

# Homework 6

*Hanao Li hl3202*

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#####Part 1: Inverse Transform Method 1.

If  $X \sim \text{Cauchy}(\alpha, \beta)$  and  $U \sim \text{Unif}(0, 1)$ , then  $X = \alpha + \beta \tan(\pi(U - 1/2))$

Since we have  $X \sim \text{Cauchy}(0, 1)$ , then  $X = \tan(\pi(U - 1/2))$

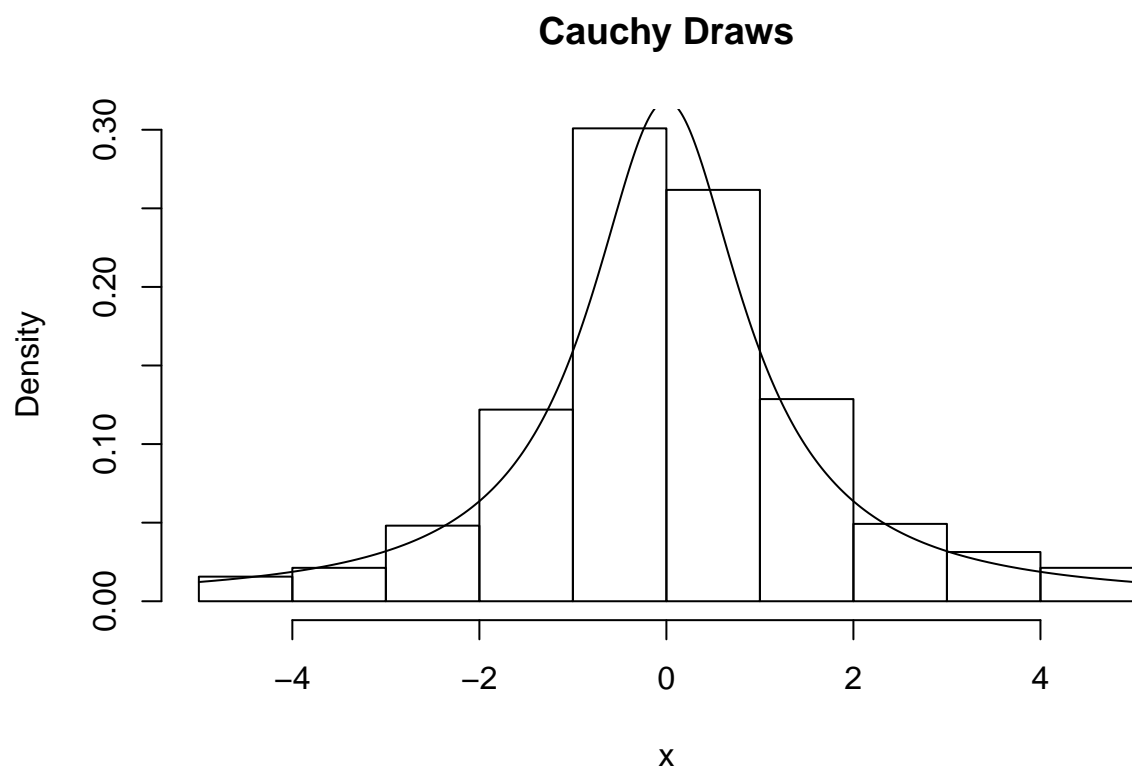
2.

```
cauchy.sim <- function(n){  
  u <- runif(n)  
  return(tan(pi*(u-0.5)))  
}  
cauchy.sim(10)
```

```
## [1] -0.8155170  1.6239486 -2.4251509  1.4982274  0.1161945  0.1935426  
## [7] -0.2744881 -0.3201585 -1.1832637 22.6364942
```

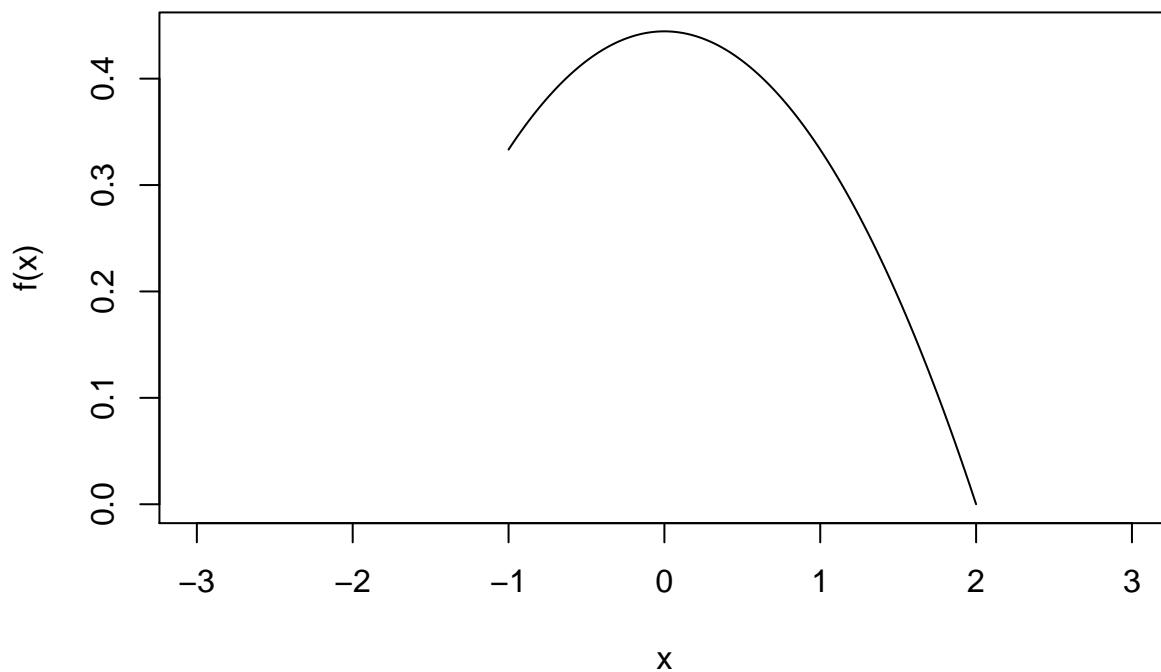
3.

```
cauchy.draws <- cauchy.sim(1000)  
hist(cauchy.draws[cauchy.draws < 5 & cauchy.draws > -5], prob = T, main = "Cauchy Draws", xlab = "x")  
x <- seq(-5, 5, 0.01)  
lines(x, 1/pi*(1+x^2))
```



