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

### **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



#### **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

-	iciai i lacciforaci s	
>>>	np.zeros((3,4))	Create an array of zeros
>>>	np.ones((2,3,4),dtype=np.int16)	Create an array of ones
>>>	d = np.arange(10, 25, 5)	Create an array of evenly
		spaced values (step value)
>>>	np.linspace(0,2,9)	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("myarray.txt", a, delimiter=" ")
```

#### Data Types

١	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

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

### Asking For Help

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

#### Array Mathematics

#### **Arithmetic Operations**

>>> g = a - b	Subtraction
array([[-0.5, 0. , 0. ], [-3. , -3. , -3. ]])	
>>> np.subtract(a,b)	Subtraction
>>> b + a	Addition
array([[ 2.5, 4. , 6. ], [ 5. , 7. , 9. ]])	
>>> np.add(b,a)	Addition
>>> a / b	Division
array([[ 0.66666667, 1. , 1. ], [ 0.25 , 0.4 , 0.5 ]]	)
>>> np.divide(a,b)	Division
>>> a * b	Multiplication
array([[ 1.5, 4., 9.], [ 4., 10., 18.]])	
>>> np.multiply(a,b)	Multiplication
>>> np.exp(b)	Exponentiation
>>> np.sqrt(b)	Square root
>>> np.sin(a)	Print sines of an array
>>> np.cos(b)	Element-wise cosine
>>> np.log(a)	Element-wise natural logarithn
>>> e.dot(f) array([[ 7., 7.],	Dot product
[ 7 7.11)	1

#### Comparison

>>> a == b array([[False, True, True],	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

riggicgate i anetions	
>>> 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
>>> nn etd(h)	Standard deviation

### **Copying Arrays**

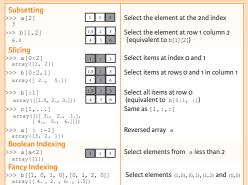
>>> np.copy(a)	Create a view of the array with the same data Create a copy of the array Create a deep copy of the array
>>> n = a.copy()	Create a deep copy or the array

### Sorting Arrays

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

### Subsetting, Slicing, Indexing

#### Also see List



### **Array Manipulation**

Transposing Array	
>>> i = np.transpose(b)	Permute arra
>>> i.T	Permute arra
Changing Array Shape	

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

## Adding/Removing Elements

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

#### **Combining Arrays**

array([ 1, 2, 3, 10, 15, 20]
>>> np.vstack((a,b))
array([[ 1. , 2. , 3. ],
>>> 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]

Л	Splitting Arrays
	>>> np.hsplit(a,3)
	[array([1]),array([2]),array([3])]
	>>> np.vsplit(c,2)
	[array([[[ 1.5, 2., 1. ], [ 4., 5., 6. ]]]),
	array([[[ 3., 2., 3.],
	[ 4., 5., 6.]]])]

rmute array dimensions rmute array dimensions

Select a subset of the matrix's rows and columns

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

Stack arrays vertically (row-wise)

>>> np.concatenate((a,d),axis=0) Concatenate arrays

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 index Split the array vertically at the 2nd index



# **Data Wrangling**

with pandas **Cheat Sheet** http://pandas.pydata.org

# **Syntax** – Creating DataFrames

	2	5	8	11		
	3	6	9	12		
df = pd	DataF	rame(				
	{"a	" : [	4,5,	6],		
	"t	)" : [	7, 8,	9],		
	"(	:":[	10, 1	l, 12]	},	
	index	= [1	, 2, 3	3])		
Specify v	alues fo	or each	column			
df = pd	.DataF	rame(				
[[4	1, 7,	10],				
[ ]	5, 8,	11],				
Īέ	5. 9.	12]],				
index=[1, 2, 3],						
columns=['a', 'b', 'c'])						
Specify values for each row.						
		а	b	С		
		а		•		
	n v					

		2	5	8	11	
	е	2	6	9	12	
df = pd.DataFrame(						
		"a"				
		"b"	: [7	, 8,	9],	
		"c"	-	-	-	
index =	pd.	Mult:	iInd	ex.f	rom_t	tuples(
	[	('d'	,1),	('d'	,2),	('e',2)],
		nai	mes=	['n'	,'v'	])))
Create D	ataFr	ame v	vith a	Multi	iIndex	

# **Method Chaining**

Most pandas methods return a DataFrame so that another pandas method can be applied to the result. This improves readability of code.

