

## Actividad 2: Manejo de RStudio

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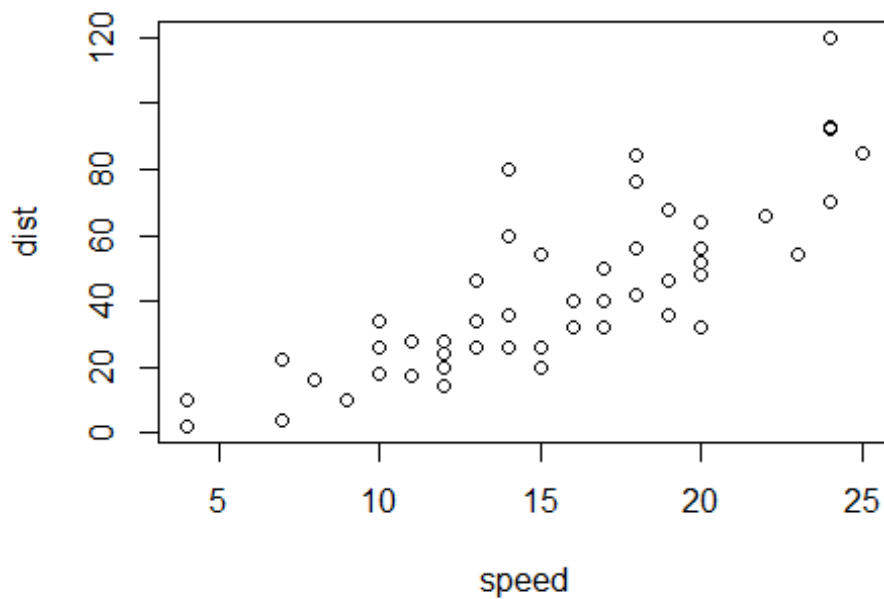
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Primera clase

```
plot(cars)
```



```
#install.packages("ggplot2")
```

```
library(ggplot2)
```

```
x<-35
```

```
y<-10
```

```
x
```

```
## [1] 35
```

```
y
```

```
## [1] 10
x*y
## [1] 350
x/y
## [1] 3.5
x>y #mayor que
## [1] TRUE
x<y #menor que
## [1] FALSE
x>=35 #mayor o igual que
## [1] TRUE
y==10 #igual
## [1] TRUE
x!=10 #diferente
## [1] TRUE
#formas de asignación de variables
v1<-100 ; v1
## [1] 100
v2<<-200 ; v2
## [1] 200
v3=300 ; v3
## [1] 300
1000->v4 ; v4
## [1] 1000
v5<-(1:10) ; v5
## [1] 1 2 3 4 5 6 7 8 9 10
v6 <- c(1:10) ; v6
## [1] 1 2 3 4 5 6 7 8 9 10
```

```

#matriz (vector numérico, row, column)
m1<- matrix(1:9,3,3) ; m1

##      [,1] [,2] [,3]
## [1,]    1    4    7
## [2,]    2    5    8
## [3,]    3    6    9

print("hola mundo")

## [1] "hola mundo"

v10<- c("IBT","BIOTECNOLOGIA","INGENIERIA") ; v10 #concatenación

## [1] "IBT"          "BIOTECNOLOGIA" "INGENIERIA"

v11<- (1:3)/3 ; v11

## [1] 0.3333333 0.6666667 1.0000000

v12<- (v11*3); v12

## [1] 1 2 3

v13<- v12/v12; v13

## [1] 1 1 1

x<-(1:3)
y<-c("IBT","BIOTECNOLOGIA","INGENIERIA")
z<-c(TRUE, TRUE, FALSE)
m2<-data.frame(x,y,z) ; m2 #data frame para almacenamiento de datos
heterogéneos

##   x      y      z
## 1 1      IBT  TRUE
## 2 2 BIOTECNOLOGIA  TRUE
## 3 3  INGENIERIA  FALSE

head(iris)

##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1          5.1         3.5          1.4          0.2  setosa
## 2          4.9         3.0          1.4          0.2  setosa
## 3          4.7         3.2          1.3          0.2  setosa
## 4          4.6         3.1          1.5          0.2  setosa
## 5          5.0         3.6          1.4          0.2  setosa
## 6          5.4         3.9          1.7          0.4  setosa

tail(iris)

##   Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
## 145          6.7         3.3          5.7          2.5 virginica
## 146          6.7         3.0          5.2          2.3 virginica

```

```
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1      1.8 virginica
```

```
x<-iris ; x #asignación
```

```
##      Sepal.Length Sepal.Width Petal.Length Petal.Width  Species
## 1      5.1      3.5      1.4      0.2    setosa
## 2      4.9      3.0      1.4      0.2    setosa
## 3      4.7      3.2      1.3      0.2    setosa
## 4      4.6      3.1      1.5      0.2    setosa
## 5      5.0      3.6      1.4      0.2    setosa
## 6      5.4      3.9      1.7      0.4    setosa
## 7      4.6      3.4      1.4      0.3    setosa
## 8      5.0      3.4      1.5      0.2    setosa
## 9      4.4      2.9      1.4      0.2    setosa
## 10     4.9      3.1      1.5      0.1    setosa
## 11     5.4      3.7      1.5      0.2    setosa
## 12     4.8      3.4      1.6      0.2    setosa
## 13     4.8      3.0      1.4      0.1    setosa
## 14     4.3      3.0      1.1      0.1    setosa
## 15     5.8      4.0      1.2      0.2    setosa
## 16     5.7      4.4      1.5      0.4    setosa
## 17     5.4      3.9      1.3      0.4    setosa
## 18     5.1      3.5      1.4      0.3    setosa
## 19     5.7      3.8      1.7      0.3    setosa
## 20     5.1      3.8      1.5      0.3    setosa
## 21     5.4      3.4      1.7      0.2    setosa
## 22     5.1      3.7      1.5      0.4    setosa
## 23     4.6      3.6      1.0      0.2    setosa
## 24     5.1      3.3      1.7      0.5    setosa
## 25     4.8      3.4      1.9      0.2    setosa
## 26     5.0      3.0      1.6      0.2    setosa
## 27     5.0      3.4      1.6      0.4    setosa
## 28     5.2      3.5      1.5      0.2    setosa
## 29     5.2      3.4      1.4      0.2    setosa
## 30     4.7      3.2      1.6      0.2    setosa
## 31     4.8      3.1      1.6      0.2    setosa
## 32     5.4      3.4      1.5      0.4    setosa
## 33     5.2      4.1      1.5      0.1    setosa
## 34     5.5      4.2      1.4      0.2    setosa
## 35     4.9      3.1      1.5      0.2    setosa
## 36     5.0      3.2      1.2      0.2    setosa
## 37     5.5      3.5      1.3      0.2    setosa
## 38     4.9      3.6      1.4      0.1    setosa
## 39     4.4      3.0      1.3      0.2    setosa
## 40     5.1      3.4      1.5      0.2    setosa
## 41     5.0      3.5      1.3      0.3    setosa
## 42     4.5      2.3      1.3      0.3    setosa
```

## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor
## 64	6.1	2.9	4.7	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 66	6.7	3.1	4.4	1.4	versicolor
## 67	5.6	3.0	4.5	1.5	versicolor
## 68	5.8	2.7	4.1	1.0	versicolor
## 69	6.2	2.2	4.5	1.5	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 71	5.9	3.2	4.8	1.8	versicolor
## 72	6.1	2.8	4.0	1.3	versicolor
## 73	6.3	2.5	4.9	1.5	versicolor
## 74	6.1	2.8	4.7	1.2	versicolor
## 75	6.4	2.9	4.3	1.3	versicolor
## 76	6.6	3.0	4.4	1.4	versicolor
## 77	6.8	2.8	4.8	1.4	versicolor
## 78	6.7	3.0	5.0	1.7	versicolor
## 79	6.0	2.9	4.5	1.5	versicolor
## 80	5.7	2.6	3.5	1.0	versicolor
## 81	5.5	2.4	3.8	1.1	versicolor
## 82	5.5	2.4	3.7	1.0	versicolor
## 83	5.8	2.7	3.9	1.2	versicolor
## 84	6.0	2.7	5.1	1.6	versicolor
## 85	5.4	3.0	4.5	1.5	versicolor
## 86	6.0	3.4	4.5	1.6	versicolor
## 87	6.7	3.1	4.7	1.5	versicolor
## 88	6.3	2.3	4.4	1.3	versicolor
## 89	5.6	3.0	4.1	1.3	versicolor
## 90	5.5	2.5	4.0	1.3	versicolor
## 91	5.5	2.6	4.4	1.2	versicolor
## 92	6.1	3.0	4.6	1.4	versicolor

## 93	5.8	2.6	4.0	1.2	versicolor
## 94	5.0	2.3	3.3	1.0	versicolor
## 95	5.6	2.7	4.2	1.3	versicolor
## 96	5.7	3.0	4.2	1.2	versicolor
## 97	5.7	2.9	4.2	1.3	versicolor
## 98	6.2	2.9	4.3	1.3	versicolor
## 99	5.1	2.5	3.0	1.1	versicolor
## 100	5.7	2.8	4.1	1.3	versicolor
## 101	6.3	3.3	6.0	2.5	virginica
## 102	5.8	2.7	5.1	1.9	virginica
## 103	7.1	3.0	5.9	2.1	virginica
## 104	6.3	2.9	5.6	1.8	virginica
## 105	6.5	3.0	5.8	2.2	virginica
## 106	7.6	3.0	6.6	2.1	virginica
## 107	4.9	2.5	4.5	1.7	virginica
## 108	7.3	2.9	6.3	1.8	virginica
## 109	6.7	2.5	5.8	1.8	virginica
## 110	7.2	3.6	6.1	2.5	virginica
## 111	6.5	3.2	5.1	2.0	virginica
## 112	6.4	2.7	5.3	1.9	virginica
## 113	6.8	3.0	5.5	2.1	virginica
## 114	5.7	2.5	5.0	2.0	virginica
## 115	5.8	2.8	5.1	2.4	virginica
## 116	6.4	3.2	5.3	2.3	virginica
## 117	6.5	3.0	5.5	1.8	virginica
## 118	7.7	3.8	6.7	2.2	virginica
## 119	7.7	2.6	6.9	2.3	virginica
## 120	6.0	2.2	5.0	1.5	virginica
## 121	6.9	3.2	5.7	2.3	virginica
## 122	5.6	2.8	4.9	2.0	virginica
## 123	7.7	2.8	6.7	2.0	virginica
## 124	6.3	2.7	4.9	1.8	virginica
## 125	6.7	3.3	5.7	2.1	virginica
## 126	7.2	3.2	6.0	1.8	virginica
## 127	6.2	2.8	4.8	1.8	virginica
## 128	6.1	3.0	4.9	1.8	virginica
## 129	6.4	2.8	5.6	2.1	virginica
## 130	7.2	3.0	5.8	1.6	virginica
## 131	7.4	2.8	6.1	1.9	virginica
## 132	7.9	3.8	6.4	2.0	virginica
## 133	6.4	2.8	5.6	2.2	virginica
## 134	6.3	2.8	5.1	1.5	virginica
## 135	6.1	2.6	5.6	1.4	virginica
## 136	7.7	3.0	6.1	2.3	virginica
## 137	6.3	3.4	5.6	2.4	virginica
## 138	6.4	3.1	5.5	1.8	virginica
## 139	6.0	3.0	4.8	1.8	virginica
## 140	6.9	3.1	5.4	2.1	virginica
## 141	6.7	3.1	5.6	2.4	virginica
## 142	6.9	3.1	5.1	2.3	virginica

## 143	5.8	2.7	5.1	1.9	virginica
## 144	6.8	3.2	5.9	2.3	virginica
## 145	6.7	3.3	5.7	2.5	virginica
## 146	6.7	3.0	5.2	2.3	virginica
## 147	6.3	2.5	5.0	1.9	virginica
## 148	6.5	3.0	5.2	2.0	virginica
## 149	6.2	3.4	5.4	2.3	virginica
## 150	5.9	3.0	5.1	1.8	virginica

```
y<-data.frame(a=iris$Petal.Length, b=iris$Species); y
```

