

Computational statistics

Week 1

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1 Exercise 1.22

1.1 Suppose that $X \sim f(x|\theta)$, with prior distribution $\pi(\theta)$, an interest is in the estimation of the parameter $h(\theta)$.

a) Using the loss function $L(\delta, h(\theta))$, show that the estimator that minimizes the Bayes risk $\int \int L(\delta, h(\theta)) f(x|\theta) \pi(\theta) dx d\theta$, is given by the estimator θ that minimizes (for each x)

$$\int L(\delta, h(\theta)) \pi(\theta|x) d\theta.$$

Solution:

The Bayes risk can be written :

$$R_\pi = \int_{\Theta} \int_{\chi} L(\delta, h(\theta)) f(x|\theta) \pi(\theta) dx d\theta$$

with $x \in \chi$ and $\theta \in \Theta$.

Suppose χ and Θ are σ -finite measure spaces. By definition, $L(\delta, h(\theta)) \geq 0$ so $L(\delta, h(\theta)) f(x|\theta) \pi(\theta)$ is measurable and non-negative. In this context, we can use the Tonelli's theorem to invert the ordering of the two integrals.

$$R_\pi = \int_{\chi} \int_{\Theta} L(\delta, h(\theta)) f(x|\theta) \pi(\theta) d\theta dx$$

The Bayes theorem (see slide 16 or page 12 of the book) states that:

$$f(x|\theta)\pi(\theta) = Z(x)\pi(\theta|x)$$

with $Z(x) = \int f(x|\theta)\pi(\theta)d\theta$ is the marginal density of X (or Bayesian evidence).

Hence,

$$\begin{aligned} R_\pi &= \int_{\mathcal{X}} \int_{\Theta} L(\delta, h(\theta)) Z(x) \pi(\theta|x) d\theta dx \\ &= \int_{\mathcal{X}} \phi(\pi\theta|x) Z(x) dx \end{aligned}$$

with $\phi(\pi\theta|x) = \mathbb{E} [L(\delta, h(\theta))|x] = \int_{\Theta} L(\delta, h(\theta)) \pi(\theta|x) d\theta$, the posterior loss.

We can see that minimizing the Bayes risk for each x is equivalent to minimizing the posterior loss $\phi(\pi\theta|x) = \int_{\Theta} L(\delta, h(\theta)) \pi(\theta|x) d\theta$.

To conclude the estimator that minimizes the Bayes risk is given by the estimator θ that minimizes (for each x)

$$\int_{\Theta} L(\delta, h(\theta)) \pi(\theta|x) d\theta.$$

b) For $L(\delta, \theta) = \|h(\theta) - \delta\|^2$, show that the Bayes estimator of $h(\theta)$ is $\delta^* = \mathbb{E}^* [h(\theta)|x]$.

Solution:

The Bayes estimator minimizes, for each x , $\phi(\pi\theta|x)$.

$$\begin{aligned} \phi(\pi\theta|x) &= \int_{\Theta} L(\delta, h(\theta)) \pi(\theta|x) d\theta \\ &= \mathbb{E} [L(\delta, h(\theta))|x] \\ &= \mathbb{E} [\|h(\theta) - \delta\|^2 |x] \\ &= \mathbb{E} [\|h(\theta)\|^2 + \|\delta\|^2 - 2 \langle \delta, h(\theta) \rangle |x] \\ &= \mathbb{E} [\|h(\theta)\|^2 + \delta^2 - 2 \langle \delta, h(\theta) \rangle |x] \\ &= \mathbb{E} [\|h(\theta)\|^2 |x] + \delta^2 - 2 \langle \delta, \mathbb{E} [h(\theta)|x] \rangle \end{aligned}$$

The minimum is such that:

$$\phi^{*'}(\pi\theta|x) = 2\delta^* - 2\mathbb{E}^* [h(\theta)|x] = 0$$

$$\Rightarrow \delta^* = \mathbb{E}^* [h(\theta)|x].$$

It is a minimum since $\phi^{*''}(\pi\theta|x) > 0$.

c) For $L(\delta, \theta) = |h(\theta) - \delta|$, show that the Bayes estimator of $h(\theta)$ is the median of the posterior distribution.

Solution:

The Bayes estimator minimizes, for each x , $\phi(\pi\theta|x)$.

$$\begin{aligned}\phi(\pi\theta|x) &= \int_{\Theta} L(\delta, h(\theta)) \pi(\theta|x) d\theta \\ &= \int_{\Theta} |h(\theta) - \delta| \pi(\theta|x) d\theta \\ &= \int_{-\infty}^{\delta} (\delta - h(\theta)) \pi(\theta|x) d\theta + \int_{\delta}^{+\infty} (h(\theta) - \delta) \pi(\theta|x) d\theta \\ &= \delta \int_{-\infty}^{\delta} \pi(\theta|x) d\theta - \int_{-\infty}^{\delta} h(\theta) \pi(\theta|x) d\theta + \int_{\delta}^{+\infty} h(\theta) \pi(\theta|x) d\theta - \delta \int_{\delta}^{+\infty} \pi(\theta|x) d\theta.\end{aligned}$$

$$\begin{aligned}\Rightarrow \phi'(\pi\theta|x) &= \int_{-\infty}^{\delta} \pi(\theta|x) d\theta + h(\delta) \pi(\delta|x) - h(\delta) \pi(\delta|x) - h(\delta) \pi(\delta|x) - \int_{\delta}^{+\infty} \pi(\theta|x) d\theta + h(\delta) \pi(\delta|x) \\ &= \int_{-\infty}^{\delta} \pi(\theta|x) d\theta - \int_{\delta}^{+\infty} \pi(\theta|x) d\theta \\ &= \mathbb{P}(h(\theta) \leq \delta|x) - \mathbb{P}(h(\theta) > \delta|x) \\ &= 2\mathbb{P}(h(\theta) \leq \delta|x) - 1\end{aligned}$$

Hence, $\phi^{*'}(\pi\theta|x) = 0$ if $\mathbb{P}(h(\theta) \leq \delta^*|x) = \frac{1}{2}$.

To conclude the Bayes estimator of $h(\theta)$ is such that $\mathbb{P}(h(\theta) \leq \delta^*|x) = \frac{1}{2}$, i.e., δ^* is the conditional median of $h(\theta)$ given x .

