MCMC

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```
require("HI")
```

```
## Loading required package: HI
require("MCMCpack")

## Loading required package: MCMCpack

## Loading required package: coda

## Loading required package: MASS

## ## ## Markov Chain Monte Carlo Package (MCMCpack)

## ## Copyright (C) 2003-2018 Andrew D. Martin, Kevin M. Quinn, and Jong Hee Park

## ##
## ## Support provided by the U.S. National Science Foundation
```

We know that:

##

$$x|\delta, \mu_1, \mu_2, \sigma_1^2, \sigma_2^2 \sim \delta N(\mu_1, \sigma_1^2) + (1 - \delta)N(\mu_2, \sigma_2^2)$$

The prior distributions for five parameters are:

(Grants SES-0350646 and SES-0350613)

$$\pi(\delta) = 1$$

$$\mu_1 \sim N(0, 100)$$

$$\mu_2 \sim N(0, 100)$$

$$\sigma_1^2 \sim IG(0.5, 10)$$

$$\sigma_2^2 \sim IG(0.5, 10)$$

The full conditional distribution :

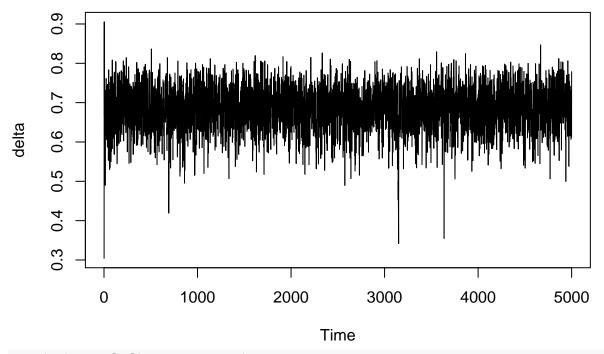
$$f(\delta, \mu_1, \mu_2, \sigma_1^2, \sigma_2^2 | \mathbf{x}) \propto \prod_{i=1}^n \left(\delta \frac{1}{\sqrt{2\pi\sigma_1^2}} exp \left\{ -\frac{(x_i - \mu_1)^2}{2\sigma_1^2} \right\} + (1 - \delta) \frac{1}{\sqrt{2\pi\sigma_2^2}} exp \left\{ -\frac{(x_i - \mu_2)^2}{2\sigma_2^2} \right\} \right)$$

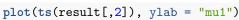
$$\times exp \left\{ -\frac{\mu_1^2}{200} \right\} \times exp \left\{ -\frac{\mu_2^2}{200} \right\} \times (\sigma_1^2)^{-1.5} exp \left\{ -\frac{(\sigma_1^2)^2}{10} \right\} \times (\sigma_2^2)^{-1.5} exp \left\{ -\frac{(\sigma_2^2)^2}{10} \right\}$$

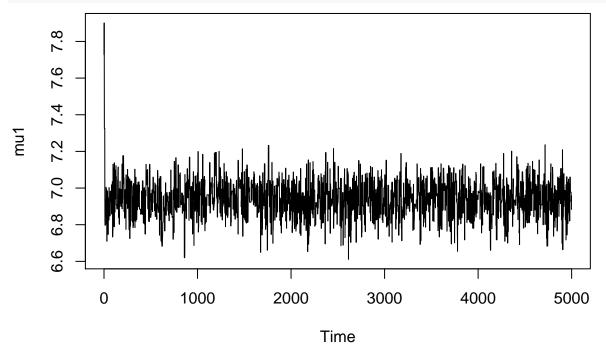
Take log on both side to get the loglikelihood function.

