Anaconda Setup

Anaconda - Individual Edition

"The open-source Individual Edition (Distribution) is the easiest way to perform Python/R data science and machine learning on a single machine. Developed for solo practitioners, it is the toolkit that equips you to work with thousands of open-source packages and libraries."

"Individual Edition is an open source, flexible solution that provides the utilities to build, distribute, install, update, and manage software in a cross-platform manner. Conda makes it easy to manage multiple data environments that can be maintained and run separately without interference from each other."

Anaconda presentation page

Anaconda setup

You can follow the guides found at https://docs.anaconda.com/anaconda/install/ for all platforms.

We'll cover the steps from the Anaconda documentations for Windows and Linux users here. We will only be working with Python 3.

<u>Installing on Windows</u>

1. Download the Anaconda installer



Choose your Operating System (most likely 64-Bit)

2. Install

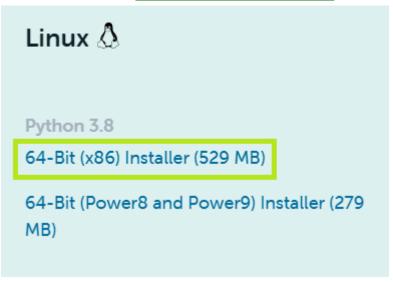
- > Double click the installer to launch
- > Click Next
- > Agree to licensing terms
- > Install for "Just Me"
- > Select destination folder and click Next
- > Recommended Do not add Anaconda3 to PATH environment variable (we will use the Anaconda Prompt each time when working with Anaconda)
- > Recommended Register Anaconda3 as default Python <python_version>
- > Click Install
- > Next
- 3. Open Anaconda Prompt



You will be using this instead of cmd (or you can use PyCharm virtual environments)

<u>Installing on Linux</u>

1. In your browser, download the Anaconda installer for Linux.



Choose the first 64-Bit Installer

- 2. Install
 - > bash ~/Downloads/Anaconda3-2020.02-Linux-x86 64.sh (replace
 - ~/Downloads/ with the path to where you downloaded/moved your .sh file)
 - > Press Enter to view license terms
 - > Agree to license terms

- > Recommended Accept default install location
- > Recommended Choose "yes" when asked if installer should run conda init
- > Close and open your terminal or just type source ~/.bashrc

Conda environments

Anaconda vs Conda

conda is both a command line tool, and a python package.

Miniconda installer = Python + conda

Anaconda installer = Python + conda + meta package anaconda

Conda environments

Anaconda environments help maintain separate projects with the help of different separate virtual environments.

Different projects/applications likely have different package dependencies. Some dependencies from one project may be incompatible with other dependencies from a different project. This is why the ideal solution is to create one environment for each project. The packages inside virtual environments can be installed either with pip or conda. (running pip install while inside a conda environment will install the package only inside that environment).

Managing environments

Conda environments documentation -

 $\underline{https://docs.conda.io/projects/conda/en/latest/user-guide/tasks/manage-environments.html}$

In order to work with conda environments you will always be using the *conda* command line tool.

Create conda environments

1. Empty environment creation

```
conda create --name myenv
press y when asked "proceed ([y]/n)?"
```

replace myenv with whatever name you want to give your environment (could be the name of the project for which you'll be using the environment)

2. Environment with specific python version

```
conda create -n myenv python=3.6
```

3. Environment with specific python version and multiple packages (not necessary, you can just create your environment and install the packages later)

```
conda create -n myenv python=3.6 scipy=0.15.0 astroid
babel
```

(-n and --name mean the same thing, specify the name of your environment)

Activate conda environments

Simply creating the environment will not be sufficient, you will need to also activate it in order to change your current environment to the newly created one.

```
conda activate myenv
```

After running the command, you should be able to see to the left of your command line that you are indeed in the desired environment. By default, the active environment—the one you are currently using—is shown in parentheses () or brackets [] at the beginning of your command prompt.

```
(myenv) $
```

