

# Computer Lab 3

Tim Yuki Washio

11/19/2020

## Question 1: Stable Distribution

1.

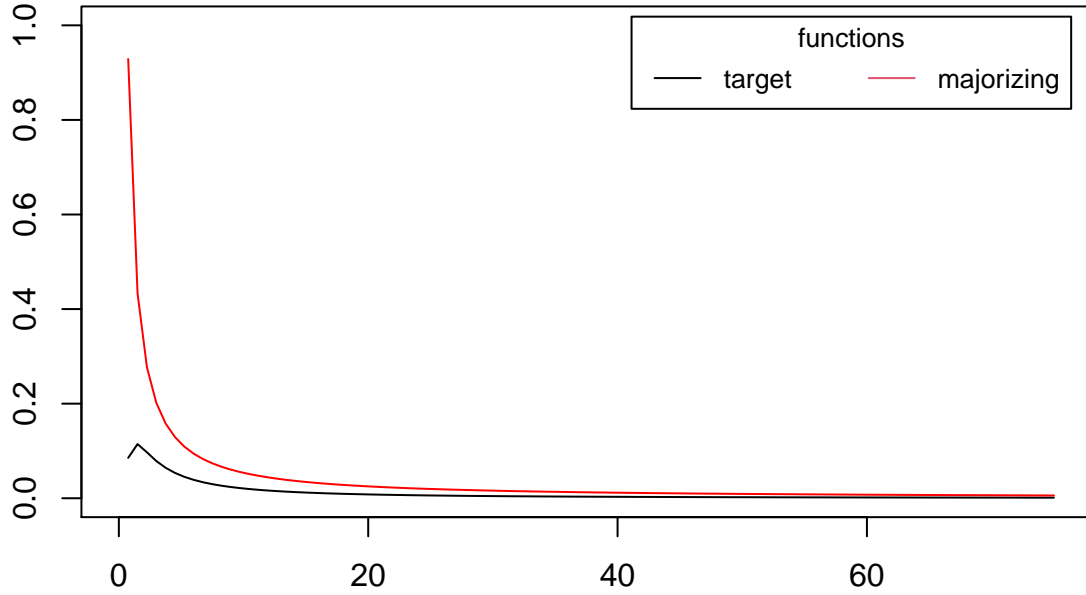
Plotting  $f(x)$  with  $c = 2$  and  $f_p(x)$  with  $\alpha = 2$  and  $T_{min} = 2$

```
c = 2
t_min = 1.5
alpha = 1.1

plot(1, xlim = c(0,75), ylim = c(0, 1), type = "n", xlab = "", ylab = "", main = "f(x) ~ f_p(x)")
curve(eval(c) * (sqrt(2 * pi)^(-1)) * exp(-eval(c)^(2) / (2 * x)) * x^(-3/2), from=0, to=75, add=TRUE, col="blue")
curve((eval(alpha) - 1 / eval(t_min)) * (x / eval(t_min))^(alpha), from=0, to=75, add=TRUE, col="red")

legend("topright", inset=.02, title="functions",
      c("target","majorizing"), horiz=TRUE, cex=0.8, col = 1:2, lty = 1)
```

$$f(x) \sim f_p(x)$$



The power-law distribution should not be used by itself because for small values of  $x$   $f_p(x) = \infty$ . If we sample from our majorizing distribution in this area, we will get a huge number of rejections. It might be better to combine the power-law distribution with another distribution that majorizes our target function in that area. We could use a uniform distribution with support  $(0, T_{min})$  in addition to our given distribution with support  $(T_{min}, \infty)$ .

