



# Getting Started on the FASRC clusters with Command Line Interface

# Learning objectives

- Log in via `ssh` to Cannon and FASSE
- How to start an interactive job with `salloc`
- How to submit a batch job with `sbatch`
- Check job status
- Cluster software modules

# Login to Cannon and FASSE – ssh

Documentation: <https://docs.rc.fas.harvard.edu/kb/terminal-access/>



Mac: Terminal, iTerm2



Linux: Xterm or Terminal

Windows



SSH client: Putty



Bash emulator: Git bash

## Cannon

```
$ ssh jharvard@login.rc.fas.harvard.edu
Password:
Verification code:
```

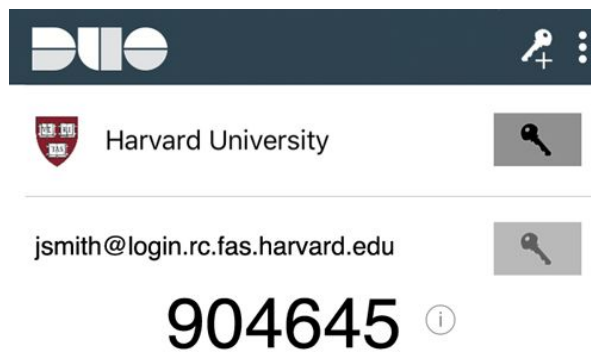
## FASSE

```
$ ssh jharvard@fasselogin.rc.fas.harvard.edu
Password:
Verification code:
```

# Login to Cannon and FASSE – 2 factor authentication

- Execute the ssh command, then:
  - Type your password (*cursor won't move!*), press enter
  - Type the 6-digit verification code (2-Factor Authentication)
    - Separate from HarvardKey
    - Updates token every 30 seconds
    - You can only use a token once

Java desktop app



# Login to Cannon and FASSE – at login node (1)

## Cannon

## FASSE

```

jharvard@local-machine $ ssh jharvard@login.rc.fas.harvard.edu
(jharvard@login.rc.fas.harvard.edu) Password:
(jharvard@login.rc.fas.harvard.edu) verificationCode:
Last failed login: Mon Sep 18 18:54:44 EDT 2023 from 132.248.81.29 on ssh:notty
There were 3 failed login attempts since the last successful login.
Last login: Tue Aug 22 12:57:45 2023
!!!!!!!!!!!!!!!!!!!! Cannon Cluster !!!!!!!!!!!!!!!!!!!!!
Welcome to Cannon, a secure HPC resource for the research community,
hosted by Research Computing at HU's Faculty of Arts and Sciences.

+----- Helpful Documentation: -----+
| https://docs.rc.fas.harvard.edu/kb/quickstart-guide/ |
| https://docs.rc.fas.harvard.edu/kb/running-jobs/ |
| https://docs.rc.fas.harvard.edu/kb/convenient-slurm-commands/ |
+-----+

+----- NEWS & UPDATES: -----+
+ OFFICE HOURS: Wednesdays noon-3pm, see website for details +
+ Check our training schedule at: https://www.rc.fas.harvard.edu/upcoming-training/ +
+-----+

NEXT MAINTENANCE: OCTOBER 2ND 7-11AM
https://www.rc.fas.harvard.edu/maintenance

ROCKY 8: Welcome to the new operating system! For more on the
changes on the cluster see:
https://docs.rc.fas.harvard.edu/kb/rocky-8-transition-guide/

OFFICE HOURS: Are held on Zoom from 12-3PM EST on Wednesdays.
See https://www.rc.fas.harvard.edu/training/office-hours/ for details.
[jharvard@holylgin03 ~]$

