# Computational Physics/ Structured Programming Lecture IV: Graphics & Visualization

Dr. Ian Kaniu

Department of Physics University of Nairobi

May 2024

Graphs

- 2 Scatter Plots
- Oensity Plots

Graphs

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- Oensity Plots

Graphs

- 2 Scatter Plots
- Oensity Plots

Graphs

- 2 Scatter Plots
- 3 Density Plots

## Graphics & Visualization

- So far we have created programs that print out words and numbers, but often we will also want our programs to produce graphics, meaning pictures of some sort.
- There are two main types of computer graphics used in physics.
  - 1) Most common of scientific visualizations, the graph: a depiction of numerical data displayed on calibrated axes.
  - 2) Scientific diagrams and animations: depictions of the arrangement or motion of the parts of a physical system, which can be useful in understanding the structure or behavior of the system.

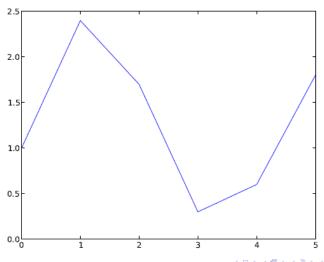
- A number of Python packages include features for making graphs.
- One such package which is powerful, easy-to-use, and popular is pylab.
- Pylab contains features for generating graphs of many different types. We will concentrate of three types that are especially useful in physics: ordinary line graphs, scatter plots, and density (or heat) plots.

• To create an ordinary graph in Python we use the functions plot and show from the pylab package.

```
from pylab import plot,show
y = [ 1.0, 2.4, 1.7, 0.3, 0.6, 1.8 ]
plot(y)
show()
```

After importing the two functions from pylab, we create the list
of values to be plotted, create a graph of those values with
plot(y), then display that graph on the screen with show().

If we run the program above, it produces a new window on the screen with a graph in it like this:

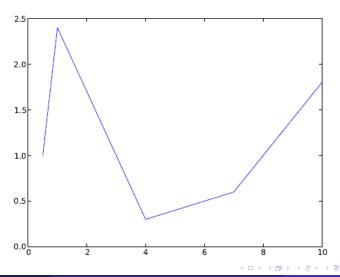


• When we want to specify both the x- and y-coordinates for the points in the graph, we use a plot statement with two list arguments, thus:

```
from pylab import plot,show
x = [ 0.5, 1.0, 2.0, 4.0, 7.0, 10.0 ]
y = [ 1.0, 2.4, 1.7, 0.3, 0.6, 1.8 ]
plot(x,y)
show()
```

- The two lists must have the same number of entries, as here. If they do not, you'll get an error message and no graph.
- Once the graph is displayed on the screen you can do other things with it i.e. zoom in on portions of the graph, move your view around the graph, or save the graph as an image file on your computer.

If we run the program above, it produces a new window on the screen with a graph in it like this:

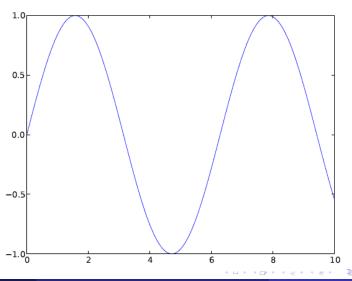


- Let us apply the plot and show functions to the creation of a slightly more interesting graph, a graph of the function sinx from x = 0 to x = 10.
- To do this we first create an array of the x values, then we take the sines of those values to get the y-coordinates of the points:

```
from pylab import plot,show
from numpy import linspace,sin
x = linspace(0,10,100)
y = sin(x)
plot(x,y)
show()
```

- The linspace function from numpy (see Handout I, section 2.5!) is used to generate the array of x-values.
- The sin function from numpy, which is a special version of sine that works with arrays—it just takes the sine of every element in the array (Alternatively we could have used the ordinary sin function from the math package).

