Introduction to High Performance Computing and Orchestra

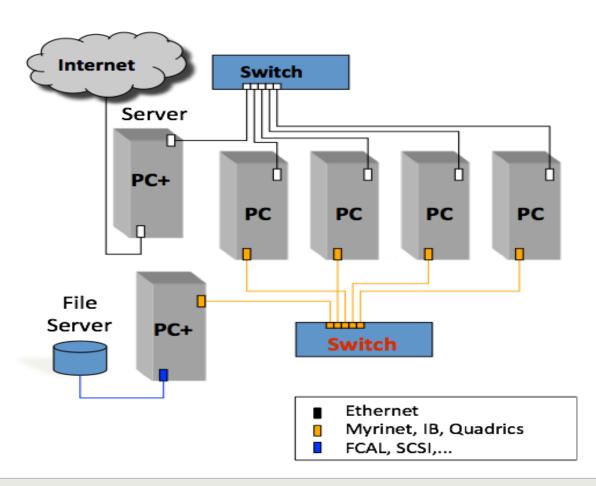
HMS Research Computing Fall 2015

What is Orchestra?

- Wiki page: https://wiki.med.harvard.edu/Orchestra
- Tech spec:
 - Over 500 compute nodes
 - Over 7000 cores
 - 10GigE interconnection
 - Over 40TB RAM
- CentOS 6 Linux
- LSF scheduler
- Total 25+PB storage

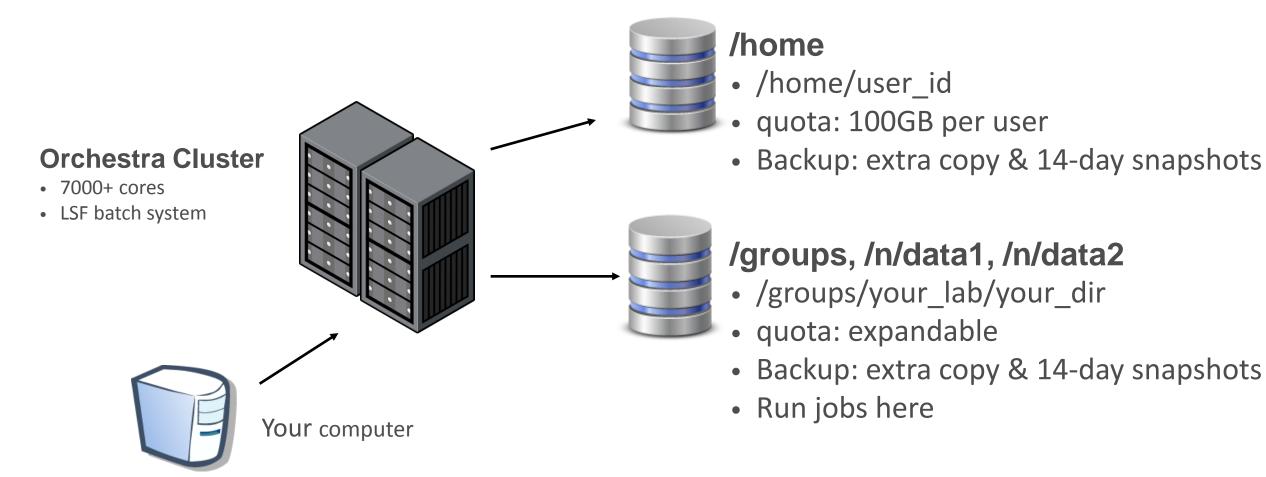


Generic Cluster Architecture



- Login nodes are for data management, job submission, code development, etc.
- Compute nodes are for production runs
- Your storage space is centralized network storage system
- Hundreds users are sharing the resources, so please be nice each other.
- Job scheduler works based on complex algorithms of fair-share, priority management, load balancing, etc.

Orchestra Storage



https://rc.hms.harvard.edu/

Scratch: Coming Soon

Scratch1 (DDN ExaScalar)

- High-performance parallel file system: Lustre
- More than 1PB of disk space
- No quota, no backup, data will be purged
- Ideal for large-scale job running

Scratch 2 (DDN GridScalar)

- High-performance parallel file system: GPFS
- More than 2PB of space
- Quota policy, for long-term storage
- Ideal for large-scale job running
- Available for large-data producing labs



Checking Storage Usage

To check your storage available:

mfk8@loge:~\$ quota

Home directory: you get 100 GB.

Group directories: space varies, can increase.

