CM7

July 18, 2021

1 [CM7] Interpretability

1.1 The implementation is done for the best hyper-parameters of naive bayes and decision trees using the original features

[28]:
```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import cross_val_score,train_test_split
from sklearn.naive_bayes import GaussianNB
```

[31]:
```
covid_dataset = pd.read_csv("cleaned_normalized_coviddata.csv")
x,x_test,y,y_test = train_test_split(covid_dataset.iloc[:,2:-3],covid_dataset.
 ↪iloc[:,-3:].astype(int),test_size=0.2,random_state=98)
```

1.1.1 Naive Bayes Implementation

[34]:
```
#Label: Recovered
classifier_gaussian_recovered = GaussianNB(var_smoothing = 1e-1)
classifier_gaussian_recovered.fit(x,y.loc[:,'Recovered'])
scores_nb_recovered = classifier_gaussian_recovered.score(x_test,y_test.loc[:
 ↪,'Recovered'])
#LAbel: Deaths
classifier_gaussian_deaths = GaussianNB(var_smoothing = 0.75)
classifier_gaussian_deaths.fit(x,y.loc[:,'Deaths'])
scores_nb_deaths = classifier_gaussian_deaths.score(x_test,y_test.loc[:
 ↪,'Deaths'])
#Label : Confirmed
classifier_gaussian_confirmed = GaussianNB(var_smoothing = 1)
classifier_gaussian_confirmed.fit(x,y.loc[:,'Confirmed'])
scores_nb_confirmed = classifier_gaussian_confirmed.score(x_test,y_test.loc[:
 ↪,'Confirmed'])
```

[35]:
```
print(scores_nb_recovered.mean())
print(scores_nb_deaths.mean())
print(scores_nb_confirmed.mean())
```

0.7210144927536232
0.894927536231884
0.9456521739130435

1.1.2 Decision Tree Implementation

[56]:
```
#Label : Recovered
classifier_DecisionTree_recovered = DecisionTreeClassifier(max_depth = 3)
classifier_DecisionTree_recovered.fit(x,y.loc[:,'Recovered'])
scores_d_recovered = classifier_DecisionTree_recovered.score(x_test,y_test.loc[:
↪,'Recovered'])
# Label : Deaths
classifier_DecisionTree_deaths= DecisionTreeClassifier(max_depth = 3)
classifier_DecisionTree_deaths.fit(x,y.loc[:,'Deaths'])
scores_d_deaths = classifier_DecisionTree_deaths.score(x_test,y_test.loc[:
↪,'Deaths'])
#Label : Confirmed
classifier_DecisionTree_confirmed = DecisionTreeClassifier(max_depth = 3)
classifier_DecisionTree_confirmed.fit(x,y.loc[:,'Confirmed'])
scores_d_confirmed = classifier_DecisionTree_confirmed.score(x_test,y_test.loc[:
↪,'Confirmed'])
```

[57]:
```
print(scores_d_recovered.mean())
print(scores_d_deaths.mean())
print(scores_d_confirmed.mean())
```

0.7862318840579711
0.9057971014492754
0.9565217391304348

1.1.3 From the above accuracy values, it can be inferred that decision tree performs almost similar to gaussian naive bayes

1.2 Learning Parameters of NB for data Interpretation and comparision

[58]:
```
var = pd.DataFrame()
var[0]=(list(list(classifier_gaussian_recovered.sigma_)[0]))
var[1]=(list(list(classifier_gaussian_recovered.sigma_)[1]))
var.columns = ['True','False']
var.index = list(x.columns)
var.T
```

[58]:

	Lat	Long_	Active	Incident_Rate	Total_Test_Results	\
True	2.102807	1.903569	2.025485	0.807028	1.571866	
False	0.526993	0.603241	0.315986	1.233058	0.625900	

	Case_Fatality_Ratio	Testing_Rate	Resident Population 2020 Census	\
True	1.141637	1.445363	1.801357	

```
False             1.064626      0.914632                          0.496507

       Population Density 2020 Census  Density Rank 2020 Census  SexRatio
True                          0.354314                  1.047409  1.550491
False                         1.680216                  1.092018  0.881031
```

1.2.1 These represent the variance between two classes of Label recovered

[61]:
```
mean = pd.DataFrame()
mean[0]=(list(list(classifier_gaussian_recovered.theta_)[0]))
mean[1]=(list(list(classifier_gaussian_recovered.theta_)[1]))
mean.columns = ['True','False']
mean.index = list(x.columns)
mean.T
```

[61]:
```
            Lat     Long_    Active  Incident_Rate  Total_Test_Results  \
True  -0.222705 -0.274367  0.277679      -0.182613            0.192396
False  0.156457  0.143668 -0.210219       0.134183           -0.146766

       Case_Fatality_Ratio  Testing_Rate  Resident Population 2020 Census  \
True              0.079321      0.113771                          0.233726
False            -0.074567     -0.033171                         -0.173113

       Population Density 2020 Census  Density Rank 2020 Census  SexRatio
True                         -0.080190                 -0.237497 -0.119118
False                         0.065903                  0.149274  0.107242
```

