

HPC User Environment 2

Jason Li

HPC User Services

LSU HPC / LONI

sys-help@loni.org

Louisiana State University, Baton Rouge

Feb 7, 2024



▪ HPC User Environment 1

1. Intro to HPC
2. Getting started
3. Into the cluster
4. Software environment (modules)

▪ HPC User Environment 2

1. Basic concepts
2. Preparing my job
3. Submitting my job
4. Managing my jobs

- **HPC User Environment 2**

1. Basic concepts
2. Preparing my job
3. Submitting my job
4. Managing my jobs

■ HPC User Environment 2

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

- **HPC User Environment 2**

1. Basic concepts

- 1) Previously on HPC User Environment 1...
- 2) Job & Job schedulers

2. Preparing my job

- 1) Basic principles
- 2) Job duration (wall time)
- 3) Number of nodes & cores
- 4) Job queues

3. Submitting my job

- 1) Interactive job
- 2) Batch job

4. Managing my jobs

- 1) Useful commands
- 2) Monitoring job health

- **HPC User Environment 2**

1. Basic concepts

- 1) Previously on HPC User Environment 1...
- 2) Job & Job schedulers

2. Preparing my job

- 1) Basic principles
- 2) Job duration (wall time)
- 3) Number of nodes & cores
- 4) Job queues

3. Submitting my job

- 1) Interactive job
- 2) Batch job

4. Managing my jobs

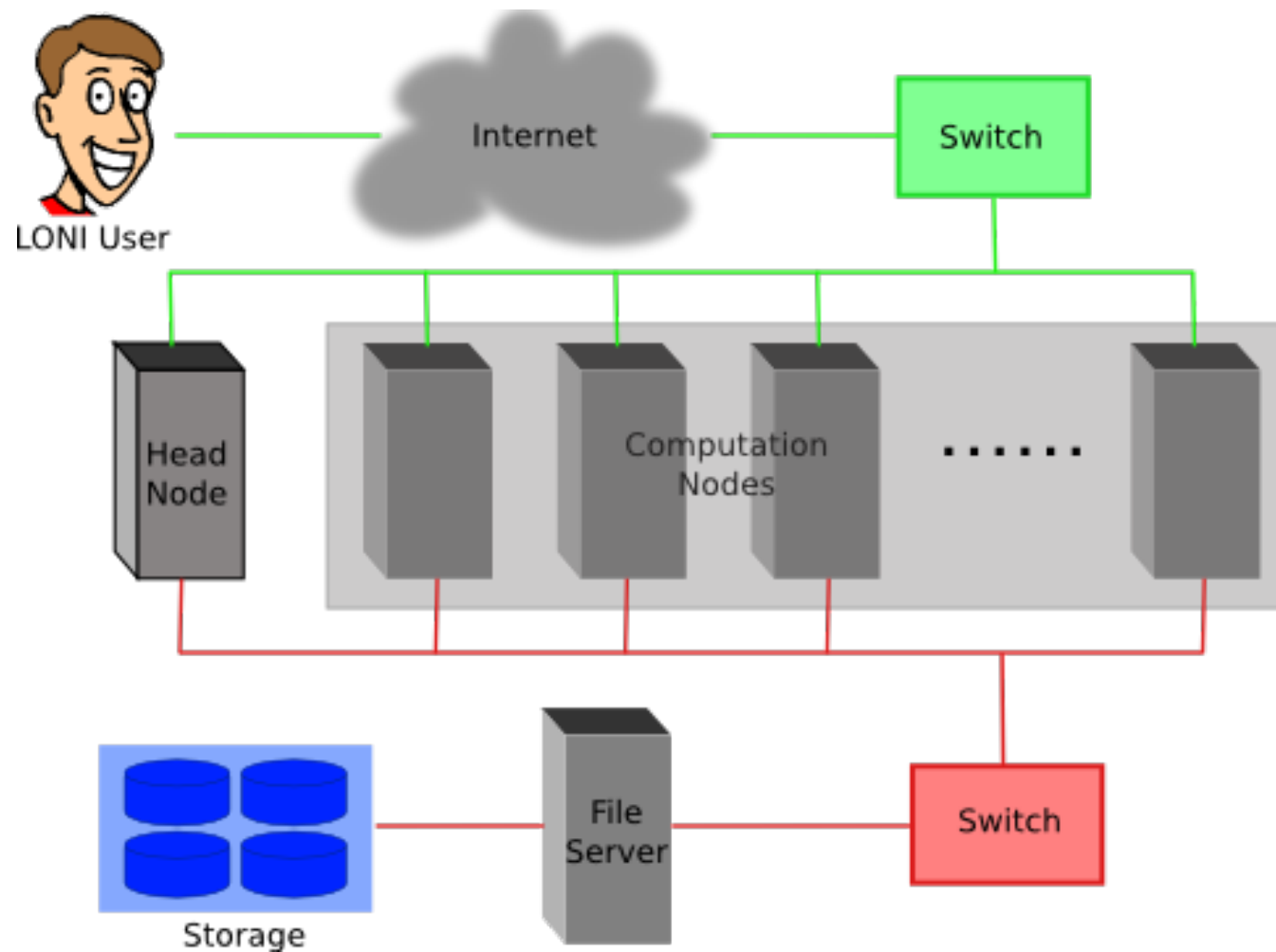
- 1) Useful commands
- 2) Monitoring job health

Two things needed to run jobs on our clusters:

1) Account

2) Allocation

1) Previously on HPC User Environment 1...



1) Previously on HPC User Environment 1...

Run my code on **all** the
resources you have,
however long it takes

sudo!
yum!
apt-get!
...

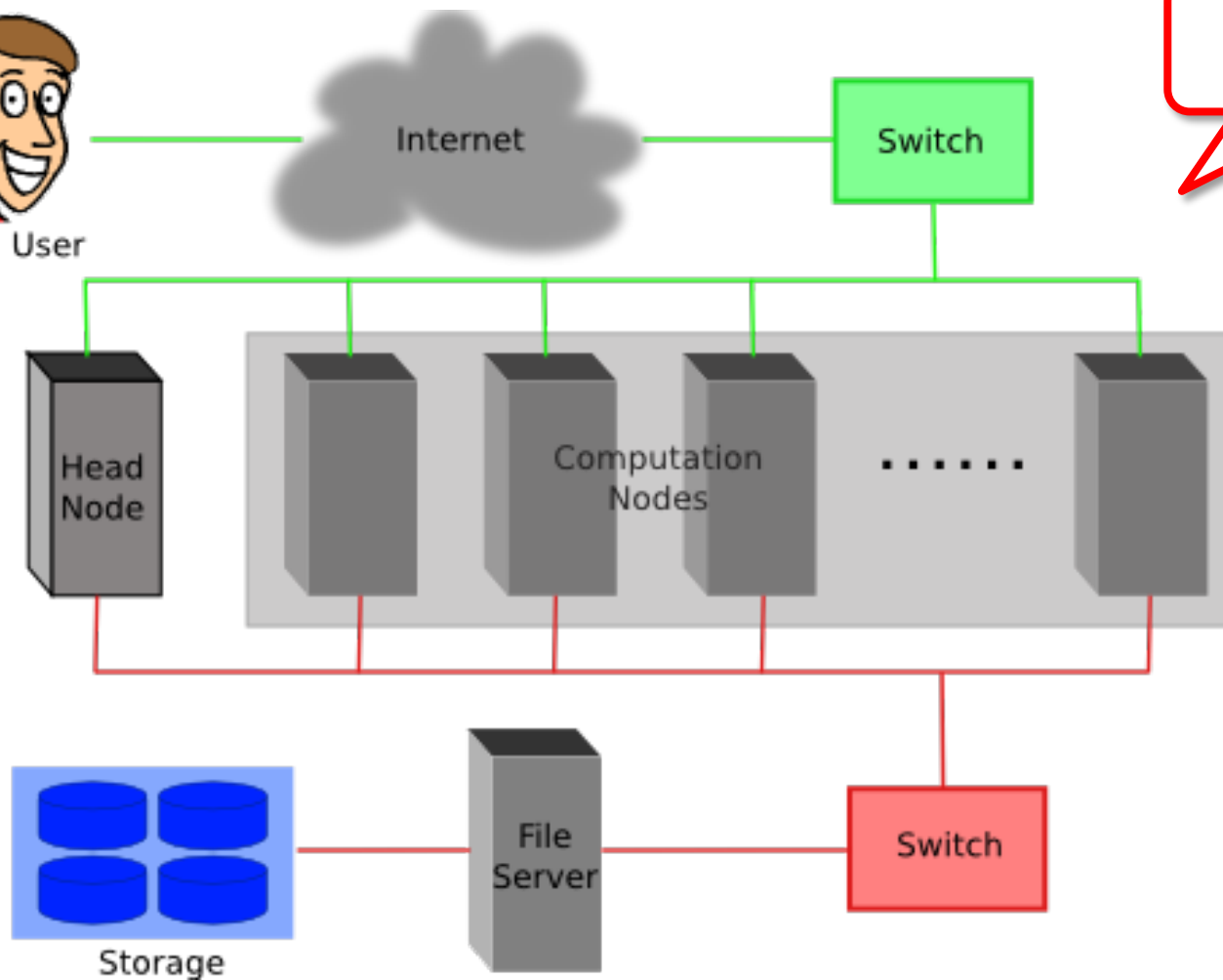


Yes, my
master!

1) Previously on HPC User Environment 1...

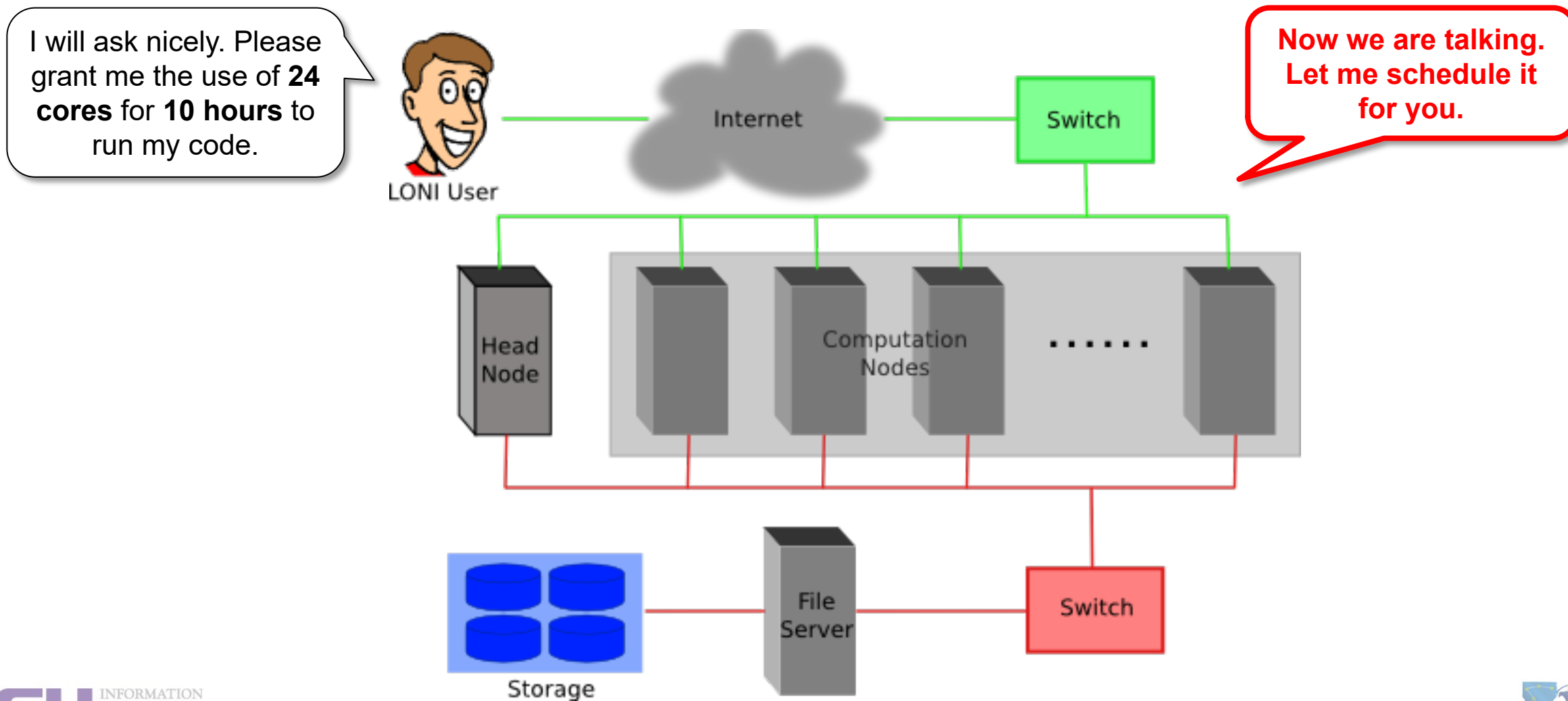
Run my code on **all** the resources you have, **however long** it takes

sudo!
yum!
apt-get!
...



