

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import accuracy_score
import matplotlib.pyplot as plt
from sklearn import datasets
```

```
pip install datawig
```

```
Looking in indexes: https://pypi.org/simple, https://us-python.pkg.dev/colab-wheels/
Requirement already satisfied: datawig in /usr/local/lib/python3.7/dist-packages (0.1.0)
Requirement already satisfied: typing==3.6.6 in /usr/local/lib/python3.7/dist-packages (from datawig) (3.6.6)
Requirement already satisfied: mxnet==1.4.0 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.4.0)
Requirement already satisfied: pandas==0.25.3 in /usr/local/lib/python3.7/dist-packages (from datawig) (0.25.3)
Requirement already satisfied: scikit-learn[alldeps]==0.22.1 in /usr/local/lib/python3.7/dist-packages (from datawig) (0.22.1)
Requirement already satisfied: graphviz<0.9.0,>=0.8.1 in /usr/local/lib/python3.7/dist-packages (from datawig) (0.8.1)
Requirement already satisfied: requests>=2.20.0 in /usr/local/lib/python3.7/dist-packages (from datawig) (2.25.1)
Requirement already satisfied: numpy<1.15.0,>=1.8.2 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.19.5)
Requirement already satisfied: pytz>=2017.2 in /usr/local/lib/python3.7/dist-packages (from datawig) (2018.9)
Requirement already satisfied: python-dateutil>=2.6.1 in /usr/local/lib/python3.7/dist-packages (from datawig) (2.8.1)
Requirement already satisfied: scipy>=0.17.0 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.5.4)
Requirement already satisfied: joblib>=0.11 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.0.1)
Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.16.0)
Requirement already satisfied: urllib3!=1.25.0,!1.25.1,<1.26,>=1.21.1 in /usr/local/lib/python3.7/dist-packages (from datawig) (1.25.11)
Requirement already satisfied: certifi>=2017.4.17 in /usr/local/lib/python3.7/dist-packages (from datawig) (2021.10.8)
Requirement already satisfied: idna<3,>=2.5 in /usr/local/lib/python3.7/dist-packages (from datawig) (2.10)
Requirement already satisfied: chardet<4,>=3.0.2 in /usr/local/lib/python3.7/dist-packages (from datawig) (3.0.4)
```

```
import datawig
```

```
path = "/content/app_data.csv"
```

```
df = pd.read_csv(path)
df
```

	Age	BMI	Sex	Height	Weight	AlvaradoScore	PediatricAppendicitis
0	12.531143	16.494601	male	159.0	41.7	7	
1	12.410678	12.595222	female	152.0	29.1	8	
2	10.537988	15.991247	male	133.5	28.5	3	
3	10.425736	16.185025	male	146.0	34.5	4	
4	13.270363	20.449137	female	164.0	55.0	2	
...

✓ 0s completed at 11:40 PM



426	12.528405	29.316297	male	152.3	68.0	7
427	12.013689	28.906250	male	160.0	74.0	5
428	7.739904	22.038188	female	120.5	32.0	5
429	10.157426	21.017920	female	142.2	42.5	9

430 rows × 41 columns



```
#df.info()
```

```
#column dropping considering y3= AppendicitisComplications
```

```
df.drop(['DiagnosisByCriteria','TreatmentGroupBinar'],axis=1,inplace=True)
```

```
# Ultrasound
```

```
df.drop(['AppendixOnSono','AppendixDiameter','AppendixWallLayers','Kokarde','TissuePerfusi  
'BowelWallThick','Ileus','Enteritis','Peritonitis'],axis=1,inplace=True)
```

```
#df.info()
```

```
df_numerical = df.filter(['Age','BMI','Height','Weight','AlvaradoScore','PediatricAppendic  
'AppendixDiameter','BodyTemp','WBCCount','NeutrophilPerc','CRPEntery'],
```

```
#df_numerical.info()
```

```
df_categorical = df.filter(['Sex','KetonesInUrine','ErythrocytesInUrine','WBCInUrine',  
'Peritonitis','AppendixWallLayers','TissuePerfusion'],axis=1).c
```

```
#df_categorical.info()
```

```
#df_categorical.head()
```

```
df_boolean = df.filter(['AppendixOnSono','MigratoryPain','LowerAbdominalPainRight','Reboun  
'Nausea','AppetiteLoss','Dysuria','FreeFluids','Kokarde',  
'SurroundingTissueReaction','PathLymphNodes','MesentricLymphadenitis',  
'FecalImpaction','Meteorism','Enteritis','AppendicitisComplications',  
'PsoasSign','Stool'],axis=1).copy()
```

```
#df_boolean.info()
```

```
#df_boolean.sample(10)

#pandas profiling
#from pandas_profiling import ProfileReport

#profile = ProfileReport(df)
#profile.to_file(output_file = "AppendicitisComplications_profiling.html")

#perform label Encoding for categorical data

from sklearn.preprocessing import LabelEncoder
from pandas import Series
df_categorical = df_categorical.apply(lambda series:pd.Series(
    LabelEncoder().fit_transform(series[series.notnull()]),
    index = series[series.notnull()].index
))

#df_categorical.info()

#df_categorical.head()

#concatanation two dataframe
df_new = pd.concat([df_numerical,df_categorical],axis=1)

#df_new.info()

# Datawig imputation

from datawig import SimpleImputer

# impute missing values using Datawig
df_dw_imputed = datawig.SimpleImputer.complete(df_new)

#df_dw_imputed.head()

#df_dw_imputed.info()

#df_dw_imputed.isnull()

#perform labelEncoding for Boolean data
```

```
df_boolean = df_boolean.apply(lambda series:pd.Series(
    LabelEncoder().fit_transform(series[series.notnull()]),
    index = series[series.notnull()].index
))
```

```
#df_boolean.head()
```

```
df_boolean = df_boolean.fillna(df_boolean.mode().iloc[0])
```

```
#df_boolean.sample(20)
```

```
#df_boolean.info()
```

```
#concatanation two dataframe
```

```
df_final = pd.concat([df_dw_imputed,df_boolean],axis=1)
```

```
df_final.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 430 entries, 0 to 429
Data columns (total 29 columns):
Age                                430 non-null float64
BMI                                430 non-null float64
Height                            430 non-null float64
Weight                            430 non-null float64
AlvaradoScore                     430 non-null float64
PediatricAppendicitisScore        430 non-null float64
BodyTemp                          430 non-null float64
WBCCount                          430 non-null float64
NeutrophilPerc                    430 non-null float64
CRPEntry                          430 non-null float64
Sex                               430 non-null float64
KetonesInUrine                    430 non-null float64
ErythrocytesInUrine               430 non-null float64
WBCInUrine                        430 non-null float64
MigratoryPain                     430 non-null int64
LowerAbdominalPainRight           430 non-null float64
ReboundTenderness                 430 non-null float64
CoughingPain                      430 non-null float64
Nausea                            430 non-null int64
AppetiteLoss                      430 non-null float64
Dysuria                           430 non-null float64
FreeFluids                        430 non-null float64
PathLymphNodes                    430 non-null float64
MesentricLymphadenitis            430 non-null float64
FecalImpaction                    430 non-null float64
Meteorism                         430 non-null float64
AppendicitisComplications         430 non-null int64
PsoasSign                         430 non-null float64
```

