

* Exercise 1: Decision Boundaries of Generative Classifiers

- Task 1: Gaussian Discriminant Analysis

(a) QDA:

We assume $A \not\approx B$ with Gaussian conditionals

$$p(x|y=k) = \frac{1}{\sqrt{(2\pi)^N \det \Sigma_k}} \exp\left[-\frac{1}{2} (x - \mu_k)^T \Sigma_k^{-1} (x - \mu_k)\right]$$

& Also $\pi_k = p(y=k)$. The MAP-discriminant is,

$$\begin{aligned} g_k(x) &= \log p(x|y=k) + \log \pi_k \\ &= -\frac{1}{2} \log \det \Sigma_k - \frac{1}{2} (x - \mu_k)^T \Sigma_k^{-1} (x - \mu_k) + \log \pi_k \end{aligned}$$

Setting $g_A(x) = g_B(x)$

$$-\frac{1}{2} \log \det \Sigma_A - \frac{1}{2} (x - \mu_A)^T \Sigma_A^{-1} (x - \mu_A) + \log \pi_A = -\frac{1}{2} \log \det \Sigma_B - \frac{1}{2} (x - \mu_B)^T \Sigma_B^{-1} (x - \mu_B) + \log \pi_B$$

Taking them to one side & expanding the quadratic forms:

$$-\frac{1}{2} x^T (\Sigma_A^{-1} - \Sigma_B^{-1}) x + (\Sigma_A^{-1} \mu_A - \Sigma_B^{-1} \mu_B)^T x + \left[-\frac{1}{2} (\mu_A^T \Sigma_A^{-1} \mu_A - \mu_B^T \Sigma_B^{-1} \mu_B) - \frac{1}{2} \ln \frac{\det \Sigma_A}{\det \Sigma_B} + \ln \frac{\pi_A}{\pi_B} \right] = 0$$

$$\text{Let, } \Lambda = -\frac{1}{2} (\Sigma_A^{-1} - \Sigma_B^{-1}), w = \Sigma_A^{-1} \mu_A - \Sigma_B^{-1} \mu_B, b = -\frac{1}{2} (\mu_A^T \Sigma_A^{-1} \mu_A - \mu_B^T \Sigma_B^{-1} \mu_B) - \frac{1}{2} \ln \frac{\det \Sigma_A}{\det \Sigma_B} + \ln \frac{\pi_A}{\pi_B}$$

So the Decision Boundary is,

$$x^T \Lambda x + w^T x + b = 0$$

⇒ Here $x^T \Lambda x = \sum_{ij} \Lambda_{ij} x_i x_j$ is a 2nd degree func. of the components of x .

Upon solving the eqn, we'll get set of points satisfying a degree-2 polynomial equation - i.e. A conic in 2D. Hence quadratic term indicates non-linear decision boundary.

(b) LDA:

$$\text{Let } \Sigma_A = \Sigma_B = \Sigma$$

$$g_k(x) = \log p(x|y=k) + \log \pi_k$$

$$= -\frac{1}{2} (x - \mu_k)^T \Sigma^{-1} (x - \mu_k) + \log \pi_k$$

& setting $g_A(x) = g_B(x)$,

$$-\frac{1}{2} [(x - \mu_A)^T \Sigma^{-1} (x - \mu_A) - (x - \mu_B)^T \Sigma^{-1} (x - \mu_B) + \log \frac{\pi_A}{\pi_B}] = 0$$

Expanding & canceling the common Σ^{-1} terms,

$$x \left[\Sigma^{-1} (\mu_A - \mu_B)^T - \frac{1}{2} (\mu_A^T \Sigma^{-1} \mu_A - \mu_B^T \Sigma^{-1} \mu_B) + \log \frac{\pi_A}{\pi_B} \right] = 0$$

\therefore Decision boundary can be written as,

$$w^T x + b = 0$$

$$w = \Sigma^{-1} (\mu_A - \mu_B)$$

$$b = -\frac{1}{2} [\mu_A^T \Sigma^{-1} \mu_A - \mu_B^T \Sigma^{-1} \mu_B] + \log \frac{\pi_A}{\pi_B}$$

- Task 2 :