Power-law distribution values:

$$\frac{df(x)}{dx} = \frac{ce^{(-c^2)/2x}(c^2 - 3x)}{2\sqrt{(2\pi)}x^{7/2}}$$

This has one root for  $c \neq 0$ ,  $x = \frac{c^2}{3}$ .

$$\max(f(x)) = f\left(\frac{c^2}{3}\right)$$

$$f_p(x) = f\left(\frac{c^2}{3}\right)$$

$$x = T_{min}$$

## Question 2: Laplace Distribution

**Write a code generating random numbers from the double exponential distribution**

The double exponential distribution is given by the following formula:

$$DE(\mu, \alpha) = \frac{\alpha}{2} \exp(-\alpha |x - \mu|)$$

To get to the inverse CDF,  $F^{-1}(p)$ , of this function we first need to get the CDF  $F(x)$

$$F(x) = \int_{-\infty}^x f(u) du = \begin{cases} \frac{1}{2} e^{\frac{x-\mu}{\alpha}} & \text{if } x < \mu \\ 1 - \frac{1}{2} e^{\frac{x-\mu}{\alpha}} & \text{if } x \geq \mu \end{cases}$$

After transformation, we get

$$F(x) = \frac{1}{2} + \frac{1}{2} \operatorname{sgn}(x - \mu) \left( 1 - \exp\left(-\frac{|x - \mu|}{\alpha}\right) \right)$$

Now we calculate the inverse CDF  $F^{-1}(p)$  by using the CDF  $F(x)$ . We set:

$$y = \frac{1}{2} + \frac{1}{2} \operatorname{sgn}(x - \mu) (1 - \exp(-\alpha |x - \mu|))$$

And we now solve for  $x$  to obtain the inverse CDF. We have two cases. When  $x \geq \mu$  where we then have:

$$x = \mu - \frac{1}{\alpha} \ln(2 - 2y)$$

and the case when  $x < \mu$  when we have:

$$x = \mu + \frac{1}{\alpha} \ln(2y)$$

We combine both expressions and express them with respect to the sign of the difference between  $x$  and  $\mu$ . We obtain the following:

$$F^{-1}(y) = \mu + \frac{1}{\alpha} \ln(1 + \operatorname{sgn}(x - \mu) - \operatorname{sgn}(x - \mu) 2y)$$

However, we would like an expression that does not depend on the sign of  $x - \mu$ . We try to find a quantity related to it but with respect to  $y$ . We investigate the critical value of  $y$  when the sign of  $x - \mu$  changes (i.e. we look for  $\operatorname{sgn}(x - \mu) \frac{1}{\alpha} \ln(1 + \operatorname{sgn}(x - \mu) - \operatorname{sgn}(x - \mu) 2y) = x - \mu$  when  $x$  is greater and smaller than  $\mu$ .)

We find that  $\operatorname{sgn}(x - \mu) = \operatorname{sgn}(y - \frac{1}{2})$ . We can simplify the expression of  $F^{-1}(u)$  and finally obtain:

$$F^{-1}(u) = \mu - \alpha \operatorname{sgn}(u - \frac{1}{2}) \ln(1 - 2|u - \frac{1}{2}|)$$

```
# Given density function
laplace_density <- function(x, mu, alpha) {
  result = (alpha/2)*exp(-alpha*abs(x-mu))
  return(result)
}

# Calculated inverse CDF for Laplace function
```

```

laplace_inv_cdf = function(p, mu, alpha){
  result = mu-alpha*sign(p-0.5)*log(1-2*abs(p-0.5))
  return(result)
}

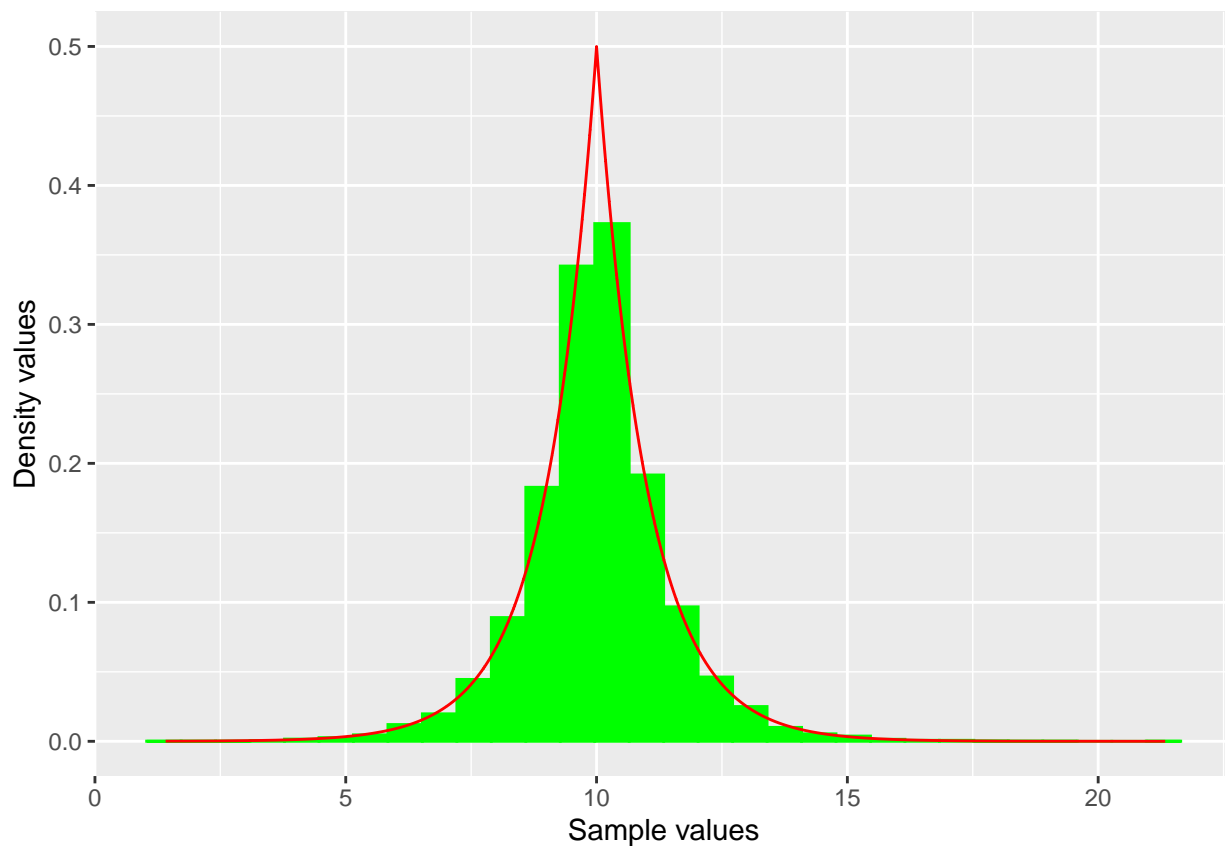
# Random number generation with n=10000, mu=10, alpha=1
n = 10000
unif_sample = runif(n)
sample = laplace_inv_cdf(unif_sample, 10, 1)
sample_density = laplace_density(sample, 10, 1)

# Plotting the histogram
hist_plot <- ggplot() +
  geom_histogram(aes(x = sample, y = ..density..), col = "green", fill="green") +
  geom_line(aes(x = sample, y = sample_density), col = "red") +
  xlab("Sample values") + ylab("Density values")

hist_plot

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

```



From this graph, we can see that our sampled numbers follow the distribution.

## Acceptance rejection method using DE(0,1) as a majorizing density for N(0,1).

The main task to solve this exercise is to find the majorizing constant  $c$  such that

$$c \cdot f_M(x) \geq f_T(x)$$

Where  $f_M(x)$  is the majorizing density and  $f_T(x)$  is the density we wish to sample from (target density). This inequality must hold true for all  $x$  that are on the support of the target function. We must choose  $c$  to be large enough for the inequality to hold true, but not so large that the rejection rate for the acceptance/rejection algorithm becomes too great. Setting the parameters in both densities to be (1,0) and setting our inequality, we have:

$$c \cdot \frac{1}{2} \exp -|x| \geq \frac{1}{\sqrt{2\pi}} \exp \left( -\frac{x^2}{2} \right)$$