```

```

jharvard@local-machine $ ssh jharvard@fasselgin.rc.fas.harvard.edu
(jharvard@fasselgin.rc.fas.harvard.edu) Password:
(jharvard@fasselgin.rc.fas.harvard.edu) VerificationCode:
Last login: Tue Jun 20 11:02:27 2023
!!!!!!!!!!!!!!!!!!!! FASSE Cluster !!!!!!!!!!!!!!!!!!!!!
Welcome to FASSE, a secure HPC resource for the research community,
hosted by Research Computing at HU's Faculty of Arts and Sciences.

+----- Helpful Documentation: -----+
| https://docs.rc.fas.harvard.edu/kb/quickstart-guide/ |
| https://docs.rc.fas.harvard.edu/kb/running-jobs/ |
| https://docs.rc.fas.harvard.edu/kb/convenient-slurm-commands/ |
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OFFICE HOURS: Are held on Zoom from 12-3PM EST on Wednesdays.
See https://www.rc.fas.harvard.edu/training/office-hours/ for details.
[jharvard@fasselgin01 ~]$

```

# Login to Cannon and FASSE – at login node (2)

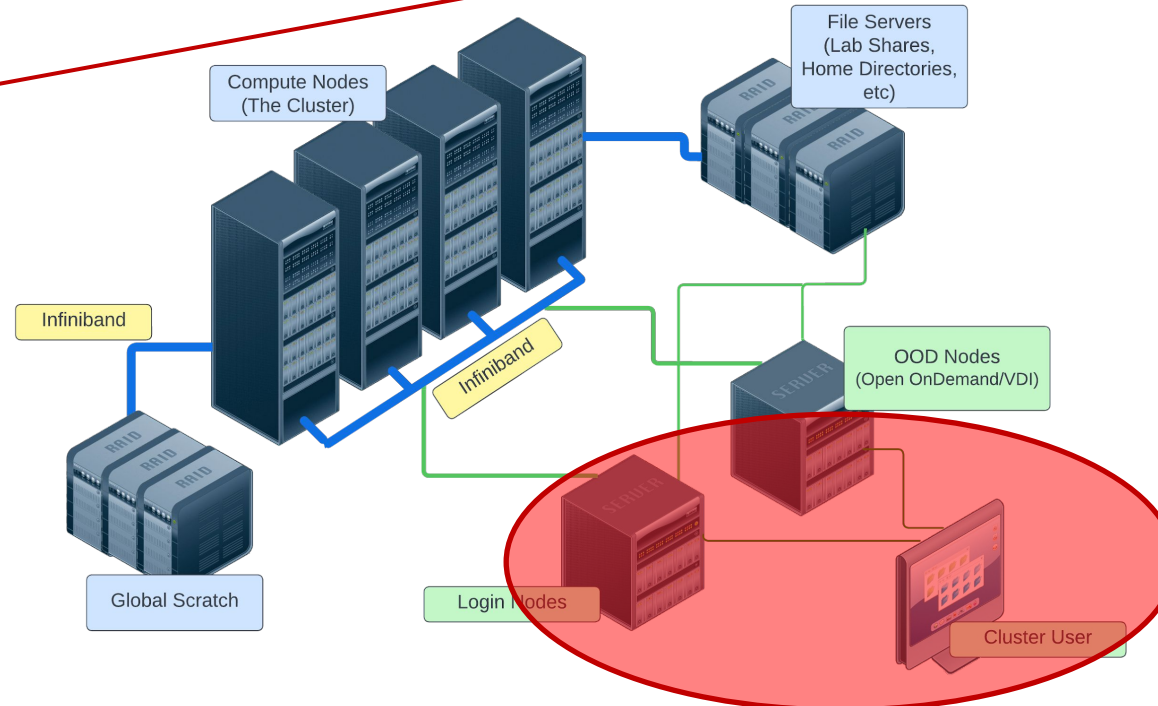
## Cannon

```
[jharvard@boslogin01 ~]$
```

## FASSE

```
[jharvard@fassellogin01 ~]$
```

Name of the login node  
assigned to you



# Login vs. compute nodes

- Login nodes
  - limited to 1 core and 4G of memory
  - not designed for analysis
  - not anything compute- or memory-intensive
  - best practice is to request a compute node as soon as you login
- Compute node via interactive job
  - work a compute node interactively – testing, debugging, installing software
  - request resources from slurm using `salloc` command
  - session will only last as long as the network connection is active
  - cannot be idle for more than 1h, session will freeze