```
df = (pd.melt(df)
        .rename(columns={
                 'variable' : 'var',
                 'value' : 'val'})
        .query('val >= 200')
```

# **Data** – A foundation for wrangling in pandas





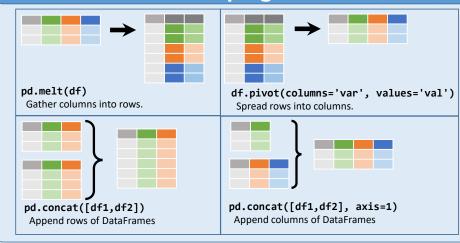


Tidy data complements pandas's vectorized operations, pandas will automatically preserve observations as you manipulate variables. No other format works as intuitively with pandas.



#### Each variable is saved Each observation is in its own column saved in its own row

# Reshaping Data – Change the layout of a data set



- df.sort values('mpg')
- Order rows by values of a column (low to high).
- df.sort values('mpg',ascending=False) Order rows by values of a column (high to low).
- df.rename(columns = {'y':'year'}) Rename the columns of a DataFrame
- df.sort index()
- Sort the index of a DataFrame
- df.reset index()

Reset index of DataFrame to row numbers, moving index to columns.

df.drop(columns=['Length', 'Height']) Drop columns from DataFrame

# **Subset Observations** (Rows)

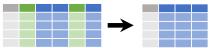


- df[df.Length > 7]
  - Extract rows that meet logical criteria.
- df.drop\_duplicates() Remove duplicate rows (only considers columns).
- df.head(n) Select first n rows.
- df.tail(n) Select last n rows.

- df.sample(frac=0.5)
  - Randomly select fraction of rows.
- df.sample(n=10)
- Randomly select n rows.
- df.iloc[10:20]
  - Select rows by position.
- df.nlargest(n, 'value') Select and order top n entries.
- df.nsmallest(n, 'value')
  - Select and order bottom n entries.

Logic in Python (and pandas)			
<	Less than	!=	Not equal to
>	Greater than	df.column.isin(values)	Group membership
==	Equals	pd.isnull( <i>obj</i> )	Is NaN
<=	Less than or equals	pd.notnull( <i>obj</i> )	Is not NaN
>=	Greater than or equals	&, ,~,^,df.any(),df.all()	Logical and, or, not, xor, any, all

# **Subset Variables** (Columns)



- df[['width','length','species']]
  - Select multiple columns with specific names.
- df['width'] or df.width
  - Select single column with specific name.
- df.filter(regex='reaex')
  - Select columns whose name matches regular expression regex.

regex (Regular Expressions) Examples		
'\.'	Matches strings containing a period '.'	
'Length\$'	Matches strings ending with word 'Length'	
'^Sepal'	Matches strings beginning with the word 'Sepal'	
'^x[1-5]\$'	Matches strings beginning with 'x' and ending with 1,2,3,4,5	
''^(?!Species\$).*'	Matches strings except the string 'Species'	

- df.loc[:,'x2':'x4']
  - Select all columns between x2 and x4 (inclusive).
- df.iloc[:,[1,2,5]]
  - Select columns in positions 1, 2 and 5 (first column is 0).
- df.loc[df['a'] > 10, ['a','c']]
  - Select rows meeting logical condition, and only the specific columns .

http://pandas.pydata.org/ This cheat sheet inspired by Rstudio Data Wrangling Cheatsheet (https://www.rstudio.com/wp-content/uploads/2015/02/data-wrangling-cheatsheet.pdf) Written by Irv Lustig, Princeton Consultants

# **Summarize Data**

df['w'].value\_counts()

Count number of rows with each unique value of variable

len(df)

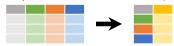
# of rows in DataFrame.

df['w'].nunique()

# of distinct values in a column.

df.describe()

Basic descriptive statistics for each column (or GroupBy)



pandas provides a large set of summary functions that operate on different kinds of pandas objects (DataFrame columns, Series, GroupBy, Expanding and Rolling (see below)) and produce single values for each of the groups. When applied to a DataFrame, the result is returned as a pandas Series for each column. Examples:

sum()

