

# 904971914\_stats102c\_hw2

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## Question 1

(a)

```
set.seed(904971914)
qla<-function(){
  #This function generates one value from the distribution X in problem 1
  #it has no arguments
  #it returns this value

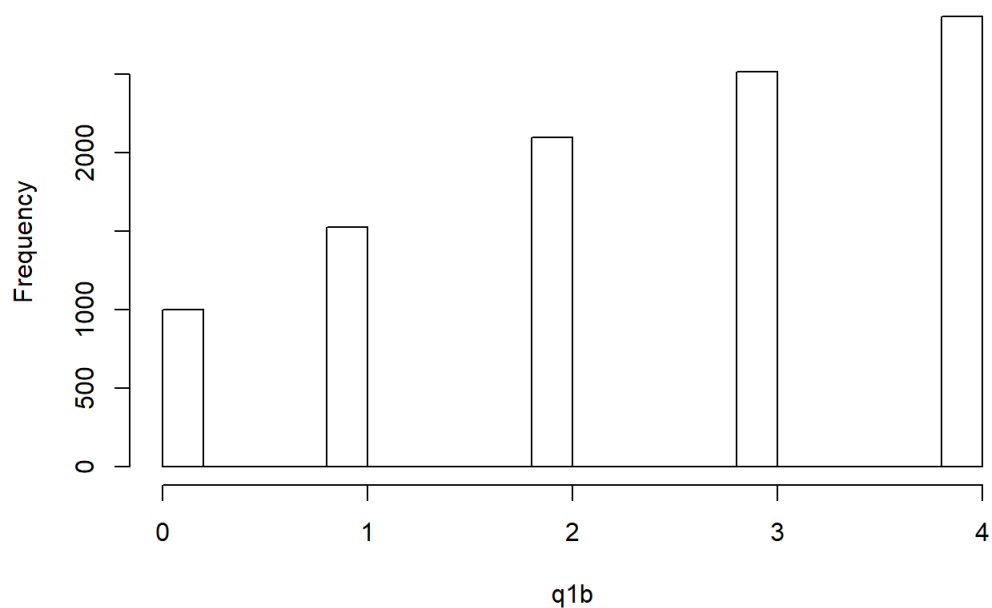
  u<-runif(1)

  if(u<=0.1){
    return(0);
  }
  else if(u<=0.25){
    return(1);
  }
  else if(u<=0.45){
    return(2);
  }
  else if(u<=0.7){
    return(3);
  }
  else{
    return(4);
  }
}
```

(b)

```
qlb<-c()
for(i in 1:10000){
  qlb<-c(qlb,qla())
}
hist(qlb,freq=T)
```

Histogram of q1b



(c)

```
sum(q1b==0)/10000
```

```
## [1] 0.0999
```

```
sum(q1b==1)/10000
```

```
## [1] 0.1524
```

```
sum(q1b==2)/10000
```

```
## [1] 0.2098
```

```
sum(q1b==3)/10000
```

```
## [1] 0.2515
```

```
sum(q1b==4)/10000
```

```
## [1] 0.2864
```

The sample relative frequencies are very close to the theoretical probabilities.

## Question 2

(a)

Pseudocode: 1 Generate  $u$  from  $\text{uniform}(0,1)$

2 while loop:

if  $u \geq F(x)$

increment  $x$  by 1

generate  $p(x)$  from poisson

increment  $F(X)$  by adding  $p(x)$

3. Stop and return  $x$

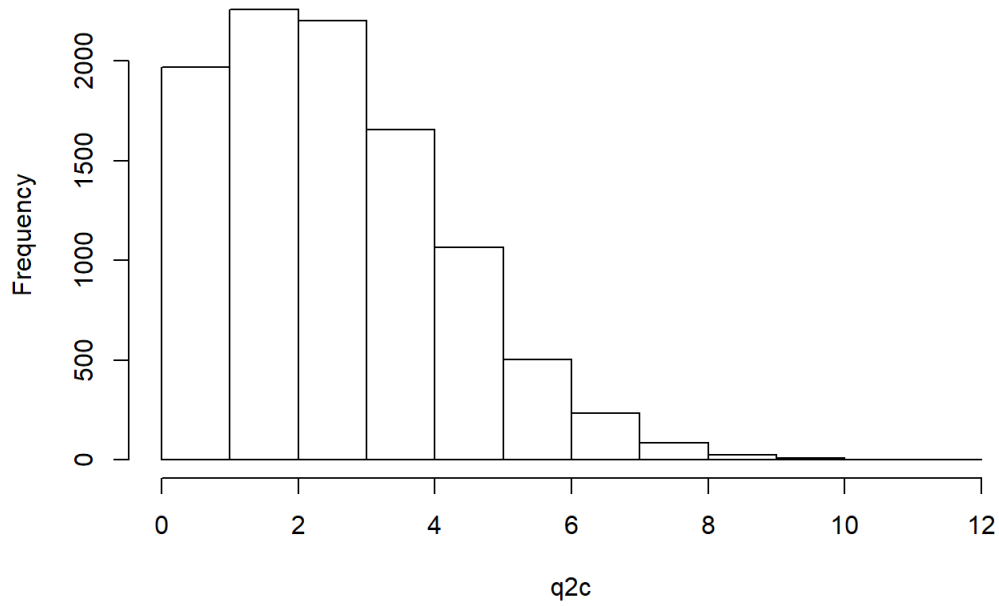
(b)

```
q2b<-function(lambda) {  
  #This function generates a value from a poisson distribution with lambda  
  #the argument lambda gives the value of the parameter  
  #the function returns the value generated  
  u<-runif(1)  
  x<-0  
  cdf<-((lambda^x)*exp(-lambda))/factorial(x)  
  while(u >=cdf) {  
    x<-x+1  
    px<-((lambda^x)*exp(-lambda))/factorial(x)  
    cdf<-cdf+px  
  }  
  return (x)  
}
```

(c)

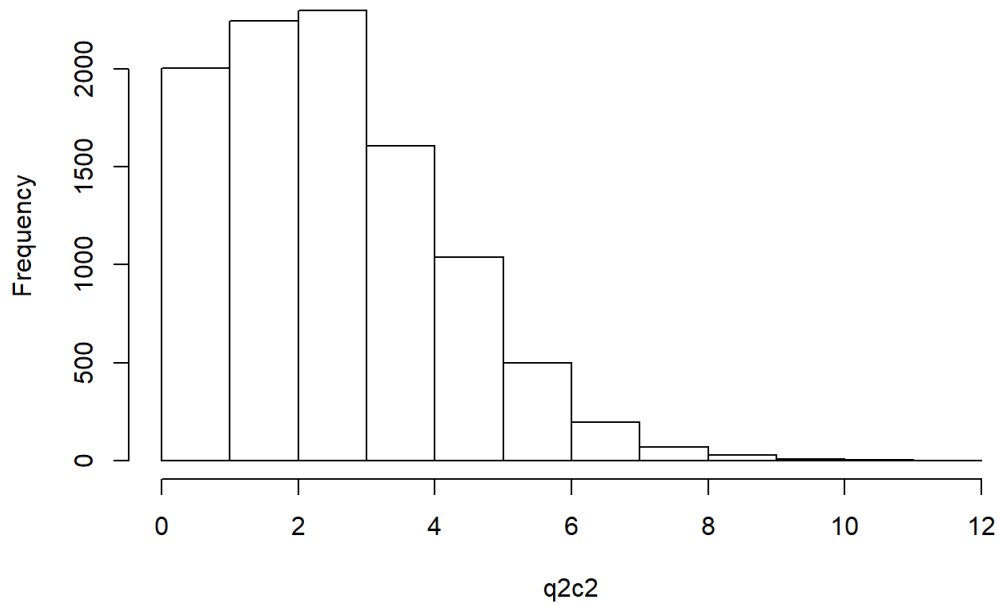
```
q2c<-c()  
for(i in 1:10000) {  
  q2c<-c(q2c,q2b(3))  
}  
  
q2c2<-rpois(n=10000,lambda=3)  
hist(q2c)
```

**Histogram of q2c**



```
hist(q2c2)
```

**Histogram of q2c2**



The results from the two

functions are very similar in terms of distribution.