/groups/groupname

/n/data1

/n/data2



Storage Policies

- /home: 14 day snapshots + 60 day full backup
- /groups, /n/data1, /n/data2: 14 day snapshots + 60 day full backup
- New tape system for long-term storage

Snapshots

- Snapshots are retained for up to 60 days: recover data
- mfk8@clarinet001:~\$ cd .snapshot
- mfk8@clarinet001:~\$ Is

Orchestra_home_daily_2015-10-02-02-00

Orchestra_home_daily_2015-10-01-02-00

- mfk8@clarinet001:~\$ cd Orchestra_home_daily_2015-10-02-02-00
- mfk8@clarinet001:~\$ cp MyRetreivedFile ~



Create an Orchestra Account

- http://rc.hms.harvard.edu/#orchestra
 - Click the red button and fill out the form!
- Your username will be your eCommons ID, with your eCommons password.

LSF: Fair Sharing

- Load Sharing Facility: distributes jobs across Orchestra fairly
- Ensures that no one user or core monopolizes Orchestra
- Users are assigned dynamic priorities
- Queues also have priorities
- Submitting lots of jobs reduces your fairshare priority
- Even if many jobs are pending, your jobs will start quickly provided you have not submitted many jobs

Logging Into Orchestra: Mac

- Open a terminal (search "terminal") ssh –I username orchestra.med.harvard.edu
- For graphics (X11 forwarding) Install Xquartz (google it) and have it running ssh **-X** -I username orchestra.med.harvard.edu

Logging Into Orchestra: Windows

- Install PuTTY (google it)
- In box under Host Name (or IP address)
 - orchestra.med.harvard.edu
- For graphics (X11 Forwarding)
 - Install Xming (google it) and have it running
 - In PuTTY, under Connection -> SSH -> X11:
 - Check "Enable X11 Forwarding" THEN log in

Logging Into Orchestra: Linux

Open a terminal (search: "terminal")

ssh –I username orchestra.med.harvard.edu

For graphics (X11 Forwarding)

ssh -X -I username orchestra.med.harvard.edu

Welcome to Orchestra!

 Where are you in Orchestra? See your terminal!

mfk8@mezzanine: ~\$

- You are logged into a "shell login server." Usually "mezzanine" or "loge." These are not meant for heavy lifting!
- You are in your home directory. This is symbolized by the "tilde " (~). This is shorthand for: /home/username
- You are in a bash environment. "\$" means "ready to accept your commands!"



Interactive Sessions

The login servers are not designed to handle intensive processes. Using 40% of a CPU results in a process being terminated. It is sometimes better to use an "interactive session." These last up to 12 hours. Start by entering your first job! This will (usually) log you into a "clarinet!"

mfk8@mezzanine:~\$ bsub -Is -q interactive -n 1 bash

"bsub" is how jobs are submitted to Orchestra LSF

- -Is (capital eye, s) tells Orchestra you want an interactive shell
- -q is the "queue" to enter (interactive)
- -n is the number of cores requested (1) (20 maximum)

"bash" opens the bash session

mfk8@clarinet001:~/\$



Listing a Folder's Contents

 To see the contents of the current folder you are in (~ means "/home/username/"), type list (ls):

mfk8@clarinet001:~\$ Is

- To get the details of a folder's contents, add "-I" mfk8@clarinet001:~\$ Is -I
- You don't have to be in a directory to see its contents

mfk8@clarinet001:~\$ ls /groups/rc-training/introtohpc

Viewing File Contents

- "less" to view file contents
- Navigate up/down, search
- "q" to quit

mfk8@clarinet001:~\$ less /groups/rc-training/introtohpc/testfile.txt

Making a Folder (Directory)

- "mkdir" stands for "make directory."
- Create a new directory for this exercise
- Spaces are discouraged. (Underscores are fine!) Case counts in Linux.

mfk8@clarinet001:~\$ mkdir MyTestDir

Moving Around: Change Directory

- "cd" stands for "change directory"
- 1 period "."means "current directory"
- 2 periods ".." means "the directory above"

mfk8@clarinet001:~\$ cd MyTestDir

Notice how the prompt tells you where you are!

mfk8@clarinet001:~/MyTestDir\$ cd ...

mfk8@clarinet001:~\$

Creating a Simple Text File

- "Nano," "vi", "emacs" are simple command-line editors available.
- To create a new file, type the editor you want, then the name of the new file. To edit an existing file, do the same.

```
mfk8@clarinet001:~$ nano myfile.txt
   This is my new file text.
    (Control-X to save and exit.)
mfk8@clarinet001:~$
mfk8@clarinet001:~$ Is
   myfile.txt
```

Copying Files

 "cp" to copy a file from a destination to a new destination. "cp" "from" "to"

mfk8@clarinet001:~\$ cp ./myfile.txt ./MyTestDir/

 You can copy a file to the current folder or to a new folder with a different name by specifying a different name (rename)

```
mfk8@clarinet001:~$ cp ./myfile.txt ./MyTestDir/myfilecopy.txt
```

```
mfk8@clarinet001:~$ cp ./myfile.txt ./mycopy2.txt
```

mfk8@clarinet001:~\$ cp /groups/rc-training/introtohpc/testfile.txt /home/mfk8/MyTestDir/



