

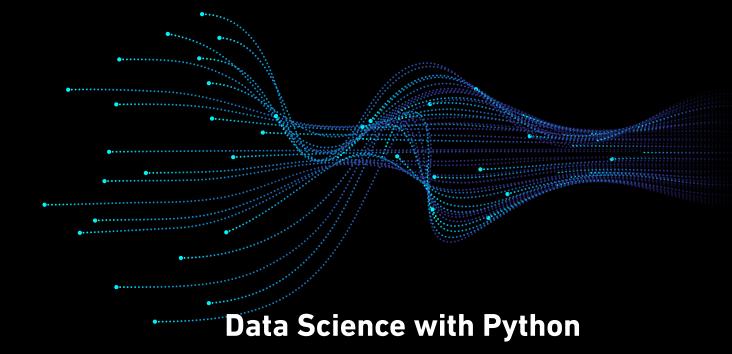
# Cheat Sheets for Al

Neural Networks,
Machine Learning,
DeepLearning &
Big Data

The Most Complete List of Best Al Cheat Sheets

BecomingHuman.Al

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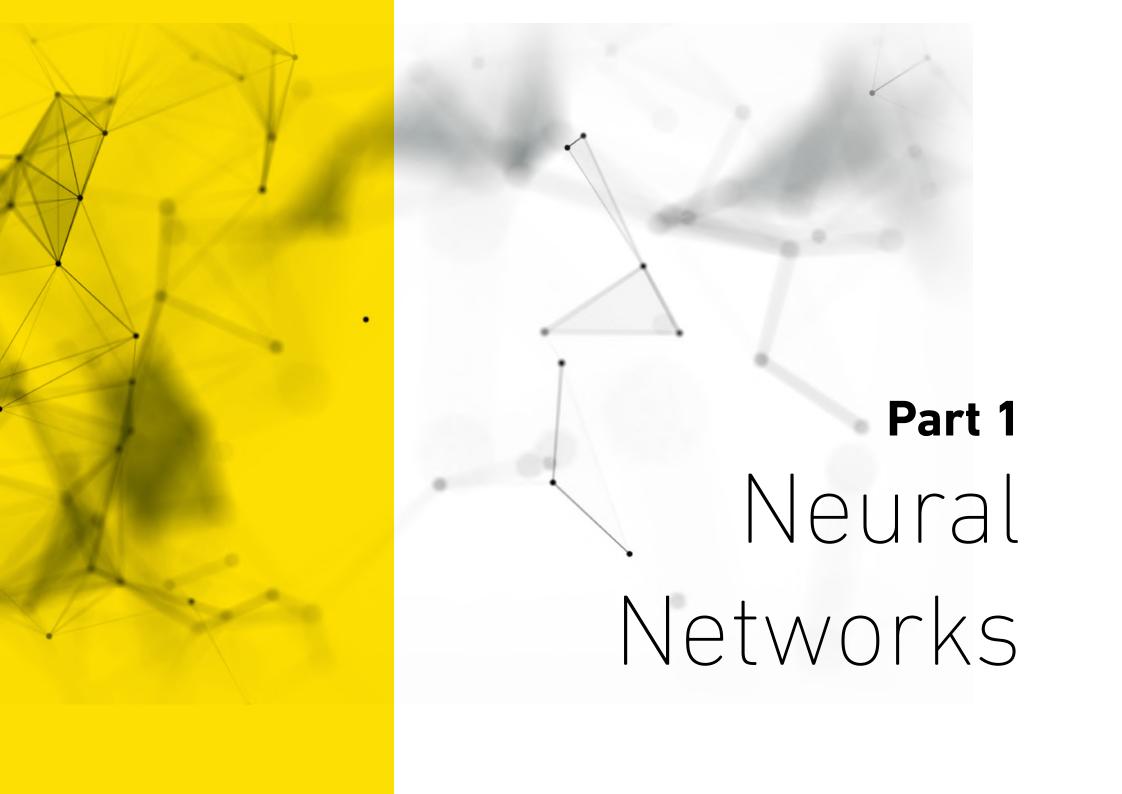
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# Neural Networks Basic

**Cheat Sheet** 

# **BecomingHuman.Al**

Backfed Input Cell

Noisy Input Cell

Spiking Hidden Cell

Hidden Cell

Output Cell

Recurrent Cell

Memory Cell

Kernel

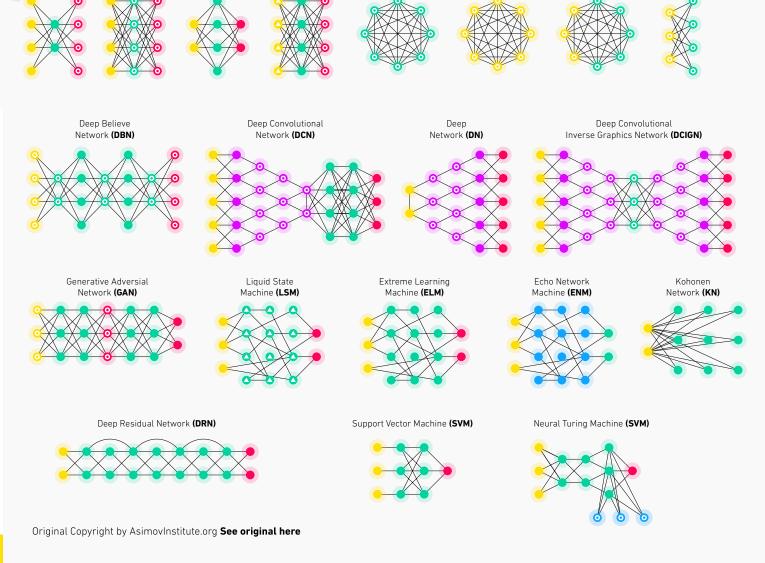
Different Memory Cell

Convolutional or Pool

Input Cell

Index

# Probablisticc Hidden Cell Match Input Output Cell



Long / Short Term

Memory (LSTM)

Hopfield

Network (HN)

Gated Recurrent

Unit (GRU)

Boltzman

Machine (BM)

Restricted

BM (RBM)

Radial Basis

Network (RBF)

Sparse

AE (SAE)

Deep Feed

Forward (DFF)

Denoising

AE (DAE)

Recurrent Neural

Network (RNN)

Markov

Chain (MC)

Feed

Forward (FF)

Variational

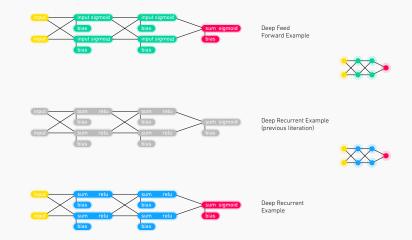
AE (VAE)

Perceptron (P)

Encorder (AE)

# Neural Networks Graphs Cheat Sheet

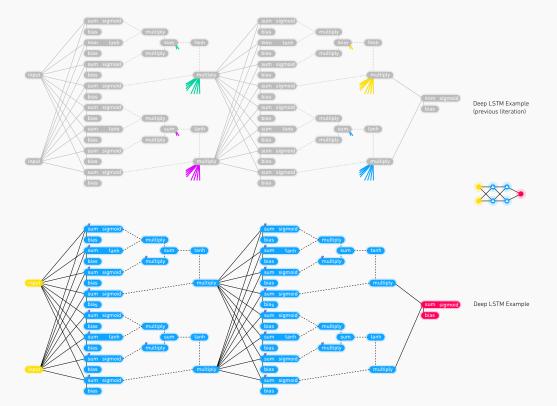
**BecomingHuman.Al** 

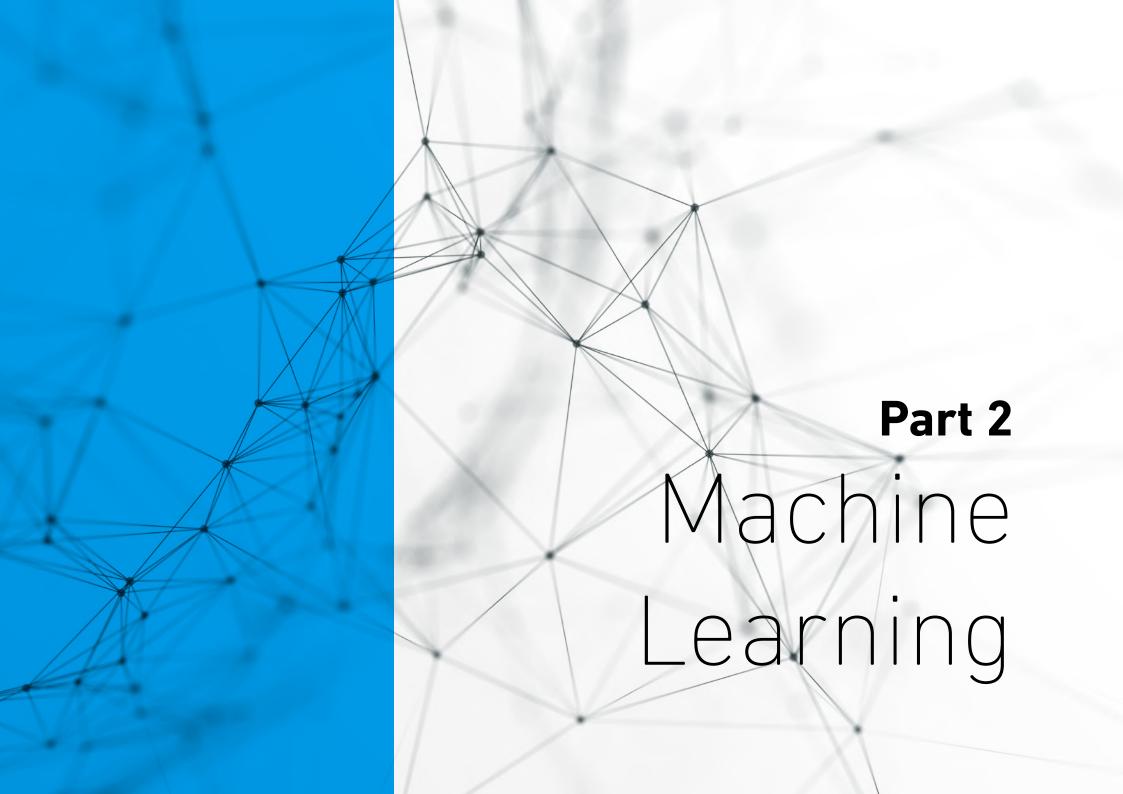












# MachineLearning Overview **MACHINE LEARNING IN EMOJI**

# **BecomingHuman.Al**







### **BASIC REGRESSION**





linear model.LinearRegression() Lots of numerical data





human builds model based on input / output

human input, machine output human utilizes if satisfactory

human input, machine output human reward/punish, cycle continues

# **CLUSTER ANALYSIS**





Similar datum into groups based on centroids



covariance.EllipticalEnvelope()







