

Setting up the environment

# Conda

- Some python applications may need different versions of packages than the ones you are currently using
- The correct way to handle such situations is by using **environments**
- We will use a popular environment manager called **conda**
- Conda is also a package manager (allows you to install additional packages)
- Conda also downloads and installs python for you

# Installing conda

- Conda installer: miniconda / anaconda
  - Miniconda: includes conda (package and environment manager) and few basic packages
  - Anaconda: includes conda, many scientific packages and a GUI
- Platform: multiple OS - 64 / 32 bit
- Python version: install conda for python 3.8 or above

# Anaconda

- Anaconda is the most popular Python data science platform
- Anaconda is a distribution of the Python and R programming languages for data science and machine learning related applications
- It also installs the Jupyter Notebook
- Includes a collection of over 1,000 open source data science packages
- Package versions are managed by the package management system conda

# Installing Anaconda

- Go to <https://www.anaconda.com/products/individual>
- Download the Python 3.x version

## Anaconda Installers

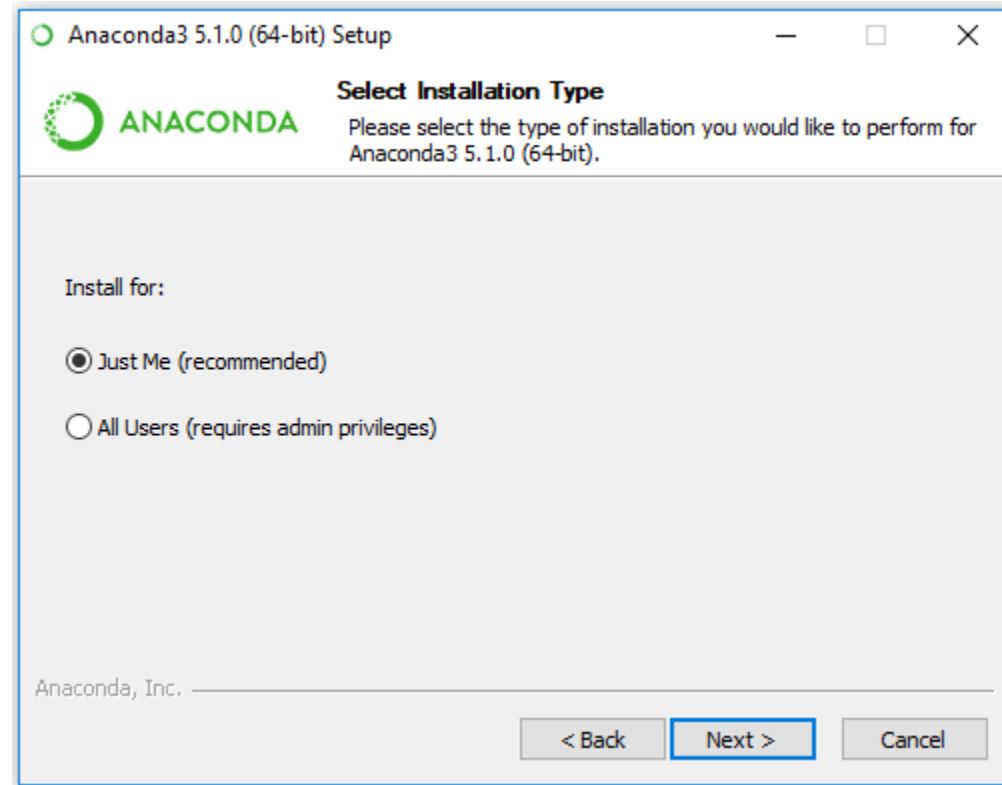
| Windows  | MacOS    | Linux       |
|---|---|--|
| Python 3.8<br>64-Bit Graphical Installer (457 MB)<br>32-Bit Graphical Installer (403 MB)  | Python 3.8<br>64-Bit Graphical Installer (435 MB)<br>64-Bit Command Line Installer (428 MB) | Python 3.8<br>64-Bit (x86) Installer (529 MB)<br>64-Bit (Power8 and Power9) Installer (279 MB) |