## Question 3

(a)

$$\begin{aligned}
 Y &= \left(\frac{X}{\alpha}\right)^\beta \\
 P(Y \leq y) &= P\left(\left(\frac{X}{\alpha}\right)^\beta \leq y\right) \\
 &= P(X \leq \alpha y^{\frac{1}{\beta}}) \\
 &= F_X(\alpha y^{\frac{1}{\beta}}) \\
 &= 1 - e^{-\frac{\alpha y^{\frac{1}{\beta}}}{\alpha}} \\
 &= 1 - e^{-y} \sim \exp(1)
 \end{aligned}$$

So it follows an exponential distribution with lambda = 1.

(b)

```

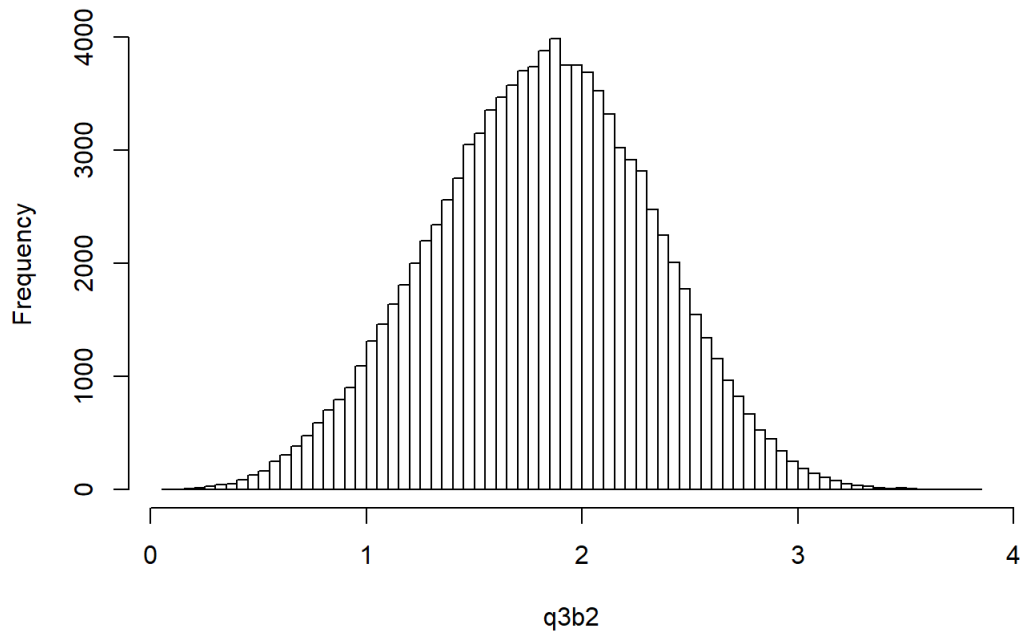
q3b<-function (n) {
  #This function generates n values from distribution of X in problem 3.
  #It takes one argument n which specifies the number of values to e generated.
  #It returns the values
  raw <-rexp(n)
  y <-(raw^(1/4))*2
  y

}

q3b2 <-q3b(100000)
hist(q3b2,breaks=100)

```

**Histogram of q3b2**



## Question 4

Assume that  $g(x)$  is the envelope function and  $n$  is total number of points.

$$N = \text{Total number of points in}(x, x + \Delta x) = n \times g(x) \times \Delta x$$

$$p = \text{Acceptance probability in}(x, x + \Delta x) = \frac{f(x)}{Mg(x)}$$

$$\text{Number of points remained in}(x, x + \Delta x) = N \times p = \frac{n}{M} f(x) \Delta x$$

$$\text{total points remained} = \sum \frac{n}{M} f(x) \Delta x = \frac{n}{M} \sum_{\text{bin number}} f(x) \Delta x = \frac{n}{M}$$

$$P(x \in (x, x + \Delta x)) = \frac{\frac{n}{M} f(x) \Delta x}{\frac{n}{M}} = f(x) \Delta x$$

$$f(x) = \frac{P(x \in (x, x + \Delta x))}{\Delta x}$$

## Question 5

(a) Take  $g(x)$  as  $\text{Unif}(0, 1)$ ,  $f(x)$  as  $\text{Beta}(3, 2)$  [

$$x \in [0, 1]$$

$$g(x) = 1$$

$$\frac{x^{\alpha-1}(1-x)^{\beta-1}}{\frac{2! * 1!}{4!}}$$

$$f(x) =$$

$$f(x) = 12 \times x^2 \times (1-x)$$

$$f'(x) = 0$$

$$x = 0 \text{ or } x = \frac{2}{3}$$

$$M = \max(g(x)) = \frac{f(\frac{2}{3})}{1} = \frac{16}{9}$$

]

