```
import matplotlib.pyplot as plt
          import seaborn as sns
          from sklearn.datasets import load_iris
          from sklearn.model_selection import train_test_split
          from sklearn.preprocessing import StandardScaler
          from sklearn.neighbors import KNeighborsClassifier
          from sklearn.metrics import classification_report, confusion_matrix
          # Load Iris dataset from sklearn
          iris = load_iris()
In [2]:
          # Create DataFrame from the dataset
          df = pd.DataFrame(data=iris.data, columns=iris.feature_names)
          df['target'] = iris.target
          # Map target names to target labels
          target_names = iris.target_names
          df['species'] = df['target'].map(lambda x: target_names[x])
          # Display first few rows of the DataFrame
          df.head()
Out[2]:
            sepal length (cm) sepal width (cm) petal length (cm) petal width (cm) target species
                       5.1
                                      3.5
                                                    1.4
          0
                                                                  0.2
                                                                              setosa
         1
                       4.9
                                      3.0
                                                    1.4
                                                                  0.2
                                                                             setosa
          2
                       4.7
                                      3.2
                                                    1.3
                                                                  0.2
                                                                          0
                                                                              setosa
                                                    1.5
         3
                       4.6
                                      3.1
                                                                  0.2
                                                                          0
                                                                              setosa
                       5.0
                                      3.6
                                                    1.4
                                                                   0.2
                                                                              setosa
In [3]:
          # Pairplot to visualize relationships between features
          sns.pairplot(df, hue='species')
          plt.show()
           sepal length (cm)
            4.5
            4.0
         sepal width (cm)
            2.0
                                                                                                                                     species
                                                                                                                                      setosa
                                                                                                                                      versicolor
                                                                                                                                      virginica
             1 .
            2.5
         2.0 betal width (cm)
            0.5
            0.0
            2.0
           1.5
          target
10
            0.5
            0.0
                                                                                                                0.5
                                                                                                                    1.0
                                                                                                                         1.5 2.0
                    sepal length (cm)
                                           sepal width (cm)
                                                                  petal length (cm)
                                                                                         petal width (cm)
                                                                                                                    target
In [4]:
          # Split data into features (X) and target (y)
          X = df.drop(['target', 'species'], axis=1)
          y = df['target']
          # Split data into train and test sets
          X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
In [5]:
          # Standardize features
          scaler = StandardScaler()
          X_train_scaled = scaler.fit_transform(X_train)
          X_test_scaled = scaler.transform(X_test)
In [6]:
          # Create K-Nearest Neighbors classifier
          knn = KNeighborsClassifier(n_neighbors=3)
          knn.fit(X_train_scaled, y_train)
          KNeighborsClassifier(n_neighbors=3)
Out[6]:
In [10]:
          # Predictions on the test set
          y_pred = knn.predict(X_test_scaled)
          # Classification report and confusion matrix
          print("Classification Report:\n", classification_report(y_test, y_pred))
          print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))
          Classification Report:
                         precision
                                       recall f1-score
                                                           support
                                        1.00
                                                   1.00
                     0
                              1.00
                                                                10
                                                                9
                     1
                              1.00
                                        1.00
                                                   1.00
                              1.00
                                        1.00
                                                   1.00
                                                                11
                                                   1.00
                                                                30
              accuracy
             macro avg
                              1.00
                                        1.00
                                                   1.00
                                                                30
                              1.00
                                        1.00
                                                   1.00
                                                                30
          weighted avg
          Confusion Matrix:
          [[10 0 0]
           [ 0 9 0]
           [ 0 0 11]]
In [12]:
          cm=confusion_matrix(y_test,y_pred)
          sns.heatmap(cm, annot=True)
          plt.xlabel('Actual Label')
          plt.ylabel('Predicate Label')
          plt.title('Confusion Matrix for the classification')
         Text(0.5, 1.0, 'Confusion Matrix for the classification')
Out[12]:
                 Confusion Matrix for the classification
                                                       - 10
                   10
                                            0
```

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2

Actual Label

In [1]:

import pandas as pd
import numpy as np