a)

```

muA = np.array([-1.0, -1.0])
muB = np.array([1.0, 1.0])
SigmaA = np.array([[1.0, 0.3],
                  [0.3, 1.0]])
SigmaB = np.array([[1.5, -0.2],
                  [-0.2, 1.5]])

piA, piB = 0.5, 0.5

# Inverses
invA = np.linalg.inv(SigmaA)
invB = np.linalg.inv(SigmaB)

# QDA parameters
Lambda = -0.5 * (invA - invB)
w_qda = invA.dot(muA) - invB.dot(muB)
b_qda = (
    -0.5 * (muA.T.dot(invA).dot(muA) - muB.T.dot(invB).dot(muB))
    - 0.5 * np.log(np.linalg.det(SigmaA) / np.linalg.det(SigmaB))
    + np.log(piA / piB)
)

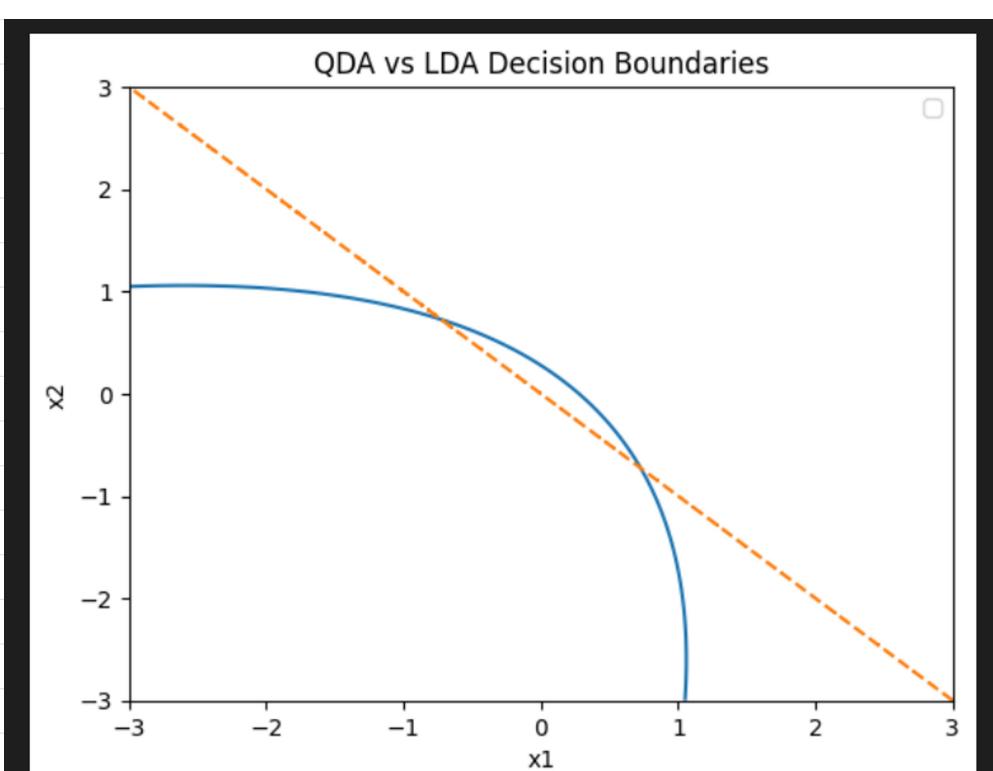
# LDA parameters
Sigma_pooled = 0.5 * (SigmaA + SigmaB)
inv_pooled = np.linalg.inv(Sigma_pooled)
w_lda = inv_pooled.dot(muA - muB)
b_lda = (
    -0.5 * muA.T.dot(inv_pooled).dot(muA) + 0.5 * muB.T.dot(inv_pooled).dot(muB)
    + np.log(piA / piB)
)

print("QDA parameters:")
print("Lambda:", Lambda)
print("w_qda:", w_qda)
print("b_qda:", b_qda)
print("\nLDA parameters:")
print("w_lda:", w_lda)
print("b_lda:", b_lda)
print("0.1s")

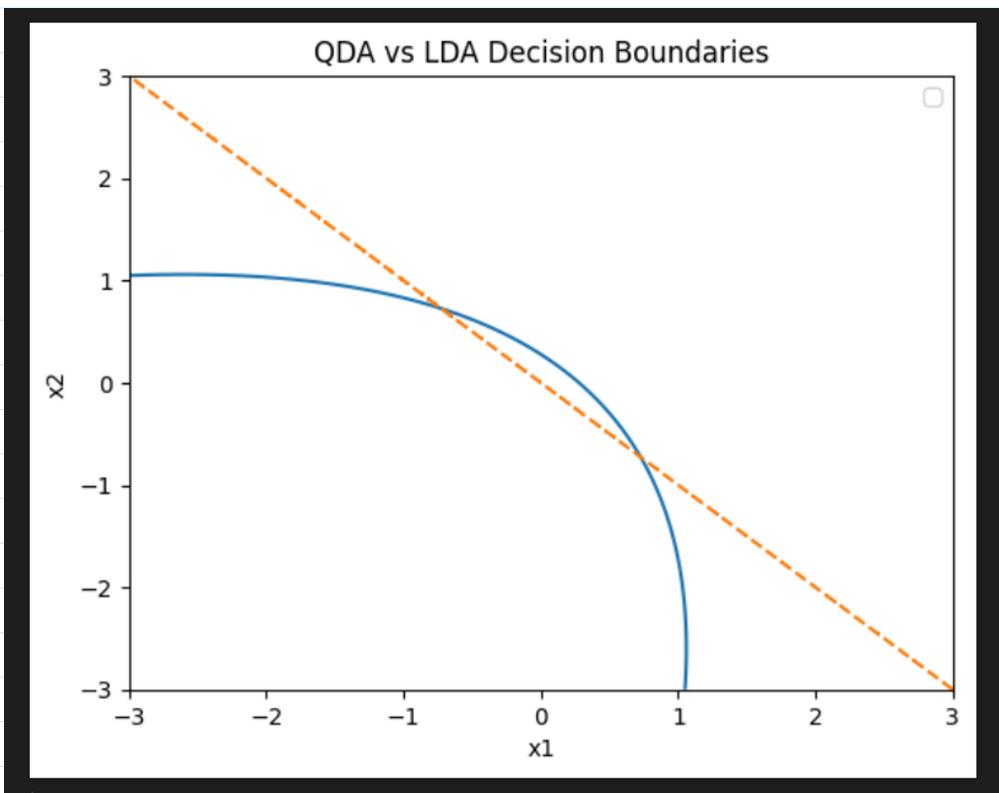
```

QDA parameters:
Lambda:
[-0.21008403 0.21008403]
[0.21008403 -0.21008403]
w_qda:
[-1.53846154 -1.53846154]
b_qda:
0.44365159750045136

LDA parameters:
w_lda:
[-1.53846154 -1.53846154]
b_lda:
0.0



b)



c) when $\Sigma_A \neq \Sigma_B$, each class has its own "shape" & orientation in feature space. QDA plugs both Σ_A^{-1} & Σ_B^{-1} into discriminant, so the boundary retains those 2nd order ($x_i x_j$) terms. Geometrically, this means the decision curve can bend, twist even from ellipses or hyperbola, following exactly how each Gaussian Contours differ.

On other hand, LDA forces $\Sigma_A = \Sigma_B = \Sigma$, we get

$$w^T x + b = 0$$

a single straight line that ignores any per-class anisotropy.

- QDA can be preferred when we have ample training samples & strong evidence that Σ_A & Σ_B really differ.
- LDA can be preferred when data is limited or maybe classes have same covariance struct. or we need linear rule

Machine Learning Essentials SS25 - Exercise Sheet 2

Instructions

- TODO 's indicate where you need to complete the implementations.
- You may use external resources, but **write your own solutions**.
- Provide concise, but comprehensible comments to explain what your code does.
- Code that's unnecessarily extensive and/or not well commented will not be scored.

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_digits
from sklearn.decomposition import PCA
from sklearn.model_selection import train_test_split
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
from scipy.stats import multivariate_normal
np.random.seed(42)
```

Exercise 1 - Part 2

Task 2a

```
In [2]: muA = np.array([-1.0, -1.0])
muB = np.array([1.0, 1.0])
SigmaA = np.array([[1.0, 0.3],
                  [0.3, 1.0]])
SigmaB = np.array([[1.5, -0.2],
                  [-0.2, 1.5]])

piA, piB = 0.5, 0.5

# Inverses
invA = np.linalg.inv(SigmaA)
invB = np.linalg.inv(SigmaB)

# QDA parameters
Lambda = -0.5 * (invA - invB)
w_qda = invA.dot(muA) - invB.dot(muB)
b_qda = (
    -0.5 * (muA.T.dot(invA).dot(muA) - muB.T.dot(invB).dot(muB))
    -0.5 * np.log(np.linalg.det(SigmaA) / np.linalg.det(SigmaB))
    + np.log(piA / piB)
)

