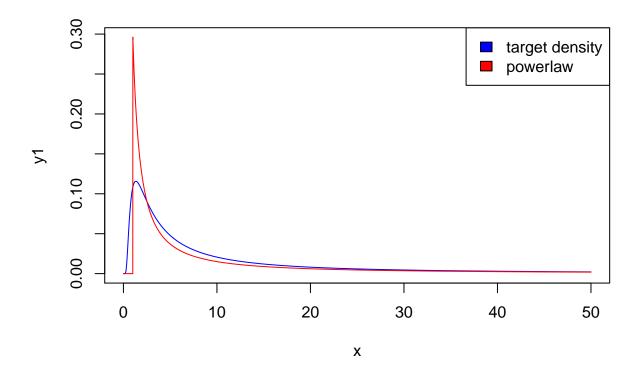
Computer Lab 3

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Question 1

```
distrib1 <- function(x,c){</pre>
  if(x>0){
    return(c*(sqrt(2*pi)^(-1))*(exp(-(c^2)/(2*x))) * (x^(-3/2)))
  else return(0)
powerlaw <- function(x,a,t){</pre>
  if(x>t){
    return(((a-1)/t) * ((x/t)^{-(-a)}))
  else return(0)
}
x \leftarrow seq(0,50, by=0.01)
c <- 2
y1 <- sapply(x, distrib1, c=c)</pre>
y2 \leftarrow sapply(x, powerlaw, a = 1.3, t = 1)
ymax \leftarrow max(y1,y2)
b \leftarrow \min(c(y1,y2))
e \leftarrow \max(c(y1,y2))
ax <- seq(b,e,by=(e-b)/200)
plot(x,y1, type="l", col = "blue", ylim = c(0,ymax))
lines(x,y2, col = "red")
legend("topright", c("target density", "powerlaw"), fill=c("blue", "red"))
```



```
# hist(y1, breaks = ax,

# col = "red",

# main = "Comparison of rnorm() with our rNorm()",

# xlab = "values",

# xlim = range(y1,y2))

# hist(y2, breaks = ax, col = "blue", xlim =range(y1,y2), add = TRUE)
```

power-law distribution cannot be used just by itself because it does nt support range from 0 to T_{min} . parameters from 0 to 1 can be taken from another distribution

lets assume that $T_{min} = 1$

probability that number will be in 0-1 region is:

$$\int_0^1 c\sqrt{2\pi}^{-1} e^{-\frac{c^2}{2x}} x^{\frac{-3}{2}} = \frac{\Gamma\left(\frac{1}{2}, \frac{c^2}{2}\right)}{\sqrt{\pi}}$$

lets assume that other distribution will be uniform(0,1)

```
majDensity <- function(x){
   if(x>1){
      return(powerlaw(x, 1.6, 1))
   }
   else {return(dunif(x,0,1))}
}
# y3 <- sapply(x, target)
# plot(x,y3, type = "l")</pre>
```

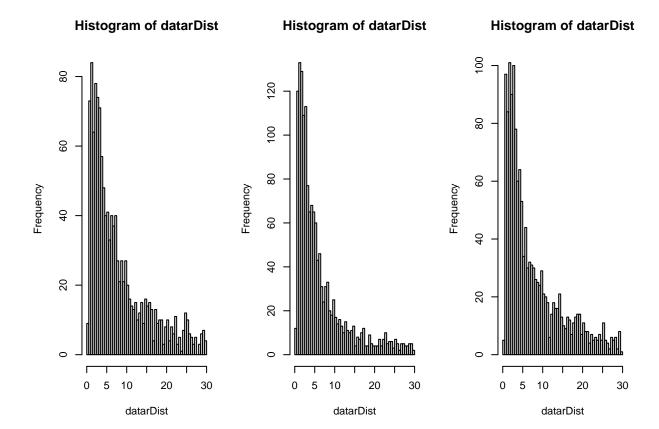
```
# #lines(x,dnorm(x,3,1.2), col ="red")
# lines(x,y1, col ="blue")
# #lines(x,y1, col = "red")
```

2

```
library(poweRlaw)
```

Warning: package 'poweRlaw' was built under R version 4.0.3

```
randomnumber <- function(prob){</pre>
  numb <- runif(1)</pre>
  if(numb<= prob){</pre>
    return(runif(1,0,1))
  else{
    return(rplcon(1,1,1.5))
}
CompleteDist <- function(c, rej){</pre>
  z <- TRUE
  res <- 0
  while (z == TRUE) {
    y <- randomnumber(0.5)
    u <- runif(1)
    if(u <= distrib1(y, 2) / (c*majDensity(y))){</pre>
      res <- y
      z <- FALSE
    }
    if(rej){
    rejected <<- rejected + 1
  }
  return(res)
}
rDist <- function(n,c ,rej = FALSE){</pre>
return(replicate(n, CompleteDist(c, rej)))
}
```



Question 2

1.

