

Introduction To Python Programming

III: Numpy, SciPy and Matplotlib

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The Libraries

- NumPy: Base library. Fast arrays and matrices, FFT, linear algebra, dates, random numbers, etc
- SciPy: Uses NumPy. Large collection of high-quality implementations of numerical algorithms: stats, ODE solvers, integration, optimization, signal processing...
- Matplotlib: Data visualization for Numpy and Scipy. 2-D and 3-D plots, animations, and interactive visualizations.

SymPy:	Symbolic computation
Pandas:	Time series analysis
scikit-learn:	Machine learning

A quick example

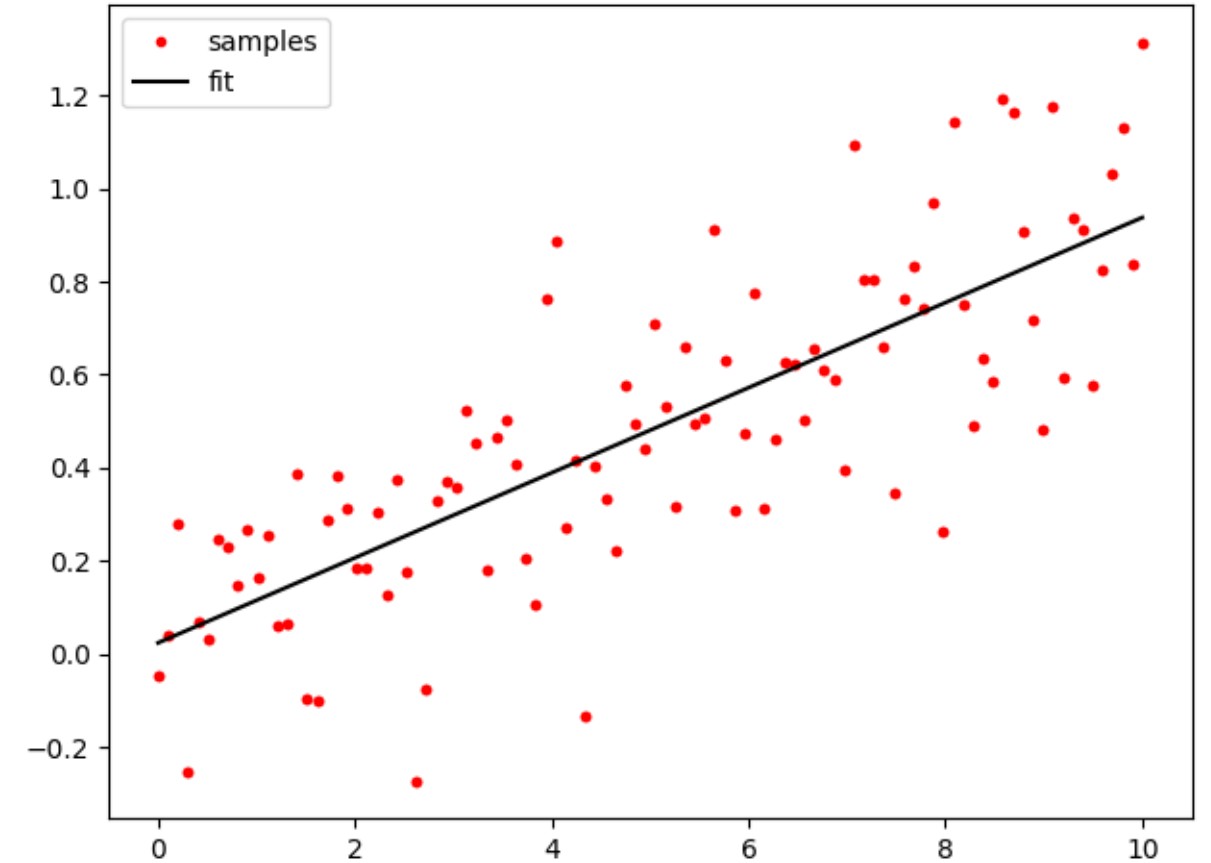
```
import numpy as np
import matplotlib.pyplot as plt
from scipy import stats

x = np.linspace(0,10,100)
y = np.random.normal(x/10, 0.2)

p = stats.linregress(x,y)

plt.figure()
plt.plot(x,y, 'r.', label='samples')
plt.plot(x, x*p.slope+p.intercept,
        "k", label='fit')

plt.legend()
```



A quick example

```
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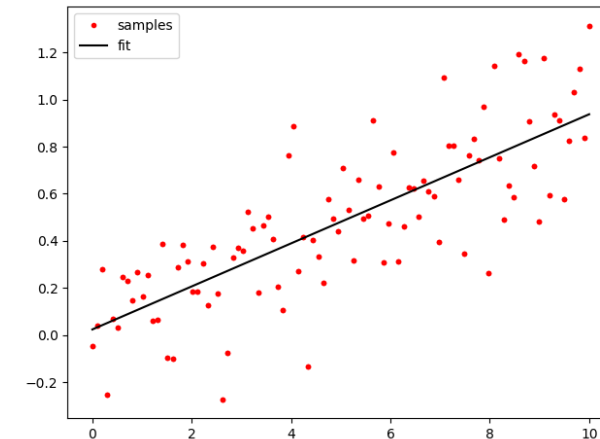
plt.legend()
```

Import numpy, name it “np”

Import matplotlib.pyplot, name it “plt”

- We often rename modules for ease of use; “np” and “plt” are standard names.

Import the “stats” module from scipy



A quick example

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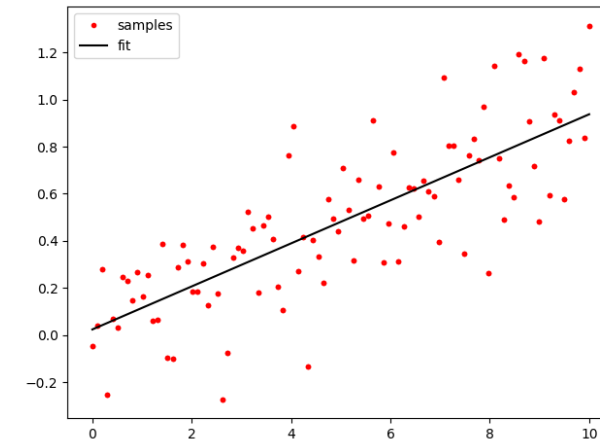
plt.figure()
plt.plot(x,y, 'r.', label='samples')
plt.plot(x, x*p.slope+p.intercept,
         "k", label='fit')

plt.legend()
```

Get an *array* of 100 numbers from 0 to 10

Get an array of normal random numbers, with the mean at the “x/10” values, and variance 0.2

Linear regression to x and y



A quick example

```
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import matplotlib.pyplot as plt
from scipy import stats

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p = stats.linregress(x,y)

plt.figure()
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plt.plot(x, x*p.slope+p.intercept,
        "k", label='fit')

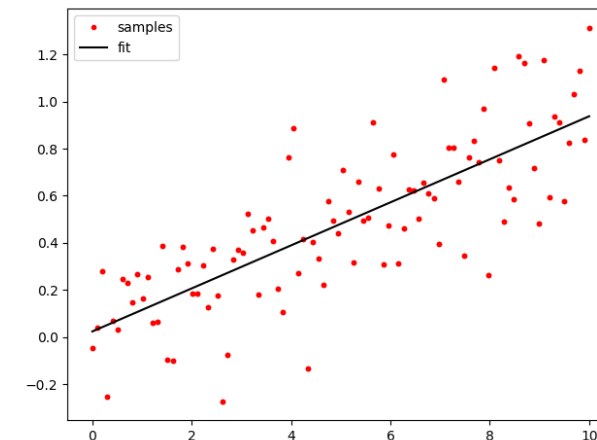
plt.legend()
```

Create a figure to plot into

Plot the (x,y) points in red ("r") dots (".") with label "samples"

Plot a line with slope p.slope and and zero at p.intercept, name it "fit"

Add the legend to the plot



Arrays

The base Numpy data structure is the *array*:

```
# 1-d vector from a list
a = np.array( [1,2,3] )

# 2-d array from a list of lists
b = np.array([[1,2], [3,4]])
b
array([[1, 2],
       [3, 4]])

a = np.outer([1,2,3], [1,2,3])
b = np.zeros_like(a)+2

a*b                                # elementwise
array([[ 2,  4,  6],
       [ 4,  8, 12],
       [ 6, 12, 18]])
```

Arrays

- Stored as a single memory area
- All elements are the same type (there's an exception...)
- very efficient, compatible with C and Fortran arrays
- array is an *iterable*:
 - can convert to/from lists
 - can loop over the elements

Arrays

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# 2-d array from a list of lists
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array([[1, 2],
       [3, 4]])

a = np.outer([1,2,3], [1,2,3])
b = np.zeros_like(a)+2

a*b                                # elementwise

array([[ 2,  4,  6],
       [ 4,  8, 12],
       [ 6, 12, 18]])
```

Why Arrays?

- The storage is very efficient
- They are fast
- You can quickly operate on all elements in an array
- You can treat them as matrices
 - same as Matlab matrices
 - many high-performance operations available

Array layout and shapes

```
b = np.arange(1,10).reshape(3,3)

b.size          # nr elements
9
b.ndim          # dimensions
2
b.shape         # size in each dimension
(3, 3)

b.reshape(2,4) # ERROR 2×4 != 9

x=np.ravel(b)
array([ 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

in memory:

b.data=

1	2	3	4	5	6	7	8	9
---	---	---	---	---	---	---	---	---

b =

1	2	3
4	5	6
7	8	9

Diagram illustrating the memory layout of a 3x3 array `b`. The array is shown as a 3x3 grid of elements (1 to 9). The memory layout is shown as a 1x9 row of elements (1 to 9). The array is labeled `b` and its shape is `b.shape`. The first dimension is `b.shape[0]` (3) and the second dimension is `b.shape[1]` (3). Red arrows indicate the row-major traversal order: 1, 2, 3, 4, 5, 6, 7, 8, 9.

`b[1,2]`

Diagram illustrating the indexing of a 3x3 array `b`. The array is shown as a 3x3 grid of elements (1 to 9). The first dimension is `b.shape[0]` (3) and the second dimension is `b.shape[1]` (3). Red arrows indicate the row-major traversal order: 1, 2, 3, 4, 5, 6, 7, 8, 9. The element at row 1, column 2 is `b[1,2]`. The labels `row` and `column` are shown below the array.

Array layout and shapes

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b = np.arange(1,10).reshape(3,3)

b.size          # nr elements
9
b.ndim          # dimensions
2
b.shape         # size in each dimension
(3, 3)

b.reshape(2,4) # ERROR 2×4 != 9

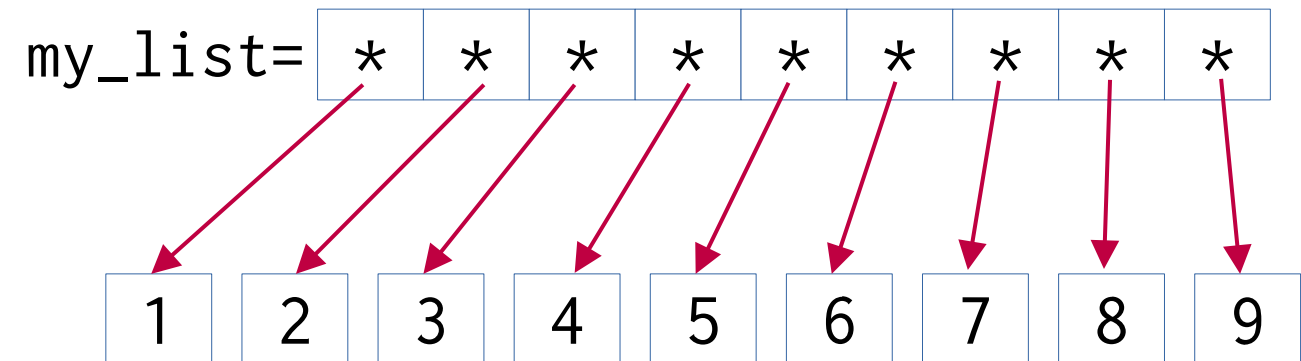
x=np.ravel(b)
array([ 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

in memory:

b.data=

1	2	3	4	5	6	7	8	9
---	---	---	---	---	---	---	---	---

compare with list:



Arrays and matrices

What about 'matrix'?

- Really the same as an array:
 - Same representation
 - mostly same methods
 - very cheap to convert with "asmatrix"
- But, multiplication is a matrix multiplication, not elementwise
- Also, can use MATLAB-like init

Most code uses array in practice

