

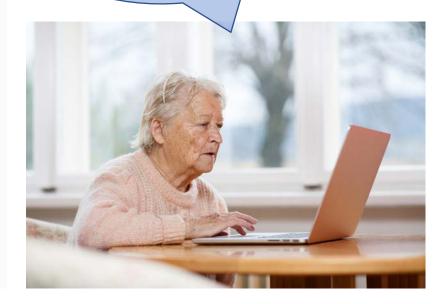
Intro to HPC*

*HPC – high-performance computing

- Why use HPC?
- Anatomy of the HPC cluster
- Basics of working with the Linux shell
- Submitting jobs
- Being a good HPC user



It's still running...



Problem

Computation takes too long

Computation is too big

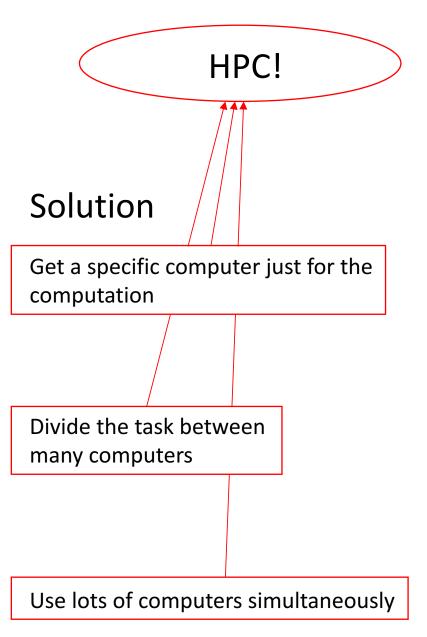
Too many computations

Problem

Computation takes too long

Computation is too big

Too many calculations



Modern instrument for High-Performance Computing is a **cluster**, consisting of lots of connected individual computers (nodes).

Supercomputer is a commonly used nickname.



