

IDS 400

Programming for Data Science in Business

Tentative schedule

Date	Lecture Number	Topics
08/24	Lecture 1	Introduction
08/31	Lecture 2	Basic
09/07	Lecture 3	Condition
09/14	Lecture 4	Loop
09/21	Lecture 5	String + Quiz 1
09/28	Lecture 6	Type
10/05	Lecture 7	Function
10/12	Lecture 8	File + Quiz 2
10/19	Lecture 9	Pandas
10/26	Lecture 10	Numpy
11/02	Lecture 11	Machine Learning
11/09	Lecture 12	Visualization
11/16	Lecture 13	Web Scraping & Deep Learning
11/23	<i>Thanksgiving</i>	<i>No lecture</i>
11/30	Final presentation	In class presentation
12/05	Project submission due	

Overview

Steps for preparing data:

- Installing/Importing required packages
- Create a Dataframe using Pandas
- Dealing with missing data (remove/replace)
- Removing duplicate data (depends on what unique variables you would like to keep!)
- Filtering/Querying Data
- Using lambda function (rows/columns)
- Merging different datasets (left/right/outer/inner)

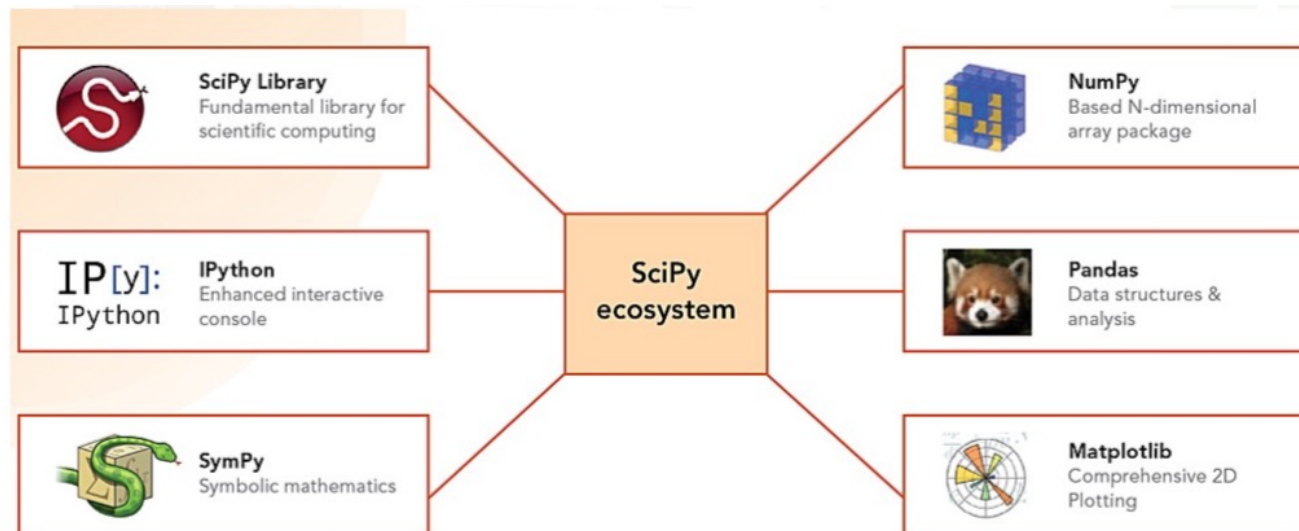
Scientific Applications

- There are a few packages available for scientific computing that extend Python's basic math module:
 - **NumPy** - numerical and scientific function libraries.
 - **Numba** - Python compiler that support JIT compilation.
 - **ALGLIB** - numerical analysis library.
 - **PyGSL** - Python interface for GNU Scientific Library.
 - **ScientificPython** - collection of scientific computing modules.

SciPy

- By far, the most commonly used packages are those in the SciPy stack.

SciPy is a Python-based ecosystem of open-source software for mathematics, science, and engineering. This stack can let you do scientific computing in Python. The six most important packages found in the SciPy stack include:



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SciPy is a Python-based ecosystem of open-source software for mathematics, science, and engineering. This stack can let you do scientific computing in Python. The six most important packages found in the SciPy stack include:

- **NumPy** - fundamental package for scientific computing.
- **SciPy** - efficient numerical routines.
- **Matplotlib** - plotting library.
- **IPython** – interactive computing.
- **SymPy** – symbolic computation library.
- **Pandas** – data analysis library.

NumPy

- The fundamental package for scientific computing with Python. It contains:
 - A powerful N-dimensional array (ndarray) object.
 - Sophisticated (broadcasting/universal) functions.
 - Tools for integrating C/C++ and Fortran code.
 - Useful linear algebra, Fourier transform, and random number capabilities.
- Besides its obvious scientific uses, NumPy can also be used as an efficient multi-dimensional container of generic data.

