

# Script-tarea.R

Usuario

2020-02-20

```
# Athziri Nalleli Charles Rosas  
# 1871521  
# 20/02/20
```

```
# Ejercicio #1 -----
```

```
library(plyr)  
accidentes <- c(0,1,0,2,2,0,1,4,3,0,1,5,1,2,3,4,0,1,1,3,4)  
acc <- count(accidentes)  
acc
```

```
##   x freq  
## 1 0    5  
## 2 1    6  
## 3 2    3  
## 4 3    3  
## 5 4    3  
## 6 5    1
```

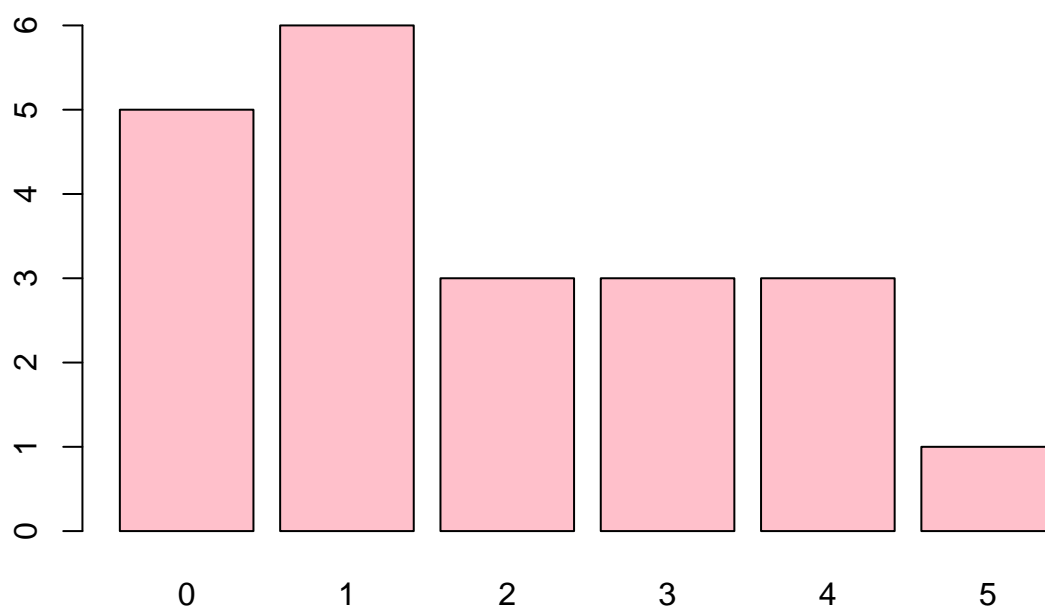
```
(acc$freq/sum(acc$freq)*100)
```

```
## [1] 23.809524 28.571429 14.285714 14.285714 14.285714  4.761905
```

```
acc$rf <- acc$freq/sum(acc$freq)*100
```

```
barplot(acc$freq, names.arg = acc$x, main = "Accidentes en el aserradero", col = "pink")
```

## Accidentes en el aserradero



```
mean(accidentes)
```

```
## [1] 1.809524
```

```
sum(accidentes)
```

```
## [1] 38
```

```
barplot(accidentes)
```

