Aim

Objective:

Implement and evaluate a Single Layer Perceptron for binary classification on both linearly separable and non-linearly separable datasets.

Workflow:

Generate synthetic datasets (one linearly separable using make_classification and one non-linearly separable using make_circles), train the Perceptron on these datasets (as well as a real-world dataset like Breast Cancer), and assess its performance using accuracy and visualizations.

Algorithm

1. Data Preparation and Exploration:

- Generate a synthetic 2D dataset using make_classification for a linearly separable scenario.
- Explore the dataset using descriptive statistics, correlation matrices, distribution plots, and pair plots.

2. Perceptron Implementation:

- Define a custom Percept ron class with an initialization of weights and bias.
- Implement an activation function (step function), a fit method for training with weight updates using stochastic gradient descent and a learning rate decay, and a predict method.

3. Model Training and Evaluation on Synthetic Data:

- Split the synthetic data into training and testing sets.
- Train the Perceptron on the training data and evaluate its performance on the test set using accuracy.

4. Evaluation on Real-World and Non-Linearly Separable Data:

- Train and evaluate the Perceptron on the Breast Cancer dataset (after scaling) to observe performance on real-world data.
- Generate a non-linearly separable dataset using make_circles, train the Perceptron, and assess its accuracy.

5. Visualization:

 Visualize the data distributions, correlations, and decision boundaries (if applicable) to gain insights into the data and model behavior.

Algorithm Description

• Single Layer Perceptron:

The Perceptron is a simple linear classifier that updates its weights and bias based on the error between the predicted and actual outputs. It uses a step activation function that outputs a binary class label. During training, the algorithm adjusts its

parameters using a learning rate (which may decay over epochs) to minimize misclassifications.

Handling Linearly Separable vs. Non-Linearly Separable Data:

- For linearly separable data (e.g., generated via make_classification), the
 Perceptron converges to a solution that perfectly separates the classes, achieving high accuracy.
- For non-linearly separable data (e.g., generated via make_circles), the inherent limitation of a single linear decision boundary is highlighted, often resulting in lower performance.

Evaluation and Visualization:

The performance of the Perceptron is quantified using accuracy metrics on both synthetic and real-world datasets. Visualization techniques (scatter plots, pair plots, and correlation matrices) are employed to understand data distributions and the effectiveness of the classifier.

Results

1. Synthetic Linearly Separable Data (make_classification)

- **Accuracy:** ~92.5%
- The Perceptron effectively learned the decision boundary on the synthetic, linearly separable dataset, achieving high accuracy.

2. Real-World Data (Breast Cancer)

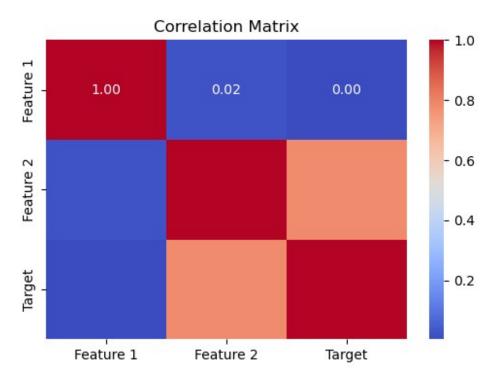
- Unscaled Data Accuracy: ~95.32%
- Scaled Data Accuracy: ~94.15%
- The Perceptron performed robustly on the Breast Cancer dataset. While the unscaled data yielded slightly higher accuracy, scaling the features still resulted in strong performance.

3. Non-Linearly Separable Data (Circles)

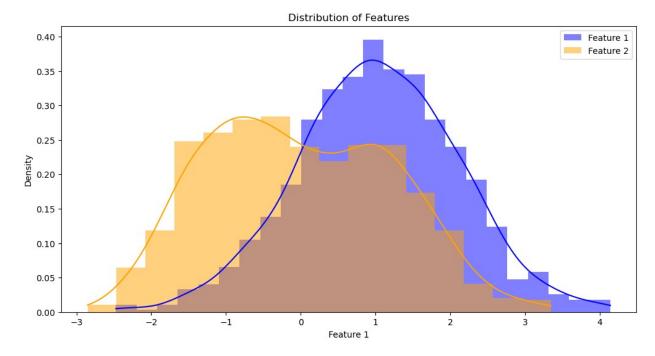
- **Accuracy:** ~70.0%
- The Perceptron struggled with the non-linearly separable circle dataset, highlighting its limitation with data that requires a non-linear decision boundary.