It is a minimum since $\phi^{*''}(\pi\theta|x) = 2\pi(\delta^*|x) \geq 0$.

2 Exercice 2.7

2.1 Establish the properties of the Box–Muller algorithm of Example 2.8. If U_1 and U_2 are iid $\mathcal{U}[0, 1]$, show that:

a) The transforms $X_1 = \sqrt{-2\log(U_1)} \cos(2\pi U_2)$, $X_2 = \sqrt{-2\log(U_1)} \sin(2\pi U_2)$, are iid $\mathcal{N}(0, 1)$.

Solution:

First of all, we can rewrite U_1 and U_2 as :

$$X_1^2 + X_2^2 = -2\log(U_1) \Rightarrow U_1 = \exp \left[-\frac{1}{2}(X_1^2 + X_2^2) \right]$$

and,

$$\frac{X_1}{X_2} = \tan(2\pi U_2) \Rightarrow U_2 = \frac{\arctan(\frac{X_1}{X_2})}{2\pi}.$$

Let us define $h : (U_1, U_2) \mapsto (X_1, X_2)$, an homeomorphism between $[0, 1]^2$ and \mathcal{R}^2 . We have $h^{-1} : (X_1, X_2) \mapsto (U_1, U_2)$ with : $h^{-1} : (X_1, X_2) = \left\{ \exp[-\frac{1}{2}(X_1^2 + X_2^2)], \frac{\arctan(\frac{X_1}{X_2})}{2\pi} \right\}$.

In this context, the joint density of (X_1, X_2) can be expressed in terms of the Jacobian of h^{-1} :

$$\begin{aligned} f_{X_1, X_2} &= |J_{h^{-1}}(X_1, X_2)| \\ &= \left| \begin{array}{cc} \frac{\partial h_1^{-1}}{\partial x_1} & \frac{\partial h_1^{-1}}{\partial x_2} \\ \frac{\partial h_2^{-1}}{\partial x_1} & \frac{\partial h_2^{-1}}{\partial x_2} \end{array} \right| \\ &= \left| \begin{array}{cc} x_1 \exp \left[-\frac{X_1^2 + X_2^2}{2} \right] & -x_2 \exp \left[-\frac{X_1^2 + X_2^2}{2} \right] \\ \frac{1}{2\pi} \frac{-X_2}{X_1^2 + X_2^2} & \frac{1}{2\pi} \frac{X_1}{X_1^2 + X_2^2} \end{array} \right| \\ &= \frac{1}{2\pi} \exp \left[-\frac{(X_1^2 + X_2^2)}{2} \right] \end{aligned}$$

Hence, we have :

$$\begin{aligned} f_{X_1, X_2} &= \frac{1}{\sqrt{2\pi}} \exp \left[-\frac{x_1^2}{2} \right] \times \frac{1}{\sqrt{2\pi}} \exp \left[-\frac{x_2^2}{2} \right] \\ &= f_{X_1} \times f_{X_2} \end{aligned} \tag{1}$$

We can clearly see that X_1 et X_2 are iid $\mathcal{N}(0, 1)$.

b) The polar coordinates are distributed as:

i): $r^2 = (X_1^2 + X_2^2) \sim \chi_2^2$,

ii): $\theta = \arctan \frac{X_2}{X_1} \sim \mathcal{U}[0, 2\pi]$.

Solution:

i): The distribution of the sum of the squares of 2 independent standard normal random variables is a chi-square with 2 degree of freedom.

In our case, X_1^2 and X_2^2 are independent since X_1 and X_2 are iid (question a). Therefore, r^2 is distributed as a chi-square random variable with 2 degree of freedom:

$$r^2 \sim \chi_2^2$$

Proof:

1. Let's prove that the square of a standard normal random variable is a chi-square random variable with 1 degree of freedom.

Let X be a standard normal random variable with a probability density function $f_X(x)$. We define Z such that $Z = X^2$.

- For $z < 0$, the distribution function of Z , $P(Z \leq z) = 0$ since Z is a square. Therefore, $f_Z(z) = 0$

- For $z \geq 0$, the distribution function of Z is :

$$\begin{aligned} P(Z \leq z) &= P(X^2 \leq z) \\ &= P(-z^{\frac{1}{2}} \leq X \leq z^{\frac{1}{2}}) \\ &= \int_{-z^{\frac{1}{2}}}^{z^{\frac{1}{2}}} f_X(x) dx \end{aligned} \tag{2}$$

Therefore, for $z \geq 0$, the density of Z is:

$$\begin{aligned} f_Z(z) &= \frac{dP(Z \leq z)}{dz} \\ &= \frac{\int_{-z^{\frac{1}{2}}}^{z^{\frac{1}{2}}} f_X(x) dx}{dz} \\ &= f_X(z^{\frac{1}{2}}) \frac{dz^{\frac{1}{2}}}{dz} - f_X(-z^{\frac{1}{2}}) \frac{d(-z^{\frac{1}{2}})}{dz} \\ &= \frac{1}{\sqrt{2\pi}} \exp \left[-\frac{1}{2}(z^{\frac{1}{2}})^2 \right] \times \frac{1}{2} z^{-\frac{1}{2}} - \frac{1}{\sqrt{2\pi}} \exp \left[-\frac{1}{2}(-z^{\frac{1}{2}})^2 \right] \times \frac{-1}{2} z^{-\frac{1}{2}} \\ &= \frac{1}{\sqrt{2\pi}} z^{\frac{1}{2}} \exp \left[-\frac{1}{2}z \right] \\ &= \frac{1}{2^{\frac{1}{2}} \Gamma(\frac{1}{2})} z^{\frac{1}{2}-1} \exp \left[-\frac{1}{2}z \right] \text{ because } \Gamma(\frac{1}{2}) = \sqrt{\pi}. \end{aligned} \tag{3}$$

We recognize the density function of a chi-square random variable with 1 degree of freedom.

