Task 3

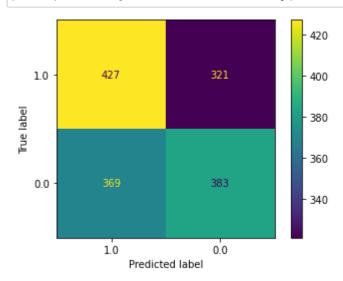
· Part B - Principal Component Analysis

```
In [1]: # PCA
        #Importing libraries
        import pandas as pd
        import numpy as np
        import matplotlib.pyplot as plt
In [2]: # Importing and reading csv file
        df=pd.read csv("E:HIGGS 6M.csv")
In [3]: # Anaylsing the distribution of classes
        # data["column_name"].value_counts(), returns unique values in that column
        print(df['1.000000000000000000e+00'].value counts())
        print('Zeros', round(df['1.00000000000000000e+00'].value counts()[0]/len(df) * 1
        print('Ones', round(df['1.000000000000000000e+00'].value_counts()[1]/len(df) * 10
        1.0
               3178344
        0.0
               2821655
        Name: 1.000000000000000000e+00, dtype: int64
        Zeros 47.03 % of the dataset
        Ones 52.97 % of the dataset
In [4]: Ones_df = df.loc[df['1.000000000000000000e+00'] == 1][0:2500] # smaples which had
        Zeros_df = df.loc[df['1.00000000000000000e+00'] == 0][0:2500] #
        normal distributed df = pd.concat([Ones df, Zeros df])
        # Shuffle dataframe rows
        df_new= normal_distributed_df.sample(frac=1, random_state=100)
        print(df new['1.00000000000000000e+00'].value counts()/len(df))
        0.0
               0.000417
        1.0
               0.000417
        Name: 1.000000000000000000e+00, dtype: float64
In [5]: # separating dependent and independent feature
        X = df new.drop("1.00000000000000000e+00",axis=1)
        y = df_new["1.000000000000000000e+00"]
In [6]: #Splitting the dataset into the training and test sets
        from sklearn.model_selection import train_test_split
        X_train, X_test, y_train, y_test = train_test_split(X,y, test_size = 0.3, random)
```

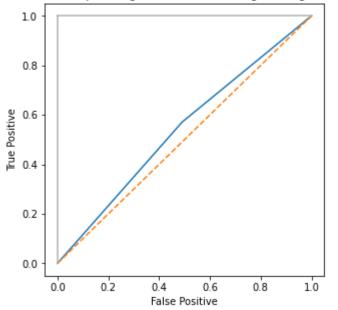
```
In [7]: #Putting the values in the same scale
         from sklearn.preprocessing import StandardScaler
         scalar = StandardScaler()
         X train = scalar.fit transform(X train)
         X_test = scalar.transform(X_test)
 In [8]: # Applying PCA
         from sklearn.decomposition import PCA
         # put none to n_componenets to create explained variance vector
         # ( contain the percentage of variance explained by each of the principal compone
         pca = PCA(n components=2)
         X_train = pca.fit_transform(X_train)
         X_test = pca.transform(X_test)
         expained variance = pca.explained variance ratio
         print(expained_variance)
         [0.14717072 0.06634403]
 In [9]: #Fitting Logistic Regression to the training set
         from sklearn.linear model import LogisticRegression
         logmodel = LogisticRegression(random_state = 0)
         logmodel.fit(X_train, y_train)
Out[9]: LogisticRegression(random state=0)
In [10]: #Prdicting the test set results
         y_pred = logmodel.predict(X_test)
In [11]: from sklearn.metrics import roc curve
```

false_positive, true_positive, threshold1 = roc_curve(y_test, y_pred)

```
In [12]: # Plotting
         # Confusion Matrix
         from sklearn.metrics import accuracy score, classification report, confusion matr
         class_labels = df_new['1.000000000000000e+00'].unique()
         cm = confusion_matrix(y_test, y_pred, labels=class_labels)
         disp = ConfusionMatrixDisplay(confusion matrix=cm, display labels=class labels)
         disp.plot()
         # ROC Plot
         plt.subplots(1, figsize=(5,5))
         plt.title('Receiver Operating Characteristic - Logistic regression')
         plt.plot(false positive, true positive)
         plt.plot([0, 1], ls="--")
         plt.plot([0, 0], [1, 0], c=".7"), plt.plot([1, 1], c=".7")
         plt.ylabel('True Positive')
         plt.xlabel('False Positive')
         plt.show()
         # Calculating model accuarcy
         accuracy=accuracy_score(y_test,y_pred)
         print("Accuracy Obtained= ",accuracy)
```







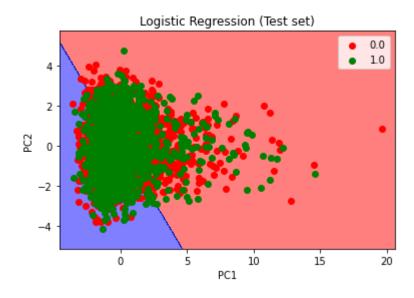
Accuracy Obtained= 0.54

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In [13]: #Visualizing the train set results from matplotlib.colors import ListedColormap X_set, y_set = X_train, y_train X1, X2 = np.meshgrid(np.arange(start = X_set[:,0].min() - 1, stop = X_set[:, 0].min() - 1 np.arange(start = X_set[:,1].min() - 1, stop = X_set[:, 1].min() - 1 plt.contourf(X1, X2, logmodel.predict(np.array([X1.ravel(), X2.ravel()]).T).resh alpha = 0.5, cmap = ListedColormap(('red') plt.xlim(X1.min(), X1.max()) plt.ylim(X2.min(), X2.max()) for i, j in enumerate(np.unique(y_set)): plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1], c = ListedColormap(('red', 'green', 'blue'))(i), label = j) plt.title('Logistic Regression (Test set)') plt.xlabel('PC1') plt.ylabel('PC2') plt.legend() plt.show() plt.show()

c argument looks like a single numeric RGB or RGBA sequence, which should be avoided as value-mapping will have precedence in case its length matches with * x* & *y*. Please use the *color* keyword-argument or provide a 2-D array with a single row if you intend to specify the same RGB or RGBA value for all point s.