# LDA parameters
Sigma_pooled = 0.5 * (SigmaA + SigmaB)
inv_pooled = np.linalg.inv(Sigma_pooled)
w_lda = inv_pooled.dot(muA - muB)
b_lda = (
    -0.5 * muA.T.dot(inv_pooled).dot(muA) + 0.5 * muB.T.dot(inv_pooled).dot(
        muB
    )
    + np.log(piA / piB)
)
```

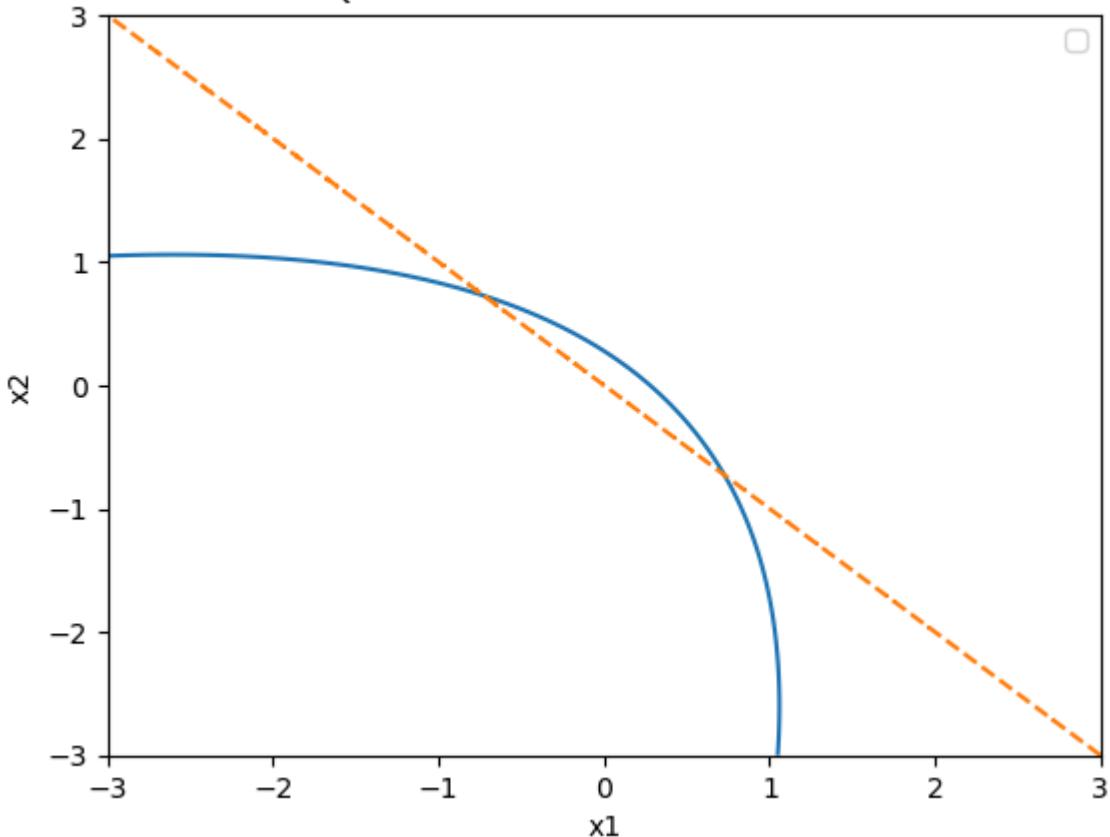
```
)  
  
print("QDA parameters:")
print("Lambda:\n", Lambda)
print("w_qda:\n", w_qda)
print("b_qda:\n", b_qda)
print("\nLDA parameters:")
print("w_lda:\n", w_lda)
print("b_lda:\n", b_lda)
```

```
QDA parameters:  
Lambda:  
[[ -0.21008403  0.21008403]  
 [ 0.21008403 -0.21008403]]  
w_qda:  
[-1.53846154 -1.53846154]  
b_qda:  
0.44365159750045136  
  
LDA parameters:  
w_lda:  
[-1.53846154 -1.53846154]  
b_lda:  
0.0
```

Task 2b

```
In [3]: invP = np.linalg.inv(Sigma_pooled)  
def g_qda(x):  
    return (-0.5*np.log(np.linalg.det(SigmaA))  
           -0.5*(x-muA)@invA@(x-muA)  
           +np.log(piA)  
           ) - (  
           -0.5*np.log(np.linalg.det(SigmaB))  
           -0.5*(x-muB)@invB@(x-muB)  
           +np.log(piB)  
           )  
  
def g_lda(x):  
    w = invP@(muA-muB)  
    b = -0.5*(muA@invP@muA - muB@invP@muB) + np.log(piA/piB)  
    return w@x + b  
  
xs = np.linspace(-3,3,400)  
ys = np.linspace(-3,3,400)  
X, Y = np.meshgrid(xs, ys)  
Z_qda = np.array([g_qda([x,y]) for x,y in zip(X.ravel(), Y.ravel())]).reshape(400,400)  
Z_lda = np.array([g_lda([x,y]) for x,y in zip(X.ravel(), Y.ravel())]).reshape(400,400)  
  
# Plot  
plt.contour(X, Y, Z_qda, levels=[0], colors='C0')  
plt.contour(X, Y, Z_lda, levels=[0], linestyles='--', colors='C1')  
plt.xlabel('x1')  
plt.ylabel('x2')  
plt.legend(['QDA', 'LDA'])  
plt.title('QDA vs LDA Decision Boundaries')  
plt.show()
```

QDA vs LDA Decision Boundaries



In []:

```
In [4]: import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_digits
from sklearn.decomposition import PCA
from sklearn.model_selection import train_test_split
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
from sklearn.model_selection import KFold
from scipy.stats import multivariate_normal
np.random.seed(42)
```

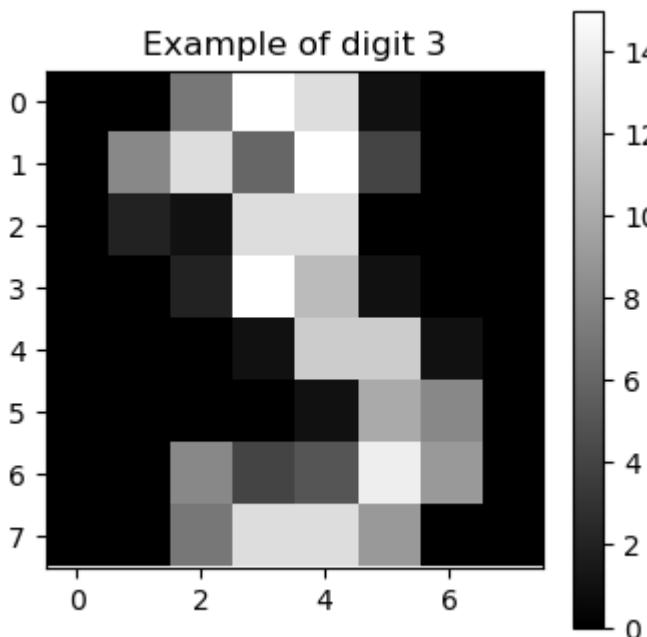
Exercise 2 - Implementing LDA

Task 1

