



The Institute for Quantitative Social Science



Getting Started on the FASRC
Clusters with Open OnDemand



Faculty of Arts and Sciences Research Computing

Learning objectives

- 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

- **OOD**: Open On Demand
- **Cluster**: large group of servers with lots of memory and processors
- **Cannon**: cluster that handles level 2 data. Named after the 19th century Harvard astronomer Annie Jump Cannon.
- **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



Cisco AnyConnect | vpn.rc.fas.harvard.edu



Please enter your RC username and password.

!!! IF YOU HAVE ISSUES UPGRADING YOUR VPN CLIENT:
Please manually download and install by logging into <https://vpn.rc.fas.harvard.edu>

This system is for authorized users at Harvard University.
No other use is permitted.

SUM1 VPN

Username:

Password:

Two-Step Verification Code:

Cancel

OK

Cisco AnyConnect | vpn.rc.fas.harvard.edu



Please enter your RC username and password.

!!! IF YOU HAVE ISSUES UPGRADING YOUR VPN CLIENT:
Please manually download and install by logging into <https://vpn.rc.fas.harvard.edu>

This system is for authorized users at Harvard University.
No other use is permitted.

SUM1 VPN

Username:

Password:

Two-Step Verification Code:

Cancel

OK

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`

Please Sign In

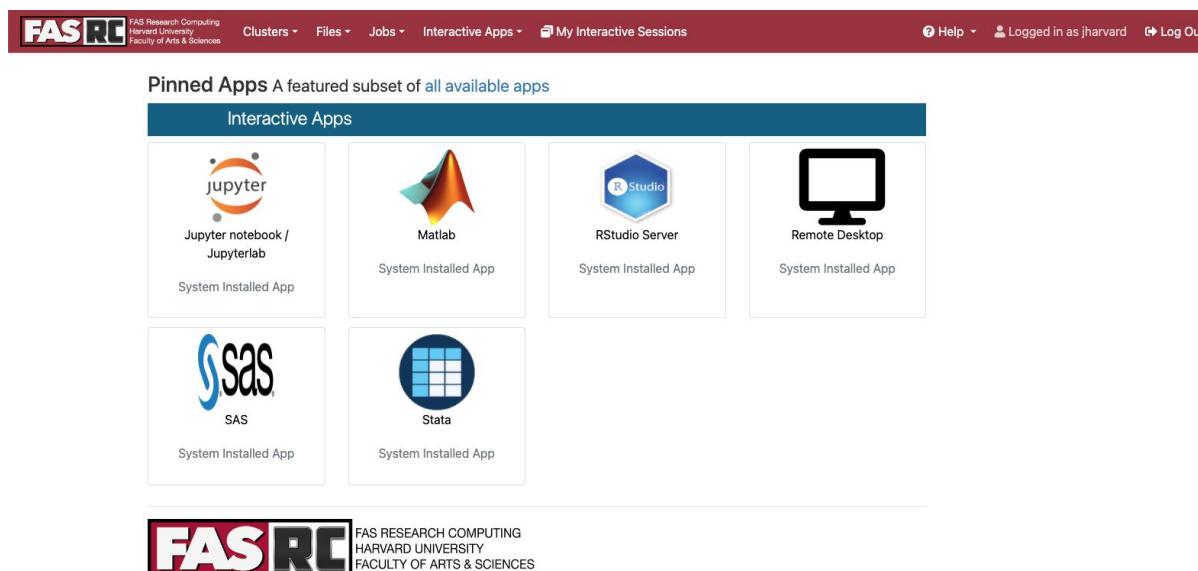
<https://rcood.rc.fas.harvard.edu>

Username:

Password:

OOD dashboard on Cannon and FASSE

Cannon



Welcome to FAS-RC Cluster

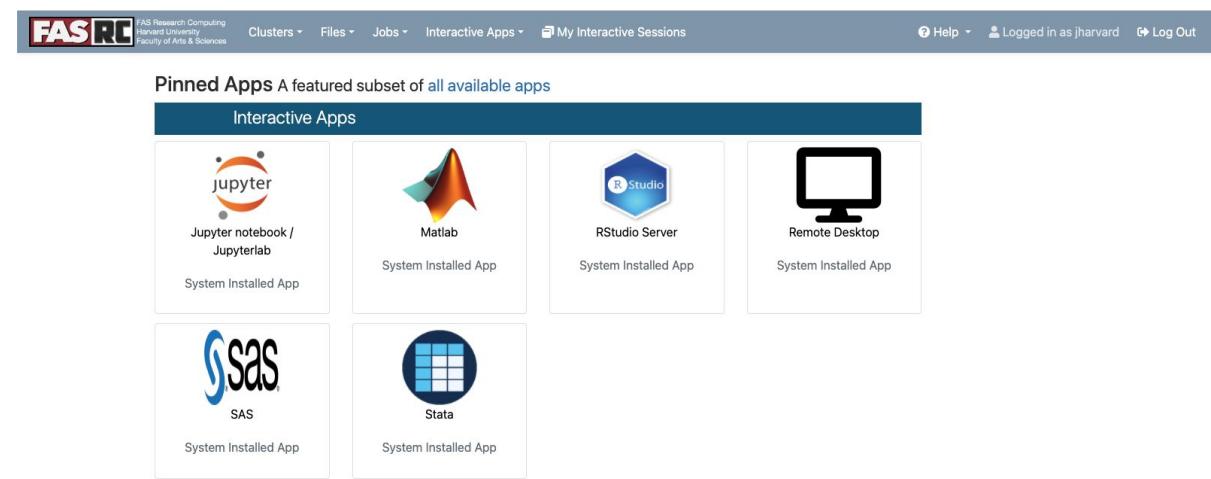
The Computing Cluster is a resource for the research community, hosted by Research Computing at Harvard University's Faculty of Arts and Sciences.

To apply for an account please refer to [this webpage](#).

From this web service you can submit your jobs, check running jobs, and open interactive graphical sessions to run your favorite applications.

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

FASSE



Welcome to FASSE

The Computing Cluster is a resource for the research community, hosted by Research Computing at Harvard University's Faculty of Arts and Sciences.

