

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

**1º Semestre - 2020**

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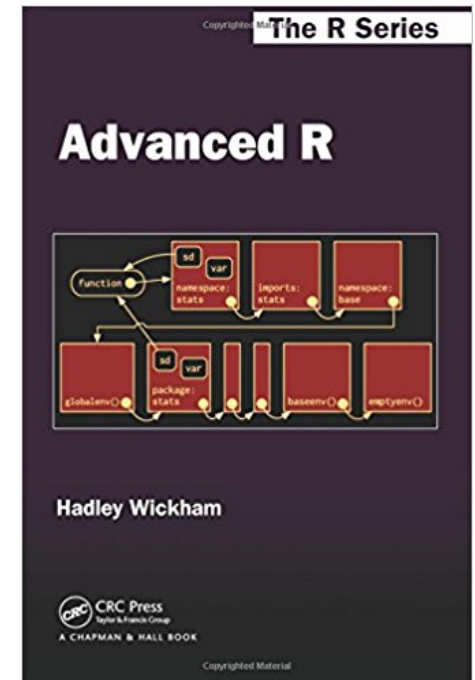
Esse material é fortemente baseado no livro

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

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

Seguindo o roteiro do Prof. Roger Peng  
<http://www.biostat.jhsph.edu/~rpeng/>

Quando chegarmos nos tópicos de modelagem e Data Science novas referências serão selecionadas.





# Overview and History of R

Roger D. Peng, Associate Professor of Biostatistics  
Johns Hopkins Bloomberg School of Public Health

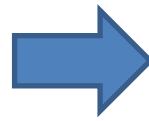
# What is R?

What is R?

R is a dialect of the S language.

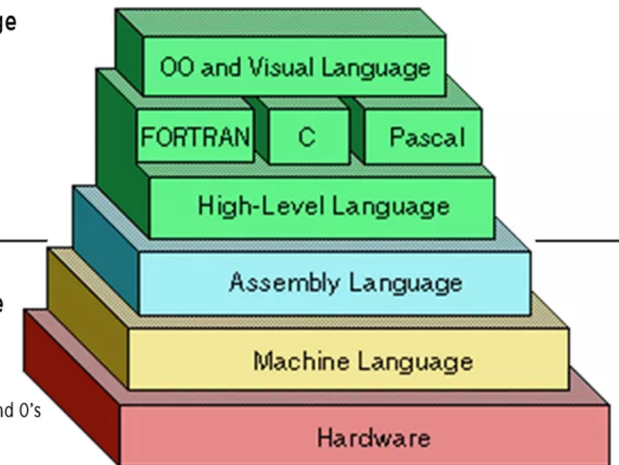
# What is S?

- S is a language that was developed by John Chambers and others at Bell Labs.
- S was initiated in 1976 as an internal statistical analysis environment—originally implemented as Fortran libraries.
- Early versions of the language did not contain functions for statistical modeling.
- In 1988 the system was rewritten in C and began to resemble the system that we have today (this was Version 3 of the language). The book *Statistical Models in S* by Chambers and Hastie (the white book) documents the statistical analysis functionality.
- Version 4 of the S language was released in 1998 and is the version we use today. The book *Programming with Data* by John Chambers (the green book) documents this version of the language.



## High Level Language

- Easy for Programmers to understand
- Contains English Words



justcode.me

# S Philosophy

In “Stages in the Evolution of S”, John Chambers writes:

“[W]e wanted users to be able to begin in an interactive environment, where they did not consciously think of themselves as programming. Then as their needs became clearer and their sophistication increased, they should be able to slide gradually into programming, when the language and system aspects would become more important.”

<http://www.stat.bell-labs.com/S/history.html>

# What is R?

- 1991: Created in New Zealand by Ross Ihaka and Robert Gentleman. Their experience developing R is documented in a 1996 *JCGS* paper.
- 1993: First announcement of R to the public.
- 1995: Martin Machler convinces Ross and Robert to use the GNU General Public License to make R free software.
- 1996: A public mailing list is created (R-help and R-devel)
- 1997: The R Core Group is formed (containing some people associated with S-PLUS). The core group controls the source code for R.
- 2000: R version 1.0.0 is released.
- 2013: R version 3.0.2 is released on December 2013.



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## The Comprehensive R Archive Network

### Download and Install R

Precompiled binary distributions of the base system and contributed packages, **Windows and Mac** users most likely want one of these versions of R:

- [Download R for Linux](#)
- [Download R for \(Mac\) OS X](#)
- [Download R for Windows](#)

R is part of many Linux distributions, you should check with your Linux package management system in addition to the link above.

