```
import numpy as np
In [17]:
        import pandas as pd
        import matplotlib.pyplot as plt
        from sklearn.linear_model import LogisticRegression
        from sklearn.metrics import confusion_matrix, precision_score, recall_score, f1_sc
        # The data file is being read in this cell
In [3]:
        df = pd.read_csv("Logistic-Regression-data-2-class-v0.csv")
        print(df.head())
In [4]:
                 x1
                           x2 yclass
        0 -12.304702
                      3.499240
        1 -21.302900 17.983794
        2 -6.320254 29.639092
                                    0
            2.259775 26.227155
                                    0
        4 -14.777150 19.536615
In [5]: X = df[['x1', 'x2']] # Features 'x1' and 'x2'
        y = df['yclass'] # Target variable 'yclass'
        # 1. Create a scatter plot to visualize the data
        plt.xlabel('x1')
        plt.ylabel('x2')
        plt.legend()
        plt.show()
                                                                        Class 0
                                                                        Class 1
             40
             20
         Ø
              0
            -20
            -40
                                       -20
                                                                  20
                          -40
                                                      0
                                               x1
```

Most of the points with same class are clustered together. That is points with class 1 are in the lower half plane whereas points with class 0 are in the upper half of the plane. The plot seems to be divided about a straight line.

```
logist_regr = LogisticRegression()
In [6]:
        logist_regr.fit(X, y)
Out[6]:
        ▼ LogisticRegression
        LogisticRegression()
In [7]:
        pred = logist_regr.predict(X)
        logist_regr.score(df[['x1','x2']], df['yclass'])
        0.9763888888888889
Out[7]:
        plt.scatter(X[y == 0]['x1'], X[y == 0]['x2'], c='blue', label='Class 0 (True)')
In [8]:
        plt.scatter(X[y == 1]['x1'], X[y == 1]['x2'], c='red', label='Class 1 (True)')
        plt.scatter(X[pred != y]['x1'], X[pred != y]['x2'], c='green', label='Misclassified
        plt.xlabel('x1')
        plt.ylabel('x2')
        plt.legend()
        plt.show()
                                                                         Class 0 (True)
                                                                         Class 1 (True)
              40
                                                                         Misclassified
             20
         Š
               0
            -20
            -40
                            -40
                                           -20
                                                           0
                                                                         20
```

Majority of the wrongly classified points are at the border where the two cluster meet. Also another discrepancy arises in the classification of the blue point which was present amongst the cluster of red points. The model ends up classifying it as a red point as it is present deep in the cluster of the red points.

x1

The score of the model is 0.976388888888888, which is a good and accurate score

```
In [10]: confusion_matrix_result = confusion_matrix(y, pred)
    precision = precision_score(y, pred)
    recall = recall_score(y, pred)
    f1 = f1_score(y, pred)

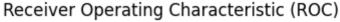
TP = confusion_matrix_result[1, 1]
```