# Installing Anaconda

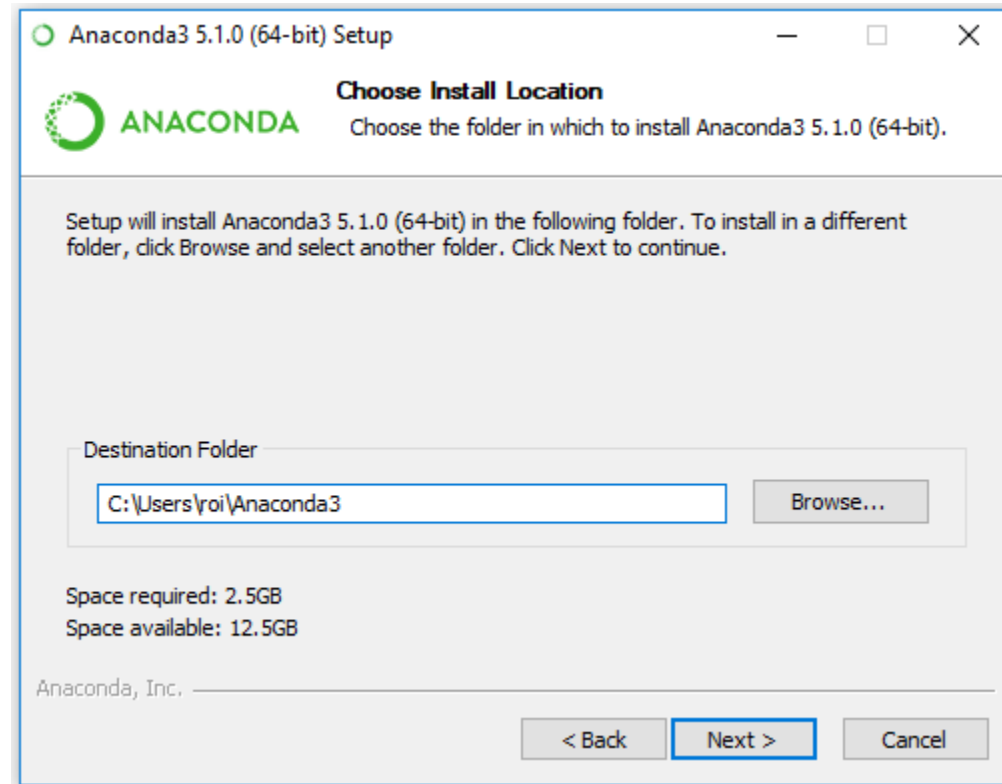
- Double click the executable file to start the installation



