Diabetes Prediction Using Decision Tree and Multinomial Naive Bayes Classifiers

June 21, 2024

- 1 PIMA Indian Diabetes Prediction Analysis Using Decision Tree and Multinomial Naive Bayes Classifiers
- 2 Import the Required Libraries

```
[83]: # Data analysis and wrangling
      import numpy as np
      import pandas as pd
      # Visualization
      import matplotlib.pyplot as plt
      import statsmodels.api as sm
      import seaborn as sns
      from sklearn.tree import DecisionTreeClassifier # Import Decision Tree
       \hookrightarrowClassifier
      from sklearn.model_selection import train_test_split # Import train_test_split_
      from sklearn import metrics #Import scikit-learn metrics module for accuracy_
       \hookrightarrow calculation
      from sklearn.metrics import confusion matrix, classification report
      # suppress the warning adding the following lines to the imports of your program
      import warnings
      #warnings.simplefilter(action='ignore', category=FutureWarning)
      # ignore all warnings
      warnings.filterwarnings('ignore')
      # sns.set(rc={'figure.figsize': [20, 20]}, font_scale=1.4)
      sns.set_theme(color_codes=True)
```

3 Load the Dataset

```
[84]: col_names = ['pregnant', 'glucose', 'bp', 'skin', 'insulin', 'bmi', 'pedigree', _
      # load dataset
     pima = pd.read_csv("data/pima-indians-diabetes.csv", header=None,
       →names=col_names)
[85]: # let's look at the datatype of each column
     pima.info()
     <class 'pandas.core.frame.DataFrame'>
     RangeIndex: 768 entries, 0 to 767
     Data columns (total 9 columns):
                   Non-Null Count Dtype
          Column
          _____
                   -----
          pregnant 768 non-null
      0
                                   int64
          glucose
                   768 non-null
                                   int.64
      1
      2
         bp
                   768 non-null
                                  int64
                   768 non-null
      3
          skin
                                  int64
      4
          insulin 768 non-null
                                  int64
      5
                   768 non-null
                                   float64
          bmi
      6
          pedigree 768 non-null
                                   float64
      7
          age
                   768 non-null
                                   int64
          label
                   768 non-null
                                   int64
     dtypes: float64(2), int64(7)
     memory usage: 54.1 KB
[86]: #let's see if any column has any null values
     pima.isna().sum()
[86]: pregnant
     glucose
                 0
     bр
                 0
     skin
                 0
     insulin
                 0
     bmi
     pedigree
                 0
     age
     label
     dtype: int64
        Feature Selection
[87]: #split dataset in features and target variable
     feature_cols = ['pregnant', 'insulin', 'bmi', 'age', 'glucose', 'bp', 'pedigree']
     X = pima[feature_cols] # Features
```

```
y = pima.label # Target variable
```

5 Split the Dataset

6 Correlation Heatmap for the Dataset

A correlation heatmap is a graphical tool that displays the correlation between multiple variables as a color-coded matrix. It's like a color chart that shows us how closely related different variables are.

In a correlation heatmap, each variable is represented by a row and a column, and the cells show the correlation between them. The color of each cell represents the strength and direction of the correlation, with darker colors indicating stronger correlations.

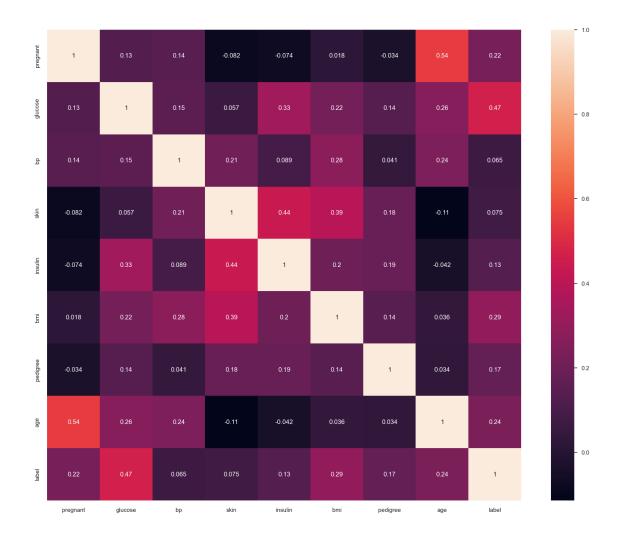
How to Read a Correlation Heatmap? In this section, we will delve into how to read a correlation heatmap, an effective visual tool for discerning the strength and direction of relationships between variables:

- Look at the color of each cell to see the strength and direction of the correlation.
- Darker colors indicate stronger correlations, while lighter colors indicate weaker correlations.
- Positive correlations (when one variable increases, the other variable tends to increase) are usually represented by warm colors, such as red or orange.
- Negative correlations (when one variable increases, the other variable tends to decrease) are usually represented by cool colors, such as blue or green.

