

R for Data Science (V): Programming

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Outline: Introduction to programming

- Introduction
- Control structures
- The apply family
- User defined functions

*Based on this presentation: **Programacion en R** by *Antonio Miñarro* Universitat de Barcelona.

Introduction

- We have introduced R as a *a language (a tool), to manage and analyze data.*
- It is also a *programming language*
 - It is simple and versatile
 - The user can create new functions that adapt to their needs
 - It is widely used (2nd most widely used in Data Science)
 - Users provide the community with a high variety of solutions
 - As a programming language it is not, however, very efficient

Example 1: Why may we need programming

- Assume we have the diabetes dataset and want to make a summary of every variable it contains.

```
library(readxl)
diabetes <- read_excel("datasets/diabetes.xls")
summary(diabetes)
```

- There are categorical variables but the system cannot recognize them.

##	numpacie	mort	tempsviu	edat
##	Min. : 1.00	Length:149	Min. : 0.00	Min. :31.00
##	1st Qu.: 38.00	Class :character	1st Qu.: 7.30	1st Qu.:43.00
##	Median : 75.00	Mode :character	Median :11.60	Median :50.00
##	Mean : 75.01		Mean :10.52	Mean :52.17
##	3rd Qu.:112.00		3rd Qu.:13.90	3rd Qu.:60.00
##	Max. :149.00		Max. :16.90	Max. :86.00
##	bmi	edatdiag	tabac	sbp
##	Min. :18.20	Min. :26.00	Length:149	Min. : 98.0
##	1st Qu.:26.60	1st Qu.:38.00	Class :character	1st Qu.:124.0
##	Median :31.20	Median :45.00	Mode :character	Median :138.0
##	Mean :31.78	Mean :45.99		Mean :139.1
##	3rd Qu.:35.20	3rd Qu.:53.00		3rd Qu.:152.0
##	Max. :59.70	Max. :81.00		Max. :222.0
##	dbp	ecg	chd	
##	Min. : 58.00	Length:149	Length:149	
##	1st Qu.: 74.00	Class :character	Class :character	
##	Median : 80.00	Mode :character	Mode :character	
##	Mean : 90.04			
##	3rd Qu.: 88.00			
##	Max. :862.00			

- A simple solution: Convert text variables into factors.

```
library(forcats)
diabetes$mort <- as_factor(diabetes$mort)
diabetes$tabac <- as_factor(diabetes$mort)
diabetes$ecg <- as_factor(diabetes$ecg)
diabetes$chd <- as_factor(diabetes$chd)
```

```
summary(diabetes)
```

```
##      numpacie      mort      tempsviu      edat
## Min.   : 1.00    Vivo  :124    Min.   : 0.00    Min.   :31.00
## 1st Qu.:38.00    Muerto: 25    1st Qu.: 7.30    1st Qu.:43.00
## Median :75.00                    Median :11.60    Median :50.00
## Mean   :75.01                    Mean   :10.52    Mean   :52.17
## 3rd Qu.:112.00                   3rd Qu.:13.90    3rd Qu.:60.00
## Max.   :149.00                   Max.   :16.90    Max.   :86.00
##      bmi      edatdiag      tabac      sbp
## Min.   :18.20    Min.   :26.00    Vivo  :124    Min.   : 98.0
## 1st Qu.:26.60    1st Qu.:38.00    Muerto: 25    1st Qu.:124.0
## Median :31.20    Median :45.00                    Median :138.0
## Mean   :31.78    Mean   :45.99                    Mean   :139.1
## 3rd Qu.:35.20    3rd Qu.:53.00                    3rd Qu.:152.0
## Max.   :59.70    Max.   :81.00                    Max.   :222.0
##      dbp      ecg      chd
## Min.   : 58.00    Normal  :111    No:99
## 1st Qu.: 74.00    Frontera: 27    Si:50
## Median : 80.00    Anormal  : 11
## Mean   : 90.04
```

- But how should we proceed if there were dozens or hundreds of variables that need to be changed?
- What if, besides, these variables had different names at every new file?
- The solution consists of providing some way to indicate that “any” character variable is transformed into a factor.
- This will be an example of a “program”,

Changing the flow of execution

Scripts are executed “lineally”

- R, as most ordinary programming languages, is executed lineally, that is from the first to last line.
- Sometimes this needs to be changed.
 - Taking alternative flows according to certain conditions
 - Repeating some instructions while certain condition holds, or a fixed number of times, . . .
- This can be accomplished using *Flow control structures*

Loop controlled by a counter: for instruction

- Loops are used in programming to repeat a specific block of code made by one or more instructions.
- Syntax of for loops:

```
for (val in sequence)
{
  statement
}
```

- Here, sequence is a vector and val takes on each of its value during the loop. In each iteration, statement is evaluated.

Example of for loop

- A for loop can be used to change the selected columns in the diabetes dataset.

```
diabetes <- data.frame(read_excel("datasets/diabetes.xls"))
are_char <- c(2,7,10,11)
for (i in are_char) {
  diabetes[,i]<-as_factor(diabetes[,i])
  cat(colnames(diabetes)[i], class(diabetes[,i]), "\n")
}
```

```
## mort factor
## tabac factor
## ecg factor
## chd factor
```

```
summary(diabetes)
```

```
##      numpacie      mort      tempsviu      edat
## Min.   : 1.00    Vivo  :124    Min.   : 0.00    Min.   :31.00
## 1st Qu.:38.00    Muerto: 25    1st Qu.: 7.30    1st Qu.:43.00
## Median :75.00                    Median :11.60    Median :50.00
## Mean   :75.01                    Mean   :10.52    Mean   :52.17
## 3rd Qu.:112.00                   3rd Qu.:13.90    3rd Qu.:60.00
## Max.   :149.00                   Max.   :16.90    Max.   :86.00
##      bmi      edatdiag      tabac      sbp
## Min.   :18.20    Min.   :26.00    No fumador:57    Min.   : 98.0
## 1st Qu.:26.60    1st Qu.:38.00    Fumador   :51    1st Qu.:124.0
## Median :31.20    Median :45.00    Ex fumador:41    Median :138.0
## Mean   :31.78    Mean   :45.99                    Mean   :139.1
## 3rd Qu.:35.20    3rd Qu.:53.00                    3rd Qu.:152.0
## Max.   :59.70    Max.   :81.00                    Max.   :222.0
##      dbp      ecg      chd
## Min.   : 58.00    Normal  :111    No:99
## 1st Qu.: 74.00    Frontera: 27    Si:50
## Median : 80.00    Anormal  : 11
## Mean   : 90.04
```

Exercise

- Create a `for` loop that reads all `.csv` filenames in your datasets directory (or the directory you decide) and prints the name of the file and the column names in the screen.

Conditional statements: if / if - else.

- Conditional statements allow different coding blocks to be executed depending on whether a certain condition is TRUE or FALSE.
- syntax of if statement is:

```
if (test_expression) {  
  statement  
}
```

- If the test_expression is TRUE, the statement gets executed. But if it's FALSE, nothing happens.
- Here, test_expression can be a logical or numeric vector, but only the first element is taken into consideration.
- In the case of numeric vector, zero is taken as FALSE, rest as TRUE.

Conditional statements: if - else.

- syntax of if-else statement is:

```
if (test_expression) {  
    statement_1  
}else{  
    statement_2  
}
```

- If the test_expression is TRUE, then statement_1 gets executed.
- If it's FALSE then statement_2 gets executed.