

## Class Preparation

### Introduction to Data Science for Economists

Before we begin class, you need to install Anaconda and some Python packages. This guide will tell you how to do it, step by step.

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1. Folder setup
2. Installing Anaconda
3. Using command-line interface and creating a package environment
4. Installing packages/libraries
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#### 1. Folder setup

1. Create a folder 'Data\_Science' on the desktop. This folder will be used to store data, code, and resources for this class.
  - a. Mac users: /Users/[[Your username]]/Desktop/Data\_Science
  - b. Windows users: C:\Users\[[Your username]]\Desktop\Data\_Science
2. Save material for the class in this folder.
  - a. You can rename or move this folder after the class ends but we will keep a consistent location and name for now so everyone is working on the same base directory.
3. Create sub-folders for each class session to keep your data and code organized.

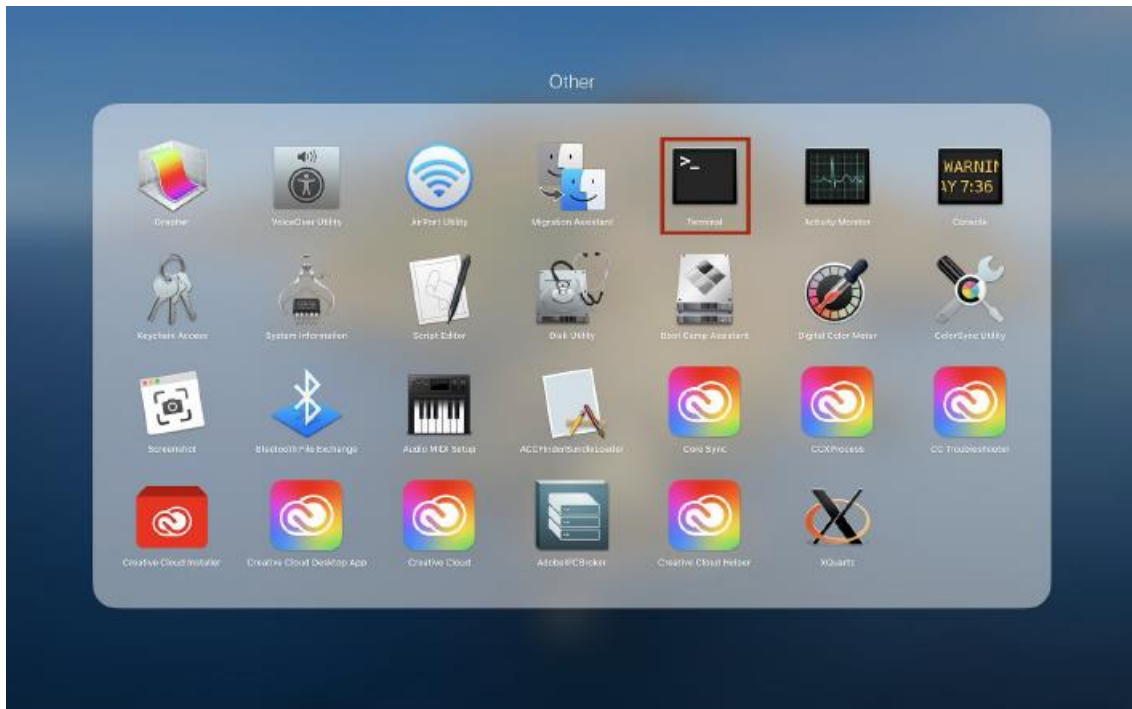
#### 2. Install Anaconda

1. What is Anaconda?
  - a. **Anaconda** is an open source data science and artificial intelligence distribution platform for Python and R programming languages. It is the most popular way to learn and use Python.
  - b. It hosts **Conda**, an open-source, cross-platform, language-agnostic package manager and environment management system. It was originally developed to solve package management challenges faced by Python data scientists, and today is a popular package manager for Python and R. It comes pre-loaded with a large number of libraries and packages.
    - i. What is a '**package**'? Package are sets of software or code that allow you to do different operations in Python or R. You have to download and load them to be able to use them in your code. '**Libraries**' are similar. You can use Conda to install, remove, and upgrade packages.
    - ii. What is an '**environment**'? An environment is a directory that contains all the files needed for a particular application, such as Python interpreter, packages, and configuration files. It is essentially the set of software and packages you are using to write code. You can use Conda to create separate environments for different projects with different package versions. Note that an environment can also refer to the set of objects you have loaded or created in your code.
  - c. What are the advantages of using Anaconda for coding?

