## Introduction to R programming for data science – day 1

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## R programming

#### R is:

- One of the leading programming languages for statistics and data science
- Useful for basic and advanced data analyses
- Prerequisite for job positions in life science and biomedicine

#### Why?

- · Its is free
- It supports reproducibile analyses
- It provides packages for data analysis

#### In this course...

We will learn the **basics of R programming** through lectures and hands-on computer exercises using Rstudio:

- variables, data types, and data structures
- data manipulation and visualization
- operations
- reading and writing files
- running and repeating tasks using functions and control statements.

Disclaimer: this is **not** a course in statistics or bioinformatics.

#### Course info

Course material: https://github.com/FFinotello/Rcourse

Note: to download a text file, right-click on Raw and select Save link as.

#### Exam

- Written test: multiple-choice questions on theory and small example codes like those in the slides (at the MUI after the course, without the laptop)
- Hands-on project: data analysis project to be described in a short report (at home, to be sent by e-mail)

## Installation

Install first R and then Rstudio

To install R:

https://www.r-project.org

See "To download R, please choose your preferred CRAN mirror".

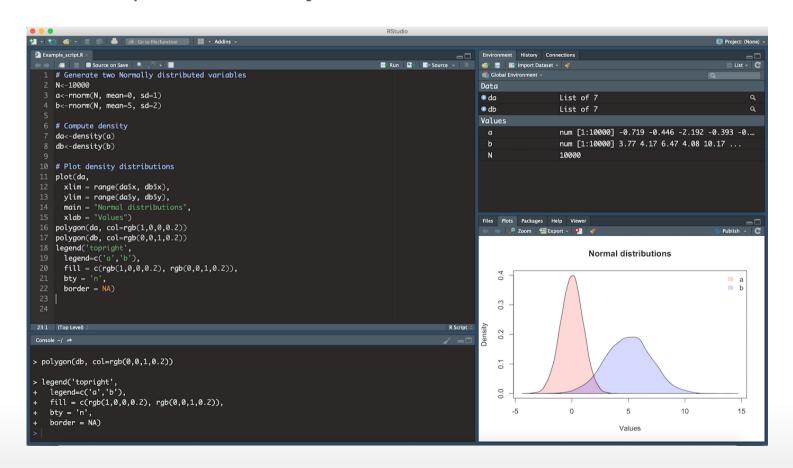
To install **RStudio Desktop**:

https://www.rstudio.com/products/rstudio/download/#download

## **RStudio**

#### RStudio console

RStudio is an integrated development environment (IDE) for R. It has four main panels with adjustable sizes:



## RStudio console

In the console you can type commands and see outputs.

```
(12^2 - 20 * 2 + 1) / 5
```

## [1] 21

log10(100)

## [1] 2

sqrt(9) / 3

## [1] 1

Have a look at the History panel: what do you see?

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## RStudio scripts

Instead of using the console, you can save your code in a .R file (e.g. *R\_script\_day1.R*)

To open a **new file**: File > New File > New R Script

To **save** as R script: File > Save as

To set the **working directory**: Session > Set Working Directory > To Source File Location

To **open** an R script: File > Open File (e.g. Example\_script.R)

You can run the code in a script altogether with Source, or in chunks using Control+Enter.

Remember to save your code regularly!

# Variables and data types

### **Variables**

Variables are symbolic names used to store data that can be manipulated in a computer program

In R, variables are assigned with <-

```
height <- 1.90

weight <- 80

# Body mass index

BMI <- weight/height^2
```

Have a look at the **Environment** panel: what do you see?

## Case-sensitive programming

R is case-sensitive

BMI

## [1] 22.16066

bmi

## Error in eval(expr, envir, enclos): object 'bmi' not found

#### Variable names

- Short, but descriptive
- Must start with a letter
- Must not contain special characters
- · Cannot be reserved words (if, function, NA, TRUE, ...)

