STA360 Homework 6 (Ken Ye)

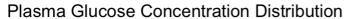
```
library(latex2exp)
library(coda)
library(ggplot2)
set.seed(0)
```

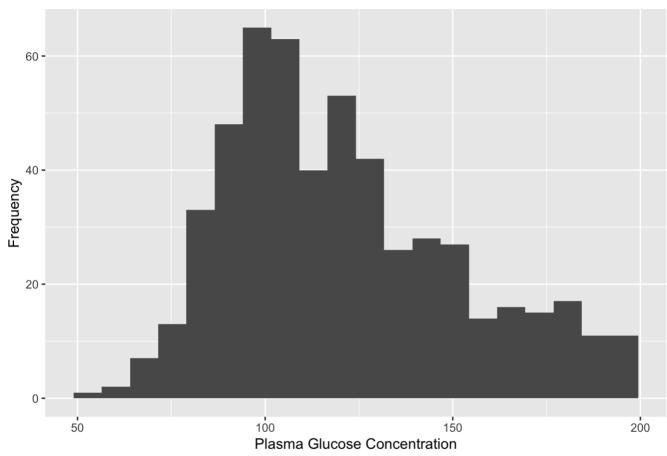
Exercise 6.2

```
# load data
glucose <- read.table("glucose.dat.txt")</pre>
```

Part a

```
# plot historgram
glucose |>
    ggplot(aes(x = V1)) +
    geom_histogram(bins = 20) +
    labs(title = "Plasma Glucose Concentration Distribution") +
    xlab("Plasma Glucose Concentration") +
    ylab("Frequency")
```





This empirical plasma glucose concentration distribution is not normal, as it"s bimodal and significantly right-skewed.

Part b

(See hand-written page)

Part c

```
Y <- glucose$V1
n <- length(Y)</pre>
nsim <- 5e4
burnin <- 1e4
# priors
a <- 1
b <- 1
mu0 <- 120
t20 <- 200
s20 <- 1000
nu0 <- 10
# storage vectors
THETA1 <- numeric(nsim - burnin)</pre>
THETA2 <- numeric(nsim - burnin)</pre>
Y.gibb <- numeric(nsim - burnin)
# starting values
p < -0.5
theta1 <- mean(Y)</pre>
theta2 <- mean(Y)</pre>
s21 < - var(Y)
s22 <- var(Y)
# Gibbs sampling
for (s in 1:nsim) {
  # simulate X
  p.x1 <- p * dnorm(Y, theta1, sqrt(s21))</pre>
  p.x2 \leftarrow (1 - p) * dnorm(Y, theta2, sqrt(s22))
  p.xi <- p.x1 / (p.x1 + p.x2)
  X \leftarrow rbinom(n, 1, p.xi)
  # calcuate group-specific summary statistics
  n1 <- sum(X)
  n2 <- n - n1
```

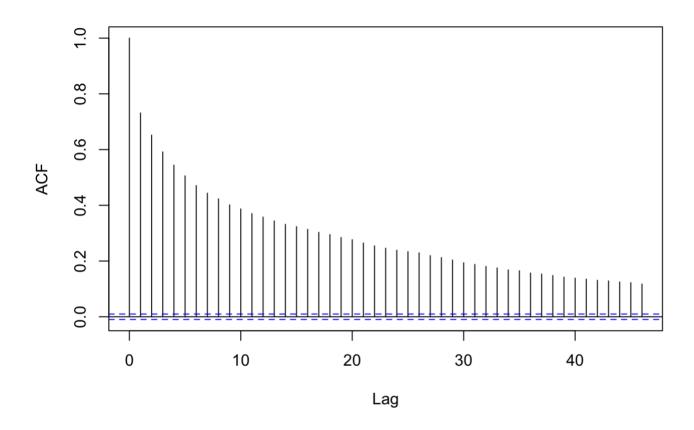
```
y1 < - Y[X == 1]
y2 < - Y[X == 0]
ybar1 <- mean(y1)</pre>
ybar2 <- mean(y2)</pre>
yvar1 <- var(y1)</pre>
yvar2 <- var(y2)</pre>
# simulate p
p < - rbeta(1, a + n1, b + n2)
# simulate thetal
t2n1 < -1 / (1 / t20 + n1 / s21)
mun1 < (mu0 / t20 + n1 * ybar1 / s21) / (1 / t20 + n1 / s21)
theta1 <- rnorm(1, mun1, sqrt(t2n1))</pre>
# simulate theta2
t2n2 < -1 / (1 / t20 + n2 / s22)
mun2 < - (mu0 / t20 + n2 * ybar2 / s22) / (1 / t20 + n2 / s22)
theta2 <- rnorm(1, mun2, sqrt(t2n2))</pre>
# simulate sigma^2 1
nun1 <- nu0 + n1
s2n1 \leftarrow (nu0 * s20 + (n1 - 1) * yvar1 + n1 * (ybar1 - theta1)^2) / nun1
s21 <- 1 / rgamma(1, nun1 / 2, s2n1 * nun1 / 2)
# simulate sigma^2 2
nun2 <- nu0 + n2
s2n2 \leftarrow (nu0 * s20 + (n2 - 1) * yvar2 + n2 * (ybar2 - theta2)^2) / nun2
s22 <- 1 / rgamma(1, nun2 / 2, s2n2 * nun2 / 2)
# simulate posterior
x.gibb <- runif(1) < p
y.gibb <- ifelse(x.gibb,</pre>
                 rnorm(1, theta1, sqrt(s21)),
                 rnorm(1, theta2, sqrt(s22)))
# store values
if (s > burnin){
```

```
THETA1[s - burnin] <- theta1
THETA2[s - burnin] <- theta2
Y.gibb[s - burnin] <- y.gibb
}
</pre>
```

```
THETA1S <- pmin(THETA1, THETA2)
THETA2S <- pmax(THETA1, THETA2)

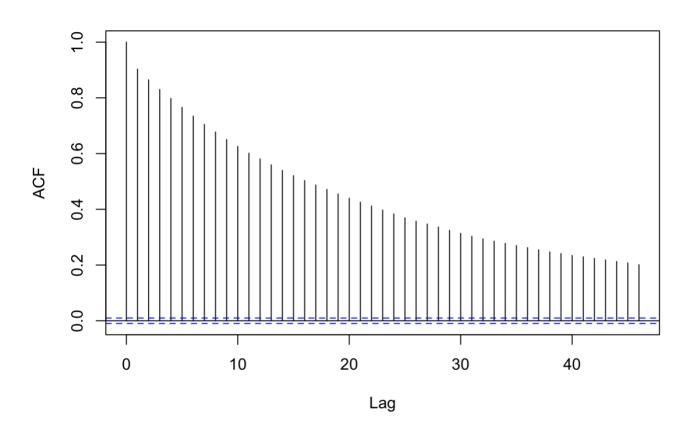
# plot autocorrelation
acf(THETA1S)
```

