

Time Series Project

Erling F Steen

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```
library(R.matlab)
```

```
## R.matlab v3.7.0 (2022-08-25 21:52:34 UTC) successfully loaded. See ?R.matlab for help.
```

```
##
```

```
## Attaching package: 'R.matlab'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      getOption, isOpen
```

```
library(depmixS4)
```

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

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

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

```
## Loading required package: nlme
```

```
data <- readMat("Mouse28-140313_BS0150_HMMready.mat")
```

```
angdata <- as.vector(data$resampledAwakeHeadAngleData)
```

```
# Transpose thet data so that there are 71 columns, one column for each cell, and 15243 rows, one for e
```

```
celldata <- t(data$celldata)
```

```
# Exclude cells that are active less than 100 times overall
```

```
col_idx = which(colSums(celldata)<100)
```

```
celldata = celldata[,-col_idx]
```

```
# Reduced to 59 cells
```

```
dim(celldata)
```

```
## [1] 15243    59
```

```

cell_data <- data.frame(celldata)

# Response and list must be on list-form

# Response has to be in a list
res <- list()

# List of families. One for each cell
families <- list()

# Number of columns(cells)
n_cells <- ncol(cell_data)

for (i in 1:n_cells) {
  form <- as.formula(paste(colnames(cell_data)[i], "~ 1",sep=""))
  res[[i]] = form
  families[[i]] = poisson()
}

# Model with 5 states
model5 <- depmix(response = res, nstates = 5, data= cell_data, family = families)
fit5 <- fit(model5)

```

```
## converged at iteration 34 with logLik: -948315.1
```

```
summary(fit5)
```

```

## Initial state probabilities model
## pr1 pr2 pr3 pr4 pr5
## 0 0 0 1 0
##
## Transition matrix
##      toS1 toS2 toS3 toS4 toS5
## fromS1 0.915 0.001 0.052 0.032 0.000
## fromS2 0.001 0.902 0.001 0.042 0.055
## fromS3 0.030 0.000 0.913 0.000 0.056
## fromS4 0.036 0.075 0.000 0.889 0.000
## fromS5 0.000 0.045 0.046 0.000 0.909
##
## Response parameters
## Resp 1 : poisson
## Resp 2 : poisson
## Resp 3 : poisson
## Resp 4 : poisson
## Resp 5 : poisson
## Resp 6 : poisson
## Resp 7 : poisson
## Resp 8 : poisson
## Resp 9 : poisson
## Resp 10 : poisson
## Resp 11 : poisson
## Resp 12 : poisson

```

