### **SVM Analysys**

#### Import the data

In [2]: # Import data import pandas as pd df = pd.read\_csv("diabetes\_binary\_5050split\_health\_indicators\_BRFSS2021.csv")

0.0

0.0

0.0

```
df.head()
Out[2]:
           Diabetes_binary HighBP HighChol CholCheck BMI Smoker Stroke HeartDiseaseorAttack PhysActivity Fruits ... AnyHealthcare NoDocbcCost GenHlth MentHlth PhysHlth DiffWalk Sex Age Education Income
        0
                    0.0
                                    0.0
                                               1 33.0
                                                          0.0
                                                                0.0
                                                                                  0.0
                                                                                              1 1 ...
                                                                                                                              0.0
                                                                                                                                     2.0
                                                                                                                                             15.0
                                                                                                                                                      0.0
                                                                                                                                                                                6.0
                                                                                                                                                                                       9.0
                                                                                                                                                              1.0
                                                                                                                                                                  1 7
                    0.0
                                    1.0
                                               1 27.0
                                                          1.0 0.0
                                                                                  0.0
                                                                                              1 0 ...
                                                                                                                              0.0
                                                                                                                                     2.0
                                                                                                                                              1.0
                                                                                                                                                      2.0
                                                                                                                                                              0.0 1 7
                                                                                                                                                                                6.0
                                                                                                                                                                                       6.0
```

1

1

0.0

0.0

0.0

3.0

3.0

2.0

30.0

0.0

0.0

0.0 1 13

0.0 0 11

0 5

0.0

4.0

5.0

5.0

3.0

7.0

3.0

0.0

0.0

0.0

0 0 ...

1 1 ...

1 ...

5 rows × 22 columns

0.0

0.0

0.0

0

0

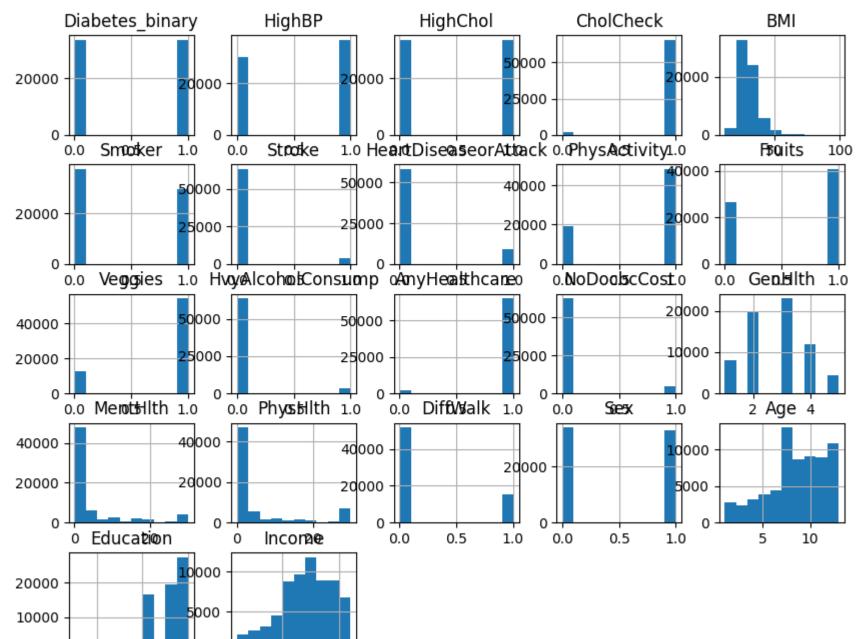
1.0

0.0

0.0

2

In [3]: # Import the libraries import numpy as np from sklearn.model\_selection import train\_test\_split, GridSearchCV from sklearn.svm import SVC from sklearn.metrics import accuracy\_score, classification\_report, roc\_curve, auc import matplotlib.pyplot as plt # Visualize the variables df.hist(figsize=(10, 8)) plt.show()



1.0 0.0

1.0 0.0

0.0

0.0

1 26.0

1 19.0

1 37.0

In [4]: X = df.drop('Diabetes\_binary', axis=1) y = df['Diabetes\_binary']

In [5]: # Split the data into training and testing sets X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.2, random\_state=42)

### Linear Kernel

Out[7]: ▼

In [6]: # Create an SVM model svm\_linear = SVC(C=0.01, kernel='linear') # Linear kernel

In [7]: # Train the model

svm\_linear.fit(X\_train, y\_train)

SVC

SVC(C=0.01, kernel='linear')

In [8]: # Set the ROC Curve results y\_test\_score = svm\_linear.decision\_function(X\_test)

> fpr\_l, tpr\_l, \_ = roc\_curve(y\_test, y\_test\_score)  $roc_auc_l = auc(fpr_l, tpr_l)$

In [14]: # Performs the Classification Report y\_pred\_l = svm\_linear.predict(X\_test)

> report\_l = classification\_report(y\_test, y\_pred\_l) print(report\_l) recall f1-score support precision

6614 0.75 0.69 0.72 6814 0.73 13428 accuracy 0.74 0.73 0.73 13428 macro avg weighted avg 0.73 13428

### Non-linear Kernel

In [10]: # Create an SVM model svm\_rbf = SVC(C=100, gamma = 0.001, kernel='rbf') # Non-linear kernel