Understanding correlation heatmaps can help us identify patterns and relationships between multiple variables. So next time you analyze data with many variables, think like an artist and use a correlation heatmap to see the colors of the relationships!

```
[89]: # Correlation Heatmap (print the correlation score each variables)
plt.figure(figsize=(20, 16))
sns.heatmap(pima.corr(), fmt='.2g', annot=True)
```

[89]: <Axes: >



7 1. Decision Tree Classifier in Scikit-learn

8 Bulid the Decision Tree Model

Evaluate the Decision Tree Model

Confusion Matrix and computing Sensitivity and Specificity

```
[91]: cm_dtree = confusion_matrix(y_test, y_pred)
      # Calculating accuracy, sensitivity and specificity from confusion matrix
      total_dtree=sum(sum(cm_dtree))
      accuracy_dtree=(cm_dtree[0,0]+cm_dtree[1,1])/total_dtree
      print('Accuracy : ', accuracy_dtree)
      sensitivity_dtree = cm_dtree[0,0]/(cm_dtree[0,0]+cm_dtree[0,1])
      print('Sensitivity : ', sensitivity_dtree)
      specificity_dtree = cm_dtree[1,1]/(cm_dtree[1,0]+cm_dtree[1,1])
      print('Specificity : ', specificity_dtree)
      tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel()
      #tn = cm_dtree[0][0]
      #fn = cm_dtree[1][0]
      #tp = cm_dtree[1][1]
      #fp = cm_dtree[0][1]
      print("True Negative = ",tn)
      print("False Negative = ",fn)
      print("True Positive = ",tp)
      print("False Positive = ",fp)
      print(classification_report(y_test, y_pred, zero_division=1))
      sns.set context ("poster")
      # Confusion matrix and derived metrics Display - New Format
      plt.figure(figsize=(5,5))
      from sklearn.metrics import ConfusionMatrixDisplay
      _ = ConfusionMatrixDisplay.from_estimator(dtree, X_test , y_test)
      plt.ylabel('Actual label')
      plt.xlabel('Predicted label')
      #all_sample_title = 'Accuracy Score for Decision Tree Classifier is {0}'.
       ⇔format(dtree.score(X_test, y_test))
      #plt.title(all_sample_title, size = 15)
      plt.show()
```

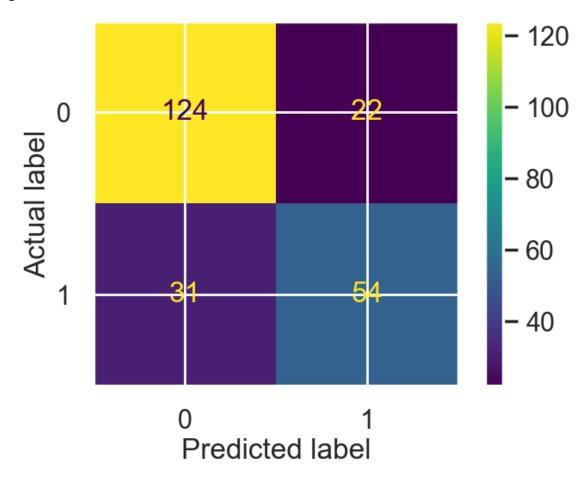
Accuracy: 0.7705627705627706 Sensitivity: 0.8493150684931506 Specificity: 0.6352941176470588

True Negative = 124

False Negative = 31 True Positive = 54 False Positive = 22

	precision	recall	f1-score	support
0	0.80	0.85	0.82	146
1	0.71	0.64	0.67	85
accuracy			0.77	231
macro avg	0.76	0.74	0.75	231
weighted avg	0.77	0.77	0.77	231