1.2.2 These represent the mean of the two classes of label : recovered

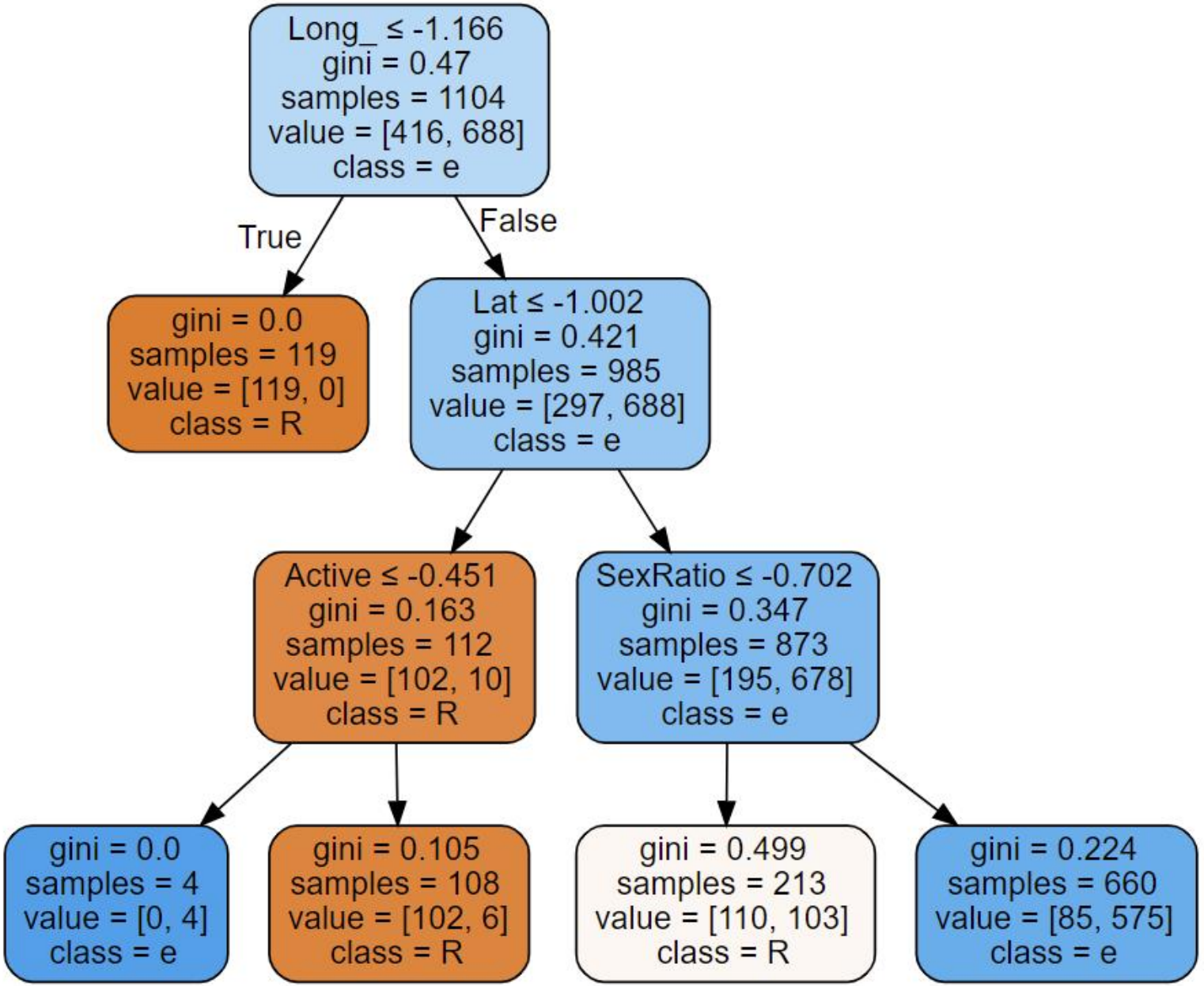
[ ]:
```
import graphviz
from sklearn import tree
dot_data = tree.export_graphviz(classifier_DecisionTree_recovered,out_file =␣
 ↪None,
                                feature_names = x.columns,
                                class_names = 'Recovered',
                                filled = True, rounded = True,
                                special_characters = True)
graph = graphviz.Source(dot_data)
graph
```

1.3 Inferences from learned parameters

1.3.1 The Long__ feature of 'False' class has very high variance and low mean. This plays a very crutial role in decision making to classify 'False' from the other parameters. Similarly from the above graph of decision trees, the Long__ becomes the head node and child nodes are created for the 'False' category. Thus from the above parameters it can be inferred that the decision making process of naive bayes and decision trees are similar though the parameters and calculations are different

[ ]:

Out[9]:


Long_ ≤ -1.166
gini = 0.47
samples = 1104
value = [416, 688]
class = e
True
False
gini = 0.0
samples = 119
value = [119, 0]
class = R
Lat ≤ -1.002
gini = 0.421
samples = 985
value = [297, 688]
class = e
Active ≤ -0.451
gini = 0.163
samples = 112
value = [102, 10]
class = R
SexRatio ≤ -0.702
gini = 0.347
samples = 873
value = [195, 678]
class = e
gini = 0.0
samples = 4
value = [0, 4]
class = e
gini = 0.105
samples = 108
value = [102, 6]
class = R
gini = 0.499
samples = 213
value = [110, 103]
class = R
gini = 0.224
samples = 660
value = [85, 575]
class = e