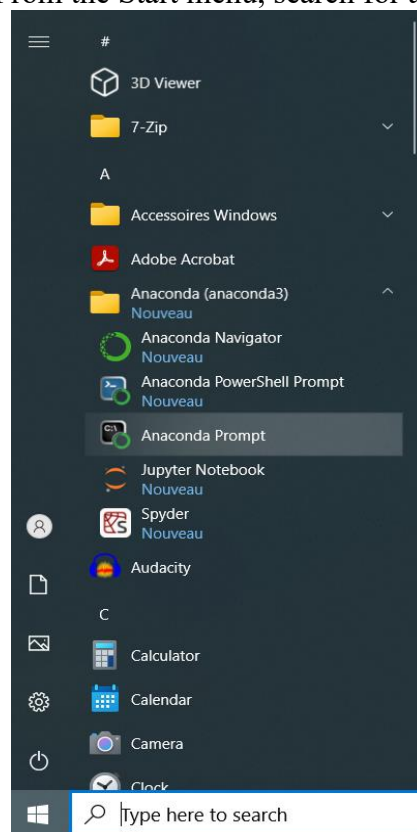
- i. It is easy to install and relatively easy to use for installing and managing packages.
  - ii. It includes many features and libraries you can use for your projects.
  - iii. You can get started quickly with coding using the included Jupyter Notebook.
  - iv. You can do many operations through the Anaconda Navigator graphical interface, avoiding the need to use the command line interface.
2. Refer to the installation guides:
  - a. Mac: <https://docs.anaconda.com/anaconda/install/mac-os/>
  - b. Windows: <https://docs.anaconda.com/anaconda/install/windows/>
3. Download installers
  - a. Go to <https://www.anaconda.com/products/individual> and click 'DOWNLOAD.'
  - b. Download an appropriate graphical installer according to the OS you are using (Mac or Windows). If you are a Windows user, you can check whether your machine is 32-bit or 64-bit following this guide: <https://support.microsoft.com/en-us/help/15056/windows-32-64-bit-faq>
  - c. After completing the download, you can follow the installation guide.
4. Note that Anaconda installs Python automatically! You do not need to install Python separately.

### 3. Create a new environment

1. Why create a new environment?
  - a. You can think of an environment as a workspace for a specific purpose (e.g. a class, a research project, etc.) The reason we recommend you to make a new environment for this class is that there can be some version conflicts among packages you might use for your future projects. Have a set environment ensures consistency across what everyone is working with.
2. We will primarily use a command-line interface for package installation, environment management, and launching python editors.
  - a. What is a **command-line interface**? A 'CLI' is a means of interacting with a computer program by inputting lines of text called command-lines. Most computer users rely on graphical user interfaces ("GUIs") instead of CLIs. However, many programs and operating system utilities lack GUIs, and are intended to be used through CLIs.
  - b. It may seem daunting but you will only use it for specific tasks with clearly indicated commands.
3. Open a command-line interface.
  - a. On Mac: Open Launchpad, then click the Terminal icon

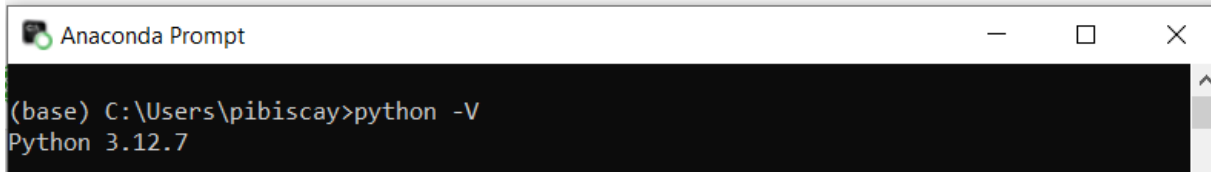


- b. On Windows: From the Start menu, search for an open ‘Anaconda Prompt.’



4. You will now see a command interface as below. By default, we are in an environment named ‘base.’ You will also see the default directory.
- On Mac, a ‘%’ symbol indicates where you can enter commands at the end of a command line.
  - On Windows, a ‘>’ symbol indicates where you can enter commands.

5. As a first test of the command line, type `python -V` and click enter to display what version of Python was installed with your Anaconda installation. It will display the response on the next line, before showing another command line.
  - a. Note that certain libraries and packages in python that we will use in this class do not work in the newest version of python. For this class, we will use python 3.9 to ensure all packages we want are installable.