We solve for  $c$  and obtain:

$$c \geq \sqrt{\frac{2}{\pi}} \exp (|x| - x^2/2)$$

We can find a solution for  $c$  for  $x > 0$  because we have an even function on the right-hand side and the other maximum will be attained at  $x = -x_{positive}$  with the same value for  $c$ . The expression maximizes for  $x = 1$  and yields:

$$c_{major} = \sqrt{\frac{2e}{\pi}}$$

```
# Takes as arguments the following:
#       · The number n of observations to generate;
#       · The target density f (standard normal distribution);
#       · The majorizing density g (standard laplace distribution);
#       · The function rg to generate random numbers according to the density g;
#       · The majorizing constant k.

# random_norm_AR <- acc_rej_norm(2000, dnorm, dlaplace, rlaplace, k)
# random_norm_st <- rnorm(2000)

set.seed(12345)
c_major <- sqrt( (2*exp(1)) / pi )

accept_reject <- function(n){
  results <- rep(0,times = n)
  counter <- 0

  for(i in 1:n){
    reject <- TRUE
    while(reject == TRUE){
      counter <- counter + 1
      random_major <- laplace_inv_cdf(runif(1), 0, 1)
      random_uniform <- runif(1)
      if(random_uniform <= dnorm(random_major)/(c_major*laplace_density(random_major, 0, 1))){
        results[i] <- random_major
      }
    }
  }
}
```

```

    reject <- FALSE
  }
}
}
return(list(rn = results, draws = counter))
}

```

```

n = 2000
unif_sample = rnorm(n)
sample = accept_reject(n)

```

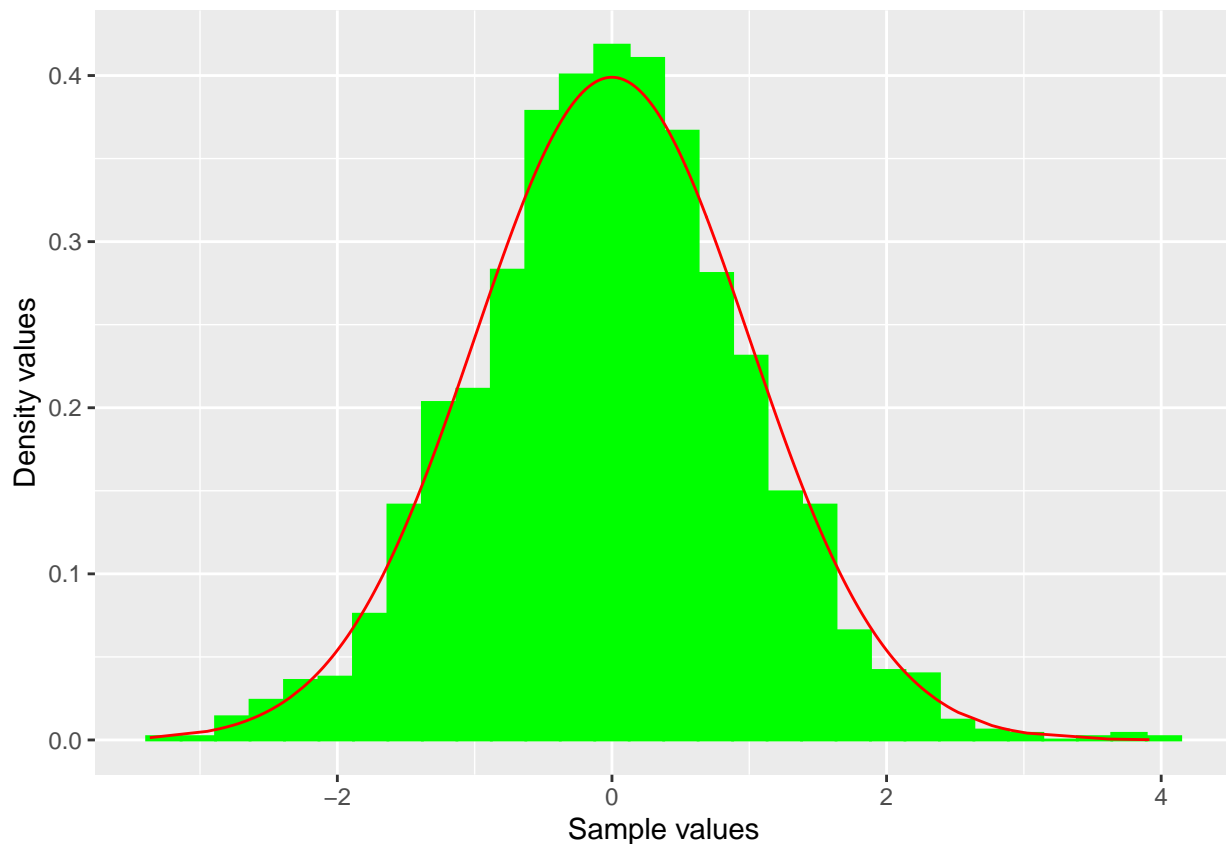
```

# Plotting the histogram
p1 <- ggplot() +
  geom_histogram(aes(x = sample$rn, y = ..density..), col = "green", fill="green") +
  geom_line(aes(x = sample$rn, y = dnorm(sample$rn)), col = "red") +
  xlab("Sample values") + ylab("Density values")

p1

