

Intelligent Systems

1. Decision Tree
2. Naïve Bayes
3. Linear Discriminant Analysis
4. Ensemble Learning
 - Bagging: Random Forest
 - Boosting:
 - Gradient Boosting
 - Ada Boosting
 - Stacking

Decision Tree

Regression algorithm

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

dataset = pd.read_csv("F:/Intelligent systems/petrol_consumption.csv") #change
dataset

dataset.shape

dataset.head(20) #show first 20 raw

X = dataset.drop('Petrol_Consumption', axis=1)
y = dataset['Petrol_Consumption']


from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.20,
random_state=1)

X_train


from sklearn.tree import DecisionTreeRegressor

regressor = DecisionTreeRegressor ()

regressor.fit(X_train, y_train)

y_pred = regressor.predict(X_test)

df= pd.DataFrame({'Actual':y_test, 'Predicted':y_pred})
```

```
df

from sklearn import metrics

print('Mean Absolute Error:', metrics.mean_absolute_error(y_test, y_pred))

print('Mean Squared Error:', metrics.mean_squared_error(y_test, y_pred))

print('Root Mean Squared Error:', np.sqrt(metrics.mean_squared_error(y_test,
y_pred)))
```

Classification algorithm

Load libraries

```
import pandas as pd

from sklearn.tree import DecisionTreeClassifier # Import Decision Tree Classifier

from sklearn.model_selection import train_test_split # Import train_test_split
function

from sklearn import metrics #Import scikit-learn metrics module for accuracy
calculation
```

load dataset

```
pima = pd.read_csv("F:\Intelligent systems\diabetes.csv")

pima.head()
```

#split dataset in features and target variable

```
X = pima.drop('Outcome', axis=1)

y = pima['Outcome']
```

Split dataset into training set and test set

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3,  
random_state=1) # 70% training
```

```
from sklearn.tree import DecisionTreeClassifier # Import Decision Tree Classifier
```

Create Decision Tree classifier object

```
clf = DecisionTreeClassifier ()
```

Train Decision Tree Classifier

```
clf = clf.fit(X_train,y_train)
```

#Predict the response for test dataset

```
y_pred = clf.predict(X_test)
```

```
y_pred
```

Model Accuracy, how often is the classifier correct?

```
print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
```

Create Decision Tree classifier object

```
clf = DecisionTreeClassifier(criterion="entropy", max_depth=3)
```

Train Decision Tree Classifier

```
clf = clf.fit(X_train,y_train)
```

#Predict the response for test dataset

```
y_pred = clf.predict(X_test)
```

Model Accuracy, how often is the classifier correct?

```
print("Accuracy:",metrics.accuracy_score(y_test, y_pred))
```

```
df=pd.DataFrame({'Actual':y_test, 'Predicted':y_pred})
```

```
df
```

```
*****
```

Naive Bayes

load the iris dataset

```
from sklearn.datasets import load_digits
```

```
iris = load_digits ()
```

store the feature matrix (X) and response vector (y)

```
X = iris.data
```

```
y = iris.target
```

splitting X and y into training and testing sets

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split (X, y, test_size=0.4,  
random_state=1)
```

training the model on training set

```
from sklearn.naive_bayes import GaussianNB
```

```
gnb = GaussianNB ()
```

```
gnb.fit(X_train, y_train)
```

making predictions on the testing set

```
y_pred = gnb.predict(X_test)
```

```
y_pred [:20]
```

```
y_test[:20]
```

```
# comparing actual response values (y_test) with predicted response values  
(y_pred)
```

```
from sklearn import metrics
```

```
print("Gaussian Naive Bayes model accuracy(in %):",  
      metrics.accuracy_score(y_test, y_pred)*100)
```

LDA

#Import Libraries

```
from sklearn.datasets import load_breast_cancer  
from sklearn.model_selection import train_test_split  
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis  
from sklearn.metrics import confusion_matrix  
import seaborn as sns  
import matplotlib.pyplot as plt
```

#load breast cancer data

```
BreastData = load_breast_cancer()
```

#X Data

```
X = BreastData.data
```

#y Data

```
y = BreastData.target
```

#Splitting data

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.33,  
random_state=44)
```



```
LDAModel = LinearDiscriminantAnalysis ()
```

```
LDAModel.fit (X_train, y_train)
```

#Calculating Details

```
print('LDAModel Train Score is : ', LDAModel.score(X_train, y_train))
```

```
print('LDAModel classea are : ', LDAModel.classes_)
```

```
print('-----')
```

```
y_pred = LDAModel.predict(X_test)
```

```
print('Predicted Value for LDAModel is : ', y_pred[:20])
```

#Calculating Confusion Matrix

```
CM = confusion_matrix(y_test, y_pred)
```

```
print('Confusion Matrix is : \n', CM)
```

drawing confusion matrix

```
sns.heatmap(CM, center = True)
```

```
plt.show()
```

