

Integrating AGC pipeline at BNL facility

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Introduction and Overview

- This work is really all thanks to BNL team
 - Doug Benjamin
 - Ofer Rind
 - Chris Hollowell
- Ongoing process, but today just showing the first (unoptimized) steps that came together pleasantly quickly
- Today showing examples of running at BNL's SDCC Jupyter instance

BNL SDCC Jupyter Launcher: custom images!

jupyterhub Home Token atlas_feickert Logout

SDCC Jupyter Launcher

HTC / Standard HTCondor Pool IC / HPC Systems

Run a notebook on a standard interactive HTCondor submit-node

Select JupyterLab Environment

Default
 Default HPC
 USATLAS

Singularity Container

None
 Custom

Start

Allows for running custom images as Singularity containers
Can pull from public image registries or from CVMFS unpacked

Development image: analysis-systems-base

- As we can use custom images at BNL created the `analysis-systems-base` image (<https://github.com/iris-hep/analysis-systems-base>)
- Images are hosted on **OSG Harbor** under `iris-hep.org`
 - `hub.opensciencegrid.org/iris-hep/analysis-systems-base`
 - Thanks Brian Lin for making this happen!

analysis-systems-base

Base Docker image for Analysis Systems environment

Get Image

Open Science Grid Harbor registry

The images are stored on the [Harbor image registry](#)

```
docker pull hub.opensciencegrid.org/iris-hep/analysis-systems-base:latest
```

CVMFS Unpacked

The images are also available through CVMFS unpacked and are available on CVMFS instances under the path

```
/cvmfs/unpacked.cern.ch/hub.opensciencegrid.org/iris-hep/analysis-systems-base:<tag>
```

Usage

Running as non-root user

Run the container with configuration options

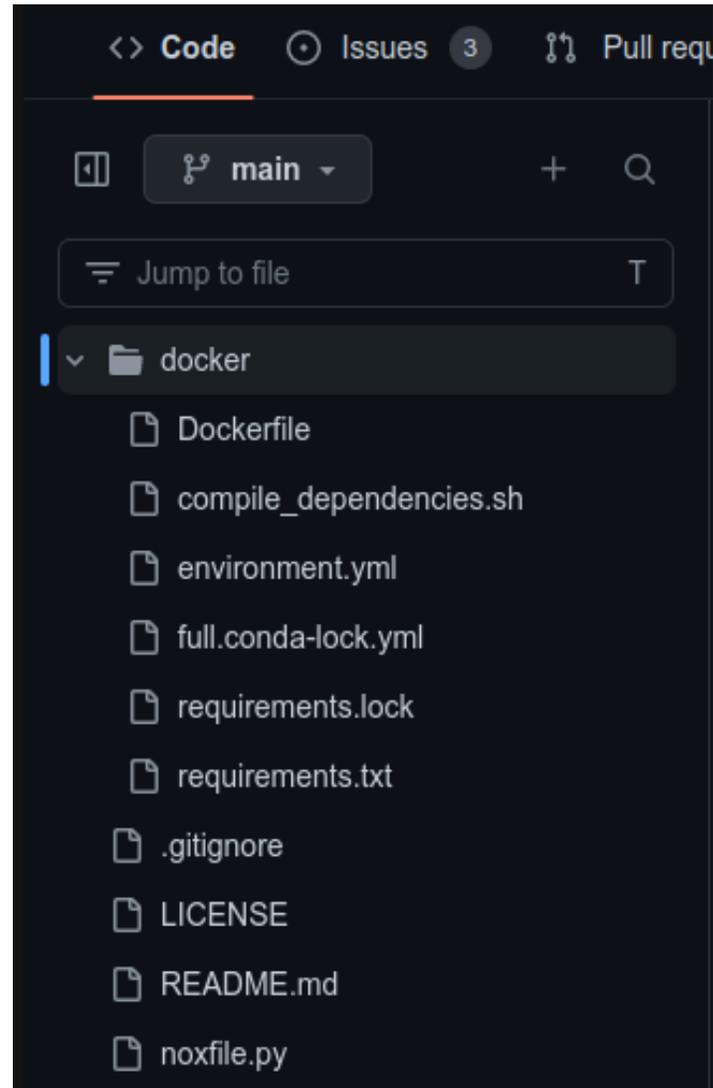
```
docker run \  
  --rm \  
  -ti \  
  --publish 8888:8888 \  
  --user $(id -u $USER):$(id -g) \  
  hub.opensciencegrid.org/iris-hep/analysis-systems-base:latest
```

