

Univariate Series fitting of HMM

```
> hmmData<-read.table("C:/Users/bill/Desktop/HMMdata.txt",header=TRUE)
```

```
> y<-hmmData[2]
```

```
> library(depmixS4)
```

Loading required package: nnet

Loading required package: MASS

Loading required package: Rsolnp

```
> set.seed(1)
```

```
> ModelData<-data.frame(y)
```

```
> colnames(ModelData)<-"y"
```

```
> HMM<-depmix(y~1,data=ModelData,nstates=2,family=gaussian())
```

```
> HMMfit<-fit(HMM,verbose = FALSE)
```

converged at iteration 40 with logLik: -487.9615

```
> print(HMMfit)
```

Convergence info: Log likelihood converged to within tol. (relative change)

'log Lik.' -487.9615 (df=7)

AIC: 989.9229

BIC: 1013.011

```
> summary(HMMfit)
```

Initial state probabilities model

pr1 pr2

0 1

Transition matrix

toS1 toS2

fromS1 0.994 0.006

fromS2 0.074 0.926

Response parameters

Resp 1 : gaussian

Re1.(Intercept) Re1.sd

St1 94.548 2.730

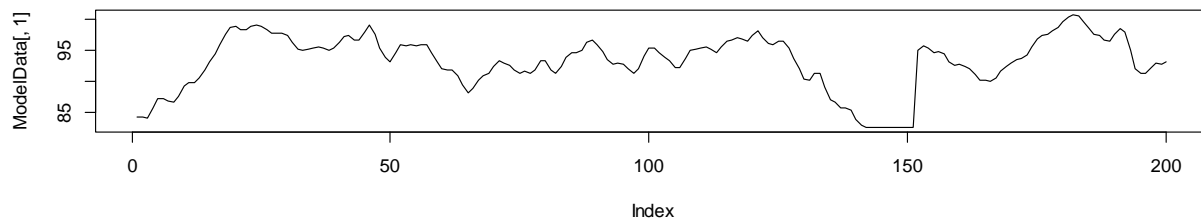
St2 84.705 2.033

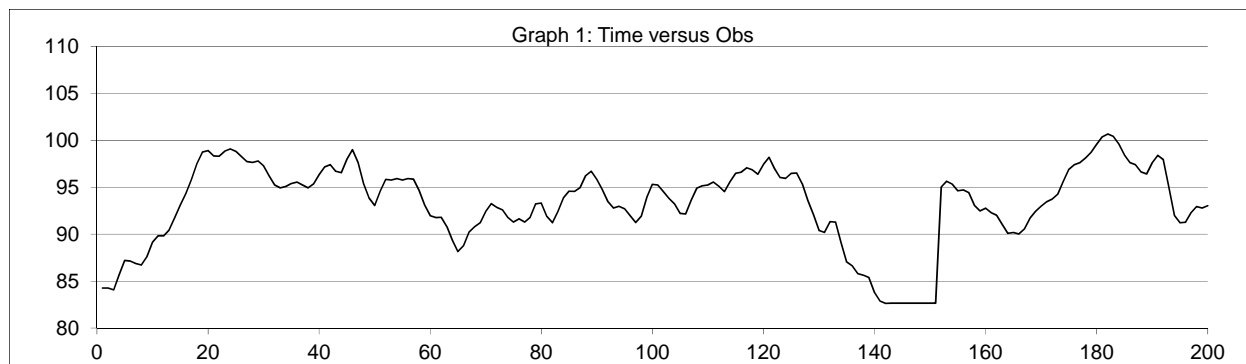
```
> HMMpost<-posterior(HMMfit)
```

```
> head(HMMpost)
```

