



R Basics

Marco Torchiano

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Introduction

What is R?



<http://cran.r-project.org/>

R is a free software environment for statistical computing and graphics.

Available on several different platform

Basic features

- CLI
 - Command Line Interface
 - Immediate evaluation of expression
- Scripts
- Extensive help system <https://www.rseek.org/>
- Large resource set online <https://stackoverflow.com/>

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IDE

- Several graphical front-ends (GUI)
- RStudio is a full IDE for R



<http://www.rstudio.com>

- Also cloud version: <https://rstudio.cloud>

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R Console

Basic text based REPL

- Read: from the user keyboard
 - Or from a script
- Evaluate the R language expression
- Print the result of the evaluation
- Loop
 - Until `quit()`

Objects are stored in a common environment

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R environment

- Shared global memory space where all objects stored
 - Variables
 - Functions
- Can be inspected at any time
- Every time a command assign a value to a variable, it is placed inside the environment
- All valuee in the environment are available to any later statement

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R script

- A text file containing commands intended to be executed as a whole
- It is possible to execute the statements one by one
 - The result is the same
- Execution means taking a statement from the script instead of reading it from the keyboard
 - Accesses the global environment

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R package

- Library of functions designed to work together
 - Include documentation
- Can be installed from R official repository (CRAN)
 - From CLI: `install.packages("ggplot2")`
 - From GUI: Tools > Install packages...
- Must be loaded before use
 - `library("ggplot2")`

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R help

- All built-in and package functions are documented
- Help system is integrated in
 - Console
 - Help on function `? log`
 - Search for topic: `?? logarithm`
 - R Studio
 - Help pane

R elements

- Statements
 - assignment
 - expression
 - control
- Functions
- Variables
- Data types
 - Primitive
 - Compound

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Statements

Statements can be terminated by

- a *new-line* : most common
- a `;`
 - to avoid ambiguities
 - to put multiple statements on a single line

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Comments

On any line from `#` until end of line is considered comments. Typical usage:

- `#` as first character: comment line
- `#` after statement: comment specific statement

```
#----- Define constants -----  
#  
PI <- 7/22 # a reasonable approximation
```

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Assignment

The global environment stores objects, e.g. values

Operator `<-` is used to store an object with a name

```
answer <- "fortytwo"
```

Variables are not *typed*

- i.e. you can (re-)assign any type of value

```
answer <- 42
```

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Assignment

Assignment operator `<-` copies the value of an expression into the environment and assign a name

- Operator `=` can be used instead
- Non recommended to avoid confusion

An assignment overwrites the value previously linked to that name

- Be careful with names

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Names

- Variable names
 - Must start with a letter,
 - Can't contain spaces
- Style recommendations:
 - Use lowercase characters
 - Use an underscore (`_`) to separate words
 - Avoid using names that are predefined

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Expression

- When an expression is entered, R evaluates it and prints the result
- Uses the names to retrieve values from the environment

```
answer
```

```
## [1] 42
```

- it is possible to explicitly force printing an expression with `print()`

```
print( (answer / 3) %% 11)
```

```
## [1] 3
```

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Primitive types

numeric

- Default type (also for integer values)
- Uses standard IEEE-754 (ISO/IEC 60559)
 - E.g., `1.2` , `1`

integer

- Used to force integer arithmetic
- Suffix letter “`L`” to force integer
 - E.g., `42L`

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Primitive types

complex

- Allow complex number operations

```
sqrt( -1 + 0i )
```

```
## [1] 0+1i
```

logical

- Keywords: `TRUE` | `FALSE`
- Also predefined variables: `T` and `F`

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Primitive types

character

- String of characters
- Can be described using both
 - `'single'` and
 - `"double"` quotes

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Data types

Type-related functions:

- Type of variable: `class(x)`
- Check type: `is.type(x)`
- Conversion: `as.type(x)`

