# **Python Basics**

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# Variables and Data Types

# Variable Assignment

>>>	x=5
>>>	X
5	

## Calculations With Variables

>>> x+2	Sum of two variables
7 >>> x-2	Subtraction of two variables
3 >>> x*2	Multiplication of two variables
10 >>> x**2	Exponentiation of a variable
25 >>> x%2	Remainder of a variable
1 >>> x/float(2)	Division of a variable
2.5	2.1.5.5 5. 4. 14.14516

# Types and Type Conversion

str()	'5', '3.45', 'True'	Variables to strings
int()	5, 3, 1	Variables to integers
float()	5.0, 1.0	Variables to floats
bool()	True, True, True	Variables to booleans

# **Asking For Help**

>>> help(str)

# Strings

```
>>> my_string = 'thisStringIsAwesome'
>>> my_string
'thisStringIsAwesome'
```

# **String Operations**

```
>>> my_string * 2
 'thisStringIsAwesomethisStringIsAwesome'
>>> my_string + 'Innit'
 'thisStringIsAwesomeInnit'
>>> 'm' in my_string
True
```

### Lists

```
>>> a = 'is'

>>> b = 'nice'

>>> my_list = ['my', 'list', a, b]

>>> my_list2 = [[4,5,6,7], [3,4,5,6]]
```

## Selecting List Elements

### Index starts at o

Also see NumPy Arrays

### Subset

Jub	366	
>>>	my_	_list[1]
>>>	my_	list[-3]
Slic	e ¯	

- >>> my\_list[1:3]
  >>> my\_list[1:]
  >>> my\_list[:3]
  >>> my\_list[:]
- Subset Lists of Lists
  >>> my\_list2[1][0]
  >>> my list2[1][:2]
- my\_list[list][itemOfList]

Copy my list

Select item at index 1
Select 3rd last item

Select items at index 1 and 2

Select items after index o

Select items before index 3

### **List Operations**

```
>>> my_list + my_list
['my', 'list', 'is', 'nice', 'my', 'list', 'is', 'nice']
>>> my_list * 2
['my', 'list', 'is', 'nice', 'my', 'list', 'is', 'nice']
>>> my_list2 > 4
```

### **List Methods**

>>>	<pre>my_list.index(a)</pre>	Get the index of an item
>>>	<pre>my_list.count(a)</pre>	Count an item
>>>	<pre>my_list.append('!')</pre>	Append an item at a time
>>>	<pre>my_list.remove('!')</pre>	Remove an item
>>>	del(my_list[0:1])	Remove an item
>>>	<pre>my_list.reverse()</pre>	Reverse the list
>>>	<pre>my_list.extend('!')</pre>	Append an item
>>>	<pre>my_list.pop(-1)</pre>	Remove an item
>>>	<pre>my_list.insert(0,'!')</pre>	Insert an item
>>>	<pre>my_list.sort()</pre>	Sort the list

# **String Operations**

### Index starts at o

```
>>> my_string[3]
>>> my_string[4:9]
```

# **String Methods**

- 4	ourning intentions	
>	>> my_string.upper()	String to uppercase
>	>> my string.lower()	String to lowercase
>	>> my string.count('w')	Count String elements
>	<pre>&gt;&gt;&gt; my string.replace('e', 'i')</pre>	Replace String elements
>	>> my string.strip()	Strip whitespaces

### Libraries

### **Import libraries**

>>> import numpy

>>> import numpy as np
Selective import

>>> from math import pi

# pandas $\lim_{y,t=\beta'x_u+\mu_t+\epsilon_u} \lim_{y,t=\beta'x_u+\mu_t+\epsilon_u} \lim_{y,t=\beta'x_u+$



Machine learning

NumPy \*matplotlib
Scientific computing 2D plotting

## Install Python



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Create and share documents with live code, visualizations, text, ...

# Numpy Arrays

### Also see Lists

```
>>> my_list = [1, 2, 3, 4]
>>> my_array = np.array(my_list)
>>> my_2darray = np.array([[1,2,3],[4,5,6]])
```

# Selecting Numpy Array Elements

### Index starts at o

```
Subset
>>> my_array[1]
Select item at index 1
```

# Slice

```
>>> my_array[0:2]
    array([1, 2])

Subset 2D Numpy arrays
>>> my_2darray[:,0]
    array([1, 4])
```

Select items at index 0 and 1

my\_2darray[rows, columns]

## Numpy Array Operations

```
>>> my_array > 3
    array([False, False, False, True], dtype=bool)
>>> my_array * 2
    array([2, 4, 6, 8])
>>> my_array + np.array([5, 6, 7, 8])
    array([6, 8, 10, 12])
```

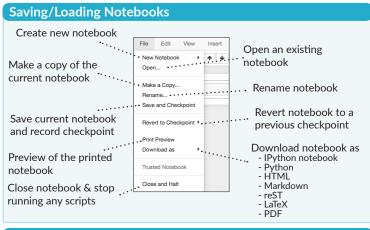
## **Numpy Array Functions**

```
>>> my array.shape
                                      Get the dimensions of the array
>>> np.append(other array)
                                      Append items to an array
>>> np.insert(my array, 1, 5)
                                      Insert items in an array
>>> np.delete(my array,[1])
                                      Delete items in an array
>>> np.mean(my array)
                                      Mean of the array
>>> np.median(my array)
                                      Median of the array
>>> my array.corrcoef()
                                      Correlation coefficient
>>> np.std(my array)
                                      Standard deviation
```

# Python For Data Science Cheat Sheet Jupyter Notebook

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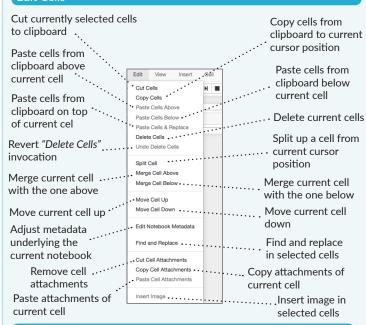
Code and text are encapsulated by 3 basic cell types: markdown cells, code cells, and raw NBConvert cells.

### Edit Cells

**Insert Cells** 

current one

Add new cell above the

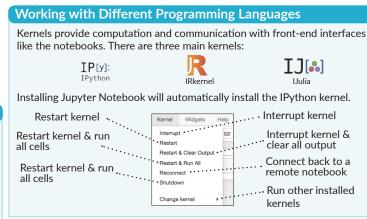


Cell

Insert Cell Relow

Add new cell below the

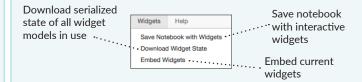
current one



### Widgets

Notebook widgets provide the ability to visualize and control changes in your data, often as a control like a slider, textbox, etc.

You can use them to build interactive GUIs for your notebooks or to synchronize stateful and stateless information between Python and JavaScript.

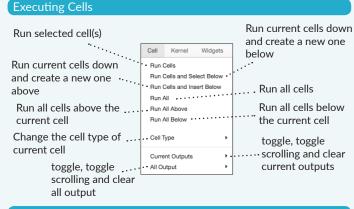


### **Command Mode:**

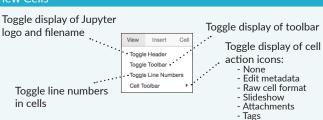




In [ ]: |



### View Cells

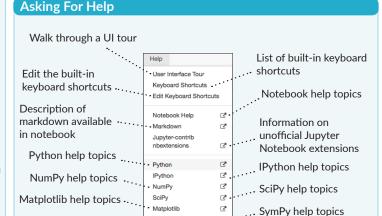


- 1. Save and checkpoint
- 2. Insert cell below
- 3. Cut cell

Pandas help topics ....

- 4. Copy cell(s)
- 5. Paste cell(s) below
- 6. Move cell up
- 7. Move cell down
- 8. Run current cell

- 9. Interrupt kernel
- 10. Restart kernel11. Display characteristics
- **12**. Open command palette
- 13. Current kernel
- 14. Kernel status
- 15. Log out from notebook server





About Jupyter Notebook

# **Python For Data Science** *Cheat Sheet*SciPv - Linear Algebra

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# SciPy

The **SciPy** library is one of the core packages for scientific computing that provides mathematical algorithms and convenience functions built on the NumPy extension of Python.