You are not my boss, buddy!

1) Previously on HPC User Environment 1...



- **HPC User Environment 2**

1. Basic concepts

- 1) Previously on HPC User Environment 1...
- 2) Job & Job schedulers

2. Preparing my job

- 1) Basic principles
- 2) Job duration (wall time)
- 3) Number of nodes & cores
- 4) Job queues

3. Submitting my job

- 1) Interactive job
- 2) Batch job

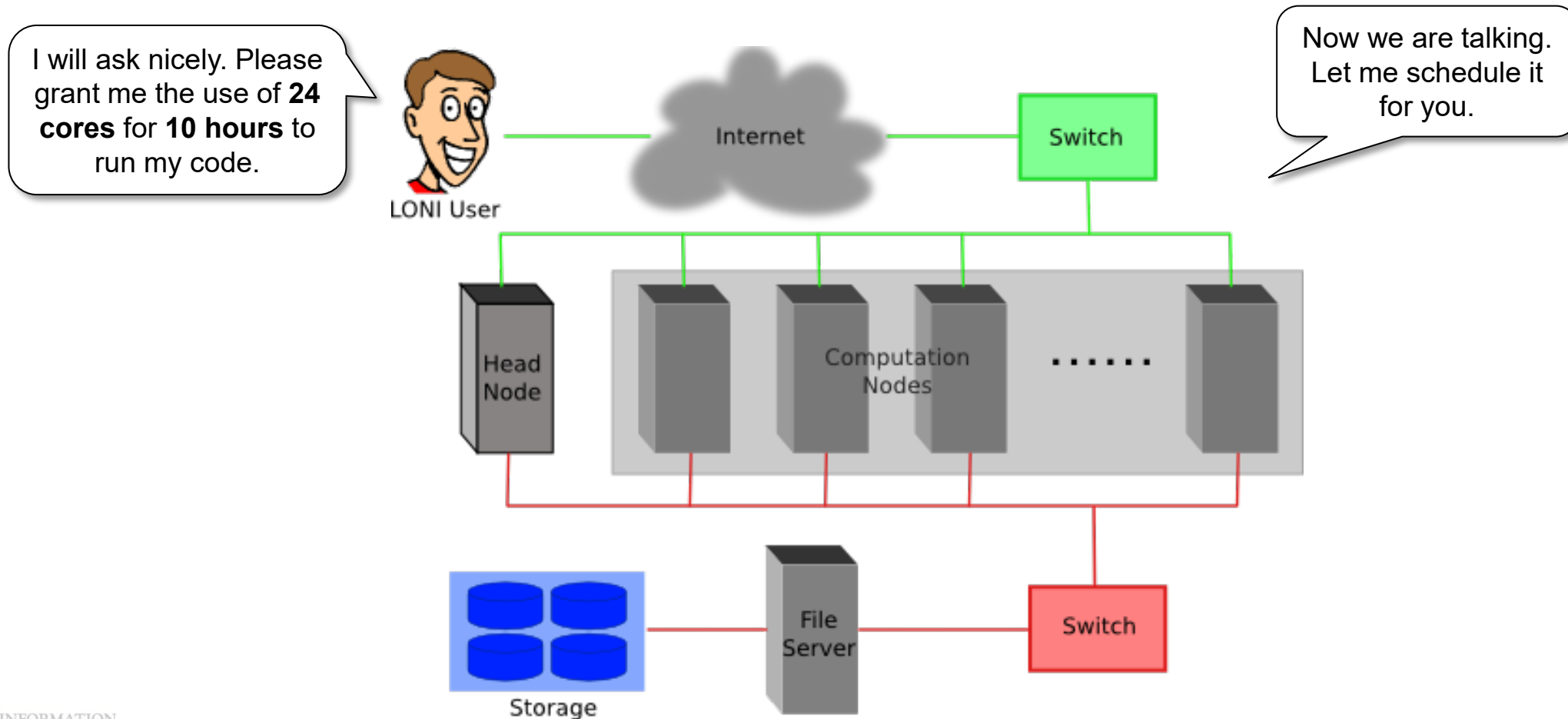
4. Managing my jobs

- 1) Useful commands
- 2) Monitoring job health

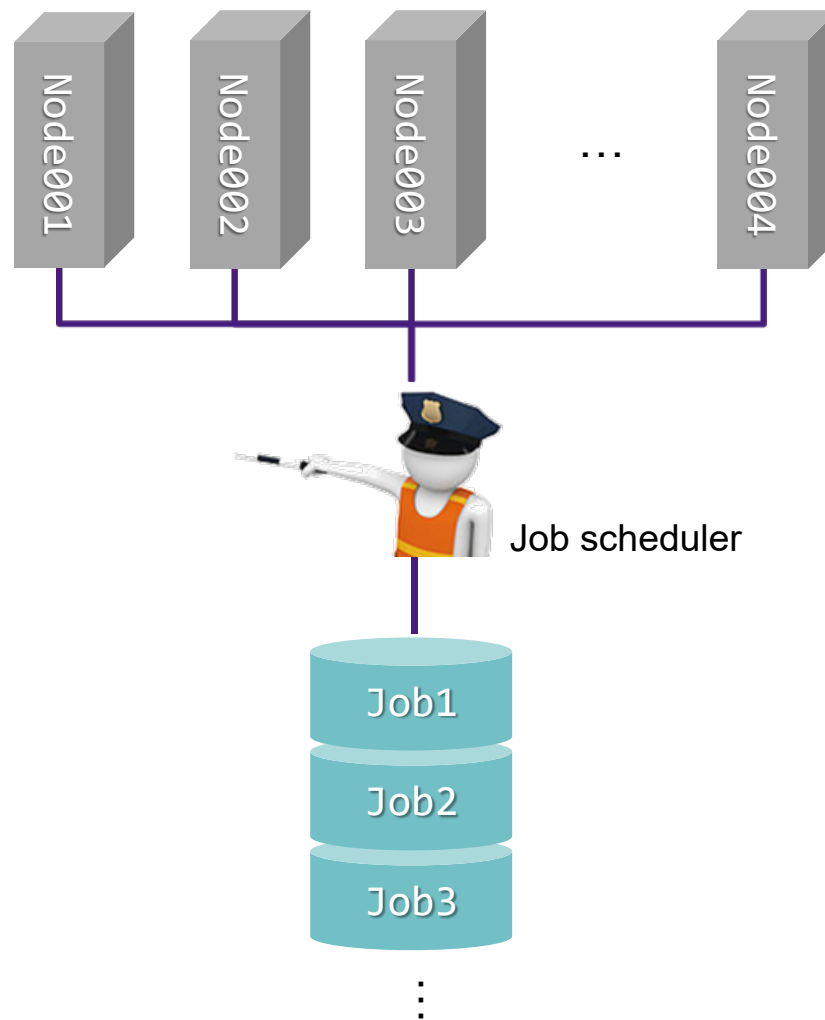
a) What's a "job"?

- A user's request to use **a number of nodes/cores** for **a certain amount of time** on a cluster.
- Calculation **MUST** be done via jobs (**NO** heavy calculation on head nodes!!)
- SUs deducted from allocations based on actual usage of each job.
 - Example:
 - My allocation: 50,000 SU
 - Running a job: 24 core * 10 hours = 240 SU
 - Balance: 49,760 SU

b) What's a "job scheduler"?

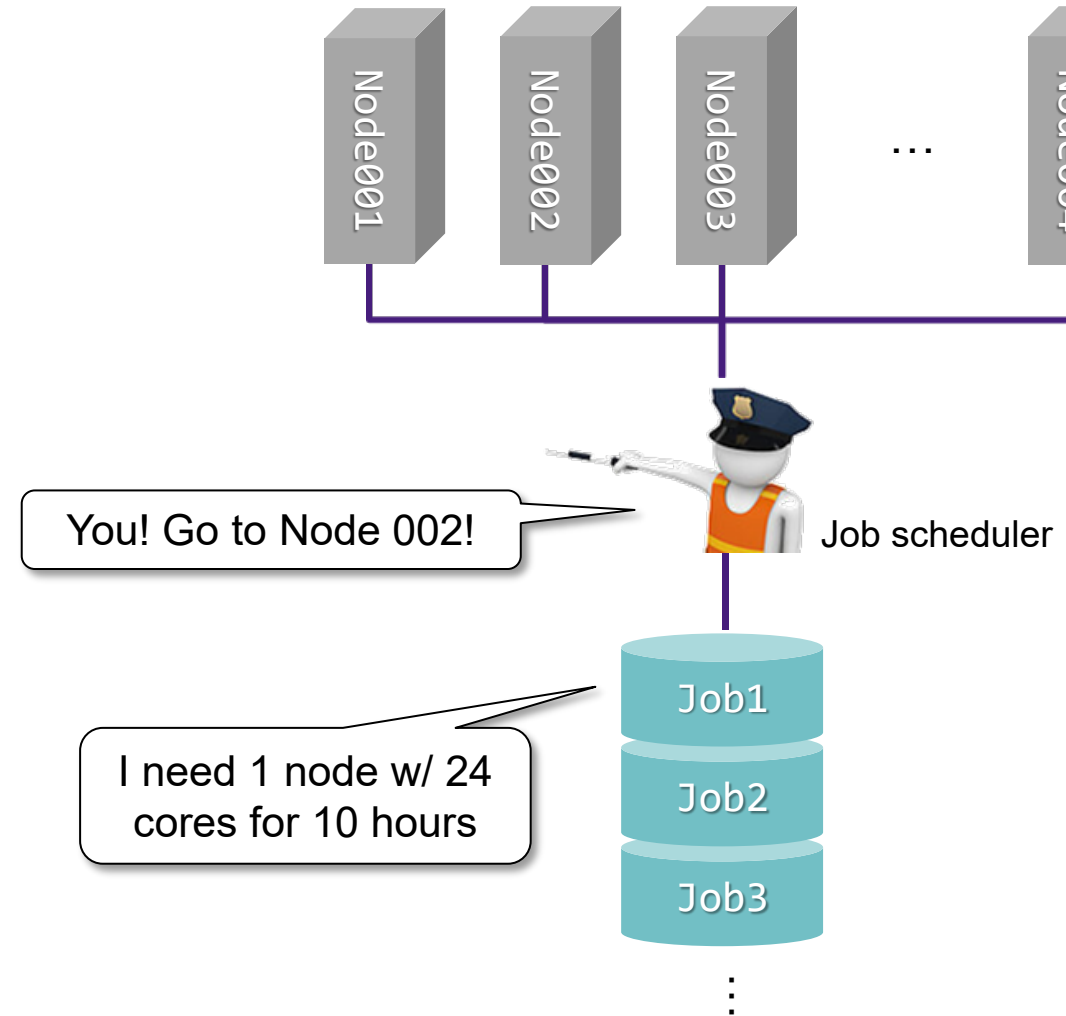


b) What's a "job scheduler"?