Import NumPy

```
import numpy
import numpy as np
from numpy import * # import all functions

dir(np) # show all the functions contained in Numpy
```

```
['ALLOW_THREADS',
 'AxisError',
 'BUFSIZE',
 'CLIP',
 'ComplexWarning',
 'DataSource',
 'ERR_CALL',
 'ERR_DEFAULT',
 'ERR_IGNORE',
 'ERR_LOG',
 'ERR_PRINT',
 'ERR_RAISE',
 'ERR_WARN',
 'FLOATING_POINT_SUPPORT',
 'FPE_DIVIDEBYZERO',
 'FPE_INVALID',
 'FPE_OVERFLOW',
 'FPE_UNDERFLOW',
 'False_',
```

```
help(np.ndarray) # Help on class ndarray in module numpy
```

Help on class ndarray in module numpy:

```
class ndarray(builtins.object)
|   ndarray(shape, dtype=float, buffer=None, offset=0,
|           strides=None, order=None)
|
|   An array object represents a multidimensional, homogeneous array
|   of fixed-size items. An associated data-type object describes the
|   format of each element in the array (its byte-order, how many bytes it
|   occupies in memory, whether it is an integer, a floating point number,
|   or something else, etc.)
|
|   Arrays should be constructed using `array`, `zeros` or `empty` (refer
|   to the See Also section below). The parameters given here refer to
|   a low-level method (`ndarray(...)`) for instantiating an array.
|
|   For more information, refer to the `numpy` module and examine the
|   methods and attributes of an array.
```


NumPy Data Types

- By default, Python have these data types:
 - **strings** - used to represent text data, the text is given under quote marks. eg. "ABCD"
 - **integer** - used to represent integer numbers. eg. -1, -2, -3
 - **float** - used to represent real numbers. eg. 1.2, 42.42
 - **boolean** - used to represent True or False.
 - **complex** - used to represent a number in complex plain. eg. $1.0 + 2.0j$, $1.5 + 2.5j$

NumPy Data Types

- NumPy has some extra data types:
 - bool_, int_, intc, intp, u/int8, u/int16, u/int32, u/int64, float_, float16, float32, float64 complex_, complex64, complex128, ...

<https://numpy.org/devdocs/user/basics.types.html>

- NumPy numerical types are instances (data-type) objects:
`numpy.dtype(object, align, copy)`

NumPy Data Types

```
# np.float is an alias for python float type.  
# np.float32 is numpy specific 32-bit float types.
```

```
x = np.float32(1.0)  
x
```

```
1.0
```

```
y = np.int_([1,2,4])  
y
```

```
array([1, 2, 4])
```

```
y.dtype
```

```
dtype('int32')
```

```
z = np.arange(3, dtype=np.uint8) # uint8: Unsigned integer (0 to 255)  
z
```

```
array([0, 1, 2], dtype=uint8)
```

```
z.dtype
```

```
dtype('uint8')
```

arange is a function in numpy which generates an array with evenly spaced values.

NumPy Arrays

- The main feature of NumPy is an array object:
 - Arrays can be N-dimensional.
 - Array elements have to be the same type.
 - Array elements can be accessed, sliced, and manipulated in the same way as the lists.
 - The number of elements in the array is fixed.
 - Shape of the array can be changed.
- Built-in NumPy array creation:
`array()`, `arange()`, `ones()`, `zeros()`, ...

NumPy Arrays

```
np.array([2,3,1,0])
```

```
array([2, 3, 1, 0])
```

```
np.array([[1,2.0],[0,0],[1+1j,3.]])
```

```
array([[1.+0.j, 2.+0.j],  
       [0.+0.j, 0.+0.j],  
       [1.+1.j, 3.+0.j]])
```

```
l = np.array([[1,2.0],[0,0],[1+1j,3.]])
```

```
l.shape
```

```
(3, 2)
```

```
np.zeros((2,3))
```

```
array([[0., 0., 0.],  
       [0., 0., 0.]])
```

```
np.zeros((2,3,4))
```

```
array([[[0., 0., 0., 0.],  
        [0., 0., 0., 0.],  
        [0., 0., 0., 0.]],  
       [[0., 0., 0., 0.],  
        [0., 0., 0., 0.],  
        [0., 0., 0., 0.]])
```

NumPy Arrays

```
np.arange(10)
```

```
array([0, 1, 2, 3, 4, 5, 6, 7, 8, 9])
```

```
np.arange(2,10,dtype=np.float)
```

```
array([2., 3., 4., 5., 6., 7., 8., 9.])
```

```
np.arange(2,3,0.1)
```

```
array([2. , 2.1, 2.2, 2.3, 2.4, 2.5, 2.6, 2.7, 2.8, 2.9])
```

NumPy Arrays

```
a = np.arange(3)
```

```
a
```

```
array([0, 1, 2])
```

```
print(a)
```

```
[0 1 2]
```

Reshape function change the dimension of a numpy array

```
np.arange(9).reshape(3,3)
```

```
array([[0, 1, 2],  
       [3, 4, 5],  
       [6, 7, 8]])
```

```
np.arange(24).reshape(2,3,4)
```

```
array([[[ 0,  1,  2,  3],  
       [ 4,  5,  6,  7],  
       [ 8,  9, 10, 11]],  
      [[12, 13, 14, 15],  
       [16, 17, 18, 19],  
       [20, 21, 22, 23]])]
```

NumPy Arrays

- `linspace(start, stop[, num, endpoint, retstep, dtype])`
creates arrays with a specified number of elements, and spaced equally between the specified beginning and end values.
- `random.random([size])`
creates arrays with random floats over the interval [0.,1.).
- `random.randint(low[, high, size, dtype])`
creates arrays with random integers from low (inclusive) to high (exclusive).

