575_HW3

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```
library(coda)
library(ggplot2)
```

a.

```
library(coda)
library(ggplot2)
set.seed(69)
nSchools <- 8
school_data <- vector("list", nSchools)</pre>
for (i in 1:nSchools) {
  fileName <- paste0("school", i, ".dat")</pre>
  school_data[[i]] <- scan(fileName, quiet = TRUE)</pre>
}
n_i <- sapply(school_data, length)</pre>
ybar <- sapply(school_data, mean)</pre>
all_data <- unlist(school_data)</pre>
N <- length(all_data)</pre>
       <- 7
mu0
sigma2_mu0 <- 5
nu0
     <- 2
sigma20 <- 15
eta0 <- 2
tau20 <- 10
n_iter <- 10000
burn_in <- 1000
theta_samples <- matrix(NA, nrow = n_iter, ncol = nSchools)</pre>
mu_samples
            <- numeric(n_iter)</pre>
sigma2_samples <- numeric(n_iter)</pre>
tau2_samples <- numeric(n_iter)</pre>
R_samples
            <- numeric(n_iter)</pre>
```

```
mu_current <- mu0</pre>
sigma2_current <- sigma20</pre>
tau2 current <- tau20
theta_current <- ybar
for (iter in 1:n_iter) {
  for (i in 1:nSchools) {
    vi <- 1 / (n_i[i] / sigma2_current + 1 / tau2_current)</pre>
    mi <- vi * (n_i[i] * ybar[i] / sigma2_current + mu_current / tau2_current)
    theta_current[i] <- rnorm(1, mean = mi, sd = sqrt(vi))</pre>
  }
  v_mu <- 1 / (nSchools / tau2_current + 1 / sigma2_mu0)</pre>
  m_mu <- v_mu * (sum(theta_current) / tau2_current + mu0 / sigma2_mu0)</pre>
  mu_current <- rnorm(1, mean = m_mu, sd = sqrt(v_mu))</pre>
  nu_new <- nu0 + N</pre>
  sum_sq <- 0
  for (i in 1:nSchools) {
    sum_sq <- sum_sq + sum((school_data[[i]] - theta_current[i])^2)</pre>
  scale_sigma2 <- (nu0 * sigma20 + sum_sq) / nu_new</pre>
  sigma2_current <- nu_new * scale_sigma2 / rchisq(1, df = nu_new)</pre>
  nu tau <- eta0 + nSchools
  sum_theta_sq <- sum((theta_current - mu_current)^2)</pre>
  scale_tau2 <- (eta0 * tau20 + sum_theta_sq) / nu_tau</pre>
  tau2_current <- nu_tau * scale_tau2 / rchisq(1, df = nu_tau)</pre>
  R_samples[iter] <- 1 + sigma2_current / tau2_current</pre>
  theta_samples[iter, ] <- theta_current</pre>
  mu_samples[iter]
                       <- mu_current
  sigma2_samples[iter] <- sigma2_current</pre>
  tau2_samples[iter] <- tau2_current</pre>
theta_samples <- theta_samples[-(1:burn_in), ]</pre>
mu_samples <- mu_samples[-(1:burn_in)]</pre>
sigma2_samples <- sigma2_samples[-(1:burn_in)]</pre>
tau2_samples <- tau2_samples[-(1:burn_in)]</pre>
            <- R_samples[-(1:burn_in)]</pre>
R_samples
mu_mcmc
           <- mcmc(mu_samples)</pre>
sigma2_mcmc <- mcmc(sigma2_samples)</pre>
tau2_mcmc <- mcmc(tau2_samples)</pre>
ess_mu <- effectiveSize(mu_mcmc)</pre>
ess_sigma2 <- effectiveSize(sigma2_mcmc)</pre>
ess_tau2 <- effectiveSize(tau2_mcmc)</pre>
```

```
cat("Effective Sample Size for mu:", round(ess_mu, 1), "\n")

## Effective Sample Size for mu: 7789

cat("Effective Sample Size for sigma2:", round(ess_sigma2, 1), "\n")

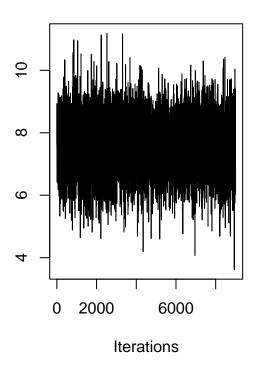
## Effective Sample Size for sigma2: 8484.7

cat("Effective Sample Size for tau2:", round(ess_tau2, 1), "\n")

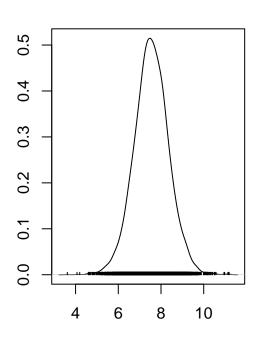
## Effective Sample Size for tau2: 6837

plot(mu_mcmc)
```

Trace of var1



Density of var1



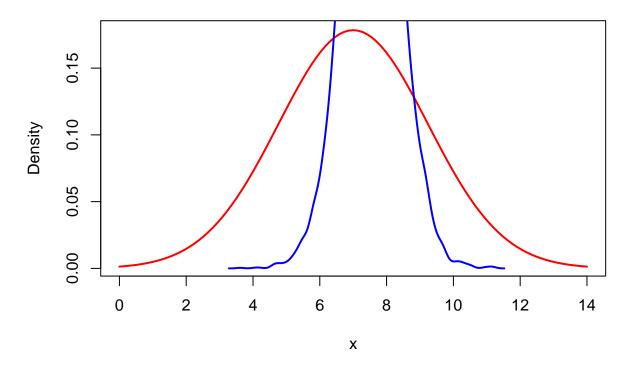
N = 9000 Bandwidth = 0.132