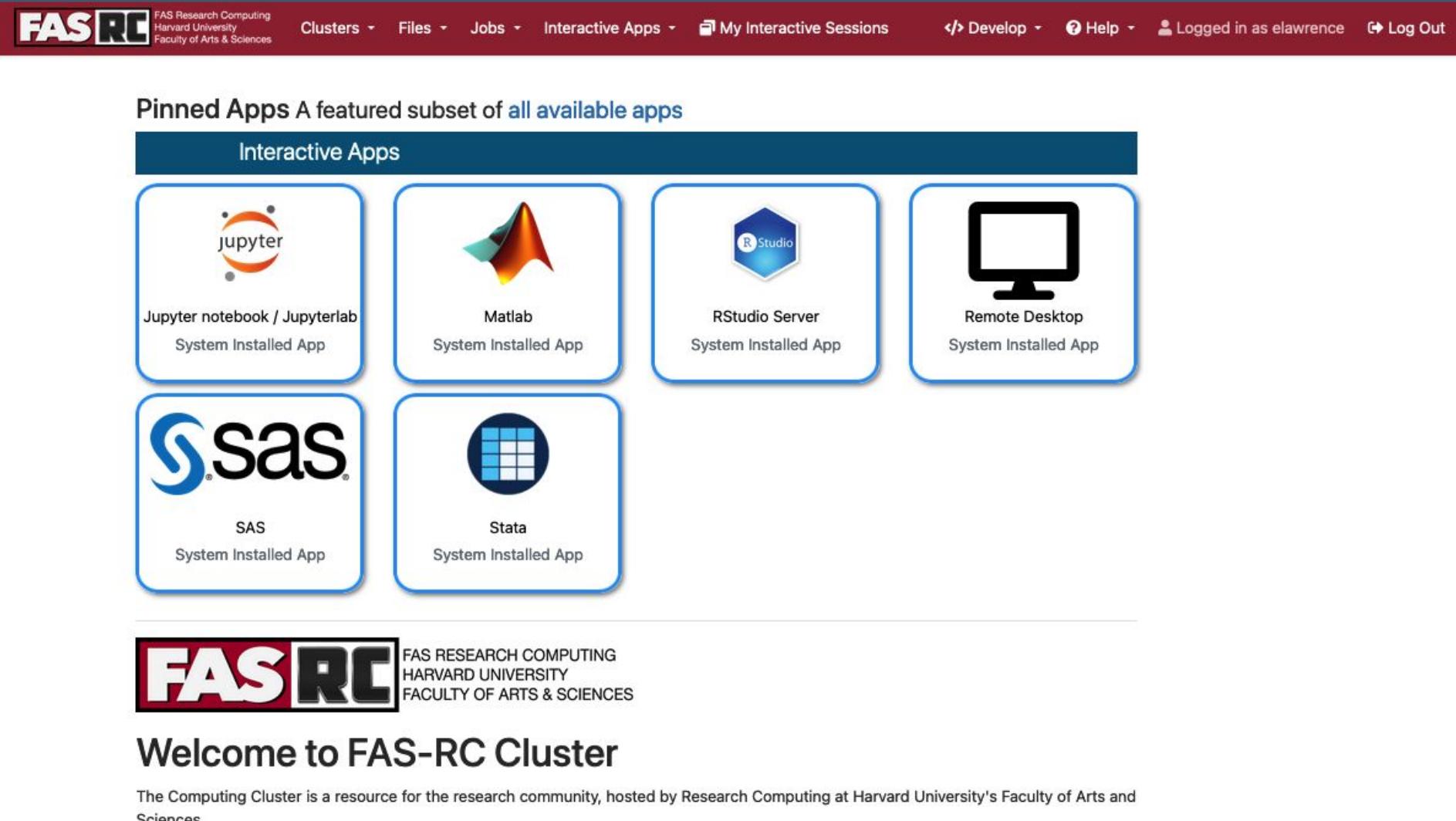
To apply for an account please refer to [this webpage](#).

From this web service you can submit your jobs, check running jobs, and open interactive graphical sessions to run your favorite applications.

These are some examples of the things you will be able to do :

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

Launching an app from the Dashboard



Pinned Apps A featured subset of [all available apps](#)

