

Quick Start for Anaconda and Jupyter

We will be using Anaconda and Jupyter Notebook for this semester. This tutorial will walk you through the installation process.

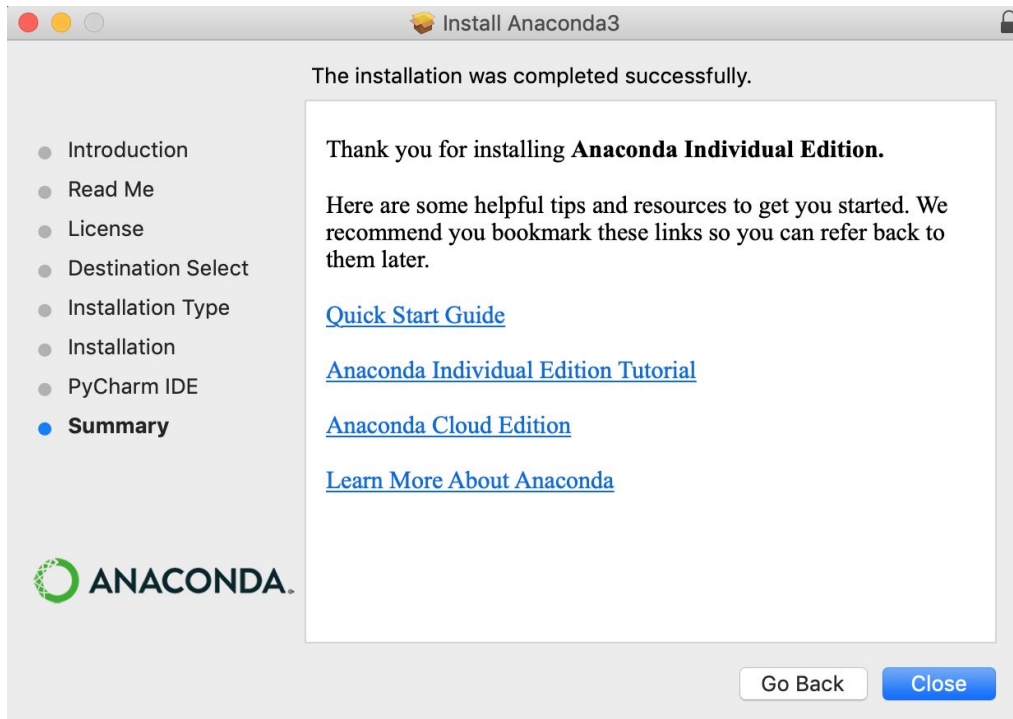
Introduction

Anaconda is an open source package management system and environment management system that runs on Windows, macOS and Linux. It helps us create and switch between various Python environments and install packages within a specific Python environment on your computer. With the help of Anaconda, we can easily handle compatibility and dependency issues of various Python packages.

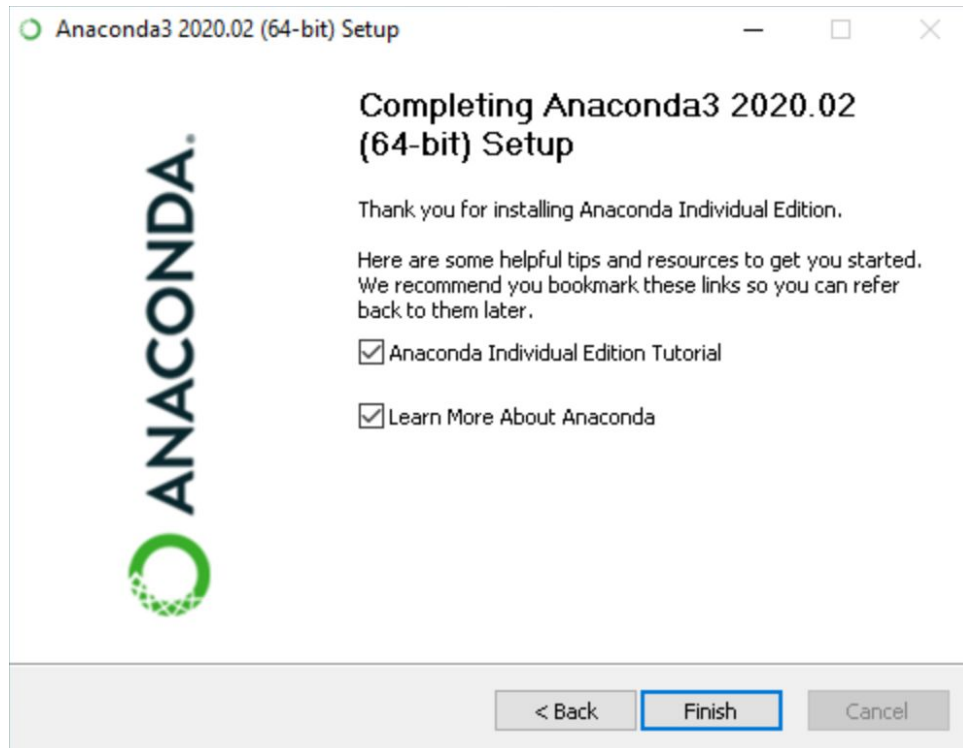
Generally speaking, one can install packages in Anaconda in two ways - Anaconda Navigator (desktop graphical user interface) and Conda (command line interface tool).