#####Part 2: Reject-Accept Method 4.

```
f <- function(x){  
  return(ifelse((x < -1 | x > 2), 0, (4-x^2)/9))  
}  
x <- seq(-1,2,0.01)  
plot(x, f(x), xlim = c(-3,3), type = 'l')
```



5.

$f'(x) = -\frac{2x}{9} = 0$ , then  $X_{max} = 0$ ,  $f_{max} = \frac{4}{9}$

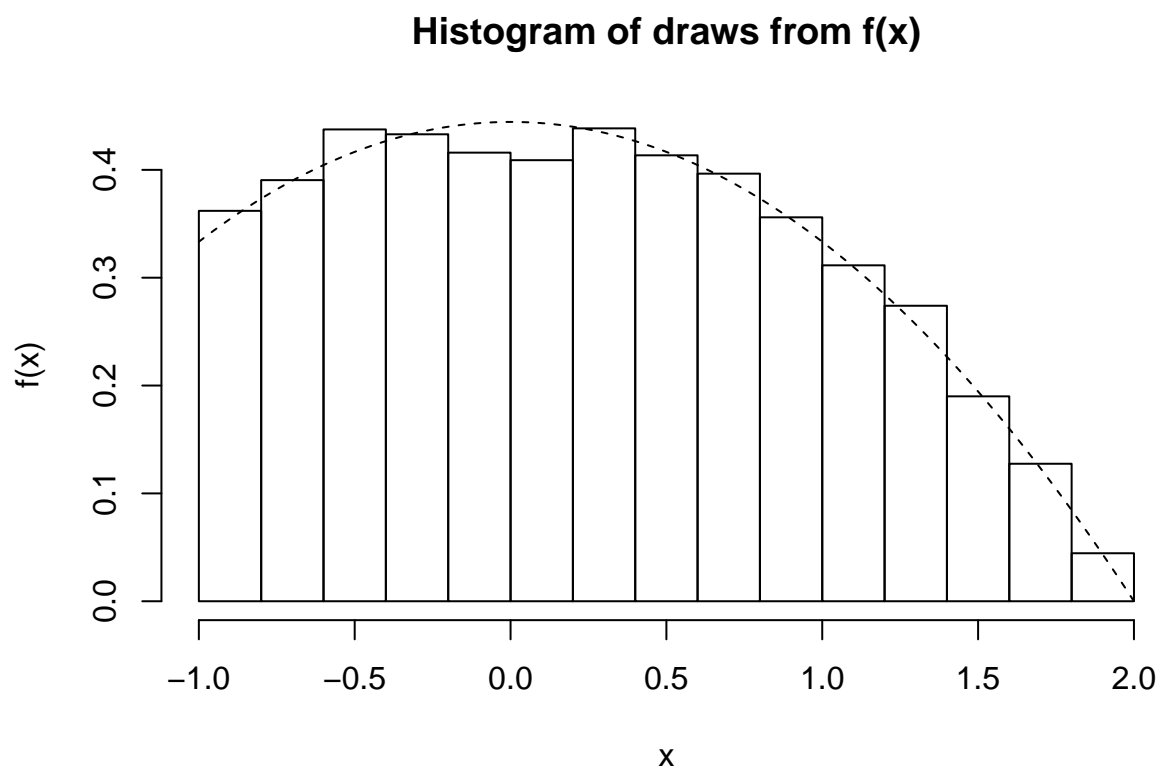
```
x.max = 0
f.max = 4/9
e <- function(x){
  return(ifelse((x < -1 | x > 2), Inf, f.max))
}
```

6.

```
n.samps <- 10000
n <- 0
samps <- numeric(n.samps)
while (n < n.samps){
  y <- runif(1,-1,2)
  u <- runif(1)
  if (u < f(y)/e(y)){
    n <- n + 1
    samps[n] <- y
  }
}
f.draws <- samps
```

7.