Sum values of each object.

count()

Count non-NA/null values of each object.

median()

Median value of each object.

quantile([0.25,0.75])

Quantiles of each object. apply(function)

Apply function to each object.

min()

Minimum value in each object. max()

Maximum value in each object. mean()

Mean value of each object.

var() Variance of each object.

std()

Standard deviation of each object.

# **Handling Missing Data**

df.dropna()

Drop rows with any column having NA/null data.

df.fillna(value)

Replace all NA/null data with value.

# Make New Columns



df.assign(Area=lambda df: df.Length\*df.Height) Compute and append one or more new columns.

df['Volume'] = df.Length\*df.Height\*df.Depth Add single column.

pd.qcut(df.col, n, labels=False) Bin column into n buckets.



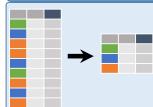
pandas provides a large set of vector functions that operate on all columns of a DataFrame or a single selected column (a pandas Series). These functions produce vectors of values for each of the columns, or a single Series for the individual Series. Examples:

max(axis=1) Element-wise max. min(axis=1) Element-wise min.

clip(lower=-10,upper=10) abs()

Trim values at input thresholds Absolute value.

# **Group Data**



df.groupby(by="col")

Return a GroupBy object, grouped by values in column named "col".

df.groupby(level="ind")

Return a GroupBy object, grouped by values in index level named "ind".

All of the summary functions listed above can be applied to a group. Additional GroupBy functions:

Windows

size()

Size of each group.

agg(function)

Aggregate group using function.

The examples below can also be applied to groups. In this case, the function is applied on a per-group basis, and the returned vectors are of the length of the original DataFrame.

shift(1)

Copy with values shifted by 1. rank(method='dense')

Ranks with no gaps.

rank(method='min')

Ranks. Ties get min rank.

rank(pct=True)

Ranks rescaled to interval [0, 1].

rank(method='first') Ranks. Ties go to first value. shift(-1)

Copy with values lagged by 1.

cumsum()

Cumulative sum.

cummax()

Cumulative max.

cummin()

Cumulative min.

cumprod()

Cumulative product.

# **Plotting**

## df.expanding()

Return an Expanding object allowing summary functions to be applied cumulatively.

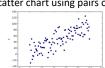
df.rolling(n)

Return a Rolling object allowing summary functions to be applied to windows of length n.

df.plot.hist()

Histogram for each column

df.plot.scatter(x='w',v='h') Scatter chart using pairs of points



# **Combine Data Sets**

adf bdf x1 x2 x1 x3 A 1 A T B 2 C 3 D T

#### Standard Joins

A 1

B 2

x1 x2

C 3

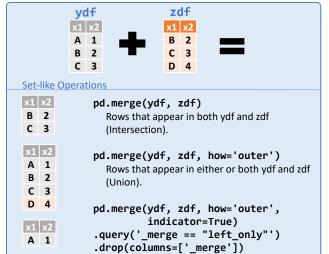
x1 x2 x3 pd.merge(adf, bdf, A 1 T how='left', on='x1') 2 F Join matching rows from bdf to adf. 3 NaN x3 pd.merge(adf, bdf, A 1.0 T how='right', on='x1') B 2.0 Join matching rows from adf to bdf. D NaN T pd.merge(adf, bdf, how='inner', on='x1') Join data. Retain only rows in both sets. pd.merge(adf, bdf, how='outer', on='x1') 2 Join data. Retain all values, all rows. C 3 NaN D NaN T Filtering Joins x1 x2 adf[adf.x1.isin(bdf.x1)]

All rows in adf that have a match in bdf.

All rows in adf that do not have a match in bdf.

Rows that appear in ydf but not zdf (Setdiff).

adf[~adf.x1.isin(bdf.x1)]



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

#### Seaborn

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#### 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")
>>> sns.set style("whitegrid") Step 2
>>> g = sns.lmplot(x="tip", y="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(a)
```

