Homework 6: Metropolis-Hastings Algorithm Logit

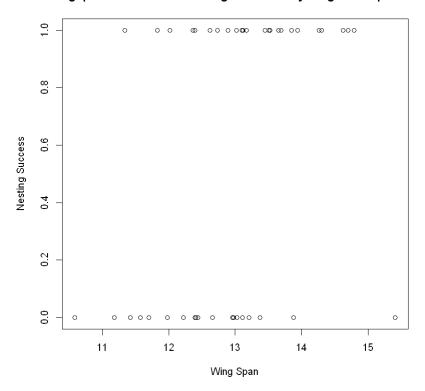
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```
sparrow.dat <- read.table("msparrownest.txt", header = F)
y <- sparrow.dat[,1]; wspan <- as.vector(sparrow.dat[,2])</pre>
```

```
plot(wspan, y,
          type="p",
          xlab = "Wing Span",
          ylab = "Nesting Success",
          main = "Wingspan in relation to Nesting Success of young male Sparrows"
          )
```

Wingspan in relation to Nesting Success of young male Sparrows



In the above relation, we observe that young male sparrows who's wingspans are small tend to have a lower sucess rate at nesting than those with larger wingspands. The above model outlined should have the capabilities at cpaturing that relationship.

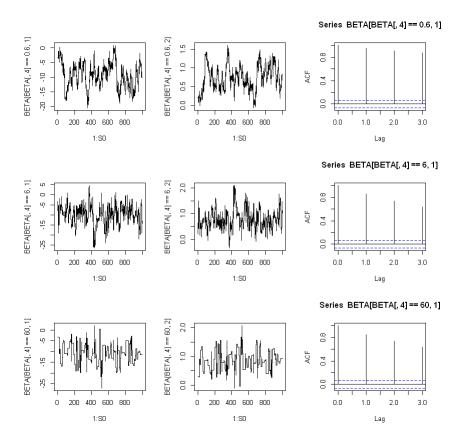
```
X <- as.matrix(cbind(rep(1, length(wspan)), wspan))</pre>
```

```
# Prior specificaations
beta \leftarrow as.matrix(c(-3.35, 0.2), nrow=3); sigma.prior \leftarrow matrix(data = c(200,0,0,100), nrow
# proposal distribution
variances <-c(0.6,6,60)
# Metropolis Set up
S0 = 1000
BETA <- matrix(rep(NaN, S0*4*3), nrow = S0*3, ncol = 4)
set.seed(42)
row.index <- 1
for (variance in variances) {
  acceptance <- 0
  sig_proposal <- variance * solve(t(X) %*% X)</pre>
  beta.current <- beta
  for (iter in 1:S0) {
    beta.proposal <- t(rmvnorm(1, mean = beta.current, sigma = sig_proposal))</pre>
    log_ratio <- log.ratio(beta.proposal = beta.proposal, beta.previous = beta.current, y = ;</pre>
    if (log(runif(1)) < log_ratio) {</pre>
      beta.current <- beta.proposal</pre>
      acceptance <- acceptance + 1
    }
    BETA[row.index, 1:2] <- beta.current</pre>
    BETA[row.index, 3] <- log_ratio</pre>
    BETA[row.index, 4] <- variance</pre>
    row.index <- row.index + 1</pre>
  }
  # Compute acceptance rate
  acceptance_rate <- acceptance / S0</pre>
  print(paste("Acceptance rate for variance", variance, ":", acceptance_rate))
par(mfrow = c(3,3))
plot(1:S0, BETA[BETA[,4] == 0.6, 1], type="l")
plot(1:S0, BETA[BETA[,4] == 0.6, 2], type = "1")
```

```
acf(BETA[BETA[,4] == 0.6,1], lag.max = 3)

plot(1:S0, BETA[BETA[,4] == 6,1], type="1")
plot(1:S0, BETA[BETA[,4] == 6,2], type ="1")
acf(BETA[BETA[,4] == 6,1], lag.max = 3)

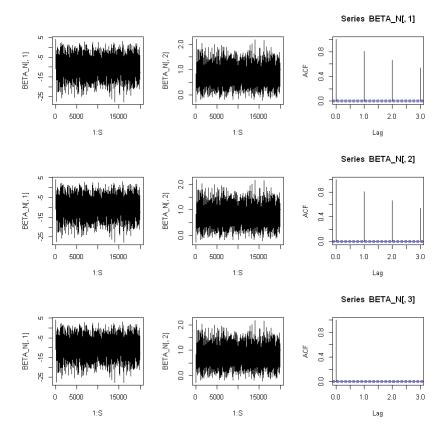
plot(1:S0, BETA[BETA[,4] == 60,1], type="1")
plot(1:S0, BETA[BETA[,4] == 60,2], type ="1")
acf(BETA[BETA[,4] == 60,1], lag.max = 3)
```



Looking at the acceptance rate ratios, I would pick the acceptance rate ratio of 50% corresponding to the proposal variance of 6.