• If we run the program above, it produces a new window on the screen with a graph in it like this:



- Notice that our plot consists of a finite set of points a hundred of them in this case - and the computer draws straight lines joining these points.
- So the end result is not actually curved; it's a set of straight-line segments.
- This is a useful and widely used trick for making curves in computer graphics: choose a set of points spaced close enough together that when joined with straight lines the result looks like a curve even though it really isn't.

 Suppose we have some experimental data in a computer file values.txt, stored in two columns, like this:

```
0 12121.71
1 12136.44
2 12226.73
3 12221.93
4 12194.13
5 12283.85
6 12331.6
7 12309.25
```

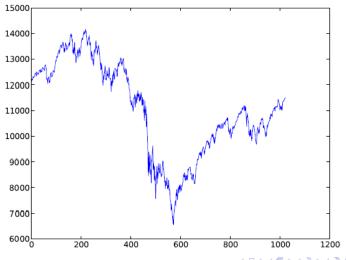
We can make a graph of these data as follows:

```
from numpy import loadtxt
from pylab import plot,show

data = loadtxt("values.txt",float)
x = data[:,0]
y = data[:,1]
plot(x,y)
show()
```

- The loadtxt function from numpy (see Handout I, section 2.4.3) is used to read the values in the file and put them in an array.
- The Python's array slicing facilities is used to extract the first and second columns of the array and put them in separate arrays x and y for plotting.

If we run the program above, it produces a new window on the screen with a graph in it like this:



 You can also vary the style in which the computer draws the curve on the graph. To do this a third argument is added to the plot function:

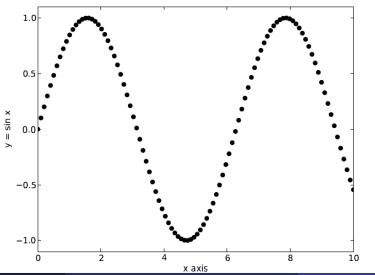
- The first letter of the string tells the computer what color to draw the curve with.
- Allowed letters are r, g, b, c, m, y, k, and w, for red, green, blue, cyan, magenta, yellow, black, and white, respectively.
- The remainder of the string says what style to use for the line.
- There are many options, i.e "-" for a solid line (like the ones we've seen so far), "--" for a dashed line, "o" to mark points with a circle, and "s" to mark points with a square.



 For example, the modification below plots our sine wave as a set of black circular points:

```
plot(x,y,"ko")
ylim(-1.1,1.1)
xlabel("x axis")
ylabel("y = sin x")
show()
```

If we run the program above, it produces a new window on the screen with a graph in it like this:



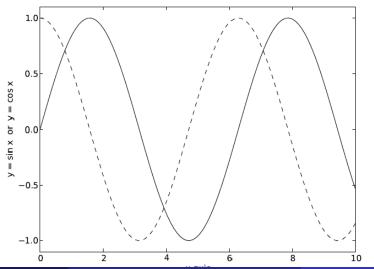
 We will often need to plot more than one curve or set of points on the same graph. This can be achieved by using the plot function repeatedly.

```
from pylab import plot,ylim,xlabel,ylabel,show
from numpy import linspace,sin,cos

x = linspace(0,10,100)
y1 = sin(x)
y2 = cos(x)
plot(x,y1,"k-")
plot(x,y2,"k-")
ylim(-1.1,1.1)
xlabel("x axis")
ylabel("y = sin x or y = cos x")
show()
```

 The program plots both the sine function and the cosine function on the same graph, one as a solid curve, the other as a dashed curve.

If we run the program above, it produces a new window on the screen with a graph in it like this:



## Class Activity 5: 20 Mins

## **Group Work (Plotting Experimental Data)**

You have been given a file called sunspots.txt, which contains the observed number of sunspots on the Sun for each month since January 1749. The file contains two columns of numbers, the first being the month and the second being the sunspot number.