NumPy Arrays

```
np.linspace(1.,4.,6)
```

```
array([1. , 1.6, 2.2, 2.8, 3.4, 4. ])
```

```
np.random.random((2,3))
```

```
array([[0.48724427, 0.87441327, 0.45243135],  
       [0.0660517 , 0.32073882, 0.30583646]])
```

```
np.random.randint(1,7,(2,6))
```

```
array([[4, 5, 6, 2, 4, 2],  
       [2, 2, 5, 6, 5, 4]])
```

Statistics in NumPy

```
a = np.random.randint(1,7,(2,3))  
a
```

```
array([[6, 1, 4],  
       [4, 3, 5]])
```

Column-wise: axis= 0

Row-wise: axis = 1

- Then try the following command:

- **a.max()**
- **a.min()**
- **a.argmax()**
- **a.argmin()**
- **np.amax(a,0)**
- **np.amax(a,1)**
- **np.mean(x)**
- **np.std(x)**
- **np.var(x)**

```
# Returns the indices of the maximum values.  
a.argmax()  
# Returns the indices of the maximum values along an axis.  
np.argmax(a, 0)
```

```
array([0, 1, 1], dtype=int64)
```

```
#Return the maximum of an array or maximum along an axis.  
np.amax(a, 0)
```

```
array([6, 3, 5])
```

```
np.amax(a, 1)
```

```
array([6, 5])
```

Sorting in NumPy

```
a = np.random.randint(1,7,(2,3))
```

a

```
array([[6, 1, 4],  
       [4, 3, 5]])
```

Then try the following command:

- `np.sort(a,axis=1)`
- `np.sort(a,axis=0)`

Column-wise: axis= 0

Row-wise: axis = 1

```
np.sort(a,axis=0)
```

```
array([[4, 1, 4],  
       [6, 3, 5]])
```

```
np.sort(a,axis=1)
```

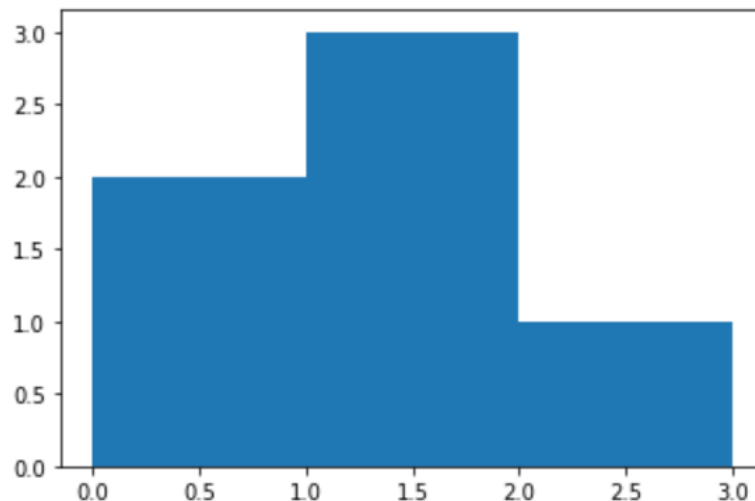
```
array([[1, 4, 6],  
       [3, 4, 5]])
```

Histogram using matplotlib

```
import matplotlib.pyplot as plt
import numpy as np
x = np.array([0.5, 0.7, 1.0, 1.2, 1.3, 2.1])
bins1 = np.array([0, 1, 2, 3])
print("ans=\n", np.histogram(x, bins1))
```

```
ans=
(array([2, 3, 1], dtype=int64), array([0, 1, 2, 3]))
```

```
plt.hist(x, bins=bins1)
plt.show()
```



This part returns information about the histogram

For histogram() and hist(), first parameter is data, second parameter is number of bins

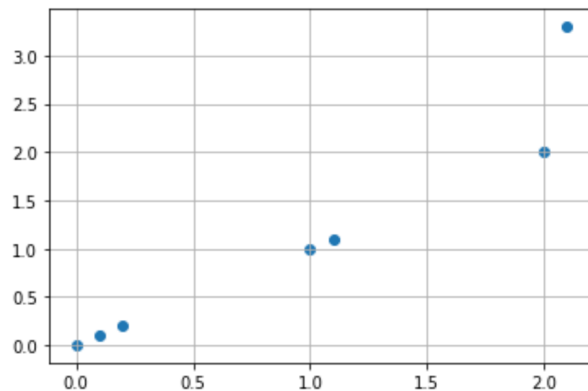
2D Histogram using matplotlib

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

xedges= [0, 1, 2, 3]
yedges= [0, 1, 2, 3, 4]
x = np.array([0, 0.1, 0.2, 1., 1.1, 2., 2.1])
y = np.array([0, 0.1, 0.2, 1., 1.1, 2., 3.3])
H, xedges, yedges= np.histogram2d(x, y, bins=(xedges, yedges))
print("ans=\n", H)

