





Getting Started on the FASRC clusters with Open OnDemand





Where to access the slides

https://docs.rc.fas.harvard.edu/kb/training-materials/





Learning objectives

- o What is Open OnDemand (OOD)?
- Accessing OOD
- Launching apps
- RStudio Server
- Jupyter Notebook
 - Create conda environment (i.e., jupyter kernel)
- Files tab
- Jobs tab
- Remote Desktop
- FASSE proxy





Some definitions

- o OOD: Open On Demand
- o FASRC: Faculty of Arts and Sciences Research Computing
- o Cluster: large group of servers with lots of memory and processors
- o **Cannon**: cluster that handles level 2 data. Named after the 19th century Harvard astronomer Annie Jump Cannon.
- o FASSE: cluster that handles level 3 data. FAS Secure Enclave.
 - Glossary of these terms: docs.rc.fas.harvard.edu/kb/glossary





What is Open OnDemand (OOD)?

- Open-source web portal to access clusters
- Web-based
 - Uses modern browser like Google Chrome, Mozilla Firefox, or Microsoft Edge.
 - Safari does not support all of OOD's features
 - No software other than a browser needs be installed on your local laptop/desktop
- Easy to learn and use
- Very similar to desktop applications
- The easiest way to run graphical applications remotely on a cluster







How to access OOD on FASRC Clusters

- 1. Get an account
 - You can choose a username. The default is first initial, last name. I'll use jharvard as an example
 - This is NOT necessarily the same as your HarvardKey username.
- 2. Log onto the FASRC VPN. This is NOT the generic Harvard VPN.
 - vpn.rc.fas.harvard.edu
 - username is
 - jharvard@fasrc (Cannon)
 - jharvard@fasse (FASSE)
- 3. Log into Open On Demand for your cluster
 - username is just jharvard, nothing else

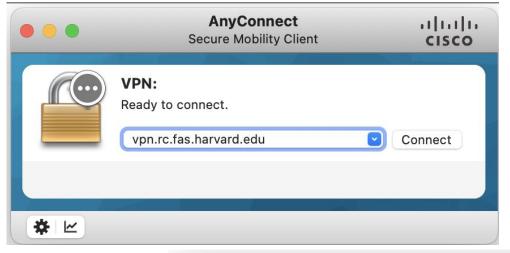
Cannon link: https://rcood.rc.fas.harvard.edu

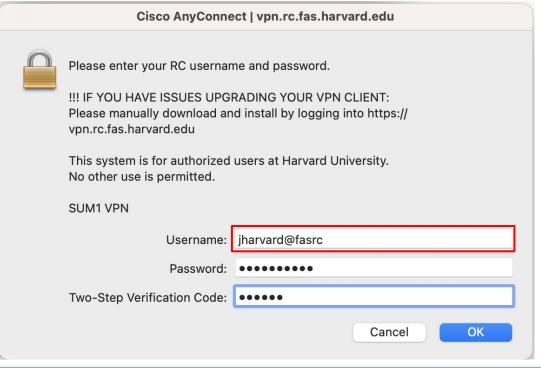
FASSE link: https://fasseood.rc.fas.harvard.edu

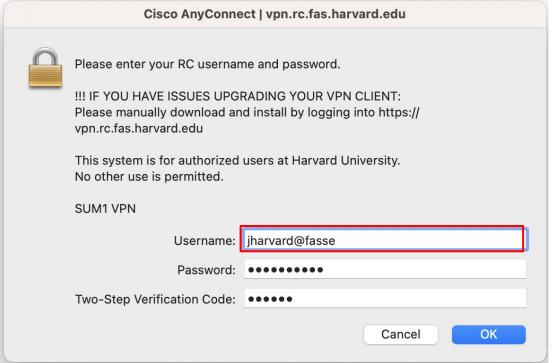




Connecting to VPN











Signing in to the OOD Dashboard

When you load the site

https://rcood.rc.fas.harvard.edu

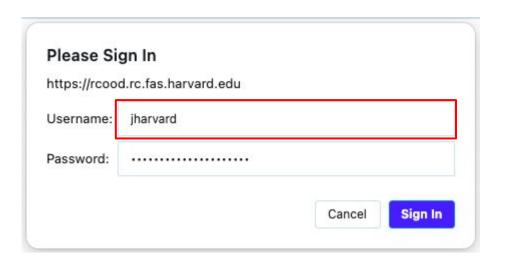
or

https://fasseood.rc.fas.harvard.edu

You will be prompted to log in.