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Special values

- **NA** : is generally interpreted as a missing, does not exist
 - Stands for Not Available
 - tested with `is.na()`
- **NULL** : is for empty object
 - tested with `is.null()`
- **NaN** : the result is not a number, e.g. `log(-1)`
 - Stands for **Not-a-N**
 - tested with `is.nan()`
- **Inf** numeric infinity ∞ , e.g. `1/0`

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Operators

- Arithmetic on numeric: `+`, `-`, `*`, `/`, `^`
 - integer `%%` (modulo)
- Comparison: `==`, `!=`, `<`, `<=`, `>`, `>=`
 - works also on strings

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Character operations

- `nchar()` : length of the string
- `paste(..., sep=" ")`: concatenates with separator
 - `paste0(...)`: no separator, i.e. `sep=""`

```
nchar("Visualization")
```

```
## [1] 13
```

```
paste("Visualization", "of", "Quantitative", "Informatic
```

```
## [1] "Visualization of Quantitative Information"
```

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Character operations

- `substr()` : extract and replaces portion of a string

```
title <- "Visualization of Quantitative Information"  
substr(title, 15, 16)
```

```
## [1] "of"
```

```
substr(title, 15, 16) <- "OF"  
title
```

```
## [1] "Visualization OF Quantitative Information"
```

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Block statements

A series of statements can be gathered in a block using the `{ ... }` syntax.

- they are treated as a single (compound) statement
- non new environment is created

Block statement are used as branches or bodies of structured control statements.

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Control statements

- `if(cond) .. else`
- `while(cond)`
- `for(var in seq)`

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Conditional

Use the usual syntax: `if(cond) ... else ...`

- else clause is optional

```
a <- 10
if( a < 0){
  "negative"
}else{
  "positive"
}
```

```
## [1] "positive"
```

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While loop

Use the `while(cond) ...` syntax

```
a <- 10
while( a > 1){
  print(a)
  a <- a / 2;
}
```

```
## [1] 10
## [1] 5
## [1] 2.5
## [1] 1.25
```

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Functions definition

Using the keyword `function`

```
percentage <- function(part,whole){
  part/whole*100
}
```

- return evaluation of last expression
- or can use `return()` statement

Can provide default values:

```
percentage <- function(part=1, whole=1){
  return( part/whole*100 )
}
```

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Function invocation

Usual invocation (positional)

```
percentage(3, 4)
```

```
## [1] 75
```

Named arguments:

```
percentage(whole=4, part=3)
```

```
## [1] 75
```

Leverage default values:

```
percentage(part=0.75)
```

```
## [1] 75
```

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Exercise 1

Define a function `pythagoras()` accepting three values (a,b,c) one of which can be missing and is computed using the Pythagorean theorem.

```
pythagoras(3, 4)
```

```
## [1] 5
```

```
pythagoras(c=5, a=3)
```

```
## [1] 4
```

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Vectors

Vectors

- All values in R are considered as vectors
 - Possibly with dimension 1 for scalar values
- When printed
 - if spread on many lines, the index of the first element printed on the line is shown in `[]`
 - for a scalar, `[1]` is shown indicating the index of the first and only element
- All elements in a vector must have the same type
 - Type coercion can be applied

Vector creation

- With *combine* function `c()` by enumeration of elements

```
v <- c(2,4,5)
```

- Remember also scalars are vector: `1 == c(1)`

- With `vector()` function, with type and length, creates a zero-ed vector

```
w <- vector("numeric",3)
```

```
w
```

```
## [1] 0 0 0
```

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Ranges

Range operator `:` generates an integer vector

```
1:3
```

```
## [1] 1 2 3
```

- equivalent to

```
c(1L, 2L, 3L)
```

```
## [1] 1 2 3
```

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Vector operations

- Merging:

```
c( 1:3, 7:9)
```

```
## [1] 1 2 3 7 8 9
```

- Type coercion can be applied

- Length with function `length()`

```
length( 1:10 )
```

```
## [1] 10
```

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Vector operations

Arithmetic operators

- Pair-wise on same-index elements

```
1:3 + 3:1
```

```
## [1] 4 4 4
```

- Recycling if different size

```
1:3 + 1
```

```
## [1] 2 3 4
```

- Longest length must be multiple of shortest

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Empty vectors

- Using primitive types function to create empty (typed) vectors

```
empty_numeric <- vector("numeric",0)
length(empty_numeric)
```

```
## [1] 0
```

```
empty_numeric
```

```
## numeric(0)
```