<Figure size 500x500 with 0 Axes>



11 Compute the other measures or metrics for the Decision Tree Model

Accuracy Score of Decision Tree Classifier is 77.06 % Accuracy Score of Decision Tree Classifier is 0.7705627705627706 Training Score of Decision Tree Classifier:76.35009310986965% Test score of Decision Tree Classifier:77.05627705627705%

```
[93]: from sklearn.metrics import accuracy_score, f1_score, precision_score,
      recall_score, jaccard_score, log_loss
      dtree_f1_score = f1_score(y_test, y_pred, average='micro')
      dtree_precision_score = precision_score(y_test, y_pred, average='micro')
      dtree_recall_score = recall_score(y_test, y_pred, average='micro')
      dtree_jaccard_score = jaccard_score(y_test, y_pred, average='micro')
      dtree_log_loss = log_loss(y_test, y_pred)
      print('Scores Calculation using average: micro')
      print('F-1 Score of Decision Tree Classifier is ', dtree_f1_score)
      print('Precision Score of Decision Tree Classifier is ', dtree precision score)
      print('Recall Score of Decision Tree Classifier is ', dtree_recall_score)
      print('Jaccard Score of Decision Tree Classifier is ', dtree_jaccard_score)
      print('Log Loss of Decision Tree Classifier is ', dtree_log_loss)
      # Manual Caluclation
      from sklearn.metrics import confusion_matrix
      tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel()
      dtree_accuracy_calculated = (tp + tn) / (tp + fp + fn + tn)
      dtree precision calculated = tp / (tp + fp)
      dtree_recall_calculated = tp / (tp + fn)
```

Scores Calculation using average: micro

F-1 Score of Decision Tree Classifier is 0.7705627705627706
Precision Score of Decision Tree Classifier is 0.7705627705627706
Recall Score of Decision Tree Classifier is 0.7705627705627706
Jaccard Score of Decision Tree Classifier is 0.6267605633802817
Log Loss of Decision Tree Classifier is 8.269755972394846
Scores Calculation using Manual Method

F-1 Score of Decision Tree Classifier is 0.6708074534161491 Precision Score of Decision Tree Classifier is 0.7105263157894737 Recall Score of Decision Tree Classifier is 0.6352941176470588 Accuracy Score of Decision Tree Classifier is 0.7705627705627706

12 Visualize the Decision Tree

```
[94]: #! pip install graphviz
[95]: #! pip install pydotplus
```

12.1 export_graphviz function converts decision tree classifier into dot file and pydotplus convert this dot file to png or displayable form on Jupyter.

```
graph.write_png('diabetes.png')
               Image(graph.create_png())
[96]:
                                                                                                             glucose ≤ 127.5
                                                                                                             entropy = 0.926
                                                                                                              samples = 537
                                                                                                             value = [354, 183]
class = 0
                                                                                                    True
                                                                                        bmi ≤ 26.45
                                                                                                                                      bmi ≤ 28.15
                                                                                      entropy = 0.72
                                                                                      samples = 342
                                                                                                                                     samples = 195
                                                                                      value = [274, 68]
                                                                                                                                    value = [80, 115]
                                                                                         class = 0
                                                                                                                                        class = 1
                                                                                                                                                                        glucose ≤ 158.5
                                                                                         age ≤ 27.5
                                                                                                                                     glucose ≤ 145.5
                                                  entropy = 0.201
samples = 96
                                                                                      entropy = 0.833
samples = 246
                                                                                                                                                                       entropy = 0.9
samples = 152
value = [48, 104]
                                                                                                                                     entropy = 0.82
                                                   value = [93, 3]
                                                                                      value = [181, 65]
                                                                                                                                     value = [32, 11]
                                                                                         class = 0
                                                                                                                                        class = 0
                                                                                                                                                                                                entropy = 0.544
samples = 56
value = [7, 49]
class = 1
                          entropy = 0.918
                                                                          entropy = 0.544
                                                                                                  entropy = 0.958
                                                                                                                          entropy = 0.402
                                                                                                                                                  entropy = 1.0
                                                                                                                                                                        entropy = 0.985
                                                   entropy = 0.088
                                                   samples = 90
value = [89, 1]
                                                                                                                           samples = 25
value = [23, 2]
                            samples = 6
                                                                                                  samples = 134
value = [83, 51]
                                                                                                                                                  samples = 18
value = [9, 9]
                                                                                                                                                                        samples = 96
value = [41, 55]
                            value = [4, 2]
                                                                          value = [98, 14]
                              class = 0
                                                                                                      class = 0
                                                                                                                                                     class = 0
                                                                                                                                                                           class = 1
```

