

# TMA4300: Exercise 1

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1/29/2022

## Problem A

### A.1

The exponential distribution has cumulative density function (CDF)

$$F(x) = 1 - e^{-\lambda x},$$

with rate parameter  $\lambda$ . By defining  $u := F(x)$ , we can express  $x$  as

$$x = -\frac{1}{\lambda} \ln(1 - u) =: F^{-1}(u).$$

This means that we can use the *inversion method* to simulate from the exponential distribution. I.e., we let  $U \sim \mathcal{U}_{[0,1]}$  and calculate  $X = F^{-1}(U)$ . Then,  $X \sim \text{Exp}(\lambda)$ . The function which simulates the exponential distribution is given below.

```
sim.exp <- function(rate, n){  
  u <- runif(n,0,1)  
  return(-1/rate * log(1 - u))  
}
```

Next, we need to check if this gives reasonable results by comparing our simulated values to the theoretical knowledge.

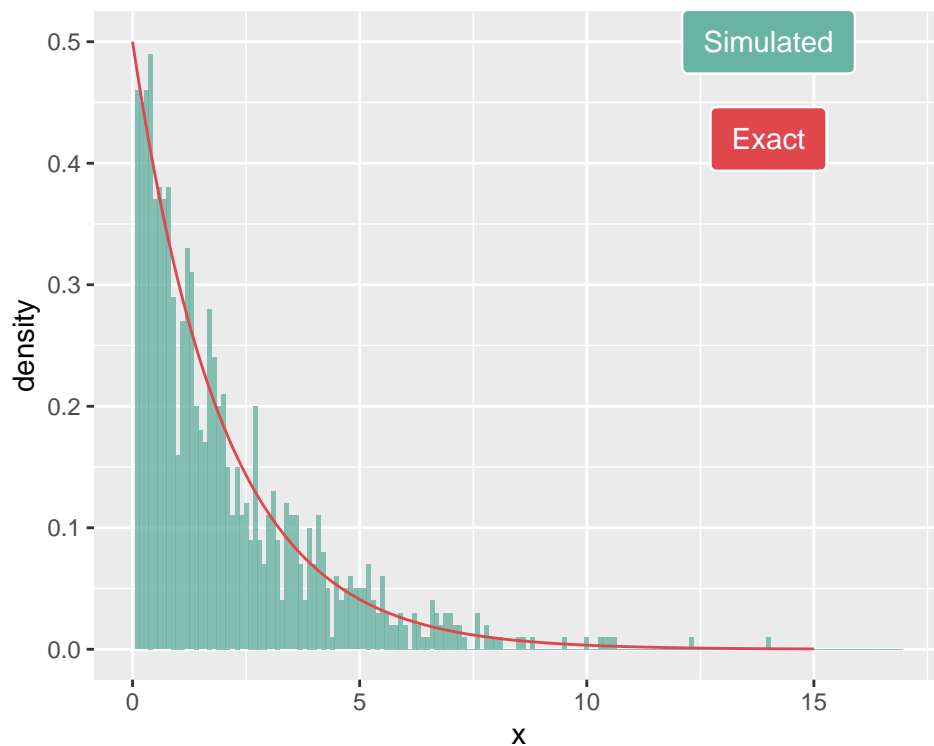
```
rate <- 0.5  
n <- 1000  
sim <- data.frame(x = sim.exp(rate, n))  
x = seq(from = 0, to = 15, by = 0.1)  
exact <- data.frame(x = x, y = rate*exp(-rate*x))  
  
ggplot(sim) +  
  geom_histogram(aes(x = x, y = ..density..),  
                 alpha = 0.8, fill = "#69b3a2", binwidth = 0.1) +  
  geom_line(data = exact, aes(x = x, y = y), color = "#e0474c") +  
  geom_label(  
    label="Simulated",  
    x=14,  
    y=0.5,  
    label.padding = unit(0.55, "lines"), # Rectangle size around label
```

```

    label.size = 0.35,
    color = "white",
    fill= "#69b3a2"
  ) +
  geom_label(
    label="Exact",
    x=14,
    y=0.42,
    label.padding = unit(0.55, "lines"), # Rectangle size around label
    label.size = 0.35,
    color = "white",
    fill = "#e0474c"
  ) +
  xlim(0,17)