# **Interacting With NumPy**

### Also see NumPy

```
>>> import numpy as np
>>> a = np.array([1,2,3])
>>> b = np.array([(1+5j,2j,3j), (4j,5j,6j)])
>>> c = np.array([[(1.5,2,3), (4,5,6)], [(3,2,1), (4,5,6)]])
```

### **Index Tricks**

>>> np.mgrid[0:5,0:5]	Create a dense meshgrid
>>> np.ogrid[0:2,0:2]	Create an open meshgrid
>>> np.r [3,[0]*5,-1:1:10j]	Stack arrays vertically (row-wise)
>>> np.c [b,c]	Create stacked column-wise arrays

## Shape Manipulation

>>> np.transpose(b)	Permute array dimensions
	Flatten the array
>>> np.hstack((b,c))	Stack arrays horizontally (column-wise)
	Stack arrays vertically (row-wise)
>>> np.hsplit(c,2)	Split the array horizontally at the 2nd index
>>> np.vpslit(d,2)	Split the array vertically at the 2nd index

## Polynomials

>>>	from numpy	import polyld	
>>>	p = poly1d	([3,4,5])	Create a polynomial object

# **Vectorizing Functions**

```
>>> def myfunc(a):
    if a < 0:
        return a*2
    else:
        return a/2
>>> np.vectorize(myfunc)
Vectorize functions
```

## Type Handling

>>> np.imag(b) >>> np.real_if_close(c,tol=1000)	Return the real part of the array elements Return the imaginary part of the array elemen Return a real array if complex parts close to o Cast object to a data type
---	--

## Other Useful Functions

>>>	np.angle(b,deg=True)	Return the angle of the complex argument
>>>	g = np.linspace(0,np.pi,num=5)	Create an array of evenly spaced values
>>>	g [3:] += np.pi	(number of samples)
>>>	np.unwrap(g)	Unwrap
>>>	np.logspace(0,10,3)	Create an array of evenly spaced values (log scale)
>>>	np.select([c<4],[c*2])	Return values from a list of arrays depending on
		conditions
>>>	misc.factorial(a)	Factorial
>>>	misc.comb(10,3,exact=True)	Combine N things taken at k time
>>>	misc.central_diff_weights(3)	Weights for Np-point central derivative
>>>	misc.derivative(myfunc, 1.0)	Find the n-th derivative of a function at a point

# Linear Algebra Also see NumPy

```
You'll use the linalg and sparse modules. Note that scipy.linalg contains and expands on numpy.linalg.
```

>>> from scipy import linalg, sparse

# **Creating Matrices**

```
>>> A = np.matrix(np.random.random((2,2)))
>>> B = np.asmatrix(b)
>>> C = np.mat(np.random.random((10,5)))
>>> D = np.mat([[3,4], [5,6]])
```

### **Basic Matrix Routines**

### Inverse

>>>	A.I
>>>	linalg.inv(A)

# Transposition >>> A.T

### >>> A.н **Trace**

>>> np.trace(A)

### Norm

>>>	linalg.norm(A)
>>>	linalg.norm(A,1)
>>>	<pre>linalg.norm(A,np.inf)</pre>
_	

### Rank

>>> np.linalg.matrix\_rank(C)

### **Determinant**

>>> linalg.det(A)

# Solving linear problems

>>>	linalg.solve(A,b)
>>>	E = np.mat(a).T
>>>	<pre>E = np.mat(a).T linalg.lstsq(F,E)</pre>

### **Generalized inverse**

>>>	linalg.pinv(C)
>>>	linalg piny2(C)

### Inverse Inverse

Tranpose matrix Conjugate transposition

### Trace

Frobenius norm
L1 norm (max column sum)
L inf norm (max row sum)

### Matrix rank

Determinant

### Solver for dense matrices Solver for dense matrices Least-squares solution to linear matrix equation

# Compute the pseudo-inverse of a matrix (least-squares solver)

# Compute the pseudo-inverse of a matrix (SVD)

# **Creating Sparse Matrices**

>>>	F = np.eye(3, k=1)	Create a 2X2 identity matrix
>>>	<pre>G = np.mat(np.identity(2))</pre>	Create a 2x2 identity matrix
>>>	C[C > 0.5] = 0	
>>>	<pre>H = sparse.csr_matrix(C)</pre>	Compressed Sparse Row matrix
>>>	<pre>I = sparse.csc_matrix(D)</pre>	Compressed Sparse Column matrix
	<pre>J = sparse.dok_matrix(A)</pre>	Dictionary Of Keys matrix
>>>	E.todense()	Sparse matrix to full matrix
>>>	sparse.isspmatrix_csc(A)	Identify sparse matrix

### Sparse Matrix Routines

### Inverse

ı	>>>	sparse.linalg.inv(1)
	No	orm
	>>>	sparse.linalg.norm(I)

# Solving linear problems >>> sparse.linalg.spsolve(H, I)

Sparse Matrix Function

# Inverse

Norm

## Solver for sparse matrices

### **Sparse Matrix Functions**

sparse.linalg.expm(I) Sparse matrix exponential
---

### Matrix Functions

### Addition

>>> np.add(A,D)

Subtraction

### Subtraction

>>> np.subtract(A,D)

### Division

>>> np.divide(A,D)

# Multiplication >>> A @ D

```
>>> np.multiply(D,A)
>>> np.dot(A,D)
>>> np.vdot(A,D)
>>> np.inner(A,D)
>>> np.outer(A,D)
>>> np.tensordot(A,D)
>>> np.kron(A,D)
```

### **Exponential Functions**

```
>>> linalg.expm(A)
>>> linalg.expm2(A)
>>> linalg.expm3(D)
```

### **Logarithm Function**

>>> linalg.logm(A)

### **Trigonometric Functions**

	TTHATE STIME (D)
>>>	linalg.cosm(D)
>>>	linalg.tanm(A)

### **Hyperbolic Trigonometric Functions**

```
>>> linalg.sinhm(D)
>>> linalg.coshm(D)
>>> linalg.tanhm(A)
```

# Matrix Sign Function

>>> np.signm(A)

# **Matrix Square Root**

>>> linalg.sqrtm(A)

# Arbitrary Functions

>>> linalg.funm(A, lambda x: x\*x)

# Decompositions

# **Eigenvalues and Eigenvectors**

```
>>> la, v = linalg.eig(A)
>>> l1, l2 = la
>>> v[:,0]
>>> v[:,1]
>>> linalg.eigvals(A)
```

# Singular Value Decomposition

>>> U,s,Vh = linalg.svd(B) >>> M,N = B.shape >>> Sig = linalg.diagsvd(s,M,N)

# LU Decomposition

>>> P,L,U = linalg.lu(C)

# Solve ordinary or generalized eigenvalue problem for square matrix Unpack eigenvalues First eigenvector Second eigenvector Unpack eigenvalues

Addition

Subtraction

Multiplication operator

Division

(Python 3)

Multiplication

Inner product

Outer product

decomposition)

Matrix sine Matrix cosine

Matrix tangent

Hypberbolic matrix sine

Matrix sign function

Evaluate matrix function

Matrix square root

Hyperbolic matrix cosine

Hyperbolic matrix tangent

Vector dot product

Tensor dot product

Kronecker product

Matrix exponential

Matrix logarithm

Matrix exponential (Taylor Series)

Matrix exponential (eigenvalue

Dot product

# Singular Value Decomposition (SVD)

Construct sigma matrix in SVD

## LU Decomposition

## **Sparse Matrix Decompositions**

>>>	<pre>la, v = sparse.linalg.eigs(</pre>	F,1)
>>>	sparse.linalg.svds(H, 2)	

Eigenvalues and eigenvectors

# Asking For Help

>>> help(scipy.linalg.diagsvd)
>>> np.info(np.matrix)

# NumPy Basics

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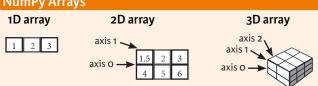
# NumPy

The **NumPy** library is the core library for scientific computing in Python. It provides a high-performance multidimensional array object, and tools for working with these arrays.