##	a	b
## 1	1.4	setosa
## 2	1.4	setosa
## 3	1.3	setosa
## 4	1.5	setosa
## 5	1.4	setosa
## 6	1.7	setosa
## 7	1.4	setosa
## 8	1.5	setosa
## 9	1.4	setosa
## 10	1.5	setosa
## 11	1.5	setosa
## 12	1.6	setosa
## 13	1.4	setosa
## 14	1.1	setosa
## 15	1.2	setosa
## 16	1.5	setosa
## 17	1.3	setosa
## 18	1.4	setosa
## 19	1.7	setosa
## 20	1.5	setosa
## 21	1.7	setosa
## 22	1.5	setosa
## 23	1.0	setosa
## 24	1.7	setosa
## 25	1.9	setosa
## 26	1.6	setosa
## 27	1.6	setosa
## 28	1.5	setosa
## 29	1.4	setosa
## 30	1.6	setosa
## 31	1.6	setosa
## 32	1.5	setosa
## 33	1.5	setosa
## 34	1.4	setosa
## 35	1.5	setosa
## 36	1.2	setosa
## 37	1.3	setosa
## 38	1.4	setosa

```
## 39 1.3      setosa
## 40 1.5      setosa
## 41 1.3      setosa
## 42 1.3      setosa
## 43 1.3      setosa
## 44 1.6      setosa
## 45 1.9      setosa
## 46 1.4      setosa
## 47 1.6      setosa
## 48 1.4      setosa
## 49 1.5      setosa
## 50 1.4      setosa
## 51 4.7 versicolor
## 52 4.5 versicolor
## 53 4.9 versicolor
## 54 4.0 versicolor
## 55 4.6 versicolor
## 56 4.5 versicolor
## 57 4.7 versicolor
## 58 3.3 versicolor
## 59 4.6 versicolor
## 60 3.9 versicolor
## 61 3.5 versicolor
## 62 4.2 versicolor
## 63 4.0 versicolor
## 64 4.7 versicolor
## 65 3.6 versicolor
## 66 4.4 versicolor
## 67 4.5 versicolor
## 68 4.1 versicolor
## 69 4.5 versicolor
## 70 3.9 versicolor
## 71 4.8 versicolor
## 72 4.0 versicolor
## 73 4.9 versicolor
## 74 4.7 versicolor
## 75 4.3 versicolor
## 76 4.4 versicolor
## 77 4.8 versicolor
## 78 5.0 versicolor
## 79 4.5 versicolor
## 80 3.5 versicolor
## 81 3.8 versicolor
## 82 3.7 versicolor
## 83 3.9 versicolor
## 84 5.1 versicolor
## 85 4.5 versicolor
## 86 4.5 versicolor
## 87 4.7 versicolor
## 88 4.4 versicolor
```



```
## 89 4.1 versicolor
## 90 4.0 versicolor
## 91 4.4 versicolor
## 92 4.6 versicolor
## 93 4.0 versicolor
## 94 3.3 versicolor
## 95 4.2 versicolor
## 96 4.2 versicolor
## 97 4.2 versicolor
## 98 4.3 versicolor
## 99 3.0 versicolor
## 100 4.1 versicolor
## 101 6.0 virginica
## 102 5.1 virginica
## 103 5.9 virginica
## 104 5.6 virginica
## 105 5.8 virginica
## 106 6.6 virginica
## 107 4.5 virginica
## 108 6.3 virginica
## 109 5.8 virginica
## 110 6.1 virginica
## 111 5.1 virginica
## 112 5.3 virginica
## 113 5.5 virginica
## 114 5.0 virginica
## 115 5.1 virginica
## 116 5.3 virginica
## 117 5.5 virginica
## 118 6.7 virginica
## 119 6.9 virginica
## 120 5.0 virginica
## 121 5.7 virginica
## 122 4.9 virginica
## 123 6.7 virginica
## 124 4.9 virginica
## 125 5.7 virginica
## 126 6.0 virginica
## 127 4.8 virginica
## 128 4.9 virginica
## 129 5.6 virginica
## 130 5.8 virginica
## 131 6.1 virginica
## 132 6.4 virginica
## 133 5.6 virginica
## 134 5.1 virginica
## 135 5.6 virginica
## 136 6.1 virginica
## 137 5.6 virginica
## 138 5.5 virginica
```

```
## 139 4.8 virginica
## 140 5.4 virginica
## 141 5.6 virginica
## 142 5.1 virginica
## 143 5.1 virginica
## 144 5.9 virginica
## 145 5.7 virginica
## 146 5.2 virginica
## 147 5.0 virginica
## 148 5.2 virginica
## 149 5.4 virginica
## 150 5.1 virginica

summary(iris)

##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
## Min.      :4.300    Min.      :2.000    Min.      :1.000    Min.      :0.100
## 1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
## Median :5.800    Median :3.000    Median :4.350    Median :1.300
## Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
## 3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
## Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500
##           Species
## setosa      :50
## versicolor:50
## virginica   :50
##
##
##

getwd()

## [1] "C:/Users/alejcd/Documents"

setwd("C:/Users/alejcd/Desktop/datasets")
```

If else

```
x = 2 #valor determinado
if (x>1 & x<7){
  print("x is between 1 and 7")
  #Si hay un valor entre los parametros establecidos se imprime la frase
}else if (x>8 & x<15){
  print("x is between 8 and 15 =D")
}

## [1] "x is between 1 and 7"
```

Como el valor es 2, se imprimio la primera frase. So hubiera sido mayor a 7, hubiera sido la segunda frase.

For

```

#Obtener el mismo valor con diferentes comandos
x = c(1,2,3,4,5) #vector con valores numéricos
y <- c(1:5)
z <- 1:5

#Llamamos variables
x

## [1] 1 2 3 4 5

y

## [1] 1 2 3 4 5

z

## [1] 1 2 3 4 5

for (i in 1:8){
  print(x[i]) #Como no hat valores después de 5, se asignan como NA
}

## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
## [1] NA
## [1] NA
## [1] NA

```

While

```

x= 2.987 #Valor asignado de x

while(x <= 4.987){
  x = x + 0.987 #Mientras x sea menor o igual a 4.987 se suman 0.987
  print(c(x,x-2,x-1)) #3 filas en resultados, una con x normal, otra con
x menos 2
  #La última es de x menos 1
}

## [1] 3.974 1.974 2.974
## [1] 4.961 2.961 3.961
## [1] 5.948 3.948 4.948

```

Break statement

```

x = 1:10
for (i in x){
  if (i ==8){
    break #Cuando i sea igual a 8 termina el operador

```

```
}  
print(i)  
}
```

```
## [1] 1  
## [1] 2  
## [1] 3  
## [1] 4  
## [1] 5  
## [1] 6  
## [1] 7
```

Next statement

```
x = 1:10  
for (i in x){  
  if (i == 8){  
    next} #Hay un skip cuando el valor es 8, es decir no se imprimira  
    print(i)  
}
```

```
## [1] 1  
## [1] 2  
## [1] 3  
## [1] 4  
## [1] 5  
## [1] 6  
## [1] 7  
## [1] 9  
## [1] 10
```

Instrucciones básicas

```
#instalación  
##install.packages("ggplot2")  
##install.packages("reshape2")  
##install.packages("readxl")  
##install.packages("scatterplot3d")  
  
#importar bibliotecas  
  
library("ggplot2") #para las gráficas.2  
library("reshape2") #para barras por grupos  
library("readxl") #para importar archivos XLSX  
library("scatterplot3d") #para scatterplot en 3D
```

Importar datos

Arrchivo CSV (Pacientes de nuevo ingreso)

```
#archivo CSV  
setwd("C:/Users/alejcd/Desktop/datasets")
```

```
mydata_csv <- read.csv("9_PACIENTES_DE_NUEVO_INGRESO.csv")
head(mydata_csv)
```

```
##      FOLIO EDAD      SEXO      ESTADO      MUNICIPIO
## 1      1    80  Femenino    GUERRERO HUITZUCO DE LOS FIGUEROA
## 2      2    71 Masculino DISTRITO FEDERAL      TLALPAN
## 3      3    45 Masculino      MORELOS      CUAUTLA
## 4      4    33 Masculino      MEXICO      ZUMPANGO
## 5      5    46 Femenino      MEXICO    NAUCALPAN DE JUAREZ
## 6      6    86 Masculino DISTRITO FEDERAL      TLALPAN
##      DESCRIPCION.DIAGNOSTICO
## 1      TUMOR MALIGNO DE LA MAMA
## 2      TUMOR MALIGNO DE LA PROSTATA
## 3      TUMOR MALIGNO DE LA PROSTATA
## 4 OTROS TUMORES MALIGNOS DE LA PIEL
## 5      TUMOR MALIGNO DE LA MAMA
## 6      TUMOR MALIGNO DEL RECTO
```

Archivo de texto (CC)

```
#archivo TXT
setwd("C:/Users/alejc/Desktop/datasets")
mydata_txt <- read.delim("cc.txt", header =FALSE)
head(mydata_txt)
```

```
##      V1      V2      V3      V4
## 1    20 0.121 0.093 0.077
## 2    50 0.283 0.252 0.162
## 3    70 0.430 0.413 0.384
## 4    80 0.455 0.488 0.487
## 5   100 0.539 0.521 0.519
## 6   130 0.705 0.697 0.721
```

