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In [ ]: # Implement K-Nearest Neighbors algorithm on diabetes.csv dataset. Compute confusion
         # matrix, accuracy, error rate, precision and recall on the given dataset.
         # Dataset link : https://www.kaggle.com/datasets/abdallamahgoub/diabetes
In [4]: import pandas as pd
         import numpy as np
         from sklearn.model_selection import train_test_split
         from sklearn.preprocessing import StandardScaler
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.metrics import confusion_matrix, accuracy_score, precision_score, recall_score, f1_score
In [5]: data=pd.read_csv("diabetes.csv")
         data
              Pregnancies Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction Age Outcome
           0
                        6
                               148
                                               72
                                                                      0 33.6
                                                                                                 0.627
                                                                                                         50
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                                85
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                                                             29
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                                                                                                 0.351
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                        8
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                               183
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                                89
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           4
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                                                                    168 43.1
                                                                                                 2.288
                                                                                                         33
         763
                       10
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                                                                                                 0.171
                                                                                                         63
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                               122
                                                             27
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         764
                                               70
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                                                                                                 0.340
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                        5
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                                                                                                 0.245
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                                                                                                         47
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         767
                                93
                                               70
                                                             31
                                                                      0 30.4
                                                                                                 0.315
                                                                                                         23
        768 rows × 9 columns
In [6]: X = data.drop("Outcome", axis=1) # Features
         y = data["Outcome"] # Target variable
In [7]: X
Out[7]:
              Pregnancies
                          Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction Age
           0
                        6
                               148
                                               72
                                                             35
                                                                      0 33.6
                                                                                                 0.627
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                                                                     94 28.1
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                                                                    168 43.1
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         763
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                                                                                                         23
        768 rows × 8 columns
In [8]: # 2. Split the dataset into training and test sets
         X_{\text{train}}, X_{\text{test}}, y_{\text{train}}, y_{\text{test}} = train_test_split(X, y, test_size=0.2, random_state=42)
In [9]: # 3. Normalize the data
```

scaler = StandardScaler()

In [10]: X train

X\_train = scaler.fit\_transform(X\_train)
X test = scaler.transform(X test)

```
Out[10]: array([[-0.52639686, -1.15139792, -3.75268255, ..., -4.13525578,
                                -0.49073479, -1.03594038],
                              [ 1.58804586, -0.27664283, 0.68034485, ..., -0.48916881,
                                 2.41502991, 1.48710085],
                              \hbox{$[\,\hbox{-0.82846011,}}\quad \hbox{0.56687102,} \ \hbox{-1.2658623 ,} \ \dots, \ \hbox{-0.42452187,}
                                 0.54916055, -0.94893896],
                              [ 1.8901091 , -0.62029661, 0.89659009, ..., 1.76054443,
                              1.981245 , 0.44308379],
[-1.13052335, 0.62935353, -3.75268255, ..., 1.34680407,
                               -0.78487662, -0.33992901],
                              \hbox{[-1.13052335, 0.12949347, 1.43720319, ..., -1.22614383,}
                                -0.61552223, -1.03594038]])
In [12]: # 4. Implement K-Nearest Neighbors (KNN)
                 k = 3 # Choose the number of neighbors (k) based on your needs
                 knn = KNeighborsClassifier(n neighbors=k)
                 knn.fit(X train, y train)
Out[12]: KNeighborsClassifier(n_neighbors=3)
In [13]: # 5. Predict and Evaluate
                 y pred = knn.predict(X test)
                y_pred
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              reduction functions (e.g. `skew`, `kurtosis`), the default behavior of `mode` typically preserves the axis it ac ts along. In SciPy 1.11.0, this behavior will change: the default value of `keepdims` will become False, the `ax
              is` over which the statistic is taken will be eliminated, and the value None will no longer be accepted. Set `ke
              epdims` to True or False to avoid this warning.
                 mode, = stats.mode( y[neigh ind, k], axis=1)
Out[13]: array([0, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 1, 0,
                              0,\ 0,\ 1,\ 1,\ 0,\ 0,\ 0,\ 1,\ 1,\ 1,\ 1,\ 1,\ 1,\ 1,\ 0,\ 0,\ 1,\ 0,\ 0,\ 1,\ 0,
                              0,\ 0,\ 0,\ 0,\ 0,\ 1,\ 0,\ 1,\ 1,\ 0,\ 0,\ 0,\ 0,\ 0,\ 1,\ 0,\ 0,\ 1,\ 1,\ 0,\ 1,
                              0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0,
                             0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 1,
                             0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1,
                             0, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0],
                            dtype=int64)
In [14]: # Compute the confusion matrix
                 conf matrix = confusion_matrix(y_test, y_pred)
                 # Calculate accuracy, error rate, precision, and recall
                 accuracy = accuracy_score(y_test, y_pred)
                 error rate = 1 - accuracy
                 precision = precision score(y test, y pred)
                 recall = recall score(y test, y pred)
                 print("Confusion Matrix:")
                 print(conf_matrix)
                 print("Accuracy:", accuracy)
                 print("Error Rate:", error_rate)
                 print("Precision:", precision)
                 print("Recall:", recall)
              Confusion Matrix:
               [[81 18]
                [27 28]]
              Accuracy: 0.7077922077922078
              Error Rate: 0.29220779220779225
              Precision: 0.6086956521739131
              Recall: 0.509090909090909
  In [ ]: # Accuracy: This measures the overall correctness of the classifier's predictions. In this case, the model is a
                 # Error Rate: The error rate is the complement of accuracy (1 - accuracy), representing the proportion of incor
                 # Precision: Precision measures the ratio of true positive predictions to the total number of positive prediction
                 # Recall: Recall measures the ratio of true positive predictions to the total number of actual positive instance
```