

Mixture Models

You ZUO

2019/10/14

ex01

```
library(mclust)
```

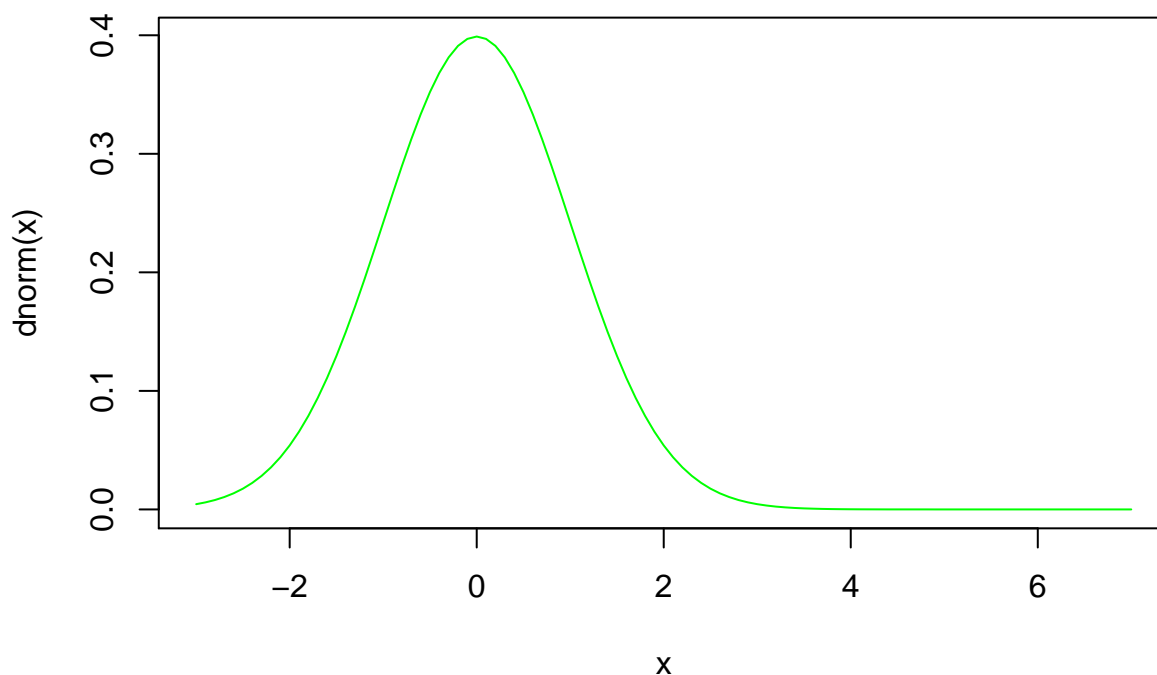
```
## Package 'mclust' version 5.4.3  
## Type 'citation("mclust")' for citing this R package in publications.
```

```
nks <- rmultinom(n = 1, size = 1000, prob = c(1/3, 2/3))
```

```
means <- c(0, 4)  
sds <- c(1, 1/2)  
sample <- mapply(function(nk, mean, sd){  
  rnorm(nk, mean, sd)}, nks, means, sds)
```

```
sample <- unlist(sample)
```