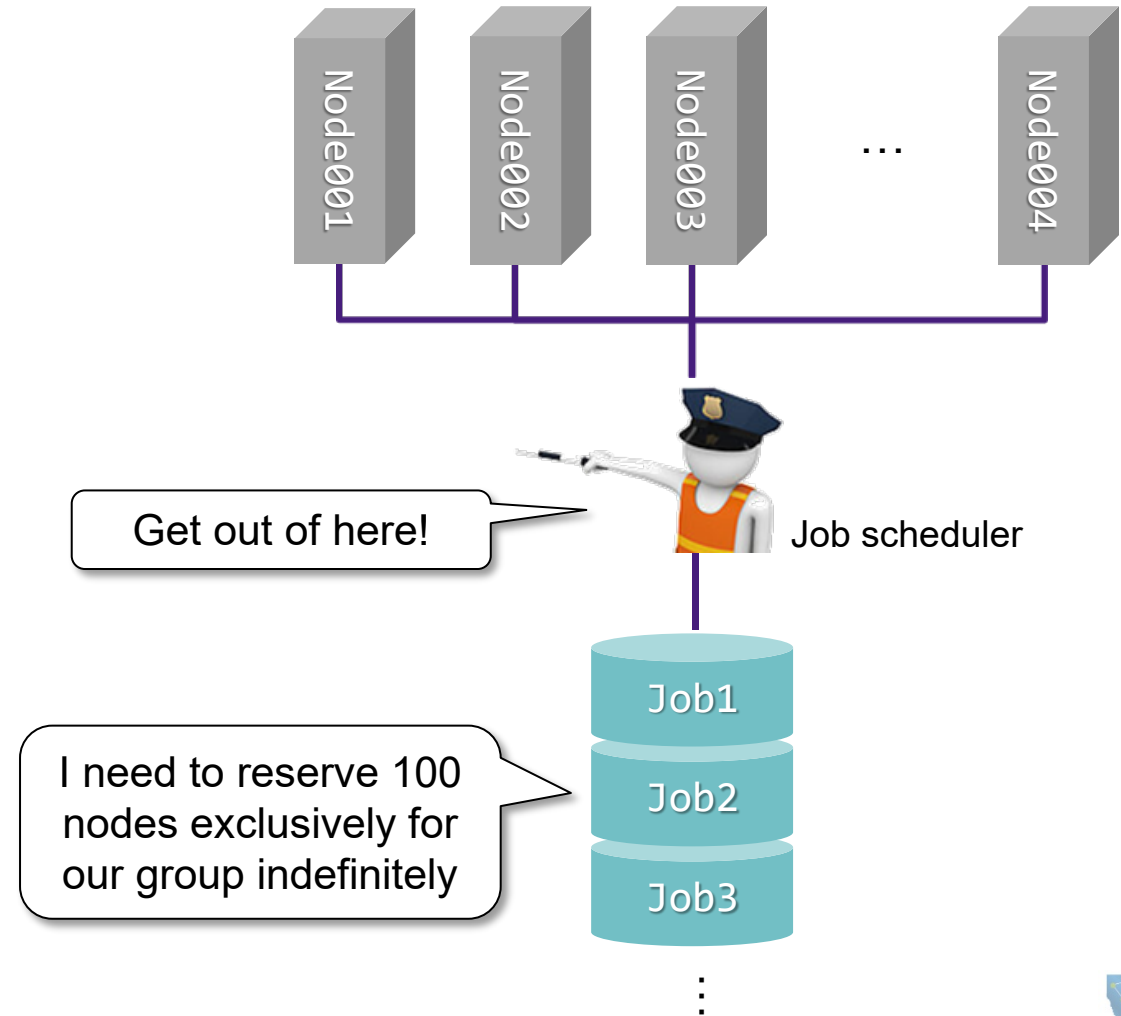
b) What's a "job scheduler"?

- i. Decides which job runs when and where



b) What's a "job scheduler"?

- i. Decides which job runs when and where
- ii. Enforces job policies





2) Job & Job scheduler

b) What's a "job scheduler"?

Job scheduler's responsibilities	Your responsibilities
<ul style="list-style-type: none">• Decides which job runs when and where• Enforces job policies	<ul style="list-style-type: none">• Decide a job's size and duration• Understand the job queuing system and policies• Submit/monitor/cancel jobs• Diagnose job health



2) Job & Job scheduler

b) What's a "job scheduler"?

i) PBS



2) Job & Job scheduler

b) What's a "job scheduler"?

i) PBS

ii) Slurm



2) Job & Job scheduler

b) What's a "job scheduler"?

	LSU HPC	LONI
i) PBS	SMIC	QB2
ii) Slurm	Deep Bayou SuperMike III	QB3



- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health



1) Basic principles

- **Two basic principles of requesting resources**
 - Number of nodes / cores, RAM size, job duration, ...

Large enough ...

Small enough ...



1) Basic principles

- **Two basic principles of requesting resources**
 - Number of nodes / cores, RAM size, job duration, ...

Large enough ...	Small enough ...
<ul style="list-style-type: none">• To successfully complete your job	<ul style="list-style-type: none">• To ensure quick turnaround• Not to waste resources for other users

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

2) Job duration (wall time)

- What is it?
 - **Real-world** (**wall**) time taken from the start to the end
 - Must tell job scheduler how long you want your job to run
 - There is a **maximum** wall time you may request (see later)

2) Job duration (wall time)

- FAQ

Q	A
<ul style="list-style-type: none">What if my command is still running when the wall time runs out?	<ul style="list-style-type: none">Job terminated, any running process killed
<ul style="list-style-type: none">What if all my commands in the job finished before the wall time runs out?	<ul style="list-style-type: none">Job exits successfully when all commands finished
<ul style="list-style-type: none">If my job exits before requested wall time, how many SUs will I be charged?	<ul style="list-style-type: none">You will be charged based on your actual time used (if less than requested)
<ul style="list-style-type: none">In that case, why don't I just request maximum wall time every time?	<ul style="list-style-type: none">Your queuing time may be long...

2) Job duration (wall time)

- Back to basic principles...

Large enough ...	Small enough ...
<ul style="list-style-type: none">• To successfully complete your job	<ul style="list-style-type: none">• To ensure quick turnaround• Not to waste resources for other users

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

3) Number of nodes & cores

- Previously in HPC User Environment 1 ...

SuperMIC		Deep Bayou		SuperMike III	
Hostname	smic.hpc.lsu.edu	Hostname	db1.lsu.edu	Hostname	mike.hpc.lsu.edu
Peak Performance/TFlops	925	Peak Performance/TFlops	257	Peak Performance/TFlops	1,285
Compute nodes	360	Compute nodes	13	Compute nodes	183
Processor/node	2 10-core	Processor/node	2 24-core	Processor/node	2 32-core
Processor Speed	2.8 GHz	Processor Speed	2.4 GHz	Processor Speed	2.6GHz
Processor Type	Intel Xeon 64bit	Processor Type	Intel Cascade Lake Xeon 64bit	Processor Type	Intel Xeon Ice Lake
Nodes with Accelerators	360	Nodes with Accelerators	13	Nodes with Accelerators	8
Accelerator Type	Xeon Phi 7120P	Accelerator Type	2 x NVIDIA Volta V100S	Accelerator Type	4 NVIDIA A100
OS	RHEL v6	OS	RHEL v7	OS	RHEL v8
Vendor		Vendor	Dell	Vendor	Dell
Memory per node	64 GB	Memory per node	192 GB	Memory per node	256/2048 GB
Detailed Cluster Description		Detailed Cluster Description		Detailed Cluster Description	
User Guide		User Guide		User Guide	
Available Software		Available Software		Available Software	

3) Number of nodes & cores

- **When submitting you job...**
 - Must tell job scheduler the number of nodes & cores you need

3) Number of nodes & cores

- FAQ

Q	A
<ul style="list-style-type: none">My code runs slow. Can I request more nodes / cores to make it faster?	<ul style="list-style-type: none">Not quite! Your code most likely is NOT using multiple nodes / cores, if:<ul style="list-style-type: none">You do not know if it is using multiple nodes / coresYou did not tell it to use multiple nodes / coresYou are not familiar with names like “MPI” / “OpenMP”Underutilization is THE most common warning received on our clusters
<ul style="list-style-type: none">How many nodes / cores should I request?	<ul style="list-style-type: none">In short: We can’t answer thatEach code / job is different. You must test to determine

3) Number of nodes & cores

- Back to basic principles...

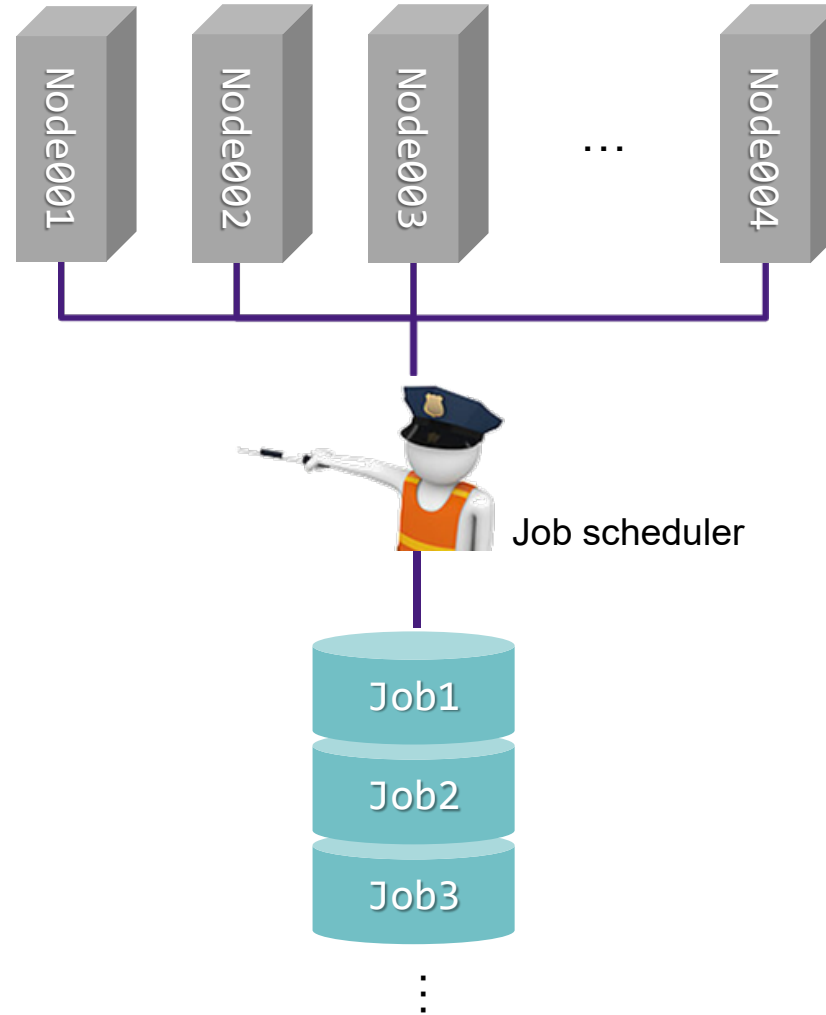
Large enough ...	Small enough ...
<ul style="list-style-type: none">• To successfully complete your job	<ul style="list-style-type: none">• To ensure quick turnaround• Not to waste resources for other users



