Some Helper Function:


```
import numpy as np

def softmax(z):
    """
    Compute the softmax function for each row of the input array.

Parameters:
    z (numpy array): A 2D numpy array where softmax is applied row-wise.

Returns:
    numpy array: Softmax probabilities.
    """
    exp_z = np.exp(z - np.max(z, axis=1, keepdims=True))
    return exp_z / np.sum(exp_z, axis=1, keepdims=True)
```

Softmax Test Case:

This test case checks that each row in the resulting softmax probabilities sums to 1, which is the fundamental property of softmax.

```
# Example test case
z_test = np.array([[2.0, 1.0, 0.1], [1.0, 1.0, 1.0]])
softmax_output = softmax(z_test)

# Verify if the sum of probabilities for each row is 1 using assert
row_sums = np.sum(softmax_output, axis=1)

# Assert that the sum of each row is 1
assert np.allclose(row_sums, 1), f"Test failed: Row sums are {row_sums}"
print("Softmax function passed the test case!")

    Softmax function passed the test case!
```

Prediction Function:

```
import numpy as np
def predict_softmax(X, W, b):
    """
    Predict the class labels for a set of samples using the trained softmax model.

Parameters:
    X (numpy.ndarray): Feature matrix of shape (n, d), where n is the number of samples and d is the number of features.
    W (numpy.ndarray): Weight matrix of shape (d, c), where c is the number of classes.
    b (numpy.ndarray): Bias vector of shape (c,).

Returns:
    numpy.ndarray: Predicted class labels of shape (n,), where each value is the index of the predicted class.
    """

# Step 1: Compute the raw scores (logits)
logits = np.dot(X, W) + b # X is (n, d), W is (d, c), so logits will be (n, c)

# Step 2: Apply the softmax function to get the probabilities
probabilities = softmax(logits)

# Step 3: Select the class with the highest probability for each sample
predicted_classes = np.argmax(probabilities, axis=1)

return predicted_classes
```

Test Function for Prediction Function:

The test function ensures that the predicted class labels have the same number of elements as the input samples, verifying that the model produces a valid output shape.

```
# Define test case
X_{\text{test}} = \text{np.array}([[0.2, 0.8], [0.5, 0.5], [0.9, 0.1]]) # Feature matrix (3 samples, 2 features)
W_{\text{test}} = \text{np.array}([[0.4, 0.2, 0.1], [0.3, 0.7, 0.5]]) \# Weights (2 features, 3 classes)
b_test = np.array([0.1, 0.2, 0.3]) # Bias (3 classes)
# Expected Output:
# The function should return an array with class labels (0, 1, or 2)
y_pred_test = predict_softmax(X_test, W_test, b_test)
# Validate output shape
assert y_pred_test.shape == (3,), f"Test failed: Expected shape (3,), got {y_pred_test.shape}"
# Print the predicted labels
print("Predicted class labels:", y_pred_test)
→ Predicted class labels: [1 1 0]
Loss Function:
def loss_softmax(y_pred, y):
    Compute the cross-entropy loss for a single sample.
    Parameters:
    y_pred (numpy.ndarray): Predicted probabilities of shape (c,) for a single sample,
                             where c is the number of classes.
    y (numpy.ndarray): True labels (one-hot encoded) of shape (c,), where c is the number of classes.
    Returns:
    float: Cross-entropy loss for the given sample.
    loss = -np.sum(y * np.log(y_pred)) # Sum over the classes (c)
    return loss
```

Test case for Loss Function:

This test case Compares loss for correct vs. incorrect predictions.

- Expects low loss for correct predictions.
- Expects high loss for incorrect predictions.

```
import numpy as np
# Define correct predictions (low loss scenario)
y_{true\_correct} = np.array([[1, 0, 0], [0, 1, 0], [0, 0, 1]]) # True one-hot labels
y_pred_correct = np.array([[0.9, 0.05, 0.05],
                           [0.1, 0.85, 0.05],
                           [0.05, 0.1, 0.85]]) # High confidence in the correct class
# Define incorrect predictions (high loss scenario)
y_pred_incorrect = np.array([[0.05, 0.05, 0.9], # Highly confident in the wrong class
                              [0.1, 0.05, 0.85],
                              [0.85, 0.1, 0.05]])
# Compute loss for both cases
loss_correct = loss_softmax(y_pred_correct, y_true_correct)
loss_incorrect = loss_softmax(y_pred_incorrect, y_true_correct)
# Validate that incorrect predictions lead to a higher loss
assert loss_correct < loss_incorrect, f"Test failed: Expected loss_correct < loss_incorrect, but got {loss_correct:.4f} >= {loss_incorrect:.
# Print results
```