Unlike when you log in to the VPN, on the website you need to use

ONLY your username, NOT username@cluster



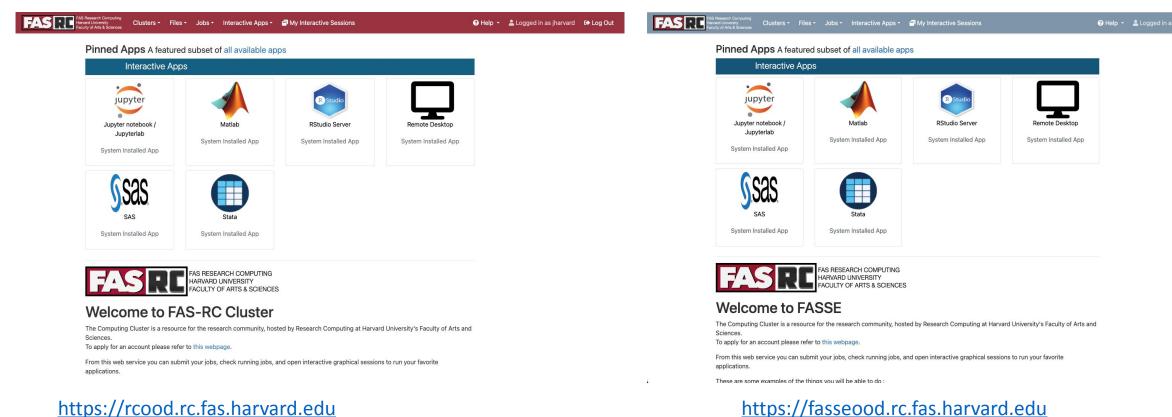




OOD dashboard on Cannon and FASSE

Cannon

FASSE



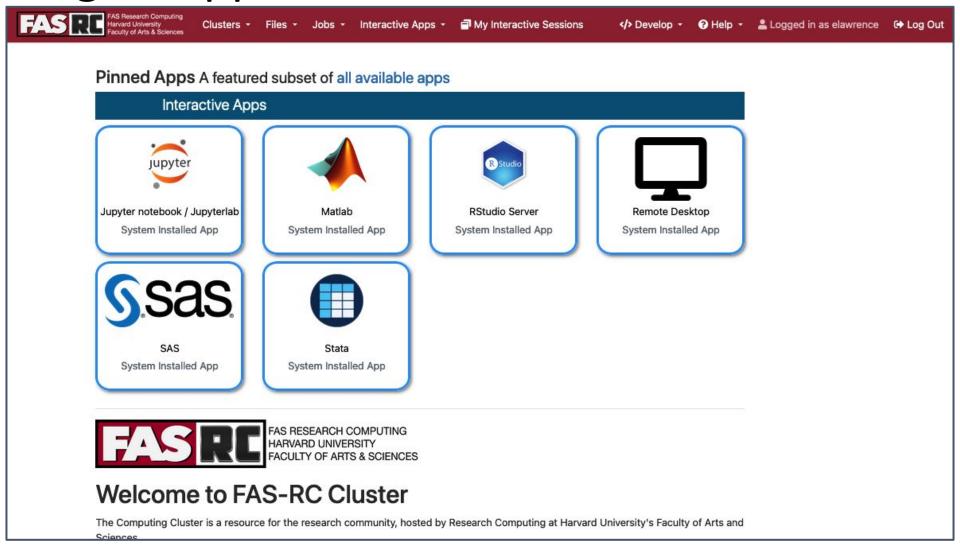
https://rcood.rc.fas.harvard.edu

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Launching an app from the Dashboard







Beginning of a form

RStudio Server

This app will launch an RStudio Server instance on a FAS RC compute node. This app provides a common software environment for FAS Informatics workshops and general-purpose single-node RStudio Server jobs.

