Ipython Notebook

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1 Notebook readability

1.1 Structure

The first cell of your notebook should contain all of your imports, so that the dependencies are clear to the reader. It also makes correction of partial pieces of your code easier.

Your cells should be as atomic as possible, meaning that one cell corresponds to an independent piece of code (for example, one question or one subquestion). Independent cells make your code easier to read. They also avoir running code uselessly: when you need to rerun a cell (to find a bug, to modify a plot, because you corrected a typo, etc.), if you have some heavy computation at the beginning, you have to execute it again which is a waste of time. If this piece of code is isolated in its own cell, you can run it only once and debug/plot in the following one.

1.2 Presenting results

You can add a cell in Markdown format, to interpret your results or explain your algorithms. Avoid using print () as much as possible (if the cell is long, it is difficult to track its orgin). Also, Markdown format supports LaTex, so you can include formulas in your text.

When you use print (), do not print too much stuff (large arrays, digits with 20 decimal).

If you are timing code, you should use the %timeit magic, which does all the work for you and format results nicely (is it really useful to have precision up to 10^{-18} s?). Which one of the following alternatives do you prefer:

```
In [1]: import time
    import numpy as np
    np.random.seed(2406)
    n, p = 1000, 5000
    X = np.random.randn(n, p)
    alpha = 1e-5
In [2]: %timeit -n 1 -r 1 np.linalg.inv(np.dot(X.T, X) + alpha * np.eye(p))
1 loop, best of 1: 4.82 s per loop

In [3]: start = time.time()
    test = np.linalg.inv(np.dot(X.T, X) + alpha * np.eye(p))
    end = time.time()
    print(end - start)
    print("total time: %f s" % (end - start))
4.829550504684448
total time: 4.829551 s
```

%timeit avoids polluting your namespace with start and end variables. You can use the n parameter to modify the numbers of times the code is run in a loop, and r to change the number of loops performed. You can use %%timeit to time the execution of a whole cell rather than a single line of code. Read the doc for more info.

1.3 import * and %pylab inline

Unless you know exactly what you are doing, I advise you against using %pylab inline or from myModule import *, because you (and the reader) end up forgetting where the variables are imported from. For example, there are differences between numpy.linalg.norm and scipy.linalg.norm, or between numpy.linalg.eig and scipy.linalg.eig; if they are just called norm and eig with no explicit import to trace them, it's confusing.

Let's see what %pylab inline imports:

```
In [1]: print(len(dir())) # dir() returns the list of names in the current local scope
23
In [2]: %pylab inline
Populating the interactive namespace from numpy and matplotlib
In [3]: print(len(dir()))
```

So there are nearly a thousand additional variables polluting your scope now (autocompletion now suggests lots of choices), and if you don't do this in the first cell, you might override some variables: common variable names such as test, clf... are now bound to variables imported by pylab.

1.4 Prototyping

Because of their interactive properties, notebooks are a very powerful tool to prototype. Everyone tends to rerun cells multiple times, change their order, change variable names... Because deleting a cell does not delete variables defined in it (which is a good thing), some variables may continue to exist even though you don't declare them anymore, giving you the false impression that your code runs fine whereas another person won't be able to run it.

 \rightarrow before handing over your notebook, restart your kernel and run all cells to check that everything is fine, even if it takes a few minutes.

2 Useful shortcuts

2.1 In command mode

Command mode is when you are not editting the content of a cell. If you are editting a cell, you have to press Esc to enable command mode.

- The most important one: h to show the shortcuts
- ctrl + c to execute a cell a keep this cell selected
- 1 + ← to execute this cell and select the next
- alt + 🗀 to execute the current cell and create an empty one below
- I, I (pressed twice) to interrupt the kernel
- x to cut the cell
- c to copy the cell
- v to paste the copied cell (below currently selected cell)
- 1 + v to paste the copied cell (above currently selected cell)
- d , d (pressed twice) to delete cell. Can be undone with ctrl + z .
- a to insert cell above
- b to insert cell **b**elow

2.2 In edit mode

Edit mode when you are editting the content of a cell (your cursor is blinking inside the cell).

- ctrl + ↑ to go to start of cell
- [ctrl] + [trl] to go to end of cell
- ctrl + 1 + to split cell (very useful but does not work on every keyboard...)
- ctrl + to go one word left. Also works with right. When used while holding ①, this allows you to quickly select portions of text.
- [ctrl] + [ctrl] + [ctrl] and [alt] + [ctrl] also work in edit mode.

When typing the name of a function, hitting $\widehat{\Box} + \longrightarrow$ (tab, not right arrow) after the function name (or inside the parenthesis of the function call) will display a short version of this function's help. This is very useful to remember the order of the parameters, their names or their default values.

2.3 Commenting and uncommenting blocks of code (more advanced)

There is no default shortcut, but you can add a custom one (in this case, ctrl + ,). Add these lines to your custom. js file:

If you are using anaconda 3 as recommended, your custom. js should be located in

```
~/anaconda3/envs/myCondaEnvName/lib/python2.7/site-packages/notebook/static/custom/custom.js
```

if you're using a conda env (python2.7 is the version of python inside the conda env, change its name accordingly), or in

```
~/anaconda3/lib/python3.5/site-packages/notebook/static/custom/custom.js
```

if you use plain conda (replace python3.5 with your python version).

Once this is done, restart jupyter. You can comment and uncomment blocks selecting them, then hitting [ctr] + [,].

3 bash

It is possible to type bash commands inside a notebook; just type! before the line of code. This is useful to avoid having to switch between your navigator and your file explorer, opening files to see the first lines, etc. You can do all this from inside your notebook:

```
In [3]: !jupyter nbconvert --to latex bash.ipynb # exporting this ipynb to LaTeX format
[NbConvertApp] Converting notebook bash.ipynb to latex
[NbConvertApp] Writing 15898 bytes to bash.tex
```

4 Frequent issues

4.1 My plots are invisible

Did you use %matplotlib inline after import matplotlib.pyplot as plt?

4.2 Source filed has changed, but nothing changes in the notebook

When you modify a source file you import in your notebook, it's safer to restart your kernel rather than just rerunning from my_source import my_function. import in Python means "if it has not already been imported, import", so rerunning import does not actually reload the function. If restarting your kernel is too expensive, have a look at the autoreload module.

4.3 Python version issue

If you have trouble with the version of Python you're using (python 2 and python 3 installed, virtualenvs, conda envs...) and you want to check which one is used, type:

```
import sys
print(sys.executable)
```

Be aware that silent issues can happen if you start jupyter notebook from inside a conda env, while ipython is not installed inside this env: the executable used is conda root instead of the one of your conda env; in that case, printing <code>sys.executable</code> will show you that you are using the wrong executable of python (which explains why you have the wrong version or you cannot import your modules).

4.4 Module and module version

If you have troubles importing a module, this StackOverflow question shows how to print a list all installed modules. myModule.__path__ and myModule.__version__ can also be useful.