### Source Code for all Platforms

Windows and Mac users most likely want to download the precompiled binaries listed in the upper box, not the source code. The sources have to be compiled before you can use them. If you do not know what this means, you probably do not want to do it!

- The latest release (2020-02-29, Holding the Windsock) [R-3.6.3.tar.gz](#), read [what's new](#) in the latest version.
- Sources of [R alpha and beta releases](#) (daily snapshots, created only in time periods before a planned release).
- Daily snapshots of current patched and development versions are [available here](#). Please read about [new features and bug fixes](#) before filing corresponding feature requests or bug reports.
- Source code of older versions of R is [available here](#).
- Contributed extension [packages](#)

### Questions About R

- If you have questions about R like how to download and install the software, or what the license terms are, please read our [answers to frequently asked questions](#) before you send an email.

What are R and CRAN?

<https://www.r-project.org/>



# The R Project for Statistical Computing

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## R Foundation

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## Help With R

## Getting Started

R is a free software environment for statistical computing and graphics. It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS. To [download R](#), please choose your preferred [CRAN mirror](#).

If you have questions about R like how to download and install the software, or what the license terms are, please read our [answers to frequently asked questions](#) before you send an email.

## News

- [R version 3.6.3 \(Holding the Windsock\)](#) has been released on 2020-02-29.
- useR! 2020 will take place in St. Louis, Missouri, USA.
- [R version 3.5.3 \(Great Truth\)](#) has been released on 2019-03-11.
- The R Foundation Conference Committee has released a [call for proposals](#) to host useR! 2020 in North America.
- You can now support the R Foundation with a renewable subscription as a [supporting member](#)
- The R Foundation has been awarded the Personality/Organization of the year 2018 award by the professional association of German market and social researchers.



# Features of R

- Syntax is very similar to S, making it easy for S-PLUS users to switch over.
- Semantics are superficially similar to S, but in reality are quite different (more on that later).
- Runs on almost any standard computing platform/OS (even on the PlayStation 3)
- Frequent releases (annual + bugfix releases); active development.
- Quite lean, as far as software goes; functionality is divided into modular packages
- Graphics capabilities very sophisticated and better than most stat packages.
- Useful for interactive work, but contains a powerful programming language for developing new tools (user -> programmer)
- Very active and vibrant user community; R-help and R-devel mailing lists and Stack Overflow

# Drawbacks of R

- Essentially based on 40 year old technology.
- Little built in support for dynamic or 3-D graphics (but things have improved greatly since the “old days”).
- Functionality is based on consumer demand and user contributions. If no one feels like implementing your favorite method, then it's *your* job!
  - (Or you need to pay someone to do it)
- Objects must generally be stored in physical memory; but there have been advancements to deal with this too
- Not ideal for all possible situations (but this is a drawback of all software packages).



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# RStudio

## Take control of your R code

RStudio is an integrated development environment (IDE) for R. It includes a console, syntax-highlighting editor that supports direct code execution, as well as tools for plotting, history, debugging and workspace management. [Click here to see more RStudio features.](#)

RStudio is available in **open source** and **commercial** editions and runs on the desktop (Windows, Mac, and Linux) or in a browser connected to RStudio Server or RStudio Server Pro (Debian/Ubuntu, Red Hat/CentOS, and SUSE Linux).

There are two versions of RStudio:



### RStudio Desktop

Run RStudio on your desktop



### RStudio Server

Centralize access and computation



### Open Source Edition

#### Overview

- Access RStudio locally
- Syntax highlighting, code completion, and smart indentation
- Execute R code directly from the source editor
- Quickly jump to function definitions
- Easily manage multiple working directories using projects
- Integrated R help and documentation
- Interactive debugger to diagnose and fix errors quickly
- Extensive package development tools