```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



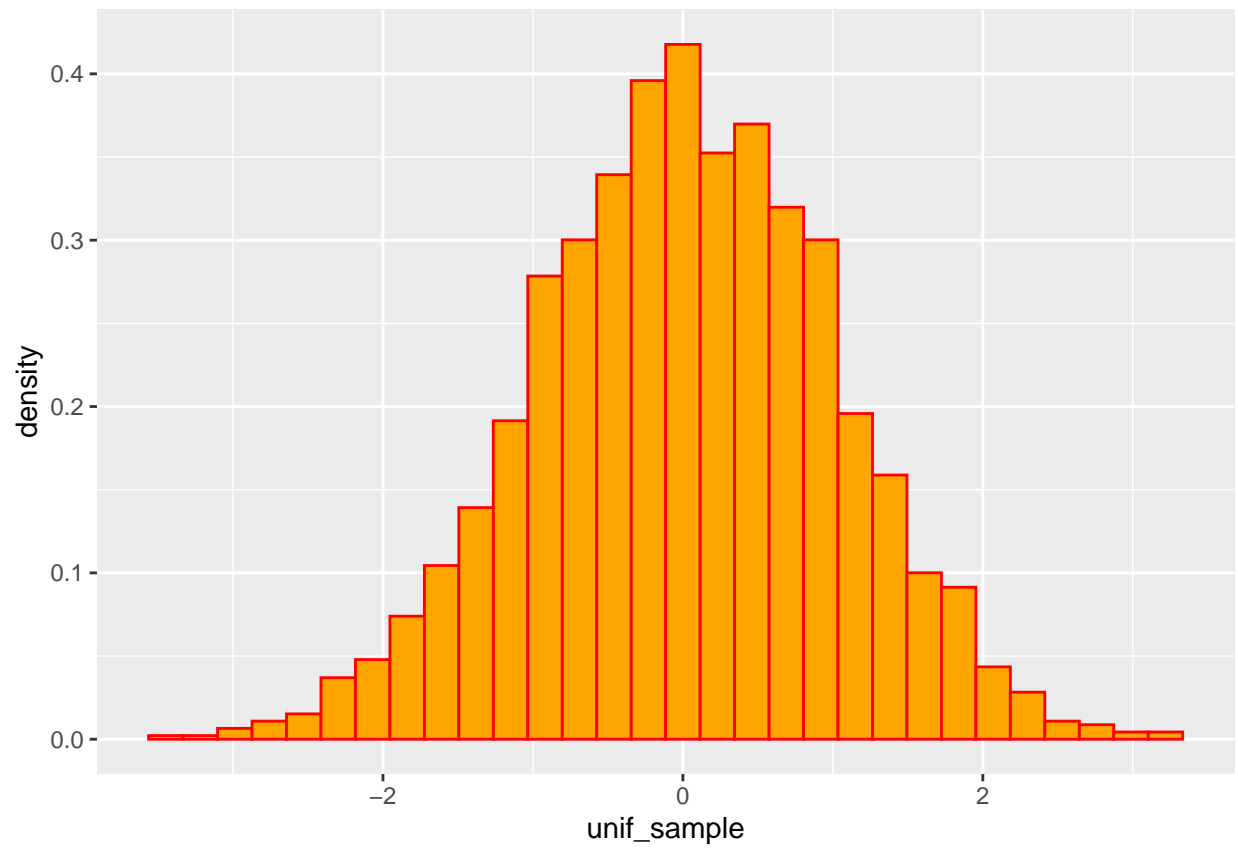
```

# histogram using rnorm
p2 <- ggplot() +
  geom_histogram(aes(x= unif_sample, y= ..density..), col = "red", fill = "orange")

p2

