

# DOBAD Package: Gibbs Sampling MCMC of Linear Birth-Death Chain with Partial Data

Charles Doss

September 2009

## Part I

# Estimating Rates for Linear Birth-Death chain via Gibbs Sampler MCMC by Exact Conditional Simulation

We are demonstrating the use of the `DOBAD` package's capability to do Bayesian estimation of the rate parameters for a linear Birth-Death chain, given partial observations, using the methods of Doss et al. (2010). Call the chain  $\{X(t)\}_{t \in \mathbb{R}}$ , and its birth rate  $la$  and its death rate  $\mu$ . We fix  $\beta \in \mathbb{R}$  and constrain  $\nu$ , the immigration rate, to be  $\nu = \beta la$ . We will denote  $\theta = (la, \mu)$ . The data is the value of the process at a finite number of discrete time points. That is, for some fixed times  $0 = t_0, t_1, \dots, t_n$ , we see the state of the process,  $X(t_i)$ . Thus the data,  $D$ , is 2 parts: a vector of the times  $t_i$ ,  $i = 0, \dots, n$  and a vector of states at each of those times,  $s_i$ , for  $i = 0, \dots, n$  (where  $X(t_i) = s_i$ ). The gamma prior is the conjugate prior if we observed the chain continuously instead of partially. The way we proceed, then, is to use independent Gamma priors on the  $\lambda$  and  $\mu$  and augment the state space for our MCMC to include the entire chain  $\{X_t\}_{t \in [0, t_n]}$  by conditionally sampling  $\{X_t\}_{t \in [0, t_n]}; \theta | D$ .

First we generate the underlying process and the “data”, set our prior parameters, and compute some summary statistics of the fully observed and partially observed processes.

```
> library(DOBAD)
> initstate = 7
> set.seed(112)
> T = 5
> L <- 0.2
> mu <- 0.4
> beta.immig <- 0.987
```

```

> trueParams <- c(L, mu, beta.immig)
> names(trueParams) <- c("lambda", "mu", "beta")
> dr <- 1e-10
> n.fft <- 1024
> delta <- 1
> dat <- birth.death.simulant(t = T, lambda = L, mu = mu, nu = L *
+   beta.immig, X0 = initstate)
> fullSummary <- BDsummaryStats(dat)
> fullSummary

      Nplus   Nminus Holdtime
12.00000 14.00000 26.73947

> MLEs <- M.step.SC(EMsuffStats = fullSummary, T = T, beta.immig = beta.immig)
> MLEs

lambdahat   muhat
0.3788540 0.5235706

> partialData <- getPartialData(seq(0, T, delta), dat)
> observedSummary <- BDsummaryStats.PO(partialData)
> observedSummary

      Nplus   Nminus Holdtime
        3         5        28

> L.mean <- 1
> M.mean <- 1.1
> aL <- 0.02
> bL <- aL/L.mean
> aM <- 0.022
> bM <- aM/M.mean
> print(paste("Variances are", aL/bL^2, "and", aM/bM^2))

```

```
[1] "Variances are 50 and 55"
```

```
> N = 100
```

```
> burn = 0
```

Now we run the MCMC. It is set to run only 100 iterations, which is obviously not enough for estimation, but does demonstrate the code.

```
> timer <- system.time(theMCMC <- BD.MCMC.SC(Lguess = L.mean, Mguess = M.mean,  
+   alpha.L = aL, beta.L = bL, alpha.M = aM, beta.M = bM, beta.immig = beta.immig,  
+   data = partialData, burnIn = burn, N = N))
```

```
[1] "BD.MCMC.SC: On the 30 th iteration params are 0.50525582705533 0.643698845152378"
```

```
[1] "BD.MCMC.SC: On the 60 th iteration params are 0.408368213696412 0.406325363782436"
```

```
[1] "BD.MCMC.SC: On the 90 th iteration params are 0.271395764173334 0.513698779398225"
```

```
> mean(theMCMC[, 1])
```

```
[1] 0.4252525
```

```
> mean(theMCMC[, 2])
```

```
[1] 0.5536474
```

```
> L
```

```
[1] 0.2
```

```
> mu
```

```
[1] 0.4
```

```
> timer
```

```
   user  system elapsed  
11.253   0.060   13.002
```

```

> options(continue = " ")

> hist(theMCMC[, 1], freq = FALSE, breaks = 20, xlab = "Lambda",
      ylab = "Density", main = "Posterior of Lambda")
> Lmean <- mean(theMCMC[, 1])
> abline(col = "red", v = Lmean)
> abline(col = "purple", v = L.mean)
> x <- seq(from = 0, to = 1, by = 0.01)
> y <- dgamma(x, shape = aL, rate = bL)
> lines(x, y, col = "blue")

> hist(theMCMC[, 2], freq = FALSE, breaks = 20, xlab = "Mu", ylab = "Density",
      main = "Posterior of Mu")
> Mmean <- mean(theMCMC[, 2])
> abline(col = "red", v = Mmean)
> abline(col = "purple", v = M.mean)
> x <- seq(from = 0, to = 1, by = 0.01)
> y <- dgamma(x, shape = aM, rate = bM)
> lines(x, y, col = "blue")

```

## References

Doss, C., Suchard, M., Holmes, I., Kato-Maeda, M., and Minin, V. (2010). Great Expectations: EM Algorithms for Discretely Observed Linear Birth-Death-Immigration Processes. *Arxiv preprint arXiv:1009.0893*.