```
posterior_summary <- function(samples) {
  mean_val <- mean(samples)
  ci <- quantile(samples, c(0.025, 0.975))
  list(mean = mean_val, CI = ci)
}

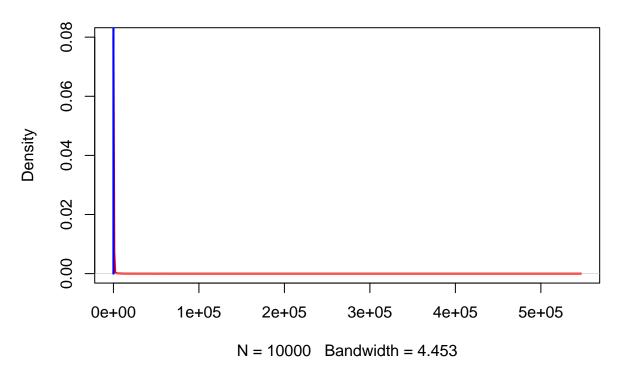
mu_summary <- posterior_summary(mu_samples)
sigma2_summary <- posterior_summary(sigma2_samples)</pre>
```

```
tau2_summary<- posterior_summary(tau2_samples)</pre>
cat("Posterior summary for mu:\n")
## Posterior summary for mu:
print(mu_summary)
## $mean
## [1] 7.550256
## $CI
##
       2.5%
            97.5%
## 5.936771 9.162568
cat("Posterior summary for sigma2:\n")
## Posterior summary for sigma2:
print(sigma2_summary)
## $mean
## [1] 14.46881
##
## $CI
##
       2.5%
               97.5%
## 11.76766 17.72840
cat("Posterior summary for tau2:\n")
## Posterior summary for tau2:
print(tau2_summary)
## $mean
## [1] 5.553585
## $CI
        2.5%
##
                 97.5%
## 1.895186 14.813303
curve(dnorm(x, mean = mu0, sd = sqrt(sigma2_mu0)),
      from = 0, to = 14, col = "red", lwd = 2, ylab = "Density",
      main = "mu: Prior (red) vs Posterior")
lines(density(mu_samples), col = "blue", lwd = 2)
```

mu: Prior (red) vs Posterior

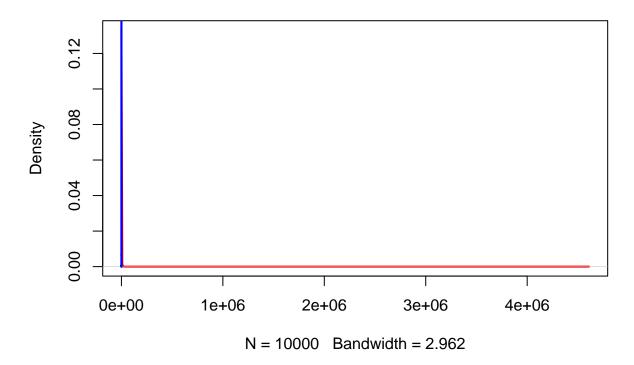


sigma2: Prior (red) vs Posterior

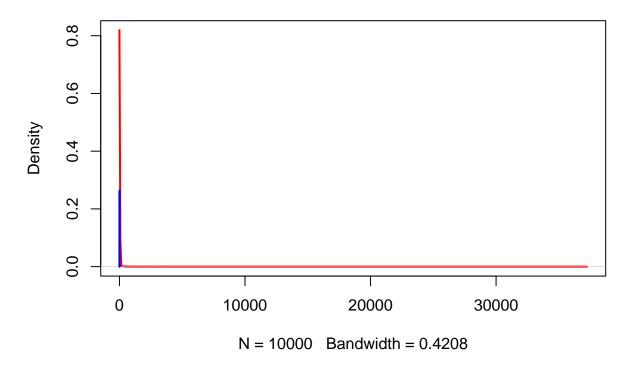


```
tau2_prior <- eta0 * tau20 / rchisq(n_prior, df = eta0)
plot(density(tau2_prior), col = "red", lwd = 2,
    main = "tau2: Prior (red) vs Posterior", ylab = "Density")
lines(density(tau2_samples), col = "blue", lwd = 2)</pre>
```

tau2: Prior (red) vs Posterior



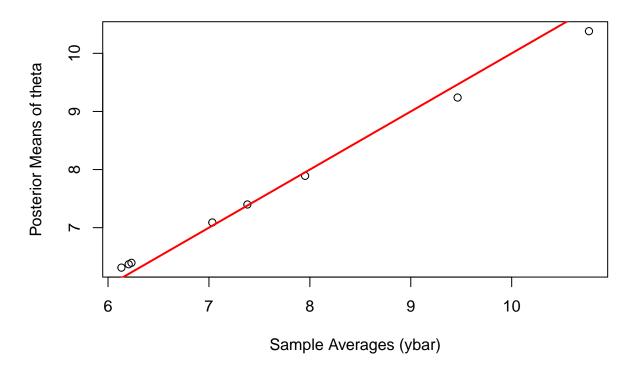
R = 1 + sigma2/tau2: Prior (red) vs Posterior



Im clearly doing something wrong here ^ but not sure what it is...

abline(0, 1, col = "red", lwd = 2)

Raw Sample Averages vs. Posterior Expectations



```
overall_ybar <- mean(all_data)
posterior_mu_mean <- mean(mu_samples)
cat("Overall sample mean:", round(overall_ybar, 3), "\n")

## Overall sample mean: 7.691

cat("Posterior mean of mu:", round(posterior_mu_mean, 3), "\n")</pre>
```

Posterior mean of mu: 7.55