



# Cheat Sheets for AI

Neural Networks,  
Machine Learning,  
DeepLearning &  
Big Data

**The Most Complete List  
of Best AI Cheat Sheets**

BecomingHuman.AI

# Table of Content



## Data Science with Python

### Machine Learning

- 06 Machine Learning Basics
- 07 Scikit Learn with Python
- 08 Scikit Learn Algorithm
- 09 Choosing ML Algorithm

11 Tensor Flow

12 Python Basics

13 PySpark Basics

14 Numpy Basics

15 Bokeh

16 Karas

17 Pandas

18 Data Wrangling with Pandas

19 Data Wrangling with dplyr & tidyr

20 SciPi

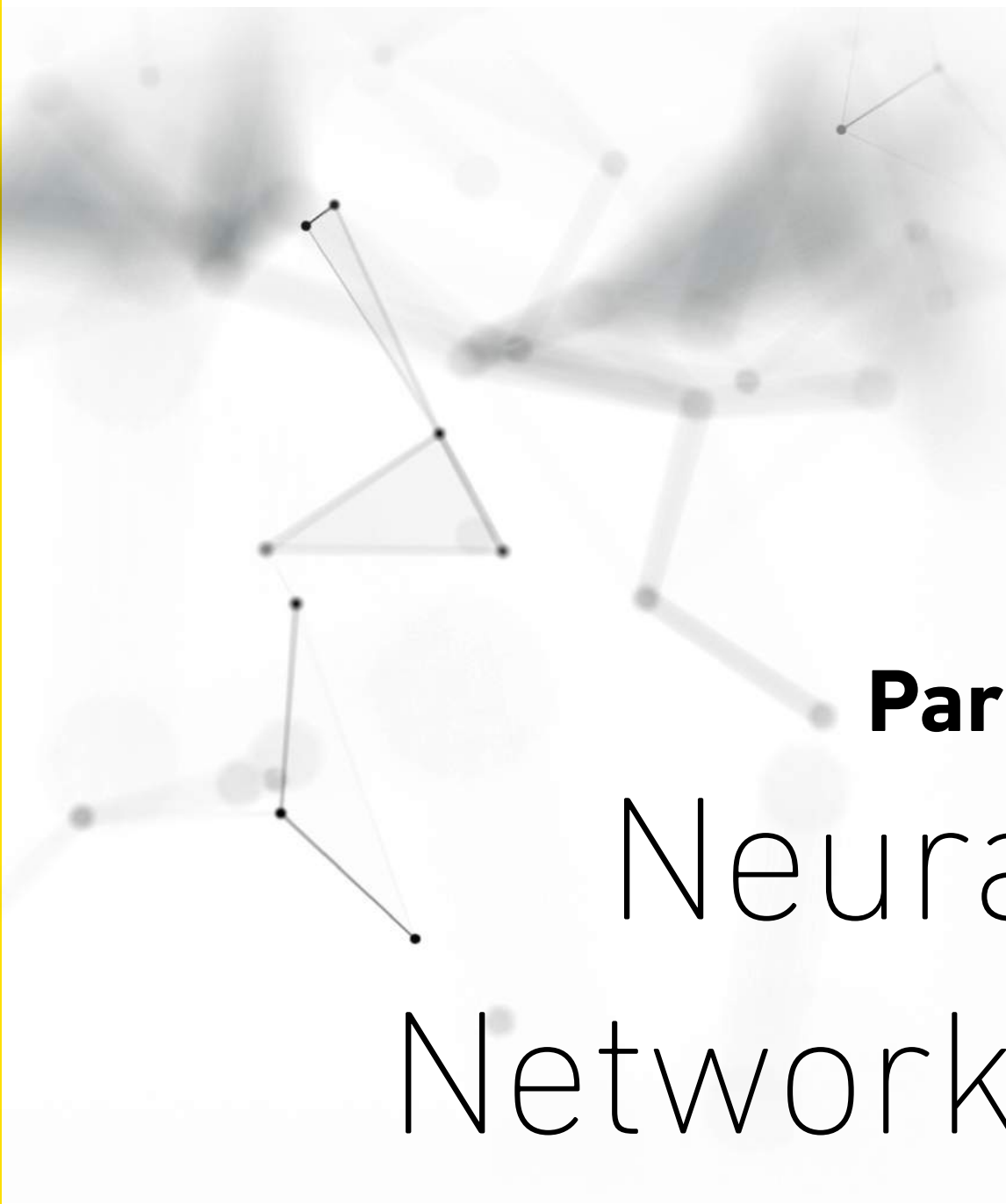
21 Matplotlib

22 Data Visualization with ggplot

23 Big-O

### Neural Networks

- 03 Neural Networks Basics
- 04 Neural Network Graphs
















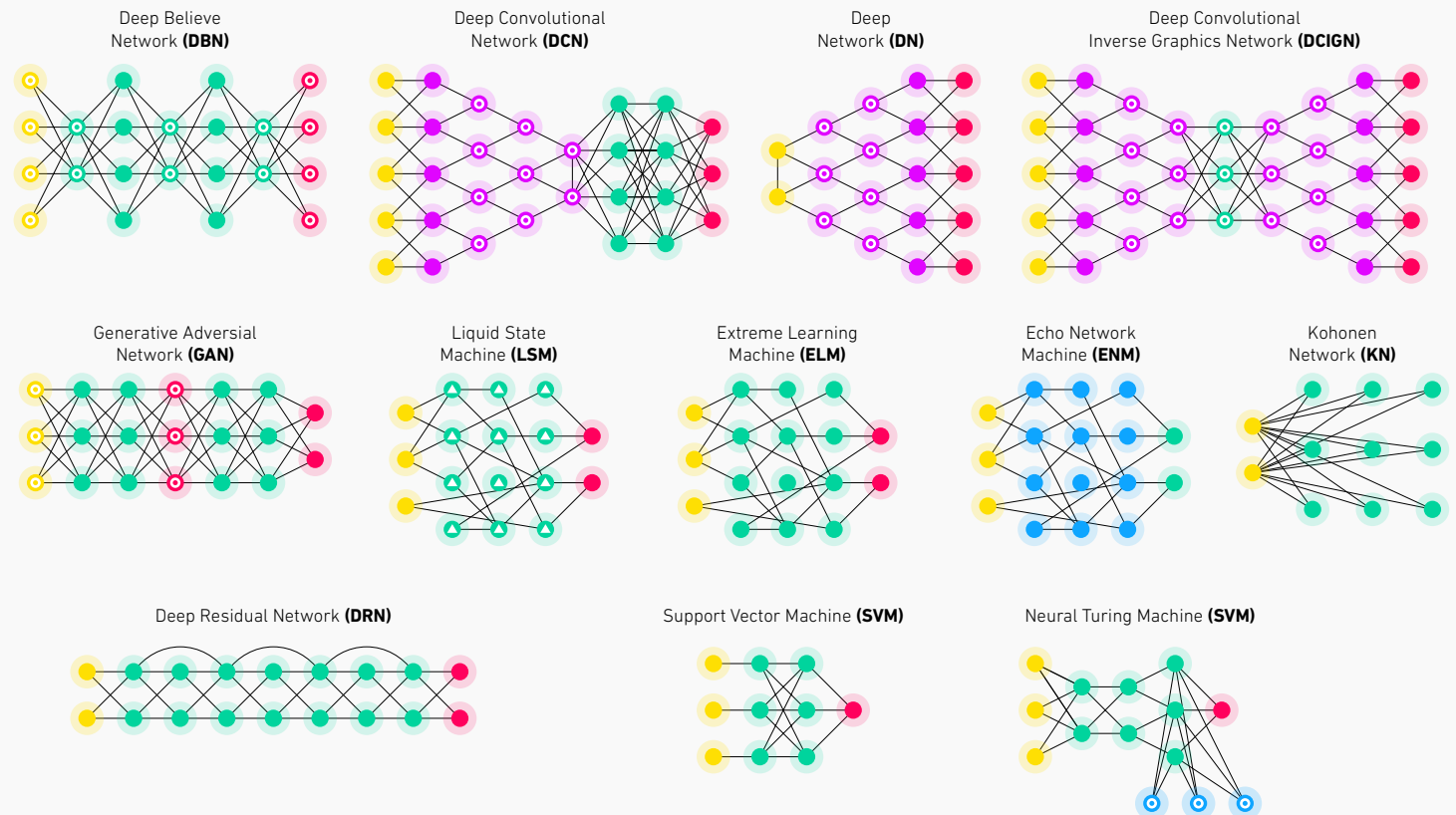
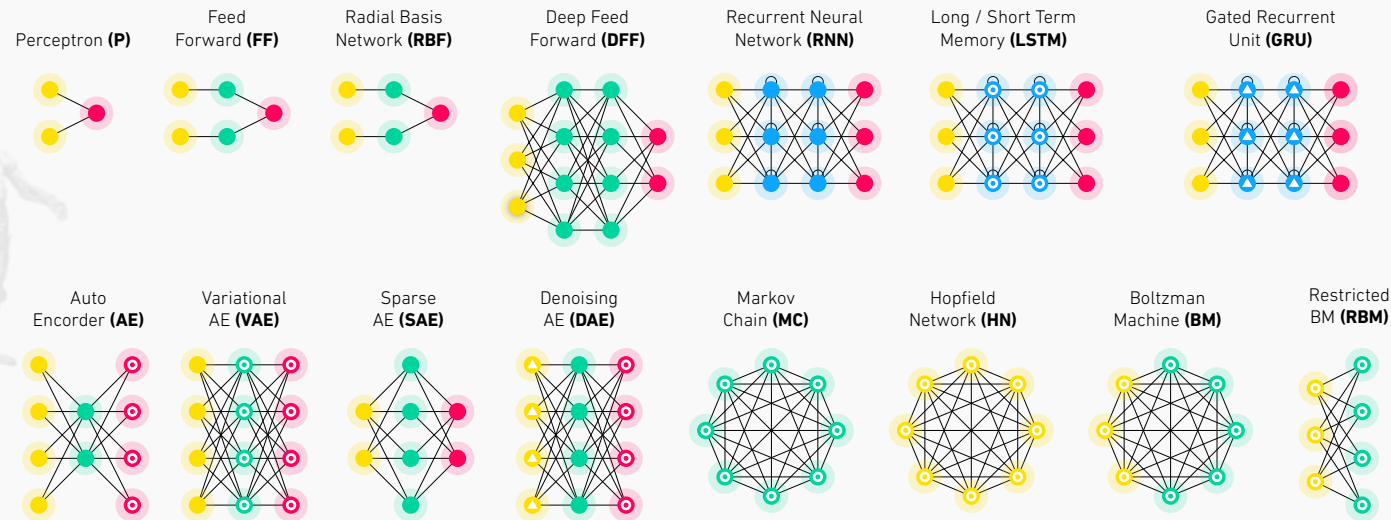
# **Part 1** Neural Networks

