

An Introduction to High Performance Computing on the Minerva Cluster

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Plan of the Course

Part 1: Basics

Part 2: HPC Facilities

Part 2: Using HPC

10:00 WELCOME

11:30-11:45 Break

13:00-14:00 LUNCH

15:30-15:45 Break

16:30 CLOSE

Welcome

- ► Stuart Rankin Research Computing User Services Team
- ▶ Paul Browne Research Computing Platforms Team
- ► Course files can be downloaded from: www.csd3.cam.ac.uk
- ▶ Please ask questions and let us know if you need assistance.



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Basics: Outline

Why Buy a Big Computer?

Inside a Modern Computer

How to Build a Supercomputer

Programming a Multiprocessor Machine

Summary



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Basics: Compute Intensive Problems

- ▶ Distribute the work for a single problem across multiple CPUs to reduce the execution time as far as possible.
- ▶ Program workload must be *parallelised*:

Parallel programs split into copies (processes or threads).

Each process/thread performs a part of the work on its own CPU, concurrently with the others.

A well-parallelised program will fully exercise as many CPUs as there are processes/threads.

- ► The CPUs typically need to exchange information rapidly, requiring specialized communication hardware.
- ► Many use cases from Physics, Chemistry, Engineering, Astronomy, Biology...
- ► The traditional domain of HPC and the Supercomputer.

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Basics: Why Buy a Big Computer?

What types of big problem might require a "Big Computer"?

Compute Intensive: A single problem requiring a large amount of computation.

Memory Intensive: A single problem requiring a large amount of memory.

Data Intensive: A single problem operating on a large amount of data.

High Throughput: Many unrelated problems to be executed in bulk.



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Basics: Scaling & Amdahl's Law

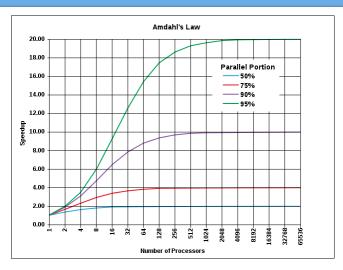
- ▶ Using more CPUs is not necessarily faster.
- ► Typically parallel codes have a scaling limit.
- ► Partly due to the system overhead of managing more copies, but also to more basic constraints;
- ► Amdahl's Law (idealized):

$$S(N) = \frac{1}{\left(1 - p + \frac{p}{N}\right)}$$

where

S(N) is the fraction by which the program has sped up relative to N=1 p is the fraction of the program which can be parallelized N is the number of CPUs.

Basics: Amdahl's Law



http://en.wikipedia.org/wiki/File:AmdahlsLaw.svg



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Basics: Data Intensive Problems

- ▶ Distribute the data for a single problem across multiple CPUs to reduce the overall execution time.
- ▶ The *same* work may be done on each data segment.
- ► Rapid movement of data to and from disk is more important than inter-CPU communication.
- ▶ Big Data problems of great current interest -
- ► Hadoop/MapReduce
- ► Life Sciences (genomics) and elsewhere.

The Bottom Line

- ► Parallelisation requires effort:
 - ► There are libraries to help (e.g. OpenMP, MPI).
 - ► First optimise performance on one CPU, then make *p* as large as possible.
- ► The scaling limit: eventually using more CPUs becomes detrimental instead of helpful.



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Basics: High Throughput

- ▶ Distribute independent, multiple problems across multiple CPUs to reduce the overall execution time.
- ▶ Workload is trivially (or *embarrassingly*) parallel:
 - * Workload breaks up naturally into independent pieces.
 - * Each piece is performed by a separate process/thread on a separate CPU (concurrently).
 - * Little or no inter-CPU communication.
- ► Emphasis is on throughput over a period, rather than on performance on a single problem.
- ► Compute intensive capable ⇒ high throughput capable
- ► Compute intensive capable ≠ high throughput capable





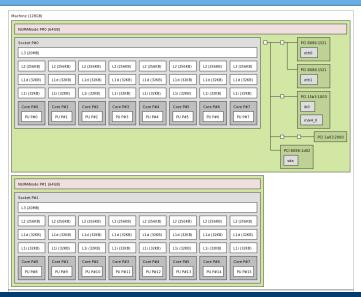
Basics: Memory Intensive Problems

- ► Require aggregation of large memory into a single system image (i.e. a single computer running Linux).
- ► Technically more challenging to build machines (very fast, low latency interconnection between all CPUs and all memory).
- Coding/porting easier (memory appears seamless).
- ▶ Optimisation harder (memory is actually highly nonuniform).
- ► Historically, the arena of large SGI systems.
- ▶ Nowadays, similar complexity inside single commodity servers.



