## Data Analytics Mini Project - 2

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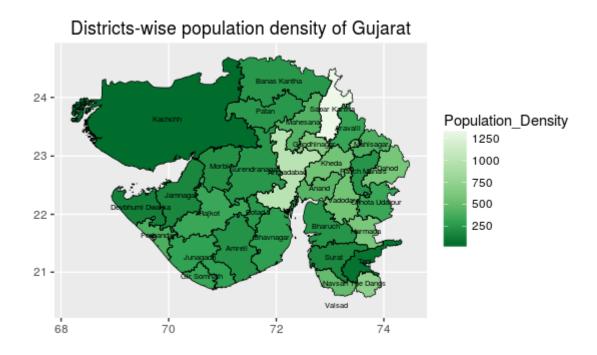
Instructor:

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## 0.1 Use R to make a map of Gujarat using an R package showing district-wise characteristic

Solution  $\rightarrow$  Given spatial data, in this case "District-wise Population Density", it can be incorporated with R map-plotting libraries to generate graphical representation of data, which is more easy to understand and gives a clear idea about the data.



#### References

- http://censusindia.gov.in/2011-prov-results/data\_files/gujarat/statement-1.xls
- https://en.wikipedia.org/wiki/List\_of\_districts\_of\_Gujarat

Some data is not available in 1<sup>st</sup> reference.

# 0.2 Mixture models form one of the most fundamental classes of generative models for clustered data.

For example, run times on unix servers in 100 universities. The following are run times in seconds.

Using mixture normal distribution (a bimodal distribution)

$$0.7*N\{\mu_1, \sigma_1^2\}+0.3*N\{\mu_2, \sigma_2^2\}$$

## A. Find the maximum likelihood estimates of the unknown parameters and their standard errors.

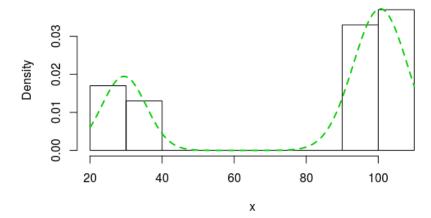
The basic idea of maximum likelihood is to minimize the negative log likelihood inorder to fit the function to the given data, by tweaking the values of parameters, which in this case are,  $\mu_1, \sigma_1^2, \mu_2, \sigma_2^2$ .

### **Output:**

B. Draw a histogram of the data and superimpose the density of the above mixture normal distribution using maximum likelihood estimates of the unknown parameters.

Output:

## Histogram of x



## 0.3 Appendix

### Question 1

```
# R code for Gujrat State
#for Population Density Data:
library (ggplot2)
library (plyr)
library (raster)
library (rgdal)
library (rgeos)
library (sp)
population_density <- read.csv("density.csv", sep = ",")
#fetching map of India from gadm
India <- getData("GADM", country = "India", level = 2)
Gujarat <- subset (India, NAME_1 == "Gujarat")
#converting map of gujarat into a data matrix
map <- fortify (Gujarat);
#casting value of id in map into integer and modifying map
map$id <- as.integer(map$id);</pre>
dat <- data.frame(id = 133:165, district = Gujarat@data$NAME_2);
#joining gujarat map dataframe with the provided population density data
map_df <- join (map, population_density, by ="id", type="inner")
centers <- data.frame(gCentroid(Gujarat, byid = TRUE));
centers$state <- dat$district;</pre>
#plotting the map
ggplot() +
  #creating map having logitude on x-axis & latitude on y-axis,
  #filled with populaiton density data
  geom_polygon(data = map_df, aes(x = long, y = lat, group = group,
                                   fill = Population_Density)) +
  geom_polygon(data = map_df, aes(x = long, y = lat, group = group),
                fill = NA, color = "black", size = 0.25) +
  scale_fill_distiller(palette = "Greens")+
  geom_text(data = centers, aes(label = state, x = x, y = y), size = 2)
  coord_map() + labs(x = "", y = "", title =
  "Districts-wise population density of Gujarat")
Question 2
Part A
library (mclust)
#library used to fit a function to a data
x = c(30.16, 30.36, 97.83, 101.59, 106.42, 30.75, 100.10, 103.30,
101.73, 25.48, 98.90, 31.41, 26.33, 32.35, 96.52, 31.93, 108.32,
99.72, 101.11, 103.92, 97.87, 97.83, 99.22, 97.51, 103.24, 29.31,
29.82, 98.42, 34.28, 27.12, 99.28, 103.77, 102.61, 27.22, 97.71,
105.96, 102.41, 30.38, 101.73, 98.59, 100.14, 99.09, 27.44, 100.37,
99.84, 97.34, 101.17, 99.14, 97.41, 99.92, 101.31, 104.61, 100.71,
```

```
30.62, 103.57, 28.35, 108.12, 100.05, 31.84, 28.80, 98.47, 27.99,
105.05, 33.33, 100.09, 23.57, 101.68, 95.62, 102.10, 98.77, 100.93,
98.68, 27.00, 102.04, 100.88, 98.79, 102.58, 27.40, 29.01, 29.57,
97.16, 96.60, 105.35, 97.74, 100.97, 101.88, 96.75, 29.01, 98.08,
99.63, 99.41, 101.96, 26.70, 31.66, 98.29, 103.51, 99.28, 99.10,
33.36, 100.36)
#BIC for parameterized Gaussian mixture models
#fitted by EM algorithm initialized by model-based
#hierarchical clustering.
#Here V indicates that data is univariate
BIC = mclustBIC(x, modelNames="V")
#Determines the best model via
#mclustBIC for a given set of model parameterizations
#and numbers of components
Model = mclustModel(x, yBIC)
#printing mu_1 and mu_2 as output
Model$parameters$mean
# printing variance_1 & variance_2
Model$parameters$variance$sigmasq
Part B
#add two more lines written below to above code
hist (x, prob=TRUE)
curve (0.3*dnorm(x,29.41,6.14)+0.7*dnorm(x,100.57,7.49),
col=3, lty=2, lwd=2, add=TRUE)
```