operations per second (FLOPS) such a system can carry out (http://top500.org). HPC Challenge and HPCG are broader, more interesting metrics.
- High Performance Computer (HPC) is any computer system whose architecture allows for above average performance.
- High Throughput Computing (HTC) is an architecture for maximum job completion; capability vs capacity computing.

Part 1: Clusters

- Clustered computing is when two or more computers serve a single resource. This improves performance and provides redundancy in case of failure system. Typically commodity systems with a highspeed local network.
- Research computing refers to the concepts and software applications used by the scientific community to aid research. It does not necessarily equate with high performance computing, or the use of clusters. It is whatever researchers use and do. Not issues of producibility and environments.

Part 1: Parallel Computing

- With a cluster architecture, applications can be more easily parallelised across them.
 - Data parallel: running the same task in parallel over different inputs, sometimes referred to as embarrassingly parallel.
 - Task parallel: running a single task in parallel with communication. e.g. sequence alignment; molecular dynamics.
- Software needs to specially written to be task parallel.
- Within node vs between node parallelism.
- WEHI systems are designed with data parallelism and within node parallelism in mind.

Part 1: HPC Cluster and Parallel Processing Components

- A chassis or rack, containing multiple computer system units, interconnect, and providing power.
- Computer system units or nodes, containing memory, local disk, and sockets.
- Sockets contain a processor
 - which has several cores
 - each core processes two logical streams of instructions (hyperthreads)
- In day to day use the term Central Processing Unit, *CPU*, nearly always refers to a hyperthread.

Part 1: A rack



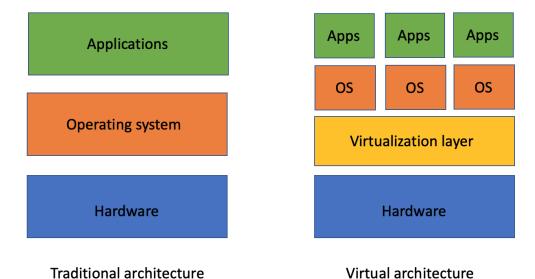
Part 1: Cloud computing

- Infrastructure as a Service (laaS)
 - Compute, storage and network
 - Located in managed datacentres
 - Available on demand
 - Pay by the minute
- Commercial Clouds
 - Amazon Web Services (AWS)
 - Google Cloud Platform
 - Microsoft Azure
- Research Clouds
 - National eResearch Collaboration Tools and Resources project (Nectar)
- Private clouds
 - WEHI Research cloud Milton!

Part 1: Virtualization

- Modern data centres present resources in virtual machines (VMs)
- Software running in a VM sees CPUs, memory, disk, network as though they are real
- A physical node may run serveral virtual machines
 - Node and machine are used interchangeably
- This has a lot of advantages:
 - different applications and users can be isolated from each other
 - Multiple operating systems can run in the same hardware
 - Machines can be easily moved between different hardware to provide the best resource utilisation
- Milton uses VMWare data centre software suit to provide virtualization.

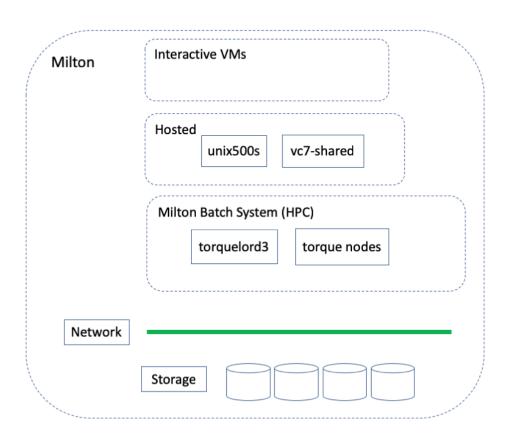
Part 1: Virtualization



Part 1: Milton Hardware

- In 2016 WEHI purchased an *on premise* High Performance Computer called Milton.
- nearly 2500 cores (2 hyperthreads per core)
 - A mix of Skylake and Broadwell Intel Xenon processors
 - Up to 64 cores in a single physical node
- 40TB RAM
 - configurations from 128GB to 1.5TB (can be reconfigured if required)
- A mix of 40GbE and 10GbE network connnections (gigabits per second)
- 20 nVidia P100 GPUs