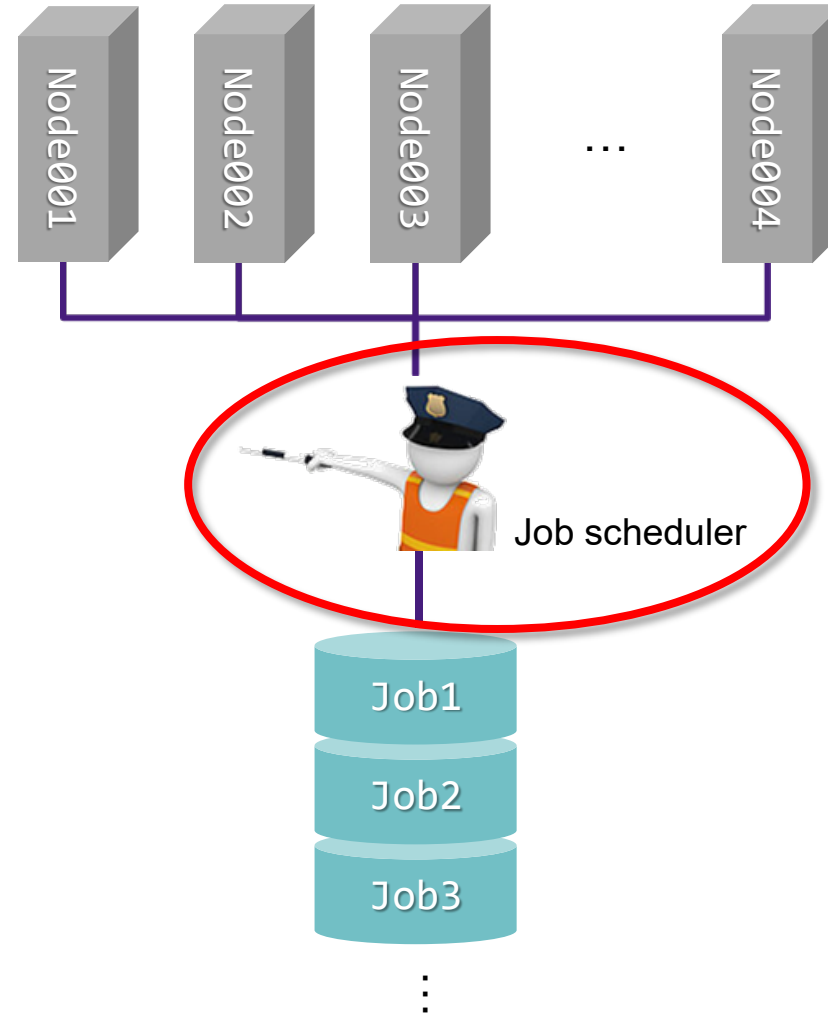
- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

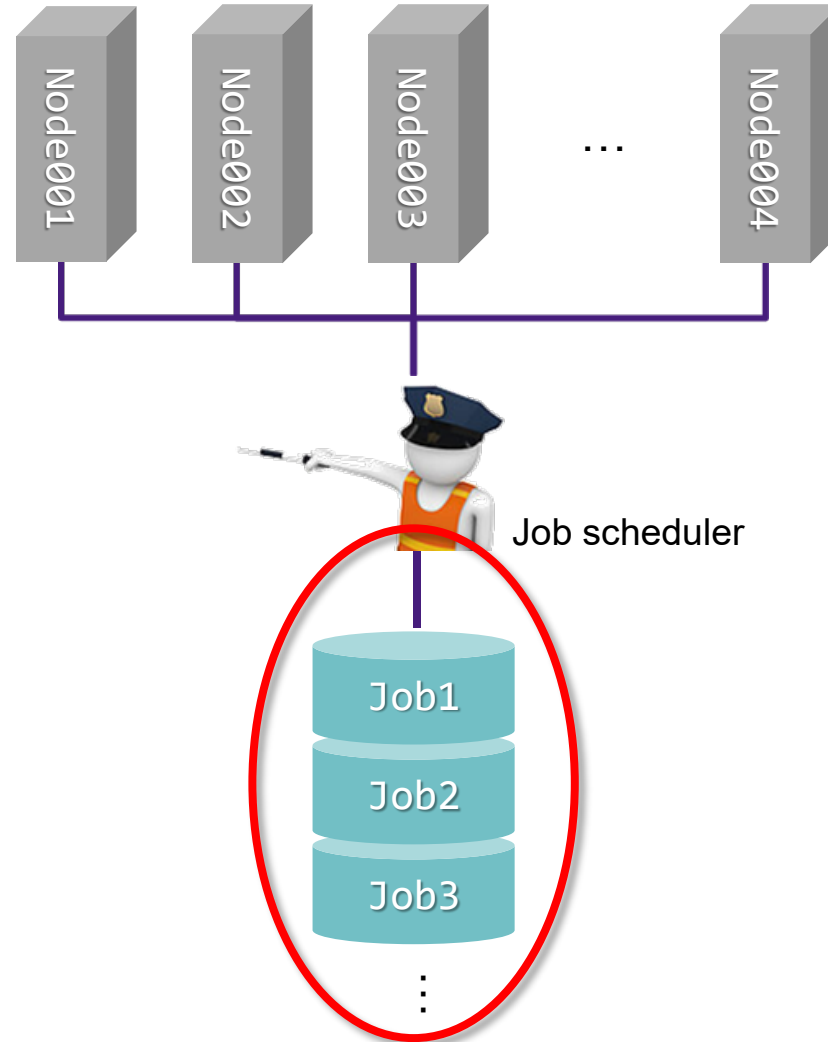
4) Job queues



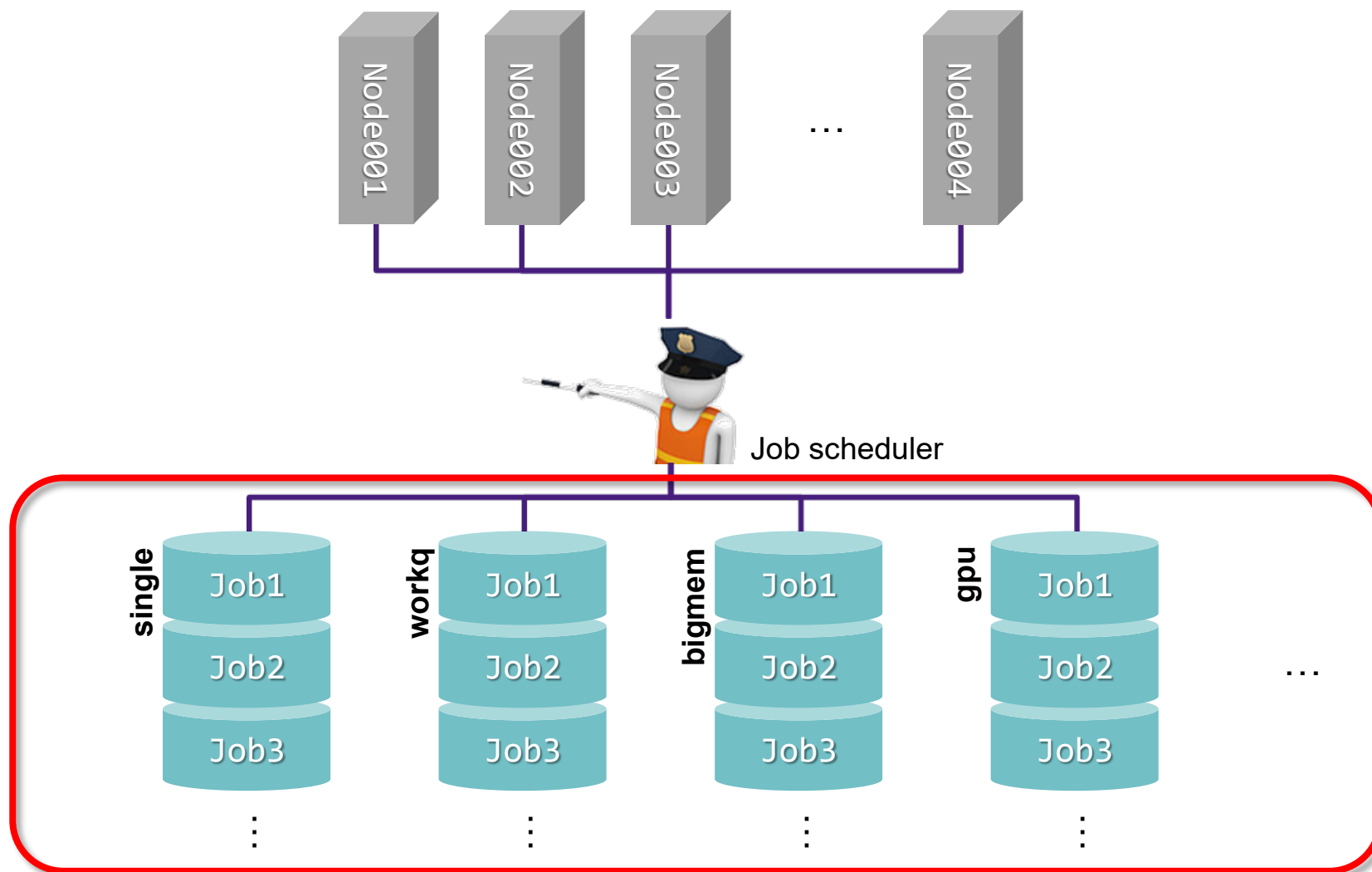
4) Job queues



4) Job queues

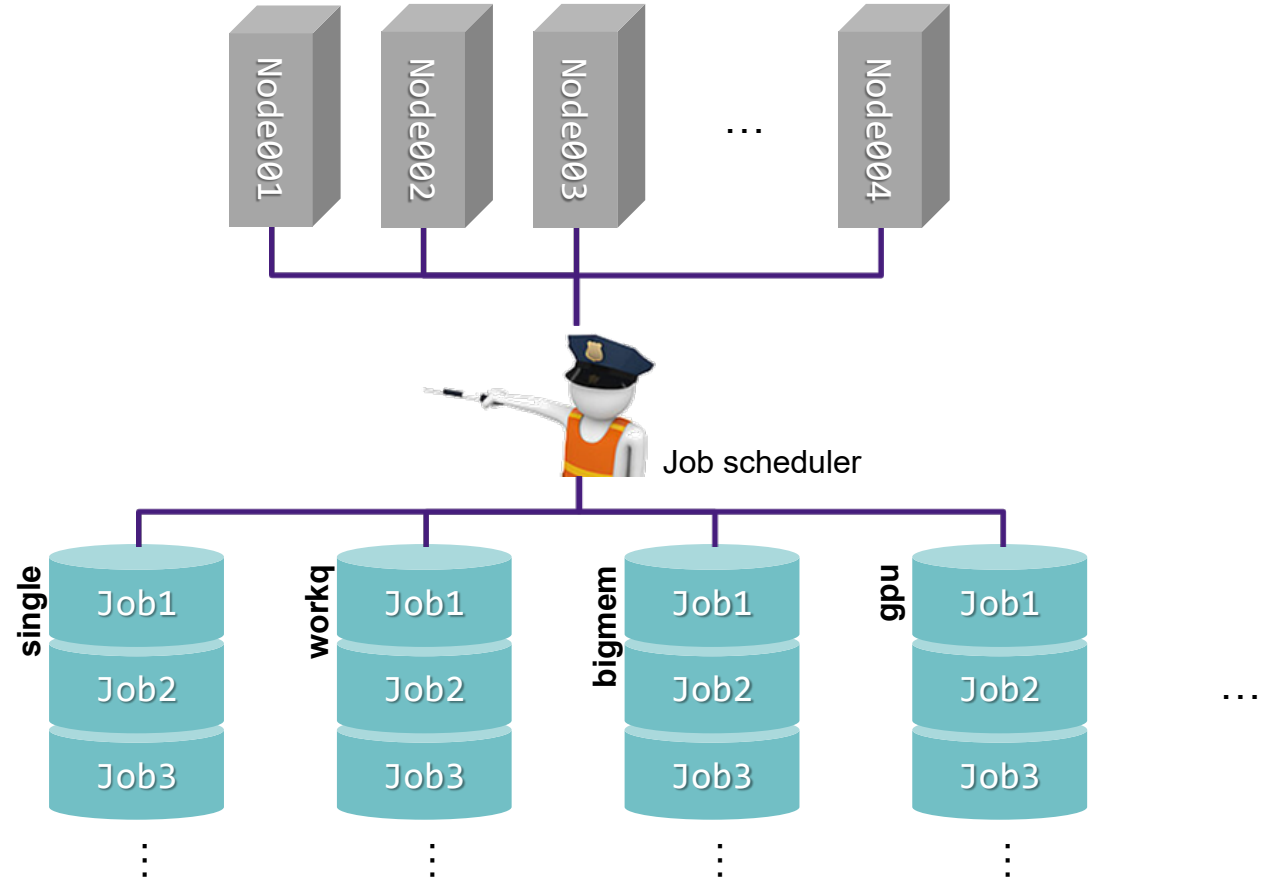


4) Job queues



a) Definition

- Different groups / lines where jobs are being grouped into
- Must pick one queue to submit job
- Goal: Use the resources more efficiently