Finding outliers through grouping

# **CLASSIFICATION**





neural network.MLPClassifier()

Complex relationships. Prone to overfitting Basically magic.





neighbors.KNeighborsClassifier()

Group membership based on proximity





tree.DecisionTreeClassifier()

If/then/else. Non-contiguous data. Can also be regression.





ensemble.RandomForestClassifier()

Find best split randomly Can also be regression





svm.SVC() svm.LinearSVC()

Maximum margin classifier. Fundamental Data Science algorithm





GaussianNB() MultinominalNB() BernoulliNB

Updating knowledge step by step with new info

# **FEATURE REDUCTION**

#### **T-DISTRIB STOCHASTIC NEIB EMBEDDING**



manifold.TSNE()

Visual high dimensional data. Convert similarity to joint probabilities

#### **PRINCIPLE COMPONENT ANALYSIS**



decomposition.PCA()

Distill feature space into components that describe greatest variance

#### **CANONICAL CORRELATION ANALYSIS**



decomposition.CCA()

Making sense of cross-correlation matrices

#### LINEAR **DISCRIMINANT ANALYSIS**





Linear combination of features that separates classes

# **OTHER IMPORTANT CONCEPTS**

**BIAS VARIANCE TRADEOFF** 

**UNDERFITTING / OVERFITTING** 

**INERTIA** 

**ACCURACY FUNCTION** 

PRECISION FUNCTION

SPECIFICITY FUNCTION

SENSITIVITY FUNCTION

# **Cheat-Sheet Skicit learn** Phyton For Data Science

BecomingHuman.Al DataCamp



# **Skicit Learn**

Skicit Learn is an open source Phyton library that implements a range if machine learning, processing, cross validation and visualization algorithm using a unified

#### A basic Example

- >>> from sklearn import neighbors, datasets, preprocessing
- >>> from sklearn.cross validation import train test split
- >>> from sklearn.metrics import accuracy score
- >>> iris = datasets.load \_iris() >>> X, y = iris.data[:, :2], iris.target
- >>> Xtrain, X test, y\_train, y test = train\_test\_split (X, y, random stat33)
- >>> 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)

# **Prediction**

#### Supervised Estimators

>>> y\_pred = svc.predict(np.random.radom((2,5))) >>> v pred = lr.predict(X test) >>> y\_pred = knn.predict\_proba(X\_test)

Unsupervised Estimators

Predict labels

Predict lahels

Predict labels in clustering algos

# **Loading the Data**

Your data beeds to be nmueric and stored as NumPy arrays or SciPy sparse matric, other types that they are comvertible to numeric arrays, such as Pandas Dataframe, are also

>>> import numpy as np >> X = np.random.random((10,5)) >>> y = np . array ( PH', IM', 'F', 'F' , 'M', 'F', 'NI', 'tvl' , 'F', 'F', 'F' )) >>> X [X < 0.7] = 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)

#### **Encoding Categorical Features**

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

#### 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

- >>> from sklearn.metrics import accuracy\_score
- >>> accuracy\_score(y\_test, y\_pred)

#### Classification Report

>>> from sklearn.metrics import classification\_report >>> print(classification\_report(y\_test, y\_pred))

#### Confusion Matrix

- >>> from sklearn.metrics import confusion matrix
- >>> print(confusion matrix(v test. v 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)

#### R<sup>2</sup> Score

>>> from sklearn.metrics import r2 score >>> r2 score(y true, y pred)

# **Clustering Metrics**

>>> 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))

# **Model Fitting**

#### Supervised learning

- >>> lr.fit(X, y) >>> knn.fit(X\_train, y\_train)
- >>> svc.fit(X train. v train)

#### Unsupervised Learning

>>> pca\_model = pca.fit\_transform(X\_train)

Fit the model to the data Fit to data, then transform it

Fit the model to the data

# **Create Your Model**

#### **Supervised Learning Estimators**

#### Linear Regression

>>> from sklearn.linear\_model import LinearRegression >>> Ir = 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()

Estimator score method

Precision recall f1-score

and support

- >>> 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)

# **Training And Test Data**

- >> from sklearn.cross validation import train\_test\_split
- >> X train, X test, y train, y test train\_test\_split(X,
  - - random state-0)

# **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, v 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),
- "weights": ["uniform", "distance"]}

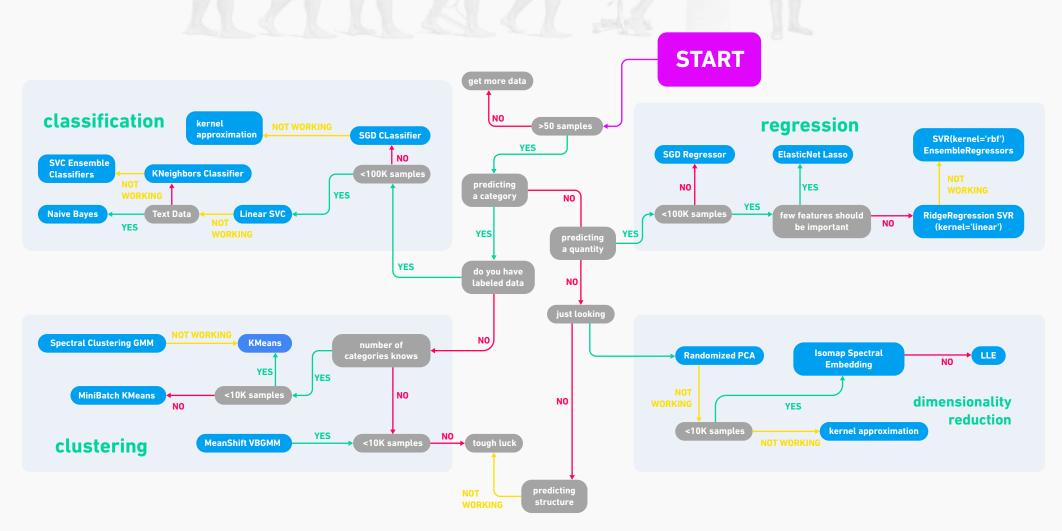
>>> rsearch = RandomizedSearchCV(estimator=knn, param distributions=params,

> n\_iter=8, random state=5)

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

# **Skicit-learn Algorithm**

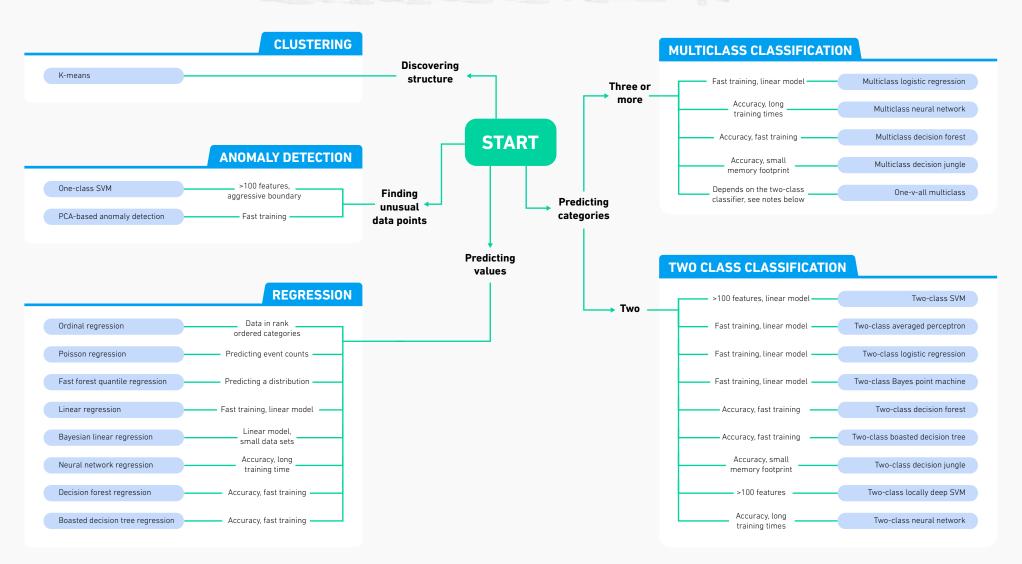
# **BecomingHuman.Al**

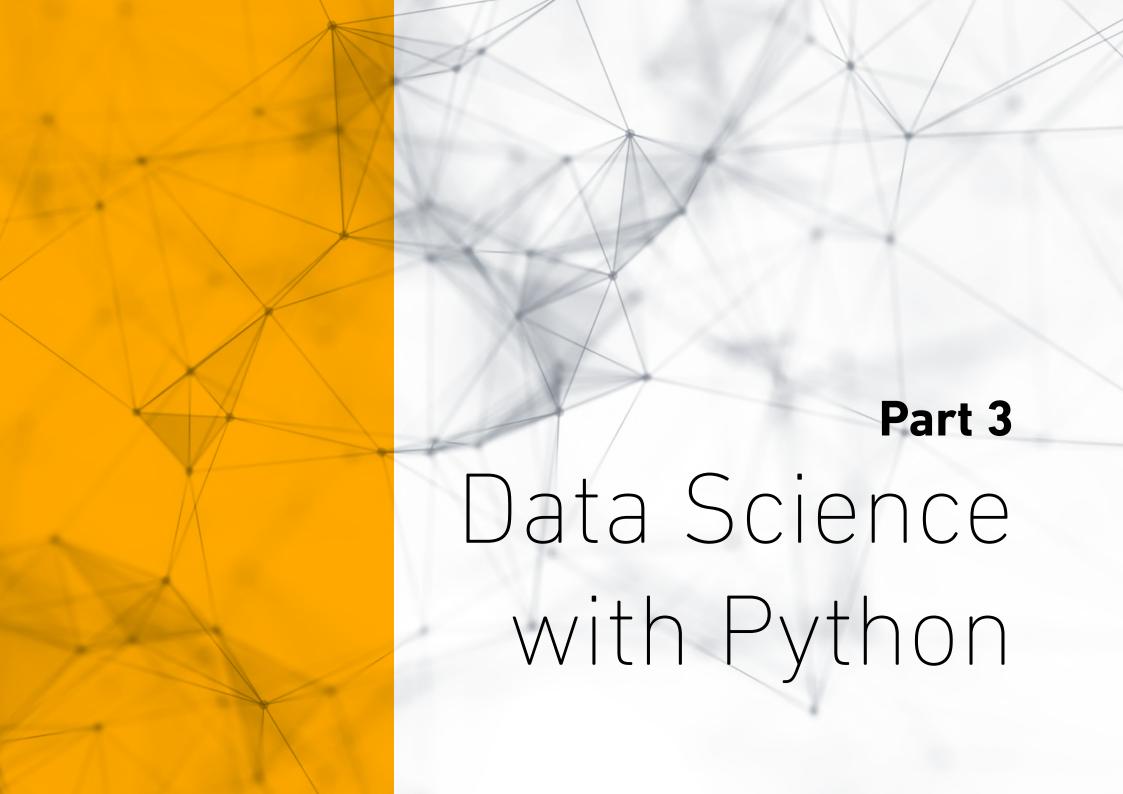


# **Algorithm** Cheat Sheet

# **BecomingHuman.Al**

This cheat sheet helps you choose the best Azure Machine Learning Studio algorithm for your predictive analytics solution. Your decision is driven by both the nature of your data and the question you're trying to answer.