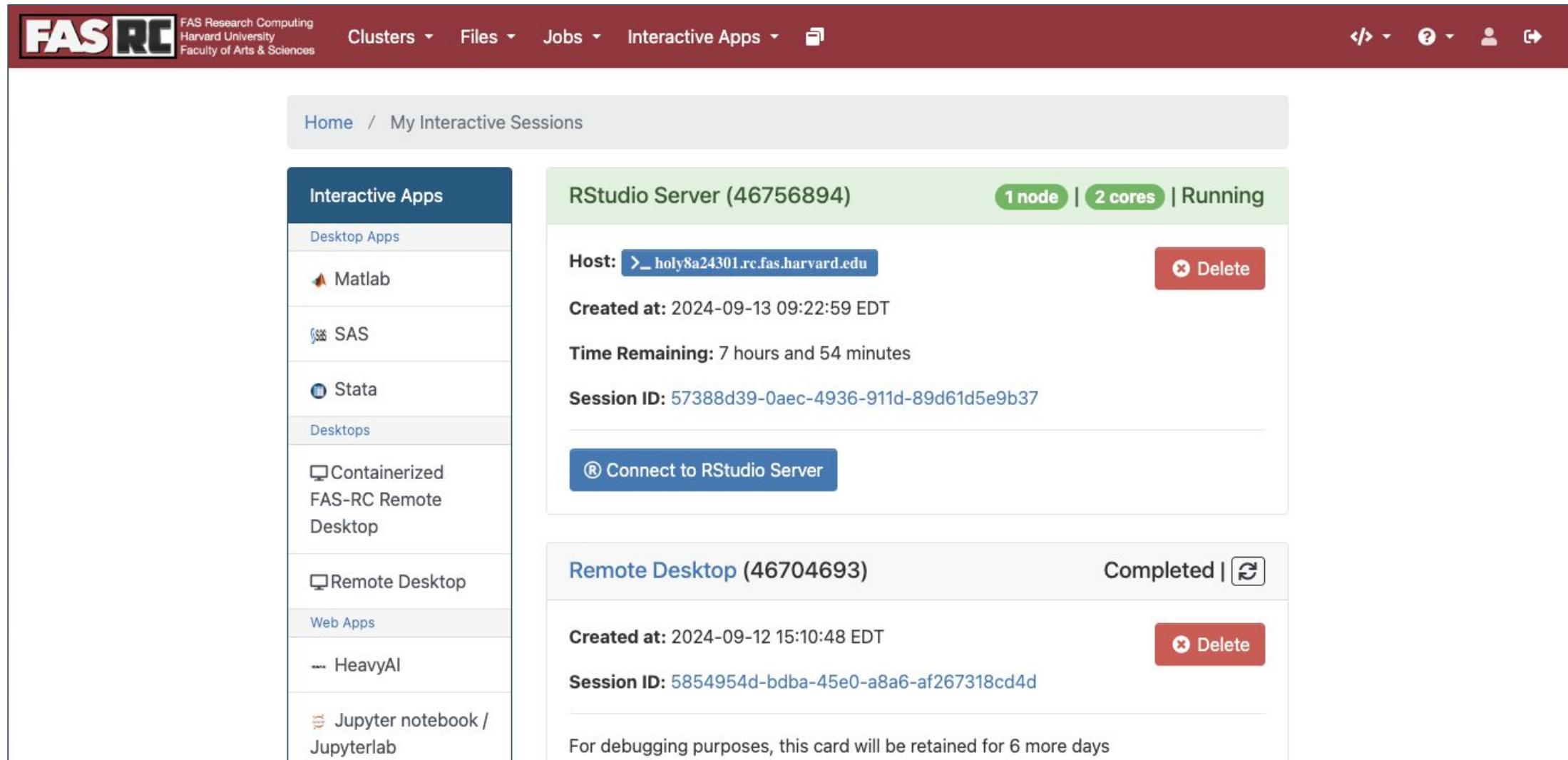
Interactive Apps

-  **Jupyter notebook / Jupyterlab**
System Installed App
-  **Matlab**
System Installed App
-  **RStudio Server**
System Installed App
-  **Remote Desktop**
System Installed App
-  **SAS**
System Installed App
-  **Stata**
System Installed App

FAS RC FAS RESEARCH COMPUTING
HARVARD UNIVERSITY
FACULTY OF ARTS & SCIENCES

Welcome to FAS-RC Cluster

The Computing Cluster is a resource for the research community, hosted by Research Computing at Harvard University's Faculty of Arts and Sciences



The screenshot shows the FAS RC interface for managing interactive sessions. On the left, a sidebar lists various desktop applications and services. The main area displays two active sessions: an RStudio Server session and a Remote Desktop session.

Interactive Apps

- Desktop Apps
 - Matlab
 - SAS
 - Stata
- Desktops
 - Containerized FAS-RC Remote Desktop
 - Remote Desktop
- Web Apps
 - HeavyAI
 - Jupyter notebook / Jupyterlab

RStudio Server (46756894) 1 node | 2 cores | Running

Host: > [holy8a24301.rc.fas.harvard.edu](#) ✖ Delete

Created at: 2024-09-13 09:22:59 EDT

Time Remaining: 7 hours and 54 minutes

Session ID: 57388d39-0aec-4936-911d-89d61d5e9b37

[② Connect to RStudio Server](#)

Remote Desktop (46704693) Completed | ⏪

Created at: 2024-09-12 15:10:48 EDT ✖ Delete

Session ID: 5854954d-bdba-45e0-a8a6-af267318cd4d

For debugging purposes, this card will be retained for 6 more days

Remote Desktop

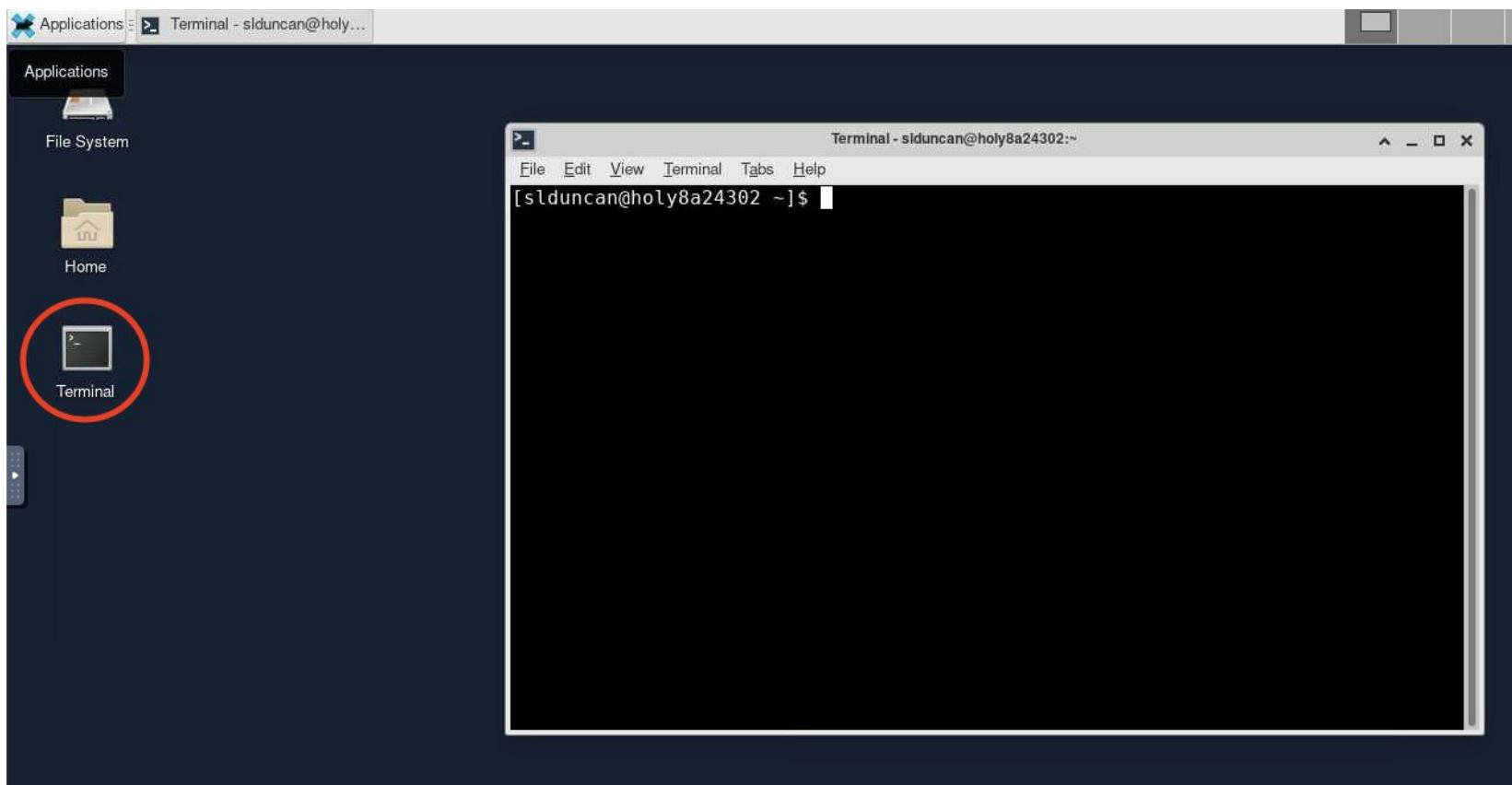
- Not as necessary as it used to be
- **For running long Jupyter Notebook sessions**
- Terminal
- Can also be used to open multiple applications in a single window
- Choose the defaults for resolution