Archivo de excel (Cons energía)

```
#archivo XLSX
setwd("C:/Users/alejc/Desktop/datasets")
mydata_xlsx <- read_excel("Cons_Energia.xlsx")
head(mydata_xlsx)
```

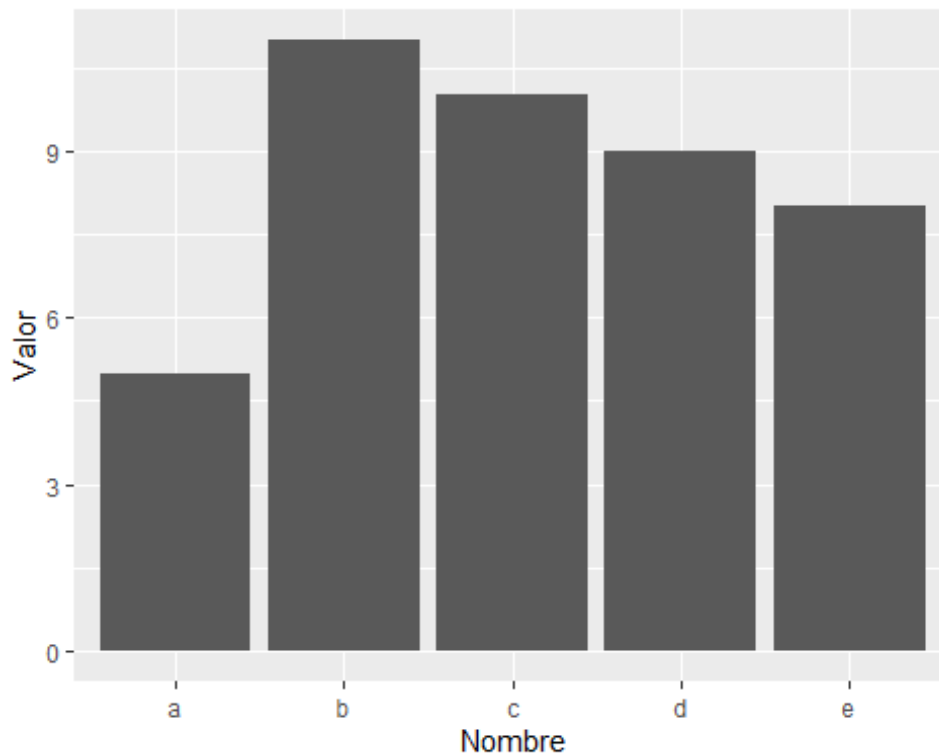
```
## # A tibble: 6 × 3
##   Cons_Energ atip_sup Atip_Sup_Inf
##   <dbl>     <dbl>     <dbl>
## 1     2.97     2.97         2
## 2     6.8     6.8         6.8
## 3     7.73     7.73         7.73
## 4     8.61     8.61         8.61
## 5     9.6     9.6         9.6
## 6    10.3    10.3        10.3
```

Gráficos con ggplot2

## Gráfica de barras

```
####Barras
#Creamos un dataframe
data <- data.frame(
  Nombre = letters[1:5], #variable dependiente
  Valor = sample(seq(4,15),5), #variable independiente RANDOM
  sd = c(1,0.2,3,2,4) #error prediseñado
)

#Solo barras, diseño simple -----
ggplot(data, aes(Nombre,Valor)) + geom_bar(stat = "identity")
```



## Barras con error

```
#Barras de error -----
b <- ggplot(data) +
  geom_bar(aes(x = Nombre, y = Valor), stat = "identity", fill= "red",
  alpha = 0.7) + #barras
  geom_errorbar(aes(x=Nombre, ymin=Valor-sd, ymax=Valor+sd), width=0.4,
  colour= "brown",
  alpha=0.9, size=1.3) #margen de error

## Warning: Using `size` aesthetic for lines was deprecated in ggplot2
3.4.0.
## i Please use `linewidth` instead.
```

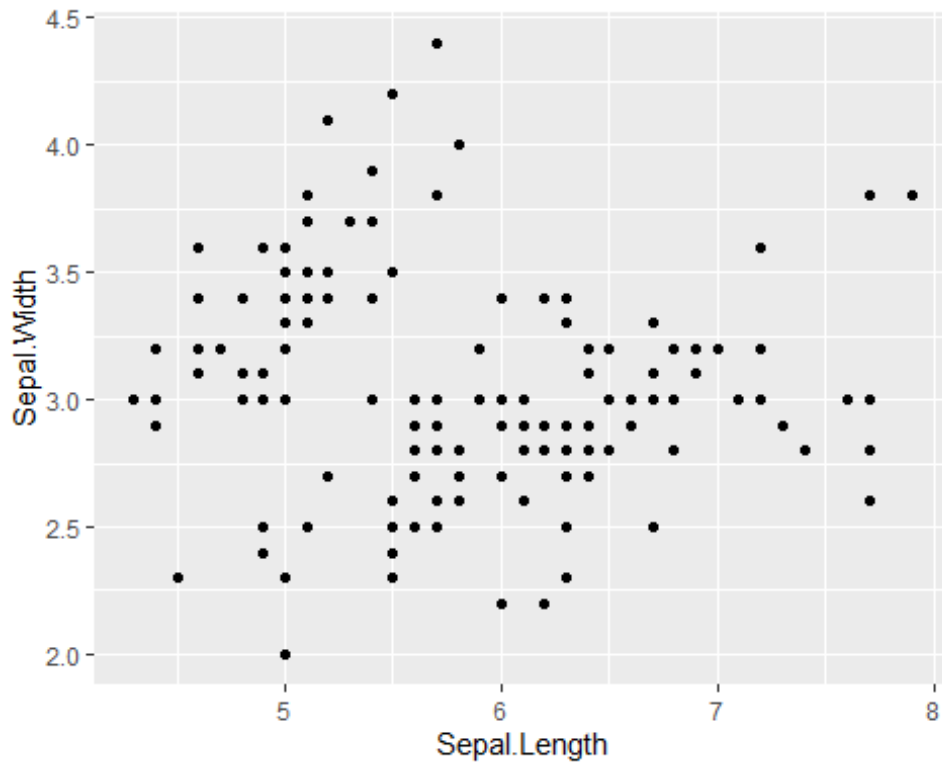
```
b + ggtitle("Bar plot with errorbars") + xlab("Dependent variable")+
ylab("Independent variable")
```



*#texto añadido al gráfico*

Scatterplots Simple

```
####Scatterplots -----
#simple
ggplot(iris,aes(x = Sepal.Length, y = Sepal.Width)) +
  geom_point()
```