```
# MATLAB-like initialization
m = np.mat('1, 2; 3, 4')

m1 = np.asmatrix(a)
m2 = np.asmatrix(b)

m1*m2                # dot product

matrix([[12., 12., 12.],
        [24., 24., 24.],
        [36., 36., 36.]])

a.dot(b)             # also dot product

matrix([[12., 12., 12.],
        [24., 24., 24.],
        [36., 36., 36.]])
```

Create Arrays

```
# Create arrays:
a = np.array([[1,2,3], [2,3,4]])
b = np.ones_like(a) # same shape with '1'
c = np.zeros((2,3,4)) # 3d array of '0'

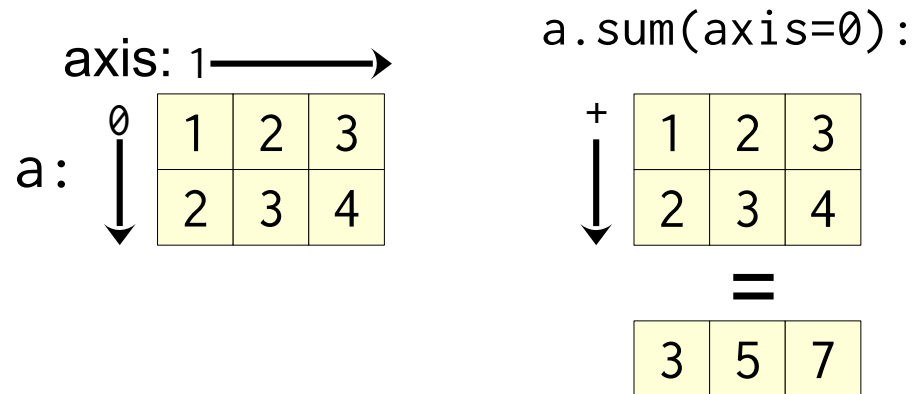
I = np.eye(3)          # identity
I
array([[1., 0., 0.],
       [0., 1., 0.],
       [0., 0., 1.]])

# you can specify the numerical type
ia = np.ones((2,2), dtype=np.complex64)
ia
array([[1.+0.j, 1.+0.j],
       [1.+0.j, 1.+0.j]], dtype=complex64)
```

- give a sequence of elements
- create pre-filled array of specific size
- Special array creation:
 - np.eye, np.diag, np.random ...
 - many python modules can convert to/from numpy arrays
- You can specify the data type

Array operations

- Many functions operate on the elements of an array
 - `max()`, `min()`, `+`, `cumsum()`, `sin()`, `conjugate()`, `round()`, `prod()` ...
- You can set an axis for operations:



- operations give you a new array

```
a = np.array([[1,2,3], [2,3,4]])

a.sum()
15
a.sum(axis=0) # operate along one axis
array([3, 5, 7])

np.log(a) # all can be called like this
array([[0.        , 0.69314, 1.09861],
       [0.69314, 1.09861, 1.38629]])

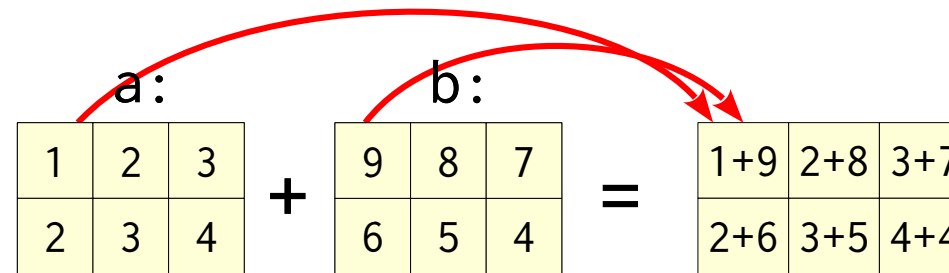
b = a.copy()
b.T # Transposition
array([[1, 2],
       [2, 3],
       [3, 4]])

b.dot(a.T) # dot product
array([[14, 20],
       [20, 29]])
```

Arrays: Broadcasting

Operations are elementwise:

```
a = np.array([[1,2,3], [2,3,4]])  
b = np.array([[9,8,7], [6,5,4]])  
a+b  
array([[10, 10, 10],  
       [ 8,  8,  8]])
```



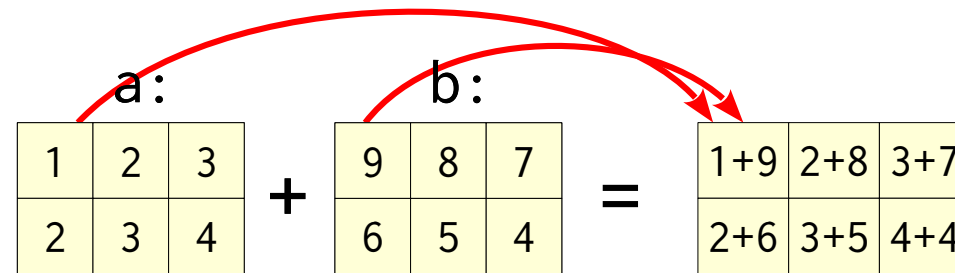
The operation happens on each pair of elements in the two arrays

→ the arrays have to be the same size

Arrays: Broadcasting

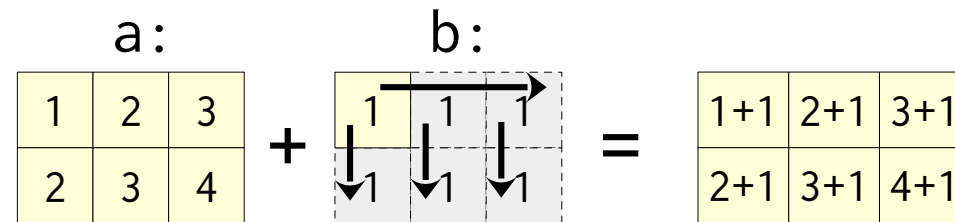
Operations are elementwise:

```
a = np.array([[1,2,3], [2,3,4]])  
b = np.array([[9,8,7], [6,5,4]])  
a+b  
array([[10, 10, 10],  
       [ 8,  8,  8]])
```



What about this?

```
a = np.array([[1,2,3], [2,3,4]])  
b = 1  
a+b  
array([[2, 3, 4],  
       [3, 4, 5]])
```

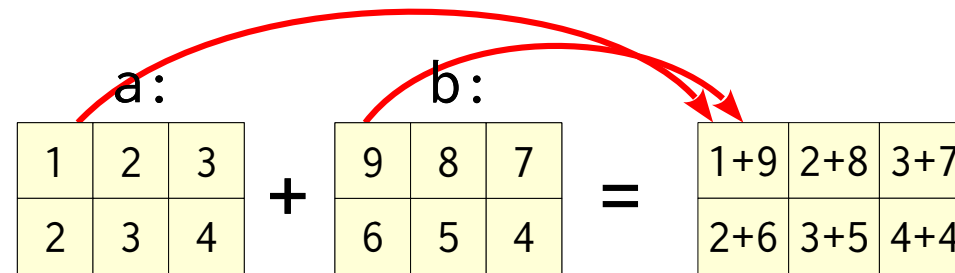


Broadcasting fills in the single value into an array of the right size, so that the elementwise operation can proceed.

Arrays: Broadcasting

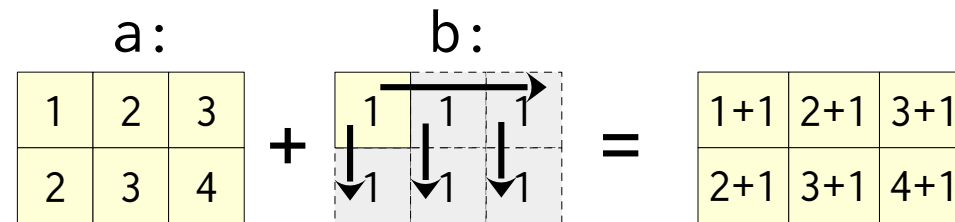
Operations are elementwise:

```
a = np.array([[1,2,3], [2,3,4]])  
b = np.array([[9,8,7], [6,5,4]])  
a+b  
array([[10, 10, 10],  
       [ 8,  8,  8]])
```



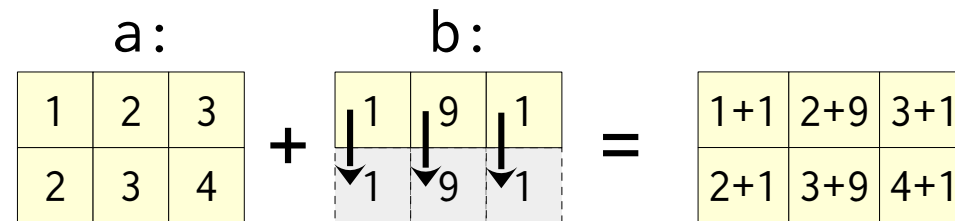
What about this?

```
a = np.array([[1,2,3], [2,3,4]])  
b = 1  
a+b  
array([[2, 3, 4],  
       [3, 4, 5]])
```



And what about this?

```
a = np.array([[1,2,3], [2,3,4]])  
b = np.array([1,9,1])  
a+b  
array([[2, 11, 4],  
       [3, 12, 5]])
```



Find information

How to find out more

- `lookfor("text")` looks for the text anywhere in numpy
- `function?` gives you documentation and examples for a function in ipython
- `np.info(function)` is another way to get the same documentation
- Documentation:

<https://numpy.org/doc/1.18/>

```
np.lookfor("median")
Search results for 'median'
-----
numpy.median
    Compute the median along the specified axis.
numpy.nanmedian
    Compute the median along the specified axis, ...
...
numpy.quantile
    Compute the q-th quantile of the data along ...

np.median?
Signature: np.median(a, axis=None, out=None, overw ...
Docstring:
Compute the median along the specified axis.

Returns the median of the array elements.

Parameters
-----
a : array_like
    Input array or object that can be converted to ...
```

Visualisation Example: sin()

calculate and plot $\sin(x)$ and $\sin(x/2)$

- `np.linspace(s,e,n)` generates `n` points between start and end
- `plt.ion()` for interactive mode
- `figure()`: use `figsize (8,6 inches)` and `dpi (128)` to set the plot size.
 - 8,6 inches × 128 dpi = 1024×768 pixels
- `plt.plot()`:
 - arrays with `x` and `y` coordinates
 - set line width, color, markers etc.

