

Ejemplos

Lizbeth Naranjo Albarrán y Luz Judith Rodríguez Esparza

Paper: *Modelos ocultos de Markov:*

una aplicación de estimación Bayesiana para series de tiempo financieras

Authors: Lizbeth Naranjo Albarrán & Luz Judith Rodríguez Esparza

Journal: Mixba'al

Year: 2023

<https://github.com/lizbethna/HMMBayes.git>

Este archivo muestra las instrucciones para correr los códigos de R y Stan.

Cadenas de Markov

```
library(ggplot2)
library(extraDistr)
library(rstan)
```

Calcular probabilidades

```
### Datos
N = 100 #tamaño de muestra
K = 4 # estados
A = matrix(0,4,4) # matriz de probabilidades de transicion
A[1,] = c(0.3, 0.3, 0, 0.4) # simplex: acelerar
A[2,] = c(0.2, 0.4, 0, 0.4) # simplex: constante
A[3,] = c(0.7, 0, 0.3, 0) # simplex: reposo
A[4,] = c(0.4, 0.1, 0.4, 0.1) # simplex: freno
rowSums(A) # renglones suman 1
```

```
[1] 1 1 1 1
```

```
di1 = c(0,0,0,1) # probabilidades del estado oculto inicial
```

```
# Funcion para calcular la distribucion estacionaria delta1
distr_estac = function(A){
  n = nrow(A)
  B = A - diag(n) # Subtract the identity to the input matrix
  B[,1] = rep(1,n) # Replace a column of ones
```

```

b = c(1,rep(0,n-1)) # Create the output vector (1,0,0,...,0)
di1 = solve(t(B),b) # Solve the system for di1
return(di1)
}

```

```

# distribucion estacionaria
(estac = distr_estac(A))

```

```
[1] 0.3634476 0.2253375 0.1495327 0.2616822
```

```

# tiempo medio de recurrencia
(tiempo = 1/estac)

```

```
[1] 2.751429 4.437788 6.687500 3.821429
```

```

# probabilidad de observaciones
prob_obs <- function(x1,A,di1){
  n = length(x1)
  px1 = rep(NA,n)
  px1[1] = di1[x1[1]]
  for(i in 2:n){
    px1[i] = A[x1[i-1],x1[i]]
  }
  prod(px1)
}
x1 = c(4,4,4,1,1,4,2,4)
prob_obs(x1,A,di1)

```

```
[1] 1.92e-05
```

Simular datos

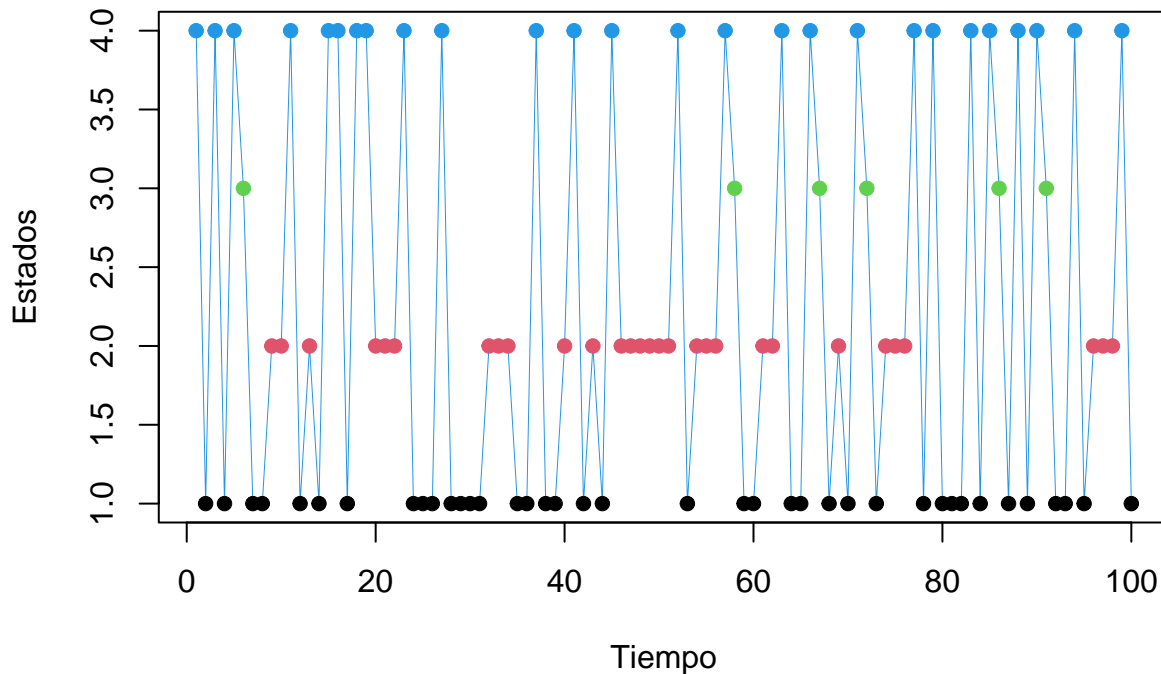
```

N = 100 # tamaño de muestra

# Generar muestra de una cadena de Markov
# T = tamaño de la cadena de Markov
# A = matriz de transicion
CM_genera <- function(N,A,di1) {
  K = ncol(A) #= nrow(A)
  z <- vector("numeric", N)
  z[1] <- sample(1:K, size = 1, prob = di1)
  for (t in 2:N)
    z[t] <- sample(1:K, size = 1, prob = A[z[t - 1], ])
  list(z = z,
       theta = list(di1 = di1, A = A))
}

cadena = CM_genera(N,A,di1)
plot(cadena$z, type="o",col=cadena$z,lwd=0.1,pch=19,
     xlab="Tiempo", ylab="Estados")

