The Gaussian class conditional pdf for 3D real valued random vector x is

$$P(x) = P(L=0) P(x|L=0) + P(L=1) P(x|L=1) + P(L=2) P(x|L=2) + P(L=3) P(x|L=3)$$

Since the class priors are uniform, P(L=0) = P(L=1) = P(L=2) = P(L=3) = 0.25

for class label 0: P(L=0) = 0.25

$$m_0 = \begin{bmatrix} 2.2 \\ 0 \\ 0 \end{bmatrix} \qquad C_0 = \begin{bmatrix} 1 & -0.5 & 0.3 \\ -0.5 & 1 & -0.5 \\ 0.3 & -0.5 & 1 \end{bmatrix}$$

for class label 1: P(L=1) = 0.25

$$M_1 = \begin{bmatrix} 2.2 \\ 2.2 \\ 0 \end{bmatrix} \qquad C_1 = \begin{bmatrix} 1 & 0.5 & 0 \\ 0.5 & 1 & 0.5 \\ 0 & 0.5 & 1 \end{bmatrix}$$

for class label 2: P(L=2) = 0.25

$$P(x|L=2) = g(x|m_2, C_2)$$

$$m_2 = \begin{bmatrix} 2.2 \\ 0 \\ 2.2 \end{bmatrix} \qquad C_2 = \begin{bmatrix} 1 & 0.3 & -0.2 \\ 0.3 & 1 & 0.3 \\ -0.2 & 0.3 & 1 \end{bmatrix}$$
For class label 3:

$$P(L=3) = 0.25$$

$$P(X|L=3) = g(X|m_3, C_3)$$

$$m_3 = \begin{bmatrix} 0 \\ 2.2 \\ 0 \end{bmatrix}$$

$$C_3 = \begin{bmatrix} 1 & 0.3 & 0.6 \\ 0.3 & 1 & 0.3 \\ 0.6 & 0.3 & 1 \end{bmatrix}$$

Hence, we have generated a 3-dimensional real value random vector X from 4 different classes having uniform class priors.

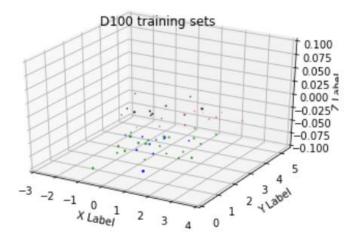
While choosing the parameters (mean & covariance), the MAP classifier that uses the true data pdf obtained the following probability of error with the given number of input samples:

Number of Samples	Probability of Error
100	0.17
200	0.15
500	0.18
1000	0.186
2000	0.1745
5000	0.1732
100000 (Test dataset)	0.17567

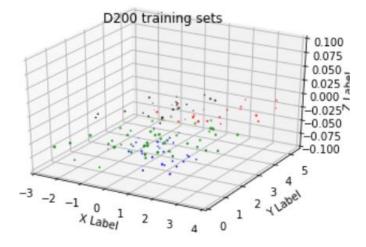
Therefore, from the table above, we can say that the probability of error lies within the 10-20 % range for the different number of samples for the set means and covariance matrices.

For the training datasets, having 100, 200, 500, 1000, 2000 and 5000 number of samples, 3-dimensional scatter plots have been generated. A scatter plot for the 100k-test dataset has also been generated and shown below.

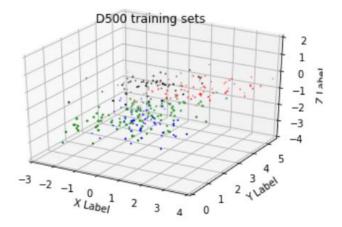
1. D100 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



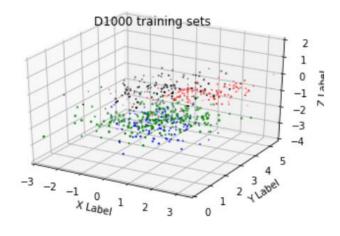
2. D200 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



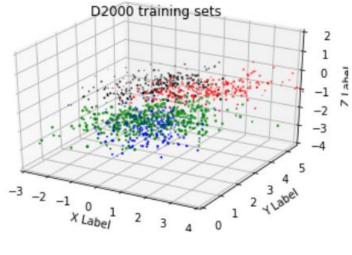
3. D500 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



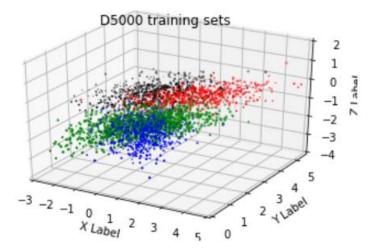
4. D1000 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



