

# HMM-depmix

*Jai Vrat Singh*

Let us generate data for training and see if depmix can uncover the pattern

some references: [https://eeecon.uibk.ac.at/psychoco/2011/slides/Visser\\_hdt.pdf](https://eeecon.uibk.ac.at/psychoco/2011/slides/Visser_hdt.pdf)

<https://quantstrattrader.wordpress.com/2016/10/05/the-problem-with-depmix-with-online-prediction/>

<https://www.quantstart.com/articles/hidden-markov-models-for-regime-detection-using-r>

<https://cran.r-project.org/web/packages/depmixS4/depmixS4.pdf>

## HMM fitting using depmix

Try fitting now..

```
library(depmixS4)
```

```
## Loading required package: nnet
```

```
## Loading required package: MASS
```

```
## Loading required package: Rsolnp
```

```
hmm <- depmix(obs~1, family = gaussian(), nstates = 2, data = datF)
```

```
hmmfit <- fit(hmm)
```

```
## iteration 0 logLik: -2569.649
```

```
## iteration 5 logLik: -2538.16
```

```
## iteration 10 logLik: -2495.075
```

```
## iteration 15 logLik: -2472.191
```

```
## iteration 20 logLik: -2456.098
```

```
## iteration 25 logLik: -2451.935
```

```
## iteration 30 logLik: -2451.653
```

```
## iteration 35 logLik: -2451.645
```

```
## converged at iteration 38 with logLik: -2451.645
```

```
#the first column has the viterbi states, the other columns have the  
# delta probabilities, see Rabiner (1989)
```

```
post <- hmmfit$posterior
```

```
post2 <- posterior(hmmfit)
```

```
#> head(post)
```

```
# state      S1      S2
```

```
#1      2 0.00000000 1.0000000
```

```
#2      2 0.08230401 0.9176960
```

```
#Must be TRUE as they are same
```

```
identical(post, post2)
```

```
## [1] TRUE
```

```
#
```

```
head(datF)
```

```
## state      obs
```

```
## 1      2 7.3191971
```