Laptop



Supercomputer



Laptop



Supercomputer



Laptop

Personal

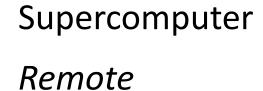


Supercomputer *Shared*



Laptop

Local







Laptop

Interactive



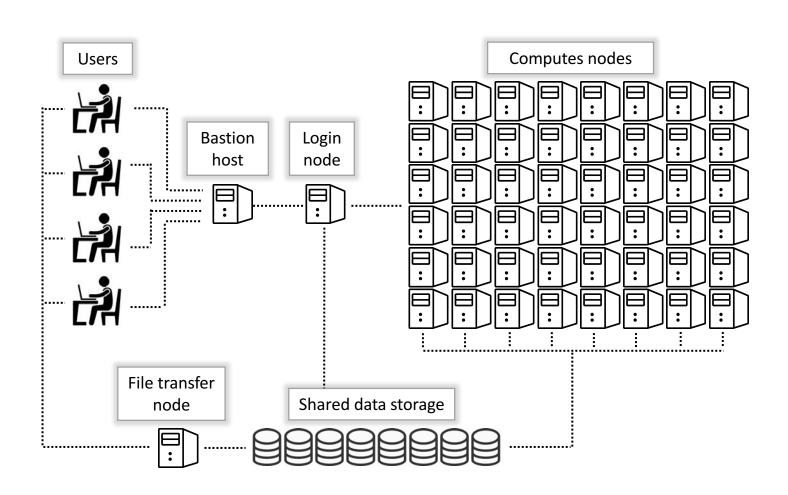
Supercomputer *Batch*



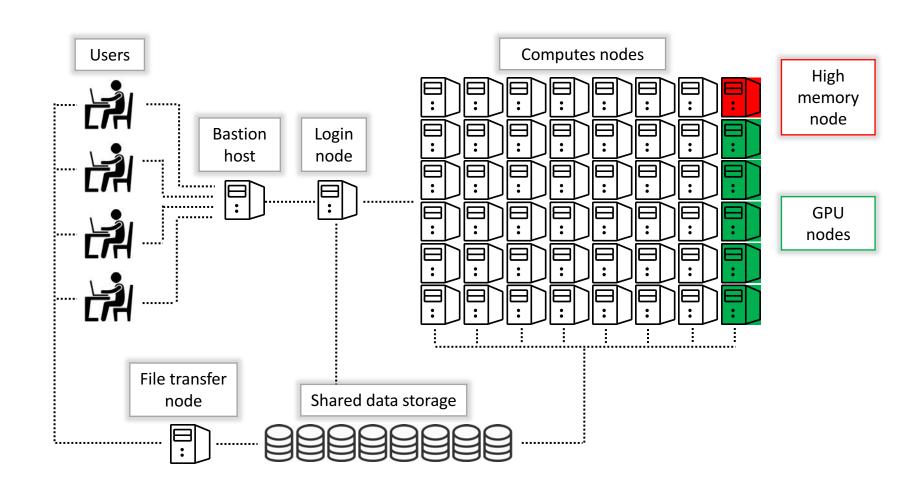
Ocelote



The diagram of the UA HPC cluster

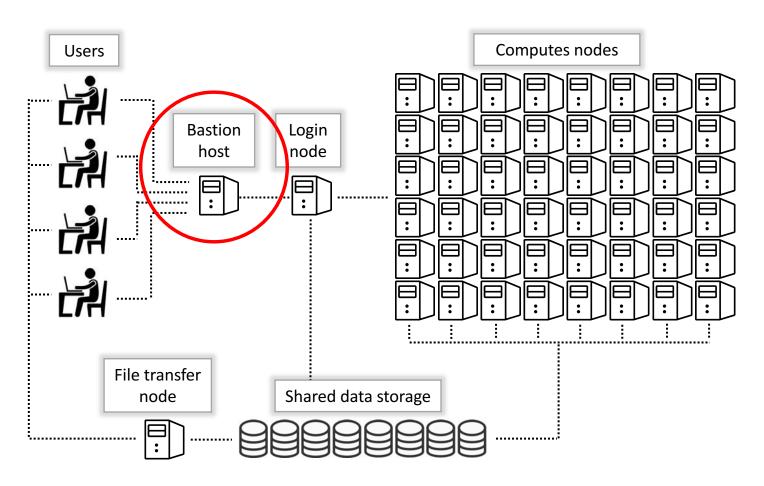


The diagram of the UA HPC cluster



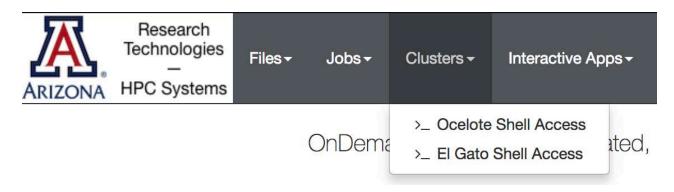
Connecting to Ocelote

ssh hpc.arizona.edu



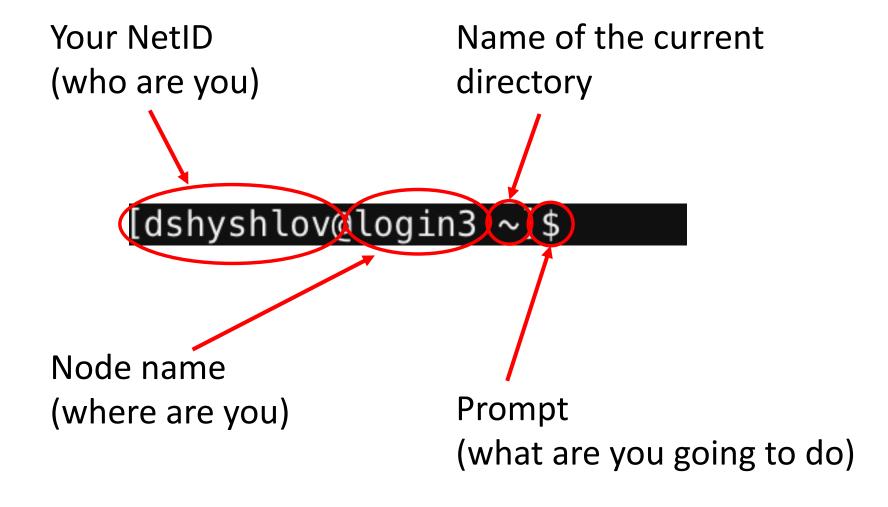
Connecting to Ocelote (web browser)

- Open ood.hpc.arizona.edu in your web browser and login with your NetID and password.
- From the "Clusters" drop-down menu choose which HPC cluster you would like to access:

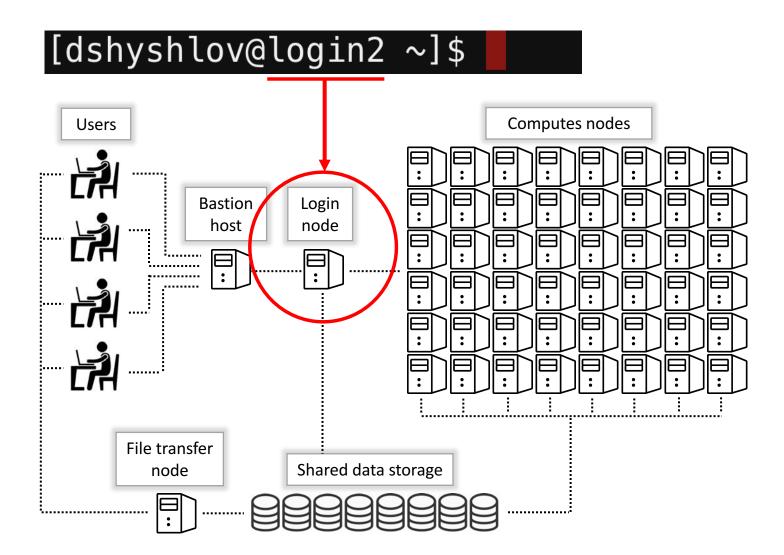


Exercise - connect to Ocelote.

Command line



Login node



Login node

Login node



Compute nodes



Login node

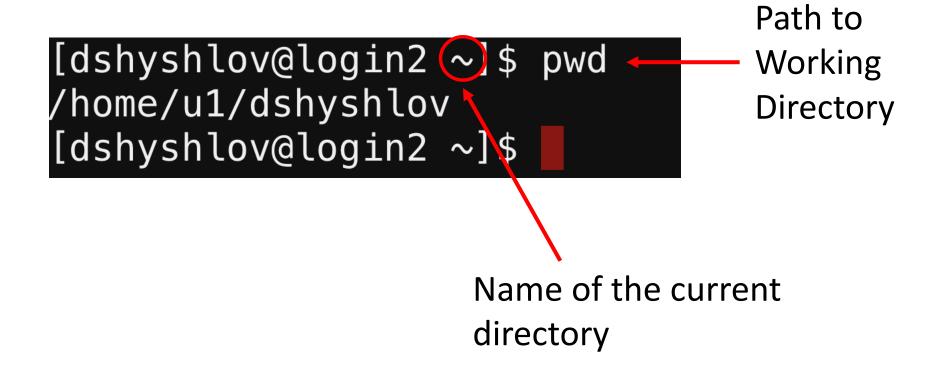
- Login node is a computer intended for users to prepare and manage computations:
 - submit jobs
 - edit files
 - compile codes
 - manage files
 - small-scale testing
- DO NOT run any calculations on the login node



Command

```
[dshyshlov@login2 ~]$ whoami
dshyshlov
[dshyshlov@login2 ~]$
```

Output



~ is a shortcut for your /home directory

List all the files and directories

[dshyshlov@login2 ∼]\$ ls

Make a directory

[dshyshlov@login2 ~]\$ mkdir Intro_to_HPC

List all the files and directories again

[dshyshlov@login2 ~]\$ ls

Change directory

```
[dshyshlov@login2 ~]$ cd Intro_to_HPC
[dshyshlov@login2 Intro_to_HPC]$
```

Go back a level

```
[dshyshlov@login2 Intro_to_HPC]$ cd ..
[dshyshlov@login2 ~]$
```

Change directory using absolute path

```
[dshyshlov@login2 ~]$ cd ~/Intro_to_HPC/
[dshyshlov@login2 Intro_to_HPC]$ ■
```

