

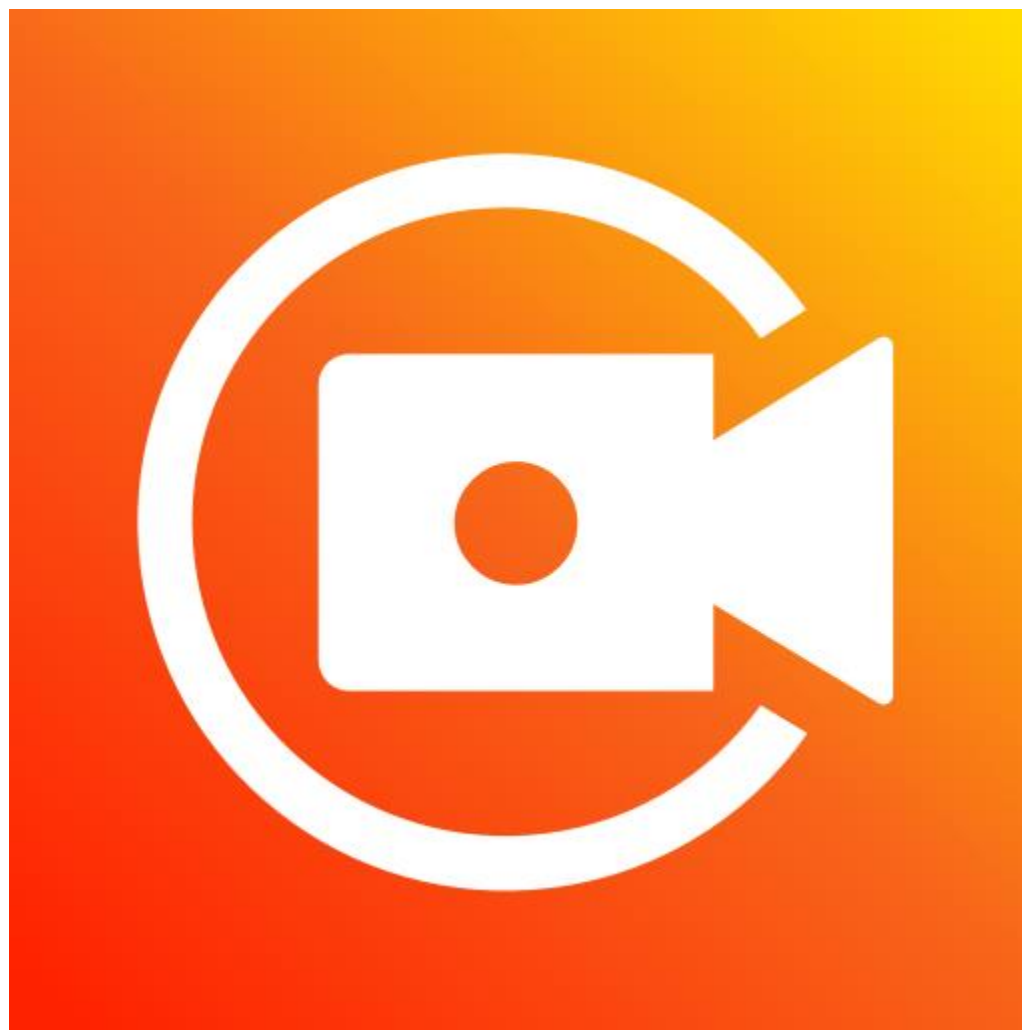
# **MAP 2112 – Introdução à Lógica de Programação e Modelagem Computacional**

**1º Semestre - 2020**

**Prof. Dr. Luis Carlos de Castro Santos**

**lsantos@ime.usp.br**

**NÃO ESQUEÇA DE INICIAR A GRAVAÇÃO**



# **MAP 2112 – Introdução à Lógica de Programação e Modelagem Computacional**

**1º Semestre - 2020**

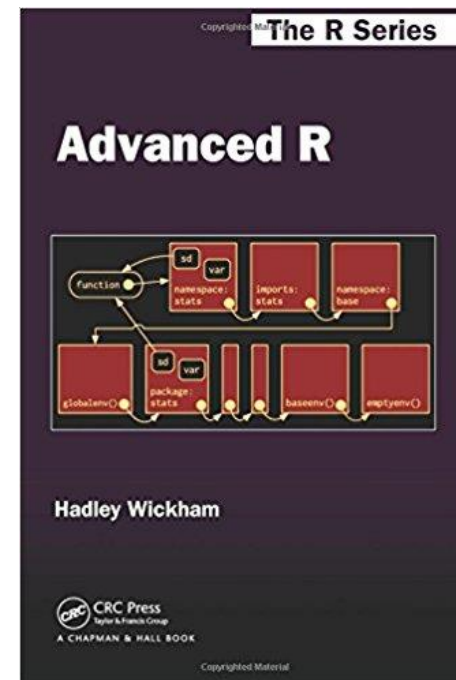
**Prof. Dr. Luis Carlos de Castro Santos**