Use the following import convention: >>> import numpy as np



# NumPy Arrays



# **Creating Arrays**

```
>>> a = np.array([1,2,3])
>>> b = np.array([(1.5,2,3), (4,5,6)], dtype = float)
>>> c = np.array([[(1.5,2,3), (4,5,6)], [(3,2,1), (4,5,6)]],
                 dtype = float)
```

### Initial Placeholders

>>> np.zeros((3,4)) >>> np.ones((2,3,4),dtype=np.int16)	
>>> d = np.arange(10,25,5)	Create an array of evenly
>>> np.linspace(0,2,9)	spaced values (step value) Create an array of evenly spaced values (number of samples)
>>> e = np.full((2,2),7)	Create a constant array
>>> f = np.eye(2)	Create a 2X2 identity matrix
>>> np.random.random((2,2))	Create an array with random values
>>> np.empty((3,2))	Create an empty array

## 1/0

# Saving & Loading On Disk

```
>>> np.save('my_array', a)
>>> np.savez('array.npz', a, b)
>>> np.load('my array.npy')
```

## Saving & Loading Text Files

>>>	np.loadtxt("myfile.txt")
>>>	np.genfromtxt("my file.csv", delimiter=',')
>>>	np.savetxt("mvarrav.txt", a, delimiter=" ")

# **Data Types**

>>> np.int64	Signed 64-bit integer types
>>> np.float32	Standard double-precision floating point
>>> np.complex	Complex numbers represented by 128 floats
>>> np.bool	Boolean type storing TRUE and FALSE values
>>> np.object	Python object type
>>> np.string_	Fixed-length string type
>>> np.unicode_	Fixed-length unicode type

## Inspecting Your Array

>>>	a.shape	Array dimensions
>>>	len(a)	Length of array
>>>	b.ndim	Number of array dimensions
>>>	e.size	Number of array elements
>>>	b.dtype	Data type of array elements
>>>	b.dtype.name	Name of data type
>>>	b.astvpe(int)	Convert an array to a different type

# **Asking For Help**

>>> np.info(np.ndarray.dtype)

# **Array Mathematics**

# **Arithmetic Operations**

>>> g = a - b array([[-0.5, 0., 0.],	Subtraction
[-3., -3., -3.]]) >>> np.subtract(a,b) >>> b + a array([[ 2.5, 4., 6.],	Subtraction Addition
[ 5. , 7. , 9. ]]) >>> np.add(b,a) >>> a / b array([[ 0.66666667, 1. , 1. ], [ 0.25 , 0.4 , 0.5 ]]	Addition Division
[ 0.25 , 0.4 , 0.5 ]] >>> np.divide(a,b) >>> a * b array([[ 1.5, 4., 9.],	Division Multiplication
<pre>[ 4., 10., 18.]]) &gt;&gt;&gt; np.multiply(a,b) &gt;&gt;&gt; np.exp(b) &gt;&gt;&gt; np.sqrt(b)</pre>	Multiplication Exponentiation Square root
>>> np.sin(a) >>> np.cos(b) >>> np.log(a) >>> e.dot(f)	Print sines of an array Element-wise cosine Element-wise natural logarithn Dot product
array([[ 7., 7.],	

# Comparison

<pre>&gt;&gt;&gt; a == b array([[False, True, True],</pre>	Element-wise comparison
<pre>[False, False, False]], dtype=bool) &gt;&gt;&gt; a &lt; 2 array([True, False, False], dtype=bool)</pre>	Element-wise comparison
>>> np.array equal(a, b)	Array-wise comparison

# **Aggregate Functions**

>>> a.sum()	Array-wise sum
>>> a.min()	Array-wise minimum value
>>> b.max(axis=0)	Maximum value of an array row
>>> b.cumsum(axis=1)	Cumulative sum of the elements
>>> a.mean()	Mean
>>> b.median()	Median
>>> a.corrcoef()	Correlation coefficient
>>> np.std(b)	Standard deviation

# **Copying Arrays**

>>> h = a.view()	Create a view of the array with the same data
>>> np.copy(a)	Create a copy of the array
>>> h = a.copy()	Create a deep copy of the array

# **Sorting Arrays**

>>> a.sort()	Sort an array
>>> c.sort(axis=0)	Sort the elements of an array's axis

# Subsetting, Slicing, Indexing

Subsetting

>>> a[2]

>>> b[1,2]

>>> a[0:2]

>>> b[:1]

array([1, 2])

array([ 2., 5.])

>>> b[0:2,1]

>>> a[ : :-1]

>>> a[a<2]

array([1])

**Fancy Indexing** 

array([3, 2, 1]) **Boolean Indexing** 

6.0 Slicina

# Also see Lists

### 1 2 3 Select the element at the 2nd index 1.5 2 3 Select the element at row o column 2 (equivalent to b[1][2])

Select items at index 0 and 1

Select items at rows 0 and 1 in column 1 4 5 6

Select all items at row o (equivalent to b[0:1, :]) array([[1.5, 2., 3.]]) >>> c[1,...] Same as [1,:,:] array([[[ 3., 2., 1.], [ 4., 5., 6.]]])

1 2 3

Reversed array a

Select elements from a less than 2

Select elements (1,0), (0,1), (1,2) and (0,0)

Select a subset of the matrix's rows and columns

# **Array Manipulation**

>>> b[[1, 0, 1, 0],[0, 1, 2, 0]]

>>> b[[1, 0, 1, 0]][:,[0,1,2,0]] 

array([ 4. , 2. , 6. , 1.5])

### Transposing Array >>> i = np.transpose(b) >>> i.T

### **Changing Array Shape** >>> b.ravel()

>>> g.reshape(3,-2)

# Adding/Removing Elements

>>> h.resize((2,6)) >>> np.append(h,g) >>> np.insert(a, 1, 5) >>> np.delete(a,[1])

### Combining Arrays >>> np.concatenate((a,d),axis=0)

array([ 1, 2, 3, 10, 15, 20]) >>> np.vstack((a,b)) array([[ 1. , 2. , 3. ], [ 1.5, 2. , 3. ], [ 4. , 5. , 6. ]]) >>> np.r [e,f] >>> np.hstack((e,f)) array([[ 7., 7., 1., 0.], [ 7., 7., 0., 1.]]) >>> np.column stack((a,d)) array([[ 1, 10], 2, 15], [ 3, 20]]) >>> np.c [a,d]

# **Splitting Arrays**

>>> np.hsplit(a,3) [array([1]),array([2]),array([3])] >>> np.vsplit(c,2) 

Permute array dimensions Permute array dimensions

Flatten the array Reshape, but don't change data

Return a new array with shape (2,6) Append items to an array Insert items in an array

Delete items from an array

Concatenate arrays

Stack arrays vertically (row-wise)

Stack arrays vertically (row-wise) Stack arrays horizontally (column-wise)

Create stacked column-wise arrays

Create stacked column-wise arrays

Split the array horizontally at the 3rd

Split the array vertically at the 2nd index



# **Pandas Basics**

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## **Pandas**

The **Pandas** library is built on NumPy and provides easy-to-use data structures and data analysis tools for the Python programming language. pandas !!!!!

Use the following import convention:

>>> import pandas as pd

### **Pandas Data Structures**

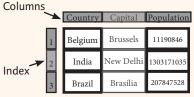
### Series

A one-dimensional labeled array capable of holding any data type



>>> s = pd.Series([3, -5, 7, 4], index=['a', 'b', 'c', 'd'])

## DataFrame



A two-dimensional labeled data structure with columns of potentially different types

```
>>> data = {'Country': ['Belgium', 'India', 'Brazil'],
           'Capital': ['Brussels', 'New Delhi', 'Brasília'],
           'Population': [11190846, 1303171035, 207847528]}
>>> df = pd.DataFrame(data,
                     columns=['Country', 'Capital', 'Population'])
```

# **Asking For Help**

>>> help(pd.Series.loc)

# Selection

Also see NumPy Arrays

# Getting

```
>>> s['b']
>>> df[1:]
   Country
             Capital Population
 1 India New Delhi 1303171035
 2 Brazil
            Brasília 207847528
```

Get one element

Get subset of a DataFrame

# Selecting, Boolean Indexing & Setting

### **By Position**

```
>>> df.iloc([0],[0])
 'Belgium'
>>> df.iat([0],[0])
 'Belgium'
```

Select single value by row & column

### **By Label**

```
>>> df.loc([0], ['Country'])
>>> df.at([0], ['Country'])
 'Belgium'
```

Select single value by row & column labels

### **By Label/Position**

>>> df.ix[2]

, , , or , Till C	1
Country	Brazil
Capital	Brasília
Population	207847528
>>> df.ix[:	,'Capital']
0 Bruss	els
1 New De	elhi
2 Brasí	.lia
>>> df.ix[1	,'Capital']
'New Delhi'	

Select single row of subset of rows

Select a single column of subset of columns

Select rows and columns

# **Boolean Indexing**

>>> s['a'] = 6

>>>	s[~(s > 1)]
>>>	s[(s < -1)   (s > 2)]
>>>	df[df['Population']>12000000
_	

Series s where value is not >1 s where value is <-1 or >2 001 Use filter to adjust DataFrame

Setting

Set index a of Series s to 6

### Read and Write to CSV

```
>>> pd.read csv('file.csv', header=None, nrows=5)
>>> pd.to csv('myDataFrame.csv')
```

### Read and Write to Excel

```
>>> pd.read excel('file.xlsx')
>>> pd.to excel('dir/myDataFrame.xlsx', sheet name='Sheet1')
 Read multiple sheets from the same file
```