2. *Let's prove that the sum of independent chi-square random variables is a chi-square random variable.*

Let $Z = \sum_{i=1}^k Z_i$ and $M_Z(t)$ its moment generating function. Suppose that the random variable Z_i is a chi-square random variable with n_i degrees of freedom and all the random variables are independent, thus:

$$\begin{aligned} M_Z(t) &= \prod_{i=1}^k M_{Z_i}(t) \\ &= \prod_{i=1}^k (1 - 2t)^{-\frac{n_i}{2}} \\ &= (1 - 2t)^{-\sum_{i=1}^k \frac{n_i}{2}} \\ &= (1 - 2t)^{-\frac{n}{2}} \end{aligned} \tag{4}$$

We can see that the moment generating function of Z is the moment generating function of a chi-square random variable with $n = \sum_{i=1}^k n_i$ degree of freedom.

3. *Combining the two facts above, we obtain that the sum of squares of 2 independent standard normal random variables is a chi-square random variable with 2 degree of freedom.*

ii): From question a, we know that $U_2 = \frac{\arctan(\frac{X_1}{X_2})}{2\pi} = \frac{\theta}{2\pi} \Rightarrow \theta = 2\pi U_2$.

Since $U_2 \sim \mathcal{U}[0, 1]$, we have:

$$\theta \sim \mathcal{U}[0, 2\pi]$$

.

c) Establish that $\exp(-\frac{r^2}{2}) \sim \mathcal{U}[0, 1]$, and so r^2 and θ can be simulated directly.

Solution:

From a), we know that: $X_1^2 + X_2^2 = -2\log(U_1)$.

From b), we have that: $r^2 = X_1^2 + X_2^2$.

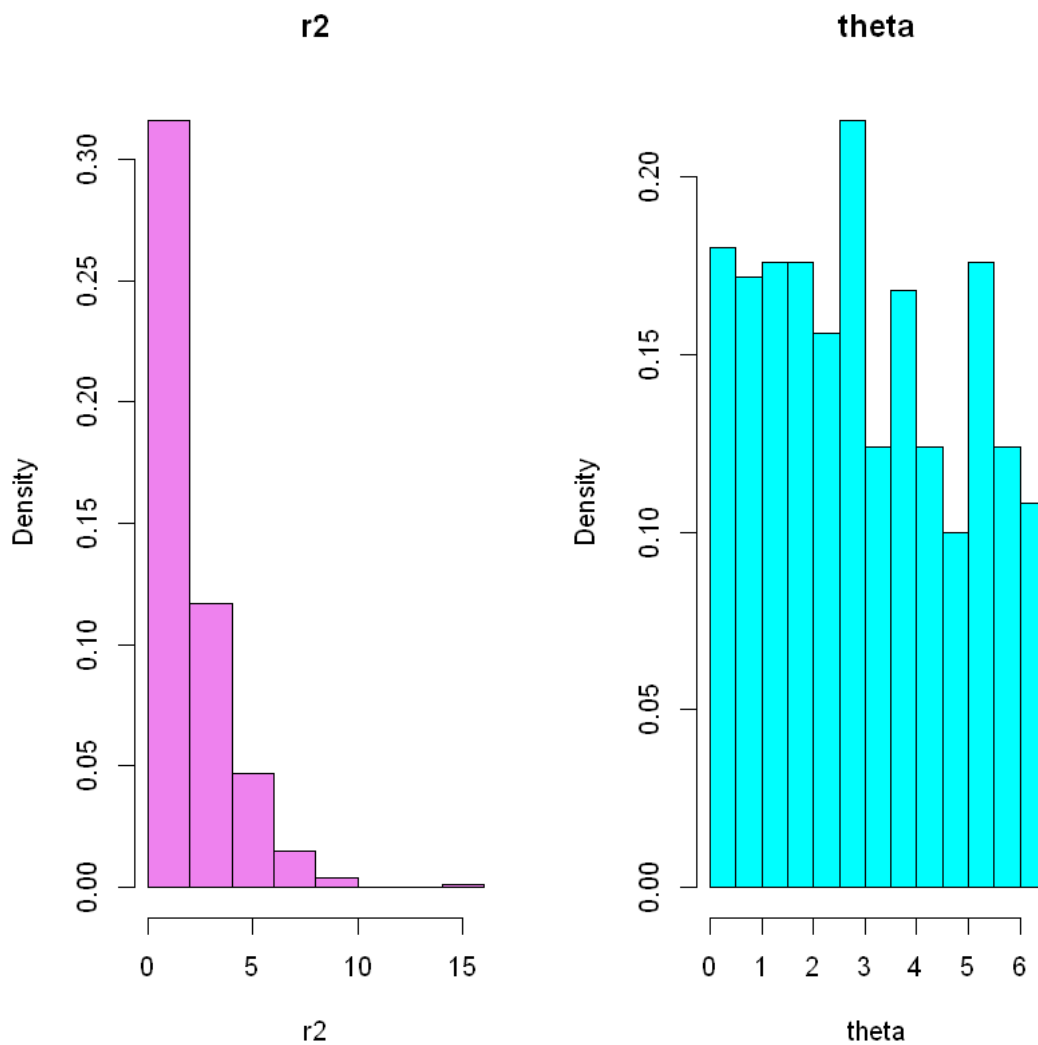
Hence, $r^2 = -2\log(U_1) \Rightarrow \exp(-\frac{r^2}{2}) = U_1$ and consequently, $\exp(-\frac{r^2}{2}) \sim \mathcal{U}[0, 1]$.

To conclude, we can simulate r^2 and θ directly from the uniform distribution on $[0, 1]$ since:

- $r^2 = -2\log(U_1)$ and $\theta = 2\pi U_2$, and
- U_1 and U_2 are iid $\mathcal{U}[0, 1]$.

Example of a simulation using R:

```
In [1]: U1 = runif(500,0,1)
        U2 = runif(500,0,1)
        r2 = -2*log(U1)
        theta = 2*pi*U2
        par(mfrow=c(1,2))
        hist(r2, freq=F, col = "violet", main="r2")
        hist(theta, freq=F, col= "cyan", main="theta")
```



3 Exercise 2.2

- 3.1 a) Generate a binomial $\text{Bin}(n, p)$ random variable with $n = 25$ and $p = .2$. Make a histogram and compare it to the binomial mass function, and to the R binomial generator.