```
import numpy as np
import matplotlib.pyplot as plt

x = np.linspace(0, 2*np.pi, 50) # 50 pts in [0..2*pi)
s1 = np.sin(x)
s2 = np.sin(x/2)

plt.ion() # interactive mode
plt.figure(figsize=(8,6), dpi=128)

plt.plot(x,s1,"b-", lw=2) # lw=linewidth
plt.plot(x,s2, ".-", c="red") # red dot-dash line
```

Visualisation Example: sin()

calculate and plot $\sin(x)$ and $\sin(x/2)$

- `plt.ion()` for interactive mode
- `figure()`: use `figsize` (8,6 inches) and `dpi` (128) to set the plot size.
- `plt.plot()`:
 - arrays with x and y coordinates
 - set line width, color, markers etc.
- `plt.xlabel()` sets label for x axis
- `plt.tight_layout()` removes extra space
- `plt.savefig()` saves figure to file
- `plt.show()` shows plot if not interactive

```
import numpy as np
import matplotlib.pyplot as plt

x = np.linspace(0, 2*np.pi, 50) # 50 pts in [0..2*pi)
s1 = np.sin(x)
s2 = np.sin(x/2)

plt.ion() # interactive mode
plt.figure(figsize=(8,6), dpi=128)

plt.plot(x,s1,"b-", lw=2) # lw=linewidth
plt.plot(x,s2, ".-", c="red") # red dot-dash line

plt.xlabel("angle")
plt.ylabel("awesomness")

plt.tight_layout() # remove extra space
plt.savefig("awesome.png") # svg, pdf, other formats
plt.show() # show on screen
```

Array indexing

arrays are indexed:

`a[row, col, depth ...]`

axis: 1 $\longrightarrow$

a: $\downarrow$ 0

1	2	3	4
5	6	7	8

- begins with 0
- negative counts from the end
- can also use the “list of lists” way to index:

`a[row][col][depth]`

- a list as index:
 - list of elements to pick in order
 - creates a copy of the data

```
a = np.array([[1,2,3,4], [5,6,7,8]])

a[0, 0]      # row, column from 0,0
1
a[0][0]      # same thing
1
a[1]         # single index=entire row
array([ 5,  6,  7,  8])

a[0, -1]     # -1 = last element
4

a[0, [0,3,2]] # select elements
array([1, 4, 3])
```

Array slices

Slice: a “cut-out” piece of an array

- Slices are *not* copies
- specify slices with ‘:’
[first elem:last elem+1]

a:

1	2	3	4	5	6	7	8
---	---	---	---	---	---	---	---

b:

1	2	3	4
5	6	7	8

a[:3] = [0:3] # start of array:3
a[2:] = [2:8] # 2:end of array
a[:] = [0:8] # all elements

```
a = np.arange(1,9)
array([1, 2, 3, 4, 5, 6, 7, 8])

a[2:4]            # start:end+1
array([3, 4])

a[2]              # 1 value
3

a[2:3]            # 1-element array
array([3])        # 2:3 is DIFFERENT from 2!

b = np.array([[1,2,3,4], [5,6,7,8]])

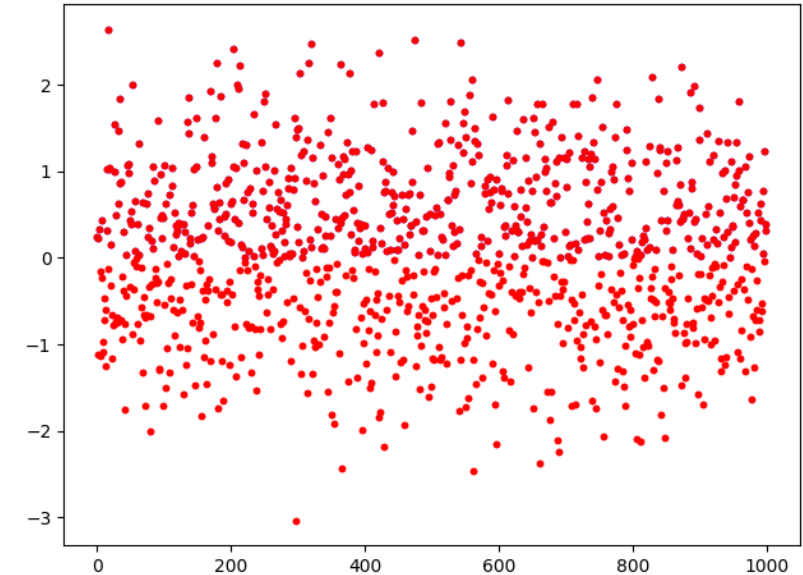
b[:, 1]
array([2, 6])    # 1d array

b[:, 1:2]
array([[2],
       [6]])    # 2d array
```

Array indexing

Using boolean indexes:

```
a = np.random.normal(size=1000)
x = np.arange(a.size)
plt.plot(x, a, "r.")
```



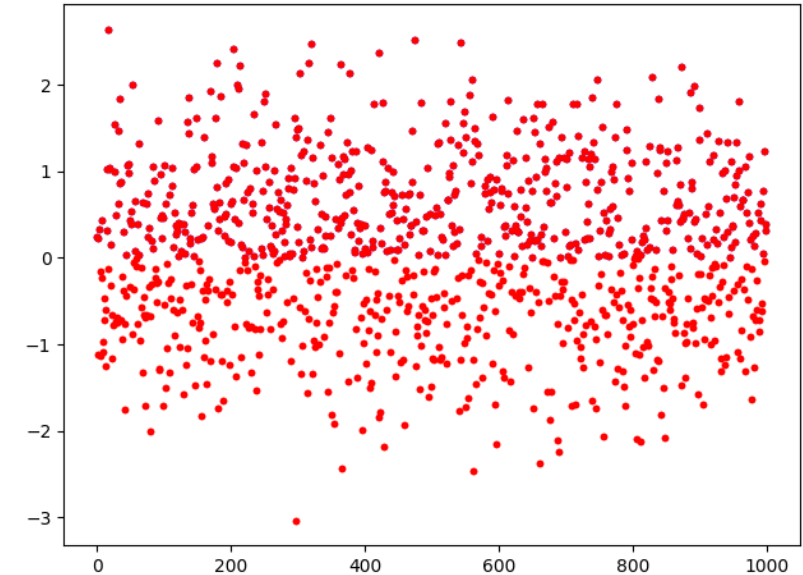
Use a test to pick what values to use

For example: pick and plot values of 'a' that are greater than or equal to 0

Array indexing

Using boolean indexes:

```
a = np.random.normal(size=1000)
x = np.arange(len(a))
plt.plot(x, a, "r.")
apos = (a>=0)
array([ True, False,  True,  True,  True ...
```



- Compare each element in a.
- Return a new array with "True" or "False" in each position.

Array indexing

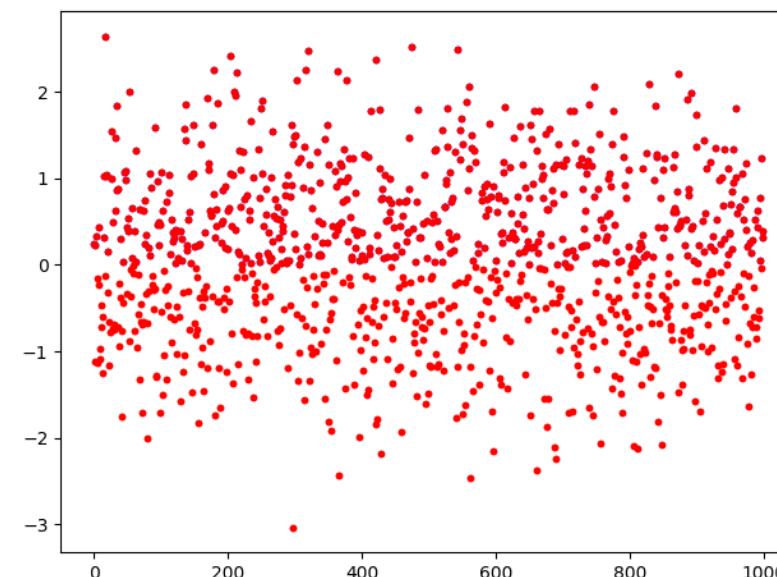
Using boolean indexes:

```
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x = np.arange(len(a))
plt.plot(x, a, "r.")
apos = (a>=0)
array([ True, False,  True,  True,  True ...
```

```
## Small example ##
```

```
b = np.random.randint(-5,5,8)
array([-2,  4, -3,  1, -1,  2,  1,  4])
```

```
bpos = (b>=0)
b[bpos]
array([4, 1, 2, 1, 4])
```

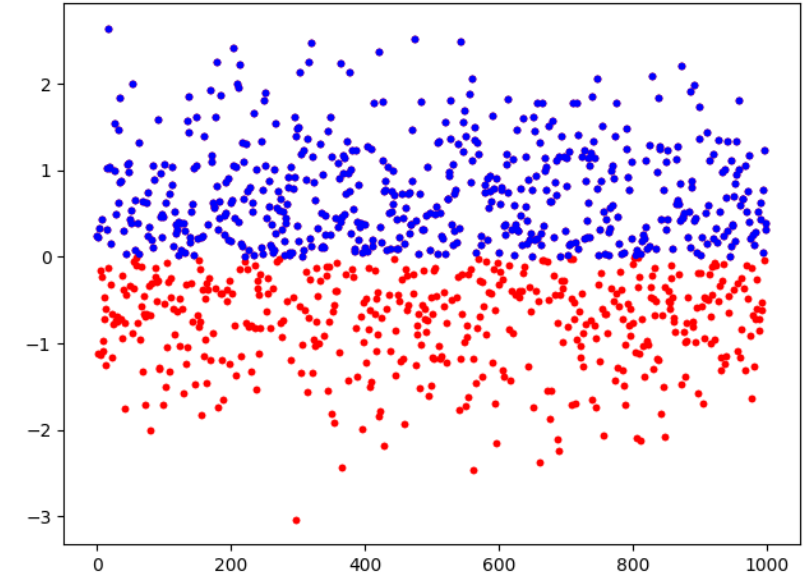


- Compare each element in a.
- Return a new array with "True" or "False" in each position.
- A boolean index returns the elements that are "True"

Array indexing

Using boolean indexes:

```
a = np.random.normal(size=1000)
x = np.arange(len(a))
plt.plot(x, a, "r.")
apos = (a>=0)
array([ True, False,  True,  True,  True ...
plt.plot(x[apos], a[apos], "b.")
```



- Compare each element in a.
- Return a new array with "True" or "False" in each position.
- A boolean index returns the elements that are "True"

Array indexing

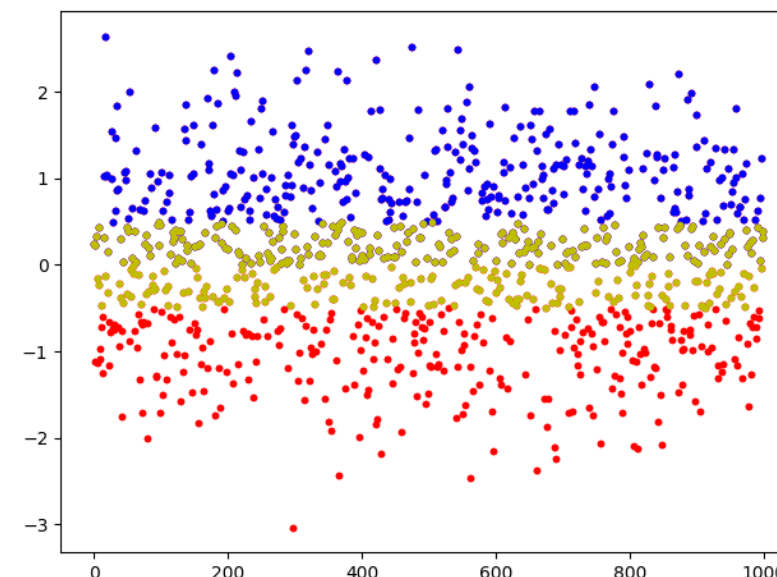
Using boolean indexes:

```
a = np.random.normal(size=1000)
x = np.arange(len(a))
plt.plot(x, a, "r.")
apos = (a>=0)
array([ True, False,  True,  True,  True ...

plt.plot(x[apos], a[apos], "b.")