```

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

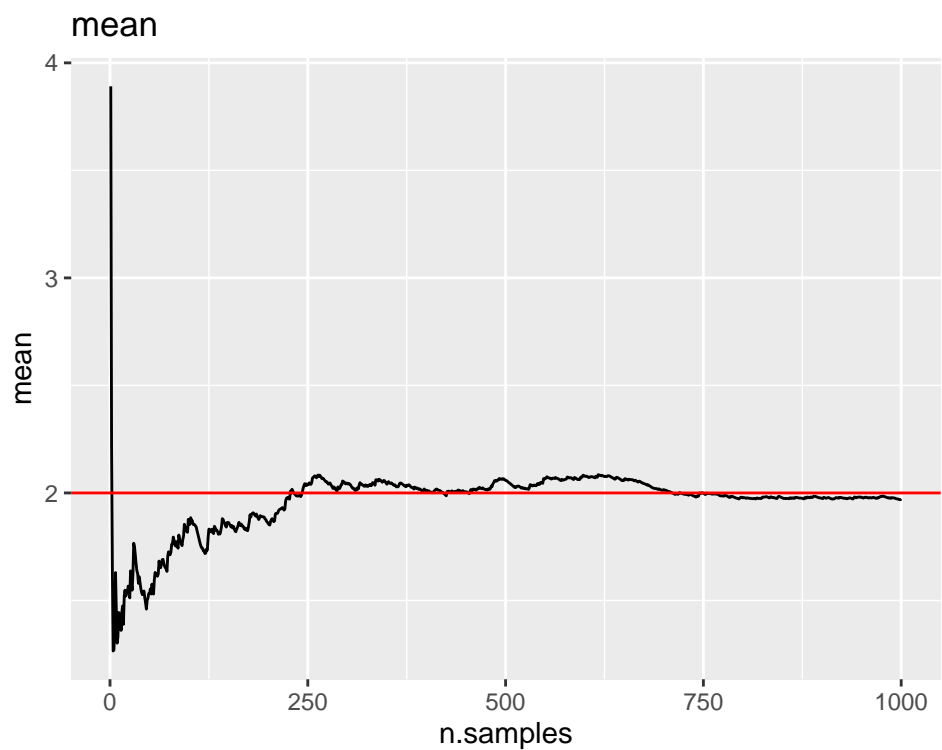


We also compare the estimated mean and variance to first and second central moments of the exponential distribution

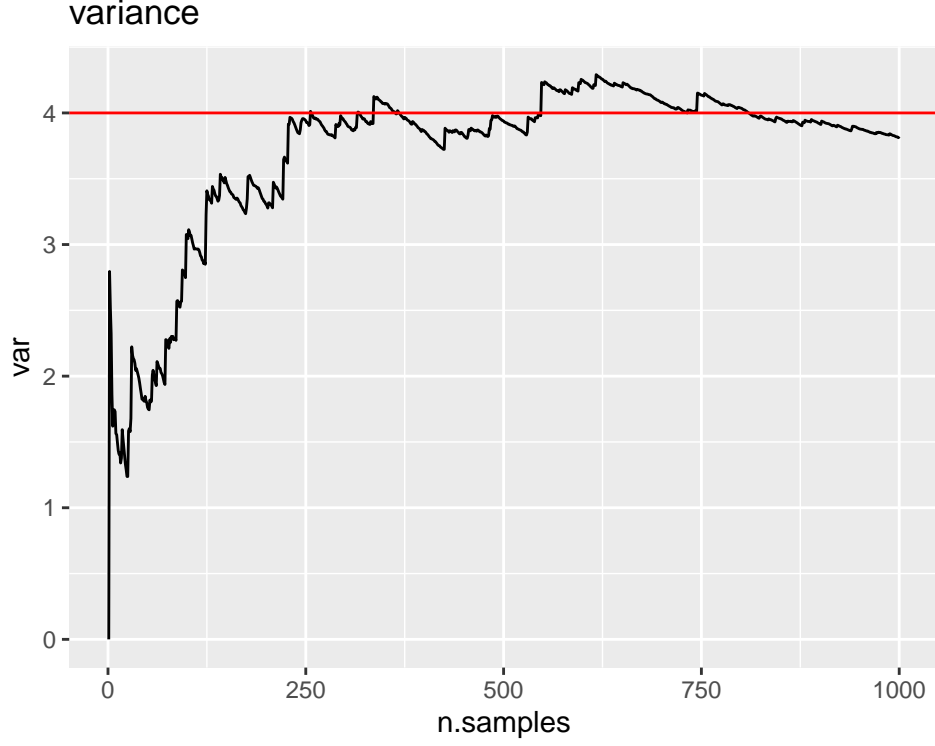
```

# mean
data.frame(n.samples = 1:n,
           mean = cumsum(sim$x)/(1:n)) %>%
  ggplot() + geom_line(aes(n.samples, mean)) +
  geom_hline(yintercept = 1/rate, color = "red") +
  ggtitle("mean")