```

## Resp 13 : poisson
## Resp 14 : poisson
## Resp 15 : poisson
## Resp 16 : poisson
## Resp 17 : poisson
## Resp 18 : poisson
## Resp 19 : poisson
## Resp 20 : poisson
## Resp 21 : poisson
## Resp 22 : poisson
## Resp 23 : poisson
## Resp 24 : poisson
## Resp 25 : poisson
## Resp 26 : poisson
## Resp 27 : poisson
## Resp 28 : poisson
## Resp 29 : poisson
## Resp 30 : poisson
## Resp 31 : poisson
## Resp 32 : poisson
## Resp 33 : poisson
## Resp 34 : poisson
## Resp 35 : poisson
## Resp 36 : poisson
## Resp 37 : poisson
## Resp 38 : poisson
## Resp 39 : poisson
## Resp 40 : poisson
## Resp 41 : poisson
## Resp 42 : poisson
## Resp 43 : poisson
## Resp 44 : poisson
## Resp 45 : poisson
## Resp 46 : poisson
## Resp 47 : poisson
## Resp 48 : poisson
## Resp 49 : poisson
## Resp 50 : poisson
## Resp 51 : poisson
## Resp 52 : poisson
## Resp 53 : poisson
## Resp 54 : poisson
## Resp 55 : poisson
## Resp 56 : poisson
## Resp 57 : poisson
## Resp 58 : poisson
## Resp 59 : poisson
##      Re1.(Intercept) Re2.(Intercept) Re3.(Intercept) Re4.(Intercept)
## St1          0.036          0.224          -2.696          -2.602
## St2         -0.014          0.181          -3.555          -2.131
## St3         -0.078          0.103          -0.845          -2.347
## St4         -0.144          0.362          -1.382          -2.315
## St5          0.100          0.039          -2.590          -1.997
##      Re5.(Intercept) Re6.(Intercept) Re7.(Intercept) Re8.(Intercept)

```

##	St1	-2.802	-2.100	-5.844	-4.197
##	St2	-4.307	-1.763	-6.881	-0.707
##	St3	-2.494	-1.738	-0.344	-0.950
##	St4	-2.244	-2.906	-6.871	-3.780
##	St5	-0.609	-3.568	-2.304	-5.337
##	Re9.(Intercept)	Re10.(Intercept)	Re11.(Intercept)	Re12.(Intercept)	
##	St1	1.798	2.082	1.694	-2.042
##	St2	1.217	1.942	1.562	-1.962
##	St3	1.355	2.022	1.666	-2.458
##	St4	1.810	2.013	1.574	-1.479
##	St5	1.279	1.931	1.348	-3.651
##	Re13.(Intercept)	Re14.(Intercept)	Re15.(Intercept)	Re16.(Intercept)	
##	St1	0.041	0.009	-3.250	0.536
##	St2	-1.309	-0.507	-2.334	0.325
##	St3	-0.953	-0.598	-0.156	0.647
##	St4	-0.923	0.845	-3.339	0.518
##	St5	-1.541	-0.271	-3.213	0.358
##	Re17.(Intercept)	Re18.(Intercept)	Re19.(Intercept)	Re20.(Intercept)	
##	St1	-3.659	-2.487	-3.521	-1.211
##	St2	-1.360	-3.962	-5.513	-3.065
##	St3	-3.845	-2.909	-0.879	-1.598
##	St4	-3.456	-3.438	-6.177	-2.930
##	St5	-3.764	-3.592	-2.264	-2.719
##	Re21.(Intercept)	Re22.(Intercept)	Re23.(Intercept)	Re24.(Intercept)	
##	St1	0.308	1.737	-3.213	-1.288
##	St2	-0.240	1.788	0.427	1.152
##	St3	-0.588	1.828	-3.260	-0.943
##	St4	1.098	1.626	-2.425	-0.897
##	St5	-0.412	1.512	-2.877	0.274
##	Re25.(Intercept)	Re26.(Intercept)	Re27.(Intercept)	Re28.(Intercept)	
##	St1	1.693	-2.364	-2.063	1.910
##	St2	1.511	-1.194	-2.039	1.775
##	St3	1.878	-2.594	-2.192	1.886
##	St4	1.722	-2.614	-2.148	2.035
##	St5	1.684	-2.579	-2.180	1.817
##	Re29.(Intercept)	Re30.(Intercept)	Re31.(Intercept)	Re32.(Intercept)	
##	St1	1.915	1.315	0.814	-1.757
##	St2	1.471	1.288	0.717	-1.772
##	St3	1.750	1.301	0.865	-1.512
##	St4	1.771	1.282	0.716	-1.513
##	St5	1.467	1.241	0.907	-1.602
##	Re33.(Intercept)	Re34.(Intercept)	Re35.(Intercept)	Re36.(Intercept)	
##	St1	-0.483	-1.224	0.519	-0.863
##	St2	-2.525	-1.382	0.579	-1.379
##	St3	-1.367	-1.128	0.605	-1.295
##	St4	-3.971	-1.163	0.479	-1.139
##	St5	-2.736	-1.346	0.425	-1.909
##	Re37.(Intercept)	Re38.(Intercept)	Re39.(Intercept)	Re40.(Intercept)	
##	St1	1.972	-1.139	-2.890	-4.467
##	St2	1.976	-0.819	-2.935	0.134
##	St3	1.910	-1.993	-2.314	-3.944
##	St4	1.975	-0.314	-0.433	-2.204
##	St5	1.801	-1.364	-2.596	-3.421
##	Re41.(Intercept)	Re42.(Intercept)	Re43.(Intercept)	Re44.(Intercept)	

