June 1, 2023

[1]: import numpy as np

```
import matplotlib.pyplot as plt
     from mpl_toolkits.mplot3d import Axes3D
     from scipy.stats import multivariate_normal
     %matplotlib widget
[2]: # we're working with 3-dimensional data
     # this 3-D random vector X takes values from a mixture of four Guassians
     # the first gaussian is the class-conditional pdf for class 1, and antoher is \Box
      ⇔for class 2
     # the remaining 2 gaussian components are the class-conditional pdfs for class_{\sqcup}
      →3 with equal weights
     # the class priors are p1 = 0.3, p2 = 0.3, p3 = 0.4
     # we want to set the distances between the means of pairs of gaussians to twice_
     → the average standard deviation
     # set up the parameters of the class-conditioned Gaussian pdfs
     # the distances between the means of pairs of gaussians are set to twice the
      →average standard deviation
     sigma_1 = np.eye(3)
     sigma_2 = np.eye(3)
     sigma_3 = np.eye(3)
     sigma_4 = np.eye(3)
     avg_std = np.sqrt(np.trace(sigma_1) + np.trace(sigma_2) + np.trace(sigma_3) +__
      →np.trace(sigma_4)) / 4
    mult = 1.5 * avg_std
    mu_1 = np.array([mult, mult, mult])
     mu_2 = np.array([0, mult, mult])
     mu_3 = np.array([mult, 0, mult])
     mu_4 = np.array([mult, mult, 0])
     # set up the parameters of the class priors
     p_1 = 0.3
```

```
The distance between mu_1 and mu_2 is 1.299038105676658
The distance between mu_2 and mu_3 is 1.8371173070873836
The distance between mu_3 and mu_4 is 1.8371173070873836
The distance between mu_4 and mu_1 is 1.299038105676658
The distance between mu_1 and mu_3 is 1.299038105676658
The distance between mu_2 and mu_4 is 1.8371173070873836
Twice the average standard deviation is 1.7320508075688772
```

0.0.1 Part A

1. Generate 10000 samples from this data distribution and keep track of the true class labels

```
[3]: # generate 10000 samples from this data distribution and keep track of the true
      ⇔class labels
     samples = 10000
     gaussian_1 = np.random.multivariate_normal(mu_1, sigma_1, samples)
     gaussian_2 = np.random.multivariate_normal(mu_2, sigma_2, samples)
     gaussian_3 = np.random.multivariate_normal(mu_3, sigma_3, samples)
     gaussian_4 = np.random.multivariate_normal(mu_4, sigma_4, samples)
     data = []
     labels = []
     # use the class priors to generate the labels
     for i in range(samples):
         # class 1
         if np.random.rand() < p_1:</pre>
             data.append(gaussian_1[i])
             labels.append(1)
         # class 2
         elif np.random.rand() < p_1 + p_2:</pre>
             data.append(gaussian_2[i])
             labels.append(2)
         # class 3
         elif np.random.rand() < p_1 + p_2 + p_3:
```

```
# class 3 data originates from a mixture of gaussians 3 and 4 with_
equal weights
if np.random.rand() < 0.5:
          data.append(gaussian_3[i])
          labels.append(3)
else:
          data.append(gaussian_4[i])
          labels.append(3)

# convert the data and labels to numpy arrays
data = np.array(data)
labels = np.array(labels)</pre>
```

2. Specify the decision rule that acheives minimum probability of error (use 0-1 loss). Implement this classifier with true data distribution knowledge. Classify the 10k samples and count the samples corresponding to each decision-label pair to empirically estimate the confusion matrix.

```
[4]: # likelihood estimation to use in the decision rule
     # the likelihood of a sample x is the probability of x given the class label
     # the likelihood of x given class 1 is the pdf of the multivariate qaussian_{\sqcup}
      ⇒with mean mu_1 and covariance sigma_1 evaluated at x
     # the likelihood of x given class 2 is the pdf of the multivariate gaussian
      \rightarrowwith mean mu_2 and covariance sigma_2 evaluated at x
     # the likelihood of x given class 3 is the pdf of the multivariate gaussians,
      with means mu 3, mu 4 and covariances sigma 3, sigma 4 equally weighted
      \rightarrowevaluated at x
     def likelihood(x: np.array, label: int) -> float:
         Given a sample x and a class label, return the likelihood of x given the
      ⇔class label
         Args:
             x (np.array): a sample
             label (int): a class label
             ValueError: if the label is not 1, 2, or 3
         Returns:
             float: the likelihood of x given the class label
         .....
         if label == 1:
             return multivariate_normal.pdf(x, mu_1, sigma_1)
         elif label == 2:
             return multivariate_normal.pdf(x, mu_2, sigma_2)
         elif label == 3:
```