Series THETA1S



acf(THETA2S)

Series THETA2S



```
# print effective size
print("The effective size of theta (1) (s) is: ")

## [1] "The effective size of theta (1) (s) is: "

print(effectiveSize(THETA1S))

## var1
## 1280.576
```

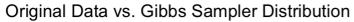
```
print("The effective size of theta (2) (s) is: ")

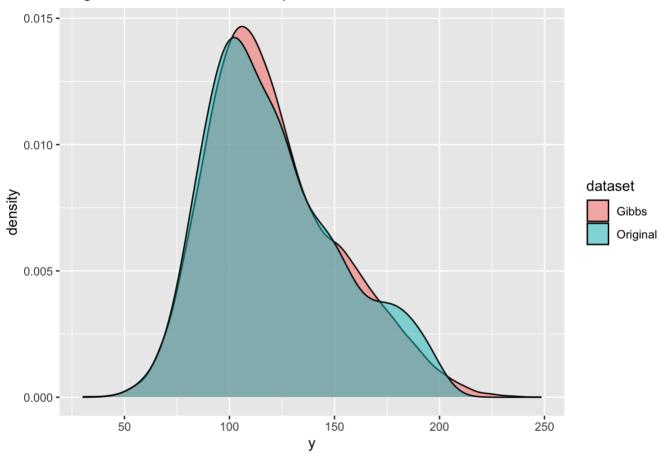
## [1] "The effective size of theta (2) (s) is: "

print(effectiveSize(THETA2S))

## var1
## 875.6929
```

Part d





Based on the graph, the densities of the original and the Gibbs sampler are very close, indicating that this two-component mixture model is a good fit for the glucose data.

Exercise 6.3

```
# load data
divorce <- read.table("divorce.dat.txt")</pre>
```

```
# function for simulating from a constrained normal distribution with mean mean and standard deviation sd, constr
ained to lie in the interval (a,b)
rcnorm <- function(n, mean = 0, sd = 1, a = -Inf, b = Inf){
  u <- runif(n, pnorm((a - mean) / sd), pnorm((b - mean) / sd) )
  mean + sd * qnorm(u)
}
```

Part a

(See hand-written page)

Part b

(See hand-written page)

Part c

```
Y <- divorce$V1
X <- divorce$V2</pre>
n <- length(Y)</pre>
nsim < -5e4
burnin <- 1e4
# priors
t2b <- 16
t2c <- 16
# storage vectors
Z <- rep(list(0 * length(X)), times = nsim - burnin)</pre>
BETA <- numeric(nsim - burnin)</pre>
C <- numeric(nsim - burnin)</pre>
# starting values
z \leftarrow rep(0, n)
beta <- 0
c <- 0
# Gibbs sampling
for (s in 1:nsim) {
  # simulate beta
  mubn <- sum(z*X) / (sum(X^2) + 1/ (t2b))
  t2bn <- 1 / (sum(X^2) - 1 / t2b)
  beta <- rnorm(1, mubn, sqrt(t2bn))</pre>
  # simulate c
  z0 <- subset(z, Y == 0)
  z1 <- subset(z, Y == 1)
  a <- max(z0)
  b \le min(z1)
  c <- rcnorm(1, 0, sqrt(t2c), a, b)</pre>
  # simulate z
  for (i in 1:n){
```

```
if(Y[i] == 1){
    z[i] <- rcnorm(1, beta * X[i], 1, c, Inf)
}
else{
    z[i] <- rcnorm(1, beta * X[i], 1, -Inf, c)
}

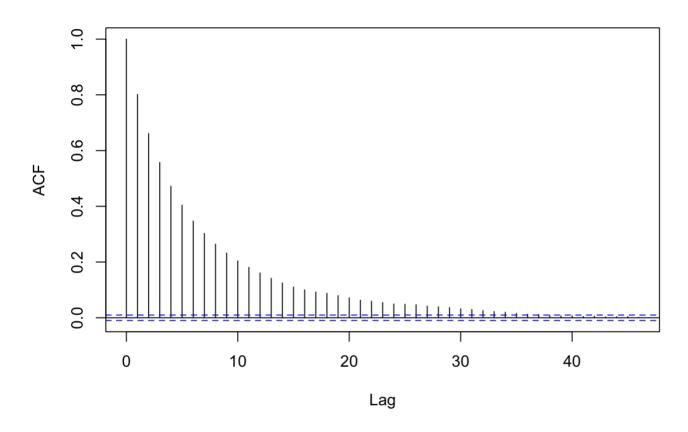
# store values
if (s > burnin) {
    Z[[s - burnin]] <- z
    BETA[s - burnin] <- beta
    C[s - burnin] <- c
}</pre>
```

beta effective size and autocorrelation
effectiveSize(BETA)

```
## var1
## 3338.234
```

acf(BETA)