```
In [5]: # TODO: Load digits dataset, visualize one example image of digit 3
digits = load_digits()
X, y = digits.data, digits.target

digit_3_indices = np.where(y == 3)[0]
example_3_idx = digit_3_indices[0] # Take the first occurrence

example_3 = X[example_3_idx].reshape(8, 8)
plt.figure(figsize=(4, 4))
plt.imshow(example_3, cmap='gray', interpolation='nearest')
plt.title(f"Example of digit 3")
plt.colorbar()
plt.show()
```



Task 2

```
In [6]: # TODO: Filter the dataset to keep only digits 3 and 9, split into training
mask_3_9 = np.logical_or(y == 3, y == 9)
X_filtered = X[mask_3_9]
y_filtered = y[mask_3_9]

# Transform labels to binary (3 -> -1, 9 -> 1)
y_binary = np.where(y_filtered == 3, -1, 1)

# Split the data with ratio 3:2 for train:test
X_train, X_test, y_train, y_test = train_test_split(
    X_filtered, y_binary, test_size=0.4, random_state=42
)

print(f"Number of samples for digit 3: {np.sum(y_binary == -1)}")
print(f"Number of samples for digit 9: {np.sum(y_binary == 1)}")
print(f"Training set size: {X_train.shape[0]}")
print(f"Test set size: {X_test.shape[0]}")
```

Number of samples for digit 3: 183
 Number of samples for digit 9: 180
 Training set size: 217
 Test set size: 146

Task 3

```
In [7]: def features_2d(x):
    """
    This function takes the 64x1 feature vectors and returns a 2D representation
    """
    # Select the two most discriminative features based on variance
    feature_variances = np.var(X_train, axis=0) # Compute variance for each
    most_discriminative_indices = np.argsort(feature_variances)[-2:] # Get the two most discriminative indices

    feature1 = x[:, most_discriminative_indices[0]]
    feature2 = x[:, most_discriminative_indices[1]]

    return np.column_stack((feature1, feature2))
```

```
# Create an embedded dataset using the selected features
X_train_2d = features_2d(X_train)
X_test_2d = features_2d(X_test)

print(f"Shape of transformed training data: {X_train_2d.shape}")
print(f"Shape of transformed test data: {X_test_2d.shape}")
```

Shape of transformed training data: (217, 2)
 Shape of transformed test data: (146, 2)

Task 4

```
In [8]: def pca_rep(x):
    """
    This function takes the 64x1 feature vectors and returns a 2D representation
    """
    # Standardize the data
    pca = PCA(n_components=2)
    return pca.fit_transform(x)

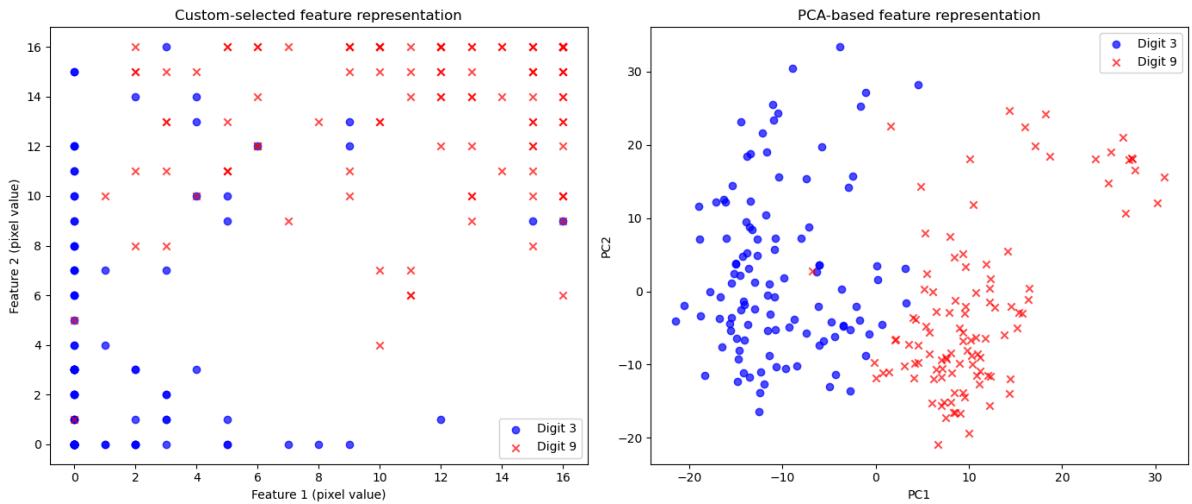
# TODO: Create a PCA-embedded dataset. Visualize & compare the embeddings. Etc.
pca = PCA(n_components=2)
pca.fit(X_train)
X_train_pca = pca.transform(X_train)
X_test_pca = pca.transform(X_test)

# Plot both feature representations
plt.figure(figsize=(14, 6))

plt.subplot(1, 2, 1)
for label, marker, color in zip([-1, 1], ['o', 'x'], ['blue', 'red']):
    plt.scatter(
        X_train_2d[y_train == label, 0],
        X_train_2d[y_train == label, 1],
        marker=marker,
        color=color,
        alpha=0.7,
        label=f'Digit {"3" if label == -1 else "9"}'
    )
plt.title("Custom-selected feature representation")
plt.xlabel("Feature 1 (pixel value)")
plt.ylabel("Feature 2 (pixel value)")
plt.legend()

plt.subplot(1, 2, 2)
for label, marker, color in zip([-1, 1], ['o', 'x'], ['blue', 'red']):
    plt.scatter(
        X_train_pca[y_train == label, 0],
        X_train_pca[y_train == label, 1],
        marker=marker,
        color=color,
        alpha=0.7,
        label=f'Digit {"3" if label == -1 else "9"}'
    )
plt.title("PCA-based feature representation")
plt.xlabel("PC1")
plt.ylabel("PC2")
plt.legend()
```

```
plt.tight_layout()
plt.show()
```



Task 5

```
In [9]: def fit_lda(training_features, training_labels):
    """
    Compute LDA parameters.
    """

    # TODO: Implement LDA
    # Filter out "dead" pixels (features with very low variance)
    variances = np.var(training_features, axis=0)
    active_indices = np.where(variances > 0.001)[0]
    features = training_features[:, active_indices]