# Neural Networks Basic Cheat Sheet

BecomingHuman.AI

## Index

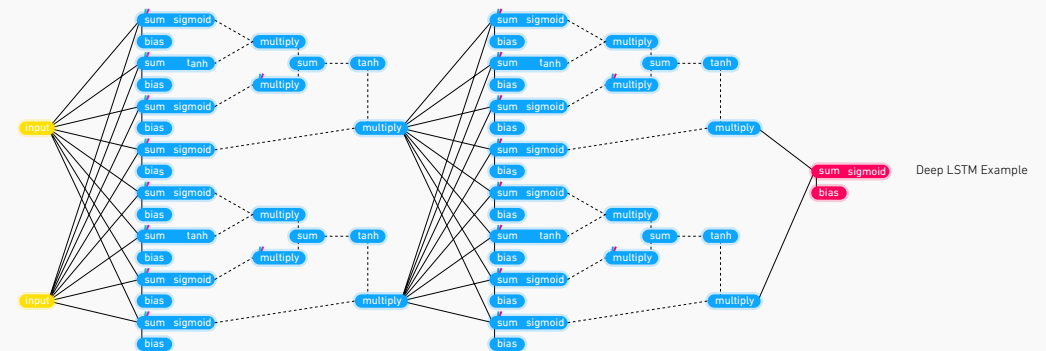
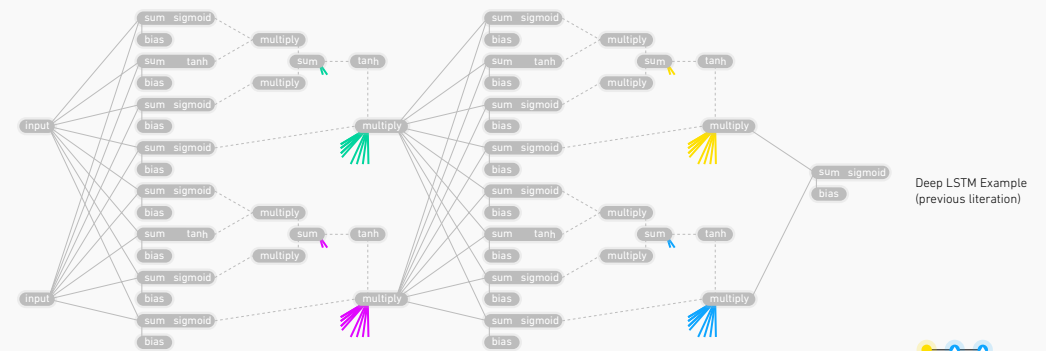
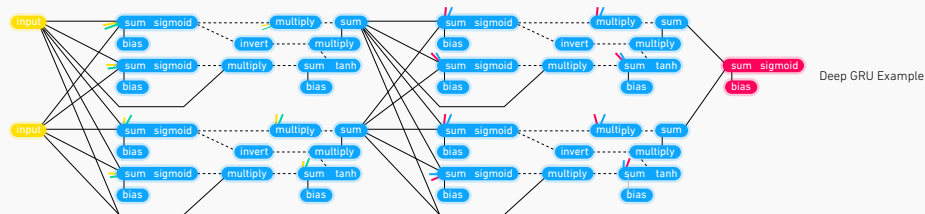
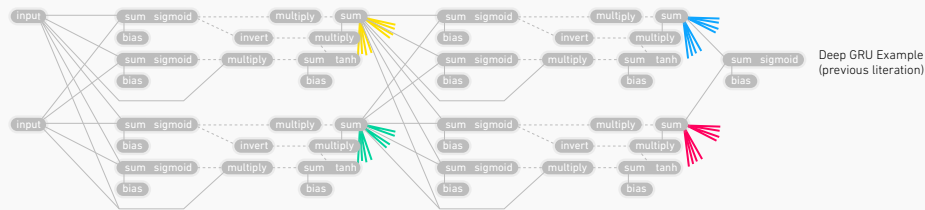
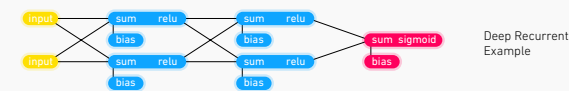
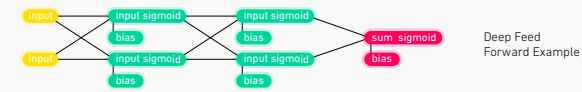
-  Backfed Input Cell
-  Input Cell
-  Noisy Input Cell
-  Hidden Cell
-  Probablistic Hidden Cell
-  Spiking Hidden Cell
-  Output Cell
-  Match Input Output Cell
-  Recurrent Cell
-  Memory Cell
-  Different Memory Cell
-  Kernel
-  Convolutional or Pool



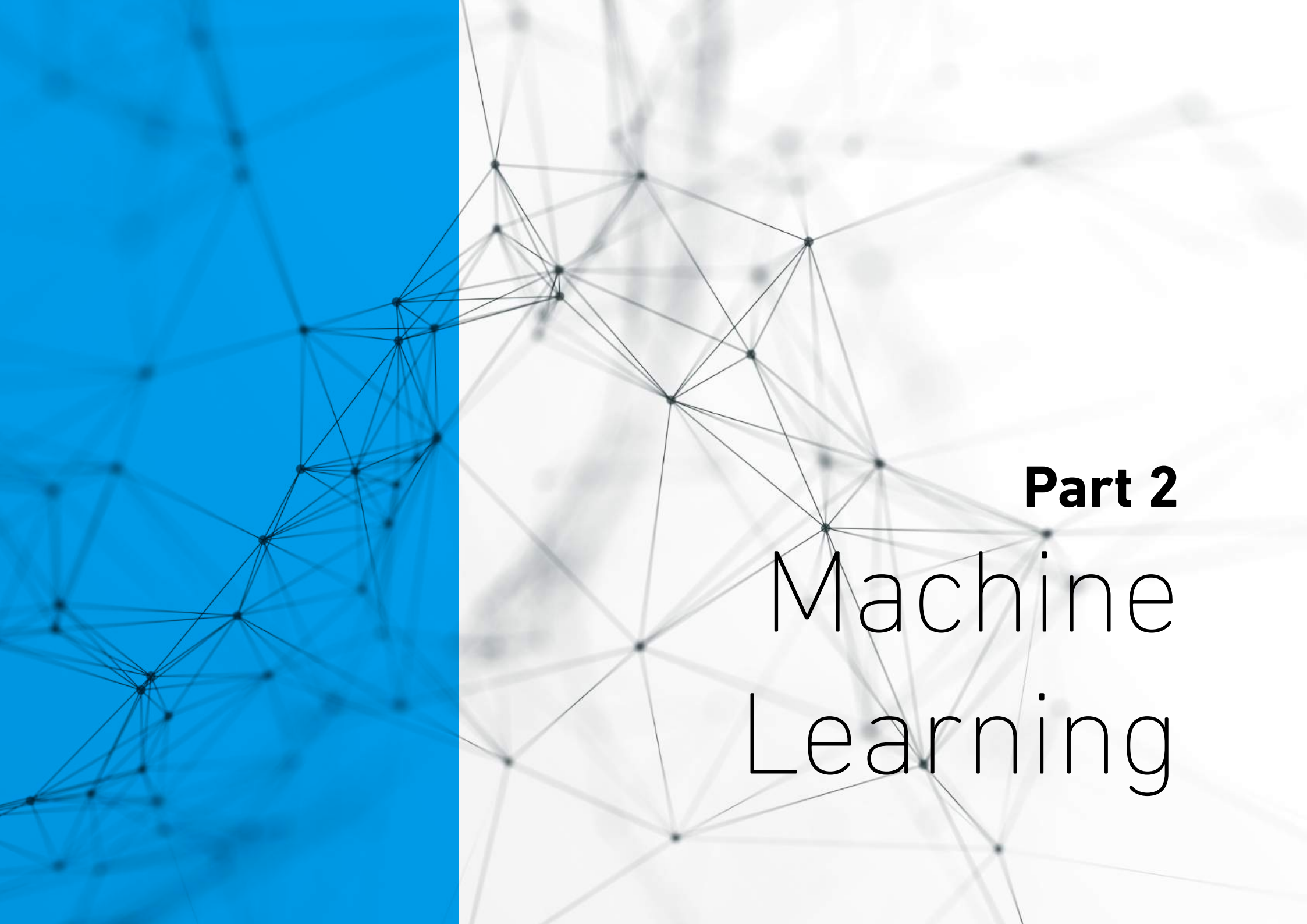
Original Copyright by AsimovInstitute.org [See original here](#)

# Neural Networks Graphs Cheat Sheet

## Becoming Human.AI







# **Part 2** Machine Learning

# Machine Learning Overview

## MACHINE LEARNING IN EMOJI

### Becoming Human.AI

#### SUPERVISED

human builds model based on input / output

#### UNSUPERVISED

human input, machine output  
human utilizes if satisfactory

#### REINFORCEMENT

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

#### BASIC REGRESSION

##### LINEAR

`linear_model.LinearRegression()`

Lots of numerical data



##### LOGISTIC

`linear_model.LogisticRegression()`

Target variable is categorical



#### CLUSTER ANALYSIS

##### K-MEANS

`cluster.KMeans()`

Similar datum into groups based on centroids



##### ANOMALY DETECTION

`covariance.EllipticalEnvelope()`

Finding outliers through grouping



#### CLASSIFICATION

##### NEURAL NET

`neural_network.MLPClassifier()`

Complex relationships. Prone to overfitting  
Basically magic.