Series BETA

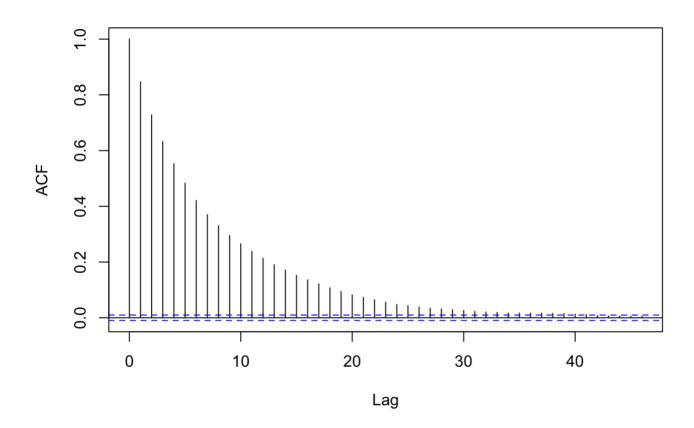


c effective size and autocorrelation effectiveSize(C)

var1 ## 2611.212

acf(C)

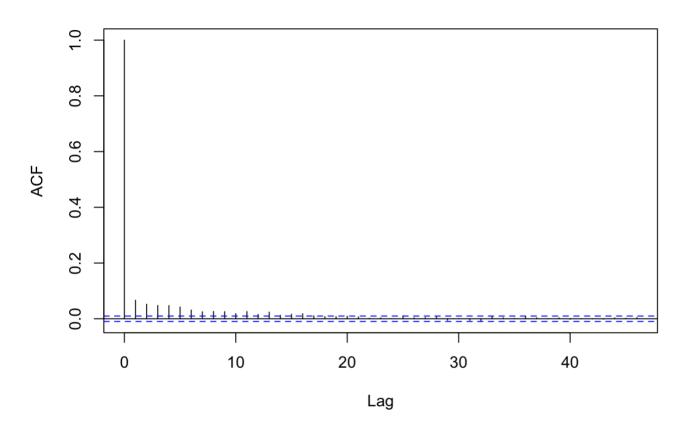
Series C

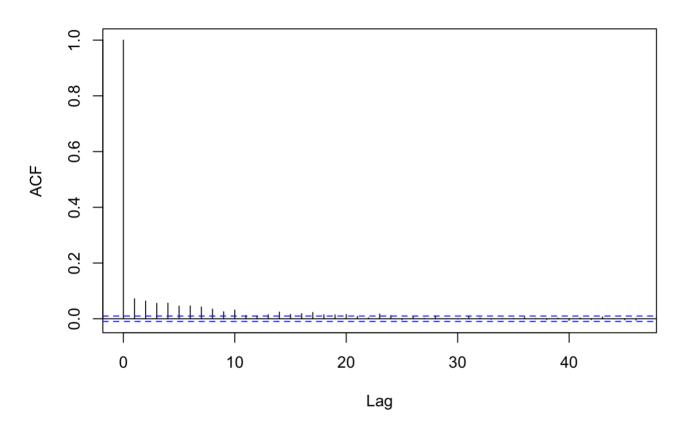


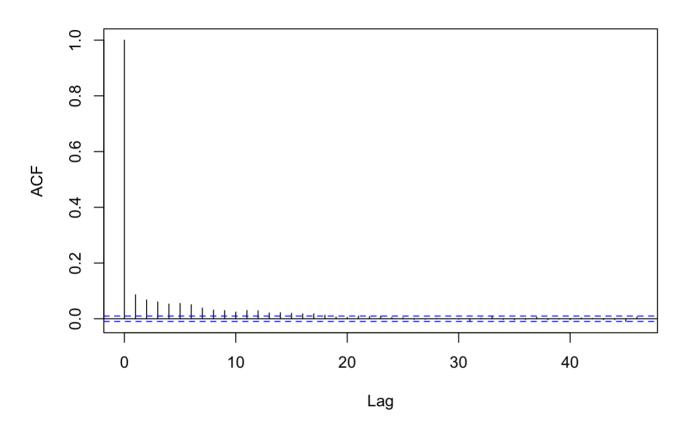
```
# zi's effective size
Zis <- rep(list(0 * length(Z)), times = n)
for (i in 1:length(Z)){
    for (j in 1:n){
        Zis[[j]][i] <- Z[[i]][j]
    }
}
Zis.eff <- rep(0, times = length(Zis))
for (i in 1:length(Zis)){
    Zis.eff[i] <- effectiveSize(unlist(Zis[[i]]))
}
Zis.eff</pre>
```

