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Introduction to R

This is *not* a systematic or comprehensive introduction to R, it's just a quick look how R works. You are going to learn R more systematically during this course, especially in the exercises.

The following books (as pdf-files on ILIAS, in the folder Software: R and Jupyter Notebook -> Books) offer a more comprehensive introduction to R.

- Dalgaard Peter: Introductory Statistics with R (Springer)
 Peter Dalgaard is a developer of R. It's a systematic introduction to R. Recommended.
- Jim Albert, Maria Rizzo: R by Example (Springer)
 As the title says, the content of this book consists entirely of examples. Recommended, although some examples are quite advanced.
- Andre de Vries, Joris Meys: R for Dummies
 If you like the Dummies series, this book is for you.

1 What is R?

The programming language R was *specifically* developed to solve statistical problems. For information about the history of R, see

```
https://en.wikipedia.org/wiki/R_(programming_language)
```

R has the advantage that it is very widespread in the statistics community. In a way R is the state-of-the-art software for solving statistical problems. Furthermore, the software is open source and therefore available for free.

For beginners, the handling of R needs somewhat to get used to (like for any software), but once that happened you'll find that R is a very powerfull software.

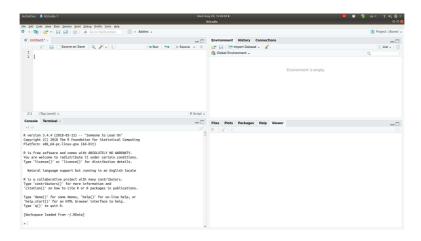
2 Installation

The installation instructions are on ILIAS (folder Software: R and Jupyter Notebook -> Installation R and Jupyter Notebook).

3 R with RStudio

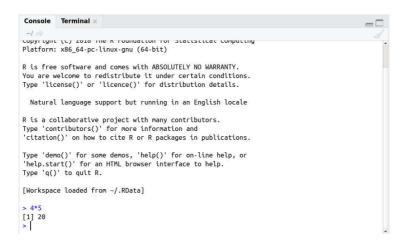
R itself has only limited (or cumbersome to be more correct) capabilities for entering and handling commands via a terminal. To get around this limitation, we recommend to use an user interface. We're going to use the interface RStudio which is the most widespread and popular.

If you open RStudio, for four panes appear.



- 1. Upper left pane: This is our working pane, i.e. for writing commands.
- 2. Lower left pane: This is the heart of R. The commands are executed and the results displayed (except graphics) in this pane.
- 3. Lower right pane: Used among others for displaying graphics.
- 4. Upper right pane: This pane lists all defined objects.

Commands can be written and executed directly in the lower left pane. For example, we type 4*5 after the >-prompt, press Return and the result is shown.



However, it is better to use the upper left pane to write the commands, because the content in this pane can be saved and then edited later.

To execute commands in this pane, highlight the commands and press ctrl+enter (pressing enter alone just produces a linebreak within the editor).

The result is again displayed in the lower left pane.



4 First steps

4.1 R as calculator

As we have just seen, we can use R as a calculator:

```
1+1

## [1] 2

or

2*4

## [1] 8
```

```
exp(3)
## [1] 20.08554
```

You can add comments to the source code, which is highly recommended. If you start a line with a hashtag #, then the text in this line is ignored during execution of the commands.

```
## [1] 2
```

Additional empty spaces are ignored during execution of the commands

```
# A first example
1 + 1
## [1] 2
```

Empty spaces are often used to enhance the readability of the code.

4.2 Assignments

Now it gets a little more interesting. We can assign values to variables. This is done with the assignment operator <-.

```
a <- 3
bb <- 4

a

## [1] 3

bb

## [1] 4
```

The value 3 is assigned to the variable a and 4 is assigned to the variable bb. Now, we can use these variables for further calculations.

```
2.7 * a

## [1] 8.1

# The ^-Operator is used for the calculation of powers
a^2

## [1] 9

a / bb

## [1] 0.75
```

Remarks:

1. The operator <- is read as one character and can't be separated.

- 2. As we have seen, variable names can consist of several characters. They *cannot* start with a number or a dot.
- 3. R is case-sensitiv, that means that the variable a and the variable A are different.

```
a <- 7
a
## [1] 7
A
## Error in eval(expr, envir, enclos): object 'A' not found</pre>
```

Variable A is not recognized by R because this variable hasn't been defined before.

4. We can also use the equality operator = for assignments.

```
a = 3
b = 4

a
## [1] 3
b
## [1] 4
```

However, this is for several reasons not recommended and we will only use the assignment operator <-.

