

# Introduction to High-Performance Computing (HPC)

<http://tinyurl.com/QMB-orchestra-slides>

Radhika Khetani, PhD

Lead Bioinformatics Trainer

Harvard Chan Bioinformatics Core (HBC)

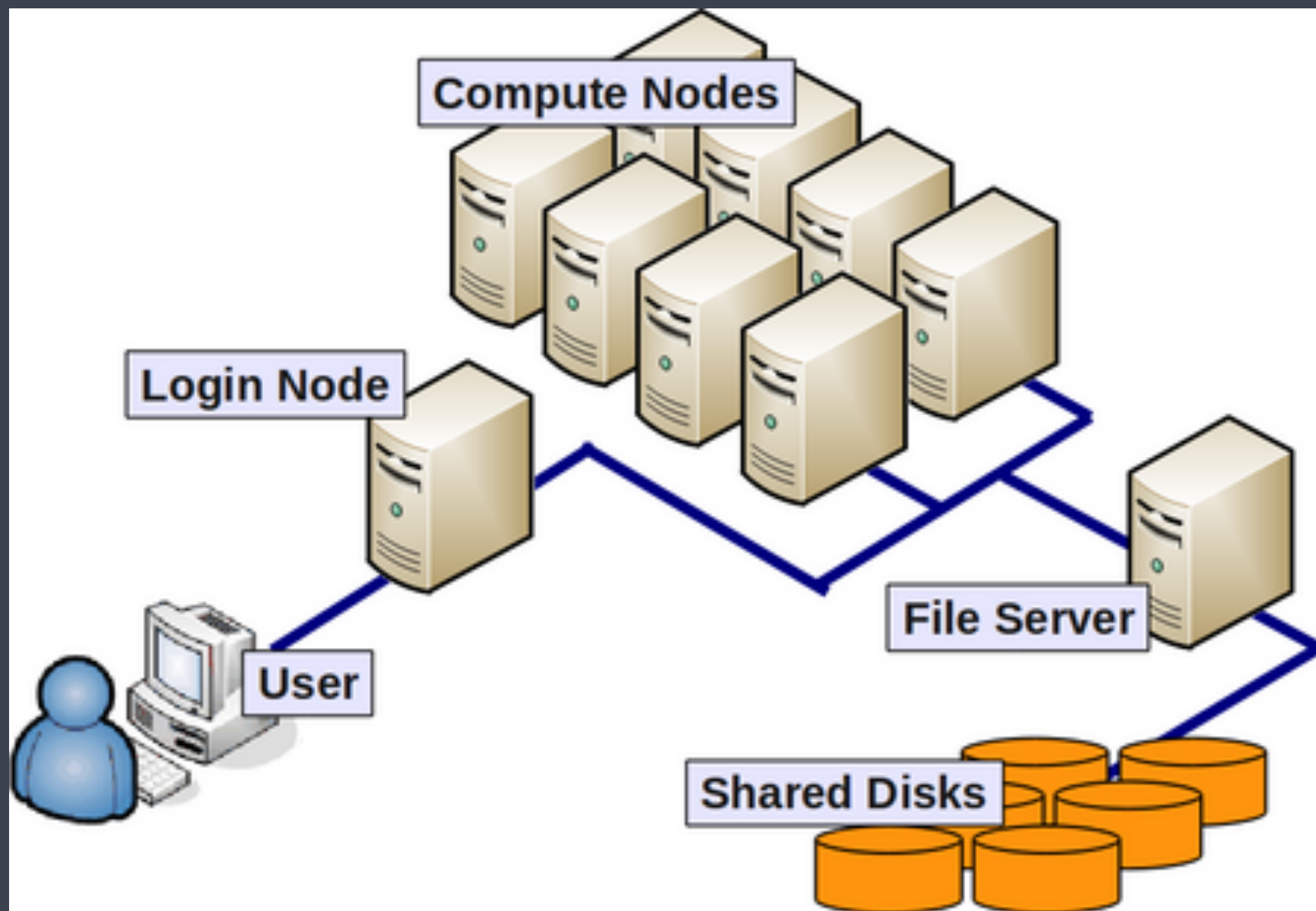
# HPC cluster

*“High Performance Computing most generally refers to the practice of aggregating computing power in a way that delivers much higher performance than one could get out of a typical desktop computer or workstation in order to solve large problems in science, engineering, or business.”*