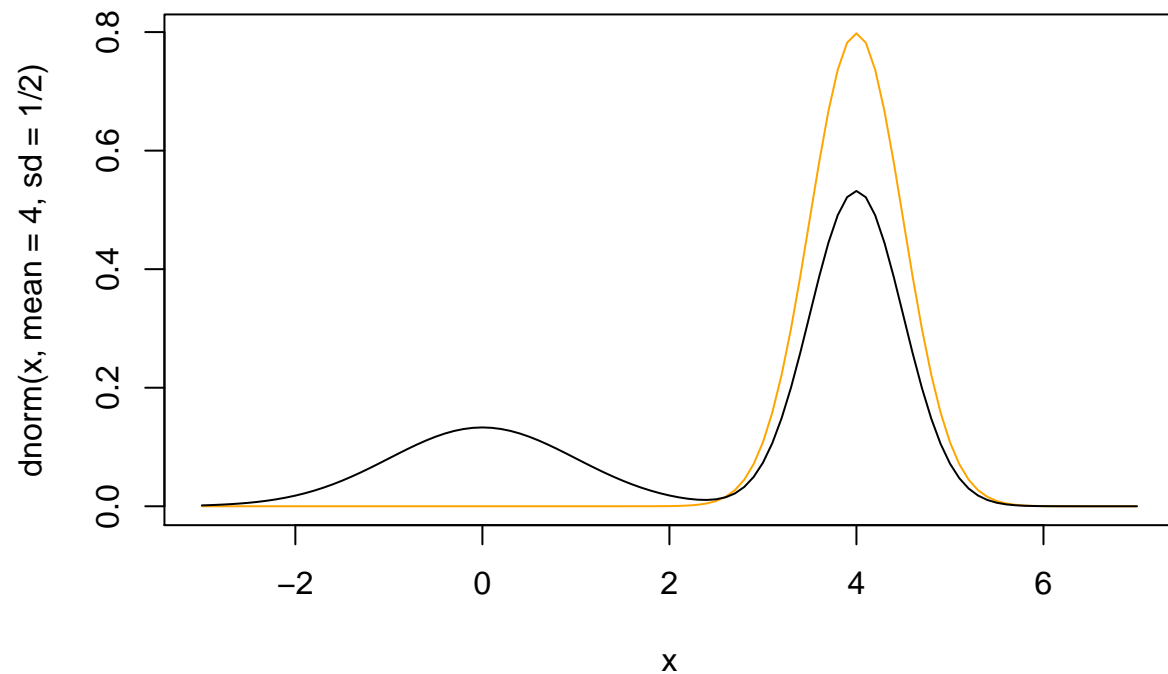
```
curve(dnorm(x), from = -3, to = 7, col = "green")
```



```
curve(dnorm(x, mean = 4, sd = 1/2), from = -3, to = 7, col = "orange")
```

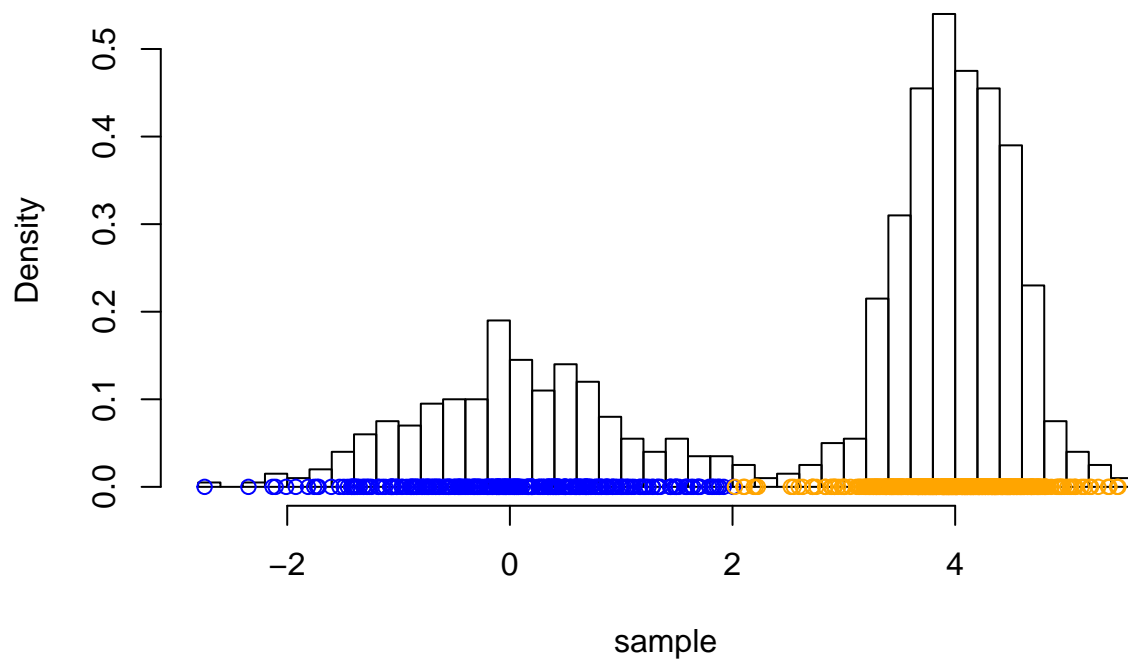
```
mixture <- function(x){  
  1/3*dnorm(x) + 2/3 * dnorm(x,4,1/2)  
}
```

```
curve(mixture(x), -3, 7, add = T)
```