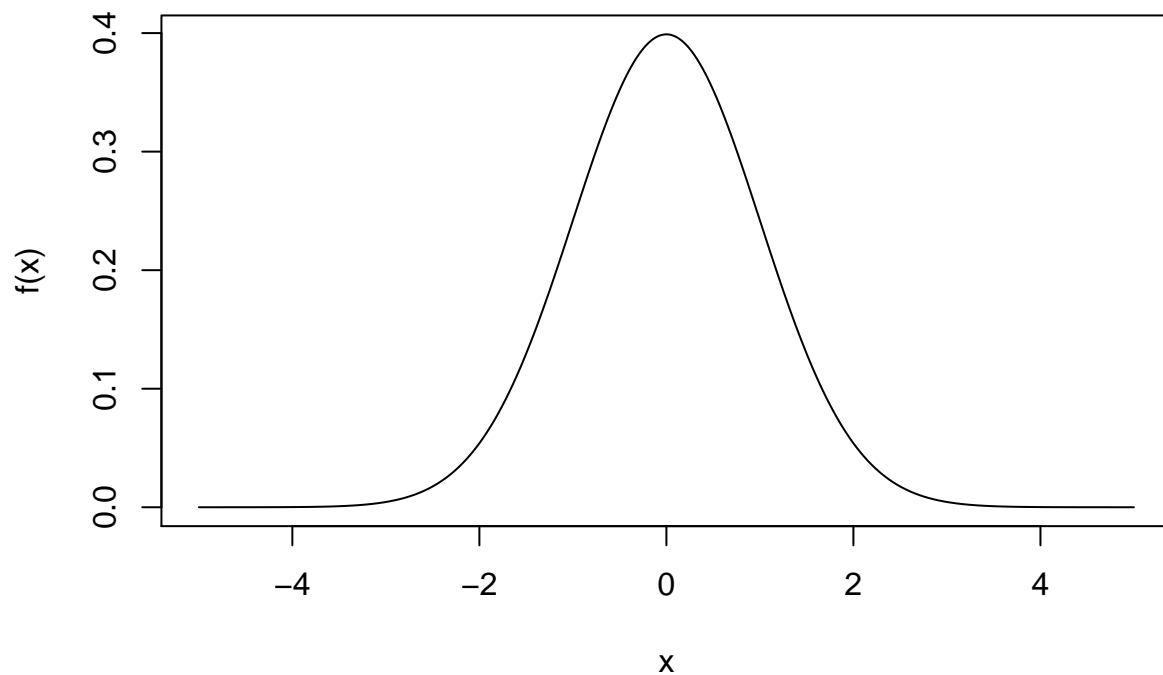
```
hist(f.draws, prob = T, ylab = "f(x)", xlab = "x", main = "Histogram of draws from f(x)")
lines(x, f(x), lty = 2)
```



#####Problem 3: Reject-Accept Method Continued 8.

```
f <- function(x){  
  return(1/sqrt(2*pi)*exp(-x^2/2))  
}  
x <- seq(-5, 5, 0.01)  
plot(x, f(x), type = 'l', main = "Standard Normal Distribution")
```