13 Plot ROC Cure of the Decision Tree Model

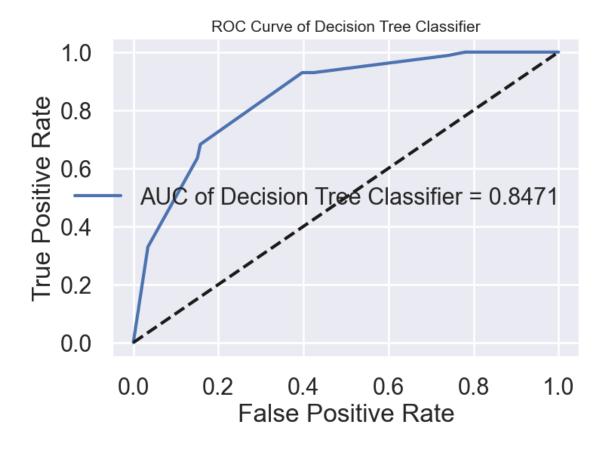
```
[97]: from sklearn.metrics import roc_curve, roc_auc_score
      y_pred_proba = dtree.predict_proba(X_test)[:][:,1]
      df_actual_predicted = pd.concat([pd.DataFrame(np.array(y_test),__

¬columns=['y_actual']), pd.DataFrame(y_pred_proba,

columns=['y_pred_proba'])], axis=1)
      df_actual_predicted.index = y_test.index
      \# Calculate and print the ROC (Receiver Operating Characteristic) curve and AUC_{f U}
       ⇔ (Area under the ROC Curve)
      fpr_dtree, tpr_dtree = roc_curve(df_actual_predicted['y_actual'],__
       →df_actual_predicted['y_pred_proba'])
      auc dtree = roc auc score(df actual predicted['y actual'],

¬df_actual_predicted['y_pred_proba'])
      plt.plot(fpr_dtree, tpr_dtree, label='AUC of Decision Tree Classifier = %0.4f'_
       plt.plot(fpr_dtree, fpr_dtree, linestyle = '--', color='k')
      plt.xlabel('False Positive Rate')
      plt.ylabel('True Positive Rate')
      plt.title('ROC Curve of Decision Tree Classifier', size = 15)
      plt.legend()
```