## Need for HPC cluster?

- **Resources:** Your computer does not have the resources to run the desired NGS analysis
- **Speed:** You want to produce results faster than your computer can
- **Software:** You need software that is unavailable or unusable on your computer

# HPC cluster



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# HPC cluster components

**Nodes:** Individual computers in the cluster

**Cores:** individual processing units available within each CPU of each Node

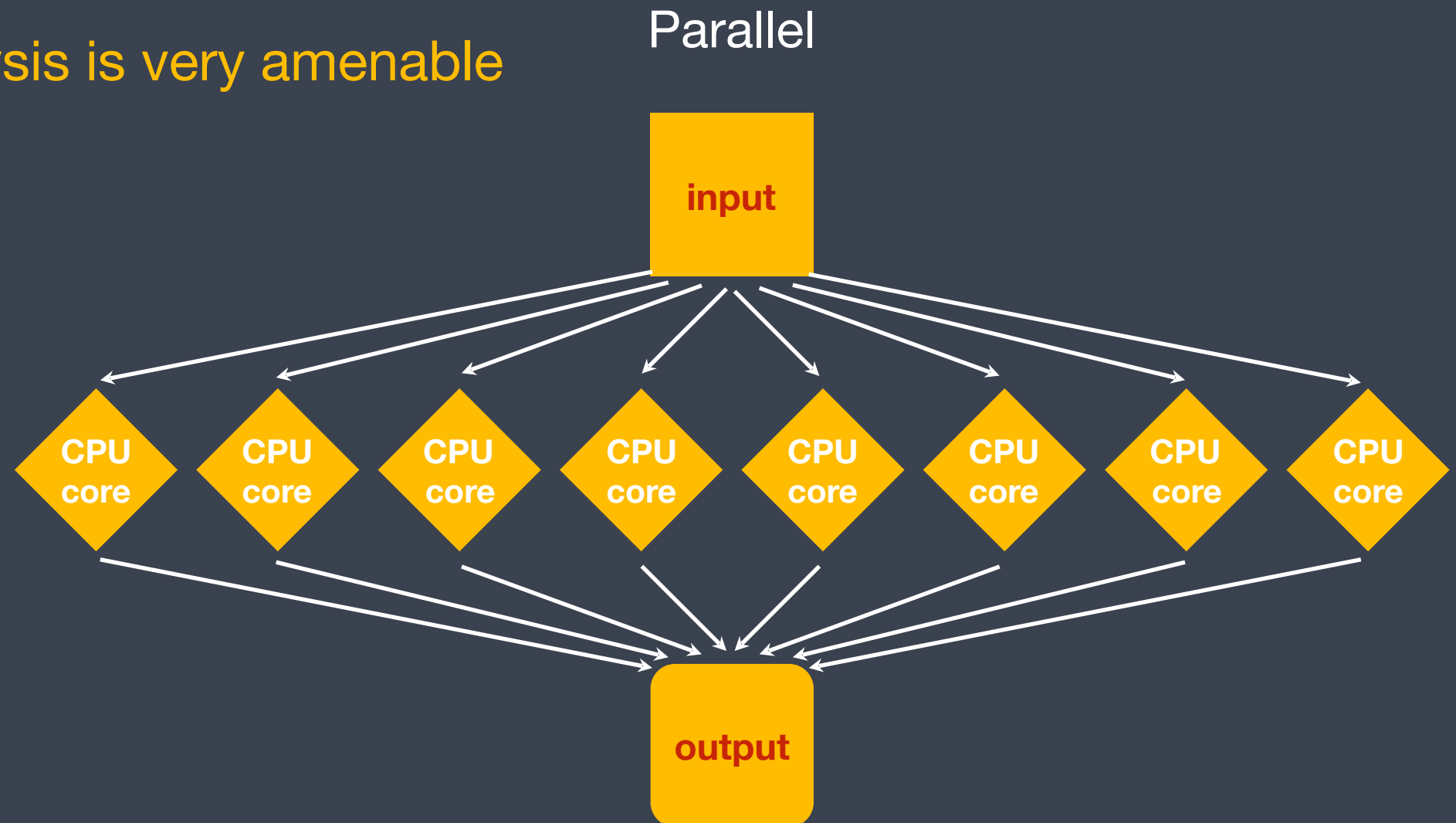
*e.g. a “Node” with eight “quad”-core CPUs = 32 cores for that node.*