**lsantos@ime.usp.br**

# ROTEIRO (continuação)

Material Prof. Roger Peng

<http://www.biostat.jhsph.edu/~rpeng/>



Esse material é fortemente baseado no livro

**Advanced R (Chapman & Hall/CRC The R Series)**

de Hadley Wickham (<http://hadley.nz/>) o Cientista-chefe do RStudio



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*Data structures*

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	Homogeneous	Heterogeneous
1d	Atomic vector	List
2d	Matrix	Data frame
nd	Array	

---

# Subsetting

---

# *Subsetting*

---

R's subsetting operators are powerful and fast. Mastery of subsetting allows you to succinctly express complex operations in a way that few other languages can match. Subsetting is hard to learn because you need to master a number of interrelated concepts:

- The three subsetting operators.
- The six types of subsetting.
- Important differences in behaviour for different objects (e.g., vectors, lists, factors, matrices, and data frames).
- The use of subsetting in conjunction with assignment.

# Subsetting

There are a number of operators that can be used to extract subsets of R objects.

- `[]` always returns an object of the same class as the original; can be used to select more than one element (there is one exception)
- `[[` is used to extract elements of a list or a data frame; it can only be used to extract a single element and the class of the returned object will not necessarily be a list or data frame
- `$` is used to extract elements of a list or data frame by name; semantics are similar to that of `[[`.



# Subsetting

```
> x <- c("a", "b", "c", "c", "d", "a")
> x[1]
[1] "a"
> x[2]
[1] "b"
> x[1:4]
[1] "a" "b" "c" "c"
> x[x > "a"]
[1] "b" "c" "c" "d"
> u <- x > "a"
> u
[1] FALSE TRUE TRUE TRUE TRUE FALSE
> x[u]
[1] "b" "c" "c" "d"
```

## Atomic vectors

```
x <- c(2.1, 4.2, 3.3, 5.4)
```

- Positive integers return elements at the specified positions:

```
x[c(3, 1)]
#> [1] 3.3 2.1
x[order(x)]
#> [1] 2.1 3.3 4.2 5.4
```

```
> x <- c(2.1,4.2,3.3,5.4)
> order(x)
[1] 1 3 2 4
> |
```

```
# Duplicated indices yield duplicated values
```

```
x[c(1, 1)]
#> [1] 2.1 2.1
```

```
# Real numbers are silently truncated to integers
```

```
x[c(2.1, 2.9)]
#> [1] 4.2 4.2
```

```
x <- c(2.1, 4.2, 3.3, 5.4)
```

- **Negative integers** omit elements at the specified positions:

```
x[-c(3, 1)]  
#> [1] 4.2 5.4
```

You can't mix positive and negative integers in a single subset:

```
x[c(-1, 2)]  
#> Error: only 0's may be mixed with negative subscripts
```

```
x <- c(2.1, 4.2, 3.3, 5.4)
```

- **Logical vectors** select elements where the corresponding logical value is TRUE. This is probably the most useful type of subsetting because you write the expression that creates the logical vector:

```
x[c(TRUE, TRUE, FALSE, FALSE)]  
#> [1] 2.1 4.2  
x[x > 3]  
#> [1] 4.2 3.3 5.4
```

If the logical vector is shorter than the vector being subsetted, it will be *recycled* to be the same length.

```
x[c(TRUE, FALSE)]  
#> [1] 2.1 3.3  
# Equivalent to  
x[c(TRUE, FALSE, TRUE, FALSE)]  
#> [1] 2.1 3.3
```

```
x <- c(2.1, 4.2, 3.3, 5.4)
```

A missing value in the index always yields a missing value in the output:

```
x[c(TRUE, TRUE, NA, FALSE)]  
#> [1] 2.1 4.2 NA
```

- **Nothing** returns the original vector. This is not useful for vectors but is very useful for matrices, data frames, and arrays. It can also be useful in conjunction with assignment.

```
x[ ]  
#> [1] 2.1 4.2 3.3 5.4
```

- **Zero** returns a zero-length vector. This is not something you usually do on purpose, but it can be helpful for generating test data.

```
x[0]  
#> numeric(0)
```

```
x <- c(2.1, 4.2, 3.3, 5.4)
```

If the vector is named, you can also use:

- **Character vectors** to return elements with matching names.

```
(y <- setNames(x, letters[1:4]))
```

```
#>  a  b  c  d
```

```
#> 2.1 4.2 3.3 5.4
```

```
y[c("d", "c", "a")]
```

```
#>  d  c  a
```

```
#> 5.4 3.3 2.1
```

```
# Like integer indices, you can repeat indices
```

```
y[c("a", "a", "a")]
```

```
#>  a  a  a
```

```
#> 2.1 2.1 2.1
```

```
# When subsetting with [ names are always matched exactly
```

```
z <- c(abc = 1, def = 2)
```

```
z[c("a", "d")]
```

```
#> <NA> <NA>
```

```
#>  NA  NA
```

### 3.1.2 Lists

Subsetting a list works in the same way as subsetting an atomic vector. Using `[]` will always return a list; `[[` and `$`, as described below, let you pull out the components of the list.

# Subsetting Lists

```
> x <- list(foo = 1:4, bar = 0.6)
> x[1]
$foo
[1] 1 2 3 4

> x[[1]]
[1] 1 2 3 4

> x$bar
[1] 0.6
> x[["bar"]]
[1] 0.6
> x["bar"]
$bar
[1] 0.6
```

```
Console Terminal x
~/ ↩
> x <- list(foo=1:4, bar=0.6)
> a <- x[1]
> a
$foo
[1] 1 2 3 4

> class(a)
[1] "list"
> b<- x[[1]]
> b
[1] 1 2 3 4
> class(b)
[1] "integer"
> |
```



# Subsetting Lists

```
> x <- list(foo = 1:4, bar = 0.6, baz = "hello")
> x[c(1, 3)]
$foo
[1] 1 2 3 4

$baz
[1] "hello"
```

# Subsetting Lists

The `[[` operator can be used with *computed* indices; `$` can only be used with literal names.

```
> x <- list(foo = 1:4, bar = 0.6, baz = "hello")
> name <- "foo"
> x[[name]] ## computed index for 'foo'
[1] 1 2 3 4
> x$name    ## element 'name' doesn't exist!
NULL
> x$foo
[1] 1 2 3 4 ## element 'foo' does exist
```

# Subsetting Nested Elements of a List

The `[[` can take an integer sequence.

```
> x <- list(a = list(10, 12, 14), b = c(3.14, 2.81))
> x[[c(1, 3)]]
[1] 14
> x[[1]][[3]]
[1] 14

> x[[c(2, 1)]]
[1] 3.14
```

```
Console Terminal x
~/ ↵
> x <- list(a = list(10, 12, 14), b = c(3.14, 2.81))
> aa <- x[1]
> aa
$a
$a[[1]]
[1] 10

$a[[2]]
[1] 12

$a[[3]]
[1] 14

> class(aa)
[1] "list"
> bb <- x[2]
> bb
$b
[1] 3.14 2.81

> class(bb)
[1] "list"
> cc <- x[[2]]
> cc
[1] 3.14 2.81
> class(cc)
[1] "numeric"
> |
```

```
Console Terminal x
~/ ↵
> x <- list(a=list(10,12,14),b=c(3.14,2.81))
> x[[c(1,3)]]
[1] 14
> x[c(1,3)]
$a
$a[[1]]
[1] 10

$a[[2]]
[1] 12

$a[[3]]
[1] 14

$<NA>
NULL

> |
```

O que aconteceu ?

### 3.1.3 Matrices and arrays

You can subset higher-dimensional structures in three ways:

- With multiple vectors.
- With a single vector.
- With a matrix.

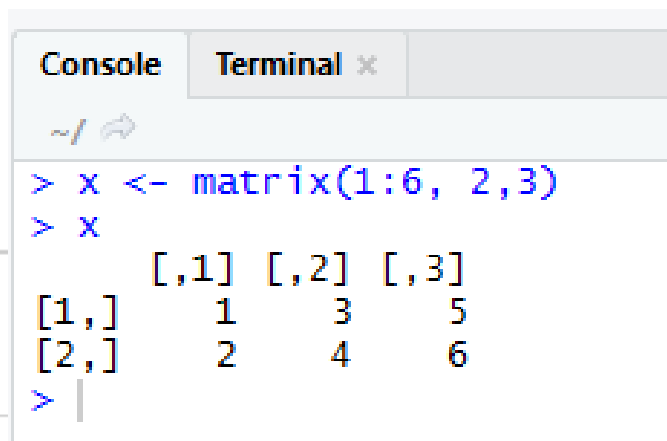
# Subsetting a Matrix

Matrices can be subsetting in the usual way with  $(i,j)$  type indices.

```
> x <- matrix(1:6, 2, 3)
> x[1, 2]
[1] 3
> x[2, 1]
[1] 2
```