Try assign the value of 10 to a variable called:

- myVar
- my\_var
- my.var
- my-var
- 1var
- · var1

## **Operations**

Once variables are assigned, they can be used to perform operations

```
x <- 6
```

y <- 2

x + y

## [1] 8

x^y

## [1] 36

x == y

## [1] FALSE

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## Arithmetic operators

Operator	Description
+	Addition
-	Subtraction
*	Multiplication
/	Division
٨	Exponent
%%	Modulus (remainder from division)
%/%	Integer Division

## Relational operators

Operator	Description
<	Less than
>	Greater than
<=	Less than or equal to
>=	Greater than or equal to
==	Equal to
!=	Not equal to

## **Logical operators**

Operator	Description
!	Logical NOT
&	Element-wise logical AND
&&	Logical AND
	Element-wise logical OR
	Logical OR

## Atomic vector types

In R, there are several types of atomic vectors (also called *classes*):

- Character
- Numeric
- Logical
- · Complex
- Integer

## Data type examples

```
Name <- "Maria"
class(Name)
## [1] "character"
PhD <- TRUE
class(PhD)
## [1] "logical"
yearsSincePhD <- 5</pre>
class(yearsSincePhD)
## [1] "numeric"
```

## Missing Values

In R, missing values are usually coded with NA

```
Name <- "Michael"
PhD <- FALSE
yearsSincePhD <- NA
is.na(Name)
## [1] FALSE
is.na(yearsSincePhD)
## [1] TRUE
Note: NA are not strings (no quotation marks)
```

## Other indefinite values

Not a number: NaN

```
( var1 <- 0/0 )

## [1] NaN

Infinite: Inf

( var2 <- 9/0 )

## [1] Inf

var3 <- -10/0</pre>
```

What is the value of *var3*?

#### Data conversion/coercion

Class conversion is possible, but it must be handled with care

```
var1 <- 1
( var2 <- as.character(var1) )</pre>
## [1] "1"
class(var2)
## [1] "character"
( var3 <- as.logical(var1) )</pre>
## [1] TRUE
```

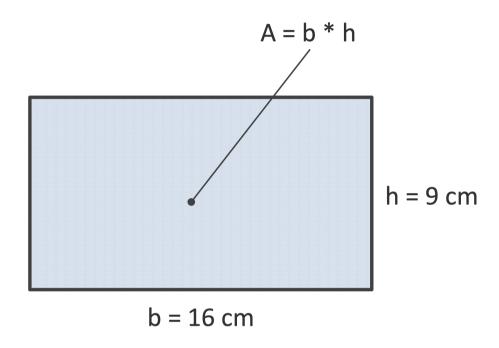
## Data type conversion

Conversion to	Function	Rules
numeric	as.numeric	$\mathtt{FALSE} \to 0$
		$\texttt{TRUE}  \to  1$
		"1", "2", $\ldots \rightarrow 1, 2, \ldots$
		$"A", \ldots \rightarrow NA$
logical	as.logical	$0 \rightarrow \mathtt{FALSE}$
		other numbers $\rightarrow$ TRUE
		"FALSE", "F" $ ightarrow$ FALSE
		"TRUE", "T" $ ightarrow$ TRUE
		other characters $\rightarrow NA$
character	as.character	$1, 2, \ldots \rightarrow$ "1", "2", $\ldots$
		$\texttt{FALSE} \to \texttt{"FALSE"}$
		$TRUE  \to  "TRUE"$

Source: https://cran.r-project.org/doc/contrib/Paradis-rdebuts\_en.pdf

## **Exercises**

#### Ex. 1.1



- Save the length of the base and heigth of the rectangle depited above into two numeric variables called, respectively, *b* and *h*.
- Compute the area of the rectangle and save it into a numeric variable called *A*.
- Check if the area is bigger than  $100 \text{ cm}^2$ .

