### HW 10

#### Jared Brotamonte

12/6/2023

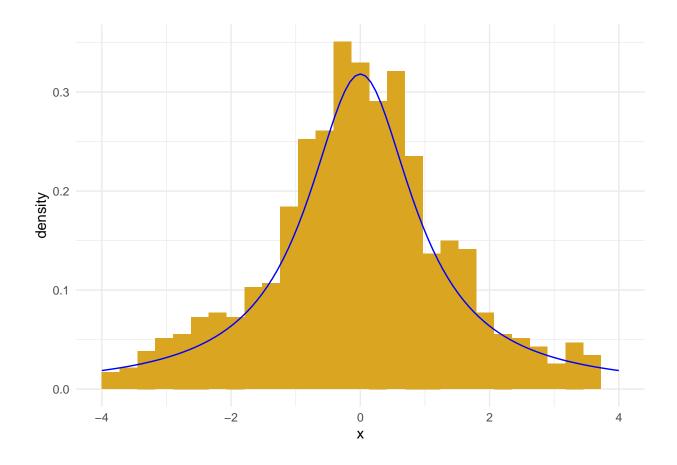
### Exercise 1

(a)

```
# Function to draw random samples from Cauchy distribution using Inverse CDF
draw_cauchy_samples <- function(n) {</pre>
  # get n random values from U(0,1)
 probs <- runif(n)</pre>
  # Inverse CDF for Cauchy distribution
 x <- qcauchy(probs, location = 0, scale = 1)
 return(x)
# Number of samples
n_{samples} \leftarrow 1000
# Draw 1000 random samples from Cauchy distribution
cauchy_samples <- data.frame(x = draw_cauchy_samples(n_samples))</pre>
# Plot the histogram and the true Cauchy density
cauchy_samples %>%
  ggplot() +
  geom_histogram(aes(x = x, y = ..density..), fill = "goldenrod", bins = 30) +
  stat_function(fun = dcauchy, color = "blue", args = list(location = 0, scale = 1)) +
 xlim(c(-4, 4)) + # Set x-axis limits to [-4, 4]
 theme_minimal()
```

```
## Warning: Removed 153 rows containing non-finite values (stat_bin).
```

<sup>##</sup> Warning: Removed 2 rows containing missing values (geom\_bar).