- a) Write a program that reads in the data and makes a graph of sunspots as a function of time.
- b) Modify your program to display only the first 1000 data points on the graph.
- c) Modify your program further to calculate and plot the running average of the data, defined by  $Y_k$

$$Y_k = \frac{1}{2r} \sum_{m=-r}^r y_{k+m}$$

where r = 5 in this case (and the  $y_k$  are the sunspot numbers). Have the program plot both the original data and the running average on the same graph, again over the range covered by the first 1000 data points.

- In an ordinary graph, such as those of the previous section, there is one independent variable, usually placed on the horizontal axis, and one dependent variable, on the vertical axis.
- The graph is a visual representation of the variation of the dependent variable as a function of the independent one.
- In other cases, however, we measure or calculate two dependent variables.
- A classic example in physics is the temperature and brightness also called the magnitude of stars.
- Typically we might measure temperature and magnitude for each star in a given set and we would like some way to visualize how the two quantities are related.

 A standard approach is to use ascatter plot, a graph in which the two quantities are placed along the axes and we make a dot on the plot for each pair of measurements, i.e., for each star in this case

- Pylab provides the function scatter, which is designed specifically for making scatter plots.
- It works in a similar fashion to the plot function: you give it two lists or arrays one containing the x-coordinates of the points and the other containing the y-coordinates, and it creates the corresponding scatter plot: scatter(x,y)
- You do not have to give a third argument telling scatter to plot the data as dots - all scatter plots use dots automatically but to display it you need to use the function show.

 Suppose, for example, that we have the temperatures and magnitudes of a set of stars in a file called stars.txt on our computer, like this:

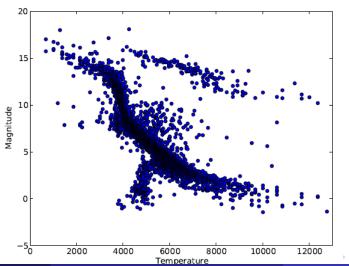
```
4849.4 5.97
5337.8 5.54
4576.1 7.72
4792.4 7.18
5141.7 5.92
6202.5 4.13
```

- The first column is the temperature and the second is the magnitude.
- Here's a Python program to make a scatter plot of these data:

```
# Plot of the Hertzsprung-Russell diagram
from pylab import scatter,xlabel,ylabel,xlim,ylim,show
from numpy import loadtxt

data = loadtxt("stars.txt",float)
x = data[:,0]
y = data[:,1]
scatter(x,y)
xlabel("Temperature")
ylabel("Magnitude")
xlim(0,13000)
ylim(-5,20)
show()
```

If we run the program above, it produces a new window on the screen with a scatter plot in it like this:



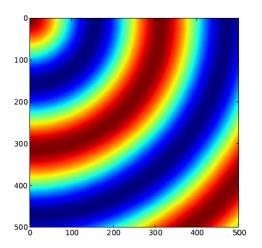
- xlabel and ylabel were used to label the temperature and magnitude axes, and xlim and ylim to set the ranges of the axes.
- You can also change the size and style of the dots and many other things, as well as use scatter two or more times in succession to plot two or more sets of data on the same graph, etc.
- See the on-line manual at matplotlib.org for more details.

- Two-dimensional data are harder to visualize on a computer screen than the one-dimensional lists of values that go into an ordinary graph.
- One tool that is helpful in many cases is the density plot, a two-dimensional plot where color or brightness is used to indicate data values.
- In Python density plots are produced by the function imshow from pylab.

