Week 2: Computing in python I, Some useful packages and tools MSc/MRes CMEE 2014-15

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A FEW THINGS FIRST

- Any issues with version control and pushing your Week 1? DO NOT clone cmee2014masterepo into cmeecoursework
- If you don't have ipdb (let me know if you still don't get ipdb):

sudo apt-get install python-ipdb

- Let's discuss the dictionary.py and yesterday's Regex exercises in the afternoon
- We will do a quick recap of what we did in the whole week tomorrow morning (back in CPB!)

USEFUL PYTHON PACKAGES

- These are always available as standard libraries (just require import from within python or ipython):
 - io: file input-output with *.csv, *.txt, etc.
 - subprocess: to run other programs, including multiple ones at the same time, including operating system-dependent functionality
 - sqlite3: for manipulating and querying sqlite databases
 - math: for mathematical functions
- These you will have to install in terminal using sudo apt-get install python-packagename (as you did for ipdb earlier)):
 - scipy: for scientific computing
 - matplotlib: for plotting (very matlab-like, requires scipy) (all packaged in pylab)
 - scrapy: for writing web spiders that crawl web sites and extract data from them
 - beautifulsoup: for parsing HTML and XML (can do what scrapy does)
 - biopython: for biological computation, including bioinformatics

USING python TO PATCH TOGETHER WORK-FLOWS, I

- You can use python to build an automated work-flow that involves multiple applications (R, LATEX, bash, etc.)
- For example, you could, in theory, write a single Python script to generate and update your MSc/MRes dissertation, tables, plots, and all!
- the subprocess modules will enable you to do this
- Let's try first launch ipython, then cd to your python code directory, and type:

```
import subprocess
subprocess.os.system("geany boilerplate.py")
subprocess.os.system("gedit ../Data/TestOaksData.csv")
subprocess.os.system("python boilerplate.py") # A bit silly!
```

Easy as pie!

USING python TO PATCH TOGETHER WORK-FLOWS, II

Similarly, to compile your Latex document (using pdflatex in this case):

```
subprocess.os.system("pdflatex yourlatexdoc.tex")
```

• You can also do this (instead of using subprocess.os):

```
subprocess.Popen("geany boilerplate.py", shell=True).wait()
```

- You can also run $\mathbb R$ in a similar way (More on this in Python Week 2)
- Why am I telling you all this now? Because of the next two weeks (esp. GIS)

Using python to patch together work-flows,

- You can also use subprocess.os to make your code OS (Linux, Windows, Mac) independent
- For example to assign paths:

```
subprocess.os.path.join('directory', 'subdirectory', 'file')
```

- The result would be appropriately different on Windows!
- Note that in all cases you can "catch" the output of subprocess so that you can then use the output within your python script:

```
MyPath = subprocess.os.path.join('directory', 'subdirectory', 'file')
```

• Explore what subprocess can do by tabbing subprocess. (so also for submodules, e.g., subprocess.os. an then tab)