5. Like most programming languages, R doesn't understand the expression 3a.

```
3a
## Error: <text>:1:2: unexpected symbol
## 1: 3a
## ^
```

For multiplications, we have to use the multiplication operator * (3*a).

4.3 Vectors

4.3.1 Generate vectors

Vectors are very important in \mathbb{R} . A vector¹ is a so-called *onedimensionial array*. In more familiar terms, a vector is simply a ordered list² of objects. Vectors are generated with the command $\mathfrak{c}(\ldots)$ (concatenate).

```
z_1 <- c(3, 4, 4.5, -2, 7)

z_1

## [1] 3.0 4.0 4.5 -2.0 7.0
```

¹The expression "vector" is in R more general than the one used in mathematics, where the components consist generally just of numbers. See the vector mixed in the example above.

²We have to be careful here as the expression "list" is also used in R and has a different, more general meaning. But in the beginning, you can think of vectors as ordered lists.

Remarks:

- 1. Each entry in a vector is called a *component* of this vector. In this document, we'll use component and entry interchangeably.
- 2. Colloquially, the c(...) command glues all the entries together and we can regard the vector as *one* object consisting of other objects.
- 3. As we have seen with the vector mixed, vectors can be very general. The entries doesn't have to be numbers and the same entries can occur several times.
- 4. There is no difference between '...' and "...".
- 5. The order of the components in a vector is important. In the following example the vectors **a** und **b** are *not* the same because the entries 1 and 2 are in a different order.

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

# The command "identical" checks whether two object are equal
# (result "true") or not (result "false")

identical(a, b)

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

6. Vectors like z_2, where the components are successive numbers occur quite often in R. There is a shortcut command for this kind of vectors.

```
z_2 <- 1:5
z_2
```

```
## [1] 1 2 3 4 5
a <- -3:4
a
## [1] -3 -2 -1 0 1 2 3 4
```

7. An important remark. If you look at online examples, you'll often see the variables names like z.1, i.e. dots are used within variable names. This is perfectly acceptable. It is a convention in R to use dots within variable names.

However, using dots within variable names in Python are *not* acceptable because the dot has a special use in Python.

To avoid any confusion with Python codes, it's better to use the subscript dash _ within variable names in R.

4.3.2 Operations with vectors

Simple operations

The components of the vectors z_1 and z_2 are numbers (which is often, but certainly not always the case in R) and we can perform the usual arithmetical operations with these vectors.

```
z_1 + z_2

## [1] 4.0 6.0 7.5 2.0 12.0

4 * z_1

## [1] 12 16 18 -8 28

z_1 * z_2

## [1] 3.0 8.0 13.5 -8.0 35.0
```

The result of the last multiplication of these two vectors is a vector³. The first component of the resulting vector is the product of the first components of the vectors, the second component is the product of the second components and so on.

³I.e. this product is *not* to be mistaken with the dot product.

Of course, a multiplication with the vector mixed doesn't make any sense and R produces an error message:

```
3 * mixed
## Error in 3 * mixed: non-numeric argument to binary operator
```

First statistical operations

As an example for a different operation on vectors, we wish to calculate the average (or mean) of the values in the vector z_1 . We use the R-command mean(...).

```
mean(z_1)
## [1] 3.3
```

Or we can calculate the sum of the values of the vector z_1 (or the minimum or the maximum).

```
sum(z_1)
## [1] 16.5

min(z_1)
## [1] -2

max(z_1)
## [1] 7
```

Of course, we could have done this by just taking a look at the vectors as there are just 5 components. It's a completely different story, if a vector has 100 000 entries.

The command length(...) determines the number of components in a vector.

```
length(mixed)
## [1] 6
```

The vector mixed has 6 entries.

It is often useful to access a specific component of a vector which is done by putting square brackets after the variable with a number for the corresponding component within the brackets.

```
mixed[2]
## [1] "sad"
```

The second entry of the vector mixed is sad.

Now, we want to know what the first and the third entries are. How can we do this? The obvious, but wrong idea, runs as follows:

```
mixed[1, 3]
## Error in mixed[1, 3]: incorrect number of dimensions
```

Because a vector is a one-dimensional object, the components are accessed just by *one* number within the square brackets. If we want to access several components at the same time, we exchange the *one* number in the square brackets with *one* other object and that is in this case the vector c(1, 3):

```
mixed[c(1, 3)]
## [1] "2" "5.3"
```

If we want to access components 2, 3, 4, 5, we can use, as we have seen, a shortcut command: 2:5.

```
mixed[2:5]
## [1] "sad" "5.3" "YES" "beautiful"
```

4.3.3 R -commands with options

To illustrate a command with an option in R, we create the following vector:

```
z_3 < c(5, 2, NA, 4)
```

The "value" NA (not available) occurs often in data science. These are "values", which for some reason are unknown. For example, in a survey some persons didn't want to indicate their age. These are the NA's in this case.