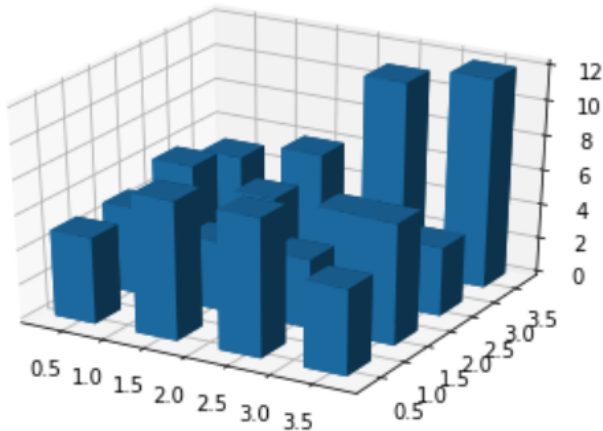
plt.scatter(x, y)
plt.grid()
plt.show()
```

```
ans=
[[3. 0. 0. 0.]
 [0. 2. 0. 0.]
 [0. 0. 1. 1.]
```



If you want to plot a 3D histogram

```
from mpl_toolkits.mplot3d import Axes3D
import matplotlib.pyplot as plt
import numpy as np
np.random.seed(19680801)
fig = plt.figure()
ax = fig.add_subplot(111, projection='3d')
x, y = np.random.rand(2, 100) * 4
hist, xedges, yedges = np.histogram2d(x, y, bins=4, range=[[0, 4], [0, 4]])
xpos, ypos = np.meshgrid(xedges[:-1] + 0.25, yedges[:-1] + 0.25, indexing="ij")
xpos = xpos.ravel()
ypos = ypos.ravel()
zpos = 0
dx = dy = 0.5 * np.ones_like(zpos)
dz = hist.ravel()
ax.bar3d(xpos, ypos, zpos, dx, dy, dz, zsort='average')
plt.show()
```



NumPy Linear Algebra

- All linear algebra routines expect an object that can be converted into a 2-dimensional array.
- The output is also a two-dimensional array.
 - `dot(a, b[, out])` - dot product of two arrays.
 - `trace(a[, offset, axis1, axis2, dtype, out])` - returns the sum along diagonals of the array.
 - `inv(a)` - computes the inverse of a matrix.
 - `eig(a)` - eigenvalues and right eigenvectors of a square array.
 - `solve(a, b)` - solves a linear matrix equation, or system of linear scalar equations.

NumPy LinAlg

```
from numpy import *  
from numpy.linalg import *  
a = array([[1.0, 2.0], [3.0, 4.0]])  
print(a)
```

```
[[1. 2.]  
 [3. 4.]]
```


```
a.transpose()
```

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

```
inv(a) #inverse
```

```
array([[ -2. ,  1. ],  
       [ 1.5, -0.5]])
```

$$\begin{bmatrix} a & b \\ c & d \end{bmatrix}^{-1} = \frac{1}{ad-bc} \begin{bmatrix} d & -b \\ -c & a \end{bmatrix}$$


determinant

NumPy LinAlg

```
#unit 2x2 matrix; "eye"~"I"  
u = eye(2)  
u
```

```
array([[1., 0.],  
       [0., 1.]])
```

```
trace(u)
```

```
2.0
```

```
j = array([[0.0, -1.0], [1.0, 0.0]])  
dot(j,j) # matrix product
```

```
array([[ -1.,  0.],  
       [ 0., -1.]])
```

```
eig(j) # get eigenvalues & eigenvectors
```

```
(array([0.+1.j, 0.-1.j]),  
 array([[0.70710678+0.j, 0.70710678-0.j],  
        [0.      -0.70710678j, 0.      +0.70710678j]]))
```

The trace of a square matrix A is defined to be the sum of elements on the main diagonal (from the upper left to the lower right) of A .

Let A be a matrix, with

$$A = \begin{pmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{33} \end{pmatrix} = \begin{pmatrix} -1 & 0 & 3 \\ 11 & 5 & 2 \\ 6 & 12 & -5 \end{pmatrix}$$

Then

$$\text{tr}(A) = \sum_{i=1}^3 a_{ii} = a_{11} + a_{22} + a_{33} = -1 + 5 + (-5) = -1$$

"Dot Product"

$$\begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix} \times \begin{bmatrix} 7 & 8 \\ 9 & 10 \\ 11 & 12 \end{bmatrix} = \begin{bmatrix} 58 \\ \end{bmatrix}$$

About eigenvalue:

[https://mathworld.wolfram.com/Eigenvalue.html#:~:text=Entries%20%3E%20Interactive%20Demonstrations%20%3E-,Eigenvalue,144\).](https://mathworld.wolfram.com/Eigenvalue.html#:~:text=Entries%20%3E%20Interactive%20Demonstrations%20%3E-,Eigenvalue,144).)

NumPy LinAlg

- There are two alcohol solutions: 50% & 90%.
- How many gallons of each solution to be mixed to get 10 gallons of 74% alcohol solution?

- $x_1 + x_2 = 10$

$$AX = Y$$

- $0.5x_1 + 0.9x_2 = 0.74 \cdot 10 = 7.4$