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

- μ location parameter
- b > 0 scale parameter

inverse CDF of DE:

Source - https://en.wikipedia.org/wiki/Laplace_distribution

$$F^{-1}(p) = \mu - bsgn(p - 0.5)ln(1 - 2|p - 0.5|)$$

```
where b = \frac{1}{\alpha}

rLaplace <- function(n, mean = 0, alpha = 1){

b <- 1/alpha

u <- runif(n)

res <- mean - (b*sign(u-0.5) * log(1-(2*abs(u-0.5))))

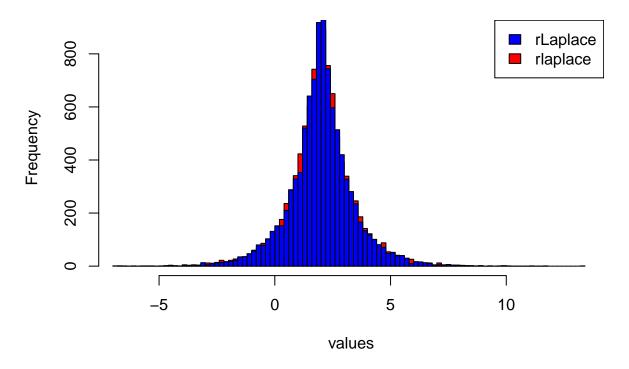
return(res)
```

meaning:

- 1. calculate b.
- 2. take n random variables from uniform distribution [0,1]

3. calculate random numbers from inverse CDF of laplace distribution where \mathbf{x} is a random variable from uniform distribution

Comparison of rlaplace function from rmutil with our rLaplace



2.

```
DE <- function(x, mean = 0,alpha = 1){
    return((0.5*alpha)*exp((-alpha)*abs(x-mean)))
}

genNorm <- function(c, rej){
    z <- TRUE
    res <- 0
    while (z == TRUE) {
        y <- rLaplace(1)
        u <- runif(1)
        if(u <= pnorm(y) / (c*DE(y))){
            res <- y
            z <- FALSE
        }
        if(rej){
            rejected <<- rejected + 1
        }
    }
    return(res)
}</pre>
```

```
rNorm <- function(n,c,rej = FALSE){
  return(replicate(n, genNorm(c, rej)))
}</pre>
```

algorithm:

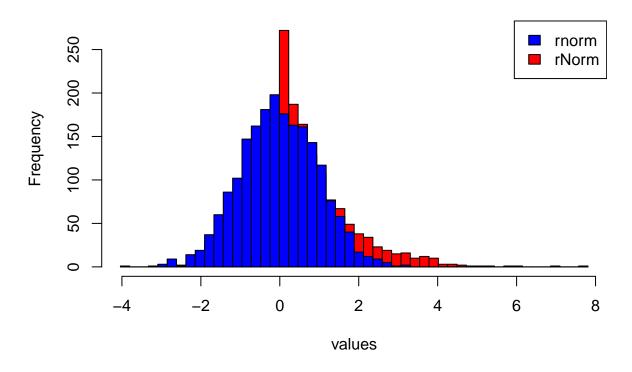
- 1. write Laplace probability function
- 2. assign 0 to result value res
- 3. generate random number y from rLaplace function
- 4. generate random number u from uniform distribution
- 5. check if u is less or equal to probability of y in normal distribution / c * probability of y in laplace distribution
 - a) if yes, return y
 - b) repeat steps from 3

c should be > 0; $sup_x(f(x)/g(x)) \le c$

Estimated optimal c - 1.3154892

 $source-http://www.columbia.edu/\sim ks 20/4703-Sigman/4703-07-Notes-ARM.pdf$

Comparison of rnorm() with our rNorm()



	mean	variance
rNorm()	0.3648458	1.3887557
rnorm()	-0.0158433	0.9534873

rejection rate: 1.249, expected rejection rate = c = 1.3154892, difference - -0.0664892