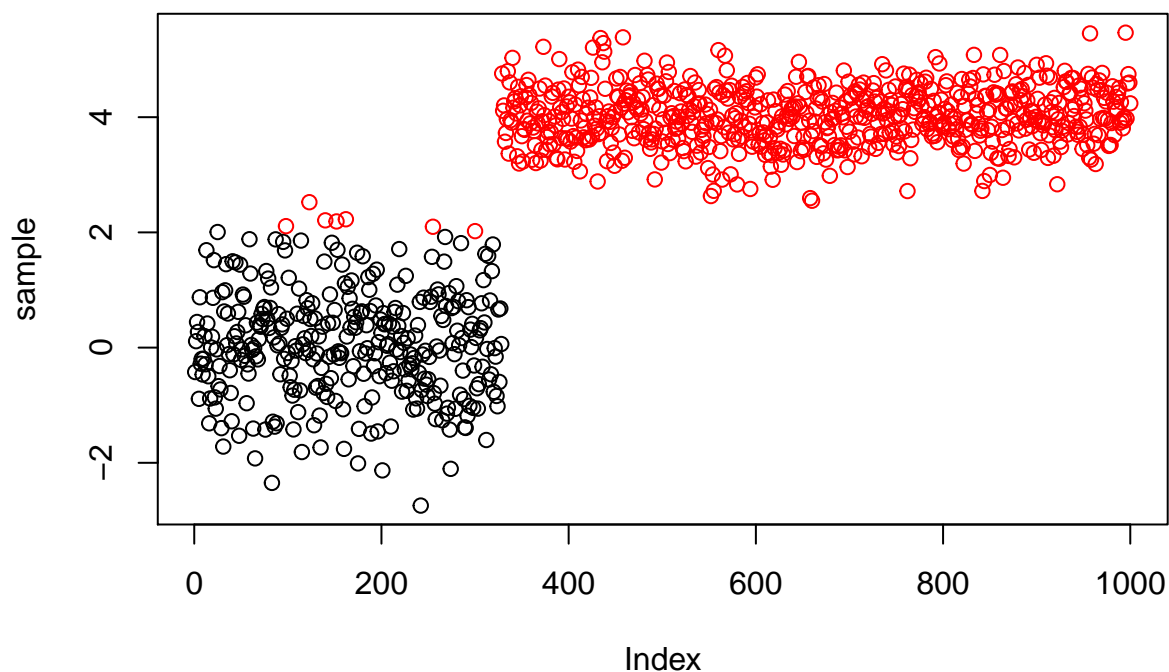


```
cl <- kmeans(x = sample, centers = 2, nstart = 20)
hist(x = sample, probability = T, breaks = 30)
points(sample, rep(0, 1000), col = c("blue", "orange")[cl$cluster])
```

Histogram of sample



```
plot(x = sample, col = cl$cluster)
```



```
list_of_samples <- split(sample, cl$cluster)
lapply(list_of_samples, mean)
```

```
## $`1`
## [1] 0.0273904
##
## $`2`
## [1] 3.995648
```

```
lapply(list_of_samples, sd)
```

```
## $`1`
## [1] 0.9030341
##
## $`2`
## [1] 0.5251815
```

```
# E variances partagées, variances libres
```

```
modelE <- Mclust(data = sample, G = 2, modelNames = "E", warn = F)
```

```
modelV <- Mclust(data = sample, G = 2, modelNames = "V", warn = F)
```

```
summary(modelE)
```

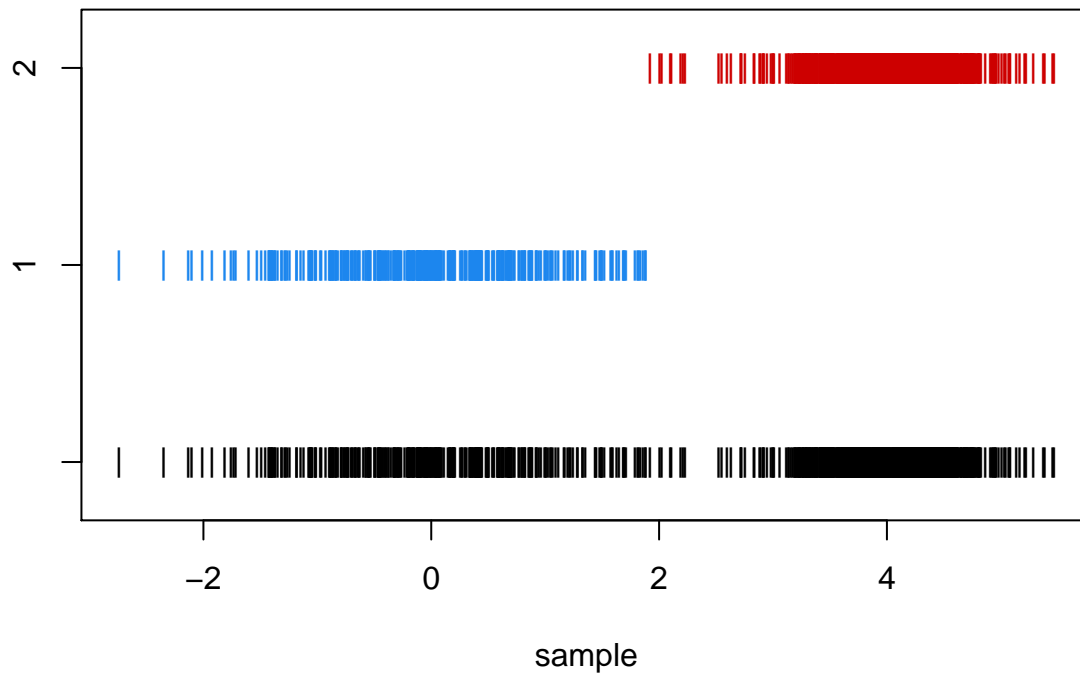
```
## -----
## Gaussian finite mixture model fitted by EM algorithm
## -----
##
## Mclust E (univariate, equal variance) model with 2 components:
##
##   log-likelihood    n df         BIC          ICL
##      -1638.218 1000   4  -3304.067  -3316.516
##
## Clustering table:
##    1    2
```

```
## 319 681
```