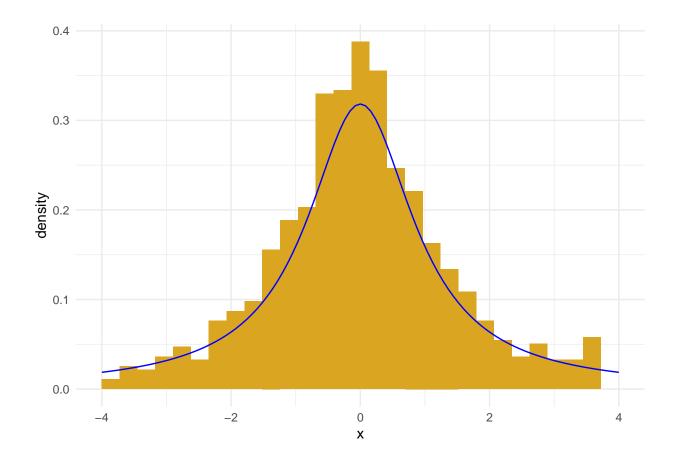


# (b)

```
# Load required libraries
library(ggplot2)
# Function to draw random samples from Cauchy distribution
draw_cauchy_samples_accept_reject <- function(n) {</pre>
  # define Cauchy density
  dcauchy_custom <- function(x) {</pre>
    return(dcauchy(x, location = 0, scale = 1))
  }
  # define parameters
  M \leftarrow 2.6
  n_samples \leftarrow n
  # create a vector to store samples
  samples <- c()</pre>
  count <- 0
  while(length(samples) < n_samples) {</pre>
    count <- count + 1
    # sample at random from x in the interval [-4, 4]
```

```
x \leftarrow runif(1, -4, 4)
    # calculate f(x)
    fx <- dcauchy_custom(x)</pre>
    \# calculate M * g(x)
    Mgx \leftarrow M * dunif(x, -4, 4) # Using a uniform distribution for g(x) in the interval [-4, 4]
    # calculate acceptance probability
    prob <- fx / Mgx</pre>
    # draw random sample from U(0, 1)
    q <- runif(1)
    # accept or reject sample
    if (q < prob) {
      samples <- append(samples, x)</pre>
    }
  }
  return(samples)
# Number of samples
n_samples \leftarrow 1000
# Draw 1000 random samples from Cauchy distribution using Accept-Reject method
cauchy_samples_accept_reject <- draw_cauchy_samples_accept_reject(n_samples)</pre>
# Plot the histogram and the true Cauchy density
data.frame(x = cauchy_samples_accept_reject) %>%
  ggplot() +
  geom_histogram(aes(x = x, y = ..density..), fill = "goldenrod", bins = 30) +
  stat_function(fun = dcauchy, color = "blue", args = list(location = 0, scale = 1)) +
  xlim(c(-4, 4)) + # Set x-axis limits to [-4, 4]
 theme_minimal()
```

## Warning: Removed 2 rows containing missing values (geom\_bar).



(c)

The histograms of part a and b look largely similar.

# Exercise 2

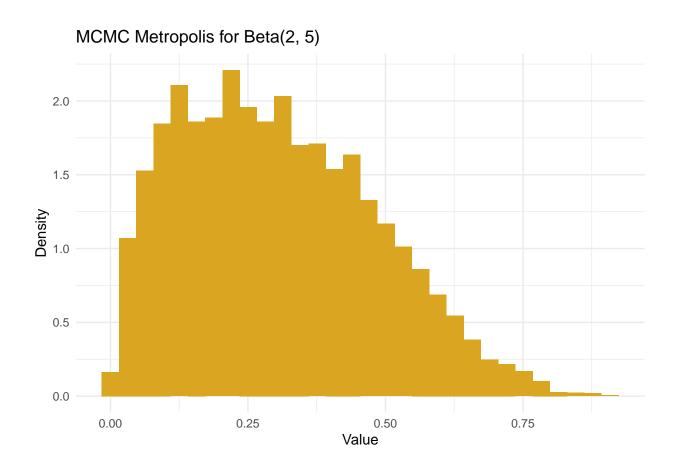
(a)

```
# Define the Beta distribution function
dbeta_custom <- function(x) {
   return(dbeta(x, shape1 = 2, shape2 = 5))
}

# MCMC Metropolis algorithm for Beta distribution
n_samples <- 10000
count <- 0
samples <- c()
burn_in <- 100

# Initial point
x <- runif(1)</pre>
```

```
while (length(samples) < (n_samples + burn_in)) {</pre>
  count <- count + 1</pre>
  # Calculate f(x)
  fx <- dbeta_custom(x)</pre>
  # Generate candidate point at random from a uniform distribution
  x.star <- runif(1)</pre>
  # Calculate f(x.star)
  fx.star <- dbeta_custom(x.star)</pre>
  # Accept or reject sample
  if (fx.star > fx) {
    samples <- append(samples, x.star)</pre>
    x <- x.star
  } else {
    # Draw\ random\ sample\ from\ U(0,1)
    q <- runif(1)
    if (q < (fx.star / fx)) {</pre>
      samples <- append(samples, x.star)</pre>
      x <- x.star
    }
  }
}
# Remove burn-in
samples <- samples[(burn_in + 1):length(samples)]</pre>
# Plot the sample
data.frame(x = samples) %>%
  ggplot() +
  geom_histogram(aes(x = x, y = ..density..), fill = "goldenrod", bins = 30) +
  labs(title = "MCMC Metropolis for Beta(2, 5)", x = "Value", y = "Density") +
  theme_minimal()
```



(b)

```
# Define the Beta distribution function
dbeta_custom <- function(x) {
   return(dbeta(x, shape1 = 2, shape2 = 5))
}

# Plot histogram and overlay PDF
data.frame(x = samples) %>%
   ggplot() +
   geom_histogram(aes(x = x, y = ..density..), fill = "goldenrod", bins = 30) +
   stat_function(fun = dbeta_custom, color = "blue", size = 1.5) +
   labs(title = "Histogram and PDF Overlay", x = "Value", y = "Density") +
   theme_minimal()
```