# Installing Anaconda



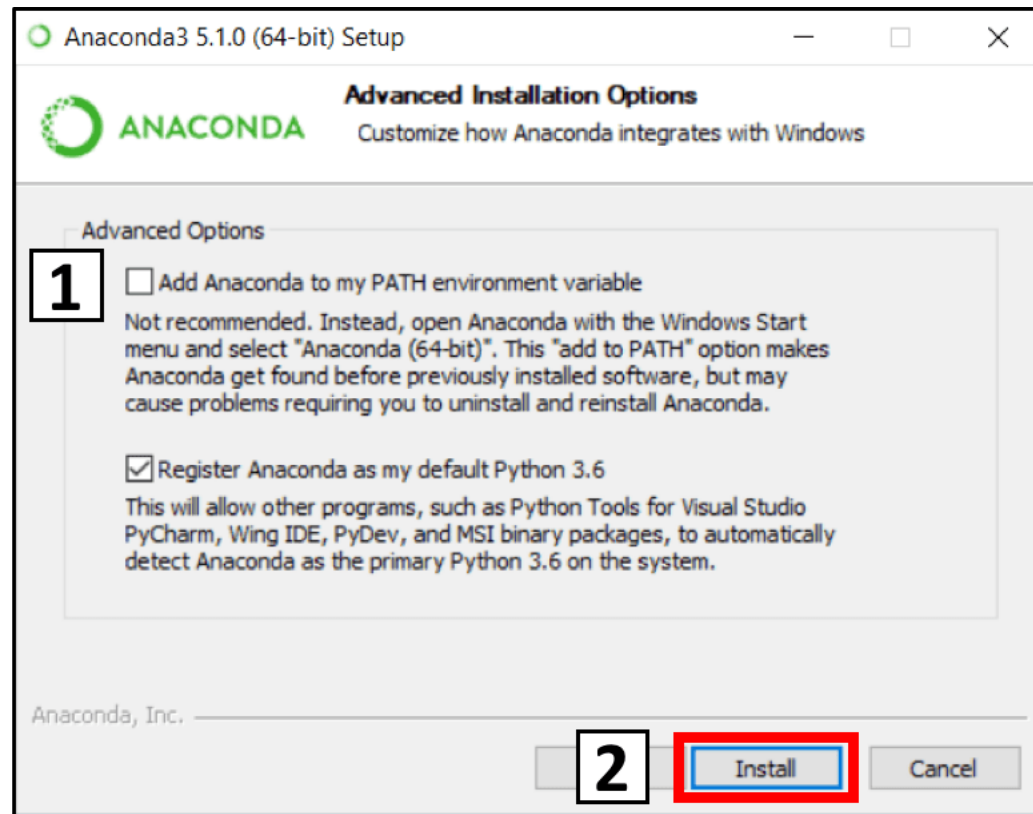
# Installing Anaconda



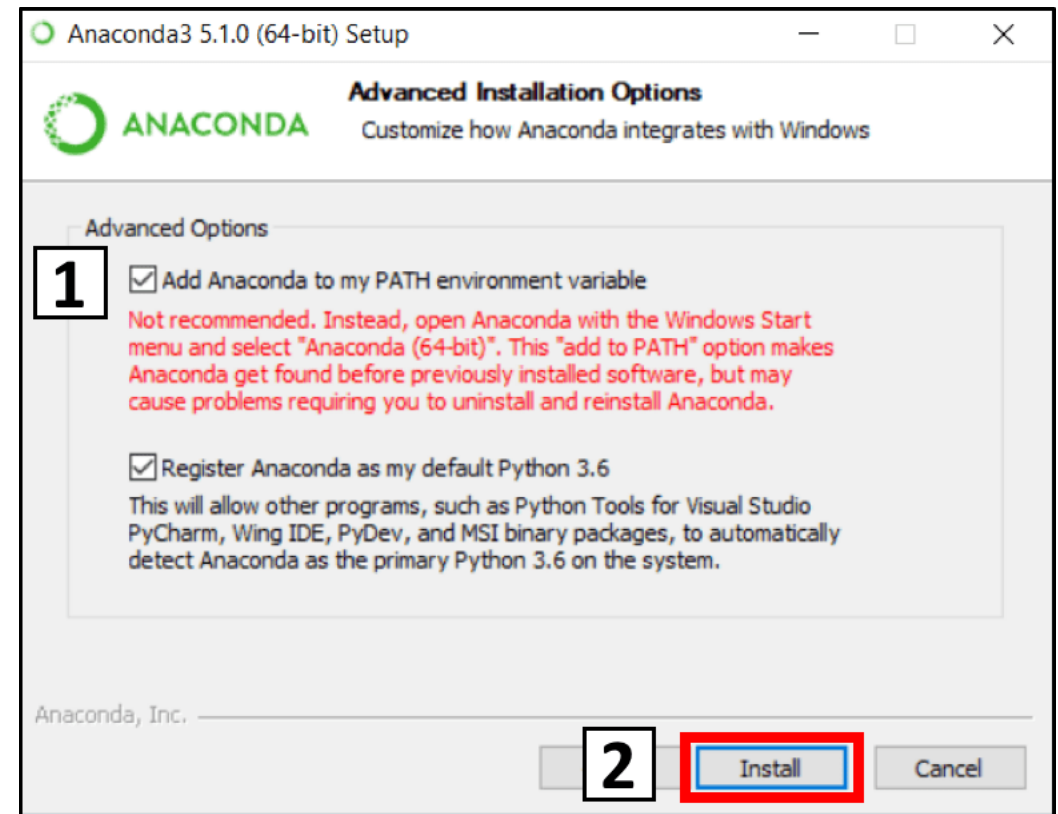


# Installing Anaconda

## Recommended Approach



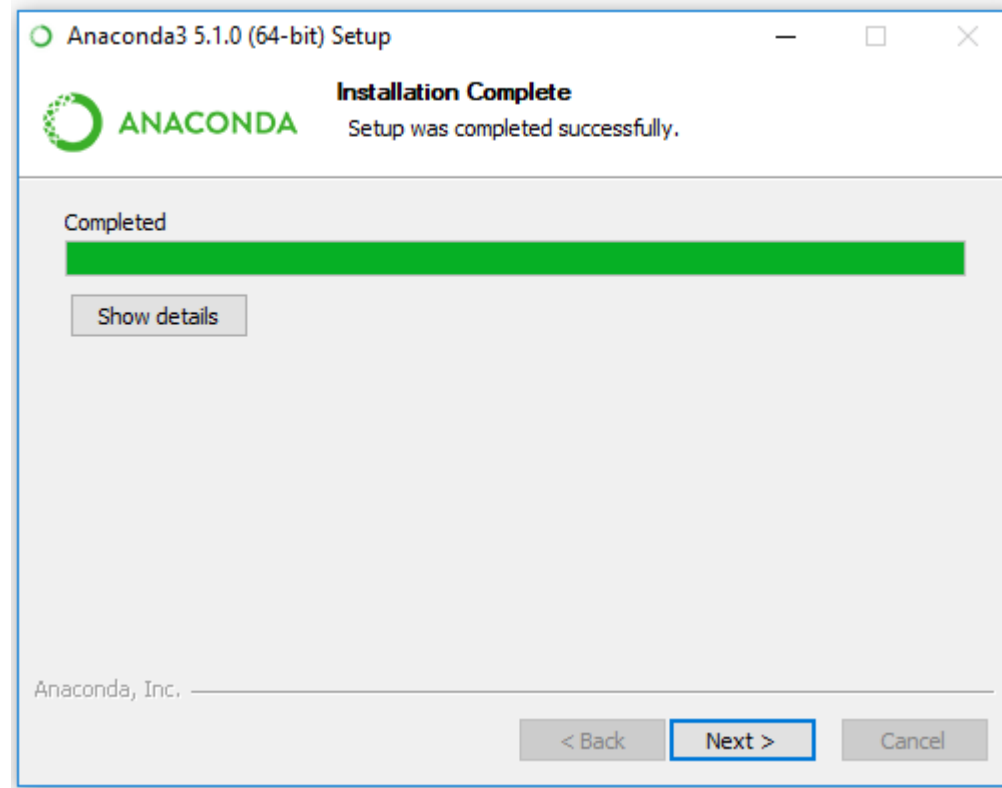
## Alternative Approach



# Installing Anaconda

- The recommended approach is to not check the box to add Anaconda to your path. This means you will have to use Anaconda Navigator or the Anaconda Command Prompt (located in the Start Menu under "Anaconda") when you wish to use Anaconda (you can always add Anaconda to your PATH later if you don't check the box)
- If you want to use Anaconda in your command prompt (or git bash, terminal, etc), please use the alternative approach and check the box

# Installing Anaconda



# Conda

- Miniconda / Anaconda sets up two things for you: **Conda** and the **root environment**
- You can check conda works by typing conda in your shell (command prompt / terminal)

```
(base) → ~ conda
usage: conda [-h] [-V] command ...
```

conda is a tool for managing and deploying applications, environments and packages.

Options:

positional arguments:

|           |  |
|-----------|--|
| command   |  |
| clean     | Remove unused packages and caches.   |
| config    | Modify configuration values in .condarc. This is modeled after the git config command. Writes to the user .condarc file (/Users/alon/.condarc) by default. |
| create    | Create a new conda environment from a list of specified packages.  |
| help      | Displays a list of available conda commands and their help strings.  |
| info      | Display information about current conda install.   |