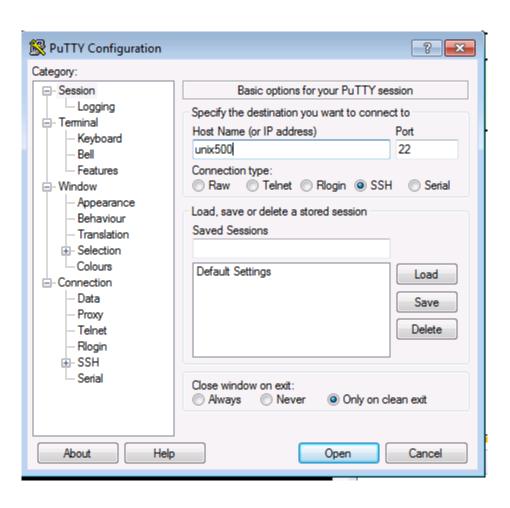
Part 1: Milton overview



Part 1: Logging In

- To log on to Milton you need a WEHI username that's is!
- Mac OS 10.x.
 - Open a terminal
 - o ssh unix500
 - ssh <username>@unix500
 - Type your password
- For MS-Windows users, the free PuTTY client is recommended (http://putty.org).
- You can login from multiple terminal windows!
- Note: Logging into other systems may be more complicated.
 - e.g. ssh eat@spartan.hpc.unimelb.edu.au
 - o <username>@<fully qualified system name>

Part 1: Logging in from Windows



Part 1: Logging in from Windows

```
vc7hpc-347.hpc.wehi.edu.au - PuTTY
                                                                   - - X
🚜 login as: thomas.e
thomas.e@unix500's password:
```

Part 1: SSH Keys

- Consider using an _ssh/config file and using passwordless SSH by creating a keypair and adding to your _ssh/authorized_keys file on Milton or other target system.
- SSH Keys will make your life easier. Follow the instructions here: https://dashboard.hpc.unimelb.edu.au/faq/#how-do-setup-passwordless-ssh-login

Part 2: It's a Linux Command Line World



- The command line interface (CLI) provides a great deal more power and is very resource efficient.
- Linux scales and does so with stability and efficiency.
- Critical software such as the Message Parsing Interface (MPI) and nearly all scientific programs are built for Linux.
- The operating system and many applications are provided as "free and open source" which are better placed to improve, optimize and maintain.

Part 2: File System Hierarchy

- Files are organised in a hierarchy of directories (or folders)
- When a user logs in on a Linux or other UNIX-like system on the command line, they start in their *home* directory (/home/<username>).
 - WEHI user names are usually lastname.i, e.g. thomas.e
- "Everything in the UNIX system is a file" (Kernighan & Pike, 1984, 41).
- See https://swcarpentry.github.io/shell-novice/fig/standardfilesystem-hierarchy.svg

Part 2: Exploring The Environment

Command	Explanation	
whoami	"Who Am I?; prints the effective user id	
pwd	"Print working directory"	
ls	"List" directory listing	

Part 2: Command Options - the ls command

• Linux commands often come with options expressed as:

```
<command> --<option[s]>
```

Options can be expressed as full words or abbreviated characters.

Command	Explanation
ls -lart	long, all, reverse time
ls —lah	long, all, human readable size

Part 2: Getting command help

Linux commands come with "man" (manual) pages, which provide a terse description of the meaning and options available to a command. A verbose alternative to man is info.

Command	Explanation
<command/> help	Command help summmary (usually)
man <command/>	Display the manual entry for the command
<pre>info <command/></pre>	A verbose description of the command
whatis <command/>	A terse description of the command

Command line editing

- up and down arrow keys navigate through your command line history
 - history command
- left and right arrrow keys move the text on the command line

Key stroke	Explanation
up arrow	previous command
down arrow	next command
ctrl-a	beggining of the line
ctrl-e	end of the line
ctrl-u	delete to the beginning of the line
ctrl-k	delete to the end of the line

Tab to completion

- Hit <tab> and the current text will complete if there is a unique completion
- Hit <tab> twice to get a list of possible completions
- Try it: ls -l
 /stornext/HPCScratch/evan/training/Lesson_1.md

Part 2: the less command

- used for conviently viewing large amounts of text
- try less /stornext/HPCScratch/evan/training/Lesson_1.md

Key stroke	Explanation
<space bar=""></space>	page down
u	page up
/ <text></text>	find forwards
? <text></text>	find backwards
/,?	repeat find
g	goto to top
G	goto to bottom
q	quit
h	help

Part 2: Pipes

Linux also have very useful 'pipes' and redirect commands. To pipe one command through another use the '|' symbol.

