

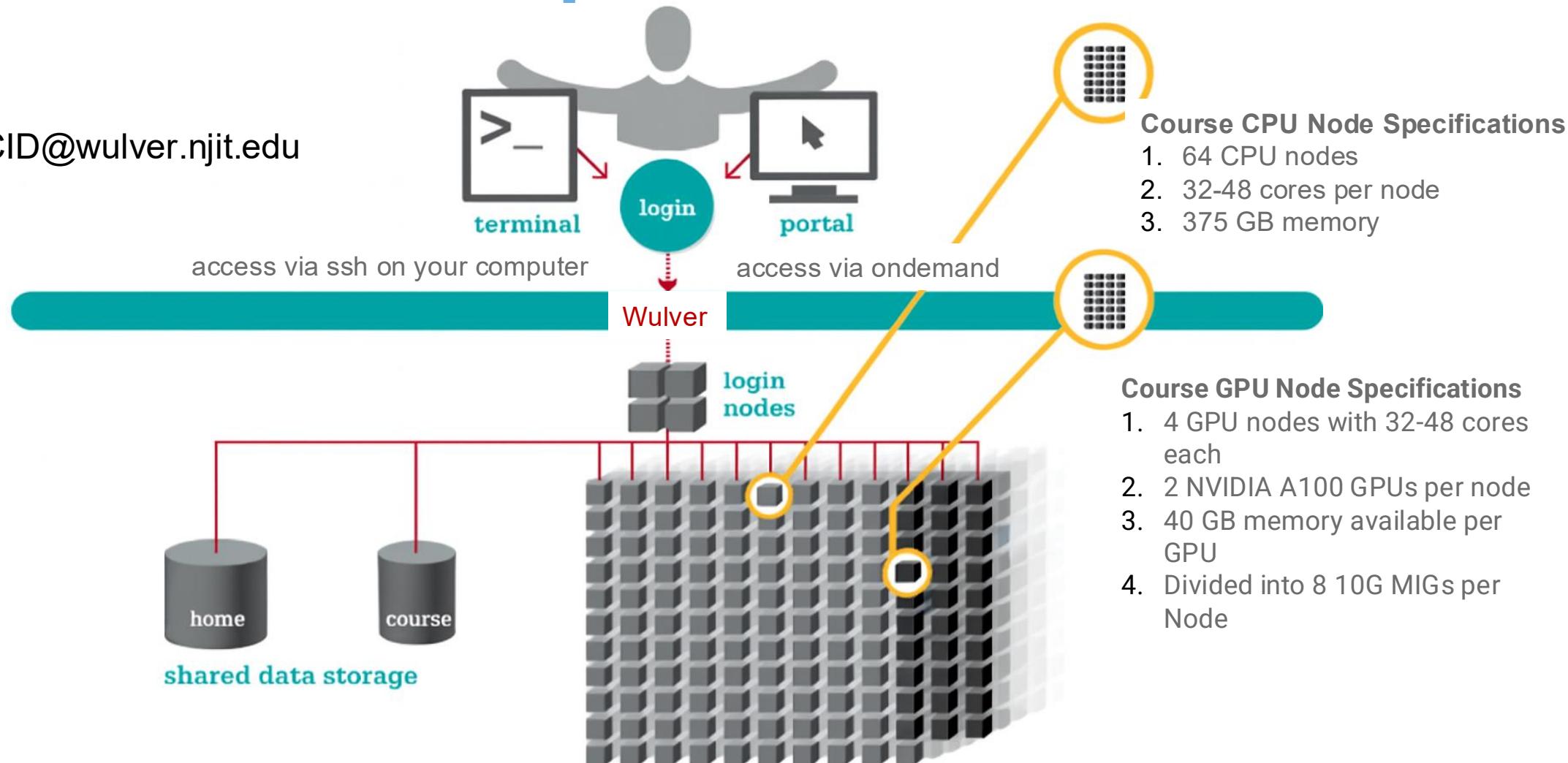


High Performance Computing



Wulver Cluster Specifications

ssh \$UCID@wulver.njit.edu

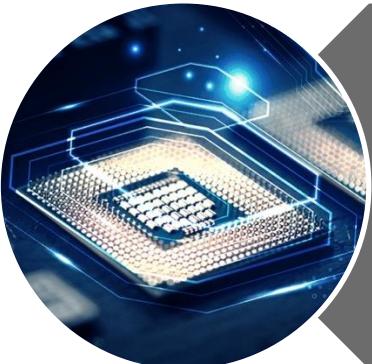


HPC Allocations



Storage

- Home (~50GB/user) - Limited quota: not intended for primary storage
- Course (/course/2026) - Primary storage