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**Shared disk:** storage that can be shared (and accessed) by all nodes

# Multi-threading

NGS data analysis is very amenable to this strategy.



Faster and more efficient

# HPC cluster

- multi-user, shared resource
- lots of nodes = lots of processing capacity + lots of memory
- a system like this requires constant maintenance and upkeep.



Please see our [Orchestra status](#) page for known issues.

The **Orchestra** platform provides UNIX-based high performance computing, web hosting, and database hosting services at Harvard Medical School.

**Orchestra** and its associated services are managed by the [Research Computing Group](#), part of the HMS [Information Technology Department](#).

Please [submit a request](#) via the RC web site for help with **Orchestra** or feedback. You can also subscribe to the [mailing list](#) for Orchestra users.

Wiki page: <https://wiki.med.harvard.edu/Orchestra>



# Introduction to High Performance Computing and Orchestra

HMS Research Computing  
2016

*(Thanks to Kris Holton for sharing HMS-RC slides)*



# What is Orchestra?

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- Tech specs:
  - Over 550 compute nodes
  - Over 7,500 cores
  - Over 42TB RAM
- CentOS 6 Linux
- LSF scheduler
- Total 28+PB storage



# Using Orchestra!

# 1. Logging in to remote machines (securely)

- When logging in we used the “ssh” command,  
**ssh stands for Secure Shell**
- **ssh** is a protocol for data transfer that is secure, i.e the data is encrypted as it travels between your computer and the cluster (remote computer)
- Commonly used commands that use the **ssh** protocol for data transfer are, **scp** and **sftp**

## 2. Using & installing software

# Using Software Environment Modules

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- Most “software” on Orchestra is installed as an environment module.
- An environment module makes the necessary modifications in \$PATH to make sure the program runs without any issues.
- Allows for clean, easy loading, including most dependencies, and switching versions.

# Using Software Environment Modules

---

```
$ module avail
```

```
$ module avail seq/
```

```
$ module avail stats/
```

```
$ module avail seq/bowtie/
```

What software  
environment modules are  
available on Orchestra?

- Loading modules

```
$ module load seq/bowtie/2.0.6
```

- See all modules that have been loaded

```
$ module list
```

- Unloading modules

```
$ module unload seq/bowtie/2.0.6
```

### 3. The Job Scheduler, LSF



# LSF: Fair Sharing

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- **Load Sharing Facility:** distributes jobs across Orchestra fairly
- Ensures that no single 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

Choosing the proper resources for your job with  
the appropriate `bsub` options

# The “bsub”

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```
$ bsub -q queue -W hr:min job/command
```

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

# bsub options

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`-W 3:00`            *#time requested*  
`-q multicore`    *#name of the queue*  
`-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 %J.err`        *#send errors to file*  
`-o %J.out`        *#send screen output to file*  
`-N`                *#notify when job completes*

# Runtime Limit ( $b_{sub} - W$ )

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- 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

# Multithreading (`bsub -n`)

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- 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`!

# CPU Limit

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- 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 (-p 8) but only 1 core requested (no -n), job killed in 1 hour (8/8)

# Reserving Memory (`bsub -R`)

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- 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



# Shared Queues (`bsub -q`)

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- ***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.

<https://wiki.med.harvard.edu/Orchestra/ChoosingAQueue>

Job submission scripts

# Shell Scripts: The Basics

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```
#!/bin/sh                #always at the top

#commands 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`

Run this script in an interactive session as `sh myshellscript.sh`

OR

Submit it as a job to the LSF scheduler