If we want to determine the minimum of the vector **z_3**, R produces the following result:

```
min(z_3)
## [1] NA
```

R tries to find the minimum value in this vector, but it doesn't know what to do with the NA, so it returns NA.

But we can still try to find the minimum value of the *existing* numbers in this vector. This is done with a so-called *option* which we put additionally within the round brackets of the function call:

```
min(z_3, na.rm = TRUE)
## [1] 2
```

In this case the option is na.rm = ... and means *na remove*. If we set this option TRUE, then all NA's are removed for the calculation of the minimum, if FALSE then the original vector is used, including the NA.

All options have a default setting. In our example, the default setting is na.rm = FALSE (or simply F). That means, if we don't add the option to the command, R sets na.rm = FALSE.

```
min(z_3)

## [1] NA

min(z_3, na.rm = FALSE)

## [1] NA
```

Most of the time, commands have several options. We can check which options are available for a command, including the default setting, using

?min

or

help(min)

Extremes {base} R Documentation

Maxima and Minima

Description

Returns the (regular or parallel) maxima and minima of the input values.

pmax*() and pmin*() take one or more vectors as arguments, recycle them to common length and return a single vector giving the 'parallel' maxima (or minima) of the argument vectors.

Usage

```
max(..., na.rm = FALSE)
min(..., na.rm = FALSE)
pmax(..., na.rm = FALSE)
pmin(..., na.rm = FALSE)
pmax.int(..., na.rm = FALSE)
pmin.int(..., na.rm = FALSE)
```

Arguments

... numeric or character arguments (see Note).

na.rm a logical indicating whether missing values should be removed.

4.4 Data sets (twodimensional)

A lot of data are given in the form of twodimensional data sets (tables). The following table consist of fictional temperatures in several Swiss cities during several month.

	Luzern	Basel	Chur	Zurich
Jan	2	5	-3	4
Feb	5	6	1	0
Mar	10	11	13	8
Apr	16	12	14	17
May	21	23	21	20
Jun	25	21	23	27

Such data sets are usually stored in files. In this case the file is called **weather.csv** (on ILIAS), which you can download and save in a suitable folder⁴.

A huge source for data sets is:

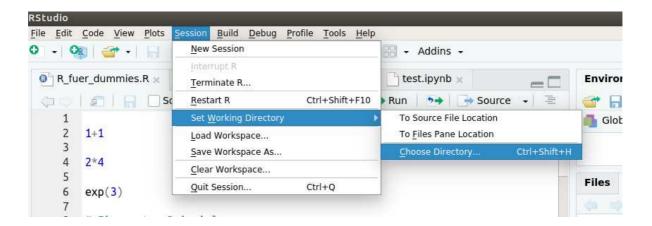
 $^{^4}$ The folder path should't contain empty spaces or umlauts as ä, ö, ü or other special characters. R handles these badly.

4.4.1 Loading (importing) data sets

To load data sets, it's recommended that you set your working directory first. This can be done within RStudio or manually. We highly recommend that you do this with RStudio, especially for Windows user.

RStudio

Go to (see figure below) Session, Set Working Directory, Choose directory, then click your way through to folder you want to work with, then Open. RStudio will set the working directory automatically.



Manually

With the setwd(...)-command ("set working directory"), we specify the working directory in which the file weather.csv has been saved. This is in *my* case /home/bl/Dropbox/Statistics/Software_R_Python/R.

setwd("/home/bl/Dropbox/Statistics/Software_R_Python/R")

Important for Windows user

Windows uses backslashes \ instead of slashes / in their pathes. So your path looks something like

C:\Users\Documents\folder\R

This path doesn't work in R. To get around this, you have to double the backslashes:

setwd("C:\\Users\\Documents\\folder\\R")

Loading data sets

Once we have set the working directory, we can load (import) the data set with the command read.csv(...).

```
temp <- read.csv("weather.csv")</pre>
temp
  Luzern Basel Chur Zurich
## Jan 2 5 -3
## Feb
        5
             6
                  1
                        0
## Mar
        10 11
                        8
                 13
        16 12 14
## Apr
                       17
        21 23
                 21
                       20
## May
## Jun
         25
             21
                  23
                       27
```

We can also add the file path directly to the file name:

```
temp <- read.csv("/home/bl/Dropbox/Statistics/Software_R_Python/R/weather.csv")
temp</pre>
```

In both cases the content of the table was assigned to the variable temp (for temperature). The assignment of data sets to variables is very useful as we will see shortly.