| init      | Initialize conda for shell interaction. [Experimental]   |
| install   | Installs a list of packages into a specified conda environment.  |
| list      | List linked packages in a conda environment.   |
| package   | Low-level conda package utility. (EXPERIMENTAL)  |
| remove    | Remove a list of packages from a specified conda environment.  |
| uninstall | Alias for conda remove.  |
| run       | Run an executable in a conda environment. [Experimental]   |
| search    | Search for packages and display associated information. The input is a MatchSpec, a query language for conda packages. See examples below.                 |
| update    | Updates conda packages to the latest compatible version.   |
| upgrade   | Alias for conda update.  |

# Creating an environment

- You can install packages inside one environment without affecting other environments
- Environments are especially useful if you are working on several projects with different dependencies

```
(base) → ~ conda create --name ml
Collecting package metadata (current_repodata.json): done
Solving environment: done

## Package Plan ##

  environment location: /Users/alon/anaconda3/envs/ml

Proceed ([y]/n)? y

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

(base) → ~ conda activate ml
(ml) → ~
```

# Environment management

- You don't have to use environments (but its recommended)
- If you are missing a package, just google "conda package\_name"
- All packages we will use in this course are available through conda
- For now, make sure the following packages are installed in the new environment you just created:
  - jupyter
  - numpy
  - pandas
  - matplotlib

