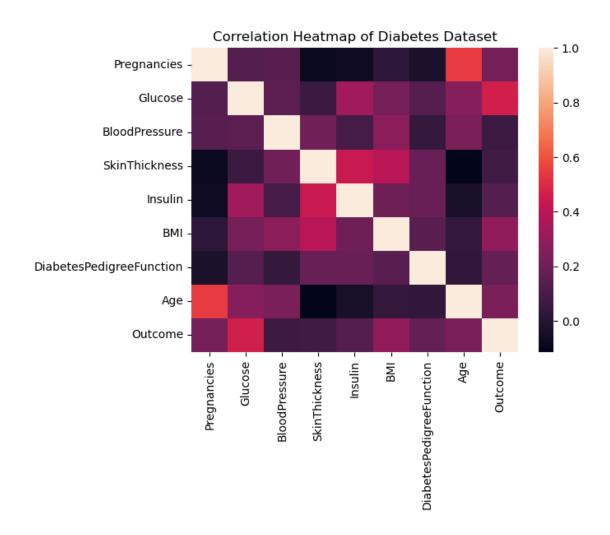
Diabetes Detection using DT and KNN - 21F21484 VIVEIK CATARAM SAICHANDRA

January 6, 2025

```
[1]: #Importing Necessary Libraries
     import numpy as np
     import pandas as pd
     import matplotlib.pyplot as plt
     import seaborn as sns
[2]: #Importing and Reading the Dataset
     dataset = pd.read_csv('diabetes.csv')
     dataset.head()
[2]:
        Pregnancies
                     Glucose BloodPressure SkinThickness
                                                             Insulin
                                                                        BMI
                         148
                                                         35
                                                                       33.6
     0
                  6
                          85
                                                         29
                                                                    0 26.6
     1
                  1
                                          66
     2
                  8
                         183
                                          64
                                                          0
                                                                    0 23.3
     3
                  1
                          89
                                          66
                                                         23
                                                                   94
                                                                       28.1
     4
                  0
                         137
                                          40
                                                                  168 43.1
                                                         35
        DiabetesPedigreeFunction
                                  Age
                                        Outcome
     0
                           0.627
                                    50
                           0.351
                                              0
     1
                                    31
                           0.672
     2
                                    32
                                              1
                                              0
     3
                           0.167
                                    21
     4
                           2.288
                                    33
                                              1
[3]: #Plotting Heatmap of the Dataset
     plt.figure(1)
     sns.heatmap(dataset.corr())
     plt.title('Correlation Heatmap of Diabetes Dataset')
     plt.show()
```



```
[4]: # Replace O values with the median of the feature's distribution to handle Zerous(O) values

columns_with_zeros = ['Pregnancies', 'Glucose', 'BloodPressure',us'SkinThickness', 'Insulin', 'BMI'] #List contatining parameters with O values

for column in columns_with_zeros:
    median = dataset[column].median()
    dataset[column] = dataset[column].replace(0, median)

dataset.head()
```