Remarks:

1. Once the table is imported, the table is then referred in R-terminology as *data frame*. That's the reason, why data frames are sometimes assigned to the variable df.

```
df <- ...
```

You see this quite often in blogs or newsgroups. It's important keep in mind that df is a variable and *not* a command.

- 2. There exist several commands for loading files. By now, a lot of data sets are available in the .csv-format, so we can use the command read.csv(...).
- 3. There is a more general command read.table(...), which can load other file formats as well. Let's try this out for our file weather.csv:

```
temp <- read.table("weather.csv")
temp</pre>
```

```
##
         V1
## 1 Luzern , "Basel", "Chur", "Zurich"
## 2
        Jan
                             ,2,5,-3,4
## 3
        Feb
                             ,5,6,1,0
## 4
        Mar
                           ,10,11,13,8
## 5
        Apr
                         ,16,12,14,17
## 6
                         ,21,23,21,20
        May
## 7
        Jun
                          ,25,21,23,27
```

This doesn't looks right, something must have gone wrong. The table doesn't look "nice". There are two reasons for this behaviour:

a) The column names are V1 and V2, which is not correct. These should be Luzern, Basel, Chur and Zurich.

We have to force read.table(...) to accept the first line as column names. This is done with the option header = T

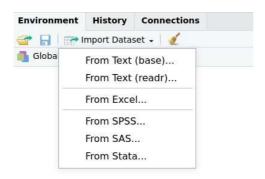
```
temp <- read.table("weather.csv", header = T)</pre>
temp
##
     Luzern X..Basel...Chur...Zurich.
        Jan
                              ,2,5,-3,4
## 1
## 2
        Feb
                               ,5,6,1,0
## 3
        Mar
                            ,10,11,13,8
                           ,16,12,14,17
## 4
        Apr
## 5
        May
                           ,21,23,21,20
## 6
        Jun
                           ,25,21,23,27
```

b) The table still doesn't look right. The entries appear to be separated by commas, but read.table(...) expects by default that the entries in the table are separated by empty spaces.

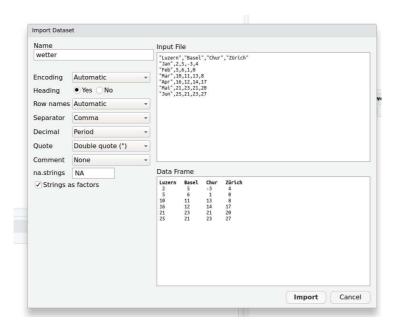
However, it is possible to load files of data sets which are separated by commas. This is done with th option sep = ",".

```
temp <- read.table("weather.csv", sep = ",")</pre>
temp
      Luzern Basel Chur Zurich
          2 5
                    -3
## Jan
## Feb
          5
               6
                    1
                            0
                    13
          10
                            8
## Mar
               11
          16
               12
                           17
## Apr
                    14
               23 21
## May
          21
                           20
```

4. There is another possibility to import (load) files, which doesn't work properly for non-.csv-files⁵. In RStudio click on in the upper right pane Import Dataset, then click on the first entry (might look different for different operating systems).



Then click through to your folder and click on the desired file. The following window appears.



On the upper right side, the original table appears (Input File). The table on lower right side (Data Frame) looks as it should be. If that's not the case, you have to change the parameters on the left hand side (change for example Separator and see what happens).

⁵csv: comma separated values

4.4.2 Accessing colums and rows

An important remark at the beginning: the first row and first column are *not* part of a data frame. These are used to access of the individual data, whole (or parts of) rows and columns.

A very useful command is the head(...) command. The output are by default the first 6 rows (in our example a bit useless because the table has only 6 rows). With this command we can check whether the file has been correctly imported.

```
head(temp)
    Luzern Basel Chur Zurich
## Jan 2 5 -3
## Feb
       5
            6
                1
                      0
        10 11
## Mar
                13
        16 12 14
## Apr
                     17
## May
        21
            23
                21
                     20
        25
                23
                     27
## Jun
            21
```

The size of a data set is often unknown. If we want to know the number of rows and columns in a data set, we use the command dim(...):

```
dim(temp)
## [1] 6 4
```

Important: The first value is the number of rows, the second value the number of columns. In the example above, we have 6 rows and 4 columns.