Resolution

width	1024	px	height	768	px
-------	------	----	--------	-----	----

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
```

Creating your own kernel for Jupyter Notebook

- You can create your own kernels, but some command line needed
- Note: a kernel is the same as a conda, python, or mamba environment

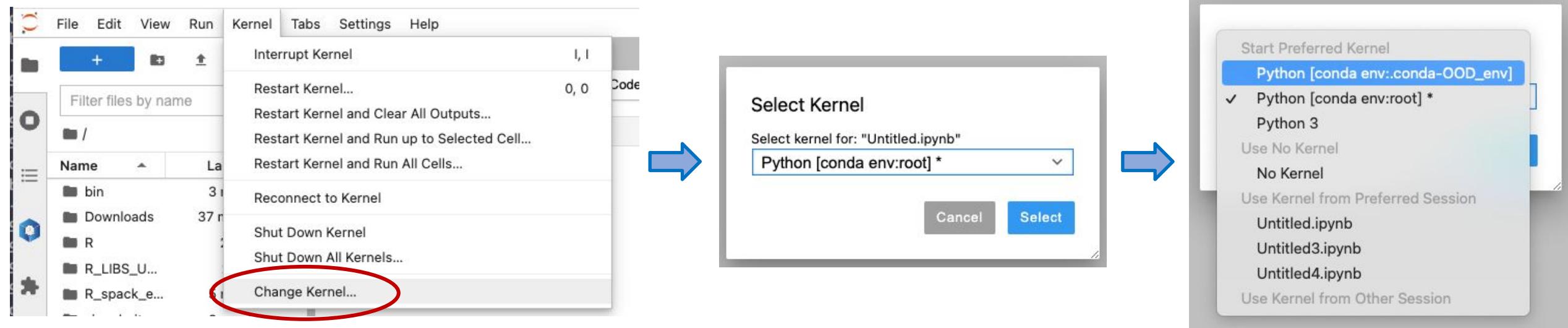
1. Open a terminal in the “Remote Desktop” app
⇒Don’t create mamba environments inside Jupyter Notebook/Lab!
2. Create mamba environment and install package `ipykernel` and `nb_conda_kernels`
- 3.

```
[jharvard@holly7c02111 ~]$ module load python
[jharvard@holly7c02111 ~]$ mamba create -n OOD_env python=3.11 pip wheel numpy
[jharvard@holly7c02111 ~]$ mamba activate OOD_env
(OOD_env) [jharvard@holly7c02111 ~]$ mamba install ipykernel nb_conda_kernels
```

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

Using your new environment in Jupyter Notebook

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



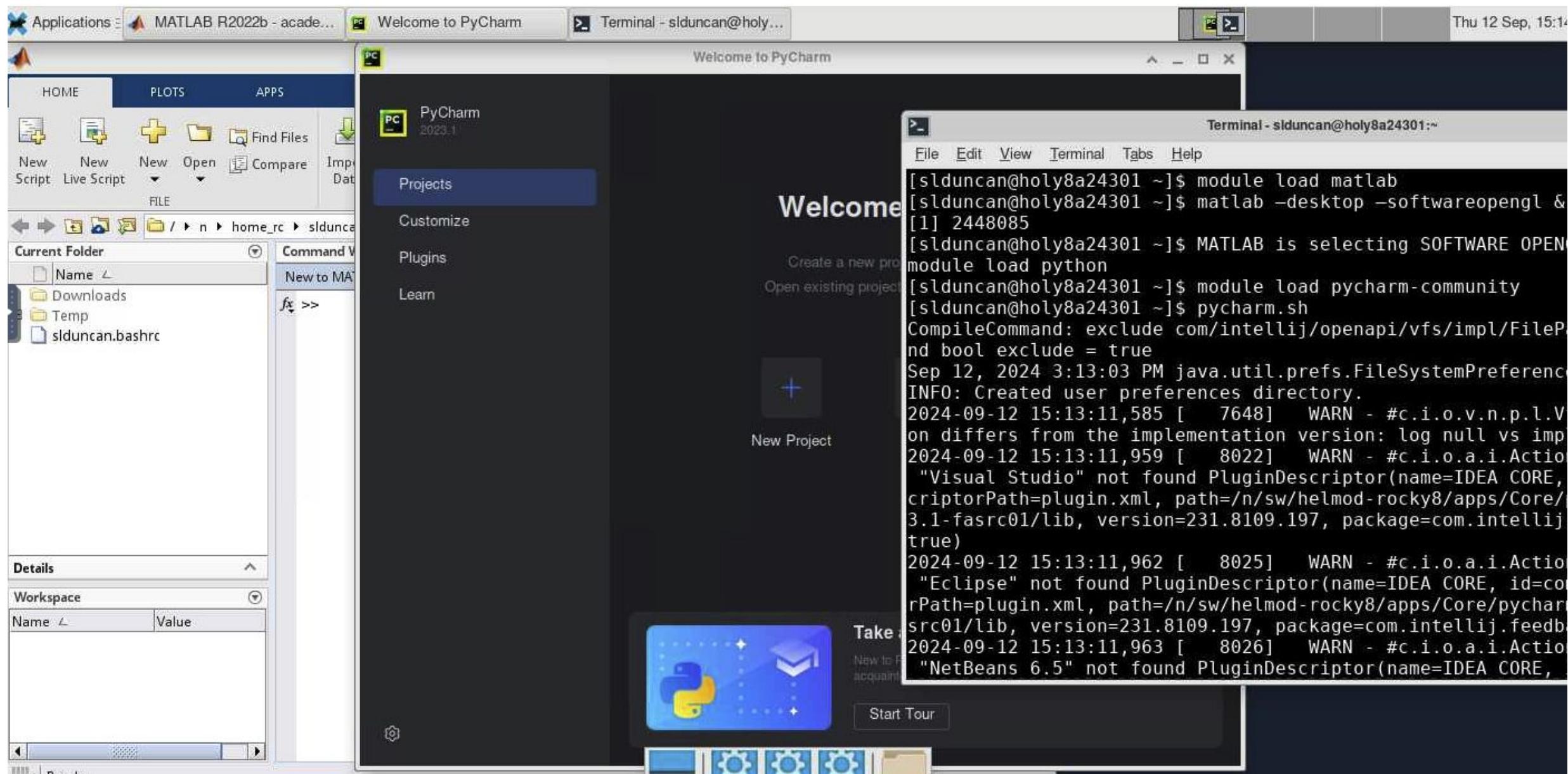
Opening Multiple Applications in Remote Desktop

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

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