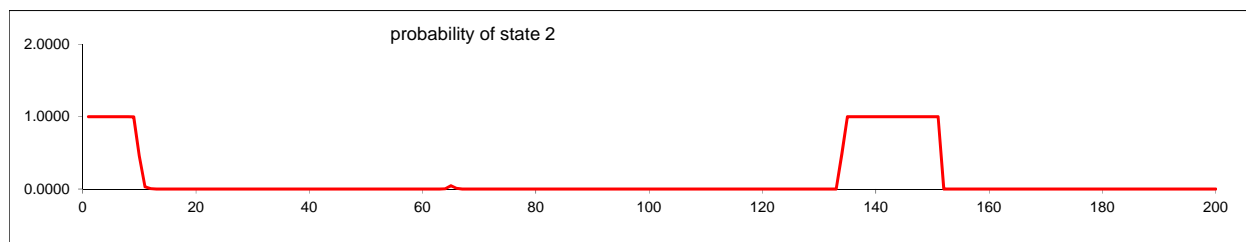
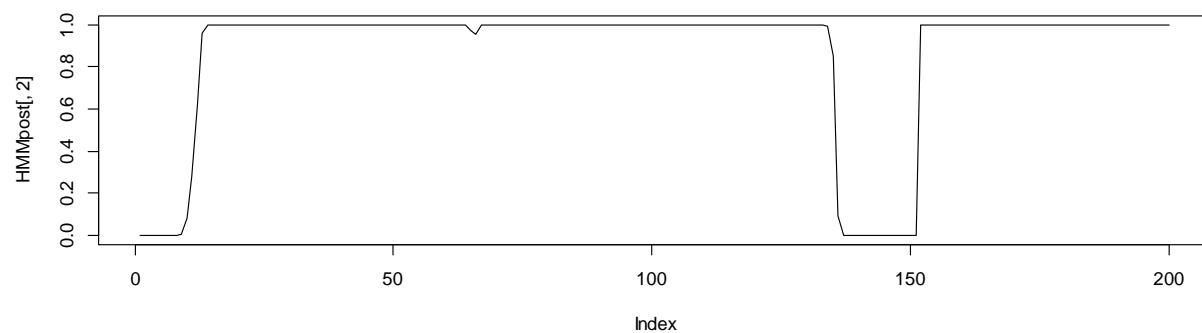
	state	S1	S2
1	2	0.000000e+00	1.0000000
2	2	5.104915e-05	0.9999490
3	2	3.926793e-05	0.9999607
4	2	3.301796e-04	0.9996698
5	2	3.429852e-03	0.9965701
6	2	2.999896e-03	0.9970001

```
> plot(ModelData[,1],type="l")
```





```
> plot(HMMpost[,2],type="l")
```



Fitting a Multivariate Normal HMM

```
> hmmData<-read.table("C:/Users/bill/Desktop/MHMM.txt",header=TRUE)
> x<-hmmData[2]
> y<-hmmData[3]
> library(depmixS4)
Loading required package: nnet
Loading required package: MASS
Loading required package: Rsolnp
> ModelData<-data.frame(x,y)
> z<-data.matrix(ModelData)

> rModels<-list()
> rModels[[1]]<-list(MVNresponse(z~1))
> rModels[[2]]<-list(MVNresponse(z~1))
> trstart=c(0.9,0.1,0.1,0.9)
> transition<-list()
> transition[[1]]<-transInit(~1,nstates=2,data=data.frame(1),pstart=c(trstart[1:2]))
> transition[[2]]<-transInit(~1,nstates=2,data=data.frame(1),pstart=c(trstart[3:4]))
> instart=runif(2)
> inMod<-transInit(~1,ns=2,ps=instart,data=data.frame(1))
> mod<- makeDepmix(response=rModels,transition=transition,prior=inMod)

> fm<-fit(mod,emc=em.control(random=FALSE))
iteration 0 logLik: -1033.003
iteration 5 logLik: -814.8728
converged at iteration 9 with logLik: -814.8716
```

> summary(fm)

Initial state probabilities model

St1 St2

(Intercept) 0 10.165

Transition matrix

toS1 toS2
fromS1 0.981 0.019
fromS2 0.037 0.963

Transition probabilities		
	1	2
1	0.96	0.04
2	0.03	0.97

Response parameters

Re1.coefficients1 Re1.coefficients2 Re1.Sigma1 Re1.Sigma2 Re1.Sigma3
St1 8.082 10.070 3.715 -2.944 4.512
St2 9.982 4.664 15.147 6.805 3.854

Estimated mean vector, Covariance, Correlation matrix

State 1

$$\mu_1 = \begin{bmatrix} 8.082 \\ 10.070 \end{bmatrix}, \Sigma_1 = \begin{bmatrix} 3.715 & -2.944 \\ -2.944 & 4.512 \end{bmatrix}, P_1 = \begin{bmatrix} 1 & -0.71892 \\ -0.71892 & 1 \end{bmatrix}$$

$$\sigma_1 = 1.927434 \quad \rho = -0.71892 \quad \sigma_2 = 2.124618$$

State 2

$$\mu_2 = \begin{bmatrix} 9.982 \\ 4.664 \end{bmatrix}, \Sigma_2 = \begin{bmatrix} 15.147 & 6.805 \\ 6.805 & 3.854 \end{bmatrix}, P_2 = \begin{bmatrix} 1 & 0.890423 \\ 0.890423 & 1 \end{bmatrix}$$

$$\sigma_1 = 3.891915 \quad \rho = 0.890423 \quad \sigma_2 = 1.96367$$

Parameters

State	μ_1	σ_1	μ_2	σ_2	ρ
1	10	4	5	2	0.9
2	8	2	10	2	-0.7

> fmpost<-posterior(fm)

> plot(fmpost[,2],type="l")