##### K-NN

`neighbors.KNeighborsClassifier()`

Group membership based on proximity



##### DECISION TREE

`tree.DecisionTreeClassifier()`

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



##### RANDOM FOREST

`ensemble.RandomForestClassifier()`

Find best split randomly  
Can also be regression



##### SVM

`svm.SVC()` `svm.LinearSVC()`

Maximum margin classifier. Fundamental Data Science algorithm



##### NAIVE BAYES

`GaussianNB()` `MultinomialNB()` `BernoulliNB()`

Updating knowledge step by step with new info



#### FEATURE REDUCTION

##### T-DISTRI 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

`lda.LDA()`

Linear combination of features that separates classes



#### OTHER IMPORTANT CONCEPTS

##### BIAS VARIANCE TRADEOFF

##### UNDERFITTING / OVERFITTING

##### INERTIA

##### ACCURACY FUNCTION

$(TP+TN) / (P+N)$

##### PRECISION FUNCTION

`manifold.TSNE()`

##### SPECIFICITY FUNCTION

$TN / (FP+TN)$

##### SENSITIVITY FUNCTION

$TP / (TP+FN)$

# Cheat-Sheet Skicit learn Phyton For Data Science BecomingHuman.AI



## 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 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(random.random((2,5)))
>>> y_pred = lr.predict(X_test)
>>> y_pred = knn.predict_proba(X_test)
```

Predict Labels  
Predict Labels  
Estimate probability of a label

### Unsupervised Estimators

```
>>> y_pred = k_means.predict(X_train)
```

Predict labels in clustering algo

## Loading the Data

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

```
>>> 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

```
>>> knn.score(X_test, y_test)
>>> from sklearn.metrics import accuracy_score
>>> accuracy_score(y_test, y_pred)
```

Estimator score method  
Metric scoring functions

#### Classification Report

```
>>> from sklearn.metrics import classification_report
>>> print(classification_report(y_test, y_pred))
```

Precision, recall, f1-score  
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)
```

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

```
>>> 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))
```

## Model Fitting

### Supervised learning

```
>>> lr.fit(X, y)
>>> knn.fit(X_train, y_train)
>>> svc.fit(X_train, y_train)
```

Fit the model to the data

### Unsupervised Learning

```
>>> k_means.fit(X_train)
>>> pca_model = pca.fit_transform(X_train)
```

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

## 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 = GaussianNB()
```

#### KNN

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

### Unsupervised Learning Estimators

#### Principal Component Analysis (PCA)

```
>>> from sklearn.decomposition import PCA
>>> pca = PCA(n_components=0.95)
```

#### K Means

```
>>> from sklearn.cluster import KMeans
>>> k_means = KMeans(n_clusters=3, random_state=0)
```

## Training And Test Data

```
>> from sklearn.cross_validation import train_test_split
>> X_train, X_test, y_train, y_test = train_test_split(X,
                                                         y,
                                                         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, y_train)
>>> print(grid.best_score_)
>>> print(grid.best_estimator_.n_neighbors)
```

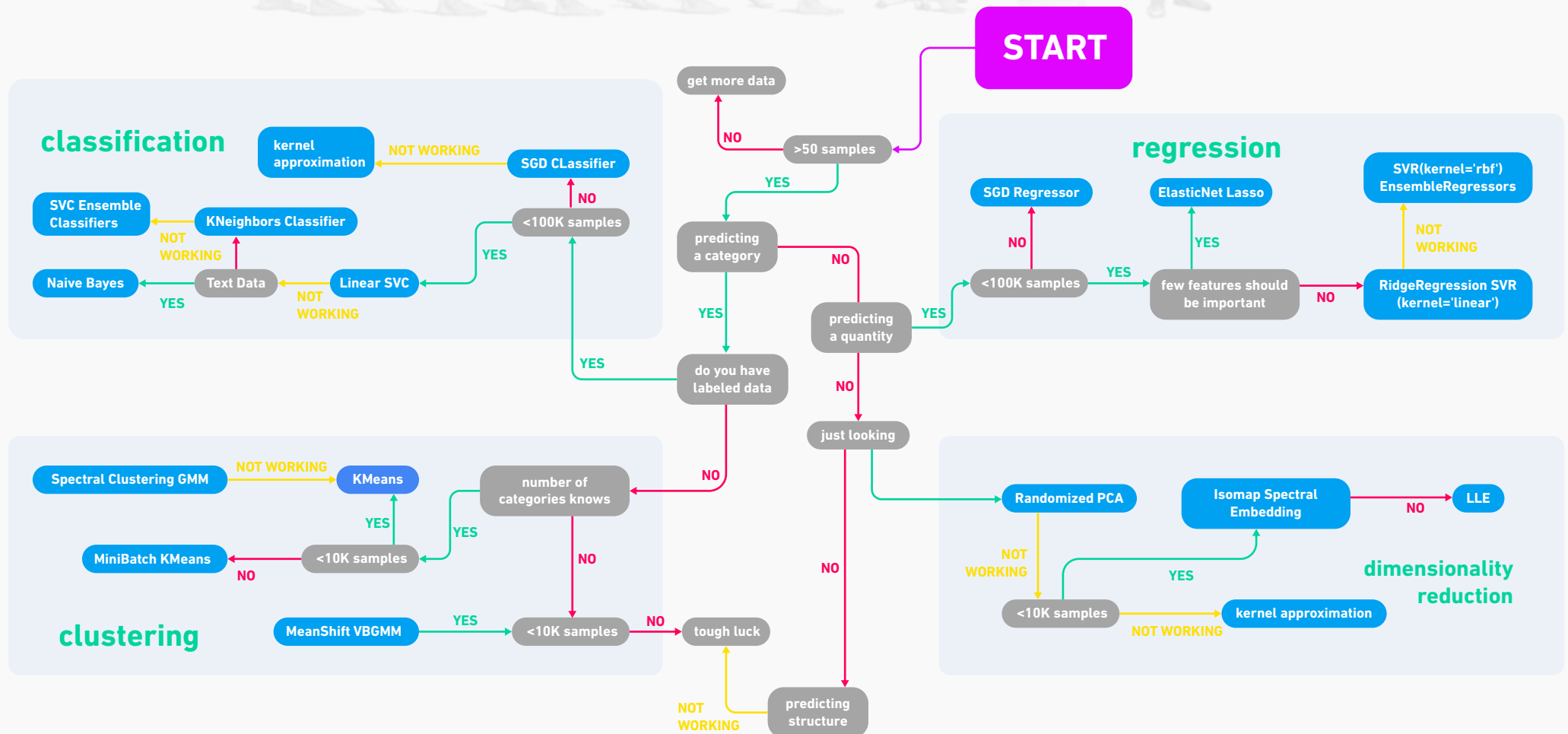
### Randomized Parameter Optimization

```
>>> from sklearn.grid_search import RandomizedSearchCV
>>> params = [{"n_neighbors": range(1,5),
              "weights": ["uniform", "distance"]}
>>> rsearch = RandomizedSearchCV(estimator=knn,
                                param_distributions=params,
                                cv=4,
                                n_iter=8,
                                random_state=5)
>>> rsearch.fit(X_train, y_train)
>>> print(rsearch.best_score_)
```



# Skicit-learn Algorithm

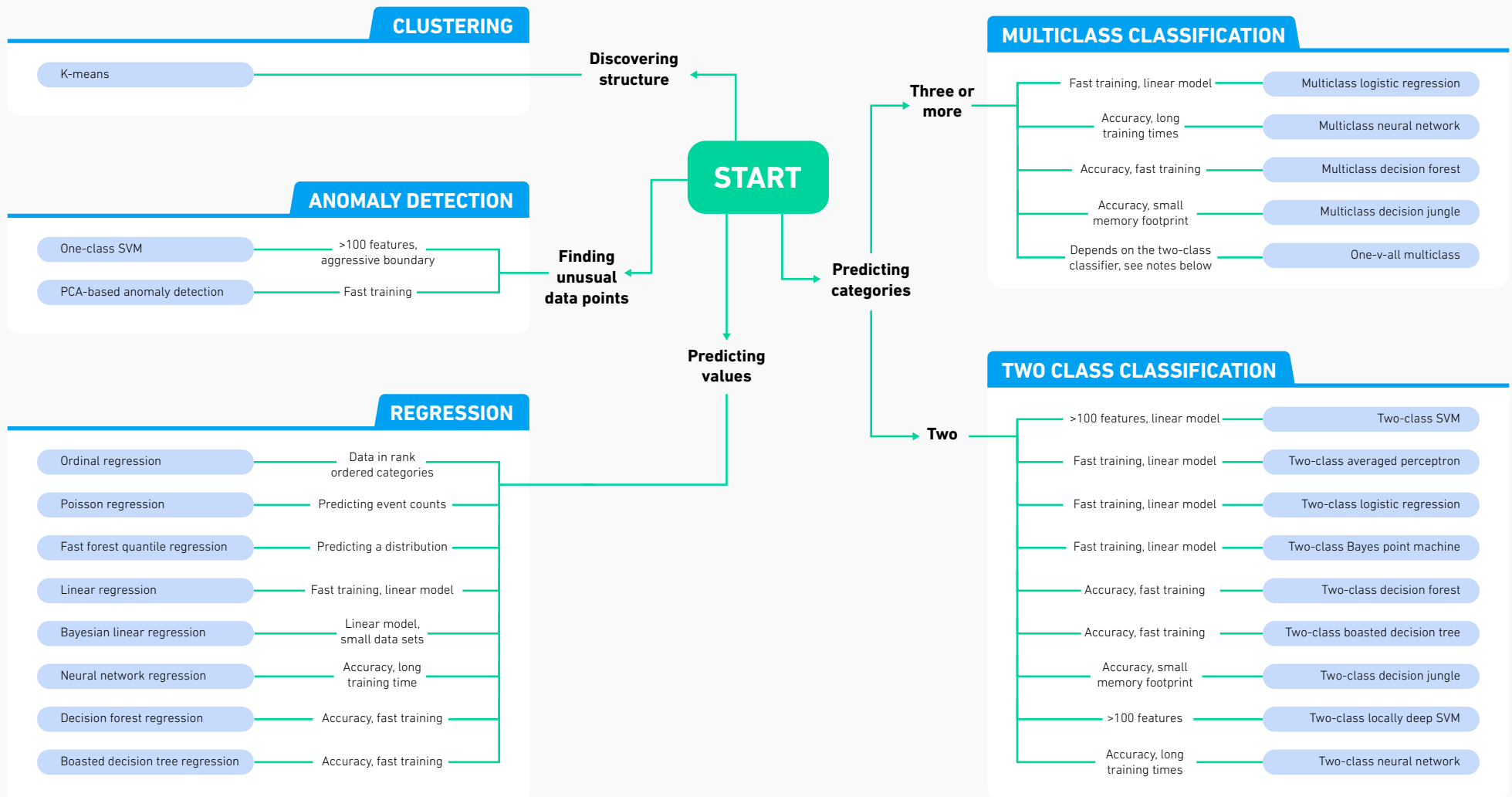
## Becoming Human.AI



# Algorithm Cheat Sheet

## Becoming Human.AI

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.



The background is split vertically. The left half is a solid orange color with a faint, dark orange geometric pattern of interconnected lines and dots. The right half is a light grey color with a faint, dark grey geometric pattern of interconnected lines and dots, similar to the one on the left but less dense.

**Part 3**

# Data Science with Python

# Tensor Flow Cheat Sheet

## Becoming Human.AI



In May 2017 Google announced the second-generation of the TPU, as well as the availability of the TPUs in Google Compute Engine.[12] The second-generation TPUs deliver up to 180 teraflops of performance, and when organized into clusters of 64 TPUs provide up to 11.5 petaflops.

### Info

#### TensorFlow

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.

#### 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.

#### Skflow

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.