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
import warnings
warnings.filterwarnings('ignore')
```

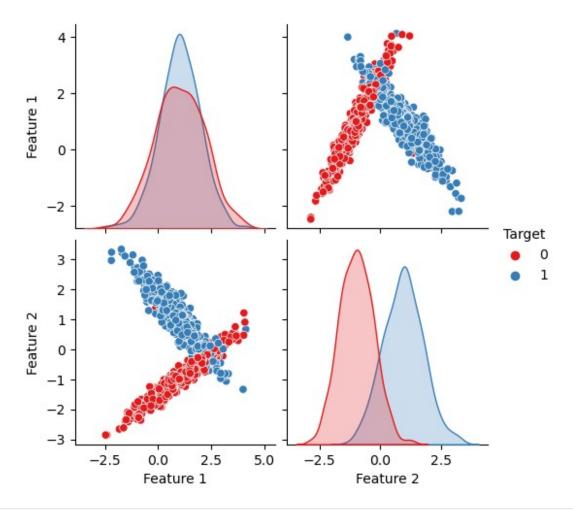
```
X, y = make classification(n samples=1000, n features=2,
n informative=2, n redundant=0, n clusters per class=1,
random_state=42)
df = pd.DataFrame(X, columns=["Feature 1", "Feature 2"])
df["Target"] = y
df
     Feature 1
                Feature 2
                           Target
0
      0.601034
                 1.535353
                                 1
1
      0.755945
                 -1.172352
                                 0
2
                                 0
      1.354479
                -0.948528
3
      3.103090
                 0.233485
                                 0
4
      0.753178
                 0.787514
                                 1
                 0.451639
995
      1.713939
                                 1
                                 0
996
      1.509473
                -0.794996
                                 1
997
      2.844315
                 0.211294
                                 1
998
     -0.025876
                 1.619258
                 0.756925
                                 0
999
      3.641478
[1000 \text{ rows } x \text{ 3 columns}]
df.describe()
         Feature 1
                       Feature 2
                                       Target
count
       1000.000000
                    1000.000000
                                  1000.000000
                       -0.012693
mean
          1.025840
                                     0.499000
std
          1.071457
                       1.225378
                                     0.500249
         -2.472718
min
                       -2.850971
                                     0.000000
25%
          0.307209
                       -0.984268
                                     0.000000
          1.023750
                       -0.102945
50%
                                     0.000000
75%
          1.724713
                        0.973550
                                     1.000000
          4.138715
                        3.342864
                                     1.000000
max
plt.figure(figsize=(6, 4))
sns.heatmap(df.corr(), annot=True, cmap='coolwarm', fmt='.2f')
plt.title('Correlation Matrix')
plt.show()
```



```
plt.figure(figsize=(12, 6))
sns.histplot(df["Feature 1"], kde=True, color="blue", label="Feature
1", stat="density", linewidth=0)
sns.histplot(df["Feature 2"], kde=True, color="orange", label="Feature
2", stat="density", linewidth=0)
plt.title('Distribution of Features')
plt.legend()
plt.show()
```



sns.pairplot(df, hue="Target", palette="Set1")
plt.show()



```
class Perceptron:
    def __init__(self, learning_rate:float = 0.00001, epoch:int =
100_000, decay_rate: float = 0.975):
        self.learning_rate = learning_rate
        self.epoch = epoch
        self.decay_rate = decay_rate
        self.weight = None
        self.bias = None

def activation_function(self, x):
        return 1 if x >=0 else 0

def fit(self, X:np.ndarray, y:np.ndarray):
        self.weight = np.random.randn(X.shape[1]) * 0.01
        self.bias = 0
        prev_weights = np.copy(self.weight) # To monitor convergence
        for epoch in range(1, self.epoch+1):
```

```
indices = np.random.permutation(len(X))
            X shuffled = X[indices]
            y shuffled = y[indices]
            total error = 0
            for i in range(len(X shuffled)):
                # Forward Pass
                weighted_sum = np.dot(X_shuffled[i], self.weight) +
self.bias
                predicted = self.activation function(weighted sum)
                # Calculate error
                error = y_shuffled[i] - predicted
                total error += abs(error)
                # Update weights
                self.weight += self.learning_rate * error *
X shuffled[i]
                self.bias += self.learning_rate * error
            self.learning rate *= self.decay_rate
            if epoch % 1000 == 0:
                print(f"Epoch {epoch}/{self.epoch}, Total error:
{total error}", end='\r')
            if np.all(np.abs(self.weight - prev weights) < le-100):</pre>
                print(f"Convergence reached at epoch {epoch}.")
                break
            prev weights = np.copy(self.weight)
    def predict(self, X:np.ndarray):
        results = []
        for i in range(len(X)):
            weighted sum = np.dot(X[i], self.weight) + self.bias
            pred = self.activation function(weighted sum)
            results.append(pred)
        return np.array(results)
X_train, X_test, y_train, y_test = train_test_split(
    Χ,
    у,
    test size=0.25,
```