# **Tensor Flow** Cheat Sheet

# BecomingHuman.Al

# Installation

# How to install new package in Python

Example: pip install requests

#### How to install tensorflow?

python\_version = cp27/cp34

sudo pip install

https://storage.googleapis.com/tensorflow/linux/\$device/ten-

#### How to install Skflow

Helpers

Get object type

str(object)

variables.

local variables

dir( huiltin ) Other built-in functions

signatures and so on)

#### How to install Keras

update ~/.keras/keras.json - replace "theano" by "tensorflow"

Python helper Important functions

Get help for object (list of available methods, attributes,

Return the dictionary containing the current scope's global

Update and return a dictionary containing the current scope's

Return the identity of an object. This is guaranteed to be unique

Get list of object attributes (fields, functions)

Transform an object to string object?

Shows documentations about the object

among simultaneously existing objects.

# Info

#### **TensorFlow**

to 11.5 petaflops.

TensorFlow™ is an open source software library created by Google for numerical computation and large scale computation. Tensorflow bundles together Machine Learning, Deep learning models and frameworks and makes them useful by way of common metaphor.

In May 2017 Google

second-generation of

announced the

TensorFlow the TPU, as well as

the availability of the TPUs in

up to 180 teraflops of perfor-

Google Compute Engine.[12] The

second-generation TPUs deliver

mance, and when organized into

clusters of 64 TPUs provide up

#### Keras

#### Keras is an open sourced neural networks library, written in

Python and is built for fast experimentation via deep neural networks and modular design. It is capable of running on top of TensorFlow, Theano, Microsoft Cognitive Toolkit, or PlaidML.

Scikit Flow is a high level interface base on tensorflow which can be used like sklearn. You can build you own model on your own data quickly without rewriting extra code provides a set of high level model classes that you can use to easily integrate with your existing Scikit-learn pipeline code.

#### Some useful functions

tf.get default graph()

Tensor Flow

Main classes

tf.reset\_default\_graph()

ops.reset\_default\_graph()

tf.convert to tensor(value)

#### TensorFlow Optimizers

AdadeltaOptimizer

AdamOptimizer

RMSPropOptimizer

#### Reduction

reduce\_max

reduce\_all

accumulate r

#### **Activation functions**

relu

relu6 elu

softplus

softsign

dropout

bias\_add

sigmoid

tanh

sigmoid\_cross\_entropy\_with\_logits

softmax

log softmax

softmax cross entropy with logits sparse\_softmax\_cross\_entropy\_with\_logits

weighted\_cross\_entropy\_with\_logits

# Skflow

#### Main classes

TensorFlowClassifier

TensorFlowRegressor

TensorFlowDNNRegressor

#### Each classifier and regressor have following fields n\_classes=0 (Regressor), n\_classes are expected to be input (Classifier)

TensorFlowRNNClassifier - there is 50

learning rate=0.1.

#### Each class has a method fit

fit(X, v, monitor=None, loadir=None

X: matrix or tensor of shape [n samples, n features...]. Can be iterator that returns arrays of features. The training input samples for fitting the model.

Y: vector or matrix [n\_samples] or [n\_samples, n\_outputs]. Can be iterator that returns array of targets. The training target values (class labels in classification, real numbers in

monitor: Monitor object to print training progress and invoke early stopping

logdir: the directory to save the log file that can be used for optional visualization.

predict (X, axis=1, batch\_size=None)

X: array-like matrix, [n\_samples, n\_features...] or iterator. axis: Which axis to argmax for classification.

By default axis 1 (next after batch) is used. Use 2 for sequence

hatch size: If test set is too hig use hatch size to solit it into mini batches. By default the batch\_size member variable is

y: array of shape [n\_samples]. The predicted classes or

# Phyton For Data Science

# **Cheat-Sheet Phyton Basic**

# BecomingHuman.Al



# **Variables and Data Types**

#### Variable Assignment

>>> x=5 >>> x

#### **Calculations With Variables**

>>> x+2 Sum of two variables Subtraction of two variables >>> x\*2 Multiplication of two variables >>> x\*\*7 Exponentiation of a variable 25 >>> x%2 Remainder of a variable >>> x/float(2 Division of a variable 2.5

#### **Calculations With Variables**

str() '5', '3,45', 'True' Variables to strings int() 5. 3. 1 Variables to integers float() 5.0, 1.0 Variables to floats hool() True True True Variables to booleans

# **Asking For Help**

>>> help(str)

# Lists

Subset

>>> my\_list[1]

#### Also see NumPy Arrays

>>> h = 'nice' >>> my list = ['my', 'list', a, b] >>> my list2 = [[4,5,6,7], [3,4,5,6]]

#### **Selecting List Elements**

# Select item at index 1

my list[list][itemOfList]

>>> my\_list[-3] Select 3rd last item Slice >>> my\_list[1:3] Select items at index 1 and 2 Select items after index 0 >>> my\_list[1:] >>> my\_list[:3] Salact items before index 3 >>> my\_list[:] Copy my\_list

Subset Lists of Lists

>>> my list2[1][0] >>> my list2[1][:2]

#### **List Operations**

>>> mv list + mv 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**

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

# **Numpy Arrays**

#### Also see Lists

>>> my\_list = [1, 2, 3, 4] >>> my\_array = np.array(my\_list) >>> mv 2darrav = np.array([[1,2,3],[4.5.6]])

### **Selecting Numpy Array Elements**

#### Index starts at 0

Subset

>>> my\_array[1]

Select item at index 1

Slice

>>> my\_array[0:2]

Select items at index 0 and 1

Subset 2D Numpy arrays

>>> my\_2darray[:,0]

my 2darray[rows, columns]

#### **Numpy Array Operations**

>>> my\_array > 3 array([Fa >>> my\_array \* 2 >>> my\_array + np.array([5, 6, 7, 8])
array([6, 8, 10, 12])

#### Numpy Array Operations

>>> 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(mv arrav) Mean of the array >>> np.median(my\_array) Median of the array >>> my\_array.corrcoef() Correlation coefficient Standard deviation >>> np.std(my\_array)

# Strings

Also see NumPy Arrays

>>> my\_string = 'thisStringIsAwesome' >>> my\_string

#### String Operations

>>> my\_string \* 2 >>> my\_string + 'Innit' >>> 'm' in my\_string

#### String Operations

Index starts at

>>> my\_string[3] >>> my string[4:9]

#### String Methods

>>> my\_string.upper() String to uppercase >>> my\_string.lower() String to lowercase Count String elements >>> my\_string.count('w') >>> my\_string.replace('e', 'i') Replace String elements >>> my\_string.strip() Strip whitespaces

# Libraries

#### Import libraries

>>> import numpy >>> import numpy as np Selective import >>> from math import pi

# **Install Python**



Leading open data science platform powered by Pytho



Free IDE that is included with Anaconda



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

# Python For Data Science Cheat Sheet

# **PySpark - RDD Basics**

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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]')

#### **Calculations With Variables**

Retrieve SparkContext version >>> sc.pythonVer Retrieve Python version Path where Spark is installed on >>> str(sc.sparkHome) Retrieve name of the Snark User >>> str(sc.sparkUser()) running SparkContext >>> sc.appName Return application name >>> sc.applicationId Retrieve application ID >>> sc.defaultParallelism Return default level of parallelism >>> 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", "1q")) >>> sc = SparkContext(conf = conf)

# Configuration

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

\$ ./bin/spark-shell --master local[2]

\$./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 whole Text Files().

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

# **Selecting Data**

### Getting

>>> rdd.collect() Return a list with all RDD elements [('a', 7), ('a', 2), ('b', 2)] >>> rdd.take(2) Take first 2 RDD elements >>> rdd.first() Take first RDD element >>> rdd.top(2) Take top 2 RDD elements Sampling

>>> rdd3.sample(False, 0.15, 81),collect() [3,4,27,31,40,41,42,43,60,76,79,80,86,97

Return sampled subset of rdd3

#### Filtering

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

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

# Iterating

#### Getting

>>> def g(x): print(x) >>> rdd.foreach(g)

# Retrieving RDD Information

#### **Basic Information**

>>> rdd.getNumPartitions() List the number of partitions >>> rdd.count() Count RDD instances >>> rdd.countByKey()
defaultdict(<type 'int'>,{'a':2,'b':1}) Count RDD instances by key >>> rdd.countByValue() Count RDD instances defaultdict(<type 'int'>,{('b',2):1,('a',2):1,('a',7):1}) >>> rdd.collectAsMap() Return (key,value) pairs as a >>> rdd3.sum() Sum of RDD elements Sum of RDD elements >>> sc.parallelize([]).isEmpty() Check whether RDD is empty

#### Summary

>>> rdd3.max() Maximum value of RDD elements >>> rdd3.min() Minimum value of RDD elements Mean value of RDD elements >>> rdd3.mean() >>> rdd3.stdev() 28.866070047722118 Standard deviation of RDD elements >>> rdd3.variance() 833.25 Compute variance of RDD elements >>> rdd3.histogram(3) ([0,33,66,99],[33,33,34]) Compute histogram by bins >>> rdd3.stats() Summary statistics (count, mean,

# **Reshaping Data**

#### Reducing

>>> rdd.reduceByKey(lambda x,y : x+y) .collect() each key [('a',9),('b',2)] Merge the rdd values for >>> rdd.reduce(lambda a, b: a + b)
('a',7,'a',2,'b',2) Merge the rdd values

#### Grouping by

>>> rdd3.groupBy(lambda x: x % 2) .mapValues(list) Return RDD of grouped values .collect() >>> rdd.groupByKey() Group rdd by key .mapValues(list) .collect() [('a',[7,2]),('b',[2])]