4) Job queues

a) Definition



b) Available queues

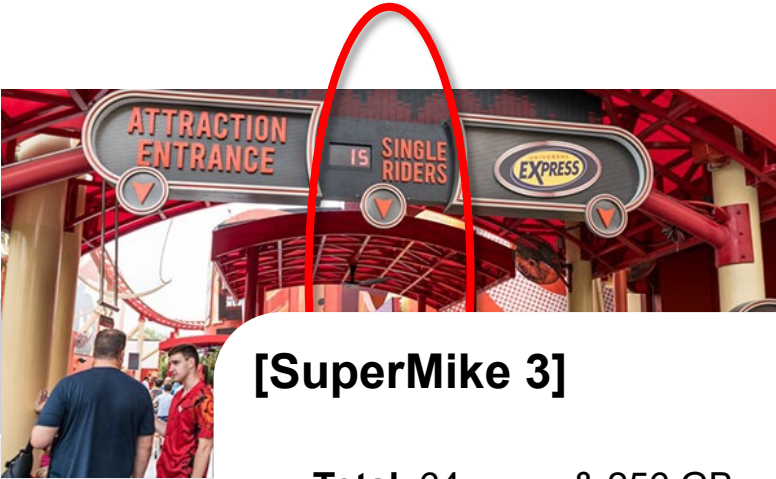
i. **workq** / **ckpt**

Description		<ul style="list-style-type: none">• General purposes• Most likely your default queue• Difference: non-preemptable (workq) vs. preemptable (ckpt)
Names		<ul style="list-style-type: none">• All clusters: workq / ckpt
Resource availability	Nodes	<ul style="list-style-type: none">• One or multiple• Up to a maximum
	Cores	<ul style="list-style-type: none">• All cores on the node(s)
	Memory	<ul style="list-style-type: none">• All memory on the node(s)
Max duration		<ul style="list-style-type: none">• 72 hours (3 days)

b) Available queues

ii. **single**

Description		• Only need a portion of one node
Names		• All clusters: single
Resource availability	Nodes	• A portion of one node
	Cores	• PBS: 1/2/4/6/8 • Slurm: 1 ~ all cores
	Memory	• A portion, proportional to the number of requested cores
Max duration		• 168 hours (7 days)



[SuperMike 3]

- **Total:** 64 cores & 256 GB memory
→ 4 GB / core
- **Request:** 10 cores
→ 40 GB memory

b) Available queues

iii. **bigmem**

Description		<ul style="list-style-type: none">Your job needs large memory
Names		<ul style="list-style-type: none">All clusters: bigmem
Resource availability	Nodes	<ul style="list-style-type: none">One or multiple
	Cores	<ul style="list-style-type: none">All cores on the node
	Memory	<ul style="list-style-type: none">All memory on the node
Max duration		<ul style="list-style-type: none">72 hours (3 days)

b) Available queues

iv. GPU

Description		<ul style="list-style-type: none"> Your job needs GPU 	
Names		<ul style="list-style-type: none"> SMIC: v100 gpu, nvlink Deep Bayou (*): gpu SuperMike 3 (*): gpu QB3: gpu 	<ul style="list-style-type: none"> Deep Bayou (*) SuperMike 3 (*): gpu
Resource availability	Nodes	<ul style="list-style-type: none"> One or multiple 	<ul style="list-style-type: none"> Portion of one node
	Cores	<ul style="list-style-type: none"> All cores on the node 	<ul style="list-style-type: none"> Portion of one node
	Memory	<ul style="list-style-type: none"> All memory on the node 	<ul style="list-style-type: none"> Portion of one node
	GPU	<ul style="list-style-type: none"> All GPU devices on the node 	<ul style="list-style-type: none"> 1 ~ all GPU devices
Max duration		<ul style="list-style-type: none"> 72 hours (3 days) 	

[SuperMike 3]

- **Total:** 64 cores & 4 GPUs
→ 16 cores / GPU
- **Request:** 3 GPUs
→ 48 cores

c) Queues by clusters (LSU HPC)

Cluster	Queue	Cores per node (ppn)	Max running jobs	Max nodes per user
SuperMIC	workq	20	45 (global)	86
	checkpt			
	single	1,2,4,6,8,16		2
	v100	36		
	bigmem	28		
DeepBayou	gpu	24,48	4 (global)	4
	nvlink	12,24,36,48		2
SuperMike3	workq	64	32 (global)	96
	checkpt			
	single	1 ~ 64		4
	gpu	16,32,48,64		
	bigmem	64		

c) Queues by clusters (LONI)

Cluster	Queue	Cores per node (ppn)	Max running jobs	Max nodes per user
QB-2	workq	20	32 (global)	128
	checkpt			
	single	1,2,4,6,8		1
	bigmem	48		
QB-3	workq	48	32 (global)	96
	checkpt			
	single	1 ~ 48		
	gpu	48		4
	bigmem	48		2

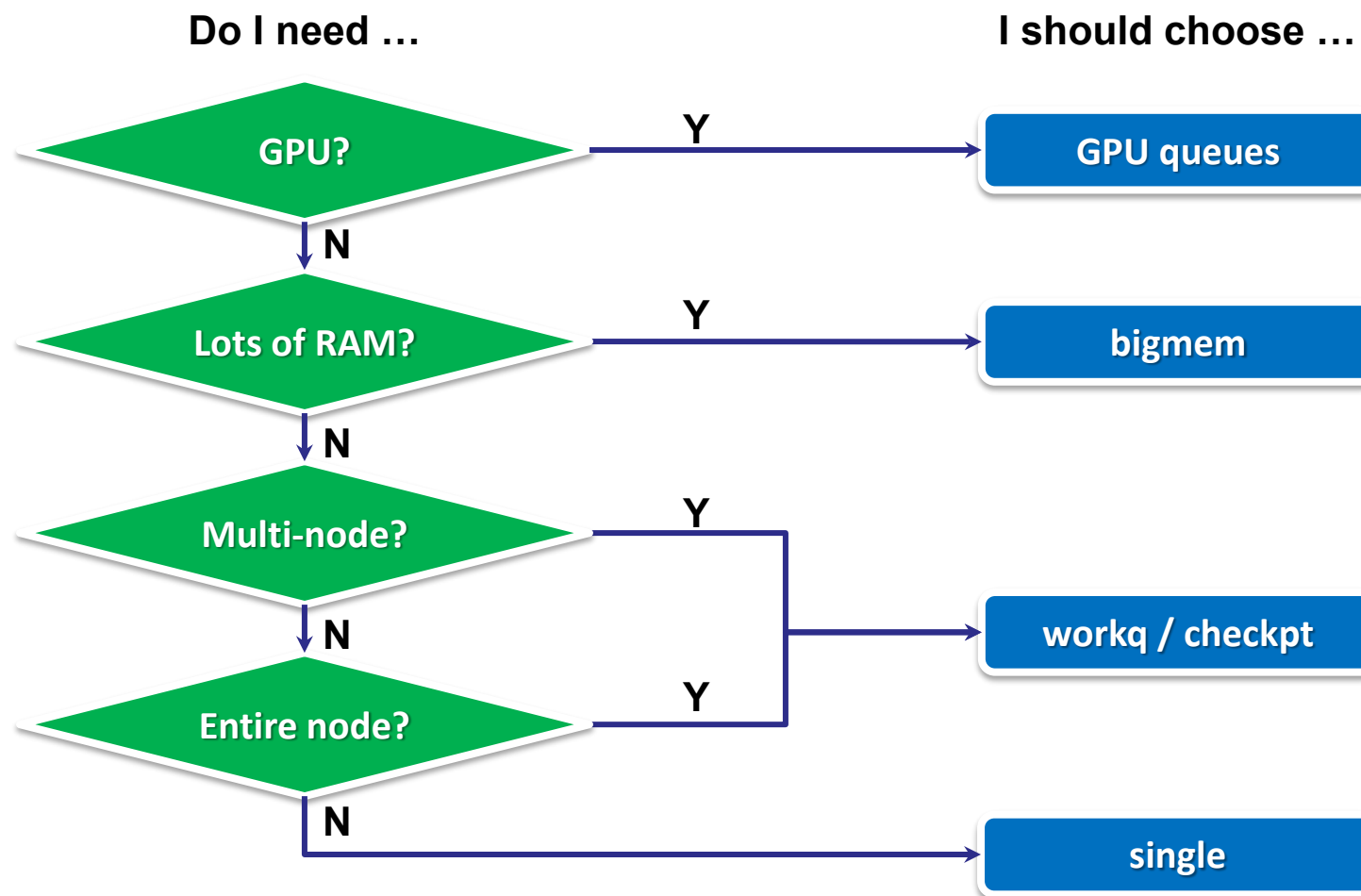


4) Job queues

d) Choose your queue

Large enough ...	Small enough ...
<ul style="list-style-type: none">To successfully complete your job	<ul style="list-style-type: none">To ensure quick turnaroundNot to waste resources for other users

d) Choose your queue



d) Choose your queue

Test

My job ...	Queue choice? (include number of nodes / cores)
<ul style="list-style-type: none">• SMIC• MPI code, needs 100 CPU cores<ul style="list-style-type: none">- Hint: SMIC has 20 cores / node	workq / checkpoint (5 nodes, 20 cores per node)
<ul style="list-style-type: none">• SuperMike 3• Uses 3 GPUs to train a neural network<ul style="list-style-type: none">- Hint: SuperMike 3 has 64 cores / node, 4 GPUs / node → 16 cores / GPU	gpu (1 node, 48 cores per node)
<ul style="list-style-type: none">• QB-3• Single-core serial code• Needs to store and process 30 GB data in RAM<ul style="list-style-type: none">- Hint: QB-3 has 192 GB RAM / node, 4 GB RAM / core	single (1 node, 8 cores per node)

e) Useful commands to check queues

- i. **qstat -q** : All queue information

```
(base) [jasonli3@mike2 ~]$ qstat -q
```

Queue	Memory	CPU	Time	Walltime	Node	Run	Que	Lm	State
admin	--	--	--	--	--	0	0	--	E R
single	--	--	--	168:00:00	1	0	0	--	E R
checkpt	--	--	--	72:00:00	--	3	0	--	E R
workq	--	--	--	72:00:00	--	12	0	--	E R
bigmem	--	--	--	72:00:00	--	0	0	--	E R
gpu	--	--	--	72:00:00	--	0	0	--	E R
						15	0		

e) Useful commands to check queues

- ii. **qfree** : Free nodes in each queue

```
(base) [jasonli3@mike2 ~]$ qfree
PBS total nodes: 183, free: 120, busy: 58, down: 2, use: 31%
PBS workq nodes: 171, free: 108, busy: 54, queued: 0
PBS single nodes: 171, free: 108, busy: 0, queued: 0
PBS checkpoint nodes: 171, free: 108, busy: 4, queued: 0
PBS bigmem nodes: 4, free: 4, busy: 0, queued: 0
PBS gpu nodes: 8, free: 8, busy: 0, queued: 0
```

e) Useful commands to check queues

iii. **sinfo** (Slurm only) : Detailed node health information of all queues

```
(base) [jasonli3@mike2 ~]$ sinfo
PARTITION AVAIL  TIMELIMIT  NODES  STATE NODELIST
single*    up 7-00:00:00      2  inval mike[035,138]
single*    up 7-00:00:00      1   comp mike144
single*    up 7-00:00:00     58  alloc mike[008-026,031-034,036-044,046-050,141-143,148-162,167-169]
single*    up 7-00:00:00    108   idle mike[001-007,027-030,045,051-137,139,145-146,163-166,170-171]
single*    up 7-00:00:00      2   down mike[140,147]
checkpt    up 3-00:00:00      2  inval mike[035,138]
checkpt    up 3-00:00:00      1   comp mike144
checkpt    up 3-00:00:00     58  alloc mike[008-026,031-034,036-044,046-050,141-143,148-162,167-169]
checkpt    up 3-00:00:00    108   idle mike[001-007,027-030,045,051-137,139,145-146,163-166,170-171]
checkpt    up 3-00:00:00      2   down mike[140,147]
workq      up 3-00:00:00      2  inval mike[035,138]
workq      up 3-00:00:00      1   comp mike144
workq      up 3-00:00:00     58  alloc mike[008-026,031-034,036-044,046-050,141-143,148-162,167-169]
workq      up 3-00:00:00    108   idle mike[001-007,027-030,045,051-137,139,145-146,163-166,170-171]
workq      up 3-00:00:00      2   down mike[140,147]
bigmem     up 3-00:00:00      4   idle mike[172-175]
gpu        up 3-00:00:00      8   idle mike[176-183]
```