- The python package/library scipy can help you do serious number crunching:
 - Linear algebra
 - Numerical integration
 - Fourier transforms
 - Interpolation
 - Special functions (Incomplete Gamma, Bessel, etc.)
 - Random numbers and statistical functions
- Let's have a quick look at scipy
- In the following, we will use the array class in scipy for data manipulations and calculations (I mentioned this on the first day)
- Scipy arrays are similar in some respects to python lists, but are more naturally multidimensional, homogeneous in type, and allow efficient manipulations

```
In [1]: import scipy
In [2]: a = scipy.array(range(5))
In [3]: a
Out[3]: array([0, 1, 2, 3, 4])
In [4]: a = scipy.array(range(5), dtype = float)
In [5]: a
Out[5]: array([ 0., 1., 2., 3., 4.])
In [6]: a.dtype
Out[6]: dtype('float64')
In [7]: x = scipy.arange(5)
In [8]: x
Out[8]: array([0, 1, 2, 3, 4])
In [9]: x = scipy.arange(5.)
In [10]: x
Out[10]: array([ 0., 1., 2., 3., 4.])
```

```
In [11]: x.
                            x.fill
x.T
              x.conj
x.nbytes
             x.round
                           x.take
x.all
           x.conjugate ...
x.searchsorted x.tofile
             x.conjugate x.flags
x.ndim
x.anv
             x.copy
                           x.flat
x.newbyteorder x.setfield x.tolist
             x.ctypes x.flatten
x.argmax
x.nonzero
             x.setflags x.tostring
             x.cumprod
                           x.getfield
x.argmin
x.prod
             x.shape
                           x.trace
x.argsort x.cumsum
                            x.imaq
      x.size
x.ptp
                            x.transpose
x.astype x.data
                           x.item
           x.sort
x.put
                            x.var
           x.diagonal x.itemset
x.squeeze x.view
x.base
            x.squeeze
x.ravel
x.byteswap
           x.dot
                           x.itemsize
x.real
             x.std
           x.dtype
x.choose
                            x.max
           x.strides
x.repeat
x.clip
           x.dump
                            x.mean
x.reshape
             x.sum
x.compress x.dumps
                            x.min
x.resize
              x.swapaxes
In [11]: x.tolist()
Out[11]: [0.0, 1.0, 2.0, 3.0, 4.0]
In [12]: x.shape
Out[12]: (5,)
```

```
In [14]: mat = scipy.array([[0, 1], [2, 3]])
In [16]: mat.shape
Out[16]: (2, 2)
In [17]: mat[1]
Out[17]: array([2, 3])
In [18]: mat[0,0]
Out[18]: 0
In [19]: mat.ravel() # flatten!
Out[19]: array([0, 1, 2, 3])
In [20]: scipy.ones((4,2))
Out [20]:
array([[ 1., 1.],
     [ 1., 1.],
      [ 1., 1.],
      [ 1., 1.]])
In [21]: scipy.zeros((4,2))
Out [21]:
array([[ 0., 0.],
    [ 0., 0.],
      [ 0., 0.],
      [ 0., 0.]])
```

```
In [22]: scipy.identity(4)
Out [22]:
array([[ 1., 0., 0., 0.],
     [ 0., 1., 0., 0.],
      [ 0., 0., 1., 0.],
      [ 0., 0., 0., 1.]])
In [23]: m = scipy.identity(4)
In [24]: m.reshape((8, 2))
Out [24]:
array([[ 1., 0.],
   [ 0., 0.],
      [ 0., 1.],
      [ 0., 0.],
     [ 0., 0.],
      [ 1., 0.],
      [ 0., 0.],
      [ 0., 1.11)
In [25]: m.fill(16)
In [26]: m
Out [26]:
array([[ 16., 16., 16., 16.],
    [ 16., 16., 16., 16.],
     [ 16., 16., 16., 16.],
     [ 16., 16., 16., 16.]])
```

Let's perform some common operations on arrays:

```
In [1]: import scipy
In [2]: mm = scipy.arange(16)
In [3]: mm = mm.reshape(4,4)
In [4]: mm
Out [4]:
array([[ 0, 1, 2, 3],
     [4, 5, 6, 7],
      [8, 9, 10, 11].
      [12, 13, 14, 15]])
# NOTE: Rows first
In [5]: mm.transpose()
Out [51:
array([[ 0, 4, 8, 12],
      [ 1, 5, 9, 13],
      [2, 6, 10, 14],
       [ 3, 7, 11, 15]])
In [6]: mm + mm.transpose()
Out [6]:
array([[ 0, 5, 10, 15],
     [ 5, 10, 15, 20],
      [10, 15, 20, 25],
      [15, 20, 25, 30]])
```

```
In [7]: mm - mm.transpose()
Out [7]:
arrav([[0, -3, -6, -9],
     [ 3, 0, -3, -6],
      [6, 3, 0, -3],
      [ 9, 6, 3, 011)
In [8]: mm * mm.transpose()
## Elementwise!
Out [8]:
array([[ 0, 4, 16, 36],
     [ 4, 25, 54, 91],
      [ 16, 54, 100, 154],
       [ 36, 91, 154, 225]])
In [9]: mm / mm.transpose()
Warning: divide by zero encountered in divide
# Note the integer division
Out [9]:
array([[0, 0, 0, 0],
   [4, 1, 0, 0],
      [4, 1, 1, 0],
      [4, 1, 1, 1]])
```