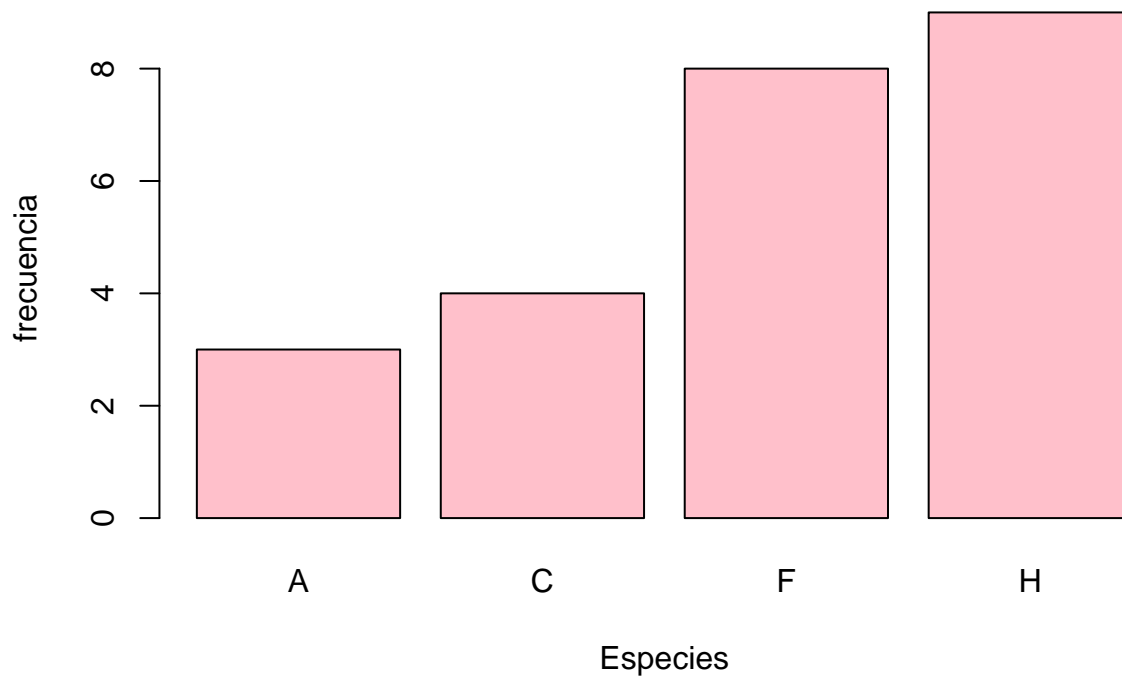
```
# Ejercicio #2 -----

especies <- c("F","H","F","C","F","A","H","F","H","C","A","C","F","H","H","H","F","H","A","C","F","H","H")

.sp <- count(especies)
.sp$rf <- .sp$freq/sum(.sp$freq)*100
.sp

##      x freq      rf
## 1 A      3 12.50000
## 2 C      4 16.66667
## 3 F      8 33.33333
## 4 H      9 37.50000

barplot(.sp$freq, names.arg = .sp$x, col = "pink", ylab = "frecuencia", xlab = "Especies")
```



```
# Ejercicio #3 -----

library(repmis)
conjunto <- source_data("https://www.dropbox.com/s/hmsf07bbayxv6m3/cuadro1.csv?dl=1")

## Downloading data from: https://www.dropbox.com/s/hmsf07bbayxv6m3/cuadro1.csv?dl=1
## SHA-1 hash of the downloaded data file is:
## 2bdde4663f51aa4198b04a248715d0d93498e7ba

.vc <- table(conjunto$Vecinos, conjunto$Especie)
.vc1 <- addmargins(as.table(.vc))
.vc1

##
##      C  F  H Sum
## 0    1  0  2   3
## 1    1  2  1   4
## 2    3  2  1   6
## 3    5  3  5  13
## 4    5  5  3  13
## 5    5  1  0   6
## 6    2  1  2   5
## Sum 22 14 14  50

# Ejercicio #4 -----

dbh <- conjunto$Diametro
```

```

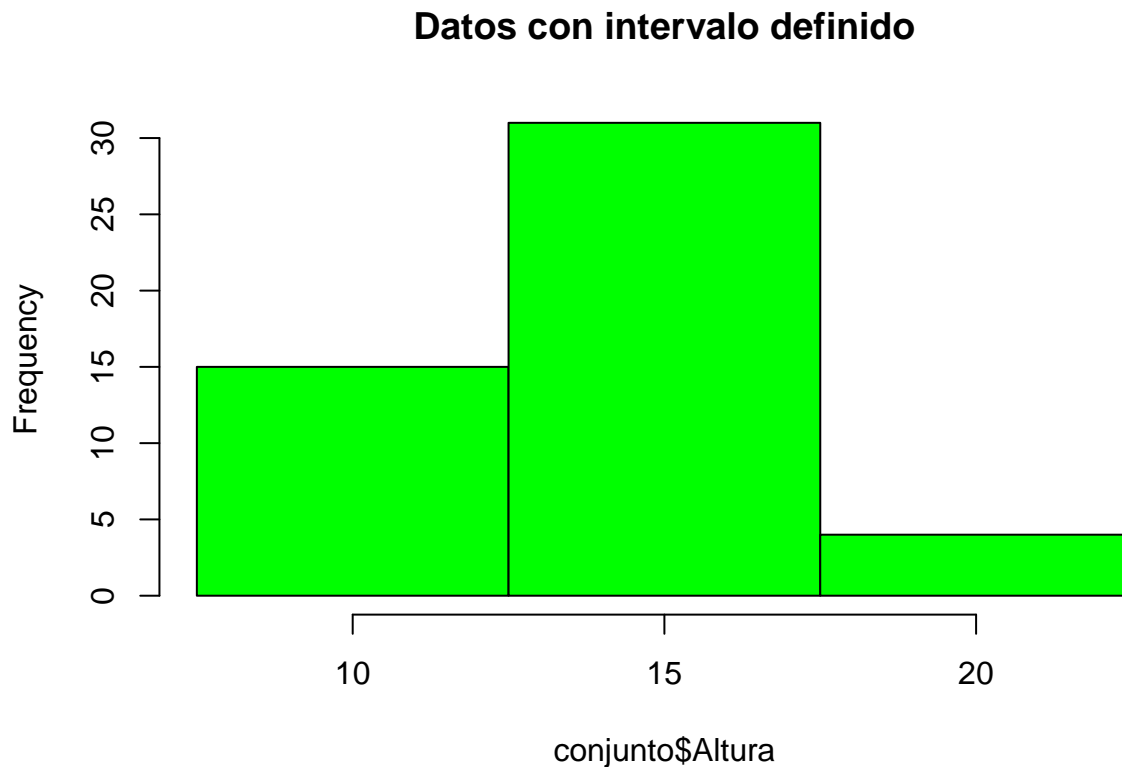
range(dbh)

## [1] 7.7 22.7
intervalo <- seq(7.5, 22.5, by=5)
intervalo

## [1] 7.5 12.5 17.5 22.5
dbh.table <- cut(dbh, intervalo)
table(dbh.table)

## dbh.table
## (7.5,12.5] (12.5,17.5] (17.5,22.5]
##          6          27          16
hist(conjunto$Altura, breaks = intervalo, main = "Datos con intervalo definido", col = "green")