- The combine function without arguments gives `NULL`
 - The reason is that no type is specified

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Vector access

- Operator `[]`
- Uses an index to access an element

In R, indexes start at 1!!!

```
s = c("aa", "bb", "cc", "dd", "ee")
s[1]
```

```
## [1] "aa"
```

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Vector access

- With index 0 returns an empty vector

```
s[0]
```

```
## character(0)
```

- Out of bound returns `NA`

```
s[6]
```

```
## [1] NA
```

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Vector slicing

Slicing allows extracting a subset of the vector elements

- Using a vector of indexes

```
s[ c(1,3) ]
```

```
## [1] "aa" "cc"
```

- Indexes can be repeated

```
s[c(5,1,1)]
```

```
## [1] "ee" "aa" "aa"
```

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Vector slicing

- Using a vector of logicals

```
l <- c(TRUE, FALSE, FALSE, FALSE, TRUE)
```

```
s[ l ]
```

```
## [1] "aa" "ee"
```

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For-Loops

For-loop syntax: `for(variable in vector)`

- in each iteration the `variable` will assume all the consecutive values in the `vector`.

```
min <- 100;
for( d in c(7,2,5,10,20,12,3) ){
  if( d < min)
    min <- d
}
min
```

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

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For-Loops

Iteration on a vector can be implemented also with an index over a range

```
min <- 100;
numbers <- c(7,2,5,10,20,12,3)
for( i in 1:length(numbers) ){
  if( numbers[i] < min)
    min <- numbers[i]
}
min
```

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

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Loop control

- **break**: steps out of the loop skip rest of body
- **next**: skips remaining of the body and start new iteration

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Named vectors

- Elements of a vector can be named

```
days<-c(Jan=31, Feb=28, Mar=31, Apr=30, May=31, Jun
        Jul=31, Aug=31, Sep=30, Oct=31, Nov=30, Dec
```

- When printed, names are reported above the values

```
days
```

```
## Jan Feb Mar Apr May Jun Jul Aug Sep Oct Nov Dec
##  31  28  31  30  31  30  31  31  30  31  30  31
```

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Named vectors

- Names can be used instead of indexes

```
days[ "Feb" ]
```

```
## Feb
##  28
```

- Also for slicing purposes

```
days[ c( "Feb", "Dec" ) ]
```

```
## Feb Dec
##  28  31
```

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Named vectors

Function `names()` access names

- Allows getting and setting names

```
names(days)
```

```
## [1] "Jan" "Feb" "Mar" "Apr" "May" "Jun" "Jul" "Au"
## [9] "Sep" "Oct" "Nov" "Dec"
```

```
triplet <- 1:3
names(triplet) <- c("one", "two", "three")
triplet
```

```
## one two three
## 1 2 3
```

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Exercise 2

Modify the `pythagoras()` function so that it returns a vector with three elements named 'a', 'b', and 'c' according to the Pythagorean theorem.

```
pythagoras(3,4)
```

```
## a b c
## 3 4 5
```

```
pythagoras(c=5,a=4)
```

```
## a b c
## 4 3 5
```

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Character vector

- `strsplit(s, split)`: creates a list of vectors of strings by splitting at given separator

```
strsplit(title, " ")
```

```
## [[1]]  
## [1] "Visualization" "OF" "Quantitative"  
## [4] "Information"
```

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Character vector

Function `paste(..., sep=" ", collapse)`:

- first concatenates strings at corresponding indexes (w/recycling) with separator
- then concatenates elements of the resulting vector

```
paste(1:3, c("one", "two", "three"), ".")
```

```
## [1] "1 one ." "2 two ." "3 three ."
```

```
paste(1:3, c("one", "two", "three"), ".", collapse=" - ")
```

```
## [1] "1 one . - 2 two . - 3 three ."
```