```



```
# variance
data.frame(n.samples = 1:n,
           mean = cumsum(sim$x)/(1:n),
           mean2 = cumsum(sim$x^2)/(1:n)) %>%
  mutate(var = mean2-mean^2) %>%
  ggplot() + geom_line(aes(n.samples, var)) +
  geom_hline(yintercept = 1/rate^2, color = "red") +
  ggtitle("variance")
```



We observe that the the sampled mean and variance approach the theoretical values as the number of samples grows larger.

### A.3

a)

To find the value of  $c$ , we integrate the density over the entire domain and equate the result to 1:

$$1 = \int_{-\infty}^{\infty} f(x) dx = c \int_{-\infty}^{\infty} \frac{e^{\alpha x}}{(1 + e^{\alpha x})^2} dx.$$

To progress from here, we introduce the substitution,  $v = e^{\alpha x}$ , which gives

$$1 = \frac{c}{\alpha} \int_0^{\infty} \frac{dv}{(1 + v)^2} = \frac{c}{\alpha} \left( -\frac{1}{1 + v} \right) \Big|_0^{\infty} = \frac{c}{\alpha}.$$

Consequently,  $c = \alpha$ .

b)

The CDF is defined as follows,

$$\begin{aligned} F(x) &= \int_{-\infty}^x f(z) dz = \int_0^{\exp(\alpha x)} \frac{dv}{(1 + v)^2} \\ &= 1 - \frac{1}{1 + \exp(\alpha x)} = \frac{\exp(\alpha x)}{1 + \exp(\alpha x)}, \end{aligned}$$

where we have used the same substitution as earlier, namely  $v = e^{\alpha z}$ . We notice that  $F(x)$  is the Sigmoid function, which has the well known logit-function as its inverse. I.e.,

$$F^{-1}(x) = \frac{1}{\alpha} \ln \left( \frac{x}{1-x} \right).$$

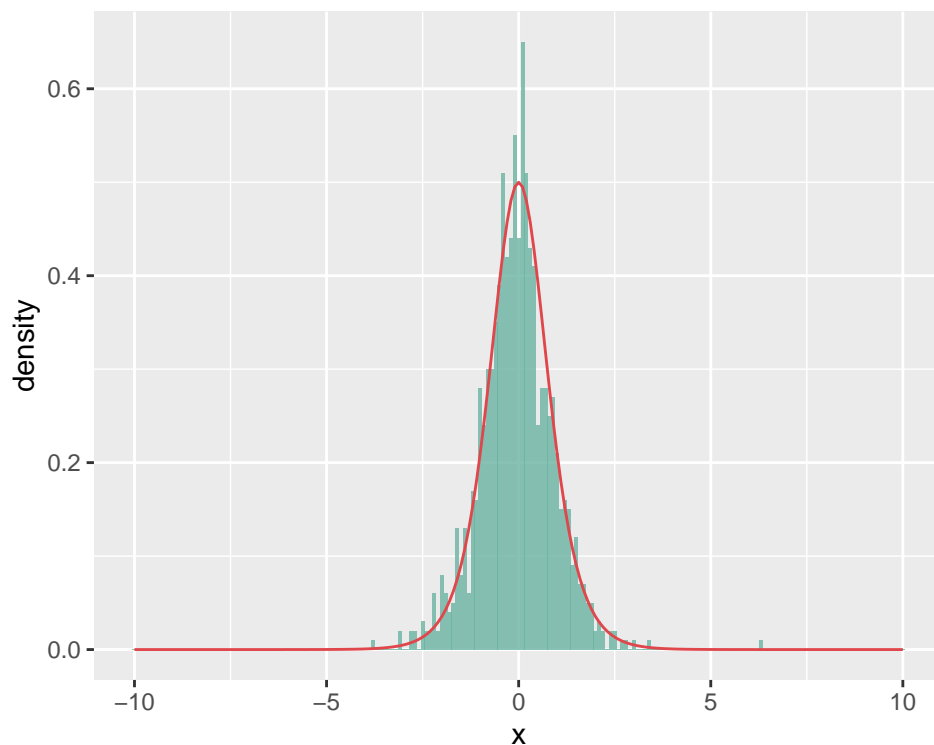
### c)