# Interactive job on Cannon (1)

## Requesting an interactive job

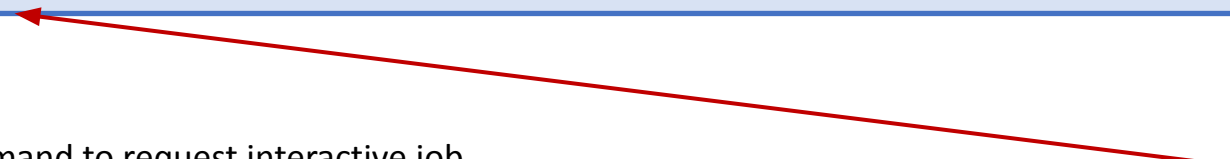
```
[jharvard@boslogin01 ~]$ salloc --partition test --mem-per-cpu 1G --time 01:00:00
salloc: Pending job allocation 2741096
salloc: job 2741096 queued and waiting for resources
salloc: job 2741096 has been allocated resources
salloc: Granted job allocation 2741096
salloc: Nodes holy7c02410 are ready for job
[jharvard@holy7c02410 ~]$
```

`salloc` - slurm command to request interactive job

`--partition test` - requesting a compute node in a specific partition

`--mem-per-cpu 1G` - memory requested in GB (if no unit is specified, the default is MB)

`--time 00:01:00` - time requested (1 hour, format HH:MM:SS or D-HH:MM)