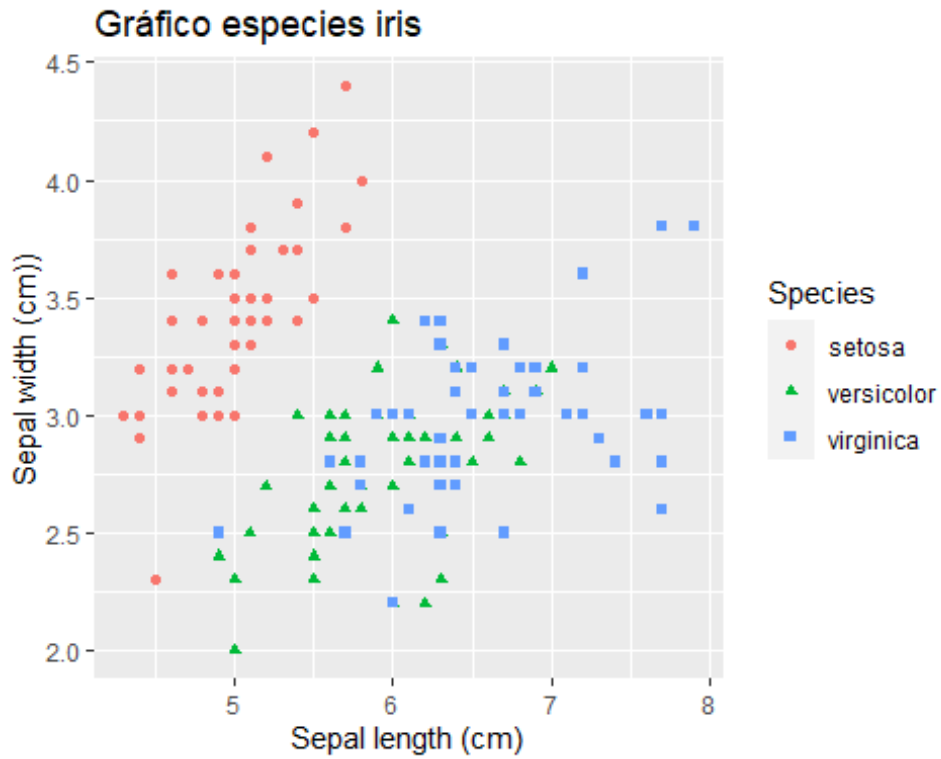
Scatterplot

más complejo que simple Usando set de datos iris

```
#"Complejo"
c <- ggplot(data = iris, aes(x=Sepal.Length, y = Sepal.Width))+
#Asignamos valores
  geom_point(aes(color=Species, shape=Species))

c + ggtitle ("Gráfico especies iris") + xlab("Sepal length (cm)") +
  ylab("Sepal width (cm)")
```



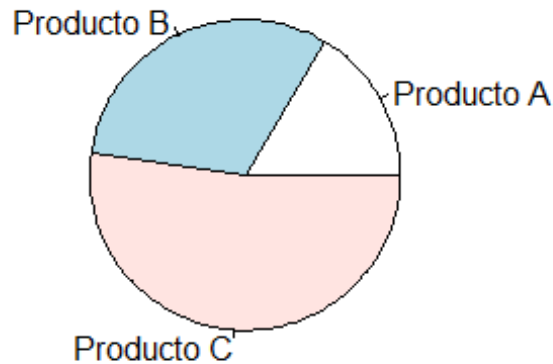


*#Añadimos texto al gráfico*

Gráfico PIE

```
#Pie -----
x <- c(11,21,35) #Hacemos una compilación
lbls <- c("Producto A", "Producto B", "Producto C") #Nombramos Labels
pie(x, labels = lbls, main = "Pie chart of products") #Realizamos gráfico
```

## Pie chart of products



De barras a PIE

```
#Otro ----- (No entendí bien, pero sale un PIE y tarda en cargar)
df = data.frame("Products" = c("Product A", "Product B", "Product C"),
"porciento" = c(11,21,35))

#Basic bar
pC = ggplot(df, aes(x= "", y = porciento, fill = Products)) +
geom_bar(stat = "identity", width = 1, alpha =0.7)

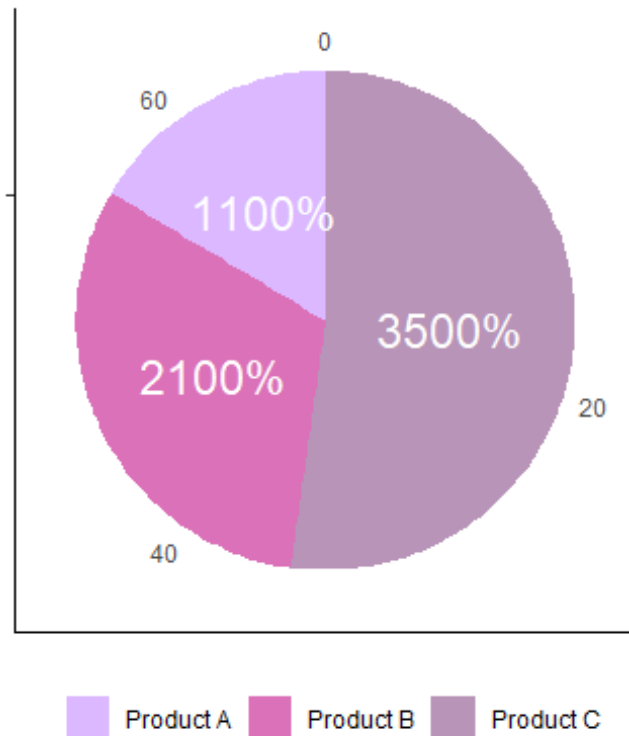
#Conversión a pie
pC = pC+coord_polar("y", start = 0) + geom_text(aes(label =
paste0(round(porciento*100),
"%")),
position = position_stack(
vjust = 0.5), size = 6, color = "white")
#Añade color
pC = pC + scale_fill_manual(values=c("#CC99FF", "#CC3399", "#996699"))
#Proporcionamos colores de manera manual

#Remueve labels + titulo
pC = pC + labs (x= NULL, y = NULL, fill = NULL)

#Limpieza final
pC = pC + theme_classic()
```

```
pC <- pC + theme(legend.position = "bottom")
```

```
pC #Llamamos al gráfico final
```



Scatterplots en 3D, usando dataset iris

```
#Scatterplots-----
```

```
#3D instalación
```

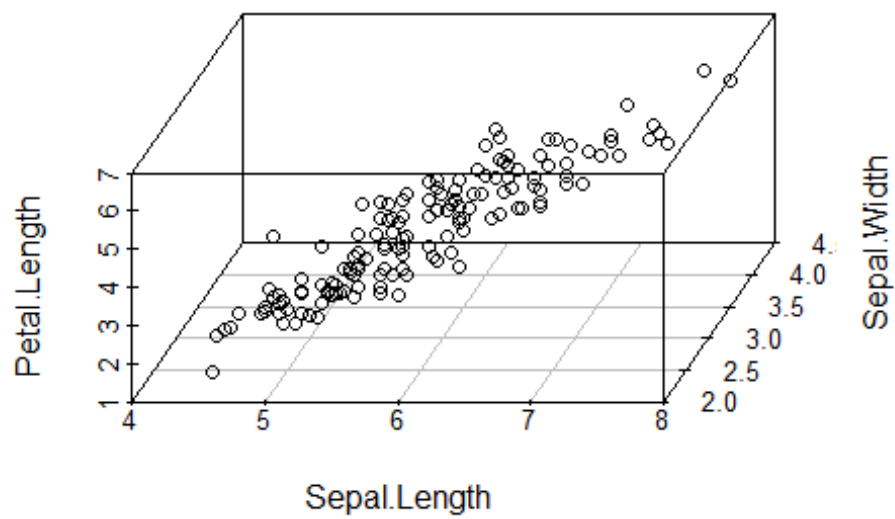
```
install.packages("scatterplot3d")
```

```
## Warning: package 'scatterplot3d' is in use and will not be installed
```

```
library("scatterplot3d")
```

```
#Gráfico simple
```

```
scatterplot3d(iris[,1:3], angle = 75)
```