Indices can also be missing.

```
> x[1, ]
[1] 1 3 5
> x[, 2]
[1] 3 4
```



```
Console Terminal x
~/
> x <- matrix(1:6, 2, 3)
> x
      [,1] [,2] [,3]
[1,]    1    3    5
[2,]    2    4    6
> |
```

## Subsetting a Matrix

By default, when a single element of a matrix is retrieved, it is returned as a vector of length 1 rather than a  $1 \times 1$  matrix. This behavior can be turned off by setting `drop = FALSE`.

```
> x <- matrix(1:6, 2, 3)
> x[1, 2]
[1] 3
> x[1, 2, drop = FALSE]
     [,1]
[1,] 3
```



## Subsetting a Matrix

Similarly, subsetting a single column or a single row will give you a vector, not a matrix (by default).

```
> x <- matrix(1:6, 2, 3)
> x[1, ]
[1] 1 3 5
> x[1, , drop = FALSE]
      [,1] [,2] [,3]
[1,]    1    3    5
```

### 3.1.4 Data frames

Data frames possess the characteristics of both lists and matrices: if you subset with a single vector, they behave like lists; if you subset with two vectors, they behave like matrices.

```
Console Terminal x
~/ ↩
> df <- data.frame(x = 1:3, y=3:1, z = letters[1:3])
> df
  x y z
1 1 3 a
2 2 2 b
3 3 1 c
> df[df$x == 2,]
  x y z
2 2 2 b
> df[c(1,3)]
  x z
1 1 a
2 2 b
3 3 c
> |

> a <- df[df$x == 2,]
> class(a)
[1] "data.frame"
> b <- df[c(1,3)]
> class(b)
[1] "data.frame"
> |
```

```
# There are two ways to select columns from a data frame
```

```
# Like a list:
```

```
df[c("x", "z")]
```

```
#>   x z
```

```
#> 1 1 a
```

```
#> 2 2 b
```

```
#> 3 3 c
```

```
# Like a matrix
```

```
df[, c("x", "z")]
```

```
#>   x z
```

```
#> 1 1 a
```

```
#> 2 2 b
```

```
#> 3 3 c
```

```
# There's an important difference if you select a single  
# column: matrix subsetting simplifies by default, list  
# subsetting does not.
```

```
str(df["x"])
```

```
#> 'data.frame':   3 obs. of  1 variable:
```

```
#> $ x: int  1 2 3
```

```
str(df[, "x"])
```

```
#> int [1:3] 1 2 3
```

# Partial Matching

Partial matching of names is allowed with `[` and `$`.

```
> x <- list(aardvark = 1:5)
> x$a
[1] 1 2 3 4 5
> x[["a"]]
NULL
> x[["a", exact = FALSE]]
[1] 1 2 3 4 5
```

# Removing NA Values

A common task is to remove missing values (**NA**s).

```
> x <- c(1, 2, NA, 4, NA, 5)
> bad <- is.na(x)
> x[!bad]
[1] 1 2 4 5
```

```
x <- c(1, 2, NA, 4, NA, 5)
> is.na(x)
[1] FALSE FALSE TRUE FALSE TRUE FALSE
> bad <- is.na(x)
> bad
[1] FALSE FALSE TRUE FALSE TRUE FALSE
> !bad
[1] TRUE TRUE FALSE TRUE FALSE TRUE
> x[!bad]
[1] 1 2 4 5
```

## Removing NA Values

What if there are multiple things and you want to take the subset with no missing values?

```
> x <- c(1, 2, NA, 4, NA, 5)
> y <- c("a", "b", NA, "d", NA, "f")
> good <- complete.cases(x, y)
> good
[1] TRUE TRUE FALSE TRUE FALSE TRUE
> x[good]
[1] 1 2 4 5
> y[good]
[1] "a" "b" "d" "f"
```

# Removing NA Values

```
> airquality[1:6, ]
  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4  67    5    1
2    36    118  8.0  72    5    2
3    12    149 12.6  74    5    3
4    18    313 11.5  62    5    4
5    NA     NA 14.3  56    5    5
6    28     NA 14.9  66    5    6

> good <- complete.cases(airquality)
> airquality[good, ][1:6, ]
  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4  67    5    1
2    36    118  8.0  72    5    2
3    12    149 12.6  74    5    3
4    18    313 11.5  62    5    4
7    23    299  8.6  65    5    7
```



Files Plots Packages Help Viewer

R: The R Datasets Package ▾ Find in Topic

# The R Datasets Package

---

Documentation for package 'datasets' version 3.5.2

- [DESCRIPTION file.](#)

## Help Pages

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[datasets-package](#) The R Datasets Package