```
$ bsub -q mcore -W 12:00 -n 4 -N -e %J.err -o %J.out sh myshellscript.sh
```

# Creating a Job Submission Script

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```
#!/bin/sh
#BSUB -q mcore
#BSUB -W 12:00
#BSUB -o %J.out
#BSUB -e %J.err
#BSUB -N
#BSUB -n 4
#BSUB -R "rusage[mem=12000]"

module load seq/tophat2.1.1

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

Save as **myshellscript.lsf**

Run as **\$ bsub < myshellscript.lsf**

Orchestra will notify you when the job is done, or if there is an error

Managing jobs and getting information about  
submitted/running jobs

# Monitoring Jobs

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- List info about jobs/their status:

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

-r (running jobs)

-p (pending jobs)

-l (command entered, long form)

# Terminating Jobs

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- Terminate a job (jobid given at submission)

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

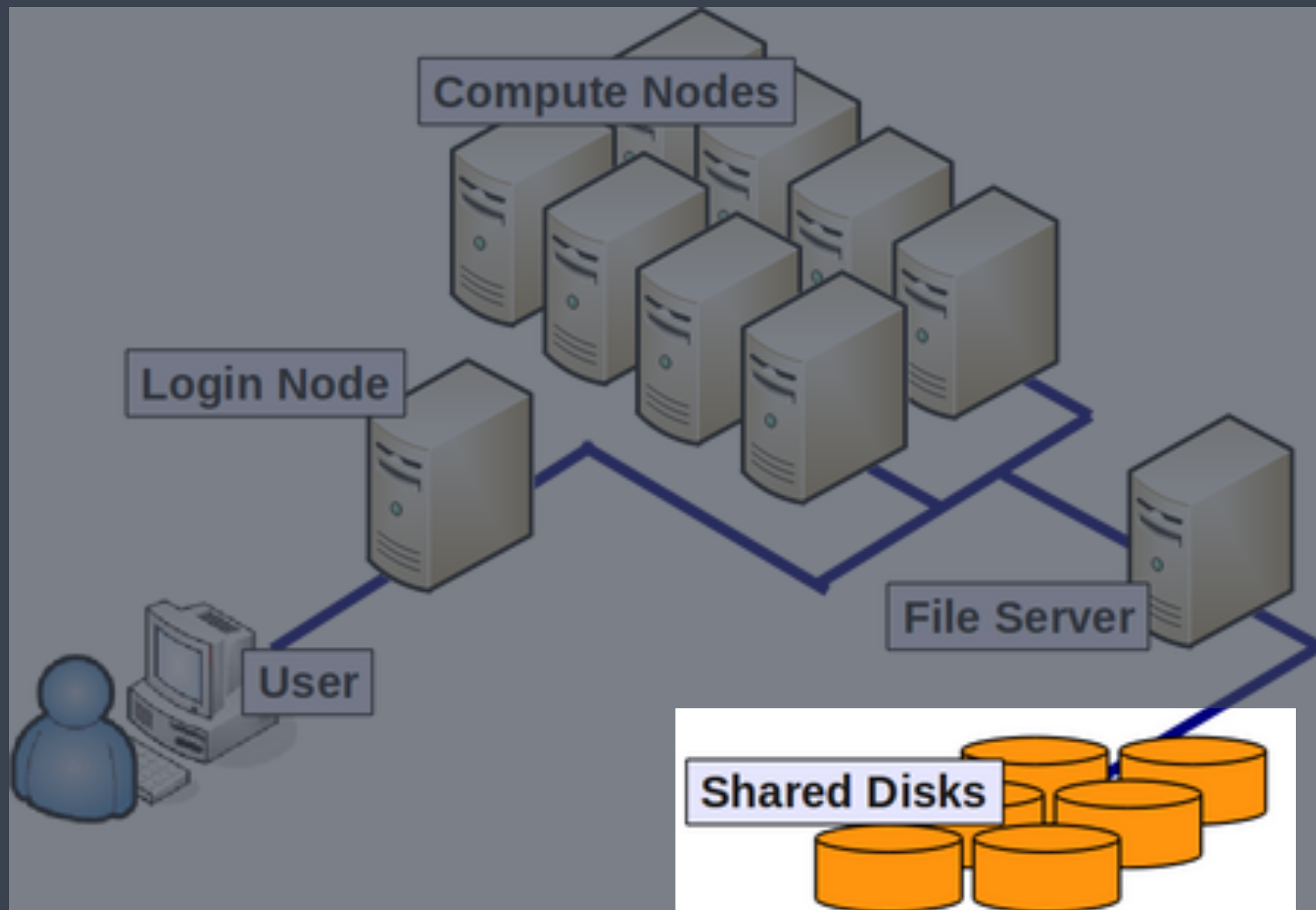
- Terminate all jobs (be careful!)