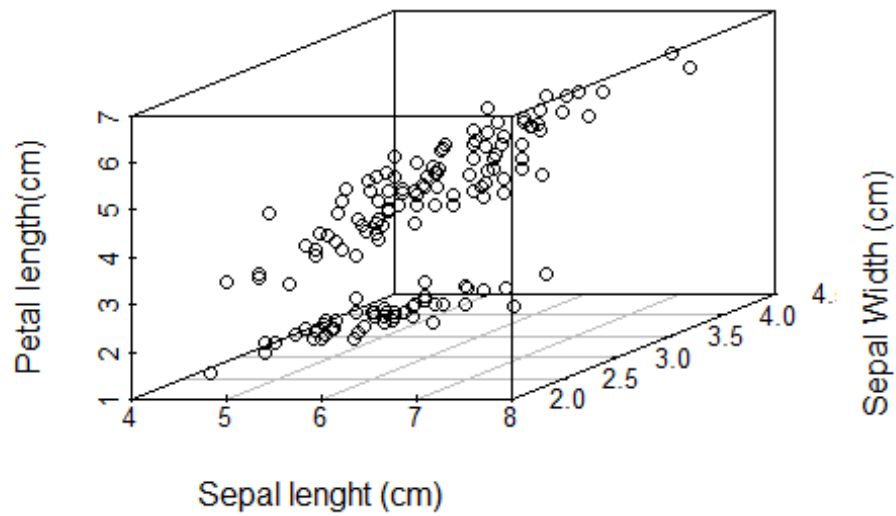


Scatterplot 3D

con ejes titulados

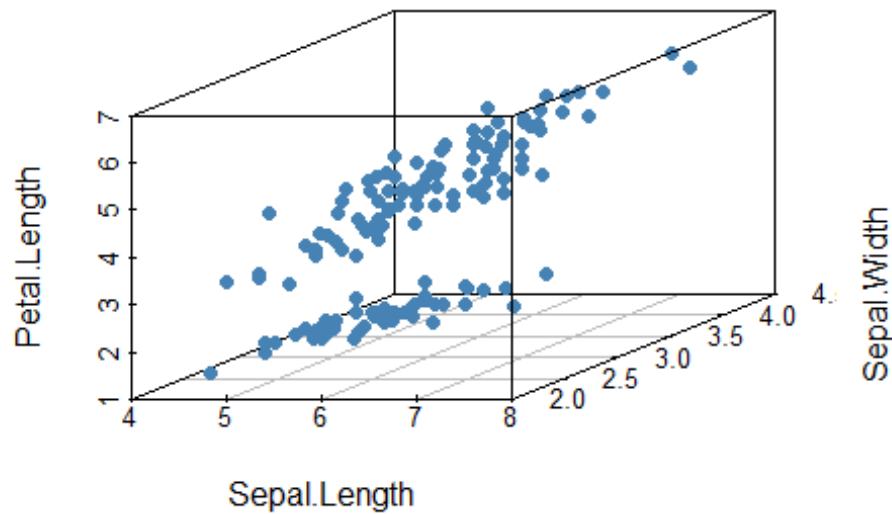
```
#Especificaciones para cada label  
scatterplot3d(iris[,1:3],  
              main = "3D scatterplot",  
              xlab = "Sepal lenght (cm)",  
              ylab = "Sepal Width (cm)",  
              zlab = "Petal length(cm)")
```

### 3D scatterplot



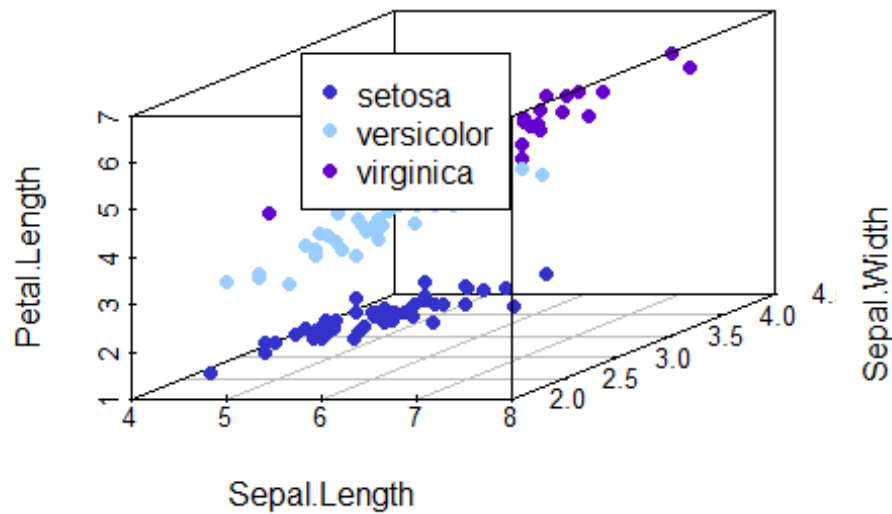
Cambiamos color de marcadores

```
#Cambiar color marcadores  
scatterplot3d(iris[,1:3], pch = 16, color = "steelblue") #Asignamos color azul
```



Asignamos colores específicos dependiendo de la especie

```
#Especies c/ color
colors <- c("#3333CC", "#99CCFF", "#6600CC") #Asignamos colores
colors <- colors[as.numeric(iris$Species)] #Colores asignados por especie
s3d <- scatterplot3d(iris[,1:3], pch = 16, color = colors)
#Especificaciones gráfico
legend(s3d$xyz.convert(6.3, 4.5), legend = levels(iris$Species), col = c(
  "#3333CC", "#99CCFF", "#6600CC"), pch = 16) #Para Leyendas
```