Installation

- Installing on macOS
 1. Download the graphical macOS installer for your version of Python (Python 3 recommended) at <https://www.anaconda.com/products/individual#macos>
 2. Double-click the installer and click continue to start the installation.
 3. A successful installation message:



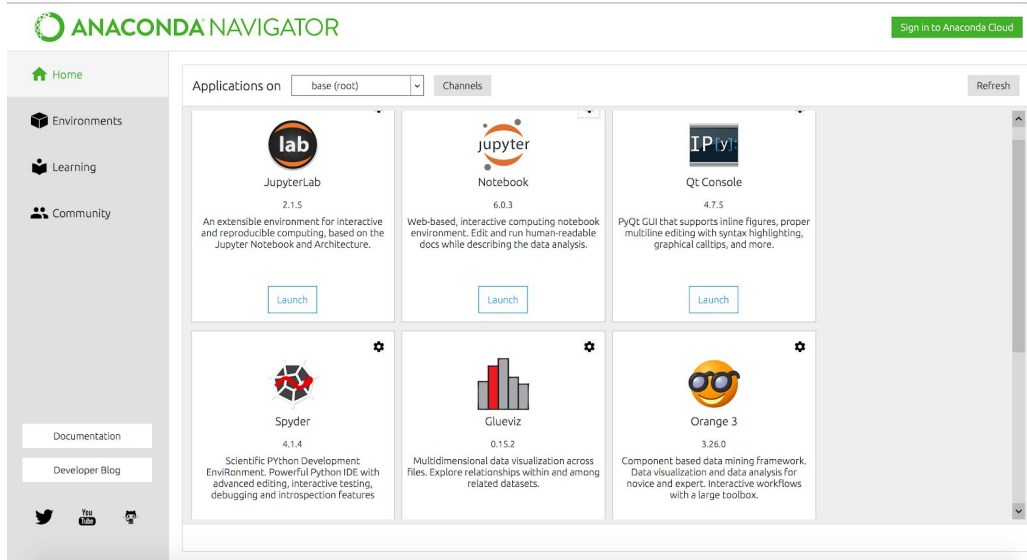
- Installing on Windows
 1. Download the graphical macOS installer for your version of Python (Python 3 recommended) at <https://www.anaconda.com/products/individual#windows>
 2. Double-click the installer and click continue to start the installation.
 3. A successful installation message:



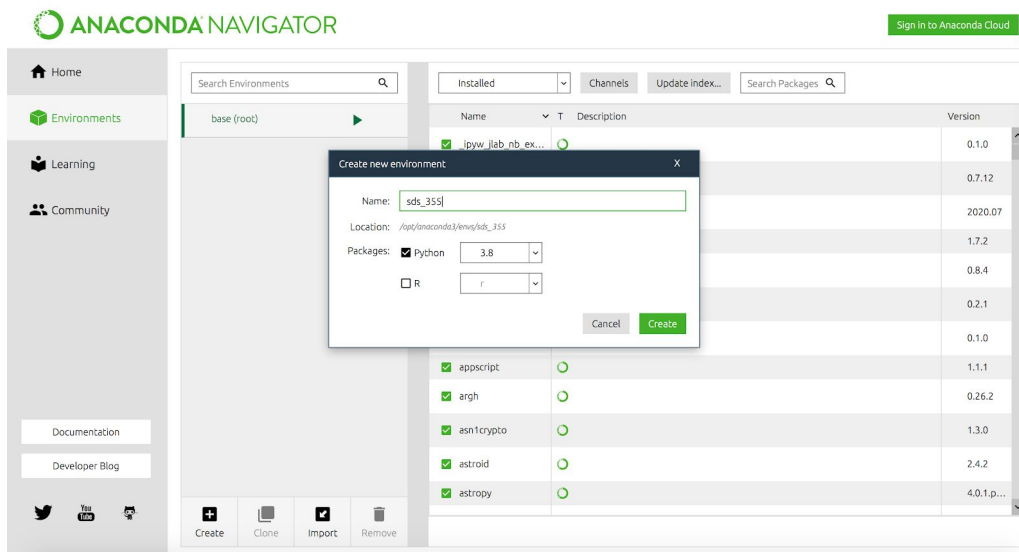
User Guide for Anaconda

With Navigator, we can create separate Python environments in which Python packages and their dependencies will not interact with other environments. After installing all the packages needed, we will launch the Jupyter Notebook to write and run Python scripts.

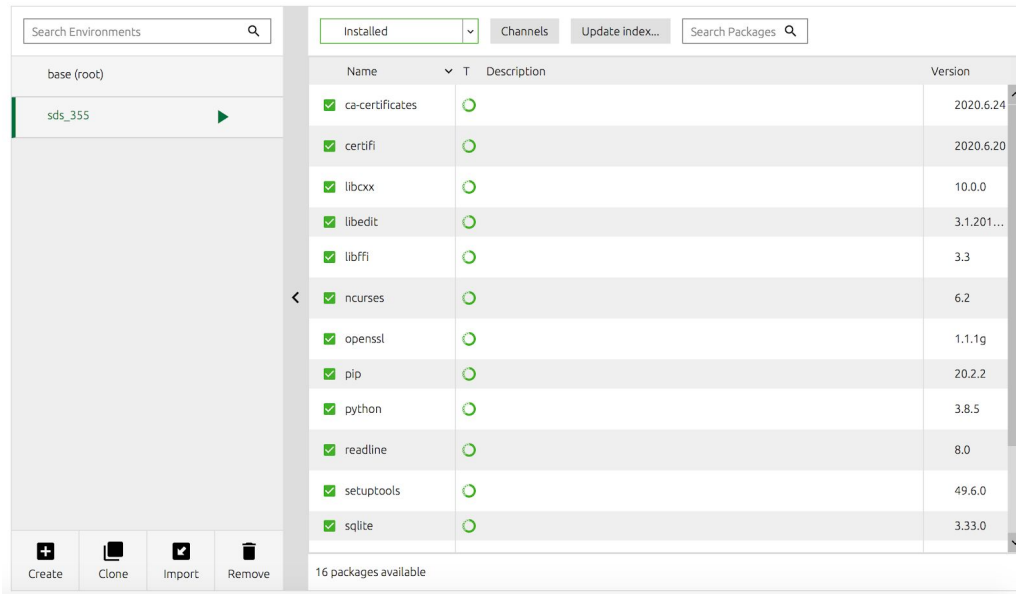
1. **Open the application:** open the Anaconda Navigator application either through Mac Launchpad or Windows Start menu.



