

Evaluations on Classifiers on Breast Cancer Dataset

April 15, 2021

```
[1]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_breast_cancer
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import precision_score, recall_score,
    ↪classification_report, confusion_matrix
from sklearn.metrics import roc_curve, auc, plot_precision_recall_curve

[2]: # Load the data
data = load_breast_cancer()
X = data.data
y = data.target

[3]: # Split the dataset into training set and test set (80, 20).
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2)

[4]: def model_evaluation(clf):
    # (1). The accuracy of your model on the test data
    accuracy = clf.score(X_test, y_test)
    print("1. accuracy of the model on the test data: " + str(accuracy) + "\n")

    # (2). The precision and recall values
    y_pred = clf.predict(X_test)
    # precision value
    precisions = precision_score(y_test, y_pred, average=None)
    print('2. \n1) precision values: ' + str(precisions))
    # recall value
    recalls = recall_score(y_test, y_pred, average=None)
    print('2) recall values: ' + str(recalls))

    # (3). A classification report (scikit-learn has a function that can create
    ↪this for you)
    target_names = list(data.target_names)
    class_report = classification_report(y_test, y_pred,
    ↪target_names=target_names)
```

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print('\n3. classification report\n' + class_report)

# (4). The confusion matrix for this experiment
print('\n4. confusion matrix')
print(confusion_matrix(y_test, y_pred))

# (5). An ROC curve
fpr, tpr, _ = roc_curve(y_test, y_pred)
roc_auc = auc(fpr, tpr)

plt.figure()
lw = 2
plt.plot(fpr, tpr, color='darkorange',
         lw=lw, label='ROC curve (area = %0.2f)' % roc_auc)
plt.plot([0, 1], [0, 1], color='navy', lw=lw, linestyle='--')
plt.xlim([0.0, 1.0])
plt.ylim([0.0, 1.05])
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('5. Receiver operating characteristic (ROC curve)')
plt.legend(loc="lower right")
plt.show()

# (6). A Precision/Recall curve
disp = plot_precision_recall_curve(clf, X_test, y_test)
disp.ax_.set_title('6. 2-class Precision-Recall curve')

```

1 Use scikit-learn's DecisionTreeClassifier to train a supervised learning model

```

[5]: from sklearn import tree
      clf0 = tree.DecisionTreeClassifier()
      clf0 = clf0.fit(X_train, y_train)
      model_evaluation(clf0)

```

1. accuracy of the model on the test data: 0.9298245614035088

2.

1) precision values: [0.95238095 0.91666667]

2) recall values: [0.86956522 0.97058824]

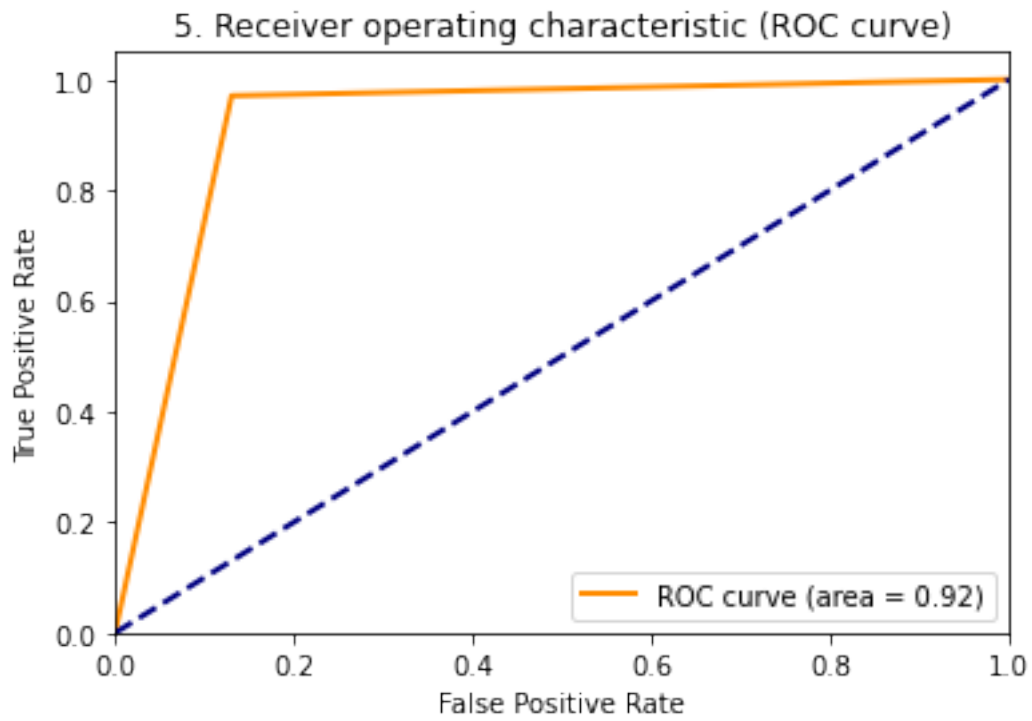
3. classification report

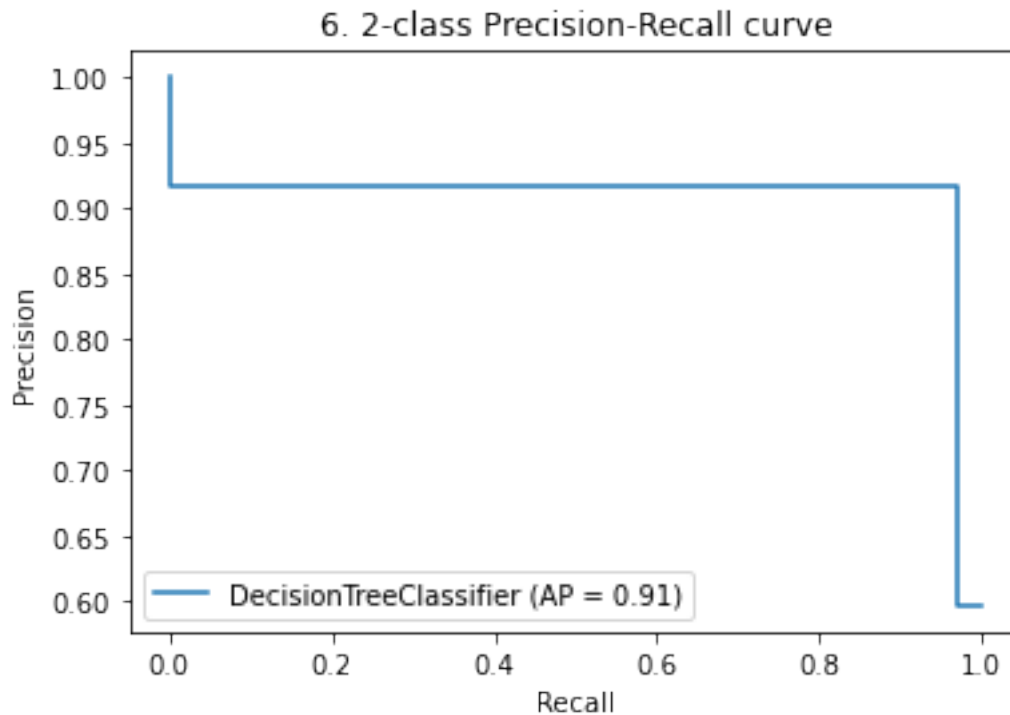
	precision	recall	f1-score	support
malignant	0.95	0.87	0.91	46
benign	0.92	0.97	0.94	68

accuracy			0.93	114
macro avg	0.93	0.92	0.93	114
weighted avg	0.93	0.93	0.93	114

4. confusion matrix