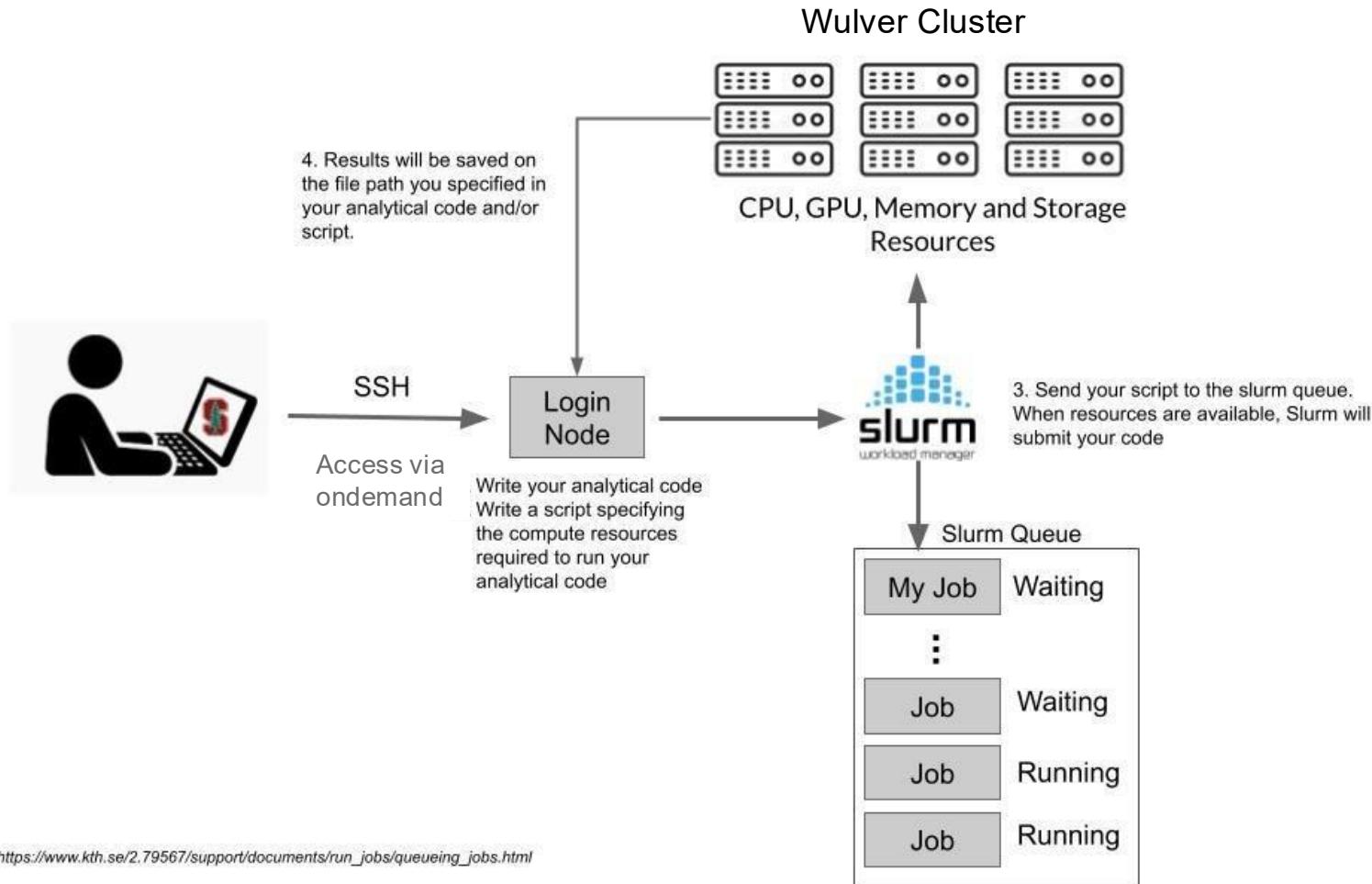


Computing Time

- 1 SU = Number of CPUs x Walltime in hours x usage factor
- SU allocation – 2500 SUs per user

Use `quota_info` to check the allocations

Why do supercomputers use queuing?



Manage Jobs – Options

Mandatory Options

Directive	Options	Description
--account=	account	Check your email or use quota_info
--partition=	<u>Partition</u>	Request a partition of resources for job allocation (queue)
--time=	<u>Time</u> [[d-]h:]m[:s]	Minimum time limit on job allocation
--qos=	Job Priorities	Define the job priority. Use --qos=course

see <https://slurm.schedmd.com/srun.html> for more details

HPC Partitions

- Example of SU charges: (20 cores with 1 MIG for 8 hours without --mem)
 - $SU = (20 + 2) * 8 = 176$
- Example of SU charges: (20 cores with 1 MIG for 8 hours with --mem = 128G)
 - $SU = [2 + \text{MAX}(20, 128/4=32)] * 8 = (2 + 32) * 8 = 34 * 8 = 272$