```
# Matlab
[jharvard@holly7c02111 ~]$ module load matlab
[jharvard@holly7c02111 ~]$ matlab -desktop -softwareopengl &

#PyCharm
[jharvard@holly7c02111 ~]$ module load python
[jharvard@holly7c02111 ~]$ module load pycharm-community
[jharvard@holly7c02111 ~]$ pycharm.sh
```



Inactivity lock out

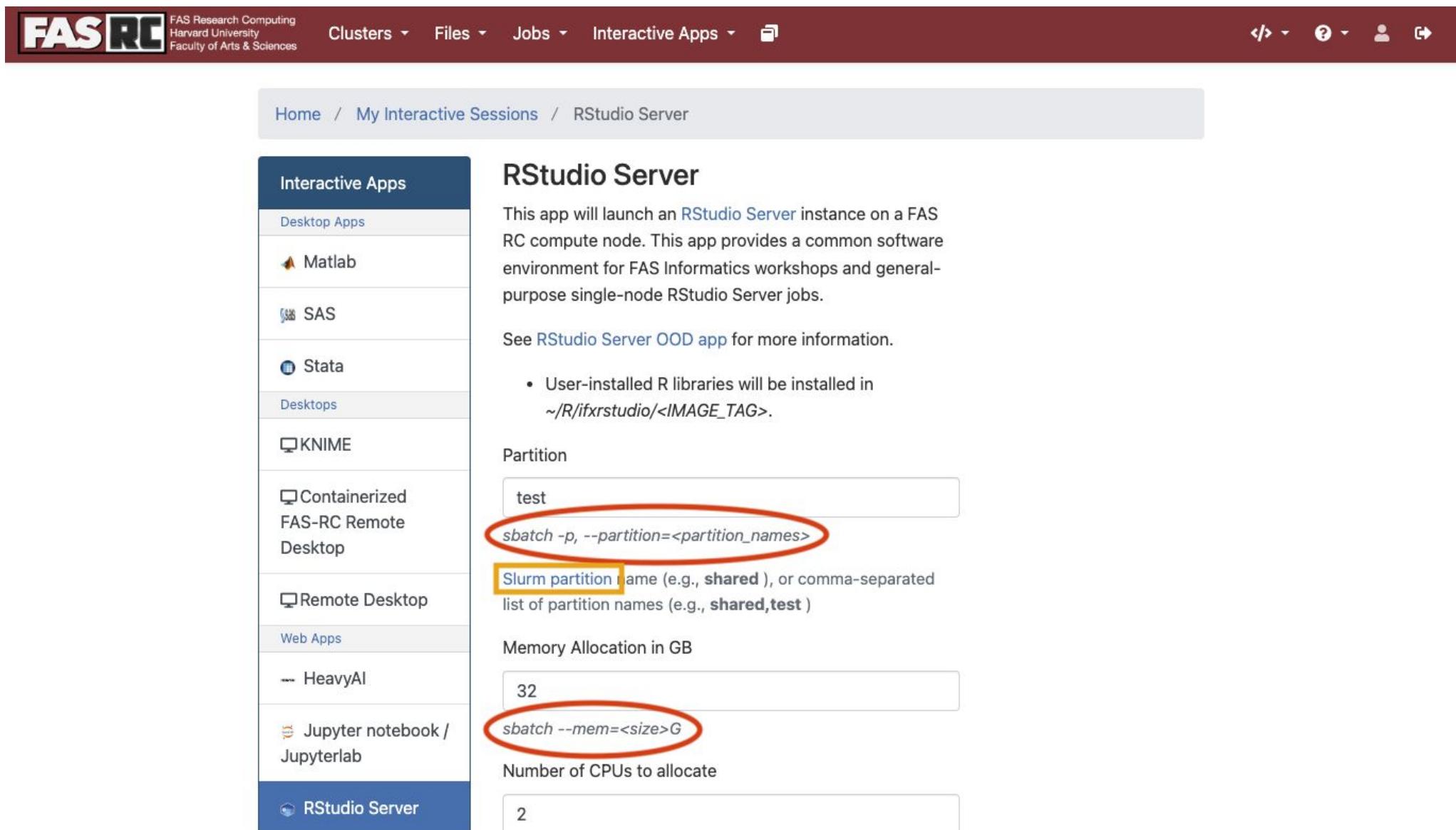
- It may lock out due to inactivity
- Use your FASRC password to unlock



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 [Cannon](#) vs [FASSE](#)
 - 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



Interactive Apps

Clusters ▾ Files ▾ Jobs ▾ Interactive Apps ▾ 

Home / My Interactive Sessions / RStudio Server

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

`32`

`sbatch --mem=<size>G`

Number of CPUs to allocate

`2`

Slurm Partitions

Partition is the term that Slurm uses for queues. Partitions can be thought of as a set of resources and parameters around their use (See also: [Convenient Slurm Commands](#)). You can find out what partitions you have access to using the `spart` command. FASSE has [different partitions than Cannon](#).

Note: In the case where no resources have been requested explicitly, default resources that get allocated to a job on Cannon or FASSE are, `serial_requeue` for the partition, 10 minutes for the time, 1 core, and 100 MB for the memory.

Partition	Nodes	Cores per Node	CPU Core Types	Mem per Node (GB)	Time Limit	Max Jobs	Max Cores	GPU Capable?	/scratch size (GB)
sapphire	186	112	Intel "Sapphire Rapids"	990	3 days	none	none	No	396
shared	310	48	Intel "Cascade Lake"	184	3 days	none	none	No	68
bigmem	4	112	Intel "Sapphire Rapids"	1988	3 days	none	none	No	396
bigmem_intermediate	3	64	Intel "Ice Lake"	2000	14 days	none	none	No	396
gpu	36	64	Intel "Ice Lake"	990	3 days	none	none	Yes (4 A100/node)	396
intermediate	12	112	Intel "Sapphire Rapids"	990	14 days	none	none	No	396
unrestricted	8	48	Intel "Cascade Lake"	184	none	none	none	No	68
test	18	112	Intel "Sapphire Rapids"	990	12 Hours	5	112	No	396
gpu_test	14	64	Intel	487	12	5	64	Yes (8 A100 MIG)	172

Table of Contents

Overview: The FASRC Cluster Uses Slurm to Manage Jobs