Command	Explanation	
who -u less	"Who" shows who is logged on and how long they've been idle.	
ps afux	"ps" provides a list of current processes. Check	
less	man ps	

Part 2: Redirects

To redirect output use the '>' symbol. To redirect input (for example, to feed data to a command) use the '<'. Concatenation is achieved through the use of '>>' symbol.

Command	Explanation
w > list.txt	'w' is a combination of who, uptime and ps -a, redirected
w >> list.txt	Same command, concatenated

Part 2: Files and Text Editing I

- Linux filenames can be constructed of any characters except the
 forward slash, which is for directory navigation. However it is best
 to avoid punctuation marks, non-printing characters (e.g., spaces).
 It is *much* better to use underscores or CamelCase instead of
 spaces, newlines etc (including in job names).
- Linux is case-sensitive with its filenames (e.g., list.txt, LIST.txt lisT.txT are different).

Part 2: Navigating the file system

Command	Explanation
cd	change directory
pwd	present working directory

- Files are located via paths a series of directory names seperated
 by /
- Paths can be
 - absolute the start with a '/'
 - relative to the working directory without a leading '/'
 - means the current directory
 - means up one directory
- Demonstrate on the terminal

Part 2: Files and Text Editing II

- Files do not usually require a program association suffix, although you may find this convenient (a C compiler like files to have .c in their suffix, for example).
- The type of file can be determined with the file command. The type returned will usually be text, executable binaries, archives, or a catch-all "data" file.
 - file
 /stornext/HPCScratch/evan/training/Lesson_1.md
 file
 /stornext/HPCScratch/evan/training/random.bam
- There are three text editors usually available on Linux systems on the command-line. These are nano (1989, as pico) and vim (or vi), and or emacs (both 1976). See https://www.vimgolf.com/
 - nano is probably the easiest to start with.

Part 2: Copying Files to a Local System

To get a copy of the files from an external source to your home directory, you will probably want to use wget, or git, or scp.

Command	Explanation
wget URL	Non-interactive download of files over http, https, ftp etc.
git clone URL	Clone a repository into a new directory.

Part 2: Copying Files Within a Local System

To copy a file from within a system use the cp command. Common options include -r to copy an entire directory

Command	Explanation
cp source destination	Copy a file from source to destination
<pre>cp -r source destination</pre>	Recursive copy (i.e., a directory)
<pre>cp -a source destination</pre>	Recursive copy as archive (permissions, links)

Part 2: Copying

```
Try
cp list.txt list2.txt
cp -r /stornext/HPCScratch/evan/training .
```

Part 2: Copying Files Between Systems

To copy files to between systems desktop use SCP (secure copy protocol) or SFTP (secure file transfer protocol), combining the ssh and cp functionality. The cp options can also be used. The source or destination address should also require a remote shell login.

Command	Explanation
<pre>scp source.address:/path/</pre>	Copies files across
<pre>destination.address:/path/</pre>	a network

Part 2: Synchronising Files and Directories

- The rsync utility provides a fast way to keep two collections of files "in sync" by tracking changes.
- The source or destination address should also require a remote shell login.

```
For example; rsync -avz --update lev@spartan.hpc.unimelb.edu.au:files/workfiles .
```

Part 2: Synchronising Files and Directories II

Command	Explanation
rsync source destination	General rsync command
rsync -avze ssh	With ssh
<pre>username@remotemachine:/path/to/source .</pre>	encryption

Part 2: Synchronising Files and Directories III

- The rsync -avz command ensures that it is in archive mode (recursive, copies symlinks, preserves permissions), is verbose, and compresses on transmission.
- The —update option restricts the copy only to files that are newer than the destination.
- Note that rsync is "trailing slash sensitive". A trailing / on a source means "copy the contents of this directory". Without a trailing slash it means "copy the directory".