Comment: We have seen in lecture 1 that if $U \sim U_{[0,1]}$, then the random variable $F^{-1}(U)$ has the distribution F (Probability Integral Transform). It is the starting point of the inverse transform method that I am going to use: to generate a random variable $x \sim F$, I generate a random variable $u \sim U_{[0,1]}$ and then I use the transform $x = F^{-1}(u)$.

```
In [2]: n = 25
        p = 0.2

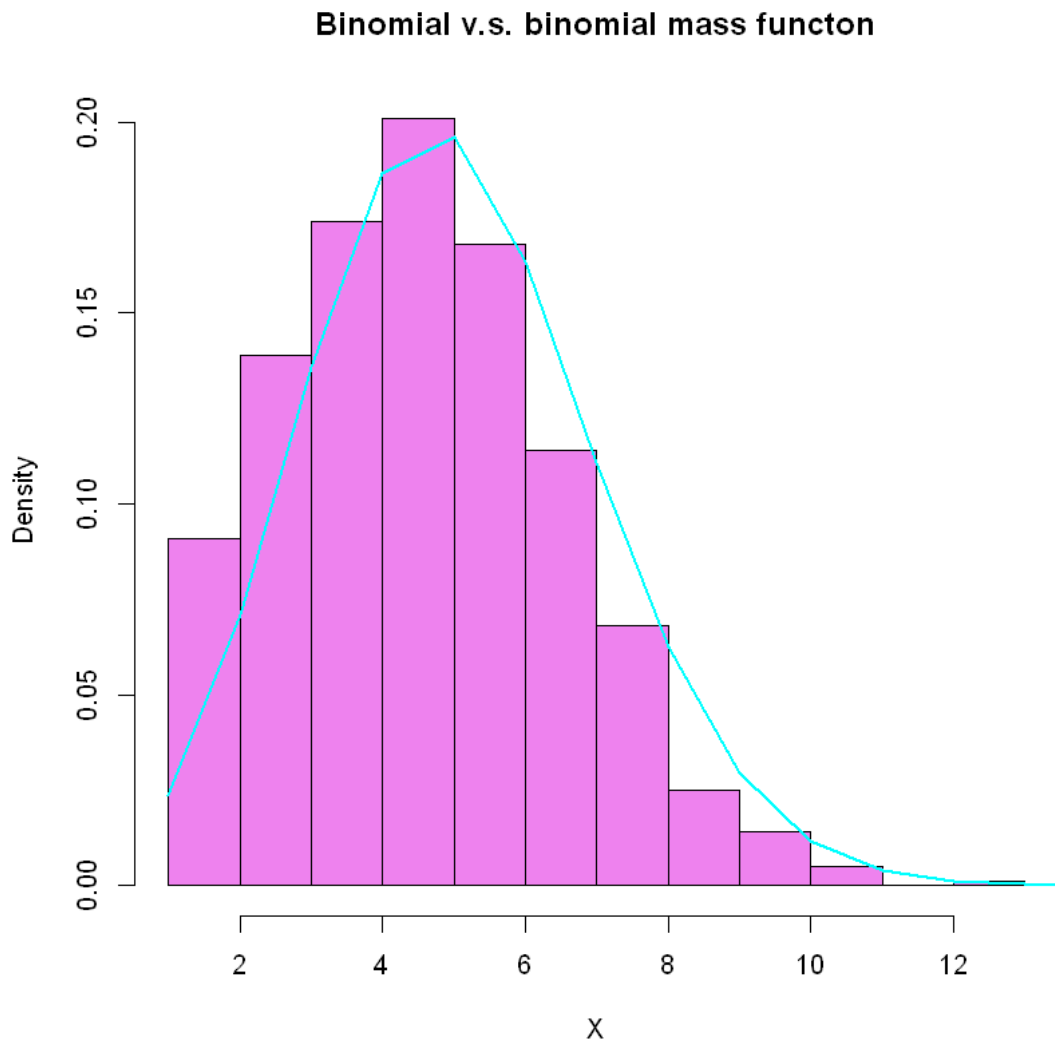
        N = 1000 # number of simulations

        # 1) I use the inverse transform method for generating binomial random variables
        Binomial = function(N,n,p){
          proba = pbinom(0:n,n,p)
          X = rep(0,N)
          for (i in 1:N){
            u = runif(1)
            X[i] = 0 + sum(proba<u)
          }
          return(X)
        }

        # 2) I use the R binomial generator for generating binomial random variables
        rbino = rbinom(N, n, p)

In [3]: # Histogram : binomial (inverse transform method) v.s. binomial mass function

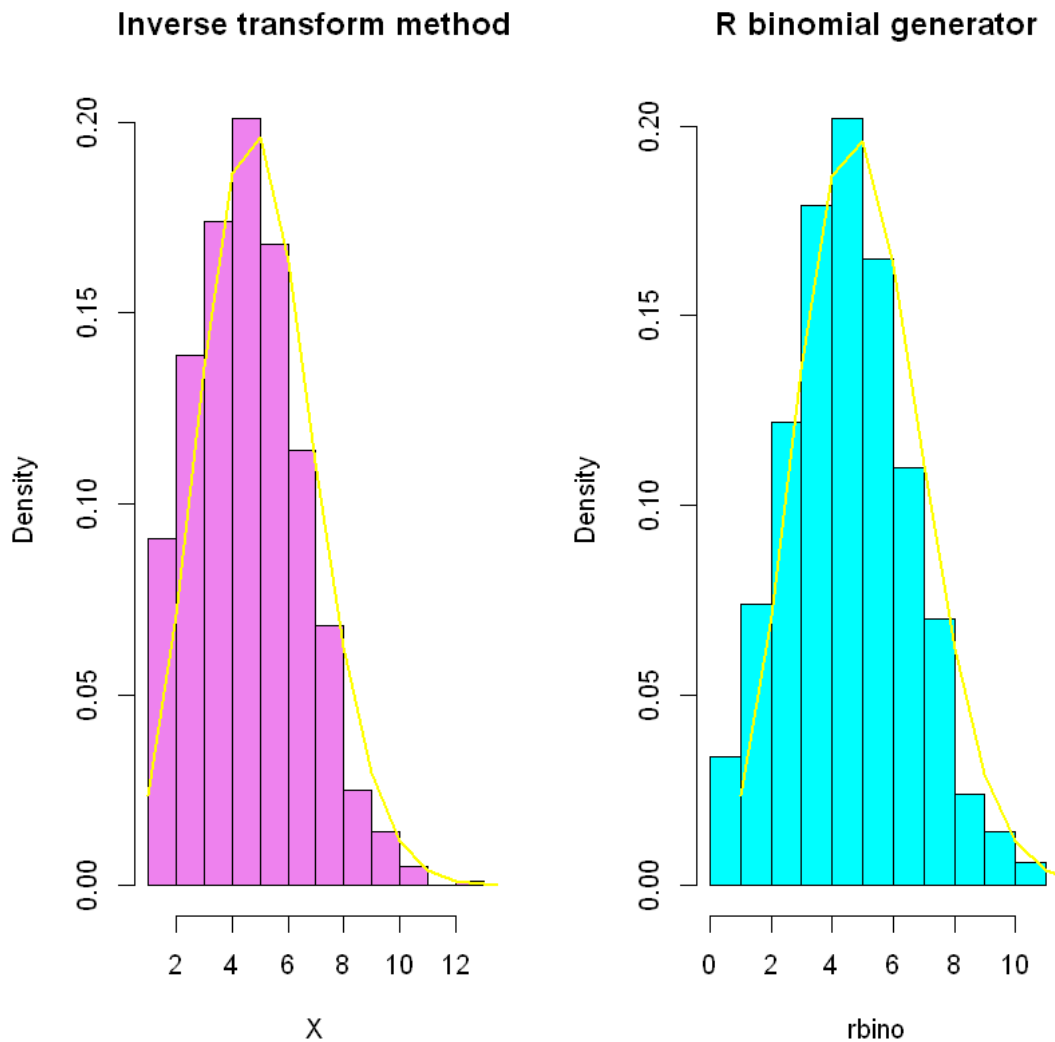
        X = Binomial(N,n,p)
        hist(X, col="violet", freq=F, main="Binomial v.s. binomial mass functon")
        # Mass function
        lines(1:n, dbinom(1:n,n,p), lwd = 2, col="cyan")
```