#### Aggregating

>>> rdd foldBvKev(f) add)

.collect()

>>> segOp = (lambda x.v; (x[0]+v,x[1]+1)) Aggregate RDD elements of each partition and then the results >>> comb0p = (lambda x,y:(x[0]+y[0],x[1]+y[1]))Aggregate values of each RDD key >>> rdd3.aggregate((0,0),seqOp,combOp) Aggregate the elements of each 4950 partition, and then the results >>> rdd.aggregateByKey((0,0),seqop,combop) .collect() [('a',(9,2)), ('b',(2,1))] Merge the values for each key >>> rdd3.fold(0.add)

[('a',9),('b'.2)] >>> rdd3.kevBv(lambda x: x+x) Create tunies of RDD elements by collect() applying a function

# **Applying Functions**

.collect() [('a',x'),('a',y'),('a',z'),('b',p'),('b',r')]

>>> rdd.map(lambda x: x+(x[1],x[0])) Apply a function to each RDD element .collect() [('a',7,7,'a'),('a',2,2,'a'),('b',2,2,b')] >>> rdd5 = rdd.flatMap(lambda x: Apply a function to each RDD x+(x[1],x[0])ent and flatten the result >>> rdd5.collect()
['a',7,7,'a',a',2,2,'a',b',2,2,'b']

>>> rdd4.flatMapValues(lambda x: x) Apply a flatMap function to each (kev.value)pair of rdd4 without

of rdd and rdd2

# **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() [('d', 1)] with no matching key in rdd >>> rdd.cartesian(rdd2).collect() Return the Cartesian product

# Sort

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

# **Reshaping Data**

>>> rdd.repartition(4) New RDD with 4 partitions >>> rdd.coalesce(1) Decrease the number of partitions in the

# 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

# NumPy Basics Cheat Sheet

# BecomingHuman.Al





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.

1D array







# **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))	Create an array of zero
>>> np.ones((2,3,4),dtype=np.int16)	Create an array of one
>>> d = np.arange(10,25,5)	Create an array of evenly space values (step valu
>>> np.linspace(0,2,9)	Create an array of even spaced values (number of sample
>>> e = np.full((2,2),7)	Create a constant arra
>>> f = np.eye(2)	Create a 2X2 identity matr
>>> np.random.random((2,2))	Create an array with random value
>>> np.empty((3,2))	Create an empty arra

# 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=" ")

# **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.astype(int)	Convert an array to a different type

# **Data Types**

	A TOTAL CONTRACTOR OF THE PARTY AND ADDRESS.
>>> 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
>>> np.object	Python object type values
>>> np.string_	Fixed-length string type
>>> np.unicode_	Fixed-length unicode type

# Asking For Help

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

# **Array Mathematics**

# **Arithmetic Operations**

>>> g = a - b array([[-0.5, 0. , 0. ], [-3. , -3. , -3. ]])	Subtraction
[-3., -3., -3.]]) >>> np.subtract(a,b)	Subtraction
>>> b + a array([[ 2.5, 4. , 6. ], [ 5. , 7. , 9. ]])	Addition
>>> np.add(b,a)	Addition
>>> a / b array([[ 0.66666667, 1. , 1. ], [ 0.25 , 0.4 , 0.5 ]])	Division
>>> np.divide(a,b)	Division
>>> a * b array([[ 1.5, 4. , 9. ], [ 4. , 10. , 18. ]])	Multiplication
>>> 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 logarithm
>>> e.dot(f) array([[ 7., 7.], [ 7., 7.]])	Dot product

#### Comparison

Element-wise comparison	>>> a == b array([[False, True, True], [False, False, False]], dtype=bool)
Element-wise comparison	>>> a < 2 array([True, False, False], dtype=bool)
Array-wise comparison	>>> np.array_equal(a, b)

# **Aggregate Functions**

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

# **Copying Arrays**

Create a view of the array with the same data >>> h = a.view() >>> 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 Sort the elements >>> c.sort(axis=0) of an array's axis

# Subsetting, Slicing, Indexing

Subsetting	
>>> a[2]	1 2 3 Select the element at the 2nd index
>>> b[1,2] 6.0	1.5 2 3 Select the element at row 1 column 2 (equivalent to b[1][2])
Slicing	
>>> a[0:2] array([1, 2])	1 2 3 Select items at index 0 and 1
>>> b[0:2,1] array([ 2., 5.])	1.5 2 3 4 5 6
>>> b[:1] array([[1.5, 2., 3.]])	1.5 2 3 Select all items at row 0 (equivalent to b[0:1, :])
>>> c[1,] array([[[ 3., 2., 1.], [ 4., 5., 6.]]])	Same as [1,]
>>> a[::-1] array([3, 2, 1])	Reversed array a
Boolean Indexing	
>>> a[a<2] array([1])	1 2 3 Select elements from a less than 2
Fancy Indexing	
>>> b[[1, 0, 1, 0],[0, 1, 2, 0]] array([ 4. , 2. , 6. , 1.5])	Select elements (1,0),(0,1),(1,2) and (0,0)
>>> b[[1, 0, 1, 0]][.[0,12,0]] array[[[4, 1, 5, .6, .4, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,	Select a subset of the matrix's rows and columns

# **Array Manipulation**

# **Transposing Array**

>>> i = np.transpose(b)	Permute array dimensions
>>> i.T	Permute array dimensions

# Adding/Removing Elements

Return a new array with shape (2,6	>>> h.resize((2,6))
Append items to an arra	>>> np.append(h,g)
Insert items in an arra	>>> np.insert(a, 1, 5)
Delete items from an arra	>>> np.delete(a,[1])

# **Splitting Arrays**

vise sum		
um value	>>> np.hsplit(a,3) [array([1]),array([2]),array([3])] index	Split the array horizontally
rray row	, , ,	at the 3rd
elements	>>> np.vsplit(c,2) Split the array vertically	at the 2nd index
Mean	[array([[ 1.5, 2., 1.], [ 4., 5., 6.]]]),	at the zha maex
Madian	[ 4. , 5. , 6. ]]]),	

# **Changing Array Shape**

Flatten the array	>>> b.ravel()
Reshape, but don't change data	>>> g.reshape(3,-2)

#### Combining Arraya

S	Combining Array
s=0) Concatenate arrays	>>> np.concatenate((a,d),ax array([ 1, 2, 3, 10, 15, 20])
tack arrays vertically (row-wise)	>>> np.vstack((a,b)) array([[ 1. , 2. , 3. ],
tack arrays vertically (row-wise)	>>> np.r_[e,f] 5
Stack arrays horizontally (column-wise)	>>> np.hstack((e,f)) array([[ 7., 7., 1., 0.], [ 7., 7., 0., 1.]])
Create stacked column-wise arrays	>>> np.column_stack((a,d)) array([[ 1, 10],

>>> np.c\_[a,d] Create stacked column-wise arrays



# **Bokeh** Cheat Sheet

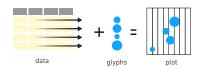
# BecomingHuman.Al



# Data Types

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

- 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
- >>> from bokeh.plotting import figure >>> from bokeh.io import output\_file, show >>> x = [1, 2, 3, 4, 5] >>> v = [6, 7, 2, 4, 5] >>> p = figure(title="simple line example", step 2 x axis label='x'. y\_axis\_label='y') >>> p.line(x, y, legend="Temp.", line\_width=2) <step 3 >>> output file("lines html") sten ( >>> show(p) \_\_\_\_step 5

# Data

Also see Lists, NumPv & Pandas

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

>>> import numpy as np >>> import pandas as pd

>>> df = pd.DataFrame(np.array([[33.9,4,65, 'US'],

[32.4,4,66, 'Asia'], [21.4,4,109, 'Europe']]), columns=['mpg',cyl', 'hp', 'origin'], index=['Toyota', 'Fiat', 'Volvo'])

>>> from hokeh models import ColumnDataSource

>>> cds\_df = ColumnDataSource(df)

# **Plotting**

>>> from bokeh.plotting import figure >>> p1 = figure(plot\_width=300, tools='pan,box\_zoom') >>> p2 = figure(plot\_width=300, plot\_height=300, x\_range=(0, 8), y\_range=(0, 8)) >>> p3 = figure()

# Show or Save Your Plots

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

# **Renderers & Visual Customizations**

#### **Glyphs**



#### Scatter Markers

>>> p1.circle(np.array([1,2,3]), np.array([3,2,1]), fill color='white') >>> p2.square(np.array([1.5,3.5,5.5]), [1,4,3],

color='blue', size=1)

#### Line Glyphs

>>> p1.line([1,2,3,4], [3,4,5,6], line\_width=2) >>> p2.multi line(pd.DataFrame([[1,2,3],[5,6,7]]), pd.DataFrame([[3,4,5],[3,2,1]]),

#### **Rows & Columns Layout**

#### Rows

>>> from bokeh.lavouts import row >>> layout = row(p1,p2,p3)

#### Columns

>>> from bokeh.layouts import columns

>>> layout = column(n1 n2 n3)

Nesting Rows & Columns

# >>>layout = row(column(p1,p2), p3)

>>> from bokeh.layouts import gridplot

>>> row1 = [p1,p2]>>> row2 = [n3]

**Grid Lavout** 

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

#### Legends

#### Legend Location

#### Inside Plot Area

>>> p.legend.location = 'bottom\_left'

#### 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])

>>> legend = Legend(items=[("One", [p1, r1]),("Two", [r2])], location=(0, -30))

>>> p.add\_layout(legend, 'right')

#### **Customized Glyphs**

# Selection and Non-Selection Glyphs

>>> p = figure(tools='box\_select' >>> p.circle('mpg', 'cyl', source=cds\_df, selection color='re

nonselection\_alpha=0.1)

#### Hover Glyphs

>>> hover = HoverTool(tooltips=None, mode='vline') >>> p3.add tools(hover)



>>> color\_mapper = CategoricalColorMapper(

factors=['US', 'Asia', 'Europe'], palette=['blue', 'red', 'green'])

>>> p3.circle('mpg', 'cyl', source=cds\_df, color=dict(field='origin',

transform=color\_mapper),

# legend='Origin'))

#### **Linked Plots**

#### Also see data

Also see data

#### 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')

# Tabbed Layout

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

>>> tab1 = Panel(child=p1, title="tab1") >>> tab2 = Panel(child=p2, title="tab2")

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

### **Legend Orientation**

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

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

#### Legend Background & Border

>>> p.legend.border line color = "navv" >>> p.legend.background\_fill\_color = "white"

# 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()

#### Standalone HTML

>>> from bokeh.embed import file html >>> html = file html(p, CDN, "my\_plot")