Since we have an analytic expression for the inverse, we can again use the *inversion method* to sample from  $f$ . The sampling-function is given below.

```
sim.sig<- function(alpha, n){
  u <- runif(n,0,1)
  return(1/alpha * log(u/(1-u)))
}

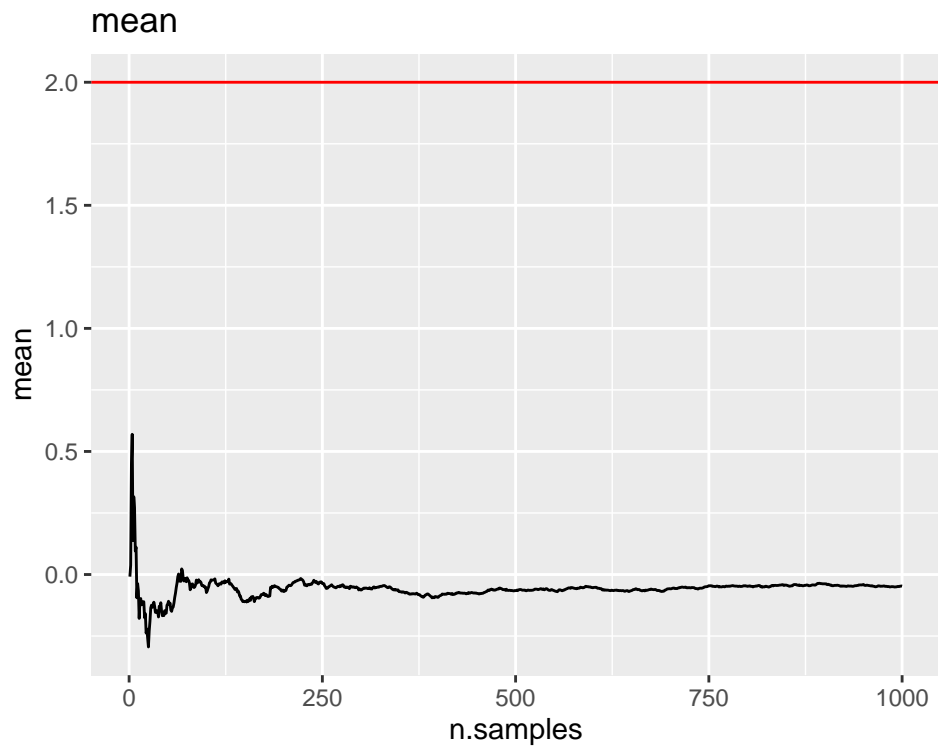
alpha <- 2
n <- 1000
sim <- data.frame(x = sim.sig(alpha, n))
x = seq(from = -10, to = 10, by = 0.1)
exact <- data.frame(x = x, y = alpha*exp(alpha*x)/(1 + exp(alpha*x))^2)

ggplot(sim) +
  geom_histogram(aes(x = x, y = ..density..),
                 alpha = 0.8, fill = "#69b3a2", binwidth = 0.1) +
  geom_line(data = exact , aes(x = x, y = y), color = "#e0474c")
```

