Analyzing Financial and Economic Data with R - Online Version

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Analyzing Financial and Economic Data with R

by Marcelo Scherer Perlin

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About New Edition

My plan is to keep updating the content of this book every two years. Its been fun to see how much can change in the R ecosystem in just a couple of years. Here are the main updates in this revision:

- New pipeline operator A new pipeline operator (|>) was introduced in R version 4.1.0. While the old pipeline from {magrittr} can still be found in the wild, my best bet is that, given its native quality and ease of use, the new pipeline will dominate the scene.
- New packages Many of the packages used in previous editions have changed over the years. A couple of packages were dropped from CRAN, and others were substituted by upgraded versions.
- New book package I rewrote all functions in package {afedR3} towards a modular approach, facilitating the future maintenance of the book content. It also includes a testing framework, which will make sure all content in the book is presented as it should.

I hope you enjoy this new edition. Its been a great pleasure to evolve with the book, and I hope I can keep maintaining it over the next decades.

Marcelo S. Perlin,

Porto Alegre, Brazil, 2023-03-08

Preface

Since you are reading this book, you are likely a data analyst looking for alternative and more efficient ways to add value to your organization, an undergraduate or graduate student in the first steps of learning data science, or an experienced researcher, looking for new computational tools. In any case, be assured that you are in the right place. This book will teach you how to use R and RStudio for data analysis in finance and economics.

The first version of the book originates from the class material I teach my postgraduate students in my university. By observing students learning and using R in the classroom, I frequently see the positive impact of technology on their careers. They spend less time doing repetitive and soul-crushing spreadsheet data chores, and more time thinking about their analysis and learning new tools. This book my humble attempt to go beyond the local classroom and reach an international audience.

Another motivation for writing this book is my personal experience using code from other researchers. Usually, the code is not well-organized, lacks clarity, and, possibly, only works in the computer of its author! After being constantly frustrated, I realized the work required to figure out the code of other researchers would take more time than writing the procedure myself. These cases hurt the development of science, as one of its basic principles is the **reproducibility of experiments**. As researchers are expected to be good writers, it should also be expected that their code is in a proper format and readable by other people. With this book, I will tackle this problem by presenting a code structure focused on scientific reproducibility, organization, and usability.

In this book, we will not work on the advanced uses of R. The content will be limited to simple and practical examples. One challenge I had while writing

this book was defining the boundary between introductory and advanced material. Wherever possible, I gradually dosed the level of complexity. For readers interested in learning advanced features and inner workings of R, I suggest the book Venables et al. (2004), Teetor (2011) and Wickham (2019).

This is what you'll learn from this book:

- Using R and RStudio In chapter 01 we will discuss the use of R as a programming platform designed to solve data-related problems in finance and economics. In chapter 02 we will explore basic commands and functionalities that will increase your productivity as a data analyst.
- Importing financial and economic data In chapters 04 and 05 we will learn to import data from local files, such as an Excel spreadsheet, or the internet, using specialized packages that can download financial and economic data such as stock prices, economic indices, the US yield curve, corporate financial statements, and many others.
- Cleaning, structuring and analyzing the data with R In chapters 06 and 07 we will concentrate our study on the ecosystem of basic and advanced classes of objects within R. We will learn to manipulate objects such as numeric vectors, dates and whole tables. In chapters 08 and 09 we'll learn to use the programming to solve data-related problems such as cleaning and structuring messy data. In chapter 11 we will learn applications of the most common econometric models used in finance and economics including linear regression, generalized linear model, Arima model and others.
- Creating a visual analysis of data In chapter 10 we'll learn to use functions from package {ggplot2} to create clever visualizations of our datasets, including the most popular applications in finance and economics, time series and statistical plots.
- Reporting your results In chapter 12 we will see how to report our data analysis using specialized packages and the *RMarkdown* technology. It includes the topic of presenting and exporting tables, figures and models to a written report.
- Writing better and faster code In the last chapter of the book we discuss best programming practices with R. We will look at how to profile code and search for bottlenecks and improving execution time with caching strategies using package {memoise}, C++ code with {Rcpp} and parallel computing with {furrr}.

Conventions

The format of the book was chosen to maximize learnability and memorization. Here are the conventions used throughout the text:

Packages Every R package used in the text will have the textual format of {package}. The first time a R package shows up in the text, a formal citation will also be available.

Functions Functions are formatted as glimpse(), with the information of which package the function belongs to. This notation is simply a copy of real R code, that is, you can call functions using the same structure. The first time the function is referenced, the package name will be included, except for packages that are pre-loaded in a R session ({base}, {utils} and others).

Code All R code will be presented in boxes, with the code output prefixed by string R>. Inline comments are set with the symbol #. Anything on the right side of # is not evaluated by R. Here's an example, showing the contents of a list in R:

```
# create a list
my_list <- list('xx', 1:5, 'dec')

# print list
print(my_list)

R> [[1]]
R> [1] "xx"
R>
R> [[2]]
R> [1] 1 2 3 4 5
R>
R> [[3]]
R> [1] "dec"
```

Supplement Material

All the material used in the book, including code examples separated by chapters, is publicly available on the internet and distributed with an R package called {afedR3}. It includes data files and several functions that can make it easier to run the examples of the book. If you plan to write some code as you read the book, this package will greatly help your journey.

In order to install the book package in your computer, you need to execute a couple of lines of code in R. For that, copy and paste the following commands into RStudio prompt (bottom left of screen, with a ">" sign) and press enter for each command. Be aware you'll need R and RStudio installed in your computer (see section 1.4 for details).

```
# install devtools dependency
install.packages('devtools')

# install book package
devtools::install_github('msperlin/afedR3')
```

What this code will do is to install package {devtools}, a required dependency for installing a package from Github, which is where the book bundle is hosted. After that, a call to devtools::install_github('msperlin/afedR3') will install the package in your computer. You can safely ignore any warning messages about long paths during installation.

After installing package {afedR3}, you can, but its not necessary, to copy all book files to a local folder by executing the following command in R:

```
afedR3::bookfiles_get(path_to_copy = '~/afedR3')
```

The previous code will unzip the book file into your "Documents/afedR3" folder, as the tilda (~) is a shortcut to your "Documents" directory¹. If you prefer the old-fashioned way of using an internet page, you can find and download the package zip file from github².

A suggestion, before you read the rest of the book: go to the book website and search for the related links page at the bottom. There you will find all internet addresses highlighted in the text, including the links for the installation of R and RStudio.

Content for Instructors

If you are an R instructor, you'll find plenty of material you can use with your classes. I made sure you get everything you need:

Over 100 exercises Every chapter in this book includes exercises that your students can practice, with solutions available in the web version of the book. Also, all exercises are available in the exams for-

¹In R, you can type path.expand('~') to see exactly where is your "Documents" folder located.

²https://github.com/msperlin/afedR3

mat, meaning that you can compile the same exercises in pdf or html. Moreover, you can export the exercises to e-learning platforms such as Moodle and Blackboard. See this blog post³ for instructions on how to use it with your students.

Web version The first seven chapters of the book are freely available at link https://www.msperlin.com/afedr, which is more than enough material for an introductory class on R and data analysis.

All of this content is released with the MIT license, so feel free to use and abuse it, as long as you give the credits to the original author. You can find the content within the book package {afedR3} (see previous instructions on installation) or directly at the book site⁴.

I hope you enjoy this book and find it useful for your work.

Good reading!

Marcelo S. Perlin

³https://www.msperlin.com/post/2023-03-10-compiling-exercises-afedR3/

⁴https://www.msperlin.com/publication/2020 book-afedr-en/



Introduction

In the digital era, information is abundant and accessible. From the ever-changing price of financial contracts to the unstructured data of social media websites, the high volume of information creates a strong need for data analysis in the workplace. A company or organization benefit immensely when it can create a bridge between raw information from its environment and making strategic decisions. Undoubtedly, this is a prolific time for professionals skilled in using the right tools for acquiring, storing, and analyzing data.

In particular, datasets related to Economics and Finance are widely available to the public. International and local institutions, such as central banks, government research agencies, financial exchanges, and many others, provide their data publicly, either by legal obligation or to foment research. Whether you are looking into statistics for a particular country or a company, most information is just two clicks away.

Not surprisingly, it is expected that a graduate student or a data analyst has learned at least one programming language that allows him/her to do his work more efficiently. **Learning how to program is becoming a requisite for the job market**. This is where the role and contribution of R comes into play. In the next sections, I will explain what R is and why you should use it.

1.1 What is R

R is a programming language specially designed to resolve statistical problems and display graphical representations of data. R is a modern version of S, a programming language originally created in Bell Laboratories (formerly AT&T, now Lucent Technologies). The base code of R was developed by two academics, **Ross Ihaka** and **Robert Gentleman**, resulting in the programming platform we have today. For anyone curious about the name, the letter R was chosen due to the common first letter of the name of their creators.

Today, R is almost synonymous with data analysis, with a large user base and consolidated modules. It is likely that researchers from various fields, from economics to biology, find in R significant preexisting code that facilitates their analysis. On the business side, large and established companies, such as *Google* and *Microsoft*, already adopted R as the internal language for data analysis. R is maintained by the **R Foundation**¹ and the **R Consortium**², a collective effort to fund projects for extending the programming language.

1.2 Why Choose R

Learning a new programming language requires a lot of time and effort. Perhaps you're wondering why you should choose R and invest time in learning it. Here are the main arguments.

First, R is a mature and stable platform, continuously supported and intensively used in the industry. When choosing R, you will have the computational background not only for an academic career but also to work as a data analyst in private organizations. Due to its open license, you can use R anywhere. Also, the strong support from the community means it is very unlikely the R platform will ever fade away or be substituted. Depending on your career choices, R might be the only programming language you ever need to learn.

Learning R is easy. My experience in teaching R allows me to confidently state that students, even those with no programming experience, have no problem learning the language and using it to create their own code. The language is intuitive and certain rules and functions can be extended to different cases. Once you understand how the software expects you to think, it become easy to traverse over different modules and functionalities.

The engine of R and the interface of RStudio creates a highly productive environment. The graphical interface provided by RStudio facilitates the use of R and increases productivity by introducing new features to the platform. By combining both, the user has at his disposal many

¹https://www.r-project.org/foundation/

²https://www.r-consortium.org/

tools that facilitate the development of research scripts and other projects.

R Packages allow the user to do many different things with R. We will soon learn that R offers several modules that can be installed over the internet whenever necessary. These modules extend the basic language of R and enable the most diverse functionalities. Besides basic data tasks such as reading and writing, you can, for example, use R to build and publish a blog, send emails, create exams, write random jokes and poems (seriously!), and many other features. The existing external modules in R are truly an impressive achievement of the community.

R is compatible with different operating systems and it can interface with different programming languages. If you need to execute code from another programming language, such as C++, Python, Julia, it is easy to integrate it with R. Therefore, the user is not restricted to a single programming language and can easily use features and functions from others. For example, the C++ code is well known for its superior speed in numerical tasks. From an R script, you can use package $\{Rcpp\}$ to write a C++ function and effortlessly use it within your R code.

R is free! The main software and all its packages are free. A generous license motivates the adoption of the R language in a business environment, where obtaining individual and collective licenses of commercial software can be costly. This means you can take R anywhere you go, regardless of whether you have a budget for software or not.

1.3 What Can You Do With R and RStudio?

R is a fairly complete programming language and any computational problem can be solved based on it. Given the adoption of R for different areas of knowledge, the list is extensive. With finance and economics, I can highlight the following possibilities:

- Substitute and improve data-intensive tasks from spreadsheet-like software;
- Develop routines for managing investment portfolios and executing financial orders;
- Creating tools for calculating and reporting economic indices such as inflation and unemployment;

- Performing empirical data research using statistical techniques, such as econometric models and hypothesis testing;
- Create dynamic websites with the {shiny} package, allowing anyone in the world to use a computational tool created by you;
- Automate the process of writing technical reports with the RMarkdown and Quarto technology;

Moreover, public access to packages developed by users further expands these capabilities. The CRAN views website³ offers a *Task Views* panel for the topic of Finance⁴ and Econometrics⁵. There you can find the main packages to perform specific operations such as importing financial data from the internet, estimating econometric models, calculation of different risk estimates, among many other possibilities. Reading this page and the knowledge of these packages is essential for those who intend to work in Finance and Economics.

1.4 Installing R

Before going any further, let's install the required software on your computer. The most direct and practical way to install R is to direct your favourite internet browser to R website⁶ and click the *Download* link in the left side of the page, as shown in Figure 1.1.