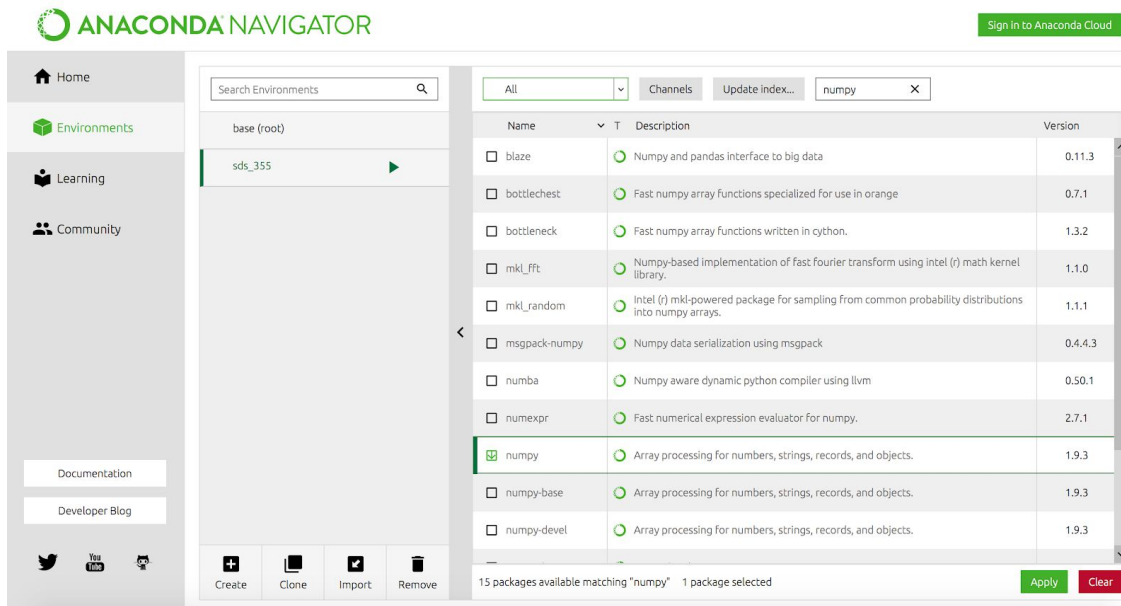
2. **Create new environments:** Create a new environment named `sds_355` by clicking the Environments tab and clicking the Create button.

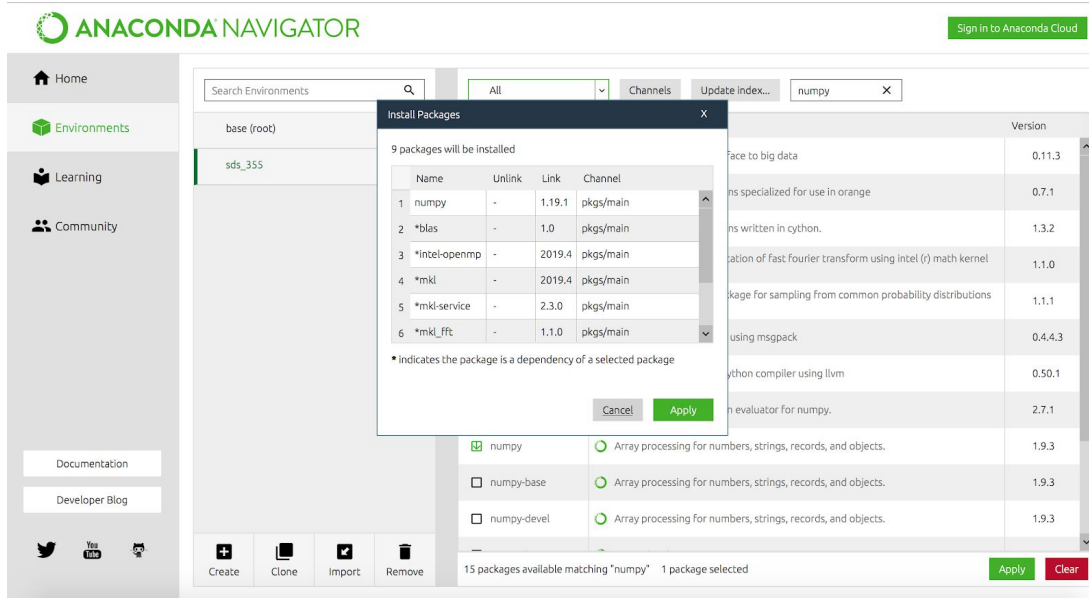


3. **Activate the environment:** Now you should be able to see two Python environments - `base(root)` and `sds_355`. Activate the `sds_355` environment by clicking the `sds_355` tab. The green triangle represents the activated environment.