### >>> xlsx = pd.ExcelFile('file.xls') >>> df = pd.read excel(xlsx, 'Sheet1')

# Read and Write to SQL Query or Database Table

```
>>> from sqlalchemy import create engine
>>> engine = create engine('sglite:///:memory:')
>>> pd.read sql("SELECT * FROM my table;", engine)
>>> pd.read sql table('my table', engine)
>>> pd.read sql query("SELECT * FROM my table;", engine)
read sql() is a convenience wrapper around read sql table() and
```

>>> pd.to sql('myDf', engine)

read sql query()

# Dropping

>>>	s.drop(['a', 'c'])	Drop values from rows (axis=0)
>>>	<pre>df.drop('Country', axis=1)</pre>	Drop values from columns(axis=1)

### **Sort & Rank**

```
>>> df.sort index()
                                        Sort by labels along an axis
>>> df.sort values(by='Country')
                                        Sort by the values along an axis
>>> df.rank()
                                        Assign ranks to entries
```

# **Retrieving Series/DataFrame Information**

### **Basic Information**

```
>>> df.shape
                             (rows,columns)
>>> df.index
                             Describe index
>>> df.columns
                             Describe DataFrame columns
>>> df.info()
                            Info on DataFrame
                            Number of non-NA values
>>> df.count()
```

### Summary

```
Sum of values
>>> df.sum()
>>> df.cumsum()
                                Cummulative sum of values
                                Minimum/maximum values
>>> df.min()/df.max()
                               Minimum/Maximum index value
>>> df.idxmin()/df.idxmax()
>>> df.describe()
                                Summary statistics
                                Mean of values
>>> df.mean()
>>> df.median()
                                Median of values
```

# **Applying Functions**

```
>>> f = lambda x: x*2
>>> df.apply(f)
                            Apply function
                            Apply function element-wise
>>> df.applymap(f)
```

# **Data Alignment**

# Internal Data Alignment

NA values are introduced in the indices that don't overlap:

```
>>> s3 = pd.Series([7, -2, 3], index=['a', 'c', 'd'])
>>> s + s3
       10.0
       NaN
       5.0
 С
       7.0
 d
```

# **Arithmetic Operations with Fill Methods**

You can also do the internal data alignment yourself with the help of the fill methods:

```
>>> s.add(s3, fill value=0)
 a 10.0
 b
     -5.0
 С
     5.0
 d
     7.0
>>> s.sub(s3, fill value=2)
>>> s.div(s3, fill value=4)
>>> s.mul(s3, fill value=3)
```



# Scikit-Learn

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### Scikit-learn

Scikit-learn is an open source Python library that implements a range of machine learning, preprocessing, cross-validation and visualization algorithms using a unified interface.



## A Basic Example

```
>>> from sklearn import neighbors, datasets, preprocessing
>>> from sklearn.model selection import train test split
>>> from sklearn.metrics import accuracy score
>>> iris = datasets.load iris()
>>> X, y = iris.data[:, :2], iris.target
>>> X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=33)
>>> scaler = preprocessing.StandardScaler().fit(X train)
>>> X train = scaler.transform(X train)
>>> X test = scaler.transform(X test)
>>> knn = neighbors.KNeighborsClassifier(n neighbors=5)
>>> knn.fit(X train, y train)
>>> y pred = knn.predict(X test)
>>> accuracy score(y test, y pred)
```

# **Loading The Data**

Also see NumPy & Pandas

Your data needs to be numeric and stored as NumPy arrays or SciPy sparse matrices. Other types that are convertible to numeric arrays, such as Pandas DataFrame, are also acceptable.

```
>>> import numpy as np
>>> X = np.random.random((10,5))
>>> X[X < 0.7] = 0
```

# Training And Test Data

```
>>> from sklearn.model_selection import train_test_split
>>> X train, X test, y train, y test = train test split(X,
                                                  random state=0)
```

# **Create Your Model**

### Supervised Learning Estimators

### Linear Regression

```
>>> from sklearn.linear model import LinearRegression
>>> lr = LinearRegression(normalize=True)
```

### Support Vector Machines (SVM)

```
>>> from sklearn.svm import SVC
>>> svc = SVC(kernel='linear')
```

### Naive Baves

>>> from sklearn.naive bayes import GaussianNB >>> gnb = GaussianNB()

### KNN

>>> from sklearn import neighbors >>> knn = neighbors.KNeighborsClassifier(n neighbors=5)

### Unsupervised Learning Estimators

### Principal Component Analysis (PCA)

>>> from sklearn.decomposition import PCA >>> pca = PCA(n components=0.95)

### K Means

>>> from sklearn.cluster import KMeans >>> k means = KMeans(n clusters=3, random state=0)

## **Model Fitting**

### Supervised learning

>>> lr.fit(X, y) >>> knn.fit(X train, y train) >>> svc.fit(X train, y train)

### Unsupervised Learning

>>> k means.fit(X train)

>>> pca model = pca.fit transform(X train) | Fit to data, then transform it

### Fit the model to the data

Fit the model to the data

# Prediction

### **Supervised Estimators**

>>> y pred = svc.predict(np.random.random((2,5))) >>> y pred = lr.predict(X test)

>>> y pred = knn.predict proba(X test)

## Unsupervised Estimators

>>> y pred = k means.predict(X test)

### Predict labels Predict labels Estimate probability of a label

## Predict labels in clustering algos

# **Preprocessing The Data**

### Standardization

- >>> from sklearn.preprocessing import StandardScaler
- >>> scaler = StandardScaler().fit(X train) >>> standardized X = scaler.transform(X train)
- >>> standardized X test = scaler.transform(X test)

### Normalization

- >>> from sklearn.preprocessing import Normalizer >>> scaler = Normalizer().fit(X train) >>> normalized X = scaler.transform(X train)
- >>> normalized X test = scaler.transform(X test)

### Binarization

- >>> from sklearn.preprocessing import Binarizer >>> binarizer = Binarizer(threshold=0.0).fit(X)
- >>> binary X = binarizer.transform(X)

# **Encoding Categorical Features**

- >>> from sklearn.preprocessing import LabelEncoder
- >>> enc = LabelEncoder()
- >>> y = enc.fit transform(y)

# Imputing Missing Values

- >>> from sklearn.preprocessing import Imputer
- >>> imp = Imputer(missing values=0, strategy='mean', axis=0) >>> imp.fit transform(X train)

# Generating Polynomial Features

- >>> from sklearn.preprocessing import PolynomialFeatures
- >>> poly = PolynomialFeatures(5)
- >>> poly.fit transform(X)

## **Evaluate Your Model's Performance**

### **Classification Metrics**

### **Accuracy Score**

- >>> knn.score(X test, y test)
- >>> from sklearn.metrics import accuracy score Metric scoring functions

Estimator score method

>>> accuracy score(y test, y pred)

### Classification Report

>>> from sklearn.metrics import classification report Precision, recall, fi-score >>> print(classification report(y test, y pred)) and support

### Confusion Matrix

>>> from sklearn.metrics import confusion matrix >>> print(confusion matrix(y test, y pred))

### Regression Metrics

### Mean Absolute Error

- >>> from sklearn.metrics import mean absolute error >>> y true = [3, -0.5, 2]
- >>> mean\_absolute\_error(y\_true, y\_pred)

# Mean Squared Error

- >>> from sklearn.metrics import mean squared error
- >>> mean squared error(y test, y pred)

- >>> from sklearn.metrics import r2 score
- >>> r2 score(y true, y\_pred)

# Clustering Metrics

### **Adjusted Rand Index**

>>> from sklearn.metrics import adjusted rand score >>> adjusted rand score(y true, y pred)

### Homogeneity

- >>> from sklearn.metrics import homogeneity score
- >>> homogeneity score(y true, y pred)

### V-measure

>>> from sklearn.metrics import v measure score >>> metrics.v measure score(y true, y pred)

### **Cross-Validation**

- >>> from sklearn.cross validation import cross val score
- >>> print(cross val score(knn, X train, y train, cv=4)) >>> print(cross val score(lr, X, y, cv=2))

# **Tune Your Model**

## **Grid Search**

- >>> from sklearn.grid search import GridSearchCV >>> params = {"n neighbors": np.arange(1,3),
- "metric": ["euclidean", "cityblock"]}
- >>> grid = GridSearchCV(estimator=knn, param grid=params)
- >>> grid.fit(X train, y train) >>> print(grid.best score )
- >>> print(grid.best\_estimator .n neighbors)

# Randomized Parameter Optimization

- >>> from sklearn.grid search import RandomizedSearchCV >>> params = {"n neighbors": range(1,5),
- - n iter=8,
  - random state=5)
  - >>> rsearch.fit(X train, y train) >>> print(rsearch.best score )



# Keras

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### Keras

Keras is a powerful and easy-to-use deep learning library for Theano and TensorFlow that provides a high-level neural networks API to develop and evaluate deep learning models.

