

## Corrección\_Sarha.R

Sarha

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```
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# 03.03.21

# Parte I Importar datos csv -----
--

conjunto <- read.csv("Cuadro1.csv", header = TRUE)
head(conjunto)

##   Arbol Fecha Especie Posicion Vecinos Diametro Altura
## 1     1    12      F        C        4     15.3   14.78
## 2     2    12      F        D        3     17.8   17.07
## 3     3     9      C        D        5     18.2   18.28
## 4     4     9      H        S        4      9.7    8.79
## 5     5     7      H        I        6     10.8   10.18
## 6     6    10      C        I        3     14.1   14.90

tail(conjunto)

##   Arbol Fecha Especie Posicion Vecinos Diametro Altura
## 45    45    24      C        I        4     10.2   13.93
## 46    46    23      F        I        3     14.4   12.68
## 47    47    24      C        S        6      7.7   10.00
## 48    48    25      C        S        5      9.9    8.69
## 49    49    25      H        D        1     20.4   16.73
## 50    50    24      H        D        3     20.9   16.25

# Datos en consola -----
--

dbh <- c(16.5, 25.3, 22.1, 17.2, 16.1, 8.1, 34.3, 5.4, 5.7, 11.2, 24.1,
        14.5, 7.7, 15.6, 15.9, 10, 17.5, 20.5, 7.8, 27.3, 9.7, 6.5,
        23.4,
        8.2, 28.5, 10.4, 11.5, 14.3, 17.2, 16.8)

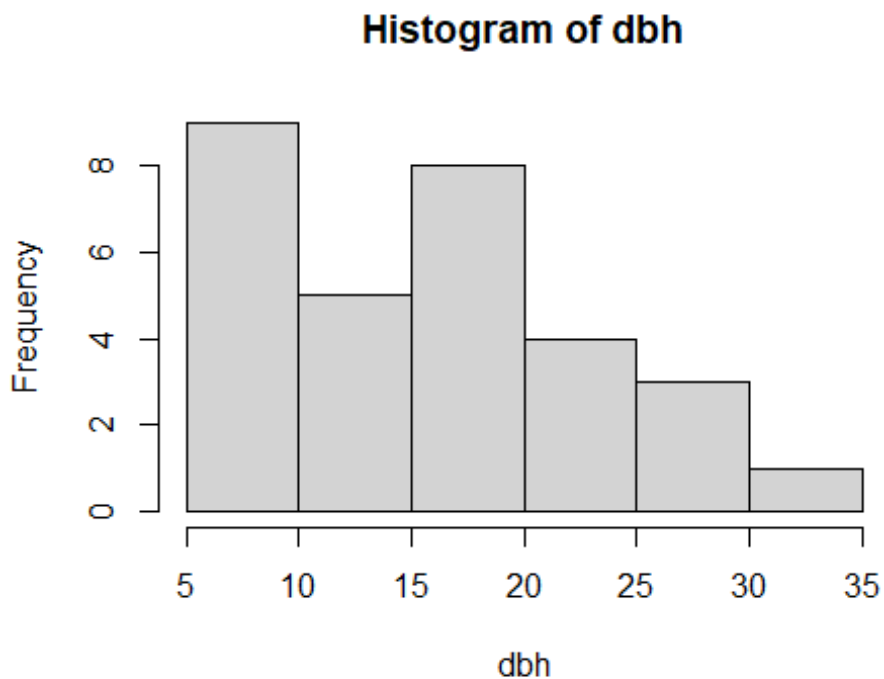
sum(dbh)

## [1] 469.3

prod(dbh)

## [1] 2.125828e+34
```

```
hist(dbh)
```



```
# Datos de URL -----
--

prof_url <-
"http://www.profepa.gob.mx/innovaportal/file/7635/1/accionesInspeccionfoa
np.csv"
profepa <- read.csv(prof_url)
head(profepa)

