2019-01-17 Lab 1

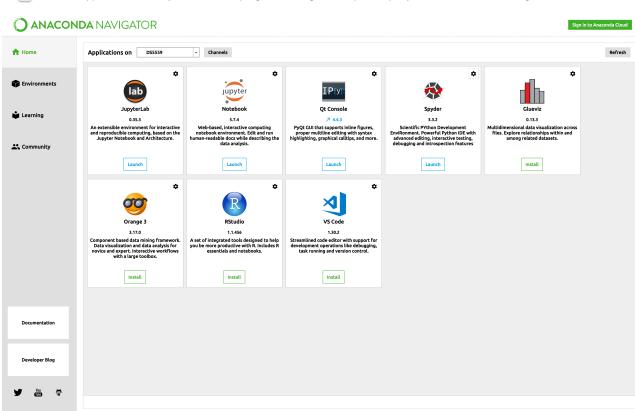
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Exercise 1: Install Anaconda

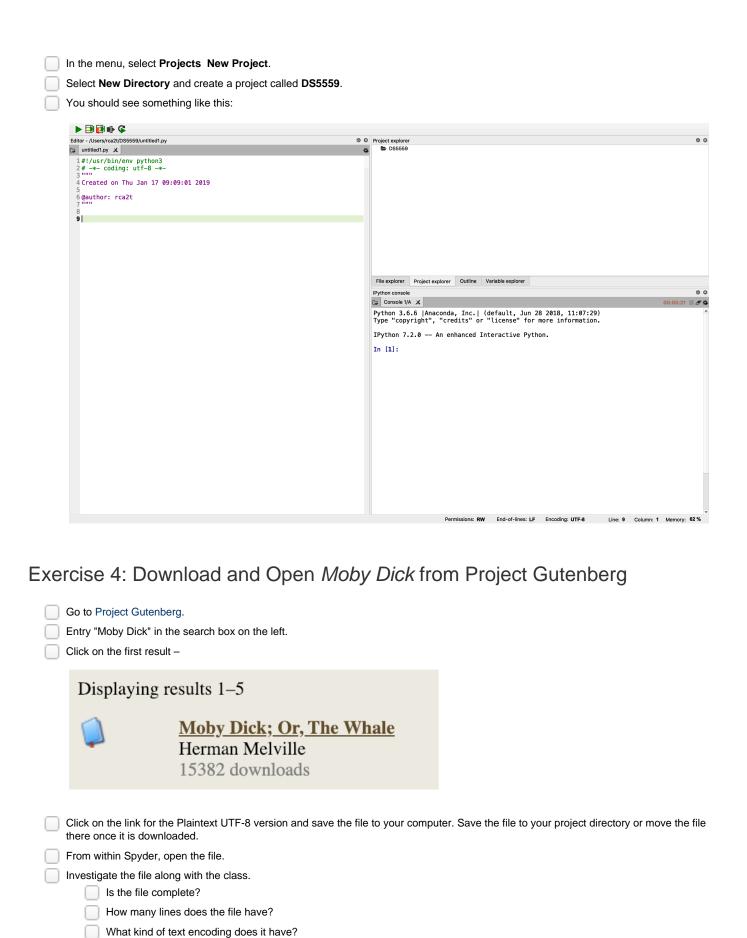
- Go to the Anaconda download page, find the distribution for your operating system, and download it.

 Install it once it is downloaded.
- Find the application icon in your toolbar or program manager and open it up you should see something like this:



Exercise 2: Install Spyder and JupyterLabs

- In Anaconda Navigator, locate the cards for Jupyterlab and Spyder and click the Install button for each. You may also install Jupyter Notebook if you'd like.
 Once installed, click the Launch button to run it.
 Spyder should appear.
- Optionally configure the panels as instructed.



What kind of line endings does it have?

Does the file come with metadata?
What content do we want to keep?
Where does the cruft begin and end?

Exercise 5: Open File in Python and Convert to Text From 1

Create a new Python file called moby.py.

Follow instructor for specific commands.

Code

```
# -*- coding: utf-8 -*-
import re
import pandas as pd
# Identify the source text (F0)
src_file = '2701-0.txt'
# Import the text as list of lines
lines = open(src_file, 'r', encoding='utf-8').readlines()
# Trim the cruft we identified
lines = lines[340:21964]
# Convert the lines into one big line, preserving line breaks
bigline = ''.join(lines)
# Split the bigline into paragraphs
paras = re.split(r'\n\n+', bigline)
# Break line into paragraphs
# Split by non-character
paras2 = []
for para in paras:
    tokens = re.split(r'\W+', para)
    paras2.append(tokens)
# Split by non-character but keep them
paras3 = []
for para in paras:
    tokens = re.split(r'(\W+)', para)
    paras3.append(tokens)
# Split by non-character using list comprehension
```

```
paras4 = [re.split(r'\W+', para) for para in paras]
# Try in Pandas
# Import paragraphs into a data frame
df = pd.DataFrame(paras, columns=['line'])
df.index.name = 'line_id'
df.line = df.line.str.strip()
# Tokenize using this one trick
df2 = df.line.str.split(r'\W+', expand=True)\
    .stack()\
    .to_frame()\
    .rename(columns={0: 'token'})
df2.index.names = ['line_id', 'token_id']
# Do a simple normalization
df2['norm'] = df2.token.str.lower()
# Get top N tokens
N = 30
df2['norm'].value_counts().head(N).sort_values().plot(kind='barh')
```

```
# Visualize dispersion plots of 'ahab' and 'whale'
(df2['norm'] == 'whale').astype('int').plot(figsize=(10, 1))
(df2['norm'] == 'ahab').astype('int').plot(figsize=(10, 1))
```