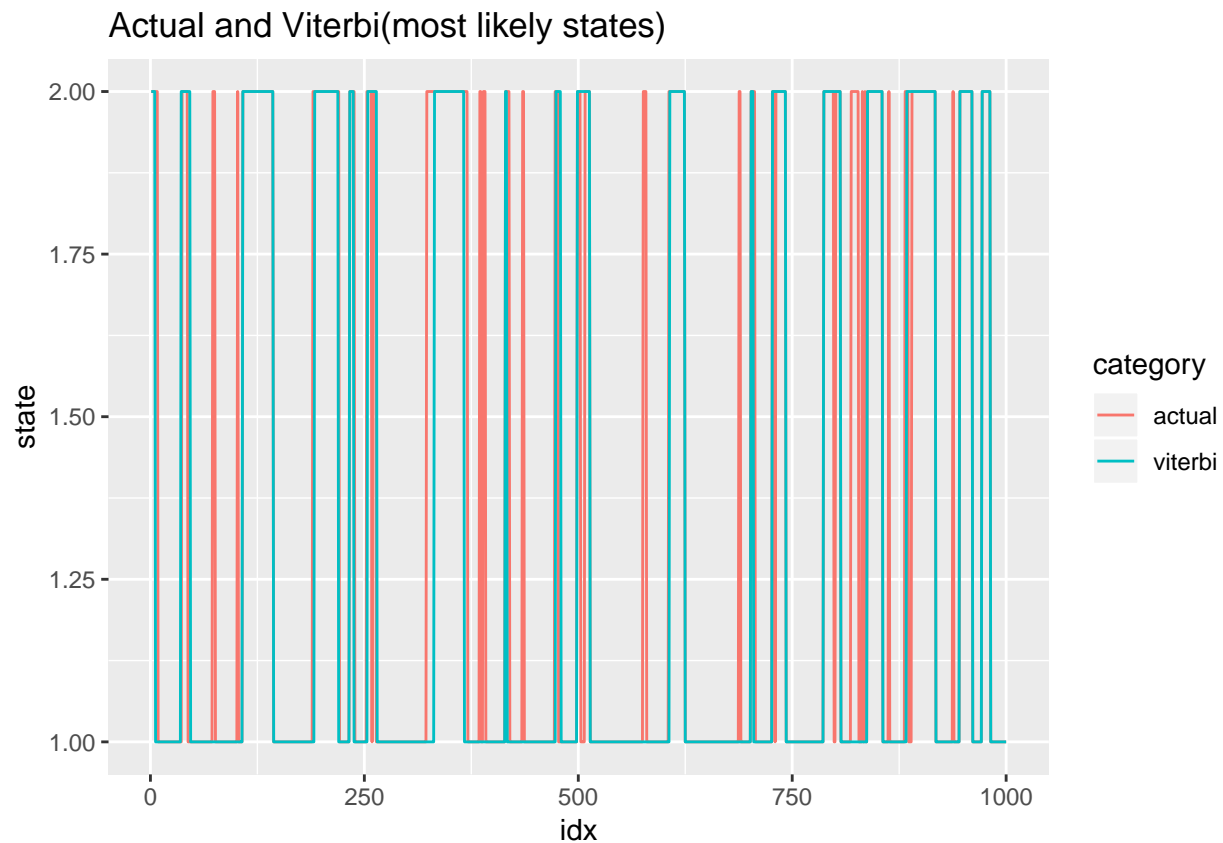
```
## 2      2  8.3811232
## 3      2 -4.1598002
## 4      2  3.9497162
## 5      2  1.9769313
## 6      2  0.7784465
```

```
library(ggplot2)
library(reshape2)
temp <- data.frame("idx" = 1:dim(datF)[1], "state" = datF$state, "category" = "actual")
head(temp)
```

```
##   idx state category
## 1    1     2  actual
## 2    2     2  actual
## 3    3     2  actual
## 4    4     2  actual
## 5    5     2  actual
## 6    6     2  actual
```

```
temp <- rbind( temp,
               data.frame("idx" = 1:dim(post)[1], "state" = post$state, "category" = "viterbi"))
```

```
# Map to color
ggplot(data=temp, aes(x=idx, y=state, group=category, colour=category)) +
  geom_line() +
  ggtitle("Actual and Viterbi(most likely states)")
```



Response

```

hmmfit@response

## [[1]]
## [[1]][[1]]
## Model of type gaussian (identity), formula: obs ~ 1
## Coefficients:
## (Intercept)
##      -1.042706
## sd  2.056819
##
##
## [[2]]
## [[2]][[1]]
## Model of type gaussian (identity), formula: obs ~ 1
## Coefficients:
## (Intercept)
##      1.747935
## sd  4.145138

```

```

#mean
hmmfit@response[[1]][[1]]@parameters$coefficients

```

```

## (Intercept)
##      -1.042706

```

```

#2.047001
#sd
hmmfit@response[[1]][[1]]@parameters$sd

```

```

##      sd
## 2.056819

```

```

#3.989855

#mean
hmmfit@response[[2]][[1]]@parameters$coefficients

```

```

## (Intercept)
##      1.747935

```

```

#-0.9999505
#sd
hmmfit@response[[2]][[1]]@parameters$sd

```

```

##      sd
## 4.145138

```

```

#1.986069

```

We can see that Model is able to uncover the states in graph. Let us plot probabilities.

```

temp <- data.frame("idx"= 1:dim(post)[1], post[,2:3])
temp.melt <- melt(temp, id.vars = "idx")
colnames(temp.melt) <- c("idx", "state", "probability")
# Map to color
ggplot(data=temp.melt, aes(x=idx, y=probability, group=state, colour=state)) +
  geom_line() +
  ggtitle("posterior probabilities")

```

