



Python - Package Installation & Execution on FASRC Clusters

Learning objectives

- Python using CLI (Command Line Interface)
 - Interactive
 - Sbatch
- Python Package installation
 - Interactive
 - Sbatch
- Python using OOD (Open On Demand)
- Jupyter Notebook
 - Create conda environment (i.e., jupyter kernel)

Python Programming Language

[Python Programming Language – FASRC DOCS](#)

- High-level, general-purpose, and object-oriented programming language with emphasis on code readability and use of significant indentation.
- Ideal for scripting and rapid application development given its dynamic typing, elegant syntax, and automatic memory management (garbage collection).
- Has a comprehensive standard library. Also known as “batteries-included” language.
- Python’s implementation is mostly in C.
 - Python’s core interpreter, CPython, written in C.
- Interpreted language, hence slower than compiled languages, like C and Fortran.
 - Compiled generates executable
 - Interpreted executes instructions directly on the fly without compiling a program into machine language

Training Material

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

```
# Check current location & change if desired for this training: pwd
cd <desired-location>
```

```
# Clone FASRC User Codes repository:
```

```
https://github.com/fasrc/User\_Codes/tree/master
```

```
SSH - git clone git@github.com:fasrc/User_Codes.git
```

```
HTTPS - git clone https://github.com/fasrc/User\_Codes.git
```

```
# Create a training folder & go to that folder:
```

```
mkdir python-training
```

```
cd python-training
```

```
# Copy Python folders from the User Codes directory:
```

```
cp -r ../User_Codes/Languages/Python .
```

```
cp -r ../User_Codes/Parallel_Computing/Python/Python-Multiprocessing-Tutorial .
```

Python using CLI - Interactive

```
# Login to Cannon
ssh <username>@login.rc.fas.harvard.edu

# Go to a compute node on the test partition:
salloc --partition test --nodes=1 --cpus-per-task=2 --mem=12GB --time=00:30:00

# Check Python modules available on Cannon:
module spider python

# Get detailed information on specific module, e.g.:
module spider python/3.10.13-fasrc01

# Load the latest (usually also the default) Python module:
module load python
```

CLI Training:

<https://docs.rc.fas.harvard.edu/wp-content/uploads/2013/10/Getting-started-on-FASRC-clusters-with-CLI.pptx.pdf>

Python using CLI - Interactive

- Check Python version: `python --version`
- Invoke Python interpreter: `python`
- Execute Python programming interactively:

```
def square(x):  
    """square a number"""  
    return x ** 2  
  
for N in range(1, 4):  
    print(N, "squared is", square(N))
```

- Exit Python: `exit()`
- Or run a python script interactively: `python myscript.py`

Python using CLI - sbatch; Example 1

https://github.com/fasrc/User_Codes/tree/master/Languages/Python/Example1

```
# Go to Example1 folder
cd Python/Example1
# Submit job
sbatch run.sbatch
```

run.sbatch: Batch-job submission script
for queuing the job

mc_pi.py: Source code for calculating
Pi using Monte-Carlo method

```
#!/bin/bash
#SBATCH -J mc_pi                # job name
#SBATCH -o mc_pi.out            # standard output file
#SBATCH -e mc_pi.err            # standard error file
#SBATCH --nodes=1               # number of nodes
#SBATCH --cpus-per-task=1       # number of cores
#SBATCH --partition=serial_requeue # partition
#SBATCH --time=0-00:30          # time in D-HH:MM
#SBATCH --mem=4000              # memory in MB

# Load required modules
module load python

# Run program
python mc_pi.py
```

Python Package Installation - Interactive

- Go to a compute node on the test partition:

```
salloc -p test --nodes=1 --cpus-per-task=2 --mem=12GB --time=01:00:00
```

- Create a vanilla mamba/conda environment (for multiprocessing exercise):

```
module load python  
mamba create --prefix=/n/hollylabs/LABS/<desired-folder>/multiproc_env python=3.11 -y
```

- Alternatively, if default *\$HOME* is desired, then do following instead:

```
module load python  
mamba create --name multiproc_env python=3.11 -y
```

- See [Python Package Installation](#)

Python Package Installation

- Activate conda/mamba environment:

```
mamba activate /n/holylabs/LABS/<desired-folder>/multiproc_env
```

- Or if \$HOME used, then:

```
mamba activate multiproc_env
```

- Install relevant python packages (Mamba recommended):

```
mamba install numpy pandas matplotlib -y  
pip install jupyterlab swifter
```

- Always pip install inside a conda environment to avoid package conflicts
- https://docs.rc.fas.harvard.edu/kb/python-package-installation/#Pip_Installs
- Deactivate the environment:

```
mamba deactivate
```