In [4]: *# Histogram : binomial (inverse transform method) v.s. the R binomial generator*

```
par(mfrow=c(1,2))
hist(X,freq=F, col="violet", main="Inverse transform method")
lines(1:n, dbinom(1:n,n,p), lwd = 2, col="yellow")

hist(rbino,freq=F, col="cyan", main="R binomial generator")
lines(1:n, dbinom(1:n,n,p), lwd = 2, col="yellow")
```



Comment: The histograms are very similar. The transform method gives satisfactory results.

3.2 b) Generate 5,000 logarithmic series random variables. Make a histogram and plot the mass function.

In [5]: *# I use the inverse transform method:*

```
logarithmic = function(N,p){
  x = seq(1,35,1)
  mass = (-(1-p)^x)/(x*log(p))
  mass_f = cumsum(mass)
  X = rep(0,N)
```

```

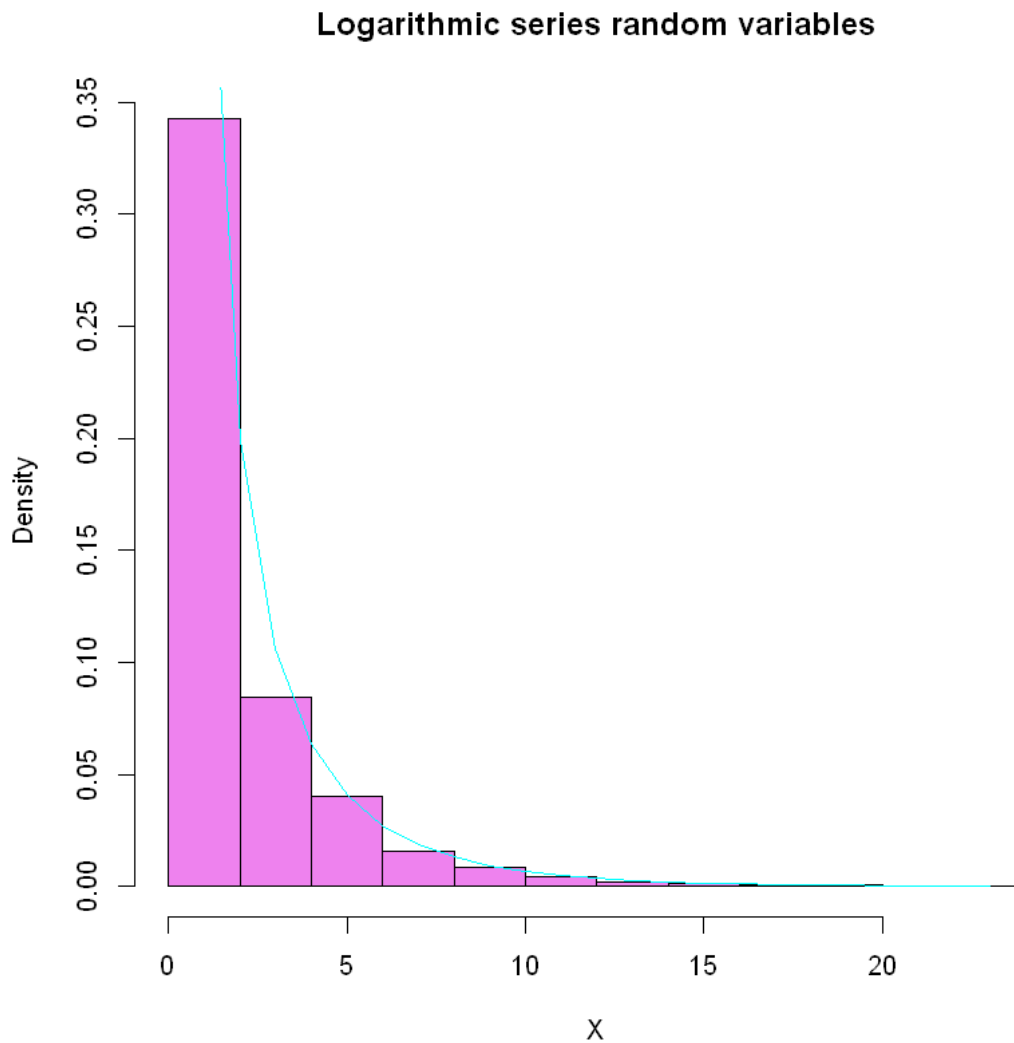
    for (i in 1:N){
      unif = runif(1)
      X[i] = x[1] + sum(mass_f < unif)
    }
    return (X)
  }
}

```

```

In [6]: N = 5000
p = 0.2
X = logarithmic(N,p)
hist(X, freq=F, col = "violet", main="Logarithmic series random variables")
X = seq(min(X), max(X), 1)
lines((-1-p)^X/(X*log(p)), lwd = 1.5, col="cyan")

```



Conclusion: To conclude, the inverse transform method gives satisfactory method with the binomial or logarithmic series random variables.