**-- A --**

<a href="#">ability.cov</a>	Ability and Intelligence Tests
<a href="#">airmiles</a>	Passenger Miles on Commercial US Airlines, 1937-1960
<a href="#">AirPassengers</a>	Monthly Airline Passenger Numbers 1949-1960
<a href="#">airquality</a>	New York Air Quality Measurements
<a href="#">anscombe</a>	Anscombe's Quartet of 'Identical' Simple Linear Regressions
<a href="#">attenu</a>	The Joyner-Boore Attenuation Data
<a href="#">attitude</a>	The Chatterjee-Price Attitude Data
<a href="#">austres</a>	Quarterly Time Series of the Number of Australian Residents

**-- B --**

<a href="#">beaver1</a>	Body Temperature Series of Two Beavers
<a href="#">beaver2</a>	Body Temperature Series of Two Beavers
<a href="#">beavers</a>	Body Temperature Series of Two Beavers



Files Plots Packages Help Viewer

R: New York Air Quality Measurements Find in Topic

## Usage

`airquality`

## Format

A data frame with 154 observations on 6 variables.

[,1] Ozone    numeric Ozone (ppb)  
 [,2] Solar.R numeric Solar R (lang)  
 [,3] Wind     numeric Wind (mph)  
 [,4] Temp    numeric Temperature (degrees F)  
 [,5] Month   numeric Month (1--12)  
 [,6] Day     numeric Day of month (1--31)

## Details

Daily readings of the following air quality values for May 1, 1973 (a Tuesday) to September 30, 1973.

- **Ozone:** Mean ozone in parts per billion from 1300 to 1500 hours at Roosevelt Island
- **Solar.R:** Solar radiation in Langleys in the frequency band 4000–7700 Angstroms from 0800 to 1200 hours at Central Park
- **Wind:** Average wind speed in miles per hour at 0700 and 1000 hours at LaGuardia Airport
- **Temp:** Maximum daily temperature in degrees Fahrenheit at La Guardia Airport.

## Source

The data were obtained from the New York State Department of Conservation (ozone data) and the National Weather Service (meteorological data).

## References

Chambers, J. M., Cleveland, W. S., Kleiner, B. and Tukey, P. A. (1983) *Graphical Methods for Data Analysis*. Belmont, CA: Wadsworth.

## Examples

```
require(graphics)
pairs(airquality, panel = panel.smooth, main = "airquality data")
```

```
Console Terminal x
~/
> head(airquality)
  Ozone Solar.R Wind Temp Month Day
1    41    190  7.4  67     5    1
2    36    118  8.0  72     5    2
3    12    149 12.6  74     5    3
4    18    313 11.5  62     5    4
5    NA     NA 14.3  56     5    5
6    28     NA 14.9  66     5    6
> tail(airquality)
  Ozone Solar.R Wind Temp Month Day
148   14     20 16.6  63     9   25
149   30    193  6.9  70     9   26
150   NA    145 13.2  77     9   27
151   14    191 14.3  75     9   28
152   18    131  8.0  76     9   29
153   20    223 11.5  68     9   30
> |
```