The next screen gives you a choice of the mirror to download the installation files. The CRAN repository (R Comprehensive Archive network) is mirrored in various parts of the world. You can choose one of the links from the nearest location to you. If undecided, just select the mirror θ -Cloud (see Figure 1.2), which will automatically take you to the nearest location.

The next step involves selecting your operating system, likely to be Windows. From now on, due to the greater popularity of this platform, we will focus on installing R in Windows. The instructions for installing R in other operating systems can be easily found online. Regardless of the underlying platform, using R is about the same. There are a few exceptions, especially when R interacts with the file system. In the content of the book, special care was taken to choose functions that work the same way in different operating systems. A few exceptions are highlighted throughout the book. So, even if

³https://cran.r-project.org/web/views

⁴https://cran.r-project.org/web/views/Finance.html

⁵https://cran.r-project.org/web/views/Econometrics.html

⁶http://www.r-project.org/



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CRAN

R Project

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Developer Pages
R Bloa

The R Project for Statistical Computing

Getting Started

R is a free software environment for statistical computing and graphi 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 sc or what the license terms are, please read our answers to frequently questions before you send an email.

News

- R version 4.2.2 (Innocent and Trusting) has been released or 10-31
- R version 4.1.3 (One Push-Up) was released on 2022-03-10.
- Thanks to the organisers of useR! 2020 for a successful online conference. Recorded tutorials and talks from the conference are available on the R Consortium YouTube channel.

Figure 1.1: Initial page for downloading R

you are using a Mac or a flavor of Linux, you can take full advantage of the material presented here.

After clicking the link $Download\ R$ for Windows, as in Figure 1.3, the next screen will show the following download options: base, contrib, old.contrib and RTools. The first (base), should be selected. It contains the download link to the executable installation file of R in Windows.

Some R packages requires local compilation of the files. For that, you need RTools, a bundle of compilers and utilities. So, you can safely install RTools from CRAN website.

After clicking the link base, the next screen will show the link to the download of the R installation file. After downloading the file, open it and follow the steps in the installation screen. At this time, no special configuration is required. I suggest keeping all the default choices and simply hit accept in the displayed dialogue screens. After the installation of R, it is strongly recommended to install RStudio, which will be addressed next.

Belnet, the Belgian research and

CRAN Mirrors

The Comprehensive R Archive Network is available at the following URLs, please che statistics on the status of the mirrors can be found here: main page, windows release

If you want to host a new mirror at your institution, please have a look at the **CRAN N**

0-Cloud

Brazil

Automatic redirection to servers https://cloud.r-project.org/ sponsored by Rstudio Argentina Universidad Nacional de La Plata http://mirror.fcaglp.unlp.edu.ar/CRAN/ Australia CSIRO https://cran.csiro.au/ https://mirror.aarnet.edu.au/pub/CRAN/ AARNET School of Mathematics and Statis https://cran.ms.unimelb.edu.au/ Melbourne https://cran.curtin.edu.au/ Curtin University Austria https://cran.wu.ac.at/ Wirtschaftsuniversität Wien Belgium https://www.freestatistics.org/cran/ Patrick Wessa

Figure 1.2: Choosing the CRAN mirror

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 (Debian, Fedora/Redhat, Ubuntu)
- Download R for macOS
- Download R for Windows

https://ftp.belnet.be/mirror/CRAN/

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

Figure 1.3: Choosing the operating system

R for Windows

Subdirectories:



Binaries for base distribution. This is what you want Binaries of contributed CRAN packages (for R >= 3 Binaries of contributed CRAN packages for outdate. Tools to build R and R packages. This is what you w Windows, or to build R itself.

Figure 1.4: Installation options



Be aware that R has a consistent release schedule. Every four months a new version of R is released, fixing bugs and implementing new solutions. There are two main types of releases, major and minor. For example, today, 2023-02-23, the latest version of R is 4.2.2. The first digit ("4") indicates the major release while all others are of the *minor* type. Generally, the *minor* changes are very specific and, possibly, will have little impact on your work. However, unlike minor releases, major releases are fully reflected in the R package ecosystem. Every time you install a new major version of R, you will have to reinstall all packages. Particularly, the problem here is that it is not uncommon that a new major release comes with package incompatibility issues. My advice is: every time a new major release of R comes out, wait a few months before installing it on your machine. Thus, the authors of the packages will have more time to update their codes, minimizing the possibility of compatibility problems.

1.5 Installing RStudio

The base installation of R includes its own GUI (graphical user interface), where we can write and execute code. However, this native interface has several limitations. RStudio Desktop substitutes the original GUI and makes access to R more practical and efficient. One way to understand this relationship is with an analogy with cars. While R is the engine of the programming language, RStudio is the body and instrument panel, which significantly improves the user experience. With RStudio you'll have code highlight, creation of projects, and much more.

The installation of RStudio is simpler than that of R. Direct your favourite

browser to Posit (formerly RStudio) website⁷ and click in *Download RStudio* and then *Download RStudio Desktop*. After that, just select the installation file relative to the operating system on which you will work. This option is probably *WINDOWS Vista* 7/8/10. Note that RStudio is also available for Mac and Linux.

I emphasize that using RStudio is not essential to develop programs in R. Other interfaces are available and can be used. However, in my experience, RStudio is the interface that offers a vast range of features for the language and is widely used, which justifies its choice. If you want to explore other programming interfaces for R, one that I really enjoy and use is Microsoft's VSCode⁸.

1.6 Resources in the Web

The R community is vivid and engaging. There are many authors, such as myself⁹, that constantly release material about R in their blogs. It includes the announcement of new packages, analysis of real world datasets, curiosities, rants, and tutorials. R-Bloggers¹⁰ is a website that aggregates these blogs, making it easier for anyone to access and participate. I strongly recommend to sign up for the R-Bloggers feed in RSS¹¹, Facebook¹² or Twitter¹³. Not only you'll be informed of what is happening in the R community, but also learn a lot by reading other people's code and articles.

Learning and using R can be a social experience. Several conferences and user groups are available in many countries. You can find the complete list in this $link^{14}$. I also suggest looking in social platforms for local R groups in your region.

1.7 Structure and Organization

This book presents a practical approach to using R in finance and economics. To get the most out of it, I suggest you first try to understand what the code does and, after that, use it on your own computer. Whenever you find a piece

⁷https://posit.co/

⁸https://code.visualstudio.com/

⁹https://www.msperlin.com

¹⁰https://www.r-bloggers.com/

¹¹https://feeds.feedburner.com/RBloggers

¹²https://www.facebook.com/rbloggers/?fref=ts

¹³https://twitter.com/Rbloggers

¹⁴https://jumpingrivers.github.io/meetingsR/index.html

of code that you do not understand, go on and study it. At first, it might seem like a daunting task but, with time, be confident that the learning process will get a lot easier as the code blocks will start to make sense and connect to each other.

Learning to program in a new platform is like learning a foreign spoken language: the use in day-to-day problems is imperative to create fluency. All the code and data used in this book is available with the installation of package {afedR3} (see the preface for instructions on how to install it). I suggest you test the code on your computer and play with it, modifying the examples and checking the effect of changes in the outputs. Whenever you have a computational problem, try using R to solve it. You'll stumble and make mistakes at first. But I guarantee that, soon enough, you'll be able to write complex data tasks effortlessly.

Throughout the book, every demonstration of code will have two parts: the R code and its output. The output is nothing more than the textual result of the commands on the screen. All inputs and outputs code will be marked in the text with a special format. See the following example:

```
# create a list
this_list <- list('abc', 1:5, 'dec')

# print list
print(this_list)

R> [[1]]
R> [1] "abc"
R>
R> [[2]]
R> [1] 1 2 3 4 5
R>
R> [[3]]
R> [1] "dec"
```

For the previous chunk of code, lines this_list <- list('abc', 1:5, 'dec') and print(this_list) are actual commands given to R. The output of this simple piece of code is the on-screen presentation of the contents of object this_list.

The code can also be spatially organized using newlines. This is a common strategy around arguments of functions. The next chunk of code is equivalent to the previous and will run the exact same way. Notice how we used a new line to vertically align the arguments of function list. You'll soon see that, throughout the book, this type of vertical alignment is constantly used.

The code also follows a well-defined structure. One decision in writing computer code is how to name objects and how to structure it. It is recommended to follow a clear pattern, so it is easy to maintain over time and be used and understood by others. For this book, a mixture of the author's personal choices with the coding style suggested by Google¹⁵ was used. The reader, however, may choose the structure he finds more efficient and aesthetically pleasing. Like many things in life, this is a choice. We will get back at discussing code structure in chapter 13.

1.8 Exercises

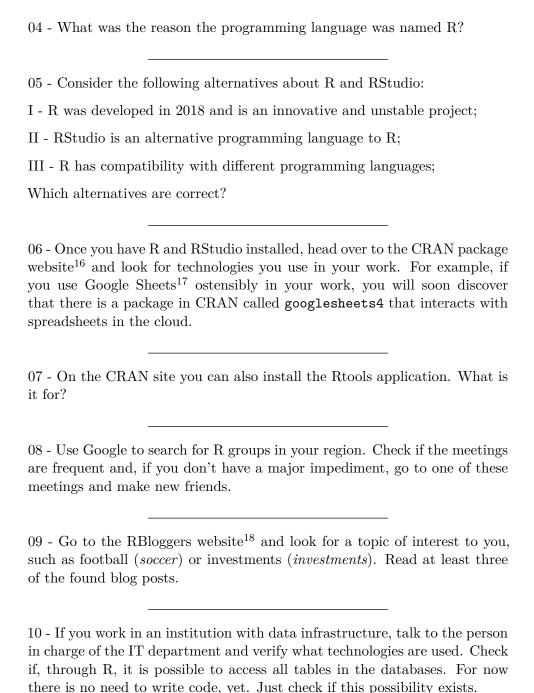
01 - The R language was developed based on what other programming language?

02 - What are the names of the two authors of R?

03 - Why is R special when comparing to other programming languages, such as Python, C++, javascript and others?

 $^{^{15} \}rm https://google.github.io/styleguide/Rguide.xml$

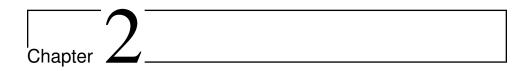
1.8. EXERCISES 29



 $^{^{16} \}rm https:$ //cloud.r- project.org/web/packages/available_packages_by_date.html

¹⁷https://www.google.com/sheets/about/

¹⁸https://www.r-bloggers.com/



Basic Operations in R

When working with R and RStudio, there are fundamental tasks (or basic operations) that you will be repeating many times over. In this chapter we will look at these basic operations with RStudio, including:

- Rstudio interface and shortcuts
- basic R commands
- working with files
- working with related file extensions
- the autocomplete feature of RStudio.

Here, we will go through the initial steps from the viewpoint of someone who has never worked with R and possibly never had contact with other programming language. Those already familiar with the software may not find novel information here and, therefore, I suggest skipping to the next section. However, I recommended that you at least check the discussed topics so you can confirm your knowledge about the features of the software and how to use them for working smarter, and not harder. This is especially true for RStudio, which offers several tools to increase your productivity.

2.1 Working With R

The greatest hurdle a new user faces when developing routines in R is the format of work – the so-called development cycle. Our interaction with computers has been simplified over the years and we are currently comfortable with the point @click format. That is, if you want to perform an operation

on the computer, just point the *mouse* to a specific location on the screen and click a button. Visual cues in a series of steps allow the execution of complex tasks. Be aware, however, that this form of interaction is just one layer above what actually happens. Behind all these *clicks*, there is a command being executed on your computer. Any common task such as opening a *pdf* file, a spreadsheet document, directing a *browser* to a web page has an underlying call to a code.

The point&click format of visual and motor interaction has its benefits in facilitating and popularizing the use of computers. However, it is not flexible and effective when working with computational procedures such as data analysis. A better approach would be to create a file containing several instructions in sequence and, in the future, simply request that the computer **execute** this file using the recorded procedures. There is no need to do a "scripted" point and click operation. You spend some time studying commands and writing the program but, in the future, it will always execute the recorded procedure in the same way.

Using *scripts* provides a significant gain in productivity when comparing to a *point&click* type of interface. Going further, the risk of human error in executing the procedure is almost nil, because the commands and their required sequence of execution are recorded in the text file and will always be executed in the same way. This is one of the main reasons why programming languages are popular in science. All the steps of a data-based research, including results, can be replicated by different people, in different computers.

While programming in R, the ideal format of work is to merge the mouse movement with commands. R and RStudio have some functionality with the *mouse*, but their capacity is optimized when we perform operations using code. When a group of commands is performed in a smart way, we have an R script that should preferably produce something important to us at the end of its execution. In finance and economics, this can be the current price of a stock, the value of an economic index such as inflation, the result of academic research, among many other possibilities.