```
Stool                                430 non-null float64
dtypes: float64(26), int64(3)
memory usage: 97.5 KB
```

#correlation and pvalue

```
from scipy import stats
corr_df=pd.DataFrame(columns=['r','p'])

for col in df_final:
    print(col)
    if pd.api.types.is_numeric_dtype(df_final[col]):
        r,p = stats.pearsonr(df_final.AppendicitisComplications,df_final[col])
        corr_df.loc[col]=[round(r,3),round(p,3)]
```

corr_df

Age
BMI
Height
Weight
AlvaradoScore
PediatricAppendicitisScore
BodyTemp
WBCCount
NeutrophilPerc
CRPEntry
Sex
KetonesInUrine
ErythrocytesInUrine
WBCInUrine
MigratoryPain
LowerAbdominalPainRight
ReboundTenderness
CoughingPain
Nausea
AppetiteLoss
Dysuria
FreeFluids
PathLymphNodes
MesentricLymphadenitis
FecalImpaction
Meteorism
AppendicitisComplications
PsoasSign
Stool

	r	p
Age	-0.098	0.043
BMI	-0.069	0.153
Height	-0.084	0.082

Weight	-0.071	0.144
AlvaradoScore	0.279	0.000
PediatricAppendicitisScore	0.255	0.000
BodyTemp	0.285	0.000
WBCCount	0.327	0.000
NeutrophilPerc	0.262	0.000
CRPEntry	0.615	0.000
Sex	-0.020	0.677
KetonesInUrine	-0.102	0.035
ErythrocytesInUrine	-0.180	0.000
WBCInUrine	-0.051	0.289
MigratoryPain	0.065	0.177
LowerAbdominalPainRight	-0.061	0.205
ReboundTenderness	0.069	0.152
CoughingPain	0.053	0.277
Nausea	0.207	0.000
AppetiteLoss	0.145	0.003
Dysuria	0.013	0.792
FreeFluids	0.112	0.021
PathLymphNodes	-0.040	0.403
MesentricLymphadenitis	0.006	0.901
FecallImpaction	0.049	0.311
Meteorism	0.013	0.794
AppendicitisComplications	1.000	0.000
PsoasSign	-0.084	0.082
Stool	-0.112	0.021

```
df_final.shape
```

```
(430, 29)
```

```
df_final['AppendicitisComplications'].value_counts()
```

```
df_final[AppendicitisComplications].value_counts()

0      379
1       51
Name: AppendicitisComplications, dtype: int64
```

1 = yes, 0 = NO

```
no = df_final[df_final.AppendicitisComplications==0]
yes = df_final[df_final.AppendicitisComplications==1]
```

```
print(no.shape)
print(yes.shape)
```

```
(379, 29)
(51, 29)
```

```
#splitting the data for training and testing
```

```
X=df_final.drop(columns='AppendicitisComplications',axis=1)
Y=df_final['AppendicitisComplications']
```

```
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=.2, stratify=Y, random
```

```
print(X.shape)
print(X_train.shape)
print(X_test.shape)
```

```
(430, 28)
(344, 28)
(86, 28)
```

```
print(Y.shape)
print(Y_train.shape)
print(Y_test.shape)
```

```
(430,)
(344,)
(86,)
```

N_estimator_Random Forest classifier

```

from sklearn.ensemble import RandomForestClassifier
forest = RandomForestClassifier(random_state = 1, n_estimators = 10, min_samples_split = 2)
forest.fit(X_train, Y_train)

RandomForestClassifier(bootstrap=True, ccp_alpha=0.0, class_weight=None,
                       criterion='gini', max_depth=None, max_features='auto',
                       max_leaf_nodes=None, max_samples=None,
                       min_impurity_decrease=0.0, min_impurity_split=None,
                       min_samples_leaf=1, min_samples_split=2,
                       min_weight_fraction_leaf=0.0, n_estimators=10,
                       n_jobs=None, oob_score=False, random_state=1, verbose=0,
                       warm_start=False)

model_score2 = forest.score(X_test, Y_test)
model_score1 = forest.score(X_train, Y_train)
print(model_score1)
print(model_score2)

0.9854651162790697
0.9186046511627907

```

Logistic Regression

```

# model training using logistic regression
from sklearn.linear_model import LogisticRegression
model = LogisticRegression()
model.fit(X_train, Y_train)

/usr/local/lib/python3.7/dist-packages/sklearn/linear_model/_logistic.py:940: Conver
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:
https://scikit-learn.org/stable/modules/preprocessing.html
Please also refer to the documentation for alternative solver options:
https://scikit-learn.org/stable/modules/linear\_model.html#logistic-regression
extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
LogisticRegression(C=1.0, class_weight=None, dual=False, fit_intercept=True,
                   intercept_scaling=1, l1_ratio=None, max_iter=100,
                   multi_class='auto', n_jobs=None, penalty='l2',
                   random_state=None, solver='lbfgs', tol=0.0001, verbose=0,
                   warm_start=False)

# accuracy score for training data and testing data
X_train_prediction=model.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)

X_test_prediction=model.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)

```



```
X_testing_accuracy, accuracy_pos, accuracy_neg, accuracy,
```

```
print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data: 0.9186046511627907
Accuracy score for testing data: 0.9418604651162791
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(model , X_train, Y_train, cv = kf)
result
```

```
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https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

```
extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
```

```
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```

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STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9070588235294117
```

```
from sklearn.model_selection import cross_val_score
```

```
from sklearn.model_selection import KFold
```

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

```
k = 10
```

```
kf = KFold(n_splits=k, random_state=None)
```

```
result = cross_val_score(model , X_test, Y_test, cv = kf)
```

```
result
```

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```

```
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STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9319444444444445
```

```
from sklearn import metrics  
import matplotlib.pyplot as plt
```

```
# make predictions
```