```
# Prior specificaations
beta <- as.matrix(c(-3.35, 0.2), nrow=3); sigma.prior <- matrix(data = c(200,0,0,100), nrow
# Metropolis Set up
S = 20000</pre>
```

```
BETA_N <- matrix(rep(NaN, S*3), nrow = S, ncol = 3)</pre>
set.seed(42)
acceptance <- 0
sig_proposal <- 6 * solve(t(X) %*% X)</pre>
beta.current <- beta
for (iter in 1:S) {
    beta.proposal <- t(rmvnorm(1, mean = beta.current, sigma = sig_proposal))</pre>
    log_ratio <- log.ratio(beta.proposal = beta.proposal, beta.previous = beta.current, y = )
    if (log(runif(1)) < log_ratio) {</pre>
      beta.current <- beta.proposal</pre>
      acceptance <- acceptance + 1
    BETA_N[iter, 1:2] <- beta.current</pre>
    BETA_N[iter, 3] <- log_ratio</pre>
acceptance_rate <- acceptance / S
par(mfrow = c(3,3))
plot(1:S, BETA_N[, 1],type="l")
plot(1:S, BETA_N[, 2], type ="1")
acf(BETA_N[,1], lag.max = 3)
plot(1:S, BETA_N[, 1],type="l")
plot(1:S, BETA_N[, 2], type ="1")
acf(BETA_N[,2], lag.max = 3)
plot(1:S, BETA_N[, 1],type="l")
plot(1:S, BETA_N[, 2],type ="1")
acf(BETA_N[,3], lag.max = 3)
```



MLE Estimate
glm(y~X[,2], family = "binomial")

Call: glm(formula = y ~ X[, 2], family = "binomial")

Coefficients:

(Intercept) X[, 2] -9.7868 0.7758

Degrees of Freedom: 42 Total (i.e. Null); 41 Residual

Null Deviance: 59.03

Residual Deviance: 53.38 AIC: 57.38

Beta.df <- as.data.frame(BETA_N[,1:2])
colnames(Beta.df) <- c("Beta1", "Beta2")</pre>

```
beta.estimates <- Beta.df %>%
  summarise(across(
   everything(),
   list(
     mean = mean,
      2.5\% = ~ quantile(.x, 0.025),
      97.5\% = ~ quantile(.x, 0.975),
     se = -sd(.x)/sqrt(n())
   ),
    .names = "{.col}_{.fn}"
  )) %>%
  pivot_longer(
    cols = everything(),
   names_to = c("beta", "stat"),
   names_sep = "_",
   values_to = "value"
  ) %>%
 pivot_wider(
   names_from = stat,
   values_from = value
  ) %>%
  column_to_rownames(var = "beta")
beta.estimates
```

The estimates for beta of the MLE and Metropolis Algorithm are very similar to one another.

```
x_vals <- seq(10, 16, length.out = 100)

logistic_function <- function(beta1, beta2, x) {
    exp(beta1 + beta2 * x) / (1 + exp(beta1 + beta2 * x))
}

predictions <- sapply(x_vals, function(x) {
    logistic_function(Beta.df$Beta1, Beta.df$Beta2, x)
})

pred_summary <- apply(predictions, 2, function(pred) {
    c(mean = mean(pred), lower = quantile(pred, 0.025), upper = quantile(pred, 0.975))
})</pre>
```

```
pred_summary_df <- data.frame(
    x = x_vals,
    mean = pred_summary["mean", ],
    lower = pred_summary["lower.2.5%", ],
    upper = pred_summary["upper.97.5%", ]
)

plot(pred_summary_df$x, pred_summary_df$mean, type = "l", col = "blue", lwd = 2,
        ylab = "Nesting Success Probability", xlab = "Wingspan", ylim = range(pred_summary_df$lelines(pred_summary_df$x, pred_summary_df$lower, col = "red", lty = 2)
lines(pred_summary_df$x, pred_summary_df$upper, col = "red", lty = 2)
legend("bottomright", legend = c("Mean", "95% Credible Interval"), col = c("blue", "red"), legend</pre>
```

