

PHYS4038/MLiS and ASI/MPAGS

Scientific Programming in



mpags-python.github.io

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Exercises 5

- 1) Only using `scipy.special.yn`, plot the zeroth-order Bessel function of the second kind, and use `fsolve` to find the value of the third root, i.e. the third lowest value of x where $Y_0(x) = 0$.
- 2) Use `quad` to find the integral of $Y_0(x)$ between $x=0$ and the third root.
- 3) Use `scipy.stats.norm.rvs` to create 100 samples from a Normal distribution for some mean and sigma. Then use `plt.hist` to create a 10-bin histogram of the samples (see the return values). Determine the centre of each bin.
- 4) Create a function `f((m, s), a, x, y)` which returns the sum of the squared residuals between the values in `y` and a Gaussian with mean `m`, sigma `s` and amplitude `a`, evaluated at `x`.
- 5) Use function you created in (4) with `scipy.optimize.minimize` to fit a Gaussian to the histogram created in (3). Plot and compare with `scipy.stats.norm.fit`.

Any questions?

- skype (spbamford): <https://join.skype.com/KpW5oCLNNijt>
- email steven.bamford@nottingham.ac.uk

An introduction to scientific programming with



Session 6:
Data handling

Databases

- Python has tools for accessing most (all?) databases
 - e.g. MySQL, SQLite, MongoDB, Postgres, ...
- Allow one to work with huge datasets
- Data can be at remote locations
- Robust and fast
- May require knowledge of DB-specific language
- But often provide Pythonic interface

Databases

- SQLite
 - Lightweight
 - No server
 - Just uses files (convenient, but less powerful)
 - Standard python module: `sqlite3`

Databases

- MySQL
 - Widely used
 - Need MySQL server installed
 - Official: `mysql-connector-python`
 - `mysqlclient`, `pymysql`, `SQLAlchemy`

Databases

- MongoDB
 - NoSQL database
 - Need Mongo database server
 - Official: pymongo

Databases

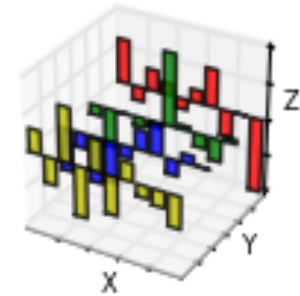
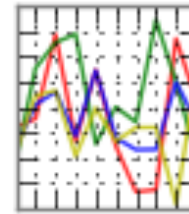
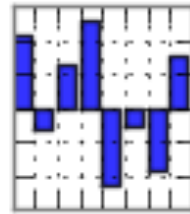
- Python has tools for accessing most (all?) databases
 - e.g. MySQL, SQLite, MongoDB, Postgres, ...
- Allow one to work with huge datasets
- Data can be at remote locations
- Fast random read and write
- Atomic transactions
- Concurrent connections

Databases

- **DB pros and cons**

- Allow one to work with huge datasets
- Data can be at remote locations
- Fast random read and write
- Concurrent, atomic transactions
- However, most databases are designed for webserver use
 - typically not optimised for data analysis
 - write once, multiple sequential reads

pandas

$$y_{it} = \beta' x_{it} + \mu_i + \epsilon_{it}$$


- Python Data Analysis Library
 - <http://pandas.pydata.org>
- Easy-to-use data structures
 - DataFrame (more friendly recarray)
 - Handles missing data (more friendly masked array)
 - read and write various data formats
 - data-alignment
 - tries to be helpful, though not always intuitive
 - Easy to combine data tables
 - Surprisingly fast!

Notebook demo...

Dask

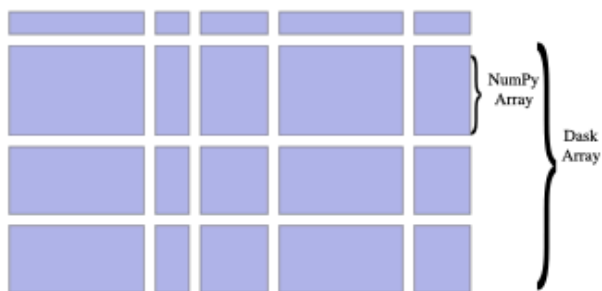


```
# Arrays implement the Numpy API
import dask.array as da
x = da.random.random(size=(10000, 10000),
                      chunks=(1000, 1000))
x + x.T - x.mean(axis=0)
```

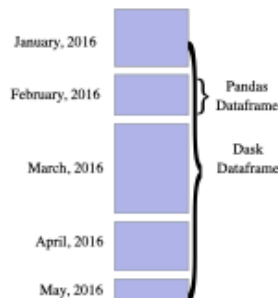
```
# Dataframes implement the Pandas API
import dask.dataframe as dd
df = dd.read_csv('s3://.../2018-*-.csv')
df.groupby(df.account_id).balance.sum()
```

```
# Dask-ML implements the Scikit-Learn API
from dask_ml.linear_model \
    import LogisticRegression
lr = LogisticRegression()
lr.fit(train, test)
```