which will then launch Jupyter Lab with corresponding option defaults

```
jupyter lab --no-browser --ip 0.0.0.0 --port 8888
```

Development image: analysis-systems-base

- As we can use custom images at BNL created the `analysis-systems-base` image (<https://github.com/iris-hep/analysis-systems-base>)
- Goal is to use lock files to make as much of the image as fully reproducible as possible and statically defined.



BNL SDCC Jupyter Launcher: custom images!

SDCC Jupyter Launcher

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Singularity Container

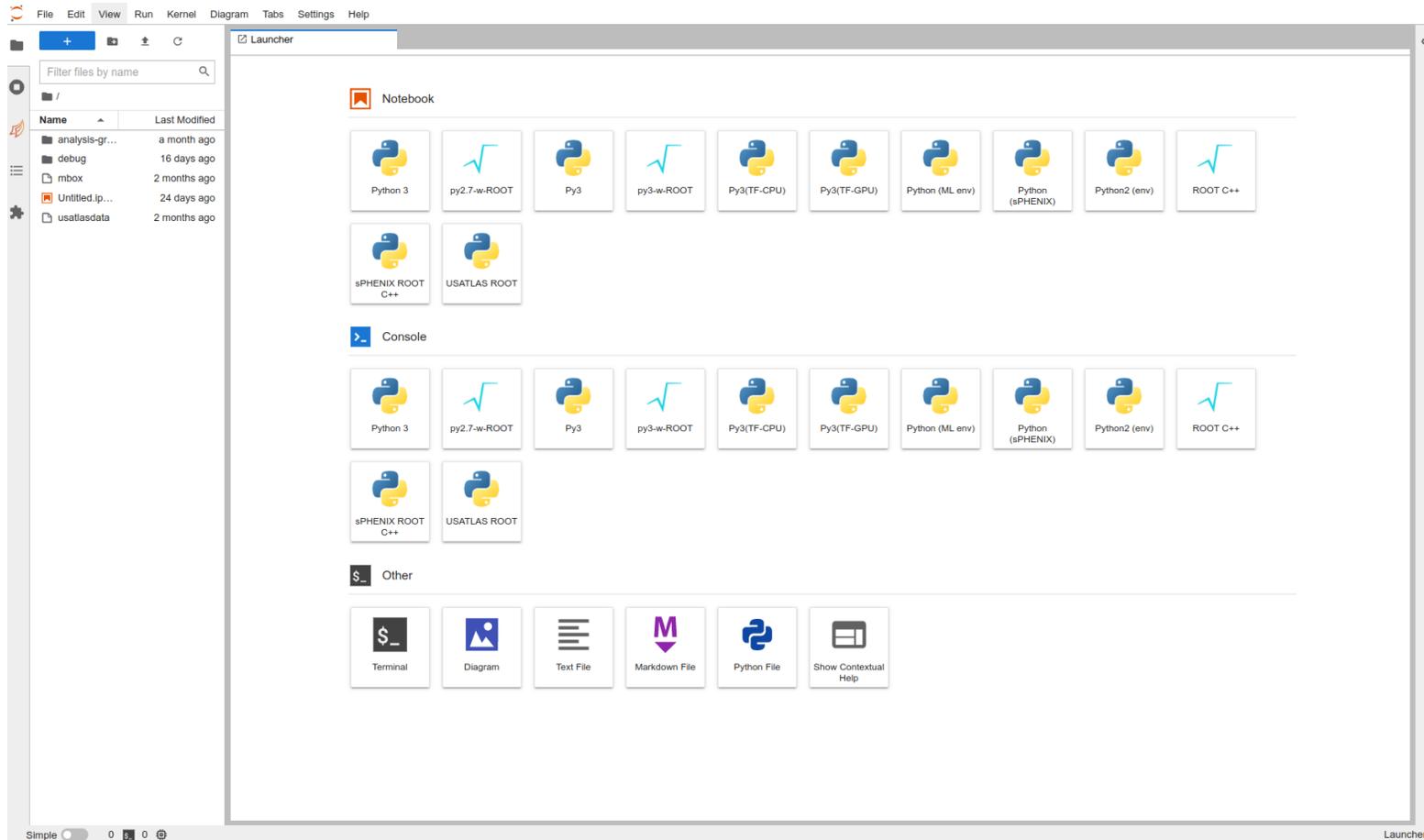
- None
- Custom

Start

Today's demo:

`/cvmfs/unpacked.cern.ch/hub.opensciencegrid.org/iris-hep/analysis-systems-base:2022-12-15`

WIP: Custom kernel discovery



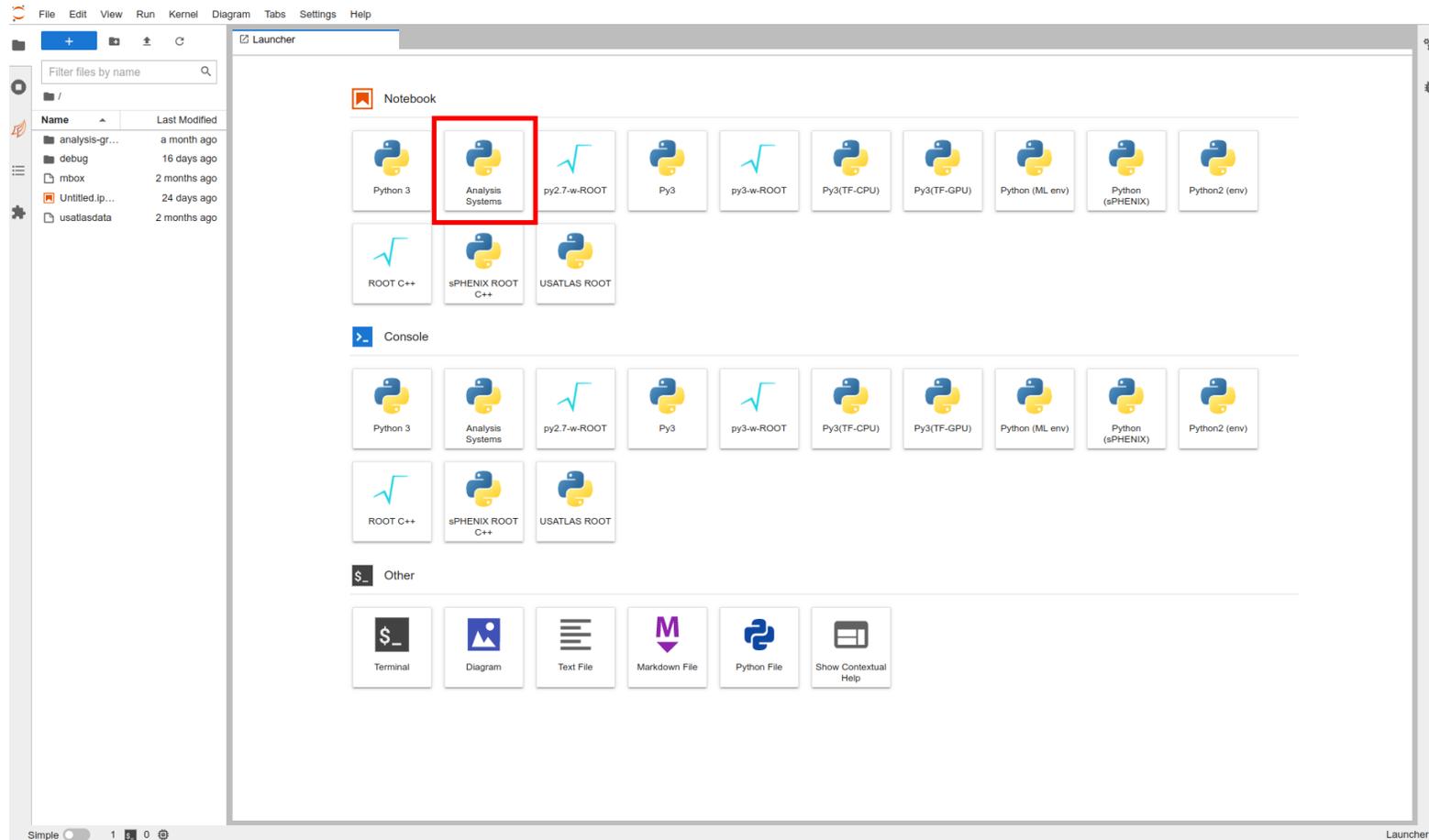
At the moment, lacking mounting (or something else) to mount custom kernels from user defined virtual environments