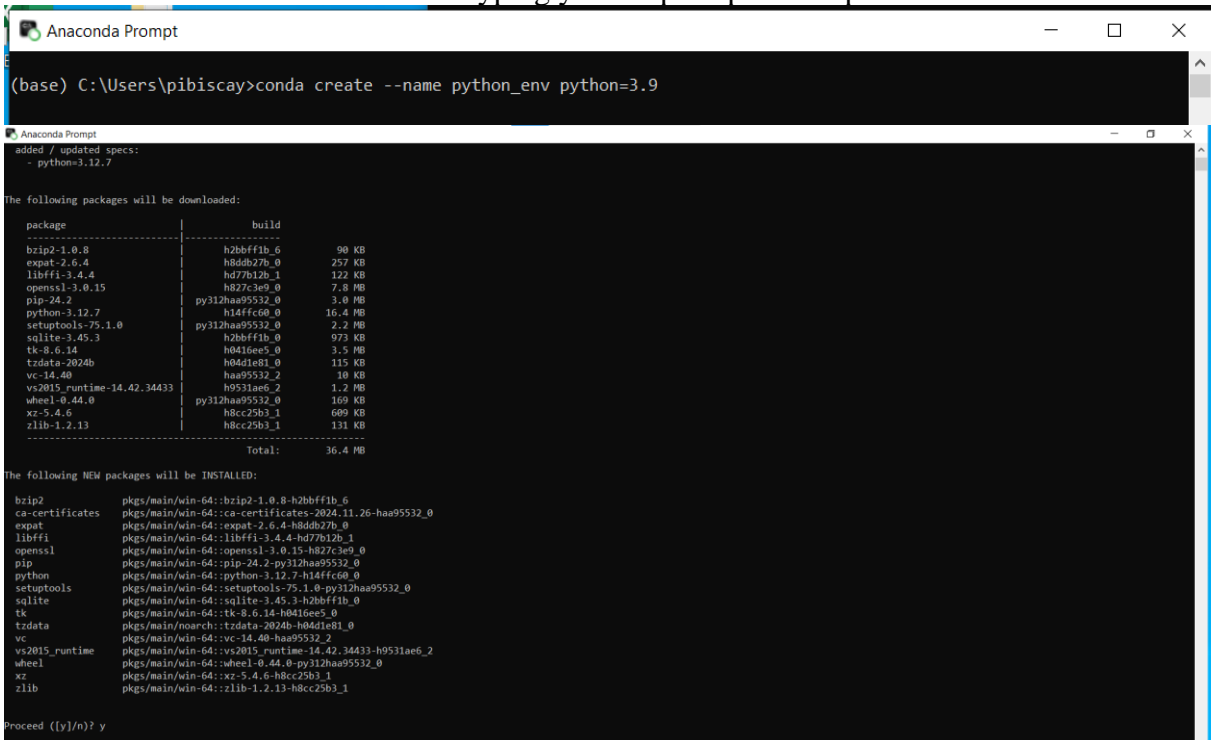


```

Anaconda Prompt
(base) C:\Users\pibiscay>python -V
Python 3.12.7

```

6. Create a new environment named 'python\_env' by entering the following commands. Type each command at the end of a command line, then press enter and wait for the interface to complete the operation.
  - a. `conda create --name python_env python=3.9`
    - i. This line creates the environment we will use for this class, and specifies the target python version.
    - ii. Proceed with typing `y` when prompted and press enter



```

Anaconda Prompt
(base) C:\Users\pibiscay>conda create --name python_env python=3.9

added / updated specs:
- python=3.12.7

The following packages will be downloaded:

```

package	build	size
bzip2-1.0.8	h2bbff1b_6	90 KB
expat-2.6.4	h8ddb27b_0	257 KB
libffi-3.4.4	hd77b12b_1	122 KB
openssl-3.0.15	h827c3e9_0	7.8 MB
pip-24.2	py312haa95532_0	3.0 MB
python-3.12.7	h14ffc60_0	16.4 MB
setuptools-75.1.0	py312haa95532_0	2.2 MB
sqlite-3.45.3	h2bbff1b_0	973 KB
tk-8.6.14	h0416ee5_0	3.5 MB
tzdata-2024b	h0416e81_0	115 KB
vc-14.40	haa95532_2	10 KB
vs2015_runtime-14.42.34433	h9531ae6_2	1.2 MB
wheel-0.44.0	py312haa95532_0	169 KB
xz-5.4.6	h8cc25b3_1	609 KB
zlib-1.2.13	h8cc25b3_1	131 KB
Total:		36.4 MB

```

The following NEW packages will be INSTALLED:

```

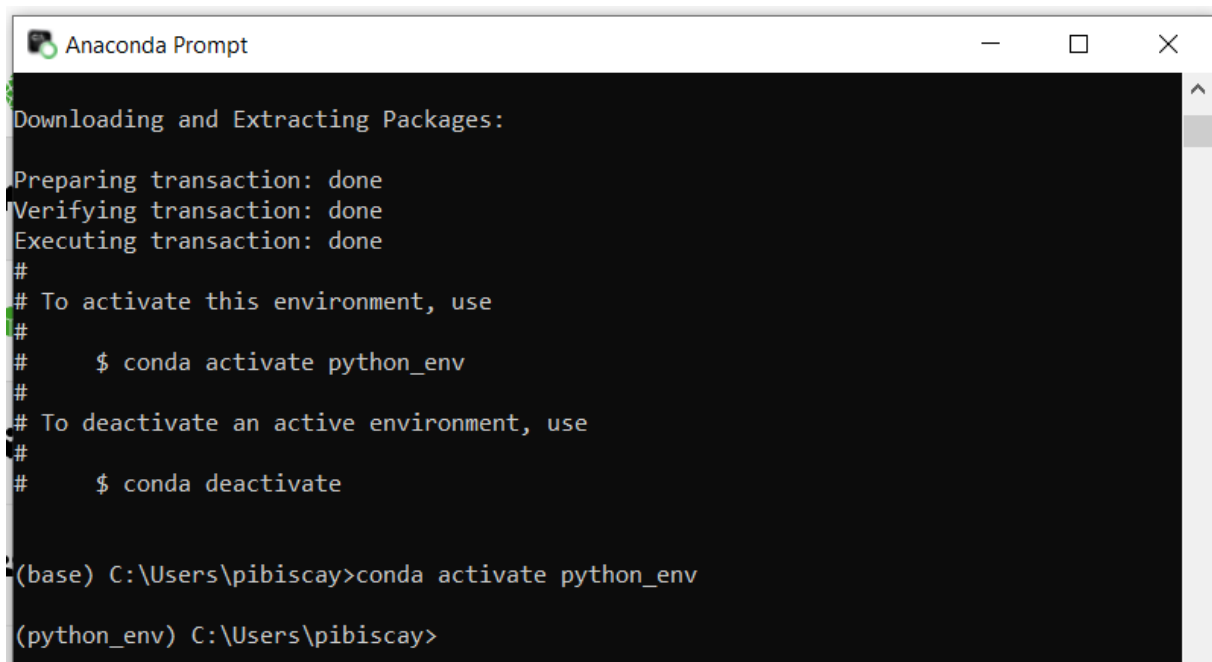
package	path
bzip2	pkgs/main/win-64::bzip2-1.0.8-h2bbff1b_6
ca-certificates	pkgs/main/win-64::ca-certificates-2024.11.26-haa95532_0
expat	pkgs/main/win-64::expat-2.6.4-h8ddb27b_0
libffi	pkgs/main/win-64::libffi-3.4.4-hd77b12b_1
openssl	pkgs/main/win-64::openssl-3.0.15-h827c3e9_0
pip	pkgs/main/win-64::pip-24.2-py312haa95532_0
python	pkgs/main/win-64::python-3.12.7-h14ffc60_0
setuptools	pkgs/main/win-64::setuptools-75.1.0-py312haa95532_0
sqlite	pkgs/main/win-64::sqlite-3.45.3-h2bbff1b_0
tk	pkgs/main/win-64::tk-8.6.14-h0416ee5_0
tzdata	pkgs/main/win-64::tzdata-2024b-h0416e81_0
vc	pkgs/main/win-64::vc-14.40-haa95532_2
vs2015_runtime	pkgs/main/win-64::vs2015_runtime-14.42.34433-h9531ae6_2
wheel	pkgs/main/win-64::wheel-0.44.0-py312haa95532_0
xz	pkgs/main/win-64::xz-5.4.6-h8cc25b3_1
zlib	pkgs/main/win-64::zlib-1.2.13-h8cc25b3_1

```

Proceed ([y]/n)? y