```
A = array([[1.0,1.0],[0.5,0.9]])  
Y = array([[10.0],[7.4]])  
solve(A,Y) #solve linear equations
```

```
array([[4.],  
       [6.]])
```

NumPy LinAlg

- A drone flying with the wind could cover in 2 hours.
- The return trip against the wind took 2.5 hours.
- How fast was the drone?
- What was the air speed?

$$2d + 2w = 60$$

$$2.5d - 2w = 60$$

Trip	Rate		Time		Distance
With wind	d + w	×	2	=	60
Against wind	d - w	×	2.5	=	60

```
A = array([[2.0, 2.0], [2.5, -2.5]])
Y = array([[60.0], [60.0]])
solve(A, Y) #solve linear equations

array([[27.],
       [ 3.]])
```

NumPy Matrix Versus Array

- NumPy matrices are strictly 2-dimensional, while NumPy arrays (ndarrays) are N-dimensional.
- Matrix objects are a subclass of ndarray, so they inherit all the attributes and methods of ndarrays.
- The main advantage of NumPy matrices is that they provide a convenient notation for matrix multiplication. e.g. If A and B are matrices, then $A*B$ is their matrix product.

NumPy Matrices

```
A = matrix('1.0 2.0; 3.0 4.0')  
A
```

```
matrix([[1., 2.],  
        [3., 4.]])
```

```
type(A)
```

```
numpy.matrix
```

```
Y = matrix('5.0; 7.0')
```

```
print(A.I) #inverse
```

```
[[ -2.   1. ]  
 [ 1.5 -0.5]]
```

```
print(A.I*Y) #multiplication
```

```
[[ -3.]  
 [  4.]]
```

```
solve(A,Y) #solve linear equations
```

```
matrix([[ -3.],  
        [  4.]])
```

Note:

$$A.X = Y$$

$$\text{Inv}(A)*Y = X \quad \text{or} \quad A.I*Y = X$$

SciPy

- A collection of mathematical algorithms and convenient functions built on the NumPy extension of Python.
- An interactive Python session for manipulating and visualizing data.
- A data-processing and system-prototyping environment rivaling systems such as MATLAB, IDL, Octave, R-lab, and SciLab.

```
>>> import scipy
```

SciPy Sub -- Modules

- **cluster** -clustering algorithms
- **integrate** - integration and ordinary differential equation solvers.
- **interpolate** - interpolation and smoothing splines
- **io** - input and output
- **linalg** - linear algebra
- **optimize** - optimization and root-finding routines
- **stats** – statistical distributions and functions

```
>>> from scipy import linalg, optimize
```

```
>>> from scipy import *
```

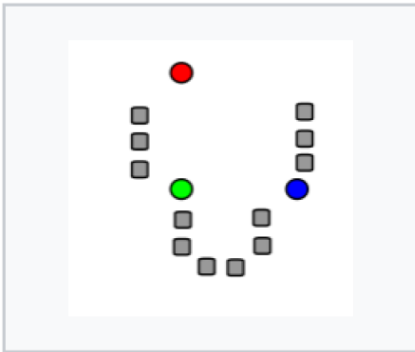
SciPy Clustering

- **Clustering** - finds clusters and cluster centers in a set of unlabeled data.
- Intuitively, a cluster comprises a group of data points whose inter-point distances are small compared to the distances to points outside of the cluster.

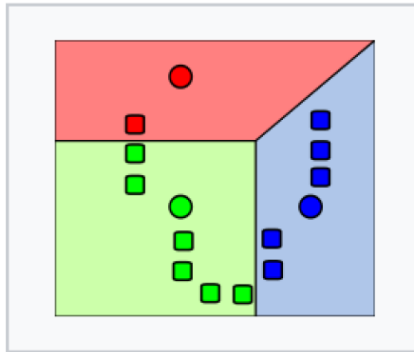
SciPy *K-means Clustering*

- **scipy.cluster.vq**
 - **kmeans(obs, k_or_guess[, iter, thresh, ...])**
 - perform k-means on a set of observation vectors forming k clusters.
 - **kmeans2(data, k[, iter, thresh, minit, ...])**
 - classify a set of observations into k clusters using the k-means algorithm.
- Given an initial set of k centers, the k-means algorithm alternates the two steps:
 - 1) For each center, we identify the subset of training points (its cluster) that is closer to it than any other center.
 - 2) The means of each feature for the data points in each cluster are computed, and this mean vector becomes the new center for that cluster.

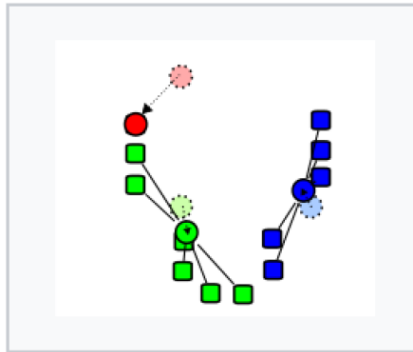
K-means Clustering (k=3)



k initial "means" (in this case $k=3$) are randomly generated within the data domain (shown in color).



k clusters are created by associating every observation with the nearest mean.



The centroid of each of the k clusters becomes the new mean.



Steps 2 and 3 are repeated until convergence has been reached.

SciPy *2-means Clustering*