##           Entidad Inspecciones Recorridos.de.vigilancia Operativos
## 1 Aguascalientes           0                1            0
## 2 Baja California           0                1            0
## 3 Baja California Sur       0                0            0
## 4 Campeche                 0                0            0
## 5 Chiapas                  0                0            0
## 6 Chihuahua                3                1            1

profepa

##           Entidad Inspecciones Recorridos.de.vigilancia
Operativos
## 1 Aguascalientes           0                1
0
## 2 Baja California           0                1
0
```

|       |                     |   |   |
|-------|---------------------|---|---|
| ## 3  | Baja California Sur | 0 | 0 |
| 0     |                     |   |   |
| ## 4  | Campeche            | 0 | 0 |
| 0     |                     |   |   |
| ## 5  | Chiapas             | 0 | 0 |
| 0     |                     |   |   |
| ## 6  | Chihuahua           | 3 | 1 |
| 1     |                     |   |   |
| ## 7  | Coahuila            | 1 | 0 |
| 0     |                     |   |   |
| ## 8  | Colima              | 0 | 0 |
| 0     |                     |   |   |
| ## 9  | Distrito Federal    | 0 | 0 |
| 0     |                     |   |   |
| ## 10 | Durango             | 0 | 0 |
| 0     |                     |   |   |
| ## 11 | Guanajuato          | 0 | 0 |
| 0     |                     |   |   |
| ## 12 | Guerrero            | 0 | 0 |
| 0     |                     |   |   |
| ## 13 | Hidalgo             | 0 | 0 |
| 0     |                     |   |   |
| ## 14 | Jalisco             | 0 | 0 |
| 0     |                     |   |   |
| ## 15 | México              | 2 | 0 |
| 0     |                     |   |   |
| ## 16 | Michoacán           | 1 | 3 |
| 1     |                     |   |   |
| ## 17 | Morelos             | 2 | 0 |
| 1     |                     |   |   |
| ## 18 | Nayarit             | 0 | 1 |
| 0     |                     |   |   |
| ## 19 | Nuevo León          | 0 | 0 |
| 0     |                     |   |   |
| ## 20 | Oaxaca              | 0 | 0 |
| 0     |                     |   |   |
| ## 21 | Puebla              | 0 | 0 |
| 0     |                     |   |   |
| ## 22 | Querétaro           | 0 | 0 |
| 0     |                     |   |   |
| ## 23 | Quintana Roo        | 0 | 0 |
| 0     |                     |   |   |
| ## 24 | San Luis Potosí     | 0 | 0 |
| 0     |                     |   |   |
| ## 25 | Sinaloa             | 0 | 0 |
| 0     |                     |   |   |
| ## 26 | Sonora              | 0 | 0 |
| 0     |                     |   |   |
| ## 27 | Tabasco             | 0 | 0 |
| 0     |                     |   |   |

```
## 28          Tamaulipas          0          0
0
## 29          Tlaxcala           4          2
0
## 30          Veracruz           0          1
0
## 31          Yucatán            0          0
0
## 32          Zacatecas          0          1
0
## 33 Oficinas Centrales          6         10
0

sum(profepa$Inspecciones)

## [1] 19

sum(profepa$Operativos)

## [1] 3

sum(profepa$Recorridos.de.vigilancia)

## [1] 21

prof_url_2 <- paste0("http://www.profepa.gob.mx/innovaportal/",
                     "file/7635/1/accionesInspeccionfoanp.csv")
profepa2 <- read.csv(prof_url_2)
head(profepa2)

##          Entidad Inspecciones Recorridos.de.vigilancia Operativos
## 1    Aguascalientes          0              1              0
## 2    Baja California          0              1              0
## 3 Baja California Sur          0              0              0
## 4         Campeche           0              0              0
## 5         Chiapas            0              0              0
## 6    Chihuahua              3              1              1

# Importar datos URL seguros -----
--

library(repmis)

## Warning: package 'repmis' was built under R version 4.0.4

conjunto.2 <-
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
```

```

head(conjunto.2)

##   Arbol Fecha Especie Clase Vecinos Diametro Altura
## 1     1    12      F     C       4     15.3   14.78
## 2     2    12      F     D       3     17.8   17.07
## 3     3     9      C     D       5     18.2   18.28
## 4     4     9      H     S       4      9.7    8.79
## 5     5     7      H     I       6     10.8   10.18
## 6     6    10      C     I       3     14.1   14.90

sum(conjunto.2$Vecinos)

## [1] 167

library(readr)

## Warning: package 'readr' was built under R version 4.0.4

file <-
paste0("https://raw.githubusercontent.com/Caarloinee/PrincipiosEstadistic
a2021/main/Cuadro1.csv")
inventario <- read.csv(file)
head(inventario)