## A Basic Example

```
>>> import numpy as np
>>> from keras.models import Sequential
>>> from keras.layers import Dense
>>> data = np.random.random((1000,100))
>>> labels = np.random.randint(2, size=(1000,1))
>>> model = Sequential()
>>> model.add(Dense(32,
                    activation='relu',
                    input dim=100))
>>> model.add(Dense(1, activation='sigmoid'))
>>> model.compile(optimizer='rmsprop',
                  loss='binary crossentropy',
                  metrics=['accuracy'])
>>> model.fit(data,labels,epochs=10,batch size=32)
>>> predictions = model.predict(data)
```

### Data

### Also see NumPy, Pandas & Scikit-Learn

Your data needs to be stored as NumPy arrays or as a list of NumPy arrays. Ideally, you split the data in training and test sets, for which you can also resort to the train test split module of sklearn.cross validation.

### Keras Data Sets

```
>>> from keras.datasets import boston_housing,
                                   cifar10,
                                   imdb
>>> (x_train,y_train),(x_test,y_test) = mnist.load data()
>>> (x train2,y train2), (x test2,y test2) = boston housing.load data()
>>> (x_train3,y_train3),(x_test3,y_test3) = cifar10.load_data()
>>> (x train4,y train4), (x test4,y test4) = imdb.load data(num words=20000)
>>> num classes = 10
```

### Other

```
>>> from urllib.request import urlopen
>>> data = np.loadtxt(urlopen("http://archive.ics.uci.edu/
ml/machine-learning-databases/pima-indians-diabetes/
pima-indians-diabetes.data"),delimiter=",")
>>> X = data[:,0:8]
>>> y = data [:,8]
```

# **Model Architecture**

# Sequential Model

```
>>> from keras.models import Sequential
>>> model = Sequential()
>>> model2 = Sequential()
>>> model3 = Sequential()
```

### Multilayer Perceptron (MLP)

### **Binary Classification**

```
>>> from keras.layers import Dense
>>> model.add(Dense(12,
                     input dim=8,
                     kernel initializer='uniform',
                     activation='relu'))
>>> model.add(Dense(8,kernel initializer='uniform',activation='relu'))
>>> model.add(Dense(1, kernel initializer='uniform', activation='sigmoid'))
Multi-Class Classification
```

```
>>> from keras.layers import Dropout
>>> model.add(Dense(512,activation='relu',input shape=(784,)))
>>> model.add(Dropout(0.2))
>>> model.add(Dense(512,activation='relu'))
>>> model.add(Dropout(0.2))
>>> model.add(Dense(10,activation='softmax'))
```

>>> model.add(Dense(64,activation='relu',input dim=train data.shape[1])) >>> model.add(Dense(1))

>>> from keras.layers import Activation,Conv2D,MaxPooling2D,Flatten

### Convolutional Neural Network (CNN)

```
>>> model2.add(Conv2D(32,(3,3),padding='same',input shape=x train.shape[1:]))
>>> model2.add(Activation('relu'))
>>> model2.add(Conv2D(32,(3,3)))
>>> model2.add(Activation('relu'))
>>> model2.add(MaxPooling2D(pool size=(2,2)))
>>> mode12.add(Dropout(0.25))
>>> model2.add(Conv2D(64,(3,3), padding='same'))
>>> model2.add(Activation('relu'))
>>> model2.add(Conv2D(64,(3, 3)))
>>> model2.add(Activation('relu'))
>>> model2.add(MaxPooling2D(pool size=(2,2)))
>>> mode12.add(Dropout(0.25))
>>> model2.add(Flatten())
>>> model2.add(Dense(512))
>>> model2.add(Activation('relu'))
>>> model2.add(Dropout(0.5))
>>> model2.add(Dense(num classes))
>>> model2.add(Activation('softmax'))
```

### Recurrent Neural Network (RNN)

```
>>> from keras.klayers import Embedding,LSTM
>>> model3.add(Embedding(20000,128))
>>> model3.add(LSTM(128,dropout=0.2,recurrent_dropout=0.2))
>>> model3.add(Dense(1,activation='sigmoid'))
```

Also see NumPy & Scikit-Learn

# Preprocessing

# Sequence Padding

```
>>> from keras.preprocessing import sequence
>>> x train4 = sequence.pad sequences(x train4, maxlen=80)
>>> x test4 = sequence.pad sequences(x test4, maxlen=80)
```

## One-Hot Encoding

```
>>> from keras.utils import to categorical
>>> Y train = to categorical(y train, num classes)
>>> Y test = to categorical(y test, num classes)
>>> Y_train3 = to_categorical(y_train3, num_classes)
>>> Y_test3 = to_categorical(y_test3, num_classes)
```

### **Train and Test Sets**

```
>>> from sklearn.model selection import train test split
>>> X train5, X test5, y train5, y test5 = train test split(X,
                                                       test size=0 33.
                                                       random state=42)
```

## Standardization/Normalization

```
>>> from sklearn.preprocessing import StandardScaler
>>> scaler = StandardScaler().fit(x train2)
>>> standardized X = scaler.transform(x train2)
>>> standardized X test = scaler.transform(x test2)
```

# Inspect Model

```
Model output shape
>>> model.output shape
>>> model.summary()
                                      Model summary representation
>>> model.get config()
                                      Model configuration
>>> model.get weights()
                                     List all weight tensors in the model
```

# **Compile Model**

```
MLP: Binary Classification
>>> model.compile(optimizer='adam',
                   loss='binary crossentropy',
                   metrics=['accuracy'])
MLP: Multi-Class Classification
>>> model.compile(optimizer='rmsprop',
                   loss='categorical crossentropy',
                   metrics=['accuracy'])
MLP: Regression
>>> model.compile(optimizer='rmsprop',
                   loss='mse',
                   metrics=['mae'])
```

optimizer='adam',

metrics=['accuracy'])

**Recurrent Neural Network** 

```
Model Training
>>> model3.fit(x train4.
             y Train4,
             batch size=32,
             epochs=15,
             verbose=1,
             validation data=(x test4, y test4))
```

# **Evaluate Your Model's Performance**

>>> model3.compile(loss='binary crossentropy',

```
>>> score = model3.evaluate(x test,
                                 y_test,
batch size=32)
```

### Prediction

```
>>> model3.predict(x test4, batch size=32)
>>> model3.predict classes(x test4,batch size=32)
```

# Save/Reload Models

```
>>> from keras.models import load model
>>> model3.save('model file.h5')
>>> my model = load model('my model.h5')
```

# **Model Fine-tuning**

## Optimization Parameters

```
>>> from keras.optimizers import RMSprop
>>> opt = RMSprop(lr=0.0001, decay=1e-6)
>>> model2.compile(loss='categorical crossentropy',
                   optimizer=opt,
                   metrics=['accuracy'])
```

## Early Stopping

```
>>> from keras.callbacks import EarlyStopping
>>> early stopping monitor = EarlyStopping(patience=2)
>>> model3.fit(x train4,
             y train4,
             batch size=32,
             epochs=15,
             validation data=(x test4, y test4),
             callbacks=[early_stopping_monitor])
```



# PySpark - RDD Basics

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# Spark

**PySpark** is the Spark Python API that exposes the Spark programming model to Python.



# **Initializing Spark**

### SparkContext

```
>>> from pyspark import SparkContext
>>> sc = SparkContext(master = 'local[2]')
```

### Inspect SparkContext

```
>>> sc.version
                                   Retrieve SparkContext version
>>> sc.pythonVer
                                   Retrieve Python version
                                   Master URL to connect to
>>> sc.master
>>> str(sc.sparkHome)
                                   Path where Spark is installed on worker nodes
                                   Retrieve name of the Spark User running
>>> str(sc.sparkUser())
                                   SparkContext
                                   Return application name
                                   Retrieve application ID
>>> sc.applicationId
                                   Return default level of parallelism
>>> sc.defaultParallelism
>>> sc.defaultMinPartitions
                                   Default minimum number of partitions for
                                   RDDs
```

## Configuration

```
>>> from pyspark import SparkConf, SparkContext
>>> conf = (SparkConf()
            .setMaster("local")
            .setAppName("My app")
            .set("spark.executor.memory", "1g"))
>>> sc = SparkContext(conf = conf)
```

# Using The Shell

In the PySpark shell, a special interpreter-aware SparkContext is already created in the variable called sc.

```
$ ./bin/spark-shell --master local[4] --py-files code.py
$ ./bin/pyspark --master local[4] --py-files code.py
```

Set which master the context connects to with the --master argument, and add Python .zip, .egg or .py files to the runtime path by passing a comma-separated list to --py-files.