### Installation

#### How to install new package in Python

`pip install <package-name>`

Example: `pip install requests`

#### How to install tensorflow?

`device = cpu/gpu`

`python_version = cp27/cp34`

`sudo pip install`

`https://storage.googleapis.com/tensorflow/linux/$device/tensorflow-0.8.0-$python_version-none-linux_x86_64.whl`  
`sudo pip install`

#### How to install Skflow

`pip install sklearn`

#### How to install Keras

`pip install keras`

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

### Helpers

#### Python helper Important functions

`type(object)`

Get object type

`help(object)`

Get help for object (list of available methods, attributes, signatures and so on)

`dir(object)`

Get list of object attributes (fields, functions)

`str(object)`

Transform an object to string object?

Shows documentations about the object

`globals()`

Return the dictionary containing the current scope's global variables.

`locals()`

Update and return a dictionary containing the current scope's local variables.

`id(object)`

Return the identity of an object. This is guaranteed to be unique among simultaneously existing objects.

`import_builtin_`

`dir(builtins)`

Other built-in functions

### Tensor Flow

#### Main classes

`tf.Graph()`

`tf.Operation()`

`tf.Tensor()`

`tf.Session()`

#### Some useful functions

`tf.get_default_session()`

`tf.get_default_graph()`

`tf.reset_default_graph()`

`ops.reset_default_graph()`

`tf.device("/cpu:0")`

`tf.name_scope(value)`

`tf.convert_to_tensor(value)`

#### TensorFlow Optimizers

`GradientDescentOptimizer`

`AdadeltaOptimizer`

`AdagradOptimizer`

`MomentumOptimizer`

`AdamOptimizer`

`FtrlOptimizer`

`RMSPropOptimizer`

#### Reduction

`reduce_sum`

`reduce_prod`

`reduce_min`

`reduce_max`

`reduce_mean`

`reduce_all`

`reduce_any`

`accumulate_n`

#### Activation functions

`tf.nn?`

`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`

etc.

### Skflow

#### Main classes

`TensorFlowClassifier`

`TensorFlowRegressor`

`TensorFlowDNNClassifier`

`TensorFlowDNNRegressor`

`TensorFlowLinearClassifier`

`TensorFlowLinearRegressor`

`TensorFlowRNNClassifier`

`TensorFlowRNNRegressor`

`TensorFlowEstimator`

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

`batch_size=32,`

`steps=200, // except`

`TensorFlowRNNClassifier - there is 50`

`optimizer='Adagrad',`

`learning_rate=0.1,`

#### Each class has a method fit

`fit(X, y, monitor=None, logdir=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 regression).

**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)`

Args:

**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 predictions.

**batch\_size:** If test set is too big, use batch size to split it into mini batches. By default the batch\_size member variable is used.

Returns:

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

# Phyton For Data Science

# Cheat-Sheet Phyton Basic

## BecomingHuman.AI



### Variables and Data Types

#### Variable Assignment

```
>>> x=5
>>> x
5
```

#### Calculations With Variables

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

#### 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
bool()	True, True, True	Variables to booleans

### Asking For Help

```
>>> help(str)
```

### Lists

Also see NumPy Arrays

```
>>> a = 'is'
>>> b = 'nice'
>>> my_list = ['my', 'list', a, b]
>>> my_list2 = [[4,5,6,7], [3,4,5,6]]
```

#### Selecting List Elements

Index starts at 0

<b>Subset</b>	
>>> my_list[1]	Select item at index 1
>>> my_list[-3]	Select 3rd last item
<b>Slice</b>	
>>> my_list[1:3]	Select items at index 1 and 2
>>> my_list[1:]	Select items after index 0
>>> my_list[:3]	Select items before index 3
>>> my_list[:]	Copy my_list
<b>Subset Lists of Lists</b>	
>>> my_list2[1][0]	my_list[list][itemOfList]
>>> my_list2[1][:2]	

#### List Operations

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

#### List Methods

>>> my_list.index(a)	Get the index of an item
>>> my_list.count(a)	Count an item
>>> my_list.append('!')	Append 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)
>>> my_2darray =
np.array([[1,2,3],[4,5,6]])
```

#### Selecting Numpy Array Elements

Index starts at 0

<b>Subset</b>	
>>> my_array[1]	Select item at index 1
2	
<b>Slice</b>	
>>> my_array[0:2]	Select items at index 0 and 1
array([1, 2])	
<b>Subset 2D Numpy arrays</b>	
>>> my_2darray[:,:0]	my_2darray[rows, columns]
array([], [])	

#### Numpy Array Operations

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

#### Numpy Array 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(my_array)	Mean of the array
>>> np.median(my_array)	Median of the array
>>> my_array.corrcoef()	Correlation coefficient
>>> np.std(my_array)	Standard deviation

### Strings

Also see NumPy Arrays

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

#### String Operations

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

#### String Operations

Index starts at 0

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

#### String Methods

>>> my_string.upper()	String to uppercase
>>> my_string.lower()	String to lowercase
>>> my_string.count('w')	Count String elements
>>> 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 Python



SPYDER

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

BecomingHuman.AI



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

>>> sc.version	Retrieve SparkContext version
>>> sc.pythonVer	Retrieve Python version
>>> sc.master	Master URL to connect to
>>> str(sc.sparkHome)	Path where Spark is installed on worker nodes
>>> str(sc.sparkUser())	Retrieve name of the Spark User 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', '1g'))
>>> 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 `wholeTextFiles()`.

```
>>> 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
[('a', 7), ('a', 2)]	
>>> rdd.first()	Take first RDD element
('a', 7)	
>>> rdd.top(2)	Take top 2 RDD elements
[('b', 2), ('a', 7)]	

#### 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)	Filter the RDD
.collect()	Return distinct RDD values
[('a',7),('a',2)]	Return (key,value) RDD's keys
>>> rdd5.distinct().collect()	
[a,2,b,7]	
>>> rdd.keys().collect()	
[a, a, b]	

### Iterating

#### Getting

```
>>> def g(x): print(x)
>>> rdd.foreach(g)
('a', 7)
('b', 2)
('a', 2)
```

### Retrieving RDD Information

#### Basic Information

>>> rdd.getNumPartitions()	List the number of partitions
>>> rdd.count()	Count RDD instances
3	
>>> rdd.countByKey()	Count RDD instances by key
defaultdict(<type 'int'>, {'a':2,'b':1})	
>>> rdd.countByValue()	Count RDD instances by value
defaultdict(<type 'int'>, {('b',2):1,('a',2):1,('a',7):1})	
>>> rdd.collectAsMap()	Return (key,value) pairs as a dictionary
{a: 2,b: 2}	
>>> rdd3.sum()	Sum of RDD elements
4950	
>>> sc.parallelize([]).isEmpty()	Check whether RDD is empty
true	

#### Summary

>>> rdd3.max()	Maximum value of RDD elements
99	
>>> rdd3.min()	Minimum value of RDD elements
0	
>>> rdd3.mean()	Mean value of RDD elements
49.5	
>>> rdd3.stdev()	Standard deviation of RDD elements
28.866070047722118	
>>> rdd3.variance()	Compute variance of RDD elements
833.25	
>>> rdd3.histogram(3)	Compute histogram by bins
[(0,33,66,99),[33,33,34)]	
>>> rdd3.stats()	Summary statistics (count, mean, stdev, max & min)

### Applying Functions

>>> 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: x+(x[1],x[0]))	Apply a function to each RDD element 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 (key,value) pair of rdd4 without changing the keys
.collect()	
[('a','x'),('a','y'),('a','z'),('b','p'),('b','r')]	

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

### Sort

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

### Reshaping Data

#### Reducing

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

#### Grouping by

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

#### Aggregating

>>> seqOp = (lambda x,y: (x[0]+y,x[1]+1))	Aggregate RDD elements of each partition and then the results
>>> combOp = (lambda x,y: (x[0]+y[0],x[1]+y[1]))	
>>> rdd3.aggregate((0,0),seqOp,combOp)	Aggregate values of each RDD key
(4950,100)	
>>> rdd.aggregateByKey((0,0),seqOp,combOp)	Aggregate the elements of each 4950 partition, and then the results
.collect()	Merge the values for each key
[('a,(9,2)), ('b,(2,1))]	
>>> rdd3.fold(0,add)	
4950	
>>> rdd.foldByKey(0, add)	
.collect()	
[('a',9),('b',2)]	
>>> rdd3.keyBy(lambda x: x+x)	Create tuples of RDD elements by applying a function
.collect()	

### Reshaping Data

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

### Saving

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

### Stopping SparkContext

```
>>> sc.stop()
```

### Execution

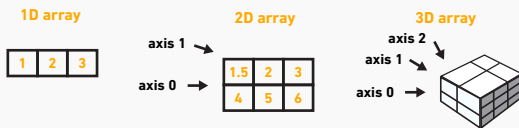
```
$ ./bin/spark-submit examples/src/main/python/pi.py
```

# NumPy Basics Cheat Sheet

## BecomingHuman.AI



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.



## 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 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
```

## I/O

### 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

```
>>> 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          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 logarithm
>>> np.dot(f)           Dot product
array([[ 7. ,  7. ]])
```

### Comparison

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

### Aggregate Functions

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

## Copying Arrays

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

## Sorting Arrays

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

## Subsetting, Slicing, Indexing

### Subsetting

```
>>> a[2]              Select the element at the 2nd index
3
>>> b[1,2]            Select the element at row 1 column 2
6.0                    (equivalent to b[1][2])
```

### Slicing

```
>>> a[0:2]            Select items at index 0 and 1
array([1, 2])
>>> b[0:2,1]          Select items at rows 0 and 1 in column 1
array([[2., 5.]])
>>> b[:1]             Select all items at row 0
array([[1.5, 2., 3.]])   (equivalent to b[0:1,:])
>>> c[1,...]          Same as [1,:,:]
array([[[ 3., 2., 1. ],
        [ 4., 5., 6. ]]])
>>> a[::-1]           Reversed array a
array([3, 2, 1])
```

### Boolean Indexing

```
>>> a[a<2]           Select elements from a less than 2
array([1])
```

### Fancy Indexing

```
>>> b[[1,0,1,0],[0,1,2,0]] Select elements (1,0),(0,1),(1,2) and (0,0)
array([ 4.,  2.,  6.,  1.5])
>>> b[[1,0,1,0]][:,[0,1,2,0]] Select a subset of the matrix's rows
array([[ 4.,  5.,  6.,  4. ],   and columns
       [ 1.5,  2.,  3.,  1.5],
       [ 4.,  5.,  6.,  4. ],
       [ 1.5,  2.,  3.,  1.5]])
```

## Array Manipulation

### Transposing Array

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

### Adding/Removing Elements

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

### Splitting Arrays

```
>>> np.hsplit(a,3)       Split the array horizontally at the 3rd
array([1]),array([2]),array([3])) index
>>> np.vsplit(c,2)       Split the array vertically at the 2nd index
array([[[ 1.5,  2. ,  1. ],
        [ 4. ,  5. ,  6. ]]]),
```

### Changing Array Shape

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

### Combining Arrays

