

# mclust analysis of the Old Faithful Dataset

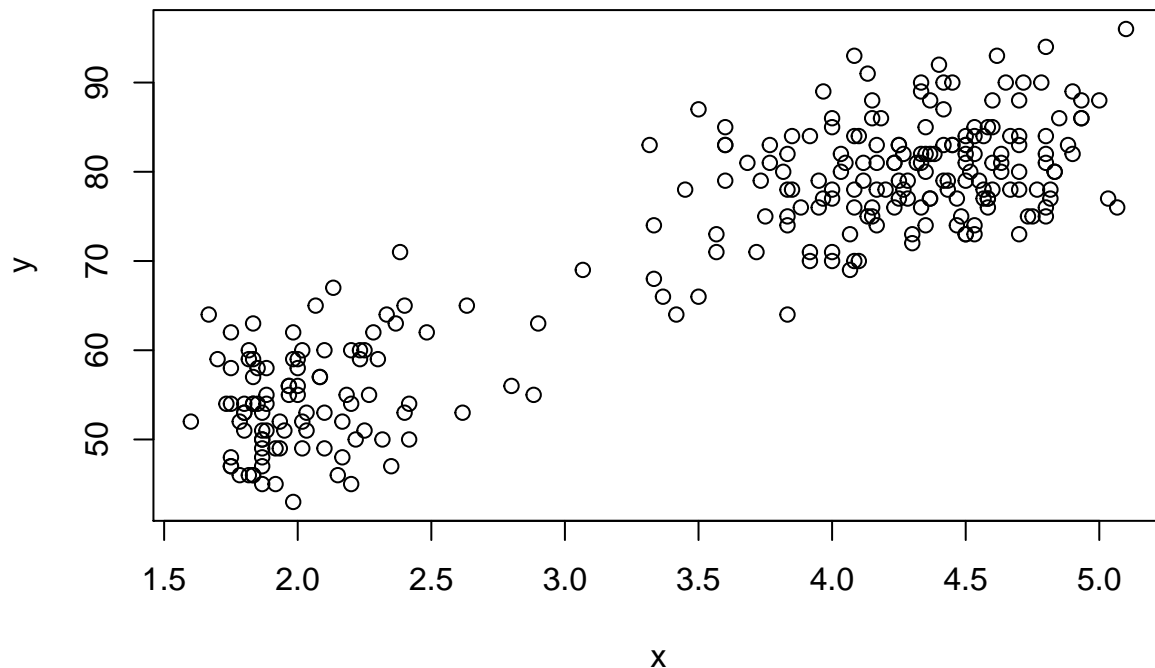
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Wed Dec 3 20:33:47 2014

```
# mclust provides functionality for cluster analysis combining model-based  
# hierarchical clustering, EM for Gaussian mixture models, and BIC.  
# First the cluster analysis estimates the number of clusters that best  
# represents this data set and also the covariance structure of the spread points.  
# This is performed through the technique called Bayesian Information Criterion (BIC)  
# that varies the number of cluster from 1 to 9. The BIC is the value of the  
# maximized loglikelihood measured with a penalty for the number of parameters  
# in the model. Then it's executed the hierarchical clustering technique (HC),  
# which doesn't require a initialization phase. The output of the HC, that is, the  
# cluster that each element belongs, is used to initialize the  
# Expectation-Maximization technique (EM).  
#  
library(mclust)
```

```
## Package 'mclust' version 4.4  
## Type 'citation("mclust")' for citing this R package in publications.
```

```
#  
x = faithful[,1] # get the first column of the faithful data set  
y = faithful[,2] # get the second column of the faithful data set  
plot(x,y) # plot before clustering
```