```
from pylab import *
from numpy import *
from numpy.random import *
from scipy.cluster.vq import *
```

```
# data generation
```

```
data = vstack((rand(100,2)+array([.5,.5]),rand(100,2)))
```

```
# computing k-means with k = 2 (2 clusters)
```

```
centroids,_ = kmeans(data,2)
```

```
# assign each sample to a cluster
```

```
index,_ = vq(data,centroids)
```

```
# some plotting using numpy's logical indexing
```

```
plot(data[index==0,0],data[index==0,1],'or',
```

```
data[index==1,0],data[index==1,1],'ob')
```

```
plot(centroids[:,0],centroids[:,1],'sg',markersize=8)
```

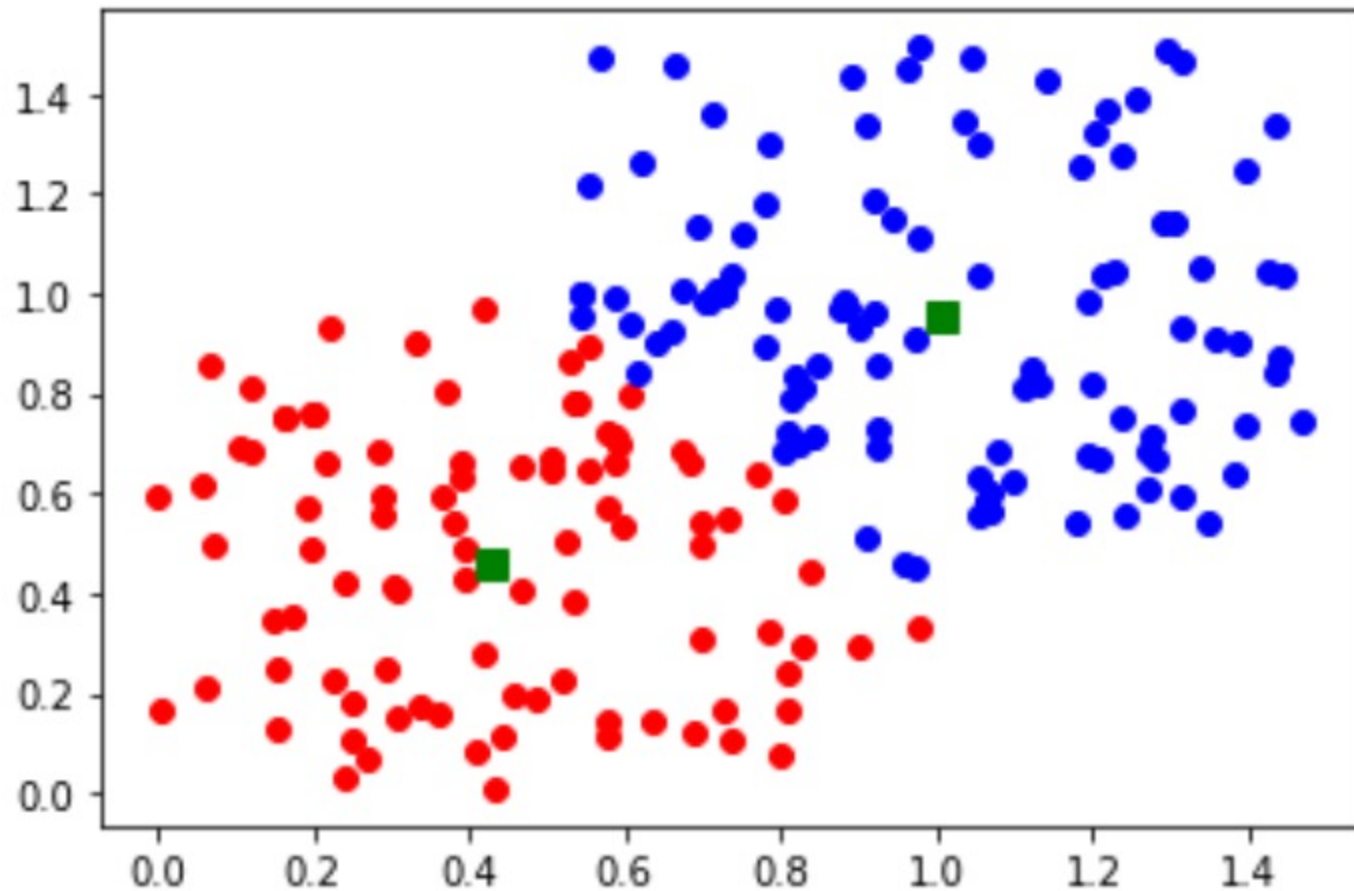
```
show()
```

'o': Use circle markers.

'r': Use red color.

's': Use square markers

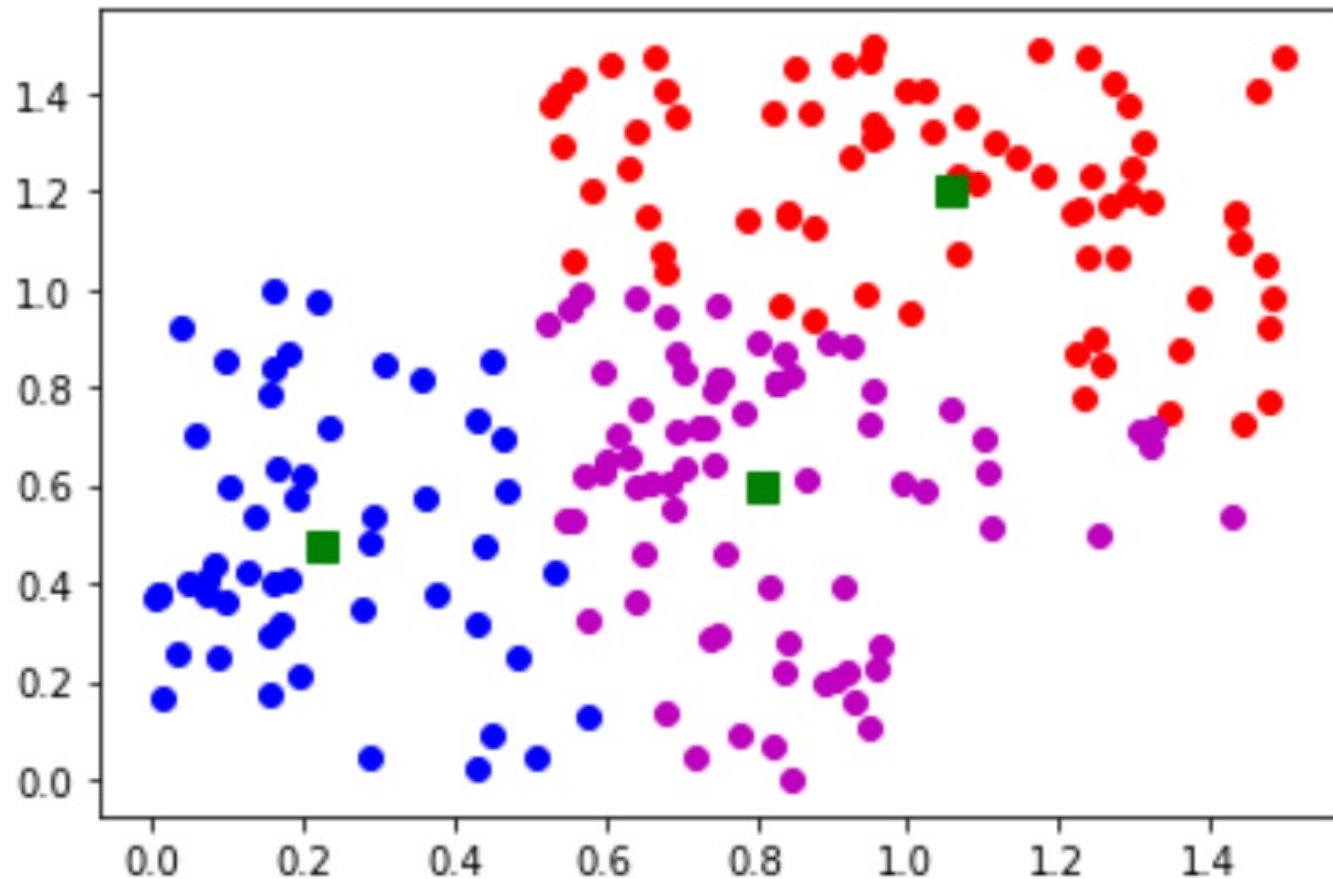
SciPy *2-means* Clustering



SciPy *3-means Clustering*

