

Setup Jupyter via Longleaf On Demand

Super Important: Make sure you complete the [First-time setup](#) section at the end of this document or you will not be able to start the coursework for CHEM 467.

Once your account is approved you will be able to access Longleaf. There are many ways to access Longleaf, but we will be using Longleaf On-Demand for this course.

Access Jupyter via Longleaf On Demand

1. Go to <https://ondemand.rc.unc.edu/pun/sys/dashboard> (You may want to bookmark this site for quick access throughout the course).
2. Sign in using your onyen.
3. Click on **Jupyter Notebook**.
4. Request resources and click **Launch**.
 - **Number of hours:** 8 hr. maximum
 - **Number of CPUs:** 1 is sufficient

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Interactive Apps

Desktops

Longleaf Desktop

Longleaf Desktop (GPU-FULL)

Longleaf Desktop (GPU)

MD Desktop

GUIs

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MATLAB

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Pymol

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SAS

Spyder

Stata

Servers

Jupyter Notebook

Jupyter Notebook (GPU)

Jupyter Notebook (GPU-FULL)

RStudio Server

Jupyter Notebook

This app will launch a Jupyter Notebook server on Longleaf.

Number of hours

8

Number of CPUs

1

Unless you are certain you are running parallel code you should specify **one** cpu.

☐ Request GPU

Check if you would like to request a GPU for this session.

Additional Job Submission Arguments

☐ legacy jupyter notebook

Check if you would like to use the legacy jupyter notebook user interface, otherwise you get the jupyterlab interface.

☐ I would like to receive an email when the session starts

Jupyter startup directory

This field allows you to specify the absolute path to the Longleaf directory where you would like the Jupyter session to start.

Launch

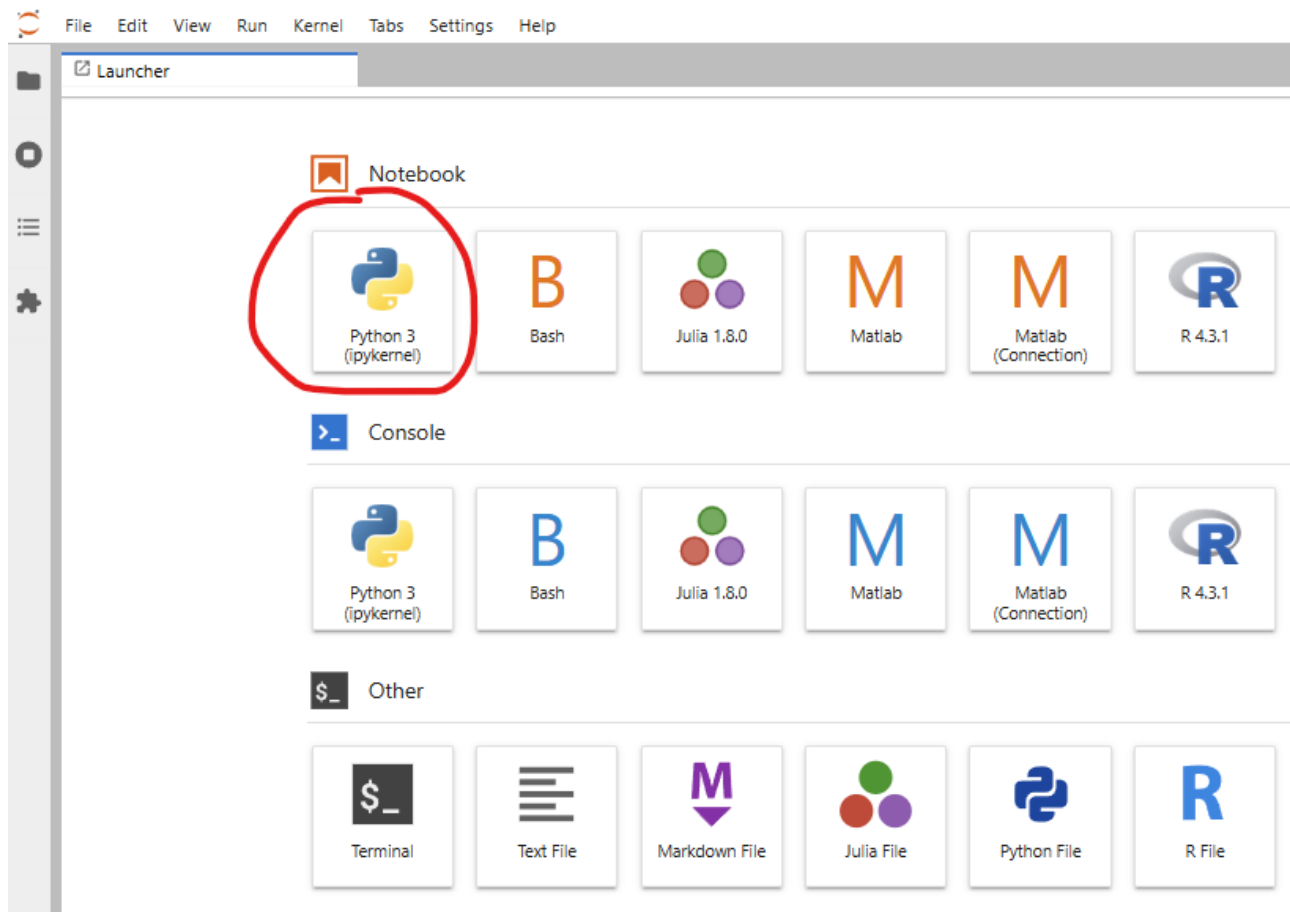
* The Jupyter Notebook session data for this session can be accessed under the [data root directory](#).