Gráficos: heatmaps

```
#Heatmaps
library("gplots")

##
## Attaching package: 'gplots'

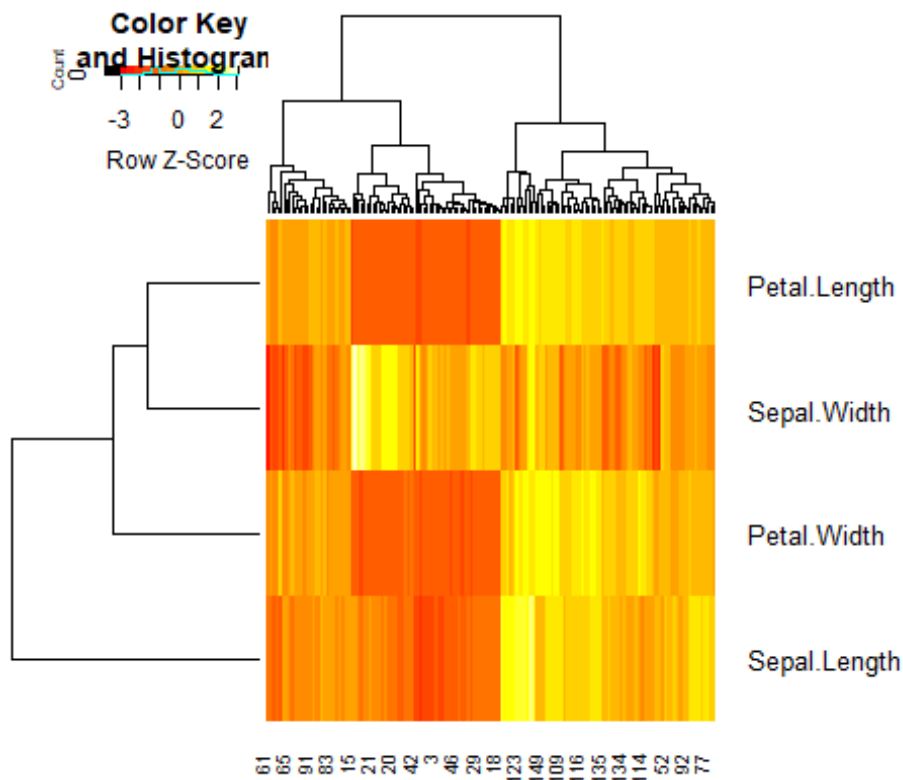
## The following object is masked from 'package:stats':
##
##     lowess

heatmap.2(t(iris[,1:4]), trace = "none", scale = "row", key = TRUE, mar=
c(2,8),
          cexRow = 1, ColsideColors = c("orange", "purple",
"magenta")[iris$Species])

## Warning in plot.window(...): "ColsideColors" is not a graphical
parameter

## Warning in plot.xy(xy, type, ...): "ColsideColors" is not a graphical
parameter

## Warning in title(...): "ColsideColors" is not a graphical parameter
```



*#Heatmap tiene sus propios colores asignados, por los que se mantienen los del preset*  
*#aunque asignemos unos en colsidecolors*

Detalles de heatmaps No encuentre ggarrange !!

Chunk no funciona

```
#Parametros extra para diseño
#library("ggarrange")
#tiff("ABI.tiff", width = 4000, height = 4000, units = 'px', res=300,
#compression = 'lzw')
#ggarrange(pC, b, box, labels = c("A", "B", "C"))
#dev.off()
```

Gráficos: boxplot

```
#Boxplot
par(mfrow=c(2,2)) #subplot

#Asignamos valores a comparar de cada caja
#Comparación 1
boxplot(Sepal.Length~Species, data = iris,
        xlab = "Species", ylab = "Sepal Length", main = "Sepal Length
comp")

#Comparación 2
boxplot(Sepal.Width~Species, data = iris,
```



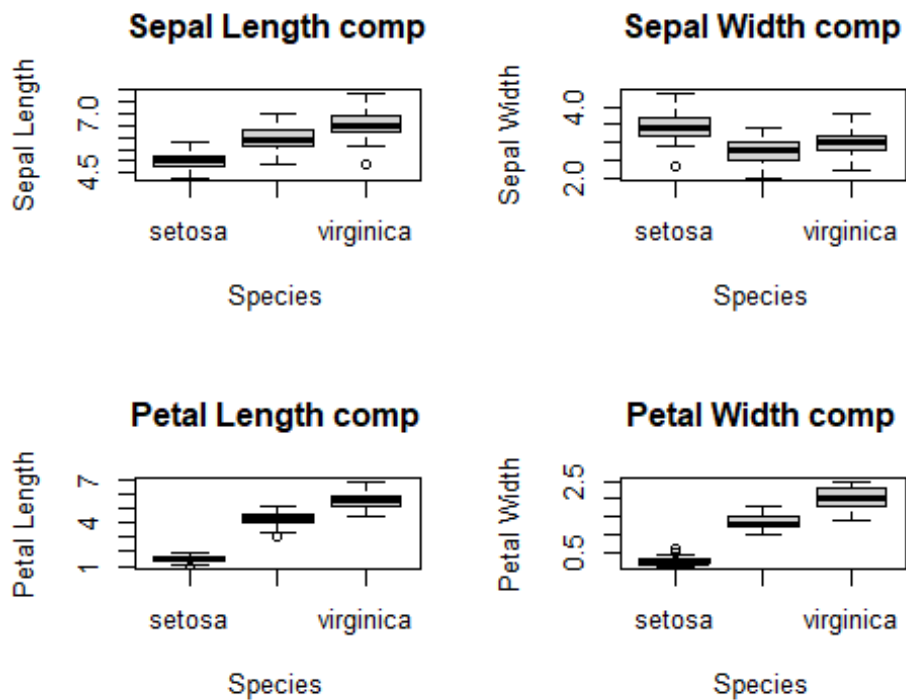
```

    xlab = "Species", ylab = "Sepal Width", main = "Sepal Width
comp")

#Comparación 3
boxplot(Petal.Length~Species,data = iris,
        xlab = "Species", ylab = "Petal Length", main = "Petal Length
comp")

#Comparación 4
boxplot(Petal.Width~Species,data = iris,
        xlab = "Species", ylab = "Petal Width", main = "Petal Width
comp")

```



```

par(mfrow=c(1,1)) #Grafica individual

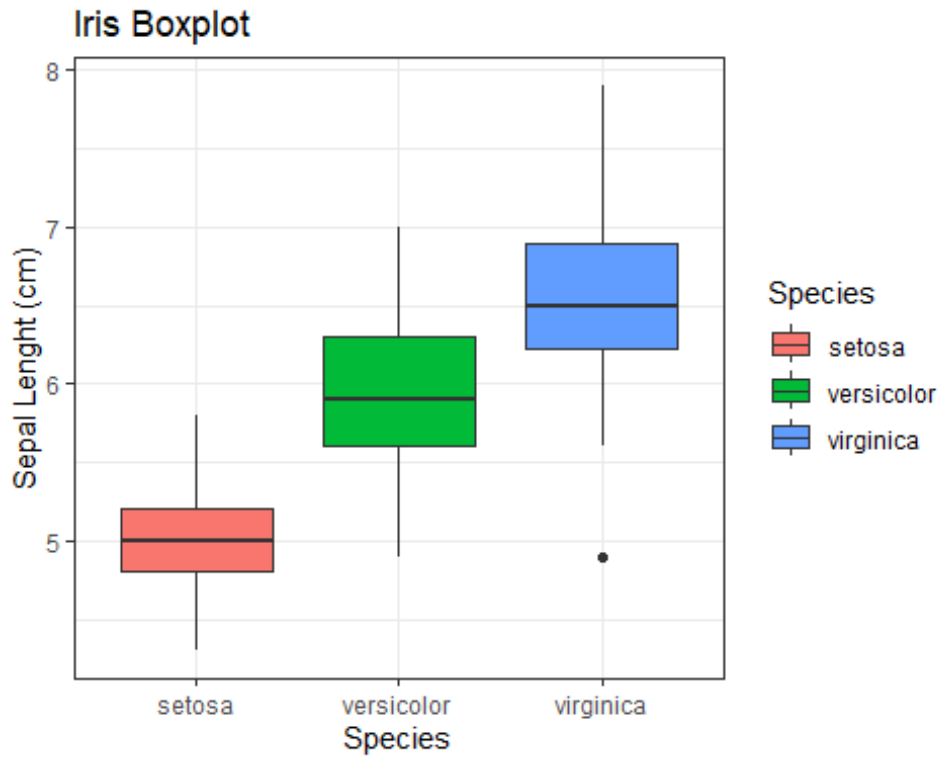
```