Like other software, R allows us to import data and export files. We can use code to import a dataset stored in a local file – or the web–, analyze it and paste the results into a technical report. Going further, we can use RStudio and the *RMarkdown* technology to write a dynamic report, where code and content are integrated. Needless to say that, by using the capabilities of R and RStudio, you will work smarter and faster.

The final product of working with R and RStudio will be an R script that

produces digital elements for a data report. A good example of a simple and polished R script can be found at this link¹. Open it and you'll see the content of a file with extension .R that will download stock prices of two companies and create a plot and a table. By the end of the book, you will understand what is going on in the code and how it gets the job done. Even better, you'll be able to improve it. Soon, you'll learn to execute the code on your own computer. If impatient, simply copy the text content of the link to a new RStudio R script, save it, and press control + shift + enter.

2.2 Objects in R

In R, everything is an object, and each type of object has its properties. For example, the daily closing prices of the IBM stock over 2023 can be represented as a numerical vector, where each element is a price recorded at the end of a trading day. Dates related to these prices can be represented as text (*string*) or as a unique Date class. Finally, we can represent the price data and the dates together by storing them in a single object of type dataframe, which is nothing more than a table with rows and columns.

While we represent data as objects in R, a special type is a function. It stores a pre-established manipulation of other objects available to the user. R has an extremely large number of functions, which enable the user to perform a wide range of operations. For example, the basic commands of R, available in the package {base}, adds up to a total of 1258 functions.

Each function has its own name and a programmer can write their own functions. For example, the **sort()** function is a procedure that sorts elements within a vector. If we wanted to sort the elements of 2, 1, 4, 3, 1, simply insert the following command in the *prompt* (left bottom side of RStudio's screen) and press *enter*:

```
my_vec <- c(2, 1, 4, 3, 1)

sorted_vec <- sort(
    x = my_vec,
    decreasing = TRUE
    )

print(sorted_vec)</pre>
```

 $^{^1} https://github.com/msperlin/afedR3/blob/main/inst/extdata/others/S_Example_Script.R$

R> [1] 4 3 2 1 1

The **sort()** function is used with start and end parentheses. These parentheses serve to highlight the entries (*inputs*), that is, the information sent to the function to produce something that will be saved in object **sorted_vec**. Note that each entry is separated by a comma, as in my_fct(input1, input2, input3, ...). We also set option decreasing = TRUE. This is a specific directive for the **sort()** function to order the value from highest to lowest.

Be aware that functions are at the heart of R and we will dedicate a large part of this book to them. You can use the available functions or write your own. You can also publish functions as a package and let other people use your code. We will discuss more about functions in chapter 8.

2.3 International and Local Formats

Before explaining the use of R and RStudio, it is important to highlight some rules of formatting numbers, latin characters and dates.

decimal: Following an international notation, the decimal point in R is defined by the period symbol (.), as in 2.5 and not a comma, as in 2,5. If this is not the standard format in your country, you'll have issues when importing local data from text files. Sometimes, such as with storing data in Microsoft Excel files, the reading function already takes care of the conversion. This, however, is generally an exception. As a general rule of using R, only use commas to separate the inputs of a function. Under no circumstances should the comma symbol be used as the decimal point separator. Always give priority to the international format because it will be compatible with the vast majority of data.

Latin characters: Due to its international standard, R has problems understanding Latin characters, such as the cedilla and accents. If you can, avoid using Latin characters in the names of your variables or files. In the content of character objects (text), you can use them without problems as long as the encoding of the script is correctly specified (e.g. UTF-8, Latin1). I strongly recommend the use of the English language for writing code and defining object names. This automatically eliminates the use of Latin characters and facilitates the usability of the code by people outside of your country.

date format: Dates in R are structured according to the ISO 8601² format. It follows the YYYY-MM-DD pattern, where YYYY is the year in four numbers (e.g. 2023), MM is the month as a number and DD is the day. An example

²https://www.iso.org/iso-8601-date-and-time-format.html

of date is 2023-03-10. This, however, may not be the case in your country. When importing local data, make sure the dates are in this format. If necessary, you can convert any date to the ISO format. Again, while you can work with your local format of dates in R, it is best advised to use the international notation. The conversion between one format and another is quite easy and will be presented in chapter 7.

If you want to learn more about your local format in R, use the following command by typing it in the prompt and pressing enter:

Sys.localeconv()

R>	decimal_point	thousands_sep	grouping
R>	"."	11 11	11 11
R>	<pre>int_curr_symbol</pre>	currency_symbol	${\tt mon_decimal_point}$
R>	"BRL "	"R\$"	","
R>	mon_thousands_sep	mon_grouping	positive_sign
R>	"."	"\003\003"	11 11
R>	negative_sign	int_frac_digits	frac_digits
R>	"_"	"2"	"2"
R>	p_cs_precedes	p_sep_by_space	n_cs_precedes
R>	"1"	"1"	"1"
R>	n_sep_by_space	p_sign_posn	n_sign_posn
R>	"1"	"1"	"1"

The output of **Sys.localeconv()** shows how R interprets decimal points and the thousands separator, among other things. As you can see from the previous output, this book was compiled using the Brazilian notation for the currency symbol, but uses the dot point for decimals.



Be careful when modifying the format that R interprets symbols. As a rule of thumb, if you need to use a specific format, do it separately within the context of the code. Avoid permanent changes as you never know where such formats are being used. That way, you can avoid unpleasant surprises in the future.

2.4 Types of Files in R

Like any other programming platform, R has a ecosystem of file extensions, where each has a different purpose. In the vast majority of cases, however, the work will focus mostly on a couple of types. Next, I describe the various file extensions you'll find in a day to day basis. The items in the list are ordered by importance. Note that I omitted graphic files such as .png, .jpg,

.gif and data storage/spreadsheet files (.csv, .xlsx, ..) among others, as they are not exclusive to R.

Files with extension .R: text files containing R code. These are the files which you will spend most of your time. They contain the sequence of commands that configures the main script and computational routines of the data research. Examples: Script-stock-research.R, R-fcts.R.

Files with extension .RData or .rds: files that store data in the native format. These files are used to save/write objects created in different sessions into your hard drive. For example, you can use a .rds file to save a table after processing and cleaning up the raw database. By freezing the data into a local file, we can later load it for subsequent analysis. Examples: cleaned-inflation-data.rds, model-results.RData.

Files with extension .Rmd and .quarto: files used for editing dynamic documents in the RMarkdown and Quarto format. Using these files allows the creation of documents where text and code output are integrated into the same document. While RMarkdown is mostly related to R, the quarto format allows for a more flexible integration of text and code for other programming languages such as Python. In chapter 12 we have a dedicated section for RMarkdown, which will explore this functionality in detail. Example: investment-report.Rmd.

Files with extension .Rproj: contain files for editing projects in RStudio, such as a new R package, a shiny application or a book. While you can use the functionalities of RStudio projects to write R scripts, it is not a hard requirement. For those interested in learning more about this functionality, I suggest the Posit manual³. Example: project-retirement.Rproj.

2.5 Explaining the RStudio Screen

After installing the R and RStudio, open RStudio by double-clicking its icon. Be aware that R also has its own interface and this often causes confusion. In Windows, you can search for RStudio link using the *Start* button and typing *Rstudio*.

After opening RStudio, the resulting window should look like Figure 2.1.

Note that RStudio automatically detected the installation of R and initialized a session on the left side of the interface.

As a first exercise, click File, New File, and R Script. A text editor should

³https://support.posit.co/hc/en-us/articles/200526207-Using-Projects

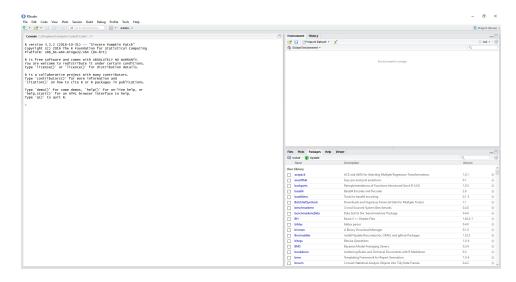


Figure 2.1: The RStudio screen

appear on the left side of the screen. It is there that we will spend most of our time developing code. Commands are executed sequentially, from top to bottom. A side note, all .R files created in RStudio are just text files and can be edited anywhere. As an exercise, use Windows's Notepad to open an R file. You'll see that the raw code is the same, but without the syntax highlighting.

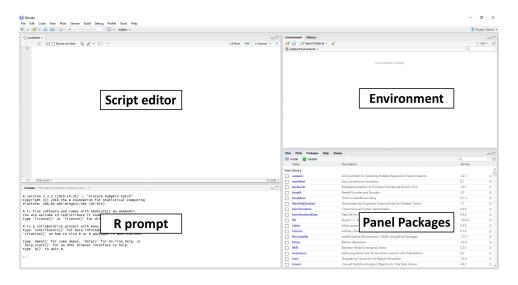


Figure 2.2: Explaining the RStudio screen



When using RStudio, a common suggestion is to change the color scheme to a **dark mode** setting. It is not just an aesthetic issue, but also a strategy for preventing eye strain. Since you will be spending a lot of time in front of the computer, it is smart to change the colors of the interface to relieve your eyes of the constant brightness of the screen. That way, you'll be able to work longer, without straining your vision. You can configure the color scheme of RStudio by going to the option *Tools*, *Global Options* and then *Appearance*. A dark color scheme that I personally like and suggest is *Ambience*.

After the previous steps in RStudio, the resulting screen should look like the image in Figure 2.2. The main items/panels are:

Script Editor: located on the left side and above the screen. This panel is used to write scripts and functions, mostly on files with the .R extension;

R prompt: on the left side and below the script editor. It displays the *prompt*, which can also be used to give commands to R. The main purpose of the *prompt* is to test code and display the results of the commands entered in the script editor;

Environment: located on the top-right of the screen. Shows all objects, including variables and functions, currently available to the user. Also note a *History* panel, which shows the history of commands previously executed by the user;

Panel Packages: shows the packages installed and loaded by R. Here you have four tabs: *Files*, to load and view system files; *Plots*, to view statistical figures created in R; *Help* to access the help system and *Viewer* to display dynamic and interactive results, such as a web page.

As an introductory exercise, let's initialize two objects in R. Inside the prompt (lower left side), insert these commands and press enter at the end of each. The <- symbol is nothing more than the result of joining < (less than) with the - (minus sign). The ' symbol represents a single quotation mark and, in most computer keyboards, it is found under the escape (esc) key (top left).

```
# set x
x <- 1
# set y
y <- 'My humble text'</pre>
```

If done correctly, notice that two objects are available in the *environment* panel, one called x with a value of 1, and another called y with the text content "My humble text". Also noticed how we used specific symbols to define objects x and y. The use of double quotes (" ") or single quotes (' ') defines objects of the class character. Numbers are defined by the value itself. As will be discussed later, understanding R object classes are important as each has a different behavior within the R code. After executing the previous commands, notice that the *history tab* has been updated.

Now, lets show the values of x on the screen. To do this, type the following command:

```
# print contents of x
print(x)
```

R> [1] 1

The **print()** function is one of the main commands for displaying values in the *prompt* of R. The text displayed as [1] indicates the index of the first line number. To verify this, enter the following command, which will show a lengthy sequence of numbers on the screen:

```
# print a sequence
print(50:100)
R>
     [1]
          50
               51
                    52
                         53
                              54
                                   55
                                       56
                                            57
                                                 58
                                                      59
                                                           60
                                                               61
                                                                    62
                                                                         63
```

```
R> [15]
           64
                65
                     66
                           67
                                68
                                     69
                                          70
                                               71
                                                    72
                                                         73
                                                              74
                                                                    75
                                                                         76
                                                                              77
R> [29]
           78
                79
                     80
                           81
                                82
                                          84
                                                    86
                                                         87
                                                              88
                                                                    89
                                                                         90
                                                                              91
                                     83
                                               85
R> [43]
                93
           92
                     94
                          95
                                96
                                     97
                                          98
                                               99 100
```

Here, we use the colon symbol in 50:100 to create a sequence starting at 50 and ending at 100. Note that, on the left side of each line, we have the values [1], [13], and [25]. These represent the index of the first element presented in the line. For example, the fifteenth element of 50:100 is 64.

2.6 R Packages

One of the greatest benefits of using R is its package collection. A package is nothing more than a group of procedures aimed at solving a particular computational problem. R has at its core a collaborative philosophy. Users provide their codes for others to use. And, most importantly, **all packages are free**. For example, consider the scenario where a user is interested in accessing data about historical inflation in the USA. He can install and use an R module that is specifically designed for importing data from central banks and research agencies.