```
>>> np.concatenate((a,d),axis=0) Concatenate arrays
array([ 1,  2,  3, 10, 15, 20])
>>> np.vstack((a,b))      Stack arrays vertically (row-wise)
array([[ 1. ,  2. ,  3. ],
       [ 1.5,  2.,  3. ],
       [ 4. ,  5. ,  6. ]])
>>> np.r_[e,f]            Stack arrays vertically (row-wise)
>>> np.hstack((e,f))      Stack arrays horizontally (column-wise)
array([[ 7.,  7.,  0.,  1. ]])
>>> np.column_stack((a,d)) Create stacked column-wise arrays
array([[ 1, 10],
       [ 2, 15],
       [ 3, 20]])
>>> np.c_[a,d]            Create stacked column-wise arrays
```



# Bokeh Cheat Sheet

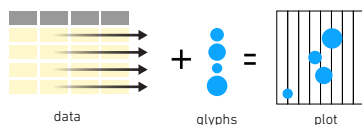
BecomingHuman.AI



## 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 interface are:

1. Prepare some data:  
Python lists, NumPy arrays, Pandas DataFrames and other sequences of values
2. Create a new plot
3. Add renderers for your data, with visual customizations
4. Specify where to generate the output
5. Show or save the results

```
>>> from bokeh.plotting import figure
>>> from bokeh.io import output_file, show
>>> x = [1, 2, 3, 4, 5]
>>> y = [6, 7, 2, 4, 5]
>>> p = figure(title='simple line example',
>>>             x_axis_label='x',
>>>             y_axis_label='y')
>>> p.line(x, y, legend='Temp', line_width=2)
>>> output_file('lines.html')
>>> show(p)
```

## Data

Also see Lists, NumPy & 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 bokeh.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)
```

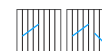
## Renderers & Visual Customizations

### 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]]),
>>>               color='blue')
```

### Rows & Columns Layout

#### Rows

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

#### Columns

```
>>> from bokeh.layouts import columns
>>> layout = column(p1, p2, p3)
```

#### Nesting Rows & Columns

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

### Grid Layout

```
>>> from bokeh.layouts import gridplot
>>> row1 = [p1, p2]
>>> row2 = [p3]
>>> 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

Also see data



**Selection and Non-Selection Glyphs**

```
>>> p = figure(tools='box_select')
>>> p.circle('mpg', 'cyl', source=cds_df,
>>>          selection_color='red',
>>>          nonselection_alpha=0.1)
```



**Hover Glyphs**

```
>>> hover = HoverTool(tooltips=None, mode='vline')
>>> p3.add_tools(hover)
```



**Colormapping**

```
>>> 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

#### 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])
```

## 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(p)
```

## Statistical Charts With Bokeh

Also see Data

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



#### Bar Chart

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



#### Box Plot

```
>>> from bokeh.charts import BoxPlot
>>> p = BoxPlot(df, values='vals', label='cyl',
>>>             legend='bottom_right')
```



#### Histogram

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



#### Scatter Plot

```
>>> from bokeh.charts import Scatter
>>> p = Scatter(df, x='mpg', y='hp',
>>>             marker='square',
>>>             xlabel='Miles Per Gallon',
```

# Keras Cheat Sheet

## BecomingHuman.AI



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

### A Basic Example

```
>>> import numpy as np
>>> from keras.models import Sequential
>>> from keras.layers import Dense
>>> data = np.random.random((1000,100))
>>> labels = np.random.randint(2,size=(1000,1))
>>> model = Sequential()
>>> model.add(Dense(32,
                    activation='relu',
                    input_dim=100))
>>> model.add(Dense(1, activation='sigmoid'))
>>> model.compile(optimizer='rmsprop',
                 loss='binary_crossentropy',
                 metrics=['accuracy'])
```

### Data

Also see NumPy, Pandas & Scikit-Learn

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

### Keras Data Sets

```
>>> from keras.datasets import boston_housing,
mnist,
cifar10,
imdb
>>> (x_train,y_train),(x_test,y_test) = mnist.load_data()
>>> (x_train2,y_train2),(x_test2,y_test2) = boston_housing.load_data()
>>> (x_train3,y_train3),(x_test3,y_test3) = cifar10.load_data()
>>> (x_train4,y_train4),(x_test4,y_test4) = imdb.load_data(num_words=20000)
>>> num_classes = 10
>>> 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/
pima-indians-diabetes.data'),delimiter=',')
>>> X = data[:,0:8]
>>> y = data[:,8]
```

### Model Architecture

#### Sequential Model

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

#### Multilayer Perceptron (MLP)

##### Binary Classification

```
>>> from keras.layers import Dense
>>> model.add(Dense(12,
                  input_dim=8,
                  kernel_initializer='uniform',
                  activation='relu'))
>>> model.add(Dense(8,kernel_initializer='uniform',activation='relu'))
>>> model.add(Dense(1,kernel_initializer='uniform',activation='sigmoid'))
```

##### Multi-Class Classification

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

##### Regression

```
>>> 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.layers 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.summary()
>>> 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 Fine-tuning

#### Optimization Parameters

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

#### Early Stopping

```
>>> from keras.callbacks import EarlyStopping
>>> early_stopping_monitor = EarlyStopping(patience=2)
>>> model3.fit(x_train4,
              y_train4,
              batch_size=32,
              epochs=15,
              validation_data=(x_test4,y_test4),
              callbacks=[early_stopping_monitor])
```

### Compile Model

#### MLP: Binary Classification

```
>>> model.compile(optimizer='adam',
                 loss='binary_crossentropy',
                 metrics=['accuracy'])
```

#### MLP: Multi-Class Classification

```
>>> model.compile(optimizer='rmsprop',
                 loss='categorical_crossentropy',
                 metrics=['accuracy'])
```

#### MLP: Regression

```
>>> model.compile(optimizer='rmsprop',
                 loss='mse',
                 metrics=['mae'])
```

#### Recurrent Neural Network

```
>>> model3.compile(loss='binary_crossentropy',
                  optimizer='adam',
                  metrics=['accuracy'])
```

### Save/ Reload Models

```
>>> from keras.models import load_model
>>> model3.save('model_file.h5')
>>> my_model = load_model('my_model.h5')
```

### Model Training

```
>>> model3.fit(x_train4,
              y_train4,
              batch_size=32,
              epochs=15,
              verbose=1,
              validation_data=(x_test4,y_test4))
```

### Evaluate Your Model's Performance

```
>>> score = model3.evaluate(x_test,
                           y_test,
                           batch_size=32)
```

### 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,
                                                         y,
                                                         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

BecomingHuman.AI



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  
data type

```
>>> 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  
types

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

	Belgium	Capital	Population
0	Belgium	Brussels	11190846
1	India	New Delhi	1303171035
2	Brazil	Brasilia	207847528

## Dropping

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

## Sort & Rank

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

## Retrieving Series/ DataFrame Information

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

### Summary

```
>>> df.sum()                   Sum of values  
>>> df.cumsum()               Cumulative 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  
-5  
>>> df[1:]                    Get subset of a DataFrame  
Country Capital Population  
1 India New Delhi 1303171035  
2 Brazil Brasilia 207847528
```

### Selecting, Boolean Indexing & Setting

```
By Position  
>>> df.iloc[[0],[0]]          Select single value by row &  
'Belgium' column  
>>> df.iat[[0],[0]]           Select single value by row &  
'Belgium' column labels
```

```
By Label  
>>> df.loc[[0], ['Country']]   Select single value by row &  
'Belgium' column labels  
>>> df.at[[0], ['Country']]    Select single value by row &  
'Belgium' column labels
```

```
By Label/Position  
>>> df.ix[2]                  Select single row of  
Country Brazil subset of rows  
Capital Brasilia  
Population 207847528  
>>> df.ix[:, 'Capital']        Select a single column of  
0 Brussels subset of columns  
1 New Delhi  
2 Brasilia  
>>> df.ix[1, 'Capital']        Select rows and columns  
'New Delhi'
```

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

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

## Asking For Help

```
>>> help(pd.Series.loc)
```

## Applying Functions

```
>>> f = lambda x: x*2  
>>> df.apply(f)                Apply function  
>>> 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  
c 5.0  
d 7.0
```

### 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  
c 5.0  
d 7.0  
>>> s.sub(s3, fill_value=2)  
>>> s.div(s3, fill_value=4)
```

## I/O

### Read and Write to CSV

```
>>> pd.read_csv('file.csv', header=None, nrows=5)  
>>> df.to_csv('myDataFrame.csv')
```

### Read and Write to Excel

```
>>> pd.read_excel('file.xlsx')  
>>> pd.to_excel('dir/myDataFrame.xlsx', sheet_name='Sheet1')
```

**Read multiple sheets from the same file**

```
>>> xlsx = pd.ExcelFile('file.xls')  
>>> df = pd.read_excel(xlsx, 'Sheet1')
```

### Read and Write to SQL Query or Database Table

```
>>> from sqlalchemy import create_engine  
>>> engine = create_engine('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.AI

### Pandas Data Structures

#### Pivot

```
>>> df3 = df2.pivot(index='Date',
                    columns='Type',
                    values='Value')
```

Spread rows into columns

	Date	Type	Value
0	2016-03-01	a	11.432
1	2016-03-02	b	13.031
2	2016-03-01	c	20.784
3	2016-03-03	a	99.906
4	2016-03-02	a	1.303
5	2016-03-03	c	20.784

	Type	a	b	c
2016-03-01		11.432	NaN	20.784
2016-03-02		1.303	13.031	NaN
2016-03-03		99.906	NaN	20.784

#### Pivot Table

```
>>> df4 = pd.pivot_table(df2,
                        values='Value',
                        index='Date',
                        columns='Type')
```

Spread rows into columns

		0	1
1	5	0.233482	0.390959
2	4	0.184713	0.237102
3	3	0.433522	0.429401

Unstacked

		1	5	0	0.233482
				1	0.390959
		2	4	0	0.184713
				1	0.237102
		3	3	0	0.433522
				1	0.429401

Stacked

#### Melt

```
>>> pd.melt(df2,
            id_vars=['Date'],
            value_vars=['Type', 'Value'],
            value_name='Observations')
```

Gather columns into rows

	Date	Type	Value
0	2016-03-01	a	11.432
1	2016-03-02	b	13.031
2	2016-03-01	c	20.784
3	2016-03-03	a	99.906
4	2016-03-02	a	1.303
5	2016-03-03	c	20.784

	Date	Variable	Observations
0	2016-03-01	Type	a
1	2016-03-02	Type	b
2	2016-03-01	Type	c
3	2016-03-03	Type	a
4	2016-03-02	Type	a
5	2016-03-03	Type	c
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

#### Selecting