Gráfica compacta de 3 colores, usando Iris como dataset

```

#Color
box <- ggplot(data=iris, aes(x=Species, y=Sepal.Length)) #Asignamos
valores
box <- box + geom_boxplot(aes(fill=Species)) +
  ylab("Sepal Length (cm)") + ggtitle("Iris Boxplot") #Título y nombre
eje Y
box <- box + theme_bw() #Asignamos tema visual del gráfico
box #Llamamos gráfico

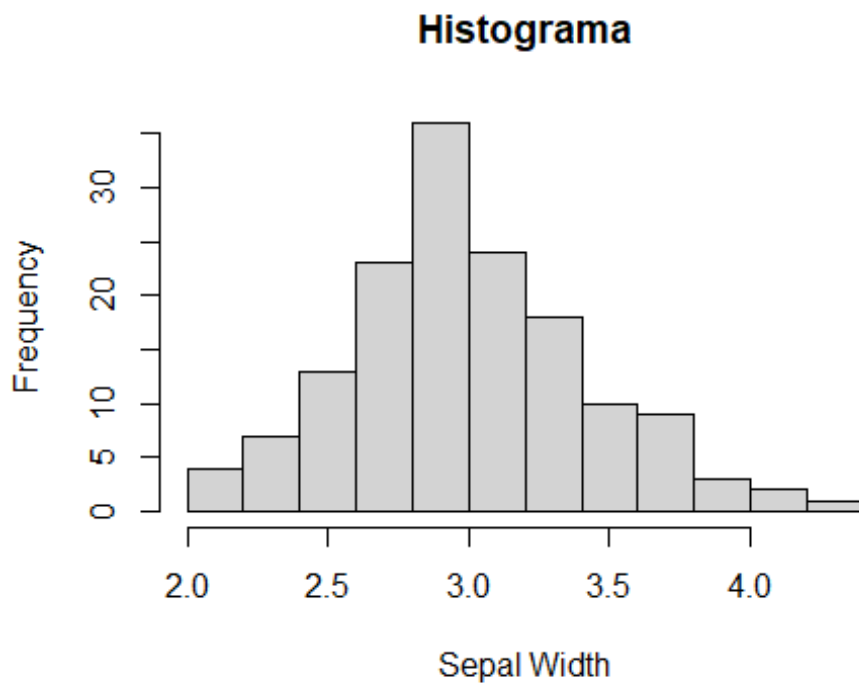
```



Gráficos:

histograma

```
#Histograma
#Simple
hist(iris$Sepal.Width, freq = NULL, density= NULL, breaks=12,
     xlab = "Sepal Width", ylab = "Frequency", main = "Histograma")
#Títulos
```

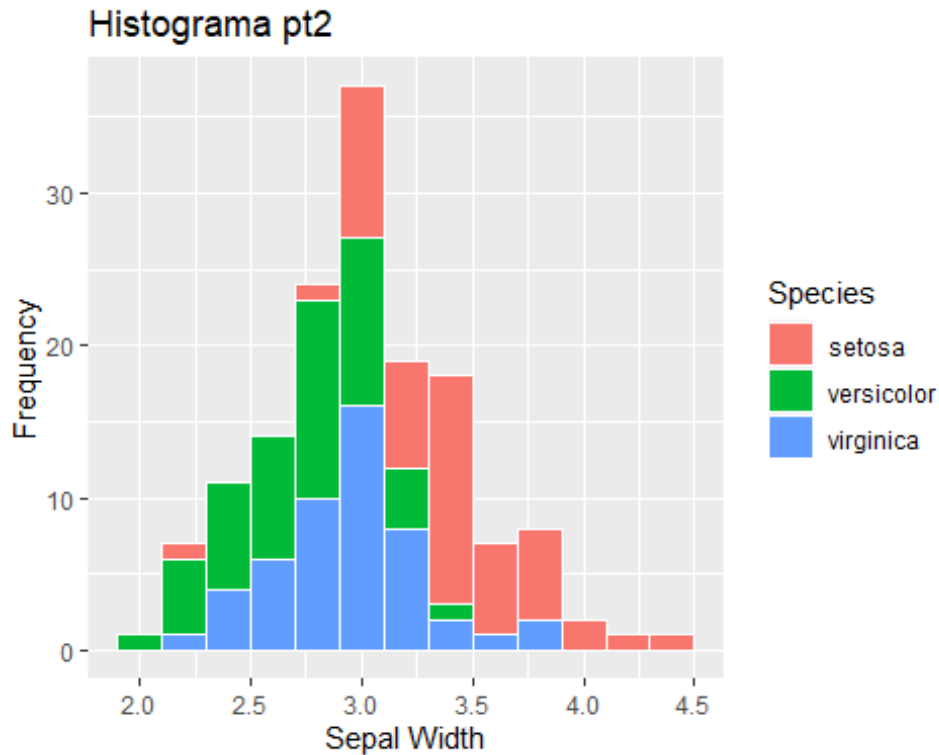


Comparación

entre 3 especies

```
#Comparando Las 3 especies
histogram <- ggplot(data=iris, aes( x=Sepal.Width))

#Asignamos valores a histograma
histogram + geom_histogram(binwidth = 0.2, color = "white",
aes(fill=Species))+
  xlab("Sepal Width") + ylab("Frequency") + ggtitle ("Histograma pt2")
```



Barras por grupos

*#Gráfica de barras por grupos*

```
install.packages("reshape2")
```

```
## Warning: package 'reshape2' is in use and will not be installed
```

```
library("reshape2")
```

*#Usamos base de datos iris para el gráfico*

```
head(iris)
```

```
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1         5.1         3.5          1.4          0.2   setosa
## 2         4.9         3.0          1.4          0.2   setosa
## 3         4.7         3.2          1.3          0.2   setosa
## 4         4.6         3.1          1.5          0.2   setosa
## 5         5.0         3.6          1.4          0.2   setosa
## 6         5.4         3.9          1.7          0.4   setosa
```

```
iris2 <- melt(iris, id.vars = "Species")
```

```
iris2[1:3,]
```

```
##   Species    variable value
## 1  setosa Sepal.Length    5.1
## 2  setosa Sepal.Length    4.9
## 3  setosa Sepal.Length    4.7
```

```

head(iris2)

##   Species      variable value
## 1  setosa Sepal.Length   5.1
## 2  setosa Sepal.Length   4.9
## 3  setosa Sepal.Length   4.7
## 4  setosa Sepal.Length   4.6
## 5  setosa Sepal.Length   5.0
## 6  setosa Sepal.Length   5.4

x <- iris

#Asignamos valores para el gráfico
bar1 <- ggplot(data=iris2, aes(x=Species, y=value, fill=variable))
bar1 + geom_bar (stat= "identity", position= "dodge") +
  #Asignamos colores del gráfico
  scale_fill_manual(values=c("#660000", "#FFCC33", "#CC0033", "#6600CC"),
                    name= "Iris\nMeasurements",
                    #Ponemos nombre/valor de Los 4 segmentos del gráfico
                    breaks = c("Sepal.Length", "Sepal.Width",
"Petal.Length",
                    "Petal.Width"),
                    labels = c("Sepal.Length", "Sepal.Width",
"Petal.Length",
                    "Petal.Width"))

```