```
## St1      -2.363      0.575      -0.883      -2.177
## St2       0.822     -2.894     -1.254     -4.293
## St3     -2.958     -0.212     -2.344      2.486
## St4     -3.017      2.079     -1.991     -6.465
## St5       1.076     -2.459     -1.571     -1.227
##      Re45.(Intercept) Re46.(Intercept) Re47.(Intercept) Re48.(Intercept)
## St1      -1.741      -0.143     -3.596     -5.587
## St2     -1.554       0.863     -4.587     -4.315
## St3     -3.439     -0.091      0.781      0.039
## St4     -0.724     -0.263     -3.400     -3.196
## St5     -1.425       0.953      0.753      0.310
##      Re49.(Intercept) Re50.(Intercept) Re51.(Intercept) Re52.(Intercept)
## St1      -5.133     -1.326      1.414     -3.076
## St2     -0.019     -4.326     -5.116      0.801
## St3     -5.766     -2.955     -4.579     -1.085
## St4     -0.859     -1.429     -1.607      1.985
## St5     -2.296     -2.805     -3.589     -0.427
##      Re53.(Intercept) Re54.(Intercept) Re55.(Intercept) Re56.(Intercept)
## St1      -0.887     -1.425     -2.845     -5.230
## St2     -5.455     -2.267     -4.010     -0.757
## St3     -4.273     -2.148     -4.006     -0.042
## St4     -1.375     -2.500     -2.511     -0.784
## St5     -3.904     -0.400     -2.910      2.277
##      Re57.(Intercept) Re58.(Intercept) Re59.(Intercept)
## St1     -1.030       0.052     -0.648
## St2     -4.318     -3.324     -0.401
## St3      2.774     -4.277     -0.822
## St4     -6.465     -2.409     -0.461
## St5     -0.841     -4.503     -0.652
```

```
AIC(model15)
```

```
## [1] 4579341
```

Fit models with more than 5 states

```
# Dette tar tid!
# Model with 10 states
mod10 <- depmix(response = res, nstates = 10, data = cell_data, family = families)
fit10 <- fit(mod10)
# Model with 15 states
mod15 <- depmix(response = res, nstates = 15, data = cell_data, family = families)
fit15 <- fit(mod15)
```

```
#summary(fit10)
```

Model selection based on AIC

```
# Model selection based on AIC
```

```
AIC(fit5)
```

```
## [1] 1897268
```

```
# AIC(fit10)
```

```
# AIC(fit15)
```

```
# AIC(fit20)
```

Indices of each state

```
# States for all time intervals
```

```
states = posterior(fit5)$state
```

```
## Warning in .local(object, ...): Argument 'type' not specified and will default  
## to 'viterbi'. This default may change in future releases of depmixS4. Please  
## see ?posterior for alternative options.
```

```
# Indices for which state is 1
```

```
idx_state1 <- which(states == 1)
```

```
idx_state2 <- which(states == 2)
```

```
idx_state3 <- which(states == 3)
```

```
idx_state4 <- which(states == 4)
```

```
idx_state5 <- which(states == 5)
```

```
#Average for of
```

```
avg_state1 <- mean(angdata[idx_state1], na.rm = TRUE)
```

```
avg_state2 <- mean(angdata[idx_state2], na.rm = TRUE)
```

```
avg_state3 <- mean(angdata[idx_state3], na.rm = TRUE)
```

```
avg_state4 <- mean(angdata[idx_state4], na.rm = TRUE)
```

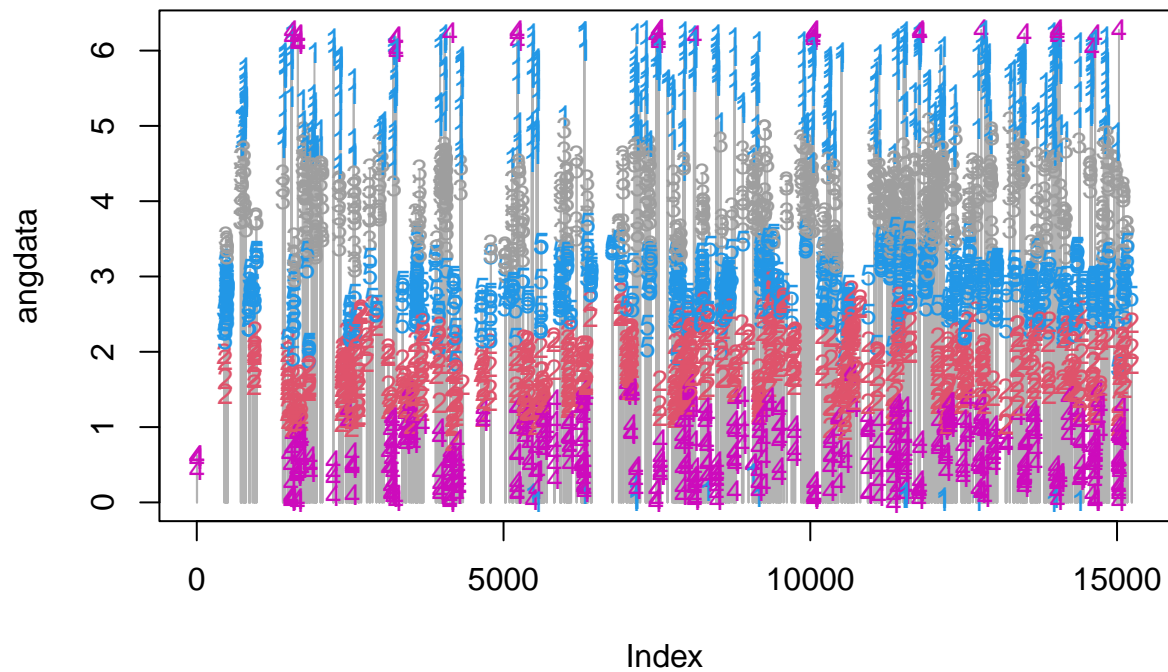
```
avg_state5 <- mean(angdata[idx_state5], na.rm = TRUE)
```