- We can do a lot more (but won't!) by importing the linalg sub-package: import scipy.linalg
- Two other particularly useful scipy sub-packages are:
 - scipy.integrate (what will I need this for?)
 - scipy.stats (why not use R for this?)



Let's take a quick spin in scipy.stats!

```
In [18]: import scipy.stats
In [19]: scipy.stats.
scipv.stats.arcsine
                                 scipv.stats.lognorm
scipy.stats.bernoulli
                                 scipy.stats.mannwhitneyu
scipy.stats.beta
                                 scipy.stats.maxwell
scipy.stats.binom
                                 scipy.stats.moment
scipy.stats.chi2
                                scipy.stats.nanstd
scipy.stats.chisgprob
                                scipy.stats.nbinom
scipy.stats.circvar
                                scipv.stats.norm
scipv.stats.expon
                                scipv.stats.powerlaw
scipy.stats.gompertz
                       scipy.stats.t
                                scipy.stats.uniform
scipy.stats.kruskal
In [19]: scipv.stats.norm.rvs(size = 10) # 10 samples from
N(0,1)
Out [19]:
array([-0.951319, -1.997693, 1.518519, -0.975607, 0.8903,
      -0.171347, -0.964987, -0.192849, 1.303369, 0.6728])
In [20]: scipy.stats.norm.rvs(5, size = 10)
# change mean to 5
Out [20]:
array([ 6.079362, 4.736106, 3.127175, 5.620740, 5.98831,
       6.657388, 5.899766, 5.754475, 5.353463, 3.243201)
```

- Let's look at an example using scipy.integrate
- Create LV1.py in your Week2/Code directory

READINGS AND RESOURCES

- www.matplotlib.org/
- For SciPy, the official documentation is best:
 docs.scipy.org/doc/scipy/reference/
 Read about the scipy modules you think will be important to you...
- Many illustrative examples at http://wiki.scipy.org/Cookbook (including Lotka-Volterra!)
- In general, good module-specific cookbooks are out there (e.g., biopython)

PRACTICAL 2 I

- Get Pracs 0 and 1 sorted out!
- Don't forget to bring today's (functional) scripts under version control LV1.py

PRACTICAL 2 II

- Convert LV1.py into another script called LV2.py that does the following:
 - Take arguments for the four LV model parameters r, a, m , e from the commandline

```
LV2.py arg1 arg2 ... etc
```

- 2 Runs the Lotka-Volterra model with prey density dependence $rx(1-\frac{x}{k'})$
- Saves the plot as .pdf in an external results directory (Week2/Results)
- The chosen parameter values should show in the plot (e.g., r = 1, a = .5, etc)
- You change time length t too
- Also include a script called run_LV2.py in Code that will run LV2.py with appropriate arguments



PRACTICAL 2 III

- Extra credit if you also choose appropriate values for the paramaters such that both predator and prey persist under in model with prey density dependence
- Extra-extra credit if you can write a recursion version of the model in discrete time (what's this?) – it should do everything that run_LV2.py does
- Extra-extra-extra credit if you can write a recursion versiaon of the model in discrete time with a random gaussian fluctuation at each time-step (use scipy.stats)

PRACTICAL 2 IV

• Complete the code *blackbirds.py* that you find in the master repo (necessary data file is also there)