(b)

```
my_beta<-function(x,a,b) {
  #this function gives the pdf of a beta distribution
  #it takes in x as the subject, a and b as parameters
  #it returns the pdf at x
  y <-(x^(a-1)) * ((1-x)^(b-1))
  z<-(factorial(a-1)) * (factorial(b-1)) / factorial(a+b-1)
  y/z
}

q5b<-function(n) {
  #This function generates a distribution from n sample points using the acceptance-rejection method
  #it takes in one argument n as the number of sample points
  #it returns a vector of accepted x
  M<-16/9
  result<-c()

  for(i in 1:n) {

    x<-runif(1)
    U<-runif(1)
    if (U < my_beta(x,3,2)/M) {
      result <-c(result,x)
    }

  }

  result
}

q5b2<-q5b(100000)
hist(q5b2)
```



(c)

```
length(q5b2) / 100000
```

```
## [1] 0.56225
```

```
1 / (16/9)
```

```
## [1] 0.5625
```

The acceptance rate is very close to 1/M.

## Question 6

(a)

[

$$\begin{aligned}
 f(x) &= \frac{1}{2} e^{-|x|} \\
 F(x) &= \int_{-\infty}^x f(x) dx \\
 &= \int_{-\infty}^x \frac{1}{2} e^{-|x|} dx \\
 \text{if } (x < 0), f(x) &= \frac{1}{2} e^x \\
 \text{if } (x \geq 0), f(x) &= \frac{1}{2} e^{-x} \\
 \text{if } (x < 0), F(x) &= \frac{1}{2} e^x \\
 \text{if } (x \geq 0), F(x) &= 1 - \frac{1}{2} e^{-x} \\
 \text{if } (x < 0), F^{-1}(u) &= \ln(2u) \\
 \text{if } (x \geq 0), F^{-1}(u) &= -\ln(2(1-u)) = -\ln(2u), \text{ as } u \sim \text{Unif}(0, 1), 1-u \sim \text{Unif}(0, 1)
 \end{aligned}$$

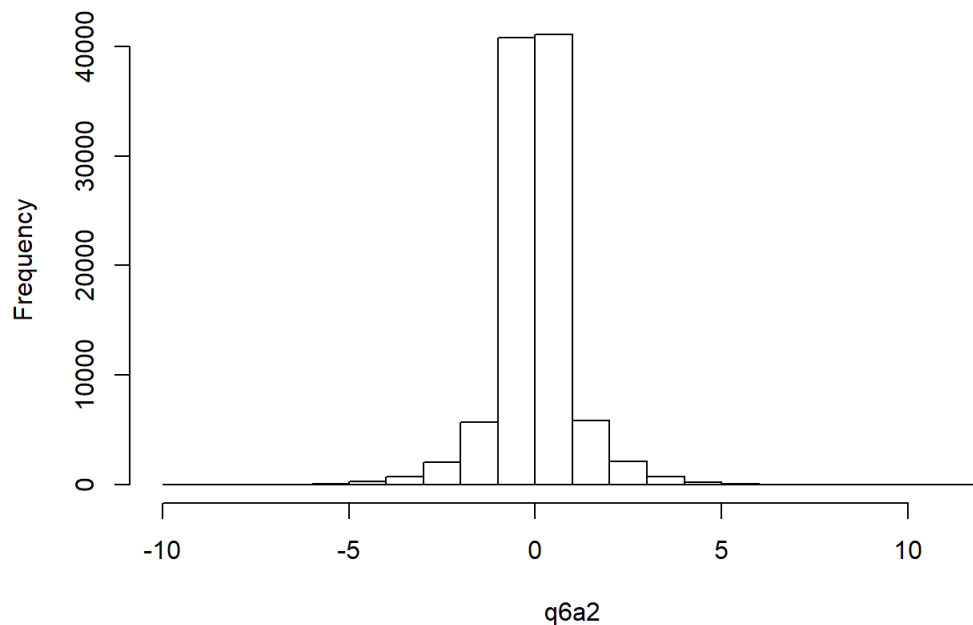
]

```
q6a<-function(n){
  #This function generates n data points from the laplace distribution
  #it takes in one argument n as the number of samples to be generated
  #it returns a vector containing all the samples.
  u1 <-runif(n/2)
  result <- -log(2*u1)
  u2 <-runif(n/2)
  result<-c(result,log(2*u2))
  result
}
q6a2<-q6a(100000)
system.time(q6a(100000))
```

```
##      user  system elapsed
##    0.02    0.00    0.01
```

```
hist(q6a2)
```

**Histogram of q6a2**



(b) Take  $g(x)$  as Norm(0,3).