In [11]: # Train the model svm\_rbf.fit(X\_train, y\_train)

Out[11]: SVC SVC(C=100, gamma=0.001)

In [12]: # Set the ROC Curve results

y\_test\_score = svm\_rbf.decision\_function(X\_test) fpr\_n, tpr\_n, \_ = roc\_curve(y\_test, y\_test\_score)  $roc_auc_n = auc(fpr_n, tpr_n)$ 

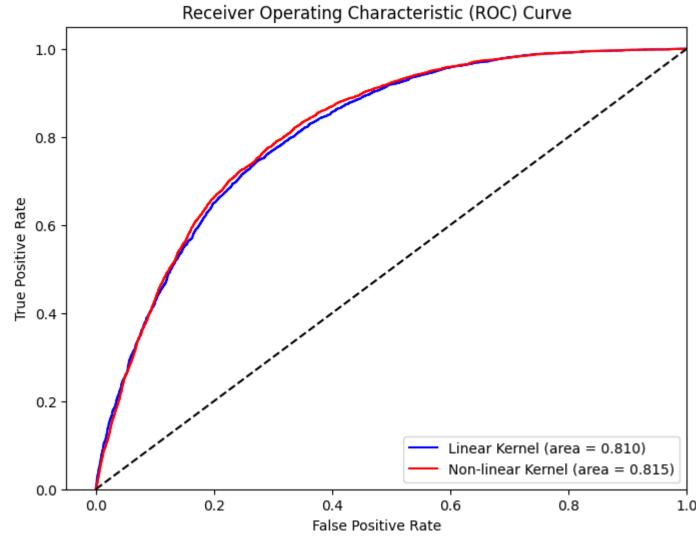
In [15]: # Performs the Classification Report y\_pred\_n = svm\_rbf.predict(X\_test)

report\_n = classification\_report(y\_test, y\_pred\_n) print(report\_n)

recall f1-score support precision 0.0 0.78 0.67 0.72 6614 6814 1.0 0.72 0.81 0.76 0.74 13428 accuracy macro avg 0.75 0.74 0.74 13428 weighted avg 0.75 0.74 0.74 13428

# **ROC Curves**

In [19]: # Plot both ROC curves on the same graph plt.figure(figsize=(8, 6)) # Linear Kernel ROC curve plt.plot(fpr\_1, tpr\_1, label='Linear Kernel (area = %0.3f)' % roc\_auc\_1, color='b') # Non-linear Kernel ROC curve plt.plot(fpr\_n, tpr\_n, label='Non-linear Kernel (area = %0.3f)' % roc\_auc\_n, color='r') # Add details to the plot plt.plot([0, 1], [0, 1], 'k--') plt.xlim([-0.05, 1.0]) plt.ylim([0.0, 1.05]) plt.xlabel('False Positive Rate') plt.ylabel('True Positive Rate') plt.title('Receiver Operating Characteristic (ROC) Curve') plt.legend(loc="lower right") plt.show()



# Grid Search - Linear Kernel

In [5]: # Select the optimal C parameter by cross-validation tuned\_parameters =  $[\{'C': [0.01, 0.1, 1, 10, 100]\}]$ clf = GridSearchCV(SVC(kernel='linear'), tuned\_parameters, cv=2, n\_jobs=-1, scoring='f1') clf.fit(X\_train, y\_train)

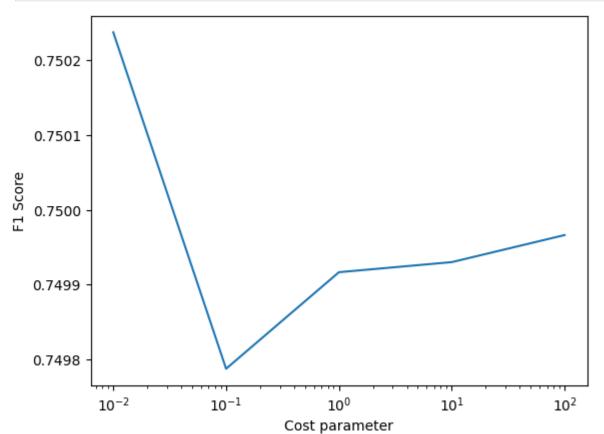
cv\_results = pd.DataFrame(clf.cv\_results\_) # Plots the results

plt.plot(cv\_results['param\_C'], cv\_results['mean\_test\_score']) plt.xscale(value='log') plt.ylabel('F1 Score')

plt.show()

plt.xlabel('Cost parameter')

# Prints the best parameters print('Best parameters : ' + str(clf.best\_params\_))



Best parameters : {'C': 0.01}

## Grid Search - Non-linear Kernel

In [6]: # Select the optimal C and gamma parameters by cross-validation tuned\_parameters =  $[\{'C': [0.01, 0.1, 1, 10, 100],$ 'gamma': [0.001, 0.01, 0.1, 1, 10]}] clf = GridSearchCV(SVC(kernel='rbf'), tuned\_parameters, cv=2, n\_jobs=-1, scoring='f1') clf.fit(X\_train, y\_train)

# Prints the best parameters
print('Best parameters : ' + str(clf.best\_params\_))

Best parameters : {'C': 100, 'gamma': 0.001}