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Basics: Inside a Modern Computer





- ► Today's commodity servers already aggregate both CPUs and memory to make a single system image in a single box.
- ► Even small computers now have multiple CPU cores per socket ⇒ each socket is a Symmetric Multi-Processor (SMP).
- ► Larger computers have multiple sockets (each with local memory):
 all CPU cores (unequally) share the node memory
 ⇒ the node is a shared memory multiprocessor
 with Non-Uniform Memory Architecture (NUMA)
 but users still see a single computer (single system image).



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Basics: How to Build a Supercomputer

- ► A supercomputer aggregates contemporary CPUs and memory to obtain increased computing power.
- ► Usually today these are clusters.
- 1. Take some (multicore) CPUs plus some memory.
 - ► Could be an off-the-shelf server, or something more special.
 - ► A NUMA, shared memory, multiprocessor building block: a node.



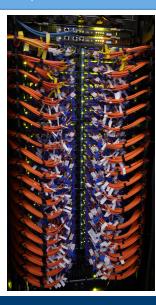


Basics: How to Build a Supercomputer

2. Connect the nodes with one or more networks. E.g.

Gbit Ethernet: 100 MB/sec Omni-Path: 10 GB/sec

Faster network is for inter-CPU communication across nodes. Slower network is for management and provisioning. Storage may use either.





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Basics: Programming a Multiprocessor Machine

- ► Non-parallel (serial) code
 - $\ast\,$ For a single node as for a workstation.
 - * Typically run as many copies per node as CPUs, assuming node memory is sufficent.
 - * Replicate across multiple nodes.
- ► Parallel code
 - * Shared memory methods within a node. E.g. pthreads, OpenMP.
 - * Distributed memory methods spanning multiple nodes. Message Passing Interface (MPI).

Basics: How to Build a Supercomputer

- 3. Logically bind the nodes
 - Clusters consist of distinct nodes (i.e. separate Linux computers) on common private network(s) and controlled centrally.
 - * Private networks allow CPUs in different nodes to communicate.
 - * Clusters are distributed memory machines: Each process/thread sees only its local node's CPUs and memory (without help).
 - * Each process/thread must fit within a single node's memory.
 - More expensive machines logically bind nodes into a single system i.e. CPUs and memory.
 - * E.g. SGI UV.
 - Private networks allow CPUs to see CPUs and memory in other nodes.
 - * These are shared memory machines.
 - * Logically a single system 1 big node but very non-uniform.
 - * A single process can span the entire system.



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Basics: Summary

- ► Why have a supercomputer?
 - ▶ Big single problems, many problems, Big Data.
- ▶ Most current supercomputers are clusters of separate nodes.
- ► Each node has multiple CPUs and non-uniform shared memory.
- ► Parallel code uses shared memory (pthreads/OpenMP) within a node, distributed memory (MPI) spanning multiple nodes.
- ► Non-parallel code uses the memory of one node, but may be copied across many.





Part II: HPC Facilities

Minerva

► Each compute node:

- * 2x16 cores, Intel Skylake 2.6 GHz32 CPUs
- * 192 GB RAM6 GB per CPU
- * 100 Gb/sec Omni-Path10 GB/sec (for MPI and storage)
- ► 16 compute nodes
- ► 1 high-memory login/head node (minerva-login1.npl.co.uk), 768 GB RAM.
- ▶ 1 GPU node, 2 × NVIDIA P100 GPU, 192 GB RAM.