Cluster Jobs are Generally Run From the Command Line

- > Cluster Applications Should Not Be Run From Login Nodes

- > Storage and Scratch on the Cluster

SLURM Resources

- > Summary of Slurm Commands

- > Slurm Limits

Slurm Partitions

- > Partition Details

Submitting Batch Jobs Using the `batch` Command

- > Notifications by Email:

- > It is important to accurately request resources, especially memory

Monitoring Job Progress with `squeue` and `sacct`

- > See Broader Queue with `showq`

- > Canceling Jobs with

Allocated Time (expressed in MM , or HH:MM:SS , or DD-
HH:MM)

02-08:00

sbatch -t, --time=<time>

R version

R 4.3.3 (Bioconductor 3.18, RStudio 2023.09.1) ▾

This defines the version of R/Bioconductor you want to use.

I would like to receive an email when the session starts

(optional) email address for job status notification

(optional) Slurm Reservation

sbatch --reservation=<name>

Leave blank if you do not have a Slurm reservation

Start rstudio with a new configuration

*Checking this box will start rstudio with a fresh
configuration.*

This is useful if you need to run different instances at the
same time with different configurations.

Leave the box unchecked if you want rstudio to start with the
default configuration you have in your `~/.rstudio`

(optional) Slurm Account

sbatch -A, --account=<account>

Leave blank if you do not have multiple Slurm accounts

Additional Slurm options

additional slurm options (**long format option only**)

Example : --constraint=intel --exclusive

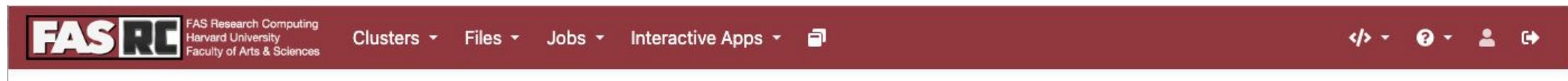
Please make sure the options you select are compatible with other fields in the form.

Also, please make sure you use **long format option only**
(e.g. --nodelist=holy7c24502 instead of -w holy7c24502)

Launch

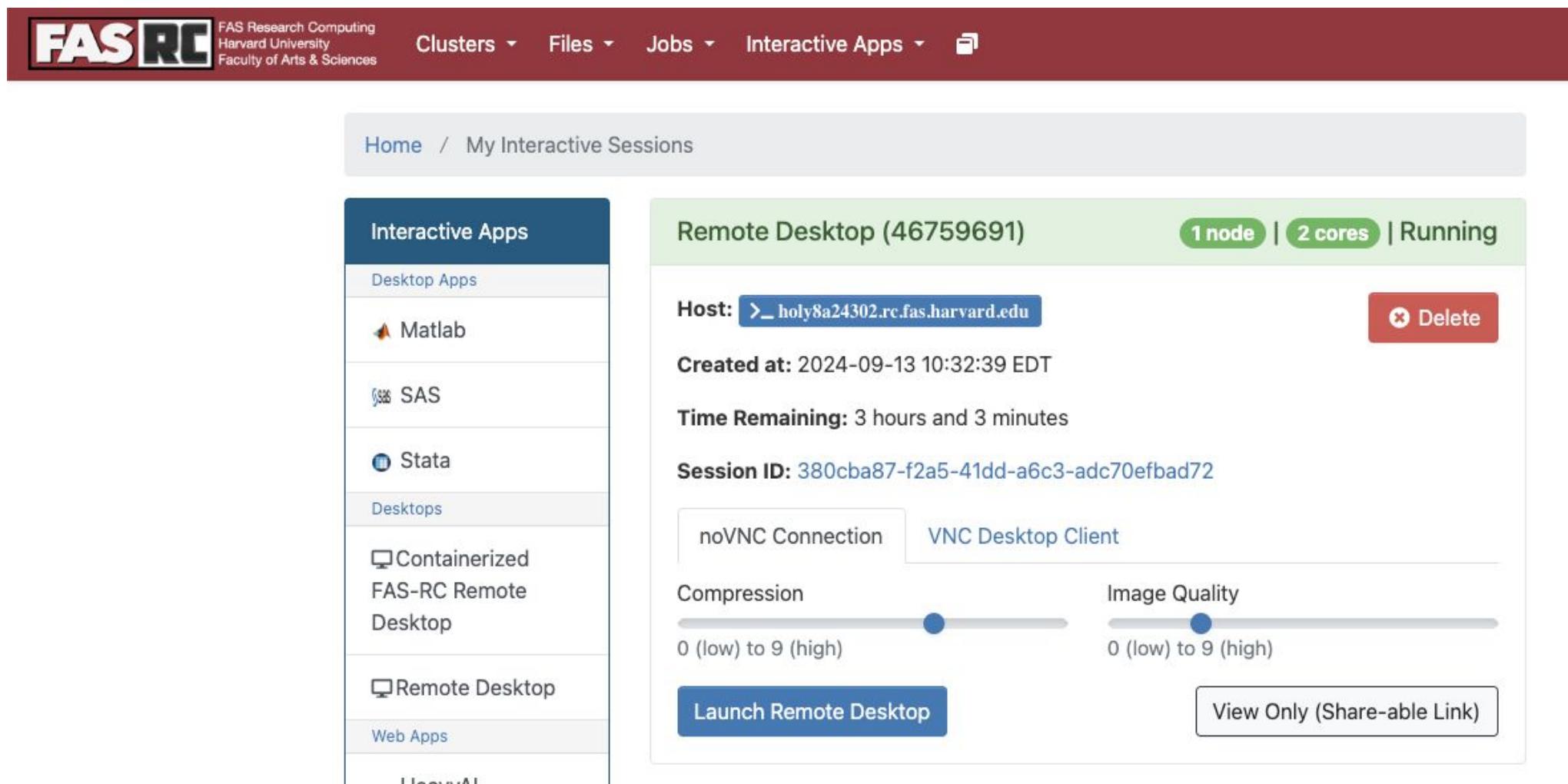
* The RStudio Server session data for this session can be accessed under the [data root directory](#).

Menu bar



- My Interactive Sessions
- Files
- Jobs
- Interactive apps

My Interactive Sessions