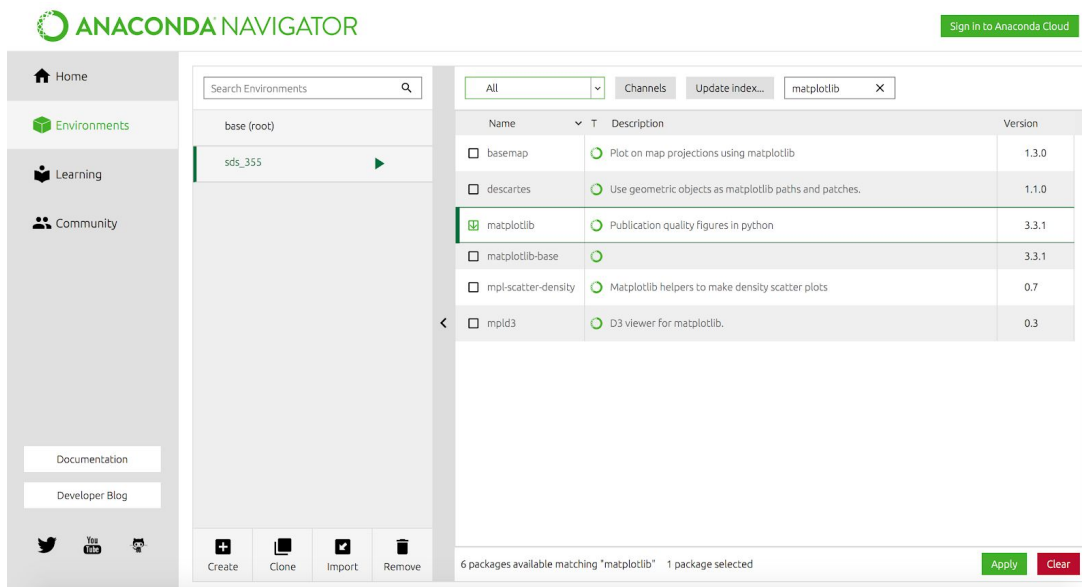


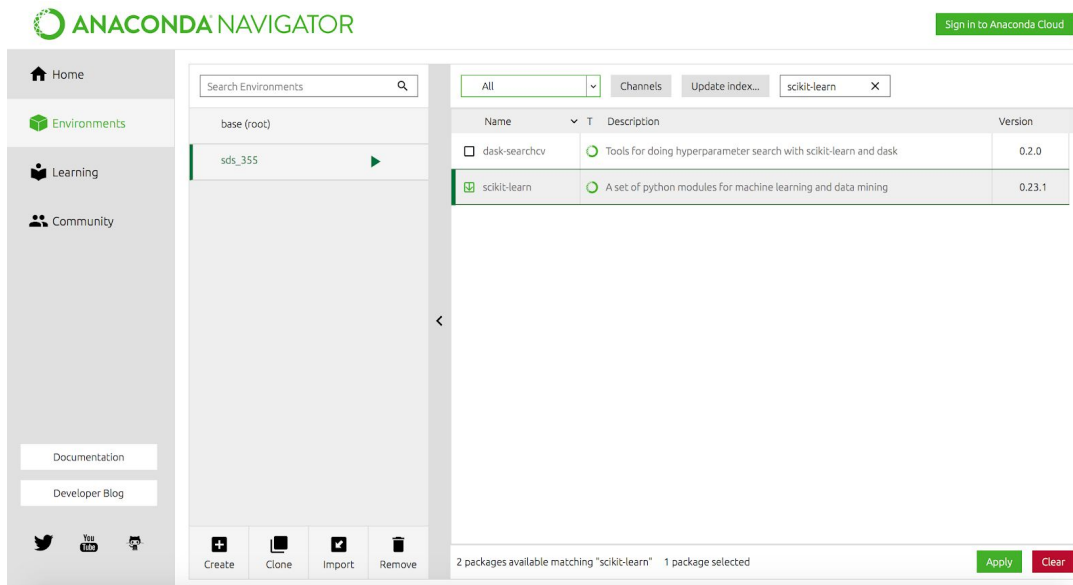
4. **Install packages:** In order to add packages to the current environment, we could click the Update Index first and put the name of the package in the search box. Among all the packages matching the search criteria, we could choose one and click apply. The packages and the dependencies could be automatically installed.