5. Wait for computing resources to be allocated to you (~1 minute).

6. Click **Connect to Jupyter**.

7. In the **Launcher** and under the section labeled **Notebook**, click **Python3 (ipkernel)** to open a notebook running Python.

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Setting up your Jupyter notebook

Jupyter notebooks are made up of two main types of cells: **Markdown cells** and **Code cells**. You run a Jupyter notebook one cell at a time by click inside the cell and press **shift+enter**.

Markdown cells

Jupyter notebooks allow us to use a language called Markdown to type notes and descriptions in markdown cells. It's a good practice to have your first cell be markdown to explain the purpose of the notebook.

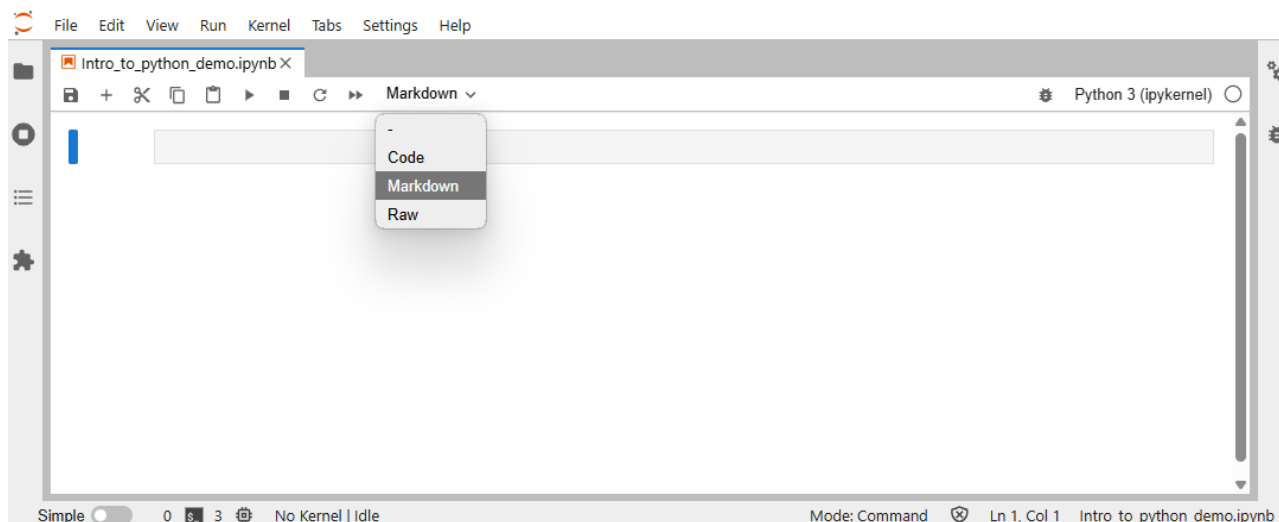
Basic markdown syntax