See RStudio Server OOD app for more information.

 User-installed R libraries will be installed in ~/R/ifxrstudio/<IMAGE_TAG>.

Partition

test

sbatch -p, --partition=<partition_names>

Slurm partition name (e.g., **shared**), or comma-separated list of partition names (e.g., **shared,test**)

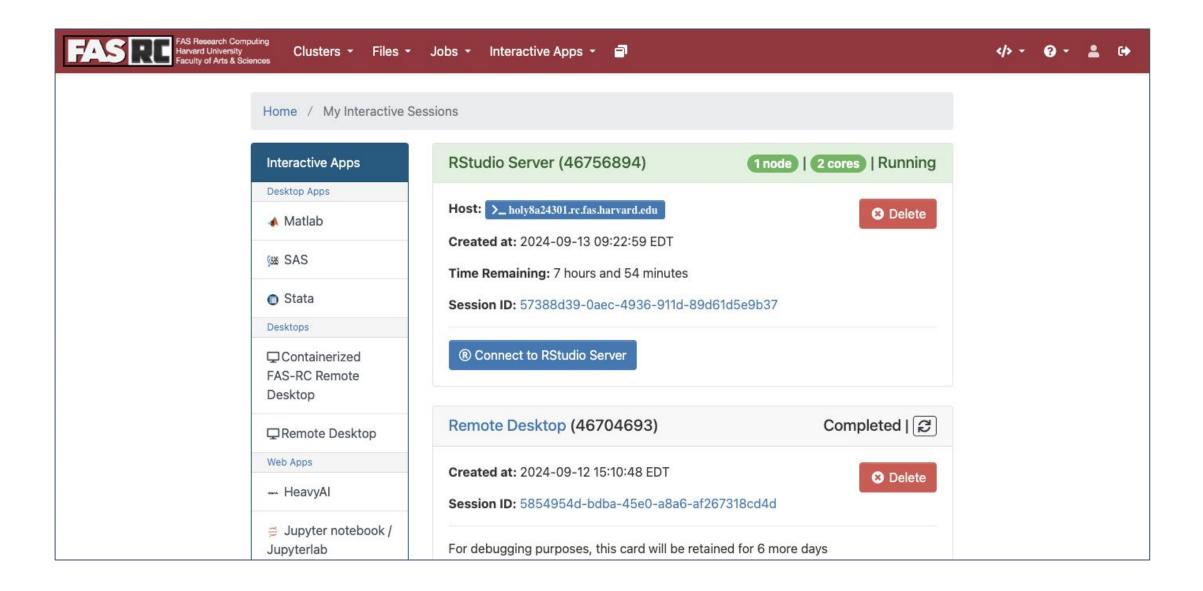
Memory Allocation in GB

8

sbatch --mem=<size>G











Jupyter Notebook (1)

- You can create your own kernels, but some command line needed
- Note: a kernel is the same as a conda, python, or mamba environment
- Open a terminal in the "Remote Desktop" app
 ⇒Don't create mamba environments inside Jupyter Notebook/Lab!
- 2. Create mamba environment and install package ipykernel and nb_conda_kernels
- [jharvard@holy7c02111 ~]\$ module load python
 [jharvard@holy7c02111 ~]\$ mamba create -n OOD_env python=3.11 pip wheel numpy
 [jharvard@holy7c02111 ~]\$ mamba activate OOD_env

