

CS Computer Lab 3

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Question 1: Cluster sampling

In this section, we used the data file **population.csv**, containing a list of Swedish cities, along with their respective populations. The aim was to select twenty cities at random for an opinion pool, where the random sampling was without replacement and with probabilities proportional to the populations of each city.

In order to do this, we normalised the population proportions so that these were represented as sub-intervals of the interval $[0,1]$. A uniform random number was then generated; whichever sub-interval this number fell in, the corresponding city was selected. The function is detailed below:

```
selectone <- function (data) {  
  interval <- c()  
  interval[1] <-  
    data$Population[1]/sum(data$Population)  
  for (i in 2:dim(data)[1]) {  
    interval[i] <- interval[i-1] +  
      data$Population[i]/sum(data$Population)  
  }  
  rand <- runif(1, 0, 1)  
  pick <- length(which(interval < rand)) + 1  
  return(data[pick, ])  
}
```

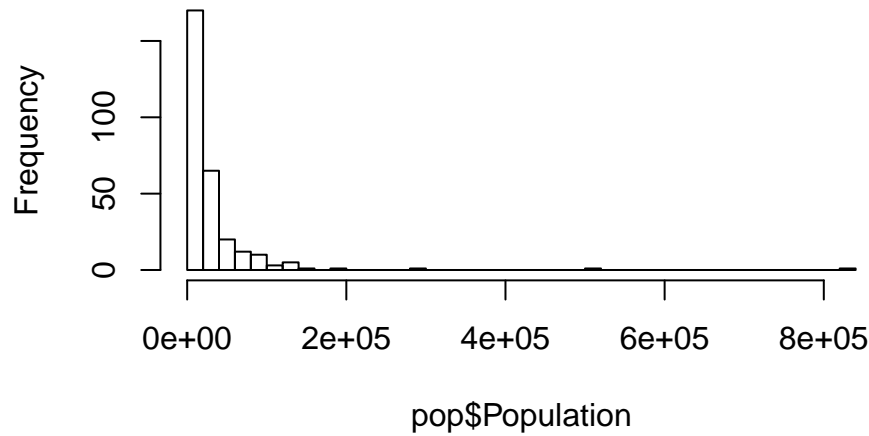
We then applied the function to the list of cities 20 times, each time removing the selected one from the list for the next iteration.

The following cities were selected:

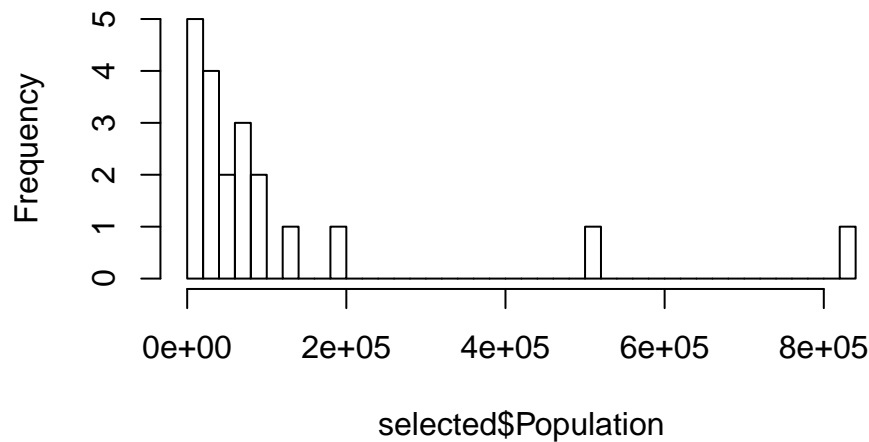
##	Municipality	Population
## 1	Karlskoga	29742
## 2	Ulricehamn	22753
## 3	Kalmar	62388
## 4	Jönköping	126331
## 5	Ljungby	27410
## 6	Täby	63014
## 7	Trelleborg	41891
## 8	Stockholm	829417
## 9	Luleå	73950
## 10	Uppsala	194751
## 11	Fagersta	12249
## 12	Göteborg	507330
## 13	Sundsvall	95533
## 14	Gävle	94352
## 15	Piteå	40860
## 16	Ovanåker	11530
## 17	Smedjebacken	10758
## 18	Katrineholm	32303
## 19	Nybro	19576
## 20	Hallsberg	15235

Generally our program selected large cities: the mean population of our selected ones was 115,569, much larger than the sample mean of 32,209. This is also shown by comparing the histograms of the two.

Populations of all 270 cities



Populations of our 20 selected cities



These histograms confirm what we observed initially; our selection process favours large cities. This makes sense when considering the approach taken here, however the people from these selected cities may be unrepresentative of Swedish people as a whole, as clearly an overwhelming proportion of people live in lots of small cities.