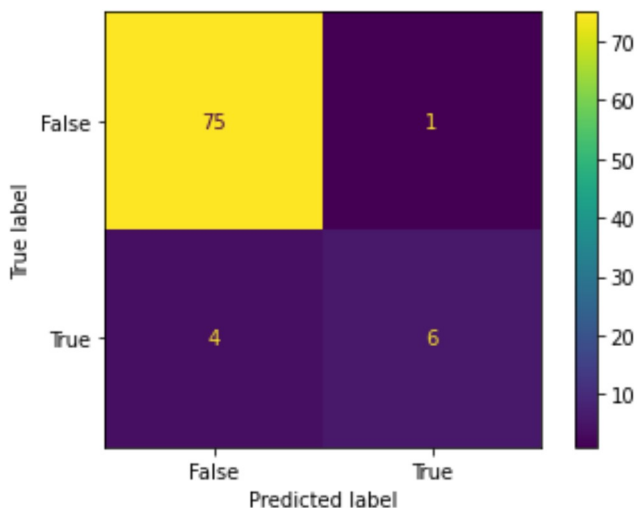
```
predicted = model.predict(X_test)
```

```
from sklearn.metrics import accuracy_score, confusion_matrix
```

```
confusion_matrix = metrics.confusion_matrix(X_test, predicted)
```

```
confusion_matrix = metrics.confusion_matrix(y_test, predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_labels=[0, 1])  
cm_display.plot()  
plt.show()
```



```
TN = confusion_matrix[0][0]  
FN = confusion_matrix[1][0]  
TP = confusion_matrix[1][1]  
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))  
specificity = (TN / float(TN + FP))  
ppv = (TP / float(TP + FP))  
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)  
print("specificity: ",specificity)  
print("PPV: ",ppv)  
print("NPV: ",npv)
```

```
Sensitivity:  0.6  
specificity:  0.9868421052631579  
PPV:  0.8571428571428571  
NPV:  0.9493670886075949
```

```
# AUROC and AUPR value
```

```
from sklearn.metrics import auc, roc_curve, precision_recall_curve
```

```
y_predictProb = model.predict_proba(X_test)
```

```
fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])  
roc_auc = auc(fpr, tpr)
```

```
precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
```

```
area = auc(recall, precision)
```

```
print("AUROC:", roc_auc)
```

```
print("AUPR:", area)
```

```
AUROC: 0.9697368421052631
```

```
AUPR: 0.7758531746031746
```

```
# AUROC graph
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
```

```
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
```

```
plt.xlabel('False Positive Rate')
```

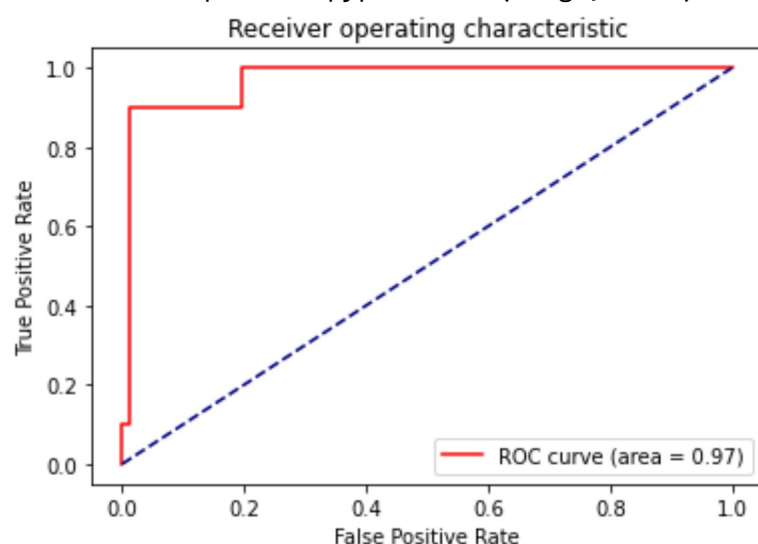
```
plt.ylabel('True Positive Rate')
```

```
plt.title('Receiver operating characteristic')
```

```
plt.legend(loc="lower right")
```

```
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```



```
# AUPR graph
```

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
```

```
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
```

```
plt.xlabel('False Positive Rate')
```

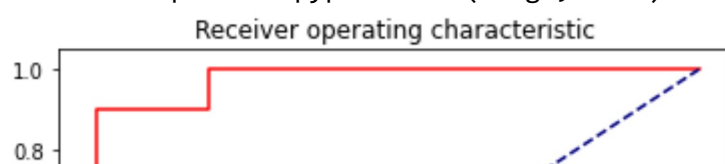
```
plt.ylabel('True Positive Rate')
```

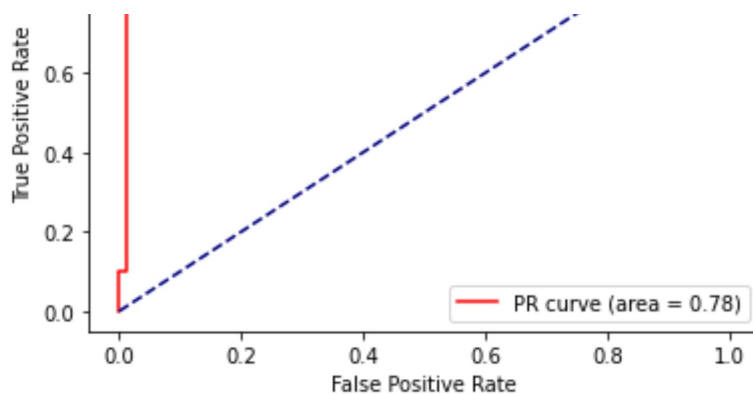
```
plt.title('Receiver operating characteristic')
```

```
plt.legend(loc="lower right")
```

```
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```





Random Forest

```
# model training Using random forest
from sklearn.ensemble import RandomForestClassifier
forest = RandomForestClassifier(random_state = 1, n_estimators = 10, min_samples_split = 2)
forest.fit(X_train, Y_train)
```

```
RandomForestClassifier(bootstrap=True, ccp_alpha=0.0, class_weight=None,
                        criterion='gini', max_depth=None, max_features='auto',
                        max_leaf_nodes=None, max_samples=None,
                        min_impurity_decrease=0.0, min_impurity_split=None,
                        min_samples_leaf=1, min_samples_split=2,
                        min_weight_fraction_leaf=0.0, n_estimators=10,
                        n_jobs=None, oob_score=False, random_state=1, verbose=0,
                        warm_start=False)
```

```
# accuracy score for training data and testing data
X_train_prediction=forest.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)
```

```
X_test_prediction=forest.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)
```

```
print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data:  0.9854651162790697
Accuracy score for testing data:  0.9186046511627907
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
```

```
result = cross_val_score(forest , X_train, Y_train, cv = kf)
result
```

```
array([0.91428571, 0.88571429, 0.91428571, 0.85714286, 0.88235294,
       0.88235294, 0.85294118, 0.85294118, 0.94117647, 0.91176471])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.8894957983193278
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(forest , X_test, Y_test, cv = kf)
result
```

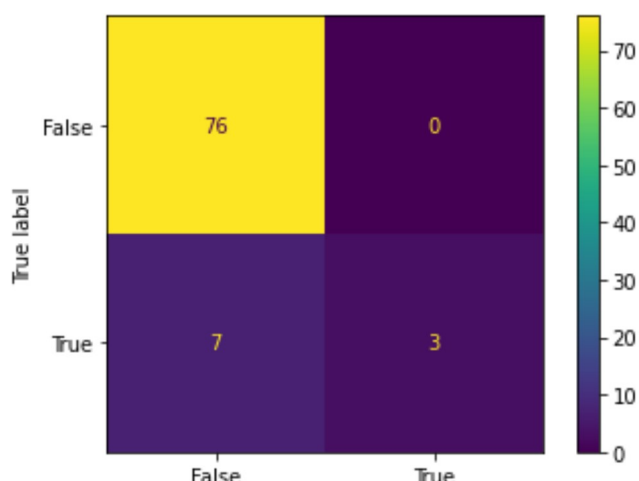
```
array([0.88888889, 1.          , 1.          , 0.77777778, 0.77777778,
       1.          , 1.          , 1.          , 0.875       , 0.75        ])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9069444444444444
```

