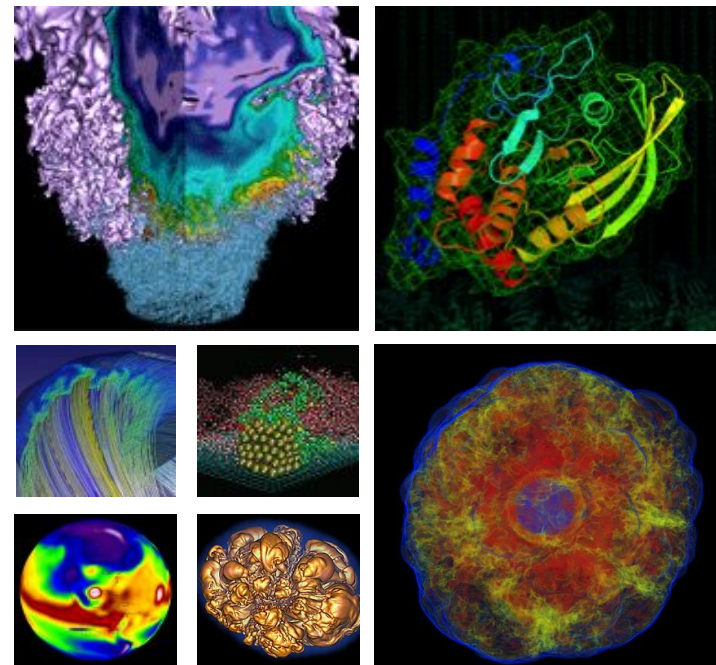


Python and Jupyter at NERSC



Rollin Thomas

Data Architect

Data and Analytics Services, NERSC

New User Training, 2019-01-25

Science via Python@NERSC

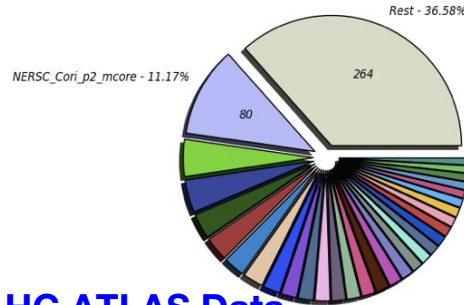


Powering Workflows to Understand Properties of Materials

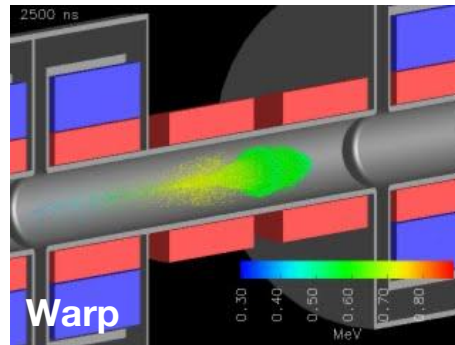
NBODYKIT
Modeling Dark Matter and Dark Energy



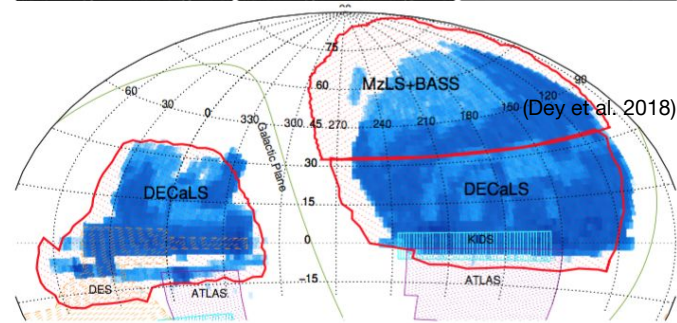
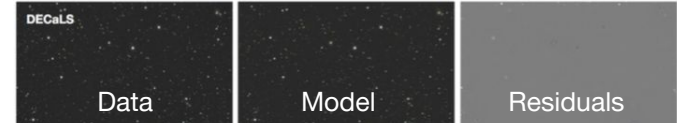
NEvents Processed in MEEvents (Million Events) (Sum: 723.00)