```
>>> df3.loc[:,(df3>1).any()]
>>> df3.loc[:,(df3>1).all()]
>>> df3.loc[:,df3.isnull().any()]
>>> df3.loc[:,df3.notnull().all()]
```

Select cols with any vals > 1  
Select cols with vals > 1  
Select cols with NaN  
Select cols without NaN

#### Indexing With isin

```
>>> df[(df.Country.isin(df2.Type))]
>>> df3.filter(items=['a','b'])
>>> df.select(lambda x: not x%5)
```

Find same elements  
Filter on values  
Select specific elements

#### Where

```
>>> s.where(s > 0)
```

Subset the data

#### Query

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

Query DataFrame

#### Setting/Resetting Index

```
>>> df.set_index('Country')
>>> df4 = df.reset_index()
>>> df = df.rename(index=str,
                  columns={'Country':'cntry',
                           'Capital':'cptl',
                           'Population':'ppltn'})
```

Set the index  
Reset the index  
Rename DataFrame

#### Reindexing

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

#### Forward Filling

```
>>> df.reindex(range(4),
               method='ffill')
```

```
>>> s3 = s.reindex(range(5),
                   method='bfill')
```

```
Country Capital Population
0 Belgium Brussels 11190846
1 India New Delhi 1303171035
2 Brazil Brasilia 207847528
3 Brazil Brasilia 207847528
```

```
0 3
1 3
2 3
3 3
```

#### Multindexing

```
>>> 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.drop_duplicates('Type', keep='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		data2	
X1	X2	X1	X3
a	11.432	a	20.784
b	1.303	b	NaN
c	99.906	d	20.784

#### Pivot

```
>>> pd.merge(data1,
            data2,
            how='left',
            on='X1')
```

	X1	X2	X3
a	a	11.432	20.784
b	b	1.303	NaN
c	c	99.906	NaN

```
>>> pd.merge(data1,
            data2,
            how='right',
            on='X1')
```

	X1	X2	X3
a	a	11.432	20.784
b	b	1.303	NaN
d	d	NaN	20.784

```
>>> pd.merge(data1,
            data2,
            how='inner',
            on='X1')
```

	X1	X2	X3
a	a	11.432	20.784
b	b	1.303	NaN

```
>>> pd.merge(data1,
            data2,
            how='outer',
            on='X1')
```

	X1	X2	X3
a	a	11.432	20.784
b	b	1.303	NaN
c	c	99.906	NaN
d	d	NaN	20.784

#### 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,
                              freq='M')
>>> 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

## BecomingHuman.AI

### Syntax Creating DataFrames

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

		a	b	c
n	v			
d	1	4	7	10
	2	5	8	11
e	2	6	9	12

```
df = pd.DataFrame(  
    {'a': [4, 5, 6],  
     'b': [7, 8, 9],  
     'c': [10, 11, 12]},  
    index = pd.MultiIndex.from_tuples(  
        [(1, 'd'), (2, 'e')],  
        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.plot.hist()**  
Histogram for each column



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



### Tidy Data A foundation for wrangling in pandas

In a tidy data set:

Each variable is saved in its own column

&

Each 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

$M * A = F$

### Reshaping Data Change the layout of a data set

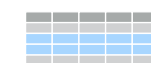
**pd.melt(df)**  
Gather columns into rows.

**df.pivot(columns='var', values='val')**  
Spread rows into columns.

**pd.concat([df1, df2])**  
Append rows of DataFrames

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

### 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	is	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	&[1,~^,df.any(),df.all]	Logical and, or, not, xor, any, all

### Windows



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

Additional GroupBy functions:

**size()** Size of each group.  
**agg(function)** 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'.

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].

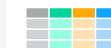
### Summarise Data

**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, 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.

### Combine Data Sets

**ydf + zdf**

#### Set Operations

**pd.merge(ydf, zdf)**  
Rows that appear in both ydf and zdf (Intersection).

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

**pd.merge(ydf, zdf, how='outer', indicator=True)**  
**.query('merge == "left\_only"')**  
**.drop(columns=['\_merge'])**  
Rows that appear in ydf but not zdf (Setdiff)

**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

### Handling Missing Data

**df.dropna()**  
Drop rows with any column having NA/null data.

**df.fillna(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.

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

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

**abs()**  
Absolute value.

#### Standard Joins

**pd.merge(adf, bdf, how='left', on='x1')**  
Join matching rows from bdf to adf.

**pd.merge(adf, bdf, how='right', on='x1')**  
Join matching rows from adf to bdf.

**pd.merge(adf, bdf, how='inner', on='x1')**  
Join data. Retain only rows in both sets.

**pd.merge(adf, bdf, 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)]**  
All rows in adf that do not have a match in bdf

# Data Wrangling with dplyr and tidyr

## Cheat Sheet

### Becoming Human.AI

#### Syntax Helpful conventions for wrangling

##### **dplyr::tbl\_df(iris)**

Converts data to tbl class. tbl's 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
1           5.1           3.5           1.4
2           4.9           3.0           1.4
3           4.7           3.2           1.3
4           4.6           3.1           1.5
5           5.0           3.6           1.4
...           ...           ...
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)

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa

##### **dplyr::%>%**

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

**x %>% f(y)** is the same as **f(x, y)**

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

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

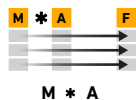
```
iris %>%
  group_by(Species) %>%
  summarise(avg = mean(Sepal.Width)) %>%
  arrange(avg)
```

#### 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



#### Reshaping Data Change the layout of a data set



**tidyr::gather(cases, "year", "n", 2:4)**  
Gather columns into rows.



**tidyr::spread(pollution, size, amount)**  
Spread rows into columns



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



**tidyr::unite(data, col, ..., sep)**  
Unite several columns into one.

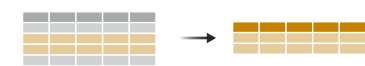
**dplyr::data\_frame(a = 1:3, b = 4:6)**  
Combine vectors into data frame (optimized).

**dplyr::arrange(mtcars, mpg)**  
Order rows by values of a column (low to high).

**dplyr::arrange(mtcars, desc(mpg))**  
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)



**dplyr::filter(iris, Sepal.Length > 7)**  
Extract rows that meet logical criteria.

**dplyr::distinct(iris)**  
Remove duplicate rows.

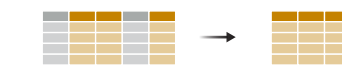
**dplyr::sample\_frac(iris, 0.5, replace = TRUE)**  
Randomly select fraction of rows.

**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).

#### Subset Variables (Columns)



**dplyr::select(iris, Sepal.Width, Petal.Length, Species)**  
Select columns by name or helper function.

##### Helper functions for select - ?select

**select(iris, contains(" "))**  
Select columns whose name contains a character string.

**select(iris, ends\_with("Length"))**  
Select columns whose name ends with a character string.

**select(iris, everything())**  
Select every column.

**select(iris, matches("1.1"))**  
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, Sepal.Length:Petal.Width)**  
Select all columns between Sepal.Length and Petal.Width (inclusive).

**select(iris, -Species)**  
Select all columns except Species.

#### Group Data

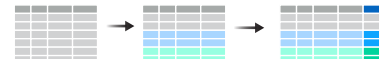
**dplyr::group\_by(iris, Species)**  
Group data into rows with the same value of Species.

**iris %>% group\_by(Species) %>% summarise(...)**  
Compute separate summary row for each group.

**dplyr::ungroup(iris)**  
Remove grouping information from data frame.



**iris %>% group\_by(Species) %>% mutate(...)**  
Compute new variables by group.



#### 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:

**dplyr::first**  
First value of a vector.

**dplyr::last**  
Last value of a vector.

**dplyr::nth**  
Nth value of a vector.

**dplyr::n**  
# of values in a vector.

**dplyr::n\_distinct**  
# of distinct values in a vector.

**IQR**  
IQR of a vector

**min**  
Minimum value in a vector.

**max**  
Maximum value in a vector.

**mean**  
Mean value of a vector.

**median**  
Median value of a vector.

**var**  
Variance of a vector.

**sd**  
Standard deviation of a vector.

#### Combine Data Sets



##### Mutating Joins

**dplyr::left\_join(a, b, by = "x1")**  
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\_join(a, b, by = "x1")**  
Join data. Retain all values, all rows.

##### Filtering Joins

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

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

#### Make New Variables



**dplyr::mutate(iris, sepal = Sepal.Length + Sepal.Width)**  
Compute and append one or more new columns.

**dplyr::mutate\_each(iris, funs(min\_rank))**  
Apply window function to each column.

**dplyr::transmute(iris, sepal = Sepal.Length + Sepal.Width)**  
Compute one or more new columns. Drop original columns



Mutate uses **window functions**, functions that take a vector of values and return another vector of values, such as:

**dplyr::lead**  
Copy with values shifted by 1.

**dplyr::lag**  
Copy with values lagged by 1.

**dplyr::dense\_rank**  
Ranks with no gaps.

**dplyr::min\_rank**  
Ranks. Ties get min rank.

**dplyr::percent\_rank**  
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.

**dplyr::cumall**  
Cumulative all

**dplyr::cumany**  
Cumulative any

**dplyr::cummean**  
Cumulative mean

**cumsum**  
Cumulative sum

**cummax**  
Cumulative max

**cummin**  
Cumulative min

**cumprod**  
Cumulative prod

**pmax**  
Element-wise max

**pmin**  
Element-wise min

##### Set Operations

**dplyr::intersect(y, z)**  
Rows that appear in both y and z.

**dplyr::union(y, z)**  
Rows that appear in either or both y and z.

**dplyr::setdiff(y, z)**  
Rows that appear in y but not z.

##### Binding

**dplyr::bind\_rows(y, z)**  
Append z to y as new rows.

**dplyr::bind\_cols(y, z)**  
Append z to y as new columns.  
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

## Becoming Human.AI



### Interacting With NumPy

Also see NumPy

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

#### Index Tricks

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

#### Shape Manipulation

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

#### Polynomials

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

#### Vectorizing Functions

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

#### Type Handling

```
>>> np.real(b)                   Return the real part of the array elements
>>> np.imag(b)>>>                Return the imaginary part of the array elements
np.real_if_close(c,tol=1000)     Return a real array if complex parts close to 0
>>> np.cast['f'](np.pi)         Cast object to a data type
```

#### Other Useful Functions

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

### Linear Algebra

Also see NumPy

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

```
>>> from scipy import linalg, sparse
```

#### Creating Matrices

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

#### Basic Matrix Routines

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

```
Transposition
>>> A.T                           Transpose 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)                 Determinant
```