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#### Ex. 1.2

```
Name <- "Maria"
Age <- "20"
PhD <- "TRUE"
```

#### What is the class of the variables above?

- 1. Name: character; Age: numeric; PhD: character
- 2. Name: character; Age: character; PhD: character
- 3. Name: character; Age: numeric; PhD: logical
- 4. None of the above

## Ex. 1.3

x <- 5

y <- 0

z <- x/y

#### What is the value of *z*?

- 1. -Inf
- 2. NA
- 3. NaN
- 4. None of the above

## Data structures

#### Data structures

In R, there are different data structures, including:

Data structure	Can contain different data types?
Vector	No
Matrix	No
Factor	No
List	Yes
Data.frame	Yes

#### Data structure: vector

Vectors can be build with the *c* function and their length can be assessed with the function *length* 

```
( x <- c(1, 2, 3, 4) )

## [1] 1 2 3 4

( y <- c("a", "b", "c") )

## [1] "a" "b" "c"

length(y)

## [1] 3</pre>
```

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#### Data structure: vector

Vectors can contain a single data type only, so beware of conversions

```
(z <- c("a", "b", 1, 2))
```

$$(v \leftarrow c(x, y))$$

What are the classes of *z* and *v*?

## Fast ways to build vectors

```
( j <- 1:4 ) # Seq. of integers
## [1] 1 2 3 4
(x \leftarrow seq(0, 1, 0.1)) # Seq. of numbers (from, to, step)
## [1] 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0
( y <- rep("A", 5) ) # Seq. of repeated numbers or characters
## [1] "A" "A" "A" "A" "A"
( z <- c(rep("A", 3), rep("B", 5)) ) # Vector combination
## [1] "A" "A" "A" "B" "B" "B" "B" "B"
```

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## Assign and query vector names

```
( age <- c(19, 30, 20) )
## [1] 19 30 20
names(age) <- c("Mary", "John", "Lisa") # Assign</pre>
age
## Mary John Lisa
     19
          30
               20
names(age) # Query
## [1] "Mary" "John" "Lisa"
```

#### Data structure: matrix

2-dimensional object that can be build with the function *matrix* 

```
( M <- matrix(1:6, ncol=3, byrow=FALSE) ) # Fill by columns (def.)

## [,1] [,2] [,3]
## [1,] 1 3 5
## [2,] 2 4 6

( N <- matrix(1:6, ncol=3, byrow=TRUE) ) # Fill by rows

## [,1] [,2] [,3]
## [1,] 1 2 3
## [2,] 4 5 6</pre>
```

## Matrix rows and columns

```
M <- matrix(1:6, nrow=2, byrow=FALSE)</pre>
dim(M)
## [1] 2 3
ncol(M)
## [1] 3
nrow(M)
## [1] 2
```

#### Matrix row and column names

```
colnames(M) <- c("Sample A", "Sample B", "Sample C") # Assign</pre>
rownames(M) <- c("Gene1", "Gene2") # Assign</pre>
Μ
       Sample A Sample B Sample C
## Gene1 1 3
## Gene2 2 4
colnames(M) # Query
## [1] "Sample_A" "Sample_B" "Sample_C"
rownames(M) # Query
## [1] "Gene1" "Gene2"
```

### Data structure: factor

Categorical variable that can be built with the function *factor* 

```
strain <- factor(c("WildType", "WildType", "Mutant", "Mutant"))</pre>
strain
## [1] WildType WildType Mutant
                                  Mutant
## Levels: Mutant WildType
length(strain)
## [1] 4
levels(strain)
## [1] "Mutant"
                  "WildType"
```

### **Factor conversion**

```
(x \leftarrow factor(c(10, 2, 2, 3)))
## [1] 10 2 2 3
## Levels: 2 3 10
( y <- as.numeric(x) ) # This does NOT work</pre>
## [1] 3 1 1 2
( z <- as.numeric(as.character(x)) ) # This works</pre>
## [1] 10 2 2 3
```

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### Data structure: list

A list is a structure that can contain different objects (including lists!)