Cost Function:

```
import numpy as np

def softmax(z):
    """
    Compute the softmax function for each row of the input array.
    """
    exp_z = np.exp(z - np.max(z, axis=1, keepdims=True)) # To improve numerical stability return exp_z / np.sum(exp_z, axis=1, keepdims=True) # Normalize to get probabilities

def loss_softmax(y_pred, y):
    """
    Compute the cross-entropy loss for a single sample.
    """
    loss = -np.sum(y * np.log(y_pred)) # Sum over the classes (c) return loss
```

Test Case for Cost Function:

The test case assures that the cost for the incorrect prediction should be higher than for the correct prediction, confirming that the cost function behaves as expected.

```
import numpy as np
def softmax(z):
    Compute the softmax function for each row of the input array.
    \exp_z = \text{np.exp}(z - \text{np.max}(z, \text{axis=1}, \text{keepdims=True})) \text{ } \# \text{ To improve numerical stability}
    return exp_z / np.sum(exp_z, axis=1, keepdims=True) # Normalize to get probabilities
def loss_softmax(y_pred, y):
    Compute the cross-entropy loss for a single sample.
    loss = -np.sum(y * np.log(y_pred)) # Sum over the classes (c)
    return loss
def cost_softmax(X, y, W, b):
    Compute the total cost (average cross-entropy loss) for the dataset.
    Parameters:
    X (numpy.ndarray): Feature matrix of shape (n, d).
    y (numpy.ndarray): True labels (one-hot encoded) of shape (n, c).
    W (numpy.ndarray): Weight matrix of shape (d, c).
    b (numpy.ndarray): Bias vector of shape (c,).
    Returns:
    float: Total cost.
    z = np.dot(X, W) + b
    y_pred = softmax(z)
    total_cost = np.mean([loss_softmax(y_pred[i], y[i]) for i in range(X.shape[0])])
    return total_cost
# Example 1: Correct Prediction (Closer predictions)
X_correct = np.array([[1.0, 0.0], [0.0, 1.0]]) # Feature matrix for correct predictions
y\_correct = np.array([[1, \ 0], \ [0, \ 1]]) \quad \text{\# True labels (one-hot encoded, matching predictions)}
W_{correct} = np.array([[5.0, -2.0], [-3.0, 5.0]]) # Weights for correct prediction
b_{correct} = np.array([0.1, 0.1]) # Bias for correct prediction
# Example 2: Incorrect Prediction (Far off predictions)
X_{incorrect} = np.array([[0.1, 0.9], [0.8, 0.2]]) # Feature matrix for incorrect predictions
y_{incorrect} = np.array([[1, 0], [0, 1]]) # True labels (one-hot encoded, incorrect predictions)
W_{incorrect} = np.array([[0.1, 2.0], [1.5, 0.3]]) # Weights for incorrect prediction
```

```
b_incorrect = np.array([0.5, 0.6]) # Bias for incorrect prediction

# Compute cost for correct predictions
cost_correct = cost_softmax(X_correct, y_correct, W_correct, b_correct)

# Compute cost for incorrect predictions
cost_incorrect = cost_softmax(X_incorrect, y_incorrect, W_incorrect, b_incorrect)

# Check if the cost for incorrect predictions is greater than for correct predictions
assert cost_incorrect > cost_correct, f"Test failed: Incorrect cost {cost_incorrect} is not greater than correct cost {cost_correct}"

# Print the costs for verification
print("Cost for correct prediction:", cost_correct)
print("Cost for incorrect prediction:", cost_incorrect)

print("Test passed!")

Cost for correct prediction: 0.0006234364133349324
Cost for incorrect prediction: 0.29930861359446115
Test passed!
```