```
mfk8$loge:~$ bkill 0
```

## 4. Filesystems and storage



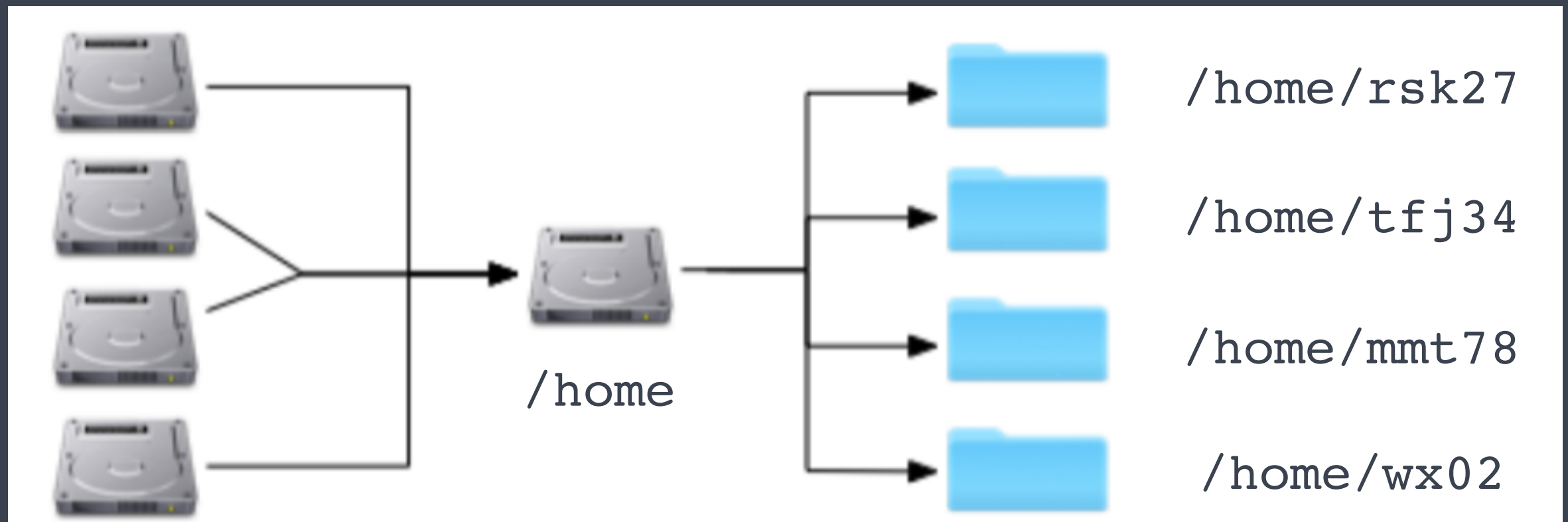
# Filesystems and storage



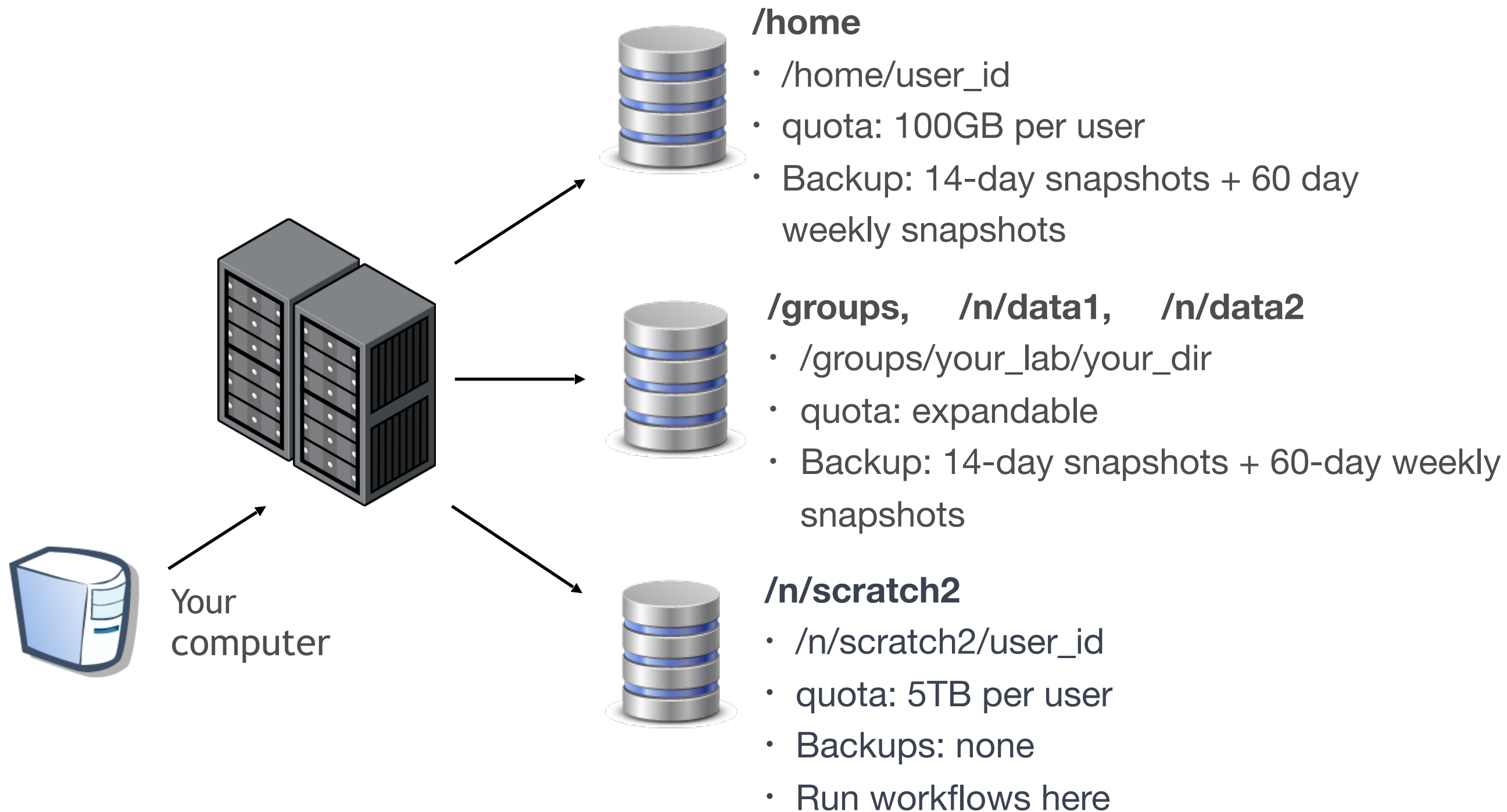
# Filesystems and storage

- Storage on HPC systems is organized differently than on your personal machine
- Physical disks are bundled together into a virtual volume; this volume may represent a single filesystem, or may be divided up, or partitioned, into multiple filesystems
- Filesystems are accessed over the internal network

# Filesystems and storage



# Storage on Orchestra



# Checking storage usage

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- To check your storage available:

`$ quota`

- Home directory (/home/username): you get 100GB.
- Group directories: space varies, can increase.

`/groups/groupname`

`/n/data1`

`/n/data2`

# For more direction:

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- <https://wiki.med.harvard.edu/Orchestra/NewUserGuide>
- <http://rc.hms.harvard.edu>
- [rchelp@hms.harvard.edu](mailto:rchelp@hms.harvard.edu)
- Office Hours: Wednesdays 1-3p Gordon Hall 500