##   Arbol Fecha Especie Posicion Vecinos Diametro Altura
## 1     1    12      F         C       4     15.3   14.78
## 2     2    12      F         D       3     17.8   17.07
## 3     3     9      C         D       5     18.2   18.28
## 4     4     9      H         S       4      9.7    8.79
## 5     5     7      H         I       6     10.8   10.18
## 6     6    10      C         I       3     14.1   14.90

# Parte II Operaciones con La base de datos -----
--

dbh

## [1] 16.5 25.3 22.1 17.2 16.1  8.1 34.3  5.4  5.7 11.2 24.1 14.5  7.7
15.6 15.9
## [16] 10.0 17.5 20.5  7.8 27.3  9.7  6.5 23.4  8.2 28.5 10.4 11.5 14.3
17.2 16.8

mean(dbh)

## [1] 15.64333

dbh < 10

## [1] FALSE FALSE FALSE FALSE FALSE  TRUE FALSE  TRUE  TRUE FALSE FALSE
FALSE
## [13]  TRUE FALSE FALSE FALSE FALSE FALSE  TRUE FALSE  TRUE  TRUE FALSE
TRUE
## [25] FALSE FALSE FALSE FALSE FALSE FALSE

```

```

sum(dbh < 10)

## [1] 8

which(dbh < 10)

## [1] 6 8 9 13 19 21 22 24

dbh.url <-
"https://raw.githubusercontent.com/mgtagle/PrincipiosEstadistica2021/main/DBH\_1.csv"
parcelas <- read.csv(dbh.url)

tree.13 <- parcelas[!(parcelas$parcela == "2"),]
tree.23 <- parcelas[!(parcelas$parcela == "1"),]
tree.12 <- parcelas[!(parcelas$parcela == "3"),]

# Media de cada parcela dbh -----
--

mean(tree.13$dbh)

## [1] 15.42

mean(tree.23$dbh)

## [1] 15.37

mean(tree.12$dbh)

## [1] 16.14

# Selección de submuestras -----
--

tree_mean <- subset(parcelas, dbh <= mean(parcelas$dbh))
tree.up <- subset(parcelas, dbh >= mean(parcelas$dbh))
mean(tree_mean$dbh); mean(tree.up$dbh)

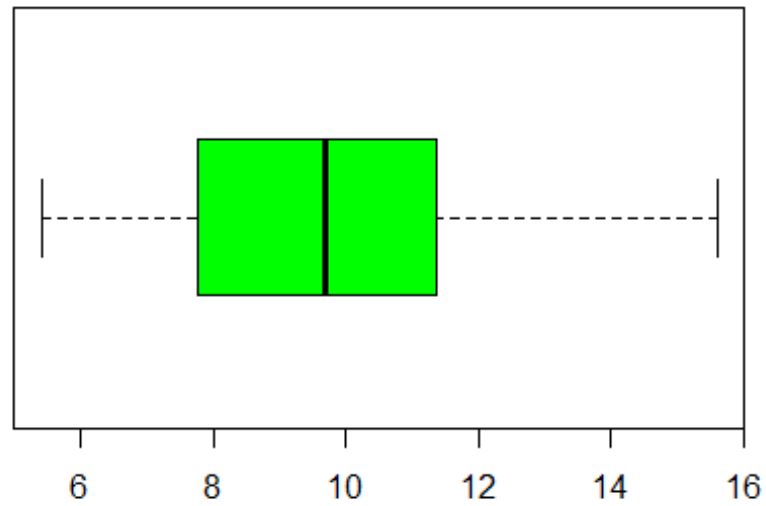
## [1] 9.773333
## [1] 21.51333

# Representación gráfica de subconjuntos -----
--

boxplot(tree_mean$dbh, main = "DBH <= media", col = "green", horizontal =
TRUE)

