Read in and manipulate data
Plotting —line data, histograms, images.
mcmaster for practice
Generating (and plotting) data Planck + density
profile

There are lots of modules to read in data from a file — try numpy modules genfromtxt or loadtxt

```
import numpy as np
help(np.genfromtxt)

x1,y1 = np.genfromtxt('dataset1.txt',unpack=True,dtype=np.float)

# that was it!
# we use unpack to tell python to throw out the two columns and we caught them with arrays x and y
# but we could have just captured whatever came out, then it just would be a merged array:
data = np.genfromtxt('dataset1.txt',dtype=np.float)
print data
print data.shape
print data[:,0]
print x1

# another nice thing, genfromtxt will read data from a URL! What?!
A = np.genfromtxt('https://raw.githubusercontent.com/jbrownlee/Datasets/master/
pima-indians-diabetes.data.csv',unpack=True, delimiter=',')
```

```
# OK, let's fit our x and y data with a straight line, first define a line function:
def myline(x, m, b):
     return m*x+b
# simple, nothing to it!
# now let's grab a function that performs a least-squares curve fit from the scipy.optimize package:
from scipy.optimize import curve fit
help(curve fit)
bestfit, covar mat = curve fit(myline, x1, y1, p0 = [1.0, 1.0])
print(bestfit)
# and overplot the best-fit:
plt.plot(x1, mylin(x1, bestfit[0], bestfit[1]),'k:')
plt.xlim((-1,21)) # limit the X range of our plot
# make a better sampled plot of our best fit line with a thicker line:
x fit = np.arange(0, 20, 0.5)
plt.plot(x fit, myline(x fit, bestfit[0], bestfit[1]), 'k--', lw=2)
# always add axis labels!
plt.xlabel('Xvalue')
plt.ylabel('Yvalue')
```

Type **python** and then enter these commands:

import numpy as np import pylab as P mu,sigma = 100, 15

x=mu + sigma*P.randn(10000)

n,bins,patches = P.hist(x,50,normed=1,histtype='stepfilled',

facecolor='green', alpha=0.75)

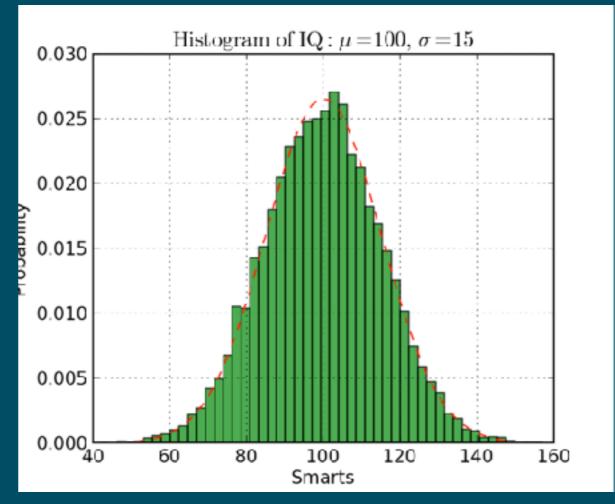
y=P.normpdf(bins,mu,sigma)

I=P.plot(bins,y,'k',linewidth=1.5)

P.xlabel('smarts')

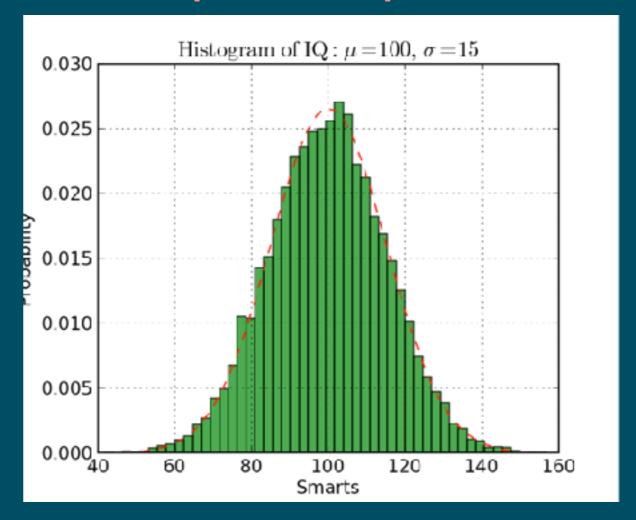
P.ylabel('Probability')

P.show()



Type **python** and then enter these commands:

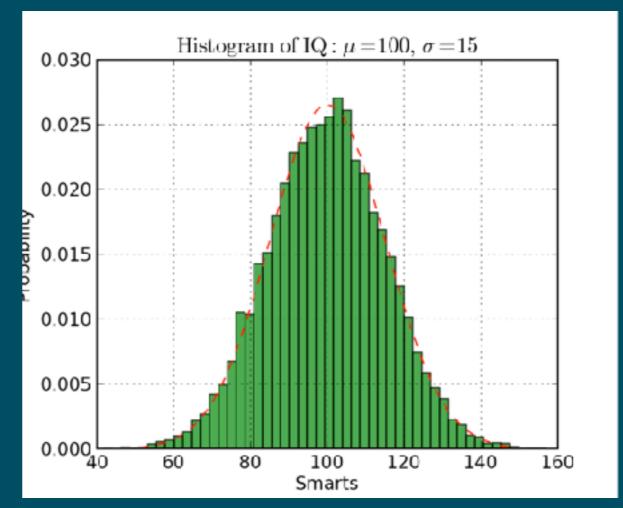
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y=P.normpdf(bins,mu,sigma)
I=P.plot(bins,y,'k',linewidth=I.5)
P.xlabel('Smarts')
P.ylabel('Probability')
P.show()
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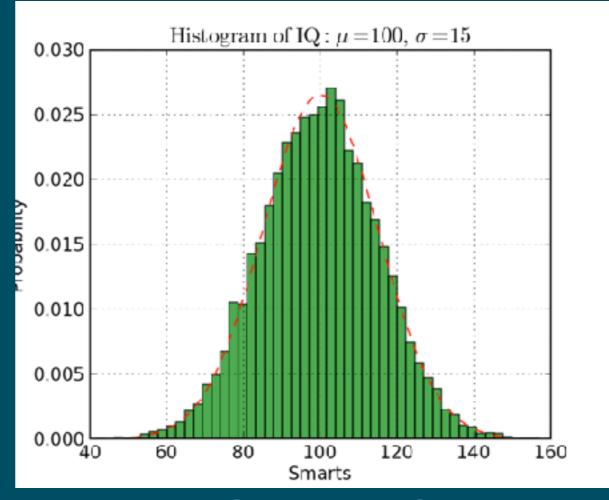
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faceco y=P.no I=P.plo

P.xlab

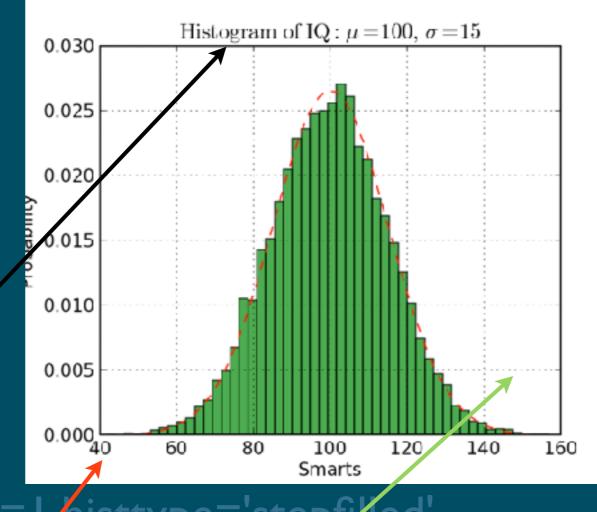
Extra bells and whistles:

Ptitle(r'\$\mathrm{Histogram/ of\ IQ:}\ \mu=100,\ \sigma=15\$')

P.axis([40, 160, 0, 0.03])

P.grid(True)

P.ylabel('Probability')
P.show()



How do we know the possible options for hist (or any python command?)

look for the documentation from the imported library -- here it's

http://matplotlib.sourceforge.net/index.html

Now you practice! Using your pretty version of mcmaster globular cluster data, make a histogram of the number of globs as function of declination

Let's make your own python script!

Write a script that will generate a file containing the Planck spectrum (wavelength and Intensity at that wavelength for many wavelengths)

$$I = \frac{2hc^2}{\lambda^5} \frac{1}{e^{hc/\lambda kT} - 1.0}$$

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intensity= ((2*h*c**2)/lambda**5)* (1.0/ (e**((h*c)/(lambda*k*T) -1.0)

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intensity= ((2*h*c**2)/wavelen**5)* (1.0/ (e**((h*c)/(wavelen*k*T) -1.0)

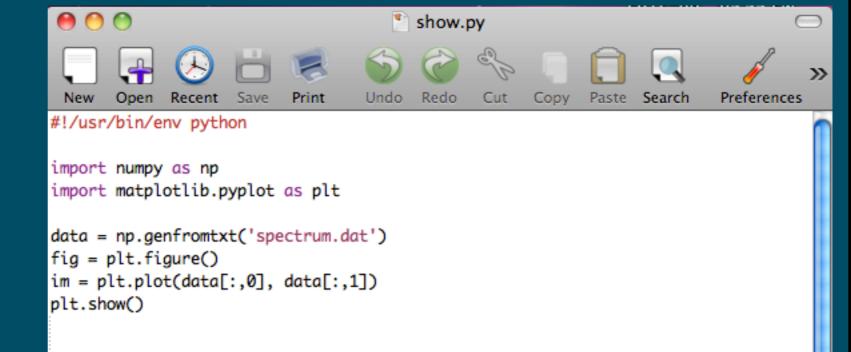
$$I = \frac{2hc^2}{\lambda^5} \frac{1}{e^{hc/\lambda kT} - 1.0}$$

