MTH 511a - Mini Project

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Due: 2nd November at 8:00pm

Please read the instructions on submission very carefully. The checking will be automatic, so if you do not follow proper directions, you will not be able to get any marks. This mini-project counts for 10% of the overall grade.

Data

Each and every one of you is provided with a unique dataset of 4 columns and 500 rows. You can load the data in R using the following command:

dat <- read.table("https://dvats.github.io/assets/data/rollnumber.txt")</pre>

where replace rollnumber with your roll number.

Model

Your goal is to fix a mixture model to the data provided to you. The data provided to you may have any number of clusters with a maximum cluster size of C = 7.

You may use any technique you like to fit the best mixture model to this fourdimensional data. Your final chosen model must be saved in a list called model. For example, in the Gaussian mixture model case, model would have list components

• Clusters = number of fitted clusters  $(C^*)$ .  $C^*$  will be a number between 2 and 7 (inclusive)

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- pi = a numeric vector of mixture probabilities of length  $C^*$
- $\mathtt{mu} = \mathtt{a} \ C^* \times \mathtt{4} \ \mathtt{matrix} \ \mathtt{of} \ \mathtt{mean} \ \mathtt{vectors} \ \mathtt{of} \ \mathtt{the} \ C^* \ \mathtt{different} \ \mathtt{mixture} \ \mathtt{components}$
- Sigma = a list of  $C^*$  4 × 4 covariance matrices; one for each component.
- log.like = negative log-likelihood value on the given data.

## Prediction

Write a function pred.loss that has arguments X.new (for test data input) and model (a fitted mixture model discussed above). This function calculates the loss on a new dataset X.new. I will input an X.new matrix of size  $n_1 \times 4$  for some  $n_1 > 0$  and the function should output loss, a scalar number.

```
pred.loss <- function(X.new, model)
{
    ...
    loss <- ...
    return(loss)
}</pre>
```

There should be no other input arguments in the function and the function should not use any other global variables aside from X.new and model. The loss is calculated based on the model you've fit.

Any external packages needed to run this function, must be loaded *inside* the function.

## Submission

At the end of your script, run the following code:

```
save(pred.loss, model, file = "rollnumer.Rdata")
```

This will save your function pred.loss and model in a file with your roll number as the name, in your <u>current working directory</u>. (To see your working directory, type getwd() in the R console).

When done, please upload your rollnumber. Rdata file in the following Dropbbox link:

https://www.dropbox.com/request/3qxQJd59SnUZUhPeSVzB

When submitting, please follow the instructions:

- Please sign out of Dropbox if you are signed in to Dropbox.
- Under "Your Name" write down your roll number
- Under "Your Email" use your iitk email id.

## Some words of caution

- Make sure your optimization routines don't consume all max.iter. If this happens, then this means you haven't converged as yet to the estimator.
- If your estimate of the covariance matrix  $\Sigma$ s is not positive-definite in any iteration, I recommend making your code run in such a way that it repeats that run with a different starting value.
- Follow the naming conventions I use here, otherwise you may not get any points. Particularly, make sure your final estimated model is stored in model and your function name is pred.loss. Do not use any other names for these two objects!
- Before submitting the rollnumber.Rdata file, make sure everything works. Change your working directory to the folder that contains rollnumber.Rdata. Run the following lines

```
rm(list = ls())
load("rollnumber.Rdata")
```

This will first clear all memory of the R session and then load your .Rdata file. After this, call pred.loss(X, model) using a dummy X matrix you create of size  $n_1 \times 4$ , for any  $n_1 > 1$ . If you get back a scalar number, then that means your function should give me no errors. The only possibility of an error is if your function uses a package that is not loaded *inside* the function.

## Evaluation

A unique X.new has been created for each and every one of you. This dataset will not be revealed to you. When you submit rollnumber.Rdata, I will load this in R and calculate

```
test.error <- pred.loss(X.new, model)</pre>
```

using your function. I will also calculate the loss under the *true model* for your dataset. (This true model is what was used to generate the data – this is unknown to you).

Finally, I will calculate

$$Score = \frac{\texttt{test.error}}{\texttt{true.error}}$$

The closer your score is to 1, the higher marks you get. Note, I will also check your pred.loss function. If there is an error in that function, then I will deduct points, and use my own version of the pred.loss function.

Good luck and SUBMIT ON TIME! Late submissions will be flagged by Dropbox and marks will be deducted.