## Standard Normal Distribution

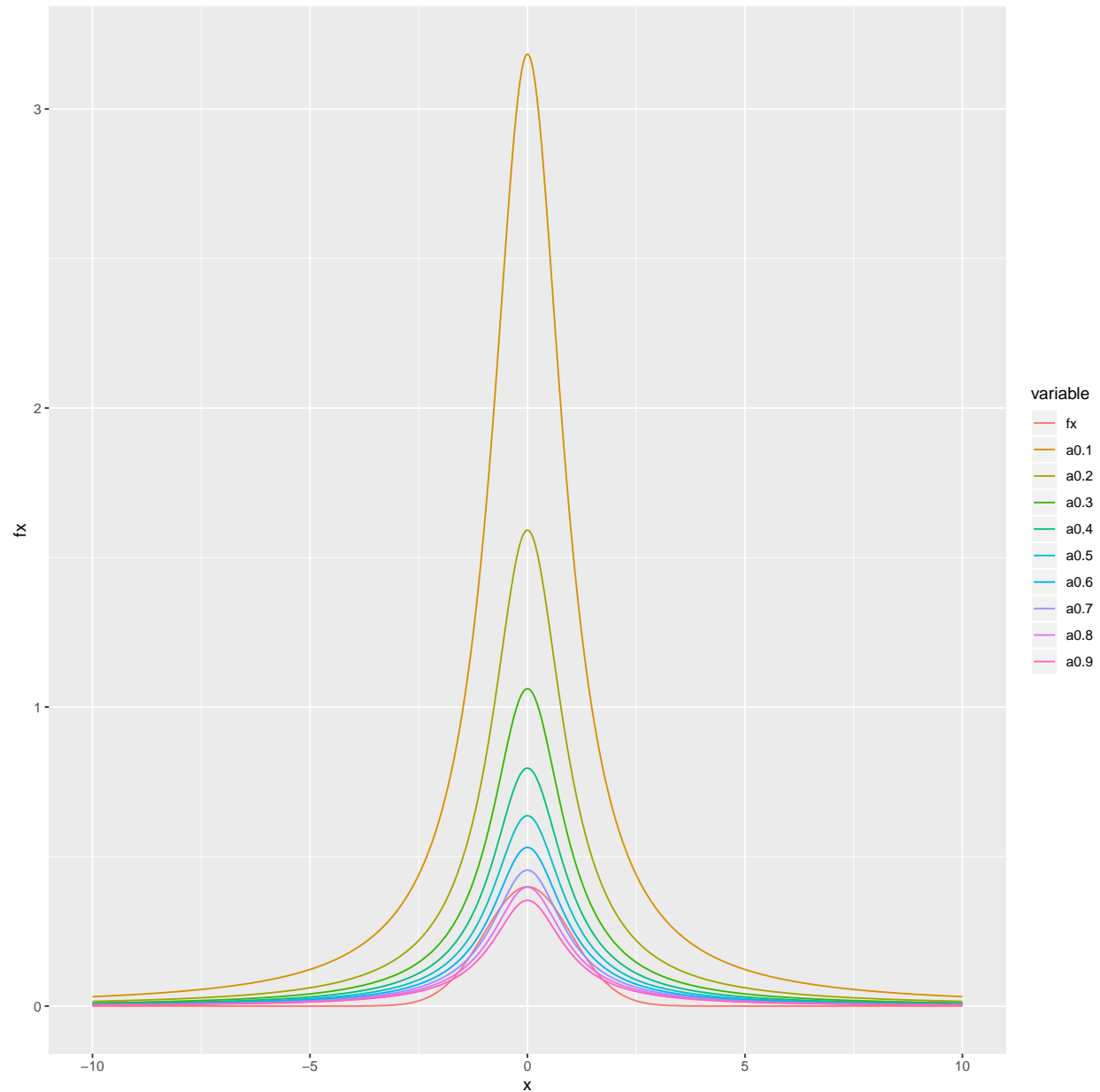


9.

```
e <- function(x,a){  
  return((1/pi*1/(1+x^2))/a)  
}
```

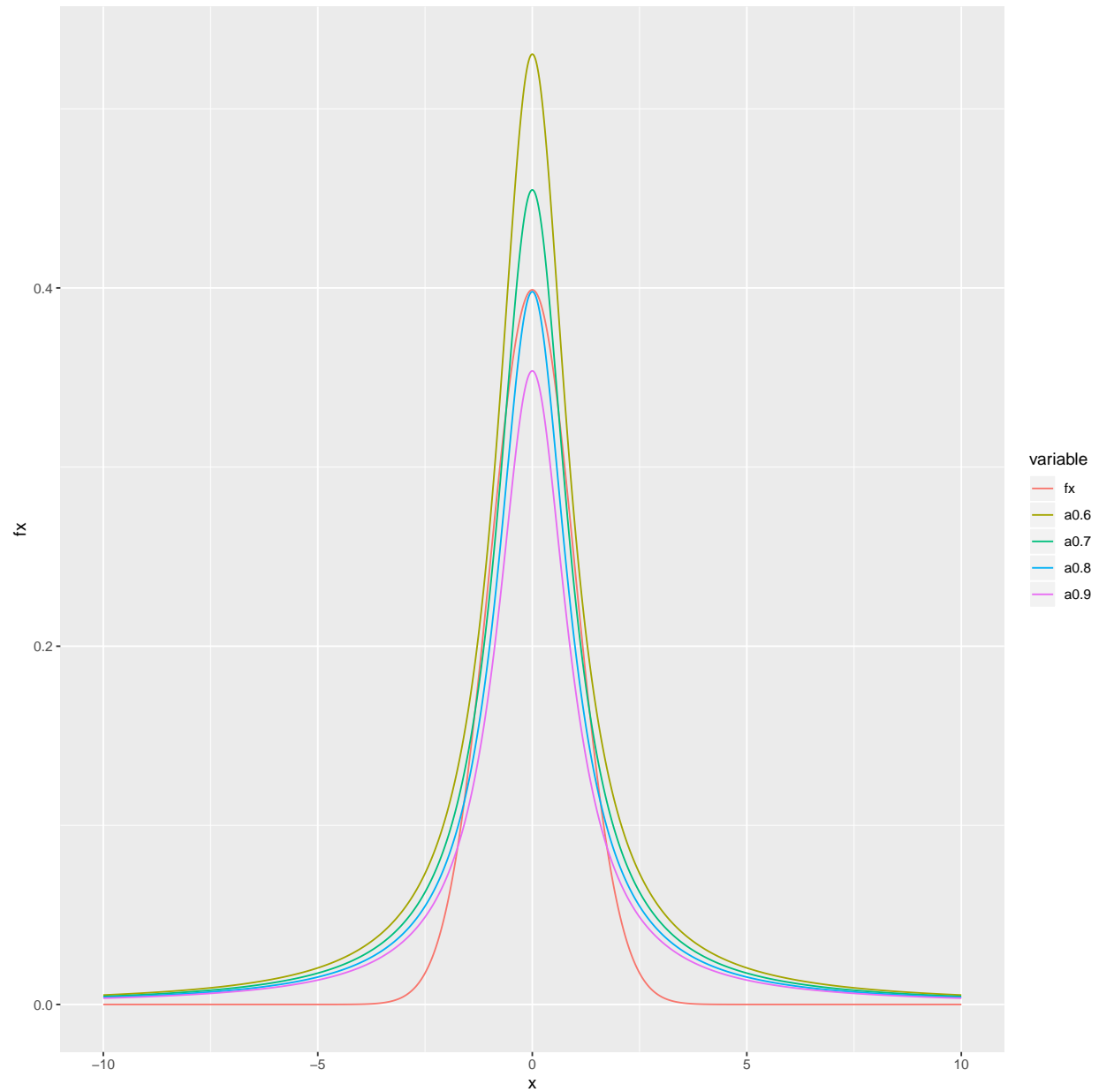
10.

```
library(ggplot2)  
suppressWarnings(library(reshape2))  
# We first choose alpha value from 0.1 to 0.9  
x <- seq(-10,10,0.01)  
fx <- f(x)  
alpha <- seq(0.1,0.9,0.1)  
m <- list()  
for (i in 1:length(alpha)){  
  nam <- paste("a", alpha[i], sep = "")  
  m[[i]] <- e(x,alpha[i])  
  names(m)[i] <- nam  
}  
m <- do.call(cbind,m)  
m <- as.data.frame(cbind(x,fx,m))  
newm <- melt(m, id.vars = "x", value.name = "fx")  
ggplot(newm, aes(x, fx, col = variable)) +  
  geom_line()
```



```
# Alpha seems to be good from 0.6 to 0.9

m <- list()
alpha <- seq(0.6,0.9,0.1)
for (i in 1:length(alpha)){
  nam <- paste("a", alpha[i], sep = "")
  m[[i]] <- e(x,alpha[i])
  names(m)[i] <- nam
}
m <- do.call(cbind,m)
m <- as.data.frame(cbind(x,fx,m))
newm <- melt(m, id.vars = "x", value.name = "fx")
ggplot(newm, aes(x, fx, col = variable)) +
  geom_line()
```