#### Support

Community forums only

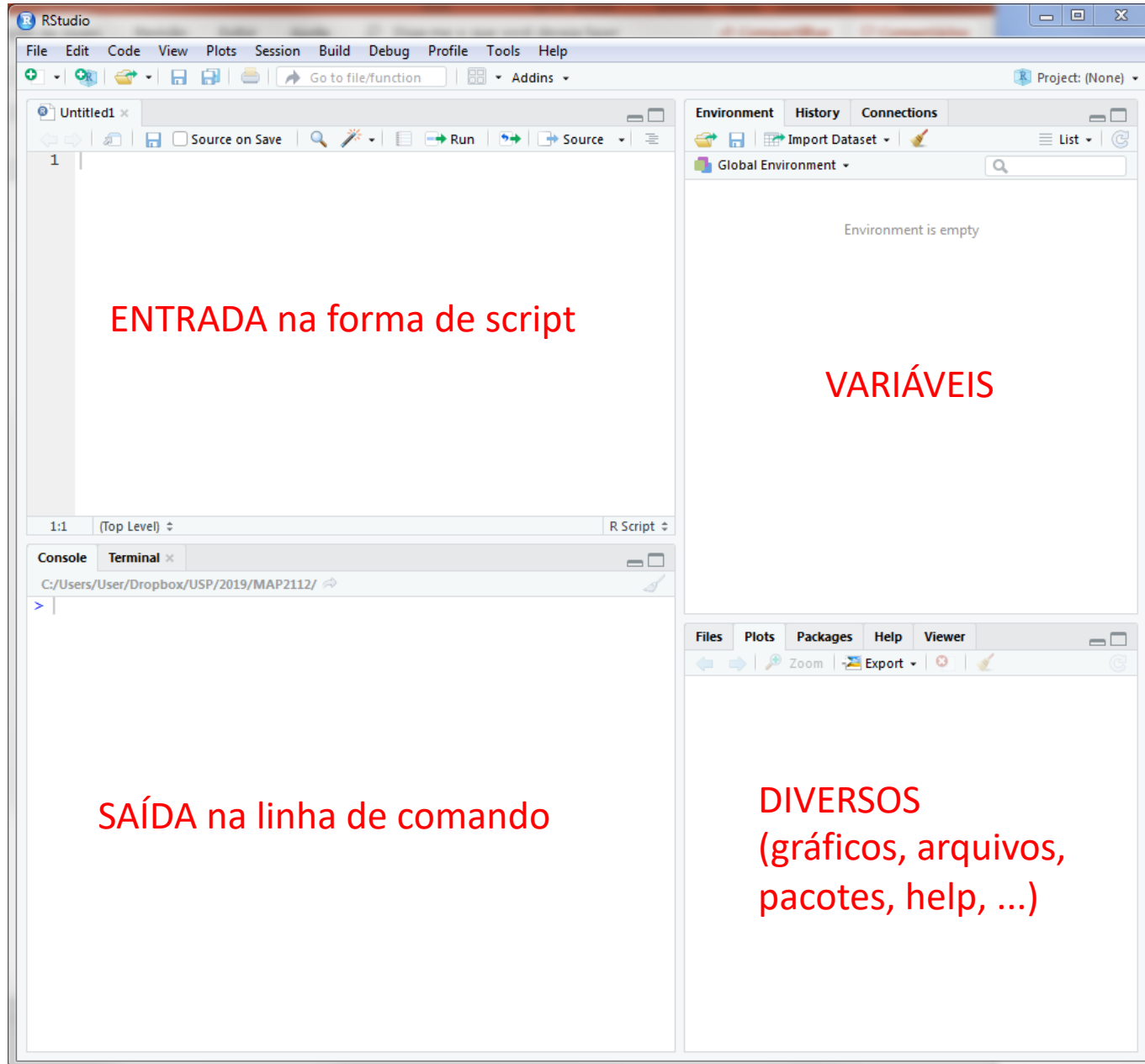
#### License

AGPL v3

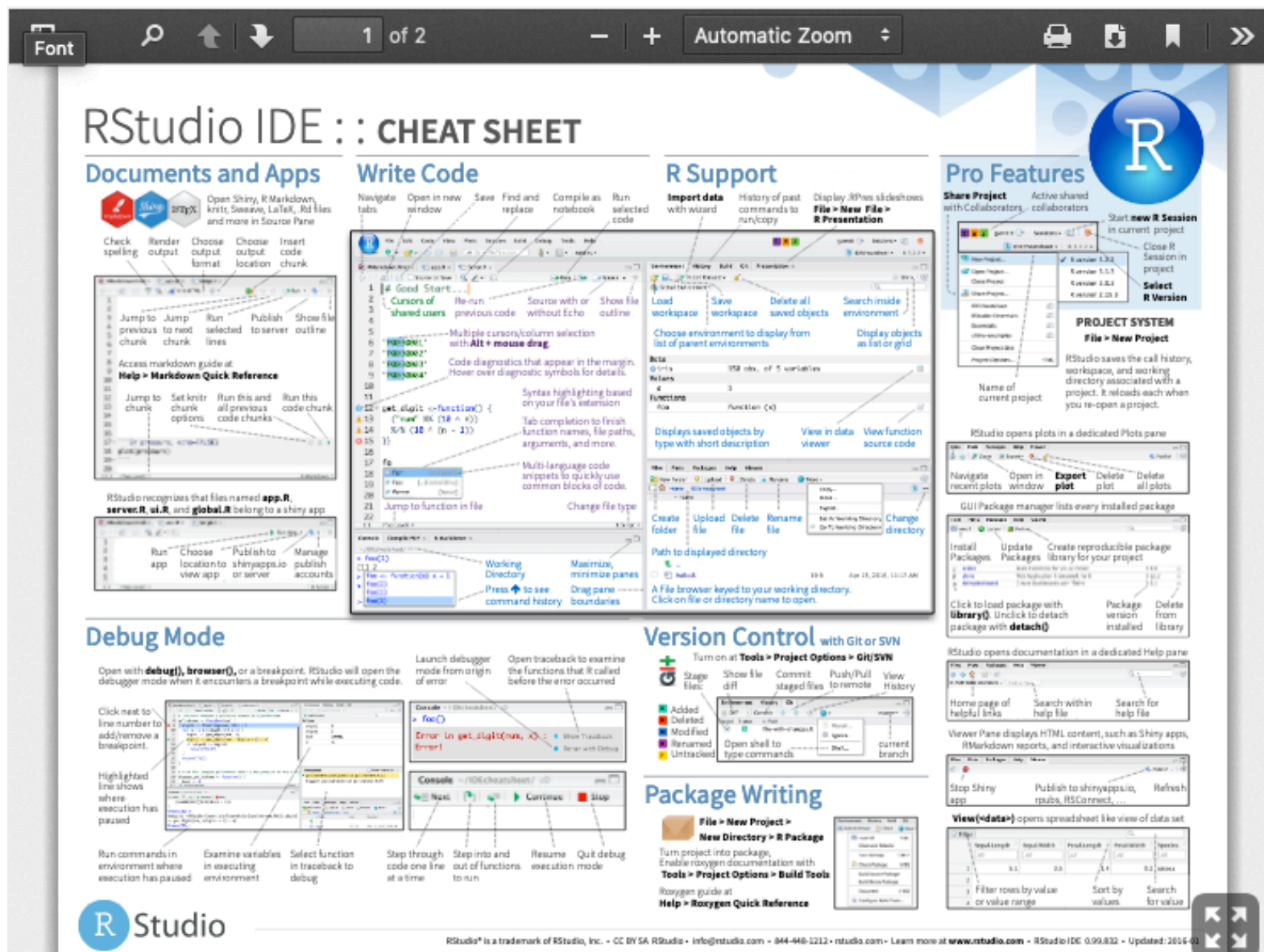
#### Pricing

Free

[DOWNLOAD RSTUDIO DESKTOP](#)



<https://resources.rstudio.com/rstudio-developed/rstudio-ide>



**RStudio IDE : : CHEAT SHEET**

**Documents and Apps**

- Open Shiny, R Markdown, Knit, Sweave, LaTeX, Rd files and more in Source Pane
- Check spelling
- Render output
- Choose output format
- Choose output location
- Insert code chunk
- Jump to previous chunk
- Jump to next chunk
- Run selected lines
- Publish to server
- Show file outline
- Access markdown guide at **Help > Markdown Quick Reference**
- Jump to chunk
- Set knitr chunk options
- Run this and all previous code chunks
- Run this code chunk

RStudio recognizes that files named **app.R**, **server.R**, **ui.R**, and **global.R** belong to a shiny app

- Run app
- Choose location to view app
- Publish to shinyapps.io or server
- Manage publish accounts

**Write Code**

- Navigate tabs
- Open in new window
- Save
- Find and replace
- Compile as notebook
- Run selected code
- Source with or without echo
- Show file outline
- Multiple cursors/column selection with **Alt + mouse drag**
- Code diagnostics that appear in the margin. Hover over diagnostic symbols for details.
- Syntax highlighting based on your file's extension
- Tab completion to finish function names, file paths, arguments, and more.
- Multi-language code snippets to quickly use common blocks of code.
- Jump to function in file
- Change file type
- Working Directory
- Press **Alt** to see command history
- Maximize, minimize panes
- Drag pane boundaries

**R Support**

- Import data with wizard
- History of past commands to run/copy
- Display R/Python slideshows
- File > New File > R Presentation**
- Load workspace
- Save workspace
- Delete all saved objects
- Search inside environment
- Display objects as list or grid
- Choose environment to display from list of parent environments
- Displays saved objects by type with short description
- View in data viewer
- View function source code
- Create folder
- Upload file
- Delete file
- Rename file
- Change directory
- Path to displayed directory
- A file browser keyed to your working directory. Click on file or directory name to open.