 (OOD_env)[jharvard@holy7c02111 ~]\$ mamba install ipykernel nb_conda_kernels

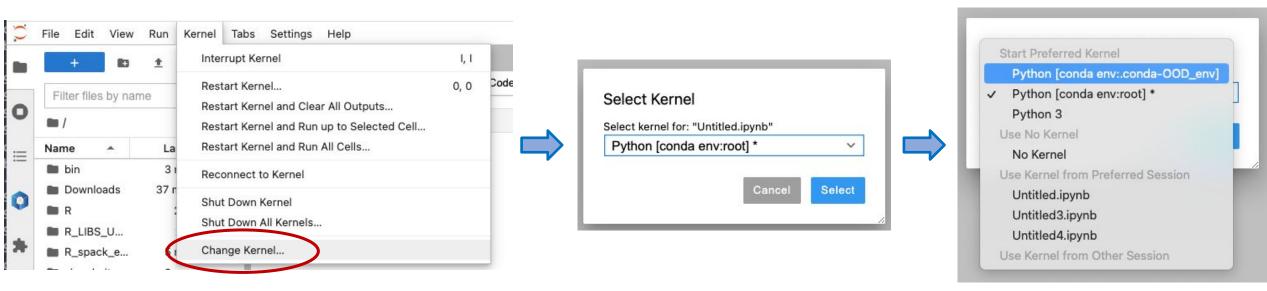
https://docs.rc.fas.harvard.edu/kb/python-package-installation/#Use_mamba_environme nt_in_Jupyter_Notebooks





Jupyter Notebook (2)

- 3. Launch new Jupyter Notebook session (existing session will not work!)
- 4. Select newly created mamba environment as the kernel
 - a. Open a notebook
 - b. On the top menu, click Kernel -> Select Kernel -> Click on OOD_env

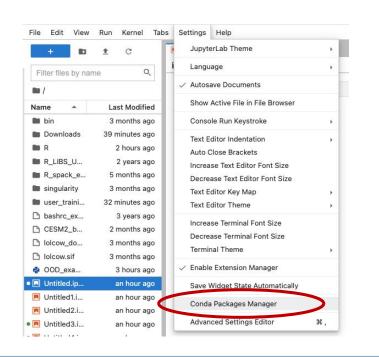


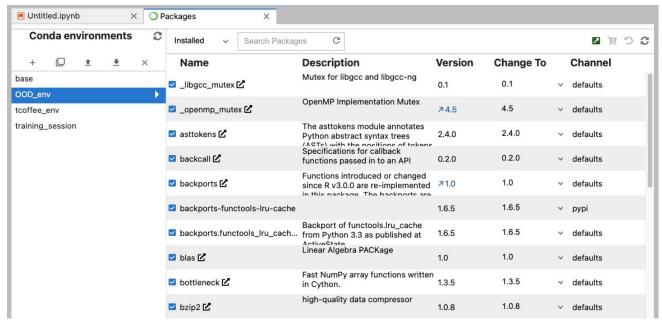




Jupyter Notebook (3)

- 5. Managing (install, uninstall, update) packages
 - a. We recommend using the command line
 - b. You can also use the conda package manager, which is the same thing as a mamba package manager: On the top menu, click Settings -> Conda Package Manager -> OOD_env



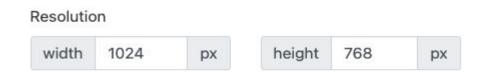






Remote Desktop

- Not as necessary as it used to be
- For running long Jupyter Notebook sessions
- Terminal
- Can also be used to open multiple applications in a single window
- Choose the defaults for resolution

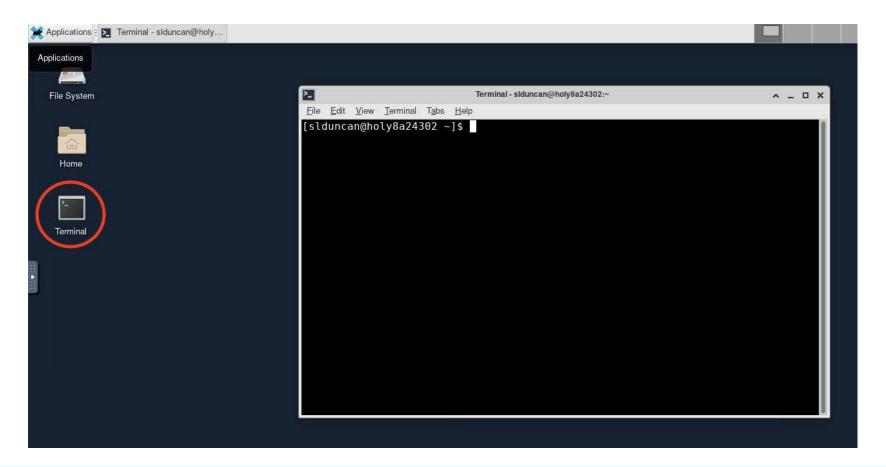






Starting a terminal

- Double click on the icon that looks like a computer screen
- From there you can type in any commands you want to







