

# Introduction to Programming

## Lecture 7-8: Introduction to R

Clément Mazet-Sonilhac  
`clement.mazet@sciencespo.fr`

Sciences Po Paris

# Disclaimer

- Most of the material is drawn from the excellent course prepared by software carpentry
- In particular, most exercises are drawn from it (If you really want to learn something, don't look up the answers)
- Other source of inspiration is the very complete QuantEcon website

# What and why?

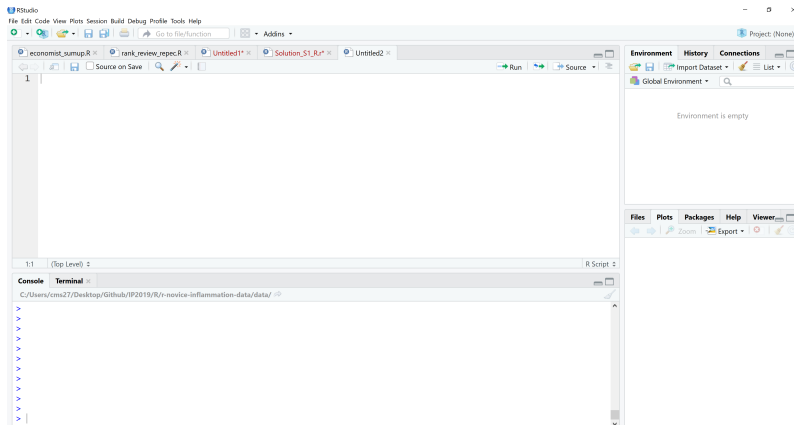
R : Let's start !

- Why are we using R?
  - ▶ **Better than Stata** by ANY metric
  - ▶ Free
  - ▶ Extremely popular amongst scientists, in particular statisticians and economists
  - ▶ Exists a large library of external packages

# What and why?

R : Let's start !

- Let's start by opening RStudio !



# Variables

## Create a variable in R

- A variable : a container with a name
- To create a variable called `weight` with value 55, just type :

```
weight <- 55 (or weight = 55)
```

# Variables

## Create a variable in R

- A variable : a container with a name
- To create a variable called `weight` with value 55, just type :
- Can treat the variable like a regular number. Try `weight + 1`
- Can change an variable's value by assigning it a new value. Just type :

```
weight <- 55 (or weight = 55)
```

```
weight <- 60
```

# Variables

## Create a variable in R

- R only stores the value, not the calculation used to create a variable ( $\neq$  Excel). Try this :

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## Create a variable in R

- R only stores the value, not the calculation used to create a variable ( $\neq$  Excel). Try this :

```
weightlb <- 2.2 * weightkg
```

```
c(weightkg, weightlb)
```

```
weightkg <- 80
```

```
c(weightkg, weightlb)
```

- `c` is also a function (probably the most used function in R), stands for combine



# Variables

Create a variable in R

- Some conventions on the name of variables
  1. start with lower case letters
  2. separate words with underscores
  3. use only lowercase letters, underscores, and numbers

# Motivating example

## Analyzing data w. R

- **The data** : We are studying inflammation in patients who have been given a new treatment for arthritis, and we need to analyze the first dozen data sets. The data sets are stored in comma-separated values (CSV) format. Each row holds the observations for just one patient. Each column holds the inflammation measured in a day, so we have a set of values in successive days.

# Motivating example

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  1. Go to the clone of my Github Repo on your computer (or to my Github repo ([github.com/CMS27/IP2019](https://github.com/CMS27/IP2019))) and download :  
`r-novice-inflammation-data`
  2. Goal : load the data, calculate the average value of inflammation per day, plot the results

# Motivating example

## Analyzing data w. R

- Loading data :
  1. Set the directory where the data is stored with `setwd()` :

```
setwd("C:/Users/YourName/.../data")
```
  2. Import data in `d` with :

```
d=read.csv(file="inflammation-01.csv", header=FALSE)
```
- both `setwd()` and `read.csv()` are functions that takes some arguments