*# Alpha is good at 0.8*

11.

```
normal.sim <- function(n){
  n.samps <- n
  n <- 0
  samps <- numeric(n.samps)
  a = 0.8
  while (n < n.samps){
    y <- cauchy.sim(1)
    u <- runif(1)
    if (u < f(y)/e(y,a)){
      n <- n + 1
      samps[n] <- y
    }
  }
}
```

```

    }
  }
  return(samps)
}
normal.sim(10)

```

```

## [1] 0.73607864 0.91088131 -0.06323431 0.11270693 0.02508774 0.56185361
## [7] -1.73153476 -0.08200524 -1.23690743 1.61518216

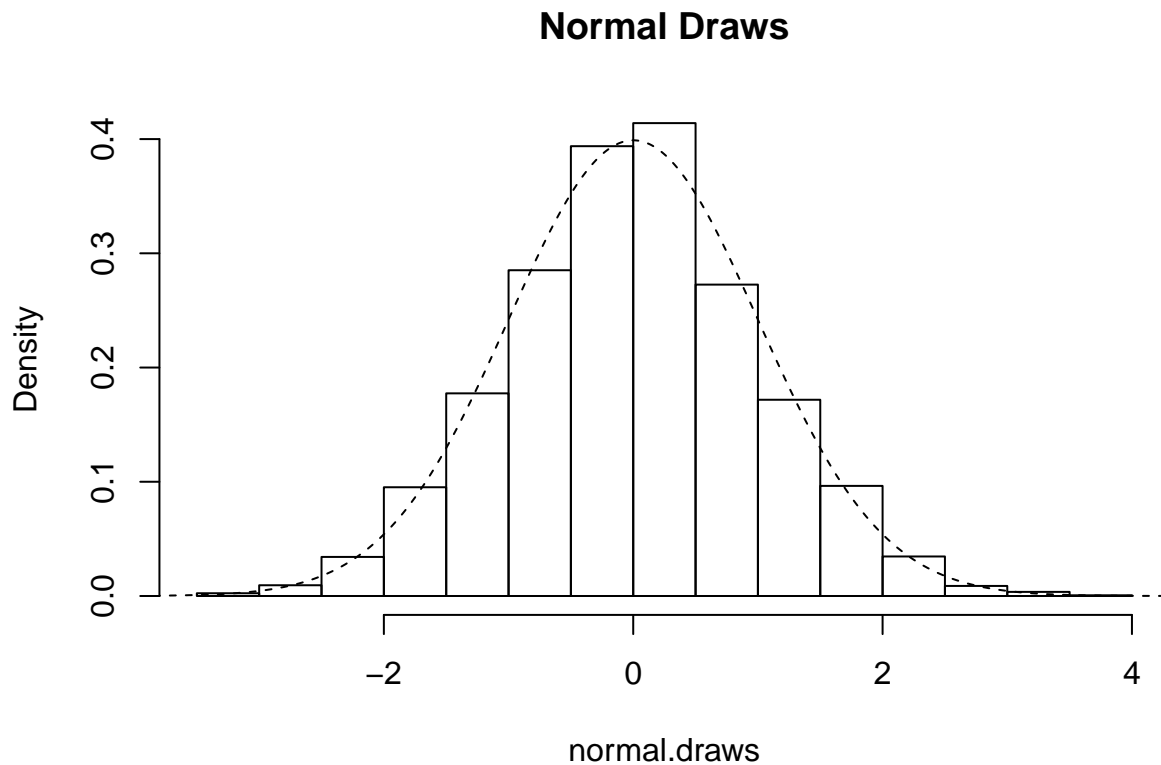
```

12.

```

normal.draws <- normal.sim(10000)
hist(normal.draws, prob = T, main = "Normal Draws")
x <- seq(-5, 5, 0.01)
lines(x, dnorm(x), lty = 2)

```



#####Part 3: Simulation with Built-in R Functions 13.

```

x <- 5
x.vals <- c(x)
while (x > 0){
  r <- runif(1, -2, 1)
  x <- x + r
  x.vals <- c(x.vals, x)
}
x.vals

```

```

## [1] 5.0000000 5.6414627 6.1594880 4.8686355 3.2703491 4.2308837 2.7291568

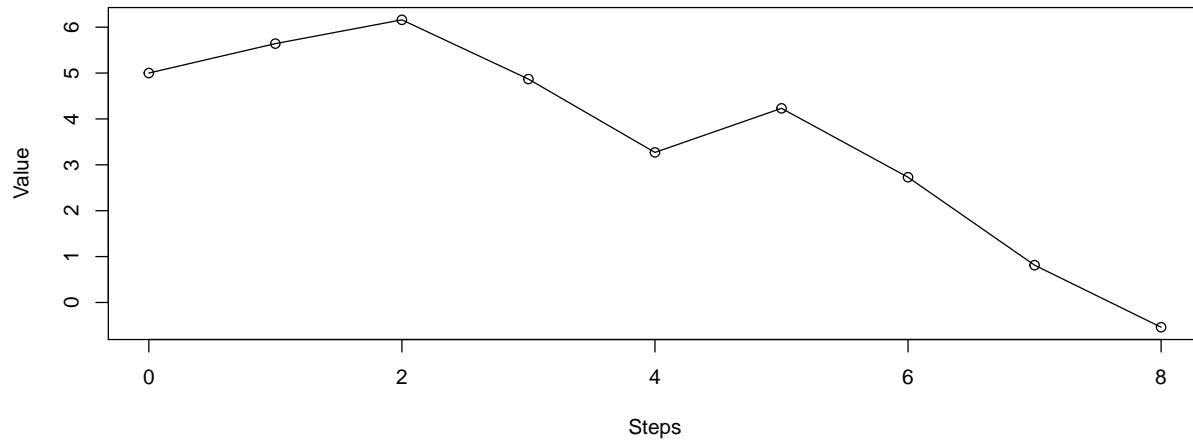
```



```
## [8] 0.8131343 -0.5392317
```