Próxima Semana

# EXERCÍCIOS NA MONITORIA

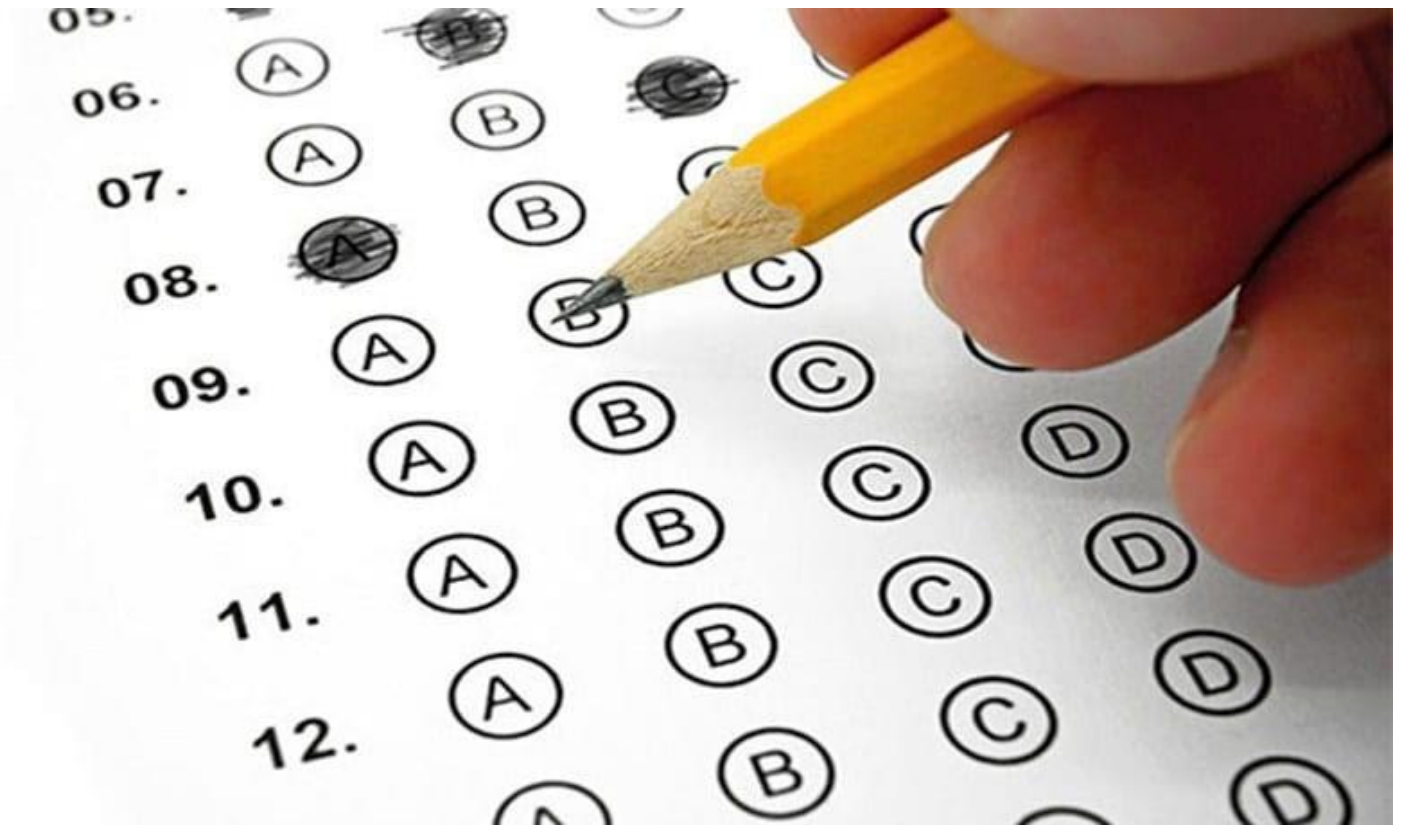
## Horário alternativo

## TBD



No dia 28/4/2020

# TESTE através do Moodle



Cheat Sheets – “cola oficial”

<https://www.rstudio.com/resources/cheatsheets/>

Base R

<http://github.com/rstudio/cheatsheets/raw/master/base-r.pdf>

# Base R Cheat Sheet

## Getting Help

### Accessing the help files

#### ?mean

Get help of a particular function.

**help.search('weighted mean')**

Search the help files for a word or phrase.

**help(package = 'dplyr')**

Find help for a package.

### More about an object

#### str(iris)

Get a summary of an object's structure.

#### class(iris)

Find the class an object belongs to.

## Using Packages

### install.packages('dplyr')

Download and install a package from CRAN.

### library(dplyr)

Load the package into the session, making all its functions available to use.

### dplyr::select

Use a particular function from a package.

### data(iris)

Load a built-in dataset into the environment.

## Working Directory

### getwd()

Find the current working directory (where inputs are found and outputs are sent).

### setwd('C://file/path')

Change the current working directory.

**Use projects in RStudio to set the working directory to the folder you are working in.**

## Vectors

### Creating Vectors

c(2, 4, 6)	2 4 6	Join elements into a vector
2:6	2 3 4 5 6	An integer sequence
seq(2, 3, by=0.5)	2.0 2.5 3.0	A complex sequence
rep(1:2, times=3)	1 2 1 2 1 2	Repeat a vector
rep(1:2, each=3)	1 1 1 2 2 2	Repeat elements of a vector

### Vector Functions

<b>sort(x)</b> Return x sorted.	<b>rev(x)</b> Return x reversed.
<b>table(x)</b> See counts of values.	<b>unique(x)</b> See unique values.

### Selecting Vector Elements

#### By Position

<b>x[4]</b>	The fourth element.
<b>x[-4]</b>	All but the fourth.
<b>x[2:4]</b>	Elements two to four.
<b>x[-(2:4)]</b>	All elements except two to four.
<b>x[c(1, 5)]</b>	Elements one and five.

#### By Value

<b>x[x == 10]</b>	Elements which are equal to 10.
<b>x[x &lt; 0]</b>	All elements less than zero.
<b>x[x %in% c(1, 2, 5)]</b>	Elements in the set 1, 2, 5.