• Here's the program that produces a density plot:

```
from pylab import imshow,show
from numpy import loadtxt
data = loadtxt("circular.txt",float)
imshow(data)
show()
```

If we run the program above, it produces a new window on the screen with a density plot in it like this:



• The file circular.txt contains a simple array of values, like this:

```
0.0050 0.0233 0.0515 0.0795 0.1075 ...
0.0233 0.0516 0.0798 0.1078 0.1358 ...
0.0515 0.0798 0.1080 0.1360 0.1639 ...
0.0795 0.1078 0.1360 0.1640 0.1918 ...
0.1075 0.1358 0.1639 0.1918 0.2195 ...
```

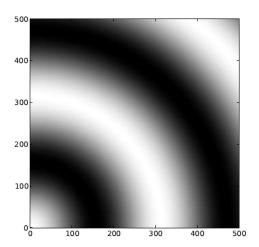
- The program reads the values in the file and puts them in the two-dimensional array data using the loadtxt function, then creates the density plot with the imshow function and displays it with show.
- The computer automatically adjusts the color-scale so that the picture uses the full range of available shades.

- Density plots with this particular choice of colors from blue to red (or similar) are sometimes called heat maps, because the same color scheme is often used to denote temperature, with blue being the coldest temperature and red being the hottest.
- To change to gray-scale, for instance, you use the function gray, which takes no arguments:

```
from pylab import imshow,gray,show
from numpy import loadtxt

data = loadtxt("circular.txt",float)
imshow(data,origin="lower")
gray()
show()
```

The gray color scheme, which runs from black for the lowest values to white for the highest:



- Pylab provides many other color schemes, which you may find useful occasionally.
- A complete list, with illustrations, is given in the on-line documentation at matplotlib.org, but here are a few that might find use in physics:

jet	The default heat-map color scheme
gray	Gray-scale running from black to white
hot	An alternative heat map that goes black-red-yellow-white
spectral	A spectrum with 7 clearly defined colors, plus black and white
bone	An alternative gray-scale with a hint of blue
hsv	A rainbow scheme that starts and ends with red

#### 3D Graphics

**Reading Material**: "3D Graphics" - Handout will be uploaded together with the lecture

#### Part Four

 VPython, the nickname for Python plus the Visual package, is particularly useful for creating 3-D solids, 2-D plots, and animations.

Here's a code that produces two plots using vpython:

```
from vpython import * # Import Vpython
graph1=graph(align='left', width=400, height=400,
  background=color.white,foreground=color.black)
  Plot1=gcurve(color=color.red) # gcurve method
for x in arange(0,8.1,0.1): # x range
     Plot1.plot(pos=(x,5*cos(2*x)*exp(-0.4*x)))
 graph2=graph(align='right', width=400, height=400,
  background=color.white,foreground=color.black,
title='2-D Plot', xtitle='x', ytitle='f(x)')
  Plot2=gdots(color=color.black) # Dots
for x in arange(-5,5,0.1):
Plot2.plot(pos=(x,cos(x))) # plot dots
```

- Notice that the plotting technique with VPython is to create first a plot object, and then to add the points one at a time to the object.
- In contrast, matplotlib.org creates a vector of points and plots the entire vector in one fell swoop.
- Here's a link where you can find more vpython examples with animations: https://www.glowscript. org/#/user/GlowScriptDemos/folder/Examples/

# Please go through the tutorial videos presented in the following links

- https:
  //www.youtube.com/watch?v=vEMCiugDnKI&list=
  PLdCdV2GBGyXOnMaPS1Bg07IOU\_00ApuMo
- https:
  //www.youtube.com/watch?v=aPnZ4TIUn08&list=
  PLdCdV2GBGyXOnMaPS1Bg07IOU\_00ApuMo&index=2
- https:
  //www.youtube.com/watch?v=IbRo-\_sTUvQ&list=
  PLdCdV2GBGyXOnMaPS1Bg07IOU\_00ApuMo&index=3
- https:
  //www.youtube.com/watch?v=Y2snMyfgDuo&list=
  PLdCdV2GBGyXOnMaPS1Bg07IOU\_00ApuMo&index=4

## Summary of Lecture

#### **Questions and Class Discussions:**

- Graphs
- Scatter Plots
- Density Plots
- 3D Plots & vPython

#### These lecture notes will be uploaded in SOMAS

 Post your questions, comments and suggestions in the Forum Discussion Page in SOMAS.