Bagging:

Random Forest

Regression

#Import Libraries

```
from sklearn.datasets import load_boston  
from sklearn.model_selection import train_test_split  
from sklearn.ensemble import RandomForestRegressor  
from sklearn.metrics import mean_absolute_error  
from sklearn.metrics import mean_squared_error  
from sklearn.metrics import median_absolute_error
```

#load boston data

```
BostonData = load_boston()
```

#X Data

```
X = BostonData.data  
print ('X Data is \n', X[:10])  
#print ('X shape is ', X.shape)  
#print ('X Features are \n', BostonData.feature_names)
```

#y Data

```
y = BostonData.target
#print('y Data is \n' , y[:10])
#print('y shape is ' , y.shape)
```

#Splitting data

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.33,
random_state=44, shuffle =True)
```

#Splitted Data

```
#print ('X_train shape is ' , X_train.shape)
#print ('X_test shape is ' , X_test.shape)
#print ('y_train shape is ' , y_train.shape)
#print ('y_test shape is ' , y_test.shape)

RFRModel = RandomForestRegressor (n_estimators=200, max_depth=3,
random_state=33)

RFRModel.fit (X_train, y_train)
```

#Calculating Details

```
print ('Random Forest Regressor Train Score is: ' , RFRModel.score(X_train,
y_train))

print ('Random Forest Regressor Test Score is: ' , RFRModel.score(X_test, y_test))

print ('Random Forest Regressor No. of features are: ' , RFRModel.n_features_)

print ('-----')
```

#Calculating Prediction

```
y_pred = RFRModel.predict(X_test)
print ('Predicted Value for Random Forest Regressor is: ', y_pred [:10])
y_test [:10]
RandomForestRegressorModel.feature_importances_
```

#Calculating Mean Absolute Error

```
MAEValue = mean_absolute_error (y_test, y_pred, multioutput=
'uniform_average')
print ('Mean Absolute Error Value is: ', MAEValue)
```

#Calculating Mean Squared Error

```
MSEValue = mean_squared_error(y_test, y_pred, multioutput='uniform_average')
print('Mean Squared Error Value is : ', MSEValue)
```

#Calculating Median Squared Error

```
MdSEValue = median_absolute_error(y_test, y_pred)
print('Median Squared Error Value is : ', MdSEValue )
```

.....

Classification

#Import Libraries

```
from sklearn.datasets import load_breast_cancer
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt
```

#load breast cancer data

```
BreastData = load_breast_cancer()
```

#X Data

```
X = BreastData.data
#print('X Data is \n' , X[:10])
#print('X shape is ' , X.shape)
#print('X Features are \n' , BreastData.feature_names)
```

#y Data

```
y = BreastData.target
#print('y Data is \n' , y[:10])
```

```
#print('y shape is ' , y.shape)
```

```
#print('y Columns are \n' , BreastData.target_names)
```

#Splitting data

```
X_train, X_test, y_train, y_test = train_test_split (X, y, test_size=0.33,  
random_state=44)
```

```
#print('X_train shape is ' , X_train.shape)
```

```
#print('X_test shape is ' , X_test.shape)
```

```
#print('y_train shape is ' , y_train.shape)
```

```
#print('y_test shape is ' , y_test.shape)
```

```
RFC = RandomForestClassifier (criterion = 'gini', n_estimators=100,  
max_depth=2, random_state=33) #criterion can be also: entropy
```

```
RFC.fit (X_train, y_train)
```

#Calculating Details

```
print(' RFC Train Score is : ' , RFC.score(X_train, y_train))
```

```
print('RandomForestClassifierModel Test Score is : ' , RFC.score(X_test, y_test))
```

```
print('RandomForestClassifierModel features importances are : ' ,  
RFC.feature_importances_)
```

```
print('-----')
```

#Calculating Prediction

```
y_pred = RFC
```

```
.predict(X_test)
```

```
print('Predicted Value for RandomForestClassifierModel is : ', y_pred[:10])
```

```
print('actual Value for RandomForestClassifierModel is : ', y_test[:10])
```

```
#Calculating Confusion Matrix
```

```
CM = confusion_matrix(y_test, y_pred)
```

```
print('Confusion Matrix is : \n', CM)
```

```
# drawing confusion matrix
```

```
sns.heatmap(CM, center = True)
```

```
plt.show()
```

Boosting:

Gradient Boosting

Classification algorithm

Import models and utility functions

```
from sklearn.ensemble import GradientBoostingClassifier
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.metrics import accuracy_score
```

```
from sklearn.datasets import load_digits
```

```
dataset=load_digits ()
```

```
X = dataset.data
```

```
y=dataset.target
```

```
X.shape
```

Splitting dataset

```
train, test_X, train_y, test_y = train_test_split (X, y, test_size = 0.25, random_state  
= 20)
```

Instantiate Gradient Boosting Regressor

