

Sheet 5

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In [1]: import numpy as np
        from matplotlib import pyplot as plt
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1 Bayes: Signal or Noise?

$$p(\text{gamma ray} \mid \text{target direction}) = \frac{p(\text{target direction} \mid \text{gamma ray}) \cdot p(\text{gamma ray})}{p(\text{background})} = \frac{0.95 \cdot 0.1}{0.9} \approx 0.11$$

2 Bayes Classifier

a)

$$L(y, \hat{y}) = \begin{bmatrix} 0 & 1 \\ 10 & 0 \end{bmatrix}$$

where:

- $y = 0$ is the true class, and $\hat{y} = 0$ or $\hat{y} = 1$ are the predicted classes.
- $y = 1$ is the true class, and $\hat{y} = 0$ or $\hat{y} = 1$ are the predicted classes.

With $P(y = 1|x) = p(x)$ and $P(y = 0|x) = 1 - p(x)$ we calculate the conditional risk $R(\hat{y}|x)$ for each possible predicted label \hat{y} :

1. **For $\hat{y} = 0$:**

$$R(0|x) = L(0,0) \cdot P(y = 0|x) + L(1,0) \cdot P(y = 1|x) = 0 \cdot (1 - p(x)) + 10 \cdot p(x) = 10 \cdot p(x)$$

2. **For $\hat{y} = 1$:**

$$R(1|x) = L(0, 1) \cdot P(y = 0|x) + L(1, 1) \cdot P(y = 1|x) = 1 \cdot (1 - p(x)) + 0 \cdot p(x) = 1 - p(x)$$

To minimize the conditional risk, choose $\hat{y} = 0$ if $R(0|x) < R(1|x)$, and $\hat{y} = 1$ otherwise.

Thus, we need:

$$10 \cdot p(x) < 1 - p(x)$$

Rearrange this inequality:

$$11 \cdot p(x) < 1 \quad \Rightarrow \quad p(x) < \frac{1}{11}$$

The optimal Bayes classifier is:

$$\hat{y} = \begin{cases} 0 & \text{if } p(x) < \frac{1}{11} \\ 1 & \text{if } p(x) \geq \frac{1}{11} \end{cases}$$

This asymmetric loss matrix is appropriate in situations where the consequences of a false negative (predicting $y = 0$ when $y = 1$) are much more severe than a false positive (predicting $y = 1$ when $y = 0$).

b)

The Bayes classifier minimizes the **expected loss** by choosing the prediction \hat{y} that minimizes the **conditional risk** $R(\hat{y}|x)$ given x . For each potential prediction $\hat{y} \in \{0, 1, \dots, k\}$, we calculate $R(\hat{y}|x)$.

1. **Conditional Risk for $\hat{y} = j$, where $j \in \{1, \dots, k\}$:** If we predict class j , the conditional risk is:

$$R(j|x) = \sum_{i=1}^k L(i, j) \cdot P(y = i|x)$$

$$R(j|x) = \sum_{\substack{i=1 \\ i \neq j}}^k P(y = i|x) = 1 - P(y = j|x)$$

2. **Conditional Risk for the "Reject" Option $\hat{y} = 0$:** If we choose to reject, the conditional risk is:

$$R(0|x) = \sum_{i=1}^k L(i, 0) \cdot P(y = i|x) = \alpha \sum_{i=1}^k P(y = i|x) = \alpha$$

since the probabilities $P(y = i|x)$ sum to 1.

To minimize the conditional risk, we compare $R(j|x)$ for $j \in \{1, \dots, k\}$ with $R(0|x) = \alpha$.

- For each class j , we will choose $\hat{y} = j$ if:

$$R(j|x) < \alpha \quad \Rightarrow \quad 1 - P(y = j|x) < \alpha \quad \Rightarrow \quad P(y = j|x) > 1 - \alpha$$

- If no class j satisfies $P(y = j|x) > 1 - \alpha$, we choose the "reject" option, $\hat{y} = 0$.

