Lab 1 Machine Learning Bianca Caissotti di Chiusano (i6245461) import pandas as pd from sklearn import tree

from sklearn.metrics import accuracy score from sklearn.model selection import train test split import matplotlib.pyplot as plt import numpy as np Lab Task 4c)

data = pd.read csv('diabetes.csv')

preg

plas

data

pres skin insu mass pedi age 0 0.627 148 72 33.6 50 tested_positive 1 85 66 29 0 26.6 0.351 31 tested_negative

28.1

32.9

23.3 0.672

43.1 2.288

36.8 0.340

26.2 0.245

30.1 0.349

30.4 0.315

We can now see that our accuracy score when using the training data is 1. This is because of the max_depth is not limited and all possible decisions for the training data can be satisfied. However, the accuracy score when using the test data did not change much, compared to when using one-level trees,

because the tree was trained using the training data, thus not all decisions for the test data can be

Training one-level and multi-level decision trees on the glass data

Si

1.12 73.03 0.64

74.55

70.16

72.88

73.21

1.41 72.64

1.54

1.00

1.25

1.71

Κ

0.59

0.00

0.12

0.59

0.69

0.37

0.73

Ca

8.77

8.53

8.43

7.59

16.19

8.32 0.0

7.97

7.94

8.54

clf glass = tree.DecisionTreeClassifier(criterion = 'entropy', max_depth = None) X train, X test, Y train, Y test = train test split(X glass, Y glass, test size = 0.34

For Task 4d: Estimate the accuracy rates of the resulting decision trees using the training set and hold-out validation. Plot the accuracy rates based on the training set and hold-out validation for min_samples_leaf

Function I called estimate. The decision tree is multi level. And the min sample lea

clf = tree.DecisionTreeClassifier(criterion = 'entropy', max depth = None, min sar

def estimate(X train, X test, Y train, Y test, num leafs, acc train, acc test):

Ba

0.0

0.0

0.0

0.0

0.0

0.0

0.0

0.0 9.32 0.0 0.00

Fe

0.00

0.00

0.00

0.00

0.24

0.00

0.14

0.00

class

'build wind float'

'vehic wind float'

'build wind float'

'build wind non-float'

'build wind non-float'

'build wind non-float'

tableware

0.00 'build wind non-float'

'vehic wind float'

tableware

0.167

0.171

32

21

63

27

47

0

23

35

48

27

23

0

31

0

94

168

180

0

112

0

class

tested_positive

tested_negative

tested_positive

tested_negative

tested_negative

tested_negative

tested_positive

23 tested_negative

We are firstly training decision trees on the diabetes data So we read the CSV file.

2 8 183 64 3 89 66 4 137 40

763 10 101 76 764 2 122 70 765 121

766 126 60 767 70 1 93 768 rows × 9 columns

output column (Vector) Y Y = data['class'] # input matrix X X = data.drop(['class'], axis = 1) Starting from the One-Level decision tree: so the max_depth parameter of DecisionTreeClassifier is set to 1. This means that we are only training one layer of the decision tree. We then randomly split the diabetes data into "training" and "test" data sets. Test_size is 0.34, because it is randomly selecting 66% of rows

for training, and the rest for testing. We use this training data to train the classifier. Following that use the trained classifier to predict the output given the same training data (that was used to train). And calculate the accuracy score by comparing the trained output with the just calculated output Yp.

clf = tree.DecisionTreeClassifier(criterion = 'entropy', max depth = 1) X train, X test, Y train, Y test = train test split(X, Y, test size = 0.34, random sta clf = clf.fit(X train, Y train) Yp = clf.predict(X train)

acc = accuracy score(Y train, Yp) print(acc) 0.7648221343873518 Now, more importantly, we want to see how good out trained clf is at predicting output for data that it has never seen, thus the test data. # Do the same but for the test data

Yp = clf.predict(X test) acc = accuracy score(Y test, Yp) print(acc) 0.7213740458015268

We can observe that the two calculated accuracies (for train and test data) are quite similar as we are only training a one-level decision trees. Following, the same exact steps are applied, however this time for a multi-level decision tree, thus setting max_depth to None in our DecisionTreeClassifier. # MULTI-LEVEL DECISION TREES

clf = tree.DecisionTreeClassifier(criterion = 'entropy', max depth = None) X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size = 0.34, random_state clf = clf.fit(X train, Y train) Yp = clf.predict(X train) acc = accuracy_score(Y_train, Yp) print(acc) Yp = clf.predict(X test) acc = acc = accuracy score(Y test, Yp)

1.0

0.732824427480916

satisfies, even without any limit on the depth of the tree.

