

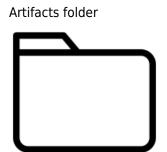
mlflow with Run:ai mlflow overview



What is needed for mlflow to run?

- 1) A Database to store information related to experiment runs
- 2) An Artifacts folder to store objects related to the runs







How does mlflow work?

1) First we start the server with a CLI command

running either of these commands will automatically create a databse in the local directory, if one doesn't exist. mlflow ui
(or)
mlflow server

we can also choose to specify the location of the database, and artifact folder, as well as the host IP, and port.

```
mlflow server \
```

- --backend-store-uri=sqlite:///abs/path/to/db/mlflow.db \
- --default-artifact-root=/abs/path/to/artifacts \
- --host=0.0.0.0\
- --port=5000



How does mlflow work?

2) call mlflow commands within the python script

you'll want to import mlflow, then set the tracking uri so that mlflow will save everything to the database and artifact folder.

Then you'll want to start your run, and at the end, you'll want to end the run.

```
import mlflow

mlflow.set_tracking_uri('0.0.0.0:5000')
mlflow.start_run()

"""

your code here
"""

mlflow.end_run()
```

mlflow with run:ai



What is needed for mlflow to run on run; ai?

- 1) A persistent directory to keep
- mlflow database
- mlflow artifacts folder

- 2) A docker image with the following installed
- mlflow
- jupyterlab*
- jupyter-server-proxy*

^{*}needed in order to access the mlflow UI

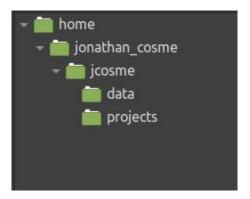
creating persistent directory



Before we start

We need to create an 'mlflow' folder on our NFS.

This is what our NFS folder structure looks like now



This is the default folder structure for our jupyter lab image

```
home
joyvyan
work
```

Before we start

We need to create an 'mlflow' folder on our NFS.

For our example:

Whenever we create a job on run:ai, we *must* always mount our NFS to the default jupyter work directory





Before we start

We need to create an 'mlflow' folder on our NFS.

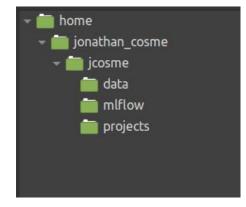
1. After we mount our NFS volume, our work directory will look like this:



2. Using Jupyter Lab, we create an 'mlflow' folder within the work directory:



3. This will cause our NFS directory to automatically look like this:



docker image



Docker used in our example

The docker image we will use is: jonathancosme/mlflow-ui

This is what is in the dockerfile:

```
RUN mamba install -c conda-forge mlflow -y && \
mamba clean --all -f -y install git (needed by mlflow)

RUN sudo apt-get update && \
sudo apt-get install -y git copy new config file, with mlflow UI access configured
```



Docker used in our example

The docker image we will use is: jonathancosme/mlflow-ui

in order to access the mlflow UI, we need to add this entry to the jupyter_server_config.py file, and replace the existing file in the image

we specify our database and artifact folder locations (this is why we must always mount our NFS directory to the default jupyter work directory)



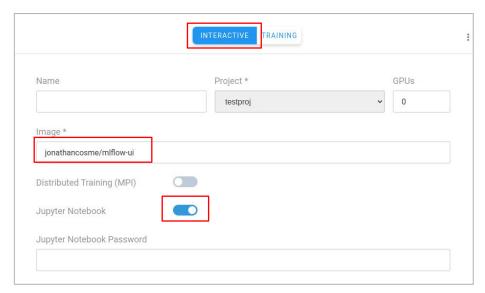
Accessing the mlflow UI



Access mlflow UI

Create a jupyter interactive job with:

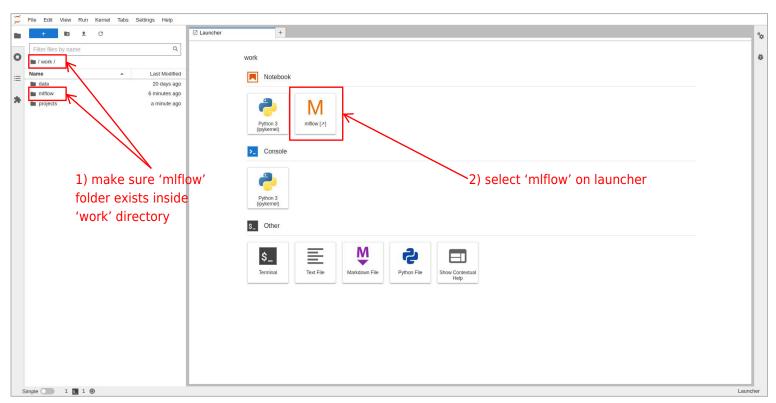
- image jonathancosme/mlflow-iu
- mounted NFS folder (with 'mlflow' folder) in default jupyter work directory





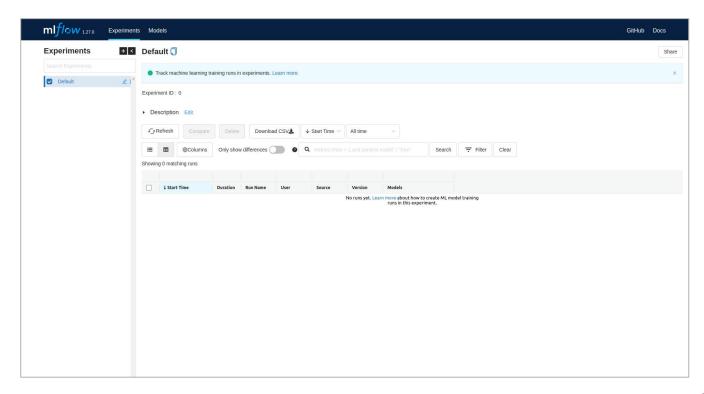


Access mlflow UI



Access mlflow UI

A new tab should appear with the mlflow UI





running mlflow experiments with run:ai



Python scripts

in your script, you should first launch the mlflow server

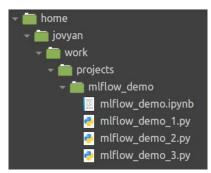
Then be sure to set the tracking uri

```
import mlflow
import os
import time
from subprocess import Popen
# first we start the mlflow server
mlflow command = 'mlflow server --backend-store-uri=sqlite:////home/jovyan/work/mlflow/
    mlflow.db --default-artifact-root=/home/jovyan/work/mlflow/artifacts --host=0.0.0.0
    --port=5000 --serve-artifacts'
print(f'running command: {mlflow command}')
proc = Popen([mlflow command], shell=True,
             stdin=None, stdout=None, stderr=None, close fds=True)
print(f'waiting 5 seconds...')
time.sleep(5) # we wait 5 seconds to give the server time to start
# create the tracking uri string
uri = 'http://0.0.0.0:5000' # in our example, this is always the same
project name = 'test project'
run name = 'run 1'
mlflow.set tracking uri(uri)
mlflow.set experiment(experiment name=project name)
mlflow.start run(run name=run name)
your code here
mlflow.end run()
```



CLI submission

Our example scrips are located here:



so our CLI command would look like this:

```
runai submit \
--project testproj \
--gpu 0 \
--job-name-prefix mlflow-demo \
--image jonathancosme/mlflow-ui \
--volume /home/jonathan_cosme/jcosme:/home/jovyan/work \
-- pythom work/projects/mlflow_demo/mlflow_demo_1.py
```

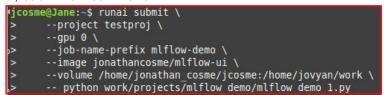
make sure you use the mlflow-ui docker image

make sure you mount the NFS to the work directory

The code to run the job must specify the python script with relative location

Example job submission

1) submit CLI command



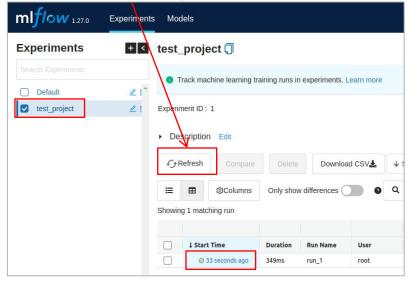
2) a new job should appear



3) wait for job to finish



4) refresh mlflow UI to see updates





Thank you!