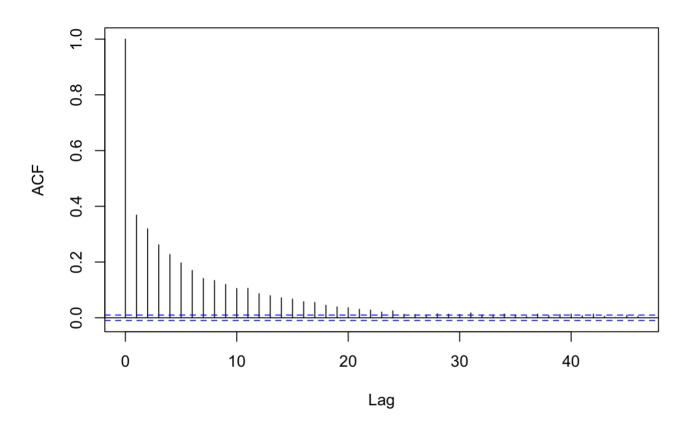
```
## [1] 19027.034 17233.683 16214.720 5989.295 17960.810 19064.730 18205.878
## [8] 6995.131 18500.697 6292.201 19794.650 3498.864 6996.895 7192.662
## [15] 6925.164 19898.214 6836.595 6516.958 17812.840 18809.866 6719.969
## [22] 19622.109 19797.833 6620.454 19957.361
```

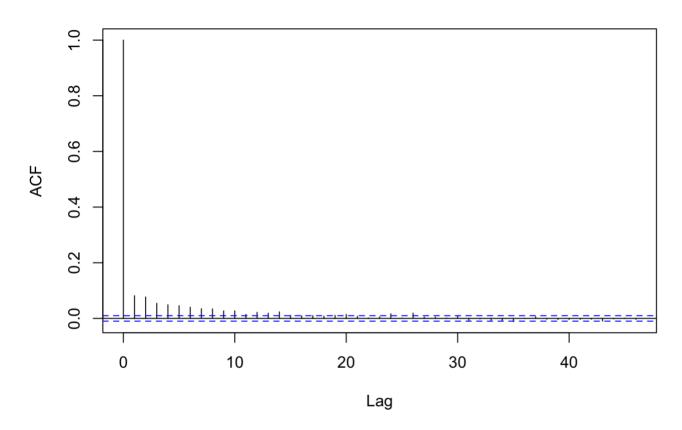
```
# zi's autocorrelation
for (i in 1:length(Zis)){
  acf(as.mcmc(Zis[[i]]))
}
```