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CSD3 - University of Cambridge

- ► Cambridge Service for Data Driven Discovery
- ► Peta4 Intel CPU cluster
- ► Wilkes2 NVIDIA GPU cluster
- ► Hadoop-based data analytic platform
- ► Burst buffer
- ► Some NPL users through CORE.

Peta4-Skylake

- ► Each compute node:
 - * 2x16 cores, Intel Skylake 2.6 GHz32 CPUs
 - * 192 GB or 384 GB RAM6 GB or 12 GB per CPU
 - * 100 Gb/sec Omni-Path10 GB/sec (for MPI and storage)
- ► 768 compute nodes
- ▶ 8 login nodes (login-cpu.hpc.cam.ac.uk)



Wilkes2-GPU

- ► CPUs are general purpose
- ► Some types of parallel workload fit vector processing well:
 - ► Single Instruction, Multiple Data (SIMD)
 - ► Think pixels on a screen
 - ► GPUs specialise in this type of work
 - ► Also competitor many-core architectures such as the Intel Phi



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Peta4-KNL (Intel Phi)

- ► Each compute node:
 - * 64 cores, Intel Phi 7210256 CPUs
 - * 96 GB RAM96 GB RAM
 - * 100 Gb/sec Omni-Path10 GB/sec (for MPI and storage)
- ► 342 compute nodes
- ► Shared login nodes with Peta4-Skylake

- ► Each compute node:
 - * 4 × NVIDIA P100 GPU4 GPUs
 - * 1x12 cores, Intel Broadwell 2.2 GHz12 CPUs
 - * 96 GB RAM96 GB RAM
 - * 100 Gb/sec (4X EDR) Infiniband.10 GB/sec (for MPI and storage)
- ▶ 90 compute nodes.
- ▶ 8 login nodes (login-gpu.hpc.cam.ac.uk).



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Cluster Storage

- ▶ Minerva uses NFS to share user directories between all cluster nodes (150 TB).
- ► CSD3 uses the Lustre cluster filesystem:
 - * Very scalable, high bandwidth.
 - * Multiple RAID6 back-end disk volumes.
 - * Multiple object storage servers.
 - * Single metadata server.
 - * Tape-backed HSM on newest filesystems.
 - * 12 GB/sec overall read or write.
 - $* \ \, \mathsf{Prefers} \,\, \mathsf{big} \,\, \mathsf{read/writes} \,\, \mathsf{over} \,\, \mathsf{small}.$





- ► Please contact the NPL IT Service Desk:
 - ► itservicedesk@npl.co.uk
 - ▶ Room F12-CS1
 - ► Tel. 6000
- ▶ Second line support is provided by the Cambridge RCS.



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Using HPC: Connecting

► SSH secure protocol only.

Supports login, file transfer, remote desktop...

Part III: Using HPC

Connecting: Windows Clients

- putty, pscp, psftp http://www.chiark.greenend.org.uk/ sgtatham/putty/download.html
- ► WinSCP http://winscp.net/eng/download.php
- ► TurboVNC (remote desktop, 3D optional) http://sourceforge.net/projects/turbovnc/files/
- ► Cygwin (provides an application environment similar to Linux) http://cygwin.com/install.html Includes X server for displaying graphical applications running remotely.
- ► MobaXterm http://mobaxterm.mobatek.net/





Connecting: Linux/MacOSX/UNIX Clients

- ► ssh, scp, sftp, rsync Installed (or installable).
- ► TurboVNC (remote desktop, 3D optional) http://sourceforge.net/projects/turbovnc/files/
- ► On MacOSX, install XQuartz to display remote graphical applications.

http://xquartz.macosforge.org/landing/



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Connecting: First time login

► The first connection to a particular hostname produces the following:

The authenticity of host 'minerva-login1.npl.co.uk (139.143.201.10)' can't be established.

ECDSA key fingerprint is SHA256:k/eB+LjcAfQW56XCzK9QptT0wVWF7j3a/CPxPRd7+lE.
ECDSA key fingerprint is MD5:18:9a:97:e2:87:4c:07:60:cb:43:46:f2:bb:d8:3d:01.