```
Solving linear problems
>>> linalg.solve(A,b)             Solver for dense matrices
>>> 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
                                   (least-squares solver)
>>> linalg.pinv2(C)               Compute the pseudo-inverse of
                                   a matrix (SVD)
```

#### Creating Matrices

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

#### 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
>>> linalg.expm(A)                Matrix exponential
>>> linalg.expm2(A)               Matrix exponential (Taylor Series)
>>> linalg.expm3(D)               Matrix exponential
                                   (eigenvalue decomposition)
```

```
Logarithm Function
>>> linalg.logm(A)                Matrix logarithm
```

```
Trigonometric Functions
>>> linalg.sinm(D)                Matrix sine
>>> linalg.cosm(D)                Matrix cosine
>>> linalg.tanm(A)                Matrix tangent
```

```
Hyperbolic Trigonometric Functions
>>> linalg.sinhm(D)               Hyperbolic matrix sine
>>> linalg.coshm(D)               Hyperbolic matrix cosine
>>> linalg.tanhm(A)               Hyperbolic matrix tangent
```

```
Matrix Sign Function
>>> np.signm(A)                   Matrix sign function
```

```
Matrix Square Root
>>> linalg.sqrtm(A)               Matrix square root
```

```
Arbitrary Functions
>>> linalg.funm(A, lambda x: x*x) Evaluate matrix function
```

#### Sparse Matrix Routines

```
Inverse
>>> sparse.linalg.inv(l)           Inverse
```

```
Norm
>>> sparse.linalg.norm(l)          Norm
```

```
Solving linear problems
>>> sparse.linalg.spsolve(H,l)     Solver for sparse matrices
```

#### Sparse Matrix Functions

```
>>> sparse.linalg.expm(l)          Sparse matrix exponential
```

#### Decompositions

##### Eigenvalues and Eigenvectors

```
>>> la, v = linalg.eig(A)          Solve ordinary or generalized
                                   eigenvalue problem for
                                   square matrix
>>> l1, l2 = la
>>> v[:,0]                         First eigenvector
>>> v[:,1]                         Second eigenvector
>>> linalg.eigvals(A)              Unpack eigenvalues
```

##### Singular Value Decomposition

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

##### LU Decomposition

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

#### Sparse Matrix Decompositions

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

### Asking For Help

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

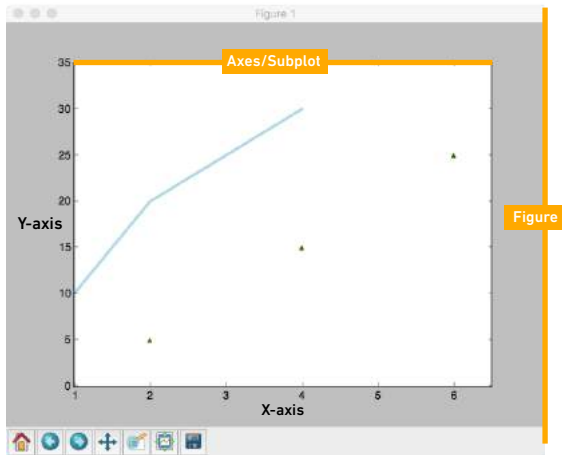
Matplotlib is a Python 2D plotting library which produces publication-quality figures in a variety of hardcopy formats and interactive environments across platforms.

# Matplotlib Cheat Sheet

## BecomingHuman.AI

### Anatomy & Workflow

#### Plot Anatomy



#### Workflow

- 01 Prepare data
- 02 Create plot
- 03 Plot
- 04 Customize plot
- 05 Save plot
- 06 Show plot

```
>>> import matplotlib.pyplot as plt
>>> x = [1,2,3,4]
>>> y = [10,20,25,30]
>>> fig = plt.figure()
>>> ax = fig.add_subplot(111)
>>> ax.plot(x, y, color='lightblue', linewidth=3)
>>> ax.scatter([2.4, 6], [5.15, 25], color='darkgreen', marker='^')
>>> ax.set_xlim(1, 6.5)
>>> plt.savefig('foo.png')
>>> plt.show()
```

### Prepare The Data

Also see Lists & NumPy

#### Index Tricks

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

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

#### Create Plot

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

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

#### Axes

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

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

### Plotting Routines

#### 1D Data

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

```
>>> fig, ax = plt.subplots()
>>> im = ax.imshow(img,
>>>                  arrays cmap='gist_earth',
>>>                  interpolation='nearest',
>>>                  vmin=-2,
>>>                  vmax=2)
```

### Customize Plot

#### Colors, Color Bars & Color Maps

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

#### Mathtext

```
>>> plt.title(r'$\sigma_i=15$', fontsize=20)
```

#### Limits, Legends & Layouts

##### Limits & Autoscaling

```
>>> ax.margins(x=0.0,y=0.1)
>>> ax.axis('equal')
>>> ax.set(xlim=[0,10.5],ylim=[-1.5,1.5])
>>> ax.set_xlim(0,10.5)
```

##### Legends

```
>>> ax.set(title='An Example Axes',
>>>          ylabel='Y-Axis',
>>>          xlabel='X-Axis')
>>> ax.legend(loc='best')
```

##### Ticks

```
>>> ax.xaxis.set(ticks=range(1,5),
>>>               ticklabels=[3,100,-12,'foo'])
>>> ax.yaxis.set(direction='inout',
>>>               length=10)
```

##### Subplot Spacing

```
>>> fig3.subplots_adjust(wspace=0.5,
>>>                       hspace=0.3,
>>>                       left=0.125,
>>>                       right=0.9,
>>>                       top=0.9,
>>>                       bottom=0.1)
```

##### Axes Spines

```
>>> ax1.spines['top'].set_visible(False)
>>> ax1.spines['bottom'].set_position(('outward',10))
```

### Save Plot

#### Save figures

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

#### Show Plot

```
>>> plt.show()
```

### Close & Clear

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

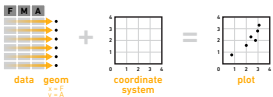


# Data Visualisation with ggplot2

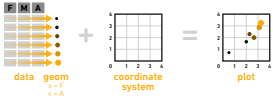
## Cheat Sheet

### 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, map variables in the data set to aesthetic properties of the geom like **size**, **color**, and **x** and **y** locations

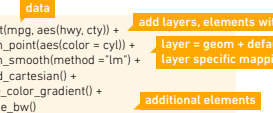


Build a graph with **qplot()** or **ggplot()**

**aesthetic mappings** **data** **geom**

**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 qplot().



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()**

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

**r + coord\_fixed**(ratio = 1/2)  
ratio, xlim, ylim  
Cartesian coordinates with fixed aspect ratio between x and y units

**r + coord\_flip**()  
xlim, ylim  
Flipped Cartesian coordinates

**r + coord\_polar**(theta = "X", direction = 1)  
theta, start, direction  
Polar coordinates

**r + coord\_trans**(trans = "sqrt")  
trans, ylim, xlim, ylim  
Transformed cartesian coordinates. Set extras and strains to the name of a window function.

**r + coord\_map**(projection = "ortho", orientation = c(41, -74, 0))  
projection, orientation, xlim, ylim  
Map projections from the maptools package (mercator (default), azequialera, lagrange, etc.)

### Geoms

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

#### One Variable

**Continuous**

**a <- ggplot(mpg, aes(hwy))**

**a + geom\_area**(stat = "bin")  
x, y, alpha, color, fill, linetype, size  
b = geom\_area(aes(geom\_type = "density"), stat = "bin")

**a + geom\_density**(kernel = "gaussian")  
x, y, alpha, color, fill, linetype, size, weight  
b = geom\_density(aes(geom\_type = "density"))

**a + geom\_dotplot**()  
x, y, alpha, color, fill

**a + geom\_freqpoly**()  
x, y, alpha, color, fill, linetype, size  
b = geom\_freqpoly(aes(geom\_type = "density"))

**a + geom\_histogram**(binwidth = 5)  
x, y, alpha, color, fill, linetype, size, weight  
b = geom\_histogram(aes(geom\_type = "density"))

**Discrete**

**b <- ggplot(mpg, aes(fill))**

**b + geom\_bar**()  
x, alpha, color, fill, linetype, size, weight

#### Graphical Primitives

**c <- ggplot(mpg, aes(long, lat))**

**c + geom\_polygon**(aes(group = group))  
x, y, alpha, color, fill, linetype, size

**d <- ggplot(economics, aes(date, unemployment))**

**d + geom\_area**(lineend = "butt", linejoin = "round", linetype = 1)  
x, y, alpha, color, fill, linetype, size

**d + geom\_ribbon**(aes(ymin = unemployment - 900, ymax = unemployment + 900))  
x, y, alpha, color, fill, linetype, size

**e <- ggplot(economics, aes(date, unemployment))**

**e + geom\_segment**(aes(xend = long + delta, long, yend = lat + delta, lat))  
x, y, alpha, color, fill, linetype, size

**e + geom\_rect**(aes(xmin = long, ymin = lat, xmax = long + delta, long, yend = lat + delta, lat))  
x, y, alpha, color, fill, linetype, size

#### Three Variables

**sealsSz <- with(seals, sqrt(delta\_long^2 + delta\_lat^2))**  
**m <- ggplot(seals, aes(long, lat))**

**m + geom\_contour**(aes(z = z))  
x, y, z, alpha, color, fill, linetype, size, weight

**m + geom\_raster**(aes(fill = z), hjust = 0.5, vjust = 0.5, interpolate = FALSE)  
x, y, alpha, fill

**m + geom\_tile**(aes(fill = z))  
x, y, alpha, color, fill, linetype, size

### Faceting

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

**t <- ggplot(mpg, aes(cty, hwy)) + geom\_point()**

**t + facet\_grid(. ~ fl)**  
facet into columns based on fl

**t + facet\_grid(year ~ .)**  
facet into rows based on year

**t + facet\_grid(year ~ fl)**  
facet into both rows and columns

**t + facet\_wrap(. ~ fl)**  
wrap facets into a rectangular layout one or more discrete variables.

Set **scales** to let axis limits vary across facets

**t + facet\_grid(~ x, scales = "free")**  
x and y 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:f	fl:g	fl:h
a	a	a	a	a	a
c	d	e	p	r	