It can be created with the function *list* 

```
v <- 1:3
M <- matrix(1:4, ncol=2)
( myList <- list(myVec=v, myMat=M) )

## $myVec
## [1] 1 2 3
##
## $myMat
## [,1] [,2]
## [1,] 1 3
## [2,] 2 4</pre>
```

## List names and length

```
length(myList)
## [1] 2
names(myList)
## [1] "myVec" "myMat"
myList <- list(v, M)</pre>
names(myList)
## NULL
```

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### Data structure: data.frame

A data.frame is a 2-dimensional object ("matrix-like") that can contain different types of data on its colums.

It can be created with the function data.frame

```
( DF <- data.frame(name=c("Mary", "John", "Lisa"),
   age=c(19, 30, 20),
   city=c("New York", "Seattle", "New York"),
   stringsAsFactors=FALSE) )</pre>
```

```
## name age city
## 1 Mary 19 New York
## 2 John 30 Seattle
## 3 Lisa 20 New York
```

### Data.frame rows and columns

```
dim(DF)
## [1] 3 3
nrow(DF)
## [1] 3
ncol(DF)
## [1] 3
colnames(DF)
## [1] "name" "age" "city"
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```

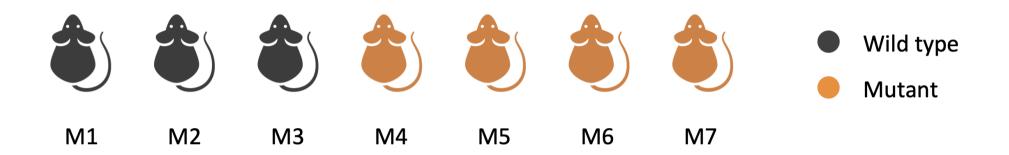
# **Exercises**

```
x <- c("a", "b", "c")
y <- seq(1,10)
z <- c(x,y)
```

#### What is the class of *z*?

- 1. Character and numeric
- 2. Numeric
- 3. Character
- 4. NA

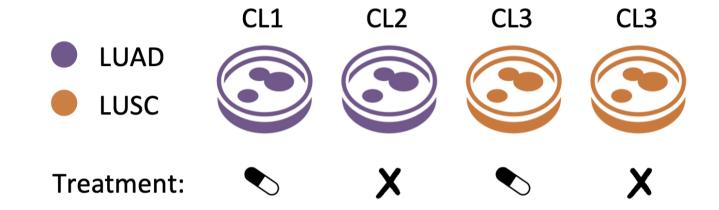
Build a numeric vector containing all even numbers greater than 0 and not exceeding 100.



- Build a vector of factors storing the information about which mice is wild type ("WT") or mutant ("MU") in the figure above (use rep).
- Assign mice identifiers (M1, M2, ...) to the same vector (use names).

- Initialize a matrix with 2 rows and 3 columns that has all "1" on the first row and all "2" on the second row.
- · Give it column and row names as you wish.
- · Check its dimensions.

- Initialize a numeric vector to store the year in which you got your driving licence and the year you got your first car (NA values are possible). Set appropriate names.
- · Initialize a character vector to store the names of your favorite cities in Europe (as many as you like).
- Save both vectors in a list using meaningful names.



- · Imagine you have an experiment with 4 lung cancer cell lines: 2 from adenocarcinomas (LUAD) and 2 from squamous cell carcinoma (LUSC).
- · One cell line from each cancer subtype is treated with a drug, the other one is untreated, as shown in the figure above.
- Save in a data.frame the info about the experiment: cell line identifier, lung-cancer subtype, and treatment.
- Check how many rows and columns the data.frame has.

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### Useful resources

R for beginners: <a href="https://cran.r-project.org/doc/contrib/Paradis-rdebuts\_en.pdf">https://cran.r-project.org/doc/contrib/Paradis-rdebuts\_en.pdf</a>

Stackoverflow (Ask a Question): https://stackoverflow.com

### **Solutions**

- **Ex. 1.1:** A=144 (>100)
- Ex. 1.2: 2) Character, character
- Ex. 1.3: 4) None of the above (Inf)
- Ex. 1.4: 3) Character