Running Jupyter Notebook in Remote Desktop

- Very important: Jupyter Notebook will not continue to run if you close the Jupyter notebook page! The cell that is running will lose the data and output files will not be written
 - 1. Solution: run Remote Desktop app and launch Jupyter Notebook from within Remote Desktop
 - 2. Documentation:

https://docs.rc.fas.harvard.edu/kb/ood-remote-desktop-how-to-open-software/#Jupyter Notebook

```
#Jupyter Notebook
[jharvard@holy7c02111 ~]$ module load python
[jharvard@holy7c02111 ~]$ mamba activate OOD_env
[jharvard@holy7c02111 ~]$ jupyter notebook
```





Opening Multiple Applications in Remote Desktop

Documentation: https://docs.rc.fas.harvard.edu/kb/ood-remote-desktop-how-to-open-software/

- It can be used to launch most GUI applications
 - Load module
 - 2. Set environmental variables (if needed)
 - Launch software
- You can have multiple applications open

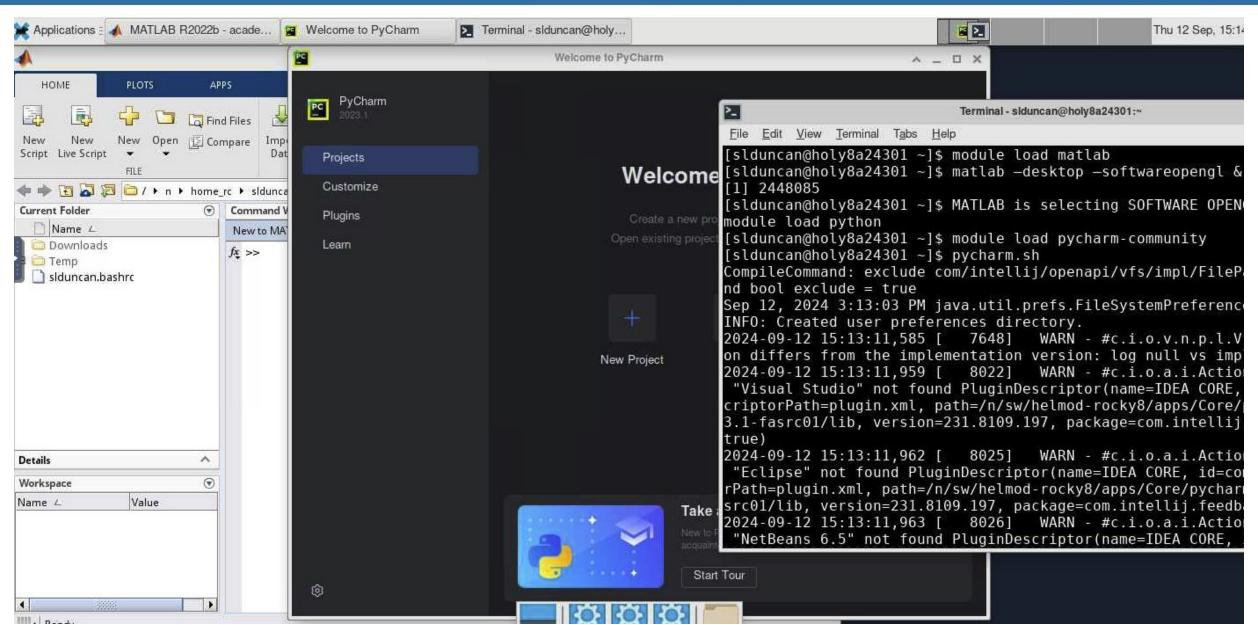
```
# Matlab
[jharvard@holy7c02111 ~]$ module load matlab
[jharvard@holy7c02111 ~]$ matlab -desktop -softwareopengl &

#PyCharm
[jharvard@holy7c02111 ~]$ module load python
[jharvard@holy7c02111 ~]$ module load pycharm-community
[jharvard@holy7c02111 ~]$ pycharm.sh
```