**Pro Features**

- Share Project with Collaboratory
- Active shared collaborators
- Start new R Session in current project
- Close R Session in project
- Select R Version
- PROJECT SYSTEM**
- File > New Project**
- RStudio saves the call history, workspace, and working directory associated with a project. It reloads each when you re-open a project.
- Name of current project
- RStudio opens plots in a dedicated Plots pane
- Navigate recent plots
- Open in window
- Export plot
- Delete plot
- Delete all plots
- GUI Package manager lists every installed package
- Install Packages
- Update Packages
- Create reproducible package library for your project
- Click to load package with **library()**. Unlick to detach package with **detach()**
- Package version installed
- Delete from library
- RStudio opens documentation in a dedicated Help pane
- Home page of helpful links
- Search within help file
- Search for help file
- Viewer Pane displays HTML content, such as Shiny apps, RMarkdown reports, and interactive visualizations
- Stop Shiny app
- Publish to shinyapps.io, rpubs, RSConnect, ...
- Refresh
- View(<data>)** opens spreadsheet like view of data set
- Filter rows by value or value range
- Sort by values
- Search for value

**Debug Mode**

- Open with **debug()**, **browse()**, or a breakpoint. RStudio will open the debugger mode when it encounters a breakpoint while executing code.
- Click next to line number to add/remove a breakpoint.
- Highlighted line shows where execution has paused
- Run commands in environment where execution has paused
- Examine variables in executing environment
- Select function in traceback to debug
- Step through code one line at a time
- Step into and out of functions to run
- Resume execution
- Quit debug mode

**Version Control with Git or SVN**

- Turn on at **Tools > Project Options > Git/SVN**
- Stage files
- Show file diff
- Commit staged files
- Push/Pull staged files to remote
- View History
- Open shell to type commands
- current branch

**Package Writing**

- File > New Project > New Directory > R Package**
- Turn project into package, enable roxygen documentation with **Tools > Project Options > Build Tools**
- Roxygen guide at **Help > Roxygen Quick Reference**

**RStudio**

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# Entering Input

At the R prompt we type expressions. The `<-` symbol is the assignment operator.

```
> x <- 1
> print(x)
[1] 1
> x
[1] 1
> msg <- "hello"
```

The grammar of the language determines whether an expression is complete or not.

```
> x <- ## Incomplete expression
```

The `#` character indicates a comment. Anything to the right of the `#` (including the `#` itself) is ignored.



# Evaluation

When a complete expression is entered at the prompt, it is evaluated and the result of the evaluated expression is returned. The result may be auto-printed.

```
> x <- 5 ## nothing printed
> x      ## auto-printing occurs
[1] 5
> print(x) ## explicit printing
[1] 5
```

The `[1]` indicates that `x` is a vector and `5` is the first element.

# Printing

```
> x <- 1:20
> x
[1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15
[16] 16 17 18 19 20
```

The `:` operator is used to create integer sequences.

# Objects

R has five basic or “atomic” classes of objects:

- character
- numeric (real numbers)
- integer
- complex
- logical (True/False)

The most basic object is a vector

- A vector can only contain objects of the same class
- BUT: The one exception is a *list*, which is represented as a vector but can contain objects of different classes (indeed, that’s usually why we use them)

Empty vectors can be created with the `vector()` function.

Empty vectors can be created with the `vector()` function.

```
> x <- vector("numeric",5)
> x
[1] 0 0 0 0 0
```

# Numbers

- Numbers in R are generally treated as numeric objects (i.e. double precision real numbers)
- If you explicitly want an integer, you need to specify the `L` suffix
- Ex: Entering `1` gives you a numeric object; entering `1L` explicitly gives you an integer.
- There is also a special number `Inf` which represents infinity; e.g. `1 / 0`; `Inf` can be used in ordinary calculations; e.g. `1 / Inf` is `0`
- The value `NaN` represents an undefined value ("not a number"); e.g. `0 / 0`; `NaN` can also be thought of as a missing value (more on that later)

# Attributes

R objects can have attributes

- names, dimnames
- dimensions (e.g. matrices, arrays)
- class
- length
- other user-defined attributes/metadata

Attributes of an object can be accessed using the `attributes()` function.

