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 the life sciences

Why?

- · Its is free
- · It supports reproducible analysis
- · 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.

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Course info

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

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

Evaluation: based on a data analysis project to be described in a short report and sent by e-mail.

How to ask questions

- · Chat
- · Collaborative Google Doc
- · E-mail

Installation

Install first R and then Rstudio

To install **R** (version \geq 4):

https://www.r-project.org

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

To install **RStudio Desktop**:

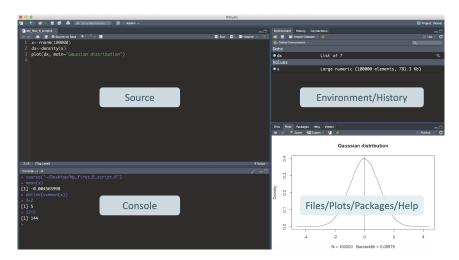
https://www.rstudio.com/products/rstudio/download/#download

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RStudio

RStudio console

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



From the toolbar: View > Panes > Show All Panes

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The RStudio console

In the console you can type commands and see outputs.

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

## [1] 21

log10(100) - sqrt(9) / 3

## [1] 1

# Any text preceded by a hashtag is not evaluated (= comment!)
```

Variables and data types

Variables

Variables: 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
```

Variable names

- · Should be short but descriptive
- · Must start with a letter
- Must not contain special characters
- · Cannot be reserved words

Reserved words: words with special meaning in R that cannot be used as identifiers. Examples: if, break, Inf, TRUE

```
de cellFrac, cell_frac, cell.frac, cellfrac_1
```

```
👎 cell-frac, 1cellfrac, TRUE
```

```
(T <- 5 * 2 )
## [1] 10
```

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Case-sensitive programming

R is case-sensitive

```
# Body mass index (BMI)
# defined as: weight/height^2
BMI <- 80/1.90^2

BMI

## [1] 22.16066

bmi

## Error in eval(expr, envir, enclos): object 'bmi' not found</pre>
```

Operations

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

```
height <- 1.90
weight <- 80
( BMI <- weight/height^2 )

## [1] 22.16066

x <- 6
y <- 2
x <- 4
x + y

## [1] 6
```

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Arithmetic operators

Operator	Description
+	Addition
-	Subtraction
*	Multiplication
1	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

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Logical operators

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

Atomic vector types

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

- · Character
- Numeric
- · Logical
- · Complex
- · Integer

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Data type examples

```
Name <- "Maria"
class(Name)

## [1] "character"

PhD <- TRUE
class(PhD)

## [1] "logical"

yearsSincePhD <- 5
class(yearsSincePhD)

## [1] "numeric"</pre>
```

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</pre>
```

Note: NA are not strings (no quotation marks)

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Other indefinite values

Not a number: NaN

```
var1 <- 0/0
var1

## [1] NaN

Infinite: Inf

( var2 <- 9/0 )

## [1] Inf

( var3 <- -10/0 )

## [1] -Inf</pre>
```

Data conversion/coercion

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

```
var1 <- 1
( var2 <- as.character(var1) )

## [1] "1"

class(var2)

## [1] "character"

( var3 <- as.logical(var1) )

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

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Data type conversion

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

Source: https://cran.r-project.org/doc/contrib/Paradis-rdebuts_en.pdf

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 *length* function

```
( 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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Vectors and data types

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

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

## [1] "a" "b" "1" "2"

(v <- c(x, y))

## [1] "1" "2" "3" "4" "a" "b" "c"
```

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

Creating numeric vectors

```
j <- 1:4  # Seq. of integers
j

## [1] 1 2 3 4

x <- seq(0, 1, 0.1) # Seq. of numbers (from, to, step)
x

## [1] 0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0</pre>
```

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Creating vectors with rep

```
y <- rep("A", 5) # Seq. of repeated numbers or characters
y

## [1] "A" "A" "A" "A" "A"

z <- c(rep("A", 3), rep("B", 5)) # Vector combination
z

## [1] "A" "A" "A" "B" "B" "B" "B" "B"</pre>
```

Vector names

```
( age <- c(19, 30, 20) )
## [1] 19 30 20

names(age) <- c("Mary", "John", "Lisa") # Assign
age
## Mary John Lisa
## 19 30 20

names(age) # Query
## [1] "Mary" "John" "Lisa"</pre>
```

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Accessing vectors

You can access the elements of a vector by using *logical* or *numeric* indexes (positive and negative) between brackets:

```
days <- c("Mon", "Tue", "Wed", "Thu", "Fri", "Sat", "Sun")

days[c(1,6,7)]

## [1] "Mon" "Sat" "Sun"

weekend <- days[c(FALSE, FALSE, FALSE, FALSE, FALSE, TRUE, TRUE)]

weekend <- days[-seq(1,5)] # Negative ind. of elements to be removed</pre>
```

Accessing vectors

To access vector elements, you must use square brackets. Round brackets are for functions.

```
age <- c(11, 12, 18, 20, 45, 2, 33)
age[1]
## [1] 11
age(1)
## Error in age(1): could not find function "age"
mean(age)
## [1] 20.14286</pre>
```

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Accessing vectors

Vector elements can be also accessed using their names, when initialized

```
age <- c(11, 12, 18, 20)
names(age) <-c("John", "Lisa", "Maria", "Markus")
age

## John Lisa Maria Markus
## 11 12 18 20

age[c("John", "Maria")]

## John Maria
## 11 18</pre>
```

Subsetting vectors

You can extract the elements of a vector that satisfy a certain condition:

```
age <- c(11, 12, 18, 20, 45, 2, 33)
age[age>=18] # Logical index

## [1] 18 20 45 33
age[which(age>=18)] # Numerical index
## [1] 18 20 45 33
```

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Manipulating vectors

You can change the elements of a vector:

```
x <- c(11, 12, -18, 20, -45, -2, 33)
x[1] <- 35 # Subsitute the first element
x
## [1] 35 12 -18 20 -45 -2 33
x[x<0] <- 0 # Set to 0 all negative elements
x
## [1] 35 12 0 20 0 0 33</pre>
```

Data structure: factor

A **factor** is categorical variable that can be built with the function **factor**

```
strain <- factor(c("WildType", "WildType", "Mutant", "Mutant"))
strain

## [1] WildType WildType Mutant Mutant
## Levels: Mutant WildType

length(strain)

## [1] 4

levels(strain)

## [1] "Mutant" "WildType"</pre>
```

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Factor conversion

```
( x <- factor(c(10, 2, 2, 3)))

## [1] 10 2 2 3

## Levels: 2 3 10

( y <- as.numeric(x) ) # This does NOT work

## [1] 3 1 1 2

( z <- as.numeric(as.character(x)) ) # This works

## [1] 10 2 2 3</pre>
```