hw8

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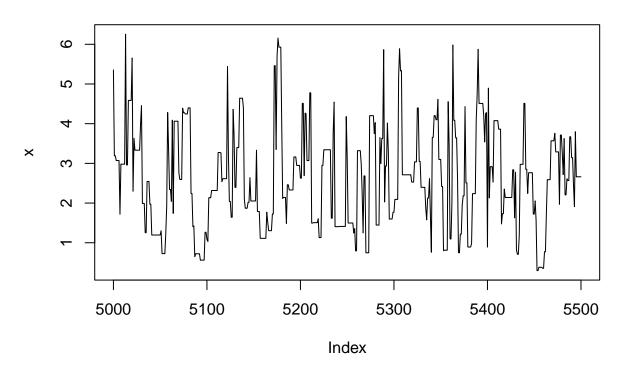
11.1

```
num_samples <- 10000</pre>
sigma <- 2
samples <- numeric(num_samples)</pre>
samples[1] \leftarrow rchisq(1, df = 1)
rejection_count <- 0</pre>
uniform_samples <- runif(num_samples)</pre>
f <- function(x, sigma) {</pre>
  if (x < 0) return(0)
  stopifnot(sigma > 0)
  return((x / sigma^2) * exp(-x^2 / (2 * sigma^2)))
for (i in 2:num_samples) {
  current_sample <- samples[i - 1]</pre>
  proposed_sample <- rchisq(1, df = current_sample)</pre>
  numerator <- f(proposed_sample, sigma) * dchisq(current_sample, df = proposed_sample)</pre>
  denominator <- f(current_sample, sigma) * dchisq(proposed_sample, df = current_sample)</pre>
  acceptance_prob <- numerator / denominator</pre>
  if (uniform_samples[i] <= acceptance_prob) {</pre>
    samples[i] <- proposed_sample</pre>
  } else {
    samples[i] <- current_sample</pre>
    rejection_count <- rejection_count + 1
  }
print(rejection_count)
```

[1] 5433

```
par(mfrow = c(1, 1))
index <- 5000:5500
subset_samples <- samples[index]
plot(index, subset_samples, type = "l", main = "Metropolis-Hastings Samples", ylab = "x", xlab = "Index")</pre>
```

Metropolis-Hastings Samples



相比于图 11.1, 更多的样本被拒绝, 有效性降低

```
11.2
```

```
num_samples <- 10000</pre>
sigma <- 4
initial_value <- 1</pre>
samples <- numeric(num_samples)</pre>
samples[1] <- initial_value</pre>
rejection_count <- 0</pre>
uniform_samples <- runif(num_samples)</pre>
target_distribution <- function(x, sigma) {</pre>
  if (x < 0) return(0)
  stopifnot(sigma > 0)
  return((x / sigma^2) * exp(-x^2 / (2 * sigma^2)))
}
for (i in 2:num_samples) {
  current_sample <- samples[i - 1]</pre>
  proposed_sample <- rgamma(1, current_sample, 1)</pre>
  numerator <- target_distribution(proposed_sample, sigma) * dgamma(current_sample, proposed_sample, 1)</pre>
  denominator <- target_distribution(current_sample, sigma) * dgamma(proposed_sample, current_sample, 1
  acceptance_prob <- numerator / denominator</pre>
```

```
samples[i] <- proposed_sample
} else {
    samples[i] <- current_sample
    rejection_count <- rejection_count + 1
}

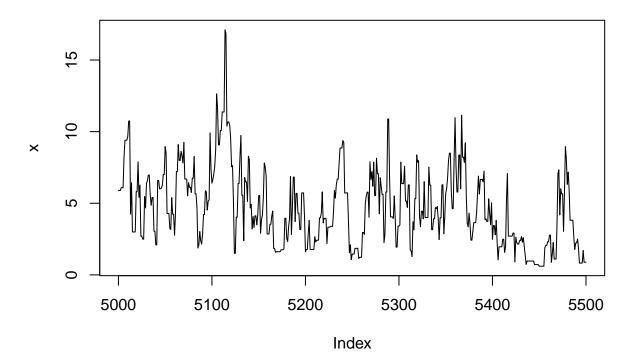
print(rejection_count)

## [1] 3083</pre>
```

if (uniform_samples[i] <= acceptance_prob) {</pre>

```
subset_index <- 5000:5500
subset_samples <- samples[subset_index]
plot(subset_index, subset_samples, type = "1", main = "Metropolis-Hastings Samples", ylab = "x", xlab =</pre>
```

Metropolis-Hastings Samples



11.3(1)