Copying Directories

- "cp" "options" "source" "destination"
- Options:
 - -r: recursively (copies all)

mfk8@clarinet001:~\$ cp -r ./MyTestDir ./MyTest2

Moving Data

• "move" "from" "to"

mfk8@clarinet001~:\$ mv ~/MyTestDir/myfilecopy.txt ~

mfk8@clarinet001~:\$ mv ./MyTestDir/ ./MyTestDir2/

Removing Files/Folders

• "rm" to remove a file

mfk8@clarinet001:~\$ rm myfile.txt

"rm –r" to remove a folder recursively

mfk8@clarinet001:~\$ rm -r ./MyTest2

Wildcard * Pattern Matching

- Useful for copying/removing/etc all files matching a certain pattern
- Example Case:

To copy "all" files ending in ".fastq":

\$ cp *.fastq ./NewFastqFolder

Getting Data Onto Orchestra

- Use an FTP client of your choice
- Mac/Windows/Linux: Filezilla (google it)
- Connect to:

orchestra.med.harvard.edu your username and password port 22



Using Software: Environment Modules

- Most "software" on Orchestra is installed as an environment module. Allows for clean, easy loading, including most dependencies, and switching versions.
 - \$ module avail
 - \$ module avail seq/
 - \$ module avail stats/
 - \$ module avail seq/bowtie/

Loading/Unloading Modules

- Loading modules
 - \$ module load seq/bowtie/2.0.6
- Which module version is loaded (if at all)?
 - \$ which bowtie
- See all modules loaded
 - \$ module list
- Unloading modules
 - \$ module unload seq/bowtie/2.0.6
- Dump all modules
 - \$ module purge



Compiling your own software

- Users can compile software in their /home or /groups directories, where they have permission
- Binaries just require "unzipping" (ie tar –zxvf .tgz)
- Common compiling libraries are found as modules:

dev/compiler/gcc-4.8.5

dev/boost/1.57.0

dev/openblas/0.2.14

dev/lapack

Common libraries are found in "utils"

Installing Software: Binary Example

- mfk8@loge:~\$ bsub -ls -q interactive bash
- mfk8@clarinet001:~\$ wget http://path/to/binary/mysoftware.tar.gz
- mfk8@clarinet001:~\$ tar -zxvf mysoftware.tar.gz
- mfk8@clarinet001:~\$ Is mysoftware/bin

Installing Software: Source

- mfk8@loge:~\$ bsub —ls —q interactive bash
- mfk8@clarinet001:~\$ module load dev/compiler/gcc-4.8.5
- mfk8@clarinet001:~\$ wget http://path/to/source/mysoftware.tar.gz
- mfk8@clarinet001:~\$ tar -zxvf mysoftware.tar.gz
- mfk8@clarinet001:~\$ cd mysoftware
- mfk8@clarinet001:~\$ less README
- mfk8@clarinet001:~\$./configure --prefix=/home/mfk8/software
- mfk8@clarinet001:~\$ make
- mfk8@clarinet001:~\$ make install



Python, R, Perl

- Users manage their own package libraries: get the version you want, when you want it!
- Python: virtual environment allows pip installs

https://wiki.med.harvard.edu/Orchestra/PersonalPythonPackages

R: set up personal R library for cran, bioconductor

https://wiki.med.harvard.edu/Orchestra/PersonalRPackages

Perl: local::lib allows cpan, cpanm

https://wiki.med.harvard.edu/Orchestra/PersonalPerlPackages

Shebangs: #!/usr/bin/env python/Rscript/perl



Submitting Jobs

- In an "interactive session", programs can be called directly.
 - mfk8@clarinet001:~\$ bowtie -n 4 hg19 file1_1.fq file1_2.fq
- From the login shell (and also interactive or any compute nodes), a program is submitted to Orchestra via a job (bsub).
 - mfk8@clarinet001:~\$ bsub < mybowtiejob.sh
- Orchestra will notify you when the job is done, or if there is an error.

The "bsub"

mfk8@clarinet001:~\$ bsub -q queue -W hr:min job

- Necessary:
 - -q (queue)
 - -W (runtime in hr:min)

https://rc.hms.harvard.edu/

Shared Queues

- mpi queue if you have an MPI parallel job
- priority queue if you have just one or two jobs to run
- mcore queue if you have multi-core jobs to run.
- short queue if your jobs will take less than 12 hours to run.
- Else: long queue.