```
# make predictions
predicted = forest.predict(X_test)
from sklearn.metrics import accuracy_score, confusion_matrix
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
cm_display.plot()
plt.show()
```



----- Predicted label -----

```
TN = confusion_matrix[0][0]
FN = confusion_matrix[1][0]
TP = confusion_matrix[1][1]
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))
specificity = (TN / float(TN + FP))
ppv = (TP / float(TP + FP))
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)
print("specificity: ",specificity)
print("PPV: ",ppv)
print("NPV: ",npv)
```

```
Sensitivity:  0.3
specificity:  1.0
PPV:  1.0
NPV:  0.9156626506024096
```

```
y_predictProb = forest.predict_proba(X_test)
```

```
fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
roc_auc = auc(fpr, tpr)
```

```
precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
area = auc(recall, precision)
```

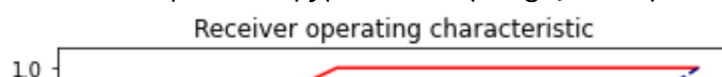
```
print("AUROC:",roc_auc)
print("AUPR:",area)
```

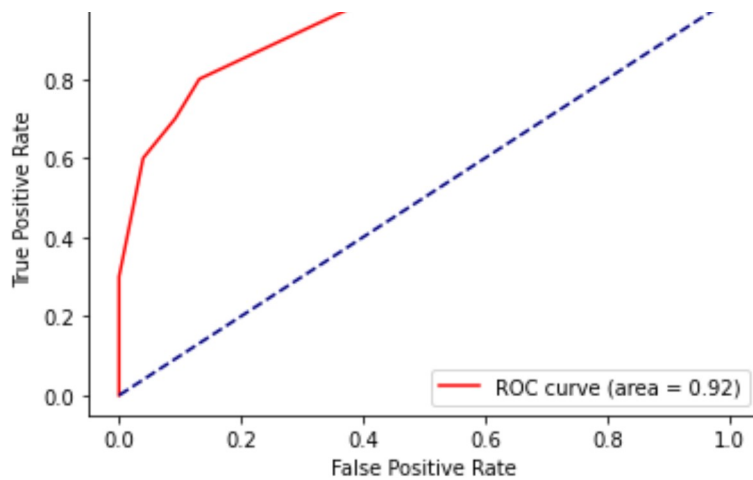
```
AUROC: 0.9223684210526314
AUPR: 0.7148664343786295
```

```
# AUROC graph
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```

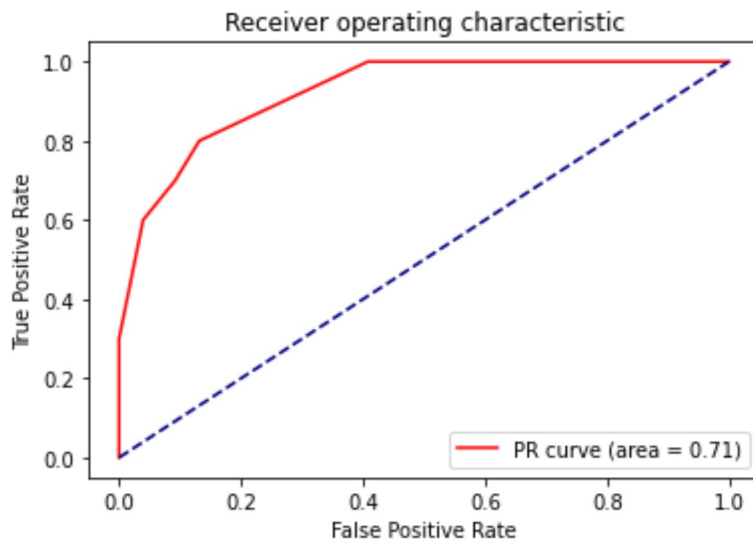




AUPR graph

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

<function matplotlib.pyplot.show(*args, **kw)>



Decision Tree

```
# using decisin tree
from sklearn.tree import DecisionTreeClassifier
dclf = DecisionTreeClassifier()
dclf.fit(X_train,Y_train)
```

```
DecisionTreeClassifier(ccp_alpha=0.0, class_weight=None, criterion='gini',
                      max_depth=None, max_features=None, max_leaf_nodes=None,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, presort='deprecated',
                      random_state=None, splitter='best')
```

```
# accuracy score for training data and testing data
```

```
X_train_prediction=dclf.predict(X_train)
```

```
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)
```

```
X_test_prediction=dclf.predict(X_test)
```

```
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)
```

```
print('Accuracy score for training data: ',X_training_accuracy)
```

```
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data: 1.0
```

```
Accuracy score for testing data: 0.8837209302325582
```

```
from sklearn.model_selection import cross_val_score
```

```
from sklearn.model_selection import KFold
```

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

```
k = 10
```

```
kf = KFold(n_splits=k, random_state=None)
```

```
result = cross_val_score(dclf , X_train, Y_train, cv = kf)
```

```
result
```

```
array([0.82857143, 0.82857143, 0.8       , 0.77142857, 0.88235294,
       0.82352941, 0.73529412, 0.79411765, 0.91176471, 0.88235294])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.8257983193277312
```

```
from sklearn.model_selection import cross_val_score
```

```
from sklearn.model_selection import KFold
```

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

```
k = 10
```

```
kf = KFold(n_splits=k, random_state=None)
```

```
result = cross_val_score(dclf , X_test, Y_test, cv = kf)
```

```
result
```

```
array([0.88888889, 0.88888889, 1.         , 0.88888889, 1.         ,
       1.         , 1.         , 1.         , 0.875       , 1.         ])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

Avg accuracy: 0.9541666666666666

```
# make predictions
```

```
predicted = dclf.predict(X_test)
```

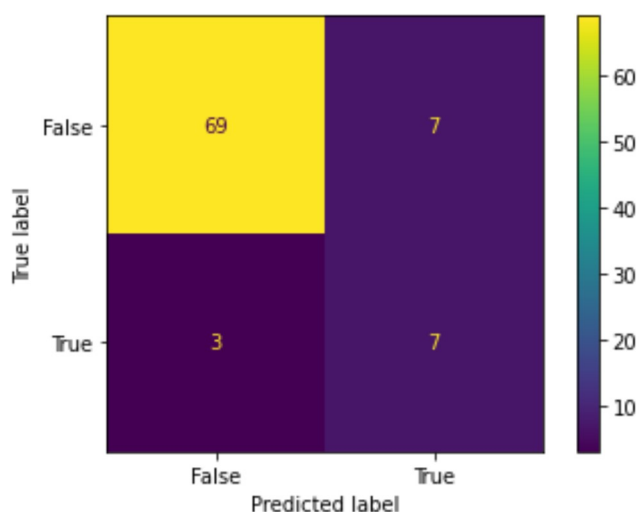
```
from sklearn.metrics import accuracy_score, confusion_matrix
```

