

## SAVIO cluster

### Register new accounts

1. create portal accounts at MyBRC User Portal (<https://mybrc.brc.berkeley.edu/>)
2. After following the on-screen instructions in the portal and registering and/or logging in, you should first review and sign the cluster [User Access Agreement Form](#) on the Home ("Welcome") page (if they haven't already done so from within the portal previously) by clicking on the "Review" button
3. Click on the "Join" button to request to join project, you can search **ic\_chem242** under project name to find our class project. Click on "join".
4. We will approve your request. And for new users there will be a processing wait time of a few days or so between the time they join their first project via the MyBRC User Portal and the time they receive the confirmation email pointing them to instructions on how they can access their new BRC user account.

### From Terminal (You'll need do this first to setup the kernel)

You only need to do this once.

- Log into cluster
  1. **ssh *your\_user\_name*@hpc.brc.berkeley.edu**
  2. fill in your password and OTP in one entry
  3. Run the following command (in one line) to create a symlink from the kernel I set to your directory:  
**ln -s ~nancy\_guan/.local/share/jupyter/kernels/pt10  
~*your\_user\_name*/.local/share/jupyter/kernels/pt10**
  4. Then run the following command and make sure you can see kernel pt10.  
**jupyter kernelspec list**  
Available kernels:  
pt10 /global/home/users/nancy\_guan/.local/share/jupyter/kernels/pt10
  5. Now you can proceed with the next session to run interactive notebooks.

### From Server (for interactive jupyter notebooks)

1. Go to [ood.brc.berkeley.edu](https://ood.brc.berkeley.edu)
2. You'll be taken to a sign in page. Fill in username, password and OTP here. OTP is a 6-digit one-time password (from Google Authenticator)

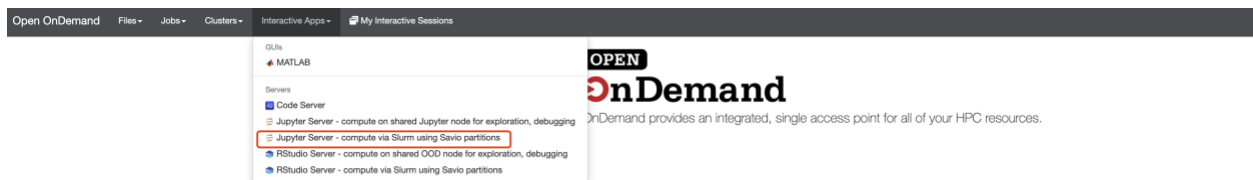
Sign in

https://ood.brc.berkeley.edu







Username

Password

3. You should see this page, and then select interactive apps – Jupyter Server Savio partitions



4. Fill in the information as shown below to request an interactive node. You can change the wall clock time to the number of hours you need.

Interactive Apps
GUIs
 MATLAB
Servers
 Code Server
 Jupyter Server - compute on shared Jupyter node for exploration, debugging
 Jupyter Server - compute via Slurm using Savio partitions
 RStudio Server - compute on shared OOD node for exploration, debugging
 RStudio Server - compute via Slurm using Savio partitions

## Jupyter Server - compute via Slurm using Savio partitions

version: 2021.pumpkin-2-g42a194c

This app will launch a [Jupyter](#) server session on the Berkeley Research Computing(BRC) Savio cluster.

### Name of the Job

### SLURM Partition

The SLURM Partition in which you want to launch this Jupyter session.

### SLURM Account/Project Name

The SLURM account (i.e., the value of the `-A` or `--account` flag used when submitting a SLURM job).

### SLURM QoS Name

The QoS you want run under. Most FCA users choose `savio_normal`. Condo users can either use their `condo` QoS or the `low-priority` QoS.

### Number of Compute Nodes

The number of nodes you want for your Jupyter session.

### Number of CPU Cores per Node

The number of CPU cores you want per node for your Jupyter session.

### Number and Type of GPUs

The GRES value in the form `'gpu:x'` or `'gpu:type:x'` where `'x'` is the number of GPUs and (optionally) `'type'` is the type of GPU. Remember to specify two CPUs for each GPU in the 'Number of CPU Cores per Node' field.

### Wall Clock Time

The maximum number of hours your Jupyter session will run for. To save FCA credits or free up resources for your condo group members, you should delete your session when you are done.

### Email Address (Optional)

Enter your email address if you would like to receive an email when the session starts. Leave blank for no email.

Launch

\* The Jupyter Server - compute via Slurm using Savio partitions session data for this session can be accessed under the [data root directory](#).

5. You should see this page when the node is ready. Click on Connect to Jupyter. If you finish earlier than the time you requested, you can click on Delete to free up the node for other people.

**Jupyter Server - compute via Slurm using Savio partitions (11701372)** 1 node | 2 cores | Running

**Host:** >n0223.savio2 Delete

**Created at:** 2022-04-16 12:17:28 PDT

**Time Remaining:** 59 minutes

**Session ID:** cbeee7da-63b6-4b14-a4e0-cd80f8ac0c39

Connect to Jupyter

6. You are now on the Jupyter Server! You can upload files here or start a new notebook. You should see a kernel with name Python3.6 PT1.0.0 after you follow previous steps to set up a symlink, which allows you to import pytorch.

The screenshot shows the JupyterLab interface. At the top, there's a 'jupyter' logo and 'Quit' and 'Logout' buttons. Below that, there are tabs for 'Files', 'Running', and 'IPython Clusters'. The 'Files' tab is active, showing a file browser with a path bar indicating the current directory is '/global/home/users/nancy\_guan'. A list of files and folders is shown, including '..', 'cgem', 'ondemand', and 'teaching'. On the right side, there's a dropdown menu for 'Notebook:' with options: 'Python 2.7', 'Python 3.6', 'Python 3.7', 'Python3.6 PT1.0.0' (highlighted with a red box), 'Python3.6 TF-1.12', 'Python3.6 pytorch/1.0.0', 'Python3.7 TF-2.1.0', 'Python3.7 TF-2.3.0', and 'pyspark'. Below this, there's an 'Other:' section with options: 'Text File', 'Folder', and 'Terminal'. A tooltip 'Create a new notebook with Python 2.7' is visible next to the 'Python 2.7' option.

## **File Transfer**

<https://docs-research-it.berkeley.edu/services/high-performance-computing/user-guide/data/transferring-data/using-scp-savio/>

Run these commands from your local terminal

- Download.  
**scp your\_user\_name@dtb.brc.berkeley.edu:/remote/path/myfile /local/path**
- Upload.  
**scp /local/path/myfile your\_user\_name@ dtb.brc.berkeley.edu:/remote/path**
- Use **scp -r** for transferring folders.
- The remote path needs to be inside your \$HOME folder, you can find this path by running **pwd** after log into cluster. In general it should look like **/global/home/users/your\_user\_name**