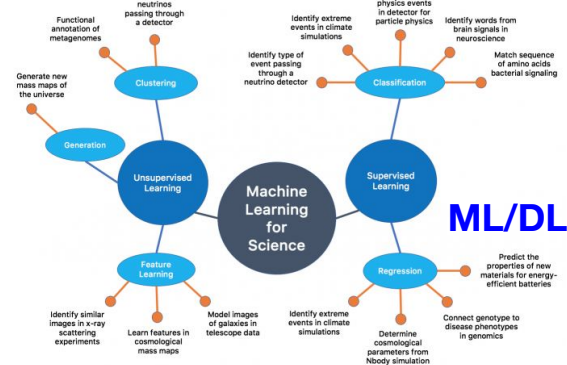
LHC ATLAS Data Processing Workflow



PIC Code for Plasmas and High Current Particle Beams



Sky Survey Catalogs for Cosmology



NERSC Python Documentation



Good docs advise on how to
use Python at NERSC.

Updates are ~continuous.

[Main page.](#)

Frequently Asked Questions

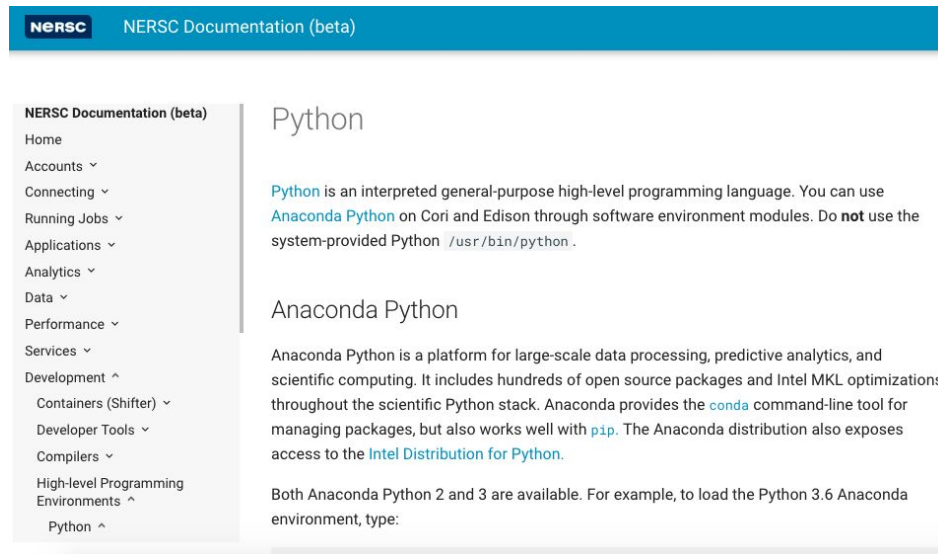
[FAQ page.](#)

Suggest new questions!

Advice/gotchas for KNL users

[KNL page.](#)

Advice on [optimizing Python.](#)



New site: docs.nersc.gov

Use Environment Modules

NERSC

Environment modules:

Environment modules project:

<http://modules.sourceforge.net/>

Always* “module load python”

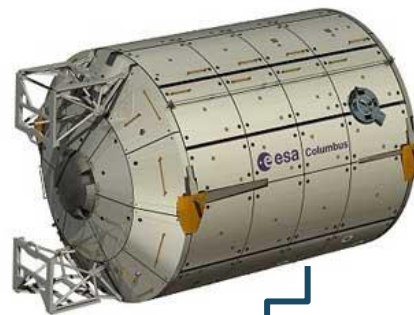
Do not use /usr/bin/python.

Using #!/usr/bin/env python is OK!

What is there?