```
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
```

```
cm_display.plot()
```

```
plt.show()
```



```
TN = confusion_matrix[0][0]
```

```
FN = confusion_matrix[1][0]
```

```
TP = confusion_matrix[1][1]
```

```
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))
```

```
specificity = (TN / float(TN + FP))
```

```
ppv = (TP / float(TP + FP))
```

```
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)
```

```
print("specificity: ",specificity)
```

```
print("PPV: ",ppv)
```

```
print("NPV: ",npv)
```

Sensitivity: 0.7

specificity: 0.9078947368421053

PPV: 0.5

NPV: 0.9583333333333333

```
# AUROC and AUPR value
y_predictProb = dclf.predict_proba(X_test)

fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
roc_auc = auc(fpr, tpr)

precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
area = auc(recall, precision)

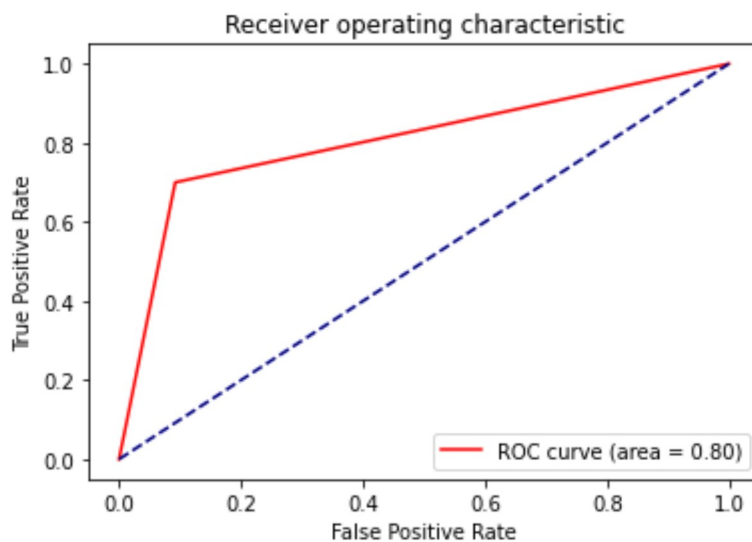
print("AUROC:",roc_auc)
print("AUPR:",area)

AUROC: 0.8039473684210526
AUPR: 0.6174418604651162
```

```
# AUROC graph
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```

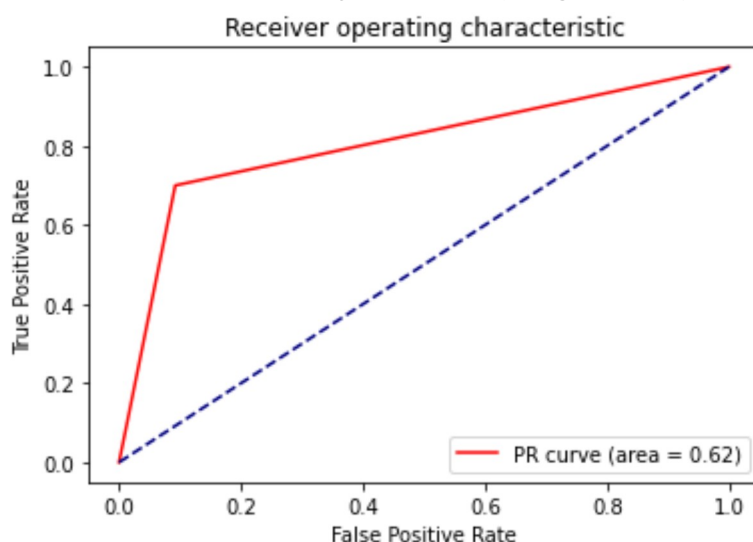


```
# AUPR graph
```

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
```

```
plt.legend(loc= 'lower right' )
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```



Gradient Bosst

```
#using GradientBoost
from sklearn.ensemble import GradientBoostingClassifier
gdb = GradientBoostingClassifier(random_state = 1, n_estimators = 10, min_samples_split =
gdb.fit(X_train,Y_train)
```

```
GradientBoostingClassifier(ccp_alpha=0.0, criterion='friedman_mse', init=None,
                           learning_rate=0.1, loss='deviance', max_depth=3,
                           max_features=None, max_leaf_nodes=None,
                           min_impurity_decrease=0.0, min_impurity_split=None,
                           min_samples_leaf=1, min_samples_split=2,
                           min_weight_fraction_leaf=0.0, n_estimators=10,
                           n_iter_no_change=None, presort='deprecated',
                           random_state=1, subsample=1.0, tol=0.0001,
                           validation_fraction=0.1, verbose=0,
                           warm_start=False)
```

```
# accuracy score for training data and testing data
X_train_prediction=gdb.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)
```

```
X_test_prediction=gdb.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)
```

```
print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data: 0.9476744186046512
Accuracy score for testing data: 0.9069767441860465
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(gdb , X_train, Y_train, cv = kf)
result
```

```
array([0.88571429, 0.94285714, 0.88571429, 0.8         , 0.91176471,
       0.85294118, 0.85294118, 0.85294118, 0.97058824, 0.91176471])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.8867226890756303
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(gdb , X_test, Y_test, cv = kf)
result
```

```
array([0.88888889, 0.88888889, 1.         , 0.88888889, 0.88888889,
       1.         , 1.         , 1.         , 0.875     , 1.         ])
```

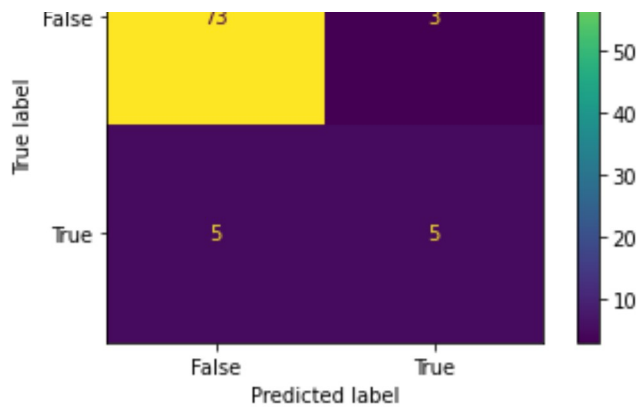
```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9430555555555555
```

```
# make predictions
predicted = gdb.predict(X_test)
from sklearn.metrics import accuracy_score, confusion_matrix
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
cm_display.plot()
plt.show()
```





```
TN = confusion_matrix[0][0]
FN = confusion_matrix[1][0]
TP = confusion_matrix[1][1]
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))
specificity = (TN / float(TN + FP))
ppv = (TP / float(TP + FP))
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)
print("specificity: ",specificity)
print("PPV: ",ppv)
print("NPV: ",npv)
```

```
Sensitivity: 0.5
specificity: 0.9605263157894737
PPV: 0.625
NPV: 0.9358974358974359
```

```
# AUROC and AUPR value
y_predictProb = gdb.predict_proba(X_test)
```

```
fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
roc_auc = auc(fpr, tpr)
```

```
precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
area = auc(recall, precision)
```

```
print("AUROC:",roc_auc)
print("AUPR:",area)
```