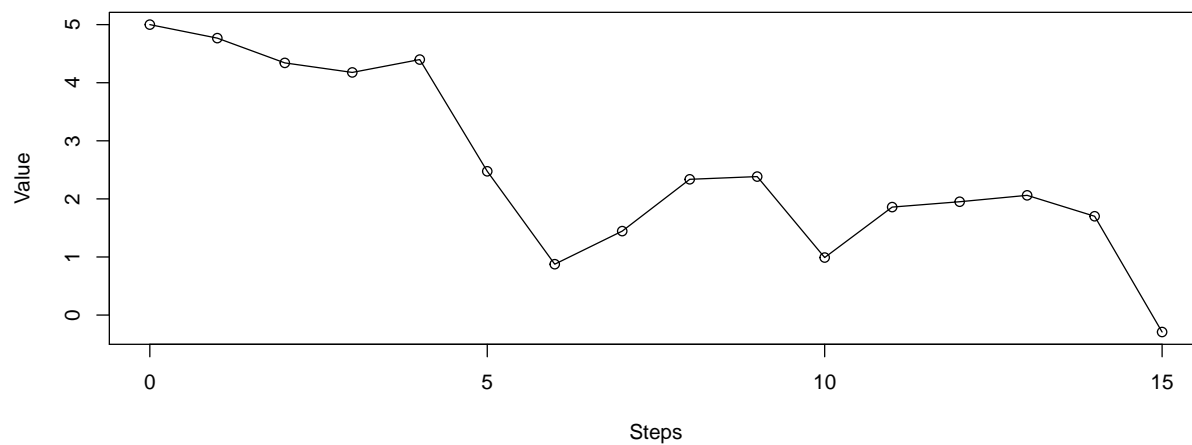
14.

```
plot(0:(length(x.vals) - 1), x.vals, type = 'o', xlab = "Steps", ylab = "Value")
```

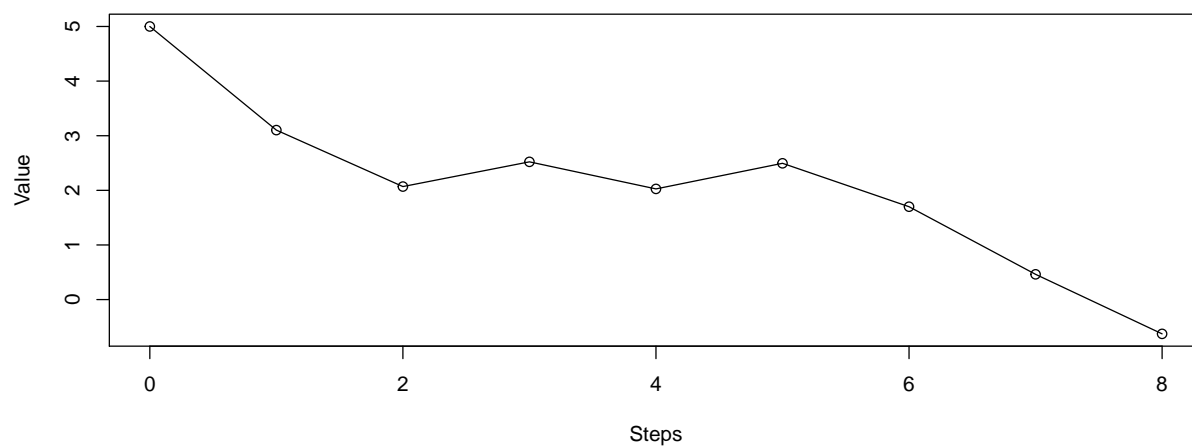


15.

```
random.walk <- function(x.start = 5, plot.walk = TRUE){  
  x.vals <- c(x.start)  
  num.steps <- 0  
  while (x.start > 0){  
    r <- runif(1, -2, 1)  
    x.start <- x.start + r  
    x.vals <- c(x.vals, x.start)  
    num.steps <- num.steps + 1  
  }  
  if (plot.walk == TRUE){  
    plot(0:num.steps, x.vals, type = 'o', xlab = "Steps", ylab = "Value")  
  }  
  output <- list(x.vals, num.steps)  
  names(output) <- c("x.vals", "num.steps")  
  
  return(output)  
}  
  
random.walk()
```



```
## $x.vals
## [1] 5.0000000 4.7677850 4.3413215 4.1772027 4.3985265 2.4749181
## [7] 0.8755790 1.4453621 2.3370404 2.3839827 0.9901354 1.8588850
## [13] 1.9510849 2.0612143 1.7010838 -0.2916823
##
## $num.steps
## [1] 15
random.walk()
```



```
## $x.vals
## [1] 5.0000000 3.1027987 2.0690005 2.5216511 2.0262637 2.4950284 1.7004206
## [8] 0.4623079 -0.6276088
##
## $num.steps
## [1] 8
random.walk(10, F)
```

```
## $x.vals
## [1] 10.0000000 8.0129886 7.5870304 6.5092611 5.2595840 4.4866026
## [7] 3.6251574 2.7411669 0.7556797 -0.2023972
##
## $num.steps
## [1] 9
```

```
random.walk(10, F)
```

```
## $x.vals
## [1] 10.0000000 10.6631103 9.6757474 9.3282607 8.6160893 7.9767263
## [7] 6.3293430 4.8584431 5.4272541 6.0685201 6.8793571 7.5588311
## [13] 7.9146708 6.5358636 6.5289751 6.0856862 4.4307243 3.7010518
## [19] 3.7430589 3.0336377 3.5115856 3.6739033 2.5657297 2.0619642
## [25] 1.1943557 0.2690885 -0.2227623
##
## $num.steps
## [1] 26
```