#### Components

>>> from bokeh.embed import components >>> script div = components(n)

# **Statistical Charts** With Bokeh

Also see Data

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



#### >>> from bokeh charts import Bar

>>> p = Bar(df, stacked=True, palette=['red;'blue'])

>>> from bokeh.charts import BoxPlot >>> p = BoxPlot(df, values='vals', label='cyl', legend='bottom\_right')



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

#### Scatter Plot

>>> from bokeh.charts import Scatter >>> p = Scatter(df, x='mpg', y ='hp', marker='square

xlabel='Miles Per Gallon',

# **Keras** Cheat Sheet

# **BecomingHuman.Al**



Keras is a powerfuland 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 no

>>> 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

input\_dim=100))

>>> model.add(Dense(1, activation='sigmoid'))

>>> model.compile(optimizer='rmsprop',

loss='binary crossentropy'.

metrics=['accuracy'])

### Data

Also see NumPv. 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,

>>> (x train.v train).(x test.v 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

>>> model.fit(data,labels,epochs=10,batch\_size=32)

>>> predictions = model.predict(data)

#### Other

>>> from urllib.request import urlopen

>>> data = np.loadtxt(urlopen("http://archive.ics.uci.edu/ ml/machine-learning-databases/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()

#### Multilaver 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))

#### **Convolutional Neural Network (CNN)**

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

>>> 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)))

>>> model2.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)))

>>> model2.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.klavers 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'))

# **Inspect Model**

>>> model.output shape

>>> model.summarv()

>>> model.get\_config()

>>> model.get weights()

Model output shape Model summary representation Model configuration

List all weight tensors in the model

# **Prediction**

>>> model3.predict(x\_test4, batch\_size=32)

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

# **Model Training**

>>> model3.fit(x train4,

**Model Fine-tuning** 

**Optimization Parameters** 

>>> from keras.optimizers import RMSprop

>>> opt = RMSprop(lr=0.0001, decay=1e-6)

>>> from keras.callbacks import EarlyStopping

y\_train4, batch\_size=32,

**MLP: Binary Classification** >>> model.compile(optimizer='adam

>>> model.compile(optimizer='rmsproi

>>> model.compile(optimizer='rmsprop',

**Recurrent Neural Network** 

Save/ Reload Models

>>> from keras.models import load model

>>> my\_model = load\_model('my\_model.h5')

>>> model3 save('model\_file.h5')

**MLP: Regression** 

>>> model3.compile(loss='binar

Early Stopping

>>> model3.fit(x\_train4,

Compile Model

>>> model2.compile(loss='categorical crossentropy',

optimizer=opt,

>>> early\_stopping\_monitor = EarlyStopping(patience=2)

metrics=['accuracy'])

epochs=15, validation\_data=(x\_test4,y\_test4),

metrics=['accuracy'])

loss='categorical\_cro metrics=['accuracy'])

metrics=['mae'])

optimizer='adam', metrics=['accuracy'])

**MLP: Multi-Class Classification** 

callbacks=[early stopping monitor])

y\_train4, batch size=32. verbose=1

validation\_data=(x\_test4,y\_test4))

# **Evaluate Your** Model's Performance

>>> score = model3.evaluate(x test

# **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)

# **Pandas Basics Cheat Sheet**

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Use the following import convention: >>> import pandas as pd

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

# **Pandas Data Structures**

#### Series

#### A one-dimensional

labeled array a capable of holding any

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

#### **Data Frame**

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



>>> data = {'Country': ['Belgium', 'India', 'Brazil'], 'Country': ['Belgium', 'India', 'Brazil'],

'Capital': ['Brussels', 'New Delhi', 'Brasília'],

'Population': [11190846, 1303171035n207847 >>> df = pd.DataFrame(data,

columns=['Country', 'Capital', 'Population'])

# **Dropping**

>>> s.drop(['a', 'c']) >>> df.drop('Country', axis=1)

Drop values from rows (axis=0) Drop values from columns(axis=1)

# Sort & Rank

>>> df.sort\_index() >>> df.sort\_values(by='Country') >>> df.rank()

Sort by labels along an axis Sort by the values along an axis Assign ranks to entries

# Retrieving Series/ **DataFrame Information**

>>> df shane (rows.columns) Describe index >>> df index >>> df.columns Describe DataFrame columns >>> df infn() Info on DataFrame >>> df count() Number of non-NA values

#### Summarv

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

#### Selection

Also see NumPy Arrays

#### Getting

>>> s['b'] Get one element >>> df[1:] Get subset of a DataFrame Population Country Capital New Delhi 1303171035

#### Selecting, Boolean Indexing & Setting

#### By Position

Select single value by row & >>> df.iloc[[0],[0]] 'Belgium' >>> df.iat([0],[0])

#### Select single value by row & By Label

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

#### By Label/Position

>>> df.ix[2] Country Capital Brasília Population 207847528 Select a single column of >>> df.ix[:,'Ca 0 Brussels 1 New Delhi 2 Brasília >>> df.ix[1,'Capital']
'New Delhi' Select rows and columns

#### **Boolean Indexing**

Series s where value is not >1 >>> s[~(s > 1)] >>> s[(s < -1) | (s > 2)] s where value is <-1 or >2 >>> df[df['Population']>1200000000] Use filter to adjust DataFrame

#### Setting

Set index a of Series s to 6 >>> s['a'] = 6

# **Asking For Help**

>>> help(pd.Series.loc)

# **Applying Functions**

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

# **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 a 10.0 b NaN

#### **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** >>> s.sub(s3, fill value=2) >>> s.div(s3, fill\_value=4)

# 1/0

#### Read and Write to CSV

>>> pd.read csv('file.csv', header=None, nrows=5) >>> df.to\_csv('mvDataFrame.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('sqlite:///: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 read\_sql\_query()

>>> pd.to\_sql('myDf', engine)

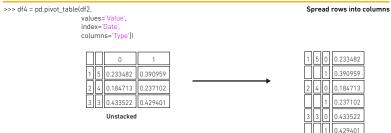
# **Pandas Cheat Sheet**

BecomingHuman.Al

# **Pandas Data Structures**

### **Pivot** >>> df3= df2.pivot(index='Date Spread rows into columns values='Value') Date Type Value Date 2016-03-2016-03-01 2016-03-03 2016-03-03

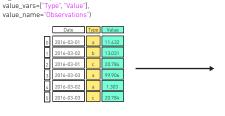
#### **Pivot Table**



#### Melt

>>> pd.melt(df2,

id\_vars=["Date"],



Gather columns into rows

Stacked

	Date	Variable	Observations
0	2016-03-01	Type	a
1	2016-03-02	Туре	b
2	2016-03-01	Туре	С
3	2016-03-03	Туре	а
4	2016-03-02	Туре	а
5	2016-03-03	Туре	С
6	2016-03-01	Value	11.432
7	2016-03-02	Value	13.031
8	2016-03-01	Value	20.784
9	2016-03-03	Value	99.906
10	2016-03-02	Value	1.303
11	2016-03-03	Value	20.784

# Advanced Indexing

#### Also see NumPy Arrays

eteeting	
>> df3.loc[:,(df3>1).any()]	Select cols with any vals >
>> df3.loc[:,(df3>1).all()]	Select cols with vals >
>> df3.loc[:,df3.isnull().any()]	Select cols with Nat
>> df3.loc[:,df3.notnull().all()]	Select cols without Nat

#### Indexing With isin

>>> df[(df.Country.isin(df2.Type))] Find same elements >>> df3.filter(items="a"."b"]) Filter on values >>> df.select(lambda x: not x%5) Select specific elements

>>> s where(s > 1)

Subset the data

>>> df6.query('second > first')

Query DataFrame

#### Setting/Resetting Index

>>> df.set_index('Country')	Set the index
>>> df4 = df.reset_index()	Reset the index
>>> df = df.rename(index=str,	Rename DataFrame
columns={"Country":"cntry",	
"Capital":"cptl",	

Forward Filling

>>> s3 = s.reindex(range(5)

#### Reindexing

>>> s2 = s.reindex(['a','c','d','e','b'])

#### Forward Filling

>>> df.reindex(range(4),

method='ffill')

0 3 Country Capital Population 0 Belgium Brussels 11190846 1 3 1 India New Delhi 2 3 2 Brazil Brasília 207847528 3 3 207847528 3 Brazil Brasília 4 3

#### MultiIndexing

>>> arrays = [np.array([1,2,3]) np.array([5,4,3])]

>>> df5 = pd.DataFrame(np.random.rand(3, 2), index=arrays)

>>> tuples = list(zip(\*arrays))

>>> index = pd.MultiIndex.from\_tuples(tuples,

names=['first', 'second'])

>>> df6 = pd.DataFrame(np.random.rand(3, 2), index=index)

>>> df2.set\_index(["Date", "Type"])

# **Duplicate Data**

>>> s3.unique()
>>> df2.duplicated('Type')
>>> df2 dron_dunlicates('Tyne' keen='last')

>>> df.index.duplicated()

Return unique values Check duplicates **Drop duplicates** Drop duplicates

# **Grouping Data**

#### Aggregation

>>> df2.groupby(by=['Date','Type']).mean()

>>> df4.groupby(level=0).sum()

>>> df4.groupby(level=0).agg({'a':lambda x:sum(x)/len(x), 'b': np.sum})

#### Transformation

>>> customSum = lambda x: (x+x%2)

>>> df4.groupby(level=0).transform(customSum)

# Missing Data

>>> df.dropna() >>> df3.fillna(df3.mean()) >>> df2.replace("a", "f")

Drop NaN value Fill NaN values with a predetermined value Replace values with others

# **Combining Data**

#### data1

X1	X2
а	11.432
b	1.303
С	99.906



#### **Pivot**

1100			
>>> pd.merge(data1, data2,	X1	X2	Х3
how='left', on='X1')	а	11.432	20.784
	b	1.303	NaN
	С	99.906	NaN
>>> pd.merge(data1, data2,	X1	X2	Х3
how='right',	а	11.432	20.784
on='X1')	b	1.303	NaN
	d	NaN	20.784
>>> pd.merge(data1, data2.	X1	X2	Х3
how='inner', on='X1')	а	11.432	20.784
OH= XT )	b	1.303	NaN
>>> pd.merge(data1, data2.	X1	X2	Х3
how='outer', on='X1')	а	11.432	20.784
on= XI)	h	1 303	NaN

#### Join

>>> data1.join(data2, how='right')

#### Concatenate

#### Vertical

>>> s.append(s2)

#### Horizontal/Vertical

>>> pd.concat([s,s2],axis=1, keys=['One','Two']) >>> pd.concat([data1, data2], axis=1, join='inner')

#### **Dates**

>>> df2['Date']= pd.to\_datetime(df2['Date'])

>>> df2['Date']= pd.date\_range('2000-1-1', periods=6,

>>> dates = [datetime(2012.5.1), datetime(2012.5.2)]

>>> index = pd.DatetimeIndex(dates)

>>> index = pd.date\_range(datetime(2012,2,1), end, freq='BM')

# **Visualization**

>>> import matplotlib.pyplot as plt

>>> s.plot() >>> plt.show()

>>> df2.plot()

>>> plt.show()

# Data Wrangling with pandas Cheat Sheet

# Syntax Creating DataFrames

	a	b	С
1	4	7	10
2	5	8	11
3	6	9	12

#### df = pd.DataFrame( {"a" : [4 ,5, 6], "b": [7, 8, 9]. "c": [10, 11, 12]},

index = [1, 2, 3]) Specify values for each column.