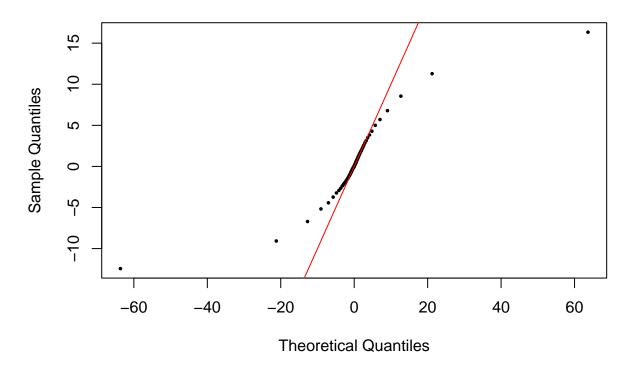
```
num_samples <- 10000
sigma <- 3

samples <- numeric(num_samples)
samples[1] <- rnorm(1, 0, sigma)
rejection_count <- 0
uniform_samples <- runif(num_samples)</pre>
```

```
# Metropolis-Hastings algorithm
for (i in 2:num_samples) {
  current_sample <- samples[i - 1]</pre>
  proposed_sample <- rnorm(1, current_sample, sigma)</pre>
  numerator <- 1 + current_sample^2</pre>
  denominator <- 1 + proposed_sample^2</pre>
  numerator <- numerator * dnorm(current sample, proposed sample, sigma)
  denominator <- denominator * dnorm(proposed_sample, current_sample, sigma)</pre>
  acceptance_prob <- numerator / denominator</pre>
  if (uniform_samples[i] <= acceptance_prob) {</pre>
    samples[i] <- proposed_sample</pre>
  } else {
    samples[i] <- current_sample</pre>
    rejection_count <- rejection_count + 1</pre>
}
# Print the number of rejections
print(rejection_count)
## [1] 4472
 (2)
num_samples <- 10000</pre>
sigma <- 3
burn_in <- 1000
samples <- numeric(num_samples)</pre>
samples[1] <- rnorm(1, 0, sigma)</pre>
rejection_count <- 0</pre>
uniform_samples <- runif(num_samples)</pre>
# Metropolis-Hastings algorithm
for (i in 2:num_samples) {
  current_sample <- samples[i - 1]</pre>
  proposed_sample <- rnorm(1, current_sample, sigma)</pre>
  numerator <- 1 + current_sample^2</pre>
  denominator <- 1 + proposed_sample^2</pre>
  numerator <- numerator * dnorm(current_sample, proposed_sample, sigma)</pre>
  denominator <- denominator * dnorm(proposed_sample, current_sample, sigma)</pre>
  acceptance_prob <- numerator / denominator</pre>
  if (uniform_samples[i] <= acceptance_prob) {</pre>
    samples[i] <- proposed_sample</pre>
  } else {
    samples[i] <- current_sample</pre>
    rejection_count <- rejection_count + 1</pre>
  }
```

```
}
print(rejection_count)
## [1] 4863
# Remove burn-in samples
post_burn_in_samples <- samples[(burn_in + 1):num_samples]</pre>
p_{values} \leftarrow seq(0.1, 0.9, 0.1)
sample_quantiles <- quantile(post_burn_in_samples, p_values)</pre>
cauchy_quantiles <- qcauchy(p_values)</pre>
quantile_comparison <- round(rbind(sample_quantiles, cauchy_quantiles), 3)</pre>
print(quantile_comparison)
                                       30%
##
                        10%
                                20%
                                               40%
                                                     50%
                                                            60%
                                                                  70%
                                                                         80%
                                                                               90%
## sample_quantiles -2.280 -1.164 -0.598 -0.232 0.029 0.341 0.763 1.434 2.995
## cauchy_quantiles -3.078 -1.376 -0.727 -0.325 0.000 0.325 0.727 1.376 3.078
p values \leftarrow seq(0.95, 1, 0.01)
sample_quantiles <- quantile(post_burn_in_samples, p_values)</pre>
cauchy_quantiles <- qcauchy(p_values)</pre>
quantile_comparison <- round(rbind(sample_quantiles, cauchy_quantiles), 3)</pre>
print(quantile_comparison)
##
                       95%
                             96%
                                     97%
                                             98%
                                                    99%
                                                           100%
## sample quantiles 5.249 6.265 7.633 9.920 13.067 27.824
## cauchy_quantiles 6.314 7.916 10.579 15.895 31.821
#QQ Plot
p_points <- ppoints(100)</pre>
sample_quantiles <- quantile(post_burn_in_samples, p_points)</pre>
cauchy_quantiles <- qcauchy(p_points)</pre>
qqplot(cauchy_quantiles, sample_quantiles, cex = 0.5, pch = 16,
       main = "Q-Q Plot: Cauchy Distribution", xlab = "Theoretical Quantiles", ylab = "Sample Quantiles
abline(0, 1, col = "red")
```

Q-Q Plot: Cauchy Distribution



11.6(1)

```
{\it \# Define the Metropolis-Hastings algorithm for Laplace distribution}
rw.Laplace <- function(N, x0, sigma) {</pre>
  x <- numeric(N)
  x[1] < -x0
  u <- runif(N)
  k <- 0
  for (i in 2:N) {
    xt <- x[i - 1]
    y <- rnorm(1, xt, sigma)
    if (u[i] \leftarrow exp(abs(xt) - abs(y))) {
      x[i] <- y
    } else {
      x[i] \leftarrow x[i - 1]
      k <- k + 1
    }
  return(list(x = x, k = k))
```

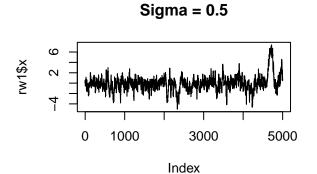
(2)