[97]: <matplotlib.legend.Legend at 0x19ef8551510>



14 2. Multinomial Naive Bayes Classifier in Scikit-learn

15 Bulid the Multinomial Naive Bayes Model

Accuracy of Multinomial Naive Bayes Classifier: 55.84415584415584

16 Evaluate the Multinomial Naive Bayes Model

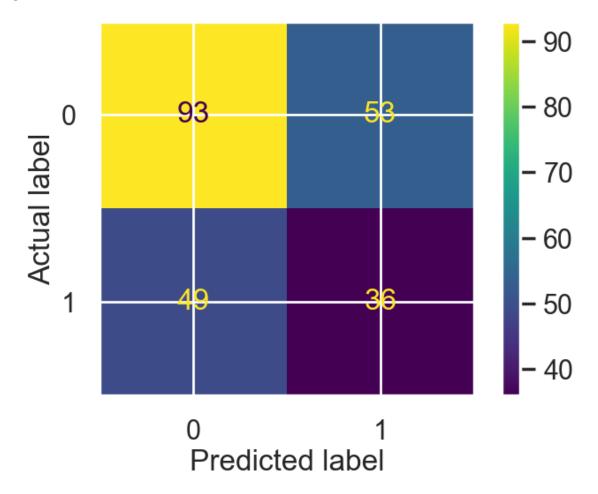
17 Confusion Matrix and computing Sensitivity and Specificity

```
[99]: cm_mnb = confusion_matrix(y_test, y_pred)
      # Calculating accuracy, sensitivity and specificity from confusion matrix
      total_mnb=sum(sum(cm_mnb))
      accuracy_mnb=(cm_mnb[0,0]+cm_mnb[1,1])/total_mnb
      print('Accuracy : ', accuracy_mnb)
      sensitivity\_mnb = cm\_mnb[0,0]/(cm\_mnb[0,0]+cm\_mnb[0,1])
      print('Sensitivity : ', sensitivity_mnb)
      specificity_mnb = cm_mnb[1,1]/(cm_mnb[1,0]+cm_mnb[1,1])
      print('Specificity : ', specificity_mnb)
      tn, fp, fn, tp = confusion matrix(y test, y pred).ravel()
      #tn = cm_mnb[0][0]
      #fn = cm_mnb[1][0]
      #tp = cm_mnb[1][1]
      #fp = cm_mnb[0][1]
      print("True Negative = ",tn)
      print("False Negative = ",fn)
      print("True Positive = ",tp)
      print("False Positive = ",fp)
      print(classification_report(y_test, y_pred, zero_division=1))
      sns.set context ("poster")
      # Confusion matrix and derived metrics Display - New Format
      plt.figure(figsize=(5,5))
      from sklearn.metrics import ConfusionMatrixDisplay
      _ = ConfusionMatrixDisplay.from_estimator(mnb, X_test , y_test)
      plt.ylabel('Actual label')
      plt.xlabel('Predicted label')
      #all_sample_title = 'Accuracy Score for Multinomial Naive Bayes Classifier is_
       \rightarrow {0}'.format(mnb.score(X_test, y_test))
      #plt.title(all_sample_title, size = 15)
      plt.show()
     Accuracy: 0.5584415584415584
     Sensitivity: 0.636986301369863
     Specificity: 0.4235294117647059
     True Negative = 93
```

False Negative = 49 True Positive = 36 False Positive = 53

	precision	recall	f1-score	support
0	0.65	0.64	0.65	146
1	0.40	0.42	0.41	85
accuracy			0.56	231
macro avg	0.53	0.53	0.53	231
weighted avg	0.56	0.56	0.56	231

<Figure size 500x500 with 0 Axes>



18 Compute the other measures for the Multinomial Naive Bayes Model

Accuracy Score of Multinomial Naive Bayes Classifier is 55.84 % Accuracy Score of Multinomial Naive Bayes Classifier is 0.5584415584415584 Training Score of Multinomial Naive Bayes Classifier:60.33519553072626% Test score of Multinomial Naive Bayes Classifier:55.84415584415584%

```
[101]: from sklearn.metrics import accuracy_score, f1_score, precision_score,
       →recall_score, jaccard_score, log_loss
      mnb f1 score = f1 score(y test, y pred, average='macro')
      mnb_precision_score = precision_score(y_test, y_pred, average='macro')
      mnb_recall_score = recall_score(y_test, y_pred, average='macro')
      mnb_jaccard_score = jaccard_score(y_test, y_pred, average='macro')
      mnb_log_loss = log_loss(y_test, y_pred)
      mnb_f1_score = f1_score(y_test, y_pred, average='micro')
      mnb_precision_score = precision_score(y_test, y_pred, average='micro')
      mnb_recall_score = recall_score(y_test, y_pred, average='micro')
      mnb_jaccard_score = jaccard_score(y_test, y_pred, average='micro')
      mnb_log_loss = log_loss(y_test, y_pred)
      print('Scores Calculation using average: micro')
      print('=======')
      print('F-1 Score of Multinomial Naive Bayes Classifier is ', mnb_f1_score)
      print('Precision Score of Multinomial Naive Bayes Classifier is ', u
       →mnb precision score)
      print('Recall Score of Multinomial Naive Bayes Classifier is ', u
       →mnb_recall_score)
      print('Jaccard Score of Multinomial Naive Bayes Classifier is ', _
        →mnb_jaccard_score)
      print('Log Loss of Multinomial Naive Bayes Classifier is ', mnb_log_loss)
```

```
# Manual Caluclation
from sklearn.metrics import confusion_matrix
tn, fp, fn, tp = confusion_matrix(y_test, y_pred).ravel()
mnb_accuracy_calculated = (tp + tn) / (tp + fp + fn + tn)
mnb_precision_calculated = tp / (tp + fp)
mnb_recall_calculated = tp / (tp + fn)
mnb_f1_score_calculated = 2 * ( (mnb_precision_calculated *_

mnb_recall_calculated) / (mnb_precision_calculated + mnb_recall_calculated) )
print('Scores Calculation using Manual Method')
print('======"")
print('F-1 Score of Multinomial Naive Bayes Classifier is ', 
 →mnb_f1_score_calculated)
print('Precision Score of Multinomial Naive Bayes Classifier is ', u
 →mnb_precision_calculated)
print('Recall Score of Multinomial Naive Bayes Classifier is ', ...
 →mnb_recall_calculated)
print("Accuracy Score of Multinomial Naive Bayes Classifier is ", u
 →mnb_accuracy_calculated)
```