    # Get unique classes
    classes = np.unique(training_labels)

    # Compute class means
    mu = {}
    for k in classes:
        mu[k] = np.mean(features[training_labels == k], axis=0)

    # Compute priors
    p = {}
    n_samples = features.shape[0]
    for k in classes:
        p[k] = np.sum(training_labels == k) / n_samples

    # Compute pooled covariance matrix
    covmat = np.zeros((features.shape[1], features.shape[1]))
    for k in classes:
        class_samples = features[training_labels == k]
        centered = class_samples - mu[k]
        covmat += centered.T @ centered

    covmat /= n_samples

    return mu, covmat, p, active_indices

    # TODO: Fit separate LDA models using your hand-crafted embedding, the PCA embedding, and the raw pixel values
    # Filter out dead pixels for the full dataset before fitting
    X_train_filtered = X_train.copy()
    X_test_filtered = X_test.copy()
```

```
# Fit LDA to the 3 different feature sets
mu_2d, covmat_2d, p_2d, active_indices_2d = fit_lda(X_train_2d, y_train)
mu_pca, covmat_pca, p_pca, active_indices_pca = fit_lda(X_train_pca, y_train)
mu_full, covmat_full, p_full, active_indices_full = fit_lda(X_train, y_train)

print(f"Number of active features in full dataset: {len(active_indices_full)}")
print(f"Shape of covariance matrix for full dataset: {covmat_full.shape}")
```

Number of active features in full dataset: 55

Shape of covariance matrix for full dataset: (55, 55)

Task 6

```
In [10]: def predict_lda(mu, covmat, p, test_features):
    """
    Predict labels using the LDA decision rule.
    """

    # TODO: Implement the LDA decision rule
    try:
        precision_matrix = np.linalg.inv(covmat)
    except np.linalg.LinAlgError:
        # Add a small regularization if matrix is singular
        precision_matrix = np.linalg.inv(covmat + 1e-6 * np.eye(covmat.shape[0]))

    w = precision_matrix @ (mu[1] - mu[-1])

    b = -0.5 * (mu[1].T @ precision_matrix @ mu[1] -
                 mu[-1].T @ precision_matrix @ mu[-1]) + np.log(p[1] / p[-1])

    # Predict labels
    scores = test_features @ w + b
    predicted_labels = np.sign(scores)

    return predicted_labels

# TODO: Perform LDA on the filtered train sets of all 3 embeddings, evaluate error rates

# Function to calculate error rate
def error_rate(y_true, y_pred):
    return np.mean(y_true != y_pred)

# Predict using our custom 2D features
X_train_2d_active = X_train_2d[:, active_indices_2d]
X_test_2d_active = X_test_2d[:, active_indices_2d]

y_train_pred_2d = predict_lda(mu_2d, covmat_2d, p_2d, X_train_2d_active)
y_test_pred_2d = predict_lda(mu_2d, covmat_2d, p_2d, X_test_2d_active)

train_error_2d = error_rate(y_train, y_train_pred_2d)
test_error_2d = error_rate(y_test, y_test_pred_2d)

# Predict using PCA features
X_train_pca_active = X_train_pca[:, active_indices_pca]
X_test_pca_active = X_test_pca[:, active_indices_pca]

y_train_pred_pca = predict_lda(mu_pca, covmat_pca, p_pca, X_train_pca_active)
y_test_pred_pca = predict_lda(mu_pca, covmat_pca, p_pca, X_test_pca_active)

train_error_pca = error_rate(y_train, y_train_pred_pca)
test_error_pca = error_rate(y_test, y_test_pred_pca)
```

```
# Predict using full feature set
X_train_full_active = X_train[:, active_indices_full]
X_test_full_active = X_test[:, active_indices_full]

y_train_pred_full = predict_lda(mu_full, covmat_full, p_full, X_train_full_active)
y_test_pred_full = predict_lda(mu_full, covmat_full, p_full, X_test_full_active)

train_error_full = error_rate(y_train, y_train_pred_full)
test_error_full = error_rate(y_test, y_test_pred_full)

print("\nError rates:")
print(f"Custom 2D features - Train: {train_error_2d:.4f}, Test: {test_error_2d:.4f}")
print(f"PCA features - Train: {train_error_pca:.4f}, Test: {test_error_pca:.4f}")
print(f"Full features - Train: {train_error_full:.4f}, Test: {test_error_full:.4f}")

Error rates:
Custom 2D features - Train: 0.0968, Test: 0.1096
PCA features - Train: 0.0369, Test: 0.0616
Full features - Train: 0.0046, Test: 0.0274
```

Task 7

In [11]:

```
# TODO: For your hand-crafted embedding, visualize the decision boundary of
def plot_decision_regions(X, y, mu, covmat, p, active_indices):
    # Define grid boundaries
    x_min, x_max = X[:, 0].min() - 1, X[:, 0].max() + 1
    y_min, y_max = X[:, 1].min() - 1, X[:, 1].max() + 1

    # Create a meshgrid
    xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.1),
                         np.arange(y_min, y_max, 0.1))

    # Flatten the grid points
    grid_points = np.c_[xx.ravel(), yy.ravel()]

    # Filter grid points based on active indices
    grid_points_active = grid_points[:, active_indices]

    # Predict classes for grid points
    Z = predict_lda(mu, covmat, p, grid_points_active)
    Z = Z.reshape(xx.shape)

    # Plot decision regions
    plt.figure(figsize=(10, 8))
    plt.contourf(xx, yy, Z, alpha=0.3, cmap=plt.cm.coolwarm)

    # Plot scatter plot of training data
    for label, marker, color in zip([-1, 1], ['o', 'x'], ['blue', 'red']):
        plt.scatter(
            X[y == label, 0],
            X[y == label, 1],
            marker=marker,
            color=color,
            alpha=0.7,
            edgecolors='k',
            label=f'Digit {"3" if label == -1 else "9"}'
        )