In the Markdown language, we create headers and lists using the following syntax.

Element	Basic Markdown Syntax
Heading	# H1
	## H2
	### H3
Ordered List	1. First item
	2. Second item
	3. Third item
Unordered List	- First item
	- Second item
	- Third item

Let's practice!

1. Open a new Jupyter notebook using the [instructions above](#).
2. Click the first cell.
3. Select **Markdown** from the cell type dropdown menu in the notebook ribbon (shown below).



4. Paste the following into the markdown cell.

```
# Python for chemists
```

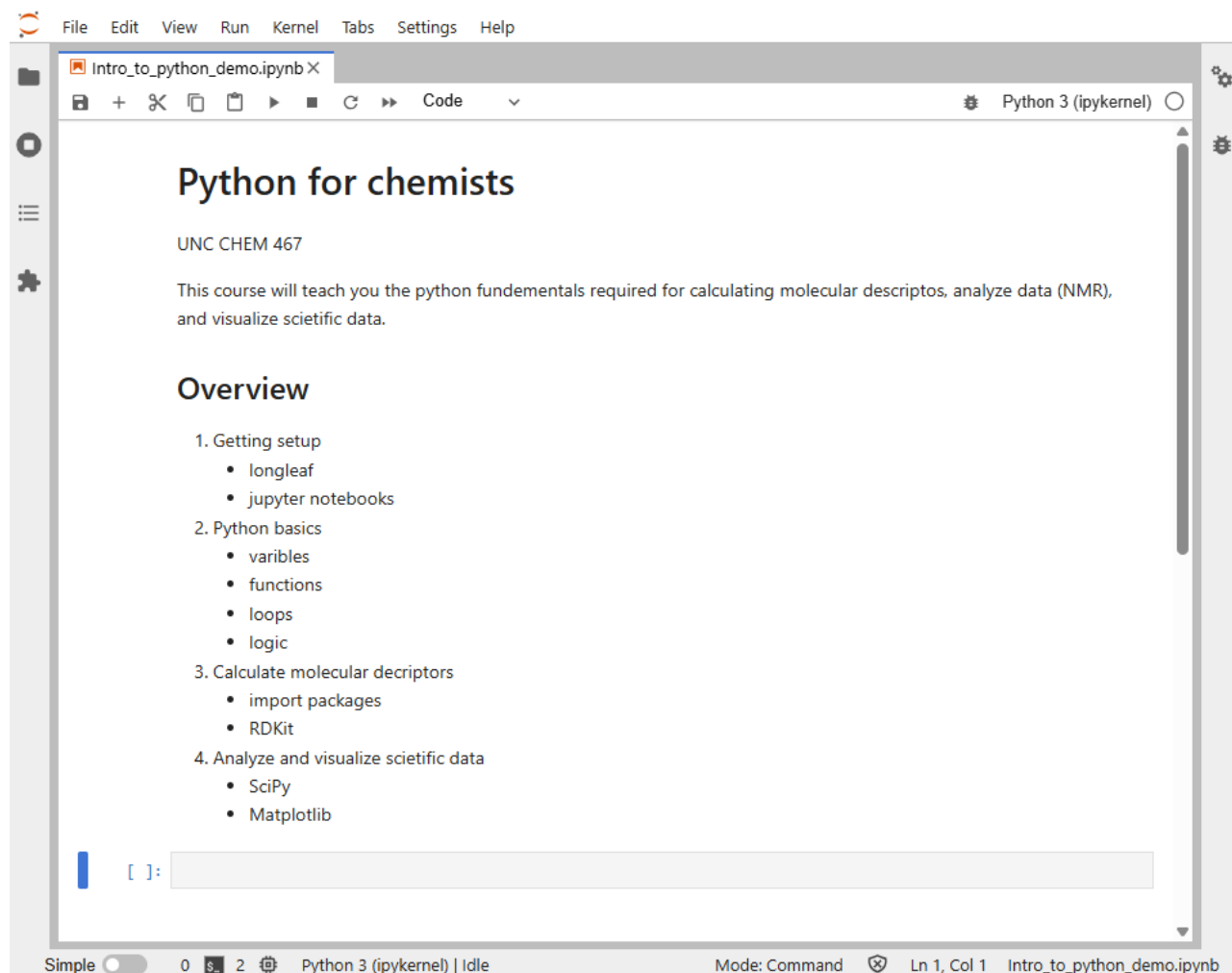
```
UNC CHEM 467
```

```
This course will teach you the python fundamentals required for  
calculating molecular descriptors, analyze data (NMR), and visualize  
scientific data.
```

```
## Overview
```