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Inactivity lock out

- It may lock out due to inactivity
- Use your FASRC password to unlock







Filling out a form to launch an app

- Request the resources that you need
 (If you don't know for a first trial run, use similar resources as your laptop/desktop)
 - Partition (Name): depends on <u>Cannon</u> vs <u>FASSE</u>
 - Memory (RAM): amount of memory in GB
 - Number of cores: recommended at least 2
 - Number of GPUs: if >= 1, make sure you **select** a gpu partition
 - Allocated time: time you would like your session to run.
 - Email for status notification: to know when job starts, ends
 - Reservation: if you have a special reservation (this requires approval from FASRC)
 - Account: use this if you have more than one PI_lab affiliation

the minimum and/or maximum values of each field depends on the selected partition





Menu bar

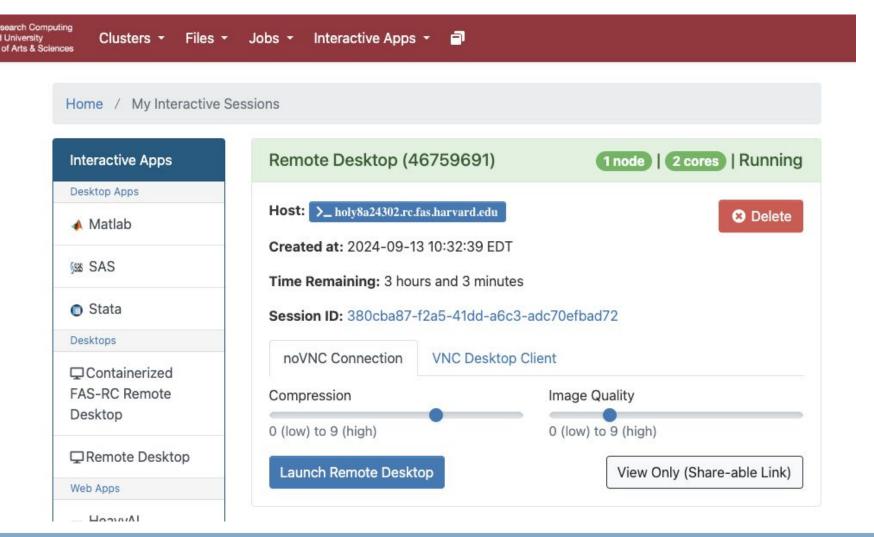


- My Interactive Sessions
- Files
- Jobs
- Interactive apps





My Interactive Sessions

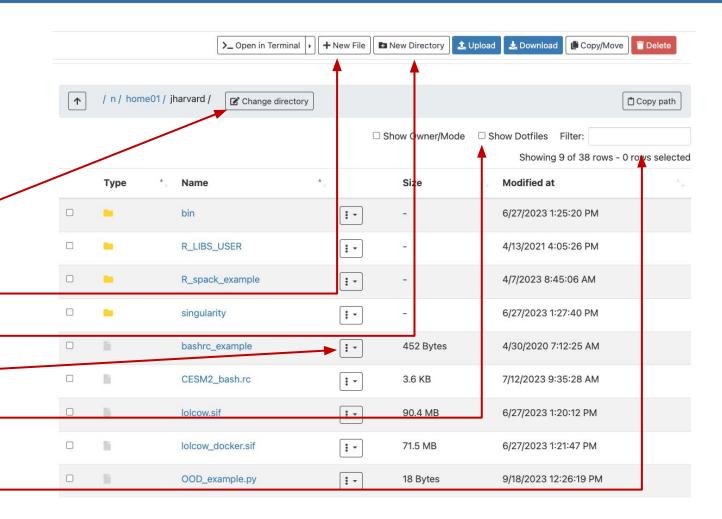






Files tab

- Default options: home directory and holyscratch
- Click on "Change directory" to go to a lab share at /n/holylabs/LABS
- Create new file
- Create new directory (i.e., folder)
- Click on three dots for options
- Check "Show Dotfiles" to see hidden files
- Filter to find files or directories in current directory