Scores Calculation using average: micro

F-1 Score of Multinomial Naive Bayes Classifier is 0.5584415584415584
Precision Score of Multinomial Naive Bayes Classifier is 0.5584415584415584
Recall Score of Multinomial Naive Bayes Classifier is 0.5584415584415584
Jaccard Score of Multinomial Naive Bayes Classifier is 0.38738738738738737
Log Loss of Multinomial Naive Bayes Classifier is 15.915379418571211
Scores Calculation using Manual Method

F-1 Score of Multinomial Naive Bayes Classifier is 0.41379310344827586 Precision Score of Multinomial Naive Bayes Classifier is 0.4044943820224719 Recall Score of Multinomial Naive Bayes Classifier is 0.4235294117647059

Accuracy Score of Multinomial Naive Bayes Classifier is 0.5584415584415584

19 Plot ROC Cure of the Multinomial Naive Bayes Model

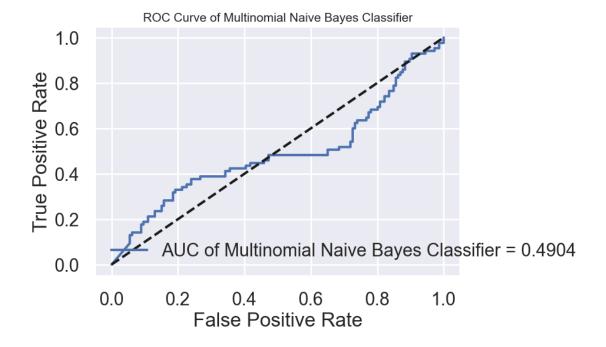
```
[102]: from sklearn.metrics import roc_curve, roc_auc_score
y_pred_proba = mnb.predict_proba(X_test)[:][:,1]

df_actual_predicted = pd.concat([pd.DataFrame(np.array(y_test),__
columns=['y_actual']), pd.DataFrame(y_pred_proba,__
columns=['y_pred_proba'])], axis=1)

df_actual_predicted.index = y_test.index

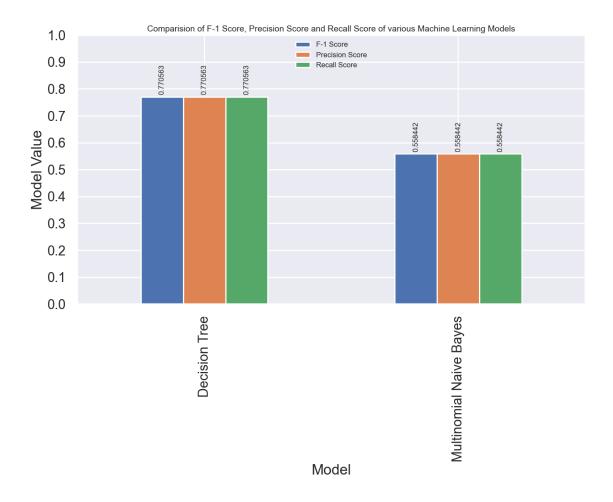
# Calculate and print the ROC (Receiver Operating Characteristic) curve and AUC_
columns = (Area under the ROC Curve)
```