```



```

dbh.prop <- cbind(table(dbh.table))
dbh.per <- round(prop.table(dbh.prop)*100,2)

```

*# Ejercicio #5 -----*

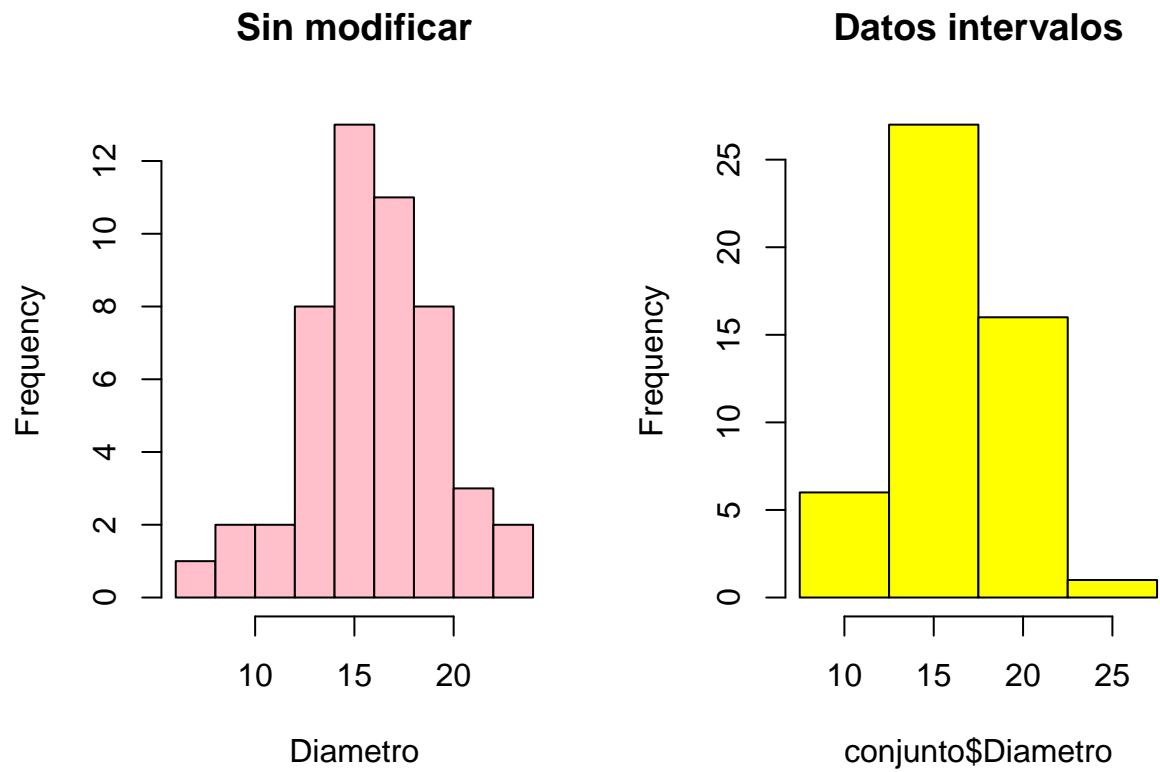
```

intervalo <- seq(7.5, 27.5, by=5)
intervalo

```

```
## [1] 7.5 12.5 17.5 22.5 27.5
```

```
par(mfrow=c(1,2))
hist(conjunto$Diametro,col = "pink", main = "Sin modificar", xlab = "Diametro")
hist(conjunto$Diametro,breaks = intervalo, col = "yellow", main = "Datos intervalos")
```



```
par(mfrow=c(1,1))
```