Shared Queues Breakdown

Queue Name	Priority	Max Cores	Max Runtime
interactive	12	12	12 hours
priority	14	12	1 month
mcore	12	12	1 month
mpi	12	512	1 month
parallel	10	512	1 month
short	8	12	12 hours
long	6	12	1 month
mini	5	1	10 minutes

Runtime Limit

- -W in hours:minutes
- Runtimes are subject to the maximum time permitted per queue (see table)
- If your job exceeds your runtime, your job will be killed ⊗
- Running many jobs that finish quickly (less than a few minutes) is suboptimal and may result in job suspension, contact RC to learn how to batch jobs

CPU Limit

- Amount of seconds the cluster works on your job (calculated by LSF)
- Ncores * Runlimit (-n * -W)
- Common error:

bsub –q short –W 8:00 tophat –p 8

tophat asks for 8 cores but only 1 requested (no -n), job killed in 1 hour

Reserving Memory

- Most nodes have 90GB memory available over all cores, some have more
- Make a resource request with
 - -R "rusage[mem=8000]" (memory requested in MB)
- Memory multiplies by cores requested, so
 - -n 4 –R "rusage[mem=16000]" reserves 64GB memory
- Asking for more memory may cause jobs to pend longer
- TERM_MEMLIMIT errors indicate that not enough memory was reserved



Multithreading

- A single CPU can execute multiple processes (threads) concurrently
- n indicates how many cores are requested
- Jobs that are overefficient (use more cores than reserved) jeopardize the health of a node
- Reserve the same amount of cores in your job and your bsub!



MPI

- Efficient way to run parallel jobs
- Dedicated queue with same-type compute nodes for optimal performance (800 cores)
- Matlab, Python, Java, R, C++, Fortran have MPI options
- Orchestra implementation: openMPI-1.8.6

Other bsub options

- -n 4 (number of cores requested)
- -R "rusage[mem=8000]" (memory requested in MB)
- -J jobname
- -a openmpi (type of mpi run)
- -Is (interactive shell)
- -e errfile (send errors to file)
- -o outfile (send screen output to file)
- -N (notify when job completes)



Monitoring Jobs

List info about jobs/their status:

```
mfk8$loge:~$ bjobs
```

- -r (running jobs)
- -p (pending jobs)
- -I (command entered, long form)
- List historical job information

```
mfk8$loge:~$ bhist jobid
```

Job Suspensions

- Jobs in the "long" queue can be suspended by LSF temporarily to allow jobs from the "short" queue (less than 12h) to run.
- If a job is suspended for >50% of the –W runlimit cumulatively, it will be automatically moved to the "no_suspend" queue where it can't be suspended anymore.

Terminating Jobs

Terminate a job (jobid given at submission)

```
mfk8$loge:~$ bkill jobid
```

Terminate all jobs (be careful!)

mfk8\$loge:~\$ bkill 0

Shell Scripts: The Basics

#! /bin/sh #always at the top

#program with options

tophat -p 4 -o ./mytophatdir1 hg19 file1_1.fastq file1_2.fastq

tophat -p 4 -o ./mytophatdir2 hg19 file2_1.fastq file2_2.fastq

#save as myshellscript.sh

Calling a Shell Script

mfk8@balcony:~\$ bsub -q mcore -W 12:00 -n 4 -e errfile.txt -o outfile.txt –N **sh** myshellscript.sh

Creating a Complete Shell Script

```
#! /bin/sh
#BSUB -q mcore
#BSUB -W 12:00
#BSUB -o outfile.txt
#BSUB -e errfile.txt
#BSUB -N
#BSUB -n 4
#BSUB -R "rusage[mem=12000]"
tophat -p 4 -o ./mytophatdir1 hg19 file1_1.fastq file1_2.fastq
#save as myshellscript2.sh
```

Calling a Complete Shell Script

mfk8@balcony:~\$ bsub < myshellscript2.sh

File Properties

 "chmod" to change who can read/write/execute files/directories chmod options file/directory

Who? user group others all (u/g/o/a)

What? read write execute (r/w/x)

Do? +/-

- chmod u+x myfile #makes a file executable to owner
- chmod o-rwx myfile #takes away permission from others to read/write execute

Class Exercises

- In your "home" directory, create a file called ".forward" using nano with this piece of information: your preferred email address for Orchestra notifications. Save.
- Copy the .bam file from /groups/rc-training/introtohpc to your "home" directory, load "samtools" module, and submit a job to the "short" queue for "1 hr" that has this command:
 - samtools sort control.bam sorted.bam
- Monitor this job, then kill it
- Delete all class files (except .forward)



For more direction:

- https://wiki.med.harvard.edu/Orchestra/NewUserGuide
- http://rc.hms.harvard.edu
- rchelp@hms.harvard.edu
- Office Hours: Wednesdays 1-3p Gordon Hall 500
- Class survey:

http://hmsrc.me/introhpc2016-survey1