```

- b. `conda activate python_env`
  - i. This enters the newly created environment. You will see (python\_env) at the start of the command line instead of (base).
  - ii. Check the python version is 3.9 as desired.



```
Anaconda Prompt

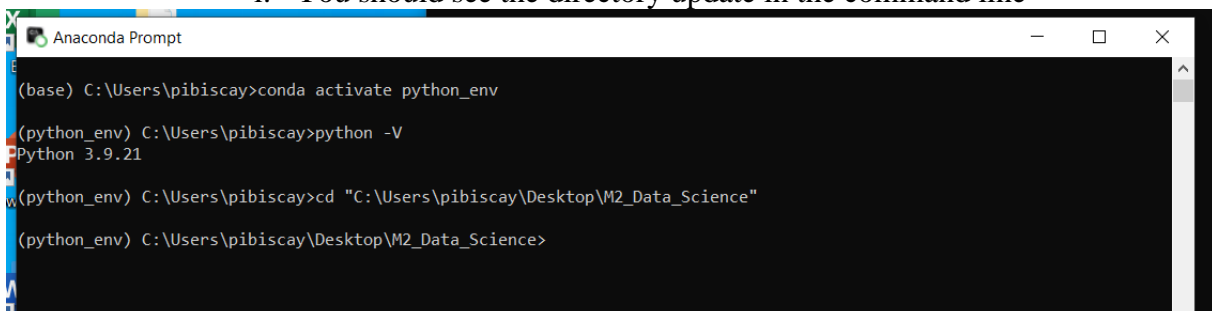
Downloading and Extracting Packages:

Preparing transaction: done
Verifying transaction: done
Executing transaction: done
#
# To activate this environment, use
#
#     $ conda activate python_env
#
# To deactivate an active environment, use
#
#     $ conda deactivate

(base) C:\Users\pibiscay>conda activate python_env

(python_env) C:\Users\pibiscay>
```

7. Set the working directory to the class folder you set up in step 1
  - a. `cd "[[your path]]"`
  - b. For example, `[[your path]] = C:\Users\[Your username]]\Desktop\M2_Data_Science`
    - i. You should see the directory update in the command line



```
Anaconda Prompt

(base) C:\Users\pibiscay>conda activate python_env

(python_env) C:\Users\pibiscay>python -V
Python 3.9.21

(python_env) C:\Users\pibiscay>cd "C:\Users\pibiscay\Desktop\M2_Data_Science"

(python_env) C:\Users\pibiscay\Desktop\M2_Data_Science>
```

#### 4. Install some libraries/packages

1. Type the following code to install selected libraries/packages we will use in this class. We will install more packages later on. As with the above, type the code in the command line and press Enter, wait for the operation to proceed, and type 'y' then Enter whenever prompted to proceed.
  - a. Note that installing packages uses Conda, the package manager that comes pre-loaded with Anaconda.
  - b. Make sure to activate your environment before installing the packages, so that they are all saved to 'python\_env' for future use.
2. `conda install jupyter`
  - a. This package is useful for working in Jupyter Notebooks, which we will use in this class.

```
Anaconda Prompt - conda install jupyter

(base) C:\Users\pibiscay>conda activate python_env

(python_env) C:\Users\pibiscay>conda install jupyter
Channels:
 - defaults
Platform: win-64
Collecting package metadata (repodata.json): done
Solving environment: done

## Package Plan ##

  environment location: C:\Users\pibiscay\AppData\Local\anaconda3\envs\python_env

  added / updated specs:
    - jupyter

The following packages will be downloaded:
```

3. Additional libraries.

- a. Make sure to run these **one line at a time!** Be careful to check for any typos.
- b. Only run the next line after the given line has completed the installation, including saying “y” when prompted.
- c. These may take some time to all install, while some others may say all requested packages have already been installed (they may be required for earlier installations, for example).
- d. If errors are returned, Google “conda install [library\_name]” and look for information for the specific working to use to call the installation in the command prompt interface.
- e. There will likely be other libraries to install during the class. Following similar language to install them to your class python environment.