16.

```
# The expected number of iterations should be  $5/(1/2) + 1 = 11$ 
steps <- c()
for (i in 1:10000){
  random.walk(, F)
  steps[i] <- random.walk(, F)[[2]]
}
mean(steps)
```

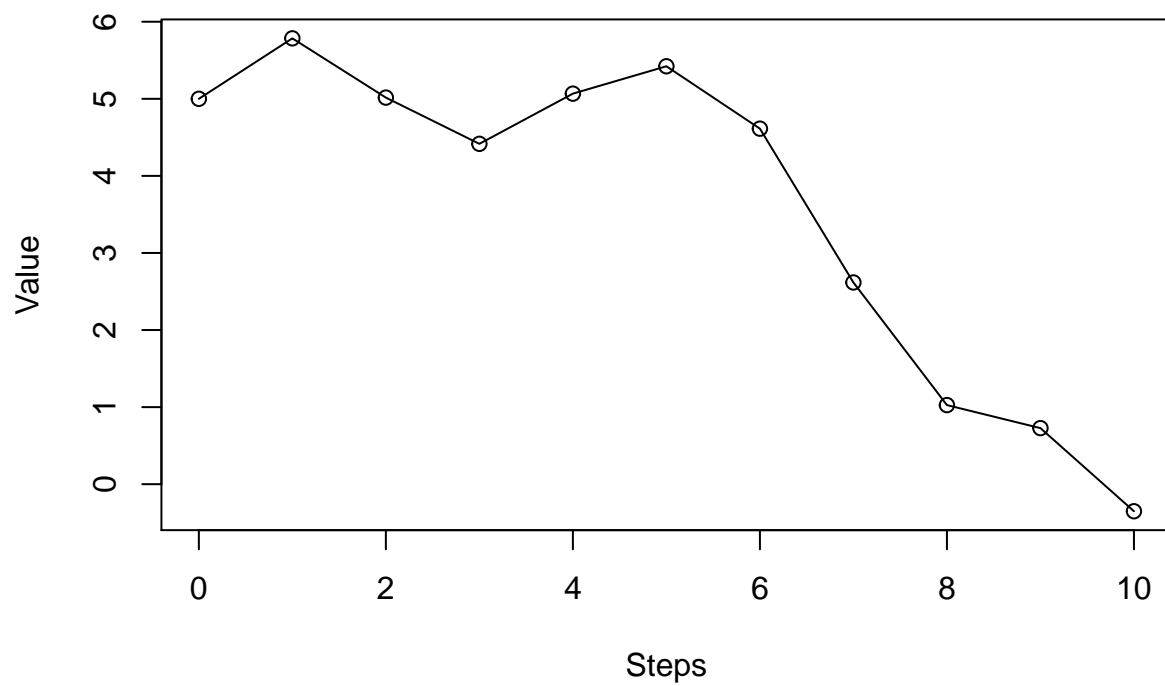
```
## [1] 11.297
```

17.

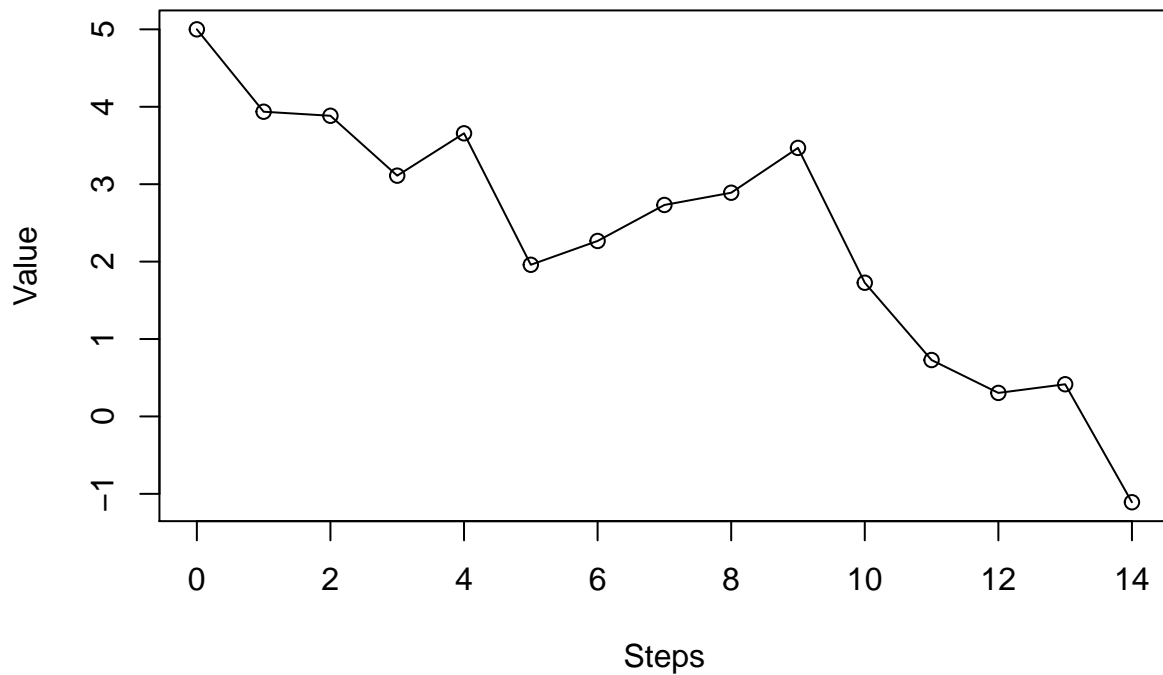
```
random.walk <- function(x.start = 5, plot.walk = TRUE, seed = NULL){
  if (is.null(seed) == FALSE){
    set.seed(seed)
  }
  x.vals <- c(x.start)
  num.steps <- 0
  while (x.start > 0){
    r <- runif(1, -2, 1)
    x.start <- x.start + r
    x.vals <- c(x.vals, x.start)
    num.steps <- num.steps + 1
  }
  if (plot.walk == TRUE){
    plot(0:num.steps, x.vals, type = 'o', xlab = "Steps", ylab = "Value")
  }
  output <- list(x.vals, num.steps)
  names(output) <- c("x.vals", "num.steps")

  return(output)
}

random.walk()
```



```
## $x.vals
## [1] 5.0000000 5.7844984 5.0157528 4.4157721 5.0668945 5.4223319
## [7] 4.6126752 2.6171137 1.0260696 0.7270525 -0.3511712
##
## $num.steps
## [1] 10
random.walk()
```



```
## $x.vals
## [1] 5.0000000 3.9360707 3.8836874 3.1099632 3.6570394 1.9589386
## [7] 2.2666793 2.7316262 2.8897985 3.4687411 1.7270853 0.7285992
## [13] 0.3037413 0.4157197 -1.1082456
##
## $num.steps
## [1] 14
```

```
random.walk(, F, 33)
```