Every function in R belongs to a package. When R initializes, packages stats, graphics, grDevices, utils, datasets, methods and base are loaded by default. This is way why we can use some functions in R without the explicit call to a library. For example, function print is from the base package and is available whenever you start R.

R packages can be accessed and installed from different sources. The main being **CRAN** (*The Comprehensive R Archive network*), and **Github**. It's worth knowing that the quantity and diversity of R packages increases every day. **CRAN** is the official repository of R and it is built and maintained by the community. One of the reasons for the success of CRAN is the quality of code. While anyone can send a package, there is an evaluation process to ensure that certain strict rules about code format and safety are respected. For those interested in creating and distributing packages in CRAN, a clear roadmap on is available on the R packages site⁴. The official (and complete) rules are available on the CRAN website⁵.



So far, I have eight package published in CRAN. In my experience, sending and publishing a package on CRAN can demand a significant amount of work, especially in the first submission. After that, it becomes a lot easier. Don't be angry if your package is rejected. My own packages were rejected several times before entering CRAN. Listen to what the maintainers tell you and try fixing all problems before resubmitting. If you're having issues that you cannot solve or find a solution on the Internet, look for help in the R-packages mailing list⁶. You'll probably be surprised at how accessible and helpful the R community can be.

The complete list of packages available on CRAN, along with a brief description of each, can be accessed at the packages section of the R site⁷. A practical way to check if there is a package that does a specific procedure is to load the previous page and search in your *browser* for a keyword of interest (e.g. "SEC data"). If there is a package that does what you want, it is very likely that the keyword is used in the description field.

Another important source for finding packages is the CRAN Task Views⁸. There you can find the collection of noteworthy packages for a given area of expertise. See the *Task Views* screen in Figure 2.3.

⁴https://r-pkgs.org/

⁵https://cran.r-project.org/web/packages/policies.html

⁷https://cran.r-project.org/web/packages/available packages by date.html

⁸https://cran.r-project.org/web/views/

CRAN Task Views Bayesian ChemPhys Bayesian Inference Chemometrics and Computational Physics ClinicalTrials Clinical Trial Design, Monitoring, and Analysis Cluster Analysis & Finite Mixture Models Cluster <u>DifferentialEquations</u> Differential Equations Probability Distribution: Econometrics Econometrics Environmetrics
Experimental Design Analysis of Ecological and Environmental Data Design of Experiments (DoE) & Analysis of Experimental Data ExtremeValue Extreme Value Analysis Empirical Finance Genetics Statistical Genetics Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization HighPerformanceComputing High-Performance and Parallel Computing with R MachineLearning Machine Learning & Statistical Learning Medical Image Analysis MedicalImaging MetaAnalysis Meta-Analysis Multivariate Statistics Natural Language Processing Natural Language Processing Numerical Mathematics
Official Statistics
Official Statistics & Survey Methodology OfficialStatistics Optimization Optimization and Mathematical Programming Analysis of Pharmacokinetic Data Phylogenetics Phylogenetics, Especially Comparative Methods Psychometric Models and Methods ReproducibleResearch Reproducible Research Robust Statistical Methods

Figure 2.3: Task View screen

A popular alternative to CRAN is Github⁹. Unlike the former, Github imposes no restrictions on the submitted code and, because of this flexibility and its version control system, it is a popular choice by developers. In practice, it is very common for developers to maintain a development version of a module on Github and the official version in CRAN. When the development version reaches a certain stage of maturity, it is then sent to CRAN.

The most interesting part about R packages is that they can be installed directly from the prompt using the internet. To find out the current amount of packages on CRAN, type and execute the following commands in the prompt:

```
# get a matrix with available packages
df_cran_pkgs <- available.packages()

# find the number of packages
n_cran_packages <- nrow(df_cran_pkgs)

# print it
print(n_cran_packages)</pre>
```

R> [1] 19278

Currently, 2023-03-10 08:20:07, there are 19278 packages available on the CRAN servers, a very impressive mark for the community of developers as a whole.

⁹https://github.com/

You can also check the amount of **locally installed packages** in R with the **installed.packages()** command:

```
# find number of packages currently installed
n_local_packages <- nrow(installed.packages())
# print it
print(n_local_packages)</pre>
```

R> [1] 628

In this case, the computer in which the book was compiled has 628 packages currently installed. Notice that, even as an experienced R programmer, I'm only using a small fraction of all packages available in CRAN!

2.6.1 Installing Packages from CRAN

Use command **install.packages()** to install a package locally. You only need to do it once for each new package. As an example, we will install a package called **{readr}**, that will be used in future chapters. Note that we defined the package name in the installation as if it were text with the use of quotation marks (" ").

```
# install package readr
install.packages("readr")
```

That's it! After executing this simple command, package {readr} will be installed and the functions related to the package will be ready for use once the package is loaded in a script. If the installed package is dependent on another package, R detects this dependency and automatically installs the missing packages. Thus, all the requirements for using the installed package are satisfied and everything will work perfectly. Be aware, however, that some modules can require external software in the level of the operating system. These cases are usually announced in the description of the package and an error informs that a requirement is missing. External dependencies for R packages are not common, but they do happen.

2.6.2 Installing Packages from Github

To install a package hosted in Github, you must first install the {devtools} package, available on CRAN:

```
# install devtools
install.packages('devtools')
```

After that, use function **install_github()** to install a package directly from Github. In the following example, we will install the development version of package {dplyr}:

```
# install ggplot2 from github
devtools::install_github("hadley/dplyr")
```

Note that the username of the developer is included in the input string. In this case, the *hadley* name belongs to the developer of **{dplyr}**, Hadley Wickham. Throughout the book, you will notice that this name appears several times. Hadley is a prolific and competent developer of several popular R packages and currently works for RStudio.



Be aware that **github packages are not moderated**. Anyone can send code there and the content is not independently checked. Never install github packages without some confidence of the author's work. Although unlikely - it never happened to me, for example - it is possible that they have malicious code.

2.6.3 Loading Packages

Within a script, use function **library()** to load a package, as in the following example.

```
# load package readr
library(readr)
```

After running this command, all functions of the package will be available in the **current** R session. Whenever you close RStudio or start a new session, you'll lose all loaded packages. This is the reason why packages are usually loaded in the top of the script: starting from a clean memory, required packages are sequentially loaded before the actual R code is executed.

If the package you want to use is not available, R will throw an error message. See an example next, where we try to load a non-existing package called unicorn.

```
library(unicorn)
```

```
R> Error in library(unicorn): There is no package called "unicorn"
```

Remember this error message. It will appear every time a package is not found. If you got the same message when running code from this book, you need to check what are the required packages of the example and install them using install.packages().

R> R>

Alternatively, if you want to use a specific function and do not want to load all functions from the same package, you can do it with the use of double colons (::), as in the following example.

```
fortunes::fortune(10)

R>
R> Overall, SAS is about 11 years behind R and S-Plus in
R> statistical capabilities (last year it was about 10 years
R> behind) in my estimation.
```

example of using a function without loading package

-- Frank Harrell (SAS User, 1969-1991)

R-help (September 2003)

Here, we use function **fortune()** from the package **{fortunes}**, which shows on screen a potentially funny phrase chosen from the R mailing list. For our example, we selected message number 10. One interesting use of package **{fortunes}** is to display a random joke every time R starts and, perhaps, lighten up your day. As mentioned before, R is fully customizable. You can find many tutorials on how to achieve this effect by searching on the web for "customizing R startup".

Another way of loading a package is by using the **require()** function. A call to **require()** has a different behavior than a call to **library()**. Whenever you try to load an uninstalled package with the **library()** function, it returns an error. This means that the script stops and no further code are evaluated. As for **require()**, if a package is not found, it returns an object with value **FALSE** and the rest of the code is evaluated. So, in order to avoid code being executed without its explicit dependencies, it is best advised to always use **library()** for loading packages in R scripts.

The use of **require()** is left for loading up packages inside of functions. If you create a custom function that requires procedures from a particular package, you must load the package within the scope of the function. For example, see the following code, where we create a new function called **fct_example** that depends on the package **{quantmod}**:

```
fct_example <- function(x){
    require(quantmod)

    df <- getSymbols(x, auto.assign = F)
    return(df)
}</pre>
```

In this case, the first time that fct_example is called, it loads up the package {quantmod} and all of its functions. Using require() inside a function is a good programming policy because the function becomes self-contained, making it easier to use it in the future. This was the first time where the complete definition of a user-created function in R is presented. Do not worry about it for now. We will explain it further in chapter 8.



Be aware that loading a package can cause a **conflict of functions**. For example, there is a function called **filter** in the **dplyr** package and also in the **stats** package. If we load both packages and call the **filter** function within the scope of the code, which one will R use?

Well, the **preference** is always for the last loaded package. This is a type of problem that can be very confusing. Fortunately, note that R itself tests for conflicts when loading a package. Try it out: start a new R session and load the dplyr package. You'll see that a message indicates that there are two conflicts with the stats package – functions filter and lag – and four with the base package.

A simple strategy to avoid bugs due to conflict of function is to call a function using the actual package name. For example, if I'm calling lag from dplyr, I can write the call as dplyr::lag. As you can see, the package name is explicit, avoiding any possible conflict.

2.6.4 Upgrading Packages

Over time, it is natural that packages available on CRAN are upgraded to accommodate new features and fix bugs. Thus, it is recommended that users update their installed packages to a new version. In R, a simple and direct way of upgrading packages is to click the button *Update* located in the package panel, lower right corner of RStudio, as shown in Figure 2.4.

The user can also update packages using commands. Simply type command update.packages() and hit *enter*, as shown below.

```
# update all installed packages
update.packages()
```

The command **update.packages()** compares the version of the installed packages with the versions available in CRAN. If it finds any difference, the new version is downloaded and installed. After running the command, all packages will be synchronized with the versions available in CRAN.

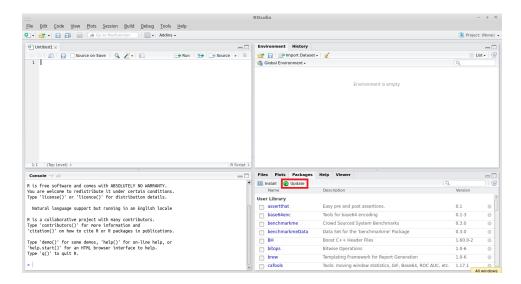


Figure 2.4: Updating R packages



Package versioning is an extremely important topic for keeping your code reproducible. Although it is uncommon to happen, a new package version can change the results of your analysis. I have a particularly unpleasant experience when a scientific article returned from a journal review and, due to the update of one of the R packages I used, I was unable to reproduce the results presented in the article. In the end everything went well, but the trauma remains.

One solution to this problem is to freeze the package versions for each project using RStudio's renv tool. In summary, renv makes local copies of the packages used in the project, which have preference over system packages. Thus, if a package is updated in the system, but not in the project, the R code will continue to use the older version and the R code will always run under the same conditions.

2.7 Running Scripts from RStudio

Now, let's copy some code into the editor's screen (upper left side). The result should look like Figure 2.5.

```
# a complete script
x <- 1
y <- "my humble text"</pre>
```

```
print(x)

R> [1] 1
print(y)
```

R> [1] "my humble text"

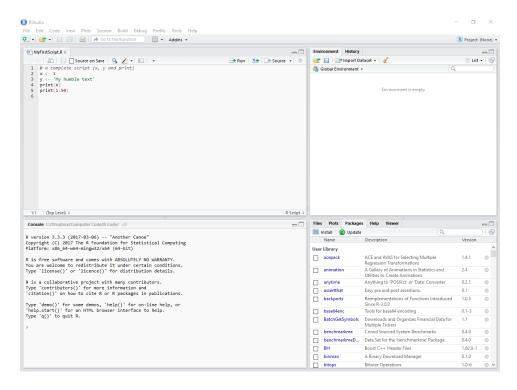


Figure 2.5: Example of an R script

After pasting all the commands in the editor, save the R file to a personal folder where you have read and write permissions. In Windows, one possibility is to save it in the <code>Documents</code> folder with a name like <code>'my_first_script.R'</code>. This saved file, which at the moment does nothing special, records the steps of a simple algorithm that creates several objects and shows their content.

2.8 Using the help files

There is no shame in seeking help. Advanced R users often seek instructions on specific tasks, whether it is to better understand the execution details of

some functions or simply to study a new procedure. The use of the R help system is part of the work and you should master it as soon as possible.