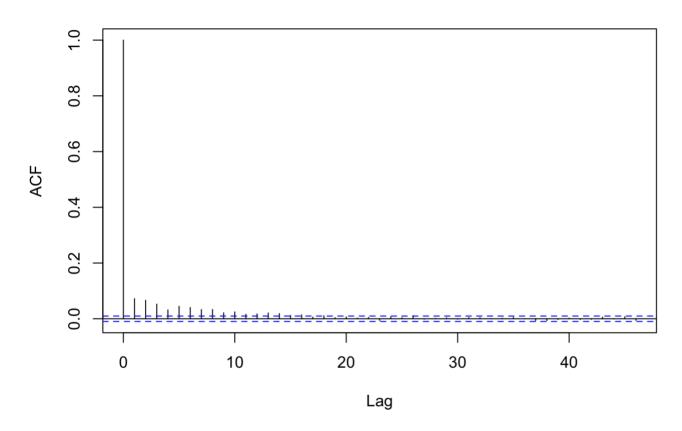


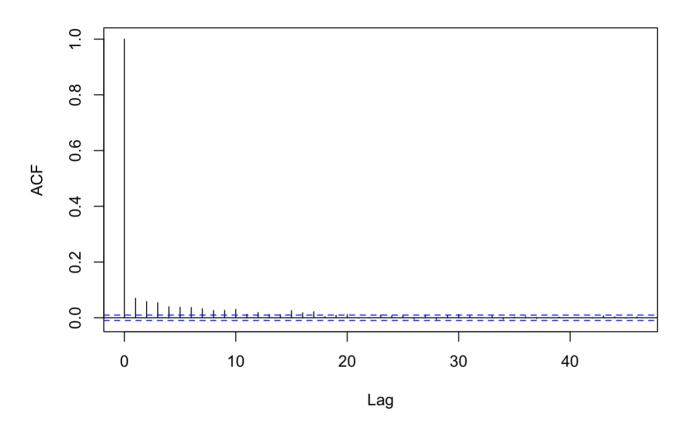


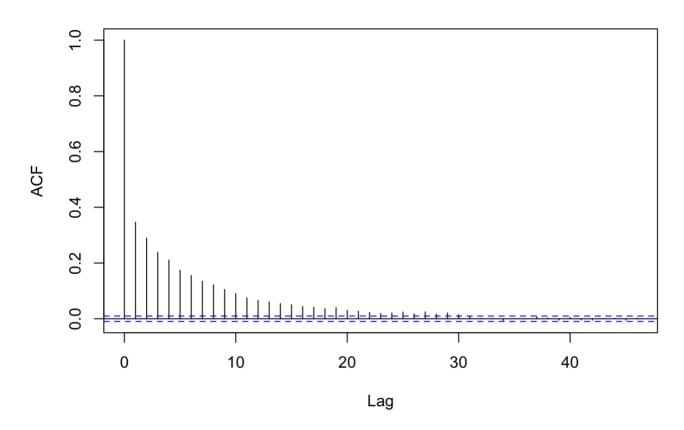


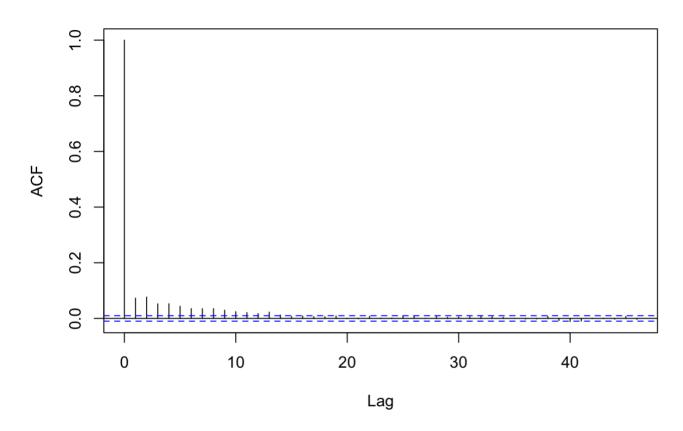


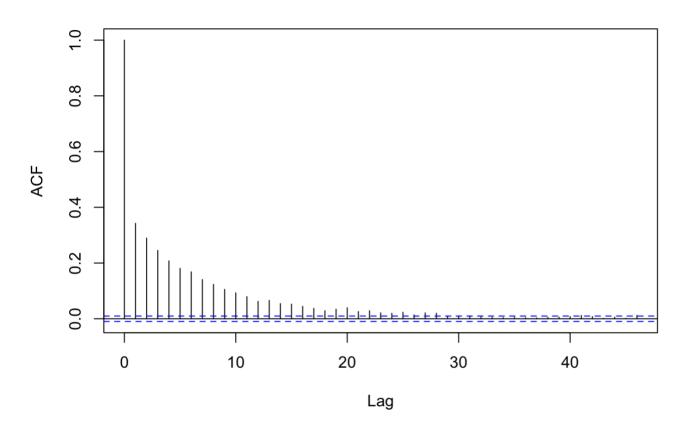


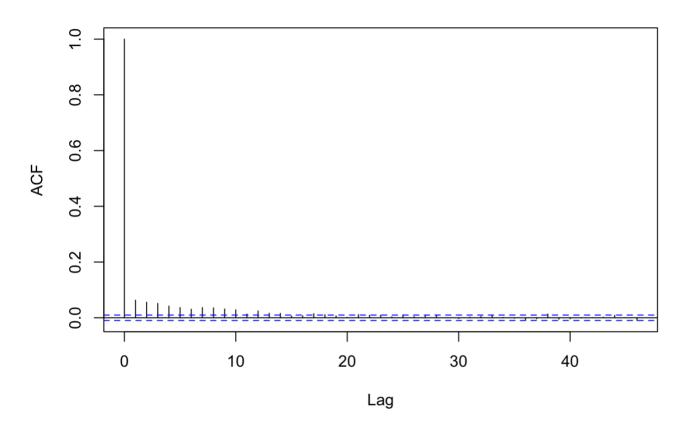


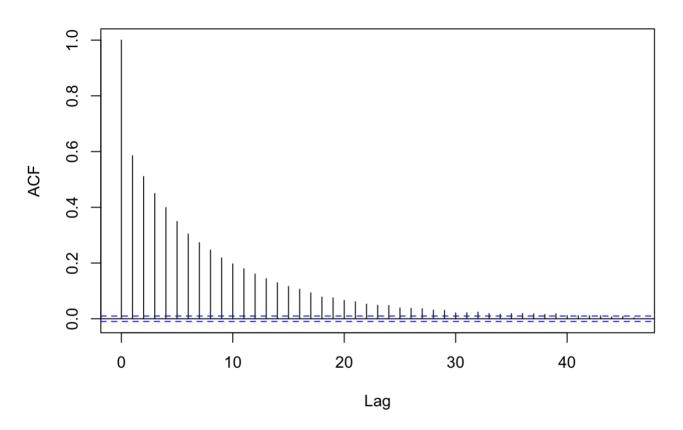


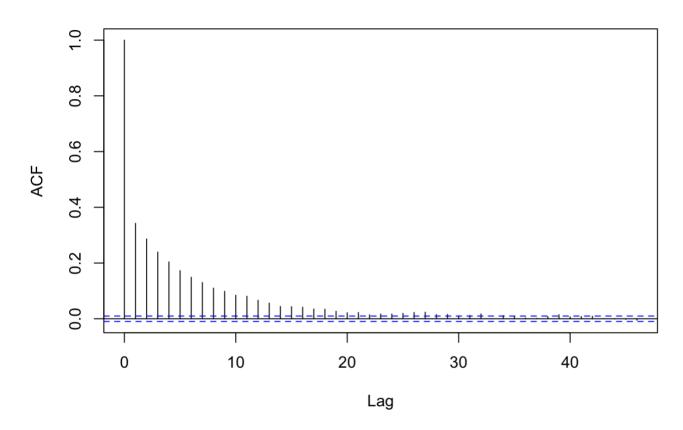


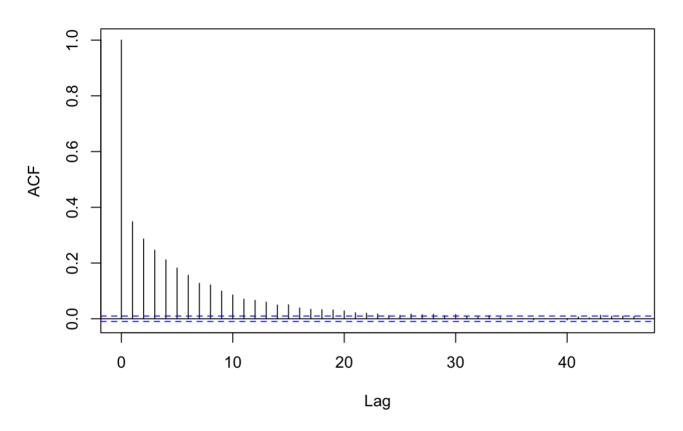