Partition	Nodes	Cores/ Node	GPU	Memory	SU charge
--partition=course	64	32-48	NA	375 GB	MAX(CPUs, Memory/4G) SU
--partition=course_gpu	4	32-48	A100 10G MIG	375 GB	2 + MAX(CPUs, Memory/4G) SU

Job Submission Time Interval Formats

- Valid time formats (with a few exceptions) for `-t` / `--time=` option

Minutes	(<code>-t 10</code> is 10 minutes)
Minutes:Seconds	(... <code>10:30</code> is 10 minutes & 30 secs)
Hours:Minutes:Seconds	(<code>1:0:0</code> is 1 hr + 0mins + 0secs)
Days-Hours:Minutes:Seconds	(<code>7-1:10:30</code> is 7days + 1hr + 10mins + 30secs)
Days-Hours	(<code>7-0</code> is 7days + 0hrs i.e. 7 days)
Days-Hours:Minutes	(<code>7-4:10</code> is 7days + 4hrs + 10mins)

- Wall time maximum - 72 hours

Manage Jobs – Options

Additional Options

Directive	Options	Description
--ntasks=	Number of cpus	Number of CPUs (tasks) to be allocated
--mem=	Memory	Total memory of the job
--gres=	Generic resources	Set the Number of gpus
--error=	File	Define standard error file
--out=	File	Define standard output file

Directive	Options	Description
--input=	File	File used for standard input
--job	Name	Define job name
--mail-type=	ALL, BEGIN, END, FAIL, REQUEUE	Notify user by email when <type> event occurs
--mail-user=	Email address	Send email to this address for events specified with mail-type option (default is submitting user).

see <https://slurm.schedmd.com/srun.html> for more details

General Application Workflow

- Log into cluster with ucid and password
- Copy input files to new directory
- Change to copied directory via command line
`cd /path/to/copied_directory`
- Copy job a template to the directory
`cp /path/to/templates/jobtemplate.job jobfile.job`
- Modify the job file:
 - Change the number of resources to desired number
 - Change the module load command based on the application name and version
 - Update command line with commands required for job
 - Update the software modules
- Submit the job file using `sbatch`

Sample Simple Job Script

```
#!/bin/bash

#SBATCH --job-name=my_job
#SBATCH --partition=course
#SBATCH --output=%x.%j.out
#SBATCH --error=%x.%j.err
#SBATCH --account=ACCOUNT
#SBATCH --qos=course
#SBATCH --time=00:20:00
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=1
#SBATCH --mail-type=ALL
#SBATCH -mail-user=ab1234@njit.edu

date
sleep 60
date
```



Job setup information
for SLURM

Commands to be run

- This runs a batch job called “my_job” to the “course” partition, with 1 task, a wall time limit of 20 minutes.
- QOS is required. Account is recommended.
- `%x.%j` expands to `JobName.JobID` and prints this into a text file

Sample MPI Job script

```
#!/bin/bash

#SBATCH --job-name=mpi_test_job
#SBATCH --partition=course
#SBATCH --output=%x.%j.out
#SBATCH --error=%x.%j.err
#SBATCH --account=ACCOUNT
#SBATCH --qos=course
#SBATCH --time=00:10:00
#SBATCH --ntasks=64

# Run application commands
srun /apps/testjobs/bin/mpihello
```

- This runs an MPI job named “mpi_test_job”, with 64 processes total. Wall time is 10 minutes.

Sample GPU Job script

```
#!/bin/bash

#SBATCH --job-name=test_gpu_job
#SBATCH --output=%x.%j.out
#SBATCH --error=%x.%j.err
#SBATCH --partition=course_gpu
#SBATCH --account=PI_UCID
#SBATCH --qos=standard
#SBATCH --time=00:20:00
#SBATCH --ntasks=2
#SBATCH --gres=gpu:a100_10g:1

# Load application environment
module load CUDA

# Run application commands
nvidia-smi
```

- This runs a GPU job named “test_gpu_job”, with 2 CPUs and 1 MIG. Wall time is 20 minutes.

Check sample job scripts in /apps/testjobs/workshop



Limitation of GPU Jobs

- You cannot run your job using multiple MIG instances.
- For example, `--gres=gpu:a100_10g:2` will allocate two instances of the 10G MIG, but it will either raise an error, or some jobs may assume it as a single MIG, even if multiple instances are requested.

