# Ejemplos

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Paper: Modelos ocultos de Markov:

una aplicación de estimación Bayesiana para series de tiempo financieras Authors: Lizbeth Naranjo Albarrán & Luz Judith Rodríguez Esparza

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

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

# Distribución Gamma-Poisson

```
## Warning: package 'ggplot2' was built under R version 4.1.2

library(extraDistr)
library(rstan)

## Loading required package: StanHeaders

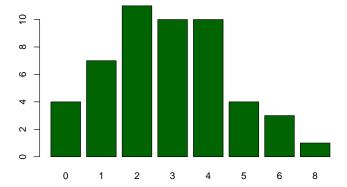
## rstan (Version 2.21.3, GitRev: 2e1f913d3ca3)

## For execution on a local, multicore CPU with excess RAM we recommend calling
## options(mc.cores = parallel::detectCores()).

## To avoid recompilation of unchanged Stan programs, we recommend calling
## rstan_options(auto_write = TRUE)
```

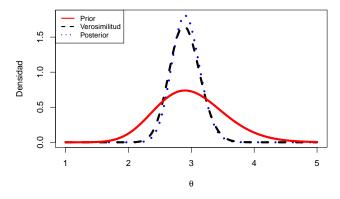
## Simular datos

```
N = 50  # tamaño de muestra
theta = 3  # parametro de media
a0 = 30; b0 = 10  # hiperparametros de la distribucion inicial
set.seed(12345)
x = rpois(N,theta)  # x ~ Poisson(theta)
barplot(table(x), nclass=10, col="darkgreen")
```

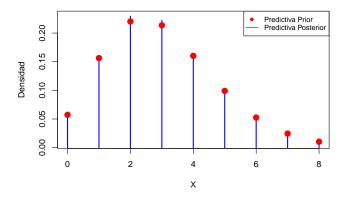


# Graficas de las distribuciones final y predictiva final

### Muestra suma X=145 N=50



#### Muestra suma X=145 N=50



datos <- list( "x"=x, "N"=N, # muestra

Chain 1: Gradient evaluation took 1.4e-05 seconds

1 / 2000 [ 0%]

Chain 1: Adjust your expectations accordingly!

Chain 1: Chain 1:

Chain 1: Iteration:

# Código Stan

```
param = c("theta", "x_star") # parametros a estimar
fit_dist <- stan("dist_poisson_gamma.stan", data=datos,</pre>
                             chains=2, warmup=1000, iter=2000, thin=2)
Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
\verb|clang -mmacosx-version-min=10.13 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -ITMAKE -I
                                                                                                                                                                                                                                               -I"/L
In file included from <built-in>:1:
In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/StanHeaders/includ
In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/
In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/
/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/includ
In file included from /Library/Frameworks/R.framework/Versions/4.1/Resources/library/RcppEigen/include/
/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 'dist_poisson_gamma' NOW (CHAIN 1).
```

"a0"=a0,"b0"=b0) # valores iniciales de la distribucion inicial

(Warmup)

Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.14 seconds.

```
Chain 1: Iteration: 200 / 2000 [ 10%]
                                         (Warmup)
                                         (Warmup)
Chain 1: Iteration: 400 / 2000 [ 20%]
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: 0.009626 seconds (Warm-up)
Chain 1:
                        0.008987 seconds (Sampling)
Chain 1:
                        0.018613 seconds (Total)
Chain 1:
SAMPLING FOR MODEL 'dist_poisson_gamma' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 3e-06 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.03 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: 0.010003 seconds (Warm-up)
Chain 2:
                        0.008384 seconds (Sampling)
Chain 2:
                        0.018387 seconds (Total)
Chain 2:
```

## Resultados

```
print(fit_dist, pars=param)

Inference for Stan model: dist_poisson_gamma.

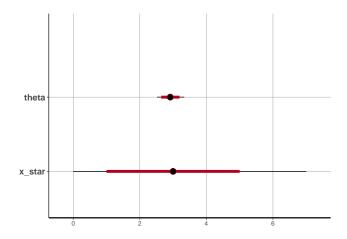
2 chains, each with iter=2000; warmup=1000; thin=2;
post-warmup draws per chain=500, total post-warmup draws=1000.

mean se_mean sd 2.5% 25% 50% 75% 97.5% n_eff Rhat
theta 2.92 0.01 0.22 2.51 2.77 2.91 3.05 3.34 630 1
```

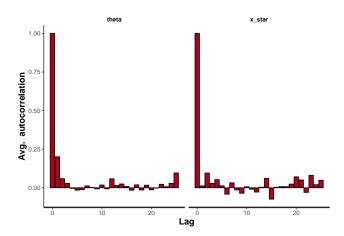
x\_star 2.83 0.07 1.72 0.00 2.00 3.00 4.00 7.00 697 1

Samples were drawn using NUTS(diag\_e) at Tue Mar 14  $18:47:04\ 2023$ . 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).

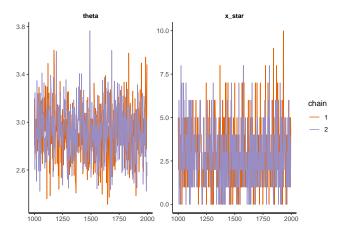
# stan\_plot(fit\_dist,pars=param)



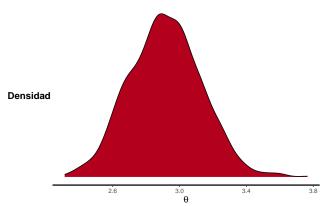
stan\_ac(fit\_dist,pars=param)



stan\_trace(fit\_dist,pars=param)



### Distribución final de $\theta$



## Distribución predictiva final de x<sup>star</sup>