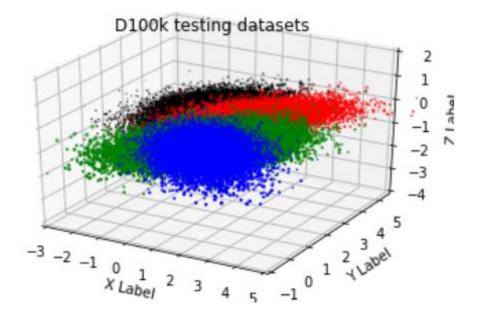
5. D2000 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



6. D5000 Training Datasets (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



7. Test Dataset 100k (Class 0 : Blue, Class 1 : Red, Class 2: Green, Class 3 : Black)



After generating the datasets and plotting the 3-D scatter plot, we must obtain the optimum perceptron number needed for the best performance (least possible error obtained). This is done by the help of K fold (in this case, k=10) cross validation. The cross-validation loop also trains the model with the MLP, and we obtain the errors obtained with their respective number of perceptron's. Once this optimum number of perceptron's are finalized, the final model is trained with

the optimal number and the training dataset. At the end, the trained model performance analysis is obtained by applying it on the test dataset (100k samples). The probability of error is also obtained and calculated to evaluate the performance of the model overall.

While calculating the optimum number of perceptron's and error obtained, the following software packages were implemented:

- K fold: This sk learn software package is used to form K folds and split the
 dataset into the training index and validation index. With the help of this,
 we divided the input (ie the matrix formed by generating random samples
 of the given number of training datasets) into training and testing.
- MLPClassifier: This stands for Multi-Layer Perceptron Classifier, and was
 used inside the loop which was formed for obtaining K Fold cross validation
 of the training dataset of different number of samples (100, 200, 500, 1000,
 2000 & 5000). There are a few values of p (ideal number of perceptron's
 needed) given in the form of a list from where it checks which 'p' gives the
 least value.

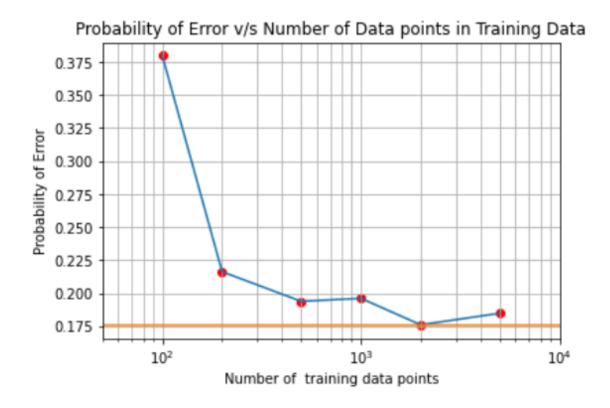
There are 2 hidden layers given (p,4), one for the number of perceptron and the other for the number of classes (which is 4 in this case). As needed, the activation function is set to 'ReLU', and the data to be trained is added to the MLP classifier using .fit on the classifier. Next, the out activation is set as 'SoftMax', as mentioned in the question. The score method is used to determine the correct and incorrect values obtained after training the neural network for the class posteriors. In this manner, by using the software packages and giving input for the activation function, number of hidden layers, output function & score calculation, we can say that it is performing the task we wanted it to.

Here is a tabular form of the optimum number of perceptron's required, along with the size of input samples and error obtained:

Number of	Optimum number of	Probability of Error
Input Samples	Perceptron's	(After doing performance analysis on
	•	100k test dataset)
100	4	0.37972
200	4	0.21622
500	6	0.19382
1000	6	0.19607
2000	8	0.176
5000	10	0.18476

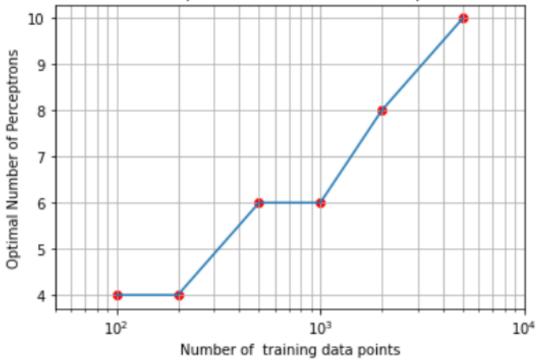
From the table we can conclude that as the size of the number of input samples increases, the probability of error decreases.

Following graph shows the output of the usage of software packages elucidated above. The probability of error v/s number of data points in training data plot again explains that the error probability decreases as the size of the dataset begins to increase. (The straight orange line depicts the empirically estimated test error for the theoretically optimal classifier)



The following graph shows the Optimal number of perceptron's number against the Number of data points. The trend we can see here is that as the number of training data points are increased, the optimum number of perceptron's required increased as well.