[

$$\begin{aligned} \frac{f(x)}{\max(g(x))} &= \frac{\frac{1}{\sqrt{3} * \sqrt{2\pi}} * e^{-\frac{x^2}{0.5 * 3}}}{\frac{1}{2} e^{-|x|}} \\ \text{if}(x > 0), (g(x))' &= \frac{\frac{x}{(3-1)} * e^{-x}}{e^{-\frac{x^2}{6}}}, \text{ take } x = 3 \\ \text{if}(x \leq 0), (g(x))' &= \frac{\frac{x}{(3+1)} * e^{-x}}{e^{-\frac{x^2}{6}}}, \text{ take } x = -3 \\ M = \max(g(x)) &= \max(g(3)) = 0.4844 \end{aligned}$$

]

```

q6b<-function(n){
  #This function generates n data points from the laplace distribution using acceptance-rejection method.
  #it takes in one argument n as the number of samples to be generated
  #it returns a vector containing all the samples accepted.
  M<-0.4844
  result<-c()

  for(i in 1:n){
    x<-rnorm(1,mean=0,sd=sqrt(3))
    f<-0.5*exp(-abs(x))
    g<-dnorm(x,mean=0,sd=sqrt(3))
    U<-runif(1)
    if (U < f/(g*M)){
      result <-c(result,x)
    }

  }
  result
}

q6b2<-q6b(100000)
system.time(q6b(100000))

```

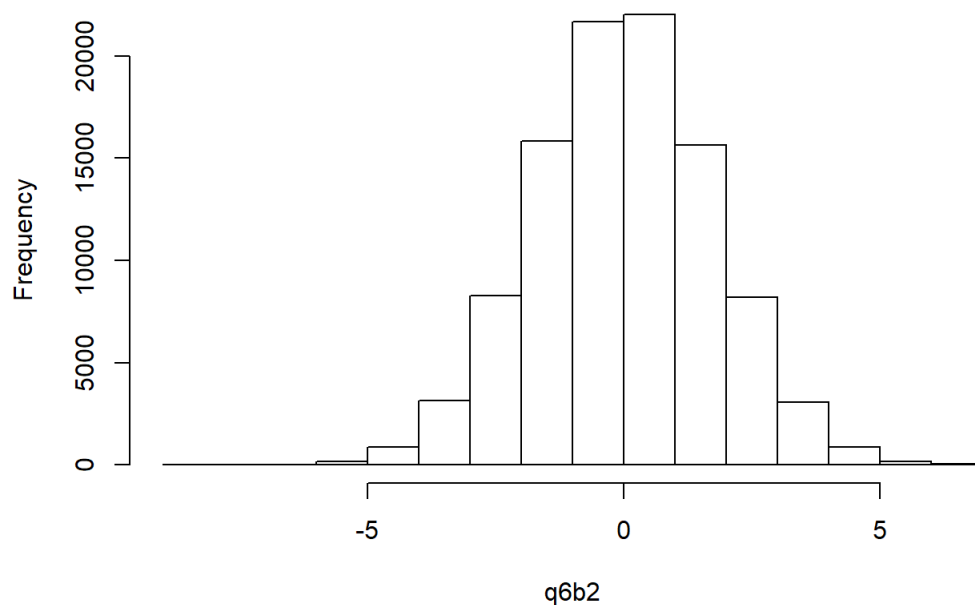
```

##      user  system elapsed
##    11.26    0.06    11.60

```

```
hist(q6b2)
```

**Histogram of q6b2**



(c) For inverse CDF, it is computationally efficient and thus takes less time. However, sometimes an explicit inverse cdf may not be available.

For acceptance-rejection, it does not require an explicit inverse cdf. However, it is sometimes hard to find an appropriate  $g(x)$  and  $M$ . Also, it has a higher computational cost than inverse CDF and thus takes more time.

Processing math: 100%