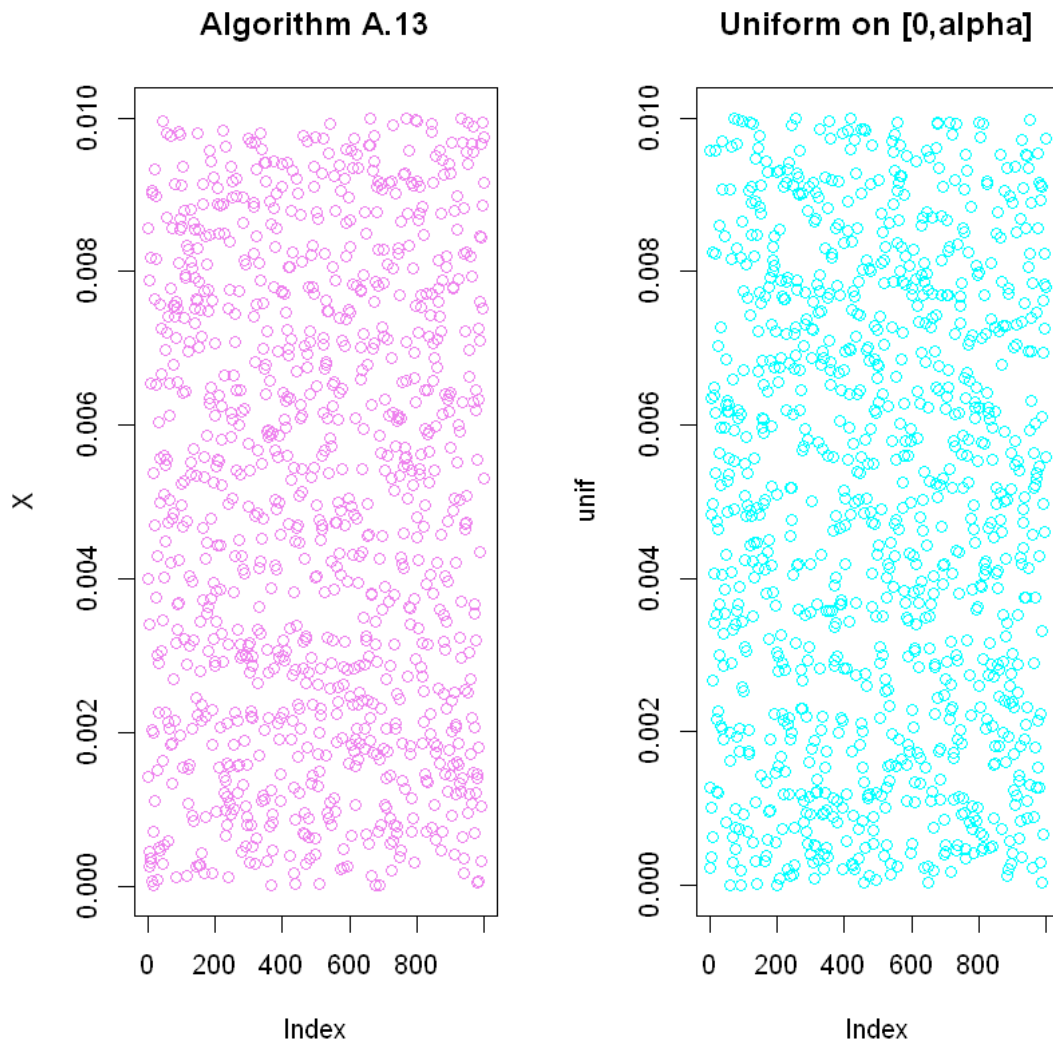
4 Exercice 2.19

4.1 For $\alpha \in [0, 1]$, show that the algorithm A.13 produces a simulation from $U([0, \alpha])$. Compare it with the transform αU , $U \sim U(0, 1)$ for values of α close to 0 and close to 1.

```
In [7]: # Algorithm A.13 :
Generateur1 = function(N, alpha){
  X = seq(1,N) # initialisation
  for (i in 1: N){
    #X[i] = runif(1)
    while (X[i] >= alpha){
      X[i] = runif(1)
    }
  }
  return(X)
}
```