CODE STARTS HERE:

#Importing all the required libraries

```
import random
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import multivariate_normal
from mpl_toolkits import mplot3d
from sklearn.neural_network import MLPClassifier
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
```

```
from sklearn.model selection import KFold
#Ignoring the output compiler warnings given while execution on Google Col
lab
import warnings
warnings.filterwarnings('ignore')
#-----
#Initializing the values of class means, covariances and class priors
#The class priors for all the 4 classes were uniform
p=0.25
#Means matrix
m0 = np.transpose(np.array([2.2,0,0]))
m1 = np.transpose(np.array([2.2,2.2,0]))
m2 = np.transpose(np.array([2.2,0,2.2]))
m3 = np.transpose(np.array([0,2.2,0]))
#Covariance matrix
c0 = np.array([[1,-0.5,0.3],[-0.5,1,-0.5],[0.3,-0.5,1]])
c1 = np.array([[1, 0.5, 0], [0.5, 1, 0.5], [0, 0.5, 1]])
c2 = np.array([[1, 0.3, -0.2], [0.3, 1, 0.3], [-0.2, 0.3, 1]])
c3 = np.array([[1, 0.3, 0.6], [0.3, 1, 0.3], [0.6, 0.3, 1]])
#-----
#Defining the MAP loss and Risk calculation function, which will then be u
sed to check whether the mean, covariance taken lie under an error of 10 t
o 20%
def risk(i,x,lam matrix):
 total = 0
 for j in range(4):
   total += lam matrix[i][j]*p[j]*multivariate normal.pdf(x,mean[j],cov[j
 return total
def MAP loss(gauss, lam matrix):
 for i in r[gauss]:
   choice = np.argmin([risk(0,i,lam matrix),risk(1,i,lam matrix),risk(2,i
,lam matrix),risk(3,i,lam matrix)])
   loss matrix[gauss][choice] = loss matrix[gauss][choice] + 1
 return loss matrix
#-----
```

```
#100 Samples
#Generation of datapoints for 100 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 100
N=100
N samples 100 = [0, 0, 0, 0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
\#Initializing the lambda and loss matrices, to be used in calculation of M
AP error
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
#Sample Generation for Class labels 0, 1, 2 & 3
for i in range(N):
  a = random.random()
  if a<0.25:
    N samples 100[0] += 1
  elif a<0.5:
    N samples 100[1] += 1
  elif a<0.75:
    N samples 100[2] += 1
  else:
    N samples 100[3] += 1
#Storing generated samples in list 'r' declared previously, which are bein
g used for MAP loss function
#for i in range (4):
  #r.append(np.random.multivariate normal(mean[i],cov[i],N samples 100[i])
#Generating random samples for all the 4 classes
r0 100 = np.random.multivariate normal(m0,c0,N samples 100[0])
r1 100 = np.random.multivariate normal(m1,c1,N samples 100[1])
r2 100 = np.random.multivariate normal(m2,c2,N samples 100[2])
r3 100 = np.random.multivariate normal(m3,c3,N samples 100[3])
r = [r0 \ 100, r1 \ 100, r2 \ 100, r3 \ 100]
#Calculating MAP to determine overall error obtained from the means and co
variances for all 4 classes (0,1,2,3)
```

```
MAP loss(0,lam matrix)
MAP loss(1,lam matrix)
MAP loss(2,lam matrix)
MAP loss(3,lam matrix)
total loss = 0
#Printing of the total loss
for i in range(4):
 for j in range(4):
   if i!=j:
     total loss += loss_matrix[i][j]
print(total loss/N)
#-----
#Plotting the 3-D scatter plot for 100 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0 100[:,0],r0 100[:,1],r0 100[:,2],c='b')
plt.scatter(r1 100[:,0],r1 100[:,1],r1 100[:,2],c='r')
plt.scatter(r2 100[:,0],r2 100[:,1],r2 100[:,2],c='g')
plt.scatter(r3_100[:,0],r3_100[:,1],r3 100[:,2],c='k')
\#Adding title and labels for x,y,z axis
plt.title("D100 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
#Setting scatter plot limits
ax.set xlim3d(-3,4)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-0.1,0.1)
plt.show()
#-----
           -----
#200 Samples
#Generation of datapoints for 200 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 200
```

```
N = 200
N \text{ samples } 200 = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
#Initializing the lambda and loss matrices, to be used in calculation of M
AP error
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
#Sample generation for class labels 0,1,2 & 3
for i in range(N):
  a = random.random()
  if a<0.25:
    N samples 200[0] += 1
  elif a<0.5:
    N samples 200[1] += 1
  elif a<0.75:</pre>
    N samples 200[2] += 1
  else:
    N samples 200[3] += 1
r0 200 = np.random.multivariate normal(m0,c0,N samples 200[0])
r1 200 = np.random.multivariate normal(m1,c1,N samples 200[1])