# Creating Vectors

The `c()` function can be used to create vectors of objects.



```
> x <- c(0.5, 0.6)      ## numeric
> x <- c(TRUE, FALSE)   ## logical
> x <- c(T, F)          ## logical
> x <- c("a", "b", "c") ## character
> x <- 9:29              ## integer
> x <- c(1+0i, 2+4i)     ## complex
```

Using the `vector()` function

```
> x <- vector("numeric", length = 10)
> x
[1] 0 0 0 0 0 0 0 0 0 0
```

`> x <- vector("logical", length = 10)`

O que seria o “default” ?

Atomic vectors are usually created with `c()`, short for combine:

```
dbl_var <- c(1, 2.5, 4.5)
# With the L suffix, you get an integer rather than a double
int_var <- c(1L, 6L, 10L)
# Use TRUE and FALSE (or T and F) to create logical vectors
log_var <- c(TRUE, FALSE, T, F)
chr_var <- c("these are", "some strings")
```

Atomic vectors are always flat, even if you nest `c()`'s:

```
c(1, c(2, c(3, 4)))
#> [1] 1 2 3 4
# the same as
c(1, 2, 3, 4)
#> [1] 1 2 3 4
```

Given a vector, you can determine its type with `typeof()`, or check if it's a specific type with an “is” function: `is.character()`, `is.double()`, `is.integer()`, `is.logical()`, or, more generally, `is.atomic()`.

```
int_var <- c(1L, 6L, 10L)
typeof(int_var)
#> [1] "integer"
is.integer(int_var)
#> [1] TRUE
is.atomic(int_var)
#> [1] TRUE
```

```
dbl_var <- c(1, 2.5, 4.5)
typeof(dbl_var)
#> [1] "double"
is.double(dbl_var)
#> [1] TRUE
is.atomic(dbl_var)
#> [1] TRUE
```

```
is.numeric(int_var)
#> [1] TRUE
is.numeric(dbl_var)
#> [1] TRUE
```

# Mixing Objects

What about the following?

```
> y <- c(1.7, "a")  ## character  
> y <- c(TRUE, 2)   ## numeric  
> y <- c("a", TRUE) ## character
```

When different objects are mixed in a vector, *coercion* occurs so that every element in the vector is of the same class.

<pre>&gt; x &lt;- c(1.7, "a") &gt; x [1] "1.7" "a"</pre>	<pre>&gt; x &lt;- c(TRUE, 2) &gt; x [1] 1 2</pre>
------------------------------------------------------------------	-----------------------------------------------------------



# Explicit Coercion

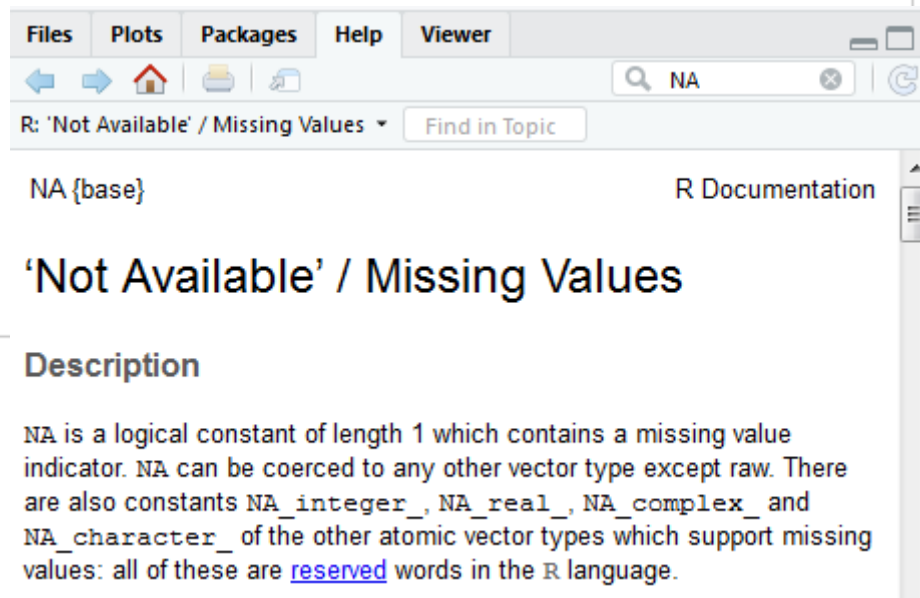
Objects can be explicitly coerced from one class to another using the `as.*` functions, if available.

```
> x <- 0:6
> class(x)
[1] "integer"
> as.numeric(x)
[1] 0 1 2 3 4 5 6
> as.logical(x)
[1] FALSE TRUE TRUE TRUE TRUE TRUE TRUE
> as.character(x)
[1] "0" "1" "2" "3" "4" "5" "6"
```