Python Package Installation - sbatch

https://github.com/fasrc/User_Codes/tree/master/Languages/Python/Example2

```
# Go to Example2 folder
cd ../Python/Example2
# Submit job
sbatch run.sbatch
```

numpy_pandas_ex.py: source code for
generating a dataframe utilizing a
mamba environment

```
#!/bin/bash
#SBATCH -J np_pandas           # job name
#SBATCH -o np_pandas.out       # standard output file
#SBATCH -e np_pandas.err       # standard error file
#SBATCH --cpus-per-task=1      # number of cores
#SBATCH --partition=test       # partition
#SBATCH --time=0-01:00         # time in D-HH:MM
#SBATCH --mem=10G              # memory in GB

# Load required modules
module load python

# Build the environment
sh build_env.sh

# Activate the environment
mamba activate my_env

# Run program
python numpy_pandas_ex.py
```

Is this always needed?

Python Package Maintenance

- Important for managing `$HOME` quota
- Remove an unused environment installed on `$HOME`:

```
[<username@computenode ~]>$ module load python  
[<username@computenode ~]>$ conda env remove --name <env-name> -y
```

- Remove unused tarballs, packages, index caches that get installed while creating an environment:

```
[<username@computenode ~]>$ module load python  
[<username@computenode ~]>$ conda clean --all -y
```

- For a specific env: <https://docs.rc.fas.harvard.edu/kb/home-directory-full/#conda>

Advanced Python & AI Packages

- Cuda-aware TensorFlow: <https://docs.rc.fas.harvard.edu/kb/tensorflow/>
- Cuda-aware PyTorch: <https://docs.rc.fas.harvard.edu/kb/pytorch/>
- Claude: <https://docs.rc.fas.harvard.edu/kb/claude/>
- OpenAI: <https://docs.rc.fas.harvard.edu/kb/openai/>
- [FASRC Guidelines for OpenAI Key and Harvard Agreement](#)

Python Using Open OnDemand (OOD)

- Open-source web portal to access clusters
- Web-based, no software needs be installed on your local laptop/desktop (except for a modern browser like Google Chrome, Mozilla Firefox)
- Easy to learn and simple to use
- Very similar to desktop applications
- The easiest way to run GUI applications remotely on a cluster
- Safari is not recommended for OOD
- OOD Training:
<https://docs.rc.fas.harvard.edu/wp-content/uploads/2013/10/Getting-started-on-FASRC-clusters-with-OOD-May2024.pdf>

How to access OOD on FASRC Clusters

- Accessing OOD from Cannon
 - Connect to FASRC VPN - [Virtual Desktop \(VDI\) through Open OnDemand – FASRC DOCS](#)
 - Then go to <https://rcood.rc.fas.harvard.edu>
- Accessing OOD from FASSE
 - Connect to FASSE VPN - [FASSE VDI Apps – FASRC DOCS](#)
 - Then go to <https://fasseood.rc.fas.harvard.edu>

FASSE proxy

Documentation: [FASSE Proxy Settings – FASRC DOCS](#)

- You may need to set FASSE proxy on
 - Firefox (web browsing)
 - Jupyter Notebook
 - Access Github
 - (Basically, anything outside of FASSE)