# combine boolean arrays with '&' and '|'
amid = (a<0.5) & (a>-0.5)

plt.plot(x[amid], a[amid], "y.")
```



- Compare each element in a.
- Return a new array with "True" or "False" in each position.
- A boolean index returns the elements that are "True"

Speed

in ipython we can use "timeit" to measure execution time

Elementwise multiplication
using Numpy:

```
a = np.ones(10000)*10  
b = np.ones_like(a)*2  
c = np.zeros_like(a)
```

```
%timeit c=a*b
```

3.35 μ s $\pm$ 3.42 ns per loop (mean $\pm$ std.
dev. of 7 runs, 100000 loops each)

Elementwise multiplication
using our own loop:

```
%%timeit  
for i in range(len(a)):  
    c[i] = a[i]*b[i]
```

4.76 ms $\pm$ 22.4 μ s per loop (mean $\pm$
std. dev. of 7 runs, 100 loops each)

Speed

in Ipython we can use "timeit" to measure execution time

Elementwise multiplication
using Numpy:

```
a = np.ones(10000)*10  
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%timeit c=a*b
```

3.35 μ s $\pm$ 3.42 ns per loop (mean $\pm$ std.
dev. of 7 runs, 100000 loops each)

Elementwise multiplication
using our own loop:

```
%timeit  
for i in range(len(a)):  
    c[i] = a[i]*b[i]
```

4.76 ms $\pm$ 22.4 μ s per loop (mean $\pm$
std. dev. of 7 runs, 100 loops each)

Our loop is **1400 $\times$** slower than Numpy!

Speed

in ipython we can use "timeit" to measure execution time

Elementwise multiplication
using Numpy:

```
a = np.ones(1000000)  
b = np.ones(1000000)  
c = np.zeros(1000000)  
  
%timeit c=a*b  
  
3.35 µs ± 3.4 µs  
dev. of 7 runs
```

Elementwise multiplication
using our own loop:

```
a)):  
    c[a*b] = 1  
  
loop (mean ±  
00 loops each)
```

Always spend as much time as possible
inside Numpy and Scipy functions!

→ If there is a function to do something,
use it. Do not write it yourself.

loading, saving and displaying

reading, writing images

- `plt.imread()` reads an image
 - 3-d array (height, width, color)
- `plt.imshow()` shows array as image
- `plt.pcolormesh()` shows as mesh
- `plt.imsave()` saves a plot as image



```
# one way to read an image: use matplotlib
img = plt.imread("shisa.jpg")

# height, width, color channels
img.shape
(200, 250, 3)

plt.imshow(i)      # show as image

# coordinates of the corners
x = np.arange(0, 251, 1)
y = np.arange(201, 0, -1)

# show mesh, green channel, grayscale
plt.pcolormesh(x,y,img[:, :, 1], cmap="gray")

plt.imsave("shisa_green.jpg")
```

Reading data

`genfromtxt()`

- Reads a text-format data file
- skips comments, empty lines
you can set what counts as comment etc.
- set the data type with "dtype="
- returns a Numpy array

```
# example data file. genfromtxt() skips '#'  
# comments and empty lines automatically
```

```
1  10 11      # whitespace is default separator  
2   9 12
```

```
np.genfromtxt("data1.txt")  
array([[ 1., 10., 11.],  
       [ 2.,  9., 12.]]) # float type
```

```
np.genfromtxt("data1.txt", dtype="int")  
array([[ 1, 10, 11],  
       [ 2,  9, 12]]) # integer
```

Reading data

`genfromtxt()`

- Reads a text-format data file
- skips comments, empty lines
you can set what counts as comment etc.
- set the data type with "dtype="
- returns a Numpy array
- "delimiter=" sets separator char
- "skip_header=" and "skip_footer=" skips lines at the beginning and end

But what if you have columns with different data types?

```
count, val1, val2      # header line
1,      10,      11      # comma-separated data
2,      9,       12
```

```
np.genfromtxt("data2.txt")
array([[nan, nan, nan],
       [nan, nan, 11.],
       [nan, nan, 12.]]) # that didn't go well

np.genfromtxt("data2.txt", skip_header=1,
              delimiter=",", dtype="int")
array([[ 1, 10, 11],
       [ 2,  9, 12]]) # much better!
```

Reading data

`genfromtxt()` can only do so much

- Matlab .mat: "scipy.io" module
- CSV: Python "CSV" module
Pandas `read_csv()`
- HDF5: Python "h5py" module
- NetCDF: "scipy.io" module
- IDL .sav: "scipy.io" module
- DICOM: Python "pydicom" module
Python "dicom-numpy" module
- ...
- ...

Scipy

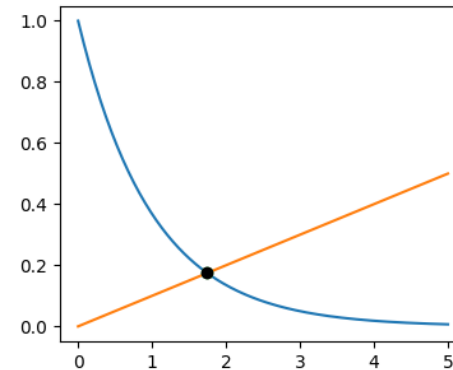
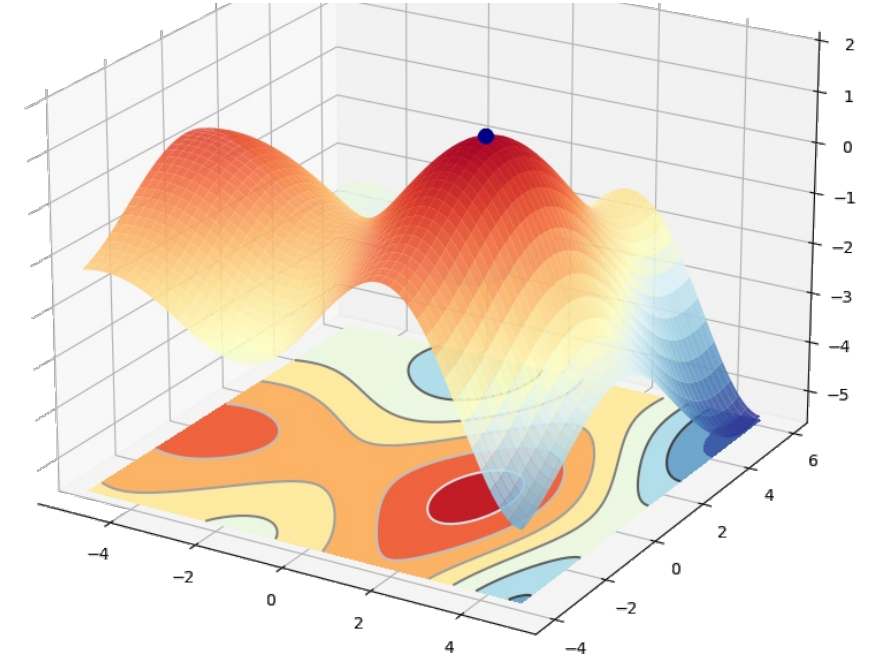
- special Defines special functions
- integrate Integration and ODE solvers
- **optimize** **find minima & roots, fit curves, LP**
- interpolate data interpolation
- fftpack FFT, discrete cosine, sine
- signal convolution, filters, spectral analysis
- linalg superset of Numpy linear algebra
- sparse sparse matrices, graphs, algebra
- spatial Voronoi, KDtree, convex hulls
- stats Statistics
- ndimage image analysis
- IO Matlab, IDL, Netcdf, other file formats

Optimization

Find the maximum/minimum/zero value of a function

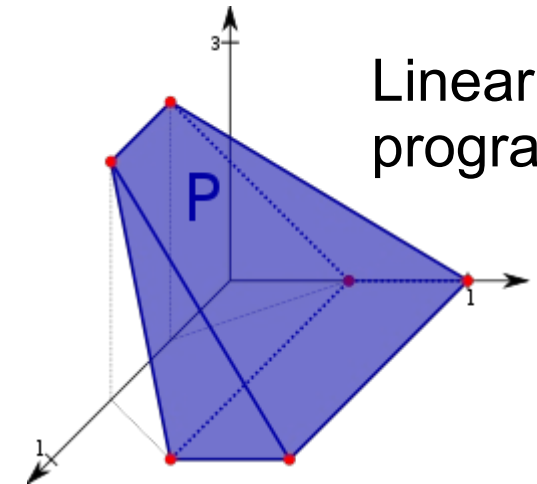
- You can optimize (almost) anything you can describe with a function:
 - simulation parameters
 - fit your model to data
 - optimize a process
 - assignment problems
 - numerical methods (finding roots, extrema, etc.)

Parameter optimization



numerical methods

Linear programming



Scipy Exercise:

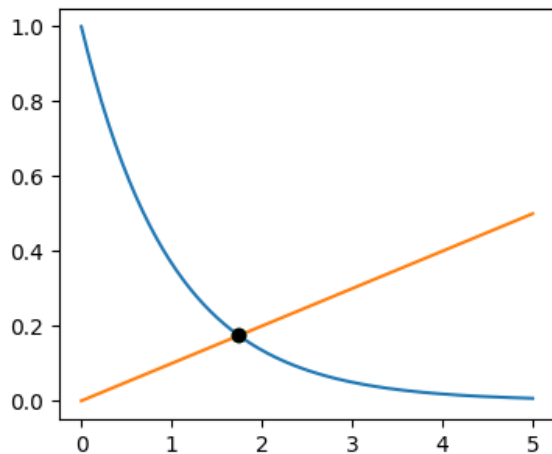
Find the root (zero) of a function

We want the intersection of two functions $\exp(-x)$ and $x/10$

1: define a python function $f(x)$

$$\exp(-x) - x/10$$

2: find where $f(x) == 0$



```
from scipy import optimize as opt

# define our function f(x)

# use a function in opt to find root of f(x)
```

see: <https://docs.scipy.org/doc/scipy/reference>

Also:

```
help(opt)
help(opt.<method>)
```

Scipy Exercise:

Find the root (zero) of a function

We want the intersection of two functions $\exp(-x)$ and $x/10$

1: define a python function $f(x)$

$$\exp(-x) - x/10$$

2: find where $f(x)=0$

Look for root finding methods:

ex: brentq, newton, fsolve

```
from scipy import optimize as opt

# define our function f(x)

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```

see: <https://docs.scipy.org/doc/scipy/reference>

Also:

`help(opt)`

`help(opt.<method>)`

Scipy Exercise:

Find the root (zero) of a function

We want the intersection of two functions $\exp(-x)$ and $x/10$

1: define a python function $f(x)$

$$\exp(-x) - x/10$$

2: find where $f(x)=0$

Look for root finding methods:

ex: brentq, newton, fsolve

```
from scipy import optimize as opt

# define our function f(x)
def f(x):
    return np.exp(-x) - x/10

# use a function in opt to find root of f(x)
opt.brentq(f, 0, 10)
1.7455280027406994
```

see: <https://docs.scipy.org/doc/scipy/reference>

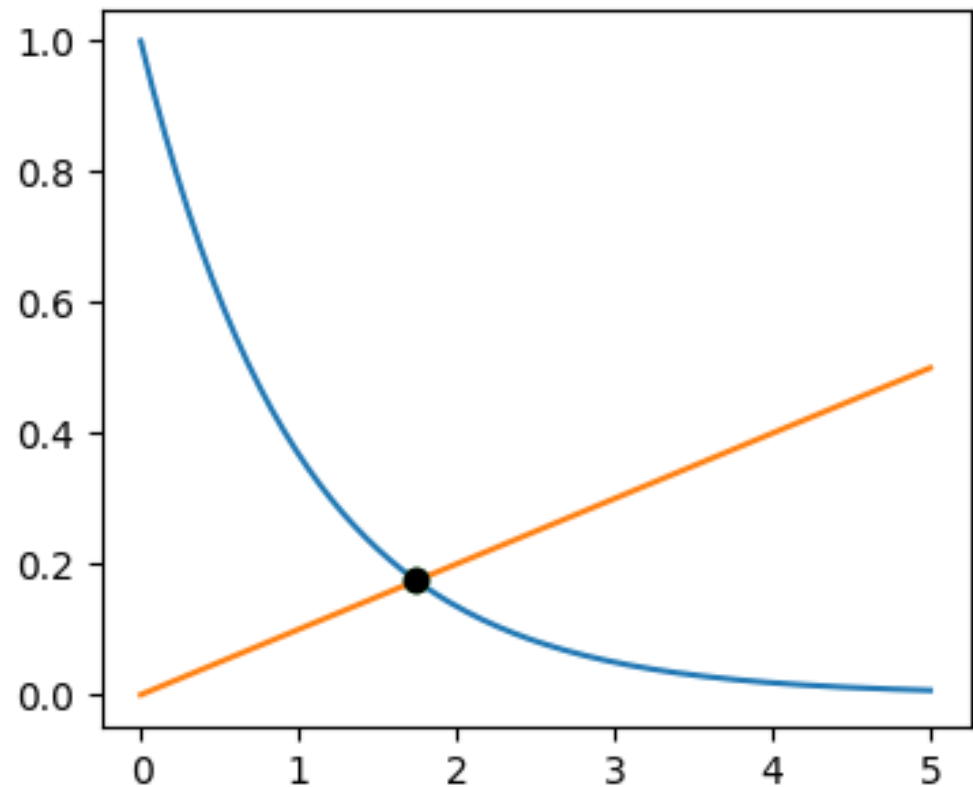
Also:

`help(opt)`

`help(opt.<method>)`

Scipy Exercise:

Find the root (zero) of a function



```
from scipy import optimize as opt

# define our function f(x)
def f(x):
    return np.exp(-x) - x/10

# use a function in opt to find root of f(x)
p = opt.brentq(f, 0, 10)

x=np.linspace(0,5,200)
plt.plot(x, np.exp(-x))
plt.plot(x, x/10)
plt.plot(p, p/10, "ko") # k=black, o=circle
```

Scipy Optimize: A few Thoughts

- `opt.minimize` is a general interface for finding function minima:

`opt.minimize(f, [x0, y0...])`
- *Many* HPC applications need parameter tuning. Tuning by hand is error prone, grid search is inefficient.
- The objective function `f` can be anything — including a separate cluster job

→ We can optimize our parameters automatically with python and a cluster

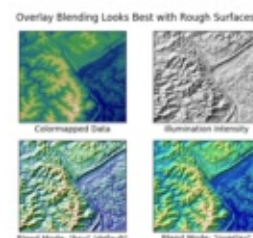
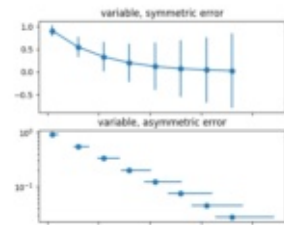
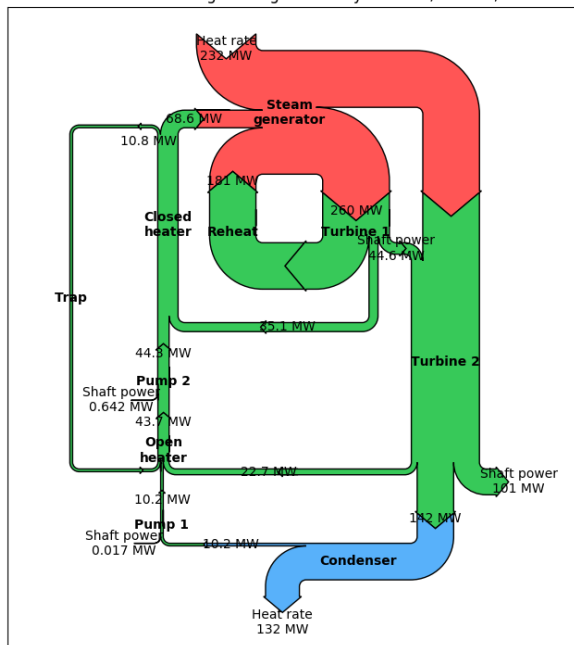
```
search_program.py

opt.minimize(f, ...)
...
f(params):
    create job.sh with params
    run ("sbatch job.sh")
    wait for job
    calc "goodness" from result
```

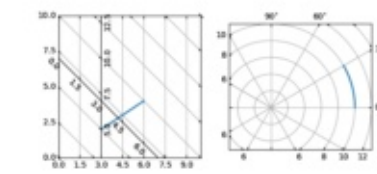
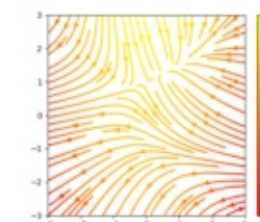
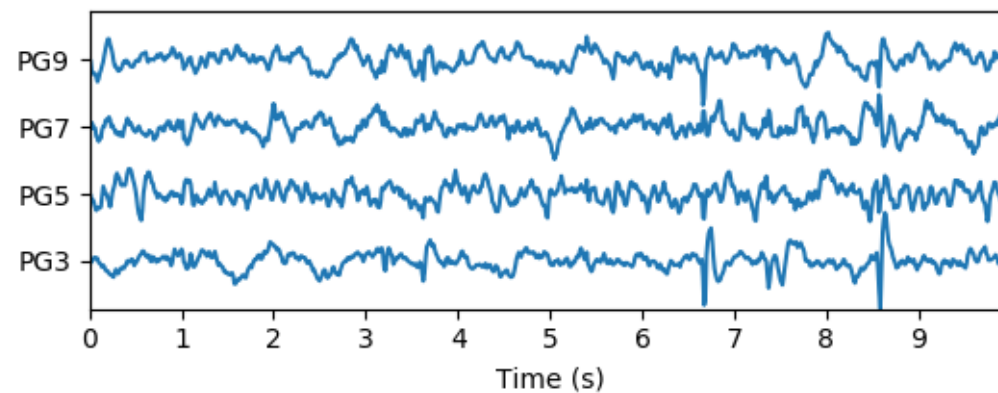
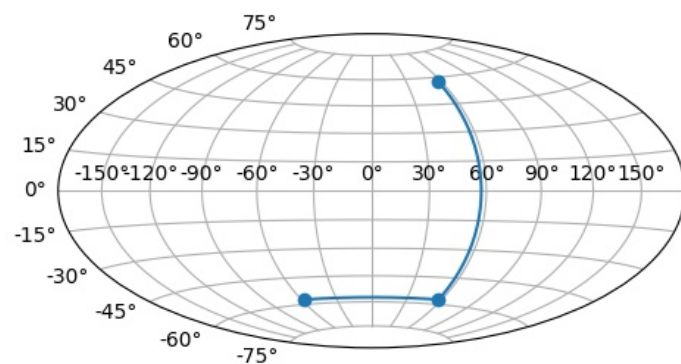
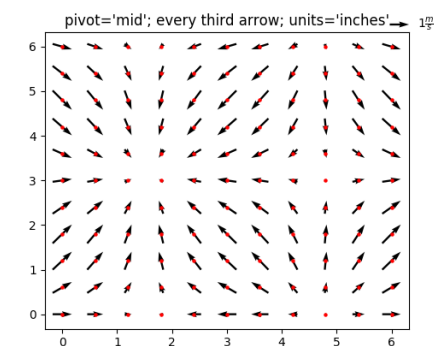
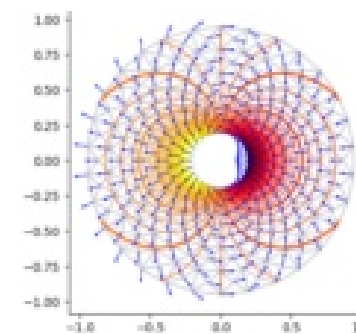
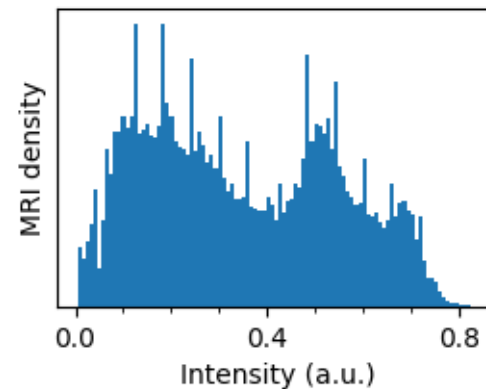
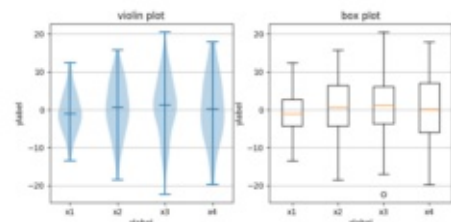
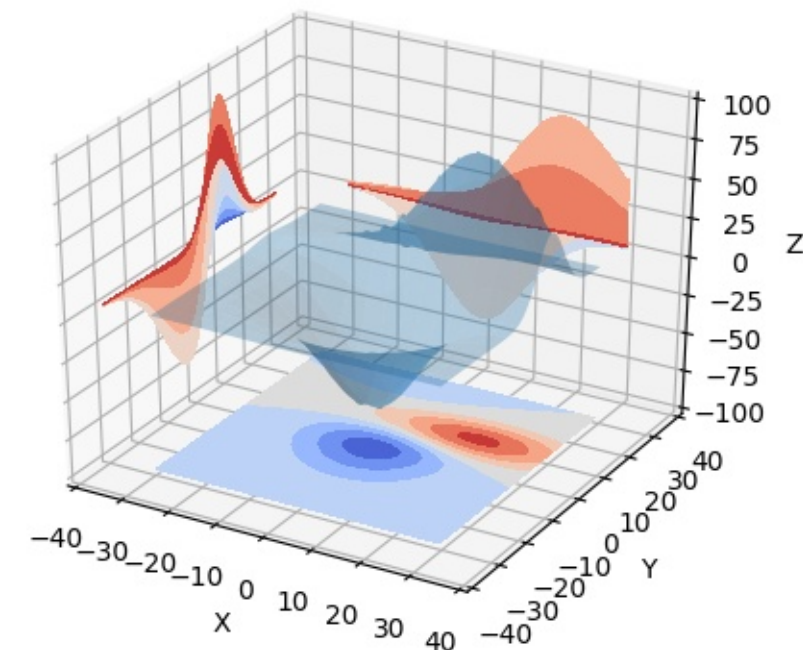
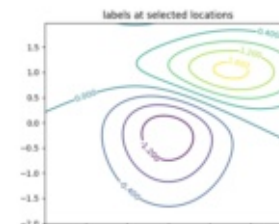
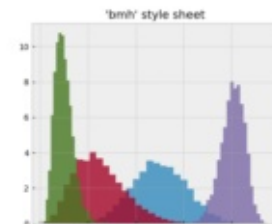
```
job.sh:

#/bin/bash
#SBATCH -t 0-1
...
```

Rankine Power Cycle: Example 8.6 from Moran and Shapiro
"Fundamentals of Engineering Thermodynamics ", 6th ed., 2008



Plots



Plots

```
plt.ion()           # interactive mode
#plt.ioff()         # stop interactive mode

plt.figure()        # create a new figure

plt.plot(x,y,...)   # plot 1-d data

# save figure to file
plt.savefig("myfig.png")

plt.show()          # if we didn't use ion()
```

- The Pyplot interface ("plt.") keeps track of figures and axes for you
- In "interactive mode" any change immediately updates the plot
 - good for interactive use
- non-interactive mode only draws the graph when you save or show the figure
 - great for programs, or when you plot lots of data

Plots

Creating the plot

```
plt.plot( ... )    # creates figure

# figure parameters:

figure(nr, figsize, dpi, facecolor)

figsize=(x, y)      # size in inches
dpi = n             # pixels per inch
facecolor = color   # background color

# colors
'r'                 # red
'red'               # red
'#ff0000'           # red
[1.0, 0.0, 0.0]     # red
```