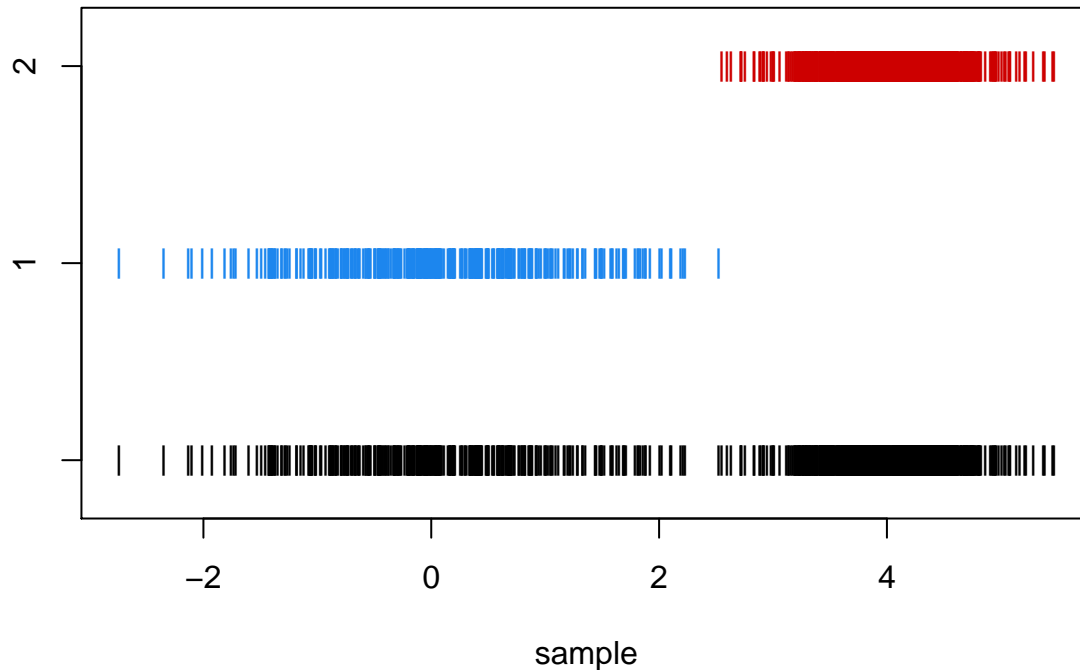
```
summary(modelV)
```

```
## -----  
## Gaussian finite mixture model fitted by EM algorithm  
## -----  
##  
## Mclust V (univariate, unequal variance) model with 2 components:  
##  
##   log-likelihood    n df         BIC          ICL  
##   -1555.845 1000   5 -3146.23 -3153.958  
##  
## Clustering table:  
##    1    2  
## 328 672
```

```
plot(modelE, what = "classification")
```



```
plot(modelV, what = "classification")
```



```
modelV$parameters$mean
```

```
##          1          2
## 0.09037535 4.01760735
```

```
modelV$parameters$variance$sigma
```

```
## [1] 0.9361713 0.2410452
```

```
nks <- rmultinom(n = 1, size = 1000, prob = c(1/3, 2/3))
```

```
means <- c(0, 4)
```

```
sds <- c(1, 1/2)
```

```
sample <- mapply(function(nk, mean, sd){
```

```
  rnorm(nk, mean, sd)}, nks, means, sds)
```

```
x <- data.frame(unlist(sample))
```

```
Init.EM <- function(x, k=2) {
```

```
  # init
```

```
  proportions <- rep(x = 1/k, times = k)
```

```
  variances <- rep(x = 1, times = k)
```

```
  means <- x[sample(1:nrow(x), k), ]
```

```
  parameters <- list(proportions = proportions, variances = variances, means = means)
```

```
  return(parameters)
```