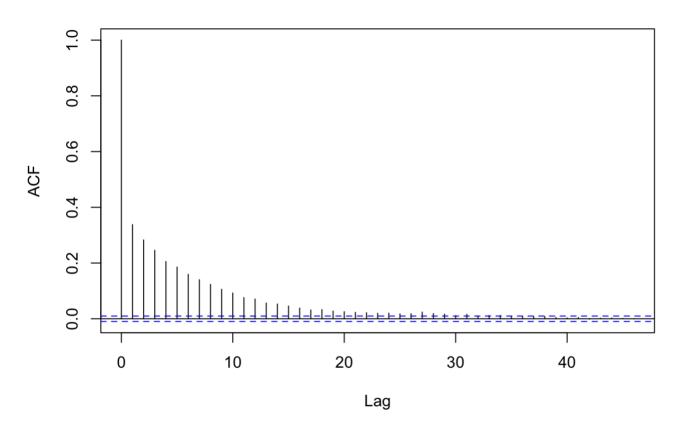


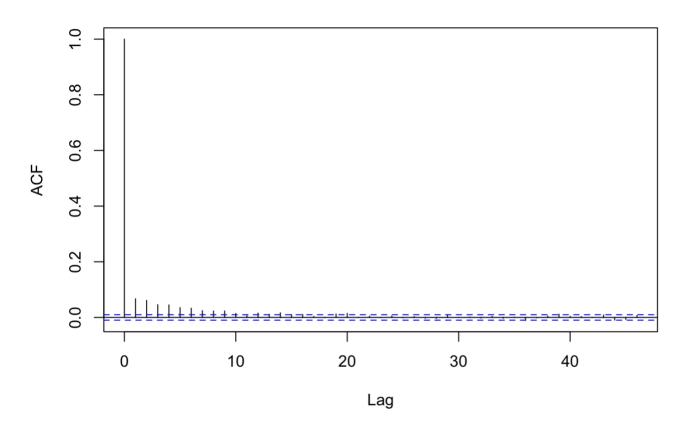


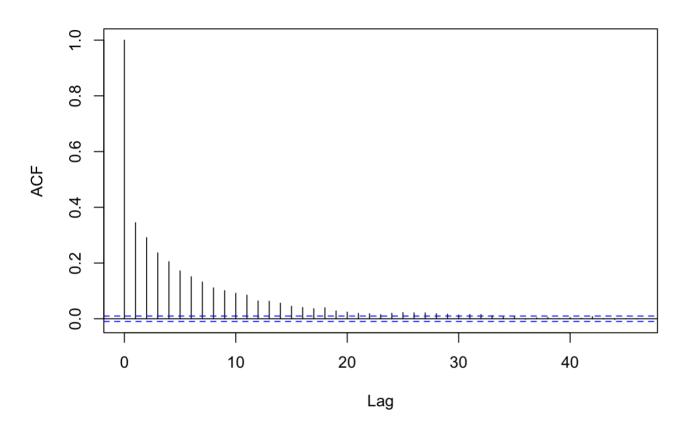


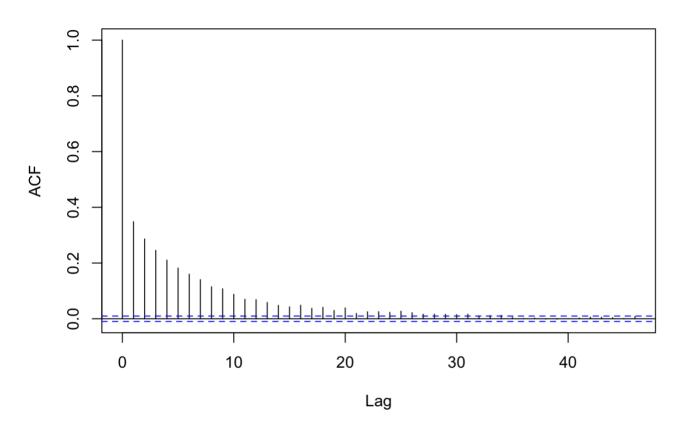


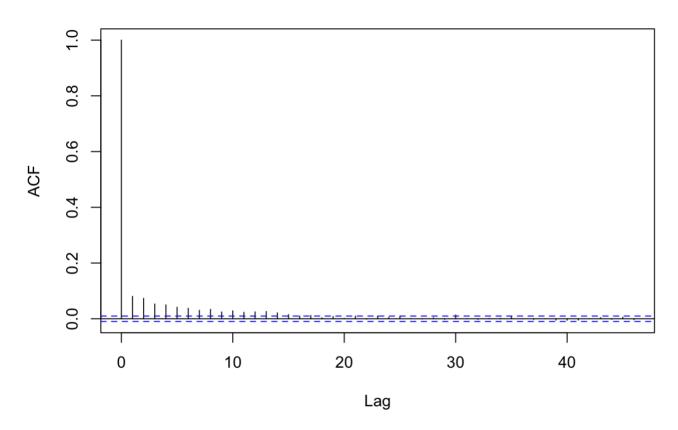


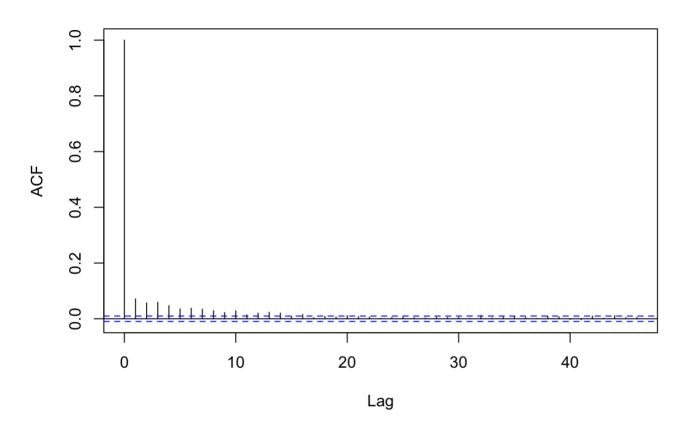


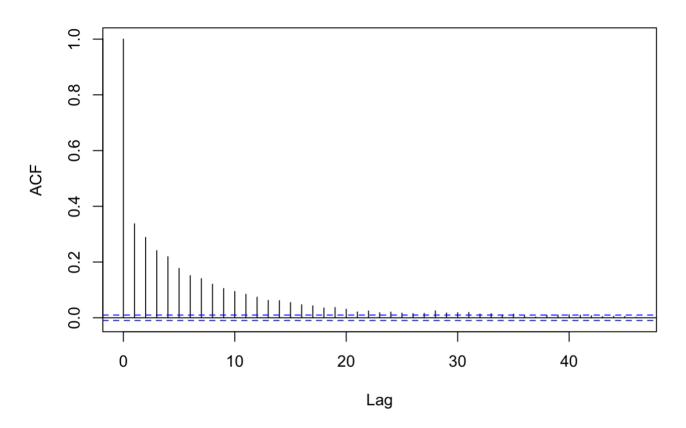


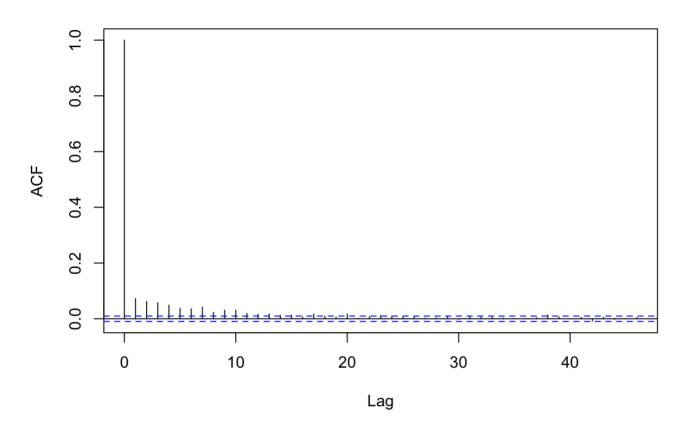