### ) Data

>>> sns.set()

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

**つ)Figure Aesthetics** 

>>> titanic = sns.load\_dataset("titanic")
>>> iris = sns.load\_dataset("iris")

### **Axis Grids**

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

Subplot grid for plotting conditional relationships

Draw a categorical plot onto a Facetgrid

Plot data and regression model fits across a FacetGrid

>>> i = sns.JointGrid(x="x",

Categorical scatterplot with

non-overlapping points

Show point estimates and

Show count of observations

Show point estimates and

confidence intervals as

rectangular bars

Boxplot

Violin plot

confidence intervals with

scatterplot glyphs

#### >>> h = sns.PairGrid(iris) >>> h = h.map(plt.scatter) >>> sns.pairplot(iris)

>>> i = i.plot(sns.regplot, sns.distplot) 

Subplot grid for plotting pairwise relationships Plot pairwise bivariate distributions Grid for bivariate plot with marginal univariate plots

Plot bivariate distribution

data=iris.

#### Categorical Plots

#### Scatterplot >>> sns.stripplot(x="species", y="petal length", Scatterplot with one categorical variable data=iris)

y="petal length". data=iris) **Bar Chart** 

data=iris)

>>> sns.barplot(x="sex", y="survived", hue="class"

>>> sns.swarmplot(x="species",

#### data=titanic) **Count Plot**

>>> sns.countplot(x="deck". data=titanic, palette="Greens d")

Point Plot >>> sns.pointplot(x="class", y="survived". hue="sex",

data=titanic, markers=["^","o"], linestyles=["-","--"])

Boxplot >>> sns.boxplot(x="alive", y="age",

hue="adult male", data=titanic) >>> sns.boxplot(data=iris.orient="h") Violinplot

>>> sns.violinplot(x="age", y="sex", hue="survived",

# Regression Plots

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

kind='kde')

#### Distribution Plots

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

### Matrix Plots

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

# **Further Customizations**

Axisgrid (	Objects	
	ine(left=True)	Remove left spine
	ylabels("Survived")	Set the labels of the y-axis
	kticklabels(rotation=45)	
>>> g.set_a	axis_labels("Survived",	Set the axis labels
	"Sex")	
>>> h.set()		Set the limit and ticks of the
7	ylim=(0,5),	x-and y-axis
2	kticks=[0,2.5,5],	

>>> 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 Adjust the limits of the y-axis >>> plt.ylim(0,100) Adjust the limits of the x-axis >>> plt.xlim(0,10) >>> plt.setp(ax,yticks=[0,5]) Adjust a plot property >>> plt.tight layout() Adjust subplot params

yticks=[0,2.5,5])

Boxplot with wide-form data

data=titanic)

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

### Seaborn styles (Re)set the seaborn default

>>> sns.set style("whitegrid") >>> sns.set style("ticks", {"xtick.major.size":8, "ytick.major.size":8} >>> sns.axes\_style("whitegrid")

Set the matplotlib parameters Set the matplotlib parameters

Return a dict of params or use with with to temporarily set the style

#### Context Functions >>> sns.set context("talk") Set context to "talk" >> sns.set context("notebook",

Set context to "notebo scale font elements and font\_scale=1.5, rc={"lines.linewidth":2.5}) scale font elements and override param mapping

>>> sns.set\_palette("hus1",3)
>>> sns.color\_palette("hus1")
>>> flatui = ["#9b59b6","#3498db","#9 >>> sns.set palette(flatui)

Define the color palette Use with with to temporarily set palette Set your own color palette

### 5) Show or Save Plot

# >>> pit.snow() >>> plt.savefig("foo.png") >>> plt.savefig("foo.png", transparent=True)

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

#### Close & Clear

### >>> plt.cla() >>> plt.clf() >>> plt.close() Clear an axis Clear an entire figure Close a window



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

#### 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, v train)
>>> y_pred = knn.predict(X_test)
 >>> accuracy score(v test, v pred)
```

#### Loading The Data

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))
```