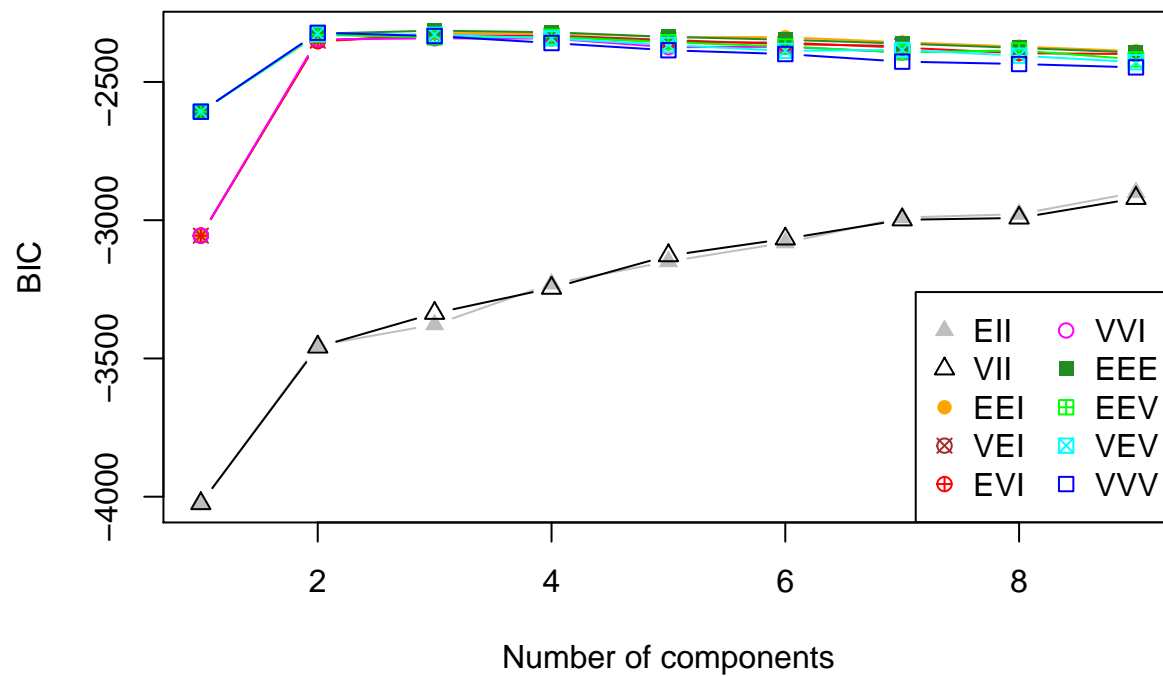
```
#  
## estimate the number of cluster (BIC), initialize (HC) and clusterize (EM)  
faithfulMclust <- Mclust(faithful)  
summary(faithfulMclust)
```

```
## -----
## Gaussian finite mixture model fitted by EM algorithm
## -----
##
## Mclust EEE (ellipsoidal, equal volume, shape and orientation) model with 3 components:
##
##   log.likelihood    n df    BIC    ICL
##             -1126 272 11 -2314 -2361
##
## Clustering table:
##    1    2    3
## 130  97  45
```

```
# In this case, the best model according to BIC is an equal-covariance model
# with 3 components or clusters. A more detailed summary including the
# estimated parameters can be obtained with the following code:
summary(faithfulMclust, parameters = TRUE)
```

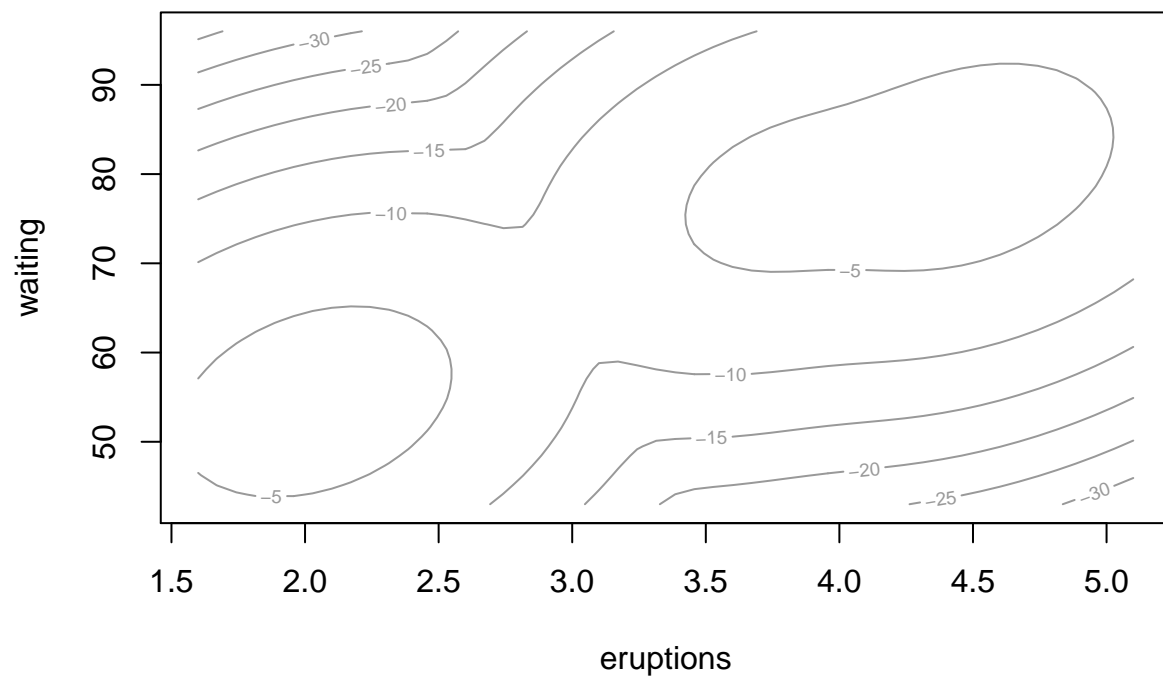
```
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##             -1126 272 11 -2314 -2361
##
## Clustering table:
##    1    2    3
## 130  97  45
##
## Mixing probabilities:
##      1      2      3
## 0.4619 0.3565 0.1816
##
## Means:
##           [,1]  [,2]  [,3]
## eruptions  4.476  2.038  3.82
## waiting   80.892 54.493 77.67
##
## Variances:
## [,,1]
##           eruptions waiting
## eruptions  0.07728  0.4765
## waiting    0.47650 33.7485
## [,,2]
##           eruptions waiting
## eruptions  0.07728  0.4765
## waiting    0.47650 33.7485
## [,,3]
##           eruptions waiting
## eruptions  0.07728  0.4765
## waiting    0.47650 33.7485
```

```
#
plot(faithfulMclust,what="BIC")
```