Name of the compute  
node assigned to you

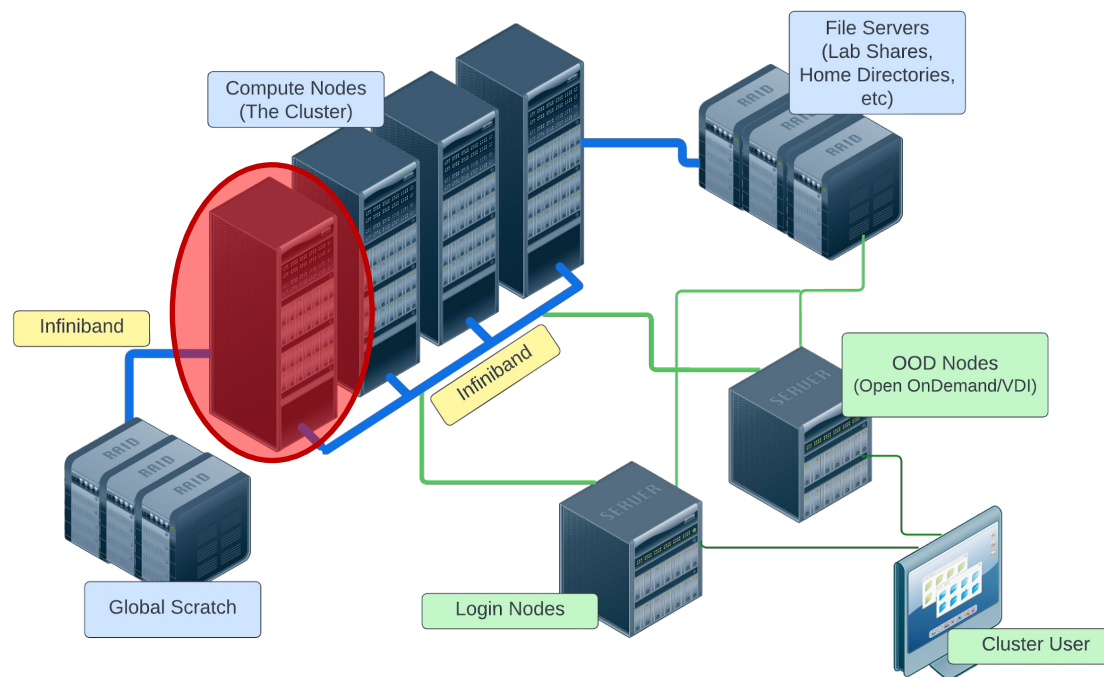


# Interactive job on Cannon (2)

Requesting an interactive job

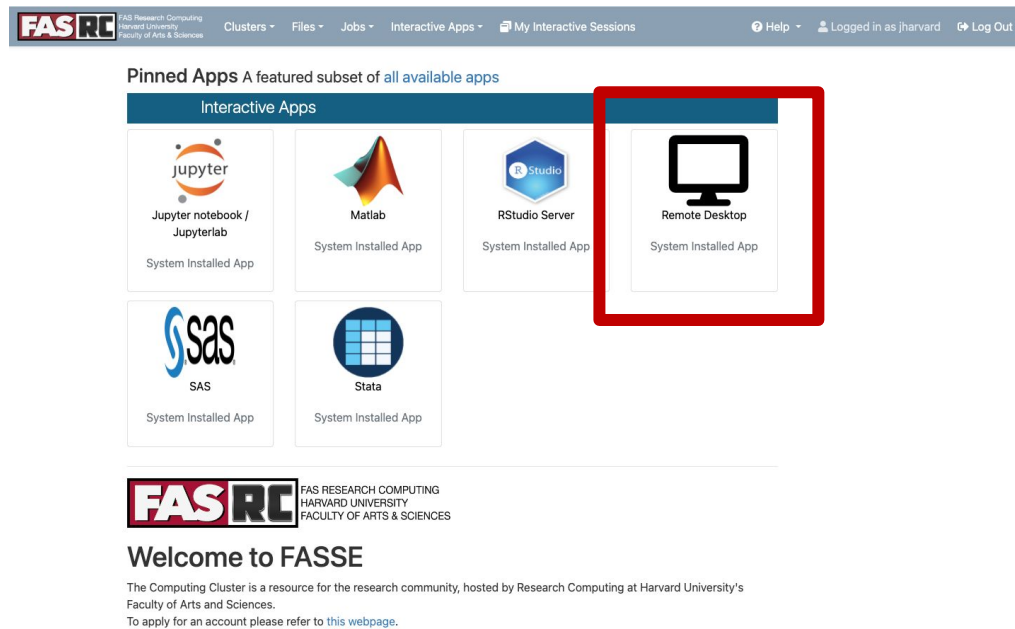
```
[jharvard@holly7c02410 ~]$
```

Name of the compute  
node assigned to you



# Interactive job on FASSE

- You cannot request an interactive job on FASSE
- You must use Remote Desktop app on Open OnDemand <https://fasseood.rc.fas.harvard.edu> and launch terminal



The screenshot shows the FAS RC Open OnDemand interface. The top navigation bar includes the FAS RC logo, user information (Logged in as jharvard), and a Log Out button. Below the navigation bar, there is a section titled "Pinned Apps A featured subset of all available apps". Underneath this, there is a sub-section for "Interactive Apps" containing six app tiles: Jupyter notebook / Jupyterlab, Matlab, RStudio Server, Remote Desktop (highlighted with a red box), SAS, and Stata. Each tile includes the app's logo, name, and "System Installed App" status. At the bottom of the page, there is a "Welcome to FASSE" message and a link to the account application page.