Computing Gradients:

```
import numpy as np
# Softmax function
def softmax(z):
   \exp_z = \text{np.exp}(z - \text{np.max}(z, \text{axis=1, keepdims=True})) # To improve numerical stability
   return exp_z / np.sum(exp_z, axis=1, keepdims=True) # Normalize to get probabilities
def compute_gradient_softmax(X, y, W, b):
   Compute the gradients of the cost function with respect to weights and biases.
   Parameters:
   X (numpy.ndarray): Feature matrix of shape (n, d).
   y (numpy.ndarray): True labels (one-hot encoded) of shape (n, c).
   W (numpy.ndarray): Weight matrix of shape (d, c).
   b (numpy.ndarray): Bias vector of shape (c,).
   Returns:
   tuple: Gradients with respect to weights (d, c) and biases (c,).
   # Number of samples
   n = X.shape[0]
   # Compute logits (linear combination of inputs and weights)
   logits = np.dot(X, W) + b # Shape (n, c)
   # Compute softmax probabilities
   y_pred = softmax(logits) # Shape (n, c)
   # Compute the gradient of the cost w.r.t. weights
   grad_W = (1 / n) * np.dot(X.T, (y_pred - y)) # Shape (d, c)
   # Compute the gradient of the cost w.r.t. biases
   grad_b = (1 / n) * np.sum(y_pred - y, axis=0) # Shape (c,)
   return grad_W, grad_b
```

Test case for compute_gradient function:

The test checks if the gradients from the function are close enough to the manually computed gradients using np.allclose, which accounts for potential floating-point discrepancies.

```
import numpy as np

# Define a simple feature matrix and true labels

X_test = np.array([[0.2, 0.8], [0.5, 0.5], [0.9, 0.1]])  # Feature matrix (3 samples, 2 features)

y_test = np.array([[1, 0, 0], [0, 1, 0], [0, 0, 1]])  # True labels (one-hot encoded, 3 classes)

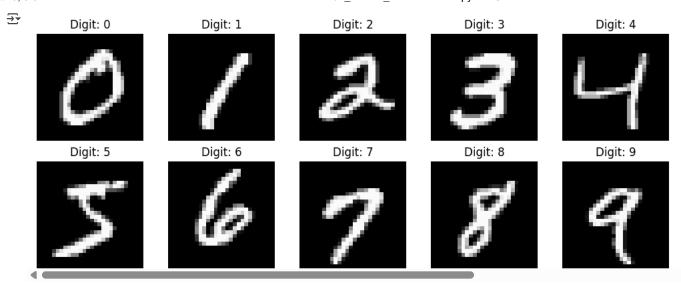
# Define weight matrix and bias vector
```

```
W_test = np.array([[0.4, 0.2, 0.1], [0.3, 0.7, 0.5]]) # Weights (2 features, 3 classes)
b_{test} = np.array([0.1, 0.2, 0.3]) # Bias (3 classes)
# Compute the gradients using the function
grad_W, grad_b = compute_gradient_softmax(X_test, y_test, W_test, b_test)
# Manually compute the predicted probabilities (using softmax function)
z test = np.dot(X test, W test) + b test
y_pred_test = softmax(z_test)
# Compute the manually computed gradients
grad_W_manual = np.dot(X_test.T, (y_pred_test - y_test)) / X_test.shape[0]
grad_b_manual = np.sum(y_pred_test - y_test, axis=0) / X_test.shape[0]
# Assert that the gradients computed by the function match the manually computed gradients
assert np.allclose(grad_W, grad_W_manual), f"Test failed: Gradients w.r.t. W are not equal.\nExpected: {grad_W_manual}\nGot: {grad_W}"
assert np.allclose(grad_b, grad_b_manual), f"Test failed: Gradients w.r.t. b are not equal.\nExpected: {grad_b_manual}\nGot: {grad_b}"
# Print the gradients for verification
print("Gradient w.r.t. W:", grad_W)
print("Gradient w.r.t. b:", grad_b)
print("Test passed!")
→ Gradient w.r.t. W: [[ 0.1031051 0.01805685 -0.12116196]
      [-0.13600547 0.00679023 0.12921524]]
     Gradient w.r.t. b: [-0.03290036 0.02484708 0.00805328]
     Test passed!
    Implementing Gradient Descent:
\label{lem:def_gradient_descent_softmax} \mbox{(X, y, W, b, alpha, n\_iter, show\_cost=False):} \\
    Perform gradient descent to optimize the weights and biases.
    Parameters:
    X (numpy.ndarray): Feature matrix of shape (n, d).
    y (numpy.ndarray): True labels (one-hot encoded) of shape (n, c).
    W (numpy.ndarray): Weight matrix of shape (d, c).
    b (numpy.ndarray): Bias vector of shape (c,).
    alpha (float): Learning rate.
    n_iter (int): Number of iterations.
    show_cost (bool): Whether to display the cost at intervals.
    Returns:
    tuple: Optimized weights, biases, and cost history.
    cost_history = []
    for i in range(n iter):
        # Compute gradients
        grad_W, grad_b = compute_gradient_softmax(X, y, W, b)
        # Update weights and biases using gradient descent
        W -= alpha * grad_W # Update weights
b -= alpha * grad_b # Update biases
        # Compute cost (optional: track cost during optimization)
        if show_cost and (i \% 100 == 0 or i == n_iter - 1): # Display cost every 100 iterations
            cost = cost\_softmax(X, y, W, b) # Compute the current cost
            cost_history.append(cost)
            print(f"Iteration {i}, Cost: {cost}") # Print the cost
    return W, b, cost_history
```