Thus, the optimal Bayes classifier is:

$$\hat{y} = \begin{cases} j & \text{if } P(y = j|x) > 1 - \alpha \text{ for some } j \in \{1, \dots, k\} \\ 0 & \text{if } P(y = j|x) \leq 1 - \alpha \text{ for all } j \in \{1, \dots, k\} \end{cases}$$

The parameter α controls the threshold for rejection:

- A **lower** α means that we are less willing to reject; we require a higher confidence in our prediction before choosing to classify. As $\alpha \rightarrow 0$, the classifier rarely chooses the "reject" option.
- A **higher** α increases the likelihood of rejection, as the threshold $1 - \alpha$ becomes lower, requiring less certainty to avoid rejection.

This approach adds flexibility, allowing for rejection when the classifier's confidence is insufficient, thus potentially reducing overall decision-making risk.

3 QDA

(a)

```
In [24]: pts = np.load('data/data1d.npy')
labels = np.load('data/labels1d.npy')

x0, x1 = pts[labels == 0], pts[labels == 1]
plt.hist(x0, bins=10, alpha=0.5, density=True, color='r')
plt.hist(x1, bins=10, alpha=0.5, density=True, color='b')

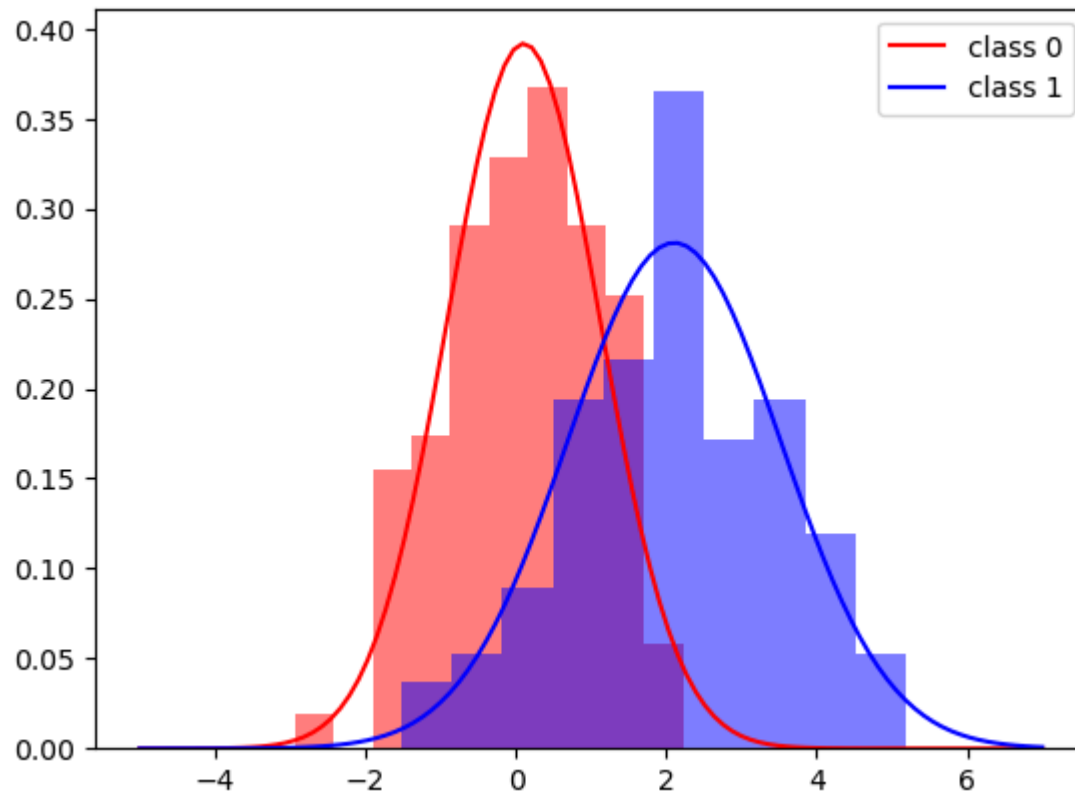
mean0, mean1 = np.mean(x0), np.mean(x1)
var0, var1 = np.var(x0), np.var(x1)

def gaussian(x, mean, var):
    return np.exp(-0.5*(x-mean)**2/var)/np.sqrt(2*np.pi*var)

x = np.linspace(-5, 7, 100)

plt.plot(x, gaussian(x, mean0, var0), label='class 0', color='r')
plt.plot(x, gaussian(x, mean1, var1), label='class 1', color='b')
plt.legend()
```

```
Out[24]: <matplotlib.legend.Legend at 0x23470f65d00>
```