Are you sure you want to continue connecting (yes/no)? yes Warning: Permanently added 'minerva-login1.npl.co.uk (139.143.201.10)' (ECDSA) to the list of known hosts.

- ▶ One should always check the fingerprint before typing "yes".
- ► Graphical SSH clients *should* ask a similar question.
- ▶ Designed to detect fraudulent servers.

Connecting: Login

► From graphical clients:

Host: minerva-login1.npl.co.uk

Username: npl\abc123 (your NPL AD account name)

► From Linux/MacOSX/UNIX (or Cygwin):

ssh -Y **npl**\\abc12@minerva-login1.npl.co.uk

Note the double backslash — this is because UNIX command interpreters treat \ as special.



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Connecting: First time login

- ► Exercise 1 Log into your Minerva account.
- ► Exercise 2 Simple command line operations.





Connecting: File Transfer

- ▶ With graphical clients, connect as before and drag and drop.
- ► From Linux/MacOSX/UNIX (or Cygwin):

```
rsync -av old_directory/
```

npl\\abc12@minerva-login1.npl.co.uk:hpc-work/new_directory

copies contents of old_directory to ~/hpc-work/new_directory.

```
rsync -av old_directory
```

copies old_directory (and contents) to

~/hpc-work/new_directory/old_directory.

- * Rerun to update or resume after interruption.
- * All transfers are checksummed.
- * For transfers in the opposite direction, place the remote machine as the first argument.
- ► Exercise 3 File transfer.



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Connecting: Remote Desktop

Remote desktop already running:

[sjr20@login-a-1 ~]\$ vncserver -list

TigerVNC server sessions:

X DISPLAY # PROCESS ID:99 130655

▶ Kill it:

[sjr20@login-a-1 ~]\$ vncserver -kill :99
Killing Xvnc process ID 130655

- ► Typically you only need one remote desktop.
- ► Keeps running until killed, or the node reboots.



Connecting: Remote Desktop

► First time starting a remote desktop:

```
[sjr20@login-a-1 ~]$ vncserver
```

You will require a password to access your desktops.

Password:

Verify:

Would you like to enter a view-only password (y/n)? n

New 'login-a-1:99 (sjr20)' desktop is login-a-1:99

Starting applications specified in /home/sjr20/.vnc/xstartup Log file is /home/sjr20/.vnc/login-a-1:99.log

- ▶ NB Choose a different password for VNC.
- ► The VNC password protects your desktop from other users.
- ► Remember the unique display number (99 here) of your desktop.



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Connecting: Remote Desktop

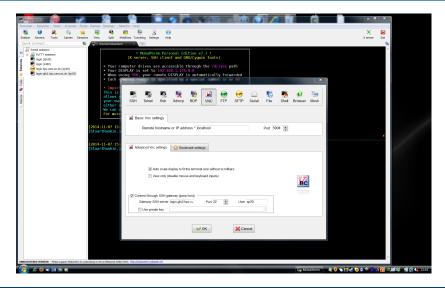
► To connect to the desktop from Linux:

vncviewer -via npl\\abc12@minerva-login1.npl.ad.local localhost:99

- ► The display number 99 will be different in general and unique to each desktop.
- ► You will be asked firstly for your AD login password, and secondly for your VNC password.
- ▶ Press F8 to bring up the control panel.
- ► Exercise 4 Remote desktop (from Windows)



Connecting: Remote Desktop (MobaXterm)





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User Environment: Filesystems

- ► /home/abc12@npl.ad.local
 - ► 50GB quota.
 - ► Visible equally from all nodes.
 - ► Single storage server.
 - ► Regular backups.
 - ► Not intended for job outputs or large/many input files.
- ~/hpc-work
 - ► Visible equally from all nodes.
 - ► Larger (1TB initial quota).
 - ► Intended for job inputs and outputs.
 - ▶ Not backed up by default.

Using HPC: User Environment

- ► CentOS Linux 7.4 (Red Hat Enterprise Linux 7.4 rebuild)
 - ▶ bash shell
 - ► Gnome or XFCE4 desktop (if you want)
 - ▶ GCC compilers and other development software.
- ▶ But you don't need to know that.