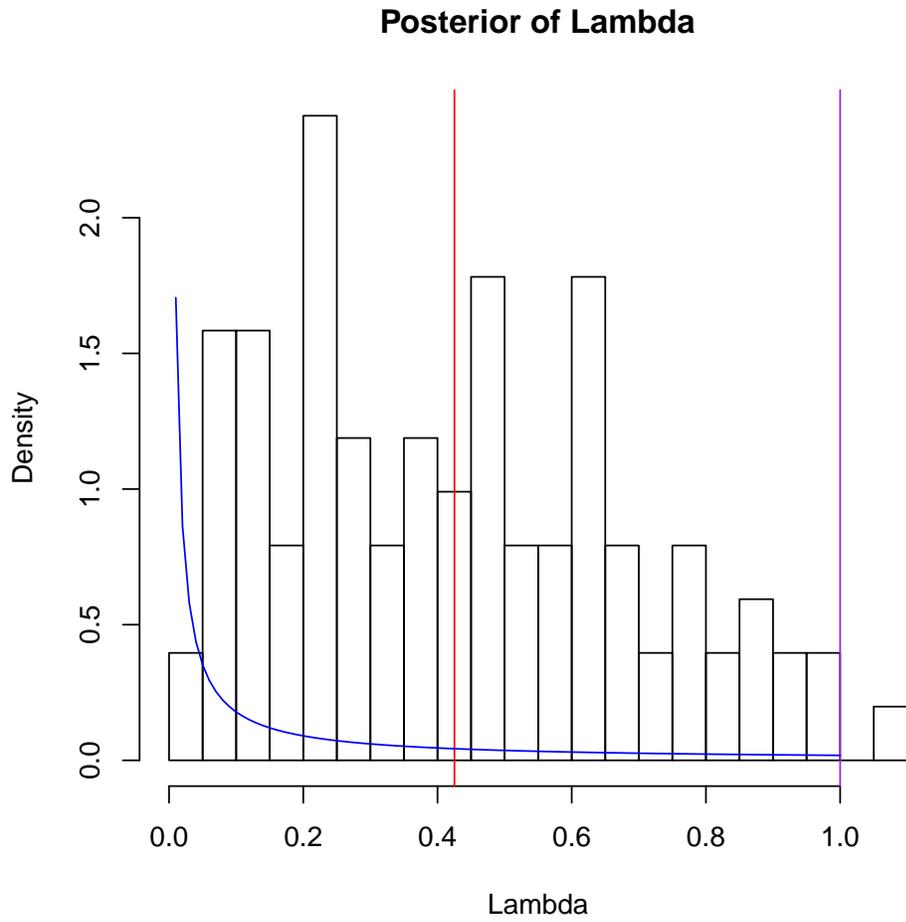


Figure 1: Posterior Density Estimation of Lambda

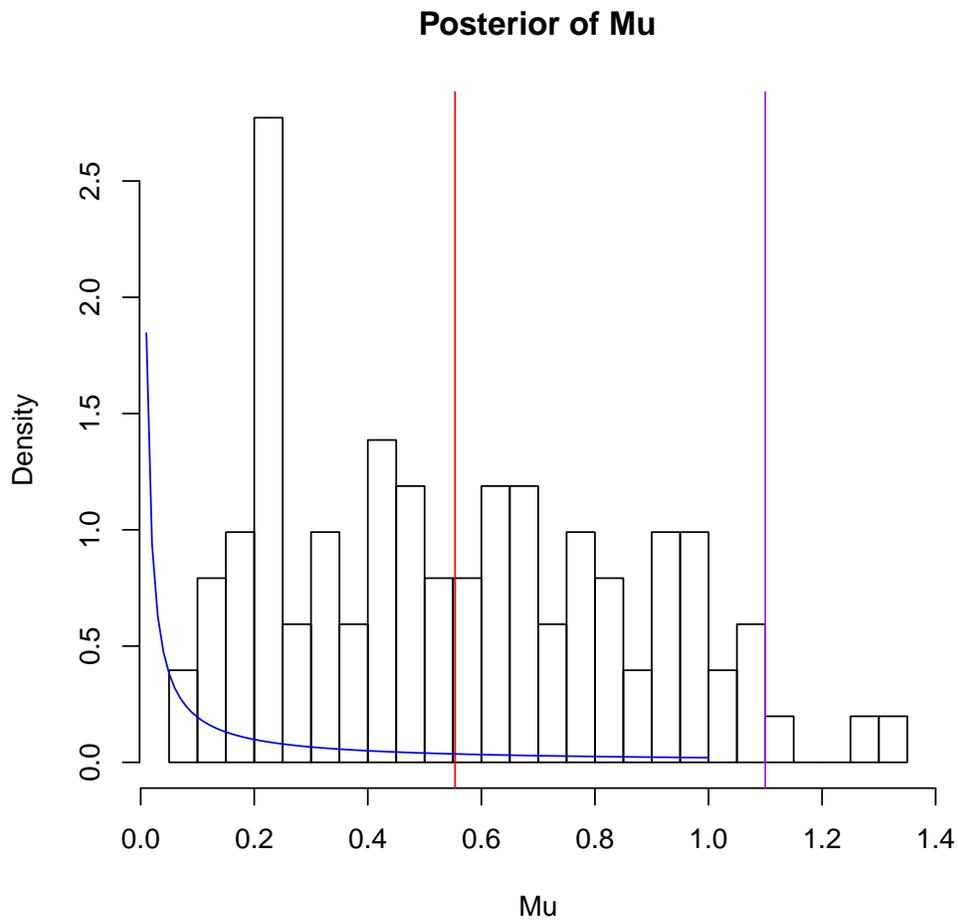


Figure 2: Posterior Density Estimation of Mu