#### Named Vectors

<b>x['apple']</b>	Element with name 'apple'.
-------------------	----------------------------

## Programming

### For Loop

```
for (variable in sequence){  
  Do something  
}
```

#### Example

```
for (i in 1:4){  
  j <- 1 + 10  
  print(j)  
}
```

### If Statements

```
if (condition){  
  Do something  
} else {  
  Do something different  
}
```

#### Example

```
if (1 > 3){  
  print('Yes')  
} else {  
  print('No')  
}
```

### While Loop

```
while (condition){  
  Do something  
}
```

#### Example

```
while (1 < 5){  
  print(1)  
  1 <- 1 + 1  
}
```

### Functions

```
function_name <- function(var){  
  Do something  
  return(new_variable)  
}
```

#### Example

```
square <- function(x){  
  squared <- x*x  
  return(squared)  
}
```

## Reading and Writing Data

Also see the **readr** package.

Input	Output	Description
<code>df &lt;- read.table('file.txt')</code>	<code>write.table(df, 'file.txt')</code>	Read and write a delimited text file.
<code>df &lt;- read.csv('file.csv')</code>	<code>write.csv(df, 'file.csv')</code>	Read and write a comma separated value file. This is a special case of read.table/write.table.
<code>load('file.Rdata')</code>	<code>save(df, file = 'file.Rdata')</code>	Read and write an R data file, a file type special for R.

#### Conditions

<b>a == b</b>	Are equal	<b>a &gt; b</b>	Greater than	<b>a &gt;= b</b>	Greater than or equal to	<b>is.na(a)</b>	Is missing
<b>a != b</b>	Not equal	<b>a &lt; b</b>	Less than	<b>a &lt;= b</b>	Less than or equal to	<b>is.null(a)</b>	Is null

## Types

Converting between common data types in R. Can always go from a higher value in the table to a lower value.

as.logical	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).
as.numeric	1, 0, 1	Integers or floating point numbers.
as.character	'1', '0', '1'	Character strings. Generally preferred to factors.
as.factor	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.

## Maths Functions

log(x)	Natural log.	sum(x)	Sum.
exp(x)	Exponential.	mean(x)	Mean.
max(x)	Largest element.	median(x)	Median.
min(x)	Smallest element.	quantile(x)	Percentage quantiles.
round(x, n)	Round to n decimal places.	rank(x)	Rank of elements.
signif(x, n)	Round to n significant figures.	var(x)	The variance.
cor(x, y)	Correlation.	sd(x)	The standard deviation.

## Variable Assignment

```
> a <- 'apple'
> a
[1] 'apple'
```

## The Environment

ls()	List all variables in the environment.
rm(x)	Remove x from the environment.
rm(list = ls())	Remove all variables from the environment.

You can use the environment panel in RStudio to browse variables in your environment.

## Matrices

```
m <- matrix(x, nrow = 3, ncol = 3)
Create a matrix from x.
```

	m[2, ] - Select a row	t(m)	Transpose
	m[, 1] - Select a column	m %*% n	Matrix Multiplication
	m[2, 3] - Select an element	solve(m, n)	Find x in: m * x = n

## Lists

```
l <- list(x = 1:5, y = c('a', 'b'))
A list is a collection of elements which can be of different types.
```

l[[2]]	l[1]	l\$x	l['y']
Second element of l.	New list with only the first element.	Element named x.	New list with only element named y.

Also see the **dplyr** package.

## Data Frames

```
df <- data.frame(x = 1:3, y = c('a', 'b', 'c'))
A special case of a list where all elements are the same length.
```

x	y
1	a
2	b
3	c

### Matrix subsetting

df[, 2]	
df[2, ]	
df[2, 2]	

### List subsetting

dfs[x]	
df[[2]]	

### Understanding a data frame

View(df)	See the full data frame.
head(df)	See the first 6 rows.

nrow(df)  
Number of rows.

ncol(df)  
Number of columns.

dim(df)  
Number of columns and rows.

cbind - Bind columns.



rbind - Bind rows.



## Strings

Also see the **stringr** package.

paste(x, y, sep = ' ')	Join multiple vectors together.
paste(x, collapse = ' ')	Join elements of a vector together.
grep(pattern, x)	Find regular expression matches in x.
gsub(pattern, replace, x)	Replace matches in x with a string.
toupper(x)	Convert to uppercase.
tolower(x)	Convert to lowercase.
nchar(x)	Number of characters in a string.

## Factors

factor(x)	Turn a vector into a factor. Can set the levels of the factor and the order.
cut(x, breaks = 4)	Turn a numeric vector into a factor by 'cutting' into sections.

## Statistics

lm(y ~ x, data=df)	Linear model.	t.test(x, y)	Perform a t-test for difference between means.	prop.test	Test for a difference between proportions.
glm(y ~ x, data=df)	Generalised linear model.	pairwise.t.test	Perform a t-test for paired data.	sdv	Analysis of variance.
summary	Get more detailed information out a model.				