Part 2: Synchronising Files and Directories IV

Rsync can be used in a synchronise mode with the --delete flag.
 Consider this with the -n , or --dry-run options first!

Command	Explanation
`rsync -avzupdate source/ username@remotemachine:/path/to/destination	Synchronise, keep older files
`rsync -avzdelete source/ username@remotemachine:/path/to/destination	Synchronise, absolutely

Part 2: Creating Directories, Copying and Moving Files

- Directories can be created with the mkdir command (e.g., mkdir braf).
- Files can be copied with the cp command
 - cd change to your home directory
 - cp /stornext/HPCScratch/evan/training/gattaca.txt
 copy a file to you home directory
 - cp gattaca.txt gattaca2.txt make another copy of the file
- Files can be moved or *renamed* with the mv command (e.g., mv gattaca2.txt braf)

Part 2: File Differences

- File differences can be determined by timestamp (e.g., ls -l gattaca.txt braf/gattaca2.txt)
- nano gattaca.txt and make a change
- Content differences can be determined by the diff command (e.g., diff gattaca.txt braf/gattaca2.txt)
- For a side-by-side representation use the command sdiff instead.
- The command comm can compare two files, lines by line (e.g., comm gattaca.txt braf/gattaca2.txt).
- cmp will simply report if there is a difference

Part 2: Searches and Wildcards

- Simple file searches
 - o ls braf/*.txt
 - ls /stornext/HPCScratch/evan/training/*.bam
- To search for files in an entire directory tree use the find command
 - find . -name *.txt'
- To search within files, use the grep command (e.g., grep -i
 NEV braf/* or grep -l)
- The most common wildcard is *, but there is also ? (single character).
- There are also range searches (e.g., [a-z] any character between a and z, inclusive)

Part 2: Deletions

- Files can be deleted with the rm command (e.g., rm gattaca.txt)
- Empty directories can be deleted with the rmdir command (e.g., rmdir braf)
- Directories, files, subdirectories etc can be delted with rm -rf
 <<dir>>
- BE VERY CAREFUL
 - ESPECIALLY WITH WILDCARDS!
 - THERE IS NO UNDO OR TRASH!
 - THERE MAY BE NO BACKUP!
- Consider the difference between rm matlab * to rm matlab*.

Part 3: A Dynamic Software Environment

- Software on Milton Linux systems (and many other HPCs) is provided by the module system
- Manages diverse software requirements in complex multi-user systems.
- Manages dependencies
- Allows multiple versions of the same program

Part 3: Module Commands I

Command	Explanation
module help	List of switches, commands and arguments for modules
module avail	Lists all the modules which are available to be loaded.
<pre>module avail <string></string></pre>	Lists all the modules starting with <string></string>
<pre>module avail 2>&1 grep <string></string></pre>	Lists all the modules containing <string></string>
<pre>module display <modulefile></modulefile></pre>	Display paths etc for modulefile

Part 3: Module Commands I

Command	Explanation
<pre>module load <modulefile></modulefile></pre>	Loads paths etc to user's environment
<pre>module unload <modulefile></modulefile></pre>	Unloads paths etc from user's environment.
module list	lists all the modules currently loaded.

Part 3: Portable Batch System I

- The Portable Batch System (or simply PBS) performs job scheduling by assigning unattended background tasks expressed as batch jobs, among the available resources.
- Originally developed by MRJ Technology Solutions under contract to NASA in the early 1990s. Released as an open-source product as OpenPBS. Forked by Adaptive Computing as TORQUE (Terascale Open-source Resource and QUEue Manager). Many of the original engineering team now part of Altair Engineering who have their own commercial version, PBSPro.
- A batch system typically consists of a resource manager (e.g., TORQUE) and a job scheduler (e.g., Maui, Moab)
- Our batch system is called Torquelord3 (TL3)

Part 3: Slurm Workload Manager

 Slurm, used at UoM on Spartan and many sites, was developed as a collaborative effort primarily by Lawrence Livermore National Laboratory, SchedMD, Linux NetworX, Hewlett-Packard, and Groupe Bull as a Free Software resource manager. As of November 2015, TOP500 list of most powerful computers in the world indicates that Slurm is the workload manager on six of the top ten systems. Slurm's design is very modular with about 100 optional plugins.