    # Plot decision boundary
    plt.contour(xx, yy, Z, [0], linewidths=2, colors='black')
```

```

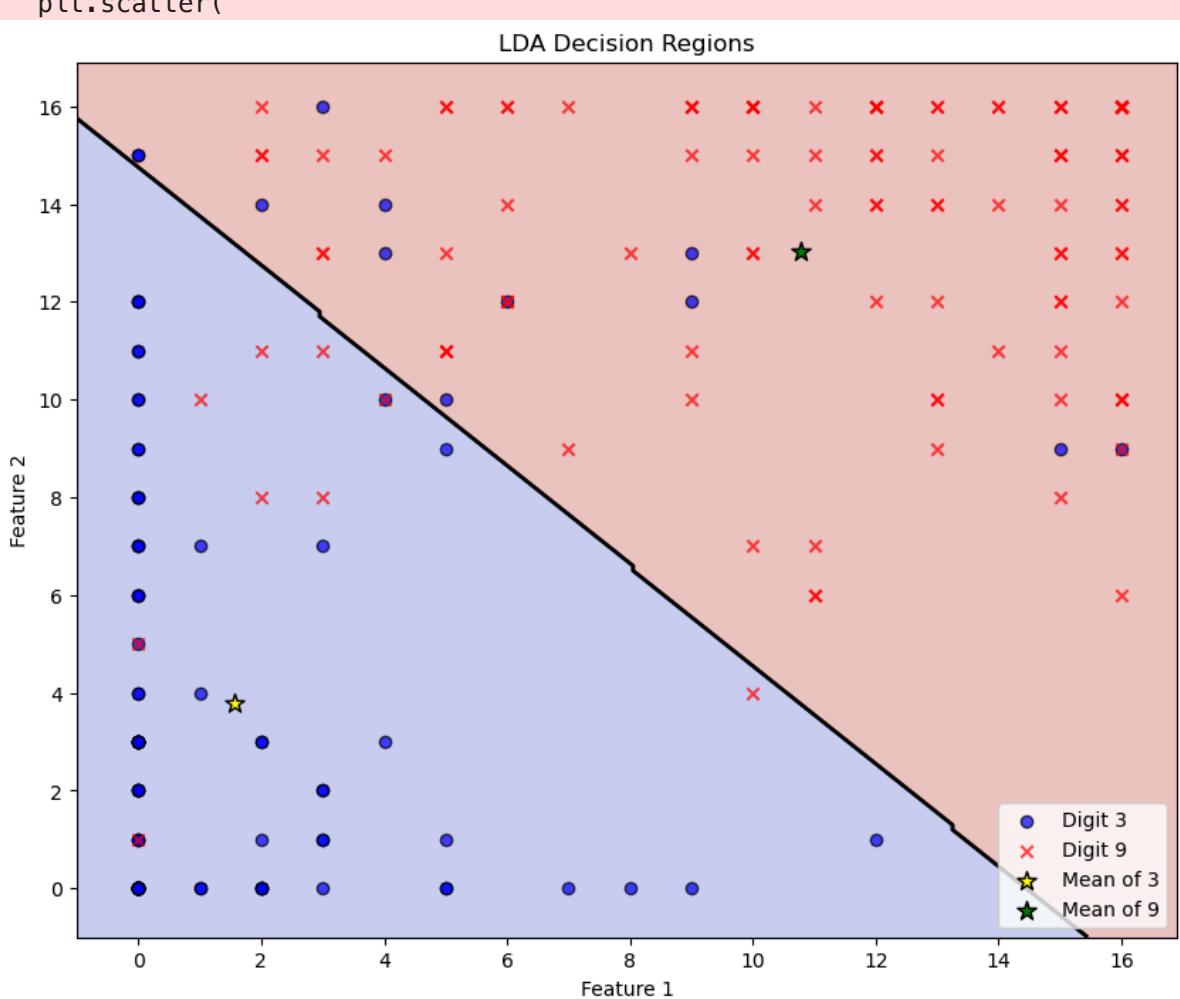
# Plot class means
for k in mu:
    if len(active_indices) > 0:
        plt.scatter(
            mu[k][0] if len(mu[k]) > 0 else 0,
            mu[k][1] if len(mu[k]) > 1 else 0,
            s=100,
            marker='*',
            color='yellow' if k == -1 else 'green',
            edgecolors='k',
            label=f'Mean of {"3" if k == -1 else "9"}'
        )

plt.title('LDA Decision Regions')
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
plt.legend()
plt.show()

# Plot decision regions for custom 2D features
plot_decision_regions(X_train_2d, y_train, mu_2d, covmat_2d, p_2d, active_ir

```

/var/folders/gl/m711nqcx42d81f7zr656_v600000gn/T/ipykernel_28106/241969234
4.py:27: UserWarning: You passed a edgecolor/edgecolors ('k') for an unfill
ed marker ('x'). Matplotlib is ignoring the edgecolor in favor of the face
color. This behavior may change in the future.



Task 8

```
In [12]: def cross_validate_lda(X, y, n_folds=10):
    # Filter out dead pixels for the full dataset
    variances = np.var(X, axis=0)
    active_indices = np.where(variances > 0.001)[0]
    X_active = X[:, active_indices]

    # Initialize cross-validation
    kf = KFold(n_splits=n_folds, shuffle=True, random_state=42)
    errors = []

    # Perform cross-validation
    for train_idx, test_idx in kf.split(X_active):
        X_train_fold, X_test_fold = X_active[train_idx], X_active[test_idx]
        y_train_fold, y_test_fold = y[train_idx], y[test_idx]

        # Fit LDA
        mu, covmat, p, active_indices = fit_lda(X_train_fold, y_train_fold)

        # Predict
        # Ensure test data uses the same active features as training data
        X_test_fold_active = X_test_fold[:, active_indices]
        y_pred = predict_lda(mu, covmat, p, X_test_fold_active)

        # Calculate error
        error = error_rate(y_test_fold, y_pred)
        errors.append(error)

    # Calculate mean and standard error
    mean_error = np.mean(errors)
    std_error = np.std(errors) / np.sqrt(n_folds)

    return mean_error, std_error, errors

# Filter digits data to include only 3 and 9
digits_data = load_digits()
X_all, y_all = digits_data.data, digits_data.target
mask_3_9 = np.logical_or(y_all == 3, y_all == 9)
X_3_9 = X_all[mask_3_9]
y_3_9 = np.where(y_all[mask_3_9] == 3, -1, 1)

# Perform 10-fold cross-validation
cv_error, cv_std, all_errors = cross_validate_lda(X_3_9, y_3_9, n_folds=10)

print("\nCross-validation results:")
print(f"Mean error: {cv_error:.4f} ± {cv_std:.4f}")
print(f"Individual fold errors: {all_errors}")
print(f"Comparison with single train/test split error: {test_error_full:.4f}
```

Cross-validation results:
 Mean error: 0.0360 ± 0.0233
 Individual fold errors: [np.float64(0.02702702702702703), np.float64(0.0), np.float64(0.0), np.float64(0.02777777777777776), np.float64(0.0), np.float64(0.0555555555555555), np.float64(0.0), np.float64(0.0), np.float64(0.0), np.float64(0.25)]
 Comparison with single train/test split error: 0.0274

```
In [ ]:
```

Exercise 3: Statistical Darts

```
In [13]: import numpy as np
import matplotlib.pyplot as plt
from matplotlib.patches import Ellipse
```

Task 1: Simulate Dart Throws

