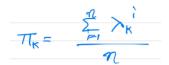
- You are given a data-set with 400 data points in {0,1}⁵⁰ generated from a mixture of some distribution in the file A2Q1.csv. (Hint: Each datapoint is a flattened version of a {0,1}^{10×5} matrix.)
 - i. (i) Determine which probabilisitic mixture could have generated this data (It is not a Gaussian mixture). Derive the EM algorithm for your choice of mixture and show your calculations. Write a piece of code to implement the algorithm you derived by setting the number of mixtures K = 4. Plot the log-likelihood (averaged over 100 random initializations) as a function of iterations.

- ii. (ii) Assume that the same data was infact generated from a mixture of Gaussians with 4 mixtures. Implement the EM algorithm and plot the log-likelihood (averaged over 100 random initializations of the parameters) as a function of iterations. How does the plot compare with the plot from part (i)? Provide insights that you draw from this experiment.
- iii. Run the K-means algorithm with K = 4 on the same data. Plot the objective of K - means as a function of iterations.
- iv. Among the three different algorithms implemented above, which do you think you would choose to for this dataset and why?

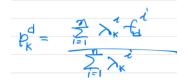
1.1)

I have used the Bernoulli model in the 4 models . each bernoulli model has different parameter p Here the graph becomes parallel to the X-axis after running some iterations .so it means if we increase the step number the error more or less remains the same.

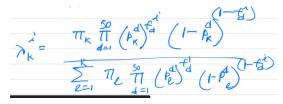
PI formula for bernoulli which gives probability that a model k is selected

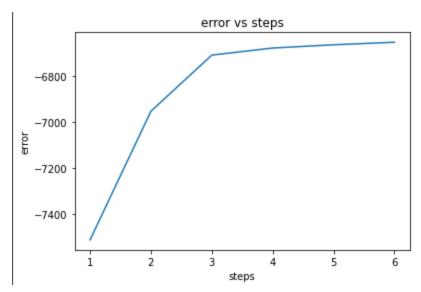


Each parameter of the model -



I have made a matrix of this parameters Each lambda value will be =





I tried to see how many points come from each of the clusters.

From cluster 1 = 122

From cluster 2= 110

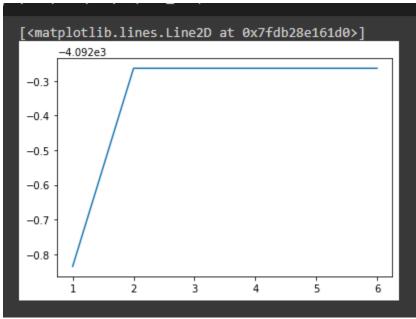
From cluster 3 = 101

From cluster 4 = 67

So the points come from more or less all 4 models.

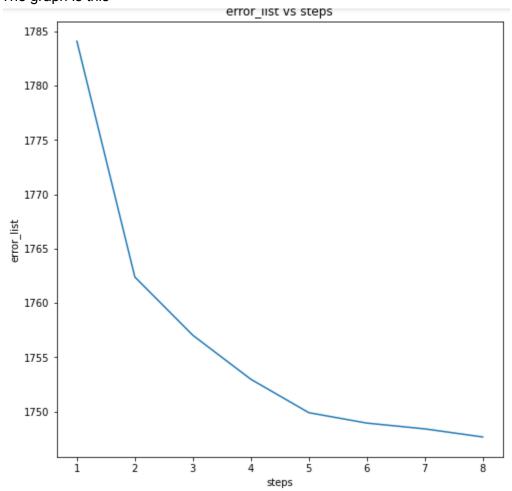
1.2)

Here we have 4 models and in each model there are 4 gaussian models with different mean and different covariance matrix . the objective function is =



Here almost all 400 points come from model 1 So it means in this case same model generates almost all data points

1.3) Here the k means algorithm is used to determine from which cluster(model) a datapoint comes from. The error function of k means is written with the iteration number of each k means . The graph is this =



But the exact graph changes each time as cluster assignment changes each time I run it. I have run this for 8 steps.

Here model1 generates = 35 points Here model2 generates = 235 points Here model3 generates = 101 points Here model4 generates = 29 points 1.4)

I have made an error function in which I am making an assignment matrix Z , which is basically the cluster number of each data point.

I am checking the distance of each point from the nearest mean and adding those -this serves as the error function.