Within RStudio, you can get help by using the *help* panel in RStudio or directly from the *prompt*. Simply enter the question mark next to the object on which you want help, as in ?mean. In this case, object means is a function and the use of the help() command will open a panel on the right side of RStudio. Another way of opening the help page of a function is the select the name of the function and press F1 in the keyboard.

In R, the help screen of a function is the same as shown in Figure 2.6. It presents a general description of the function, explains its input arguments and the format of the output. The help screen follows with references and suggestions for other related functions. More importantly, examples of usage are given last and can be copied to the prompt or script in order to accelerate the learning process.

mean {base} R Documentation

Arithmetic Mean

Description

Generic function for the (trimmed) arithmetic mean.

Usage

```
mean(x, ...)
## Default S3 method:
mean(x, trim = 0, na.rm = FALSE, ...)
```

Arguments

x An R object. Currently there are methods for numeric/logical vectors and <u>date</u>, <u>date-time</u> and <u>time interval</u> objects. Complex vectors are allowed for trim = 0, only.

trim the fraction (0 to 0.5) of observations to be trimmed from each end of x before the mean is computed. Values of trim outside that range are taken as the nearest endpoint.

na.rm a logical value indicating whether NA values should be stripped before the computation proceeds.

Figure 2.6: Help screen for function mean

If we are looking for help for a given text and not a function name, we can use double question marks as in ??"standard deviation". This operation will search for the occurrence of the term in all packages of R and it is very useful to learn how to perform a particular task. In this case, we looked for the available functions to calculate the standard deviation of a vector.

As a suggestion, the easiest and most direct way to learn a new function is by trying out the examples in the manual. This way, you can see which types of inputs the function expects and what type of output it provides back. Once you have it working in your code, read the help screen to understand if it does exactly what you expected and what are the options for its use. If the function performs the desired procedure, you can copy and paste the example for your own *script*, adjusting where necessary.

Another very important source of help is the Internet itself. Sites like stackoverflow¹⁰ and specific mailing lists and blogs, whose content is also on the Internet, are a valuable source of information. If you find a problem that could not be solved by reading the standard help files, the next logical step is to seek a solution using your error message or the description of the problem in search engines. In many cases, your problem, no matter how specific it is, has already occurred and has been solved by other users. In fact, it is more surprising not to find the solution for a programming problem on the internet, than the other way around.



Whenever you ask for help on the internet, always try to 1) describe your problem clearly and 2) add a reproducible code of your problem. Thus, the reader can easily verify what is happening by running the example on his computer, and solving the problem.

2.8.1 RStudio shortcuts

In RStudio, there are some predefined and time-saving shortcuts for running code from the editor. To execute an entire script, simply press control + shift + s. This is the *source* command. With RStudio open, I suggest testing this key combination and checking how the code saved in a R file is executed. The output of the script is shown in the prompt of R. The result in RStudio should look like Figure 2.7.

Another way of executing code is with the shortcut **control** + **enter**, which will only execute the line where the cursor is located. This shortcut is very useful in developing scripts because it allows each line of the code to be tested. As an example of usage, point the cursor to the print(x) line and press control + enter. As you will notice, only the line print(x) was executed and the cursor moves to the next line. Therefore, before running the whole script, you can test it line by line and check for possible errors.

Next, I highlight these and other RStudio shortcuts, which are also very useful.

¹⁰http://stackoverflow.com/

```
| Supplication | Supp
```

Figure 2.7: Example of a R script after execution

control + shift + s executes (source) the current RStudio file;

control + shift + enter executes the current file with echo, showing the commands on the prompt;

control + enter executes the selected line, showing on-screen commands;
 control + shift + b executes the codes from the beginning of the file to the cursor's location;

control + **shift** + **e** executes the codes of the lines where the cursor is until the end of the file.

I suggest you try to create a healthy habit by using these shortcuts from day one. Those who like to use the *mouse*, an alternate way to execute code is to click the *source* button in the upper-right corner of the code editor.

If you want to run code in a R file within another R file, you can use the **source()** command. For example, imagine that you have a main R script with your data analysis and another two scripts that performs some support operation such as importing data to R. These operations have been dismembered as a way of organizing the code.

To run the support *scripts*, just call it with function **source** in the main script, as in the following code:

```
# execute import script
source('01-import-data.R')
# execute analysis
```

```
source('02-build-tables')
```

Here, all code in 01-import-data.R and 02-build-tables.R will be executed sequentially. This equals manually opening each file and hitting control + shift + s.

2.9 Testing and Debugging Code

Developing code follows an observable cycle. At first, you will write a command line on a script, try it using control + enter, and check the result on the prompt or the content of objects. A new line of code is written once the previous line worked as expected. A moving cycle is clear, writing code is followed by line execution, followed by result checking, modify and repeat if necessary. This is a normal and expected process. You need to make sure that every line of code is correctly specified before moving to the next one. Whenever the code asks for something that is not expected (or possible), an error occurs.

When trying to find an error in a preexisting script, R offers debugging tools for controlling and assessing its execution. This is especially useful when you have a long and complicated script. The simplest and easiest tool that R and RStudio offer is **code breakpoint**. In RStudio, you can click on the left side of the script editor and a red circle will appear, as in Figure 2.8.

```
1  # set x
2  x <- 1
3
4  # set y
5  y <- 'My humble text'
6
7  # print contents of x
8  print(x)</pre>
```

Figure 2.8: Example of breakpoint in an R script

This red circle indicates a flag that will force the code to stop at that line. You can use it to test existing code and check its objects at a certain part of the execution. Pausing the code at a certain point might seem strange for a starting programmer but, for large scripts, with many functions and complex code, it is a necessity. When the execution hits the breakpoint, the prompt will change to Browse[1]> and you'll be able to try new code

and verify the content of all current objects. From the Console, you have the option to continue the execution to the next breakpoint or stop it by pressing shift+f8. The same result can be achieved using a function called browser. Have a look:

```
# set x
x <- 1

# set y
y <- 'My humble text'

browser()

# print contents of x
print(x)</pre>
```

The practical result is the same as using RStudio's red circle, but it gives you more control for the case of several commands in the same line.

2.10 Creating Simple Objects

One of the most basic and most used commands in R is the creation of objects. As shown in previous sections, you can define an object using the <- command, which is verbally translated to *assign*. For example, consider the following code:

```
# set x
x <- 123
# set my_x, my_y and my_z in one line
my_x <- 1; my_y <- 2; my_z <- 3
```

We can read this code as the value 123 is assigned to x. The direction of the arrow defines where the value is stored. For example, using 123 -> x also works, although this is not recommended as the code becomes less readable. Moreover, notice that you can create objects within the same line by separating the commands using a semi-colon.



Using an arrow symbol <- for object definition is a simple way to identify R code. The reason for this choice was that, at the time of conception of the S language, keyboards had a specific key that directly defined the arrow symbol. This means that the programmer only had to hit one key in the keyboard to create new variables. Modern keyboards, however, are different. If you find it troublesome to type this symbol, you can use a shortcut as well. In Windows, the shortcut for the symbol <- is alt plus "-".

Most programming languages uses a equality symbol (=) to define objects and, often, this creates confusion. When using R, you can also define objects with =, as in x = 123, however, no one should ever recommend it. The equality symbol has a special use within an R code as it defines function arguments, as in $my_1 \leftarrow fct(arg1 = 1, arg2 = 3)$. For now, just remember to use <- for defining objects. We will learn more about functions and using the equality symbol in a future chapter.

The name of the object is important in R. With the exception of very specific cases, you can name objects as you please. This freedom, however, can work against you. It is desirable to set short object names that make sense to the content of the script and which are simple to understand. This facilitates the understanding of the code by other users and is part of the suggested set of rules for structuring code. Note that all objects created in this book have nomenclature in English and specific format, where the white space between nouns are replaced by an underscore, as in my_x <- 1 and name_of_file <- 'my_data_file.csv'. We will get back at code structure in chapter 13.

R executes code by looking for objects available in the environment, including functions. You also need to be aware that R is case sensitive. That is, object \mathtt{m} is not the same as M. If we try to access an object that does not exist, R will return an error message and halt the execution of the rest of the code. Have a look:

print(z)

R> Error in print(z): object 'z' not found

The error occurred because the object z does not exist in the current environment. If we create a variable z as z <-321 and repeat the command print(z), we will not have the same error message.

2.11 Creating Vectors

In the previous examples, we created simple objects such as x <-1 and x <-1ABC'. While this is sufficient to demonstrate the basic commands in R, in practice, such commands are very limited. A real problem of data analysis will certainly have a larger volume of information.

When we gather many elements of the same class, such as numeric, into a single object, the result is an atomic vector. An example would be the representation of a series of daily stock prices as an atomic vector of the class numeric. Once you have a vector, you can manipulate it any way you want.

Atomic vectors are created in R using the $\mathbf{c}()$ command, which comes from the verb *combine*. For example, if we want to combine the values 1, 2 and 3 in a single object, we can do it with the following command:

```
# create numeric atomic vector
x <- c(1, 2, 3)
# print it
print(x)</pre>
```

R> [1] 1 2 3

The c() command works the same way for any other class of object, such as *character*:

```
# create character atomic vector
y <- c('text 1', 'text 2', 'text 3', 'text 4')
# print it
print(y)</pre>
```

```
R> [1] "text 1" "text 2" "text 3" "text 4"
```

The only restriction when creating vectors is that all elements must have the same class. If we insert data from different classes in a call to $\mathbf{c}()$, R will try to mutate all elements into the same class following its own logic. If the conversion of all elements to a single class is not possible, an error message is returned. Note the following example, where numeric values are set in the first and second element of \mathbf{x} and a character in the last element.

```
# a mixed vector
x <- c(1, 2, '3')
```

```
# print result of forced conversion
print(x)
```

```
R> [1] "1" "2" "3"
```

Notice that all values in x are converted to type character. The use of class() command confirms this result:

```
# print class of x
class(x)
```

R> [1] "character"

2.12 Knowing Your Environment and Objects

After using various commands, further development of an R script requires you to understand what objects are available and if their content are correct. You can find this information simply by looking at the $\it environment$ tab in the upper right corner of RStudio. However, there is a command that shows the same information in the prompt. In order to know what objects are currently available in R's memory, you can use the command ls(). Note the following example:

```
# set some objects
x <- 1
y <- 2
z <- 3
# print all objects in the environment
print(ls())</pre>
```

```
R> [1] "x" "v" "z"
```

Objects x, y and z were created and are available in the current working environment. If we had other objects, they would also appear in the output to ls().

To display the content of each object, just enter the names of objects and press enter in the *prompt*:

```
# print objects by their name
x
```

```
y
R> [1] 2
z
```

R> [1] 3

Typing the object name on the screen has the same effect as using the **print()** command. In fact, when executing the sole name of a variable in the prompt or script, R internally passes the object to the **print()** function.

In R, all objects belong to a class. As previously mentioned, to find the class of an object, simply use the **class()** function. In the following example, x is an object of the class numeric, y is a text (character) object and fct_example is a function object.

```
# set objects
x <- 1
y <- 'a'
fct_example <- function(){}

# print their classes
print(class(x))</pre>
```

```
R> [1] "numeric"
print(class(y))
```

```
R> [1] "character"
print(class(fct_example))
```

```
R> [1] "function"
```

Another way to learn more about an object is to check its textual representation. Every object in R has this property and we can find it with function **str()**:

```
# set vec
x <- 1:10
# print the textual representation of a vector
print(str(x))</pre>
```

```
R> int [1:10] 1 2 3 4 5 6 7 8 9 10 R> NULL
```

We find that object x is a vector of class int (integer). Function str() is

particularly useful when trying to understand the details of a more complex object, such as a dataframe or a list, which will be introduced in future chapter.

2.13 Finding the Size of Objects

In R, an object size can mean different things but most likely it is defined as the number of individual elements that constitute the object. This information serves not only to assist the programmer in checking possible code errors but also to know the length of iteration procedures such as *loops*, which will be treated in a later chapter of this book.

In R, the size of an object can be checked with the use of four main functions: length(), nrow(), ncol() and dim().