The screenshot shows the FAS RC web interface for managing interactive sessions. The top navigation bar includes the FAS RC logo, a search bar, and links for Clusters, Files, Jobs, Interactive Apps, and a help icon. The main content area has a breadcrumb navigation (Home / My Interactive Sessions) and a sidebar titled 'Interactive Apps' with categories: Desktop Apps (Matlab, SAS, Stata), Desktops (Containerized FAS-RC Remote Desktop, Remote Desktop), and Web Apps. The main panel displays a 'Remote Desktop (46759691)' session for Matlab, which is running on 1 node with 2 cores. The session was created at 2024-09-13 10:32:39 EDT and has 3 hours and 3 minutes remaining. The session ID is 380cba87-f2a5-41dd-a6c3-adc70efbad72. It includes configuration sliders for 'Compression' (0 to 9) and 'Image Quality' (0 to 9), and buttons for 'Launch Remote Desktop' and 'View Only (Share-able Link)'.

Home / My Interactive Sessions

Interactive Apps

Desktop Apps

- Matlab
- SAS
- Stata

Desktops

- Containerized FAS-RC Remote Desktop
- Remote Desktop

Web Apps

Matlab

Remote Desktop (46759691)

1 node | 2 cores | Running

Host: [holy8a24302.rc.fas.harvard.edu](#)

Created at: 2024-09-13 10:32:39 EDT

Time Remaining: 3 hours and 3 minutes

Session ID: [380cba87-f2a5-41dd-a6c3-adc70efbad72](#)

noVNC Connection VNC Desktop Client

Compression: 0 (low) to 9 (high)

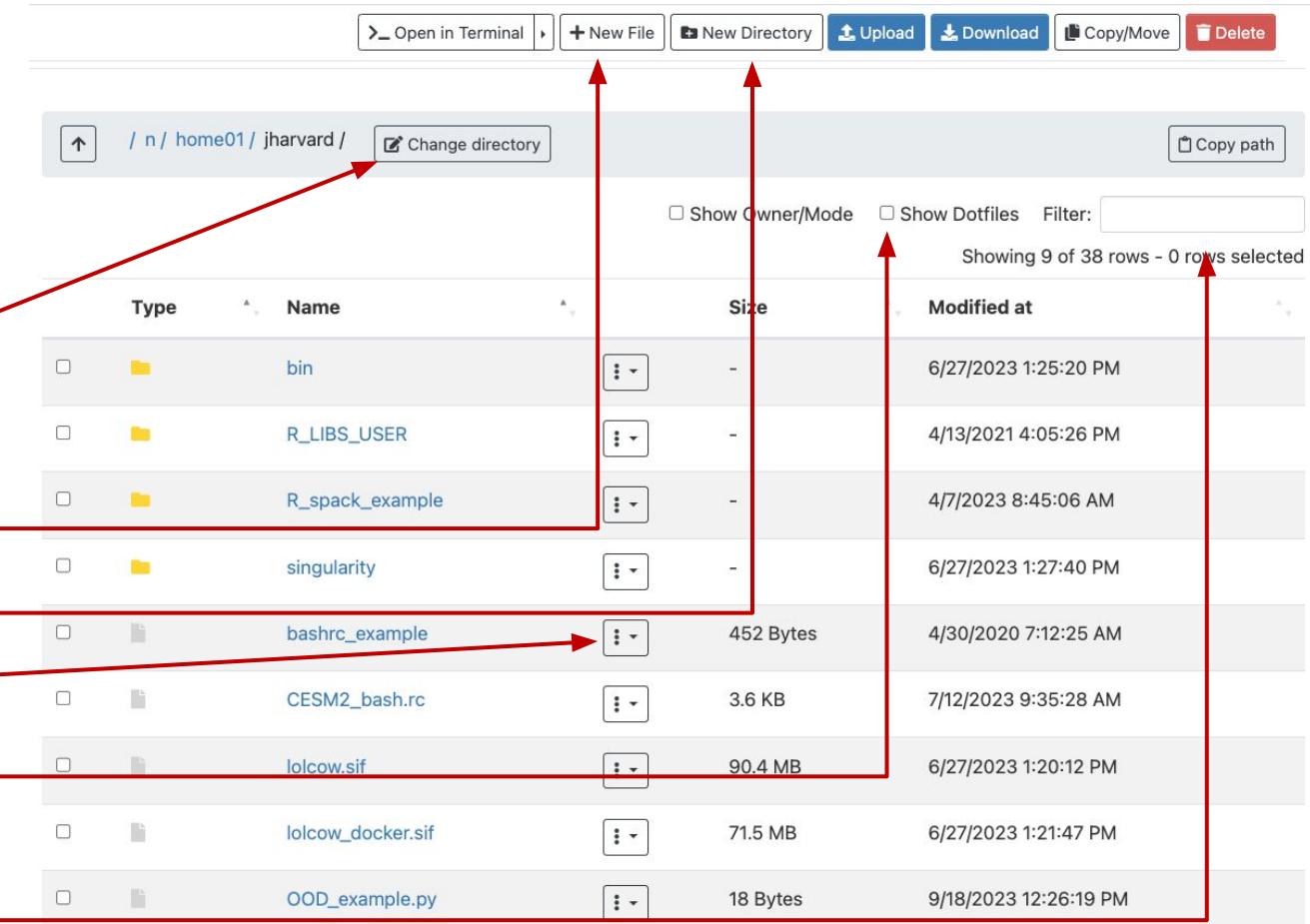
Image Quality: 0 (low) to 9 (high)

Launch Remote Desktop

View Only (Share-able Link)

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



Active Jobs and their details in the Jobs tab

Active Jobs

Show 50 entries

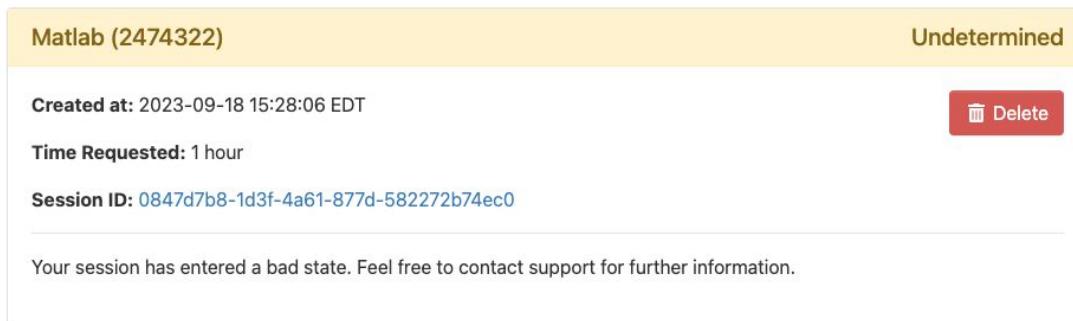
 Filter:

ID	Name	User	Account	Time Used	Queue	Status	Cluster	Actions
2469887	.fasrcood/sys/dashboard/sys/RemoteDesktop	jharvard	jharvard_lab	01:35:49	serial_requeue	Completed	Cannon Cluster	
2474168	.fasrcood/sys/dashboard/sys/Jupyter	jharvard	jharvard_lab	00:09:37	test	Running	Cannon Cluster	

Running .fasrcood/sys/dashboard/sys/Jupyter 2474168

Cluster	Cannon Cluster
Job Id	2474168
Job Name	.fasrcood/sys/dashboard/sys/Jupyter
User	jharvard
Account	jharvard_lab
Partition	test
State	RUNNING
Reason	None
Total Nodes	1