```
delta <- 0.7
n <- 100
set.seed(123)
u <- rbinom(n, prob = delta, size = 1)
x <- rnorm(n, ifelse(u == 1, 7, 10), 0.5)</pre>
```

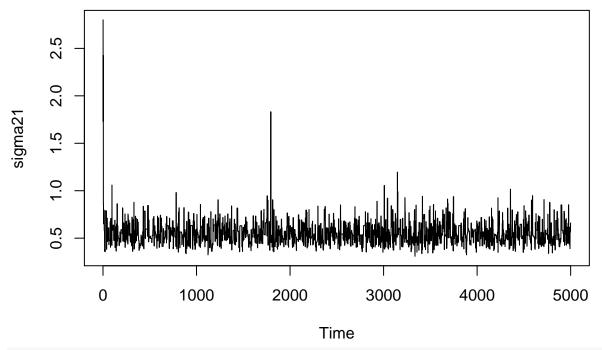
```
mylike <- function(theta, x){</pre>
  delta <- theta[1]</pre>
  mu1 <- theta[2]; sigma21 <- theta[3]</pre>
  mu2 <- theta[4]; sigma22 <- theta[5]</pre>
  return(sum(log(delta*dnorm(x, mu1, sqrt(sigma21)) + (1-delta)*dnorm(x, mu2, sqrt(sigma22)))) +
    log(dnorm(mu1, 0, 10)) + log(dnorm(mu2, 0, 10)) + log(dinvgamma(sigma21, shape = 0.5, scale = 10))
    log(dinvgamma(sigma22, shape = 0.5, scale = 10)))
mymcmc <- function(niter, thetaInit, x) {</pre>
  p <- length(thetaInit)</pre>
  thetaCurrent <- thetaInit</pre>
  out <- matrix(NA, nrow = niter, ncol = p)</pre>
  for (i in 1:niter) {
    for (j in 1:p) {
      logFC <- function(thj) {</pre>
        theta <- thetaCurrent
        theta[j] <- thj
        mylike(theta, x)
      if(j == 1){
        out[i, j] <- thetaCurrent[j] <-</pre>
        HI::arms(thetaCurrent[j], logFC,
                  function(x) ((x > 0) * (x < 1)), 1)
      } else if(j %in% c(2, 4)){
        out[i, j] <- thetaCurrent[j] <-</pre>
        HI::arms(thetaCurrent[j], logFC,