WIP: Custom kernel discovery

```
Singularity> mkdir -p ~/.local/share/jupyter/kernels
Singularity> ln --symbolic \
    /opt/micromamba/envs/analysis-systems/share/jupyter/kernels/analysis-systems \
    ~/.local/share/jupyter/kernels
Singularity>
```

Hack for time being: Create custom symlink first time
(c.f. <https://github.com/iris-hep/analysis-systems-base/issues/12>)

WIP: Custom kernel discovery



Hack for time being: Create custom symlink first time
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Example: AGC CMS Open Data $t\bar{t}$ Analysis

- Doug has moved data to
/usatlas/atlas01/atlasdisk/users/benjamin/AGC/
- Notebook runs end-to-end (good first start 🚀)
- BNL does not use Kubernetes, so not a Coffea-casa AF, so use global config

...

```
PIPELINE = "coffea" # pure coffea setup
```

```
USE_DASK = True # enable Dask
```

```
AF = "local" # local setup, not coffea-casa
```

```
AF_NAME = "bnl" # Added on Matthew's fork
```

...

Example: AGC CMS Open Data $t \bar{t}$ Analysis

- Execute the data delivery pipeline step

```
N_FILES_MAX_PER_SAMPLE = 10 # 157 GB
```

```
...
```

```
[#####] | 100% Completed | 1min 32.6s  
execution took 93.11 seconds
```

```
N_FILES_MAX_PER_SAMPLE = 50 # 678 GB
```

```
...
```

```
[#####] | 100% Completed | 6min 14.1s  
execution took 375.80 seconds
```

```
N_FILES_MAX_PER_SAMPLE = 100 # 1 TB
```

```
...
```

```
[#####] | 100% Completed | 10min 10.0s  
execution took 611.55 seconds
```

Example: AGC CMS Open Data $t \bar{t}$ Analysis

- Execute the data delivery pipeline step
- Nothing that I've done at BNL has been optimized yet (just doing defaults)
- **N.B.** Need to get better information on how scaling is done at BNL (so comparisons are not valid here yet)
 - UChicago AF is running with $AF = \text{"coffea_casa"}$ and so scaling across k8

Data	BNL AF (sec)	UChicago AF (sec)
157 GB	93	65
678 GB	375	182
1 TB	611	243

(The input files are all in the 1-2 GB range)

Comparison of unoptimized numbers (don't try to 1:1 these)

Summary

- Have a runnable environment for AGC at BNL AF (🚀)
- BNL team is working to make drop-in with custom images well specified and easy
- Things work, but to understand how scaling is working will need to improve the monitoring story
 - Currently don't have Dask dashboards detecting the Dask cluster
- Also need to try other AGC analyses and expand / modify environment

Backup