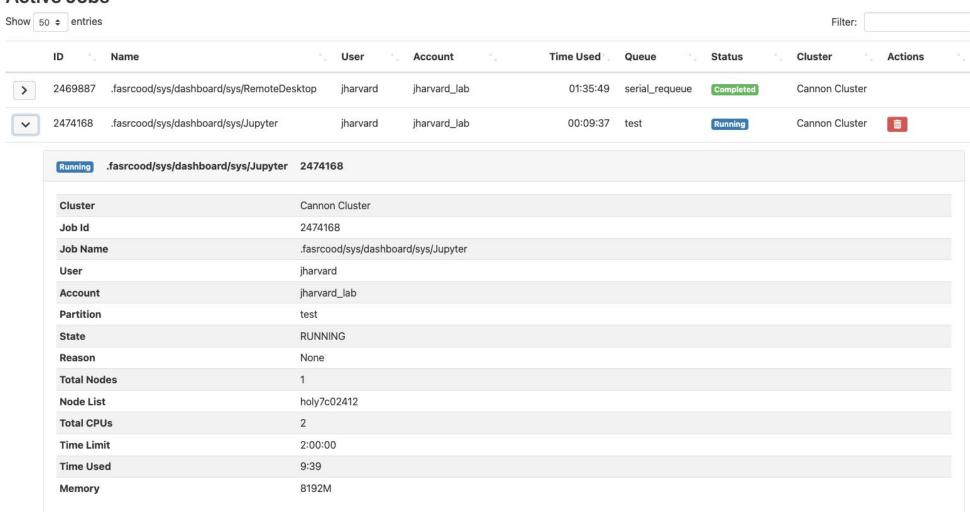






Jobs tab (1)

Active Jobs







Jobs tab (2)





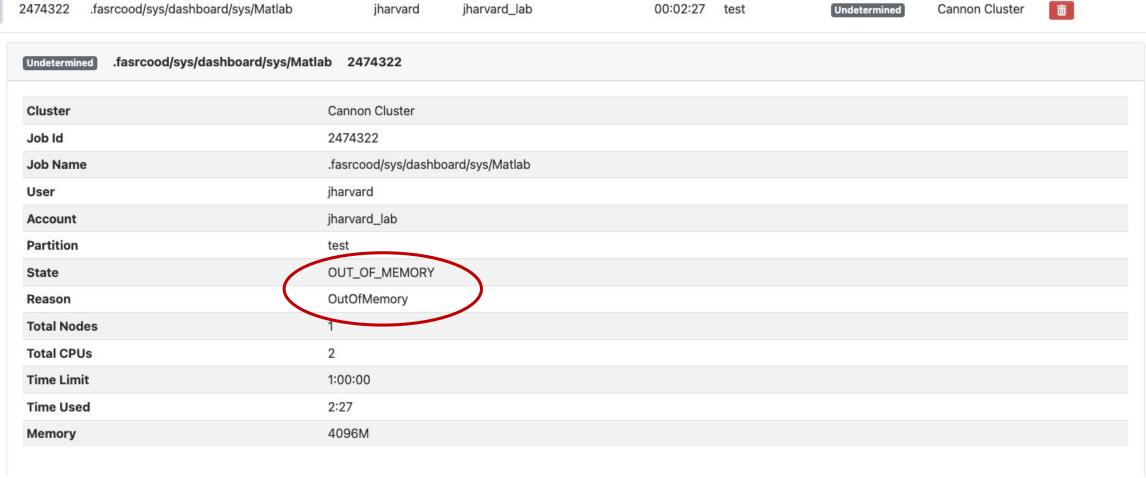
Active Jobs







Jobs tab (3)