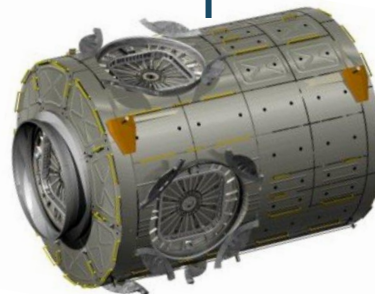
module avail python

* Unless you install your own somehow.
(Totally fine, see later in the talk.)



python/2.7-anaconda-4.4

python/3.6-anaconda-4.4



NERSC's Python is Anaconda



NERSC's builds of Python on Cori/Edison have been retired for a while.

Modules Now Leverage Anaconda Python

Distro for large-scale data analytics and scientific computing.

Handy package management and deployment (conda tool).

Conda environments replace virtualenv.

Hundreds of useful packages (400+ already installed)

Threaded Intel MKL comes for free.

Now with some ML tools too.

Additional “channels” and you can still use pip.



<https://docs.anaconda.com/anaconda/>

They are monolithic; with some add-on modules (h5py-parallel).

NERSC Python Modules



Recommended environment modules at NERSC for Python users:

```
module load python/2.7-anaconda-4.4
```

```
module load python/3.6-anaconda-4.4
```

Default Python is 2.7 up to no later than **11** months from now:

```
module load python
```

```
[= module load python/2.7-anaconda-4.4]
```

Conda Environments



Conda makes it easy to create tailored environments with the packages you need.

```
module load python/3.6-anaconda-4.4
conda create -n myenv python=2 numpy
[installation outputs]
source activate myenv
```

**And pip is OK to use too. Note,
“--no-cache-dir” is handy**

Don't bother with --user, just pip in your conda env.



Doing Things Yourself



Project-wide Anaconda installation, e.g. at `/global/common/software/<project-name>`

```
module unload python
unset PYTHONSTARTUP
```

```
wget https://repo.continuum.io/miniconda/Miniconda2-latest-Linux-x86_64.sh
/bin/bash Miniconda2-latest-Linux-x86_64.sh -b -p $PREFIX
source $PREFIX/bin/activate
    <or export PATH=$PREFIX/bin:$PATH>
conda install basemap yt...
```



Building your own mpi4py or parallel h5py?

Do not conda install ...

Do not pip install ...

Link to Cray MPICH, using compiler wrappers

```
wget https://bitbucket.org/mpi4py/mpi4py/downloads/mpi4py-3.0.0.tar.gz
tar zxvf mpi4py-3.0.0.tar.gz
cd mpi4py-3.0.0
module swap PrgEnv-intel PrgEnv-gnu
python setup.py build --mpicc=$(which cc)
python setup.py install
```



Parallelism with Python



Within a node:

Use OpenMP-threaded math libs.
Multiprocessing is OK too.



Multi-node parallelism:

Best supported by mpi4py.
Dask, PySpark work too.

Hybrid parallelism:

Best route is mpi4py + threaded math libs.

Handling MPI with mpi4py



Cluster parallelism with MPI via mpi4py:

MPI-1/2/3 specification support

OO interface ~ MPI-2 C++ bindings

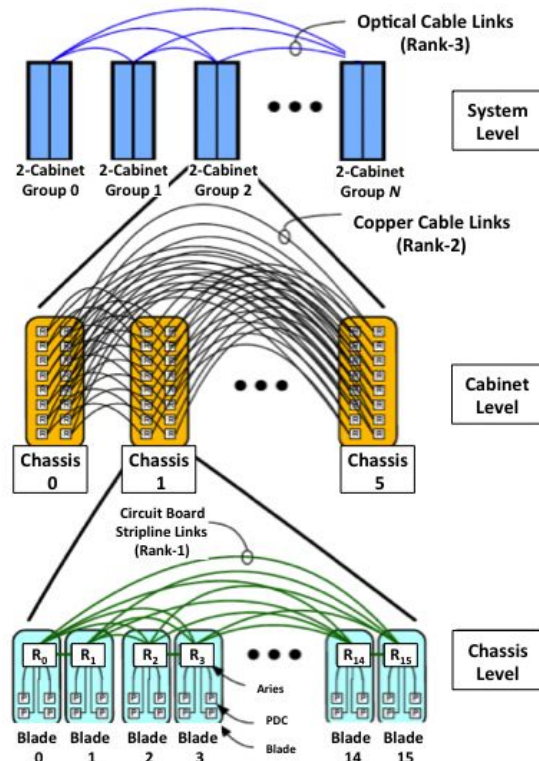
Point-to-point and collectives

Picklable Python objects & buffers

Build mpi4py & dependents with
Cray MPICH:

```
python setup.py build --mpicc=cc  
python setup.py install
```

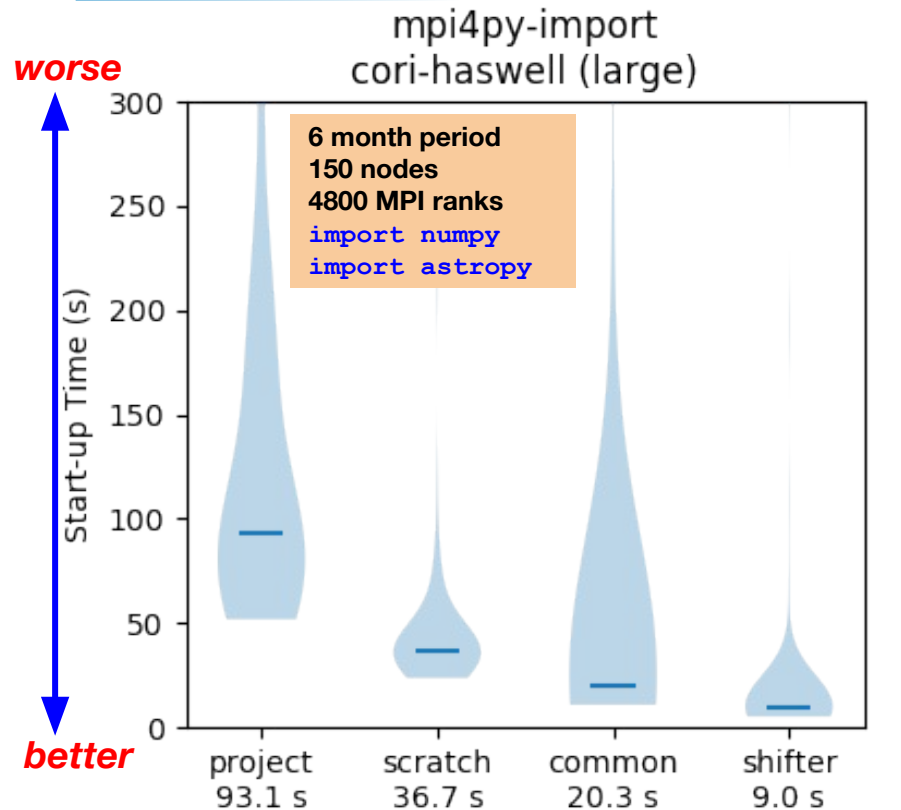
↑
Cray-provided
Compiler wrapper



Cori Aries Interconnect

Python “Slow Launch” at Scale

NERSC



[Median launch time incl. MPI_Init()]

Python’s import is metadata intensive,
⇒ catastrophic contention at scale
⇒ *it matters where you install your env*

Project (GPFS):

For sharing large data files

Scratch (Lustre):

OK, but gets purged periodically!

Common (GPFS):

RO w/Cray DVS client-side caching

Open to users now, was only staff

Shifter (Docker Containers):

Metadata lookup only on compute

Storage on compute is RAM disk

Idconfig when you build image



SHIFTER



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Profiling, Debugging



Ye olde stand-bye, `print()`!

`srunch -u python -u <script-name> ...`

Unbuffer both `srunch` and `python`.

Can be a lot of messy output to parse.