```
avg <- c(avg_state1, avg_state2, avg_state3, avg_state4, avg_state5)
```

Histogram over head direction for each state

```
plot(angdata, main="", type="h", col = gray(.7))
```

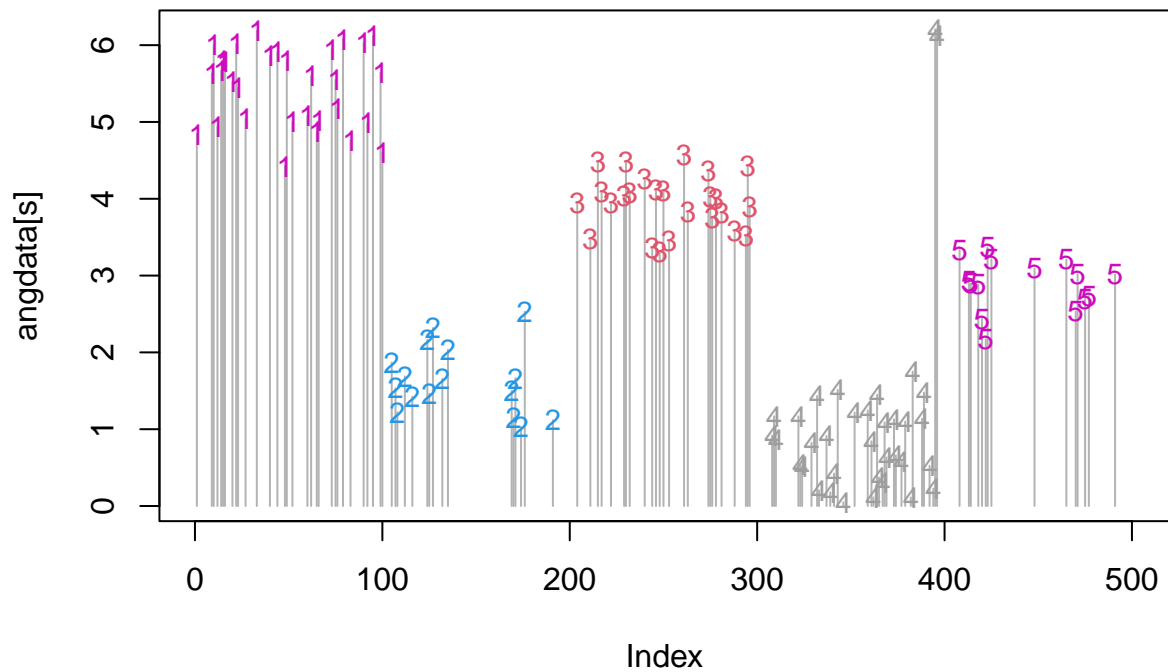
```
text(angdata, col=6*states-2, labels=states, cex=.9)
```



Difficult to see. Take a random sample of the angle data

```
# Don't plot all head angles. Take a random sample. Easier to visualize
s_1 = sample(idxx_state1, 100)
s_2 = sample(idxx_state2, 100)
s_3 = sample(idxx_state3, 100)
s_4 = sample(idxx_state4, 100)
s_5 = sample(idxx_state5, 100)
s=c(s_1, s_2, s_3, s_4, s_5)
```

```
plot(angdata[s], type = "h", col=gray(.7))
text(angdata[s], col = 6*states[s], labels = states[s], cex=.9)
```



Plotting the average head direction of each state

```
## Plot of mean head
avg
```

```
## [1] 5.255357 1.773650 4.004686 1.037438 2.848121
```

```
plot(avg, type="h")
```