[4]:	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	\
0	6	148	72	35	27	33.6	
1	1	85	66	29	27	26.6	
2	8	183	64	23	27	23.3	
3	1	89	66	23	94	28.1	
4	3	137	40	35	168	43.1	

```
0
                            0.627
                                    50
                            0.351
      1
                                    31
                                              0
      2
                            0.672
                                    32
                                              1
      3
                            0.167
                                    21
                                              0
                            2.288
                                    33
                                              1
[23]: #Splitting dataset into train-test data
      from sklearn.model_selection import train_test_split
      x = dataset.drop(columns = ['Outcome']) #Features
      y = dataset['Outcome'] #Target
      x_train, x_test, y_train, y_test = train_test_split(x, y, test_size = 0.25)__
       \hookrightarrow#Train-Test Split
[24]: #DECISION TREE (DT)
      from sklearn.tree import DecisionTreeClassifier, plot_tree
      from sklearn.metrics import accuracy_score, classification_report,_
       ⇔confusion_matrix, roc_curve
      dt_classifier = DecisionTreeClassifier() #Decision Tree Classifier model
      dt_classifier.fit(x_train, y_train) #Training the Classifier with the training_
      dt_pred = dt_classifier.predict(x_test) #Using the trained Classifier model to_
       ⇔predict the testing data
      dt_model = DecisionTreeClassifier(criterion='entropy') #Decision TreeL
       →Classifier model for calculating Information Gain
      dt_model.fit(x_train, y_train) #Training the Classifier with the training data
      importances = dt_model.feature_importances_ #Retrieving the Feature importances
      tree_structure = dt_model.tree_ #Accessing the tree structure for calculating_
       ⇔the Gini index and Entropy
      #Displaying the Gini index and Entropy for each node (split) in the tree
      for i in range(tree_structure.node_count):
          if tree_structure.children_left[i] != tree_structure.children_right[i]: __
       \hookrightarrow#If condition for the case where it is a split node
              gini index = tree_structure.impurity[i] #Gini index impurity at the
       ⇔split node
              print(f'Node {i} Gini index: {gini index}')
              entropy_value = -np.sum(tree_structure.weighted_n_node_samples[i]) * np.
       →log2(tree_structure.weighted_n_node_samples[i]) #Calculating the Entropy
              print(f'Node {i} Entropy: {entropy_value}')
      #Plotting the Decision Tree
```

DiabetesPedigreeFunction Age Outcome

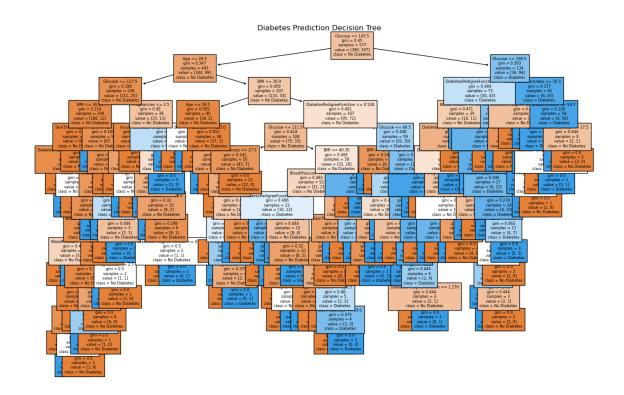
```
plt.figure(figsize=(15,10))
plot_tree(dt_classifier,feature_names = x.columns,class_names=['No_Diabetes',__
 ⇔'Diabetes'],filled=True,fontsize=6)
plt.title('Diabetes Prediction Decision Tree')
plt.show()
dt_report = classification_report(y_test, dt_pred) #Classification Report
dt_matrix = confusion_matrix(y_test, dt_pred) #Confusion Matrix
dt_accuracy = accuracy_score(y_test, dt_pred) #Accuracy Score
dt_fp_rate, dt_tp_rate, dt_threshold = roc_curve(y_test, dt_pred) #ROC Curve
dt_sensitivity = dt_matrix[0,0] / (dt_matrix[0,0] + dt_matrix[0,1])
  \hookrightarrow#Sensitivity = TP / (TP + FN)
dt_specificity = dt_matrix[1,1] / (dt_matrix[1,1] + dt_matrix[1,0])__
 \Rightarrow#Specificity = TN / (TN + FP)
dt_precision = dt_matrix[0,0] / (dt_matrix[0,0] + dt_matrix[1,0]) #Precision = ___
 \hookrightarrow TP / (TP + FP)
print(dt_report)
print(dt matrix)
print('Decision Tree Accuracy = ',dt_accuracy)
print('Decision Tree Sensitivity = ',dt_sensitivity)
print('Decision Tree Specificity = ',dt_specificity)
print('Decision Tree Precision = ',dt_precision)
plt.title('ROC Curve - Decision Tree')
plt.plot(dt_fp_rate, dt_tp_rate)
plt.plot([0, 1], ls="--")
plt.plot([0, 0], [1, 0], c=".7"), plt.plot([1, 1], c=".7")
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
Node 0 Gini index: 0.926172154055073
Node 0 Entropy: -5292.490672488443
Node 1 Gini index: 0.7664645115676525
Node 1 Entropy: -3894.485159629873
Node 2 Gini index: 0.4875229918975016
Node 2 Entropy: -1860.3037596493946
Node 3 Gini index: 0.32984607020714635
Node 3 Entropy: -1510.6126107757627
Node 4 Gini index: 0.08079313589591118
Node 4 Entropy: -664.3856189774724
Node 5 Gini index: 0.4394969869215134
Node 5 Entropy: -38.05374780501027
Node 7 Gini index: 1.0
Node 7 Entropy: -2.0
Node 11 Gini index: 0.5066503344840895
```