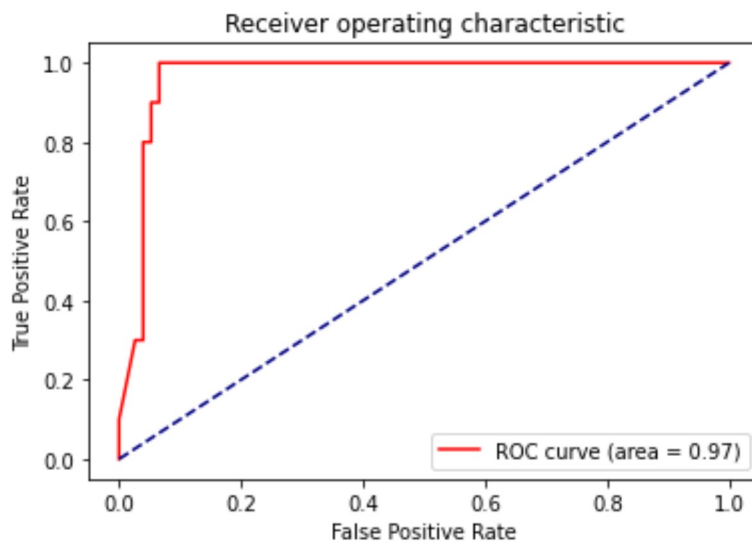
```
AUROC: 0.9657894736842105
AUPR: 0.6977647352647353
```

```
# AUROC graph
```

```
plt.figure(figsize=(10,5))
plt.plot(fpr, tpr, label='ROC curve (area = %0.2f)' % roc_auc)
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

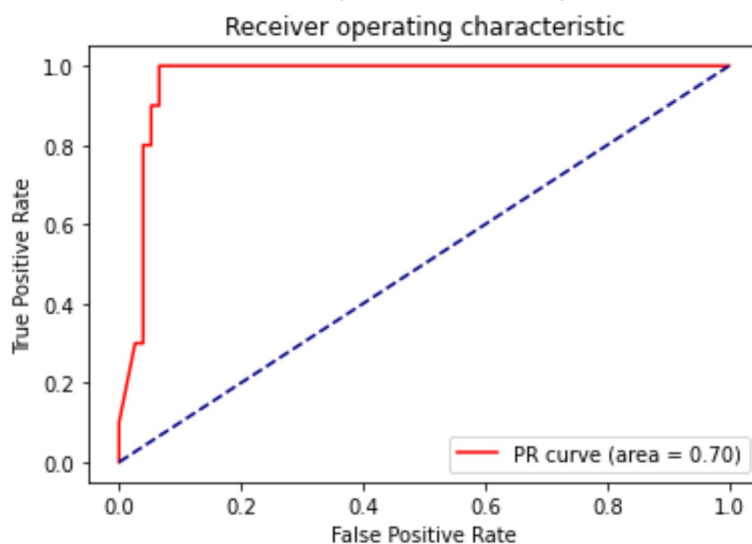
```
<function matplotlib.pyplot.show(*args, **kw)>
```



AUPR graph

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```



XGBoost

```
#using XGBClassifier
from xgboost import XGBClassifier
xgb_clf = XGBClassifier(random_state = 1, n_estimators = 10, min_samples_split = 2)
xgb_clf.fit(X_train, Y_train)
```

```
    XGBClassifier(base_score=0.5, booster='gbtree', colsample_bylevel=1,
                  colsample_bynode=1, colsample_bytree=1, gamma=0,
                  learning_rate=0.1, max_delta_step=0, max_depth=3,
                  min_child_weight=1, min_samples_split=2, missing=None,
                  n_estimators=10, n_jobs=1, nthread=None,
                  objective='binary:logistic', random_state=1, reg_alpha=0,
                  reg_lambda=1, scale_pos_weight=1, seed=None, silent=None,
                  subsample=1, verbosity=1)
```

```
# accuracy score for training data and testing data
X_train_prediction=xgb_clf.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)
```

```
X_test_prediction=xgb_clf.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)
```

```
print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data:  0.9447674418604651
Accuracy score for testing data:  0.9186046511627907
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(xgb_clf , X_train, Y_train, cv = kf)
result
```

```
array([0.91428571, 0.88571429, 0.91428571, 0.85714286, 0.91176471,
       0.88235294, 0.88235294, 0.94117647, 0.97058824, 0.91176471])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9071428571428571
```

```

from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score

```

```

k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(xgb_clf , X_test, Y_test, cv = kf)
result

```

```

array([0.88888889, 1.          , 1.          , 0.88888889, 0.88888889,
       1.          , 1.          , 1.          , 0.875      , 1.          ])

```

```
print("Avg accuracy: {}".format(result.mean()))
```

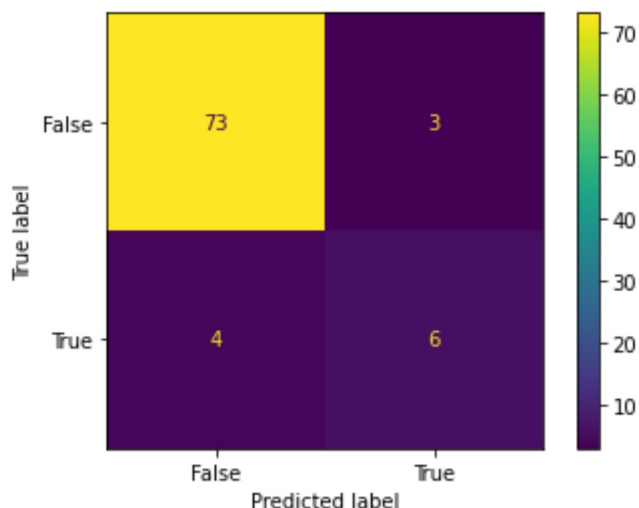
```
Avg accuracy: 0.9541666666666666
```

```

# make predictions
predicted = xgb_clf.predict(X_test)
from sklearn.metrics import accuracy_score, confusion_matrix
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)

cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
cm_display.plot()
plt.show()

```



```

TN = confusion_matrix[0][0]
FN = confusion_matrix[1][0]
TP = confusion_matrix[1][1]
FP = confusion_matrix[0][1]

```

```

sensitivity = (TP / float(TP + FN))
specificity = (TN / float(TN + FP))
nrv = (TP / float(TP + FP))

```

```
ppv = (TP / float(TP + FP))  
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)  
print("specificity: ",specificity)  
print("PPV: ",ppv)  
print("NPV: ",npv)
```

```
Sensitivity: 0.6  
specificity: 0.9605263157894737  
PPV: 0.6666666666666666  
NPV: 0.948051948051948
```

```
# AUROC and AUPR value
```

```
y_predictProb = xgb_clf.predict_proba(X_test)
```

```
fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])  
roc_auc = auc(fpr, tpr)
```