(b)

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In [26]: x = np.linspace(-10, 10, 100)

like0 = gaussian(x, mean0, var0)
like1 = gaussian(x, mean1, var1)

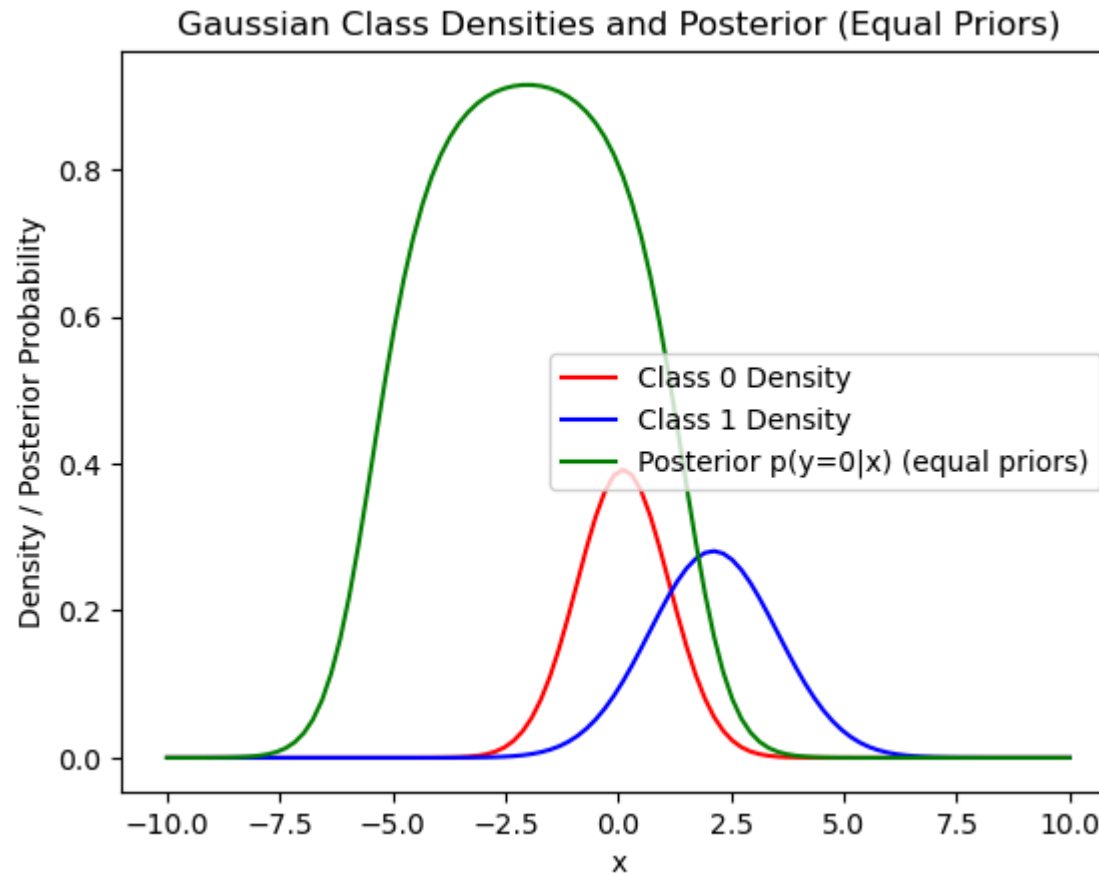
prior0, prior1 = 0.5, 0.5

post0 = like0*prior0/(like0*prior0 + like1*prior1)

plt.plot(x, like0, label="Class 0 Density", color="r")
plt.plot(x, like1, label="Class 1 Density", color="b")
plt.plot(x, post0, label="Posterior p(y=0|x) (equal priors)", color="green")
```