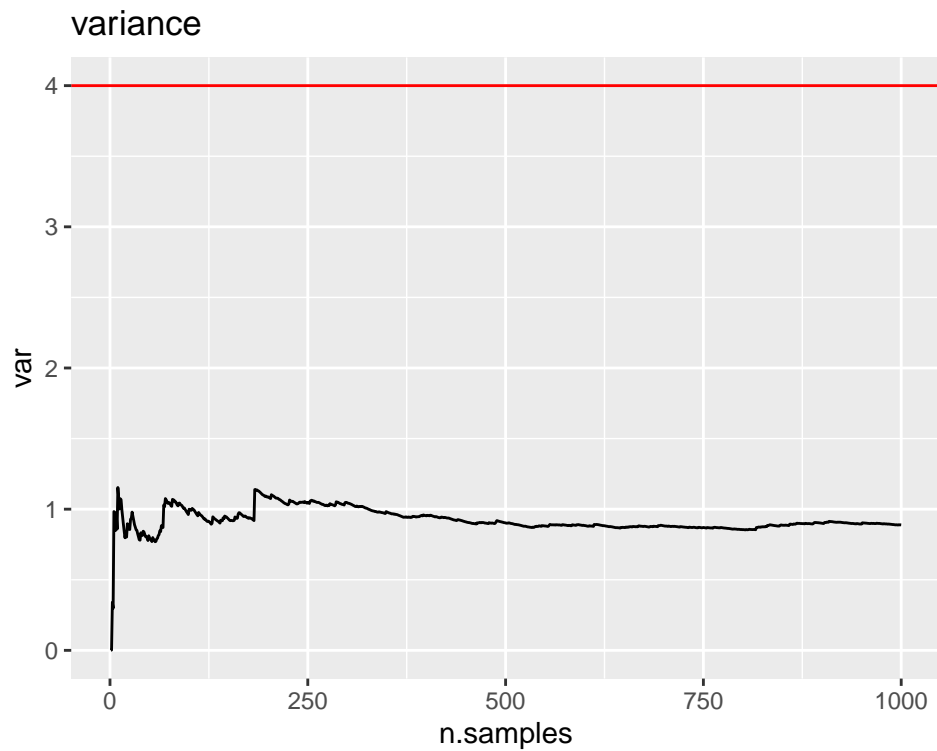


To compare the simulated values with the theoretical mean and variance, we first need to compute the expression of these moments. **Think maybe the mean is undefined??**

```
# mean
data.frame(n.samples = 1:n,
           mean = cumsum(sim$x)/(1:n)) %>%
  ggplot() + geom_line(aes(n.samples, mean)) +
  geom_hline(yintercept = 1/rate, color = "red") +
  ggtitle("mean")
```



```
# variance
data.frame(n.samples = 1:n,
           mean = cumsum(sim$x)/(1:n),
           mean2 = cumsum(sim$x^2)/(1:n)) %>%
  mutate(var = mean2-mean^2) %>%
  ggplot() + geom_line(aes(n.samples, var)) +
  geom_hline(yintercept = 1/rate^2, color = "red") +
  ggtitle("variance")
```



#### A.4

Below is our implementation of the Box-Muller algorithm.

```
box.mul <- function(n){
  odd <- FALSE
  if(n %% 2 != 0){
    odd <- TRUE
    n <- n + 1
  }

  x1 <- runif(n/2, 0, 2*pi)
  x2 <- sim.exp(0.5, n/2)

  y1 <- sqrt(x2)*cos(x1)
  y2 <- sqrt(x2)*sin(x1)

  concat <- c(y1,y2)
  if(odd){
    return(head(concat, -1))
  }
  else{
    return(concat)
  }
}
```

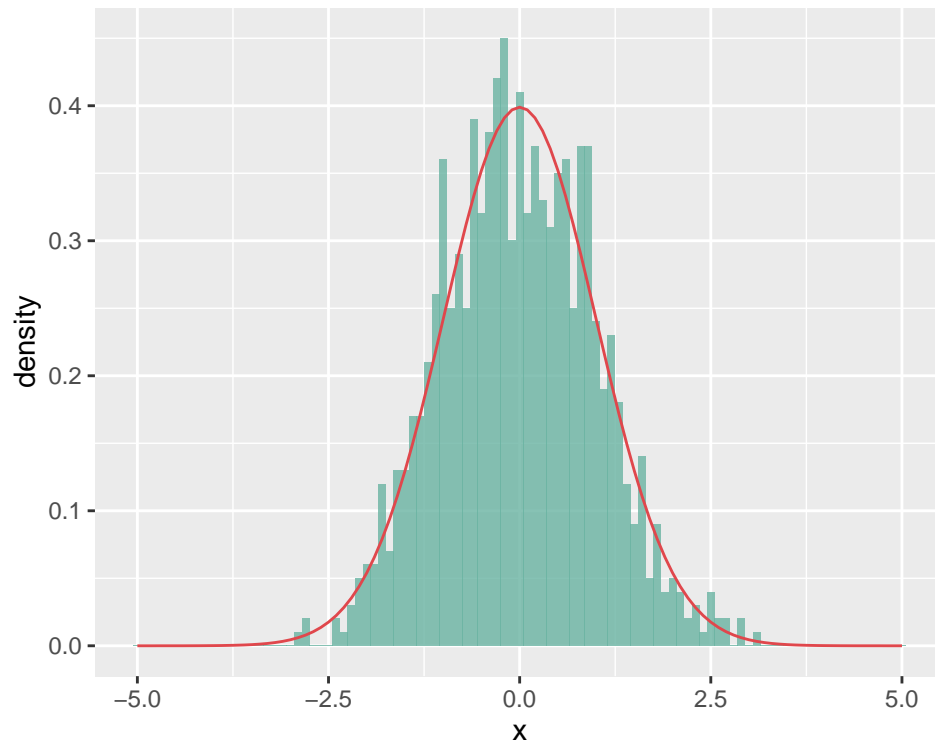
As usual, we compare the results of a simulation against the theoretical distribution.

```

n <- 1000
sim.box.mul <- data.frame(x = box.mul(n))
x = seq(from = -5, to = 5, by = 0.1)
exact <- data.frame(x = x, y = 1/sqrt(2*pi)*exp(-1/2*x^2))

ggplot(sim.box.mul) +
  geom_histogram(aes(x = x, y = ..density..),
                 alpha = 0.8, fill = "#69b3a2", binwidth = 0.1) +
  geom_line(data = exact, aes(x = x, y = y), color = "#e0474c")