1. Basic concepts

- a) How job works on clusters
- b) Job scheduler and how it works

2. Preparing my job

- a) Basic principles
 - “**large enough**” and “**small enough**”
- b) Information you need to tell job scheduler:
 - Duration
 - Number of nodes & cores
 - Job queue



- 1) Have your terminal open and ready to connect to HPC
- 2) Download our testing code (π calculation) to your /home directory
 - http://www.hpc.lsu.edu/training/weekly-materials/Downloads/pi_Jason.tar.gz
 - Hint: use *wget* command

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

3. Submitting a job

- Two types of jobs:

1) Interactive job

- Runs **in terminal** (just like using a local machine)
- **Can interact** with the job while running

2) Batch job

- Submit to server and runs **by itself**, until finished or error
- **Cannot interact** with the job while running

3. Submitting a job

- Two types of jobs:

	1) Interactive job	2) Batch job
Pros	<ul style="list-style-type: none">Can interact and monitor with job in real time	<ul style="list-style-type: none">Submit and leave it
Cons	<ul style="list-style-type: none">Waiting for human intervention is the opposite of “high performance”	<ul style="list-style-type: none">Cannot edit or interact with job while running
Ideal for	<ul style="list-style-type: none">Debugging and testingLarge compilation	<ul style="list-style-type: none">Production

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

1) Interactive job

a) Command

PBS	Slurm

1) Interactive job

a) Command

PBS	Slurm
<code>qsub -I [options]</code>	<code>srun [options] --pty <u>bash</u></code> (Or any other shell of your preference)

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre>

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre> <div>Enable X11 forwarding</div>

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre> <div>Allocation name</div>

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre>

Queue name

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre> <div>Wall time, number of nodes, number of cores</div>

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes=<# of nodes>:ppn=<# of cores PER NODE></pre> <p>Does not change with # of nodes</p>	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre> <p>Scales proportionally with # of nodes</p>

1) Interactive job

a) Command

PBS	Slurm
<pre>qsub -I \ -X \ -A <Allocation name> \ -q <Queue name> \ -l walltime=<HH:MM:SS>,nodes= nodes>:ppn=<# of cores PER NODE></pre>	<pre>run \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N <# of nodes> \ -n <# of TOTAL cores> \ --pty bash</pre>

For those who run MPI / OpenMP hybrid –

-n <# of tasks>

-c <# of cores per task>

⇒ <n> * <c> = <# of TOTAL cores>

1) Interactive job

b) Starting an interactive job

PBS

```
(base) [jasonli3@smic1 pi]$ qsub -I -A hpc_h  
n=20  
qsub: waiting for job 911565.smic3 to start  
Interactive job 911565.smic3 waiting:  
qsub: job 911565.smic3 ready
```

```
Concluding PBS prologue script - 31-Jan-2023  
-----  
(base) [jasonli3@smic045 ~]$ █
```

Slurm

```
(base) [jasonli3@mike1 pi]$ srun -A hpc_h  
srun: Job is in held state, pending sched  
srun: job 38634 queued and waiting for re  
Interactive job 38634 waiting:  
srun: job 38634 has been allocated resour  
(base) [jasonli3@mike147 pi]$ █
```

1) Interactive job

b) Starting an interactive job

PBS

```
(base) [jasonli3@smic1 pi]$ qsub -I -A hpc_h  
n=20  
qsub: waiting for job 911565.smic3 to start  
Interactive job 911565.smic3 waiting:  
qsub: job 911565.smic3 ready
```

```
Concluding PBS prologue script - 31-Jan-2023  
-----  
(base) [jasonli3@smic045 pi]$
```

Slurm

```
(base) [jasonli3@mike1 pi]$ srun -A hpc_h  
srun: Job is in held state, pending sched  
srun: job 38634 queued and waiting for re  
Interactive job 38634 waiting:  
srun: job 38634 has been allocated resour  
(base) [jasonli3@mike147 pi]$
```

Successfully started: on a computing node (**3-digit** number)

1) Interactive job

b) Starting an interactive job

PBS

```
(base) [jasonli3@smic1 pi]$ qsub -I -A hpc_h  
n=20  
qsub: waiting for job 911565.smic3 to start  
Interactive job 911565.smic3 waiting:  
qsub: job 911565.smic3 ready
```

```
Concluding PBS prologue script - 31-Jan-2023  
-----  
(base) [jasonli3@smic045 ~]$
```

Slurm

```
(base) [jasonli3@mike1 pi]$ srun -A hpc_h  
srun: Job is in held state, pending sched  
srun: job 38634 queued and waiting for re  
Interactive job 38634 waiting:  
srun: job 38634 has been allocated resour  
(base) [jasonli3@mike147 pi]$
```

PBS: Job starts in **/home** directory

Slurm: Job starts in **where the job was submitted**

c) One more thing about GPU jobs ...

PBS	Slurm
	<pre>srun \ --x11 \ -A <Allocation name> \ -p <Queue name> \ -t <HH:MM:SS> \ -N1 \ -n16 \ --gres=gpu:1 \ --pty bash</pre> <div>[GPU] Request 1 out of 4 GPUs on SuperMike 3</div>

d) Running an interactive job

- i. Serial (single-thread)
- ii. Parallel (MPI)

* **Slurm + interactive + MPI:**

\$ srun <mpi_executable>

Will hang

\$ srun **--overlap** <mpi_executable>

Will run

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

2) Batch job

- What do you need?
 - i. A **batch file** (containing job parameters and bash scripts)
 - ii. Run a **submission command** to submit this batch file

a) Batch file

PBS	Slurm

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<div>[Header] Job parameters</div> <pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <div>[Body] Commands to run after job starts</div>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <p>Shell type ("shebang")</p>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_O_WORKDIR ./pi_serial.out 100000000</pre>	<div>Allocation name</div> <pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<div>Queue name</div> <pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <div>Wall time</div>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <p>Number of nodes & cores</p>

a) Batch file

PBS ^[1]		Slurm ^[2]		Description	
#PBS -A		#SBATCH -A		Allocation name	
#PBS -q		#SBATCH -p		Queue name	
#PBS -l		#SBATCH -t		Resource request	Wall time
		#SBATCH -N			Number of nodes
		#SBATCH -n			Number of tasks
		#SBATCH -c			Number of cores per task
#PBS -o		#SBATCH -o		Standard output file	
#PBS -e		#SBATCH -e		Standard error file	
#PBS -m	a	#SBATCH --mail-type	FAIL	Send email when	Job aborts / fails
	b		BEGIN		Job begins
	e		END		Job ends
#PBS -M		#SBATCH --mail-user		Email address	
#PBS -N		#SBATCH -J		Job name	

[1] <http://www.hpc.lsu.edu/docs/pbs.php>

[2] <http://www.hpc.lsu.edu/docs/slurm.php>





2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_0_WORKDIR ./pi_serial.out 100000000</pre>	<pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p workq #SBATCH -t 1:00:00 #SBATCH -N 1 #SBATCH -n 20 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <div>[Body] Commands to run after job starts</div>

2) Batch job

a) Batch file

PBS	Slurm
<pre>#!/bin/bash #PBS -A <Allocation name> #PBS -q workq #PBS -l walltime=1:00:00 #PBS -l nodes=1:ppn=20 module load python cd \$PBS_O_WORKDIR ./pi_serial.out 100000000</pre>	<pre>#!/bin/bash #SBATCH -A <Allocation name> #SBATCH -p v #SBATCH -t 1 #SBATCH -N 1 #SBATCH -n 64 module load python cd \$SLURM_SUBMIT_DIR ./pi_serial.out 100000000</pre> <p>[Recommended] Explicitly load modules here if needed!</p>

2) Batch job

a) Batch file

PBS

```
#!/bin/bash
#PBS -A <Allocation name>
#PBS -q workq
#PBS -l walltime=1:00:00
#PBS -l nodes=1:ppn=20
```

```
module load python
```

```
cd $PBS_O_WORKDIR
./pi_serial.out 100000000
```

Slurm

```
#!/bin/bash
#SBATCH -A <Allocation name>
#SBATCH -p workq
#SBATCH -t 1:00:00
#SBATCH -N 1
#SBATCH -n 1
```

```
module load python
```

```
cd $SLURM_SUBMIT_DIR
./pi_serial.out 100000000
```

Whatever commands you need to run your jobs

2) Batch job

a) Batch file

PBS

```
#!/bin/bash
#PBS -A <Allocation name>
#PBS -q workq
#PBS -l walltime=1:00:00
#PBS -l nodes=1:ppn=20

module load python

cd $PBS_0_WORKDIR
./pi_serial.out 100000000
```

Slurm

```
#!/bin/bash
#SBATCH -A <Allocation name>
#SBATCH -p workq
#SBATCH -t 1:00:00
#SBATCH -N 1
#SBATCH -n 64

module load python

cd $SLURM_S
./pi_serial.out 100000000
```

An empty line (avoid error)



2) Batch job

b) Command

PBS	Slurm
<code>qsub <batch file name></code>	<code>sbatch <batch file name></code>

c) Useful environmental variables

PBS ^[1]	Slurm ^[2]	Description
\$PBS_JOBID	\$SLURM_JOBID	Job ID
\$PBS_O_WORKDIR	\$SLURM_SUBMIT_DIR	Job submit directory
\$PBS_NODEFILE	\$SLURM_JOB_NODELIST	A temp file, contains a list of allocated nodes' names (for MPI)
\$PBS_NUM_NODES	\$SLURM_NNODES	Number of allocated nodes
\$PBS_NP	\$SLURM_NTASKS	Number of allocated cores (tasks)
...	...	

```
#!/bin/bash
#PBS -A <Allocation name>
#PBS -q workq
#PBS -l walltime=1:00:00
#PBS -l nodes=1:ppn=20

module load python
cd $PBS_O_WORKDIR
./pt_serial.out 1000000000
```

- [1] <http://www.hpc.lsu.edu/docs/pbs.php>
 [2] <http://www.hpc.lsu.edu/docs/slurm.php>



- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

- **Running jobs on HPC \neq “Submit and done”**
 - Monitoring and managing jobs are part of the work

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

1) Useful commands

PBS ^[1]	Slurm ^[2]	Description
qstat	squeue	List all jobs

1) Useful commands

PBS ^[1]		Slurm ^[2]		Description
qstat		squeue		List all jobs
	<Job ID>		-j <Job ID>	List the job of specific ID
	-u <Username>		-u <Username>	List all jobs belong to a specific user
	<Queue name>		-p <Queue name>	List all jobs in a particular queue
qstat -n <Job ID>		scontrol show job <Job ID>		Show job details
		squeue --start		Estimated start time of queuing jobs
qdel <Job ID>		scancel <Job ID>		Cancel <Job ID>