#### df = nd DataFrame( [[4, 7, 10], [5. 8. 11]. [6. 9. 12]]. index=[1, 2, 3], columns=['a', 'b', 'c']) Specify values for each row

		а	b	С
n	v			
	1	4	7	10
d	2	5	8	11
е	2	6	9	12

#### df = pd.DataFrame( {"a":[4,5,6], "b":[7.8.9]. "c" : [10, 11, 12]}. index = pd.MultiIndex.from\_tuples( [('d',1),('d',2),('e',2)],

names=['n''v'])) Create DataFrame with a MultiIndex

# 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')

# Windows

#### 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.

# Windows

#### df nlot hist() Histogram for each column

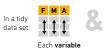
df plot scatter(x='w' v='h') Scatter chart using pairs of noints





# BecomingHuman.Al

# Tidy Data A foundation for wrangling in pandas



is saved in its

own column

Fach observation is saved in its

own row

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



# Reshaping Data Change the layout of a data set



pd.concat([df1.df2], axis=1) Append columns of DataFrames

#### 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'])

# Subset Observations (Rows



#### df[df.Length > 7]Extract rows that meet

pd.concat([df1.df2])

Append rows of DataFrames

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

of rows df.sample(n=10)

# Randomly select n rows.

df.iloc[10:20] Select rows by position

df sample(frac=0.5)

Randomly select fraction

# df.nlargest(n. 'value')

Select and order top n entries

#### df.nsmallest(n, 'value') Select and order bottom

ogic ili Fytholi (aliu	panuas)	
Less than	!=	Not equal to
Greater than	df.column.isin(values)	Group membership
Equal to	pd.isnull(obj)	Is NaN
Less than or equal to	pd.notnull(obj)	Is not NaN
Greater than or equal to	&, ,~,^,df.any(),df.all(	Logical and, or, not,
	Less than Greater than Equal to Less than or equal to	Greater than         df.column.isin(values)           Equal to         pd.isnull(obj)           Less than or equal to         pd.notnull(obj)

# 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='regex')

Select columns whose name matches regular expression regex

Logic in Python (and pandas)

#### Matches strings containing a period

Matches strings ending with word 'Length'
Matches strings beginning with the word 'Sepal'
Matches strings beginning with 'x and ending with 1,2,3,4,5
Matches strings except the string 'Species' 'Length\$ ^x[1-5]\$"

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

# **Summarise 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 GroupRy 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 values of each object.

#### count()

median()

Count non-NA/null values of each object

# 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()

var()

Mean value of each object Variance of each object

std() Standard deviation of each object.

# **Handling Missing Data**

Drop rows with any column having NA/null data.

#### Make New Columns



#### df assign(Area=lambda df: df Length\*df Height) Compute and annead one or more new column

# df['Volume'] = df.Length\*df.Height\*df.Depth

#### pd.qcut(df.col, n, labels=False)

Bin column into n buckets



pandas provides a large set of vector functions that operate on allcolumns 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) Flement-wise max

clip(lower=-10,upper=10) Trim values at input thresholds

#### min(axis=1) Flement-wise min

Vector

#### abs() Absolute value

# Combine Data Sets



B 2 Rows that appear

Rows that appear in both ydf and zdf

#### pd.merge(ydf, zdf, how='outer') Rows that appear in either or both ydf and zdf

#### pd.merge(ydf, zdf, how='outer', x1 x2 A 1 .query('\_merge == "left\_only"')

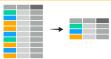
# .drop(columns=['\_merge'])

Rows that appear in ydf but not zdf (Setdiff)

cummin()

Cumulative min

# Windows



Size of each group. Aggregate group using function.

#### 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 agg(function)

shift(1)

returned vectors are of the length of the original DataFrame

#### Copy with values shifted by 1 rank(method='dense') Ranks with no gaps. rank(method='min')

Ranks. Ties get min rank. rank(nct=True) Ranks rescaled to interval [0, 1]

#### rank(method='first') Ranks. Ties go to first value.

The examples below can also be applied to groups. In this case, the function is applied on a per-group basis, and the

cummax()

Cumulative max

shift(-1) Copy with values lagged by 1. cumsum() Cumulative sum

cumprod() Cumulative product

# adf x1 x2 A 1 B 2 C 3 x1 x2 x3 dpd.merge(adf, bdf, how='left'. on='x1') Join matching rows from bdf to adf. x1 x2 x3 nd merge(adf hdf how='right' on='x1') Join matching rows from adf to bdf pd.merge(adf, bdf. how='inner', on='x1') B 2 F Join data. Retain only rows in both sets x1 x2 x3 pd.merge(adf, bdf, how='out how='outer'. on='x1') Join data Retain all values all rows

#### **Filtering Joins**



adf[adf.x1.isin(bdf.x1)] All rows in adf that have a match in bdf.

#### adf[~adf.x1.isin(bdf.x1)] x1 x2 All rows in adf that do not have a match

in bdf

# **Data Wrangling with** dplyr and tidyr **Cheat Sheet**

# **Syntax** Helpful conventions for wrangling

#### dplvr::tbl df(iris)

Converts data to thi class, this are easier to examine than data frames. R displays only the data that fits onscreen

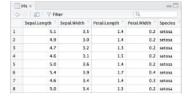
#### Source: local data frame [150 x 5] Sepal.Length Sepal.Width Petal.Length 4.9 3.0 1.4 4.7 3.2 1.3 4.6 3.1 1.5 5.0 3.6 Variables not shown: Petal.Width (dbl).

#### Species (fctr) dplyr::glimpse(iris)

Information dense summary of tbl data.

#### utils::View(iris)

View data set in spreadsheet-like display (note capital V)



#### dplvr::%>%

Passes object on lef hand side as first argument (or argument) of function on righthand side.

x %>% f(y) is the same as f(x, y)v %>% f(x, .. z) is the same as f(x, v, z)

"Piping" with %>% makes code more readable, e.g.

#### iris %>%

group\_by(Species) %>% summarise(avg = mean(Sepal.Width)) %>% arrange(avg)

# Tidy Data A foundation for wrangling in R

#### In a tidy data set:





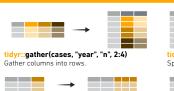


Tidy data complements R's vectorized operations. R will automatically preserve observations as you manipulate variables. No other format works as intuitively with R



# BecomingHuman.Al

Reshaping Data Change the layout of a data set



tidyr::spread(pollution, size, amount) Spread rows into columns tidyr::unite(data. col, ..., sep)

Unite several columns into one

Combine vectors into data frame (optimized). dplyr::arrange(mtcars, mpg) Order rows by values of a column (low to high) dplvr::arrange(mtcars, desc(mpg))

dplyr::data frame(a = 1:3, b = 4:6)

Order rows by values of a column (high to low) dplyr::rename(tb, y = year)

Rename the columns of a data frame

### Subset Observations (Rows

tidyr::separate(storms, date, c("y", "m", "d"))



#### dplyr::filter(iris, Sepal.Length > 7)

separate(storms, date, c("y", "m", "d")

Extract rows that meet logical criteria

#### dplyr::distinct(iris)

Remove dunlicate rows

dplyr::sample\_frac(iris, 0.5, replace = TRUE) Randomly select fraction of rows

#### dplyr::sample\_n(iris, 10, replace = TRUE) Randomly select n rows

#### dplyr::slice(iris, 10:15)

Select rows by position

#### dplyr::top\_n(storms, 2, date)

Select and order top n entries (by group if grouped data).

	Logic in R - ?	Comparison, ?base	::Logic
<	Less than		Not equal to
>	Greater than	%in%	Group membership
	Equal to	is.na	Is NA
<=	Less than or equal to	!is.na	Is not NA
50		& I I yor any all	

# Subset Variables (Columns



#### dplyr::select(iris, Sepal.Width, Petal.Length, Species)

Select columns whose name contains a character string.

Select columns whose name ends with a character string

Select every column.

Select columns whose name matches a regular expression

#### select(iris, num\_range("x", 1:5))

Select columns named x1, x2, x3, x4, x5,

#### select(iris, one\_of(c("Species", "Genus")))

Select columns whose names are in a group of names.

#### select(iris, starts\_with("Sepal"))

Select columns whose name starts with a character string.

#### select(iris Senal Length Petal Width

Select all columns between Sepal.Length and Petal.Width (inclusive).

#### select(iris, -Species)

Select all columns except Species.



Select columns by name or helper function.

#### select(iris, contains("."))

#### select(iris, ends with("Length"))

#### select(iris, everything())

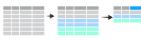
#### select(iris, matches(".t."))

# **Group Data**

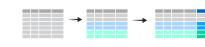
#### dplyr::group by(iris, Species) iris %>% group\_by(Species) %>% summarise(...) Group data into rows with the

#### same value of Species. dplyr::ungroup(iris)

Remove grouping information from data frame.



# iris %>% group\_by(Species) %>% mutate(...)



# **Summarise Data**



dplyr::summarise(iris, avg = mean(Sepal.Length)) Summarise data into single row of values

#### dplyr::summarise\_each(iris, funs(mean)) Apply summary function to each column.

dplyr::count(iris, Species, wt = Sepal.Length) Count number of rows with each unique value of variable (with or without weights).



Summarise uses summary functions, functions that take a vector of values and return a single value. such as:

#### dplvr::first

First value of a vector dplyr::last

Last value of a vector

# dplyr::nth

Nth value of a vector. dnlyr"n

#### # of values in a vector

dplyr::n\_distinct # of distinct values in

# a vector.

IOR of a vector

Minimum value in a vector

#### Maximum value in a vector. mean

Mean value of a vector. median

# Median value of a vector

Variance of a vector.