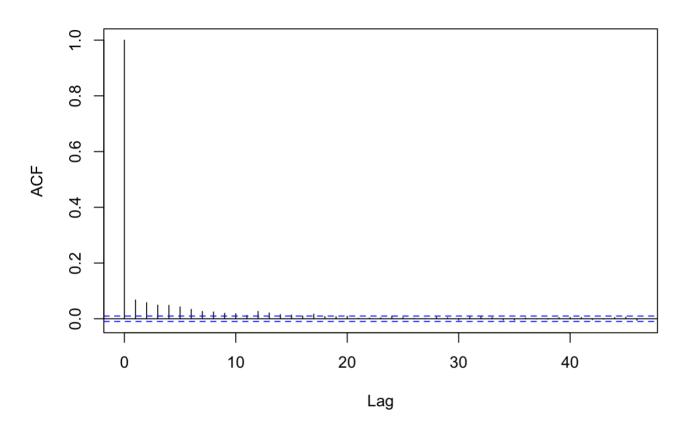


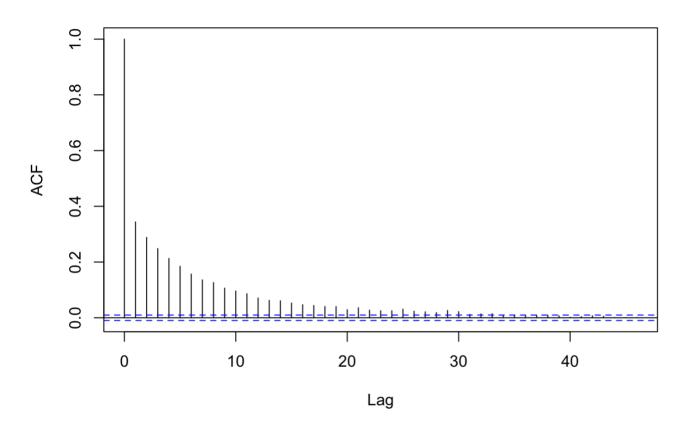




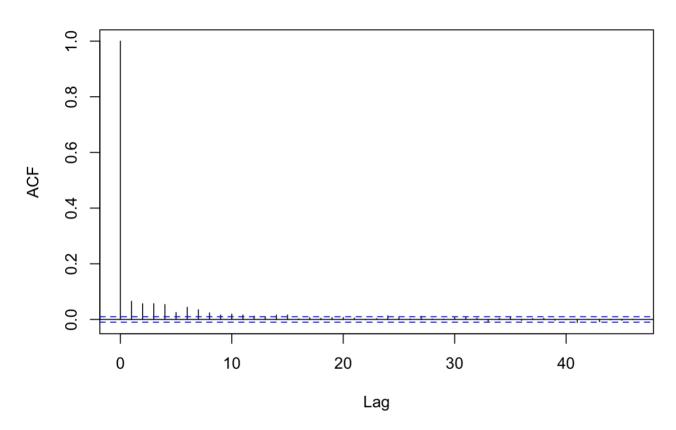








Series as.mcmc(Zis[[i]])



According to the autocorrelation plots, all values decreases and converges to 0 as we perform sufficient number of simulations, indicating that the mixing of the Markov Chain has converged to a steady distribution.