# **Loading Data**

### **Parallelized Collections**

```
>>> rdd = sc.parallelize([('a',7),('a',2),('b',2)])
>>> rdd2 = sc.parallelize([('a',2),('d',1),('b',1)])
>>> rdd3 = sc.parallelize(range(100))
>>> rdd4 = sc.parallelize([("a",["x","y","z"]), ("b",["p", "r"])])
```

### External Data

Read either one text file from HDFS, a local file system or or any Hadoop-supported file system URI with textFile(), or read in a directory of text files with wholeTextFiles().

```
>>> textFile = sc.textFile("/my/directory/*.txt")
>>> textFile2 = sc.wholeTextFiles("/my/directory/")
```

# **Retrieving RDD Information**

### **Basic Information**

```
>>> rdd.getNumPartitions()
>>> rdd.count()
>>> rdd.countByKey()
defaultdict(<type 'int'>, {'a':2,'b':1})
>>> rdd.countByValue()
defaultdict(<type 'int'>, {('b',2):1,('a',2):1,('a',7):1}
>>> rdd.collectAsMap()
 {'a': 2,'b': 2}
>>> rdd3.sum()
4950
>>> sc.parallelize([]).isEmpty()
```

List the number of partitions Count RDD instances

Count RDD instances by key

Count RDD instances by value

Return (key,value) pairs as a dictionary Sum of RDD elements

Check whether RDD is empty

### Summary

```
>>> rdd3.max()
>>> rdd3.min()
>>> rdd3.mean()
 49 5
>>> rdd3.stdev()
 28.866070047722118
>>> rdd3.variance()
 833.25
>>> rdd3.histogram(3)
 ([0,33,66,99],[33,33,34])
>>> rdd3.stats()
```

Maximum value of RDD elements

Minimum value of RDD elements Mean value of RDD elements

Standard deviation of RDD elements

Compute variance of RDD elements

Compute histogram by bins

Summary statistics (count, mean, stdev, max &

# **Applying Functions**

```
>>> rdd.map(lambda x: x+(x[1],x[0]))
        .collect()
  [('a',7,7,'a'),('a',2,2,'a'),('b',2,2,'b')]
\Rightarrow rdd5 = rdd.flatMap(lambda x: x+(x[1],x[0]))
>>> rdd5.collect()
  ['a',7,7,'a','a',2,2,'a','b',2,2,'b']
>>> rdd4.flatMapValues(lambda x: x)
  [('a', 'x'), ('a', 'y'), ('a', 'z'), ('b', 'p'), ('b', 'r')]
```

Apply a function to each RDD element Apply a function to each RDD element

Apply a flatMap function to each (key,value) pair of rdd4 without changing the keys

and flatten the result

# **Selecting Data**

# Getting

```
>>> rdd.collect()
 [('a', 7), ('a', 2), ('b', 2)]
>>> rdd.take(2)
 [('a', 7), ('a', 2)]
>>> rdd.first()
 ('a', 7)
>>> rdd.top(2)
 [('b', 2), ('a', 7)]
>>> rdd3.sample(False, 0.15, 81).collect()
```

Return a list with all RDD elements

Take first 2 RDD elements

Take first RDD element

Take top 2 RDD elements

[3,4,27,31,40,41,42,43,60,76,79,80,86,97]

### Filtering

>>> rdd.filter(lambda x: "a" in x) .collect() [('a',7),('a',2)] >>> rdd5.distinct().collect() ['a',2,'b',7] >>> rdd.keys().collect() ['a', 'a', 'b']

Return sampled subset of rdd3

## Filter the RDD

Return distinct RDD values Return (key, value) RDD's keys

# Iterating

```
>>> def g(x): print(x)
>>> rdd.foreach(g)
                                            Apply a function to all RDD elements
   ('a', 7)
   ('b', 2)
   ('a', 2)
```

# **Reshaping Data**

```
>>> rdd.reduceByKey(lambda x,y : x+y)
      .collect()
 [('a',9),('b',2)]
>>> rdd.reduce(lambda a, b: a + b)
 ('a',7,'a',2,'b',2)
```

Merge the rdd values

each kev

Merge the rdd values for

Return RDD of grouped values

### Grouping by

```
>>> rdd3.groupBy(lambda x: x % 2)
        .mapValues(list)
        .collect()
>>> rdd.groupByKey()
      .mapValues(list)
      .collect()
```

[('a',[7,2]),('b',[2])]

>>> rdd.foldByKey(0, add)

.collect()

>>> rdd3.keyBy(lambda x: x+x)

.collect()

[('a',9),('b',2)]

Group rdd by key

### Aggregating

4950

```
>>> seqOp = (lambda x, y: (x[0]+y, x[1]+1))
>>> combOp = (lambda x, y: (x[0]+y[0], x[1]+y[1]))
>>> rdd3.aggregate((0,0),seqOp,combOp)
  (4950,100)
>>> rdd.aggregateByKey((0,0),seqop,combop)
       .collect()
 [('a', (9,2)), ('b', (2,1))]
>>> rdd3.fold(0,add)
```

Aggregate RDD elements of each partition and then the results Aggregate values of each RDD key

Aggregate the elements of each partition, and then the results Merge the values for each key

> Create tuples of RDD elements by applying a function

# **Mathematical Operations**

```
>>> rdd.subtract(rdd2)
                                        Return each rdd value not contained
        .collect()
                                        in rdd2
  [('b',2),('a',7)]
>>> rdd2.subtractByKey(rdd)
                                        Return each (key,value) pair of rdd2
         .collect()
                                        with no matching key in rdd
 [('d', 1)]
>>> rdd.cartesian(rdd2).collect(
```

Return the Cartesian product of rdd and rdd2

## Sort

```
>>> rdd2.sortBy(lambda x: x[1])
                                          Sort RDD by given function
         .collect()
  [('d',1),('b',1),('a',2)]
>>> rdd2.sortByKey()
                                          Sort (key, value) RDD by key
         .collect()
  [('a',2),('b',1),('d',1)]
```

# Repartitioning

		New RDD with 4 partitions Decrease the number of partitions in the RDD to 1
///	Idd.Coalesce(I)	Decrease the number of partitions in the RDD to 1

# Saving

```
>>> rdd.saveAsTextFile("rdd.txt")
>>> rdd.saveAsHadoopFile("hdfs://namenodehost/parent/child",
                           'org.apache.hadoop.mapred.TextOutputFormat')
```

# Stopping SparkContext

>>> sc.stop()

# **Execution**

\$ ./bin/spark-submit examples/src/main/python/pi.py



# **Python For Data Science** Cheat Sheet Matplotlib

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# Matplotlib

Matplotlib is a Python 2D plotting library which produces publication-quality figures in a variety of hardcopy formats and interactive environments across platforms.



# Prepare The Data

```
>>> import numpy as np
>>> x = np.linspace(0, 10, 100)
>>> v = np.cos(x)
>>> z = np.sin(x)
```

### 2D Data or Images

```
>>> data = 2 * np.random.random((10, 10))
>>> data2 = 3 * np.random.random((10, 10))
>>> Y, X = np.mgrid[-3:3:100j, -3:3:100j]
>>> U = -1 - X**2 + Y
>>> V = 1 + X - Y**2
>>> from matplotlib.cbook import get sample data
>>> img = np.load(get sample data('axes grid/bivariate normal.npy'))
```

# Create Plot

```
>>> import matplotlib.pyplot as plt
```

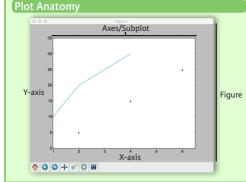
```
>>> fig = plt.figure()
>>> fig2 = plt.figure(figsize=plt.figaspect(2.0))
```

### Axes

All plotting is done with respect to an Axes. In most cases, a subplot will fit your needs. A subplot is an axes on a grid system.

```
>>> fig.add axes()
>>> ax1 = fig.add subplot(221) # row-col-num
>>> ax3 = fig.add subplot(212)
>>> fig3, axes = plt.subplots(nrows=2,ncols=2)
>>> fig4, axes2 = plt.subplots(ncols=3)
```