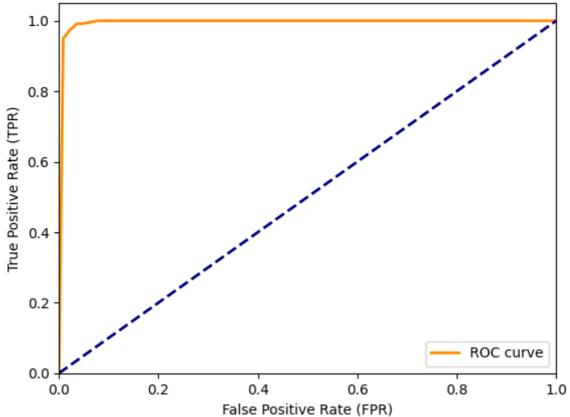
```
FP = confusion_matrix_result[0, 1]
         TN = confusion_matrix_result[0, 0]
         FN = confusion_matrix_result[1, 0]
         TPR = TP / (TP + FN)
         FPR = FP / (FP + TN)
         print("Confusion Matrix:")
         print(confusion_matrix_result)
         print("Precision:", precision)
         print("Recall:", recall)
         print("F1-score:", f1)
         print("True Positive Rate (TPR):", TPR)
         print("False Positive Rate (FPR):", FPR)
         Confusion Matrix:
         [[352 8]
          [ 9 351]]
         Precision: 0.9777158774373259
         Recall: 0.975
         F1-score: 0.9763560500695411
         True Positive Rate (TPR): 0.975
         False Positive Rate (FPR): 0.0222222222222223
In [11]: probabilities = logist_regr.predict_proba(X)
In [14]: | tpr_values = []
         fpr_values = []
         for i in range(0,11):
             # Classify observations based on the threshold
             threshold = i/10
             predicted_labels = (probabilities[:, 1] >= threshold).astype(int)
             # Calculate the confusion matrix
             confusion_matrix_result = confusion_matrix(y, predicted_labels)
             # Calculate True Positive Rate (TPR) and False Positive Rate (FPR)
             TP = confusion_matrix_result[1, 1]
             FP = confusion matrix result[0, 1]
             TN = confusion_matrix_result[0, 0]
             FN = confusion_matrix_result[1, 0]
             TPR = TP / (TP + FN)
             FPR = FP / (FP + TN)
             tpr_values.append(TPR)
             fpr values.append(FPR)
             # Print the results for the current threshold
             print(f"Threshold: {threshold:.1f}")
             print("Confusion Matrix:")
             print(confusion matrix result)
             print("True Positive Rate (TPR):", TPR)
             print("False Positive Rate (FPR):", FPR)
             print()
```

```
Threshold: 0.0
Confusion Matrix:
[[ 0 360]
[ 0 360]]
True Positive Rate (TPR): 1.0
False Positive Rate (FPR): 1.0
Threshold: 0.1
Confusion Matrix:
[[332 28]
[ 0 360]]
True Positive Rate (TPR): 1.0
False Positive Rate (FPR): 0.07777777777778
Threshold: 0.2
Confusion Matrix:
[[343 17]
[ 3 357]]
True Positive Rate (TPR): 0.991666666666667
False Positive Rate (FPR): 0.047222222222222
Threshold: 0.3
Confusion Matrix:
[[347 13]
[ 3 357]]
True Positive Rate (TPR): 0.991666666666667
False Positive Rate (FPR): 0.03611111111111111
Threshold: 0.4
Confusion Matrix:
[[350 10]
[ 7 353]]
True Positive Rate (TPR): 0.9805555555555555
False Positive Rate (FPR): 0.027777777777776
Threshold: 0.5
Confusion Matrix:
[[352 8]
[ 9 351]]
True Positive Rate (TPR): 0.975
False Positive Rate (FPR): 0.0222222222222223
Threshold: 0.6
Confusion Matrix:
[[353 7]
 [ 11 349]]
True Positive Rate (TPR): 0.96944444444444444
False Positive Rate (FPR): 0.01944444444444445
Threshold: 0.7
Confusion Matrix:
[[354 6]
[ 13 347]]
True Positive Rate (TPR): 0.9638888888888888
False Positive Rate (FPR): 0.016666666666666666
Threshold: 0.8
Confusion Matrix:
[[357
      3]
[ 18 342]]
True Positive Rate (TPR): 0.95
```

Threshold: 0.9

```
In [15]: # Plot the ROC curve
    plt.figure()
    plt.plot(fpr_values, tpr_values, color='darkorange', lw=2, label='ROC curve')
    plt.plot([0, 1], [0, 1], color='navy', lw=2, linestyle='--')
    plt.xlim([0.0, 1.0])
    plt.ylim([0.0, 1.05])
    plt.xlabel('False Positive Rate (FPR)')
    plt.ylabel('True Positive Rate (TPR)')
    plt.title('Receiver Operating Characteristic (ROC)')
    plt.legend(loc='lower right')
    plt.show()
```





From the ROC curve, we can see that the area under the curve that is AUC is comparable to the area of the square shown in the figure. Therefore, it is a very good model with accurate predictions.

```
In [18]: roc_auc = roc_auc_score(y, pred)

# Print the AUC value
print("AUC:", roc_auc)
```

AUC: 0.97638888888889