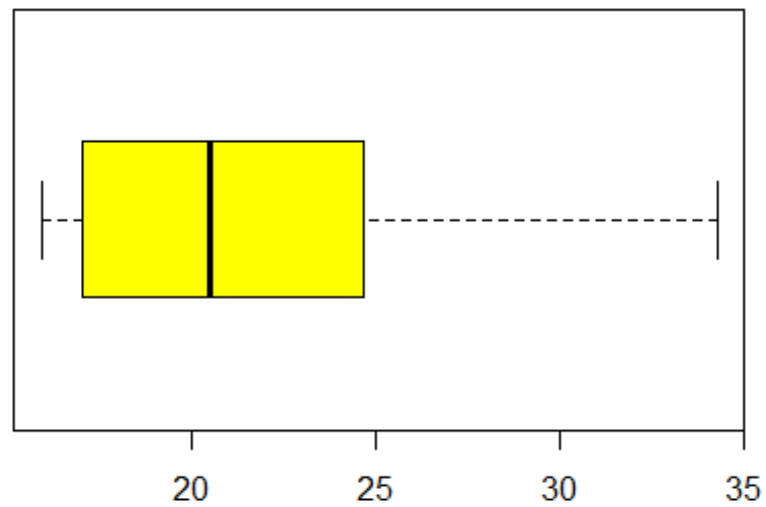
```

**DBH <= media**



```
boxplot(tree.up$dbh, main = "DBH <= media", col = "red", horizontal = TRUE)
```

**DBH <= media**



```
quantile(tree_mean$dbh, 0.5)
```

```

## 50%
## 9.7

quantile(tree_mean$dbh, 0.75)

## 75%
## 11.35

# Parte III Representación gráfica -----
--

mamiferos <- read.csv("https://www.openintro.org/data/csv/mammals.csv")
head(mamiferos)

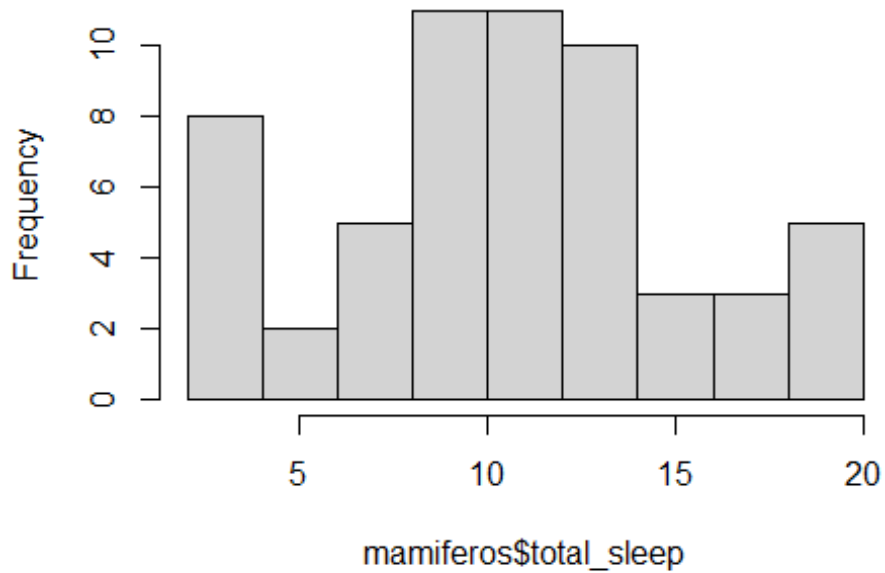
##           species  body_wt brain_wt non_dreaming dreaming
total_sleep
## 1   Africanelephant 6654.000   5712.0          NA         NA
3.3
## 2 Africangiantpouchedrat    1.000     6.6          6.3         2.0
8.3
## 3      ArcticFox    3.385    44.5          NA         NA
12.5
## 4  Arcticgroundsquirrel    0.920     5.7          NA         NA
16.5
## 5      Asianelephant 2547.000   4603.0          2.1         1.8
3.9
## 6      Baboon    10.550    179.5          9.1         0.7
9.8
##  life_span gestation predation exposure danger
## 1    38.6      645         3         5         3
## 2     4.5       42         3         1         3
## 3    14.0       60         1         1         1
## 4     NA       25         5         2         3
## 5    69.0      624         3         5         4
## 6    27.0      180         4         4         4

