### CHEATSHEET







### **Supervised Learning**

- · Decision Tree · Random Forest
- ·kNN Logistic Regression

### **Unsupervised Learning**

- · Apriori algorithm · k-means
- · Hierarchical Clustering

### **Reinforcement Learning**

- Markov Decision Process
- Q Learning

### **Python** Code

### Code

#Import Library

#Import other necessary libraries like pandas,

from sklearn import linear\_model #Load Train and Test datasets

#Identify feature and response variable(s) and #values must be numeric and numpy arrays

x\_train=input\_variables\_values\_training\_datasets y\_train=target\_variables\_values\_training\_datasets

x test=input variables values test datasets

#Create linear regression object

linear = linear\_model.LinearRegression()

#Train the model using the training sets and

#check score

linear.fit(x train, y train)

linear.score(x\_train, y\_train)

#Equation coefficient and Intercept print('Coefficient: \n', linear.coef\_)

print('Intercept: \n', linear.intercept\_)

#Predict Output

predicted= linear.predict(x\_test)

#Load Train and Test datasets

#Identify feature and response variable(s) and #values must be numeric and numpy arrays

x\_train <- input\_variables\_values\_training\_datasets</pre>

y train <- target variables values training datasets

x\_test <- input\_variables\_values\_test\_datasets x <- cbind(x\_train,y\_train)</pre>

#Train the model using the training sets and #check score

linear  $\leftarrow$  lm(y train  $\sim$  ., data = x)

summary(linear)

#Predict Output

predicted= predict(linear,x\_test)

#Import Library

from sklearn.linear\_model import LogisticRegression #Assumed you have, X (predictor) and Y (target)

x <- cbind(x\_train,y\_train)

#Train the model using the training sets and check #score

### Logistic Regression

```
#for training data set and x_test(predictor)
#of test_dataset
#Create logistic regression object
model = LogisticRegression()
#Train the model using the training sets
#and check score
model.fit(X, y)
model.score(X, y)
#Equation coefficient and Intercept
print('Coefficient: \n', model.coef_)
print('Intercept: \n', model.intercept_)
#Predict Output
predicted= model.predict(x_test)
```

```
logistic <- glm(y_train ~ ., data = x,family='binomial')
summary(logistic)
#Predict Output
predicted= predict(logistic,x_test)</pre>
```

# ecision Tree

```
#Import Library
#Import other necessary libraries like pandas, numpy...
from sklearn import tree
#Assumed you have, X (predictor) and Y (target) for
#training data set and x_test(predictor) of
#test_dataset
#Create tree object
model = tree.DecisionTreeClassifier(criterion='gini')
#for classification, here you can change the
#algorithm as gini or entropy (information gain) by
#default it is gini
#model = tree.DecisionTreeRegressor() for
#regression
#Train the model using the training sets and check
#score
model.fit(X, y)
model.score(X, y)
#Predict Output
predicted= model.predict(x_test)
```

```
#Import Library
library(rpart)
x <- cbind(x_train,y_train)
#grow tree
fit <- rpart(y_train ~ ., data = x,method="class")
summary(fit)
#Predict Output
predicted= predict(fit,x_test)</pre>
```

# SVM (Support Vector Machine)

```
#Import Library
from sklearn import svm
#Assumed you have, X (predictor) and Y (target) for
#training data set and x_test(predictor) of test_dataset
#Create SVM classification object
model = svm.svc()
#there are various options associated
with it, this is simple for classification.
#Train the model using the training sets and check
#score
model.fit(X, y)
model.score(X, y)
#Predict Output
predicted= model.predict(x_test)
```

```
#Import Library
library(e1071)
x <- cbind(x_train,y_train)
#Fitting model
fit <-svm(y_train ~ ., data = x)
summary(fit)
#Predict Output
predicted= predict(fit,x_test)</pre>
```

#Import Library from sklearn.naive\_bayes import GaussianNB

#Import Library
library(e1071)

## Vaive Baye

```
#Assumed you have, X (predictor) and Y (target) for
#training data set and x_test(predictor) of test_dataset
#Create SVM classification object model = GaussianNB()
#there is other distribution for multinomial classes
like Bernoulli Naive Bayes
#Train the model using the training sets and check
#score
model.fit(X, y)
#Predict Output
predicted= model.predict(x_test)
```

```
x <- cbind(x_train,y_train)
#Fitting model
fit <-naiveBayes(y_train ~ ., data = x)
summary(fit)
#Predict Output
predicted= predict(fit,x_test)</pre>
```

# kNN (k- Nearest Neighbors)

```
#Import Library
from sklearn.neighbors import KNeighborsClassifier
#Assumed you have, X (predictor) and Y (target) for
#training data set and x_test(predictor) of test_dataset
#Create KNeighbors classifier object model
KNeighborsClassifier(n_neighbors=6)
#default value for n_neighbors is 5
#Train the model using the training sets and check score
model.fit(X, y)
#Predict Output
predicted= model.predict(x_test)
```

```
#Import Library
library(knn)
x <- cbind(x_train,y_train)
#Fitting model
fit <-knn(y_train ~ ., data = x,k=5)
summary(fit)
#Predict Output
predicted= predict(fit,x_test)</pre>
```

### -Means

```
#Import Library
from sklearn.cluster import KMeans
#Assumed you have, X (attributes) for training data set
#and x_test(attributes) of test_dataset
#Create KNeighbors classifier object model
k_means = KMeans(n_clusters=3, random_state=0)
#Train the model using the training sets and check score
model.fit(X)
#Predict Output
predicted= model.predict(x_test)
```

```
#Import Library
library(cluster)
fit <- kmeans(X, 3)
#5 cluster solution</pre>
```

## Random Forest

```
#Import Library
from sklearn.ensemble import RandomForestClassifier
#Assumed you have, X (predictor) and Y (target) for
#training data set and x_test(predictor) of test_dataset
#Create Random Forest object
model= RandomForestClassifier()
#Train the model using the training sets and check score
model.fit(X, y)
#Predict Output
predicted= model.predict(x_test)
```

```
#Import Library
library(randomForest)
x <- cbind(x_train,y_train)
#Fitting model
fit <- randomForest(Species ~ ., x,ntree=500)
summary(fit)
#Predict Output
predicted= predict(fit,x_test)</pre>
```

## **Algorithms**

#Import Library
from sklearn import decomposition

#Assumed you have training and test data set as train and
#test

```
#Import Library
library(stats)
pca <- princomp(train, cor = TRUE)
train_reduced <- predict(pca,train)</pre>
```

```
#default value of k =min(n_sample, n_features)
#For Factor analysis
#fa= decomposition.FactorAnalysis()
#Reduced the dimension of training dataset using PCA
train_reduced = pca.fit_transform(train)
#Reduced the dimension of test dataset
test_reduced = pca.transform(test)
```

#Create PCA object pca= decomposition.PCA(n components=k)

```
test_reduced <- predict(pca,test)
```

# Gradient Boosting & AdaBoost

#Predict Output

predicted= model.predict(x\_test)

```
#Import Library
library(caret)
x <- cbind(x_train,y_train)
#Fitting model
fitControl <- trainControl( method = "repeatedcv",
+ number = 4, repeats = 4)
fit <- train(y ~ ., data = x, method = "gbm",
+ trControl = fitControl,verbose = FALSE)
predicted= predict(fit,x_test,type= "prob")[,2]</pre>
```