Copy a file

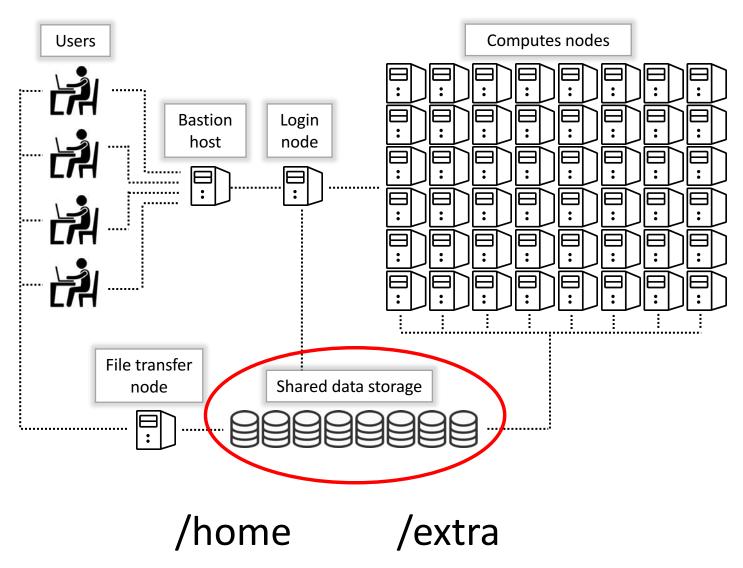
```
$ cp /tmp/first_script.pbs .
```

List all the files and directories again

```
[dshyshlov@login2 ~]$ ls
```

View contents of the file on the screen

```
$ cat first_script.pbs
```



- Every users gets two default storage locations :
 - /home
 - the default home directory
 - 15GB
 - the only backed up storage on UA HPC/extra
 - /extra
 - full path: /extra/NetID
 - 200GB
 - not backed up
 - has file count limit 600 files/GB

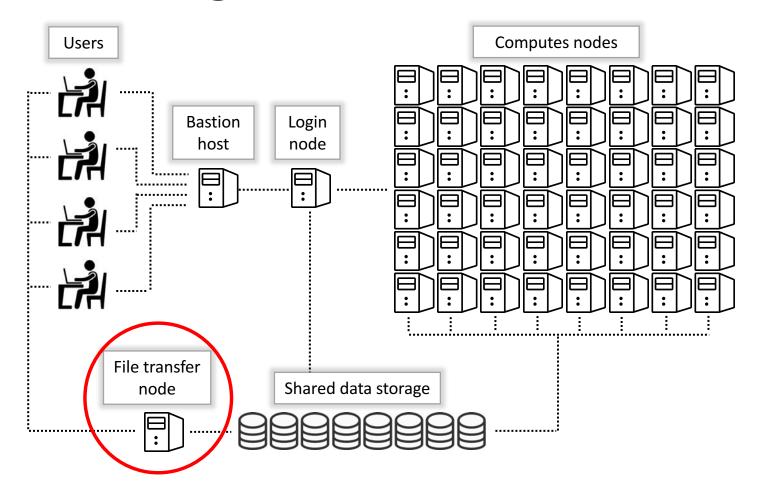
Command to list all the available storage options – uquota

[dshyshlov@login3 ~]\$ uquota						
	used	soft limit	hard limit	files/limit		
dshyshlov home & PBS	13 . 97G	14G	15G	193575		
/extra/dshyshlov	55 . 67G	200G	200G	102492/120000		

	Storage	Back-up	File limits	Speed
/home	15 GB	Nightly	None	
/extra	200 GB	None	600 files / GB	
/xdisk	200 – 1000 GB (45 day limit)	None	None	
/rsgrps	Rented space	None	600 files / GB	
/tmp	Varies ~ 800 GB (Ocelote)	None	None	Fastest (on node)



Transferring Files



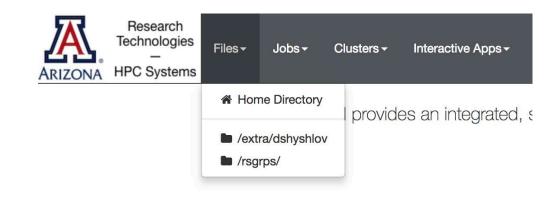
filexfer.hpc.arizona.edu

Transferring Files

- Ocelote has two specific nodes for file transfer
 - hostname filexfer.hpc.arizona.edu
- Command line options:
 - scp, sftp, rsync, irods
- GUI options
 - Windows based: WinSCP
 - Cross-platform: Cyberduck
- Parallel data transfer
 - Globus
 - best option for large files

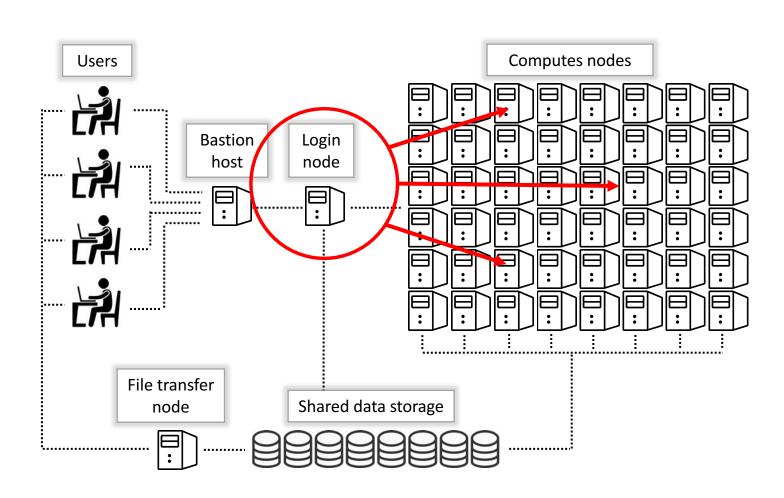
Transferring Files (web browser)

- Display and manage your files
- Edit text files
- Drag and drop files to/from the file explorer





From the login node to compute



From the login node to compute

- How do we know if there are any available nodes?
- How do we decide who gets what and when?
- How do we ensure that a task gets the resources it needs?