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Filesystems: Quotas

quota

```
[sjr20@login-a-1 ^]$ quota -s
Disk quotas for user sjr20 (uid 1004):
Filesystem space quota limit grace files quota limit grace
10.44.82.252:/hpc-work

OK 1024G 1126G 1 0 0

10.44.82.252:/home
13272K 51200M 56320M 345 0 0
```

- ► Aim to stay below the soft limit (quota).
- ► Once over the soft limit, you have 7 days grace to return below.
- ▶ When the grace period expires, or you reach the hard limit (*limit*), no more data can be written.
- ▶ It is important to rectify an out of quota condition ASAP.





Filesystems: Quotas

► quota

```
[sjr20@login-a-1 ~]$ quota -s
Disk quotas for user sjr20 (uid 1004):
Filesystem space quota limit grace files quota limit grace
10.44.82.252:/hpc-work
0K 1024G 1126G 1 0 0
10.44.82.252:/home
13272K 51200M 56320M 345 0 0
```

- ► Aim to stay below the soft limit (quota).
- ▶ Once over the soft limit, you have 7 days grace to return below.
- ▶ When the grace period expires, or you reach the hard limit (*limit*), no more data can be written.
- ▶ It is important to rectify an out of quota condition ASAP.



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User Environment: Software

- ► Free software accompanying Red Hat Enterprise Linux is (or can be) provided.
- ▶ Other software (free and non-free) is available via modules.
- Proprietary software currently available includes Matlab and COMSOL.
- ▶ New software may be possible to provide on request.
- ► Self-installed software should be properly licensed.
- ► sudo will not work. (You should be worried if it did.)

Filesystems: Permissions

- Be careful and if unsure, please ask support.
 - ► Can lead to accidental destruction of your data or account compromise.
- ▶ Avoid changing the permissions on your home directory.
 - ► Files under /home are particularly security sensitive.
 - ► Easy to break passwordless communication between nodes.



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User Environment: Environment Modules

- ► Modules load or unload additional software packages.
- ► Some are required and automatically loaded on login.
- ► Others are optional extras, or possible replacements for other modules.
- ▶ Beware unloading default modules in ~/.bashrc.
- ► Beware overwriting environment variables such as PATH and LD_LIBRARY_PATH in ~/.bashrc. If necessary append or prepend.





User Environment: Environment Modules

► Currently loaded:

module list
Currently Loaded Modulefiles:

1) dot 2) slurm centos7/global

4) centos7/default-basic

► Available:

module av



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User Environment: Environment Modules

▶ Matlab

module load matlab/r2018a

- ► Invoking matlab in batch mode: matlab -nodisplay -nojvm -nosplash command where the file command.m contains your matlab code.
- ▶ The current site license contains the Parallel Computing Toolbox.

User Environment: Environment Modules

► Whatis:

module whatis openmpi-3.0.0-gcc-4.8.5-n2hvjgm openmpi-3.0.0-gcc-4.8.5-n2hvjgm: The Open MPI Project is an open source...

► Load:

module load openmpi-3.0.0-gcc-4.8.5-n2hvjgm

► Unload:

module unload openmpi-3.0.0-gcc-4.8.5-n2hvjgm



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User Environment: Environment Modules

► Purge:

module purge

► Defaults:

module show centos7/default-basic
module load centos7/default-basic

▶ Run time environment must match compile time environment.



User Environment: Compilers

► GCC

```
gcc -03 -mtune=native code.c -o prog
gfortran -03 -mtune=native code.f90 -o prog
module load openmpi-3.0.0-gcc-4.8.5-n2hvjgm
mpicc -03 -mtune=native mpi_code.c -o mpi_prog
mpif90 -03 -mtune=native mpi_code.f90 -o mpi_prog
```

► Exercise 5: Modules and Compilers



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Using HPC: Job Submission

- ► Compute resources are managed by a scheduler: SLURM/PBS/SGE/LSF/...
- ► Jobs are submitted to the scheduler
 - analogous to submitting jobs to a print queue
 - a file (*submission script*) is copied and queued for processing.