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Sequences

Function `seq(from, to, by, length.out)` allows different combination of arguments

```
seq(1, 10) # by=1
```

```
## [1] 1 2 3 4 5 6 7 8 9 10
```

```
seq(1, 10, by=3)
```

```
## [1] 1 4 7 10
```

```
seq(1, 10, length.out=4)
```

```
## [1] 1 4 7 10
```

```
seq(1, length.out=10) # by=1
```

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Type coercion

When putting values of different type in the same vector they are (silently) coerced to the same type

- the most general type among the elements is used
- `character` > `complex` > `numeric` > `integer` > `logical`

```
c(3, "two", TRUE)
```

```
## [1] "3" "two" "TRUE"
```

```
c(22/7, 42L, FALSE)
```

```
## [1] 3.142857 42.000000 0.000000
```

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Type coercion

Coercion is performed using the conversion functions

```
as.type()
```

Not always conversion is possible, in such cases `NA` is produced

```
as.numeric(c("1", "b", "3.2"))
```

```
## Warning: NAs introduced by coercion
```

```
## [1] 1.0 NA 3.2
```

```
as.logical(c("true", "FALSE", "T", "V", "0"))
```

```
## [1] TRUE FALSE TRUE NA NA
```

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Logical to Numeric

When summing, logicals are coerced to integers

- `TRUE` → 1, `FALSE` → 0

```
thirty <- days == 30  
thirty
```

```
##   Jan   Feb   Mar   Apr   May   Jun   Jul   Aug  
## FALSE FALSE FALSE TRUE  FALSE TRUE  FALSE FALSE T  
##   Oct   Nov   Dec  
## FALSE TRUE  FALSE
```

```
sum(thirty) # how many (coercion: T->1 F->0)
```

```
## [1] 4
```

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Filtering vectors with logicals

```
names(days)[thirty]
```

```
## [1] "Apr" "Jun" "Sep" "Nov"
```

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Filtering vectors with indexes

Function `which()` returns indexes of element satisfying condition (`==TRUE`)

```
thirty.ix <- which( days==30 )  
thirty.ix
```

```
## Apr Jun Sep Nov  
##  4  6  9  11
```

```
names(days)[thirty.ix]
```

```
## [1] "Apr" "Jun" "Sep" "Nov"
```

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Sorting

Data in a vector can be sorted using function `sort()`

- Note: the original array is not modified

```
numbers <- c(3, 7, 14, 2, 5, 8)
sort(numbers)
```

```
## [1] 2 3 5 7 8 14
```

```
words <- c("There", "must", "be", "some", "kind",
           "of", "way", "out", "of", "here")
sort(words)
```

```
## [1] "be"      "here"    "kind"    "must"    "of"      "of"
## [7] "out"     "some"    "There"   "way"
```

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Ordering the indexes

Function `order()` sorts the **indexes** based on the value of the corresponding elements

- the first element of the result contains the index of the smallest element
- slicing with the ordered indexes gives a sorted vector

```
order(numbers)
```

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

```
numbers[ order(numbers) ] # slicing in order
```

```
## [1] 2 3 5 7 8 14
```

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Ranking

Function `rank()` computes the ranks of the corresponding elements

```
r <- rank(numbers)
names(r) <- numbers
r
```

```
## 3 7 14 2 5 8
## 2 4 6 1 3 5
```

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Matching

Operator `%in%` finds which element of left-hand vector are present in the right-hand one.

```
c("John", "Jane", "Mike", "Iris") %in% c("Jane", "Iris", "
```

```
## [1] FALSE TRUE FALSE TRUE
```

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Vectorization

Often it is useful to apply a function to all elements in a vector

A *vectorized* function is one that can apply the same operation to all elements of its argument

- It is much easier to use and more efficient
- Most builtin functions are vectorized