```
In [14]: def simulate_data(mu_true, Sigma_true, n_samples):
    return np.random.multivariate_normal(mu_true, Sigma_true, n_samples)
```

Task 2: Compute MLE

```
In [15]: def compute_mle(data):
    return np.mean(data, axis=0)
```

Task 3: Compute Posterior and MAP

```
In [16]: def compute_posterior(data, prior, Sigma_true):
    mu_0, Sigma_0 = prior
    N = data.shape[0]
    Sigma_true_inv = np.linalg.inv(Sigma_true)
    Sigma_0_inv = np.linalg.inv(Sigma_0)

    Sigma_post = np.linalg.inv(Sigma_0_inv + N * Sigma_true_inv)
    mu_mle = compute_mle(data)
    mu_post = Sigma_post @ (Sigma_0_inv @ mu_0 + N * Sigma_true_inv @ mu_mle)
    return mu_post, Sigma_post

def compute_map(data, prior, Sigma_true):
    mu_post, _ = compute_posterior(data, prior, Sigma_true)
    return mu_post
```

Task 4: Visualization

```
In [17]: def plot_cov_ellipse(mean, cov, ax, n_std=2.0, facecolor='none', edgecolor='black',
    vals, vecs = np.linalg.eigh(cov)
    order = vals.argsort()[:-1]
    vals, vecs = vals[order], vecs[:, order]
    theta = np.degrees(np.arctan2(*vecs[:, 0][:-1]))
    width, height = 2 * n_std * np.sqrt(vals)
    ellipse = Ellipse(xy=mean, width=width, height=height, angle=theta,
                      facecolor=facecolor, edgecolor=edgecolor, lw=2, label='Posterior')
    ax.add_patch(ellipse)

def visualize_inference(mu_true, mu_mle, mu_map, mu_post, Sigma_post, data):
    fig, ax = plt.subplots(figsize=(8, 8))
    ax.scatter(data[:, 0], data[:, 1], color='gray', s=60, label='Throws')
    ax.scatter(*mu_true, color='green', marker='*', s=200, label='True aim')
    plot_cov_ellipse(mu_true, Sigma_true, ax, n_std=2.0, facecolor='red', edgecolor='red', label='True aim')
    plot_cov_ellipse(mu_mle, Sigma_mle, ax, n_std=2.0, facecolor='blue', edgecolor='blue', label='MLE')
    plot_cov_ellipse(mu_map, Sigma_map, ax, n_std=2.0, facecolor='orange', edgecolor='orange', label='MAP')
    plot_cov_ellipse(mu_post, Sigma_post, ax, n_std=2.0, facecolor='green', edgecolor='green', label='Posterior')
```

```

ax.scatter(*mu_mle, color='blue', marker='x', s=100, label='MLE')
ax.scatter(*mu_map, color='red', marker='D', s=80, label='MAP')
ax.scatter(*mu_post, color='orange', marker='o', s=100, label='Posterior')
plot_cov_ellipse(mu_post, Sigma_post, ax, n_std=2, edgecolor='orange', 1
ax.set_title("Dart Throw Inference Visualization", fontsize=14)
ax.set_xlabel("x")
ax.set_ylabel("y")
ax.grid(True)
ax.legend()
ax.set_aspect('equal')
ax.set_xlim(-1, 1)
ax.set_ylim(-1, 1)
plt.show()

```

```

In [18]: if __name__ == "__main__":
    np.random.seed(42)

    # True parameters
    mu_true = np.array([0.0, 0.5])
    Sigma_true = np.array([[0.05, 0.02],
                          [0.02, 0.04]])

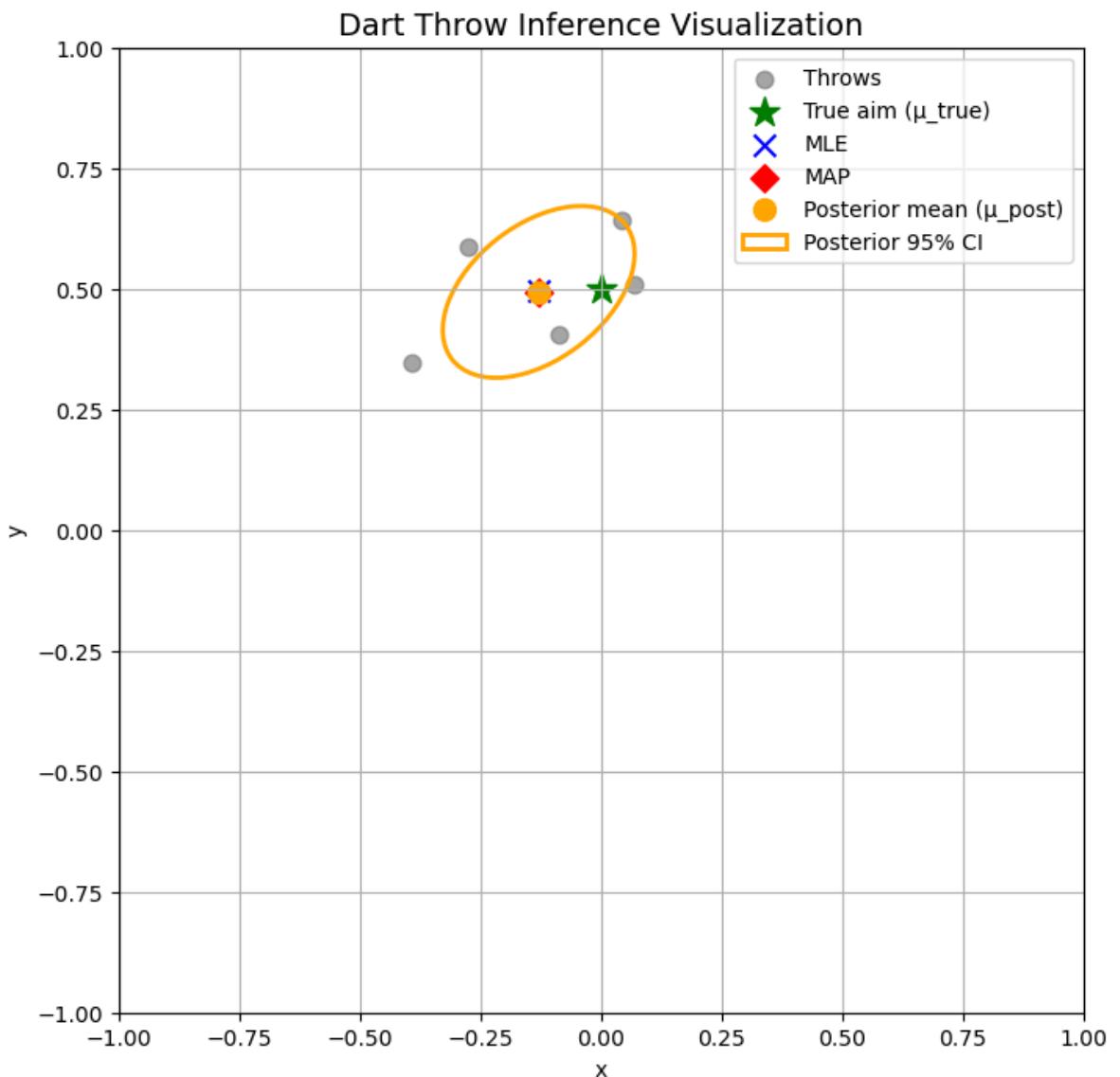
    # Prior (standard normal around bullseye)
    mu_0 = np.array([0.0, 0.0])
    Sigma_0 = np.eye(2)
    prior = (mu_0, Sigma_0)