We obtain the column names with the command colnames(...) and the row names rownames(...)

```
colnames(temp)

## [1] "Luzern" "Basel" "Chur" "Zurich"

rownames(temp)

## [1] "Jan" "Feb" "Mar" "Apr" "May" "Jun"
```

For a first overview of the data, we use the command summary(...):

```
summary(temp)
##
      Luzern
                    Basel
                                   Chur
                                                 Zurich
  Min. : 2.00 Min. : 5.00 Min. :-3.00
                                             Min. : 0.00
  1st Qu.: 6.25   1st Qu.: 7.25   1st Qu.: 4.00
##
                                             1st Qu.: 5.00
  Median :13.00
                Median :11.50
                               Median :13.50
                                             Median :12.50
##
  Mean :13.17
                Mean :13.00
                               Mean :11.50
                                             Mean :12.67
##
##
  3rd Qu.:19.75
                 3rd Qu.:18.75
                               3rd Qu.:19.25
                                             3rd Qu.:19.25
##
  Max. :25.00
                 Max. :23.00
                               Max. :23.00
                                             Max. :27.00
```

Min. is the minimal value, 1st Qu. the first (or lower) quartile (see lecture notes), Median the median (see lecture notes), Mean the mean (average), 3rd Qu. the third or upper quartile and Max.the maximal value of the corresponding columns.

Now, we want to know for example what the temperature in February in Zurich was. There a various ways to do that.

```
temp[2, 4]
## [1] 0
```

We access the value in the 2nd row and the 4th column.

Important: The first number *always* correspond to the row(s) and the second to the column(s)⁶.

However, for large data frames this method is not practicable. It's much easier to access the values with their row and column name(s):

```
temp["May", "Basel"]
## [1] 23
```

This is the temperature in May in Basel.

We can also read whole rows or columns:

```
temp["Jun", ]
## Luzern Basel Chur Zurich
## Jun 25 21 23 27
```

⁶This convention is also used for entries in matrices.

```
temp[, "Basel"]
## [1] 5 6 11 12 23 21
```

For columns, the following variant does the same (well, almost).

```
temp["Basel"]

## Basel

## Jan 5

## Feb 6

## Mar 11

## Apr 12

## May 23
## Jun 21
```

We can calculate the mean of a column:

```
mean(temp[, "Luzern"])
## [1] 13.16667
```

An important task is to save data sets or a part of it. This can be done with the command write.csv(...) for .csv-files.

```
temp1 <- temp[c("Jan", "May"), c("Chur", "Zurich")]

temp1

## Chur Zurich
## Jan -3 4
## May 21 20

write.csv(temp1, file = "weather1.csv")</pre>
```

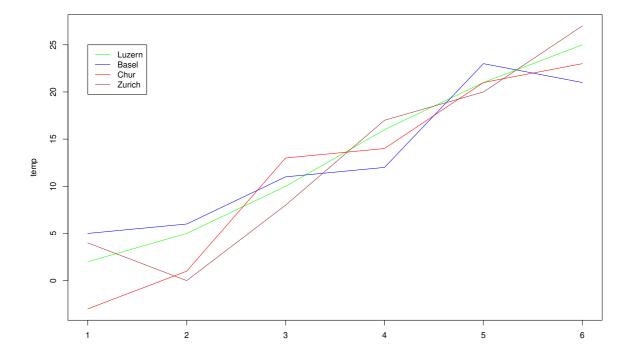
Are we interested in a specific separation of the values in the data set, we use the appropriate option sep = By default, entries in a csv file are separated with commas. For other write command the default setting is generally different. To be on the safe side, always set sep = ",".

```
write.csv(temp1, file = "weather1.csv", sep = ",")
```

4.4.3 A example for a plot

An important tool in statistics is the graphical representation of data. Here we give just a simple example, you'll learn many more applications of plots during this course.

```
color <- c('green', 'blue', 'red', 'brown')
matplot(temp, col = color, type = "l", lty = 1)
legend(x = 1, y = 25, legend = colnames(temp), col = color, lty = 1)</pre>
```



The first command

```
farben <- c('green', 'blue', 'red', 'brown')</pre>
```

defines a vector with colours as entries. We'll use this vector in the 2nd and 3th command.

The second command

```
matplot(temp, col = color, type = "1", lty = 1)
```

plots the graphs of the temperature in the columns. The option col = ... stands for *colors* which means that the graphs are colored according the vector col = color. The vector color has been defined above.

The options type = "..." and lty = ... are explained in the exercises.

The third command

```
legend(x = 1, y = 25, legend = colnames(temp), col = color, lty = 1)
```

put the legend in upper left corner. The first two arguments 1 and 25 are the coordinates of the position of the legend. The option legend = colnames(temp) designate the names for the lines, which in this case are obviously the column names of the data frame.