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Vectorization vs. loops

Specific functions are not always vectorized

```
score_to_grade <- function(score){  
  if(score<17.5) "Failed"  
  else if(score>=30.5) "30L"  
  else round(score)  
}  
scores <-c(15,24.3,32,27.5)  
score_to_grade(scores)
```

```
## Warning in if (score < 17.5) "Failed" else if (score >= 30.5) "30L" else round(score): the condition has length > 1 and only the first element will be used
```

```
## [1] "Failed"
```

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Vectorization vs. loops

A loop can be used to apply to all elements

```
grades <- numeric( length(scores) )
for( i in 1:length(grades)){
  grades[i] = score_to_grade(scores[i])
}
grades
```

```
## [1] "Failed" "24"      "30L"      "28"
```

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Vectorization functionals

A *functional* is a function that applies another function

Functional `sapply()`:

- takes a vector and a function
- applies the function to all elements of the vector
- collects the results into a vector

```
grades <- sapply(scores,score_to_grade)
grades
```

```
## [1] "Failed" "24"      "30L"      "28"
```

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Composed data types

Matrix

Construction:

```
matrix(1:9, 3, 3)
```

```
##      [,1] [,2] [,3]
## [1,]    1    4    7
## [2,]    2    5    8
## [3,]    3    6    9
```

```
A <- matrix(1:9, 3, 3, byrow=TRUE); A
```

```
##      [,1] [,2] [,3]
## [1,]    1    2    3
## [2,]    4    5    6
## [3,]    7    8    9
```

Matrix indexing

Indexes start at 1, like vectors.

```
A[2, 3] # single cell
```

```
## [1] 6
```

```
A[2, ] # row
```

```
## [1] 4 5 6
```

```
A[ , 3] # column
```

```
## [1] 3 6 9
```

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Matrix indexing

```
A[2,3] <- 66; A
```

```
##      [,1] [,2] [,3]
## [1,]    1    2    3
## [2,]    4    5   66
## [3,]    7    8    9
```

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Matrix transposition

```
B <- t(A); B
```

```
##      [,1] [,2] [,3]
## [1,]    1    4    7
## [2,]    2    5    8
## [3,]    3   66    9
```

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Matrix Composition

```
cbind(A,B) # column-wise
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6]
## [1,]    1    2    3    1    4    7
## [2,]    4    5   66    2    5    8
## [3,]    7    8    9    3   66    9
```

```
rbind(A,B) # row-wise
```

```
##      [,1] [,2] [,3]
## [1,]    1    2    3
## [2,]    4    5   66
## [3,]    7    8    9
## [4,]    1    4    7
## [5,]    2    5    8
## [6,]    3   66    9
```

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List

An array whose element can be of different types

- both primitive and compound types

Construction:

```
l <- list(c(1,2), "a"); l
```

```
## [[1]]  
## [1] 1 2  
##  
## [[2]]  
## [1] "a"
```

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List named members

Usually list members are named

```
l <- list( n=c(1,2), char="a" ) ; l
```

```
## $n  
## [1] 1 2  
##  
## $char  
## [1] "a"
```

```
names(l)
```

```
## [1] "n" "char"
```

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List access

Access to a member uses the *accessor* operator `$`, or the element indexing operator `[]`.

```
l$n; l[["n"]] ; l[[1]]
```

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

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

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

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List access

Access operators can be used to change and existing element or to add a new one if the name is not present

```
l$char = "B"  
l$logicals = c( TRUE, FALSE, TRUE )  
l
```

```
## $n  
## [1] 1 2  
##  
## $char  
## [1] "B"  
##  
## $logicals  
## [1] TRUE FALSE TRUE
```

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List slicing

Slicing return a subset of the list:

```
l[2]
```

```
## $char  
## [1] "B"
```

Indexing returns the element

```
l[[2]]
```

```
## [1] "B"
```

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Exercise 3

Modify the `pythagoras()` function so that it accepts a list with two elements named 'a', 'b', or 'c' and computes the missing one, according to the Pythagorean theorem.

```
pythagoras.list(list(a=3,b=4))
```

```
## $a  
## [1] 3  
##  
## $b  
## [1] 4  
##  
## $c  
## [1] 5
```