1. Comparison between the algorithm A.13 and simulations from $U([0, \alpha])$:

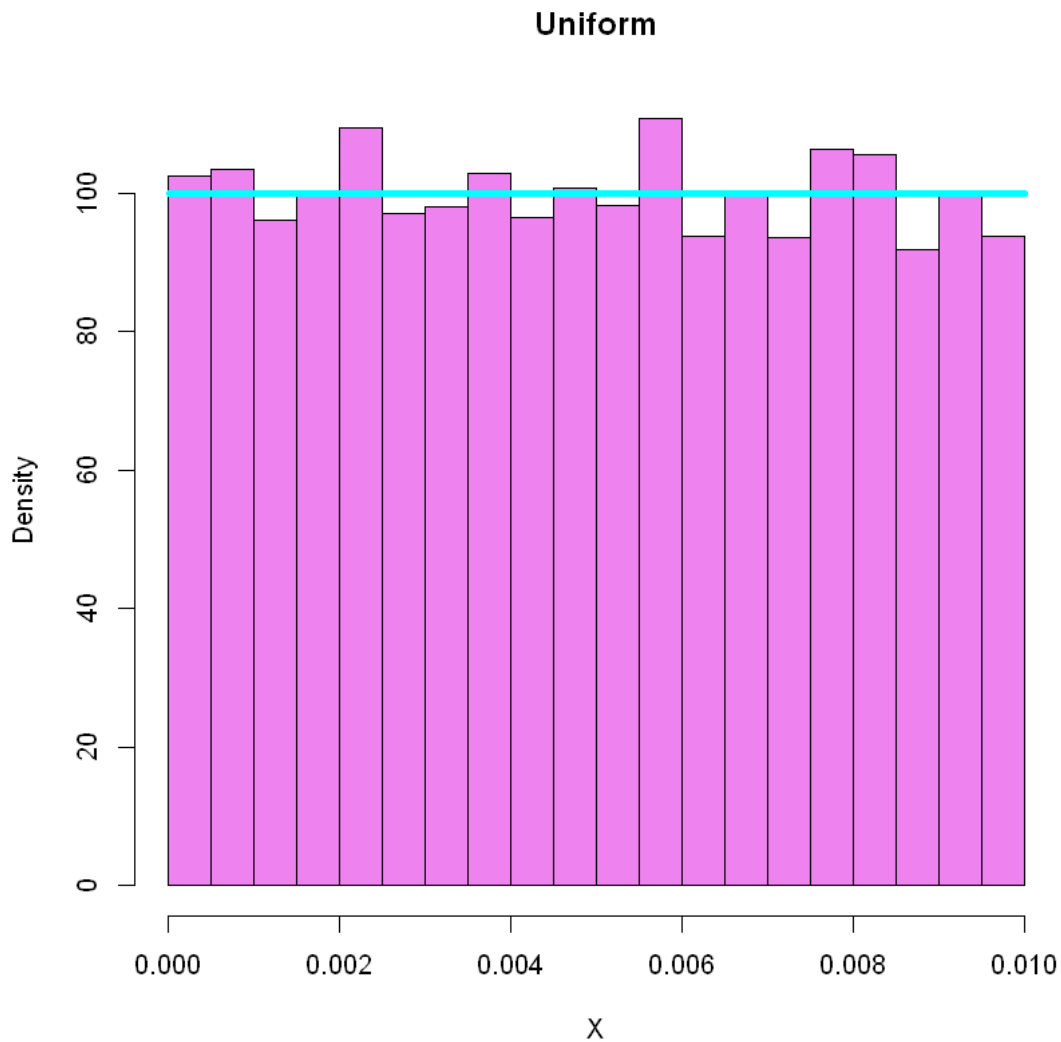
```
In [8]: N = 1000 # Number of simulations
alpha = 0.01
#alpha=runif(1) # Results are the same using alpha = runif(1)
X = Generateur1(N, alpha)
u = seq(min(X), max(X), length.out = N)
unif = runif(u,0,alpha)
par(mfrow=c(1,2))
plot(X, col = 'violet', main="Algorithm A.13")
plot(unif,col='cyan', main="Uniform on [0,alpha]")
```



The graphs are very similar.

We are now going to compare the histogram given by the algorithm A.13 with the probability density function of a random variable following a uniform distribution on $[0, \alpha]$.

```
In [9]: N = 10^4 # We increase the number of simulations in order to have a better view
X = Generateur1(N, alpha)
u = seq(min(X), max(X), length.out = N)
hist(X, freq=F, col = "violet", main="Uniform")
lines(u, dunif(u,0,alpha), lwd = 4, col="cyan")
```



Comment: To conclude, the algorithm A.13 produces a simulation from $U([0, \alpha])$. In fact, the algorithm picks the random variables with uniform distribution on $[0, 1]$ in the range of $[0, \alpha]$ where they are still uniformly distributed.

Note that results are the same using αU , $U \sim U(0, 1)$ (i.e. $\alpha = \text{runif}(1)$ in the code above) instead of $\alpha = 0.01$ (arbitrary value).

2) Comparison of the algorithm A.13 with the transform αU , $U \sim U(0, 1)$ for values of α close to 0 and close to 1

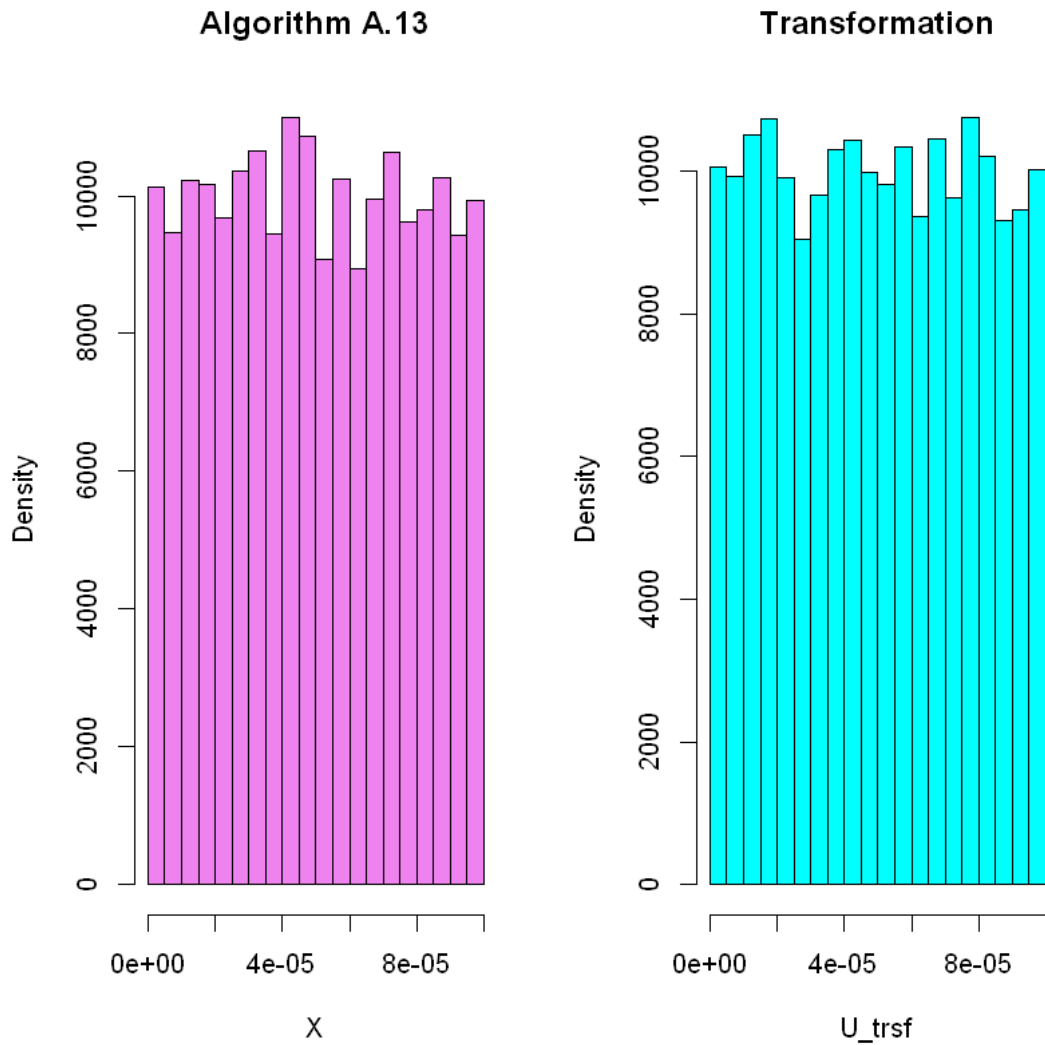
Case 1: α close to 0

```

In [10]: alpha = 0.0001
X = Generateur1(N, alpha)
U_trsf = runif(N) * alpha # Transform
Y = seq(min(U_trsf), max(U_trsf), length.out = 1000)

par(mfrow=c(1,2))
hist(X, freq=F, col = "violet", main="Algorithm A.13")
hist(U_trsf, freq=F, col= "cyan", main="Transformation")