Function **length()** is intended for objects with a single dimension, such as atomic vectors:

```
# create atomic vector
x <- c(2, 3, 3, 4, 2,1)

# get length of x
n <- length(x)

# display message
message('The length of x is ', n)</pre>
```

R> The length of x is 6

For objects with more than one dimension, such as a matrix, use functions $\mathbf{nrow}()$, $\mathbf{ncol}()$ and $\mathbf{dim}()$ (dimension) to find the number of rows (first dimension) and the number of columns (second dimension). See the difference in usage below.

```
# create a matrix
M <- matrix(1:20, nrow = 4, ncol = 5)
# print matrix
print(M)</pre>
```

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

```
# calculate size in different ways
my_nrow <- nrow(M)
my_ncol <- ncol(M)
my_n_elements <- length(M)

# display messages
message('The number of lines in M is ', my_nrow)

R> The number of lines in M is 4
message('The number of columns in M is ', my_ncol)
R> The number of columns in M is 5
```

R> The number of elements in M is 20

The dim() function shows the dimension of the object, resulting in a numeric vector as output. This function should be used when the object has more than two dimensions. In practice, however, such cases are rare as most data-related problems can be solved with a bi-dimensional representation. An example is given next:

message('The number of elements in M is ', my_n_elements)

```
# get dimension of M
my_dim <- dim(M)

# print it
print(my_dim)</pre>
```

R> [1] 4 5

In the case of objects with more than two dimensions, we can use the **array()** function to create the object and **dim()** to find its size. Have a look at the next example:

```
# create an array with three dimensions
my_array <- array(1:9, dim = c(3, 3, 3))
# print it
print(my_array)</pre>
```

```
R> , , 1
R>
R> [,1] [,2] [,3]
R> [1,] 1 4 7
```

```
R > [2,]
            2
                  5
                        8
R > [3,]
            3
                  6
                        9
R>
R> , , 2
R>
R>
         [,1] [,2] [,3]
R> [1,]
            1
                  4
                        7
R > [2,]
            2
                  5
                        8
R > [3,]
            3
                  6
                        9
R>
R> , , 3
R>
R>
         [,1] [,2] [,3]
R> [1,]
            1
R > [2,]
            2
                  5
                        8
R> [3,]
            3
                  6
# display its dimensions
print(dim(my_array))
```

R> [1] 3 3 3

2.14 Selecting Elements from an Atomic Vector

After creating an atomic vector, you can access and select portions of it. For example, suppose that a month ago you invested 1.000 USD in Apple shares. You download stock price data for previous month and want to see how much your portfolio is currently worth. In that case, you are interested only in latest price of the stocks. All price values from other dates are not relevant to our analysis and therefore could be safely ignored.

The selection of *pieces* of an atomic vector is called **indexing** and it is accomplished with the use of square brackets ([]). Consider the following example:

```
# set x
my_x <- c(1, 5, 4, 3, 2, 7, 3.5, 4.3)
```

If we wanted only the third element of my_x, we use the bracket operator as follows:

```
# get the third element of x
elem_x <- my_x[3]
# print it
print(elem_x)</pre>
```

R> [1] 4

Indexing also works using vectors containing the desired locations. If we are only interested in the last and penultimate values of my_x, we use the following code:

```
# set vector with indices
my_idx <- (length(my_x)-1):length(my_x)

# get last and penultimate value of my_x
piece_x_1 <- my_x[my_idx]

# print it
print(piece_x_1)</pre>
```

```
R> [1] 3.5 4.3
```

A cautionary note: a unique property of the R language is that if a non-existing element of an object is accessed, the program returns the value NA (not available). See the next example code, where we attempt to obtain the fourth value of a vector with only three components.

```
# set object
my_vec <- c(1, 2, 3)

# print non-existing fourth element
print(my_vec[4])</pre>
```

R> [1] NA

This is important because NA elements are contagious. That is, anything that interacts with NA will also become NA. The lack of treatment of these errors can lead to problems that are difficult to identify. In other programming languages, attempting to access non-existing elements generally returns an error and cancels the execution of the rest of the code.



In most cases, the occurrence of NA (*Not Available*) values suggests that a problem exists within the code. Always remember that NA values indicates lack of data and are contagious: anything that interacts with a NA value will turn into another NA. You should become suspicious about your code quality every time that NA values are found unexpectedly.

The use of indices is very useful when you are looking for items of a vector that satisfy some condition. For example, if we wanted to find out all values in my_x that are greater than 3, we could use the following command:

```
# find all values in my_x that is greater than 3
piece_x_2 <- my_x[my_x>3]

# print it
print(piece_x_2)
```

```
R> [1] 5.0 4.0 7.0 3.5 4.3
```

It is also possible to index elements by more than one condition using the logical operators & and | (or). For example, if we wanted the values of my_x greater than 2 and lower than 4, we could use the following command:

```
# find all values of my_x that are greater than 2 and lower then 4 piece_x_3 <- my_x[ (my_x > 2) & (my_x < 4)] print(piece_x_3)
```

```
R> [1] 3.0 3.5
```

Likewise, if we wanted all items that are lower than 3 **or** greater than 6, we use:

```
# find all values of my_x that are lower than 3 or higher than 6
piece_x_4 <- my_x[ (my_x < 3) | (my_x > 6) ]
# print it
print(piece_x_4)
```

```
R> [1] 1 2 7
```

Moreover, logic indexing also works with the interaction of different objects. That is, we can use a logical condition in one object to select items from another:

```
# set my_x and my.y
my_x <- c(1, 4, 6, 8, 12)
```

```
my_y <- c(-2, -3, 4, 10, 14)

# find all elements of my_x where my.y is higher than 0
my_piece_x <- my_x[my_y > 0 ]

# print it
print(my_piece_x)
```

```
R> [1] 6 8 12
```

Looking more closely at the indexing process, notice that we are creating a variable of the logical type. This object takes only two values: TRUE and FALSE. Have a look in the code presented next, where we create a logical object, print it and present its class.

```
# create a logical object
my_logical <- my_y > 0

# print it
print(my_logical)
```

```
R> [1] FALSE FALSE TRUE TRUE TRUE
```

```
# find its class
class(my_logical)
```

```
R> [1] "logical"
```

Logical objects are very useful whenever we are testing a particular condition on a data set. We will learn more about this and other basic classes in chapter 7.

2.15 Removing Objects from the Memory

After creating several variables, the R environment can become full of used and disposable content. In this case, it is desirable to clear the memory to erase objects that are no longer needed. Generally, this is accomplished at the beginning of a script, so that every time the script runs, the memory will be cleared before any calculation. In addition to cleaning the computer's memory, it also helps to avoid possible errors in the code.

For example, given an object x, we can delete it from memory with the command rm(), as shown next:

```
# set x
x <- 1
# remove x
rm('x')</pre>
```

After executing the command rm('x'), the value of x is no longer available in the R session. In practical situations, however, it is desirable to clean up all the memory used by all objects created in R. We can achieve this goal with the following code:

```
rm(list = ls())
```

The term list in rm(list = ls()) is a function argument of rm() that defines which objects will be deleted. The ls() command shows all the currently available objects. Therefore, by chaining together both commands, we erase all current objects available in the environment. As mentioned before, it is a good programming policy to clear the memory before running the script. However, you should only wipe out all of R's memory if you have already saved the results of interest or if you can replicate them.



Clearing memory in *scripts* is a controversial topic. Some authors argue that it is better not to clear the memory as this can erase important results. My opinion is that it is important to clear the memory at the top of the script, as long as all results are reproducible. When you start a code in a clean state – no variables or functions – it becomes easier to understand and solve possible bugs.

2.16 Displaying and Setting the Working Directory

Like other programming platforms, R always executes code in a working directory. If no directory is set, a default value is used when R starts up. It is based on the working directory that R searches for files to load data or other R scripts. It is in this directory that R saves any output if we do not explicitly define a path on the computer. This output can be a graphic file, text or a spreadsheet. That said ,it is good policy to change the working directory to the same place where the *script* is located.

The simplest way of checking the current working directory is looking at RStudio's prompt panel. At the top, in a small font and just below the word

"Console", you'll see the working path. Using code, we can check the current working directory with function **getwd()**:

```
# get current dir
my_dir <- getwd()

# display it
print(my_dir)</pre>
```

R> C:/my-books/afedR-ed4/book-content

The result of the previous code shows the folder in which this book was written and compiled.

The change of the working directory is performed with the **setwd()** command. For example, if we wanted to change our working directory to C:/My Research/, simply type in the prompt:

```
# set where to change directory
my_d <- 'C:/My Research/'
# change it
setwd(my_d)</pre>
```

After changing the directory, importing and saving files in the C:/My Research/ folder will be a lot easier.

Additionally, the easiest and most straightforward way to ensure that the working directory is the same as the R script is using **RStudio projects**. To do so, click in "File" and "New Project". Doing so will create a .Rproj file in the chosen directory of the project. The trick here is to create the R scripts in the root folder of the project. Every time you open the project, it will automatically change the working directory to where the .Rproj file is located.

Once you are working on the same path as the script, using relative paths is preferable. For example, if you are working in a folder that contains a subdirectory called data, you can enter this sub-folder with the code:

```
# change to subfolder
setwd('data')
```

Another possibility is to go to a previous level of the directory using ..., as in:

```
# change to the previous level
setwd('..')
```

So, if you are working in directory C:/My Research/ and execute the command setwd('..'), the current folder becomes C:/, which is one level below C:/My Research/.

2.17 Canceling Code Execution

Whenever R is running some code, a visual cue in the shape of a small red circle with the word stop in the right corner of the prompt will appear. This button is not only an indicator that R is busy running code but also a shortcut for canceling its execution. Another way to cancel an execution is to point the mouse to the prompt and press the escape (esc) button from the keyboard.

To try it out, run the next chunk of code in RStudio and cancel its execution using esc.

```
for (i in 1:100) {
  message('\nRunning code (please make it stop by hitting esc!)')
  Sys.sleep(1)
}
```

In the previous code, we used a for loop and function Sys.sleep to display the message '\nRunning code (please make it stop by hitting ESC!)' one hundred times, every second. For now, do not worry about the code and functions used in the example. We will discuss the use of loops in chapter 8.



Another very useful trick for defining working directories in R is to use the ~ symbol. The tilda defines the "Documents" folder in Windows, which is unique for each user. For Linux and Mac users, the tilda defines the "home" folder (e.g. "/home/USERNAME"). Therefore, by running setwd('~'), you will direct R to a personal folder in any operating system.

2.18 Code Comments

In R, comments are set using the hashtag symbol #. Any text after this symbol will not be processed. This gives you the freedom to write whatever you want within the script. An example:

```
# this is a comment (R will not parse it)
# this is another comment (R will again not parse it)
```

```
x <- 'abc' # this is an inline comment
```

Comments are an effective way to communicate any important information that cannot be directly inferred from the code. In general, you should avoid using comments that are too obvious or too generic:

```
# read CSV file
df <- read.csv('data/data_file.csv')</pre>
```

As you can see, it is quite obvious from the line df <- read.csv('..') that the code is reading a .csv file. The name of the function already states that. So, the comment was not a good one as it did not add any new information to the user. A better approach at commenting would be to set the author, description of the script and better explain the origin and last update of the data file. Have a look:

```
# Script for reproducing the results of JOHN (2019)
# Author: Mr data analyst (dontspamme@emailprovider.com)
# Last script update: 2020-01-10
#
# File downloaded from www.site.com/data-files/data_file.csv
# The description of the data goes here
# Last file update: 2020-01-10

df <- read.csv('data/data_file.csv')</pre>
```

So, by reading the comments, the user will know the purpose of the script, who wrote it and the date of the last edit. It also includes the origin of the data file and the date of the latest update. If the user wants to update the data, all he has to do is go to the referred website and download the new file.

Another productive use of comments is to set sections in the code, such as in:

```
# - remove outliers
# - remove unnecessary columns

# Create descriptive tables -----

# Estimate models -------

# Report results ------
```

The use of a long line of dashes (-) at each section of the code is intentional. It causes RStudio to identify and bookmark the sections, with a link to them at the bottom of the script editor. Test it yourself, copy and paste the above code into a new R script, save it, and you'll see that the sections appear on a button between the editor and the prompt. Such a shortcut can save plenty of time in lengthy scripts.



When you start to share code with other people, you'll soon realize that comments are essential and expected. They help transmit information that is not available from the code. This is one way of a discerning beginners from experienced programmers. Contrary to popular belief, it is likely that someone with experience in programming will be very communicative in its comments (sometimes too much!). A note here, throughout the book you'll see that the code comments are, most of the time, a bit obvious. This was intentional as clear and direct messages are important for new users, which is part of the audience of this book.

2.19 Using Code Completion with tab

A very useful feature of RStudio is $code\ completion$, an editing tool that facilitates the search of object names, packages, function arguments, and files. Its usage is very simple. After you type any first letter in the keyboard, just press tab (left side of the keyboard, above capslock) and a number of options will appear. See Figure 2.9 where, after entering the f letter and pressing tab, a small window appears with a list of object names that begin with that letter.