Good for general exploration (standard lib):

`cProfile` plus `snakeviz` or `gprof2dot`

MPI processes? [\[see an example here\]](#)

Good for a deeper dive on one function (package):

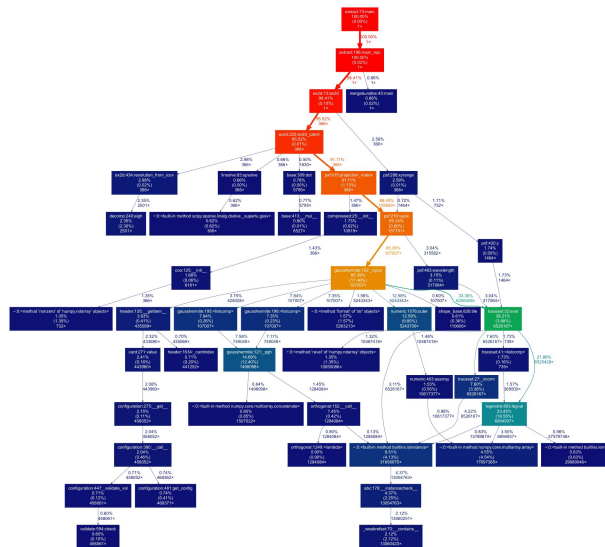
`line_profiler`

High-performance instrumented timer (mixed-language, MPI, package):

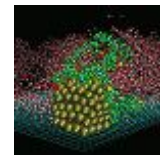
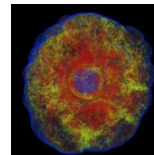
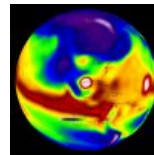
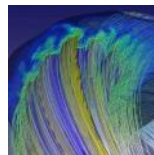
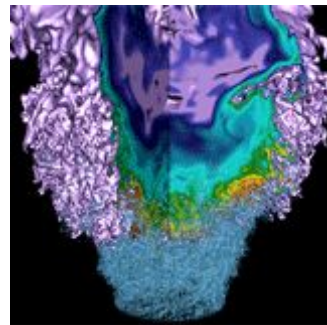
`TiMemory`

High-performance tools (mixed-language, MPI):

Intel VTune (some collection methods) on Intel Python, and Tau



Jupyter at NERSC



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Using Jupyter at NERSC



Jupyter Notebook: “Literate Computing.”
Code, text, equations, viz in a narrative.

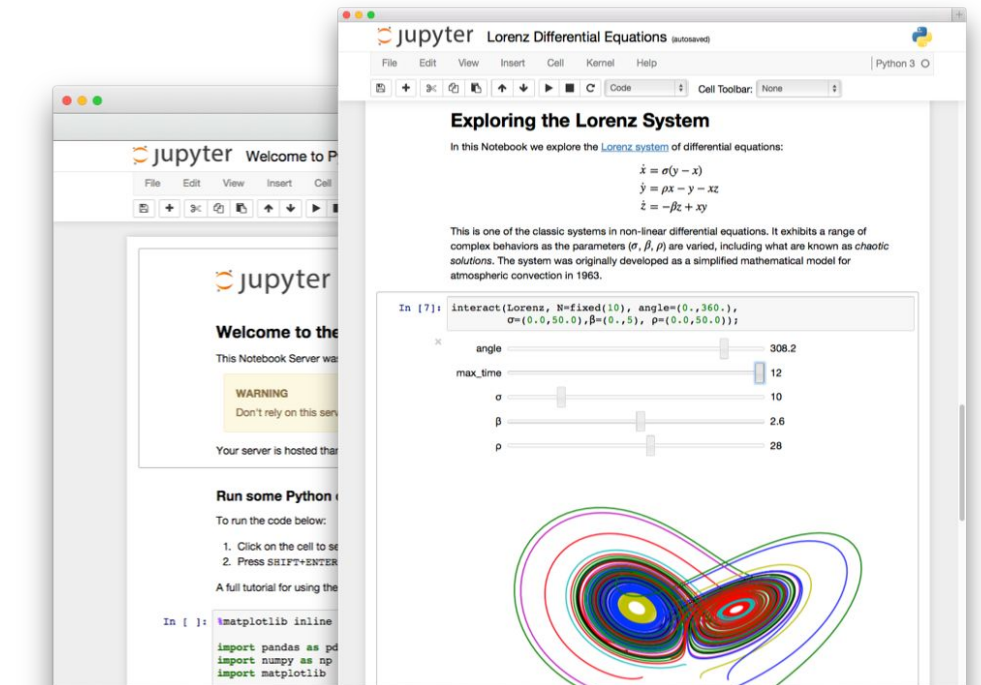
Two Hubs for Jupyter at NERSC.