With this function the bernoulli gives -1746 error

The gaussian gives = 2295 error

And the kmeans algo gives = 1746 error It means the k means and bernoulli are more or less of the same performance but gaussian model gives worse performance that them.

- (2) You are given a data-set in the file A2Q2Data_train.csv with 10000 points in (R¹⁰⁰, R) (Each row corresponds to a datapoint where the first 100 components are features and the last component is the associated y value).
 - Obtain the least squares solution w_{ML} to the regression problem using the analytical solution.
 - ii. Code the gradient descent algorithm with suitable step size to solve the least squares algorithms and plot ||w^t - w_{ML}||₂ as a function of t. What do you observe?
 - iii. Code the stochastic gradient descent algorithm using batch size of 100 and plot ||w^t − w_{ML}||₂ as a function of t. What are your observations?
 - iv. Code the gradient descent algorithm for ridge regression. Cross-validate for various choices of λ and plot the error in the validation set as a function of λ. For the best λ chosen, obtain w_R. Compare the test error (for the test data in the file A2Q2Data_test.csv) of w_R with w_{ML}. Which is better and why?

2.1)

The dataset has 10000 points and each point has 100 features. A equation of line is y=mx+c form, so here I have introduced another parameter in parameter array and it becomes size 101, and in the dataframe, I have attached one column of 1's in dataframe in every question of question 2, I have followed this.

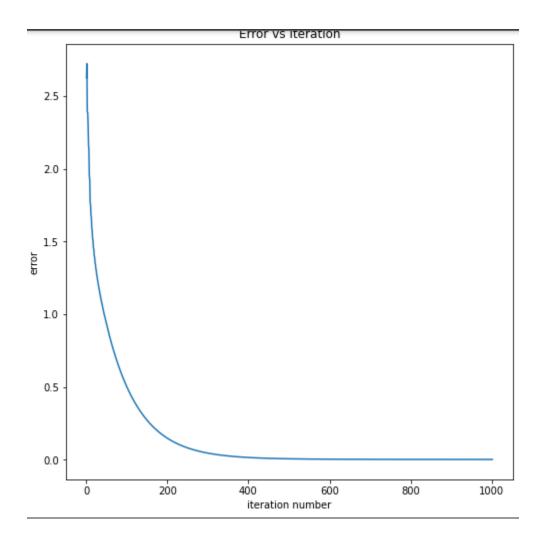
Analytical error on test data = 185.375

2.2)

In the gradient descent algorithm I have used the learning rate as the inverse of current step size.

I have run the algorithm 1000 times.

This is the graph I found -



Where in Y axis I have plotted -

$$\|\mathbf{w}^t - \mathbf{w}_{ML}\|_2$$

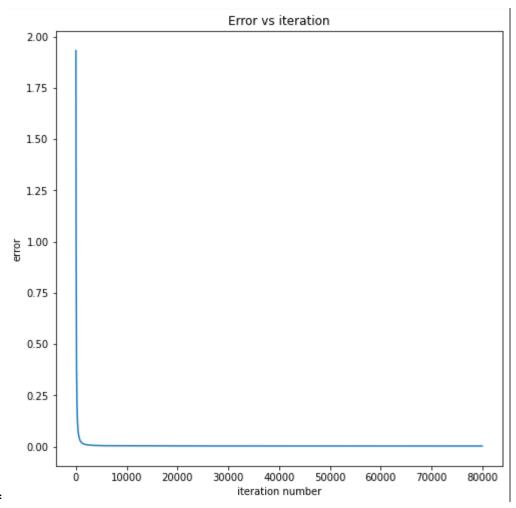
In X axis I have plotted the iteration number .

I have plotted the graph for 1000 steps.

The parameter obtained from this gives a error of = 185.055 on the test data

2.3)

In stochastic gradient descent , I have run it for batch size of 100. In test data = 183.925 In training data the error is =397.122



The graph is =

Where in Y axis I have plotted -

$$\|\mathbf{w}^t - \mathbf{w}_{ML}\|_2$$

In X axis I have plotted the iteration number .

2.4)

In ridge regression I have divided the train data into 80% and 20% and trained the data and found the parameter from the 80% and I am printing the error value on 20% of the training data . I am doing these steps for multiple lambda values. The lambda value that is giving the minimum error is the best lambda .

In this case the lambda value is = 3.42

The test data error is =184.670

The cross validation graph which shows error on y axis as a function of lambda value is =