Simple version:

```
planck_simple.py
       Open Recent Save
                                                                          Preferences
                                   Undo
                                                           Paste Search
#!/usr/bin/env python
from math import exp
# Set up some parameters
wavelen_min = 1.0e-7
                        # Minimum wavelength in meters
                        # Maximum wavelength in meters
wavelen_max = 6.0e-5
samples = 100
                        # Number of steps to output
T = 300
                        # Temperature in Kelvin
# Define some fundamental constants in SI units
c = 3.0e8
                        # Speed of light
h = 6.626e - 34
                        # Plank's constant
                         # Boltzmann's constant
k = 1.38e-23
# Open file we want to write to
file = open('spectrum.dat', 'w')
# Calculate plank spectrum and write to file in a loop
for i in range(samples):
  wavelen = wavelen_min + float(i)*(wavelen_max - wavelen_min)/float(samples)
  radiance = (2 * h * c**2 / wavelen**5) * (1.0 / (exp(h * c / (wavelen * k * T)))
) - 1.0))
  file.write("%g %g\n" % (wavelen, radiance))
# Close the file now that we're finished
file.close()
-: **- planck_simple.py All (1,0)
                                  (Python)
```

Fancy version:

```
planck_fancy.py
                                                                         Preferences
       Open Recent Save
                                                    Copy Paste Search
#!/usr/bin/env python
import sys
import numpy as np
# Set up some parameters
wavelen min = 1.0e-7
                        # Minimum wavelength in meters
wavelen_max = 6.0e-5
                        # Maximum wavelength in meters
samples = 100
                        # Number of steps to output
# Define some fundamental constants in SI units
c = 3.0e8
                        # Speed of light
h = 6.626e - 34
                        # Plank's constant
k = 1.38e-23
                        # Boltzmann's constant
def plank(T, wavelen):
 radiance = (2 * h * c**2 / wavelen**5) * (1.0 / (np.exp(h * c / (wavelen * k *
 T)) - 1.0))
  return radiance
def main():
 T = float(sys.argv[1])
                               # Get temperature as a command line parameter
 wavelen = np.arange(samples)
  wavelen = wavelen_min + (wavelen_max - wavelen_min) * wavelen / float(samples)
  radiance = plank(T, wavelen)
 data = np.column_stack( (wavelen, radiance) )
 np.savetxt('spectrum.dat', data)
if __name__ == '__main__':
  main()
-:-- planck_fancy.py Top (1,0)
                                  (Python)
Using the python shell
```



Now let's plot the data!

```
-:--- show.py All (1,0) (Python)
Using the python shell
```

Now let's plot the data (and add axes, titles, all that good stuff)...

```
lineplotdemo.py
       Open
            Recent
                   Save
                                                                 Search
                                                                         Preferences
                              lineplotdemo.py
         show.py
## To run at the command line, in the
## directory containing the data file,
## type:
## python lineplotdemo.py
## Import necessary plotting libraries.
import numpy as np
import matplotlib.pyplot as plt
## Read and unpack data from file
## into arrays x and y.
x,y = np.loadtxt('spectrum.dat',unpack=True)
## Set up figure object.
fig = plt.figure()
## Set up axes object. Anything preceded
## by ax. will be plotted on these axes.
## This becomes more important when you
## wish to plot multiple subplots on the
## same figure.
ax=fig.add_subplot(111)
## Plot the data arrays. Use plot for line
## plots, scatter for point plots.
ax.plot(x,y)
ax.set_title('Planck Spectrum')
ax.set_xlabel('wavelength')
ax.set_ylabel('Intensity')
## Display the figure. This SHOULD ONLY
## be called once at the end of your
## plotting script.
plt.show()
-:--- lineplotdemo.py All (18,0)
```

Homework for tomorrow:

Take the mcmaster* file again, and plot the positions of the globular clusters in galactic coordinates. Prettiest plot wins a prize. Put it in a keynote slide to display to the class!



Write a python script that finds the density profile of the globular cluster model in the file king.dat in Kelly's home/bootcamp/2014 directory.

Plot a two panel figure with the xy projection of the model on the left, and the density profile on the right.

Hardcopy plus source code is due by the end of class.