jupyter.nersc.gov:

External to Edison/Cori
Can't see \$SCRATCH
But can see /project, \$HOME

jupyter-dev.nersc.gov:

Spawns notebooks on Cori
Can see Cori \$SCRATCH
Same Python env as ssh login
Can submit jobs via `%sbatch`



Your Own Jupyter-dev Kernel



Most common Jupyter question:

“How do I take a conda environment and turn it into a Jupyter kernel?”

Several ways to accomplish this, here's the easy one.

```
$ module load python
$ conda create -n myenv python=3.6
$ source activate myenv
(myenv) $ conda install ipykernel <other-packages>...
(myenv) $ python -m ipykernel install --user --name myenv-jupyter
```

Point your browser to jupyter-dev.nersc.gov.

(You may need to restart your notebook server via control panel).

Kernel “myenv-jupyter” should be present in the kernel list.



This creates a
“kernel spec” file.

The kernelspec File



```
(myenv) rthomas@cori01:~> cat \  
    $HOME/.local/share/jupyter/kernels/myenv-jupyter/kernel.json  
{  
  "argv": [  
    "/global/homes/r/rthomas/.conda/envs/myenv/bin/python",  
    "-m",  
    "ipykernel_launcher",  
    "-f",  
    "{connection_file}"  
  ],  
  "display_name": "myenv-jupyter",  
  "language": "python"  
}
```

Additional Customization



```
{  
  "argv": [  
    "/global/homes/r/rthomas/.conda/envs/myenv/bin/python",  
    "-m",  
    "ipykernel_launcher",  
    "-f",  
    "{connection_file}"  
  ],  
  "display_name": "myenv-jupyter",  
  "language": "python",  
  "env": {  
    "PATH": ...,  
    "LD_LIBRARY_PATH": ...,  
  }  
}
```

Additional Customization



```
{  
  "argv": [  
    "/global/homes/r/rthomas/jupyter-helper.sh",  
    "-f",  
    "{connection_file}"  
  ],  
  "display_name": "myenv-jupyter2",  
  "language": "python",  
}
```



Meanwhile, in jupyter-helper.sh:

```
#!/bin/bash  
export SOMETHING=123  
module load texlive  
exec python -m ipykernel "$@"
```

A Shifter Kernelspec



```
{  
  "argv": [  
    "shifter",  
    "--image=continuumio/anaconda3:latest",  
    "/opt/conda/bin/python",  
    "-m",  
    "ipykernel_launcher",  
    "-f",  
    "{connection_file}"  
  ],  
  "display_name": "my-shifter-kernel",  
  "language": "python"  
}
```

Diagram illustrating the components of the Shifter Kernelspec:

- shifter**: The command to run the Shifter kernel.
- image=continuumio/anaconda3:latest**: The image name used for the kernel.
- /opt/conda/bin/python**: The path in the image to the Python interpreter.
- m**: The flag to run a module.
- ipykernel_launcher**: The module to run.
- f**: The flag to specify the connection file.
- {connection_file}**: The connection file path.