- Node 11 Entropy: -648.2415647232904
- Node 12 Gini index: 0.6442142137378307
- Node 12 Entropy: -406.42797576067073
- Node 13 Gini index: 0.5547781633412736
- Node 13 Entropy: -369.16017124398627
- Node 15 Gini index: 0.5140912790181235
- Node 15 Entropy: -361.7749775913361
- Node 16 Gini index: 0.4689955935892812
- Node 16 Entropy: -354.41343573651113
- Node 17 Gini index: 0.9709505944546686
- Node 17 Entropy: -11.60964047443681
- Node 20 Gini index: 0.3760198509692728
- Node 20 Entropy: -317.9747842438563
- Node 21 Gini index: 0.49418293484978865
- Node 21 Entropy: -192.74977452827116
- Node 22 Gini index: 0.3227569588973983
- Node 22 Entropy: -172.97373660251154
- Node 24 Gini index: 0.8112781244591328
- Node 24 Entropy: -24.0
- Node 26 Gini index: 1.0
- Node 26 Entropy: -8.0
- Node 28 Gini index: 0.9182958340544896
- Node 28 Entropy: -4.754887502163468
- Node 31 Gini index: 0.9182958340544896
- Node 31 Entropy: -4.754887502163468
- Node 36 Gini index: 0.9709505944546686
- Node 36 Entropy: -11.60964047443681
- Node 38 Gini index: 0.9182958340544896
- Node 38 Entropy: -4.754887502163468
- Node 42 Gini index: 0.9268190639645772
- Node 42 Entropy: -199.42124551085624
- Node 43 Gini index: 0.3095434291503252
- Node 43 Entropy: -75.05865002596161
- Node 44 Gini index: 0.7219280948873623
- Node 44 Entropy: -11.60964047443681
- Node 46 Gini index: 1.0
- Node 46 Entropy: -2.0
- Node 50 Gini index: 0.9709505944546686
- Node 50 Entropy: -86.43856189774725
- Node 51 Gini index: 0.9852281360342516
- Node 51 Entropy: -53.302968908806456
- Node 52 Gini index: 0.8453509366224365
- Node 52 Entropy: -38.05374780501027
- Node 53 Gini index: 0.9852281360342516
- Node 53 Entropy: -19.651484454403228
- Node 54 Gini index: 0.7219280948873623
- Node 54 Entropy: -11.60964047443681
- Node 61 Gini index: 0.9405781991505306