View list of conda environments

conda env list

```
(base) C:\Users\Andrei>conda env list
 conda environments:
                      * D:\Users\Andrei\Anaconda3
base
                         D:\Users\Andrei\Anaconda3\envs\AIF Bayes
AIF Bayes
AndreiEnvironment
                         D:\Users\Andrei\Anaconda3\envs\AndreiEnvironment
ArrowDetection
                         D:\Users\Andrei\Anaconda3\envs\ArrowDetection
BDA
                         D:\Users\Andrei\Anaconda3\envs\BDA
                         D:\Users\Andrei\Anaconda3\envs\BDA 2
BDA 2
                         D:\Users\Andrei\Anaconda3\envs\CV1
CV1
                         D:\Users\Andrei\Anaconda3\envs\DM
DM
FML_proj
                         D:\Users\Andrei\Anaconda3\envs\FML_proj
GoogleEnv
                         D:\Users\Andrei\Anaconda3\envs\GoogleEnv
```

View environment packages list

conda list

```
(AndreiEnvironment) C:\Users\Andrei>conda list
 packages in environment at D:\Users\Andrei\Anaconda3\envs\AndreiEnvironment:
                                                      Build Channel
tflow_select
                           2.3.0
                                                        mk1
absl-py
                           0.8.1
                                                     py37 0
astor
                           0.8.0
attrs
                           19.3.0
backcall
blas
bleach
ca-certificates
                           2019.11.27
certifi
                           2019.11.28
colorama
                           0.4.3
decorator
defusedxml
entrypoints
gast
google-pasta
                                             py37h351948d
grpcio
                           1.16.1
                                             py37h5e291fa
15ру
hdf5
                                                 h7ebc959 0
                           1.10.4
icc_rt
                           2019.0.0
                                                 h0cc432a
importlib_metadata
                           1.3.0
intel-openmp
                                             py37h39e3cac_0
ipykernel
python
                                             py37h39e3cac 0
                           7.11.1
 python genutils
```

Installing packages inside environments

There are 2 ways to install packages in conda environments:

1st way is by using the conda install command

```
conda install package name
```

The 2nd (and perhaps more frequent) method is by using the pip package manager pip install package name

You may have to install pip using conda first if it is not already preinstalled conda install pip

You should be all set with Anaconda and conda environments now. In short, in order to get a conda environment up and running you will have to run:

- 1. conda create --name my environment name
- 2. conda activate my_environment_name
- 3. pip install <my package name> for all the packages that you want to install

Jupyter Notebook

"Jupyter is a free, open-source, interactive web tool known as a computational notebook, which researchers can use to combine software code, computational output, explanatory text and multimedia resources in a single document."

from https://www.nature.com/articles/d41586-018-07196-1

Install Jupyter Notebook

```
conda install -c conda-forge notebook - with conda

pip install notebook - or with pip

or

Install JupyterLab

conda install -c conda-forge jupyterlab

pip install jupyterlab
```

Start Jupyter Notebook

jupyter notebook

or

Start JupyterLab

jupyter-lab

Add a conda environment to Jupyter Notebook or JupyterLab

```
conda activate cenv
conda install ipykernel
ipython kernel install --user --name=<any_name_for_kernel>
```

All set, now you can start/restart your J Notebook/Lab and should be able to see your environment



In JupyterLab when creating a new Notebook you can choose what environment (Kernel) to use.



The same thing can be done in Jupyter Notebook (choose between default Python 3 - base, or DM environment).

Experiment creation of conda environments:

1. Create a conda environment called Env1_3.7 that uses python 3.7 and install the jupyter notebook package inside of it.

Run the following code in a newly created notebook inside jupyter notebook

```
import sys
print("Python version")
print (sys.version)
```

2. Create a conda environment called Env1_3.8 that uses python 3.8.8 and install the jupyter notebook, numpy and pandas packages inside of it.

Use this environment and **proceed with the DM1.ipynb file** for this laboratory.