```



Case 2: α close to 1

```

In [11]: alpha = 0.99
X = Generateur1(N, alpha)

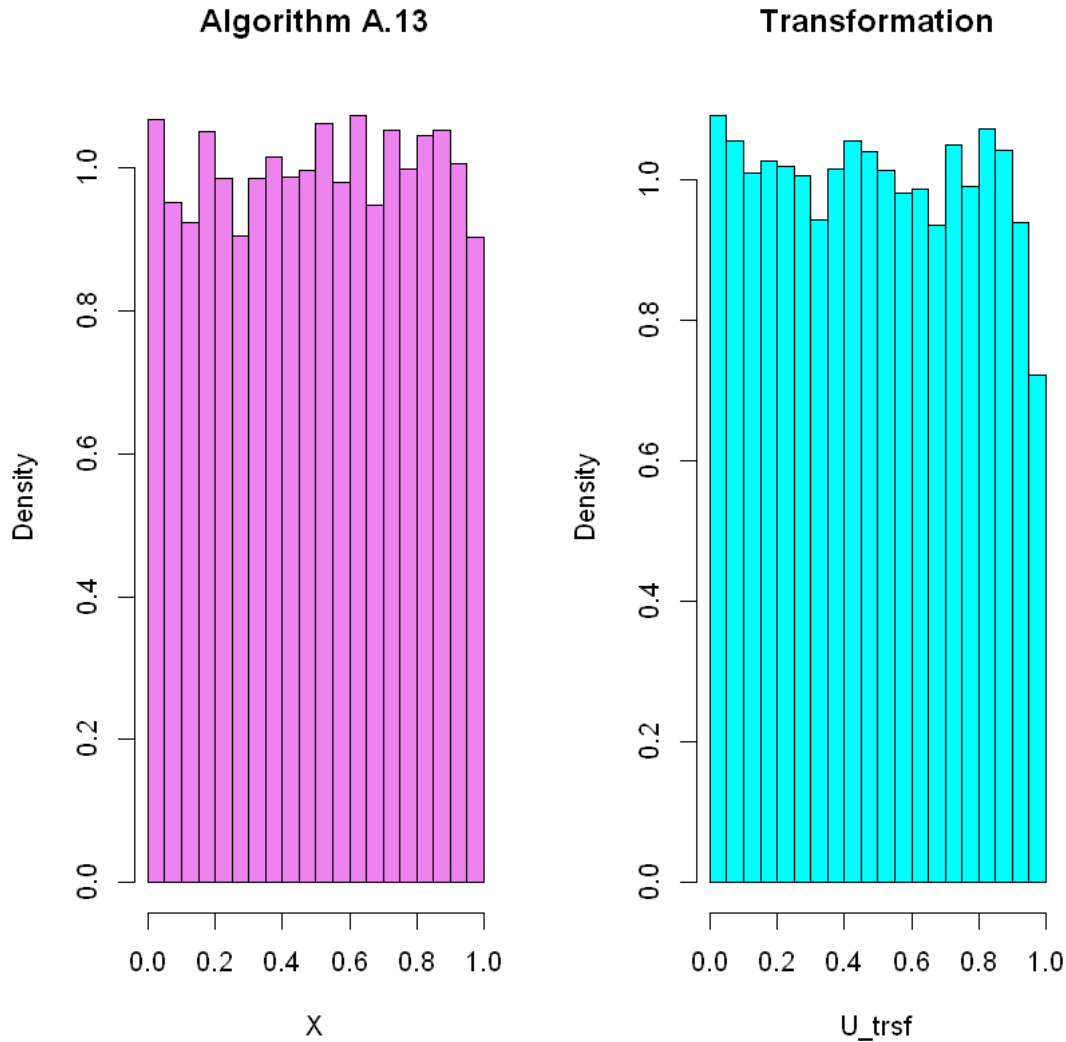
```

```

U_trsf = runif(N) * alpha # Transform
Y = seq(min(U_trsf), max(U_trsf), length.out = 1000)

par(mfrow=c(1,2))
hist(X, freq=F, col = "violet", main="Algorithm A.13")
hist(U_trsf, freq=F, col= "cyan", main="Transformation")

```



Comment: For both cases (α close to 0 or 1), the histograms from the algorithm A.13 and from the transformation are very similar. It shows that the transform αU with $U \sim U(0, 1)$ produces a simulation from $U(0, \alpha)$.

Link with the accept-reject algorithm : The simulation of the target density f is done using an instrumental density g . In our case f is the $U(0, 1)$ and g is the $U(0, \alpha)$. We have $\frac{f(X)}{Mg(X)} = \alpha$ if

$M = 1$. It means that all the points in $[0, \alpha]$ are accepted.