Standard deviation of a vector

#### Ranks rescaled to [0, 1] dplyr::row\_number

#### Ranks. Ties got to first value. dplyr::ntile

#### Bin vector into n buckets dplyr::between

#### Are values between a and b? dplyr::cume\_dist

# Cumulative distribution

# Element-wise min

### Combine Data Sets

#### **Mutating Joins**

x1 x2 x3 dplyr::lef\_join(a, b, by = "x1")

B 2 F Join matching rows from b to a Join matching rows from b to a. dplyr::right\_join(a, b, by = "x1")

Join matching rows from a to b. dplyr::inner\_join(a, b, by = "x1")

#### Join data. Retain only rows in both sets. dplyr::full ioin(a, b, by = "x1") A 1 T B 2 F C 3 NA Join data, Retain all values, all rows

dplyr::semi\_join(a, b, by = "x1")
A 1
B 2
All rows in a that have a match in All rows in a that have a match in b.

x1 x2 dplyr::anti\_join(a, b, by = "x1") All rows in a that do not have a match in b

# **Set Operations**

**Make New Variables** 

Compute and append one or more new columns

dplyr::mutate\_each(iris, funs(min\_rank))

Apply window function to each column

window

function

Copy with values shifed by 1.

Copy with values lagged by 1

dplyr::dense rank

Ranks with no gans

dplyr::percent rank

Ranks. Ties get min rank

dplyr::min rank

dplvr::lead

dplvr::lag

dplyr::mutate(iris, sepal = Sepal.Length + Sepal. Width)

dplyr::transmute(iris, sepal = Sepal.Length + Sepal. Width)

Mutate uses window functions, functions that take a vector of

Cumulative all

dplvr::cumany

Cumulative any

dplyr::cummean

Cumulative mean

Cumulative sum

Cumulative max

Cumulative min

Cumulative prod

Element-wise max

cumsum

cummax

cummin

cumprod

values and return another vector of values, such as:

Compute one or more new columns, Drop original columns

x1 x2 dplyr::intersect(y, z) B 2 Rows that appear in both y and z.



#### x1 x2 dplyr::setdiff(y, z) Rows that appear in y but not z.



# 31 32 31 32 dplyr::bind cols(v, z)

A 1 B 2
B 2 C 3
C 3 D 4 Caution: matches rows by position.

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.

# Scipy Linear Algebra **Cheat Sheet**





Also see NumPy

BecomingHuman.Al

Determinant

Create a 2X2 identity matrix

Create a 2x2 identity matrix

Compressed Sparse Row matrix

Compressed Sparse Column matrix

# **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**

Permute array dimension	>>> np.transpose(b)
Flatten the arra	>>> b.flatten()
Stack arrays horizontally (column-wis	>>> np.hstack((b,c))
Stack arrays vertically (row-wis	>>> np.vstack((a,b))
Split the array horizontally at the 2nd inde	>>> np.hsplit(c,2)
Split the array vertically at the 2nd inde	>>> np.vpslit(d,2)

#### **Polynomials**

>>> from numpy import poly1d >>> p = poly1d([3,4,5])

Create a polynomial object

#### **Vectorizing Functions**

>>> def mvfunc(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) >>> np.cast['f'](np.pi)

Return the real part of the array elements Return the imaginary part of the array elements Return a real array if complex parts close to 0 Cast object to a data type

Find the n-th derivative of a function at a point

#### **Other Useful Functions**

>>> misc.derivative(mvfunc.1.0)

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

# Linear Algebra

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**

>>> A.I	Inverse
>>> linalg.inv(A)	Inverse
Transposition	
>>> A.T	Tranpose matrix
>>> A.H	Conjugate transposition
Trace	
>>> np.trace(A)	Trace
Norm	
>>> linalg.norm(A)	Frobenius norm
>>> linalg.norm	L1 norm (max column sum)
>>> linalg.norm(A,np.inf)	L inf norm (max row sum)
Rank	
>>> np.linalg.matrix_rank(C)	Matrix rank
Determinant	

# >>> linalg.det(A)

Solving linear problems	
>>> linalg.solve(A,b)	Solver for dense matric

#### >>> E = np.mat(a).T

Solver for dense matrices >>> linalg.lstsq(F,E) Least-squares solution to linear matrix

#### Generalized inverse

>>> linalg.pinv(C) Compute the pseudo-inverse of a matrix >>> linalg.pinv2(C) Compute the pseudo-inverse of

#### **Creating Matrices**

>>> F = np.eve(3, k=1) >>> G = np.mat(np.identity(2)) >>> C[C > 0.5] = 0 >>> H = sparse.csr matrix(C)

>>> I = sparse.csc matrix(D) >>> J = sparse.dok matrix(A) >>> F todense()

Dictionary Of Keys matrix Sparse matrix to full matrix Identify sparse matrix >>> sparse.isspmatrix csc(A)

#### **Matrix Functions**

Addition

>>> np.add(A,D)	Addition
Subtraction	
>>> np.subtract(A,D)	Subtraction
Division	
>>> np.divide(A,D)	Division
Multiplication	
>>> A @ D	Multiplication operator (Python 3)
>>> np.multiply(D,A)	Multiplication
>>> np.dot(A,D)	Dot product
>>> np.vdot(A,D)	Vector dot product
>>> np.inner(A,D)	Inner product
>>> np.outer(A,D)	Outer product
>>> np.tensordot(A,D)	Tensor dot product
>>> np.kron(A.D)	Kronecker product

#### **Exponential Functions**

Matrix exponential
Matrix exponential (Taylor Series)
Matrix exponential (eigenvalue

# **Logarithm Function**

>>> linalg.logm(A)	Matrix logarithm

#### **Trigonometric Functions**

Matrix sine	>>> linalg.sinm(D)
Matrix cosine	>>> linalg.cosm(D)
Matrix tangen	>>> linalg.tanm(A)

#### **Hyperbolic Trigonometric Functions**

>>> linalg.sinhm(D)	Hypberbolic matrix sine
>>> linalg.coshm(D)	Hyperbolic matrix cosine
>>> linalg.tanhm(A)	Hyperbolic matrix tangent

#### **Matrix Sign Function**

>>> np.signm(A)	Matrix sign function
Materia Communic Donat	

#### **Arbitrary Functions**

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

Evaluate matrix function

# **Sparse Matrix Routines**

Inverse	
>>> sparse.linalg.inv(I)	Inverse
Norm	
>>> sparse.linalg.norm(I)	Norm
Solving linear problems	
>>> enarce linala encolve(H I)	Solver for sparse matrices

#### **Sparse Matrix Functions**

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

#### **Decompositions**

Eigenvalues and Eigenvectors	
>>> la, v = linalg.eig(A)	Solve ordinary or generalized

>>> l1, l2 = la	eigenvalue problem for square matrix
>>> v[:,0]	First eigenvector
>>> v[:,1]	Second eigenvector
>>> linalq.eiqvals(A)	Unpack eigenvalues

#### Singular Value Decomposition

>>> U,s,Vh = linalg.svd(B)	Singular Value Decomposition (SVD)
>>> M,N = B.shape	
>>> Sig = linalg.diagsyd(s.M.N)	Construct sigma matrix in SVD

#### **LU Decomposition**

>>> P.L.U = linalq.lu(C) I II Decomposition

#### Sparse Matrix Decompositions

>>> la, v = sparse.linalg.eigs(F,1) Eigenvalues and eigenvectors >>> sparse.linalg.svds(H. 2)

# **Asking For Help**

>>> help(scipy.linalg.diagsvd)

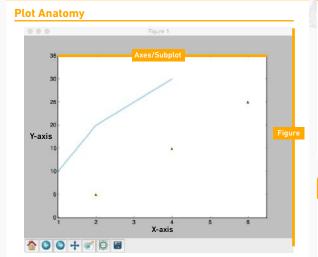
>>> np.info(np.matrix)

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

# Matplotlib Cheat Sheet

# BecomingHuman.Al

# **Anatomy & Workflow**



#### Workflow

Prepare data

Customize plot

Create plot

Plot

Show plot



# **Prepare The Data**

Also see Lists & NumPv

#### **Index Tricks**

>>> 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'))

# **Create Plot**

>>> import matplotlib.pyplot as plt

#### Figure

>>> 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)

# **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()

>>> ax.scatter(x,y,marker=".")

>>> ax.plot(x,y,marker="o")

# Linestyles

>>> 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,

Mathtext

-2.1, 'Example Graph', style='italic')

>>> ax.annotate("Sine", xy=(8, 0), xycoords='data' xytext=(10.5.0). textcoords='c arrowprops=dict(arrowstyle="-

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

#### Limits, Legends & Lavouts

#### Limits & Autoscaling

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

Add padding to a plot

>>> ax.axis('equal') Set the aspect ratio

>>> 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

>>> ax.set(title='An Example Axes'. ylabel=

xlabel='X-Axis' >>> ax.legend(loc='best')

No overlapping plot elements

Set a title and x-and

>>> ax.xaxis.set(ticks=range(1,5), ticklabels=[3,100,-12,"foo"]) direction=

Manually set x-ticks

Make y-ticks longer and go in and out

#### **Subplot Spacing**

>>> fig3.subplots adjust(wspace=0.5 hspace=0.3 left=0.125. right=0.9, top=0.9, bottom=0.1

>>> fig.tight\_layout()

#### **Axis Spines**

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

Pseudocolor plot of 2D array

Pseudocolor plot of 2D array

Plot contours

Plot 2D vector fields

>>> ax1.spines['top'=].set visible(False)

Make the top axis line for a plot invisible

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

Move the hottom

# **Plotting Routines**

#### 1D Data

>>> lines = ax plot(x v)

>>> ax.scatter(x.v) >>> axes[0,0].bar([1,2,3],[3,4,5])

>>> axes[1.0].barh([0.5.1.2.5].[0.1.2])

>>> ax.fill\_between(x,y,color='yellow')
2D Data

arrays cmap='gist\_earth', interpolation='nearest',

vmin=-2

>>> axes[1,1].axhline(0.45) >>> axes[0,1].axvline(0.65) >>> ax.fill(x,y,color='blue')

>>> fig. ax = plt.subplots()

>>> im = ax.imshow(img

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

Draw points with lines or markers connecting them

Draw a vertical line across axes Draw filled polygons

Fill between y-values and 0

Colormapped or RGB

>>> CS = plt.contour(Y,X,U) >>> axes2[2].contourf(data1)

# **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)

#### **Data Distributions**

>>> ax1.hist(y) Plot a histogram >>> ax3.boxplot(v) Make a box and whisker plot >>> ax3.violinplot(z) Make a violin plot

connectionstyle="arc3").)