    # Simulate data
    N = 5 # number of throws
    data = simulate_data(mu_true, Sigma_true, N)

    # Perform inference
    mu_mle = compute_mle(data)
    mu_post, Sigma_post = compute_posterior(data, prior, Sigma_true)
    mu_map = compute_map(data, prior, Sigma_true)

    # Visualize results
    visualize_inference(mu_true, mu_mle, mu_map, mu_post, Sigma_post, data)

```



Exercise 4

Sunday, 18 May 2025 10:32 PM

* Exercise 4 : Logistic Regression

- Task 1 :

For a single observation (x_i, y_i) , the Bernoulli model gives

$$P(y_i | x_i; \omega) = p_i^{y_i} (1-p_i)^{1-y_i}$$

where we will set $p_i = \sigma(\omega^T x_i)$

Joint likelihood for all N i.i.d obs.,

$$L(\omega) = \prod_{i=1}^N P(y_i | x_i; \omega) = \prod_{i=1}^N (p_i)^{y_i} (1-p_i)^{1-y_i}$$

Taking log,

$$\ell(\omega) = \log L(\omega) = \sum_{i=1}^N [y_i \log p_i + (1-y_i) \log (1-p_i)]$$

As $p_i = \sigma(\omega^T x_i)$

$$\ell(\omega) = \sum_{i=1}^N [y_i \log[\sigma(\omega^T x_i)] + (1-y_i) \log(1-\sigma(\omega^T x_i))]$$

This is log-likelihood for log. regres. on Bernoulli.

- Task 2 :

a] Convexity of negative log-likelihood

We know that log-likelihood is, its negative would be,

$$-\ell(\omega) = \sum_{i=1}^N [y_i \log[\sigma(\omega^T x_i)] - (1-y_i) \log(1-\sigma(\omega^T x_i))]$$

The grad. of each data pt. is,

$$\nabla_{\omega} [-\ell_i(\omega)] = [\sigma(\omega^T x_i) - y_i] x_i$$

Diff. again & we get hessian,

$$\nabla^2 [-\ell(\omega)] = \sum_{i=1}^N \sigma(\omega^T x_i) [1 - \sigma(\omega^T x_i)] x_i x_i^T$$

As $\sigma(\omega^T x_i) [1 - \sigma(\omega^T x_i)] \geq 0$ & $x_i x_i^T$ is positive semi-definite, their sum is also semi-definite.

$$\therefore \nabla^2 [-\ell(\omega)] \geq 0 \Rightarrow \ell(\omega) \text{ is convex.}$$

I believe convexity guarantees there are local minima.
 Any local min. is global. Thus, grad. descent will converge to global optimum (unique).

b) Binary cross-entropy loss :

We can see that,

$$\max_w l(w) \Leftrightarrow \min_w (-l(w)) \Leftrightarrow \min_w -\frac{1}{N} l(w)$$

Dividing by N turns negative log-likelihood into exactly the average binary cross-entropy:

$$\begin{aligned} -\frac{1}{N} l(w) &= -\frac{1}{N} \sum_{i=1}^N [y_i \log \sigma(w^T x_i) + (1-y_i) \log(1-\sigma(w^T x_i))] \\ &= L_{BCE}(w) \end{aligned}$$

Thus,

$$\arg \max_w l(w) = \arg \min_w L_{BCE}(w)$$

showing that max.-likelihood estimation under Bernoulli GLM is equal. to minimizing the binary cross-entropy loss.

* Task 3 :

a) Divergence of the MLE under linear separability.

The log-likelihood for $\{(x_i, y_i)\}_{i=1}^N$,

$$l(w) = \sum_{i=1}^N [y_i \log \sigma(w^T x_i) + (1-y_i) \log(1-\sigma(w^T x_i))]$$

$$\text{where } \sigma(z) = 1/(1+e^{-z})$$

By assumption there exists some weight w' s.t.,

$$w'^T x_i \begin{cases} > 0 & \text{if } y_i = 1, \\ < 0 & \text{if } y_i = 0 \end{cases}$$

Let's take seq. $w_k = k w' \quad - [k=1, 2, \dots]$

Then for each i ,

- if $y_i = 1$, $\omega_k^T \mathbf{x}_i = k (\omega^T \mathbf{x}_i) \rightarrow +\infty$ so
 $\sigma(\omega_k^T \mathbf{x}_i) \rightarrow 1$ & $\log \sigma(\omega_k^T \mathbf{x}_i) \rightarrow 0$
- If $y=0$, $\omega_k^T \mathbf{x}_i = k (\omega^T \mathbf{x}_i) \rightarrow -\infty$ so
 $\sigma(\omega_k^T \mathbf{x}_i) \rightarrow 0$ & $\log(1 - \sigma(\omega_k^T \mathbf{x}_i)) \rightarrow 0$
- ∴ Every term in $\ell(\omega_k)$ approaches zero &

$$\ell(\omega_k) \rightarrow 0 \text{ as } k \rightarrow \infty$$

For finite ω , $\ell(\omega) < 0$. Thus supremum of log-likelihood is 0, and it attained only in the limit $\|\omega\| \rightarrow \infty$

No finite max. exists, ∴ MLE fails to converge under linear separability.

b) Mitigation:

In order to obtain optimization with a finite sol'n one typically penalizes large weights.

Introducing L_2 or L_1 penalty to neg. log likelihood.

$$\min_{\omega} [-\ell(\omega) + \lambda \|\omega\|_2^2] \text{ or } \min_{\omega} [-\ell(\omega) + \lambda \|\omega\|_1]$$

Also if we place a gaussian (for L_2) or Laplace (L_1) prior on ω yields a MAP estimate that remains finite even when the data are linearly separable. Regularization prevents weight mag. from diverging & often improves generalization.