*conda install gdal*

*conda install pandoc*

*conda install pygeos --channel conda-forge*

*conda install numpy pandas matplotlib shapely geopandas -c conda-forge*

*conda install -c conda-forge nodejs*

*conda install -c conda-forge jupyterlab*

*conda install -c conda-forge rasterstats*

*conda install -c plotly plotly*

*conda install --channel conda-forge esda*

*conda install geopy -c conda-forge*

*conda install contextily -c conda-forge*

*conda install mapclassify -c conda-forge*

*conda install networkx -c conda-forge*

*conda install conda-forge::r-stargazer*

*conda install seaborn*

*conda install statsmodels*

*conda install xarray*

*conda install netCDF4*

*conda install rasterio*

4. Some libraries have different dependencies. This is particularly the case for some of the libraries we will use for machine learning activities. To prevent issues across libraries, we will create a **separate environment** for machine learning libraries.
  - a. Create an environment python\_ml: *conda create --name python\_ml python=3.9*
  - b. Activate the environment
  - c. Install libraries; many other libraries are installed as part of these libraries due to internal dependencies

*conda install jupyter*

*conda install scipy*

*conda install scikit-image*

*conda install numpy pygeos shapely pandas geopandas -c conda-forge*

*conda install tpot*

*conda install seaborn*

*conda install scikit-learn*

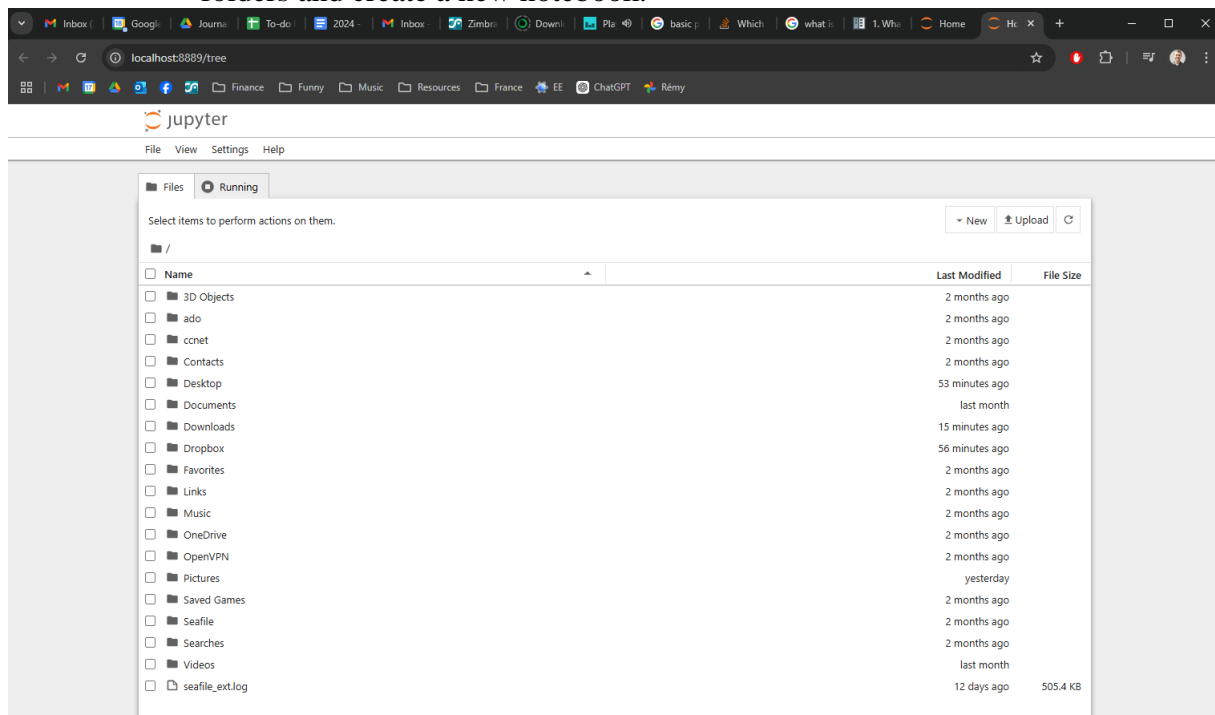
*conda install plotly::plotly*

*conda install plotly::plotly\_express*

## 5. Launch Jupyter Notebook

1. What is **Jupyter Notebook**?
  - a. Notebook documents (or “notebooks”, all lower case) are documents produced by the Jupyter Notebook App, which contain both computer code (e.g. python) and rich text elements (paragraph, equations, figures, links, etc...). Notebook documents are both human-readable documents containing the analysis description and the results (figures, tables, etc..) as well as executable documents which can be run to perform data analysis.
  - b. The **Jupyter Notebook App** is a server-client application that allows editing and running notebook documents via a web browser. The Jupyter Notebook App can be executed on a local desktop requiring no internet access (as described in this document) or can be installed on a remote server and accessed through the internet.
  - c. In addition to displaying/editing/running notebook documents, the Jupyter Notebook App has a “**Dashboard**” (Notebook Dashboard), a “control panel” showing local files and allowing to open notebook documents or shutting down their kernels.