Job tab (4)

If job no longer appears on "Active Jobs", check job status from command line with slurm job ID

slurm job ID



[jharvard@bosl	ogin01 ~]\$	sacct -j 2	2464856				
JobID	JobName	Partition	Account	AllocCPUS	State	ExitCode	
2464856	fasrcood+	test	jharvard_+	2	TIMEOUT	0:0	
2464856.bat+	batch		jharvard_+	2	CANCELLED	0:15	
2464856.ext+	extern		jharvard <u></u> +	2	COMPLETED	0:0	
[jharvard@holy	7c02111 ~]\$	\$ sacct -j	2471535				
JobID	JobName	Partition	Account	AllocCPUS	State	ExitCode	
2471535	fasrcood+	test	jharvard_+	2	OUT_OF_ME+	0:125	
2471535.bat+	batch		jharvard +	2	OUT OF ME+	0:125	
2471535.ext+	extern		jharvard +	2	COMPLETED	0:0	





Closing running OOD windows/tabs

- In most OOD apps, you can close the browser tab while the code is running, and the code will continue to run on the background
- Jupyter Notebook will not! The cell that is running will lose the data and output files will not be written
 - Solution: run Remote Desktop app and launch Jupyter Notebook from within Remote Desktop
 - Documentation:
 https://docs.rc.fas.harvard.edu/kb/ood-remote-desktop-how-to-open-software/#Jupyter_Notebook
- Because closing tabs does not end the application, it is important to cancel your job when you are done using it. Otherwise it will be charged to your lab's fairshare.





FASSE proxy

Documentation: https://docs.rc.fas.harvard.edu/kb/proxy-settings/

- You may need to set FASSE proxy on
 - RStudio server if you are unable to reach cran and download R packages
 - Stata if you are unable to load libraries via http
 - Firefox (web browsing)
 - Jupyter Notebook
 - Access Github
 - (Basically, anything outside of FASSE)





Quickstart Guides for using the FASRC Clusters

- Cannon Quickstart Guide
 - https://docs.rc.fas.harvard.edu/kb/igss-cannon-quickstart-guide
- FASSE Quickstart Guide
 - https://docs.rc.fas.harvard.edu/kb/igss-fasse-quickstart-guide
- Quickstart guides have more than just information on OOD
 - how to do text based access
 - office hours, training, tickets





FASRC documentation

- FASRC docs: https://docs.rc.fas.harvard.edu/
- GitHub User_codes: https://github.com/fasrc/User_Codes/
- Getting help
 - Office hours: https://www.rc.fas.harvard.edu/training/office-hours/
 - Ticket
 - o Email: <u>rchelp@rc.fas.harvard.edu</u> to open a ticket with us





Upcoming trainings

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

Containers on the FASRC clusters

This training covers using and managing software/workflow environments using containers. We will touch on the basic differences and how they can be used to help you build/use complex software environments on the cluster.

This training is for those who already have a FASRC account, and want to learn tools and services.





Upcoming trainings

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

Getting started on the FASRC clusters with command line interface (CLI)

- Requirement: working FASRC account with cluster access
- Audience
 - Users familiar with command-line interface
 - New to Cannon and FASSE, but familiar with HPC systems
- Content
 - Submit interactive job with salloc
 - Submit batch job with sbatch
 - Monitor jobs
 - Cluster software overview (modules, spack)





Upcoming trainings

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

Advanced Cluster Usage

This training would focus on users who are familiar with the command line interface and would like to improve job submission and management/monitoring.

Objectives:

- Submit interactive and batch jobs
- Request resources appropriate to job requirements
- Monitoring jobs, priority, when jobs will run
- Fairshare
- Scratch vs. home directory performance





Training session evaluation

Please, fill out our training session evaluation. Your feedback is essential for us to improve our trainings!

https://tinyurl.com/FASRC-training







Where to access the slides

https://docs.rc.fas.harvard.edu/kb/training-materials/







Thank you:)
FAS Research Computing