# Motivating example

## Analyzing data w. R

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  1. Set the directory where the data is stored with `setwd()` :

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setwd("C:/Users/YourName/.../data")
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  2. Import data in `d` with :

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d=read.csv(file="inflammation-01.csv", header=FALSE)
```
- both `setwd()` and `read.csv()` are functions that takes some arguments
  1. the first argument of both functions is a String  $\Rightarrow$  put quotes
  2. the second argument of `read.csv` is what we call a Boolean value (either true or false). Header : whether the first line of the file contains names for the columns of data
  3. `d` = data frame. more on this later : but basically, like an excel sheet.

# Motivating example

## Analyzing data w. R

- Manipulating the data :

1. Display the first lines of the data set with `head` :

```
head(d, n = 3L)
```

2. To take a subset of the data set, provide an index in square bracket : `[# row, # column]` :

```
d[1,1]
```

```
d[c(1, 3, 5), c(10, 20)]
```

```
d[1, 1:5]
```

```
d[, 1]
```

# Motivating example

## Analyzing data w. R

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```

2. To take a subset of the data set, provide an index in square bracket : `[# row, # column]` :

```
d[1,1] # first row, first column
```

```
d[c(1, 3, 5), c(10, 20)] # rows (1, 3 and 5), columns (10 and 20)
```

```
d[1, 1:5] # columns from (1 to 5) and row 1
```

```
d[, 1] # all rows from column 1
```

# Motivating example

## Analyzing data w. R

- In our data set, each row is a patient, each column is a day, such that `d[1,1]` is the inflammation measured on patient 1 on day 1
- **Exercise 1** : given that `min(data)`, `max(data)`, `mean(data)` are functions returning the equivalent statistics on data, find :
  1. the minimum inflammation on day 1 across all patients



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  3. the maximum inflammation on days 4, 8 and 12 across all patients

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  4. the minimum inflammation experienced by patients 3 and 6 from day 1 to 5

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  2. the maximum inflammation experienced by patient 5 (across all days)
  3. the maximum inflammation on days 4, 8 and 12 across all patients
  4. the minimum inflammation experienced by patients 3 and 6 from day 1 to 5
  5. the mean inflammation experienced by patients 2, 4 and 10 (across all days)

# Motivating example

## Analyzing data w. R

- Faster way to get some sufficient statistics (by columns) : `summary` (ex : `summary(d[, 1:5])`)
- What if we want some info, say the median, for each patient (= row)?  
No such things as `rowMedian`
- `apply` : repeat a function on all of the rows (`MARGIN = 1`) or columns (`MARGIN = 2`) of a data frame (`apply(d, 1, median)`)
- **Exercise 2** : compute in two different ways the mean for the first 10 patients of our data

# Motivating example

## Analyzing data w. R

- R plot are very nice :
- Try `plot(apply(d, 2, max), xlab = "day", ylab = "maximum", main = "maximum inflammation by day")`
- and `boxplot(d, main = "Summary")`

# Motivating example

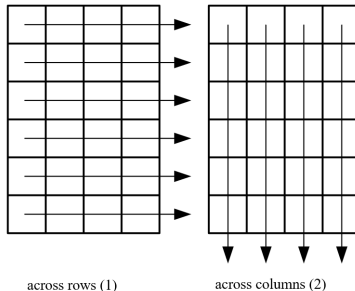
## Analyzing data w. R

- **Exercise 2** : Load the data, use the function `apply` to create a variable containing the min value each day + plot the result

# Motivating example

## Analyzing data w. R

- **Exercise 2** : Load the data, use the function `apply` to create a variable containing the min value each day + plot the result





# What and why ?

## Function in R

- Questions :
  - ▶ How do I make a function ?
  - ▶ How can I test my functions ?
  - ▶ How should I document my code ?

# What and why ?

## Function in R

- Questions :
  - ▶ How do I make a function ?
  - ▶ How can I test my functions ?
  - ▶ How should I document my code ?
- Objectives
  - ▶ Define a function that takes arguments.
  - ▶ Return a value from a function.
  - ▶ Test a function.
  - ▶ Explain why we should divide programs into small, single-purpose functions.

# What and why?

## Function in R

- Function : why is this so useful?

# What and why ?

## Function in R

- Function : why is this so useful ?
- If we only had one data set to analyse, it would probably be faster to load the file into a spreadsheet and use that to plot some simple statistics...
- ... but we have twelve files to check, and may have more in the futur !
- In this lesson, we'll learn how to write a function so that we can repeat several operations with a single command.

# Your first function

## Defining a function in R

- A function starts with a **name**, some **arguments** and an **output** :

