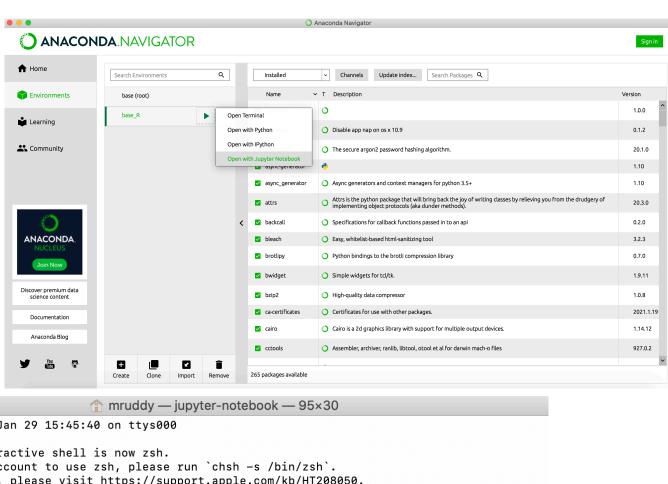
Technical Issues

Connecting to R Kernel (Mac)

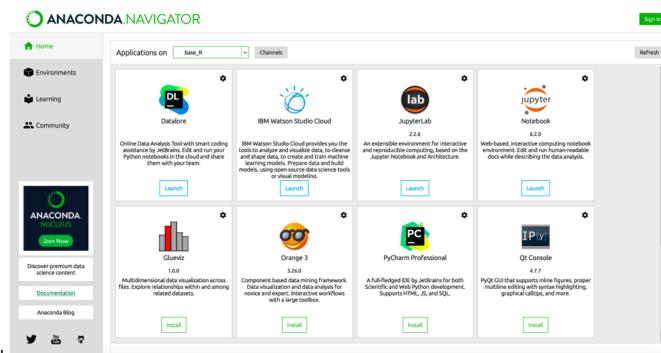
There are multiple ways to launch Jupyter Notebook after installed Anaconda



1. Environment Tab in Navigator Last login: Fri Jan 29 15:45:40 on ttys000 The default interactive shell is now zsh. To update your account to use zsh, please run `chsh -s /bin/zsh`. For more details, please visit https://support.apple.com/kb/HT208050. [(base) ml-its-902022:~ mruddy\$ conda activate base_R [(base_R) ml-its-902022:~ mruddy\$ jupyter notebook [I 15:55:47.137 NotebookApp] The port 8888 is already in use, trying another port. [I 15:55:47.138 NotebookApp] The port 8889 is already in use, trying another port. [I 15:55:47.138 NotebookApp] The port 8890 is already in use, trying another port. [I 15:55:47.139 NotebookApp] The port 8891 is already in use, trying another port. [I 15:55:47.139 NotebookApp] The port 8892 is already in use, trying another port. [I 15:55:47.258 NotebookApp] JupyterLab extension loaded from /Users/mruddy/opt/anaconda3/envs/ base_R/lib/python3.8/site-packages/jupyterlab [I 15:55:47.258 NotebookApp] JupyterLab application directory is /Users/mruddy/opt/anaconda3/en vs/base_R/share/jupyter/lab [I 15:55:47.260 NotebookApp] Serving notebooks from local directory: /Users/mruddy [I 15:55:47.260 NotebookApp] Jupyter Notebook 6.2.0 is running at: [I 15:55:47.260 NotebookApp] http://localhost:8825/?token=64799d780d483ce7e94ec47cbb867bc139e6d d0781f65121 [I 15:55:47.260 NotebookApp] or http://127.0.0.1:8825/?token=64799d780d483ce7e94ec47cbb867bc13 9e6dd0781f65121 [I 15:55:47.260 NotebookApp] Use Control-C to stop this server and shut down all kernels (twice to skip confirmation). [C 15:55:47.265 NotebookApp] To access the notebook, open this file in a browser: file:///Users/mruddy/Library/Jupyter/runtime/nbserver-364-open.html

http://localhost:8825/?token=64799d780d483ce7e94ec47cbb867bc139e6dd0781f65121

2. Command Line



3. Launch command under the Home Tab in Navigator ### !!!Known Issues!!!

Or copy and paste one of these URLs:

With the current version of Anaconda Navigator, you may have trouble connecting the R kernel when using Method 3. If you are using Method 1 or 2 to launch Jupyter notebook and are unable to connect to the R kernel, try removing and then re-creating the environment to use R. If this fails try uninstalling and reinstalling anaconda navigator (See here for details on how to do this). Otherwise please contact me.

Exporting .pdf files using pdf via LaTeX (Mac)

- Currently when running Jupyter notebook using method 1 above, attempting to export your notebook as a pdf via LaTeX returns an error (even if you have installed MiKTeX). However, it seems to work if you launch Jupyter notebook using method 2. I am still investigating why this is the case.
- In any case, I suggest a slightly longer method to create a nice .pdf of your Jupyter notebook:
 - 1. First download the notebook as an .html file
 - 2. Open the .html file with your favorite browser
 - 3. Export as this as an .html

While converting your notebook to a .pdf using LaTeX can give good results, I find that the .html sometimes looks much nicer and cleaner.

Interesting questions about R

1. Why does seq(1,10) output a sequence starting at 1? (by Ned Azar)

1. 1 2. 2 3. 3 4. 4 5. 5

In [1]:

6. 6 7. 7 8.8

seq(1,10)

9. 9 10. 10

Unlike some other programming languages (Python for instance), R starts indexing lists and sequences at 1 rather than 0. Which is "better" is subjective and depends on what it is you are trying to do.

- Fun fact: There are similar debates in mathematics for whether the "natural numbers" should start at 0 or 1.
- 1. What is the difference between seq(1,10) and seq.int(1,10)? (by Mia Kobayashi)

In [2]:

```
seq(1,10)
seq.int(1,10)
```

1. 1

2. 2 3. 3 4. 4

5. 5 6. 6 7. 7 8.8

9. 9 10. 10

1. 1 2. 2 3. 3

4. 4

5. 5 6. 6 7. 7 8.8 9. 9 10. 10

This is a non-trivial question with an interesting answer and history. It is best described by the most upvoted answer here.

TLDR: They do the same thing, but in slightly different ways. Using either is fine, but to be safe, stick with seq for now.