- Node 61 Entropy: -1592.5518002023603
- Node 62 Gini index: 0.18717625687320816
- Node 62 Entropy: -179.5249055930738
- Node 64 Gini index: 0.7219280948873623
- Node 64 Entropy: -11.60964047443681
- Node 67 Gini index: 0.9834537187362689
- Node 67 Entropy: -1277.3175378087608
- Node 68 Gini index: 0.885612871398971
- Node 68 Entropy: -762.4237512704516
- Node 70 Gini index: 0.9440870182837795
- Node 70 Entropy: -616.1313520576979
- Node 71 Gini index: 0.9736680645496201
- Node 71 Entropy: -536.9546635134159
- Node 73 Gini index: 0.9837082626231857
- Node 73 Entropy: -505.754247590989
- Node 75 Gini index: 0.9919924034538556
- Node 75 Entropy: -474.8424910217125
- Node 76 Gini index: 0.8112781244591328
- Node 76 Entropy: -43.01955000865387
- Node 77 Gini index: 0.4689955935892812
- Node 77 Entropy: -33.219280948873624
- Node 81 Gini index: 0.9652016987500656
- Node 81 Entropy: -384.0
- Node 82 Gini index: 0.905200296956048
- Node 82 Entropy: -303.57978409184955
- Node 83 Gini index: 0.9649567669505688
- Node 83 Entropy: -219.65963218934144
- Node 85 Gini index: 0.9867867202680318
- Node 85 Entropy: -192.74977452827116
- Node 86 Gini index: 0.9991421039919088
- Node 86 Entropy: -140.88144885869957
- Node 87 Gini index: 0.9828586897127056
- Node 87 Entropy: -122.21143267166839
- Node 88 Gini index: 0.6840384356390417
- Node 88 Entropy: -38.05374780501027
- Node 90 Gini index: 1.0
- Node 90 Entropy: -8.0
- Node 93 Gini index: 0.9709505944546686
- Node 93 Entropy: -58.60335893412778
- Node 95 Gini index: 1.0
- Node 95 Entropy: -43.01955000865387
- Node 96 Gini index: 0.9709505944546686
- Node 96 Entropy: -33.219280948873624
- Node 98 Gini index: 0.9182958340544896
- Node 98 Entropy: -15.509775004326936
- Node 100 Gini index: 0.9182958340544896
- Node 100 Entropy: -4.754887502163468 Node 105 Gini index: 0.5435644431995964

- Node 105 Entropy: -24.0
- Node 108 Gini index: 0.41381685030363374
- Node 108 Entropy: -43.01955000865387
- Node 111 Gini index: 0.8453509366224365
- Node 111 Entropy: -38.05374780501027
- Node 112 Gini index: 0.8112781244591328
- Node 112 Entropy: -8.0
- Node 117 Gini index: 0.934068055375491
- Node 117 Entropy: -354.41343573651113
- Node 119 Gini index: 0.885612871398971
- Node 119 Entropy: -325.2118756352258
- Node 120 Gini index: 0.97663491144401
- Node 120 Entropy: -206.1306865356277
- Node 121 Gini index: 0.9975025463691153
- Node 121 Entropy: -172.97373660251154
- Node 123 Gini index: 0.9709505944546686
- Node 123 Entropy: -147.20671786825557
- Node 124 Gini index: 0.961236604722876
- Node 124 Entropy: -48.105716335834195
- Node 126 Gini index: 0.8453509366224365
- Node 126 Entropy: -38.05374780501027
- Node 128 Gini index: 0.8112781244591328
- Node 128 Entropy: -8.0
- Node 131 Gini index: 0.7871265862012691
- Node 131 Entropy: -69.48686830125577
- Node 133 Gini index: 0.9709505944546686
- Node 133 Entropy: -33.219280948873624
- Node 135 Gini index: 0.8112781244591328
- Node 135 Entropy: -24.0
- Node 137 Gini index: 0.9182958340544896
- Node 137 Entropy: -4.754887502163468
- Node 141 Gini index: 0.3227569588973983
- Node 141 Entropy: -69.48686830125577
- Node 144 Gini index: 0.8395304981054318
- Node 144 Entropy: -946.8559515213415
- Node 145 Gini index: 0.9770012394218561
- Node 145 Entropy: -451.8571927982413
- Node 146 Gini index: 0.9575534837147482
- Node 146 Entropy: -140.88144885869957
- Node 147 Gini index: 0.8554508105601307
- Node 147 Entropy: -116.09640474436812
- Node 148 Gini index: 0.9886994082884974
- Node 148 Entropy: -64.0
- Node 150 Gini index: 0.9957274520849256
- Node 150 Entropy: -48.105716335834195
- Node 151 Gini index: 0.9709505944546686
- Node 151 Entropy: -33.219280948873624
- Node 152 Gini index: 0.9182958340544896