```
}
```

```
x <- as.matrix(x)
```

```
E.step <- function(x, parameters) {
```

```
  K <- length(parameters$means)
```

```
  Tik <- matrix(NA, nrow = nrow(x), ncol = K)
```

```
  for (k in 1:K) {
```

```
    Tik[,k] <- parameters$proportions[k] * dnorm(x[,1], mean = parameters$means[k], sd = parameters$var
```

```

}

return(Tik <- Tik/rowSums(Tik))
}

M.step <- function(x, Tik, parameters) {
  K <- length(parameters$means)
  parameters$proportions <- colSums(Tik)/nrow(x)

  for (k in 1:K) {
    parameters$means[k] <- sum(Tik[,k]*x) / sum(Tik[,k])
    parameters$variances[k] <- sum(Tik[,k]*(x - parameters$means[k])^2)/sum(Tik[,k])
  }

  return(parameters)
}

EM <- function(x, k) {

  parameters <- Init.EM(x, k)
  iter <- 0
  parameters.new <- parameters
  repeat{
    Tik <- E.step(x, parameters)
    parameters <- M.step(x, Tik, parameters)
    if((sum(unlist(parameters.new) - unlist(parameters))^2) / sum(unlist(parameters.new))^2 < 1e-20) br
    parameters.new <- parameters
  }
  return(list(parameters = parameters.new, Tik = Tik))
}

x <- data.frame(x)
# EM(x,2)

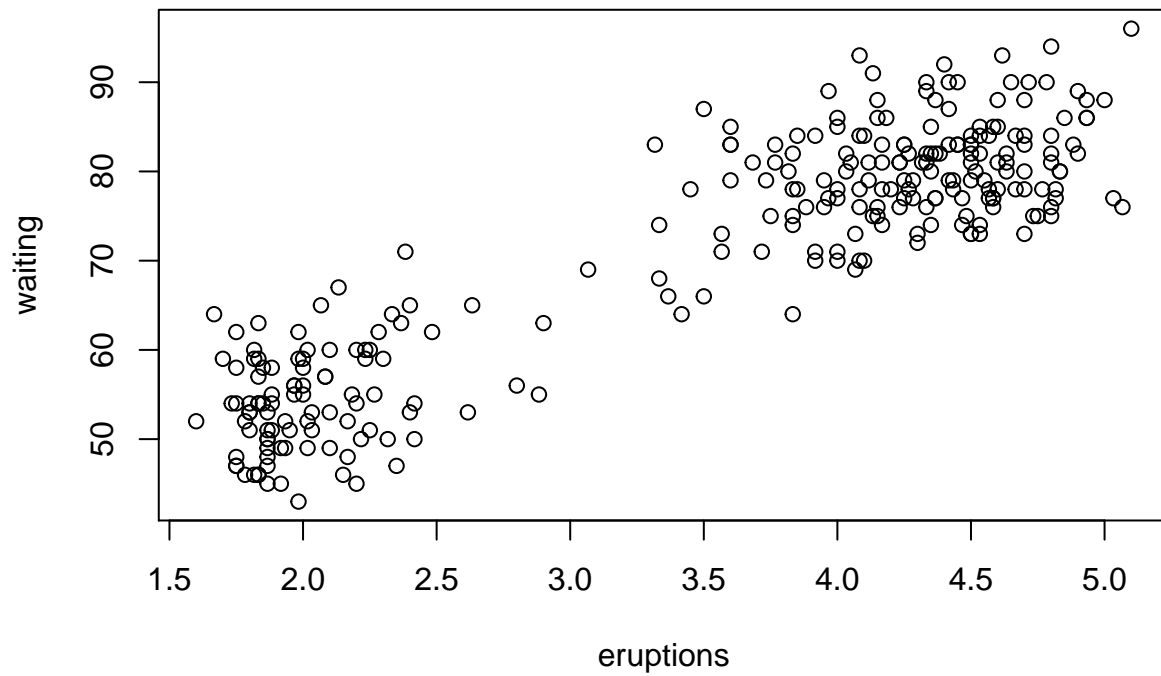
```

Exo2

```

data(faithful)
plot(faithful)

```



```
res <- Mclust(data = faithful)
plot(res)
```