# Batch job

Documentation:

<https://docs.rc.fas.harvard.edu/kb/running-jobs/>

- Automate job
- No interaction
- Can close your terminal/laptop and job will keep running
- Partitions
  - Cannon:  
<https://docs.rc.fas.harvard.edu/kb/running-jobs/>
  - FASSE:  
<https://docs.rc.fas.harvard.edu/kb/fasse/>

slurm script `runscript.sh`

slurm directives

```
#!/bin/bash
#SBATCH -J py_job           # Job name
#SBATCH -p test            # Partition(s) (separate with
                           # commas if using multiple)
#SBATCH -c 1              # Number of cores
#SBATCH -t 0-00:30:00     # Time (D-HH:MM:SS)
#SBATCH --mem=500M        # Memory
#SBATCH -o py_%j.o        # Name of standard output
                           # file
#SBATCH -e py_%j.e        # Name of standard error file

# load software environment
module load python/3.10.12-fasrc01

# print a statement
echo "This is our test slurm script"

# execute python code
python hello_world.py
```

# Test first!!

ALWAYS test the job submission script first:

- To ensure the job will complete without errors
- To ensure you understand the resource needs and have requested them appropriately

## Submitting a batch job

```
[jharvard@boslogin01 python]$ sbatch runscript.sh  
Submitted batch job 2742999  
[jharvard@boslogin01 python]$
```

# Job monitoring – sacct

## Documentation:

- o `sacct`: slurm accounting database
  - every 30 sec the node collects the amount of cpu and memory usage that all of the process ID are using for a given job. After the job ends this data is sent to slurm database
- o Common flags (i.e., options)
  - `-j jobid` or `--name=jobname`
  - `-S starttime YYYY-MM-DD` and `-E endtime YYYY-MM-DD`
  - `-o output_options`
  - See slurm docs for more options: <https://slurm.schedmd.com/sacct.html>

```
[jharvard@boslogin01 ~]$ sacct --format=JobID,Jobname,partition,state,time,start,end,elapsed,MaxRss,MaxVMSize,nnodes,ncpus,nodelist --units=G -j 2742999
```

JobID	JobName	Partition	State	Timelimit	Start	End	Elapsed	MaxRSS	MaxVMSize	NNodes	NCPUS	NodeList
2742999	py_job	test	COMPLETED	00:30:00	2023-09-21T12:03:20	2023-09-21T12:03:21	00:00:01			1	1	holy7c02410
2742999.bat+	batch		COMPLETED		2023-09-21T12:03:20	2023-09-21T12:03:21	00:00:01	0.01G	0.21G	1	1	holy7c02410
2742999.ext+	extern		COMPLETED		2023-09-21T12:03:20	2023-09-21T12:03:21	00:00:01	0.00G	0.17G	1	1	holy7c02410

# Memory usage

1. Run a test batch job
2. Check memory usage after the job has completed (with `sacct` command)

```
[jharvard@boslogin01 ~]$ sacct -j 2742999 -o ReqMem,MaxRSS
  ReqMem      MaxRSS
-----
    500M
          7512K
          4348K
[jharvard@boslogin01 ~]$ sacct -j 2742999 -o ReqMem,MaxRSS --units=G
  ReqMem      MaxRSS
-----
    0.49G
          0.01G
          0.00G
```

# Job efficiency summary – `seff`

1. Run a test batch job
2. Check job efficiency after the job has completed (with `seff` command)

```
[jharvard@boslogin01 ~]$ seff 2742999
Job ID: 2742999
Cluster: odyssey
User/Group: jharvard/jharvard_lab
State: COMPLETED (exit code 0)
Cores: 1
CPU Utilized: 00:00:00
CPU Efficiency: 0.00% of 00:00:01 core-walltime
Job Wall-clock time: 00:00:01
Memory Utilized: 7.34 MB
Memory Efficiency: 1.47% of 500.00 MB
```

```
[user@boslogin01 home]# seff 1234567
Job ID: 1234567
Cluster: odyssey
User/Group: user/user_lab
State: COMPLETED (exit code 0)
Nodes: 8
Cores per node: 64
CPU Utilized: 37-06:17:33
CPU Efficiency: 23.94% of 155-16:02:08 core-walltime
Job Wall-clock time: 07:17:49
Memory Utilized: 1.53 TB (estimated maximum)
Memory Efficiency: 100.03% of 1.53 TB (195.31
GB/node)
```