Alter jobs after submission? → **NOT allowed!**

- **HPC User Environment 2**

1. Basic concepts
 - 1) Previously on HPC User Environment 1...
 - 2) Job & Job schedulers
2. Preparing my job
 - 1) Basic principles
 - 2) Job duration (wall time)
 - 3) Number of nodes & cores
 - 4) Job queues
3. Submitting my job
 - 1) Interactive job
 - 2) Batch job
4. Managing my jobs
 - 1) Useful commands
 - 2) Monitoring job health

A job **requesting** n cores \neq A job **utilizing** n cores

- **Goal**
 - Use the allocated resources (CPU cores, RAM, time, ...) **as fully and efficiently as possible**
 - **No serious underutilizing**
 - **No serious overutilizing**
- **Things to check**
 - CPU / GPU load
 - Memory usage

2) Monitoring job health

a) Method 1: **qshow** <Job ID>

- Displays diagnostic information of a **running job**
- Can be run on **head node**

2) Monitoring job health

a) Method 1: **qshow** <Job ID>

```
(base) [jasonli3@mike4 ~]$ qshow 38581
PBS job: 38581, nodes: 1
Hostname Days Load CPU U# (User:Process:VirtualMemory:Memory:Hours)
mike145    278 64.12 6033 68 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:533M:107M:13.5 yxan:tmp_mik+:748M:128M:13.5
yxan:tmp_mik+:738M:124M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:587M:109M:13.5 yxan:tmp_mik+:743M:128M:13.5 yxan:tmp_mik+:696M:118M:13.5
yxan:tmp_mik+:528M:101M:13.5 yxan:tmp_mik+:578M:108M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:528M:106M:13.5 yxan:tmp_mik+:520M:105M:13.5
yxan:tmp_mik+:561M:106M:13.5 yxan:tmp_mik+:583M:109M:13.5 yxan:tmp_mik+:520M:103M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:738M:125M:13.5
yxan:tmp_mik+:709M:119M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:574M:107M:13.5 yxan:tmp_mik+:697M:121M:13.5 yxan:tmp_mik+:658M:115M:13.5
yxan:tmp_mik+:528M:102M:13.5 yxan:tmp_mik+:557M:108M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:515M:102M:13.5
yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:567M:108M:13.5 yxan:tmp_mik+:566M:108M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:536M:105M:13.5
yxan:tmp_mik+:519M:104M:13.5 yxan:tmp_mik+:528M:103M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5
yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:516M:101M:13.5 yxan:tmp_mik+:515M:101M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:520M:101M:13.5
yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:520M:101M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:516M:102M:13.5 yxan:tmp_mik+:587M:110M:13.5
yxan:tmp_mik+:558M:108M:13.5 yxan:tmp_mik+:524M:102M:13.5 yxan:tmp_mik+:537M:103M:13.5 yxan:tmp_mik+:572M:109M:13.5 yxan:tmp_mik+:549M:104M:13.5
yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:103M:13.5
yxan:tmp_mik+:520M:105M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:515M:104M:13.5
yxan:slurm_s+:12M:3M yxan:srun:324M:8M yxan:srun:53M:1M
PBS_job=38581 user=yxan allocation=hpc_lipidhpre queue=checkpoint total_load=64.12 cpu_hours=866.08 wall_hours=13.21 unused_nodes=0 total_nodes=1 pp
n=64 avg_load=64.12 avg_cpu=6033% avg_mem=6852mb avg_vmem=36176mb top_proc=yxan:tmp_mik+:mike145:524M:104M:13.5hr:100% toppm=yxan:tmp_mikeCpu:mik
e145:730M:125M node_processes=68
```

What to look at ...

Normal behavior ...

You should be concerned if ...

a) Method 1: **qshow** <Job ID>

```
(base) [jasonli3@mike4 ~]$ qshow 38581
PBS job: 38581, nodes: 1
Hostname Days Load CPU U# (User:Process:VirtualMemory:Memory:Hours)
mike145    278 64.12 6033 68 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:533M:107M:13.5 yxan:tmp_mik+:748M:128M:13.5
yxan:tmp_mik+:738M:124M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:587M:109M:13.5 yxan:tmp_mik+:743M:128M:13.5 yxan:tmp_mik+:696M:118M:13.5
yxan:tmp_mik+:528M:101M:13.5 yxan:tmp_mik+:578M:108M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:528M:106M:13.5 yxan:tmp_mik+:520M:105M:13.5
yxan:tmp_mik+:561M:106M:13.5 yxan:tmp_mik+:583M:109M:13.5 yxan:tmp_mik+:520M:103M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:738M:125M:13.5
yxan:tmp_mik+:709M:119M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:574M:107M:13.5 yxan:tmp_mik+:697M:121M:13.5 yxan:tmp_mik+:658M:115M:13.5
yxan:tmp_mik+:528M:102M:13.5 yxan:tmp_mik+:557M:108M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:515M:102M:13.5
yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:567M:108M:13.5 yxan:tmp_mik+:566M:108M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:536M:105M:13.5
yxan:tmp_mik+:519M:104M:13.5 yxan:tmp_mik+:528M:103M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5
yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:516M:101M:13.5 yxan:tmp_mik+:515M:101M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:520M:101M:13.5
yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:520M:101M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:516M:102M:13.5 yxan:tmp_mik+:587M:110M:13.5
yxan:tmp_mik+:558M:108M:13.5 yxan:tmp_mik+:524M:102M:13.5 yxan:tmp_mik+:537M:103M:13.5 yxan:tmp_mik+:572M:109M:13.5 yxan:tmp_mik+:549M:104M:13.5
yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:103M:13.5
yxan:tmp_mik+:520M:105M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:515M:104M:13.5
yxan:slurm_s+:12M:3M yxan:srun:324M:8M yxan:srun:53M:1M
PBS job: 38581 user=yxan allocation=hpc_lipidhpre queue=checkpoint total_load=64.12 cpu_hours=866.08 wall_hours=13.21 unused_nodes=0 total_nodes=1 pp
n=64 avg_load=64.12 avg_cpu=6033% avg_mem=6852mb avg_vmem=36176mb top_proc=yxan:tmp_mik+:mike145:524M:104M:13.5hr:100% toppm=yxan:tmp_mikeCpu:mik
e145:738M:125M_node_processes=68
```

What to look at ...	Normal behavior ...	You should be concerned if ...
avg_load	Close to allocated number of cores on the node	Consistently too low or too high

2) Monitoring job health

a) Method 1: **qshow** <Job ID>

```
(base) [jasonli3@mike4 ~]$ qshow 38581
PBS job: 38581, nodes: 1
Hostname Days Load CPU U# (User:Process:VirtualMemory:Memory:Hours)
mike145    278 64.12 6033 68 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:533M:107M:13.5 yxan:tmp_mik+:748M:128M:13.5
yxan:tmp_mik+:738M:124M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:587M:109M:13.5 yxan:tmp_mik+:743M:128M:13.5 yxan:tmp_mik+:696M:118M:13.5
yxan:tmp_mik+:528M:101M:13.5 yxan:tmp_mik+:578M:108M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:528M:106M:13.5 yxan:tmp_mik+:520M:105M:13.5
yxan:tmp_mik+:561M:106M:13.5 yxan:tmp_mik+:583M:109M:13.5 yxan:tmp_mik+:520M:103M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:738M:125M:13.5
yxan:tmp_mik+:709M:119M:13.5 yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:574M:107M:13.5 yxan:tmp_mik+:697M:121M:13.5 yxan:tmp_mik+:658M:115M:13.5
yxan:tmp_mik+:528M:102M:13.5 yxan:tmp_mik+:557M:108M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:524M:105M:13.5 yxan:tmp_mik+:515M:102M:13.5
yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:567M:108M:13.5 yxan:tmp_mik+:566M:108M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:536M:105M:13.5
yxan:tmp_mik+:519M:104M:13.5 yxan:tmp_mik+:528M:103M:13.5 yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:524M:104M:13.5 yxan:tmp_mik+:524M:104M:13.5
yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:516M:101M:13.5 yxan:tmp_mik+:515M:101M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:520M:101M:13.5
yxan:tmp_mik+:524M:103M:13.5 yxan:tmp_mik+:520M:101M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:516M:102M:13.5 yxan:tmp_mik+:587M:110M:13.5
yxan:tmp_mik+:558M:108M:13.5 yxan:tmp_mik+:524M:102M:13.5 yxan:tmp_mik+:537M:103M:13.5 yxan:tmp_mik+:572M:109M:13.5 yxan:tmp_mik+:549M:104M:13.5
yxan:tmp_mik+:519M:103M:13.5 yxan:tmp_mik+:528M:104M:13.5 yxan:tmp_mik+:520M:104M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:103M:13.5
yxan:tmp_mik+:520M:105M:13.5 yxan:tmp_mik+:528M:105M:13.5 yxan:tmp_mik+:515M:103M:13.5 yxan:tmp_mik+:515M:104M:13.5 yxan:tmp_mik+:515M:104M:13.5
yxan:slurm_s+:12M:3M yxan:srun:324M:8M yxan:srun:53M:1M
PBS_job=38581 user=yxan allocation=64.12 top_cpus=68 queue=checkpoint total_load=64.12 cpu_hours=866.08 wall_hours=13.21 unused_nodes=0 total_nodes=1 pp
n=64 avg_load=64.12 avg_cpu=603% avg_mem=6852mb avg_vmem=36176mb top_proc=yxan:tmp_mik+:mike145:524M:104M:13.5hr:100% toppm=yxan:tmp_mikeCpu:mik
e145:730M:125M node_processes=68
```

What to look at ...	Normal behavior ...	You should be concerned if ...
avg_load	Close to allocated number of cores on the node	Consistently too low or too high
ave_mem	Does not exceed total allocated memory	Exceeds total allocated memory

b) Method 2: **top**

- Displays dynamic real-time view of a **computing node**
- Must run on **computing nodes** !
 - * ssh to computing nodes while job running (cannot ssh if you do not have jobs on it)

2) Monitoring job health

b) Method 2: **top**

```
top - 02:23:58 up 278 days, 19:17,  2 users,  load average: 63.63, 39.81, 17.49
Tasks: 981 total,  65 running, 916 sleeping,   0 stopped,   0 zombie
%Cpu(s): 90.2 us,  9.2 sy,  0.0 ni,  0.0 id,  0.0 wa,  0.5 hi,  0.0 si,  0.0 st
MiB Mem : 257004.8 total, 211261.0 free,  41926.9 used,   3816.9 buff/cache
MiB Swap: 16641.0 total,  16580.7 free,    60.2 used. 212737.8 avail Mem

  PID USER      PR  NI   VIRT   RES   SHR  S  %CPU  %MEM     TIME+ COMMAND
2701318 jasonli3  20   0 595668 582356 2568  R 100.0   0.2   4:08.94 TDSE_np3_e0
2701342 jasonli3  20   0 595668 581944 2616  R 100.0   0.2   4:08.90 TDSE_np3_e0
2701249 jasonli3  20   0 595668 581792 2464  R  99.7   0.2   4:08.97 TDSE_np3_e0
2701252 jasonli3  20   0 595668 514684 2520  R  99.7   0.2   4:09.00 TDSE_np3_e0
2701261 jasonli3  20   0 595668 393828 2616  R  99.7   0.1   4:08.97 TDSE_np3_e0
2701264 jasonli3  20   0 595668 581856 2532  R  99.7   0.2   4:08.92 TDSE_np3_e0
2701270 jasonli3  20   0 595668 582480 2432  R  99.7   0.2   4:08.95 TDSE_np3_e0
2701273 jasonli3  20   0 595668 581776 2448  R  99.7   0.2   4:08.81 TDSE_np3_e0
2701276 jasonli3  20   0 595668 582160 2568  R  99.7   0.2   4:08.98 TDSE_np3_e0
2701279 jasonli3  20   0 595668 222064 2644  R  99.7   0.1   4:08.98 TDSE_np3_e0
```

What to look at ...

Normal behavior ...