- Node 152 Entropy: -15.509775004326936
- Node 159 Gini index: 0.8453509366224365
- Node 159 Entropy: -240.21499122004107
- Node 160 Gini index: 0.6892019851173655
- Node 160 Entropy: -199.42124551085624
- Node 162 Gini index: 0.8112781244591328
- Node 162 Entropy: -134.6059378176129
- Node 163 Gini index: 0.9023932827949789
- Node 163 Entropy: -98.10749561002054
- Node 164 Gini index: 0.9640787648082292
- Node 164 Entropy: -75.05865002596161
- Node 165 Gini index: 0.9709505944546686
- Node 165 Entropy: -33.219280948873624
- Node 166 Gini index: 0.8112781244591328
- Node 166 Entropy: -24.0
- Node 167 Gini index: 1.0
- Node 167 Entropy: -8.0
- Node 169 Gini index: 0.9182958340544896
- Node 169 Entropy: -4.754887502163468
- Node 174 Gini index: 0.5435644431995964
- Node 174 Entropy: -24.0
- Node 176 Gini index: 1.0
- Node 176 Entropy: -2.0
- Node 181 Gini index: 0.6500224216483541
- Node 181 Entropy: -15.509775004326936
- Node 184 Gini index: 0.46377734988775166
- Node 184 Entropy: -361.7749775913361
- Node 185 Gini index: 0.9182958340544896
- Node 185 Entropy: -4.754887502163468
- Node 188 Gini index: 0.362051251733998
- Node 188 Entropy: -339.76289771739914
- Node 189 Gini index: 0.225363639127395
- Node 189 Entropy: -317.9747842438563
- Node 190 Gini index: 0.13503620280212764
- Node 190 Entropy: -303.57978409184955
- Node 192 Gini index: 1.0
- Node 192 Entropy: -2.0
- Node 195 Gini index: 1.0
- Node 195 Entropy: -2.0
- Node 198 Gini index: 0.9182958340544896
- Node 198 Entropy: -4.754887502163468

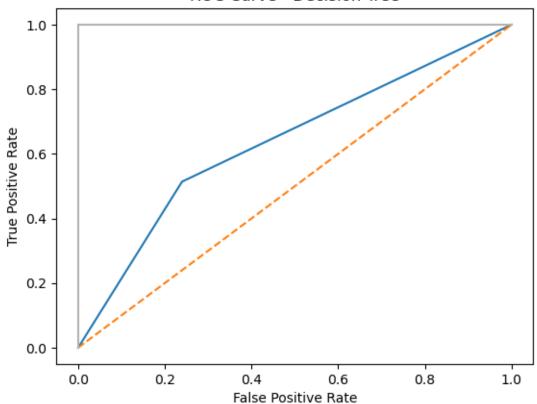


support	f1-score	recall	precision	
121	0.74	0.76	0.72	0
72	0.54	0.51	0.56	1
193	0.67			accuracy
193	0.64	0.64	0.64	macro avg
193	0.67	0.67	0.66	weighted avg

[[92 29] [35 37]]

Decision Tree Accuracy = 0.6683937823834197
Decision Tree Sensitivity = 0.7603305785123967
Decision Tree Specificity = 0.5138888888888888888
Decision Tree Precision = 0.7244094488188977