```



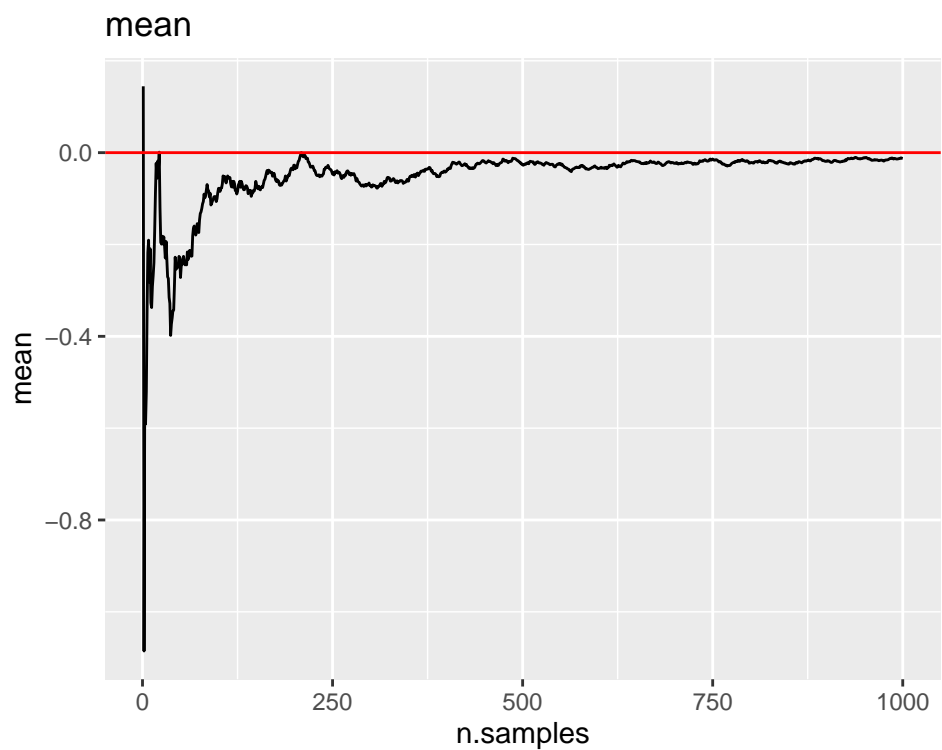
Finally, we compare the mean and variance resulting from the simulation with the theoretical values.

```

# mean
data.frame(n.samples = 1:n,
           mean = cumsum(sim.box.mul$x)/(1:n)) %>%
  ggplot() + geom_line(aes(n.samples, mean)) +
  geom_hline(yintercept = 0, color = "red") +
  ggtitle("mean")

```





```
# variance
data.frame(n.samples = 1:n,
           mean = cumsum(sim.box.mul$x)/(1:n),
           mean2 = cumsum(sim.box.mul$x^2)/(1:n)) %>%
  mutate(var = mean2-mean^2) %>%
  ggplot() + geom_line(aes(n.samples, var)) +
  geom_hline(yintercept = 1, color = "red") +
  ggtitle("variance")
```