```



Código Stan

Dada una muestra observada, se busca estimar las probabilidades de transición.

```
datos <- list( "z"=cadena$z, "N"=N, "K"=K, # muestra
              "alpha"=rep(1,K)) # valores iniciales de la distribucion inicial
param = c("gama") # parametros a estimar

fit_cm <- stan("cadenas_markov.stan", data=datos,
              chains=2, warmup=1000, iter=2000, thin=2)
```

Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c

clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/L

In file included from <built-in>:1:

In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/StanHeaders/include

In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/E

In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/E

/Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/Eigen/src/Core/util/Ma

namespace Eigen {

~

/Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/Eigen/src/Core/util/Ma

namespace Eigen {

~

;

In file included from <built-in>:1:

In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/StanHeaders/include

In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/E

/Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/Eigen/Core:96:10: fata

#include <complex>

~~~~~

3 errors generated.

make: \*\*\* [foo.o] Error 1

SAMPLING FOR MODEL 'cadenas\_markov' NOW (CHAIN 1).

Chain 1:

Chain 1: Gradient evaluation took 4.6e-05 seconds

Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.46 seconds.

Chain 1: Adjust your expectations accordingly!

Chain 1:

Chain 1:

Chain 1: Iteration: 1 / 2000 [ 0%] (Warmup)

Chain 1: Iteration: 200 / 2000 [ 10%] (Warmup)

Chain 1: Iteration: 400 / 2000 [ 20%] (Warmup)

Chain 1: Iteration: 600 / 2000 [ 30%] (Warmup)

Chain 1: Iteration: 800 / 2000 [ 40%] (Warmup)

Chain 1: Iteration: 1000 / 2000 [ 50%] (Warmup)

Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)

Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)

Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)

Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)

Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)

Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)

Chain 1:

Chain 1: Elapsed Time: 0.182916 seconds (Warm-up)

Chain 1: 0.151923 seconds (Sampling)

Chain 1: 0.334839 seconds (Total)

Chain 1:

SAMPLING FOR MODEL 'cadenas\_markov' NOW (CHAIN 2).

Chain 2:

Chain 2: Gradient evaluation took 2.1e-05 seconds

Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.21 seconds.

Chain 2: Adjust your expectations accordingly!

Chain 2:

Chain 2:

Chain 2: Iteration: 1 / 2000 [ 0%] (Warmup)

Chain 2: Iteration: 200 / 2000 [ 10%] (Warmup)

Chain 2: Iteration: 400 / 2000 [ 20%] (Warmup)

Chain 2: Iteration: 600 / 2000 [ 30%] (Warmup)

Chain 2: Iteration: 800 / 2000 [ 40%] (Warmup)

Chain 2: Iteration: 1000 / 2000 [ 50%] (Warmup)

Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)

Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)

Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)

Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)

Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)

Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)

Chain 2:

Chain 2: Elapsed Time: 0.173797 seconds (Warm-up)

Chain 2: 0.15055 seconds (Sampling)

Chain 2: 0.324347 seconds (Total)

Chain 2:

## Resultados

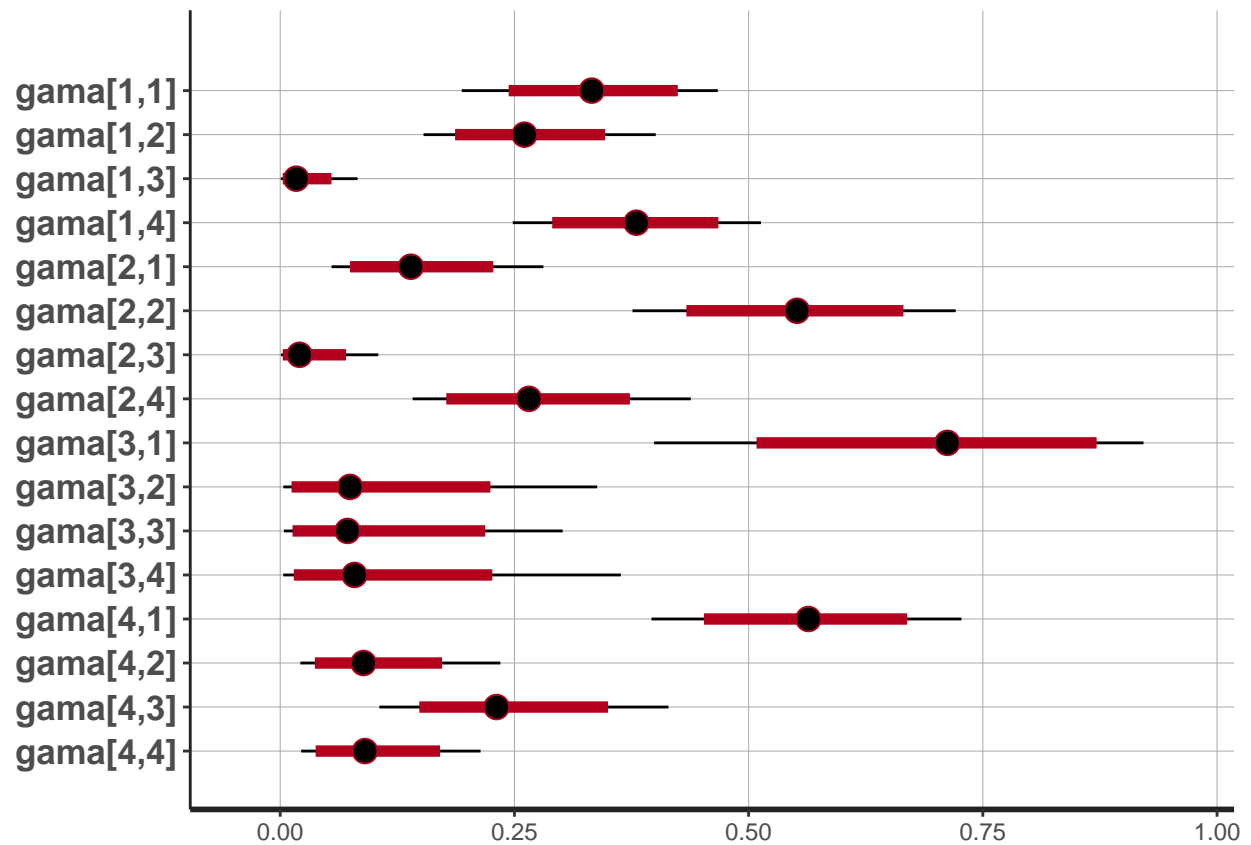
```
print(fit_cm, pars=param)
```

Inference for Stan model: cadenas\_markov.  
2 chains, each with iter=2000; warmup=1000; thin=2;  
post-warmup draws per chain=500, total post-warmup draws=1000.

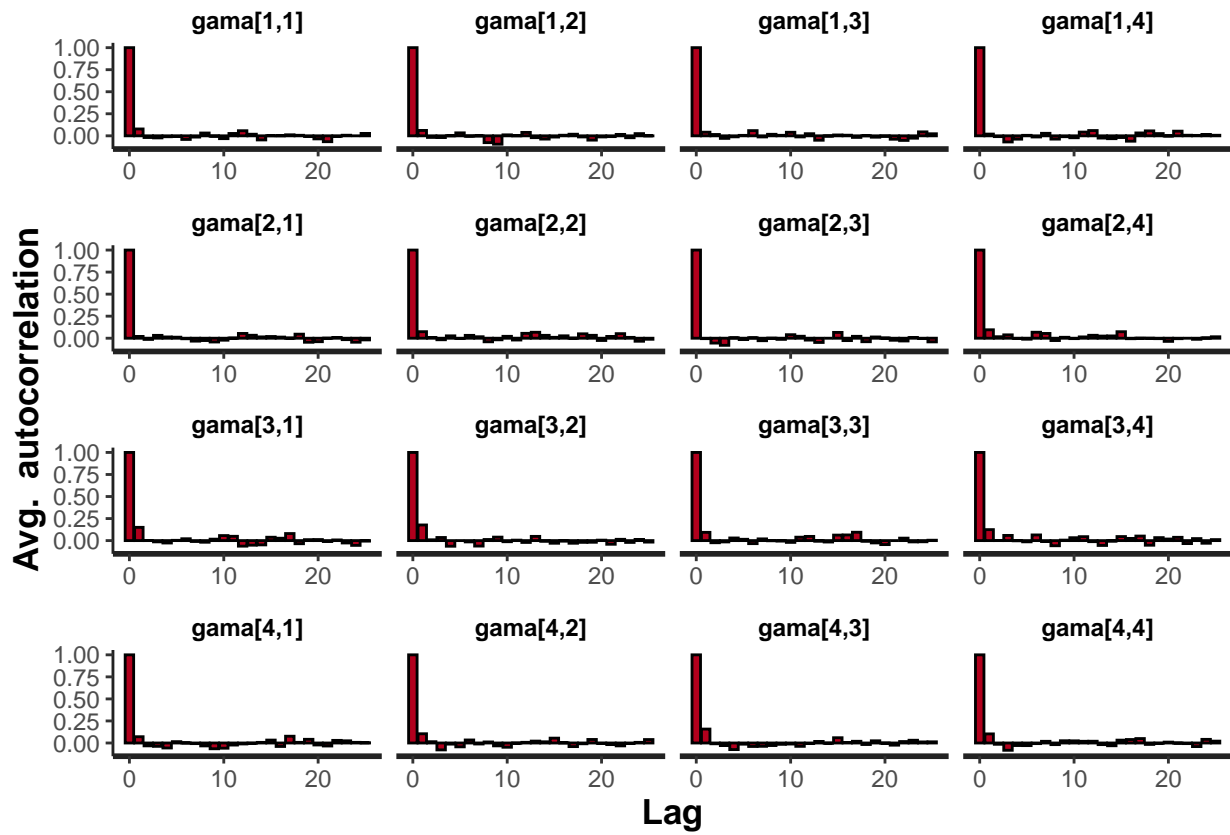
|           | mean | se_mean | sd   | 2.5% | 25%  | 50%  | 75%  | 97.5% | n_eff | Rhat |
|-----------|------|---------|------|------|------|------|------|-------|-------|------|
| gama[1,1] | 0.33 | 0.00    | 0.07 | 0.19 | 0.29 | 0.33 | 0.38 | 0.47  | 854   | 1    |
| gama[1,2] | 0.26 | 0.00    | 0.06 | 0.15 | 0.22 | 0.26 | 0.31 | 0.40  | 886   | 1    |
| gama[1,3] | 0.02 | 0.00    | 0.02 | 0.00 | 0.01 | 0.02 | 0.03 | 0.08  | 937   | 1    |
| gama[1,4] | 0.38 | 0.00    | 0.07 | 0.25 | 0.33 | 0.38 | 0.42 | 0.51  | 958   | 1    |
| gama[2,1] | 0.15 | 0.00    | 0.06 | 0.05 | 0.10 | 0.14 | 0.18 | 0.28  | 865   | 1    |
| gama[2,2] | 0.55 | 0.00    | 0.09 | 0.38 | 0.49 | 0.55 | 0.61 | 0.72  | 869   | 1    |
| gama[2,3] | 0.03 | 0.00    | 0.03 | 0.00 | 0.01 | 0.02 | 0.04 | 0.10  | 989   | 1    |
| gama[2,4] | 0.27 | 0.00    | 0.08 | 0.14 | 0.22 | 0.27 | 0.32 | 0.44  | 773   | 1    |