# Explicit Coercion

Nonsensical coercion results in **NA**s.

```
> x <- c("a", "b", "c")
> as.numeric(x)
[1] NA NA NA
Warning message:
NAs introduced by coercion
> as.logical(x)
[1] NA NA NA
> as.complex(x)
[1] NA NA NA
Warning message:
NAs introduced by coercion
```



The screenshot shows the R Documentation interface. At the top, there are tabs for 'Files', 'Plots', 'Packages', 'Help', and 'Viewer'. Below these is a search bar with the text 'NA' and a 'Find in Topic' button. The main content area is titled 'R: 'Not Available' / Missing Values' and includes a sub-header 'NA {base}'. The 'Description' section explains that NA is a logical constant of length 1 representing a missing value, and lists other related constants like NA\_integer\_, NA\_real\_, NA\_complex\_, and NA\_character\_.

Files Plots Packages Help Viewer

NA

R: 'Not Available' / Missing Values Find in Topic

NA {base} R Documentation

## 'Not Available' / Missing Values

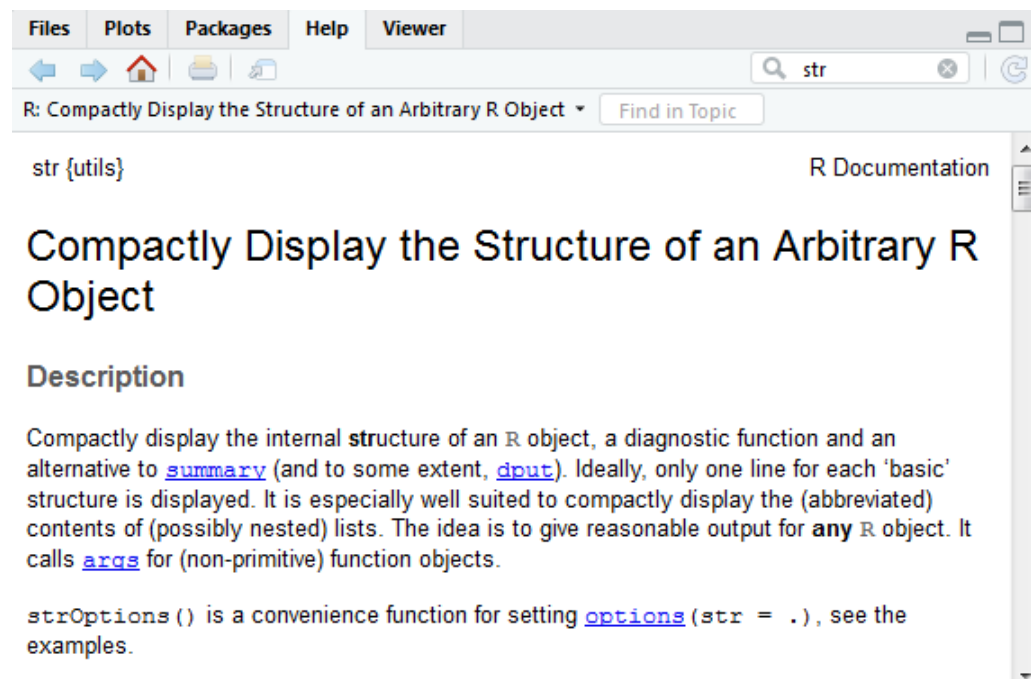
### Description

NA is a logical constant of length 1 which contains a missing value indicator. NA can be coerced to any other vector type except raw. There are also constants `NA_integer_`, `NA_real_`, `NA_complex_` and `NA_character_` of the other atomic vector types which support missing values: all of these are [reserved](#) words in the R language.

All elements of an atomic vector must be the same type, so when you attempt to combine different types they will be **coerced** to the most flexible type. Types from least to most flexible are: logical, integer, double, and character.