```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



## Appendix

```
knitr::opts_chunk$set(echo = TRUE)
library(ggplot2)
c = 2
t_min = 1.5
alpha = 1.1

plot(1, xlim = c(0,75), ylim = c(0, 1), type = "n", xlab = "", ylab = "", main = "f(x) ~ f_p(x)")
curve(eval(c) * (sqrt(2 * pi)^(-1)) * exp(-eval(c)^(2) / (2 * x)) * x^(-3/2), from=0, to=75, add=TRUE, col="red")
curve((eval(alpha) - 1 / eval(t_min)) * (x / eval(t_min))^(eval(alpha)), from=0, to=75, add=TRUE, col="green")

legend("topright", inset=.02, title="functions",
      c("target","majorizing"), horiz=TRUE, cex=0.8, col = 1:2, lty = 1)
# Given density function
laplace_density <- function(x, mu, alpha) {
  result = (alpha/2)*exp(-alpha*abs(x-mu))
  return(result)
}

# Calculated inverse CDF for Laplace function
laplace_inv_cdf = function(p, mu, alpha){
  result = mu-alpha*sign(p-0.5)*log(1-2*abs(p-0.5))
  return(result)
}

# Random number generation with n=10000, mu=10, alpha=1
n = 10000
unif_sample = runif(n)
sample = laplace_inv_cdf(unif_sample, 10, 1)
sample_density = laplace_density(sample, 10, 1)
# Plotting the histogram
hist_plot <- ggplot() +
  geom_histogram(aes(x = sample, y = ..density..), col = "green", fill="green") +
  geom_line(aes(x = sample, y = sample_density), col = "red") +
  xlab("Sample values") + ylab("Density values")

hist_plot
# Takes as arguments the following:
#       · The number n of observations to generate;
#       · The target density f (standar normal distribution);
#       · The majorizing density g (standard laplace distribution);
#       · The function rg to generate random numbers according to the density g;
#       · The majorizing constant k.

# random_norm_AR <- acc_rej_norm(2000, dnorm, dlaplace, rlaplace, k)
# random_norm_st <- rnorm(2000)

set.seed(12345)
c_major <- sqrt( (2*exp(1)) / pi )
```



```

accept_reject <- function(n){
  results <- rep(0,times = n)
  counter <- 0

  for(i in 1:n){
    reject <- TRUE
    while(reject == TRUE){
      counter <- counter + 1
      random_major <- laplace_inv_cdf(runif(1), 0, 1)
      random_uniform <- runif(1)
      if(random_uniform <= dnorm(random_major)/(c_major*laplace_density(random_major, 0, 1))){
        results[i] <- random_major
        reject <- FALSE
      }
    }
  }
  return(list(rn = results, draws = counter))
}

n = 2000
unif_sample = rnorm(n)
sample = accept_reject(n)

# Plotting the histogram
p1 <- ggplot() +
  geom_histogram(aes(x = sample$rn, y = ..density..), col = "green", fill="green") +
  geom_line(aes(x = sample$rn, y = dnorm(sample$rn)), col = "red") +
  xlab("Sample values") + ylab("Density values")

p1

# histogram using rnorm
p2 <- ggplot() +
  geom_histogram(aes(x= unif_sample, y= ..density..), col = "red", fill = "orange")

p2

```