Question 2: Different distributions

In this section we consider the double exponential distribution, with pdf:

$$DE(\mu, \alpha) = \frac{\alpha}{2} e^{-\alpha|x-\mu|} \quad (1)$$

The first goal was to generate random numbers from the $DE(0, 1)$ distribution from $Unif(0, 1)$ using the inverse CDF method. For this we need to derive the inverse CDF of $DE(0, 1)$, starting from the pdf:

$$f(x) = \frac{1}{2} e^{-|x|} \quad (2)$$

Therefore the CDF is:

$$F(x) = \int_{-\infty}^x \frac{1}{2} e^{-|u|} du \quad (3)$$

For $x \geq 0$:

$$F(x) = \int_0^x \frac{1}{2} e^{-|u|} du + \int_{-\infty}^0 \frac{1}{2} e^{-|u|} du \quad (4)$$

$$= \int_0^x \frac{1}{2} e^{-u} du + \int_{-\infty}^0 \frac{1}{2} e^u du \quad (5)$$

$$= \frac{1}{2} [e^{-u}]_0^x + \frac{1}{2} [e^u]_{-\infty}^0 \quad (6)$$

$$= 1 - \frac{1}{2} e^{-x} \quad (7)$$

For $x < 0$ (and therefore $u < 0$):

$$F(x) = \int_{-\infty}^x \frac{1}{2} e^{-|u|} du \quad (8)$$

$$= \frac{1}{2} [e^u]_{-\infty}^x \quad (9)$$

$$= \frac{1}{2} e^x \quad (10)$$

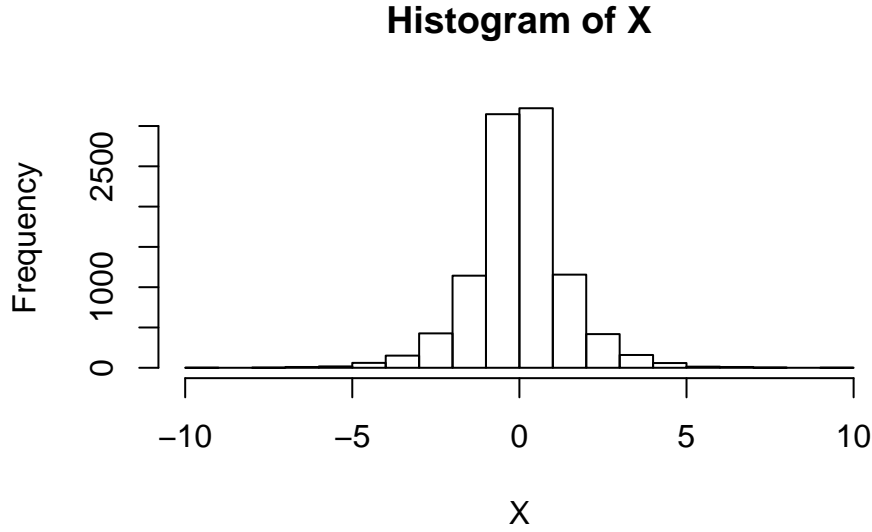
To get the inverse CDF, we set:

$$y = \begin{cases} 1 - \frac{1}{2} e^{-x}, & \text{for } x \geq 0 \\ \frac{1}{2} e^x, & \text{for } x < 0 \end{cases} \quad (11)$$

Rearranging gives us the following inverse CDF:

$$F^{-1}(y) = \begin{cases} -\log(2(1 - y)), & \text{for } y \geq \frac{1}{2} \\ \log(2y), & \text{for } y < \frac{1}{2} \end{cases} \quad (12)$$

We then wrote a function `laplace()` to generate random numbers from the $DE(0, 1)$ distribution, by generating a uniform r.v. and taking the inverse CDF of this. We generated 10000 such random numbers; the results are shown in the histogram below.



This histogram looks reasonable as it resembles the pdf of the Laplace distribution (on a much larger scale of course).

The next goal was to use the Acceptance/rejection method to generate from the Normal $N(0, 1)$ distribution (target density f_X). We were given that the previously considered $DE(0, 1)$ distribution could be used as a majorising density (f_Y). All we needed was to find a constant c such that, for all x :

$$cf_Y(x) \geq f_X(x) \quad (13)$$

Subbing in the respective pdfs:

$$\frac{c}{2}e^{-|x|} \geq \frac{1}{\sqrt{2\pi}}e^{-\frac{1}{2}x^2} \quad (14)$$

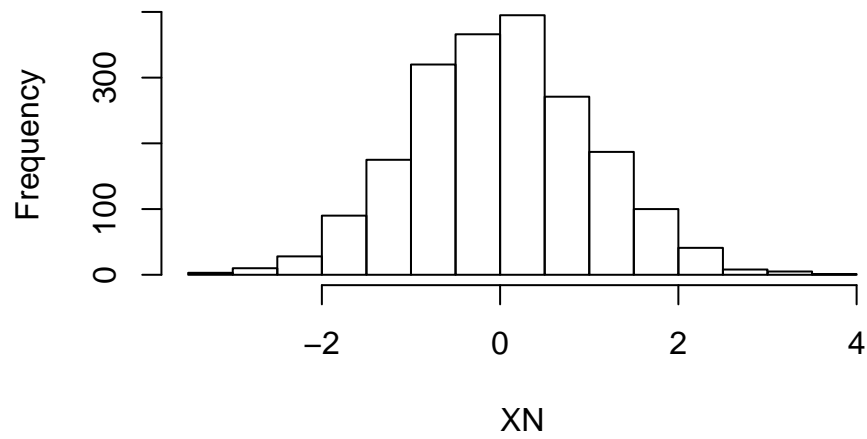
Rearranging gives us:

$$c \geq \frac{2}{\sqrt{2\pi}}e^{|x| - \frac{1}{2}x^2} \quad (15)$$

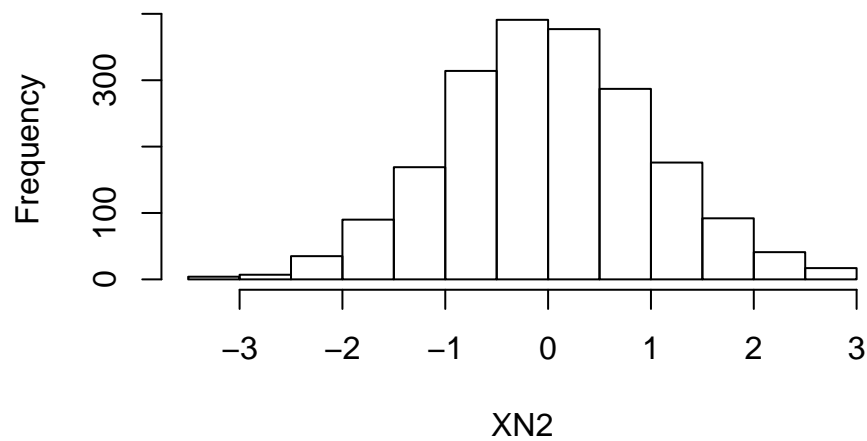
The right hand side is maximised for $x = \pm 1$ so the limiting case is $c = \frac{2}{\sqrt{2\pi}}e^{0.5}$. Using this constant c , we applied the Acceptance/rejection method to generate 2000 random numbers.

The average rejection was calculated, we obtained 0.2472714. The expected rejection rate was also found, as it is the probability that our acceptance condition is not validated, ie. $ER = 1 - 1/c$. This expected rejection rate was therefore 0.2398265, so very close to the actual rate. We also plotted the histogram of our results. Below it we have the histogram of 2000 normal random numbers using the in-built `rnorm()` function.

Random numbers generated from Acc/rej



2000 random numbers generated from rnorm()



The plots are very similar in shape, it looks like our Acceptance/rejection method was pretty effective at simulating draws from the $N(0, 1)$ distribution.

Appendix

```
knitr::opts_chunk$set(echo = FALSE, message = FALSE, fig.width = 5, fig.asp = 0.66, fig.show = "asis", )
#1.1
setwd("~/Semester2/CS/Lab3")
pop <- read.csv2(paste0(getwd(), "/population.csv"), stringsAsFactors = FALSE)

selectone <- function (data) {
  interval <- c()
  interval[1] <-
    data$Population[1]/sum(data$Population)
  for (i in 2:dim(data)[1]) {
    interval[i] <- interval[i-1] +
      data$Population[i]/sum(data$Population)
  }
  rand <- runif(1, 0, 1)
  pick <- length(which(interval < rand)) + 1
  return(data[pick, ])
}

set.seed(123456)
selected <- data.frame(Municipality=vector(length = 20), Population=vector(length = 20))
data <- pop
for (i in 1:20) {
  selected[i, ] <- selectone(data = data)
  pick <- which(data$Municipality == selected$Municipality[i])
  data <- data[ -pick, ]
}
print(selected)
meanselect <- round(mean(selected$Population))
meanpop <- round(mean(pop$Population))
hist(pop$Population, breaks = 30, main="Populations of all 270 cities")
hist(selected$Population, breaks = 30, main = "Populations of our 20 selected cities")
#2.1
rlaplace <- function(n, mu, alpha) {
  U <- runif(n, min=0, max=1)
  X <- mu - 1/alpha*(sign(U-0.5)*log(1-2*abs(U-0.5)))
  return(X)
}

set.seed(12345)
X <- rlaplace(10000, 0, 1)
hist(X)
c <- 2/sqrt(2*pi) * exp(1/2)

normgen <- function(c) {
  fY <- function(x) {
    y <- 1/2*exp(-abs(x))
    return(y)
  }
}

rej <- 0
repeat {
```

```

Y <- rlaplace(1, 0, 1)
U <- runif(1, 0, 1)
cond <- dnorm(Y, 0, 1)/(c*fY(x=Y))
if (U <= cond) {
  X <- Y
  break
}
else {
  rej <- rej+1
}

}
return(list(X=X, rej=rej))
}

XN <- c()
rej <- c()
for (i in 1:2000) {
  X <- normgen(c=c)
  XN[i] <- X$X
  rej[i] <- X$rej
}

R <- sum(rej)/(sum(rej)+2000)
ER <- 1- 1/c

hist(XN, main="Random numbers generated from Acc/rej")
XN2 <- rnorm(2000, 0, 1)
hist(XN2, main="2000 random numbers generated from rnorm()")

```

Collaborations

Methodology and results were shared and discussed with members of Group 6, Chih-Yuan Lin and Sarah Walid. Alsaadi.