- Scheduler!
- Software that manages the HPC resources and decides which computation runs where and when.

Scheduler

• Ocelote uses the scheduler PBS Pro.

• Every computation that requests resources from the scheduler is called a *job*.

• Submitting a job means requesting resources from the scheduler and giving it a list of commands to run.

Scheduler

Scheduler receives a request for resources and creates a job



Job is put in the queue, where it waits for the resources



Job is assigned to the compute nodes and performs computation



When job is finished output and error files are created

Queues

- Standard queue
 - 36,000 CPU-hours/month per group
 - higher priority

- Windfall queue
 - No time limit
 - Preemption

- Debug queue
 - 10 min limit

View the contents of a file first_script.pbs with a command

```
$ cat first_script.pbs
```

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

It's a bash shell script

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
echo 'This script is running on:'
hostname
sleep 120
```

```
#!/bin/bash
#PBS —W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

PBS directives

Name of your HPC group (usually it is your sponsor's NetID). You can find your groups name with a value command.

```
#!/bin/bash
#PBS —W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

```
#!/bin/bash
#PBS —W group list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
echo 'This script is running on:'
hostname
sleep 120
```

Type of queue

"Select statement".

Create a custom computer from Ocelote's resources for your job.

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -a standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

How long your custom computer will exist.

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem≠6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

List of commands to run on your custom computer

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb/pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

Exercise – running a batch job

Submit your first job on Ocelote.

```
$ qsub first_script.pbs
qsub: Bad GID for job execution
```

Edit first_script.pbs to correct the group name.

\$ va find the group name

```
$ qsub first_script.pbs
1827586.head1.cm.cluster
```

Exercise – running a batch job

Check on the status of the job.

\$ qstat -u NetID

 Look in the standard output and error files (use cat command).

Exercise – customizing a job

• Submit the job that uses 4 cores from 1 node.

 What parameters do you need to change in the PBS script?

Exercise – deleting a job

 You can delete the job from the queue or when it's running with a qdel command.

\$ qdel 1827579.head1.cm.cluster

You can find a full job ID with a command

\$ qstat -wa -u NetID

Interactive job

- Request compute node resources for interactive work
 - interactive data analysis or compute
 - long-running debugging
 - copying large files

\$ qsub -I first_script.pbs

 Close the interactive session with the command exit or logout.

Interactive job

```
#!/bin/bash
#PBS -W group_list=hpcteam
#PBS -q standard
#PBS -l select=1:ncpus=1:mem=6gb:pcmem=6gb
#PBS -l walltime=00:05:00
echo 'This script is running on:'
hostname
sleep 120
```

Accessing Software

GUI



Command line

\$ matlab

Or more often...

\$ matlab -noscreen < input.m > output.txt

Accessing Software

```
$ matlab
-bash: matlab: command not found
```

• To run any software in Linux the system must know where to look for appropriate binaries and libraries.

Check your environment:

```
$ echo $PATH
```

Accessing Software

• Environment modules are a convenient way to customize your environment to use software

Manage your modules with a command

\$ module

• Running *module* command without any options will open a help page.

Accessing Software: hand-on activity

- List your current modules
 - \$ module list
- List all the available modules
 - \$ module avail
- List all the available MATLAB modules
 - \$ module avail matlab
- Load a MATLAB module
 - \$ module load matlab
- List your current modules

Accessing Software: hands-on activity

Check your environment again:

\$ echo \$PATH

Anything new?

• When you run a batch job *module load* commands must be in the PBS script.

Being a good HPC user

- Things that can "break" the system:
 - heavy use of the login node
 - too many jobs
 - too many files
 - heavy I/O jobs
 - copying GB of data
 - for loops in PBS scripts

Getting help

- HPC documentation docs.hpc.arizona.edu
- HPC consulting hpc-consult@list.arizona.edu
- Visualization consulting vislab-consult@list.arizona.edu
- Statistics consulting stat-consult@list.arizona.edu