Step1: Import libraries. (Use are free to use other libraries for plotting/visualisations) ● Numpy ● Matplotlib.pyplot/Seaborne ● Train test split ● Confusion matrix

```
In [54]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
```

Step2: Generate synthetic data for two classes. Use above given parameters for samples distribution or of your choice. Also seed(0) for reproducibility.

```
In [55]: # Parameters
    total_no_of_samples = 500
    learning_rate = 0.1
    bias = 1
    max_epochs = 1000

In [56]: # Distribution 1
    mul = np.array([-2, -2])
    sigmal = np.array([[0.9, -0.0255], [-0.0255, 0.9]])
    distribution1 = np.random.multivariate_normal(mul, sigmal, total_no_of_samples // 2)

# Distribution 2
    mu2 = np.array([[0.5, 5])
    sigma2 = np.array([[0.5, 0], [0, 0.3]])
    distribution2 = np.random.multivariate_normal(mu2, sigma2, total_no_of_samples // 2)
```

Step3: Combine both the distribution (classes) and their labels to form a dataset. Use np.vstack(), np.hstack().

```
In [57]: # Combine the two distributions
data = np.vstack((distribution1, distribution2))
labels = np.hstack((np.zeros(total_no_of_samples // 2), np.ones(total_no_of_samples // 2)))
```

Step4: Include bias term by adding a column of ones to input feature matrix.

```
#adding bias term to the data
In [58]:
         data with bias = np.column stack([data,bias * np.ones(total no of samples)])
In [59]:
         data with bias
         array([[-1.53318899, -1.38367901, 1.
                                                      1,
Out[59]:
                [-1.33216563, -0.94837054, 1.
                                                      1,
                [-0.45539938, -1.7907655, 1.
                [ 3.95850867, 5.3306986 , 1.
                                                      1,
                [ 4.71294149, 5.61041534, 1.
                                                      1,
                [ 5.66140645, 4.54749005, 1.
                                                      11)
```

Step5: Split the dataset into train and test.

```
In [60]: # Split the data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(data_with_bias, labels, test_size=0.2, random_state=42)
```

Step6: Write a function to train the perceptron that will take data, labels, learning rate and max_epochs as parameters.

Step7: Define a step activation function where it will return 1 if value >= 0 else 0.

```
In [61]: def perceptron train(data, labels, learning rate, max epochs):
             # Initialize weights and bias
             num features = data.shape[1]
             weights = np.zeros(num features)
             bias = 0.0
             for epoch in range(max epochs):
                 errors = 0
                 # Iterate over each data point
                 for i in range(len(data)):
                     # Calculate the predicted output
                     prediction = predict(data[i], weights, bias)
                     # Update weights and bias based on the error
                     update = learning rate * (labels[i] - prediction)
                     weights += update * data[i]
                     bias += update
                     # Count errors for this epoch
                     errors += int(update != 0.0)
                 # Check if all data points are classified correctly
                 if errors == 0:
                     print(f"Converged in {epoch + 1} epochs.")
                     break
             return weights, bias
         def predict(data point, weights, bias):
             # Activation function (in this case, a simple step function)
             activation = np.dot(data point, weights) + bias
             return 1 if activation >= 0 else 0
```

Step8: train the perceptron on training set.

```
In [62]: # Call the perceptron_train function to train the perceptron
trained_weights, trained_bias = perceptron_train(X_train, y_train, learning_rate=0.01, max_epochs=1000)
```

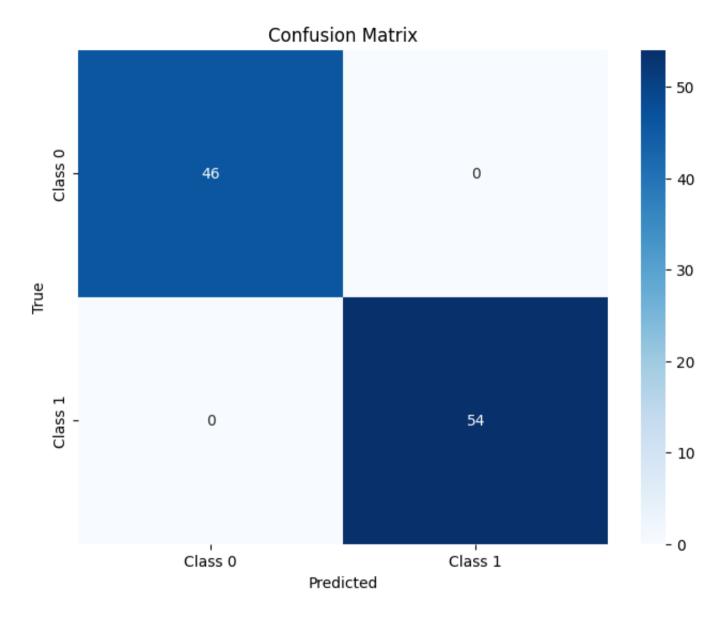
Converged in 2 epochs.

Step9: Make predictions using trained perceptron on test set. Tune the hyperparameters like learning rate, test size and find the optimal accurate perceptron model.

Step11: Plot the confusion matrix.

```
In [31]: num test samples = len(X_test)
                                  predictions = []
                                   for i in range(num test samples):
                                                  prediction = predict(X test[i], trained weights, trained bias)
                                                  predictions.append(prediction)
                                  # Calculate accuracy
                                  correct predictions = sum(prediction == y test[i] for i, prediction in enumerate(predictions))
                                  accuracy = correct predictions / num test samples
                                  print(f"Accuracy on the test set: {accuracy}")
                                   # Create confusion matrix
                                  cm = confusion matrix(y test, predictions)
                                   # Plot confusion matrix
                                  plt.figure(figsize=(8, 6))
                                  sns.heatmap(cm, annot=True, fmt="d", cmap="Blues", xticklabels=["Class 0", "Class 1"], yticklabels=["Class 0", "Class 0"], yticklabels=["C
                                   plt.title("Confusion Matrix")
                                  plt.xlabel("Predicted")
                                  plt.ylabel("True")
                                   plt.show()
```

Accuracy on the test set: 1.0



Step10: Plot the decision boundary between two classified class.

```
In [53]: weights = trained weights
         b = trained bias
         # Check if the weights are 1D (two-dimensional input space) or 2D (multi-dimensional input space)
         if len(weights) == 2:
             w1. w2 = weights
         else:
             # For a multi-dimensional perceptron, use the first two weights for plotting
             w1, w2 = weights[:2]
         # Generate x values for the decision boundary
         x decision boundary = np.linspace(min(data[:, 0]), max(data[:, 0]), 100)
         y decision boundary = (-trained weights[0] / w2) * x decision boundary - b / w2
         # Plot the decision boundary along with the scatter plot of test data
         plt.figure(figsize=(8, 6))
         # plt.scatter(X test[:, 0], X test[:, 1], c=y test, cmap='viridis', edgecolors='k', marker='o', s=100, label='Test
         # Assuming y test has values 0 and 1 for the two clusters, adjust labels accordingly
         plt.scatter(X test[y test == 0, 0], X test[y test == 0, 1], c='blue', edgecolors='k', marker='o', s=100, label='(
         plt.scatter(X test[y test == 1, 0], X test[y test == 1, 1], c='orange', edgecolors='k', marker='o', s=100, label=
         plt.plot(x decision boundary, y decision boundary, label='Decision Boundary', color='red', linewidth=2)
         plt.title('Decision Boundary and Test Data')
         plt.xlabel('Feature 1') # Corrected label for the x-axis
         plt.ylabel('Feature 2') # Corrected label for the y-axis
         plt.legend()
         plt.show()
```