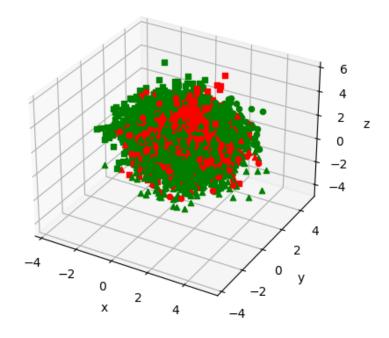
```
return 0.5 * multivariate_normal.pdf(x, mu_3, sigma_3) + 0.5 *_
multivariate_normal.pdf(x, mu_4, sigma_4)
else:
    raise ValueError("Invalid label")
```

```
[5]: # the decision rule is to choose the class label that maximizes the likelihood
     ⇔of the sample
     # the decision rule is implemented with true data distribution knowledge
     # the decision rule is implemented with true class priors
     def minimum probability error rule(x: np.array, loss: np.array) -> int:
         Given a sample x and a loss matrix, return the class label that minimizes \Box
      ⇔the probability of error
         Arqs:
             x (np.array): a sample
             loss (np.array): the loss matrix
         Returns:
             int: the class label that minimizes the probability of error
         # compute the likelihoods of x given each class label
         likelihoods = [likelihood(x, 1), likelihood(x, 2), likelihood(x, 3)]
         # calculate the posterior probabilities of x given each class label
         # the posterior probability of x given class 1 is the likelihood of x given \Box
      ⇔class 1 times the class prior of class 1
         # the posterior probability of x given class 2 is the likelihood of x given_{\sqcup}
      ⇔class 2 times the class prior of class 2
         # the posterior probability of x given class 3 is the likelihood of x given
      ⇔class 3 times the class prior of class 3
         posterior_probs = [likelihoods[0] * p_1, likelihoods[1] * p_2,__
      →likelihoods[2] * p_3]
         # calculate the expected loss of each class label
         loss_1 = posterior_probs[0] * loss[0][0] + posterior_probs[1] * loss[0][1]_u
      →+ posterior_probs[2] * loss[0][2]
         loss_2 = posterior_probs[0] * loss[1][0] + posterior_probs[1] * loss[1][1]_u
      →+ posterior_probs[2] * loss[1][2]
         loss_3 = posterior_probs[0] * loss[2][0] + posterior_probs[1] * loss[2][1]_u
      →+ posterior_probs[2] * loss[2][2]
         # return the class label that minimizes the expected loss
         return np.argmin([loss_1, loss_2, loss_3]) + 1
```

```
The confusion matrix is [[1356 610 1066] [ 669 2703 782] [ 456 325 2033]]
```

3. Provide a vizualization of the data (scatter plot in 3 dimensions). For each sample, indicate the true class label with a different marker and if it was correctly classified with a different color

```
[8]: # plot the data
     fig = plt.figure()
     ax = fig.add_subplot(111, projection='3d')
     # plot the samples
     for i in range(samples):
         # determine the color to use
         color = 'green' if labels[i] == minimum_probability_error_rule(data[i],_
      ⇔loss_1) else 'red'
         # determine the marker to use
         marker = 'o' if labels[i] == 1 else 's' if labels[i] == 2 else '^'
         # plot the sample
         ax.scatter(data[i][0], data[i][1], data[i][2], c=color, marker=marker)
     # label the axes
     ax.set xlabel('x')
     ax.set ylabel('y')
     ax.set_zlabel('z')
     # show the plot
     plt.show()
```