## Distributions

	Random Variables	Density Function	Cumulative Distribution	Quantile
Normal	rnorm	dnorm	pnorm	qnorm
Poisson	rpois	dpois	ppois	qpois
Binomial	rbinom	dbinom	pbinom	qbinom
Uniform	runif	dunif	punif	qunif

## Plotting

Also see the **ggplot2** package.

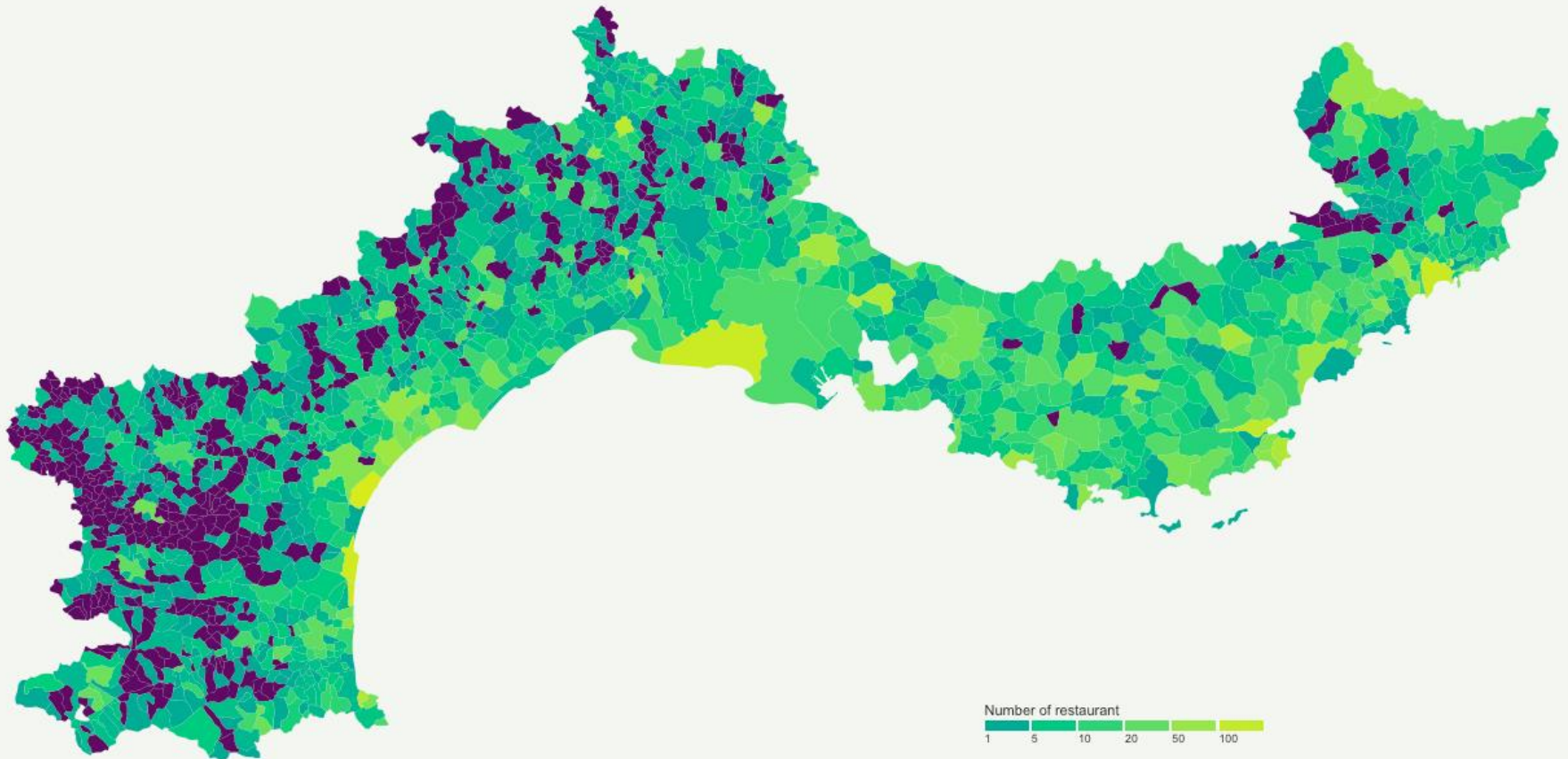
	plot(x)	Values of x in order.		plot(x, y)	Values of x against y.		hist(x)	Histogram of x.
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## Dates

See the **lubridate** package.



South of France Restaurant concentration  
Number of restaurant per city district



Data: INSEE | Creation: Yan Holtz | r-graph-gallery.com

<https://www.r-graph-gallery.com/327-chloropleth-map-from-geojson-with-ggplot2/>