Using HPC: Job Submission





-- --

Using HPC: Job Submission

- $\,\blacktriangleright\,$ Jobs are submitted from the login node
 - not itself managed by the scheduler.
- ▶ Jobs may be either non-interactive (batch) or interactive.
- ▶ Batch jobs run a shell script on the first of a list of allocated nodes.
- ► Interactive jobs provide a command line on the first of a list of allocated nodes.





Using HPC: Job Submission

Job Submission: Using SLURM

- ▶ Jobs may use part or all of one or more nodes
 - the owner can specify —exclusive to force exclusive node access.
- ► Template submission scripts are available under ~/job_templates.



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Job Submission: Show Queue

► Submitted job scripts are copied and stored in a queue:

```
[abc123@login-a-1]$ squeue -u abc123

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)

790299 skylake Test3 abc123 PD 0:00 2 (PriorityResourcesAssocGrpCPUMinsLimit)
790290 skylake Test2 abc123 R 27:56:10 2 cpu-a-[1,10]
```

▶ Prepare a shell script and submit it to SLURM:

[abc123@login-a-1]\$ sbatch slurm_submission_script Submitted batch job 790299



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Job Submission: Monitor Job

► Examine a particular job:

[abc123@login-a-1]\$ scontrol show job=790290





Job Submission: Cancel Job

► Cancel a particular job:

[abc123@login-a-1]\$ scancel 790290



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Job Submission: Accounting Commands

► How many core hours available do I have?

mybalance

User	Usage	1	Account	Usage	1	Account Limit	Available	(hours)
		+			+			
sjr20	3	Ī	SUPPORT-CPU	2,929	1	22,425,600	22,422,671	
eir20	0	1	SIIDDUBT-CDII	0	1	87 600	87 600	

▶ How many core hours does some other project or user have?

gbalance -p SUPPORT-CPU

User	Usage	į	Account	Usage	į	Account Limit	Available	(hours)
		+			+			
pfb29	2,925	П	SUPPORT-CPU	2,929		22,425,600	22,422,671	
sjr20 *	3	I	SUPPORT-CPU	2,929	I	22,425,600	22,422,671	
(Use -u for user.)								

▶ List all jobs charged to a project/user between certain times:

gstatement -	p NPL-GENE	RAL-CPU -u	xyz10 -s "2	2018-04-01-0	00:00:00" -e "2018-04	e "2018-04-30-00:00:00"				
JobID	User	Account	JobName	Partition	End	ExitCode	State	CompHrs		
263	xyz10	support-c+	_interact+	skylake	2018-04-18T19:44:40	0:0	TIMEOUT	1.0		
264	xyz10	support-c+	_interact+	skylake	2018-04-18T19:48:07	0:0	CANCELLED+	0.1		
275	xvz10	support-c+	interact+	skylake	Unknown	0:0	RUNNTNG	0.3		



Job Submission: Scripts

► SLURM

In /job_templates, see examples: slurm_submit.skylake.generic, slurm_submit.skylake.matlab.

```
#!/bin/bash
#! Name of the job:
#SBATCH -J myjob
#! Which project should be charged:
#SBATCH -A NPL-GENERAL-CPU
#! How many whole nodes should be allocated?
#SBATCH --nodes=1
#! How many tasks will there be in total? (<= nodes*32)
#SBATCH --ntasks=116
#! How much wallclock time will be required?
#SBATCH --time=02:00:00
#! Select partition:
#SBATCH -p skylake</pre>
```

- ▶ #SBATCH lines are structured comments
 - correspond to sbatch command line options.
- ► The above job will be given 1 cpu16 cpus on 1 node for 2 hours (by default there is 1 task per node, and 1 cpu per task).