```
plt.title("Gaussian Class Densities and Posterior (Equal Priors)")
plt.xlabel("x")
plt.ylabel("Density / Posterior Probability")
plt.legend()
```

Out[26]: <matplotlib.legend.Legend at 0x23470f833a0>



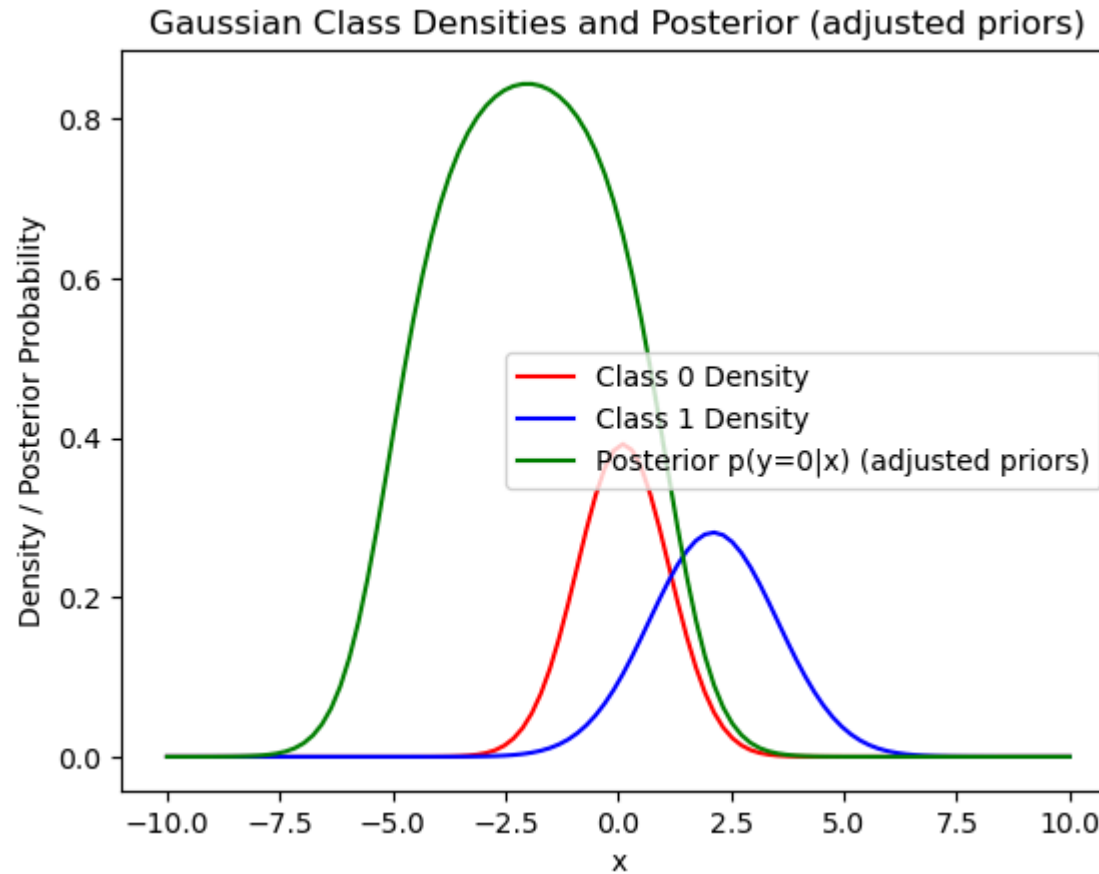
```
In [28]: prior0, prior1 = 1/3, 2/3

post0 = like0*prior0/(like0*prior0 + like1*prior1)

plt.plot(x, like0, label="Class 0 Density", color="r")
plt.plot(x, like1, label="Class 1 Density", color="b")
```