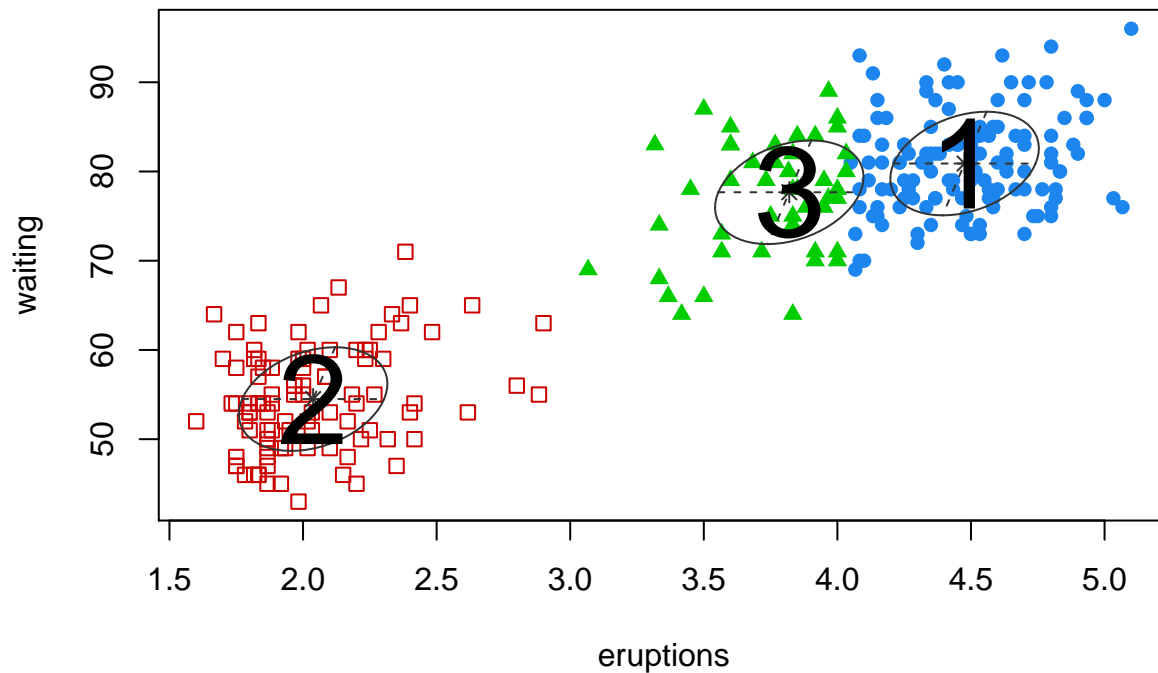
```
plot(faithfulMclust,what="density")
```

### log Density Contour Plot



```
plot(faithfulMclust,what="classification")
clustmeans<-faithfulMclust$parameters$mean
text(clustmeans[1,], clustmeans[2,], seq.int(ncol(clustmeans)), cex=4)
```

## Classification



```
#
# If the next day you experience 2 eruptions with waiting 50 and want to
# classify that value using the existing model, you would use
pp <- predict(faithfulMclust, newdata=data.frame(eruptions=2, waiting=50))
pp$classification
```

```
## [1] 2
```

```
# which gives cluster number 2
# what if you have 4 eruptions with a waiting time of 70 then
pp <- predict(faithfulMclust, newdata=data.frame(eruptions=4, waiting=70))
pp$classification
```

```
## [1] 3
```

```
# this gives cluster 3 which agrees with the plot
#
# The object produced by Mclust is a list with a components describing the
# estimated model. The names of these components can be displayed as follows:
names(faithfulMclust)
```

```
## [1] "call" "data" "modelName" "n"
```

```
## [5] "d"          "G"          "BIC"        "bic"  
## [9] "loglik"     "df"         "hypvol"     "parameters"  
## [13] "z"         "classification" "uncertainty"
```

```
#
```