```
[[40  6]
 [ 2 66]]
```





2 Train another Decision Tree Classifier - but in this case set the maximum depth of the tree to 1 (max_depth= 1).

```
[6]: clf1 = tree.DecisionTreeClassifier(max_depth=1)
      clf1 = clf1.fit(X_train, y_train)
      model_evaluation(clf1)
```

1. accuracy of the model on the test data: 0.8859649122807017

2.

1) precision values: [1. 0.83950617]

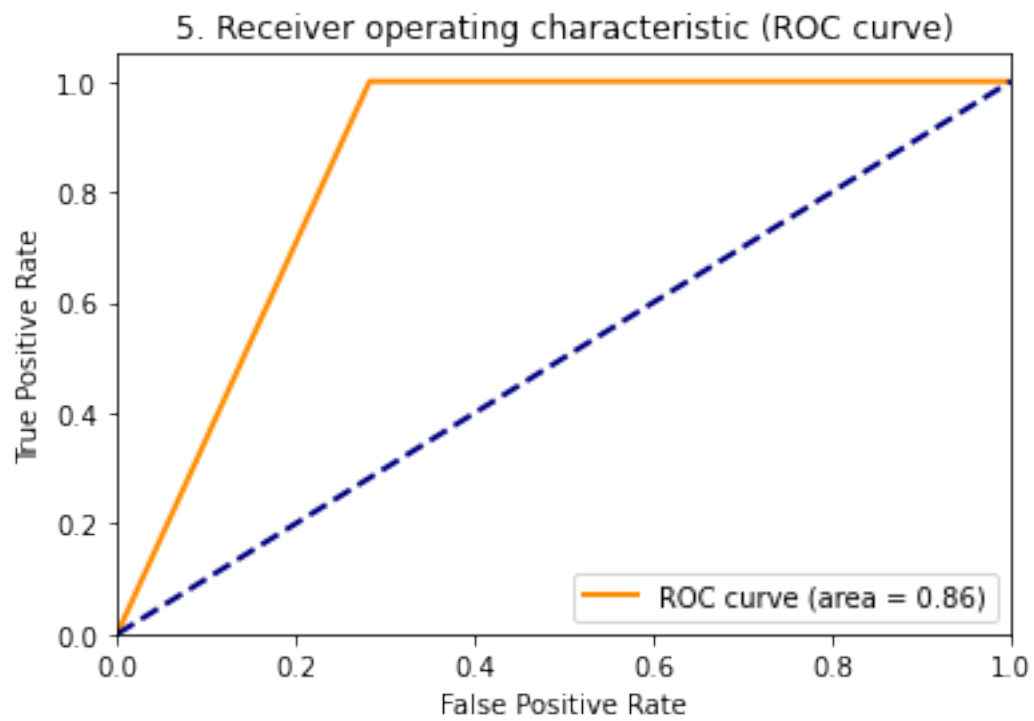
2) recall values: [0.7173913 1.]

3. classification report

	precision	recall	f1-score	support
malignant	1.00	0.72	0.84	46
benign	0.84	1.00	0.91	68
accuracy			0.89	114
macro avg	0.92	0.86	0.87	114
weighted avg	0.90	0.89	0.88	114

4. confusion matrix

```
[[33 13]  
 [ 0 68]]
```



6. 2-class Precision-Recall curve