                  function(x) ((x > -10) * (x < 20)), 1)
      } else if(j %in% c(3, 5)){
        out[i, j] <- thetaCurrent[j] <-</pre>
        HI::arms(thetaCurrent[j], logFC,
                  function(x) ((x > 0) * (x < 10)), 1)
    }
  }
  out
}
niter <- 5000
thetaInit <- c(0.5, 0, 1, 0, 1)
set.seed(1026)
result <- mymcmc(niter, thetaInit, x)</pre>
plot(ts(result[,1]), ylab = "delta")
```

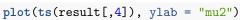


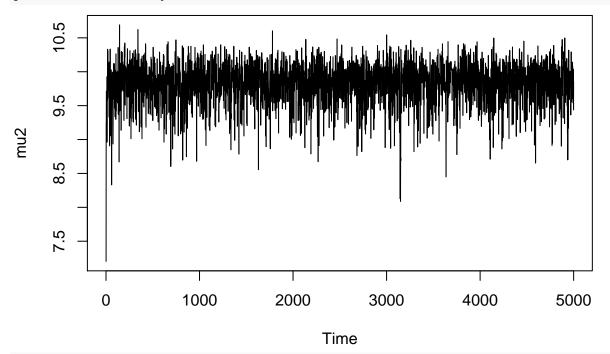




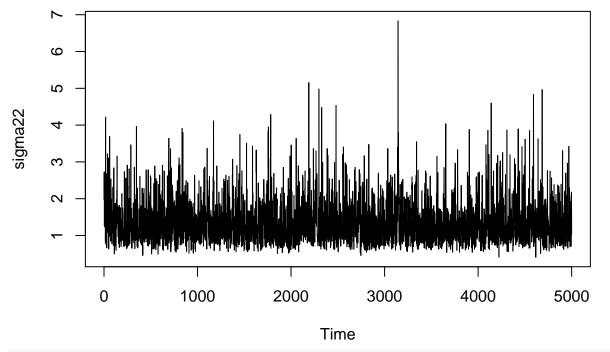
plot(ts(result[,3]), ylab = "sigma21")





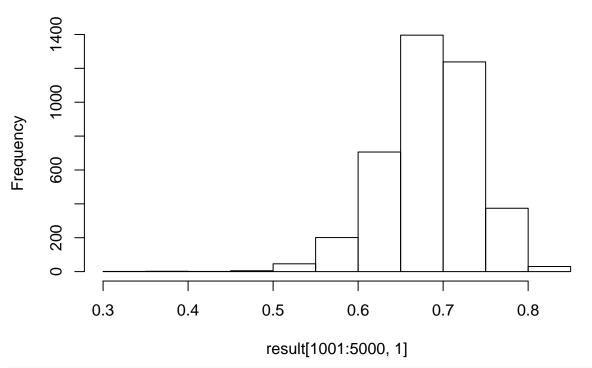


plot(ts(result[,5]), ylab = "sigma22")



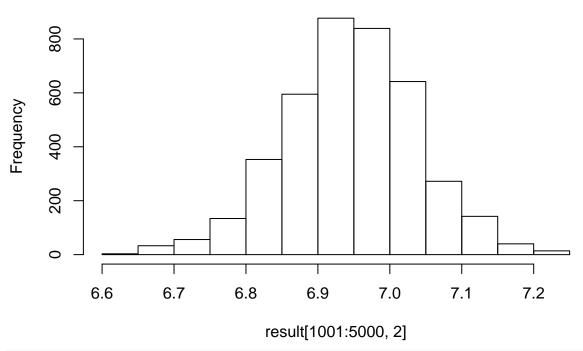
hist(result[1001:5000, 1], main = "delta")

delta



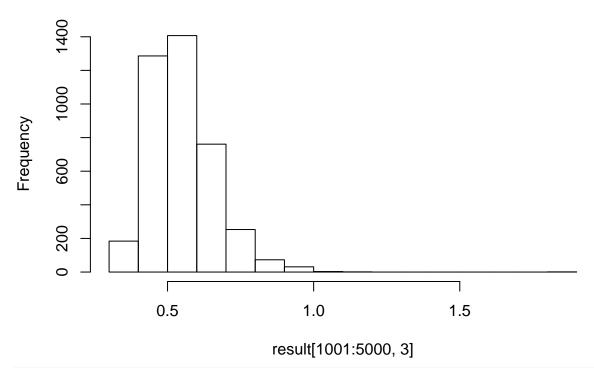
hist(result[1001:5000, 2], main = "mu1")





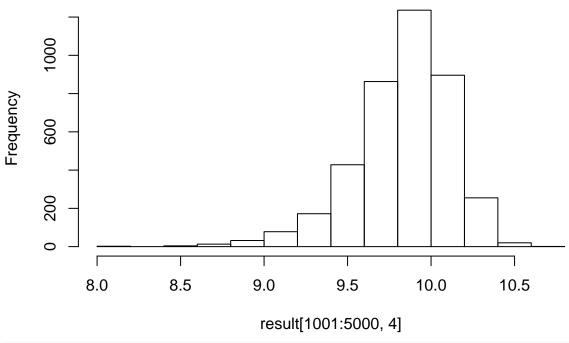
hist(result[1001:5000, 3], main = "sigma21")

sigma21



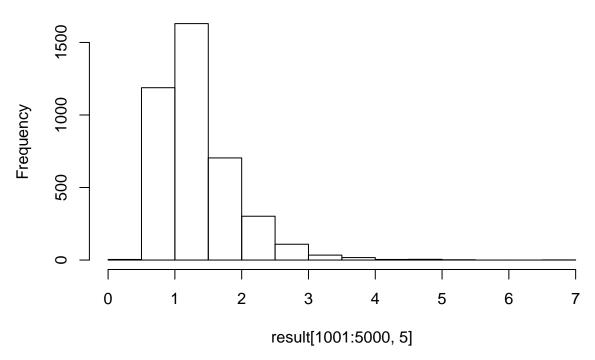
hist(result[1001:5000, 4], main = "mu2")





hist(result[1001:5000, 5], main = "sigma22")

sigma22



I only take last 4000 result in histogram and the result we gets from mcmc is similar to the true value.