1. Getting setup
 - longleaf
 - jupyter notebooks
2. Python basics
 - variables
 - functions
 - loops
 - logic
4. Calculate molecular descriptors
 - import packages
 - RDKit
5. Analyze and visualize scientific data
 - SciPy
 - Matplotlib

- Press **shift+enter** (or click the play button ► in the notebook ribbon) to run the markdown cell. After running, Markdown cell should look like the following:



Code cells

Jupyter notebooks allow us to run python code interactively and display the outputs all in one file. These features make Jupyter notebooks good for prototyping new code and for keeping a record of how you analyzed your data.

Let's practice!

- Paste the following interactive python code into the empty code cell below your Markdown cell.

```
# Example of interactive python code in Jupyter

# Import packages
import numpy as np
import matplotlib.pyplot as plt
from ipywidgets import interact

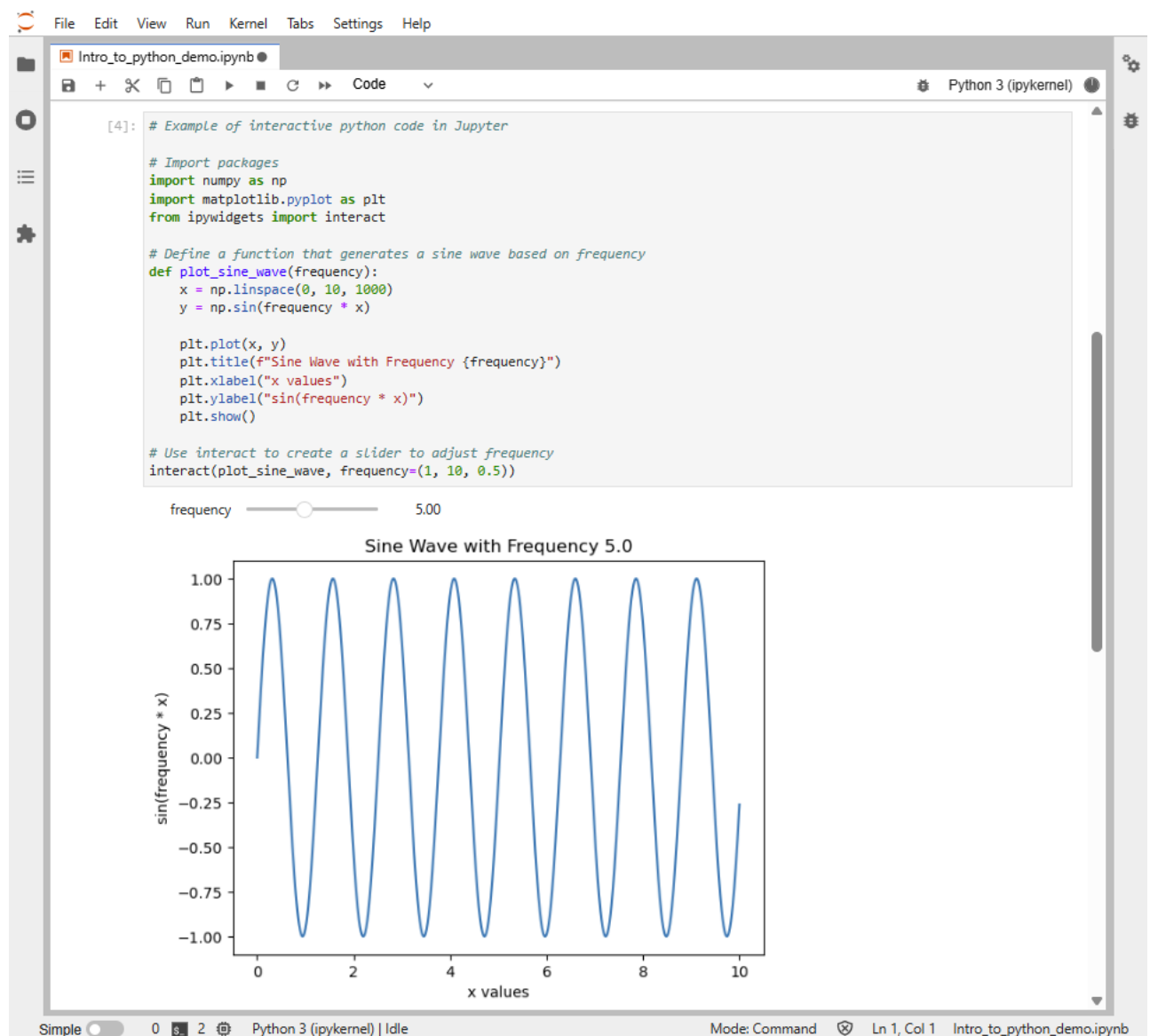
# Define a function that generates a sine wave based on frequency
def plot_sine_wave(frequency):
    x = np.linspace(0, 10, 1000)
    y = np.sin(frequency * x)
```

```
plt.plot(x, y)
plt.title(f"Sine Wave with Frequency {frequency}")
plt.xlabel("x values")
plt.ylabel("sin(frequency * x)")
plt.show()

# Use interact to create a slider to adjust frequency
interact(plot_sine_wave, frequency=(1, 10, 0.5))
```

Note: in Python (and therefore in Jupyter code cells) the # symbol is used to denote comments that will not be interpreted as code.

2. Press **shift+enter** (or click the play button ► in the notebook ribbon) to evaluate the code cell. Now your notebook should look like the following:



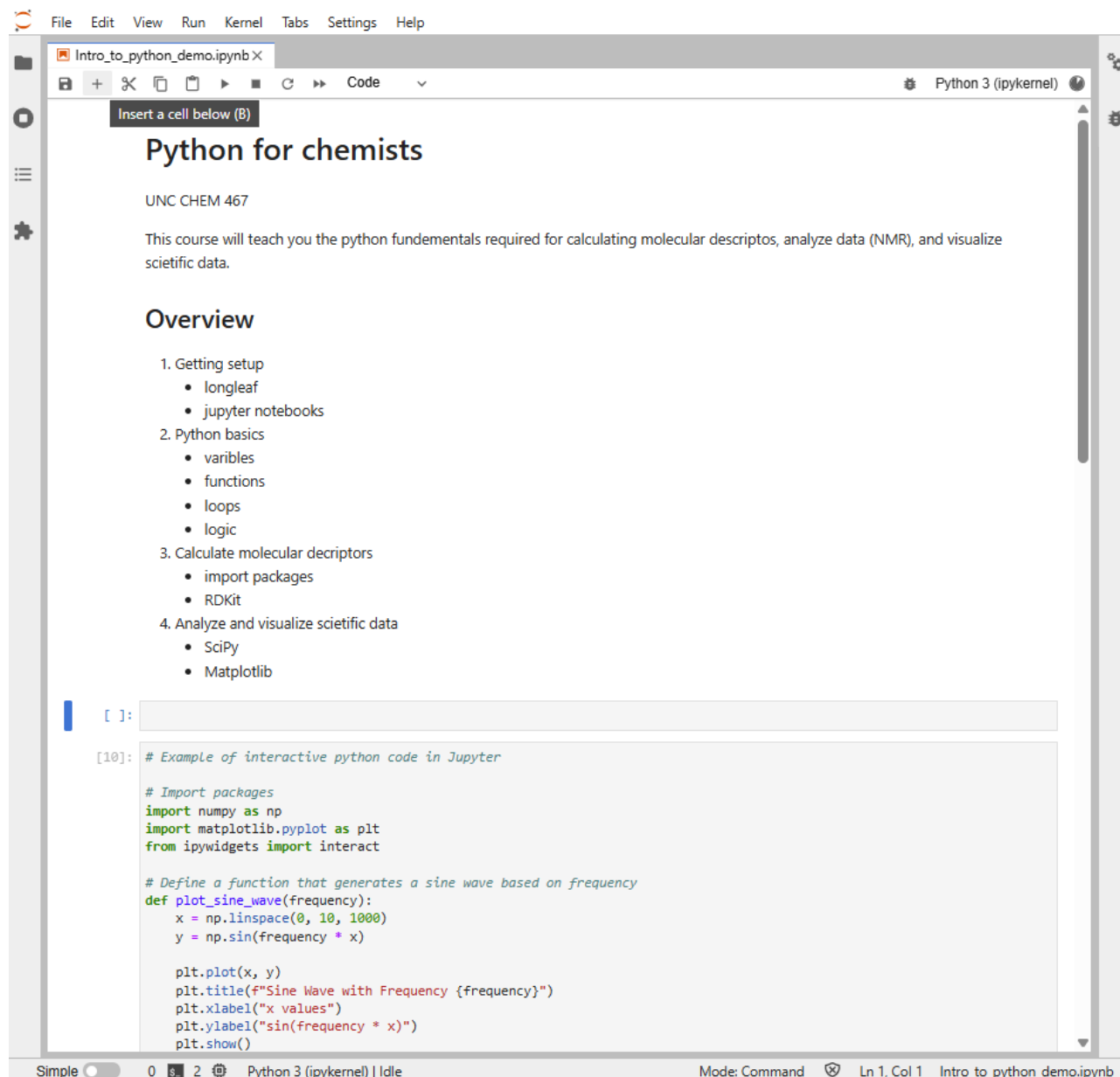
Note: move to the frequency slider to interact with the output.

How to insert and delete cells

1. Insert a new cell below your markdown cell.

- click the markdown cell
- click the **+** button in the ribbon to insert a cell below

Now your notebook should look like the following:



2. Delete the empty cell.

- right-click the cell
- click **Delete Cells**

First-Time Setup (Super important)