### Plot Anatomy & Workflow



### Workflow

```
The basic steps to creating plots with matplotlib are:
       1 Prepare data 2 Create plot 3 Plot 4 Customize plot 5 Save plot 6 Show plot
                >>> import matplotlib.pyplot as plt
                >>> x = [1,2,3,4]
               >>> y = [10, 20, 25, 30]
                >>> fig = plt.figure() < Step 2
                >>> ax = fig.add subplot(111) < Step 3
                >>> ax.plot(x, y, color='lightblue', linewidth=3) Step 3, 4
                >>> ax.scatter([2,4,6],
                                [5, 15, 25],
                                color='darkgreen',
                                marker='^')
               >>> ax.set xlim(1, 6.5)
```

# Customize Plot

### Colors, Color Bars & Color Maps

```
>>> plt.plot(x, x, x, x**2, x, x**3)
>>> ax.plot(x, y, alpha = 0.4)
>>> ax.plot(x, y, c='k')
>>> fig.colorbar(im, orientation='horizontal')
>>> im = ax.imshow(img,
                   cmap='seismic')
```

### Markers

>>>	fig, ax = plt.subplots()
>>>	<pre>ax.scatter(x,y,marker=".")</pre>
>>>	ax.plot(x, v, marker="o")

```
>>> plt.plot(x,y,linewidth=4.0)
>>> plt.plot(x,y,ls='solid')
>>> plt.plot(x,y,ls='--')
>>> plt.plot(x,y,'--',x**2,y**2,'-.')
>>> plt.setp(lines,color='r',linewidth=4.0)
```

### Text & Annotations

```
>>> ax.text(1,
            -2.1,
            'Example Graph',
           style='italic')
>>> ax.annotate("Sine",
                 xy = (8, 0),
                 xycoords='data'
                 xytext = (10.5, 0),
                 textcoords='data',
                 arrowprops=dict(arrowstyle="->",
                              connectionstyle="arc3"),)
```

### Mathtext

```
Limits, Legends & Layouts
```

>>> plt.show()

>>> plt.savefig('foo.png')

>>> plt.title(r'\$sigma i=15\$', fontsize=20)

```
Limits & Autoscaling
```

>>> ax.margins(x=0.0,y=0.1)

```
>>> ax.axis('equal')
                                                            Set the aspect ratio of the plot to 1
>>> ax.set(xlim=[0,10.5],ylim=[-1.5,1.5])
                                                            Set limits for x-and v-axis
>>> ax.set xlim(0,10.5)
                                                            Set limits for x-axis
 Leaends
                                                            Set a title and x-and y-axis labels
>>> ax.set(title='An Example Axes',
             vlabel='Y-Axis',
             xlabel='X-Axis')
>>> ax.legend(loc='best')
                                                            No overlapping plot elements
```

### Manually set x-ticks >>> ax.xaxis.set(ticks=range(1,5), ticklabels=[3,100,-12,"foo"]) Make y-ticks longer and go in and out

>>> ax.tick params(axis='y', direction='inout', length=10)

### Subplot Spacing

```
>>> fig3.subplots adjust(wspace=0.5,
                         hspace=0.3,
                         left=0.125,
                         right=0.9,
                         top=0.9,
>>> fig.tight layout()
Axis Spines
```

# Adjust the spacing between subplots

Add padding to a plot

bottom=0.1)

>>>	ax1.spines['top'].set visible(False)
>>>	<ul> <li>ax1.spines['bottom'].set position(('outward',10)</li> </ul>

**Save Plot** 

Save figures

>>> plt.savefig('foo.png')

>>> plt.savefig('foo.png', transparent=True)

Save transparent figures

# Fit subplot(s) in to the figure area

### Make the top axis line for a plot invisible ) ) Move the bottom axis line outward

# Plotting Routines

```
>>> lines = ax.plot(x, y)
>>> ax.scatter(x,y)
>>> axes[0,0].bar([1,2,3],[3,4,5])
>>> axes[1,0].barh([0.5,1,2.5],[0,1,2])
>>> axes[1,1].axhline(0.45)
>>> axes[0,1].axvline(0.65)
>>> ax.fill(x, y, color='blue')
>>> ax.fill between(x,y,color='yellow')
```

Draw points with lines or markers connecting them Draw unconnected points, scaled or colored Plot vertical rectangles (constant width) Plot horiontal rectangles (constant height) Draw a horizontal line across axes Draw a vertical line across axes

Draw filled polygons

Fill between y-values and o

### **Vector Fields**

>>>	axes[0,1].arrow(0,0,0.5,0.5)
>>>	axes[1,1].quiver(y,z)
>>>	axes[0,1].streamplot(X,Y,U,V)

Add an arrow to the axes Plot a 2D field of arrows Plot 2D vector fields

### Data Distributions

>>>	ax1.hist(y)
>>>	ax3.boxplot(y)
>>>	ax3.violinplot(z)

Plot a histogram Make a box and whisker plot Make a violin plot

# Close & Clear

>>>	plt.cla()
>>>	plt.clf()
>>>	nlt close()

Show Plot

>>> plt.show()

Clear an axis Clear the entire figure Close a window

### 2D Data or Images >>> fig, ax = plt.subplots()

>>>	im =	ax.imshow(img,
		cmap='gist earth',
		interpolation='nearest
		vmin=-2,
		vmax=2)

Colormapped or RGB arrays

>>> axes2[0].pcolor(data2) >>> axes2[0].pcolormesh(data) >>> CS = plt.contour(Y,X,U) >>> axes2[2].contourf(data1) >>> axes2[2]= ax.clabel(CS)

Pseudocolor plot of 2D array Pseudocolor plot of 2D array Plot contours Plot filled contours Label a contour plot



# Bokeh

Learn Bokeh Interactively at <a href="https://www.DataCamp.com">www.DataCamp.com</a>, taught by Bryan Van de Ven, core contributor



# **Plotting With Bokeh**

The Python interactive visualization library **Bokeh** enables high-performance visual presentation of large datasets in modern web browsers.



Bokeh's mid-level general purpose bokeh.plotting interface is centered around two main components: data and glyphs.



The basic steps to creating plots with the bokeh.plotting interface are:

1. Prepare some data:

Python lists, NumPy arrays, Pandas DataFrames and other sequences of values

- 2. Create a new plot
- 3. Add renderers for your data, with visual customizations
- 4. Specify where to generate the output
- 5. Show or save the results

# 1 ) Data

### Also see Lists, NumPy & Pandas

Under the hood, your data is converted to Column Data Sources. You can also do this manually:

# 2) Plotting

# **3** Renderers & Visual Customizations

```
Glyphs
```

### 

### Rows & Columns Layout

```
Rows
>>> from bokeh.layouts import row
>>> layout = row(p1,p2,p3)

Nesting Rows & Columns
>>>layout = row(column(p1,p2), p3)
Columns
>>> layout = column(p1,p2,p3)

Nesting Rows & Columns
>>>layout = row(column(p1,p2), p3)
```

## Linked Plots

## Linked Plots

```
Linked Axes
>>> p2.x_range = p1.x_range
>>> p2.y_range = p1.y_range
```

Linked Brushing

```
>>> p4 = figure(plot_width = 100, tools='box_select,lasso_select')
>>> p4.circle('mpg', 'cyl', source=cds_df)
>>> p5 = figure(plot_width = 200, tools='box_select,lasso_select')
>>> p5.circle('mpg', 'hp', source=cds_df)
>>> layout = row(p4,p5)
```

**Customized Glyphs** 

Hover Glyphs

Colormapping

>>> p3.add tools(hover)

Selection and Non-Selection Glyphs

>>> p.circle('mpg', 'cyl', source=cds df,

>>> color mapper = CategoricalColorMapper(

>>> p3.circle('mpg', 'cyl', source=cds df,

selection color='red',

nonselection alpha=0.1)

>>> hover = HoverTool(tooltips=None, mode='vline')

color=dict(field='origin',

factors=['US', 'Asia', 'Europe'],

palette=['blue', 'red', 'green'])

transform=color mapper),

legend='Origin'))

>>> p = figure(tools='box select')

### Legends

### Legend Location

Grid Lavout

>>> row2 = [p3]

>>> row1 = [p1,p2]

Tabbed Lavout

```
Inside Plot Area
>>> p.legend.location = 'bottom_left'
```

>>> from bokeh.layouts import gridplot

>>> layout = gridplot([[p1,p2],[p3]])

>>> tab1 = Panel(child=p1, title="tab1")

>>> tab2 = Panel(child=p2, title="tab2")

>>> layout = Tabs(tabs=[tab1, tab2])