```
precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])  
area = auc(recall, precision)
```

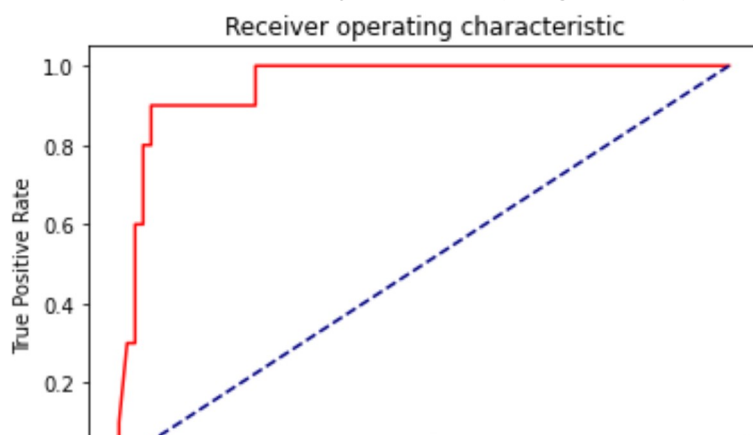
```
print("AUROC:",roc_auc)  
print("AUPR:",area)
```

```
AUROC: 0.9552631578947369  
AUPR: 0.7240671365671366
```

```
# AUROC graph
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)  
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')  
plt.xlabel('False Positive Rate')  
plt.ylabel('True Positive Rate')  
plt.title('Receiver operating characteristic')  
plt.legend(loc="lower right")  
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```

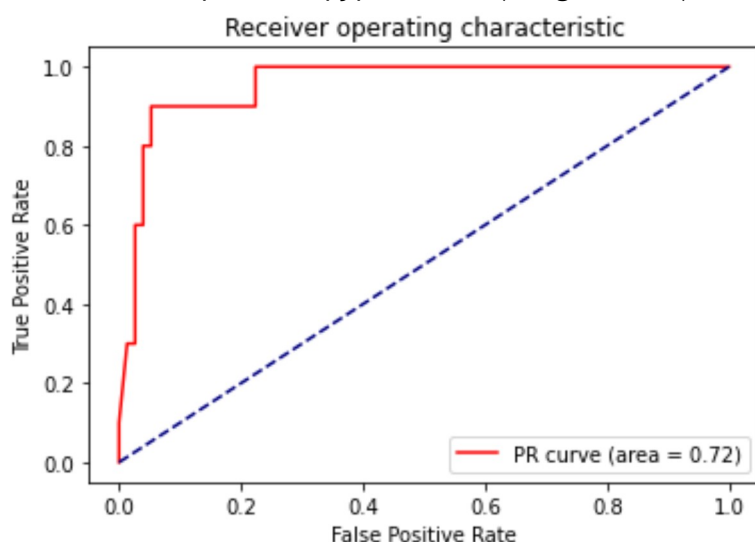




```
# AUPR graph
```

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```



Support Vector

```
#using support vector
from sklearn import svm
sv_clf = svm.SVC()
sv_clf.fit(X_train, Y_train)
```

```
SVC(C=1.0, break_ties=False, cache_size=200, class_weight=None, coef0=0.0,
    decision_function_shape='ovr', degree=3, gamma='scale', kernel='rbf',
    max_iter=-1, probability=False, random_state=None, shrinking=True,
    tol=0.001, verbose=False)
```

```
# accuracy score for training data and testing data
X_train_prediction=sv_clf.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)
```

```
X_test_prediction=sv_clf.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)

print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data: 0.9186046511627907
Accuracy score for testing data: 0.9069767441860465
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(sv_clf , X_train, Y_train, cv = kf)
result
```

```
array([0.91428571, 0.91428571, 0.91428571, 0.88571429, 0.91176471,
       0.88235294, 0.91176471, 0.91176471, 0.97058824, 0.94117647])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.915798319327731
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(sv_clf , X_test, Y_test, cv = kf)
result
```

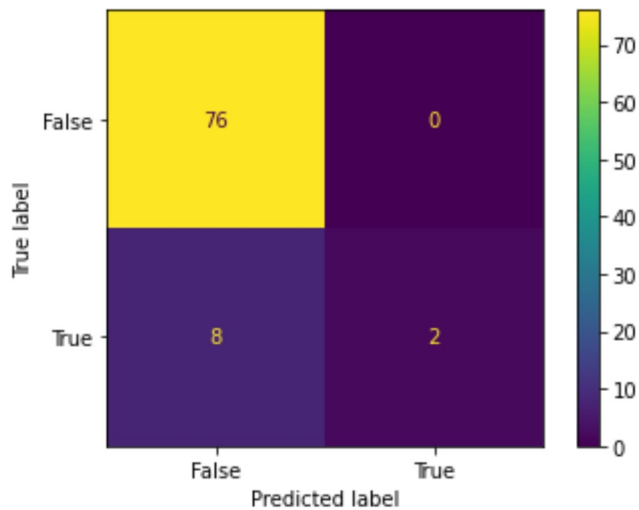
```
array([0.88888889, 1.         , 1.         , 0.88888889, 0.88888889,
       1.         , 1.         , 1.         , 0.875       , 0.875       ])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.9416666666666667
```

```
# make predictions
predicted = sv_clf.predict(X_test)
from sklearn.metrics import accuracy_score, confusion_matrix
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
cm_display.plot()
plt.show()
```



```
TN = confusion_matrix[0][0]
FN = confusion_matrix[1][0]
TP = confusion_matrix[1][1]
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))
specificity = (TN / float(TN + FP))
ppv = (TP / float(TP + FP))
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)
print("specificity: ",specificity)
print("PPV: ",ppv)
print("NPV: ",npv)
```

```
Sensitivity:  0.2
specificity:  1.0
PPV:  1.0
NPV:  0.9047619047619048
```

```
# AUROC and AUPR value
y_predictProb = sv_clf.predict_proba(X_test)
```

```
fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
roc_auc = auc(fpr, tpr)
```

```
precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
area = auc(recall, precision)
```

```
print("AUROC:",roc_auc)
print("AUPR:",area)
```

```

-----
AttributeError                                Traceback (most recent call last)
<ipython-input-116-289267775586> in <module>
      1 # AUROC and AUPR value
----> 2 y_predictProb = sv_clf.predict_proba(X_test)
      3
      4 fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
      5 roc_auc = auc(fpr, tpr)

1 frames
/usr/local/lib/python3.7/dist-packages/sklearn/svm/_base.py in _check_proba(self)
    601     def _check_proba(self):
    602         if not self.probability:
--> 603             raise AttributeError("predict_proba is not available when "
    604                                 " probability=False")
    605         if self._impl not in ('c_svc', 'nu_svc'):

AttributeError: predict_proba is not available when  probability=False