    - i. The Notebook Dashboard is the component which is shown first when you launch Jupyter Notebook App. The Notebook Dashboard is mainly used to open notebook documents, and to manage the running kernels (visualize and shutdown). The Notebook Dashboard has other features similar to a file manager, namely navigating folders and renaming/deleting files.
  - d. A notebook **kernel** is a “computational engine” that executes the code contained in a Notebook document. The ipython kernel, which we will use in this class, executes python code. Kernels for many other languages exist.

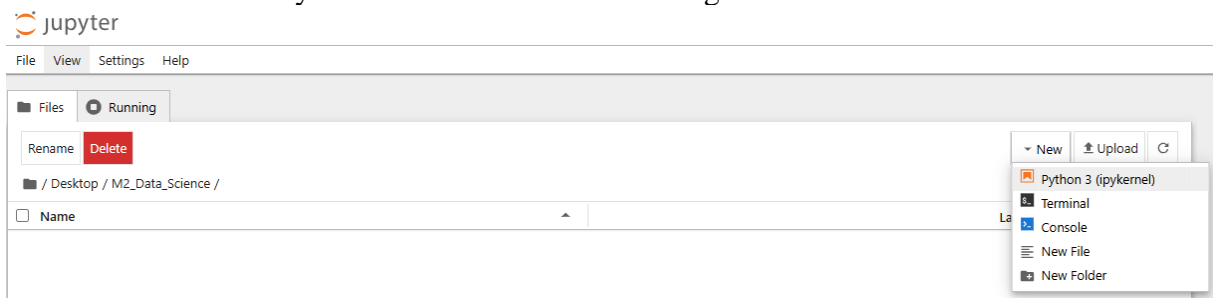
- e. When you open a Notebook document, the associated kernel is automatically launched. When the notebook is executed (either cell-by-cell or with menu Cell -> Run All), the kernel performs the computation and produces the results.
2. Why use Jupyter Notebook?
  - a. You may run Python codes using Mac terminal, Anaconda prompt, etc., however, Jupyter notebook might be one of the most convenient ways for coding in Python.
  - b. The fact that it allows for both human-readable text and code and shows analysis outputs make it very easy to navigate and read.
  - c. It is similar in many ways to R Markdown but even more easy to use (though with some differences in functionality, e.g., whether you want to hide certain chunks of code/results). It also incorporates Latex for editing math and special characters. Further, it allows you to easily export your code and results as a readable PDF.
3. How to launch Jupyter Notebook?
  - a. From the Anaconda prompt/Terminal
    - i. Open the command line interface
    - ii. `cd "[[your class folder directory]]"`
    - iii. `conda activate python_env`
    - iv. `jupyter notebook`
  - b. From the Anaconda Navigator
    - i. Go to the Environments tab and click on python\_env
    - ii. Go to Home, find Jupyter Notebook, and click Launch
  - c. Both approaches will open a new tab on your default web browser.
  - d. You will see the Notebook Dashboard, which will allow you to navigate your folders and create a new notebook.



4. Create a New Jupyter Notebook in the class folder you created in Step 1 of this document
  - a. Click on the 'Desktop' folder, then on 'M2\_Data\_Science'
  - b. On the top right, click on New, and then Python 3 (ipykernel) to create a new Jupyter Notebook using the python kernel.



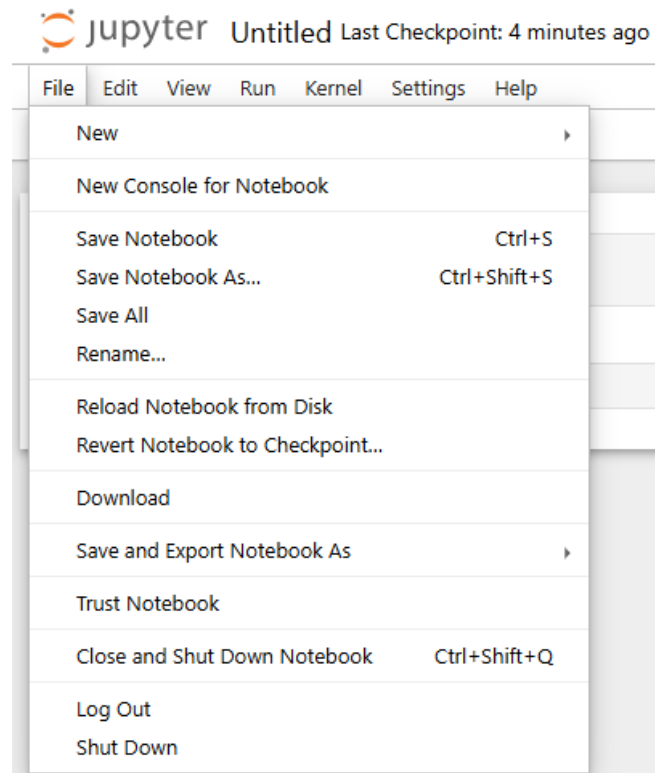
- i. Note that you can also run R in Jupyter Notebook with an R kernel, but you will have to install this using Conda first.