SHIFTER

Debugging Jupyter Stuff

NERSC

(myenv) rthomas@cori01:~> cat ~/jupyter.log ← **YOUR FRIEND**

```
[I 2018-03-19 16:00:08.175 SingleUserNotebookApp manager:40] [nb_conda_kernels] enabled, 5 kernels found
[I 2018-03-19 16:00:08.248 SingleUserNotebookApp extension:53] JupyterLab beta preview extension loaded from
/usr/common/software/python/3.6-anaconda-4.4/lib/python3.6/site-packages/jupyterlab
[I 2018-03-19 16:00:08.248 SingleUserNotebookApp extension:54] JupyterLab application directory is
/global/common/cori/software/python/3.6-anaconda-4.4/share/jupyter/lab
[I 2018-03-19 16:00:09.123 SingleUserNotebookApp handlers:73] [nb_anacondacloud] enabled
[I 2018-03-19 16:00:09.129 SingleUserNotebookApp handlers:292] [nb_conda] enabled
[I 2018-03-19 16:00:09.181 SingleUserNotebookApp __init__:35] ✓ nbpresent HTML export ENABLED
[W 2018-03-19 16:00:09.181 SingleUserNotebookApp __init__:43] ✗ nbpresent PDF export DISABLED: No module
named 'nbbrowserpdf'
[I 2018-03-19 16:00:09.186 SingleUserNotebookApp singleuser:365] Starting jupyterhub-singleuser server
version 0.8.0.rc1
[I 2018-03-19 16:00:09.190 SingleUserNotebookApp log:122] 302 GET /user/rthomas/ →
/user/rthomas/tree/global/homes/r/rthomas? (@128.55.206.24) 0.62ms
[I 2018-03-19 16:00:09.194 SingleUserNotebookApp notebookapp:1445] Serving notebooks from local directory: /
[I 2018-03-19 16:00:09.194 SingleUserNotebookApp notebookapp:1445] 0 active kernels
[I 2018-03-19 16:00:09.194 SingleUserNotebookApp notebookapp:1445] The Jupyter Notebook is running at:
[I 2018-03-19 16:00:09.194 SingleUserNotebookApp notebookapp:1445] http://0.0.0.0:56901/user/rthomas/
[I 2018-03-19 16:00:09.194 SingleUserNotebookApp notebookapp:1446] Use Control-C to stop this server and shut
down all kernels (twice to skip confirmation).
[I 2018-03-19 16:00:09.236 SingleUserNotebookApp log:122] 302 GET /user/rthomas/ →
/user/rthomas/tree/global/homes/r/rthomas? (@::ffff:10.42.245.15) 0.39ms
```



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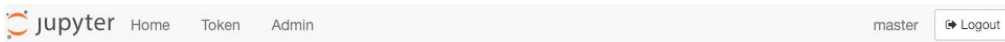


Future Jupyter Support



Working on:

- ❖ Expanding resources to support Jupyter-Dev
- ❖ New ways to launch parallel workloads managed through Jupyter
- ❖ Merging Jupyter and Jupyter-Dev into one login interface
- ❖ Expanding JupyterLab interface to:
 - Track and monitor batch jobs
 - New viewers



NERSC JupyterHub Console

	Shared CPU Node	Exclusive CPU Node	Exclusive GPU Node	Configurable
Perlmutter	start	start	start	start
Cori	start	start	start	start
Spin	start			
Resources	On a node shared with other users' notebooks but outside the batch queues.	On a node by itself within an interactive job allocation using your default repo.	One or more nodes within an interactive job allocation.	
Use Cases	Visualization and analytics that are not memory intensive and can run on just a few cores.	Visualization, analytics, machine learning that is compute or memory intensive but can be done on a single node.	Large-scale data analytics, visualization, and machine learning; reservation or non-default repository.	

Python and Jupyter at NERSC



**Python & Jupyter: integral elements of NERSC's
Data Intensive Science portfolio.**

We want users to have a:

<i>familiar</i>	Python environment
<i>productive</i>	Python experience
<i>performant</i>	Python software stack

Always looking for:

New ways to empower Python & data science users.

Feedback, advice, and even help:

<https://help.nersc.gov/>
rcthomas@lbl.gov



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Thank You



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