# Partitions

`spart` allows you to see which partitions you have access to

Documentation: <https://docs.rc.fas.harvard.edu/kb/convenient-slurm-commands/>

```
[jharvard@boslogin02 ~]$ spart
```

Partition	State	Cores	GPUs	Average Mem/Node (GB)	Nodes	Time Limit
bigmem	UP	448	0	2015	4	3-00:00:00
bigmem_intermediate	UP	192	0	2015	3	14-00:00:00
gpu	UP	2304	144	1007	36	3-00:00:00
gpu_requeue	UP	9184	698	772	156	3-00:00:00
gpu_test	UP	896	112	503	14	12:00:00
intermediate	UP	1344	0	1007	12	14-00:00:00
remoteviz	UP	32	0	377	1	3-00:00:00
sapphire	UP	21504	0	1007	192	3-00:00:00
serial_requeue	UP	88300	690	438	1457	3-00:00:00
shared	UP	13824	0	188	288	3-00:00:00
test	UP	1344	0	1007	12	12:00:00
ultramem	DRAIN	192	0	2015	3	3-00:00:00

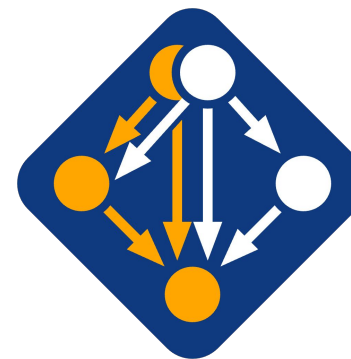


# Software – LMOD module system

- Software is loaded incrementally using modules, to set up your shell environment (e.g., `PATH`, `LD_LIBRARY_PATH`, and other environment variables)
- Why add `module load` commands in a slurm batch script? (instead of `.bashrc` file)
  - Keeps your interactive working environment simple
  - Is a record of your research workflow (reproducible research!)
  - Keep `.bashrc` module loads sparse, lest you run into software and library conflicts

```
module load matlab/R2022b-fasrc01 # recommended command
module load matlab                # loads most recent version
module list                       # show loaded modules
module purge                      # unload all loaded modules
module spider matlab              # search for modules with matlab in the name
module display matlab/R2022b-fasrc01 # show the details of the module
```

# Spack



- For software that does not have a module, you can install it with Spack:  
<https://docs.rc.fas.harvard.edu/kb/spack/>
- Install Spack in a Holyoke storage location, such as `holylabs`
  - Package installation is best done in an interactive session with 8 cores 12GB as Spack needs more resources  
`salloc --partition test --time 0-04:00 --mem 12G --cpus-per-task 8`

# Survey

Please, fill out our course survey. Your feedback is essential for us to improve our trainings!!

<http://tinyurl.com/FASRCsurvey>

# FASRC documentation

- FASRC docs: <https://docs.rc.fas.harvard.edu/>
- GitHub User\_codes: [https://github.com/fasrc/User\\_Codes/](https://github.com/fasrc/User_Codes/)
- Getting help
  - Office hours: <https://www.rc.fas.harvard.edu/training/office-hours/>
  - Ticket
    - Portal: [http://portal.rc.fas.harvard.edu/rcrt/submit\\_ticket](http://portal.rc.fas.harvard.edu/rcrt/submit_ticket) (requires login)
    - Email: [rchelp@rc.fas.harvard.edu](mailto:rchelp@rc.fas.harvard.edu)

# Upcoming training sessions

Training calendar: <https://www.rc.fas.harvard.edu/upcoming-training/>

## Getting started on the FASRC clusters with Open OnDemand

- Audience
  - New users not familiar with command-line interface
  - Wants to use a GUI
- Requirements
  - Single-node jobs
  - Working FASRC account with cluster access
- Content
  - Access Open OnDemand
  - Launch Jupyter, Rstudio Server, Remote Desktop
  - Install Rstudio Server packages
  - Install python packages for Jupyter
  - Launch software from Remote Desktop



**Thank you :)**  
**FAS Research Computing**