The autocomplete feature is self-aware and will work differently depending on where it is called. As such, it works perfectly for searching for packages. For

```
Type 'demo()' for some demos, 'help()' for on-line help, or 'help.start()' for an HTML browser interface to help. Type 'q()' to quit R.
```



Figure 2.9: Usage of autocomplete for object name

that, type library() in the prompt or editor, place the cursor in between the parentheses and press tab. The result should look something like Figure 2.10, shown next.

Figure 2.10: Usage of autocomplete for packages

Note that a description of the package or object is also offered by the code completion tool. This greatly facilitates the day to day work as the memorization of package names and R objects can be challenging. The use of the tab decreases the time to look up names, also avoiding possible errors.

The use of this tool becomes even more beneficial when objects and functions are named with some sort of pattern. In the rest of the book, you will notice that objects tend to be named with the prefix my, as my_x , my_num , my_df . Using this naming rule (or any other) facilitates the lookup for the names of objects created by the user. You can just type my_n , press tab, and a list of objects will appear.

As mentioned in the previous section, you can also find files and folders on your computer using tab. To try it, write the command $my_file <-$ "" in the prompt or a script, point the cursor to the middle of the quotes and press the tab key. A screen with the files and folders from the current working directory should appear, as shown in Figure 2.11.

```
Type 'demo()' for some demos, 'help()' for on-line help, or 'help.start()' for an HTML browser interface to help.

Type 'q()' to quit R.
```

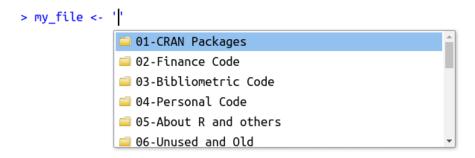


Figure 2.11: Usage of autocomplete for files and folders

The use of autocomplete is also possible for finding the name and description of function arguments. To try it out, write message() and place the mouse cursor inside the parentheses. After that, press tab. The result should be similar to Figure 2.12. By using tab inside of a function, we have the names of all arguments and their description – a mirror of the information found in the help files.

Likewise, you can also search for a function within a package with *tab*. For that, simply type the name of the package followed by two commas, as in readr::, and press *tab*. The result should be similar to Figure 2.13

Summing up, using code completion will make you more productive. You'll find names of files, objects, arguments, and packages much faster. Use it as much as you can.



Autocomplete is one of the most important tools of RStudio, helping users to find object names, locations on the hard disk, packages and functions. Get used to using the tab key and, soon enough, you'll see how much the autocomplete tool can help you write code quickly, and without typos.

```
R version 4.0.3 (2020-10-10) -- "Bunny-Wunnies Freak Out"
Copyright (C) 2020 The R Foundation for Statistical Computing
Platform: x86_64-pc-linux-gnu (64-bit)
R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.
  Natural language support but running in an English locale
R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.
> sort()
       for sort an R object with a class or a numeric, complex,
       decreasing =
                           character or logical vector. For sort.int, a numeric, complex,
                           character or logical vector, or a factor.
                           Press F1 for additional help
```

Figure 2.12: Usage of autocomplete for function arguments



Figure 2.13: Usage of autocomplete for finding functions within a package

2.20 Interacting with Files and the Operating System

As you are learning R, soon enough you'll find a data-related problem that requires some interaction with files on the computer, either by creating new folders, decompressing and compressing files, listing and removing files from the hard drive of the computer or any other type of operation. R has full support for such type of operations. You can automate any type of computer task within an R script, if so needed.

In this section we will give preference to functions from package {fs}, which provides several routines for interacting with files and folders. Be aware, however, that the base package also offers similar functions.

2.20.1 Listing Files and Folders

To list files from your computer, use function dir_ls(), where the path argument sets the directory to list the files from. For the compilation of the book, I've created a directory called data. This folder contains all the data needed to recreate the book's examples. You can check the files in the sub-folder data with the following code:

```
# list files in data folder
my_files <- fs::dir_ls(path = "data")
print(my_files)
R> data/FileWithLatinChar_ISO-8859-9.txt
```

- R> data/FileWithLatinChar_UTF-8.txt
- R> data/Financial Sample.xlsx
- R> data/MySQLiteDatabase.SQLITE
- R> data/SP500-Stocks-WithRet.rds
- R> data/SP500-Stocks_long.csv
- R> data/SP500-Stocks_wide.csv
- R> data/SP500_long_yearly_2010-01-01_2019-11-04.rds
- R> data/SQLite_db.SQLITE
- R> data/UCI_Credit_Card.csv
- R> data/price-data.csv
- R> data/pride_and_prejudice.txt
- R> data/temp.txt
- R> data/temp_file.xlsx
- R> data/top25babynames-by-sex-2005-2017.csv

There are several files with different extensions in this directory. These files contain data that will be used in future chapters.

You can also list the files recursively over inner folders, that is, list all files from all sub-folders contained in the original path. To check it, try using the following code in your computer:

```
# list all files for all subfolders (IT MAY TAKE SOME TIME...)
fs::dir_ls(path = getwd(), recurse = TRUE)
```

The previous command will list all files in the current folder and sub-folders. Depending on the current working directory, it may take some time to run it all (you can cancel it anytime by pressing esc in your keyboard).

You can also set what type of output you need. For example, if you want only the available folders, and not files, use input type = "directory":

```
# store names of directories
my_dirs <- fs::dir_ls(
    path = getwd(),
    type = "directory")

# print it
print(my_dirs)</pre>
```

- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/_book
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/blocks
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/data
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/ebook files
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/eqs
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/mem_cache
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/quandl_cache
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/resources
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/tabs
- R> /tmp/Rtmp9FKiuz/compile-book-site--12c3e0c1c8b/text-to-reuse

The output shows the directories used to write the book. It includes the directory with the data ("./data"), resources ("./resources") among others. In this same directory, you can find the chapters of the book, organized by files and based on the *RMarkdown* language (.Rmd file extension):

```
# list all files with the extension .Rmd
fs::dir_ls(glob = "*.Rmd")
```

- R> 00a-About-new-edition.Rmd
- R> 00b-Preface.Rmd
- R> 01-Introduction.Rmd
- R> 02-Basic-operations.Rmd

```
R> 03-Research-scripts.Rmd
R> 04-Importing-exporting-local.Rmd
R> 05-Importing-internet.Rmd
R> 06-Data-structure-objects--ONLINE.Rmd
R> 07-Basic-objects--ONLINE.Rmd
R> 08-Programming--ONLINE.Rmd
R> 09-Cleaning-data--ONLINE.Rmd
R> 10-Figures--ONLINE.Rmd
R> 11-Models--ONLINE.Rmd
R> 12-Reporting-results--ONLINE.Rmd
R> 13-Optimizing-code--ONLINE.Rmd
R> 14-References.Rmd
R> _Welcome.Rmd
R> afedR_ed03_ONLINE.Rmd
R> index.Rmd
```

The files presented above contain all the contents of this book, including this specific paragraph, located in file 02-Basic-operations.Rmd!

2.20.2 Deleting Files and Directories

You can also use an R session to delete files and directories from your computer. This might come in handy when dealing with disposable data files. Use these commands with responsibility. If not careful, you can easily break the operating system of your computer.

You can delete files with command file_delete():

To delete directories and all their elements, we use dir_delete():

```
# create temp dir
fs::dir_create('temp')

# create a file inside of temp
my_file <- 'temp/tempfile.csv'
write.csv(x = data.frame(x=1:10),</pre>
```

```
file = my_file)
fs::dir_delete('temp')
```

If needed, we can check if the deletion of a directory was successful with command dir_exists():

```
fs::dir_exists('temp')

R> temp
R> FALSE
```

As expected, the directory was not found.

2.20.3 Downloading Files from the Internet

We can also use R to download files from the Internet with function **download.file()**. See the following example, where we download an Excel spread-sheet from Microsoft's website:

Using download.file() is quite handy when you are working with Internet data that is constantly being updated. In the script, we can download new data in every execution, making sure that our analysis is based on current information.

One trick worth knowing is that you can also download files from cloud services such as Dropbox¹¹ and Google Drive¹². So, if you need to send a data file to a large group of people and update it frequently, just pass the file link from the cloud service. This way, any local change in the data file in your computer will be reflected for all users with the file link.

¹¹https://www.dropbox.com/

¹²https://drive.google.com/



Needless to say, **be very careful** with commands fs::file_delete and fs::dir_delete, especially when using recursion (recurse = TRUE). One simple mistake and important parts of your hard drive can be erased, breaking your operating system. Be aware that R **permanently deletes** files and folder, without any simple way to restore them.

2.20.4 Using Temporary Files and Directories

Every time a user starts an R session, a new temporary folder is created in your hard drive. This folder is used to store any disposable files and folders generated by R. The location of this directory is available with **path_temp()**:

```
windows_tempdir <- fs::path_temp()
print(windows_tempdir)</pre>
```

R> C:\Users\msperlin\AppData\Local\Temp\RtmpEGt3Q7

The name of the temporary directory, in this case 'RtmpEGt3Q7', is unique and randomly defined at the start of every new R session. This means that every R session is linked to an unique temporary folder. If two R sessions are created, as in starting two RStudio sessions, two temporary folders will be available, each with an unique name. When the computer is rebooted, all temporary directories are deleted.

The same dynamic is found for file names. If you want to use a temporary random name for some reason, use file_temp():

```
windows_tempfile <- fs::file_temp()
message(windows_tempfile)</pre>
```

R> C:\Users\msperlin\AppData\Local\Temp\RtmpEGt3Q7\file19917e5fbc7e

You can also set its extension and name:

```
windows_tempfile <- fs::file_temp(
  pattern = 'temp_',
  ext = '.csv'
)
message(windows_tempfile)</pre>
```

 $\label{local-Temp-RtmpEGt3Q7-temp_19913e1dafe5.csv} R> C:\Users\mbox{\sc Nsperlin-AppData-Local-Temp-RtmpEGt3Q7-temp_19913e1dafe5.csv} R> C:\Users\mbox{\sc Nsperlin-AppData-Local-Temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp_19913e1dafe5.csv} R> C:\Users\mbox{\sc Nsperlin-AppData-Local-Temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-temp-RtmpEGt3Q7-$

As a practical case of using temporary files and folders, let's download the

Excel worksheet from Microsoft into a temporary folder and read its content for the first five rows:

```
# set link
link_dl <- 'go.microsoft.com/fwlink/?LinkID=521962'</pre>
local_file <- fs::file_temp(ext = '.xlsx')</pre>
download.file(url = link_dl,
              destfile = local_file)
df_msft <- readxl::read_excel(local_file)</pre>
print(head(df_msft))
R> # A tibble: 6 x 16
                Country Product Disco~1 Units~2 Manuf~3 Sale ~4
R>
     Segment
     <chr>
                 <chr>
R>
                         <chr>>
                                 <chr>
                                            <dbl>
                                                    <dbl>
                                                             <dbl>
R> 1 Government Canada Carret~ None
                                            1618.
                                                         3
                                                                20
R> 2 Government Germany Carret~ None
                                            1321
                                                         3
                                                                20
                                                         3
R> 3 Midmarket France Carret~ None
                                            2178
                                                                15
R> 4 Midmarket Germany Carret~ None
                                             888
                                                         3
                                                                15
                                                         3
R> 5 Midmarket Mexico Carret~ None
                                            2470
                                                                15
R> 6 Government Germany Carret~ None
                                                         3
                                                               350
                                            1513
R> # ... with 9 more variables: `Gross Sales` <dbl>,
       Discounts <dbl>, Sales <dbl>, COGS <dbl>, Profit <dbl>,
R> #
R> #
       Date <dttm>, `Month Number` <dbl>, `Month Name` <chr>,
       Year <chr>, and abbreviated variable names
R> #
       1: `Discount Band`, 2: `Units Sold`,
R> #
```

The example Excel file contains the sales report of a company. Do notice that the imported file becomes a dataframe in our R session, a table like an object with rows and columns.

3: `Manufacturing Price`, 4: `Sale Price`

By using **file_temp()**, we do not need to delete (or worry) about the downloaded file because it will be removed from the computer's hard disk when the system is rebooted.

2.21 Exercises

R> #

⁰¹ - In RStudio, create a new script and save it in a personal folder. Now, write R commands in the script that define two objects: one holding a

sequence between 1 and 100 and the other with the text of your name (ex. 'Richard'). Execute the whole script with the keyboard shortcuts.

02 - In the previously created *script*, use function message to display the following phrase in R's *prompt*:"My name is".