You should be concerned if ...

b) Method 2: **top**

```
top - 02:23:58 up 278 days, 19:17, 2 users, load average: 63.63, 39.81, 17.49
Tasks: 981 total, 65 running, 916 sleeping, 0 stopped, 0 zombie
%Cpu(s): 90.2 us, 9.2 sy, 0.0 ni, 0.0 id, 0.0 wa, 0.5 hi, 0.0 si, 0.0 st
MiB Mem : 257004.8 total, 211261.0 free, 41926.9 used, 3816.9 buff/cache
MiB Swap: 16641.0 total, 16580.7 free, 60.2 used. 212737.8 avail Mem

  PID USER      PR  NI   VIRT   RES   SHR  S  %CPU  %MEM     TIME+ COMMAND
2701318 jasonli3  20   0 595668 582356 2568  R 100.0   0.2   4:08.94 TDSE_np3_e0
2701342 jasonli3  20   0 595668 581944 2616  R 100.0   0.2   4:08.90 TDSE_np3_e0
2701249 jasonli3  20   0 595668 581792 2464  R  99.7   0.2   4:08.97 TDSE_np3_e0
2701252 jasonli3  20   0 595668 514684 2520  R  99.7   0.2   4:09.00 TDSE_np3_e0
2701261 jasonli3  20   0 595668 393828 2616  R  99.7   0.1   4:08.97 TDSE_np3_e0
2701264 jasonli3  20   0 595668 581856 2532  R  99.7   0.2   4:08.92 TDSE_np3_e0
2701270 jasonli3  20   0 595668 582480 2432  R  99.7   0.2   4:08.95 TDSE_np3_e0
2701273 jasonli3  20   0 595668 581776 2448  R  99.7   0.2   4:08.81 TDSE_np3_e0
2701276 jasonli3  20   0 595668 582160 2568  R  99.7   0.2   4:08.98 TDSE_np3_e0
2701279 jasonli3  20   0 595668 222064 2644  R  99.7   0.1   4:08.98 TDSE_np3_e0
```

What to look at ...	Normal behavior ...	You should be concerned if ...
Load average	Close to allocated number of cores on the node	Consistently too low or too high

2) Monitoring job health

b) Method 2: **top**

```
top - 02:23:58 up 278 days, 19:17, 2 users, load average: 63.63, 39.81, 17.49
Tasks: 981 total, 65 running, 916 sleeping, 0 stopped, 0 zombie
%Cpu(s): 90.2 us, 9.2 sy, 0.0 ni, 0.0 id, 0.0 wa, 0.5 hi, 0.0 si, 0.0 st
MiB Mem : 257004.8 total, 211261.0 free, 41926.9 used, 3816.9 buff/cache
MiB Swap: 16641.0 total, 16580.7 free, 60.2 used. 212737.8 avail Mem
```

PID	USER	PR	NI	VIRT	RES	SHR	S	%CPU	%MEM	TIME+	COMMAND
2701318	jasonli3	20	0	595668	582356	2568	R	100.0	0.2	4:08.94	TDSE_np3_e0
2701342	jasonli3	20	0	595668	581944	2616	R	100.0	0.2	4:08.90	TDSE_np3_e0
2701249	jasonli3	20	0	595668	581792	2464	R	99.7	0.2	4:08.97	TDSE_np3_e0
2701252	jasonli3	20	0	595668	514684	2520	R	99.7	0.2	4:09.00	TDSE_np3_e0
2701261	jasonli3	20	0	595668	393828	2616	R	99.7	0.1	4:08.97	TDSE_np3_e0
2701264	jasonli3	20	0	595668	581856	2532	R	99.7	0.2	4:08.92	TDSE_np3_e0
2701270	jasonli3	20	0	595668	582480	2432	R	99.7	0.2	4:08.95	TDSE_np3_e0
2701273	jasonli3	20	0	595668	581776	2448	R	99.7	0.2	4:08.81	TDSE_np3_e0
2701276	jasonli3	20	0	595668	582160	2568	R	99.7	0.2	4:08.98	TDSE_np3_e0
2701279	jasonli3	20	0	595668	222064	2644	R	99.7	0.1	4:08.98	TDSE_np3_e0

What to look at ...	Normal behavior ...	You should be concerned if ...
Load average	Close to allocated number of cores on the node	Consistently too low or too high
Memory usage (not virtual memory)	Does not exceed total allocated memory	Exceeds total allocated memory

2) Monitoring job health

c) Method 3: **nvidia-smi** (for GPU only)

```
(base) [jasonli3@qbc193 ~]$ nvidia-smi
Wed Feb  1 02:38:32 2023
```

NVIDIA-SMI 510.47.03 Driver Version: 510.47.03 CUDA Version: 11.6							
GPU	Name	Persistence-M	Bus-Id	Disp.A	Volatile	Uncorr. ECC	
Fan	Temp	Perf	Pwr:Usage/Cap	Memory-Usage	GPU-Util	Compute M.	MIG M.
0	Tesla V100-PCIE...	On	00000000:3B:00.0	Off		Off	
N/A	36C	P0	54W / 250W	4155MiB / 32768MiB	72%	Default	N/A
1	Tesla V100-PCIE...	On	00000000:AF:00.0	Off		Off	
N/A	36C	P0	52W / 250W	4155MiB / 32768MiB	78%	Default	N/A

Processes:							
GPU	GI	CI	PID	Type	Process name	GPU Memory	
	ID	ID				Usage	
0	N/A	N/A	259491	C	... che/TeraChem/bin/terachem	4147MiB	
1	N/A	N/A	259491	C	... che/TeraChem/bin/terachem	4147MiB	

What to look at ...	Normal behavior ...	You should be concerned if ...
---------------------	---------------------	--------------------------------

c) Method 3: **nvidia-smi** (for GPU only)

```
(base) [jasonli3@qbc193 ~]$ nvidia-smi
Wed Feb 1 02:38:32 2023

+-----+
| NVIDIA-SMI 510.47.03      Driver Version: 510.47.03      CUDA Version: 11.6      |
+-----+-----+
| GPU  Name      Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan  Temp  Perf  Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M. |
|====+=====+====+=====+=====+=====+=====+=====+
|  0  Tesla V100-PCIE...    On   | 00000000:3B:00.0 Off |   72%   Default  |
| N/A   36C    P0      54W / 250W | 4155MiB / 32768MiB |           N/A      |
+-----+-----+
|  1  Tesla V100-PCIE...    On   | 00000000:AF:00.0 Off |   78%   Default  |
| N/A   36C    P0      52W / 250W | 4155MiB / 32768MiB |           N/A      |
+-----+-----+

+-----+
| Processes: |
| GPU   GI    CI          PID    Type    Process name                        GPU Memory |
|      ID    ID                                   |            Usage |
|====+=====+=====+=====+=====+=====+=====+
|  0   N/A   N/A       259491      C   ... che/TeraChem/bin/terachem      4147MiB |
|  1   N/A   N/A       259491      C   ... che/TeraChem/bin/terachem      4147MiB |
+-----+-----+
```

What to look at ...	Normal behavior ...	You should be concerned if ...
GPU usage	Close to 100%	Consistently too low

2) Monitoring job health

c) Method 3: **nvidia-smi** (for GPU only)

```
(base) [jasonli3@qbc193 ~]$ nvidia-smi
Wed Feb  1 02:38:32 2023

+-----+
| NVIDIA-SMI 510.47.03      Driver Version: 510.47.03      CUDA Version: 11.6     |
+-----+-----+
| GPU   Name               Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan  Temp  Perf    Pwr:Usage/Cap     |      Memory-Usage | GPU-Util  Compute M. |
|=====+=====+
|  0   Tesla V100-PCIE...    On          | 00000000:3B:00.0 Off |   4155MiB / 32768MiB | 72%      Default  |
| N/A   36C   P0      54W / 250W        |                   |              N/A     |
+-----+-----+
|  1   Tesla V100-PCIE...    On          | 00000000:AF:00.0 Off |   4155MiB / 32768MiB | 78%      Default  |
| N/A   36C   P0      52W / 250W        |                   |              N/A     |
+-----+-----+

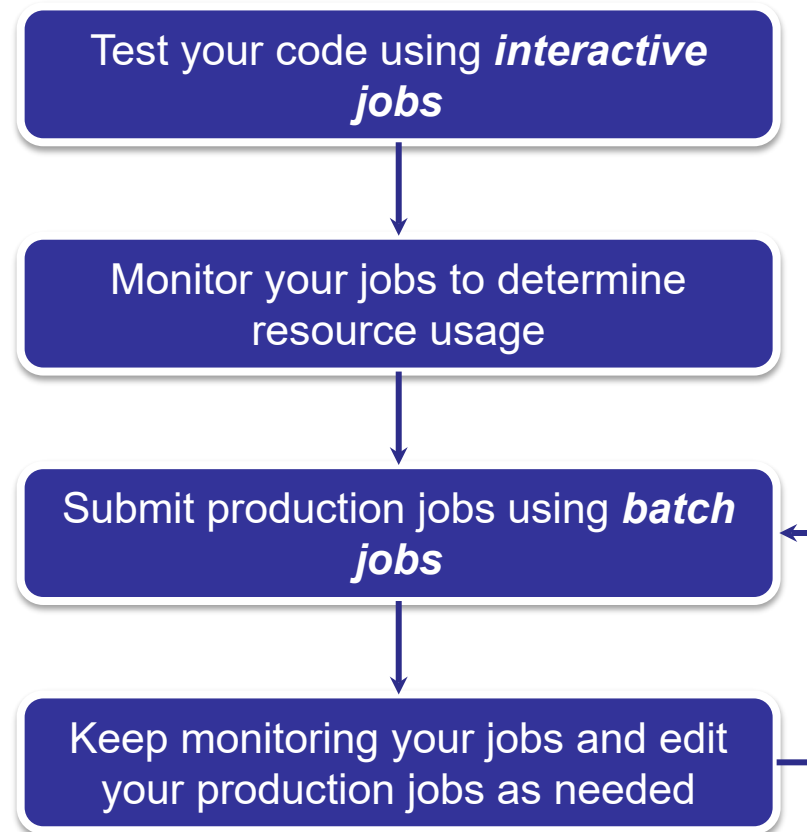
Processes:
+-----+-----+
| GPU   GI    CI          PID    Type    Process name                        GPU Memory |
|   ID   ID    ID                                  Usage    |
+-----+-----+
|  0   N/A   N/A      259491     C     ... che/TeraChem/bin/terachem      4147MiB |
|  1   N/A   N/A      259491     C     ... che/TeraChem/bin/terachem      4147MiB |
+-----+-----+
```

What to look at ...	Normal behavior ...	You should be concerned if ...
GPU usage	Close to 100%	Consistently too low
Memory usage (not virtual memory)	Not used up	Used up

d) Common issues

Issue	What would happen
Exceeded memory allocation (e.g., using more memory than allocated w/ single queue)	Terminated. Receive email notice.
Exceeded ppn/core allocation (e.g., using more cores than allocated w/ single queue)	Terminated. Receive email notice.
Seriously underutilize node CPU cores / unused nodes (e.g., Requested multiple nodes but only runs on one node)	Receive email warning. (* Killed if completely idle for a long time)
Submitting to bigmem but only using little memory	Receive email warning.
Running intensive calculation on head nodes	Terminated. Receive email notice.
Submitting too many (i.e., hundreds of) single-thread jobs	Poor parallelization and bad for server. We may reach out to you to help. (Better yet, reach out to us first)

- A typical workflow --



■ HPC User Environment 2

1. Basic concepts

1) Previously on HPC User Environment 1...

2) Job & Job schedulers → **All calculation must be submitted as jobs**

2. Preparing my job

1) Basic principles → **Large enough & small enough**

2) Job duration (wall time)

3) Number of nodes & cores

4) Job queues

3. Submitting my job

1) Interactive job → **Good for testing and debugging**

2) Batch job → **Good for production**

4. Managing my jobs

1) Useful commands

2) Monitoring job health → **How to monitor jobs health, and how to create health jobs**



Next week

- **Basic Shell Scripting**

- **Contact user services**

- Email Help Ticket: sys-help@loni.org
- Telephone Help Desk: +1 (225) 578-0900