#### Two Variables

**Continuous X, Continuous Y**

**f <- ggplot(mpg, aes(cty, hwy))**

**f + geom\_blank()**

**f + geom\_jitter**()  
x, y, alpha, color, fill, shape, size

**f + geom\_point**()  
x, y, alpha, color, fill, shape, size

**f + geom\_quantile**()  
x, y, alpha, color, fill, linetype, size, weight

**f + geom\_rug**(sides = "b")  
x, y, alpha, color, linetype, size, weight

**f + geom\_smooth**(model = lm)  
x, y, alpha, color, fill, linetype, size, weight

**f + geom\_text**(aes(label = cty))  
x, y, alpha, color, fill, linetype, size, weight

**Continuous Bivariate Distribution**

**i <- ggplot(movies, aes(year, rating))**

**i + geom\_bin2d**(binwidth = c(5, 0.5))  
x, y, alpha, color, fill, linetype, size, weight

**i + geom\_density2d**()  
x, y, alpha, color, fill, linetype, size

**i + geom\_hex**()  
x, y, alpha, color, fill, size

**Continuous Function**

**j <- ggplot(economics, aes(date, unemployment))**

**j + geom\_area**()  
x, y, alpha, color, fill, linetype, size

**j + geom\_line**()  
x, y, alpha, color, linetype, size

**j + geom\_step**(direction = "hv")  
x, y, alpha, color, linetype, size

#### Discrete X, Continuous Y

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

**g + geom\_bar**(stat = "identity")  
x, y, alpha, color, fill, linetype, size, weight

**g + geom\_boxplot**()  
lower, middle, upper, x, y, alpha, color, fill, linetype, shape, size, weight

**g + geom\_dotplot**(binaxis = "y", stackdir = "center")  
x, y, alpha, color, fill

**g + geom\_violin**(scale = "area")  
x, y, alpha, color, fill, linetype, size, weight

**Visualizing error**

**df <- data.frame(grp = c("A", "B"), fit = 4.5, se = 1.2)**  
**k <- ggplot(df, aes(grp, fit, ymin = fit - se, ymax = fit + se))**

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

**k + geom\_errorbar**()  
x, y, ymax, ymin, alpha, color, fill, linetype, size, width (also geom\_errorbarh())

**k + geom\_linerange**()  
x, y, ymax, ymin, alpha, color, fill, linetype, size

**k + geom\_pointrange**()  
x, y, ymax, ymin, alpha, color, fill, linetype, shape, size

#### Discrete X, Discrete Y

**h <- ggplot(diamonds, aes(cut, color))**

**h + geom\_jitter**()  
x, y, alpha, color, fill, linetype, size

**i <- ggplot(data, aes(map\_id = state), map = map) + expand\_limits(x = map\$long, y = map\$lat)**  
x, y, alpha, color, fill, linetype, 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\_raster**(aes(fill = z), hjust = 0.5, vjust = 0.5, interpolate = FALSE)  
Stack elements on top of one another

**s + geom\_bar**(position = "stack")  
Stack elements on top of one another

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

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

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

### Labels

**t + ggtitle**("New Plot Title")  
Add a main title above the plot

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

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

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

Use scale functions to update legend labels

### Legends

**t + theme**(legend.position = "bottom")  
Place legend at "bottom", "top", "left", or "right"

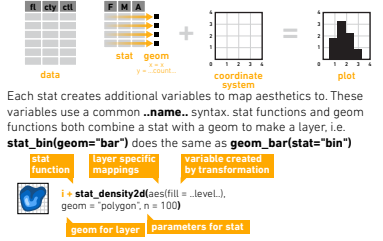
**t + guides**(color = "none")  
Set legend type for each aesthetic: colorbar, legend, or none (no legend)

**t + scale\_fill\_discrete**(name = "Title", labels = c("A", "B", "C"))  
Set legend title and labels with a scale function.

### Stats

An alternative way to build a layer

Some plots visualize a **transformation** of the original data set. Use a **stat** to choose a common transformation to visualize, e.g. **a + geom\_bar**(stat = "bin")



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

**stat** **function** **layer specific mappings** **variable created by transformation** **geom for layer** **parameters for stat**

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

**a + stat\_bin**(binwidth = 1, origin = 10)  
x, y, ..count.., ..ncount.., density, ..ndensity..

**a + stat\_binndot**(binwidth = 1, binaxis = "x")  
x, y, ..count.., ..ncount..

**a + stat\_density**(adjust = 1, kernel = "gaussian")  
x, y, ..count.., density, ..scaled..

**f + stat\_bin2d**(bins = 30, drop = TRUE)  
x, y, fill, ..count.., density, ..scaled..

**f + stat\_binhex**(bins = 30)  
x, y, fill, ..count.., density, ..scaled..

**f + stat\_density2d**(contour = TRUE, n = 100)  
x, y, color, size, ..level..

**m + stat\_contour**(aes(z = z))  
x, y, z, order, ..level..

**m + stat\_spoke**(aes(radius = z, angle = z))  
angle, radius, x, yend, y, yend, ..xend.., ..yend..

**m + stat\_summary\_hex**(aes(z = z), bins = 30, fun = mean)  
x, y, z, fill, ..value..

**g + stat\_boxplot**(coef = 1.5)  
x, y, ..lower.., ..middle.., ..upper.., ..outliers..

**g + stat\_ydensity**(adjust = 1, kernel = "gaussian", scale = "area")  
x, y, ..density.., ..scaled.., ..count.., ..n.., ..violinwidth.., ..width..

**f + stat\_ecdf**(n = 40)  
x, y, ..x.., ..y..

**f + stat\_quantile**(quantiles = c(0.25, 0.5, 0.75), formula = y ~ log(x), method = "rq")  
x, y, ..quantile.., ..x.., ..y..

**f + stat\_smooth**(method = "auto", formula = y ~ x, se = TRUE, n = 80, fullrange = FALSE, level = 0.95)  
x, y, ..se.., ..x.., ..y.., ..ymin.., ..ymax..

**f + stat\_ecdf**(n = 40)  
x, y, ..x.., ..y..

**f + stat\_quantile**(quantiles = c(0.25, 0.5, 0.75), formula = y ~ log(x), method = "rq")  
x, y, ..quantile.., ..x.., ..y..

**f + stat\_smooth**(method = "auto", formula = y ~ x, se = TRUE, n = 80, fullrange = FALSE, level = 0.95)  
x, y, ..se.., ..x.., ..y.., ..ymin.., ..ymax..

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

**f + stat\_identity**()  
values = c(3, 7)  
Shape values shown in chart on right

**ggplot()** + **stat\_qq**(aes(sample = 1:100), distribution = qt, dparams = list(df = 5))  
sample, x, y, ..x.., ..y..

**f + stat\_sum**()  
x, y, size, ..size..

**f + stat\_summary**(fun.data = "mean\_cl\_boot")  
x, y, ..x.., ..y..

**f + stat\_unique**()

### Themes

**r + theme\_bw**()  
White background with grid lines

**r + theme\_classic**()  
White background no gridlines

**r + theme\_grey**()  
Grey background (default theme)

**r + theme\_minimal**()  
Minimal theme

**ggthemes** - Package with additional ggplot2 themes

### Scales

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



**General Purpose scales**

Use with any aesthetic: alpha, color, fill, linetype, shape, size

**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 manually chosen visual values

#### X and Y location scales

Use with x or y aesthetics (x shown here)

**scale\_x\_date**(labels = date\_format("%m/%d"), breaks = date\_breaks("2 weeks"))  
~ treat x values as dates. See ?strptime for label formats.

**scale\_x\_datetime**() ~ treat x values as dates times. Use same arguments as scale\_x\_date()

**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

Use with x or y aesthetics (x shown here)

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

**n + scale\_fill\_brewer**(palette = "Blues")  
For palette choices: library(RColorBrewer) display.brewer.all()

**n + scale\_fill\_grey**(start = 0.2, end = 0.8, na.value = "red")

**o <- a + geom\_dotplot**(aes(fill = ..x..))

**o + scale\_fill\_gradient**(low = "red", high = "yellow")

**o + scale\_fill\_gradient2**(low = "white", height = "blue", mid = "white", midpoint = 25)

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

#### Shape scales

**p <- f + geom\_point**(aes(shape = fl))

**p + scale\_shape**(solid = FALSE)

**p + scale\_shape\_manual**(values = c(3, 7))  
Shape values shown in chart on right

**Size scales**

**q <- f + geom\_point**(aes(size = cyl))

**q + scale\_size\_area**(max = 6)  
Value mapped to area of circle (not radius)

### Zooming

**Without clipping (preferred)**

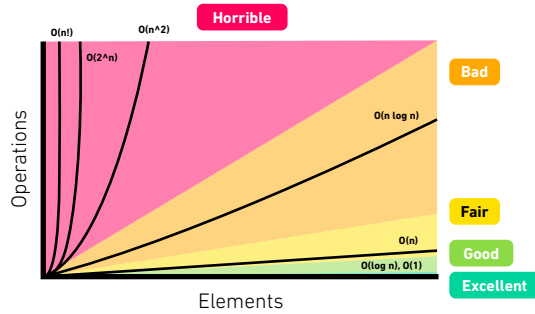
**t + coord\_cartesian**(xlim = c(0, 100), ylim = c(10, 20))

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

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

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

## Big-O Complexity Chart



## Data Structure Operation

	Time Complexity				Space Complexity			
	Average		Worst		Worst			
	Access	Search	Insertion	Deletion	Access	Search	Insertion	Deletion
Array	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$
Stack	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$
Queue	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$
Singly-Linked List	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$
Doubly-Linked List	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$	$\Theta(n)$	$\Theta(n)$	$\Theta(1)$	$\Theta(1)$
Skip List	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$	$\Theta(n \log(n))$
Hash Table	N/A	$\Theta(1)$	$\Theta(1)$	$\Theta(1)$	N/A	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$
Binary Search Tree	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$
Cartesian Tree	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	N/A	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$
B-Tree	N/A	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$
Red-Black Tree	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$
Splay Tree	N/A	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	N/A	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$
AVL Tree	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$
KD Tree	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(\log(n))$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$	$\Theta(n)$

## Array Sorting Algorithms

	Time Complexity			Space Complexity
	Best	Average	Worst	Worst
Quicksort	$\Omega(n \log(n))$	$\Theta(n \log(n))$	$O(n^2)$	$O(n \log(n))$
Mergesort	$\Omega(n \log(n))$	$\Theta(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$
Timsort	$\Omega(n)$	$\Theta(n \log(n))$	$O(n \log(n))$	$\Theta(n)$
Heapsort	$\Omega(n \log(n))$	$\Theta(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$
Bubble Sort	$\Omega(n)$	$\Theta(n^2)$	$O(n^2)$	$\Theta(n)$
Insertion Sort	$\Omega(n)$	$\Theta(n^2)$	$O(n^2)$	$\Theta(n)$
Selection Sort	$\Omega(n^2)$	$\Theta(n^2)$	$O(n^2)$	$\Omega(n^2)$
Tree Sort	$\Omega(n \log(n))$	$\Theta(n \log(n))$	$O(n^2)$	$O(n \log(n))$
Shell Sort	$\Omega(n \log(n))$	$\Theta(n(\log(n))^2)$	$O(n(\log(n))^2)$	$O(n \log(n))$
Bucket Sort	$\Omega(n+k)$	$\Theta(n+k)$	$O(n^2)$	$\Omega(n+k)$
Radix Sort	$\Omega(n+k)$	$\Theta(n+k)$	$\Omega(n+k)$	$\Omega(n+k)$
Counting Sort	$\Omega(n+k)$	$\Theta(n+k)$	$\Omega(n+k)$	$\Omega(n+k)$
Cubesort	$\Omega(n)$	$\Theta(n \log(n))$	$O(n \log(n))$	$O(n \log(n))$