```
gbc = GradientBoostingClassifier (n_estimators=300,
```

```
learning_rate=0.02,
```

```
random_state=100, max_features=5)
```


Fit to training set

```
gbc.fit (train_X, train_y)
```

Predict on test set

```
pred_y = gbc.predict(test_X)
```

accuracy

```
acc = accuracy_score(test_y, pred_y)
```

```
print ("Gradient Boosting Classifier accuracy is: {:.2f}". format (acc))
```

=====

Regression Algorithms

Import the necessary libraries

```
from sklearn.ensemble import GradientBoostingRegressor
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.metrics import mean_squared_error
```

```
from sklearn.datasets import load_diabetes
```

```
dataset=load_digits()
```

```
X = dataset.data
```

```
y = dataset.target
```

Splitting dataset

```
train_X, test_X, train_y, test_y = train_test_split (X, y, test_size = 0.25)
```

Instantiate Gradient Boosting Regressor

```
gbr = GradientBoostingRegressor (learning_rate=0.1,  
                                n_estimators=300,  
                                max_depth = 1,  
                                random_state = 30,  
                                max_features = 5)
```

Fit to training set

```
gbr.fit (train_X, train_y)
```

Predict on test set

```
pretty = gbr.predict(test_X)
```

test set RMSE

```
test_rmse = mean_squared_error(test_y, pred_y) ** (1 / 2)
```

Print rmse

```
print ('Root mean Square error: {:.2f}'.format(test_rmse))
```

ADA

```
import pandas as pd

import numpy as np

from sklearn.datasets import load_iris

from sklearn.model_selection import train_test_split

from sklearn.ensemble import AdaBoostClassifier


# Reading the dataset from the csv file

dataset = load_iris ()

X = dataset.data

y = dataset.target

Train, X_val, Y_train, Y_val = train_test_split (X, y, test_size=0.25,
random_state=28)


# Creating adaboost classifier model

adb = AdaBoostClassifier ()

adb_model = adb.fit(X_train,Y_train)

print ("The accuracy of the model on validation set is",
adb_model.score(X_val,Y_val))

y_pred=adb.predict(X_val)

y_pred

y_val
```

Stacking Methods

Regression

```
from sklearn.ensemble import StackingRegressor #Stacking regressor
```

```
from sklearn.linear_model import LinearRegression
```

```
from sklearn.datasets import load_boston
```

```
from sklearn.ensemble import RandomForestRegressor
```

```
from sklearn.tree import DecisionTreeRegressor
```

```
from sklearn.svm import LinearSVR
```

```
BostonData = load_boston()
```

#X Data

```
X = BostonData.data
```

#y Data

```
y = BostonData.target
```

```
estimators = [('Decision tree', DecisionTreeRegressor()),
```

```
              ('Random Forest', RandomForestRegressor()),
```

```
              ('SVR', LinearSVR(random_state=42))]
```

```
from sklearn.model_selection import train_test_split
```

```
stackingreg = StackingRegressor(estimators=estimators,  
final_estimator=LinearRegression())
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=42)
```

```
stackingreg.fit(X_train, y_train)
```

```
y_pred = stackingreg.predict(X_test)
```

#Model Evaluation

```
from sklearn.metrics import mean_absolute_error
```

```
from sklearn.metrics import mean_squared_error
```

```
from sklearn.metrics import median_absolute_error
```

```
print('Mean Absolute Error (MAE):', mean_absolute_error(y_test, y_pred))
```

```
print('Mean Squared Error (MSE):', mean_squared_error(y_test, y_pred))
```

```
print('Mean Squared Log Error:', median_absolute_error(y_test, y_pred))
```

Classification

```
from sklearn.ensemble import StackingClassifier
```

```
from sklearn.linear_model import LogisticRegression
```

```
from sklearn.tree import DecisionTreeClassifier
```

```
from sklearn.svm import SVC
```

```
from sklearn.datasets import load_iris
```

```
IrisData = load_iris()
```

```
#X Data
```

```
X = IrisData.data
```

```
#y Data
```

```
y = IrisData.target
```

```
estimators = [('dtrClassifier', DecisionTreeClassifier()), ('svc', SVC())]
```

```
stackingCLS = StackingClassifier(estimators=estimators,  
final_estimator=LogisticRegression())
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=42)
```

```
stackingCLS.fit(X_train, y_train)
```

```
import sklearn.metrics as accuracy_score
```

```
from sklearn.metrics import confusion_matrix
```

```
y_predict = stackingCLS.predict(X_test) #Get the splitted part for testing to be  
predicted, and check the accuray
```

#Calculating Confusion Matrix

```
CM = confusion_matrix(y_test, y_predict)
```

```
print('Confusion Matrix is : \n', CM)
```

```
print('The Score is : ',stackingCLS.score(X_test , y_test))
```