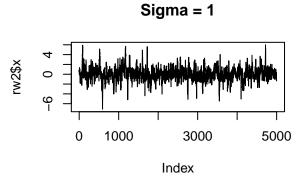
```
N <- 5000

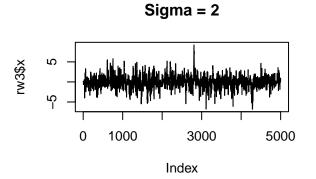
sigma <- c(0.5, 1, 2, 4)

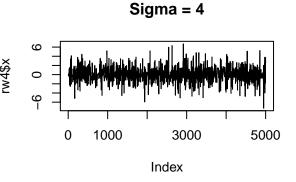
x0 <- rnorm(1)
```

```
rw1 <- rw.Laplace(N, x0, sigma[1])</pre>
rw2 <- rw.Laplace(N, x0, sigma[2])</pre>
rw3 <- rw.Laplace(N, x0, sigma[3])</pre>
rw4 <- rw.Laplace(N, x0, sigma[4])</pre>
print(c(rw1$k, rw2$k, rw3$k, rw4$k))
## [1] 834 1534 2383 3366
cat("rejection rates ", (c(rw1$k, rw2$k, rw3$k, rw4$k)/N), "\n")
## rejection rates 0.1668 0.3068 0.4766 0.6732
 (3)
# Remove burn-in samples
burn_in <- 100
y1 <- rw1$x[(burn_in + 1):N]
y2 <- rw2\$x[(burn_in + 1):N]
y3 <- rw3$x[(burn_in + 1):N]
y4 <- rw4$x[(burn_in + 1):N]
par(mfrow = c(2, 2))
plot(rw1$x, type = "l", main = paste("Sigma =", sigma[1]))
plot(rw2$x, type = "1", main = paste("Sigma =", sigma[2]))
plot(rw3$x, type = "1", main = paste("Sigma =", sigma[3]))
plot(rw4$x, type = "1", main = paste("Sigma =", sigma[4]))
```









(4)

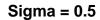
```
par(mfrow = c(2, 2))
p <- ppoints(200)
y <- qexp(p, 1)
z <- c(-rev(y), y)
fx <- 0.5 * exp(-abs(z))

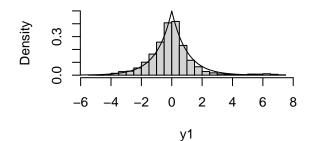
hist(y1, breaks = "Scott", freq = FALSE, ylim = c(0, 0.5), main = paste("Sigma =", sigma[1]))
lines(z, fx)

hist(y2, breaks = "Scott", freq = FALSE, ylim = c(0, 0.5), main = paste("Sigma =", sigma[2]))
lines(z, fx)

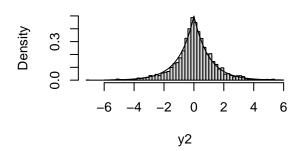
hist(y3, breaks = "Scott", freq = FALSE, ylim = c(0, 0.5), main = paste("Sigma =", sigma[3]))
lines(z, fx)

hist(y4, breaks = "Scott", freq = FALSE, ylim = c(0, 0.5), main = paste("Sigma =", sigma[4]))
lines(z, fx)</pre>
```

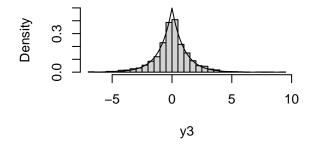




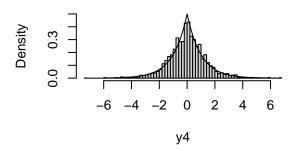
Sigma = 1



Sigma = 2



Sigma = 4



(5)

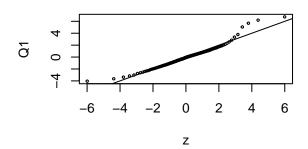
```
# Generate Q-Q plots and compare quantiles
par(mfrow = c(2, 2))
Q1 <- quantile(y1, p)
qqplot(z, Q1, cex = 0.4, main = paste("Sigma =", sigma[1]))
abline(0, 1)

Q2 <- quantile(y2, p)
qqplot(z, Q2, cex = 0.4, main = paste("Sigma =", sigma[2]))
abline(0, 1)

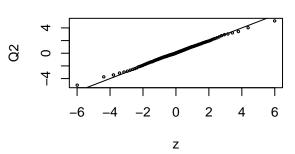
Q3 <- quantile(y3, p)
qqplot(z, Q3, cex = 0.4, main = paste("Sigma =", sigma[3]))
abline(0, 1)

Q4 <- quantile(y4, p)
qqplot(z, Q4, cex = 0.4, main = paste("Sigma =", sigma[4]))
abline(0, 1)</pre>
```

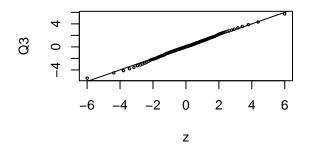
Sigma = 0.5



Sigma = 1



Sigma = 2



Sigma = 4