>>> axes2[0].pcolor(data2) >>> axes2[0].pcolormesh(data)

Plot filled contours >>> axes2[2]= ax.clabel(CS) Label a contour plot

# **Save Plot**

# Save figures

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

# Save transparent figures

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

# **Show Plot**

>>> plt.show()

# Close & Clear

>>> plt.cla()

>>> plt.clf() >>> plt.close()

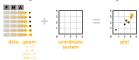
# **Data Visualisation** with applot2 **Cheat Sheet**

#### **Basics**

ggplot2 is based on the grammar of graphics, the idea that you can build every graph from the same few components: a data set, a set of geoms-visual marks that represent data points, and a coordinate system



To display data values, man variables in the data set to aesthetic properties of the geom like size color, and x and y locations



Build a graph with qplot() or qqplot()

# qplot(x = cty, y = hwy, color = cyl, data = mpg, geom = "point")

Creates a complete plot with given data, geom, and mappings. Supplies many useful defaults.

#### ggplot(data = mpg, aes(x = cty, y = hwy)) Begins a plot that you finish by adding layers to. No

defaults, but provides more control than gplot().

ggplot(mpg, aes(hwy, cty)) + geom point(aes(color = cvl)) + geom\_smooth(method ="lm") +

coord cartesian() + scale color gradient() theme bw()

Add a new layer to a plot with a geom\_\*() or stat\_\*() function. Each provides a geom, a set of aesthetic mappings, and a default stat and position adjustment.

#### last plot()

Returns the last plot

#### ggsave("plot.png", width = 5, height = 5)

Saves last plot as 5' x 5' file named "plot.png" in working directory. Matches file type to file extension.

# **Coordinate Systems**

#### r <- b + geo m\_bar()



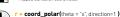
coord\_cartesian(xlim = c(0, 5)) xlim, ylim The default cartesian coordinate system



+ coord fixed(ratio = 1/2) Cartesian coordinates with fixed aspect ratio between x and y units



coord\_flip()





+ coord\_trans(vtrans = "sort") xtrans, ytrans, timx, timy Transformed cartesian coordinates. Set extras and strains to the name



+ coord map(projection = "ortho" + coord\_maptprojection = ortho, orientation=c(41, -74, 0)) projection, orientation, xlim, ylim Map projections from the mapproj package (mercator (default), azequalarea, lagrange, etc.)

# Geom's Use a geom to represent data points, use the geom's aesthetic properties to represent variables. Each function returns a layer

#### One Variable

#### Continuous











#### **Graphical Primitives**







linejoin="round", linemitre=1) x, y, alpha, color, linetyna cizo 1 + geom\_ribbon(aes(ymin=unemploy - 900, ymax=unemploy + 900)) x, ymax, ymin, alpha, color, fill, linetype, size

# + geom\_segment(aes(

xend = long + delta\_long, yend = lat + delta\_lat)) xend, v. vend, alpha, color, linetype, size

+ geom\_rect(aes(xmin = long, ymin = lat, xmax= long + delta\_long, ymax = lat + delta\_lat))

#### **Three Variables**

seals\$z <- with(seals, sqrt(delta\_long^2 + delta\_lat^2))
m <- qqplot(seals, aes(long, lat))



+ geom\_contour(aes(z = z))



geom\_raster(aes(fill = z), hjust=0.5, st=0.5, interpolate=FALSE)



geom\_tile(aes(fill = z))

# Faceting

Facets divide a nlot into subplots based on the values of one or more discrete variables.



+ facet grid(, ~ fl)



+ facet\_grid(vear ~ .)



+ facet\_grid(year ~ fl)



t + facet\_wrap(~ fl)

Set erales to let avis limits vary across facets et\_grid(y ~ x, scales = "free")

x and v axis limits adjust to individual facets

"free\_x" - x axis limits adjust "free\_y" - y axis limits adjust

Set labeller to adjust facet labels

t + facet grid( ~ fl labeller = label both) fl:c fl:d fl:e fl:p fl:r rid(. ~ fl, labeller = label both) α<sup>c</sup> α<sup>d</sup> α<sup>e</sup> α<sup>p</sup> α<sup>r</sup>

grid(. ~ fl, labeller = label\_both c d e p r

#### Two Variables

### Continuous X. Continuous Y

geom blank() geom\_jitter()







#### Discrete X, Continuous Y

g <- ggplot(mpg, aes(class, hwy))

geom\_bar(stat = "identity") geom\_boxplot()



geom\_violin(scale = "area")

# Discrete X, Discrete Y



#### Continuous Bivariate Distribution



geom\_bin2d(binwidth = c(5, 0.5))

geom\_density2d()

geom\_hex()

#### Continuous Function



geom line()

# geom\_step(direction = "hv")

### df <- data.frame(grp = c("A", "B"), fit = 4:5, se = 1:2)

k + geom\_crossbar(fatten = 2) x, y, ymax, ymin, alpha, color, fill, linetype, s



# + geom\_map(aes(map\_id = state), map = map) + expand\_limits(x = map\$long, y = map\$lat) x x alpha color fill linetyne size

# **Position Adjustments**

Position adjustments determine how to arrange geoms that would otherwise occupy the same space

# s <- ggplot(mpg, aes(fl. fill = drv))

s + geom\_bar(position = "dodge")
Arrange elements side by side



s + geom\_bar(position = "fill")
Stack elements on ton of one another norm



geom\_bar(position = "stack")



f + geom\_point(position = "jitter")
Add random noise to X and Y position of each element to avoid overplotting

Each position adjustment can be recast as a function with manual width and height arguments

colorbar, legend, or none (no legend)

s + geom\_bar(position = position\_dodge(width = 1))

# Labels

t + xlab("New X label") Change the label on the X axis

t + ylab("New Y label") Change the label on the Y axis

Leaends

guides(color = "none")

t + labs(title =" New title", x = "New x", y = "New y")
All of the above

t + scale\_fill\_discrete(name = "Title", labels = c("A", "B", "C"))

theme(legend.position = "bottom")

Use scale functions to update legend

# f + stat\_unique() Themes

f + stat identity()

f + stat sum()

dparams = list(df=5))







Stats An alternative way to build a layer

e.q. a + geom\_bar(stat = "bin")

FMA

fl cty ctl

Some plots visualize a transformation of the original data set

Each stat creates additional variables to map aesthetics to. These

variables use a common ..name.. syntax. stat functions and geom

functions both combine a stat with a geom to make a layer, i.e.

i + stat\_density2d(aes(fill = ..level..), geom = "polygon", n = 100)

a + stat\_bin(binwidth = 1, origin = 10)

f + stat\_bin2d(bins = 30, drop = TRUE)

f + stat binhex(bins = 30)

m + stat contour(aes(z = z))

+ stat\_boxplot(coef = 1.5)

stat ecdf(n = 40)

+ stat ecdf(n = 40)

+ stat\_bindot(binwidth = 1, binaxis = "x")

f + stat\_density2d(contour = TRUE, n = 100)

m+ stat\_spoke(aes(radius= z, angle = z))

ggplot() + stat\_function(aes(x = -3:3), fun = dnorm, n = 101, args = list(sd=0.5))

f + stat summary(fun.data = "mean cl boot")

m + stat\_summary\_hex(aes(z = z), bins = 30, fun = mean)

g + stat\_ydensity(adjust = 1, kernel = "gaussian", scale = "area")

stat\_quantile(quantiles = c(0.25, 0.5, 0.75), formula =  $y \sim log(x)$ ,

f + stat\_smooth(method = "auto", formula = y ~ x, se = TRUE, n = 80, fullrange = FALSE, level = 1,95) x,y1.se, x, y1.se, x, y1

 $f + stat_quantile(quantiles = c(0.25, 0.5, 0.75), formula = y \sim log(x),$ 

f + stat \_smooth(method = "auto", formula = y ~ x, se = TRUE, n = 80, fullrange = FALSE, level = 0.95) xyl\_se\_x\_y\_\_\_mn\_\_\_man\_\_

ggplot() + stat\_qq(aes(sample=1:100), distribution = qt,

a + stat\_density(adjust = 1, kernel = "gaussian")

stat\_bin(geom="bar") does the same as geom\_bar(stat="bin")

Use a stat to choose a common transformation to visualize.

theme minimal()

r + theme grev()

ggthemes - Package with additional ggplot2 themes

# Scales

Scales control how a plot mans data values to the visual values of an aesthetic. To change the mapping, add a ustom scale.



n <- b + geom\_bar(aes(fill = fl))



n + scale fill manual(



values = c("skyblue", "royalblue". "blue". "navv") limits = c("d", "e", "n", "r"), breaks =c("d", "e", "n", "r") name = "fuel", labels = c("D", "E", "P", "R"))

#### **General Purpose scales**

scale\_\*\_continuous() - map cont' values to visual values scale\_\*\_discrete() - map discrete values to visual values scale\_\*\_identity() - use data values as visual values

scale\_\*\_manual(values = c()) - map discrete values to

#### X and Y location scales

scale\_x\_date(labels = date\_format("%m/%d"), breaks = date breaks("2 weeks"))

scale\_x\_datetime() - treat x values as date times. Use scale\_x\_log10() - Plot x on log10 scale scale\_x\_reverse() - Reverse direction of x axis

#### scale x sqrt() - Plot x on square root scale Color and fill scales



n + scale fill brewer palette = "Blues") For palette choices:



display.brewer.all() + scale\_fill\_grey(



+ scale\_fill\_gradientn( colours = terrain.colors(6)) Also: rainbow(), heat.colors(), topo.colors(), cm.colors(), RColorBrewer::brewer.pal()

<- a + geom\_dotplot(

# na.value = "red")

Shape scales

aes(shape = fl))

p + scale\_shape solid = FALSE) p + scale\_shape\_manual( values = c(3:7))

0 □ 6 ▽ 12 ⊞ 18 ◆ 24 ▲ 1 ○ 7 🛛 13 🔯 19 • 25 🔻 2 △ 8 ★ 14 四 20 • • • 3 + 9 ◆ 15 ■ 21 ○

4 × 10 ⊕ 16 ● 22 ■ 0 0

5♦ 11 🕱 17 🛦 23 ♦ 0 O



scale\_size\_area(max = 6)

# Zooming

# Without clipping (preferred)

t + coord cartesian( xlim = c(0, 100), ylim = c(10, 20)

#### With clipping (removes unseen data points)



t + xlim(0, 100) + ylim(10, 20)

scale\_x\_continuous(limits = c(0, 100)) + scale\_y\_continuous(limits = c(0, 100))