Node List	holy7c02412
Total CPUs	2
Time Limit	2:00:00
Time Used	9:39
Memory	8192M

A failed job



Active Jobs

Show 50 entries

 Filter:

ID	Name	User	Account	Time Used	Queue	Status	Cluster	Actions
2469887	.fasrcod/sys/dashboard/sys/RemoteDesktop	jharvard	jharvard_lab	01:35:49	serial_requeue	Completed	Cannon Cluster	Delete
2474322	.fasrcod/sys/dashboard/sys/Matlab	jharvard	jharvard_lab	00:02:27	test	Undetermined	Cannon Cluster	Delete
2474168	.fasrcod/sys/dashboard/sys/Jupyter	jharvard	jharvard_lab	00:15:45	test	Running	Cannon Cluster	Delete

Diagnosing a failed job

<input type="checkbox"/>	2474322	.fasrcood/sys/dashboard/sys/Matlab	jharvard	jharvard_lab	00:02:27	test	Undetermined	Cannon Cluster	
Undetermined .fasrcood/sys/dashboard/sys/Matlab 2474322									
Cluster	Cannon Cluster								
Job Id	2474322								
Job Name	.fasrcood/sys/dashboard/sys/Matlab								
User	jharvard								
Account	jharvard_lab								
Partition	test								
State	OUT_OF_MEMORY								
Reason	OutOfMemory								
Total Nodes	1								
Total CPUs	2								
Time Limit	1:00:00								
Time Used	2:27								
Memory	4096M								

Detailed job information from the command line

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

slurm job ID

RStudio Server (2464856)	Completed
Created at: 2023-09-18 12:42:03 EDT	
Session ID: 743455f6-39e6-40db-85ab-4fcc9b903117	
For debugging purposes, this card will be retained for 6 more days	

```
[jharvard@boslogin01 ~]$ sacct -j 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_+	2	COMPLETED	0:0

```
[jharvard@holy7c02111 ~]$ 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/iqss-cannon-quickstart-guide>
- FASSE Quickstart Guide
 - <https://docs.rc.fas.harvard.edu/kb/iqss-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
 - Email: rchelp@rc.fas.harvard.edu to open a ticket with us

Additional trainings

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

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

Training session evaluation

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

<https://tinyurl.com/FASRCworkshop>



Thank you :)
FAS Research Computing