Preparing Dataset:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split
```

```
def load_and_prepare_mnist(csv_file, test_size=0.2, random_state=42):
    Reads the MNIST CSV file, splits data into train/test sets, and plots one image per class.
    Arguments:
                        : Path to the CSV file containing MNIST data.
    csv file (str)
    test_size (float)
                        : Proportion of the data to use as the test set (default: 0.2).
    random_state (int) : Random seed for reproducibility (default: 42).
    Returns:
    X_train, X_test, y_train, y_test : Split dataset.
    # Load dataset
    df = pd.read_csv(csv_file)
    # Separate labels and features
    y = df.iloc[:, 0].values # First column is the label
    X = df.iloc[:, 1:].values # Remaining columns are pixel values
    # Normalize pixel values (optional but recommended)
    X = X / 255.0 # Scale values between 0 and 1
    # Split data into train and test sets
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=test_size, random_state=random_state)
    # Plot one sample image per class
    plot_sample_images(X, y)
    return X_train, X_test, y_train, y_test
def plot_sample_images(X, y):
    Plots one sample image for each digit class (0-9).
    Arguments:
    X (np.ndarray): Feature matrix containing pixel values.
    y (np.ndarray): Labels corresponding to images.
    plt.figure(figsize=(10, 4))
    unique_classes = np.unique(y) # Get unique class labels
    for i, digit in enumerate(unique_classes):
        index = np.where(y == digit)[0][0] \quad \# \ Find \ first \ occurrence \ of \ the \ class
        image = X[index].reshape(28, 28) # Reshape 1D array to 28x28
       plt.subplot(2, 5, i + 1)
        plt.imshow(image, cmap='gray')
       plt.title(f"Digit: {digit}")
       plt.axis('off')
    plt.tight_layout()
    plt.show()
csv_file_path = "/content/drive/MyDrive/AI/mnist_dataset.csv" # Path to saved dataset
X_train, X_test, y_train, y_test = load_and_prepare_mnist(csv_file_path)
```



from google.colab import drive
drive.mount('/content/drive')

→ Mounted at /content/drive

A Quick debugging Step:

```
# Assert that X and y have matching lengths assert len(X_{train}) = len(y_{train}), f"Error: X and y have different lengths! X=\{len(X_{train})\}, y=\{len(y_{train})\}" print("Move forward: Dimension of Feture Matrix X and label vector y matched.")
```

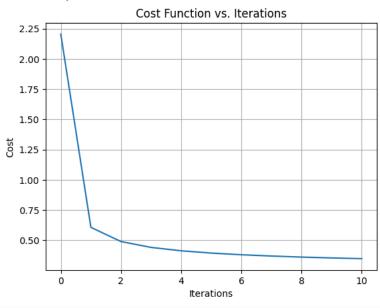
→ Move forward: Dimension of Feture Matrix X and label vector y matched.

Train the Model:

```
print(f"Training data shape: {X_train.shape}")
print(f"Test data shape: {X_test.shape}")
    Training data shape: (48000, 784)
     Test data shape: (12000, 784)
from sklearn.preprocessing import OneHotEncoder
# Check if y_train is one-hot encoded
if len(y_train.shape) == 1:
    encoder = OneHotEncoder(sparse_output=False) # Use sparse_output=False for newer versions of sklearn
    y_train = encoder.fit_transform(y_train.reshape(-1, 1)) # One-hot encode labels
    y_{test} = encoder.transform(y_{test.reshape(-1, 1)) # One-hot encode test labels
# Now y_train is one-hot encoded, and we can proceed to use it
d = X_train.shape[1] # Number of features (columns in X_train)
c = y_train.shape[1] # Number of classes (columns in y_train after one-hot encoding)
# Initialize weights with small random values and biases with zeros
W = np.random.randn(d, c) * 0.01 # Small random weights initialized
b = np.zeros(c) # Bias initialized to 0
# Set hyperparameters for gradient descent
alpha = 0.1 # Learning rate
n_iter = 1000 # Number of iterations to run gradient descent
# Train the model using gradient descent
 \textit{W\_opt, b\_opt, cost\_history = gradient\_descent\_softmax(X\_train, y\_train, W, b, alpha, n\_iter, show\_cost=True) } 
# Plot the cost history to visualize the convergence
plt.plot(cost_history)
plt.title('Cost Function vs. Iterations')
plt.xlabel('Iterations')
plt.ylabel('Cost')
plt.grid(True)
```

```
plt.show()
```

```
Titeration 0, Cost: 2.204672924174595
Iteration 100, Cost: 0.6068774717185383
Iteration 200, Cost: 0.48949204640531846
Iteration 300, Cost: 0.44099567202804435
Iteration 400, Cost: 0.4129607147662435
Iteration 500, Cost: 0.3941123485795775
Iteration 600, Cost: 0.38029851497363126
Iteration 700, Cost: 0.36959375473614575
Iteration 800, Cost: 0.3609687567122186
Iteration 900, Cost: 0.35381704925812224
Iteration 999, Cost: 0.34781113923001
```



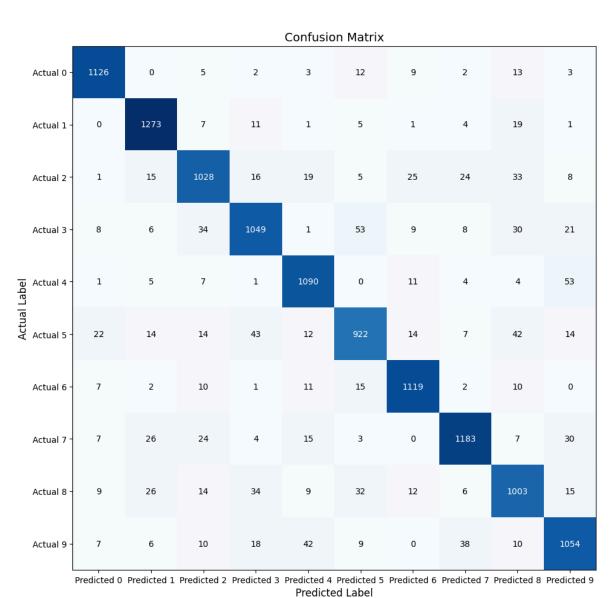
Evaluating the Model:

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.metrics import confusion_matrix, precision_score, recall_score, f1_score
# Evaluation Function
def evaluate_classification(y_true, y_pred):
    Evaluate classification performance using confusion matrix, precision, recall, and F1-score.
    y true (numpy.ndarray): True labels
    y_pred (numpy.ndarray): Predicted labels
    tuple: Confusion matrix, precision, recall, F1 score
    # Compute confusion matrix
    cm = confusion_matrix(y_true, y_pred)
    # Compute precision, recall, and F1-score
    precision = precision_score(y_true, y_pred, average='weighted')
    recall = recall_score(y_true, y_pred, average='weighted')
    f1 = f1_score(y_true, y_pred, average='weighted')
    return cm, precision, recall, f1
# Predict on the test set
y_pred_test = predict_softmax(X_test, W_opt, b_opt)
# Evaluate accuracy
y_test_labels = np.argmax(y_test, axis=1) # True labels in numeric form
# Evaluate the model
```

```
cm, precision, recall, f1 = evaluate_classification(y_test_labels, y_pred_test)
# Print the evaluation metrics
print("\nConfusion Matrix:")
print(cm)
print(f"Precision: {precision:.2f}")
print(f"Recall: {recall:.2f}")
print(f"F1-Score: {f1:.2f}")
# Visualizing the Confusion Matrix
fig, ax = plt.subplots(figsize=(12, 12))
cax = ax.imshow(cm, cmap='Blues') # Use a color map for better visualization
# Dynamic number of classes
num_classes = cm.shape[0]
ax.set_xticks(range(num_classes))
ax.set_yticks(range(num_classes))
ax.set_xticklabels([f'Predicted {i}' for i in range(num_classes)])
ax.set_yticklabels([f'Actual {i}' for i in range(num_classes)])
# Add labels to each cell in the confusion matrix
for i in range(cm.shape[0]):
    for j in range(cm.shape[1]):
        ax.text(j, i, cm[i, j], ha='center', va='center', color='white' if cm[i, j] > np.max(cm) / 2 else 'black')
# Add grid lines and axis labels
ax.grid(False)
plt.title('Confusion Matrix', fontsize=14)
plt.xlabel('Predicted Label', fontsize=12)
plt.ylabel('Actual Label', fontsize=12)
# Adjust layout
plt.tight_layout()
plt.colorbar(cax)
plt.show()
```