5. Following the same step, we could also install other packages needed in this class including matplotlib, scikit-learn, etc.

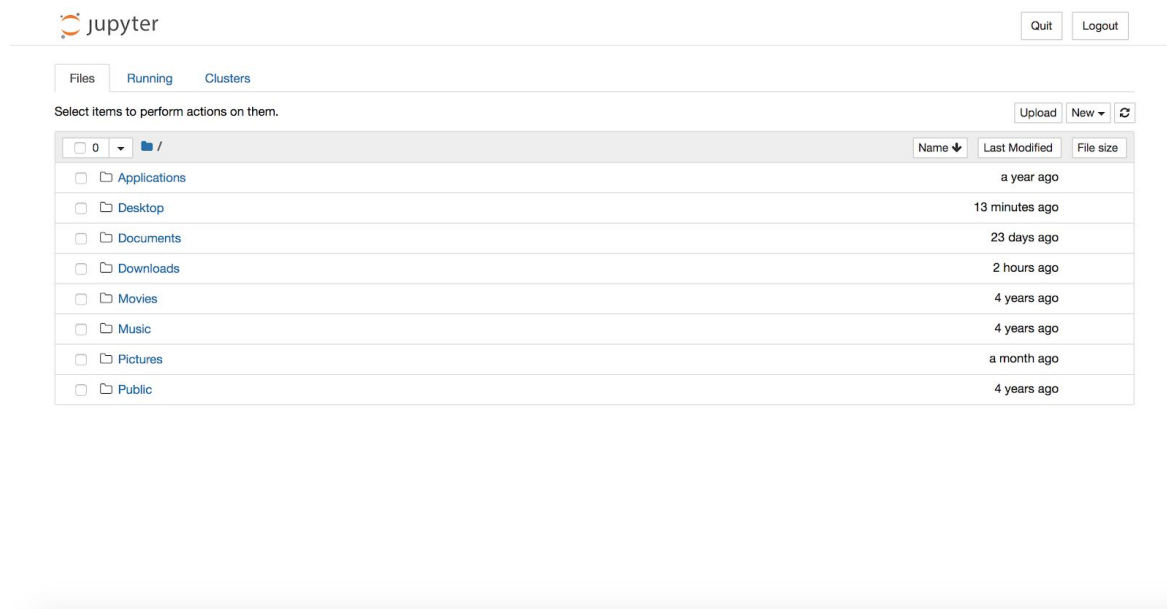




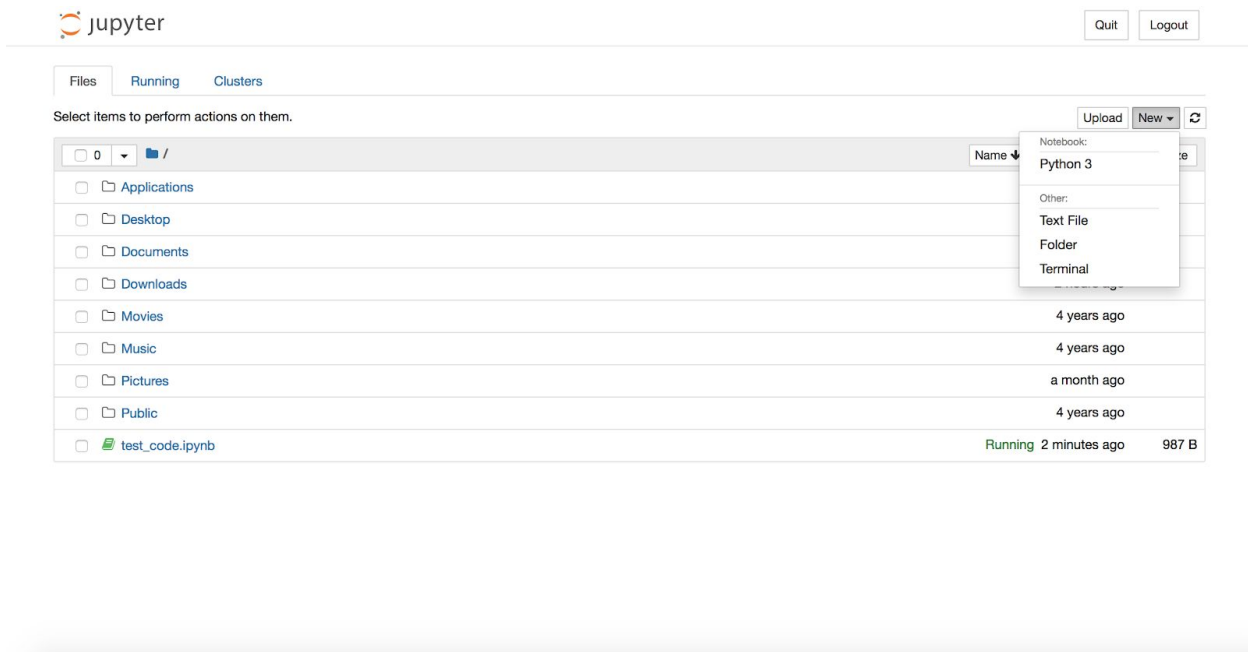
Start Coding with Jupyter

After building the Python environment, we are about to launch the Jupyter Notebook to write some Python code. The Jupyter