r2 200 = np.random.multivariate normal(m2,c2,N samples 200[2])
r3 200 = np.random.multivariate normal(m3,c3,N samples 200[3])
r = [r0 \ 200, r1 \ 200, r2 \ 200, r3 \ 200]
#Calculating loss in the given means and covariance
MAP loss(0,lam matrix)
MAP loss(1, lam matrix)
MAP loss(2, lam matrix)
MAP loss(3, lam matrix)
total loss = 0
#Showing output of total loss
for i in range(4):
  for j in range(4):
    if i!=j:
      total loss += loss_matrix[i][j]
```

```
print(total loss/N)
#-----
#Plotting the 3-D scatter plot for 200 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0 200[:,0],r0 200[:,1],r0 200[:,2],c='b')
plt.scatter(r1 200[:,0],r1 200[:,1],r1 200[:,2],c='r')
plt.scatter(r2 200[:,0],r2 200[:,1],r2 200[:,2],c='g')
plt.scatter(r3 200[:,0],r3 200[:,1],r3 200[:,2],c='k')
#Adding scatter plot title and labels for the 3 different axis
plt.title("D200 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
#Setting the scatterplot limits
ax.set x \lim 3d(-3,4)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-0.1,0.1)
plt.show()
#-----
#500 Samples
#Generation of datapoints for 500 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 500
N = 500
N \text{ samples } 500 = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
#Initializing the lambda and loss matrices, to be used in calculation of M
AP error
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
```

```
#Generating random samples for classes 0, 1, 2 & 3
for i in range(N):
  a = random.random()
  if a<0.25:
   N samples 500[0] += 1
  elif a<0.5:
    N samples 500[1] += 1
  elif a<0.75:
   N samples 500[2] += 1
  else:
    N samples 500[3] += 1
r0 500 = np.random.multivariate normal(m0,c0,N samples 500[0])
r1 500 = np.random.multivariate normal(m1,c1,N samples 500[1])
r2 500 = np.random.multivariate normal(m2,c2,N samples 500[2])
r3 500 = np.random.multivariate normal(m3,c3,N samples 500[3])
r = [r0 500, r1 500, r2 500, r3 500]
#Determining MAP loss for each class (ie 0,1,2,3)
MAP loss(0,lam matrix)
MAP loss(1, lam matrix)
MAP loss(2, lam matrix)
MAP loss(3, lam matrix)
total loss = 0
for i in range(4):
 for j in range(4):
   if i!=j:
     total loss += loss matrix[i][j]
print(total loss/N)
#-----
                             ______
#Plotting the 3-D scatter plot for 500 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0 500[:,0],r0 500[:,1],r0 500[:,2],c='b')
plt.scatter(r1_500[:,0],r1_500[:,1],r1 500[:,2],c='r')
plt.scatter(r2 500[:,0],r2 500[:,1],r2 500[:,2],c='g')
plt.scatter(r3 500[:,0],r3 500[:,1],r3 500[:,2],c='k')
```

```
#Adding label and title
plt.title("D500 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
#Setting limits of the axes
ax.set xlim3d(-3,4)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-4,2)
plt.show()
#-----
#1000 Samples
#Generation of datapoints for 1000 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 1000
N=1000
N \text{ samples } 1k = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
#Initializing the lambda and loss matrices, to be used in calculation of M
AP error
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
for i in range(N):
  a = random.random()
  if a<0.25:
   N samples 1k[0] += 1
  elif a<0.5:
   N samples 1k[1] += 1
  elif a<0.75:
    N samples 1k[2] += 1
  else:
   N samples 1k[3] += 1
r0 1k = np.random.multivariate normal(m0,c0,N samples 1k[0])
```

```
r1 1k = np.random.multivariate normal(m1,c1,N samples 1k[1])
r2 1k = np.random.multivariate normal(m2,c2,N samples 1k[2])
r3 1k = np.random.multivariate normal(m3,c3,N samples 1k[3])
r = [r0 1k, r1 1k, r2 1k, r3 1k]
MAP loss(0,lam matrix)
MAP loss(1, lam matrix)
MAP loss(2,lam matrix)
MAP loss(3, lam matrix)
total loss = 0
for i in range(4):
 for j in range(4):
   if i!=j:
     total loss += loss matrix[i][j]
print(total loss/N)
#-----
                            ______
#Plotting the 3-D scatter plot for 1000 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
#Creation of scatter plot
plt.scatter(r0 1k[:,0],r0 1k[:,1],r0 1k[:,2],c='b')
plt.scatter(r1 1k[:,0],r1 1k[:,1],r1 1k[:,2],c='r')
plt.scatter(r2 1k[:,0],r2 1k[:,1],r2 1k[:,2],c='g')
plt.scatter(r3 1k[:,0],r3 1k[:,1],r3 1k[:,2],c='k')
#Adding of title and labels of axes
plt.title("D1000 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
#Setting limits for the three axes