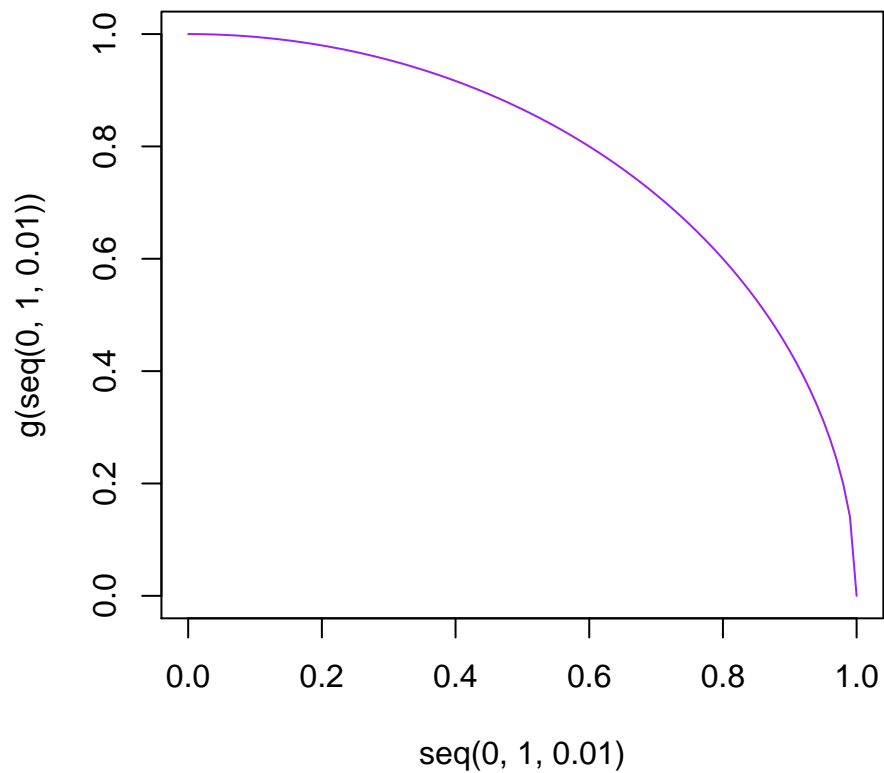
```
## $x.vals
## [1] 5.0000000 4.3378214 3.5217724 2.9729590 3.7295869 4.2612312
## [7] 3.8132800 3.1246550 2.1542497 0.2008006 -1.4452259
##
## $num.steps
## [1] 10
```

```
random.walk(, F, 33)
```

```
## $x.vals
## [1] 5.0000000 4.3378214 3.5217724 2.9729590 3.7295869 4.2612312
## [7] 3.8132800 3.1246550 2.1542497 0.2008006 -1.4452259
##
## $num.steps
## [1] 10
```

```
#####Part 4: Monte Carlo Integration 18.
```

```
g <- function(x){
  return(sqrt(1-x^2))
}
plot(seq(0, 1, .01), g(seq(0, 1, .01)), type = 'l', col = 'purple')
```



19.  $A = \frac{1}{4}\pi r^2 = \frac{1}{4}\pi * 1^2 = \frac{\pi}{4}$

20.

```
# Choose uniform distribution when perform this simulation
g.over.p <- function(x){
  return(4*sqrt(1-x^2))
}
mean(g.over.p(runif(10000000)))
```

```
## [1] 3.141666
```

```
pi
```

```
## [1] 3.141593
```

```
pi - mean(g.over.p(runif(10000000)))
```

```
## [1] -0.0002403507
```

```
1/1000 - abs(pi - mean(g.over.p(runif(1000000))))
```

```
## [1] 0.0005720405
```

```
# So we estimated the value of pi within a 1/1000 of the true value
```