Notebook is a web application where one can write code, equations and narrative text and share documents with others.

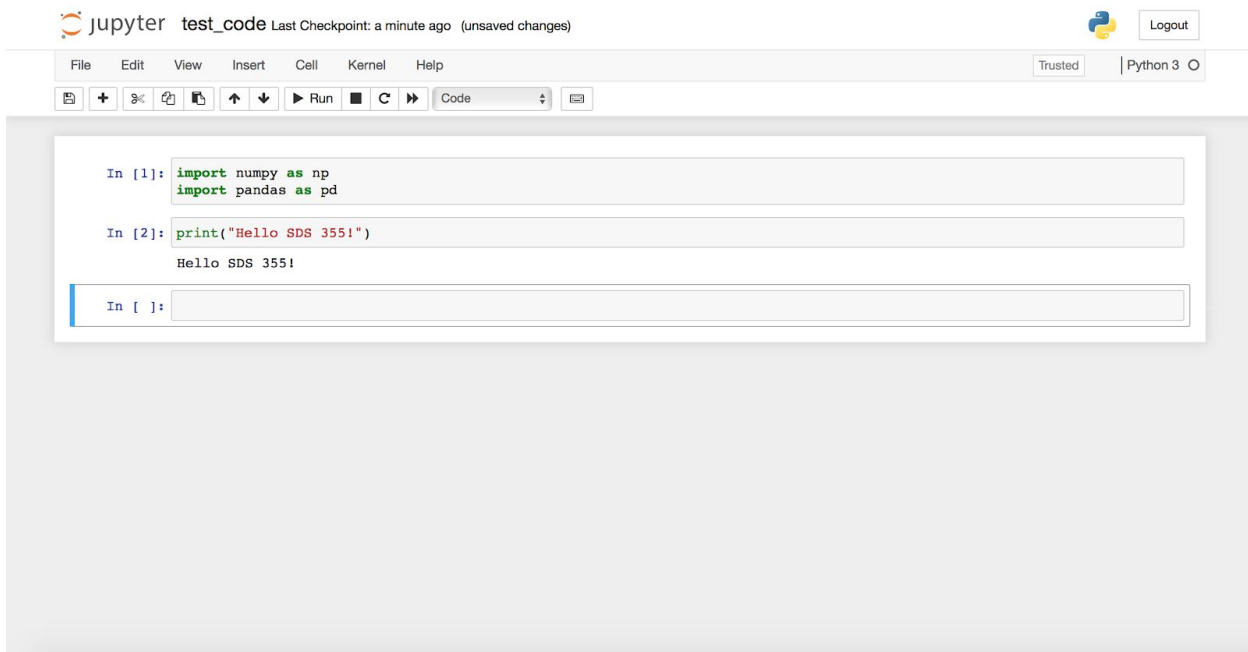
The Jupyter Notebook has been pre-installed for the base environment. For new environments, you can easily install it and launch it in the home view. You should be able to see the following webpage:



We can click New tab and create a new Python 3 notebook:



Start coding and have fun:



For people who are interested in **Conda CLI** tool:

This [cheat sheet](#) will be extremely helpful:

Conda basics

| | |
|---|--|
| Verify conda is installed, check version number | <code>conda info</code> |
| Update conda to the current version | <code>conda update conda</code> |
| Install a package included in Anaconda | <code>conda install PACKAGENAME</code> |
| Run a package after install, example Spyder* | <code>spyder</code> |
| Update any installed program | <code>conda update PACKAGENAME</code> |
| Command line help | <code>COMMANDNAME --help</code> <code>conda install --help</code> |

*Must be installed and have a deployable command, usually PACKAGENAME

Using environments

| | |
|---|--|
| Create a new environment named py35, install Python 3.5 | <code>conda create --name py35 python=3.5</code> |
| Activate the new environment to use it | WINDOWS: <code>activate py35</code> LINUX, macOS: <code>source activate py35</code> |
| Get a list of all my environments, active environment is shown with * | <code>conda env list</code> |
| Make exact copy of an environment | <code>conda create --clone py35 --name py35-2</code> |
| List all packages and versions installed in active environment | <code>conda list</code> |
| List the history of each change to the current environment | <code>conda list --revisions</code> |
| Restore environment to a previous revision | <code>conda install --revision 2</code> |
| Save environment to a text file | <code>conda list --explicit > bio-env.txt</code> |
| Delete an environment and everything in it | <code>conda env remove --name bio-env</code> |
| Deactivate the current environment | WINDOWS: <code>deactivate</code> macOS, LINUX: <code>source deactivate</code> |
| Create environment from a text file | <code>conda env create --file bio-env.txt</code> |
| Stack commands: create a new environment, name it bio-env and install the biopython package | <code>conda create --name bio-env biopython</code> |

| Installing and updating packages | | |
|---|-------------------------------------|--|
| Install a new package (Jupyter Notebook) in the active environment | | <code>conda install jupyter</code> |
| Run an installed package (Jupyter Notebook) | | <code>jupyter-notebook</code> |
| Install a new package (toolz) in a different environment (bio-env) | | <code>conda install --name bio-env toolz</code> |
| Update a package in the current environment | | <code>conda update scikit-learn</code> |
| Install a package (boltons) from a specific channel (conda-forge) | | <code>conda install --channel conda-forge boltons</code> |
| Install a package directly from PyPI into the current active environment using pip | | <code>pip install boltons</code> |
| Remove one or more packages (toolz, boltons) from a specific environment (bio-env) | | <code>conda remove --name bio-env toolz boltons</code> |
| Managing multiple versions of Python | | |
| Install different version of Python in a new environment named py34 | | <code>conda create --name py34 python=3.4</code> |
| Switch to the new environment that has a different version of Python | | Windows: <code>activate py34</code> Linux, macOS: <code>source activate py34</code> |
| Show the locations of all versions of Python that are currently in the path NOTE: The first version of Python in the list will be executed. | | Windows: <code>where python</code> Linux, macOS: <code>which -a python</code> |
| Show version information for the current active Python | | <code>python --version</code> |
| Specifying version numbers | | |
| Ways to specify a package version number for use with conda create or conda install commands, and in meta.yaml files. | | |
| Constraint type | Specification | Result |
| Fuzzy | <code>numpy=1.11</code> | 1.11.0, 1.11.1, 1.11.2, 1.11.18 etc. |
| Exact | <code>numpy==1.11</code> | 1.11.0 |
| Greater than or equal to | <code>"numpy>=1.11"</code> | 1.11.0 or higher |
| OR | <code>"numpy=1.11.1 1.11.3"</code> | 1.11.1, 1.11.3 |
| AND | <code>"numpy>=1.8, <2"</code> | 1.8, 1.9, not 2.0 |

References

<https://docs.anaconda.com/anaconda/>