```
random state=42
model = Perceptron()
model.fit(X train, y train)
Convergence reached at epoch 8607.5
pred i = model.predict(X test)
pred i
array([1, 0, 0, 0, 0, 1, 0, 1, 0, 0, 1, 1, 0, 0, 1, 0, 0, 1, 1, 1, 0,
0,
       0, 0, 0, 0, 0, 1, 1, 0, 0, 1, 0, 1, 0, 0, 1, 0, 0, 0, 0, 1, 1,
0,
       0, 0, 0, 0, 1, 0, 1, 1, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 1, 1, 1,
1,
       0, 0, 1, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1,
1,
       1, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 0, 0, 1, 1, 0, 0, 1, 0, 1, 0,
0,
       0, 1, 1, 0, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1, 1, 1, 0, 0, 0, 0, 0,
0,
       0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0, 1, 1, 0, 1, 0, 1, 0, 1, 1, 0,
0,
       0, 1, 1, 0, 1, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 1, 1,
1,
       0, 1, 0, 0, 1, 0, 0, 0, 0, 1, 1, 0, 0, 0, 1, 0, 1, 1, 0, 1, 0,
0,
       1, 1, 1, 0, 1, 0, 0, 0, 0, 1, 0, 1, 1, 0, 0, 1, 0, 0, 1, 1, 1,
0,
       1, 0, 1, 0, 1, 0, 1, 0, 0, 0, 1, 1, 0, 1, 0, 0, 1, 0, 1, 0, 0,
0,
       0, 1, 1, 0, 1, 0, 1, 0])
accuracy_score(y_true=y_test, y_pred=pred_i) * 100
89.2
accuracy = np.mean(pred i == y test)
accuracy * 100
89.2
```

Breast Cancer

```
from sklearn.datasets import load_breast_cancer
from sklearn.preprocessing import StandardScaler
```

```
data = load breast cancer()
X, y = data.data, data.target
X train, X test, y train, y test = train test split(
   Χ,
   у,
    test size=0.3,
    random state=42
)
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
model1 = Perceptron(learning rate=1000)
model1.fit(X train scaled, y train)
Convergence reached at epoch 105.
prediction = model1.predict(X test scaled)
prediction
array([1, 0, 0, 1, 1, 0, 0, 0, 1, 1, 0, 1, 1, 1, 0, 1, 1, 1, 0, 1,
       0, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 0, 0, 0, 1, 1, 0, 1,
1,
       0, 1, 1, 1, 1, 0, 0, 1, 1, 1, 1, 0, 0, 1, 1, 0, 0, 1, 1,
0,
       0, 0, 1, 1, 0, 0, 1, 0, 1, 1, 1, 0, 1, 1, 0, 1, 0, 0, 0, 0, 0,
0,
       1, 1, 1, 1, 1, 1, 1, 0, 0, 1, 0, 0, 1, 0, 0, 1, 0, 1, 0, 1,
1,
       0, 1, 0, 0, 0, 0, 1, 1, 1, 0, 1, 1, 0, 1, 0, 0, 1, 1, 0, 0,
0,
       1, 1, 1, 0, 1, 1, 1, 0, 1, 0, 1, 1, 0, 1, 0, 0, 1, 1, 0, 1, 1,
1,
       1, 0, 0, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 0, 1])
accuracy_score(y_true=y_test, y_pred=prediction) * 100
94.15204678362574
```

95.32163742690058 -> 0.00001 -> Unscaled 94.15204678362574 -> 0.01 -> Scaled

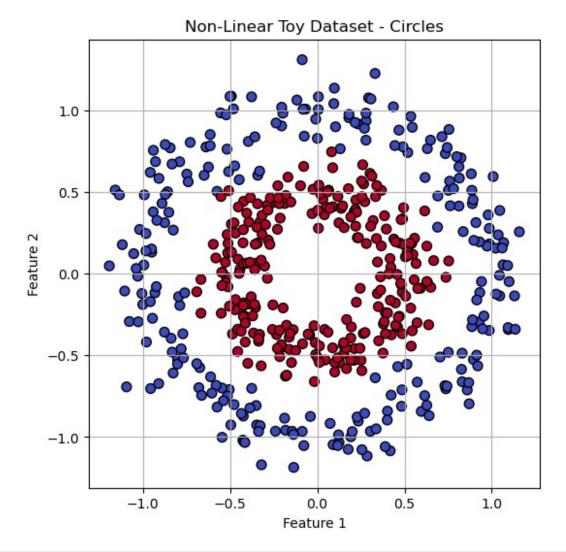
Circle Dataset

```
from sklearn.datasets import make_circles

X, y = make_circles(n_samples=500, noise=0.1, factor=0.5, random_state=42)

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

plt.figure(figsize=(6,6))
plt.scatter(X[:, 0], X[:, 1], c=y, cmap='coolwarm', s=50, edgecolors='black')
plt.title("Non-Linear Toy Dataset - Circles")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.grid(True)
plt.show()
```



```
print("Sample data points (X, y):")
print(X[:10], y[:10])
Sample data points (X, y):
[[-0.46918557 0.24791499]
 [-0.06748724 1.00676912]
 [-0.44306526
             0.02738322]
 [-0.61172505 -0.6314071 ]
 [-0.78901285 0.68451888]
 [-0.42136979 -0.25688616]
 [-0.45473954 0.08850207]
 [-0.94954151
               0.13119395]
 [-0.20468777 0.903653
 [ 0.10000474 -0.60226649]] [1 0 1 0 0 1 1 0 0 1]
model2 = Perceptron()
model2.fit(X train, y train)
```