#### 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)
```

## 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)

#### 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 Bayes**

>>> from sklearn.naive\_bayes import GaussianNB >>> gnb = GaussianNR()

#### 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)

>>> 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**

**Encoding Categorical Features** 

Generating Polynomial Features

>>> poly = PolynomialFeatures(5)

>>> poly.fit transform(X)

>>> enc = LabelEncoder()

>>> y = enc.fit transform(y)

Imputing Missing Values

y\_pred = svc.predict(np.random.random((2,5))) Predict labels pred = lr.predict(X test) y pred = knn.predict proba(X\_test)

**Unsupervised Estimators** 

>> y\_pred = k\_means.predict(X test)

>>> from sklearn.preprocessing import LabelEncoder

from sklearn.preprocessing import Imputer

>>> imp = Imputer(missing\_values=0, strategy='mean', axis=0)
>>> imp.fit transform(X train)

>>> from sklearn.preprocessing import PolynomialFeatures

Predict labels Estimate probability of a label

Predict labels in clustering algos

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

#### **Classification Metrics**

#### **Accuracy Score**

>> knn.score(X test, y test) Estimator score method >>> from sklearn.metrics import accuracy\_score Metricscoring functions >>> accuracy\_score(y\_test, y\_pred)

### **Classification Report**

>>> from sklearn.metrics import classification\_report >>> 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

>>> 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 

n\_iter=8, random\_state=5)

>>> rsearch.fit(X train, y\_train)
>>> print(rsearch.best score)





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 matplotlib platforms.

### 1) Prepare The Data

```
1D Data
 >>> import numpy as np
>>> x = np.linspace(0, 10, 100)

>>> y = 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'))

### 2 Create Plot

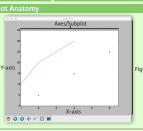
>>> import matplotlib.pyplot as plt

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

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



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] Step 1 >>> y = [10,20,25,30] >>> 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) >>> plt.savefig('foo.png' >>> plt.show()

### 4) 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')
```

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

```
>>> plt.plot(x,y,linewidth=4.0)
>>> pit.plot(x,y,!serwith=1.0)
>>> pit.plot(x,y,!ser'solid')
>>> pit.plot(x,y,!ser'--')
>>> pit.plot(x,y,'--',x**2,y**2,'-.')
>>> pit.setp(lines,color='r',linewidth=4.0)
```

>>> plt.title(r'\$sigma\_i=15\$', fontsize=20) Limits, Legends & Layouts

Limits & Autoscaling
>>> ax.margins(x=0.0,y=0.1) Add padding to a plot >>> ax.axis('equal')
>>> ax.set(xlim=[0,10.5],ylim=[-1.5,1.5]) Set the aspect ratio of the plot to 1 Set limits for x-and y-axis >>> ax.set\_xlim(0,10.5) Set limits for x-axis

Legends ax.set(title='An Example Axes', ylabel='Y-Axis', xlabel='X-Axis')

Ticks

length=10) Subplot Spacing >>> fig3.subplots\_adjust(wspace=0.5, Adjust the spacing between subplots hspace=0.3,

left=0.125, right=0.9, top=0.9, bottom=0.1) >>> fig.tight layout()

**Axis Spines** Axis spines

>>> ax1.spines['top'].set\_visible(False)

>>> ax1.spines['bottom'].set\_position(('outward',10))

Make the top axis line for a plot inv
Move the bottom axis line outward

Fit subplot(s) in to the figure area Make the top axis line for a plot invisible

Set a title and x-and y-axis labels

No overlapping plot elements

Manually set x-ticks Make y-ticks longer and go in and out

## 3) 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[1,0].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

>>	axes[0,1].arrow(0,0,0.5,0.5)	Add an arrow to the axes
>>	axes[1,1].quiver(y,z)	Plot a 2D field of arrows
>>	axes[0,1].streamplot(X,Y,U,V)	Plot 2D vector fields

>>> 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)

Label a contour plot

Pseudocolor plot of 2D array Pseudocolor plot of 2D array Plot contours Plot filled contours

### 5) Save Plot

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

#### ら)Show Plot >>> plt.show()

### Close & Clear

Clear an axis Clear the entire figure >>> plt.cla() >>> plt.clf() >>> plt.close() Close a window