Part d

```
# 95% CI for beta
CI <- quantile(BETA, c(0.025, 0.975))
print(CI)
```

```
## 2.5% 97.5%
## -2.1479751 0.7194671
```

```
# Pr(beta > 0 | y, x)
prob <- mean(BETA > 0)
print(prob)
```

[1] 0.1723

page 2. Let N = \(\Sigma\), \(\Gamma\), \(\Gamma\), \(\Gamma\) Fc of pl PLPIX, X, DA, DB, JA', JB') ON PLX, Y, DA, DB, JA', JB', P) x p(p) p(\(\frac{1}{2}|p)\) by D d dieta(p,a,b) dbiron (n.,n,p) of Pa- (1-4) ... b, (1-6) = path -1 (1-p) b+ 2-1 = dbeta (patr, btr) FC of DA : PLANX, Y, P, BB, JA', JB') X PLX, Y, BA, BB, JA', JB', P) a(Ti p(yild, on')) p(da) p(on') by () aftidnorm (Yi, A, JA) drorm (A, M., T.) d drorm (Pa, Mr., Tr.) by tout book 1989 where $M_{n_i} = \frac{\overline{\tau_{o}} M_{o} + \frac{\overline{\gamma_{i}}}{\overline{\sigma_{A}}^{2}}}{\tau_{o} M_{o} + \frac{\overline{\gamma_{i}}}{\overline{\sigma_{A}}^{2}}}$ Fr of Do BR ~ drorm (BB, Mrs, ~~~), similar to BA, where $M_{r_2} = \frac{\overline{\tau_0} H_0 + \overline{\sigma_0} V}{\overline{\tau_0} + \overline{\sigma_0} V}$ and $v_{r_2} = \left(\frac{1}{\tau_0} + \frac{r_2}{\overline{\sigma_0}}\right)^{-1}$

and $\sigma'(\theta_0) = \frac{1}{v} \left[v_0 \sigma_0 + n_1 \int_{r_0}^{r_0} (\theta_B) \right]$

Where Vn = Votrz

page 4

6.3

a)
$$z_{i}|\beta \sim N(\beta X_{i}, 1)$$
, $\varepsilon_{i}, ..., \varepsilon_{n} \sim i i i d} N(0, 1)$, $\beta \sim N(0, \tau_{p}^{*})$

$$P(\beta | y, X, z, c) \propto P(\beta, y, X, z, c)$$

$$\propto P(z | \beta, y, X, c) P(\beta)$$

$$\propto \left\{ \frac{\pi}{i z_{i}} d_{N} m(z_{i}, \beta X_{i}, 1) \right\} d_{N} m(\beta, 0, \tau_{p}^{*})$$

$$\propto \left\{ \frac{\pi}{i z_{i}} e^{-\frac{1}{2}(z_{i} - \beta X_{i})^{2}} \right\} e^{-\frac{1}{2}(\frac{\beta}{\tau_{p}})^{2}}$$

$$\propto \exp \left\{ -\frac{1}{2} \left[\sum_{x_{i}} \frac{1}{\tau_{p}} + \sum_{x_{i}} \frac{1}{\tau_{p}} - \sum_{x_{i}} \frac{1}{\tau_{p}} + \sum_{x_{i}} \frac{1}{\tau_{p}} \right] \right\}$$

$$\alpha \exp \left\{ -\frac{1}{2} \left[\sum_{x_{i}} \frac{1}{\tau_{p}} - \frac{1}{\tau_{p}} \right] \left(P - \sum_{x_{i}} \frac{1}{\tau_{p}} + \frac{1}{\tau_{p}} \right) \right\}$$

$$\sim N(\mu_{\beta_{n}}, \tau_{\beta_{n}})$$

$$where \qquad \mu_{\beta_{n}} = \frac{\sum_{x_{i}} \frac{1}{\tau_{p}} - \frac{1}{\tau_{p}}}{\sum_{x_{i}} \frac{1}{\tau_{p}} - \frac{1}{\tau_{p}}}$$

$$\alpha \sim M \quad \tau_{\beta_{n}}^{2} = \frac{1}{\sum_{x_{i}} \frac{1}{\tau_{p}} - \frac{1}{\tau_{p}}}$$

page 5 #6.3 C~ N (0, ~() b) Giver everything else, the distribution of a min only depend or y and z. IC Given Y=y and Z=z, c must > all zi's for which y; =0 and < all zi's for which yi=1 Let a = max (x; y; = 0) and b = min{z; : y; = 1}. The Fr of c is the oxp(c) but constrained to (a,b): P(C|y,x,z,p) = P(C|y,z) dp(c)p(y|z,c)= olnorm((,0, ~,) &(a,b)(() therefore, p(c/y, x, z, p) is a constrained normal density with a support on (a,b). $|Z_i|$ $z_i|\beta \sim N(\beta x_i, 1)$ Given c, observing Yi=y; will tell us that Zi must lie in the interval y; belongs to => (-00, c) it y; =0 and (c, 00) it y=1 thus, the FC of Zi is P(Z; |y, x,z.;β, c) α P(Z; |β, x;) P(y; |Z;, c) Therefore, P(Zi/Y, b, Z-i, B, c) is proportioned to a horman devisty but constrained to pither above c or below c, depending on bi.