ax.set x \lim 3d(-3,3.5)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-4,2)
plt.show()
```

```
#2000 Samples
#Generation of datapoints for 2000 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 2000
N = 2000
N \text{ samples } 2k = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
#Loss & Lambda matrix will be used to determing error and check whether it
lies between 10 and 20%
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
for i in range(N):
  a = random.random()
  if a<0.25:
    N samples 2k[0] += 1
  elif a<0.5:
    N samples 2k[1] += 1
  elif a<0.75:
    N samples 2k[2] += 1
  else:
    N samples 2k[3] += 1
r0 2k = np.random.multivariate normal(m0,c0,N samples 2k[0])
r1 2k = np.random.multivariate normal(m1,c1,N samples 2k[1])
r2 2k = np.random.multivariate normal(m2,c2,N samples 2k[2])
r3 2k = np.random.multivariate normal(m3,c3,N samples <math>2k[3])
r = [r0 \ 2k, r1 \ 2k, r2 \ 2k, r3 \ 2k]
#Function call to return error obtained
MAP loss(0, lam matrix)
MAP loss(1, lam matrix)
MAP loss(2, lam matrix)
MAP loss(3, lam matrix)
total loss = 0
#Printing total loss
```

```
for i in range(4):
 for j in range(4):
   if i!=j:
     total loss += loss matrix[i][j]
print(total loss/N)
#-----
#Plotting the 3-D scatter plot for 2000 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0 2k[:,0],r0 2k[:,1],r0 2k[:,2],c='b')
plt.scatter(r1 2k[:,0],r1 2k[:,1],r1 2k[:,2],c='r')
plt.scatter(r2 2k[:,0],r2 2k[:,1],r2 2k[:,2],c='g')
plt.scatter(r3 2k[:,0],r3 2k[:,1],r3 2k[:,2],c='k')
#Setting scatter plot title and axes labels
plt.title("D2000 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
#Setting the axes limits
ax.set xlim3d(-3,4)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-4,2)
plt.show()
#-----
#5000 Samples
#Generation of datapoints for 5000 samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 5000
N = 5000
N samples 5k = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
```

```
#Loss and Lambda matrix are being used to determine error
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
for i in range(N):
  a = random.random()
  if a<0.25:
    N samples 5k[0] += 1
  elif a<0.5:
    N samples 5k[1] += 1
  elif a<0.75:</pre>
    N samples 5k[2] += 1
  else:
    N_samples_5k[3] += 1
r0 5k = np.random.multivariate normal(m0,c0,N samples 5k[0])
r1 5k = np.random.multivariate normal(m1,c1,N samples 5k[1])
r2 5k = np.random.multivariate normal(m2,c2,N samples <math>5k[2])
r3 5k = np.random.multivariate normal(m3,c3,N samples <math>5k[3])
r = [r0 5k, r1 5k, r2 5k, r3 5k]
MAP loss(0,lam matrix)
MAP loss(1, lam matrix)
MAP loss(2, lam matrix)
MAP loss(3,lam matrix)
total loss = 0
#Printing out the error
for i in range(4):
  for j in range(4):
    if i!=j:
      total loss += loss matrix[i][j]
print(total loss/N)
#Plotting the 3-D scatter plot for 5000 Training Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0 5k[:,0],r0 5k[:,1],r0 5k[:,2],c='b')
```

```
plt.scatter(r1 5k[:,0],r1 5k[:,1],r1 5k[:,2],c='r')
plt.scatter(r2 5k[:,0],r2 5k[:,1],r2 5k[:,2],c='g')
plt.scatter(r3 5k[:,0],r3 5k[:,1],r3 5k[:,2],c='k')
#Displaying title and axes labels
plt.title("D5000 training sets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
ax.set x \lim 3d(-3, 5)
ax.set ylim3d(0,5.5)
ax.set zlim3d(-4,2)
plt.show()
#100k Test Data Samples
#Generation of datapoints for 100000 test data samples
#Checking the probability of error (and then printing the error) in the me
ans and covariances when samples are 100k
N=100000
N samples 100k = [0,0,0,0]
r = []
mean = [m0, m1, m2, m3]
cov = [c0, c1, c2, c3]
p = [0.25, 0.25, 0.25, 0.25]
loss matrix = [[0,0,0,0],[0,0,0,0],[0,0,0,0],[0,0,0,0]]
lam matrix = [[0,1,1,1],[1,0,1,1],[1,1,0,1],[1,1,1,0]]
for i in range(N):
 a = random.random()
  if a<0.25:
   N samples 100k[0] += 1
  elif a<0.5:
    N samples 100k[1] += 1
  elif a<0.75:
    N samples 100k[2] += 1
  else:
    N samples 100k[3] += 1
r0 100k = np.random.multivariate normal(m0,c0,N samples 100k[0])
r1 100k = np.random.multivariate normal(m1,c1,N samples 100k[1])
```

```
r2 100k = np.random.multivariate normal(m2,c2,N samples 100k[2])
r3 100k = np.random.multivariate normal(m3,c3,N samples 100k[3])
r = [r0_100k, r1_100k, r2\ 100k, r3\ 100k]
MAP loss(0,lam matrix)
MAP loss(1, lam matrix)
MAP loss(2,lam matrix)
MAP loss(3,lam matrix)
total loss = 0
for i in range(4):
 for j in range(4):
   if i!=j:
      total loss += loss_matrix[i][j]
print(total loss/N)
error_theo = total loss/N
#-----
#Plotting the 3-D scatter plot for 100000 Test Samples
ax = plt.axes(projection ="3d")
# Creating plot
plt.title("simple 3D scatter plot")
plt.scatter(r0_100k[:,0],r0_100k[:,1],r0 100k[:,2],c='b')
plt.scatter(r1 100k[:,0],r1 100k[:,1],r1 100k[:,2],c='r')
plt.scatter(r2_100k[:,0],r2_100k[:,1],r2_100k[:,2],c='g')
plt.scatter(r3 100k[:,0],r3 100k[:,1],r3 100k[:,2],c='k')
plt.title("D100k testing datasets")
ax.set xlabel('X Label')
ax.set ylabel('Y Label')
ax.set zlabel('Z Label')
ax.set x \lim 3d(-3,5)
ax.set ylim3d(-1,5.5)
ax.set zlim3d(-4,2)
plt.show()
```

```
#Model Order Selection
#Initializing empty list for storing the ideal number of perceptrons for e
ach training sample
perceptron number = []
#Initializing empty list to store the probability of error for each traini
ng sample
prob error = []
#Storing the perceptron values in a list to be used for MLP training
p \text{ values} = [4, 6, 8, 10, 12, 14]
#These are the labels of the test dataset (100k Samples) which will be use
d on the trained MLP model of each train dataset
#The 4 different classes are concatenated into 1 array of dimension (10000
0, 2)
r 100k = np.concatenate((r0 100k, r1 100k, r2 100k, r3 100k))
a 100k = np.array(np.zeros(N samples 100k[0]))
b 100k = np.array(np.ones(N samples 100k[1]))
c 100k = np.array(2*np.ones(N samples 100k[2]))
d 100k = np.array(3*np.ones(N samples 100k[3]))
r 100k l = np.concatenate((a 100k,b 100k,c 100k,d 100k))
r 100k labels = r 100k l.reshape(100000,1)
#100 Samples
#These are the labels of the train dataset which contains 100 samples, all
of them are concatenated into one list along with their class labels
r 100 = np.concatenate((r0 100, r1 100, r2 100, r3 100))
a 100 = np.array(np.zeros(N samples 100[0]))
b 100 = np.array(np.ones(N samples 100[1]))
c 100 = np.array(2*np.ones(N samples 100[2]))
d 100 = np.array(3*np.ones(N samples 100[3]))
r 100 l = np.concatenate((a 100,b 100,c 100,d 100))
r 100 labels = r 100 l.reshape(100,1)
#-----
                                   ._____
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
```

```
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
for p in p values:
   kf = KFold(n splits=10)
   per acc = []
   for train index, validation index in kf.split(r 100):
     clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 100[train index,:],r 100 labels[train inde
x,:])
     clf.out activation = 'softmax'
     acc = clf.score(r 100[validation index,:],r 100 labels[validation in
dex,: ])
     per acc.append(acc)
   per number.append(sum(per acc))
#The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final p 100 = p values[np.argmin(np.array(per number))]
print("Optimal number of Perceptrons: ",final p 100)
perceptron number.append(final p 100)
#After the P number is obtained, the MLP Classifier is again trained with
the optimal number of perceptrons
#The accuracy (performance asssessment) is done by applying the trained mo
del on the test data of 100k samples
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
l p 100,4),activation='relu').fit(r 100,r 100 labels)
acc 100 = clf.score(r 100k,r 100k labels)
err 100 = 1 - acc 100
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 100)
prob error.append(err 100)
#-----
#200 Samples
#These are the labels of the train dataset which contains 200 samples, all
of them are concatenated into one list along with their class labels
r 200 = np.concatenate((r0 200, r1 200, r2 200, r3 200))
a 200 = np.array(np.zeros(N samples 200[0]))
b 200 = np.array(np.ones(N samples 200[1]))
c 200 = np.array(2*np.ones(N samples 200[2]))
```

```
d 200 = np.array(3*np.ones(N samples 200[3]))
r 200 l = np.concatenate((a 200,b 200,c 200,d 200))
r 200 labels = r 200 l.reshape(200,1)
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
for p in p values:
   kf = KFold(n splits=10)
   per acc = []
    for train index, validation index in kf.split(r 200):
     clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 200[train index,:],r 200 labels[train inde
x,:])
     clf.out activation = 'softmax'
     acc = clf.score(r 200[validation index,:],r 200 labels[validation in
dex,:])
     per acc.append(acc)
   per number.append(sum(per acc))
#-----
#The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final p 200 = p values[np.argmin(np.array(per number))]