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Job Submission: Single Node Jobs

Serial jobs requiring large memory, or OpenMP codes.

```
#!/bin/bash
...
#SBATCH --nodes=1
#SBATCH --ntasks=1
# Default is 1 task per node
#SBATCH --cpus-per-task=

#SBATCH --mem=3994
# Memory per node in MB - default is pro rata by cpu number
# Increasing --mem or --cpus-per-task implicitly increases the othe
...
export OMP_NUM_THREADS= # For OpenMP across cores
$application $options
```



Job Submission: Single Node Jobs

► Serial jobs requiring large memory, or OpenMP codes.

```
#!/bin/bash
...
#SBATCH --nodes=1
#SBATCH --ntasks=1
# Default is 1 task per node
#SBATCH --cpus-per-task=1
# Default is 1 cpu (core) per task
#SBATCH --mem=3994
# Memory per node in MB - default is pro rata by cpu number
# Increasing --mem or --cpus-per-task implicitly increases the other
...
export CMP_NUM_THREADS= # For OpenMP across cores
$application $options
...
```



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Job Submission: Single Node Jobs

Serial jobs requiring large memory, or OpenMP codes.

```
#!/bin/bash
...
#SBATCH --nodes=1
#SBATCH --ntasks=1
# Default is 1 task per node
#SBATCH --cpus-per-task=8 # Half node

#SBATCH --mem=3994
# Memory per node in MB - default is pro rata by cpu number
# Increasing --mem or --cpus-per-task implicitly increases the other
...
export OMP_NUM_THREADS=8 # For OpenMP across 8 cores
$application $options
...
```

Job Submission: Single Node Jobs

► Serial jobs requiring large memory, or OpenMP codes.

```
#!/bin/bash
...
#SBATCH --nodes=1
#SBATCH --ntasks=1
# Default is 1 task per node
#SBATCH --cpus-per-task=16 # Whole node

#SBATCH --mem=3994
# Memory per node in MB - default is pro rata by cpu number
# Increasing --mem or --cpus-per-task implicitly increases the other
...
export OMP_NUM_THREADS=16 # For OpenMP across 16 cores
$application $options
...
```



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Job Submission: Single Node Jobs

Serial jobs requiring large memory, or OpenMP codes.

```
#!/bin/bash
...
#SBATCH --nodes=1
#SBATCH --ntasks=1
# Default is 1 task per node
#SBATCH --cpus-per-task=16 # Whole node

#SBATCH --mem=3994
# Memory per node in MB - default is pro rata by cpu number
# Increasing --mem or --cpus-per-task implicitly increases the other
...
export OMP_NUM_THREADS=8 # For OpenMP across 8 cores (using all memory)
$application $options
```





Job Submission: MPI Jobs

► Parallel job across multiple nodes.

```
#!/bin/bash
...
#SBATCH --nodes=4
#SBATCH --ntasks=64 # i.e. 16x4 MPI tasks in total.
#SBATCH --cpus-per-task=2
...
mpirun -np 64 $application $options
...
```

- ► SLURM-aware MPI launches remote tasks via SLURM.
- ▶ The template script uses \$SLURM_TASKS_PER_NODE to set PPN.



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Job Submission: Hybrid Jobs

► Parallel jobs using both MPI and OpenMP.

```
#!/bin/bash
...
#SBATCH --nodes=4
#SBATCH --ntasks=32  # i.e. 8x4 MPI tasks in total.
#SBATCH --cpus-per-task=2
...
export OMP_NUM_THREADS=2  # i.e. 2 threads per MPI task.
mpirun -ppn 8 -np 32 $application $options
...
```

► This job uses 64 CPUs (each MPI task splits into 2 OpenMP threads).

Job Submission: MPI Jobs

► Parallel job across multiple nodes.

```
#!/bin/bash
...
#SBATCH --nodes=4
#SBATCH --ntasks=32  # i.e. 8x4 MPI tasks in total.
#SBATCH --cpus-per-task=2
...
mpirun -ppn 8 -np 32 $application $options
...
```

- ► SLURM-aware MPI launches remote tasks via SLURM.
- ► The template script uses \$SLURM_TASKS_PER_NODE to set PPN.