Part 3: Job Submission Principles

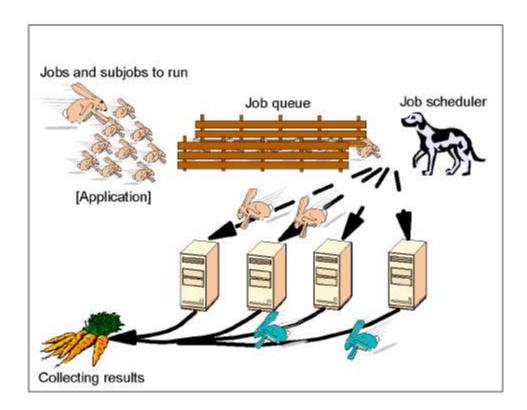
- The steps for job submission are (a) setup and launch., (b) monitor., and (c) retrieve results and analyse.
- Jobs are launched from a submit node with resource requests and, when the job scheduler decides, run on compute nodes.
 - most Milton nodes can act as submit nodes
 - o c.f. *login* node
- Job scripts are simply resource requests (understood by scheduler), a batch of commands (understood by shell) with output to files.

Part 3: Fair Share

- A cluster is a shared environment thus a a resource requesting system. Policies ensure that everyone has a "fair share" to the resources (e.g., user processor limits).
- TL3 treats all users and all resources equally.
- No single user can have mode than 75% of total processors
- Heavy users over the previous 7 hours will have lower priority compared to lighter users
- Precise details of fair share are likely to change over time.

Part 3: Backfilling

- Many schedulers and resource managers use a backfilling algorithm to improve system utilisation and maximise job throughout.
- When more resource intensive jobs are running it is possible that gaps ends up in the resource allocation. To fill these gaps a best effort is made for low-resource jobs to slot into these spaces.
- For example, on an 8-core node, an 8 core job is running, a 4 core job is launched, then an 8 core job, then another 4 core job. The two 4 core jobs will run before the second 8 core job.



From the IBM 'Red Book' on Job Submission.

Part 3: What to run and what not to run on unix500

- unix500 is a shared resource
- Primarily for
 - test and development
 - script and file preparation
- Some heavy processing is OK
 - o 8 CPUs
 - 24 GB memory
 - 24 hours elapsed time
- The batch system will be more stable and more performant

Part 3: Queues

- Jobs are submitted to queues
 - o different resource types, e.g. GPU nodes
 - different service levels, usually coupled to resource constraints
- Milton queues are:

Queue	Purpose
submit	Standard nodes, suitable for most jobs. Default if not queue is specified
submit_2xp100	Nodes with GPUs
skylake	Nodes with Intel Skylake CPUs

Part 3: Basic steps

- Develop a shell script, a text file of commands, to perform the work
 - load modules and other environmental setup
 - run commands
- Submit job with appropriate resources
- Monitor your jobs
- Check job output and expected output files.

Part 3: A script

```
#!/bin/bash
cd <my-work-dir>
module load myapp
myapp data
```

Part 3: Resources

 The system needs to know what resources your jobs needs so it can schedule your job into an appropriate node and reserves the resources you have requested

Resource	Explanation
nodes	Number of nodes
ppn	processors (i.e. CPUs) per node
mem	amount of memory
walltime	maximum elapsed time

Part 3: Submit a job - command line resources

```
qsub -l nodes=1:ppn=2,mem=8gb,walltime=01:00:00 <script>
```