```
#data generation
data = vstack((rand(100,2)+array([.5,.5]),rand(100,2)))
# computing k-means with k = 3 (3 clusters)
centroids,_ = kmeans(data,3)
# assign each sample to a cluster
index,_ = vq(data,centroids)
# some plotting using numpy's Logical indexing
plot(data[index==0,0],data[index==0,1],'or',
data[index==1,0],data[index==1,1],'ob',
data[index==2,0],data[index==2,1],'om')
plot(centroids[:,0],centroids[:,1],'sg',markersize=8)
show()
```

SciPy *3-means* Clustering





Exercise

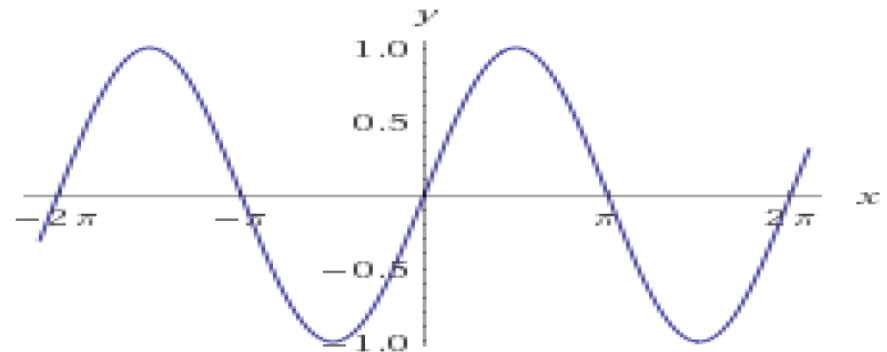
- Conduct a 4-means clustering and plot it using matplotlib

SciPy Integration

- Methods for Integrating Functions given a function object:
 - **quad** - general purpose integration
 - **dblquad** - general purpose double integration
 - **tplquad** - general purpose triple integration
 - **fixed_quad** - integrate $f(x)$ using Gaussian quadrature
 - **quadrature** - integrate with tolerance using Gaussian quadrature
 - **romberg** - integrate $f(x)$ using Romberg integration
- Methods for I.F. given a fixed set of samples:
 - **trapz** - use trapezoidal rule to compute integral
 - **cumtrapz** - use trapezoidal rule to cumulatively compute integral
 - **simps** - use Simpson's rule to compute integral
 - **romb** - use Romberg Integration to compute integral

SciPy Integration

- **np.sin** defines the sine function
- Integral $x=0$ to $x=\pi$ using **quad**



$$\int \sin(x) dx$$

$$\int_0^{\pi} \sin(x) dx = -\cos(x) \Big|_0^{\pi} = -\cos(\pi) - (-\cos(0)) = -(-1) - (-1) = 1 + 1 = 2$$

```
from scipy.integrate import *  
result = scipy.integrate.quad(np.sin, 0, np.pi)  
print(result)
```

2 with a very small error margin!

```
(2.0, 2.220446049250313e-14)
```

```
result = scipy.integrate.quad(np.sin, - np.inf, +np.inf)
```

```
print(result)  
# Integral does not converge
```

```
(0.0, 0.0)
```

SciPy Optimization

- Provides several commonly used optimization algorithms:
 - Unconstrained and constrained minimization of multivariate scalar functions (minimize) using BFGS, Nelder-Mead Simplex, Newton Conjugate Gradient, COBYLA, SLSQP, ...
 - Global (brute-force) optimization routines (e.g. basinhopping, differential_evolution)
 - Least-squares minimization (least_squares) and curve fitting (curve_fit) algorithms
 - Scalar univariate functions minimizers (minimize_scalar) and root finders (newton)
 - Multivariate equation system solvers (root) using hybrid Powell, Levenberg-Marquardt, large-scale Newton-Krylov, ...

SciPy Curve Fitting

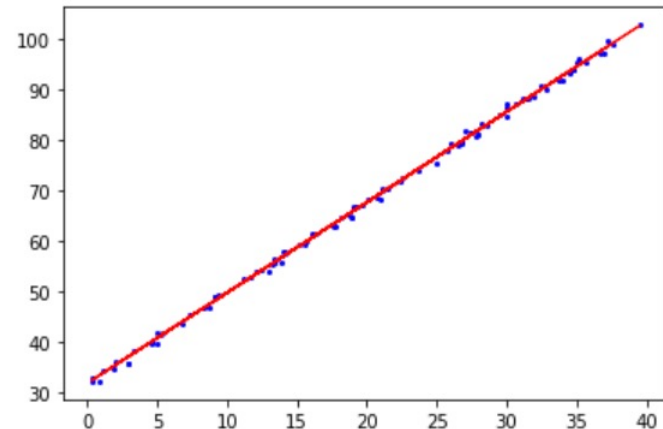
```
from pylab import *
from numpy import *
from numpy.random import *
from scipy.optimize import *

# linear regression
def linreg(x,a,b):
    return a*x+b

# data generation
input1 = randint(0,40,100)
x = input1 + rand(100)
y = (input1 * 1.8 + 32) + rand(100)

# curve fitting
attributes,variances= curve_fit(linreg,x,y)
# estimated y
y_modeled= x*attributes[0]+attributes[1]

# plot true and modeled results
plot(x,y,'ob',markersize=2)
plot(x,y_modeled,'-r',linewidth=1)
show()
```



SciPy Linear Regression

```
from pylab import *
from numpy import *
from scipy.stats import *

# data generation
input1 = random.randint(0,40,100)
x = input1+rand(100)
y = (input1*1.8+32)+rand(100)

# linear regression
slope,intercept,r_value,p_value,slope_std_error= stats.linregress(x,y)
# estimated y
y_modeled= x*slope+intercept

# plot true and modeled results
plot(x,y,'ob',markersize=2)
plot(x,y_modeled,'-r',linewidth=1)
show()
```