print("Optimal number of Perceptrons: ", final p 200)
perceptron number.append(final p 200)
#After the P number is obtained, the MLP Classifier is again trained with
the optimal number of perceptrons
#The accuracy (performance asssessment) is done by applying the trained mo
del on the test data of 100k samples
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
1 p 200,4),activation='relu').fit(r 200,r 200 labels)
acc 200 = clf.score(r 100k,r 100k labels)
err 200 = 1 - acc 200
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 200)
prob error.append(err 200)
```

```
#500 Samples
#These are the labels of the train dataset which contains 500 samples, all
of them are concatenated into one list along with their class labels
r 500 = np.concatenate((r0 500,r1 500,r2 500,r3 500))
a 500 = np.array(np.zeros(N samples 500[0]))
b 500 = np.array(np.ones(N samples 500[1]))
c 500 = np.array(2*np.ones(N samples 500[2]))
d 500 = np.array(3*np.ones(N samples 500[3]))
r 500 l = np.concatenate((a 500,b 500,c 500,d 500))
r 500 labels = r 500 l.reshape(500,1)
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
for p in p values:
   kf = KFold(n splits=10)
   per acc = []
   for train index, validation index in kf.split(r 500):
     clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 500[train index,:],r 500 labels[train inde
x,:])
     clf.out activation = 'softmax'
     acc = clf.score(r 500[validation index,:],r 500 labels[validation in
dex,: ])
     per acc.append(acc)
   per number.append(sum(per acc))
#-----
#The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final p 500 = p values[np.argmin(np.array(per number))]
print("Optimal number of Perceptrons: ",final_p_500)
perceptron number.append(final p 500)
#After the P number is obtained, the MLP Classifier is again trained with
the optimal number of perceptrons
#The accuracy (performance asssessment) is done by applying the trained mo
del on the test data of 100k samples
```

```
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
1 p 500,4),activation='relu').fit(r 500,r 500 labels)
acc 500 = clf.score(r 100k,r 100k labels)
err 500 = 1 - acc 500
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 500)
prob error.append(err 500)
#1000 Samples
#These are the labels of the train dataset which contains 1k samples, all
of them are concatenated into one list along with their class labels
r 1k = np.concatenate((r0 1k,r1 1k,r2 1k,r3 1k))
a 1k = np.array(np.zeros(N samples 1k[0]))
b 1k = np.array(np.ones(N samples 1k[1]))
c 1k = np.array(2*np.ones(N samples 1k[2]))
d 1k = np.array(3*np.ones(N samples 1k[3]))
r 1k 1 = np.concatenate((a 1k,b 1k,c 1k,d 1k))
r 1k labels = r 1k l.reshape(1000,1)
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
for p in p values:
   kf = KFold(n splits=10)
   per acc = []
    for train index, validation index in kf.split(r 1k):
      clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 1k[train index,:],r 1k labels[train index,
: ])
     clf.out activation = 'softmax'
     acc = clf.score(r 1k[validation index,:],r 1k labels[validation inde
x,:])
     per acc.append(acc)
   per number.append(sum(per acc))
                                  #The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final p 1k = p values[np.argmin(np.array(per number))]
```

```
print("Optimal number of Perceptrons: ", final p 1k)
perceptron number.append(final p 1k)
#After the P number is obtained, the MLP Classifier is again trained with
the optimal number of perceptrons
#The accuracy (performance asssessment) is done by applying the trained mo
del on the test data of 100k samples
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
l p 1k,4),activation='relu').fit(r 1k,r 1k labels)
acc 1k = clf.score(r 100k, r 100k labels)
err 1k = 1 - acc 1k
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 1k)
prob error.append(err 1k)
                             _____
#2000 Samples
#These are the labels of the train dataset which contains 2k samples, all
of them are concatenated into one list along with their class labels
r 2k = np.concatenate((r0 2k,r1 2k,r2 2k,r3 2k))
a 2k = np.array(np.zeros(N samples 2k[0]))
b 2k = np.array(np.ones(N samples 2k[1]))
c_2k = np.array(2*np.ones(N samples 2k[2]))
d 2k = np.array(3*np.ones(N samples 2k[3]))
r 2k 1 = np.concatenate((a 2k,b 2k,c 2k,d 2k))