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Factor

Represent nominal variables

- Internally stored as integer vector
- created using the `factor()` function

```
f = factor( c("Red", "Green", "Blue", "Blue",  
             "Red", "Red"))  
f
```

```
## [1] Red   Green Blue  Blue  Red   Red  
## Levels: Blue Green Red
```

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Factor

Levels:

```
levels(f)
```

```
## [1] "Blue" "Green" "Red"
```

Frequencies:

```
table(f)
```

```
## f  
## Blue Green Red  
##    2    1    3
```

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Ordered factors

```
f = factor( c("L", "M", "L", "H", "L", "H", "L"),  
           levels=c("L", "M", "H"), ordered=T)
```

```
f
```

```
## [1] L M L H L H L
```

```
## Levels: L < M < H
```

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Dataframe

It is the main data structure used to represent tabular datasets.

- Most data is processed in the form of dataframes
- Most I/O of data handle dataframes
- It is a list of vectors of equal length
- Typical semantic
 - each row is case or observation
 - each column is an attribute or variable

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Dataframe

Construction:

```
courses <- data.frame(  
  code = c("15AHM", "12BHD", "16ACF",  
          "01PNN", "01RKC", "17AXO"),  
  course= c("Chemistry", "Computer science", "Calculus  
           "Free Credits", "Linear Algebra", "Physics  
semester = c(1,1,1,2,2,2),  
credits = c(8,8,10,6,10,10)  
)
```

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Dataframe example

code	course	semester	credits
15AHM	Chemistry	1	8
12BHD	Computer science	1	8
16ACF	Calculus I	1	10
01PNN	Free Credits	2	6
01RKC	Linear Algebra	2	10
17AXO	Physics I	2	10

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Dataframe indexing

Column (attribute/variable) selection is usually performed with the *accessor* operator `$`

- list-specific syntax can be used also

```
courses$credits; courses[[4]]; courses[["credits"]]
```

```
## [1]  8  8 10  6 10 10
```

```
## [1]  8  8 10  6 10 10
```

```
## [1]  8  8 10  6 10 10
```

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Dataframe indexing and slicing

Cell indexing is similar to matrixes

```
courses[2,2]
```

```
## [1] "Computer science"
```

Dataframe slicing works like lists

```
courses[c("semester", "credits")]
```

```
##   semester credits
## 1         1       8
## 2         1       8
## 3         1      10
## 4         2       6
## 5         2      10
## 6         2      10
```

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Slicing dataframe by row

```
courses[c(1,3,6) , ]
```

##	code	course	semester	credits
## 1	15AHM	Chemistry	1	8
## 3	16ACF	Calculus I	1	10
## 6	17AXO	Physics I	2	10

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Sorting a dataframe

Order and slice

```
ord <- order( - courses$credits) # - means descending  
courses[ord, ]
```

##	code	course	semester	credits
## 3	16ACF	Calculus I	1	10
## 5	01RKC	Linear Algebra	2	10
## 6	17AXO	Physics I	2	10
## 1	15AHM	Chemistry	1	8
## 2	12BHD	Computer science	1	8
## 4	01PNN	Free Credits	2	6

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Filtering a dataframe with logicals

Use a logical *indicator* vector (TRUE for matching rows)

```
sem.2nd.ind <- courses$semester == 2  
sem.2nd.ind ## which courses are in 2nd semester
```

```
## [1] FALSE FALSE FALSE TRUE TRUE TRUE
```

```
courses.2nd <- courses[sem.2nd.ind, ]  
courses.2nd ##2nd semester courses
```

```
##      code      course semester credits  
## 4 01PNN  Free Credits         2         6  
## 5 01RKC Linear Algebra         2        10  
## 6 17AXO   Physics I           2        10
```