For example, combining a character and an integer yields a character:

```
str(c("a", 1))  
#> chr [1:2] "a" "1"
```



The screenshot shows the RStudio interface with the 'Viewer' tab selected. The search bar contains 'str'. Below the search bar, the title bar reads 'R: Compactly Display the Structure of an Arbitrary R Object'. The main content area displays the documentation for the `str` function, including its category 'str {utils}', the title 'Compactly Display the Structure of an Arbitrary R Object', and a 'Description' section. The description explains that `str` is used to compactly display the internal structure of an R object, serving as an alternative to `summary` and `debug`. It also mentions that `strOptions()` is used to set options for the function.

Files Plots Packages Help Viewer

← → Home Print Copy

str

R: Compactly Display the Structure of an Arbitrary R Object Find in Topic

str {utils} R Documentation

## Compactly Display the Structure of an Arbitrary R Object

### Description

Compactly display the internal **structure** of an R object, a diagnostic function and an alternative to [summary](#) (and to some extent, [debug](#)). Ideally, only one line for each 'basic' structure is displayed. It is especially well suited to compactly display the (abbreviated) contents of (possibly nested) lists. The idea is to give reasonable output for **any** R object. It calls [args](#) for (non-primitive) function objects.

`strOptions()` is a convenience function for setting [options\(str = .\)](#), see the examples.

When a logical vector is coerced to an integer or double, TRUE becomes 1 and FALSE becomes 0. This is very useful in conjunction with `sum()` and `mean()`

```
x <- c(FALSE, FALSE, TRUE)
as.numeric(x)
#> [1] 0 0 1
```

```
# Total number of TRUEs
sum(x)
#> [1] 1
```

```
# Proportion that are TRUE
mean(x)
#> [1] 0.3333
```

Coercion often happens automatically. Most mathematical functions (+, log, abs, etc.) will coerce to a double or integer, and most logical operations (&, |, any, etc) will coerce to a logical. You will usually get a warning message if the coercion might lose information. If confusion is likely, explicitly coerce with `as.character()`, `as.double()`, `as.integer()`, or `as.logical()`.

# Lists

Lists are a special type of vector that can contain elements of different classes. Lists are a very important data type in R and you should get to know them well.

```
> x <- list(1, "a", TRUE, 1 + 4i)
> x
[[1]]
[1] 1

[[2]]
[1] "a"

[[3]]
[1] TRUE

[[4]]
[1] 1+4i
```

Lists are different from atomic vectors because their elements can be of any type, including lists. You construct lists by using `list()` instead of `c()`:

```
x <- list(1:3, "a", c(TRUE, FALSE, TRUE), c(2.3, 5.9))
str(x)
#> List of 4
#> $ : int [1:3] 1 2 3
#> $ : chr "a"
#> $ : logi [1:3] TRUE FALSE TRUE
#> $ : num [1:2] 2.3 5.9
```

Lists are sometimes called **recursive** vectors, because a list can contain other lists. This makes them fundamentally different from atomic vectors.

```
x <- list(list(list(list())))  
str(x)  
#> List of 1  
#> $ :List of 1  
#> ..$ :List of 1  
#> .. ..$ : list()  
is.recursive(x)  
#> [1] TRUE
```

`c()` will combine several lists into one. If given a combination of atomic vectors and lists, `c()` will coerce the vectors to list before combining them. Compare the results of `list()` and `c()`:

```
x <- list(list(1, 2), c(3, 4))
y <- c(list(1, 2), c(3, 4))
str(x)
#> List of 2
#> $ :List of 2
#> ..$ : num 1
#> ..$ : num 2
#> $ : num [1:2] 3 4
str(y)
#> List of 4
#> $ : num 1
#> $ : num 2
#> $ : num 3
#> $ : num 4
```



The `typeof()` a list is `list`. You can test for a list with `is.list()` and coerce to a list with `as.list()`. You can turn a list into an atomic vector with `unlist()`. If the elements of a list have different types, `unlist()` uses the same coercion rules as `c()`.

```
Console Terminal x
C:/Users/User/Dropbox/USP/2019/MAP2112/ ↗
> x <- list("a",15,10L,TRUE)
> x
[[1]]
[1] "a"

[[2]]
[1] 15

[[3]]
[1] 10

[[4]]
[1] TRUE

> y <- unlist(x)
> y
[1] "a"      "15"     "10"     "TRUE"
> |
```

## Time-Series Calendar Heatmap

Yahoo Cloiing Price