Same steps as above are repeated for the data in glass.csv

glass_data = pd.read_csv('glass.csv')

Mg

glass_data RI Na 1.51793 12.79 3.50 **1** 1.51643 12.16 3.52 1.35 72.89 0.57 13.21 1.51793 3.48 1.51299 14.40 1.53393 12.30

1.74 0.00 209 1.51610 13.42 3.40 1.22 72.69 1.51592 12.86 3.52 2.12 72.66 211 1.51613 13.92 3.52 1.51689 2.88 12.67 2.19 1.66 72.67 0.00 1.51852 14.09

214 rows × 10 columns Y glass = glass_data['class'] X glass = glass data.drop(['class'], axis = 1) clf glass = tree.DecisionTreeClassifier(criterion = 'entropy', max_depth = 1) X train, X test, Y train, Y test = train test split(X glass, Y glass, test size = 0.34

Accuracy Score Training data clf_glass = clf_glass.fit(X_train,Y_train) Yp = clf glass.predict(X train) acc = accuracy_score(Y_train, Yp) print(acc) 0.46099290780141844 In [24]: # Accuracy Score Test data Yp = clf glass.predict(X test) acc = accuracy score(Y test, Yp) print(acc)

> clf_glass = clf_glass.fit(X_train,Y_train) Yp = clf_glass.predict(X_train) # Accuracy Score Training data acc = accuracy_score(Y_train, Yp) print(acc) 1.0 # Accuracy Score Test data

0.5616438356164384

Lab Tasks 4d)

print(acc)

Yp = clf glass.predict(X test)

acc = acc = accuracy score(Y test, Yp)

from 1 to the size of the datasets with step of 5

clf = clf.fit(X_train,Y_train)

0.4246575342465753

Multi-Level

Arrays to store Accuracy scores at each time step for both data sets acc train diabetes = [] acc_test_diabetes = [] acc train glass = []

acc_test_glass = []

In [45]:

Yp = clf.predict(X train) acc = accuracy_score(Y_train, Yp) acc_train.append(acc) Yp = clf.predict(X test)

acc = accuracy_score(Y_test, Yp) acc test.append(acc) X train, X test, Y train, Y test = train test split(X, Y, test size = 0.34, random states) X_train_glass, X_test_glass, Y_train_glass, Y_test_glass = train_test_split(X_glass, for i in range(1, data.shape[0]): estimate(X train, X test, Y train, Y test, i, acc train diabetes, acc test diabete

for i in range(1, glass data.shape[0]): estimate(X_train_glass, X_test_glass, Y_train_glass, Y_test_glass, i, acc_train_gl For both datasets, at each time step we call the estimate function. The min_sample_leaf changes from 1 to

plotted.

the size of the dataset with a time step of 5. Every accuracy score is appended to the array which is then # x axis is the time step # y axis is the accuracy plt.title("Train and Test Accuracy Diabetes") plt.ylabel("Accuracy") plt.xlabel("Time Step") plt.plot(acc train diabetes, label = "Train")

plt.show()

1.00

0.95 0.90 0.85 0.80 0.75 0.70

plt.plot(acc test diabetes, label = "Test")

300

400

Time Step

Train and Test Accuracy Glass

100

Time Step

Train Test

500

600

700

200

In both graphs, we can observe that at the beginning the training data is predicted with a greater accuracy than with the testing data, because the min_sample_leafs is small but we are still working with a multi-level tree. However as the min_sample_leafs increases than the accuracy of the training data also decreases.

800

Train and Test Accuracy Diabetes

Train Test

plt.legend(loc = 'upper center')

In [46]:

0.65 0.60 0 100 200 plt.title("Train and Test Accuracy Glass") plt.ylabel("Accuracy") plt.xlabel("Time Step") plt.plot(acc_train_glass, label = "Train") plt.plot(acc_test_glass, label = "Test")

plt.show()

1.0

0.9 0.8

0.7

Accuracy 0.6 0.5 0.4 0.3 0.2

plt.legend(loc = 'upper center')

50