# IDE (integrated development environment)

- **Sublime**
  - **Visual Studio Code**
  - PyCharm
  - Atom
  - Spyder
- 
- You are free to use any development environment / code editor you want
  - All debugging and development can be performed from Jupyter Notebook

# Jupyter notebook

- “The Jupyter Notebook is an open-source web application that allows you to create and share documents that contain live code, equations, visualizations and narrative text. Uses include: data cleaning and transformation, numerical simulation, statistical modeling, data visualization, machine learning, and much more”
- `conda install jupyter`



# Jupyter notebook

- After installing jupyter, you can activate it by running
- `$ jupyter notebook`
- This will create a local server on your machine which you can access from your browser, usually from `http://localhost:8888`

# Jupyter notebook

- Make sure jupyter is installed properly by running 'jupyter notebook'

```
(ml) → ~ jupyter notebook
[I 13:10:47.754 NotebookApp] [nb_conda_kernels] enabled, 2 kernels found
[I 13:10:48.207 NotebookApp] [jupyter_nbextensions_configurator] enabled 0.4.1
[I 13:10:48.331 NotebookApp] [nb_conda] enabled
[I 13:10:48.331 NotebookApp] Serving notebooks from local directory: /Users/alon
[I 13:10:48.331 NotebookApp] The Jupyter Notebook is running at:
[I 13:10:48.331 NotebookApp] http://localhost:8888/?token=10286d9947059e84baa6bc930efea5865ff3273ad2b77550
[I 13:10:48.331 NotebookApp] or http://127.0.0.1:8888/?token=10286d9947059e84baa6bc930efea5865ff3273ad2b77550
[I 13:10:48.331 NotebookApp] Use Control-C to stop this server and shut down all kernels (twice to skip confirmation).
[C 13:10:48.339 NotebookApp]
```

To access the notebook, open this file in a browser:

file:///Users/alon/Library/Jupyter/runtime/nbserver-59173-open.html

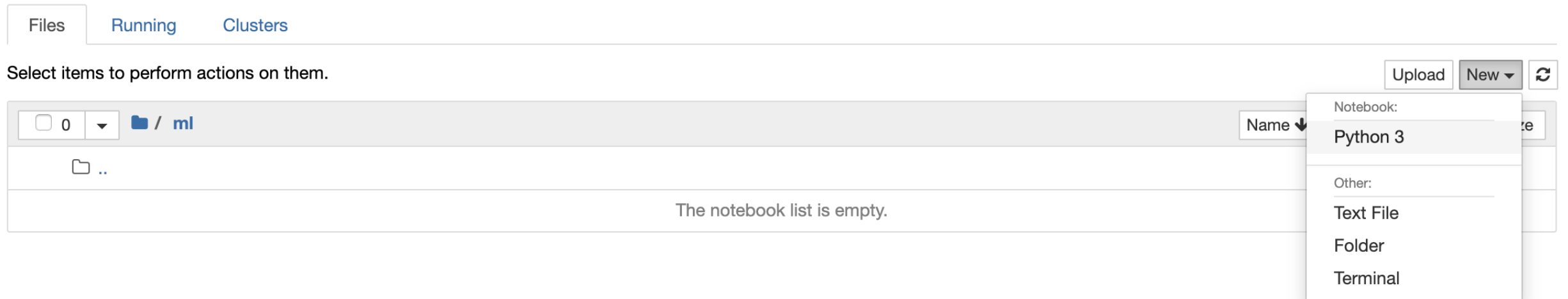
Or copy and paste one of these URLs:

http://localhost:8888/?token=10286d9947059e84baa6bc930efea5865ff3273ad2b77550

or http://127.0.0.1:8888/?token=10286d9947059e84baa6bc930efea5865ff3273ad2b77550

# Jupyter notebook

- The notebook should open automatically. Otherwise, copy and paste the link from the console to your browser (Google Chrome works best)
- Create a new notebook



# Jupyter notebook

- Type the following into the first cell and press 'shift+enter' to run:
  - import numpy as np
  - import pandas as pd
  - import matplotlib.pyplot as plt
- If the cell run without errors, you should see the following

```
In [1]: import numpy as np  
import pandas as pd  
import matplotlib.pyplot as plt
```

```
In [ ]:
```