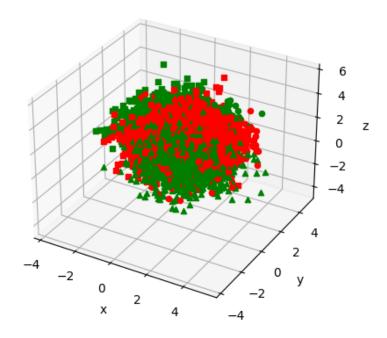


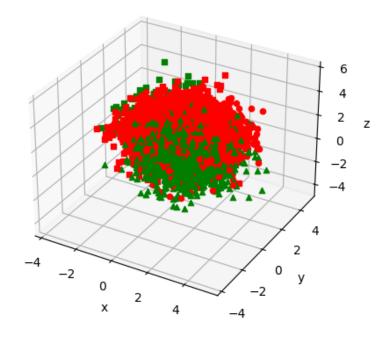
0.0.2 Part B

Repeat the exercise for the ERM classification rule with the following loss matrices. Using the same 10k samples, estimate the minimum expected risk that this optimal ERM classification rule will achieve.

```
print(confusion_matrix_10)
      confusion_matrix_100 = np.zeros((3, 3), dtype=int)
      for i in range(samples):
          guess = minimum_probability_error_rule(data[i], loss_100)
          confusion_matrix_100[labels[i] - 1][guess - 1] += 1
      # print the confusion matrix
      print("The confusion matrix for caring 100x more is")
      print(confusion_matrix_100)
     The confusion matrix for caring 10x more is
     [[ 36 102 2894]
      [ 29 909 3216]
              23 2791]]
     The confusion matrix for caring 100x more is
              4 3028]
      Γ
          0
              66 40881
      Γ
          0
              1 2813]]
[15]: # plot the data for the 10x loss
      fig = plt.figure()
      ax = fig.add_subplot(111, projection='3d')
      # plot the samples
      for i in range(samples):
          # determine the color to use
          color = 'green' if labels[i] == minimum_probability_error_rule(data[i],__
       ⇔loss_10) else 'red'
          # determine the marker to use
          marker = 'o' if labels[i] == 1 else 's' if labels[i] == 2 else '^'
          # plot the sample
          ax.scatter(data[i][0], data[i][1], data[i][2], c=color, marker=marker)
      # label the axes
      ax.set_xlabel('x')
      ax.set_ylabel('y')
      ax.set_zlabel('z')
      # show the plot
      plt.show()
      # plot the data for the 100x loss
      fig = plt.figure()
```

```
ax = fig.add_subplot(111, projection='3d')
# plot the samples
for i in range(samples):
   # determine the color to use
   color = 'green' if labels[i] == minimum_probability_error_rule(data[i],__
⇔loss_100) else 'red'
    # determine the marker to use
   marker = 'o' if labels[i] == 1 else 's' if labels[i] == 2 else '^'
   # plot the sample
   ax.scatter(data[i][0], data[i][1], data[i][2], c=color, marker=marker)
# label the axes
ax.set_xlabel('x')
ax.set_ylabel('y')
ax.set_zlabel('z')
# show the plot
plt.show()
```



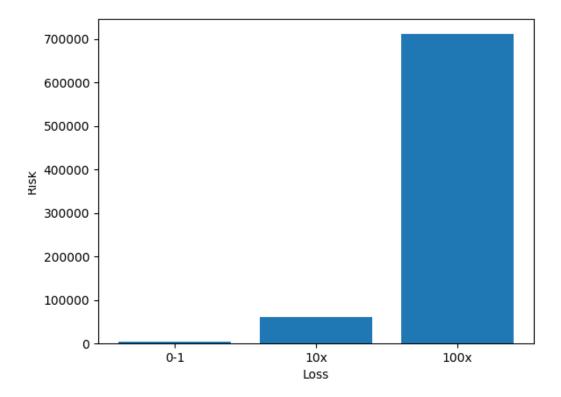


```
[14]: # estimate the minimum expected risk for each loss matrix
      risk_1 = 0
      risk_10 = 0
      risk_100 = 0
      \# use the confusion matrices to estimate the minimum expected risk
      for i in range(3):
          for j in range(3):
              risk_1 += confusion_matrix[i][j] * loss_1[i][j]
              risk_10 += confusion_matrix_10[i][j] * loss_10[i][j]
              risk_100 += confusion_matrix_100[i][j] * loss_100[i][j]
      # normalize the risks
      \# risk_1 /= samples
      \# risk_10 /= samples
      # risk_100 /= samples
      # print the risks
      print("The minimum expected risk for the 0-1 loss is", risk_1)
```

```
print("The minimum expected risk for the 10x loss is", risk_10)
print("The minimum expected risk for the 100x loss is", risk_100)

# plot the risks
fig = plt.figure()
plt.bar([1, 2, 3], [risk_1, risk_10, risk_100])
plt.xticks([1, 2, 3], ['0-1', '10x', '100x'])
plt.xlabel('Loss')
plt.ylabel('Risk')
plt.show()
```

The minimum expected risk for the 0-1 loss is 3908 The minimum expected risk for the 10x loss is 61254 The minimum expected risk for the 100x loss is 711605



We've calculated the minimum expected risk using each of the loss matrices. The notable distinction between these matrices is that they progressively care more about not making mistakes for Y = 3. As seen by the confusion matrix and the bar graph, this led to worse results as the loss matrix increased. For example, the confusion matrix for 100 has 0s all through the first column, meaning that for the true label Y = 1, there were no classifications for that label. This is because the loss matrix cared so much about not misclassifying label 3 that it would incorrectly bias towards

classifying as 3.

Another interesting note is that as the loss matrix increased, the number of correct classifications for label 3 increased. This came at a significant tradeoff, however, since it also meant that the number of incorrect classifications increased. Thus, it's important to consider all the effects of classification and not just focus on one number to determine success.