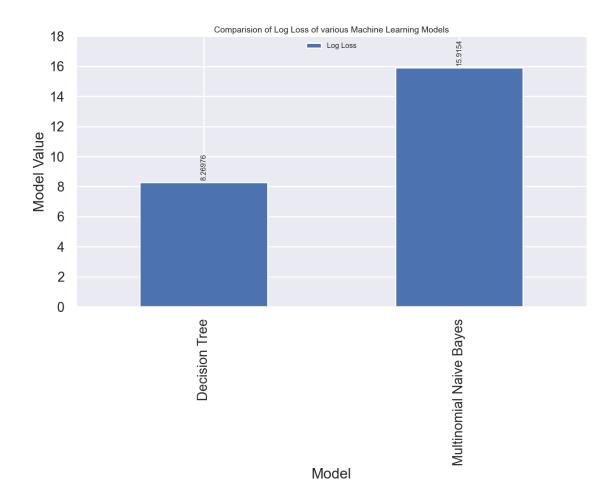
[102]: <matplotlib.legend.Legend at 0x19ef8afa350>



```
[103]: models = pd.DataFrame({
    'Model': ["Decision Tree","Multinomial Naive Bayes"],
    'F-1 Score': [dtree_f1_score, mnb_f1_score],
    'Precision Score': [dtree_precision_score, mnb_precision_score],
    'Recall Score': [dtree_recall_score, mnb_recall_score],
    'Log Loss': [dtree_log_loss, mnb_log_loss]})
sorted_models=models.sort_values(by=['F-1 Score'], ascending=False)
sorted_models
```

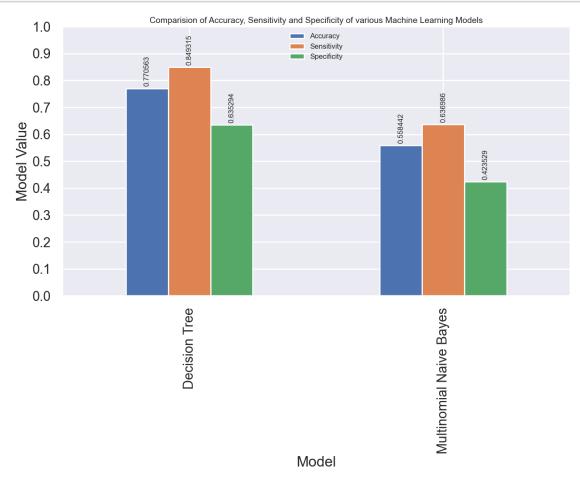
```
[103]:
                           Model F-1 Score Precision Score Recall Score \
                   Decision Tree 0.770563
                                                    0.770563
                                                                  0.770563
      0
      1 Multinomial Naive Bayes
                                   0.558442
                                                    0.558442
                                                                  0.558442
          Log Loss
         8.269756
      1 15.915379
[105]: # Plotting the graph
      axs = pd.concat([models['F-1 Score'],models['Precision Score'],models['Recall_
       Score']], axis=1).plot.bar(figsize=(15, 8))
      axs.set_title('Comparision of F-1 Score, Precision Score and Recall Score of
       →various Machine Learning Models', fontsize=14)
      axs.set_xlabel('Model')
      axs.set_ylabel('Model Value')
      axs.set_xticklabels(models['Model'])
      for container in axs.containers:
          axs.bar_label(container, padding=3, rotation=90, fontsize=12)
      plt.yticks(np.arange(0, 1.1, step=0.1))
      #axs.legend(loc='upper center', ncols=3)
      axs.legend(loc='upper center',fontsize=12)
      plt.show()
```





```
[107]: models = pd.DataFrame({
           'Model': ["Decision Tree", "Multinomial Naive Bayes"],
           'Accuracy': [accuracy_dtree, accuracy_mnb],
           'Sensitivity': [sensitivity_dtree, sensitivity_mnb],
           'Specificity': [specificity_dtree, specificity_mnb]})
       sorted_models=models.sort_values(by=['Accuracy'], ascending=False)
       sorted_models
[107]:
                            Model Accuracy
                                             Sensitivity
                                                          Specificity
       0
                    Decision Tree
                                   0.770563
                                                0.849315
                                                             0.635294
                                                             0.423529
       1 Multinomial Naive Bayes
                                   0.558442
                                                0.636986
[109]: # Plotting the graph
       axs = pd.

¬concat([models['Accuracy'],models['Sensitivity'],models['Specificity']],
□
        →axis=1).plot.bar(figsize=(15, 8))
```



20 Results and Discussion

- \bullet The Decision Tree is predicting more accurately the diabetes than the Multinomial Naive Bayes and its accuracy is 77.05%
- $\bullet\,$ The AUC value of Decision Tree is 0.847 which is higher than the AUC value of Multinomial Naive Bayes.