

* Exercise 1: Decision Boundaries of Generative Classifiers

- Task 1: Gaussian Discriminant Analysis

(a) QDA:

We assume $A \not\perp B$ 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)

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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)
)

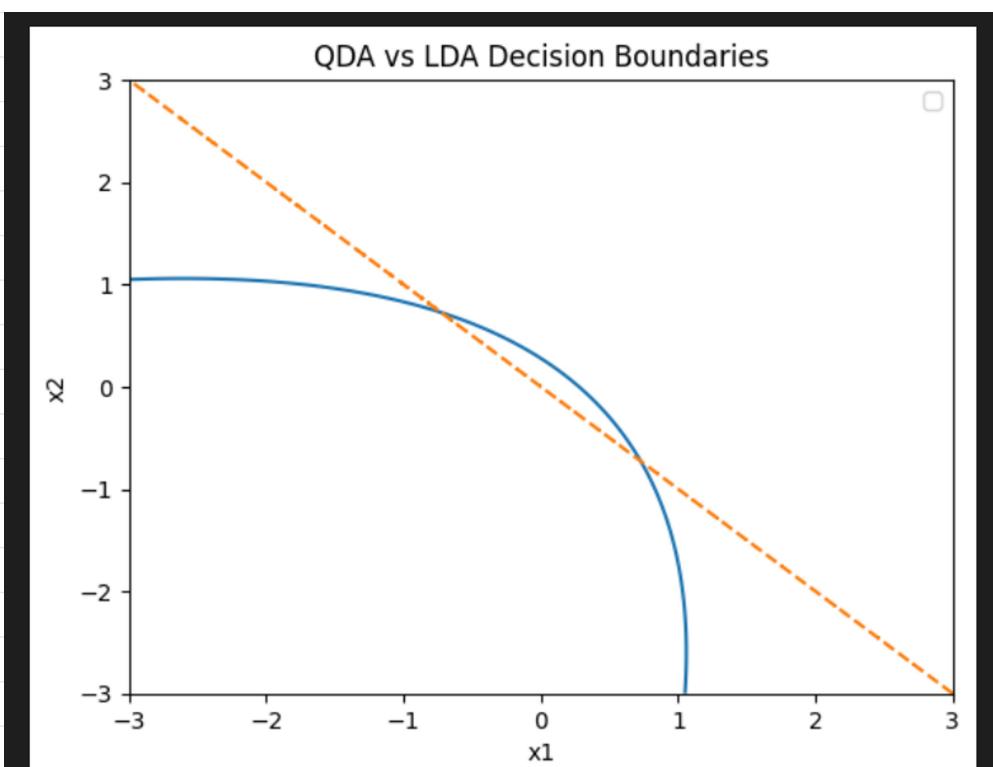
# LDA parameters
Sigma_pooled = 0.5 * (SigmaA + SigmaB)
inv_pooled = np.linalg.inv(Sigma_pooled)
w_lda = inv_pooled.dot(muA - muB)
b_lda = (
    -0.5 * muA.T.dot(inv_pooled).dot(muA) + 0.5 * muB.T.dot(inv_pooled).dot(muB)
    + np.log(piA / piB)
)

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")

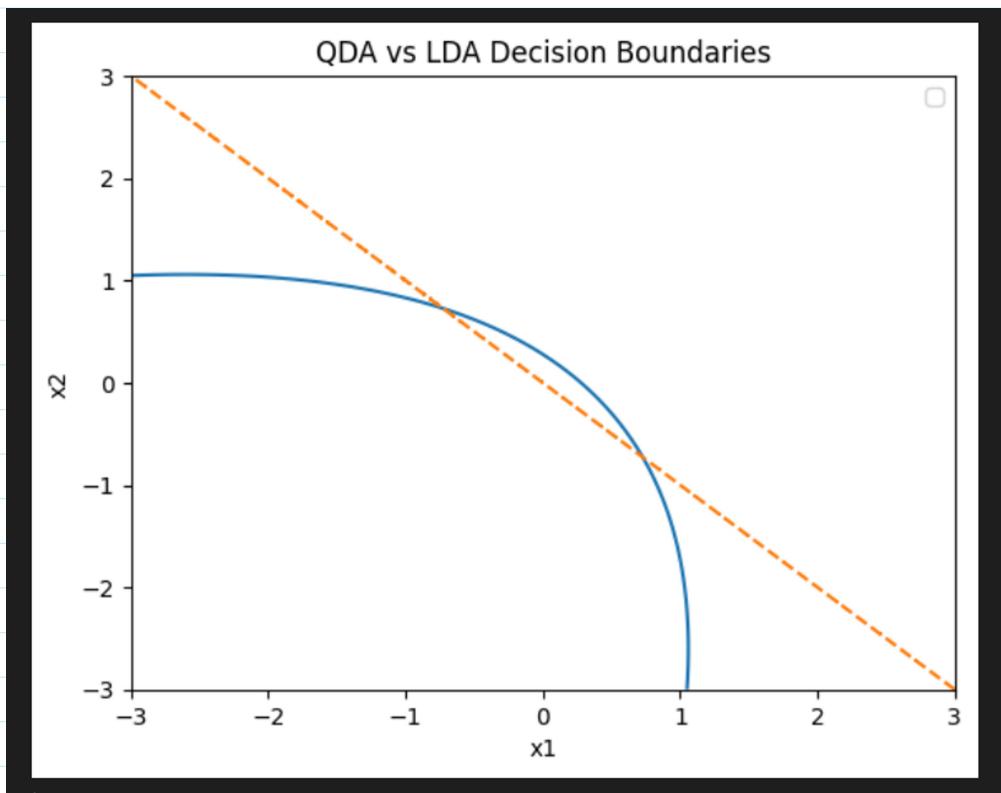
```

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