Note: commas and colons

Part 3: Submit a job - script resources

```
#!/bin/bash
#PBS -l nodes=1:ppn=2,mem=8gb,walltime=01:00:00
cd <my-work-dir>
module load myapp
myapp data
```

or

```
#!/bin/bash
#PBS -l nodes=1:ppn=2
#PBS -l mem=8gb
#PBS -l walltime=01:00:00
cd <my-work-dir>
module load myapp
myapp data
```

then

```
qsub <script>
```

Part 3: After you submit your job

- Your job is placed in a queue and job id is assigned
- Your job will have a name, which will be the script name by default
- It may take while before your job runs if the system is busy
- The job will start in your home directory not the directory where you submitted you job!
- Job output will go to the directory where the job was submitted.
- Two output files are created by default
 - < <jobname>.o<jobid> which contains program stdout
 - < <jobname>.e<jobid> which contains program stderr

Part 3: Monitoring your job

To see all the jobs in the queue

qstat

To see just your, somebody else's jobs

qstat -u <userid>

Interpreting job status

Status	Explanation
Q	Queued
R	Running
С	Complete

Part 2: Killing jobs

qdel <jobid>

or

qdel **all**

Part 4: Interactive Jobs

 An interactive job, based on the resource requests made on the command line, puts the user on to a compute node. This is typically done if they user wants to run a large script (and shouldn't do it on the login node), or wants to test or debug a job. The following command would launch one node with two processors for ten minutes.

```
qsub -I -l nodes=1:ppn=2,mem=1gb,walltime=00:10:00
```

Part 4: Worked example

```
mkdir -p ~/braf
cd ~/braf
cp /stornext/HPCScratch/evan/training/S1_R* .
cp /stornext/HPCScratch/evan/training/align.sh .
ls -l
nano align.sh
qsub -l nodes=1:ppn=1,mem=4gb,walltime=01:00:00 align.sh
qstat -u <userid>
ls align.sh.*
cat align.sh.*
```

Part 4: Exercise

Rerun the same job with 6 CPUs

- 1. Work out how to tell bowtie2 to use more CPUs
 - i. module load bowtie2
 - ii. bowtie2 --help
- 2. Modify the script
- 3. Submit with the appropriate invocation of qsub

Part 5: PBS commands: qsub

- man qsub
- google (but there are variants)

Option	explanation
-1	resources (required)
-q <queue></queue>	queue
-N <name></name>	jobname
-m <options></options>	email options
-M <email></email>	email addresses
-j eo	combine output streams
-e <path>, -o <path></path></path>	locations for output streams

Part 5: PBS environment variables

Variable	explanation
PBS_0_H0ST	execution host
PBS_0_WORKDIR	directory where qsub ran
PBS_JOBID	jobid
PBS_JOBNAME	jobname

e.g. cd \$PBS_0_WORKDIR

Part 5: PBS commands: qstat

- man qstat
- google (but there are variants)
- most qstat commands can be limited to a jobid or user

Option	explanation
-u <userid></userid>	only for that user
-n , -n1	show nodes
-f	full report
-q , -Q	show queues
-a	alternative format
-x	XML format
-e , -o	locations for output streams

Part 6: WEHI filesystems

Home areas

What does "home area" mean?

- Linux:
 - The directory you start in when you login
 - location of special files
 - \circ \$H0ME and \sim
 - These will be different on different systems

Part 6: WEHI filesystems

Home areas

- Mac:
 - The folder named with your userid on FS_ALL
 - You need to manually "connect to server ..." from the finder
- Windows
 - W: drive should be mapped automatically

Part 6: WEHI Home areas

- In most milton nodes, including unix500, the home area is
 - /home/<userid>
 - also visible as /stornext/HPCScratch/home/<userid>
- Desktop or main home area is

```
/wehisan/home/allstaff/<s>/<surname.i>/
```

- This is also the home area for the old unix3xx and unix4xx home areas
- To copy data from your desktop
 - copy to your home with the finder/explorer
 - copy from "main" home area to your "milton" home area
- Milton home areas are on HPCScratch and are not backed up

Part 6: WEHI lab areas

- All data are owned by the lab, then the division then the institute
- Most labs will have a lab area for their data.
- All valuable data should on a lab area these are backed up
- Use the Milton home area for temporary data

Slide Respository

- These slides are based on material developed by Lev Lafayette from Research Platform Services at the University of Melbourne.
- A copy of these slides and sample code is available at:

https://github.com/WEHI-ResearchComputing/MiltonIntro