Con	fusi	ion Ma	atrix	:						
[[1	126	0	5	2	3	12	9	2	13	3]
[0	1273	7	11	1	5	1	4	19	1]
[1	15	1028	16	19	5	25	24	33	8]
[8	6	34	1049	1	53	9	8	30	21]
[1	5	7	1	1090	0	11	4	4	53]
[22	14	14	43	12	922	14	7	42	14]
[7	2	10	1	11	15	1119	2	10	0]
[7	26	24	4	15	3	0	1183	7	30]
[9	26	14	34	9	32	12	6	1003	15]
[7	6	10	18	42	9	0	38	10	1054]]

Precision: 0.90 Recall: 0.90 F1-Score: 0.90



- 800 - 600 - 400

1200

- 1000

Linear Seperability and Logistic Regression:

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make_classification, make_circles
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
np.random.seed(42)
X_linear_separable, y_linear_separable = make_classification(
    n_samples=200, n_features=2, n_informative=2,
    n_redundant=0, n_clusters_per_class=1, random_state=42
)
X_train_linear, X_test_linear, y_train_linear, y_test_linear = train_test_split(
    X_linear_separable, y_linear_separable, test_size=0.2, random_state=42
logistic model linear separable = LogisticRegression()
logistic_model_linear_separable.fit(X_train_linear, y_train_linear)
X non linear separable, y non linear separable = make circles(
    n_samples=200, noise=0.1, factor=0.5, random_state=42
X_train_non_linear, X_test_non_linear, y_train_non_linear, y_test_non_linear = train_test_split(
    X_non_linear_separable, y_non_linear_separable, test_size=0.2, random_state=42
logistic_model_non_linear_separable = LogisticRegression()
logistic_model_non_linear_separable.fit(X_train_non_linear, y_train_non_linear)
def plot_decision_boundary(ax, model, X, y, title):
    h = 0.02
    x_{min}, x_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
    y_{min}, y_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
    xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))
    Z = model.predict(np.c_[xx.ravel(), yy.ravel()])
    Z = Z.reshape(xx.shape)
    ax.contourf(xx, yy, Z, alpha=0.8, cmap=plt.cm.Paired)
    ax.scatter(X[:, 0], X[:, 1], c=y, edgecolors='k', cmap=plt.cm.Paired)
    ax.set_title(title)
    ax.set_xlabel('Feature 1')
    ax.set_ylabel('Feature 2')
fig, axes = plt.subplots(2, 2, figsize=(12, 10))
plot_decision_boundary(axes[0, 0], logistic_model_linear_separable, X_train_linear, y_train_linear,
                       'Linearly Separable Data (Training)')
plot_decision_boundary(axes[0, 1], logistic_model_linear_separable, X_test_linear, y_test_linear,
                       'Linearly Separable Data (Testing)')
plot_decision_boundary(axes[1, 0], logistic_model_non_linear_separable, X_train_non_linear,
                       y_train_non_linear, 'Non-Linearly Separable Data (Training)')
plot_decision_boundary(axes[1, 1], logistic_model_non_linear_separable, X_test_non_linear,
                       y_test_non_linear, 'Non-Linearly Separable Data (Testing)')
plt.tight_layout()
plt.savefig('decision_boundaries.png')
plt.show()
```