ROC Curve - Decision Tree



```
[18]: #K-NEAREST NEIGHBOUR (KNN)
from sklearn.preprocessing import MinMaxScaler

#Normalization Preprocessing Technique
scaler = MinMaxScaler()

#Columns that require normalization
scale_columns = ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness',
'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age']
normalized_data = scaler.fit_transform(dataset[scale_columns])
normalized_dataset = pd.DataFrame(normalized_data, columns = scale_columns)
#Converting normalized data into dataset
normalized_dataset['Outcome'] = dataset['Outcome'] #Adding the unaltered Target
yvariable to the Normalized Dataset
normalized_dataset.head()
```

```
[18]: Pregnancies Glucose BloodPressure SkinThickness Insulin BMI \
0 0.3125 0.670968 0.489796 0.304348 0.015625 0.314928
1 0.0000 0.264516 0.428571 0.239130 0.015625 0.171779
```

```
2
              0.4375 0.896774
                                     0.408163
                                                     0.173913 0.015625 0.104294
      3
              0.0000 0.290323
                                                     0.173913 0.096154 0.202454
                                     0.428571
      4
              0.1250 0.600000
                                     0.163265
                                                     0.304348 0.185096 0.509202
         DiabetesPedigreeFunction
                                        Age Outcome
      0
                         0.234415 0.483333
                         0.116567 0.166667
                                                    0
      1
      2
                         0.253629 0.183333
                                                    1
      3
                         0.038002 0.000000
                                                    0
      4
                         0.943638 0.200000
                                                    1
[19]: #Splitting dataset into train-test data
      x_normal = normalized_dataset.drop(columns = 'Outcome') #Features
      y_normal = normalized_dataset['Outcome'] #Target
      x_train_normal, x_test_normal, y_train_normal, y_test_normal =_
       otrain_test_split(x_normal, y_normal, test_size = 0.25) #Train-Test Split
[20]: from sklearn.neighbors import KNeighborsClassifier
      knn_classifier = KNeighborsClassifier(n_neighbors = 5) #K-Nearest Neighbor
       \hookrightarrow Classifier model
      knn_classifier.fit(x_train_normal, y_train_normal) #Training the Classifier_u
       ⇔with the training data
      knn_pred = knn_classifier.predict(x_test_normal) #Using the trained Classifier_
      →model to predict the testing data
      knn_report = classification_report(y_test_normal, knn_pred) #Classification_
       \hookrightarrowReport
      knn_matrix = confusion_matrix(y_test_normal, knn_pred) #Confusion Matrix
      knn_accuracy = accuracy_score(y_test_normal, knn_pred) #Accuracy Score
      knn_fp_rate, knn_tp_rate, knn_threshold = roc_curve(y_test_normal, knn_pred)_
       →#ROC Curve
      knn_sensitivity = knn_matrix[0,0] / (knn_matrix[0,0] + knn_matrix[0,1])
       ⇔#Sensitivity = TP / (TP + FN)
      knn_specificity = knn_matrix[1,1] / (knn_matrix[1,1] + knn_matrix[1,0])
      \hookrightarrow#Specificity = TN / (TN + FP)
      knn_precision = knn_matrix[0,0] / (knn_matrix[0,0] + knn_matrix[1,0])
       \Rightarrow#Precision = TP / (TP + FP)
      print(knn_report)
      print(knn matrix)
      print('K-Nearest Neighbor Accuracy = ',knn_accuracy)
      print('K-Nearest Neighbor Sensitivity = ',knn_sensitivity)
      print('K-Nearest Neighbor Specificity = ',knn_specificity)
      print('K-Nearest Neighbor Precision = ',knn precision)
      plt.title('ROC Curve - K-Nearest Neighbor')
```

```
plt.plot(knn_fp_rate, knn_tp_rate)
plt.plot([0, 1], ls="--")
plt.plot([0, 0], [1, 0], c=".7"), plt.plot([1, 1], c=".7")
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
```

	precision	recall	f1-score	support
0	0.76	0.81	0.78	126
1	0.59	0.52	0.56	67
accuracy			0.71	193
macro avg	0.68	0.67	0.67	193
weighted avg	0.70	0.71	0.71	193

[[102 24] [32 35]]

K-Nearest Neighbor Accuracy = 0.7098445595854922K-Nearest Neighbor Sensitivity = 0.8095238095238095K-Nearest Neighbor Specificity = 0.5223880597014925K-Nearest Neighbor Precision = 0.7611940298507462

ROC Curve - K-Nearest Neighbor