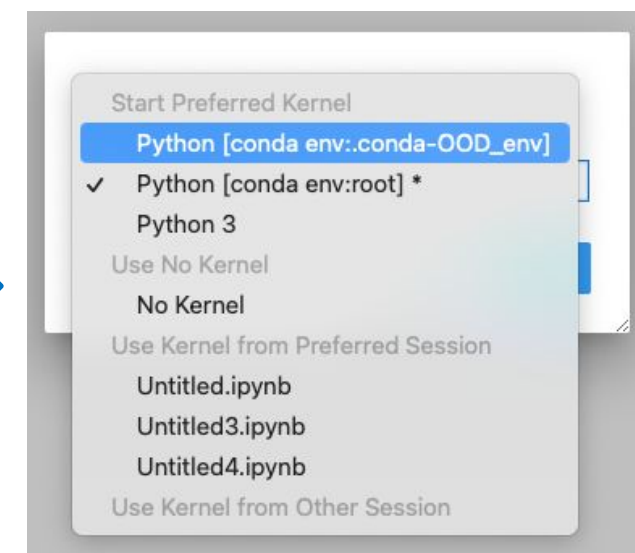
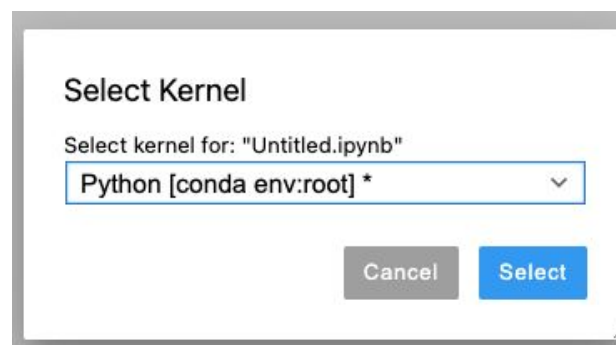
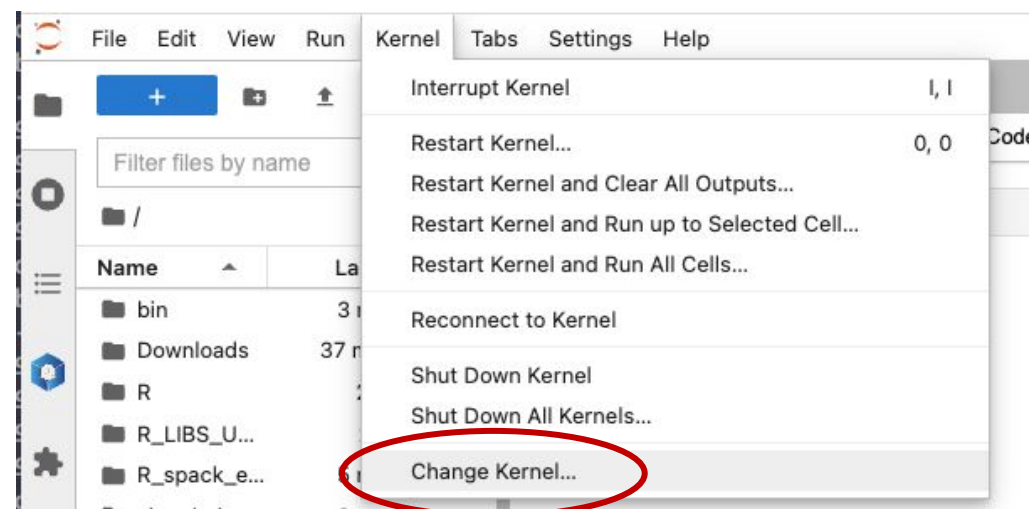
Filling a form to launch an app

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

Jupyter Notebook

- Launch **new** Jupyter Notebook session (existing session will not work!)
- Select newly created conda environment as the kernel
 - a. Open a notebook
 - b. On the top menu, click Kernel -> Select Kernel -> Click on OOD_env
 - c. Note: kernels is the same as conda, python, mamba environment



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: [Open OnDemand \(OOD/VDI\) Remote Desktop: How to open software – FASRC DOCS](#)

FASRC documentation

- FASRC docs: <https://docs.rc.fas.harvard.edu/>
- FASRC Python docs:
 - <https://docs.rc.fas.harvard.edu/kb/python/>
 - <https://docs.rc.fas.harvard.edu/kb/python-package-installation/>
- GitHub 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 (requires login)
 - Email: rchelp@rc.fas.harvard.edu

FASRC Upcoming Trainings

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

Python Multiprocessing on the FASRC cluster

Focused on some of the techniques to accelerate Python programming with emphasis on utilizing multiprocessing with numpy arrays.

Audience: Users who are familiar with basic Python, command line, HPC systems, and have attended our **Python: Package Installation & Execution on FASRC clusters training**.

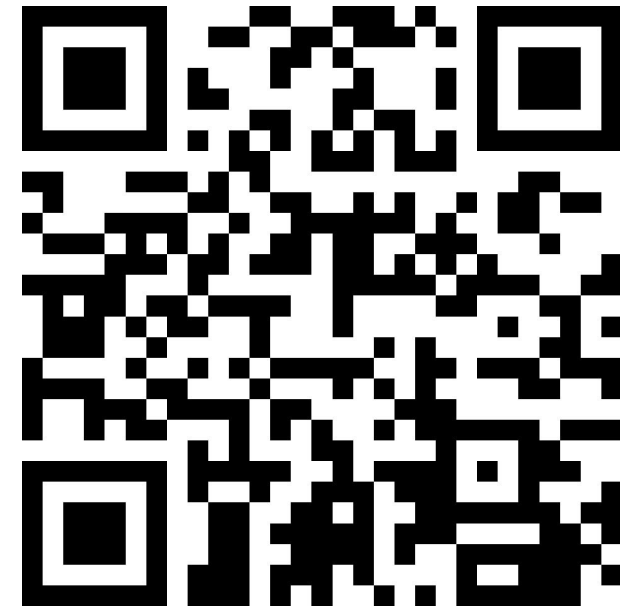
Objectives: Understanding Multiprocessing & Executing Multiprocessing on FASRC clusters

Note: All topics below are a brief overview of utilizing multiprocessing on FASRC clusters.

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>





Thank you :)
FAS Research Computing