# Stage 1: Data preparation (Day 1: Initial setup)

#### Main repository link:

• On top right corner next to the search bar link for the main repo is given or click here to see the main repo



The main repo contains various branch so make sure you select a branch which has the latest commit to see the recent updates.

#### Steps and important commands to begin-



Replace the text mentioned as <some\_txt> with your preferred choice.

#### STEP 1 Create a new conda environment

· Open an anaconda prompt and create an environment -

· Activate the environment -

```
conda activate <your_env_name>
wafer2
```

### STEP 2 Create a default structure

• Install cookiecutter template

```
pip install cookiecutter
```

· Start a new project

```
cookiecutter https://github.com/drivendata/cookiecutter-data-science
```

• After above step you'll be given options in the command line.

```
a. project_name: mlops
b. repo_name: mlops2
c. author_name: zymxiaotie
d. description: replicate mlops_main to solve app issues
e. Select open_source_license: MIT
f. s3_bucket [Optional]:
g. Select python_interpreter: python3
```

Once you are done with above step you'll see a following directory structure inside a directory by your given project\_name

```
— LICENSE
                 <- Makefile with commands like `make data` or `make train`
 — Makefile
                <- The top-level README for developers using this project.
 README.md
 — data
   └─ raw
                  <- The original, immutable data dump.
 - docs
                  <- A default Sphinx project; see sphinx-doc.org for details
                  <- Trained and serialized models, model predictions, or
- models
model summaries
                  <- Jupyter notebooks. Naming convention is a number (for
─ notebooks
ordering),
                     the creator's initials, and a short `-` delimited
description, e.g.
                     `1.0-jqp-initial-data-exploration`.
- references
                  <- Data dictionaries, manuals, and all other explanatory
materials.
 — reports
                  <- Generated analysis as HTML, PDF, LaTeX, etc.
 └─ figures
                  <- Generated graphics and figures to be used in reporting
├─ requirements.txt <- The requirements file for reproducing the analysis
environment, e.g.
                     generated with `pip freeze > requirements.txt`
├─ setup.py
                  <- makes project pip installable (pip install -e .) so src
can be imported
                  <- Source code for use in this project.
   ├─ __init__.py <- Makes src a Python module
   — data
                  <- Scripts to download or generate data
     └─ make_dataset.py
    └─ build_features.py
                  <- Scripts to train models and then use trained models to
   — models
make
                     predictions
      ├─ predict_model.py
      └─ train_model.py
   └─ visualization <- Scripts to create exploratory and results oriented
visualizations
      └─ visualize.py
```

```
└─ tox.ini <- tox file with settings for running tox; see tox.readthedocs.io
```

Now open the project in your favorite code editor.

#### STEP 3 Get the dataset

· Clone it from the dataset repository or directly-

Download Dataset

 Now extract the Prediction\_Batch\_files, Training\_Batch\_Files directory in the root directory of the project

STEP 4 Initialize git in Current working directory in your terminal, command prompt or git bash.

\$cd C:\Users\zymxi\mlops2
git init



If git is not installed in your system then download it from GIT-SCM site

## STEP 5 Install DVC and its gdrive extension

pip install dvc
pip install dvc[gdrive]

#### STEP 6 Initialize DVC

dvc init

### STEP 7 Add data into dvc for tracking

dvc add Training\_Batch\_Files/\*.csv Prediction\_Batch\_files/\*.csv

#### A

#### Warning

Above command will not work for windows users so they can create and run the following file in the root of their project

```
# NOTE: For windows user-
# This file must be created in the root of the project
# where Training and Prediction batch file as are present

import os
from glob import glob

data_dirs = ["Training_Batch_Files", "Prediction_Batch_files"]

for data_dir in data_dirs:
    files = glob(data_dir + r"/*.csv")
    for filePath in files:
        # print(f"dvc add {filePath}")
        os.system(f"dvc add {filePath}")

print("\n #### all files added to dvc ####")
```

# STEP 8 Do the first commit and push to the remote repository

run below commands on by one-

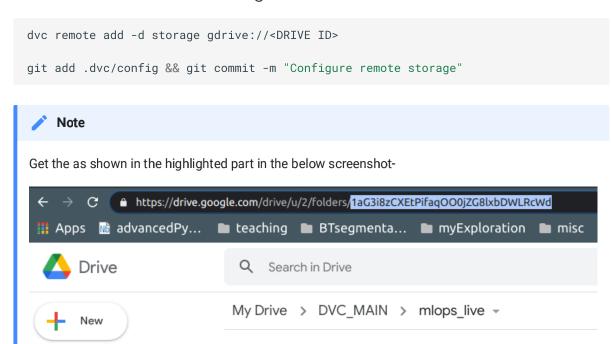


replace <USERNAME> and <REPONAME> as per you.

# STEP 9 Create and checkout a development branch for our development

```
git checkout -b dev
```

### STEP 10 Add remote storage



#### STEP 11 Push the data to the remote storage-

dvc push

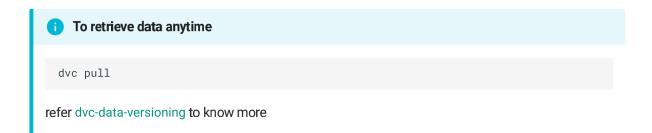
- This step will ask you to authenticate yourself by clicking on the link which will appear in the terminal.
- Once you allow dvc to read and write on gdrive it'll give an access token which you'll paste in the terminal.
- Now the copy of your data will be pushed to the gdrive
- Above step will create a gdrive credential file (Now check next step).

### STEP 12 Add Gdrive credential secrets in github repo secrets.

· Find this credentials in the given path -

```
I didn't find this file, skip step 12
.dvc >> temp >> gdrive-user-credentials.json
```

- · Now to add the secrets in your github repo -
  - · Go to settings
  - secrets
  - · Click on add secrets
  - · Give name of secretes
  - Paste the json file content from gdrive-user-credentials.json



STEP 13 Install full requirements.txt as given in the repository

```
i One line readme update and push command to dev branch-
git add README.md && git commit -m "update readme" && git push origin dev
```

STEP 14 Now you can follow along after this point as shown in the following video -