| gama[3,1] | 0.70 | 0.01    | 0.14 | 0.40 | 0.61 | 0.71 | 0.80 | 0.92  | 766   | 1    |
| gama[3,2] | 0.10 | 0.00    | 0.09 | 0.00 | 0.03 | 0.07 | 0.14 | 0.34  | 701   | 1    |
| gama[3,3] | 0.10 | 0.00    | 0.09 | 0.00 | 0.03 | 0.07 | 0.14 | 0.30  | 835   | 1    |
| gama[3,4] | 0.10 | 0.00    | 0.09 | 0.00 | 0.04 | 0.08 | 0.15 | 0.36  | 736   | 1    |
| gama[4,1] | 0.56 | 0.00    | 0.08 | 0.40 | 0.50 | 0.56 | 0.62 | 0.73  | 851   | 1    |
| gama[4,2] | 0.10 | 0.00    | 0.06 | 0.02 | 0.06 | 0.09 | 0.13 | 0.23  | 816   | 1    |
| gama[4,3] | 0.24 | 0.00    | 0.08 | 0.11 | 0.18 | 0.23 | 0.28 | 0.41  | 756   | 1    |
| gama[4,4] | 0.10 | 0.00    | 0.05 | 0.02 | 0.06 | 0.09 | 0.13 | 0.21  | 829   | 1    |

Samples were drawn using NUTS(diag\_e) at Tue Mar 14 18:46:25 2023.  
For each parameter, n\_eff is a crude measure of effective sample size,  
and Rhat is the potential scale reduction factor on split chains (at  
convergence, Rhat=1).

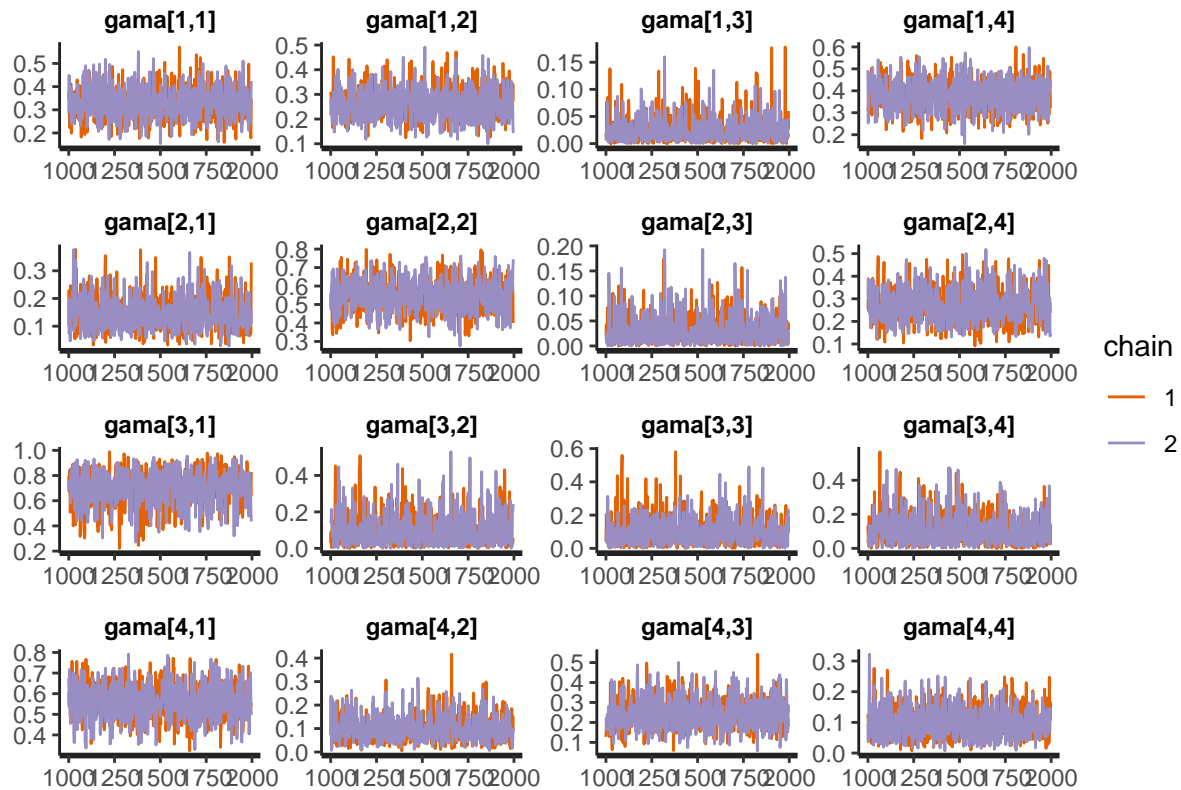
```
stan_plot(fit_cm, pars=param)
```



```
stan_ac(fit_cm, pars=param)
```



```
stan_trace(fit_cm, pars=param)
```



```
stan_dens(fit_cm, pars="gama", point_est = "mean", show_density = TRUE) +
  ggtitle(expression(paste("Distribución final de ", Gamma))) +
  ylab("Densidad") +
  theme(axis.title.x=element_text(size=14), axis.title.y=element_text(size=14),
        plot.title = element_text(size=16))
```

## Distribución final de $\Gamma$