>>> from bokeh.models.widgets import Panel, Tabs

Outside Plot Area
>>> r1 = p2.asterisk(np.array([1,2,3]), np.array([3,2,1])
>>> r2 = p2.line([1,2,3,4], [3,4,5,6])

>>> r2 = p2.line([1,2,3,4], [3,4,5,6]) >>> legend = Legend(items=[("One", [p1, r1]),("Two", [r2])], location=(0, -30)) >>> p.add\_layout(legend, 'right')

# Legend Orientation

```
>>> p.legend.orientation = "horizontal"
>>> p.legend.orientation = "vertical"
```

### Legend Background & Border

```
>>> p.legend.border_line_color = "navy"
>>> p.legend.background_fill_color = "white"
```

# 1) Output

### **Output to HTML File**

- >>> from bokeh.io import output\_file, show
- >>> output\_file('my\_bar\_chart.html', mode='cdn')

### **Notebook Output**

- >>> from bokeh.io import output\_notebook, show
- >>> output\_notebook()

# Embedding

### Standalone HTML

- >>> from bokeh.embed import file\_html
  >>> html = file html(p, CDN, "my plot")
- Components
- >>> from bokeh.embed import components
- >>> script, div = components(p)

# 5) Show or Save Your Plots

>>> show(p1)	>>> save(p1)
>>> show(layout)	>>> save(layout)

# Statistical Charts With Bokeh

# Bokeh's high-level bokeh.charts interface is ideal for quickly creating statistical charts

### Bar Chart



>>> from bokeh.charts import Bar
>>> p = Bar(df, stacked=True, palette=['red','blue'])

# Box Plot



>>> from bokeh.charts import BoxPlot

>>> p = BoxPlot(df, values='vals', label='cyl', legend='bottom\_right')

# Histogram



>>> from bokeh.charts import Histogram
>>> p = Histogram(df, title='Histogram')

### Scatter Plot



# DataCamp Learn Python for Data Science Interactively



Also see Data

# Python For Data Science Cheat Sheet (3) Plotting With Seaborn

Seaborn

Learn Data Science Interactively at www.DataCamp.com



### Statistical Data Visualization With Seaborn

The Python visualization library Seaborn is based on matplotlib and provides a high-level interface for drawing attractive statistical graphics.

Make use of the following aliases to import the libraries:

```
>>> import matplotlib.pyplot as plt
>>> import seaborn as sns
```

The basic steps to creating plots with Seaborn are:

- 1. Prepare some data
- 2. Control figure aesthetics
- 3. Plot with Seaborn
- 4. Further customize your plot

```
>>> import matplotlib.pyplot as plt
>>> import seaborn as sns
>>> tips = sns.load dataset("tips")
                                        Step 1
>>> sns.set style("whitegrid")
>>> g = sns.lmplot(x="tip",
                                        Step 3
                   v="total bill",
                   data=tips,
                   aspect=2)
>>> g = (g.set axis labels("Tip", "Total bill(USD)").
set(xlim=(0,10),ylim=(0,100))
>>> plt.title("title")
>>> plt.show(q)
```

# Data

### Also see Lists, NumPy & Pandas

```
>>> import pandas as pd
>>> import numpy as np
>>> uniform data = np.random.rand(10, 12)
>>> data = pd.DataFrame({'x':np.arange(1,101),
                          y':np.random.normal(0,4,100)})
```

### Seaborn also offers built-in data sets:

>>> sns.axes style("whitegrid")

```
>>> titanic = sns.load dataset("titanic")
>>> iris = sns.load dataset("iris")
```

### **Axis Grids**

```
>>> g = sns.FacetGrid(titanic,
                      col="survived",
                       row="sex")
>>> q = q.map(plt.hist, "age")
>>> sns.factorplot(x="pclass",
                   y="survived",
                   hue="sex",
                   data=titanic)
>>> sns.lmplot(x="sepal width",
               y="sepal length",
               hue="species",
               data=iris)
```

Subplot grid for plotting conditional relationships

Draw a categorical plot onto a Facetgrid

Violin plot

Plot data and regression model fits across a FacetGrid

```
>>> h = sns.PairGrid(iris)
                                         Subplot grid for plotting pairwise
>>> h = h.map(plt.scatter)
                                         relationships
>>> sns.pairplot(iris)
                                         Plot pairwise bivariate distributions
>>> i = sns.JointGrid(x="x",
                                         Grid for bivariate plot with marginal
                        y="y",
                                         univariate plots
                        data=data)
>>> i = i.plot(sns.regplot,
                 sns.distplot)
                                         Plot bivariate distribution
>>> sns.jointplot("sepal length"
                     "sepal width",
                    data=iris,
```

kind='kde')

### Categorical Plots

```
Scatterplot
                                                  Scatterplot with one
>>> sns.stripplot(x="species",
                                                  categorical variable
                    y="petal length",
                    data=iris)
>>> sns.swarmplot(x="species",
                                                  Categorical scatterplot with
                                                  non-overlapping points
                    y="petal length",
                    data=iris)
Bar Chart
                                                  Show point estimates and
>>> sns.barplot(x="sex",
                                                  confidence intervals with
                y="survived",
                hue="class",
                                                  scatterplot glyphs
                data=titanic)
Count Plot
                                                  Show count of observations
>>> sns.countplot(x="deck",
                  data=titanic,
                  palette="Greens d")
Point Plot
                                                  Show point estimates and
>>> sns.pointplot(x="class",
                                                  confidence intervals as
                    v="survived",
                                                  rectangular bars
                    hue="sex",
                    data=titanic,
                    palette={"male":"q",
                              "female": "m" },
                    markers=["^","o"],
                    linestyles=["-","--"])
Boxplot
>>> sns.boxplot(x="alive",
                                                  Boxplot
                 v="age",
                 hue="adult male",
                 data=titanic)
>>> sns.boxplot(data=iris,orient="h")
                                                  Boxplot with wide-form data
```

### **Regression Plots**

```
Plot data and a linear regression
>>> sns.regplot(x="sepal width",
                                         model fit
                  v="sepal length",
                  data=iris,
```

### **Distribution Plots**

```
>>> plot = sns.distplot(data.y,
                                         Plot univariate distribution
                           kde=False,
                           color="b")
```

### **Matrix Plots**

>>> sns.heatmap(uniform data, vmin=0, vmax=1) Heatmap

# **Further Customizations**

### **Axisarid Objects**

>>> g.despine(left=True)	Remove left spine
>>> g.set ylabels("Survived")	Set the labels of the y-axis
>>> g.set xticklabels(rotation=45)	Set the tick labels for x
>>> g.set_axis_labels("Survived",	Set the axis labels
"Sex")	
>>> h.set(xlim=(0,5), ylim=(0,5), xticks=[0,2.5,5],	Set the limit and ticks of the x-and y-axis
yticks=[0,2.5,5])	

### Plot

>>> plt.title("A Title")	Add plot title
>>> plt.ylabel("Survived")	Adjust the label of the y-axis
>>> plt.xlabel("Sex")	Adjust the label of the x-axis
>>> plt.ylim(0,100)	Adjust the limits of the y-axis
>>> plt.xlim(0,10)	Adjust the limits of the x-axis
>>> plt.setp(ax,yticks=[0,5])	Adjust a plot property
>>> plt.tight_layout()	Adjust subplot params

# Figure Aesthetics

Return a dict of params or use with

with to temporarily set the style

### Seaborn styles (Re)set the seaborn default >>> sns.set() Set the matplotlib parameters >>> sns.set style("whitegrid") Set the matplotlib parameters >>> sns.set style("ticks", {"xtick.major.size":8,

"vtick.major.size":8})

>>> f, ax = plt.subplots(figsize=(5,6)) Create a figure and one subplot

# Context Functions

>>> sns.violinplot(x="age",

y="sex", hue="survived",

data=titanic)

**Violinplot** 

<pre>sns.set_context("talk") sns.set_context("notebook",</pre>	Set context to "talk" Set context to "notebook' scale font elements and override param mapping

### **Color Palette**

>>>	<pre>sns.set palette("hus1",3)</pre>	Define the color palette
>>>	sns.color_palette("husl")	Use with with to temporarily set palette
>>>	flatui = ["#9b59b6","#3498db",	"#95a5a6","#e74c3c","#34495e","#2ecc71"]
>>>	sns.set_palette(flatui)	Set your own color palette

# Show or Save Plot

>>>	plt.show()
>>>	plt.savefig("foo.png")
>>>	plt.savefig("foo.png",
	transparent=True)

Show the plot Save the plot as a figure Save transparent figure

# Close & Clear

|--|