- colors can be given in many ways:

'r'	single color letter
'red'	HTML name
'#ff0000'	Red-Green-Blue triplet in hex
[1,0,0]	RGB as [0...1] triplet

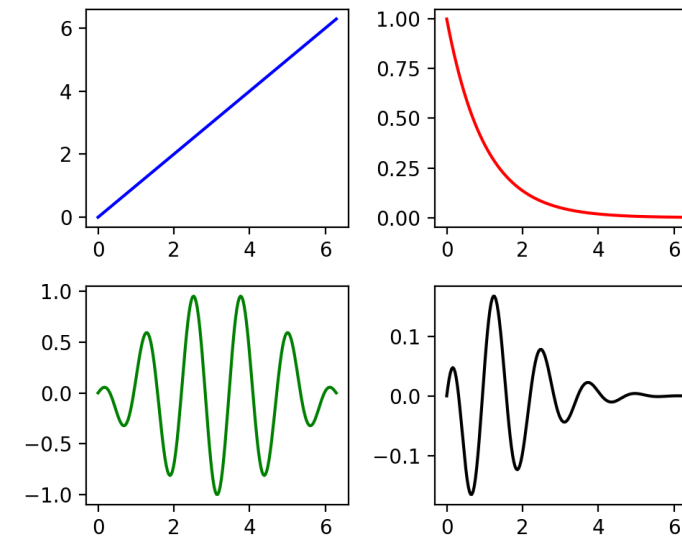
Plots

Subplots

- Make multiple plots in grids
- There are a number of ways to create subplots (including doing it "by hand")
- `plt.subplots()` is pretty flexible and not hard to use.

```
f, ax = plt.subplots(2,2)

ax[0,0].plot(x,y1, 'b') # plot each
ax[0,1].plot(x,y2, 'r') # quadrant
ax[1,0].plot(x,y3, 'g')
ax[1,1].plot(x,y4, 'k')
```



Plots

Subplots

- `sharex()` and `sharey()` makes the x or y-axis common across rows/cols or across all plots.

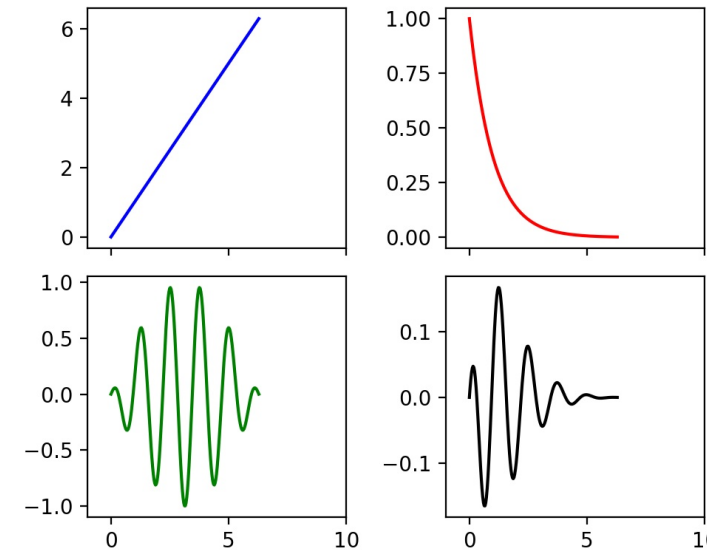
→ axis changes in one plot affects all connected plots

```
f, ax = plt.subplots(2,2, sharex=True)

ax[0,0].plot(x,y1, 'b') # plot each
ax[0,1].plot(x,y2, 'r') # quadrant
ax[1,0].plot(x,y3, 'g')
ax[1,1].plot(x,y4, 'k')

ax[1,1].clear()
plt.sca(ax[1,1]) # set current axes
plt.plot(x,y4, 'k') # can plot with
# pyplot

ax[1,1].set_xlim([-1,10]) # x axes are
# connected
```



Plot Styles

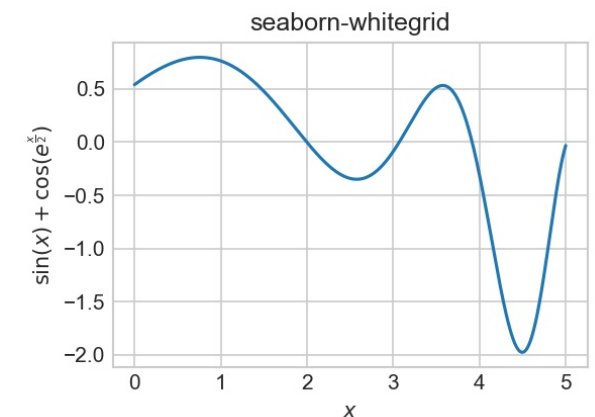
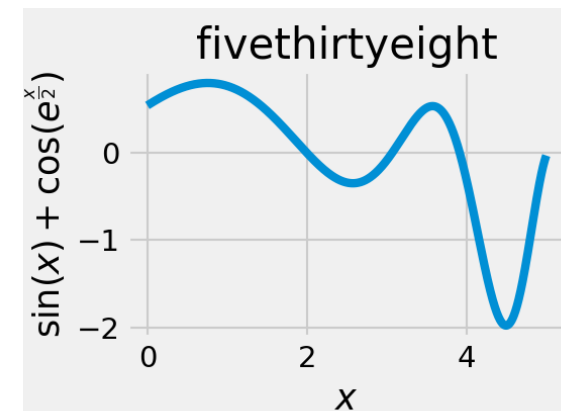
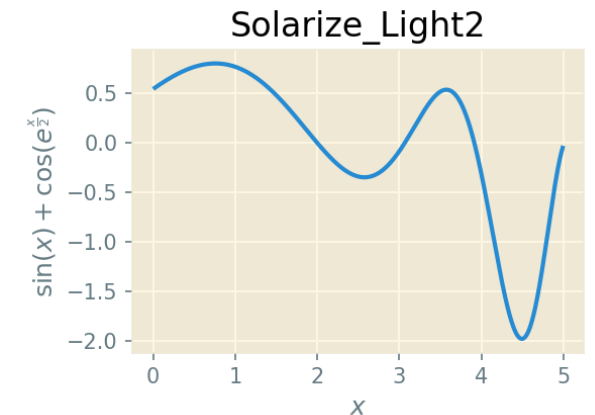
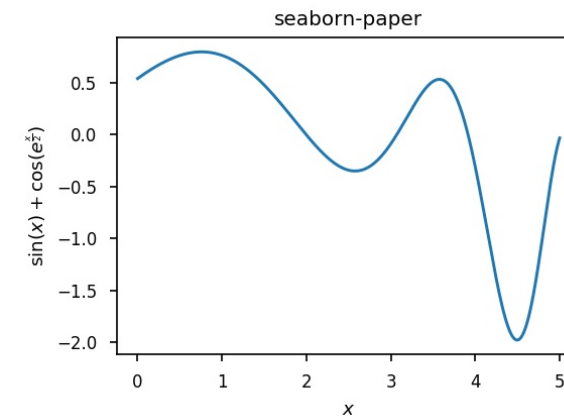
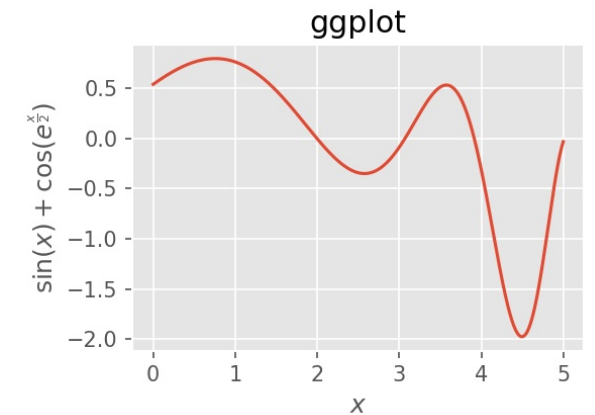
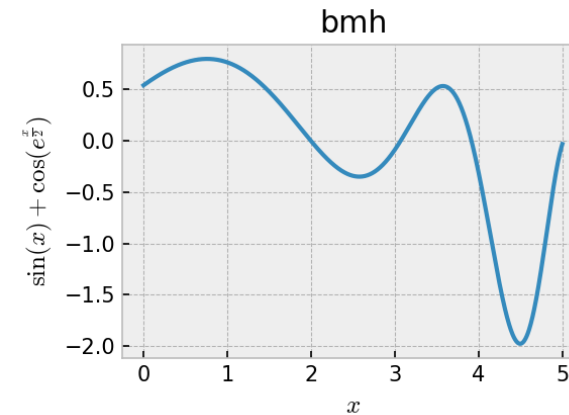
```
# styles
```

```
plt.style.use("ggplot")
```

```
print(plt.style.available)  
['seaborn-paper', 'classic', ...]
```

Styles set visual defaults

- Not all defaults are actively set, so settings from previous styles may linger.



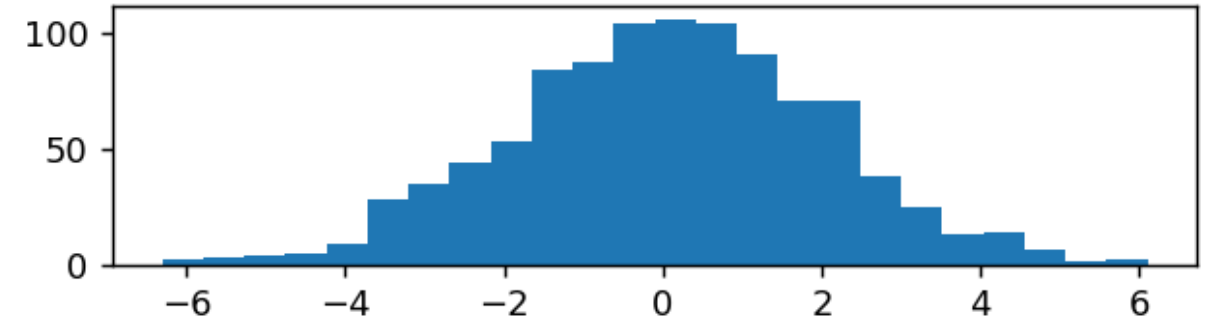
Histogram

```
p = np.random.normal(scale=2, size=1000)
plt.hist(p, bins='auto')
```

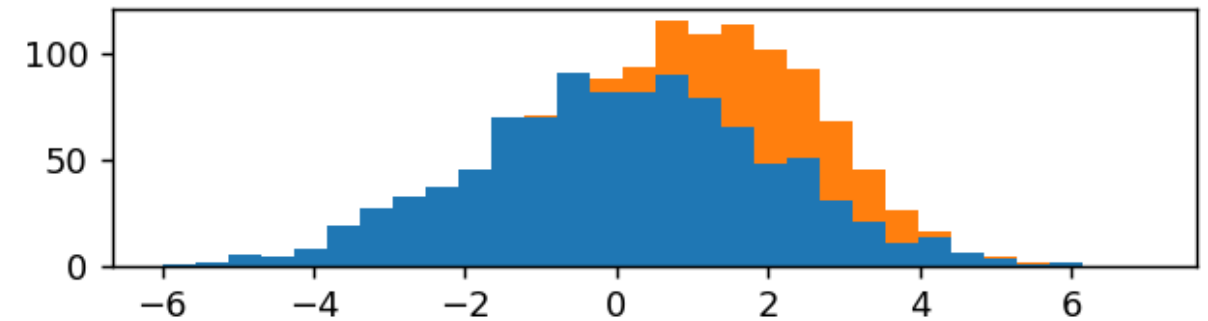
```
p2=np.random.normal(loc=2, size=300)
plt.hist((p,p2),
        histtype='barstacked', # stack bars
        bins=30,               # set bin number
        range=(-6,7))          # min and max
```

```
# can also give explicit bin positions
bins=[0.0, 1.0, ...
cumulative=True           # cumulative hist
density=True              # normalized scale
weights=[...]             # weigh points
```