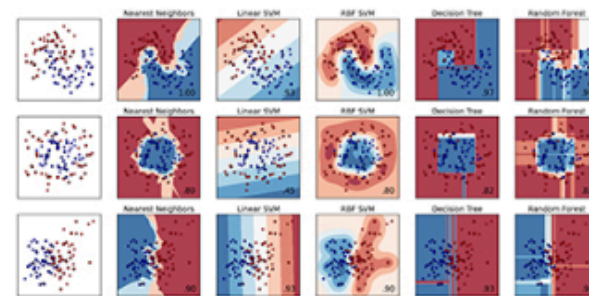
Numpy



Pandas



Scikit-Learn



PySpark



- typically for dealing with very large datasets
- distributed computing on a cluster
- need to setup infrastructure

PyTables / h5py



- <http://pytables.github.io>
- For creating, storing and analysing datasets
 - from simple, small tables to complex, huge datasets
 - standard HDF5 file format
 - incredibly fast – even faster with indexing
 - uses on the fly block compression
 - designed for modern systems
 - fast multi-code CPU; large, slow memory
- "in-kernel" – data and algorithm are sent to CPU in optimal way
- "out-of-core" – avoids loading whole dataset into memory

PyTables / h5py

- Can store many things in one HDF5 file (like FITS)
- Tree structure
- Everything in a group (starting with root group, '/')
- Data stored in leaves
- Arrays (e.g. n-dimensional images)

```
>>> from tables import *
```

```
>>> h5file = openFile("test.h5", mode = "w")
```

```
>>> x = h5file.createArray("/", "x", arange(1000))
```

```
>>> y = h5file.createArray("/", "y", sqrt(arange(1000)))
```

```
>>> h5file.close()
```

PyTables

- Tables (columns with different formats) – *better to use Pandas!*
 - described by a class
 - accessed by a row iterator

```
>>> class MyTable(IsDescription):  
        z = Float32Col()  
>>> table = h5file.createTable("/", "mytable", MyTable)  
>>> row = table.row  
>>> for i in xrange(1000):  
        row["z"] = i**(3.0/2.0)  
        row.append()  
>>> table.flush()  
>>> z = table.cols.z
```

PyTables Expr

- **Expr** enables in-kernel & out-of-core operations

```
>>> r = h5file.createArray("/", "r", np.zeros(1000))
>>> xyz = Expr("x*y*z")
>>> xyz.setOutput(r)
>>> xyz.eval()
/r (Array(1000,)) ' '
  atom := Float64Atom(shape=(), dflt=0.0)
  maindim := 0
  flavor := 'numpy'
  byteorder := 'little'
  chunkshape := None
>>> r.read(0, 10)
array([  0.          ,  1.          ,  7.999999986, 26.99999989 ,
        64.          , 124.99999917, 216.000000085, 343.000001259,
       511.99999124, 729.          ])
```


PyTables Expr

- **where** enables in-kernel selections

```
>>> r_bigish = [ row['z'] for row in  
                 table.where('(z > 1000) & (z <= 2000)' )  
  
>>> for big in table.where('z > 10000;'):  
...     print('A big z is {}'.format(big['z']))
```

- There is also a **where** in **Expr**

Multiprocessing

- Python includes modules for writing "parallel" programs:
 - `threaded` – limited by the Global Interpreter Lock
 - `multiprocessing` – generally more useful

```
from multiprocessing import Pool

def f(x):
    return x*x

pool = Pool(processes=4)      # start 4 worker processes

z = range(10)
print pool.map(f, z)  # apply f to each element of z in parallel
```