```
fname <- function(arg1, arg2, ...) {  
  output = f(arg1, arg2)  
  return(output)  
}
```

# Your first function

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- A function starts with a **name**, some **arguments** and an **output** :

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  output = f(arg1, arg2)  
  return(output)  
}
```

- **Exercise 3** : Imagine you want to convert temperatures from Fahrenheit to Kelvin. i) Create a function called `fk` that takes as argument a temperature in F and return a temperature in K (Hint :  $K = ((F - 32) \times 5/9) + 273.15$ ) and ii) test the function for value 32 and 212.

# Your first function

## Combining functions if R

- **Exercise 4** : Create a function called `kc` that takes as argument a temperature in K and return a temperature in C (celsius) (Hint :  $C = K - 273.15$ )

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# Your first function

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- Nesting functions : Try `kc(fk(32))`. What is the result ?

# Your first function

## Combining functions if R

- **Exercise 4** : Create a function called `kc` that takes as argument a temperature in K and return a temperature in C (celsius) (Hint :  $C = K - 273.15$ )
  - **Exercise 4bis** : Combine `fk` and `kc` in order to create a new function `fc` that converts a temperature in F to C.
  - Nesting functions : Try `kc(fk(32))`. What is the result ?
- ⇒ This is our first taste of how larger programs are built : we define basic operations, then combine them in ever-larger chunks to get the effect we want !

# Your first function

With words !

- Imagine you have a vector of words `vc` and a punctuation vector `vp` s.t :

```
vc <- c("Hello", "World")
```

```
and vp <- c("***")
```

⇒ Remember the combine function `c` ?

# Your first function

With words !

- Imagine you have a vector of words `vc` and a punctuation vector `vp` s.t :

```
vc <- c("Hello", "World")
```

```
and vp <- c("***")
```

⇒ Remember the combine function `c` ?

- Exercise 5** : create a function `fence` that return `"***" "Hello"`  
`"World" "***"`
- Exercise 5bis** : create a function outside that returns the first and the last element of a vector (here : `"***" "***"`). Hint : use the function `length(v)`

# Your first function

## Default value for arguments

```
mySum <- function(input1, input2 = 10) {  
  output <- input1 + input2  
  return(output)  
}
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- Q1 : what is the result of `mySum(input1 = 1,3)` ?

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- Q1 : what is the result of `mySum(input1 = 1,3)` ?
- Q2 : what is the result of `mySum(3)` ?

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## Default value for arguments

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}
```

- Q1 : what is the result of `mySum(input1 = 1,3)` ?
- Q2 : what is the result of `mySum(3)` ?
- Q3 : what is the result of `mySum(input2 = 3)` ? Why ?



# Your first function

## Working with several files

- **Exercise 6 (Difficult !)**
- Write a function called `analyze` that :
  1. takes a filename as an argument
  2. displays the three graphs produced in the previous lesson (average, min and max inflammation over time).
- Hint : `analyze("../data/inflammation-01.csv")` should produce the graphs already shown, while `analyze("../data/inflammation-02.csv")` should produce corresponding graphs for the second data set. Be sure to document your function with comments.

# Your first function

## Working with several files

- **Exercise 6\* (Very Hard !)**
- Write a function called `analyze_all` that :
  1. takes a path as an argument
  2. displays all filenames in the path
  3. use the function `analyze_all` recursively to display the three graphs produced in the previous lesson (average, min and max inflammation over time).