```
plt.plot(x, post0, label="Posterior p(y=0|x) (adjusted priors)", color="green")  
plt.title("Gaussian Class Densities and Posterior (adjusted priors)")  
plt.xlabel("x")  
plt.ylabel("Density / Posterior Probability")  
plt.legend()
```

Out[28]: <matplotlib.legend.Legend at 0x2346f84fe80>



4 Trees and Random Forests

a)

$$\text{Misclassification Rate} = 1 - \max(p(y = c))$$

$$\text{Gini} = 1 - \sum_{c=1}^2 p(y = c)^2$$

$$\text{Entropy} = - \sum_{c=1}^2 p(y = c) \log_2 p(y = c)$$

1. Node 1 (300, 100):

$$\text{Misclassification Rate} = 1 - 0.75 = 0.25$$

$$\text{Gini Impurity} = 1 - (0.75^2 + 0.25^2) = 1 - (0.5625 + 0.0625) = 0.375$$

$$\text{Entropy} = -(0.75 \log_2 0.75 + 0.25 \log_2 0.25) \approx 0.811$$

2. Node 2 (100, 300):

$$\text{Misclassification Rate} = 1 - 0.75 = 0.25$$

$$\text{Gini Impurity} = 1 - (0.25^2 + 0.75^2) = 0.375$$

$$\text{Entropy} = -(0.25 \log_2 0.25 + 0.75 \log_2 0.75) \approx 0.811$$

3. Weighted Averages for Split A:

$$\text{Misclassification Rate} = 0.25 \times \frac{400}{800} + 0.25 \times \frac{400}{800} = 0.25$$

$$\text{Gini Impurity} = 0.375 \times \frac{400}{800} + 0.375 \times \frac{400}{800} = 0.375$$

$$\text{Entropy} = 0.811 \times \frac{400}{800} + 0.811 \times \frac{400}{800} \approx 0.811$$

4. Node 1 (200, 0):

$$\text{Misclassification Rate} = 1 - 1 = 0$$

$$\text{Gini Impurity} = 1 - (1^2 + 0^2) = 0$$

$$\text{Entropy} = -(1 \cdot \log_2 1 + 0 \cdot \log_2 0) = 0$$

5. Node 2 (200, 400):

$$\text{Misclassification Rate} = 1 - 0.667 \approx 0.333$$

$$\text{Gini Impurity} = 1 - (0.333^2 + 0.667^2) = 1 - (0.111 + 0.444) = 0.444$$

$$\text{Entropy} = -(0.333 \log_2 0.333 + 0.667 \log_2 0.667) \approx 0.918$$

6. Weighted Averages for Split B:

$$\text{Misclassification Rate} = 0 \times \frac{200}{800} + 0.333 \times \frac{600}{800} \approx 0.25$$

$$\text{Gini Impurity} = 0 \times \frac{200}{800} + 0.444 \times \frac{600}{800} = 0.333$$

$$\text{Entropy} = 0 \times \frac{200}{800} + 0.918 \times \frac{600}{800} \approx 0.688$$

- Misclassification Rate: Both splits have the same misclassification rate of 0.25.
- Gini Impurity: Split B has a lower Gini impurity (0.333 vs. 0.375), so it would be preferred by the Gini criterion.
- Entropy: Split B also has a lower entropy (0.688 vs. 0.811), so it would be preferred by the entropy criterion.

Overall, Split B would be the preferred split based on both Gini impurity and entropy.

(b)

```
In [ ]: # Load the data
pts = np.load('data/data1d.npy')
labels = np.load('data/labels1d.npy')

# TODO: Sort the points to easily split them

sorted_pts = np.sort(pts)
sorted_labels = labels[np.argsort(pts)]

# TODO: Implement or find implementation for Gini impurity, entropy and misclassification rate

def probabilities(partition):
    # divide counts by size of dataset to get cluster probabilities
    return np.unique(partition, return_counts=True)[1] / len(partition)
```

```

def gini_impurity(partition):
    p = probabilities(partition)
    return 1 - np.sum(p**2)

def entropy(partition):
    p = probabilities(partition)
    return -np.sum(p * np.log2(p))

def misclassification_rate(partition):
    p = probabilities(partition)
    return 1 - np.max(p)

def compute_split_measure(l, l0, l1, method):
    p0 = probabilities(l0)
    p1 = probabilities(l1)
    p = probabilities(l)
    return method(p) - (len(l0) * method(p0) + len(l1) * method(p1)) / (len(l))

# TODO: Iterate over the possible splits, evaluating and saving the three criteria for each one

# TODO: Then, Compute the split that each criterion favours and visualize them
#       (e.g. with a histogram for each class and vertical lines to show the splits)

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In []:

(c)

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In [7]: # Load the dijet data
features = np.load('data/dijet_features_normalized.npy')
labels = np.load('data/dijet_labels.npy')

# TODO: define train, val and test splits as specified (make sure to shuffle the data before splitting it!)

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In [8]: from sklearn.ensemble import RandomForestClassifier

# TODO: train a random forest classifier for each combination of specified hyperparameters
#       and evaluate the performances on the validation set.

```

```
In [9]: # TODO: for your preferred configuration, evaluate the performance of the best configuration on the test set
```