Multiprocessing

```
from multiprocessing import Process
from time import sleep

def f(name):
    print('Hello {}, I am going to sleep now'.format(name))
    sleep(3)
    print('OK, finished sleeping')

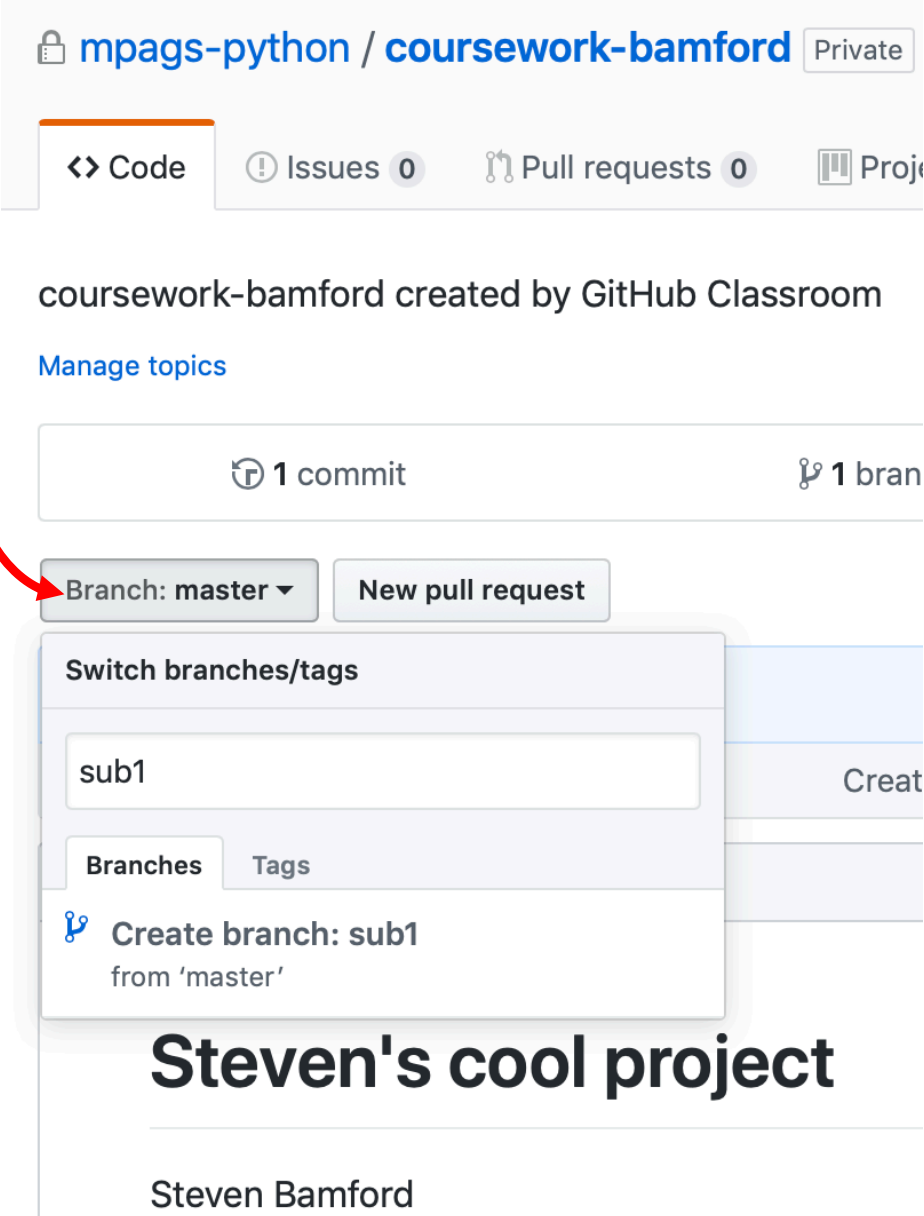
if __name__ == '__main__':
    p = Process(target=f, args=(lock, 'Steven'))
    p.start()          # start additional process
    sleep(1)           # carry on doing stuff
    print 'Wow, how lazy is that function!'
    p.join()           # wait for process to complete
```

(Really, should use a lock
to avoid writing output
to screen at same time)

```
$ python thinking.py
Hello Steven, I am going to sleep now
Wow, how lazy is that function!
OK, finished sleeping
```

Coursework submission

- Submission and feedback via your GitHub repository
- Mandatory for MLiS, optional for MPAGS
- **Create a branch called sub2**
- Should contain your draft code



The screenshot shows the GitHub interface for the repository 'mpags-python / coursework-bamford'. The repository is marked as 'Private'. Navigation tabs include 'Code', 'Issues' (0), 'Pull requests' (0), and 'Projects'. A message states 'coursework-bamford created by GitHub Classroom' with a 'Manage topics' link. Below this, it shows '1 commit' and '1 branch'. A dropdown menu for 'Branch: master' is open, displaying a 'Switch branches/tags' modal. The modal has a search bar containing 'sub1' and tabs for 'Branches' and 'Tags'. Under the 'Branches' tab, there is an option to 'Create branch: sub1 from 'master'' with a branch icon. At the bottom of the page, the text 'Steven's cool project' and the name 'Steven Bamford' are visible.

mpags-python / coursework-bamford Private

<> Code Issues 0 Pull requests 0 Projects

coursework-bamford created by GitHub Classroom

[Manage topics](#)

1 commit 1 branch

Branch: master New pull request

Switch branches/tags

sub1

Branches Tags

Create branch: sub1 from 'master'

Steven's cool project

Steven Bamford

Questions (especially about coursework code)

Any questions?

- shout and wave
- skype (spbamford)
- <https://join.skype.com/KpVW5oCLNNiJt>
- email steven.bamford@nottingham.ac.uk