```

SEARCH STACK OVERFLOW

AUROC graph

```

plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show

```

AUPR graph

```

plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show

```

Gaussian Naive Bayes

#using Naive Bayesian

```

from sklearn.naive_bayes import GaussianNB

```

```
from sklearn.naive_bayes import GaussianNB
gnb = GaussianNB()
gnb.fit(X_train, Y_train)

GaussianNB(priors=None, var_smoothing=1e-09)

# accuracy score for training data and testing data
X_train_prediction=gnb.predict(X_train)
X_training_accuracy=accuracy_score(X_train_prediction,Y_train)

X_test_prediction=gnb.predict(X_test)
X_testing_accuracy=accuracy_score(X_test_prediction,Y_test)

print('Accuracy score for training data: ',X_training_accuracy)
print('Accuracy score for testing data: ',X_testing_accuracy)
```

```
Accuracy score for training data:  0.8313953488372093
Accuracy score for testing data:  0.9418604651162791
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(gnb , X_train, Y_train, cv = kf)
result
```

```
array([0.71428571, 0.77142857, 0.77142857, 0.91428571, 0.88235294,
       0.85294118, 0.79411765, 0.82352941, 0.82352941, 0.79411765])
```

```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.8142016806722688
```

```
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import KFold
from sklearn.metrics import accuracy_score
```

```
k = 10
kf = KFold(n_splits=k, random_state=None)
result = cross_val_score(gnb , X_test, Y_test, cv = kf)
result
```

```
array([0.88888889, 0.66666667, 0.77777778, 0.44444444, 0.66666667,
       1.          , 0.875       , 0.875       , 0.75        , 0.875       ])
```



```
print("Avg accuracy: {}".format(result.mean()))
```

```
Avg accuracy: 0.7819444444444444
```

```
# make predictions
```

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

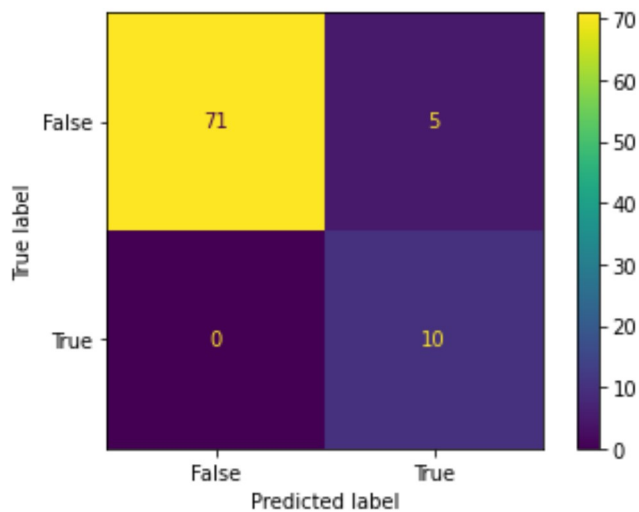
```
from sklearn.metrics import accuracy_score, confusion_matrix
```

```
confusion_matrix = metrics.confusion_matrix(Y_test,predicted)
```

```
cm_display = metrics.ConfusionMatrixDisplay(confusion_matrix = confusion_matrix, display_1
```

```
cm_display.plot())
```

```
plt.show()
```



```
TN = confusion_matrix[0][0]
```

```
FN = confusion_matrix[1][0]
```

```
TP = confusion_matrix[1][1]
```

```
FP = confusion_matrix[0][1]
```

```
sensitivity = (TP / float(TP + FN))
```

```
specificity = (TN / float(TN + FP))
```

```
ppv = (TP / float(TP + FP))
```

```
npv = (TN / float(TN + FN))
```

```
print("Sensitivity: ",sensitivity)
```

```
print("specificity: ",specificity)
```

```
print("PPV: ",ppv)
```

```
print("NPV: ",npv)
```

```
Sensitivity: 1.0
```

```
specificity: 0.9342105263157895
```

```
PPV: 0.6666666666666666
```

```
NPV: 1.0
```

```
# AUROC and AUPR value
```

```
y_predictProb = gnb.predict_proba(X_test)

fpr, tpr, thresholds = roc_curve(Y_test, y_predictProb[:,1])
roc_auc = auc(fpr, tpr)

precision, recall, thresholds = precision_recall_curve(Y_test, y_predictProb[:,1])
area = auc(recall, precision)

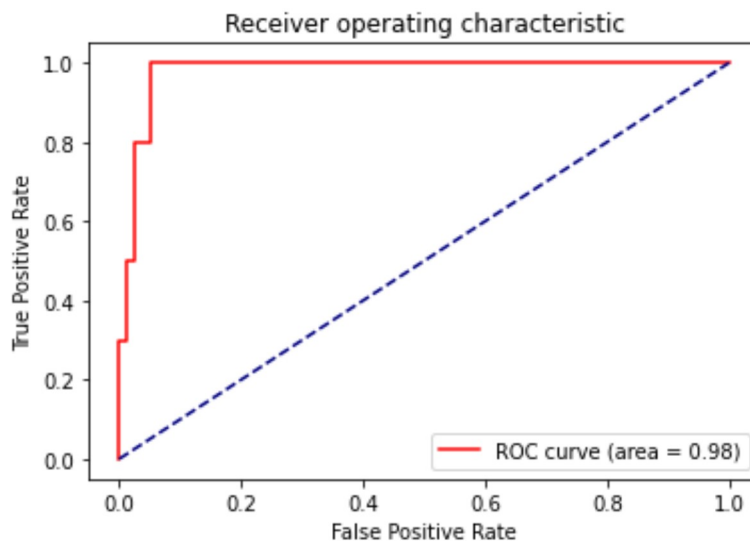
print("AUROC:",roc_auc)
print("AUPR:",area)

AUROC: 0.9789473684210527
AUPR: 0.8259371184371184
```

```
# AUROC graph
```

```
plt.plot(fpr, tpr, color='red', label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

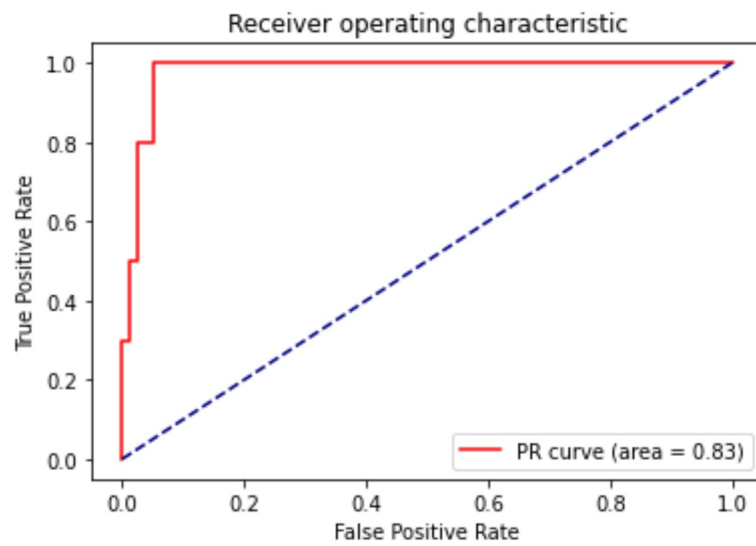
```
<function matplotlib.pyplot.show(*args, **kw)>
```



```
# AUPR graph
```

```
plt.plot(fpr, tpr, color='red', label='PR curve (area = %0.2f)' % area)
plt.plot([0, 1], [0, 1], color='navy', linestyle='--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic')
plt.legend(loc="lower right")
plt.show
```

```
<function matplotlib.pyplot.show(*args, **kw)>
```



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