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Job Submission: High Throughput Jobs

- ► Multiple serial jobs across multiple nodes.
- ▶ Use srun to launch tasks (job steps) within a job.

```
#!/bin/bash
...
#SBATCH --nodes=4
...
cd directory_for_job1
srun --exclusive -N 1 -n 1 $application $options_for_job1 > output 2> err &
cd directory_for_job2
srun --exclusive -N 1 -n 1 $application $options_for_job2 > output 2> err &
...
cd directory_for_job64
srun --exclusive -N 1 -n 1 $application $options_for_job64 > output 2> err &
wait
```

Exercise 6 - Submitting Jobs.





Job Submission: Interactive

- ► Compute nodes are accessible via SSH while you have a job running on them.
- ► Alternatively, submit an interactive job:

```
sintr -A NPL-GENERAL-CPU -N1 -n8 -t 2:0:0
```

- ▶ Within the window (screen session):
 - * Launches a shell on the first node (when the job starts).
 - * Graphical applications should display correctly.
 - * Create new shells with ctrl-a c, navigate with ctrl-a n and ctrl-a p.
 - * ssh or srun can be used to start processes on any nodes in the job.
 - * SLURM-aware MPI will do this automatically.



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Job Submission: Array Jobs (ctd)

- ► Updates can be applied to specific array elements using \${SLURM_ARRAY_JOB_ID}_\${SLURM_ARRAY_TASK_ID}
- ► Alternatively operate on the entire array via \${SLURM_ARRAY_JOB_ID}.
- ► Some commands still require the SLURM_JOB_ID (sacct, sreport, sshare, sstat and a few others).
- ► Exercise 7 Array Jobs.

Job Submission: Array Jobs

- http://slurm.schedmd.com/job_array.html
- ▶ Used for submitting and managing large sets of similar jobs.
- ► Each job in the array has the same initial options.
- ► SLURM

[abc123@login-a-1]\$ sbatch --array=1-7:21,3,5,7 -A NPL-GENERAL-CPU submit_script Submitted batch job 791609

```
[abc123@login-a-1]$ squeue -u abc123
         JOBID PARTITION
                        NAME
                                USER ST
                                           TIME NODES NODELIST(REASON)
       791609_1 skylake hpl abc123 R
                                           0:06 1 cpu-a-6
       791609_3 skylake hpl abc123 R
                                                   1 cpu-a-16
                       hpl abc123 R
                                                   1 cpu-a-7
       791609_5 skvlake
                                          0:06
                     hpl
       791609_7 skylake
                              abc123 R
                                          0:06
                                                   1 cpu-a-7
```

791609_1, 791609_3, 791609_5, 791609_7

i.e. \${SLURM_ARRAY_JOB_ID}_\${SLURM_ARRAY_TASK_ID}

SLURM_ARRAY_JOB_ID = SLURM_JOBID for the first element.



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Scheduling

- ► SLURM scheduling is multifactor:
 - ► QoS payer or non-payer?
 - Age how long has the job waited?
 Don't cancel jobs that seem to wait too long.
 - ► Fair Share how much recent usage?

 Payers with little recent usage receive boost (not implemented yet).
 - ▶ sprio -j jobid
- Backfilling
 - ▶ Promote lower priority jobs into gaps left by higher priority jobs.
 - ▶ Demands that the higher priority jobs not be delayed.
 - ▶ Relies on reasonably accurate wall time requests for this to work.
 - ▶ Jobs of default length will not backfill readily.





Wait Times

- ▶ 36 hour job walltimes are permitted.
- ▶ This sets the timescale at busy times (without backfilling).
- ► Use backfilling when possible.
- ► Short (1 hour or less) jobs have higher throughput.



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Job Submission: Scheduling Top Dos & Don'ts

- ▶ Do ...
 - ▶ Give reasonably accurate wall times (allows backfilling).
 - ► Check your balance occasionally (mybalance).
 - ► Test on a small scale first.
 - ► Implement checkpointing if possible (reduces resource wastage).
- ▶ Don't ...
 - ► Request more than you need
 - you will wait longer and use more credits.
 - ► Cancel jobs unnecessarily
 - priority increases over time.



Checkpointing

- ▶ Insurance against failures during long jobs.
- ▶ Restart from checkpoints to work around finite job length.
- ► Application native methods are best. Failing that ...



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