Manage Jobs – Submit via CLI

Submit a job script

- **\$ sbatch my_script**
- Submitted batch job 1234

Listing jobs

- For current user in Pending, Running, Suspended states:
• **\$ squeue -u \$LOGNAME**

JOBID	PARTITION	NAME	USER	ST	TIME	NODES	NODELIST (REASON)
1234	general	uname.sh	test	PD	0:00	2	(Priority)

For a more detailed query
on active job:

- **\$ scontrol show jobid=1234**

```
JobId=2 JobName=simple.job
UserId=test(1001) GroupId=test(1001) MCS_label=N/A
Priority=4294901759 Nice=0 Account=(null) QOS=standard
JobState=COMPLETED Reason=None Dependency=(null)
...
```

Canceling jobs

- **\$ scancel 1234** - Cancel job ID 1234
- **scancel --me** - Cancel all your jobs

Show information about an
active or completed job

- **\$ slurm_jobid 1234**

Common inquiries

checkq

- squeue but more details

slurm_jobid

- Show information about a running or queued job

sq

- Display pending job/queue info in a helpful way,
You can also check the last job details with sq

squeue --start

- Jobs will be listed in order expected start time
- Times are only estimates and subject to change

quota_info

- Show space and SU quotas for self or others

Some Common Problems

After using sbatch, the job disappears in 30 seconds and there's no result output.

- Check the details with slurm_jobid [JOBID], use --err and --out
- Use sq if you are unsure about the job id.

Invalid account or account/partition combination specified.

- Check --account
- Use quota_info \$LOGNAME

JOBID	PARTITION	NAME	USER	ST	TIME	NODES	NODELIST (REASON)
1234	course	uname.sh	xiss	PD	0:00	2	(ReqNodeNotAvail, Reserved for maintenance)

- Jobs that do not end before the maintenance window begins will be held until the maintenance is complete

Some Common Problems(contd)

```
JOBID PARTITION      NAME      USER ST TIME   NODES  
NODELIST (REASON)  
1234       course  uname.sh  xiss  PD 0:00      2  
(AssocGrpBillingMinutes)
```

- You have reached the limit of SU.

Error message: cggroup out of memory handler

- Check the memory requirement
- You probably need to increase memory on --mem or the number of cpus from --mem-per-cpu
- If nothing works, then the problem is likely due to incorrect setup of the problem.

Using Applications with GUI on OnDemand

Login to ondemand.njit.edu

Go to “Interactive Apps” and select the application from the list.

If you don’t find the app, select Linux Desktop

Once are you connected, select “Terminal Emulator” from “Applications” option from top left

Interactive Batch Jobs



Interactive, but handled through batch system

Resource limits same as standard batch limits



Useful for tasks forbidden on login nodes

Debug parallel programs
Quickly test your code



May not be practical when system load is high

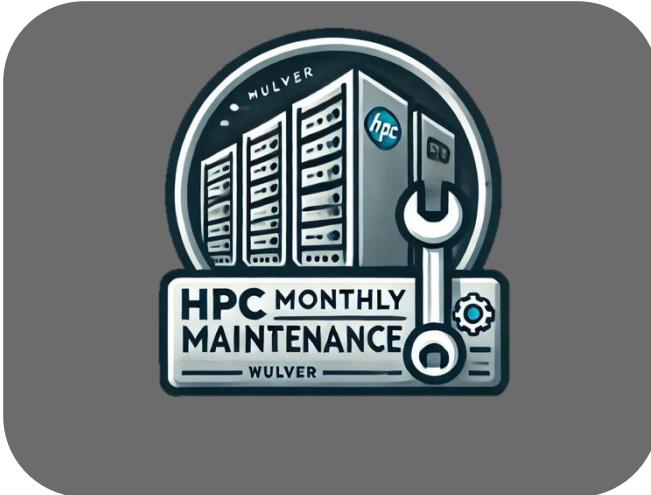
Long wait, same as standard batch job



To submit an interactive batch job (example)

- `interactive -a ACCOUNT -q QOS -p PARTITION -j JOB_TYPE`
- `interactive -h` for details

Reminder



- Wulver will be temporarily out of service for maintenance once a month, specifically on the 2nd Tuesday, to perform updates, repairs, and upgrades.
- During the maintenance period, the logins will be disabled
- Jobs that do not end before the maintenance window begins will be held until the maintenance is complete
 - Reduce the walltime in the job script to run the job



- Date: Every Monday and Wednesday
- Time: 2:00–4:00 p.m.
- Location: GITC 5320N
- Meet with our student consultants and ask any questions you have about using HPC resources.
- There's no need to create a ticket in advance.

Resources to get your questions answered

Getting Started: [Access to Wulver](#)

List of Software: [Wulver Software](#)

HOW TOs: [Conda Documentation](#)

Installing Python packages via Conda

Request Software: [HPC Software Installation](#)

Access to OnDemand [Open OnDemand](#)

MIG Information [MIG](#)

Contact: Please visit [HPC Contact](#)

Open a ticket: email to hpc@njit.edu

Consult with Research Computing Facilitator: [Facilitator Calendar Appointment](#)

System updates

- Read Message of the Day on login
- Visit [NJIT HPC News](#)





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