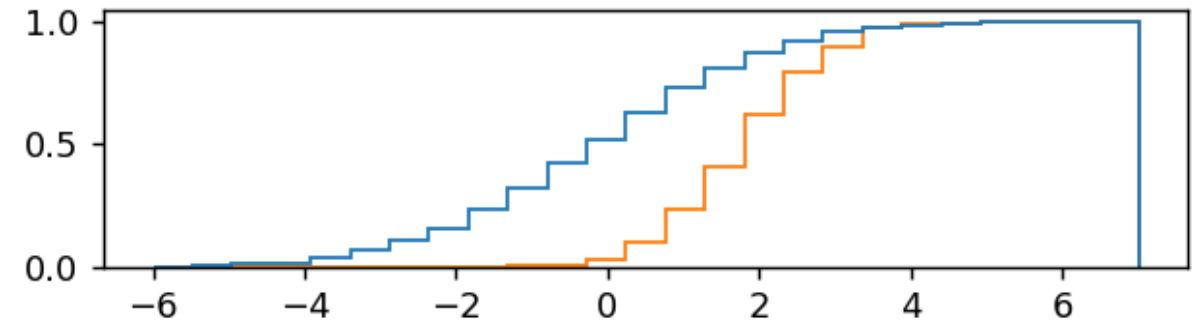
simple histogram



stacked histogram



cumulative histogram



Mesh and Contour Plots

```
# create coordinates
xc = np.arange(-5,5,0.1)
yc = np.arange(-5,5,0.1)
x,y = np.meshgrid(xc,yc)

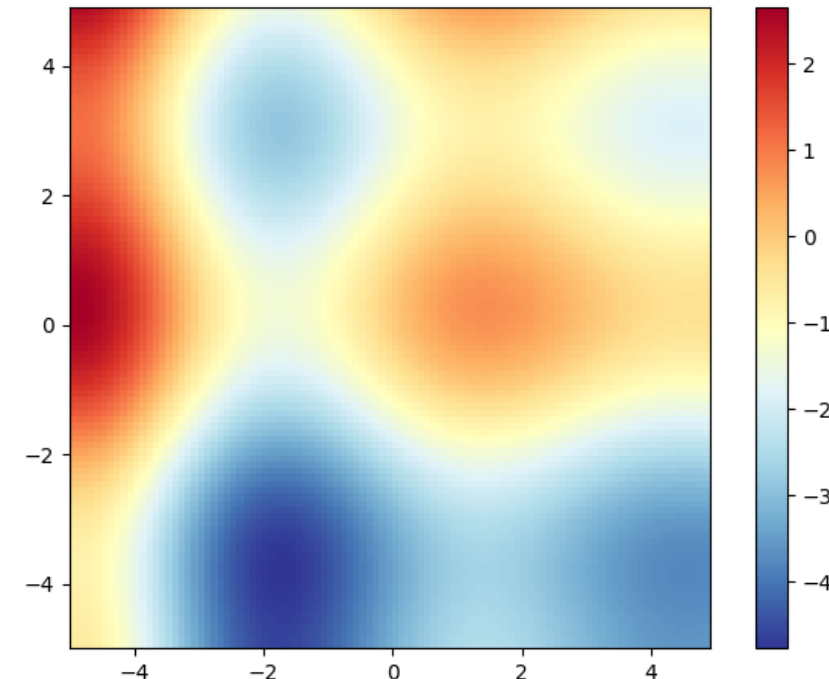
# create some 2-d data
z = np.sin(x)*np.exp(-x/5) +
    np.cos(y)-np.exp(-y/4)

f,ax = plt.subplots()
ax.set_aspect('equal')

p = ax.pcolormesh(x,y,z,      # coord, vals
                  cmap=plt.cm.RdYlBu_r)  # color map

f.colorbar(p)
```

- 2d data is an array of z-values.
- optionally give X and Y coordinates for each Z data points ("meshgrid")
- pcolormesh plots z values as colors:



Mesh and Contour Plots

```
# create coordinates
xc = np.arange(-5,5,0.1)
yc = np.arange(-5,5,0.1)
x,y = np.meshgrid(xc,yc)

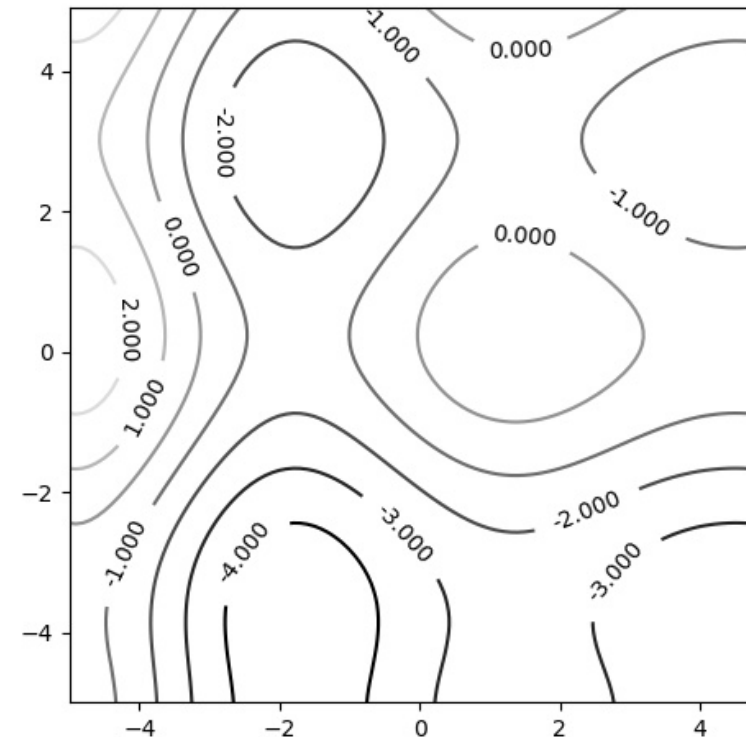
# create some 2-d data
z = np.sin(x)*np.exp(-x/5) +
    np.cos(y)-np.exp(-y/4)

f,ax = plt.subplots()
ax.set_aspect('equal')

c = ax.contour(x,y,z,      # coord, vals
              8,          # number of levels
              vmax=3,      # upper limit
              cmap=plt.cm.gray) # color map

c.clabel(colors="k")      # numerical labels
```

- 2d data is an array of z-values.
- optionally give X and Y coordinates for each Z data points ("meshgrid")
- contour plots



Mesh and Contour Plots

```
# create coordinates
xc = np.arange(-5,5,0.1)
yc = np.arange(-5,5,0.1)
x,y = np.meshgrid(xc,yc)

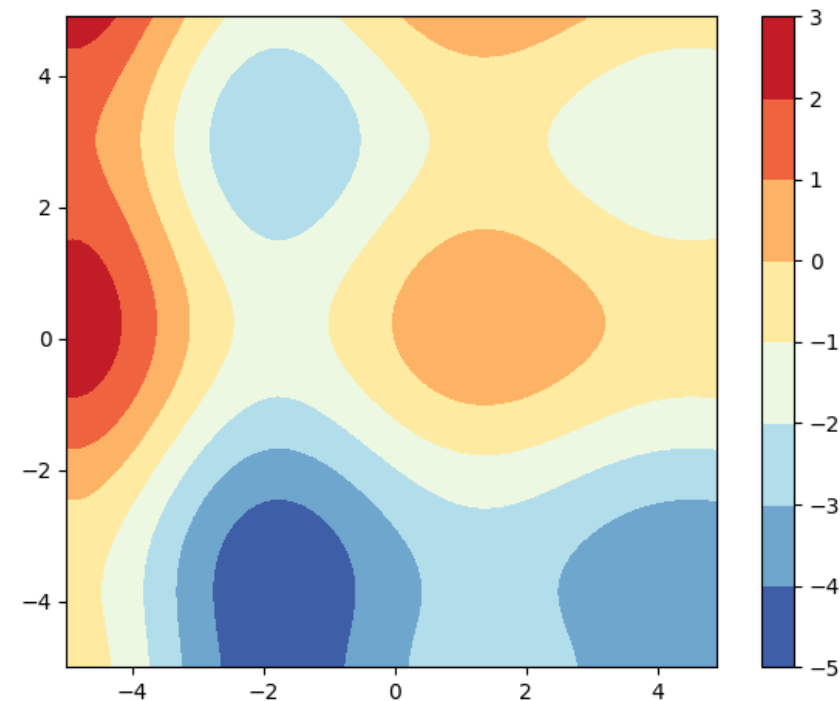
# create some 2-d data
z = np.sin(x)*np.exp(-x/5) +
    np.cos(y)-np.exp(-y/4)

f,ax = plt.subplots()
ax.set_aspect('equal')

s = ax.contourf(x,y,z, # coord, vals
               8,      # number of levels
               cmap=plt.cm.RdYlBu_r) # color map

f.colorbar(s)
```

- 2d data is an array of z-values.
- optionally give X and Y coordinates for each Z data points ("meshgrid")
- filled contour plots:



Mesh and Contour Plots

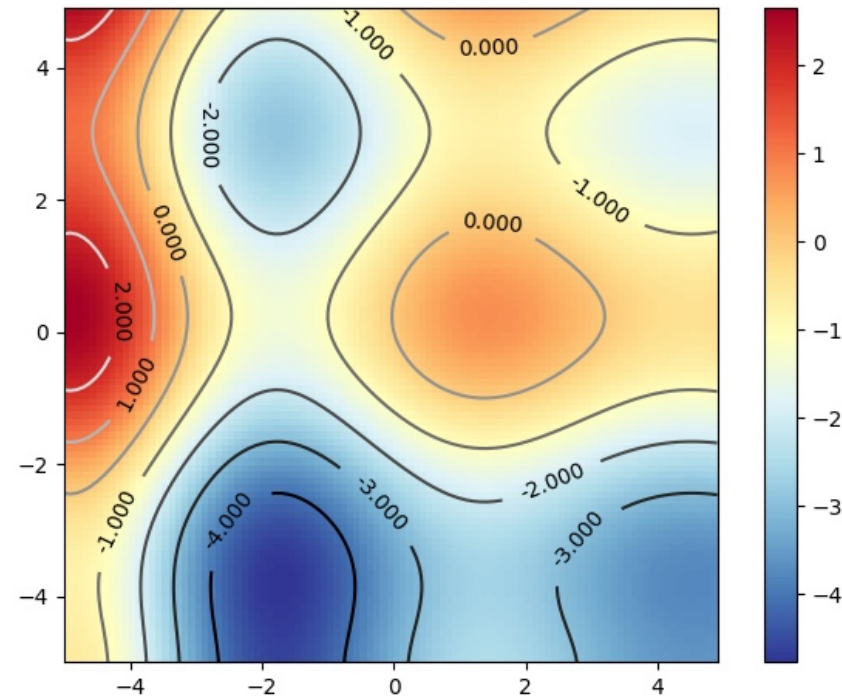
```
f, ax = plt.subplots()
ax.set_aspect('equal')

p = ax.pcolormesh(x, y, z, 8,
                  cmap=plt.cm.RdYlBu_r)

c = ax.contour(x, y, z, 8,
               vmax=3,
               cmap=plt.cm.gray)

f.colorbar(p)
c.clabel(colors="k")
```

- 2d data is an array of z-values.
- optionally give X and Y coordinates for each Z data points ("meshgrid")
- combine plots for best results:

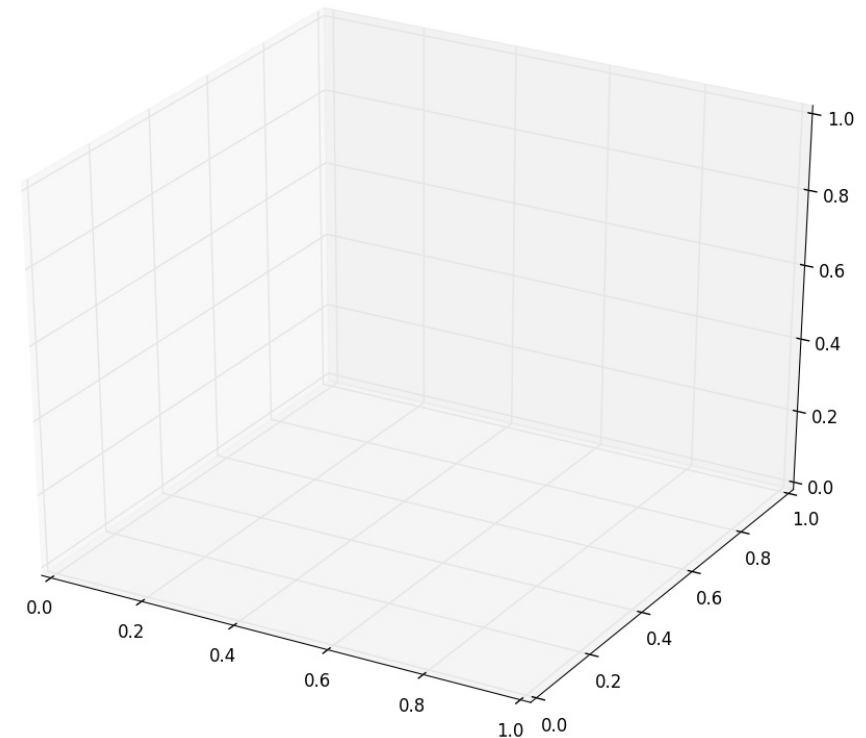


3D Plots

```
import matplotlib.pyplot as plt
import numpy as np
from mpl_toolkits.mplot3d import Axes3D

f = plt.figure(figsize=(10,8))
f.add_subplot(111, projection='3d')
ax = f.gca()
```

- import Axes3d from mpl_toolkits
 - adds 3d-related methods to matplotlib
 - still need to use older "add_subplot" instead of "subplots()"



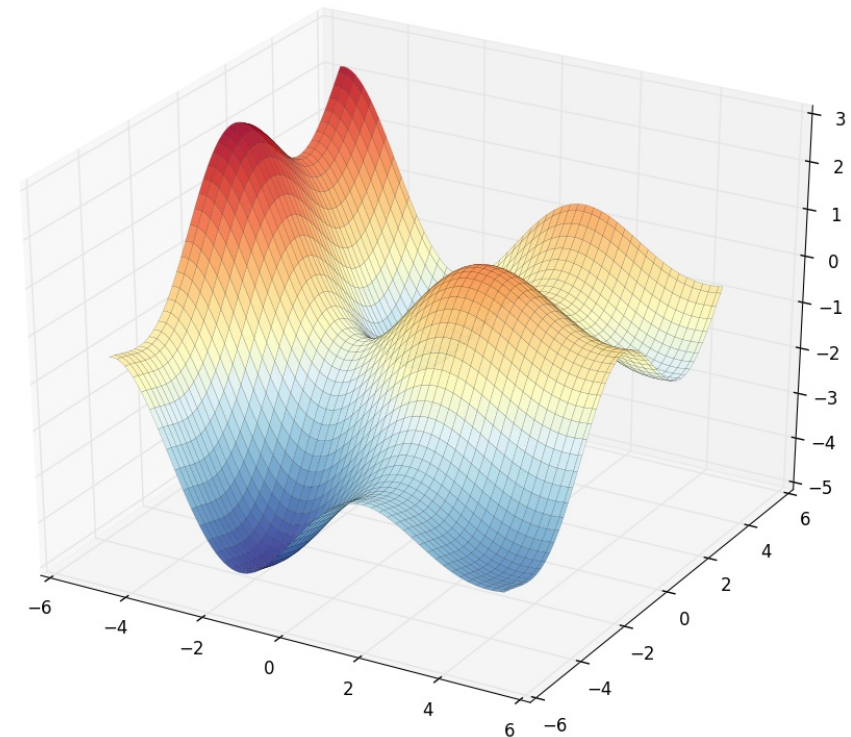
3D Plots

```
import matplotlib.pyplot as plt
import numpy as np
from mpl_toolkits.mplot3d import Axes3D

f = plt.figure(figsize=(10,8))
f.add_subplot(111, projection='3d')
ax = f.gca()

s=ax.plot_surface(x,y,z,
                 rstride=2, cstride=2,
                 lw=0.1,
                 alpha=0.9,
                 cmap=plt.cm.RdYlBu_r)
```

- `plot_surface()` plots the 3d surface
 - `rstride 2` - one gridline every 2 data rows
 - `lw` - grid line width
 - `alpha` - surface transparency



3D Plots

```
f = plt.figure(figsize=(10,8))
f.add_subplot(111, projection='3d')
ax = f.gca()

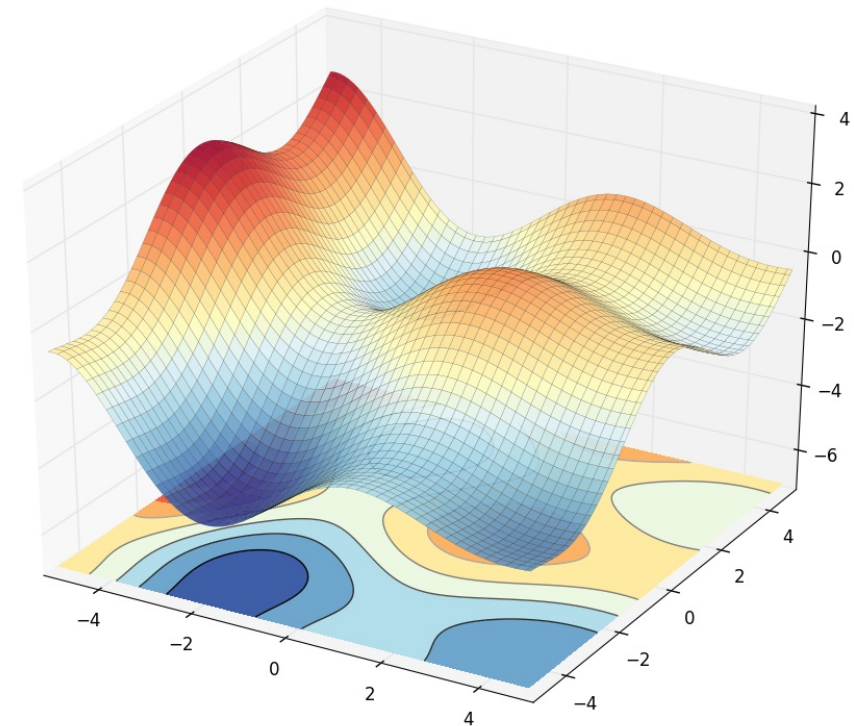
s=ax.plot_surface(x,y,z,rstride=2,
                  cstride=2,lw=0.1,alpha=0.9,
                  cmap=plt.cm.RdYlBu_r)

ax.contourf(x,y,z,
            zdir='z',
            offset=-7,
            cmap=plt.cm.RdYlBu_r)

ax.contour(x,y,z, zdir='z', offset=-7,
           lw=0.5, cmap=plt.cm.gray)

ax.set_zlim((-7,4))
```

- use `contour()` to plot the 2d contour
 - `zdir` - which data is "down"
 - `offset` - where to put the surface
 - `set_zlim()` set the floor of the graph

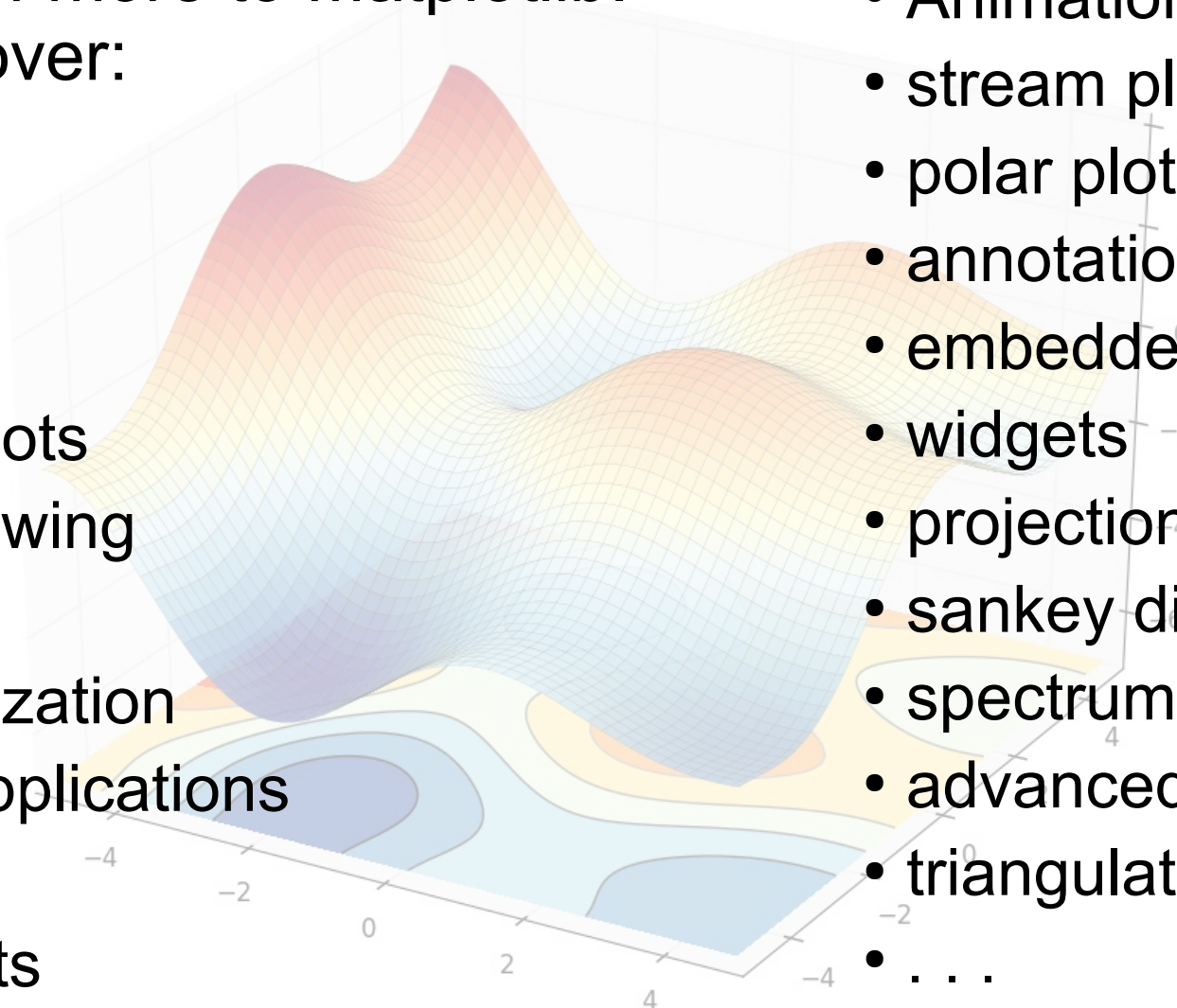


Other

- There is *much* more to Matplotlib.
We did not cover:

- Bar charts
- Pie charts
- scatterplots
- parametric plots
- free-form drawing
- boxplots
- image visualization
- Interactive applications
- Hinton plots
- output formats

- Animations
- stream plots
- polar plots
- annotations
- embedded plots
- widgets
- projections
- sankey diagrams
- spectrums and spectrograms
- advanced layouts
- triangulations
- . . .



Finally!

please fill in our feedback form!

<https://groups.oist.jp/scs/introduction-python-feedback>

Or go to SCDA → Documentation →
Introduction to Python → Feedback page