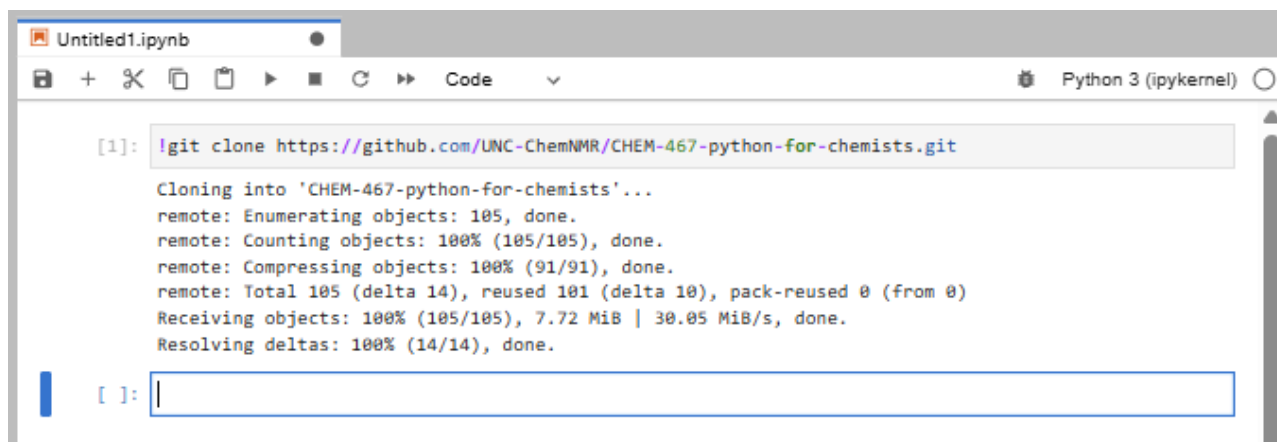
Clone the class repository

This will copy all the course material into your Longleaf storage.

1. Create a new code cell.
2. Paste this code into the code cell:

```
!git clone https://github.com/UNC-ChemNMR/CHEM-467-python-for-chemists.git
```

3. Run the code cell by pressing **shift+enter** (or click the play button ► in the notebook ribbon).
4. You should see an output that looks similar to this:

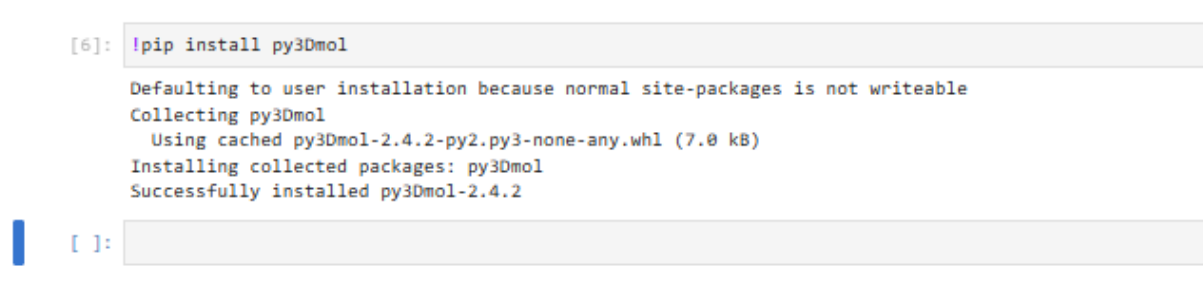


The screenshot shows a Jupyter Notebook window titled 'Untitled1.ipynb'. The code cell contains the command `!git clone https://github.com/UNC-ChemNMR/CHEM-467-python-for-chemists.git`. The output shows the cloning process: 'Cloning into 'CHEM-467-python-for-chemists'...', 'remote: Enumerating objects: 105, done.', 'remote: Counting objects: 100% (105/105), done.', 'remote: Compressing objects: 100% (91/91), done.', 'remote: Total 105 (delta 14), reused 101 (delta 10), pack-reused 0 (from 0)', 'Receiving objects: 100% (105/105), 7.72 MiB | 30.05 MiB/s, done.', and 'Resolving deltas: 100% (14/14), done.'.

Install py3Dmol

1. Create a new code cell.
2. Paste this code into the code cell:

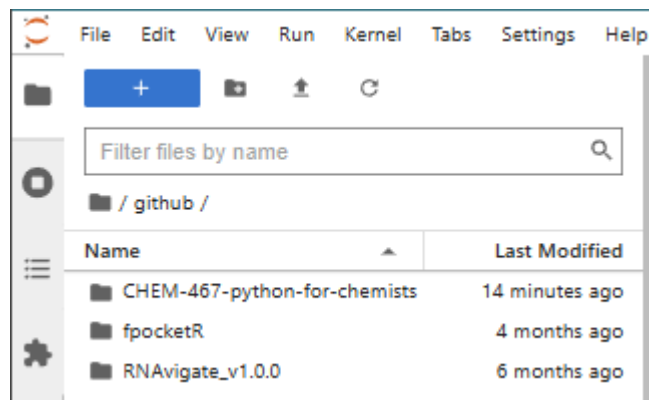
```
!pip install py3Dmol
```
3. Run the code cell by pressing **shift+enter** (or click the play button ► in the notebook ribbon).
4. You should see an output that looks similar to this:



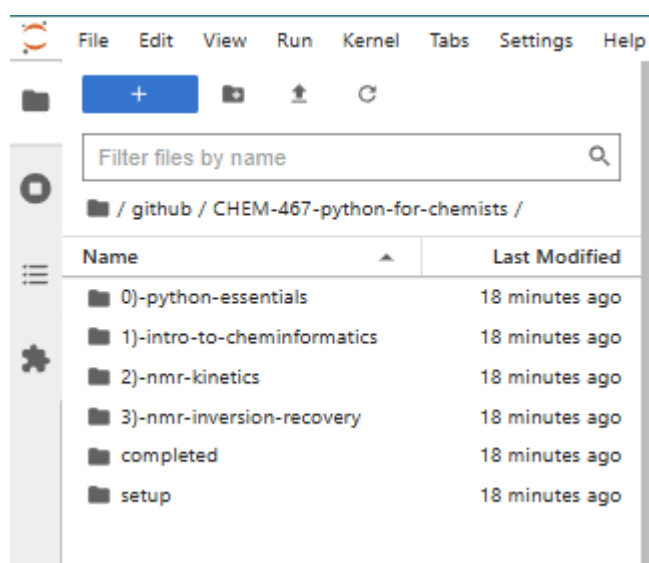
The screenshot shows a Jupyter Notebook cell with the command `!pip install py3Dmol`. The output shows the installation process: 'Defaulting to user installation because normal site-packages is not writeable', 'Collecting py3Dmol', 'Using cached py3Dmol-2.4.2-py2.py3-none-any.whl (7.0 kB)', 'Installing collected packages: py3Dmol', and 'Successfully installed py3Dmol-2.4.2'.

Navigating the course material in Jupyter

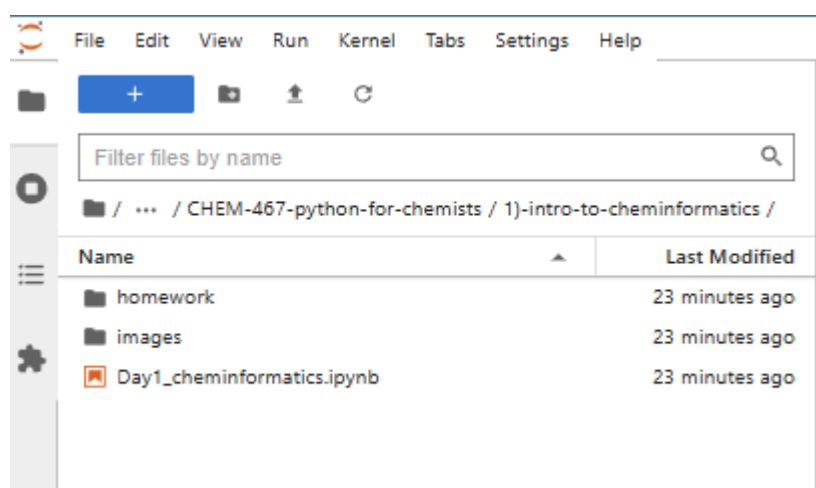
1. You should now see a folder (also called a directory) containing all the course material in the **File Browser** panel:



2. You can double-click on the course folder to open it.



3. In-class notebooks (.ipynb) and homework templates (.ipynb) are located inside the numerated sub-directories.



You are finished setting up Longleaf and Jupyter for CHEM 467!

Additional resources

Hopefully you have a good idea of how to run a Jupyter notebook in Longleaf now! If you have more questions take a look at the following resources.

- [Longleaf OnDemand](#)

- [Jupyter notebook user interface](#)