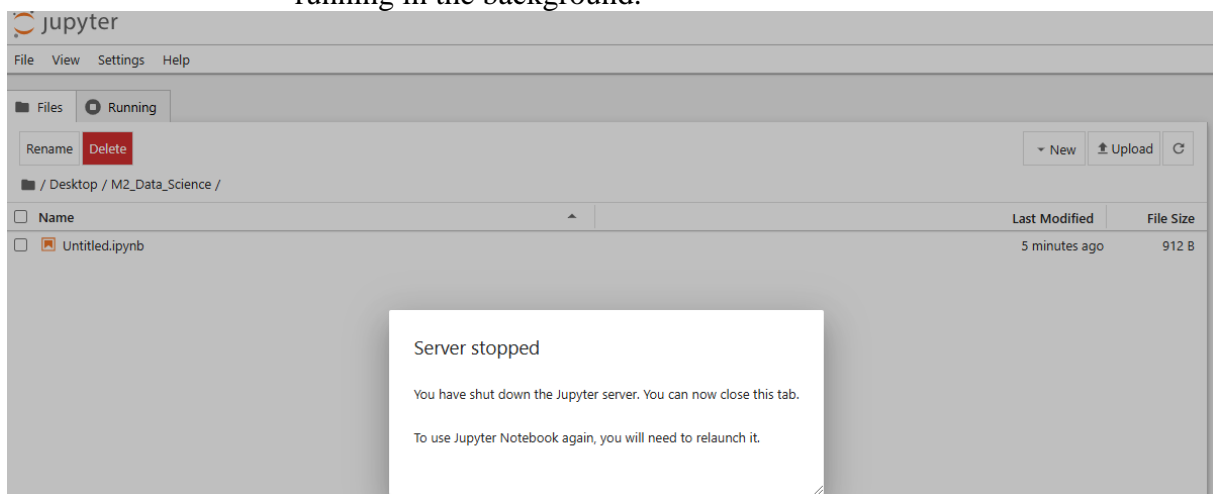
- c. A new tab will open with a blank cell.
5. Let's try our first lines of code!
    - a. Click inside the cell and type the following:
      - i. `a=1`
      - ii. `print(a)`
    - b. Navigate to the Run menu and click 'Run selected cell'



6. Close out of Jupyter Notebook.
  - a. If you wish, give the file a name and save it for future practice.
  - b. Go to the File menu and click 'Close and shut down notebook.'
    - i. Click OK when prompted.
    - ii. This will automatically close the Notebook tab.
    - iii. It is important to always do this when you are done working with a notebook to ensure your work is saved and your kernel is shut down correctly. Failure to do this can lead to problems using the kernel in the future.



- c. Go to the Jupyter Dashboard Tab. Go to the File menu and click 'Shut down'. Click "Shut down" again when prompted.
  - i. This shuts down the server hosting your Jupyter Notebook session.
  - ii. It is important to always go through this Shut down step, otherwise even if you close the Jupyter Dashboard tab the server will keep running in the background.



## Sources

<https://blog.hubspot.com/website/anaconda-python>  
[https://en.wikipedia.org/wiki/Anaconda\\_\(Python\\_distribution\)](https://en.wikipedia.org/wiki/Anaconda_(Python_distribution))  
[https://en.wikipedia.org/wiki/Command-line\\_interface](https://en.wikipedia.org/wiki/Command-line_interface)  
[https://jupyter-notebook-beginner-guide.readthedocs.io/en/latest/what\\_is\\_jupyter.html](https://jupyter-notebook-beginner-guide.readthedocs.io/en/latest/what_is_jupyter.html)