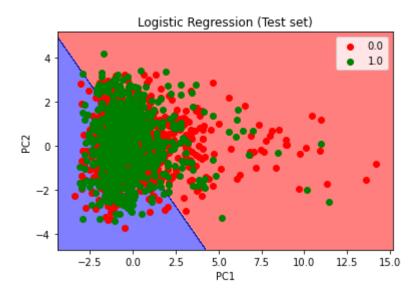
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In [14]: #Visualizing the test set results from matplotlib.colors import ListedColormap X_set, y_set = X_test, y_test X1, X2 = np.meshgrid(np.arange(start = X_set[:,0].min() - 1, stop = X_set[:, 0].min() - 1 np.arange(start = X_set[:,1].min() - 1, stop = X_set[:, 1].min() - 1 plt.contourf(X1, X2, logmodel.predict(np.array([X1.ravel(), X2.ravel()]).T).resha alpha = 0.5, cmap = ListedColormap(('red') plt.xlim(X1.min(), X1.max()) plt.ylim(X2.min(), X2.max()) for i, j in enumerate(np.unique(y_set)): plt.scatter(X_set[y_set == j, 0], X_set[y_set == j, 1], c = ListedColormap(('red', 'green', 'blue'))(i), label = j) plt.title('Logistic Regression (Test set)') plt.xlabel('PC1') plt.ylabel('PC2') plt.legend() plt.show()

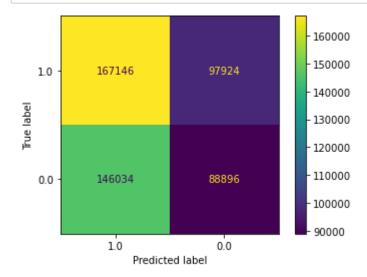
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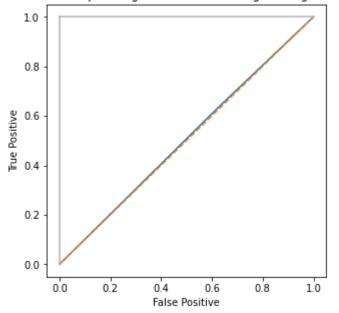
```
In [ ]:
In [15]: df_test=df[5499999:]
         # separating dependent and independent feature
         X_final_test= df_test.drop("1.00000000000000000e+00",axis=1)
         y final test = df test["1.00000000000000000e+00"]
In [16]: #Feature Normalization
         # standardization-or-mean-removal-and-variance-scaling
         X_final_test = scalar.transform(X_final_test)
         #Adding x0 = 1 to each instance for the bias term
         X_final_test = np.concatenate((np.ones((X_final_test.shape[0],1)),X_final_test),
In [17]: # Applying PCA
         from sklearn.decomposition import PCA
         # put none to n_componenets to create explained variance vector
         # ( contain the percentage of variance explained by each of the principal compone
         pca = PCA(n_components=2)
         X_final_test = pca.fit_transform(X_final_test)
         expained_variance = pca.explained_variance_ratio_
         print(expained_variance)
         [0.15009816 0.06642669]
In [18]: #Prdicting the test set results
         y_final_pred = logmodel.predict(X_final_test)
In [19]: false_positive_test, true_positive_test, threshold1 = roc_curve(y_final_test, y_f
```

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In [20]: # Plotting
         # Confusion Matrix
         class labels = df test['1.000000000000000e+00'].unique()
         cm_new = confusion_matrix(y_final_test,y_final_pred, labels=class_labels)
         disp = ConfusionMatrixDisplay(confusion_matrix=cm_new, display_labels=class_label
         disp.plot()
         # ROC Plot
         plt.subplots(1, figsize=(5,5))
         plt.title('Receiver Operating Characteristic - Logistic regression')
         plt.plot(false_positive_test, true_positive_test)
         plt.plot([0, 1], ls="--")
         plt.plot([0, 0], [1, 0], c=".7"), plt.plot([1, 1], c=".7")
         plt.ylabel('True Positive')
         plt.xlabel('False Positive')
         plt.show()
         # Calculating model accuarcy
         accuracy=accuracy_score(y_final_test,y_final_pred)
         print("Accuracy Obtained= ",accuracy)
```



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Accuracy Obtained= 0.512084

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In [21]: #Visualizing the fina test set results from matplotlib.colors import ListedColormap X_final_set, y_final_set = X_final_test, y_final_test X1 new, X2 new = np.meshgrid(np.arange(start = X final set[:,0].min() - 1, stop = np.arange(start = X_final_set[:,1].min() - 1, stop = X_final plt.contourf(X1_new, X2_new, logmodel.predict(np.array([X1_new.ravel(), X2_new.ra alpha = 0.5, cmap = ListedColormap(('red') plt.xlim(X1 new.min(), X1 new.max()) plt.ylim(X2 new.min(), X2 new.max()) for i, j in enumerate(np.unique(y_final_set)): plt.scatter(X_final_set[y_final_set == j, 0], X_final_set[y_final_set == j, 1 c= ListedColormap(('red', 'green', 'blue'))(i), label = j) plt.title('Logistic Regression (Final Test set)') plt.xlabel('PC1') plt.ylabel('PC2') plt.legend() plt.show()

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