hist(mamiferos$total_sleep)

```

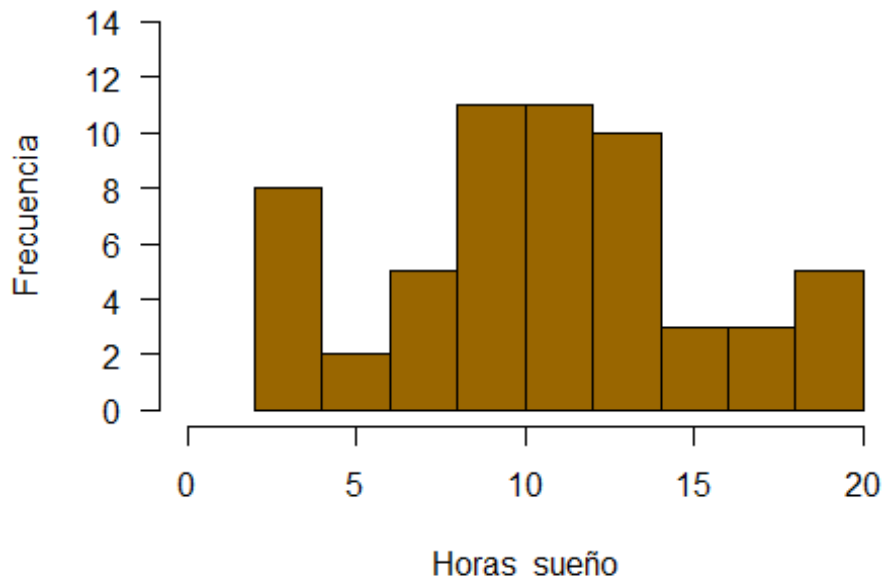


**Histogram of mamiferos\$total\_sleep**



```
hist(mamiferos$total_sleep,  
xlim = c(0,20), ylim = c(0,14),  
main = "Total de horas sueño de las 39 especies", # Cambiar el título  
xlab = "Horas sueño", # Cambiar eje de las x  
ylab = "Frecuencia", # Cambiar eje de las y  
las = 1, # Cambiar orientación de y  
col = "#996600") # Cambiar color de las barras
```

## Total de horas sueño de las 39 especies



```
fivenum(mamiferos$brain_wt)

## [1]    0.14    4.00   17.25  169.00 5712.00

# Barplot o gráfico de barra -----
--

data("chickwts")
head(chickwts[c(1:2,42:43, 62:64), ])

##   weight    feed
## 1    179 horsebean
## 2    160 horsebean
## 42   226  sunflower
## 43   320  sunflower
## 62   379   casein
## 63   260   casein

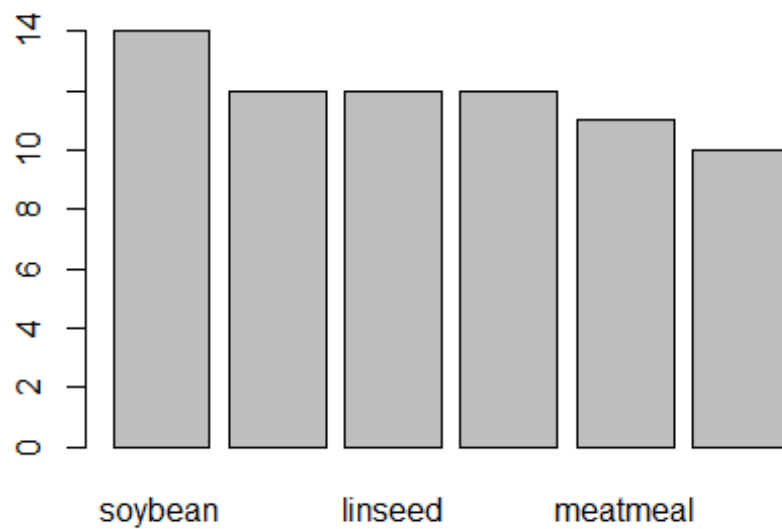
feeds <- table(chickwts$feed)
feeds

##
##   casein horsebean  linseed  meatmeal  soybean  sunflower
##      12       10      12       11      14       12

barplot(feeds)
```



```
barplot(feeds[order(feeds, decreasing = TRUE)])
```



```
barplot(feeds[order(feeds)], horiz = TRUE, las = 1, col = "blue", main = "Frecuencia por tipo de alimentaci3n", xlab = "Pollos")
```

### Frecuencia por tipo de alimentnación