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Filtering and summing

```
sum( courses.2nd$credits ) ## 2nd semester credits
```

```
## [1] 26
```

```
sum( sem.2nd.ind ) ## how many courses in 2nd semeste
```

```
## [1] 3
```

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Filtering a dataframe with indexes

Use a the function `which()`

```
sem.2nd.ix <- which( courses$semester == 2 )  
sem.2nd.ix ## indexes of courses are in 2nd semester
```

```
## [1] 4 5 6
```

```
courses.2nd <- courses[sem.2nd.ix, ]  
courses.2nd ##2nd semester courses
```

```
##      code      course semester credits  
## 4 01PNN  Free Credits         2         6  
## 5 01RKC Linear Algebra         2        10  
## 6 17AXO   Physics I           2        10
```

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Reading files

Functions `read.*`

- Read data from a file into dataframe
- Space separated: `read.table()`
- CSV: `read.csv()`
- Clipboard: `read.table(pipe(...))`
 - X11: `"clipboard"`
 - OS X: `"pbpaste"`
- Excel file: `read.xlsx()`
 - require `library(readxl)`

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R Advantages

- R is a common tool among data experts, supported wildly by both professional and academic developers
- R can be installed in any environment on any machine and used with no licensing or agreements needed
- R source code is flexible and can be adapted to specific local needs
- R can build routines straight out of a database for common and universal reporting

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R Limitations

- R is based on S, which is close to 40 years old
- R only has features that the community contributes
- Not the ideal solution to all problems
- R is a programming language and not a software package – steeper learning curve
- R can be much slower than compiled languages

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Software

- R
 - Download at: <https://cran.r-project.org>
- R-Studio Desktop
 - Download at: <https://rstudio.com/products/rstudio/>

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References

- R. Irizarry. "Introduction to Data Science - Data Analysis and Prediction Algorithms with R"
 - <https://rafalab.github.io/dsbook/>
- H.Wickham, G.Grolemund. "R for Data Science - Visualize, model, transform, tidy, and import data", O'Reilly, 2017
 - <https://r4ds.had.co.nz/index.html>

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Solutions

Solution to Exercise 1

Define a function `pythagoras()` accepting three values (a,b,c) one of which can be missing and is computed using the Pythagorean theorem.

```
pythagoras <- function(a=NULL, b=NULL, c=NULL){  
  if(is.null(a)){  
    sqrt(c^2-b^2)  
  }else if(is.null(b)){  
    sqrt(c^2-a^2)  
  }else if(is.null(c)){  
    sqrt(a^2+b^2)  
  }  
}
```

Solution to Exercise 2

Modify the `pythagoras()` function so that it returns a vector with three elements named 'a', 'b', and 'c' according to the Pythagorean theorem.

```
pythagoras <- function(a=NULL, b=NULL, c=NULL){
  nn <- is.null(a) + is.null(b) + is.null(c)
  if(nn!=1) stop("Exactly one among 'a', 'b', 'c' must be NULL")
  if(is.null(a)){
    c(a=sqrt(c^2-b^2), b=b, c=c)
  }else if(is.null(b)){
    c(a=a, b=sqrt(c^2-a^2), c=c)
  }else if(is.null(c)){
    c(a=a, b=b, c=sqrt(a^2+b^2))
  }
}
```

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Solution to Exercise 3

Modify the `pythagoras()` function so that it accepts a list with two elements named 'a', 'b', or 'c' and computes the missing one, according to the Pythagorean theorem.

```
pythagoras.list <- function(edges){
  edge_names <- c("a", "b", "c")
  edges_provided <- edge_names %in% names(edges)
  if(sum(edges_provided)!=2) stop("Wrong argument")
  if(! "a" %in% names(edges)){
    edges$a <- sqrt(edges$c^2-edges$b^2)
  }else if(! "b" %in% names(edges)){
    edges$b <- sqrt(edges$c^2-edges$a^2)
  }else if(! "c" %in% names(edges)){
    edges$c <- sqrt(edges$a^2+edges$b^2)
  }
  edges
}
```

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