r 2k labels = r 2k l.reshape(2000,1)
                                      ______
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
for p in p values:
   kf = KFold(n_splits=10)
   per acc = []
    for train index, validation_index in kf.split(r_2k):
     clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 2k[train index,:],r 2k labels[train index,
: ] )
     clf.out activation = 'softmax'
```

```
acc = clf.score(r 2k[validation index,:],r 2k labels[validation inde
x,:])
     per acc.append(acc)
   per number.append(sum(per acc))
#The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final_p_2k = p_values[np.argmin(np.array(per_number))]
print("Optimal number of Perceptrons: ",final p 2k)
perceptron number.append(final p 2k)
#After the P number is obtained, the MLP Classifier is again trained with
the optimal number of perceptrons
#The accuracy (performance asssessment) is done by applying the trained mo
del on the test data of 100k samples
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
1 p 2k,4),activation='relu').fit(r 2k,r 2k labels)
acc_2k = clf.score(r_2k,r_2k_labels)
err 2k = 1 - acc 2k
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 2k)
prob error.append(err 2k)
#-----
#5000 Samples
#These are the labels of the train dataset which contains 5k samples, all
of them are concatenated into one list along with their class labels
r_5k = np.concatenate((r0_5k,r1_5k,r2_5k,r3_5k))
a 5k = np.array(np.zeros(N samples <math>5k[0]))
b 5k = np.array(np.ones(N samples 5k[1]))
c 5k = np.array(2*np.ones(N samples <math>5k[2]))
d 5k = np.array(3*np.ones(N samples 5k[3]))
r_5k_1 = np.concatenate((a_5k,b_5k,c_5k,d_5k))
r 5k labels = r 5k l.reshape(5000,1)
#-----
#The optimal number of perceptrons are obtained via this loop with the hel
p of K fold cross validation and MLP classifier
#An empty list is created to store the accuracy, which will later be used
to obtain error
per number = []
```

```
for p in p values:
   kf = KFold(n splits=10)
   per acc = []
   for train index, validation index in kf.split(r 5k):
     clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes
=(p,4),activation='relu').fit(r 5k[train index,:],r 5k labels[train index,
: ])
     clf.out activation = 'softmax'
     acc = clf.score(r 5k[validation index,:],r 5k labels[validation inde
x,:])
     per acc.append(acc)
   per number.append(sum(per acc))
#-----
#The final optimal number of perceptrons are obtained, printed and then st
ored in the perceptron number array
final p 5k = p values[np.argmin(np.array(per number))]
print("Optimal number of Perceptrons: ",final p 5k)
perceptron number.append(final p 5k)
clf = MLPClassifier(random state=1, max iter=1000, hidden layer sizes=(fina
1 p 5k,4),activation='relu').fit(r 5k,r 5k labels)
acc 5k = clf.score(r 100k, r 100k labels)
err 5k = 1 - acc 5k
print("Error Probability after training the model and assessing performanc
e on test dataset: ",err 5k)
prob error.append(err 5k)
#-----
#Displaying lists formed by perceptron number (Optimum) & Error Probabilit
print("The ideal perceptron numbers for all training datasets is: ",percep
tron number)
print("The corresponding error probability is: ",prob error)
#-----
#Plotting graph for Probability of Error v/s Number of Data points in Trai
ning Data
data points = [100, 200, 500, 1000, 2000, 5000]
plt.grid(True, which='both')
plt.semilogx(data points, prob error)
plt.scatter(data points,prob error,c='r')
plt.xlabel("Number of training data points")
plt.ylabel("Probability of Error")
```

```
plt.title("Probability of Error v/s Number of Data points in Training Data
")
plt.plot([50,10000],[error theo,error theo])
plt.xlim(50,10000)
plt.show()
#Plotting graph for Optimal Number of Perceptrons v/s Number of Data point
s in Training Data
data points = [100, 200, 500, 1000, 2000, 5000]
plt.grid(True, which='both')
plt.semilogx(data points, perceptron number)
plt.scatter(data points, perceptron number, c='r')
plt.xlabel("Number of training data points")
plt.ylabel("Optimal Number of Perceptrons")
plt.title("Optimal Number of Perceptrons v/s Number of Data points in Trai
ning Data")
plt.xlim(50,10000)
plt.show()
#----- END OF SOLUTION 1 ------
```

I have implemented the above code using Google Collab, on the following URL:

https://colab.research.google.com/drive/1dPcA6j4fSqs8yoXjQ84Gtr0Yfuh5GRb6?usp=s haring