03 - Within the same script, show the current working directory (see function getwd, as in print(getwd())). Now, change your working directory to $Desktop\ (Desktop)$ and show the following message on the prompt screen: 'My desktop address is'. Tip: use and abuse of RStudio's autocomplete tool to quickly find the desktop folder.

04 - Use R to download the compressed zip file with the book material, available at this link¹³. Save it as a file in the temporary session folder (see function fs::file_temp()).

05 - Use the unzip function to unzip the downloaded file from previous question to a directory called 'afedR-files' inside the "Desktop" folder. How many files are available in the resulting folder? Tip: use the recursive = TRUE argument with fs::dir_ls to also search for all available subdirectories.

06 - Every time the user installs an R package, all package files are stored locally in a specific directory of the hard disk. Using command Sys.getenv('R_LIBS_USER') and fs::dir_ls, list all the directories in this folder. How many packages are available in this folder on your computer?

07 - In the same topic as previous exercise, list all files in all subfolders in the directory containing the files for the different packages (see command Sys.getenv('R_LIBS_USER')). On average, how many files are needed for each package?

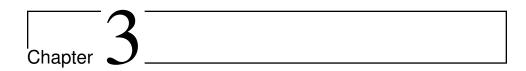
 $^{^{13} \}rm https://www.msperlin.com/files/afedr-files/afedR-code-and-data.zip$

08 - Use the install.packages function to install the yfR package on your computer. After installation, use function yf_get() to download price data for the IBM stock in the last 15 days. Tip: use function Sys.Date() to find out the current date and Sys.Date() - 15 to calculate the date located 15 days in the past.

09 - The cranlogs package allows access to downloads statistics of CRAN packages. After installing cranlogs on your computer, use the cranlogs::cran_top_downloads function to check which are the 10 most installed packages by the global community in the last month. Which package comes first? Tip: Set the cran_top_downloads function input to when = 'last-month'. Also, be aware that the answer here may not be the same as you got because it depends on the day the R code was executed.

10 - Using the devtools package, install the development version of the ggplot2 package, available in the Hadley Hickman repository. Load the package using library and create a simple figure with the code qplot(y = rnorm(10), x = 1:10).

11 - Using your programming ability check on your computer which folder, from the "Documents" directory (shortcut = \sim), has the largest number of files. Display the five folders with the largest number of files on R's prompt.



Writing Research Scripts

So far we learned how to use R for basic tasks such as interacting with the computer, creating simple vectors and downloading files from the internet. At this point, it is important to discuss the structure of a research script and, more specifically, how to organize our work in a efficient manner. As the R code base becomes larger and more complex, organization is a necessity. In this chapter, I will suggest a way to organize files and folders. So, I recommend that you follow these guidelines – or at least your own version of them – in every project you work on.

3.1 Stages of Research

Unlike other software designs, every script in data analysis follows through clear and consecutive steps to achieve its goal.

- 1. **Importation of data**: Raw (original) data is imported from a local file or the internet. At this stage, no manual data manipulation should happen. The raw data must be imported "as it is".
- 2. Cleaning and structuring the data: The raw data imported in the previous step is cleaned and structured within the needs of the analysis. Abnormal records and errors in observations can be removed or treated. The structure of the data can also be manipulated, binding (merging) different datasets and calculating variables of interest. Preferably, at the end of this stage, there should be a final collection of clean data.

- 3. Visual analysis and hypothesis testing: After cleaning and structuring the data, the work continues with the visual analysis of the data and hypothesis testing. Here, you can create graphical representations of the data for your audience and use statistical tools, such as econometric models, to test a particular hypothesis. This is the *heart* of the research and the stage most likely to take more development time.
- 4. **Reporting the results**: The final stage of a research script is reporting the results, that is, exporting selected tables and figures from R to a text processing software such as Latex, Writer (LibreOffice) or Word (Microsoft).

Each of the previous steps can be structured in a single R script or in several separate files. Using multiple files is preferable when the first steps of the research demand significant processing time. For example, when importing and organizing a large database, it is worth the trouble to separate the code in different files. It will be easier to find bugs and maintain the code. Each script will do one job, and do it well.

A practical example would be the analysis of a large dataset of financial transactions. Importing and cleaning the data takes plenty of computer time. A smart organization is to insert these primary data procedures in a R file and save the final objects of this stage in an external file. This local archive serves as a bridge to the next step, hypothesis testing, where the previously created file with clean data is imported. Every time a change is made to the hypothesis testing script, it is unnecessary to rebuild the whole dataset. This simple organization of files saves a lot of time. The underlying logic is simple, isolate the parts of the script that demand more computational time – and less development –, and connect them to the rest of the code using external data files.

If you are working with multiple files, one suggestion is to create a naming structure that informs the steps of the research in an intuitive way. An example would be to name the data importing code as O1-import-and-clean-data.R, the modeling code as O2-estimate-and-report-models.R and so on. The practical effect is that using a number in the first letter of the filenames clarifies the order of execution. We can also create a master script called O-run-it-all.R or O-main.R that runs (source) all other scripts. So, every time we make an update to the original data, we can simply run O-run-it-all.R and will have the new results, without the need to execute each script individually.

3.2 Folder Structure

A proper folder structure also benefits the reproducibility and organization of research. In simple scripts, with a small database and a low number of procedures, it is unnecessary to spend much time thinking about the organization of files. This is certainly the case for most of the code in this book. More complex programs, with several stages of data cleaning, hypothesis testing, and several sources of data, organizing the file structure is essential.

A suggestion for an effective folder structure is to create a single directory and, within it, create subdirectories for each input and output element. For example, you can create a subdirectory called data, where all the original data will be stored, a directory fig and tables, where figures and tables with final results will be exported. If you are using many custom-written functions in the scripts, you can also create a directory called r-fcts and save all files with function definitions at this location. As for the root of the directory, you should only find the main research scripts there. An example of a file structure that summarizes this idea is:

```
/Capital Markets and Inflation/
    /data/
        stock_indices.csv
        inflation_data.csv
/resources/figs/
        SP500_and_inflation.png
/tables/
        Table1_descriptive_table.tex
        Table2_model_results.tex
/r-fcts/
        fct_models.R
        fct_clean_data.R
0-run-it-all.R
1-import-and-clean-data.R
2-run-research.R
```

The research code should also be self-contained, with all files available within a sub-folder of the root directory. If you are using many different R packages, it is advisable to add a comment in the first lines of O-run-it-all.R that indicates which packages are necessary to run the code. The most friendly way to inform it is by adding a commented line that installs all required packages, as in #install.packages('pkg1', 'pkg2', ...). So, when someone receives the code for the first time, all he (or she) needs to do is uncomment the line and execute it. External dependencies and steps

for their installation should also be informed.

The organization of the code directory facilitates collaboration and reproducibility. If you need to share the code with other researchers, simply compress the directory to a single file and send it to the recipient. After decompressing the file, the structure of the folder immediately informs the user were to change the original data, the order of execution of the scripts in the root folder, and where the outputs are saved. The same benefit goes when you reuse your code in the future, say three years from now. By working smarter, you will be more productive, spending less time with repetitive and unnecessary steps for "figuring out" how the code works.

3.3 Important Aspects of a Research Script

In this section I'll be making some suggestions for how you can conduct data analysis with R. Making it clear, these are personal positions from my experience as a researcher and teacher. Many points raised here are specific to the academic environment but can be easily extended to the practice of data research in the industry. In short, these are suggestions I wish I knew when I first started my career.

Firstly, **know your data!**. I can't stress enough how this is important! The first instinct of every passionate data analyst when encountering a new set of rich information is to immediately import it into R and perform an analysis. However, a certain level of caution is needed. Every time you get your hands on a new set of data, ask yourself how much you **really** know:

- How was the data collected? To what purpose?
- What information does each column of the table represents? What are the details often missed?
- How do the available data compare with data used in other studies?
- Is there any possibility of bias within the data collection?

Furthermore, you need to remember that the ultimate goal of any research is communication. Thus, it is very likely that you will report your results to people who will have an informed opinion about the subject, including the sources and individualities of different datasets. The worst case scenario is when a research effort of three to six months in between coding and writing is nullified by a simple lapse in data checking. Unfortunately, this is not uncommon.

As an example, consider the case of analyzing the long term performance of companies in the retail business. For that, you gather a recent list of available companies and download financial records about their revenue, profit and adjusted stock price for the past twenty years. Well, the problem here is in the selection of the companies. By selecting those that are available today, you missed all companies that went bankrupt during the 20 year period. That is, by looking only at companies that stayed active during the whole period, you indirectly selected those that are profitable and presented good performance. This is a well-known effect called **survival bias**. The right way of doing this research is gathering a list of companies in the retail business twenty years ago and keep track of those that went bankrupt and those that stayed alive.

The message is clear. Be very cautious about the data you are using. Your raw tables stand at the base of the research. A small detail that goes unnoticed can invalidate your whole work. If you are lucky and the database is accompanied by a written manual, break it down to the last detail. If the information is not clear, do not be shy about sending questions to the responsible party. Likewise, if there is an inevitable operational bias in your dataset, be open and transparent about it.

The second point here is the code. After you finish reading this book, you will have the knowledge to conduct research with R. The computer will be a powerful ally in making your research ideas come true, no matter how gigantic they may be. However, a great power comes with great responsibility. Said that, you need to be aware that a single misplace line in a code can easily bias and invalidate your analysis.

Remember that analyzing data is your profession and **your reputation is your most valuable asset**. If you have low confidence in the produced code, do not publish or communicate your results. The code and its results is entirely your responsibility. Check it as many times as necessary. Always be skeptical about your own work:

- Do the descriptive statistics of the variables faithfully report the database?
- Is there any relationship between the variables that can be verified in the descriptive table?
- Do the main findings of the research make sense to the current literature of the subject? If not, how to explain it?
- Is it possible that a bug in the code has produced the results?

I'm constantly surprised by how many studies submitted to respected journals can be denied publication based on a simple analysis of the descriptive table. Basic errors in variable calculations can be easily spotted by a trained eye. The process of continuous evaluation of your research will not only make you stronger as a researcher but will also serve as practice for peer evaluation, much used in academic research. If you do not have enough confidence to report results, test your code extensively. If you have already done so and are still not confident, identify the lines of code you have doubts and seek help with a colleague or your advisor, if there is one. The latter is a strong ally who can help you in dealing with problems he/she already had.

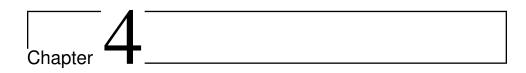
All of the research work is, to some extent, based on existing work. Today it is extremely difficult to carry out ground-breaking research. Knowledge is built in the form of blocks, one over the other. There is always a collection of literature that needs to be consulted. Therefore, you should always compare your results with the results already available in the subject. If the main results are not similar to those found in the literature, one should ask himself: could a code error have created this result?

I clarify that it is possible that the results of research differ from those of the literature, but the opposite is more likely. Knowledge of this demands care with your code. *Bugs* and code problems are quite common and can go unnoticed, especially in early versions of scripts. As a data analyst, it is important to recognize this risk and learn to manage it.

3.4 Exercises

01 - Imagine a survey regarding your household budget over time. Financial data is available in electronic spreadsheets separated by month, for 10 years. The objective of the research is to understand if it is possible to purchase a real state property in the next five years. Within this setup, detail in text the elements in each stage of the study, from importing the data to the construction of the report.

^{02 -} Based on the study proposed earlier, create a directory structure on your computer to accommodate the study. Create mock files for each subdirectory (see directory structure at section 3.2). Be aware you can create mock files and direction all in R (see functions cat and fs::dir_create).



Importing Data from Local Files

In this chapter we'll learn to import and export data available as local files in the computer. Although the task is not particularly difficult, a data analyst should understand the different characteristics of file formats and how to take advantage of it. While some data formats are best suited for sharing and collaboration, others can offer a significant boost in reading and writing speed.

Here we will draw a comprehensive list of data file formats in R, including:

- text data with comma-separated values (csv files);
- spreadsheet data from Excel (xlsx files);
- R native data files (*RData* and *rds* files);
- Lightning Fast Serialization of data frames (FST) format (fst files);
- SQLite;
- unstructured text data.

The previous packages and functions are sufficient for getting most of the work done. Nevertheless, it is worth mentioning that R can also import data from other softwares such as SPSS, Stata, Matlab, among many others. If that is your case, I suggest a thorough study of package **{foreign}}**.

4.1 The path of local files

The first lesson in importing data from local files is that the location of the file must be explicitly stated in the code. The path of the file is then passed to a function that will read the file. The best way to work with paths is to

use the *autocomplete* feature of RStudio (see section 2.19). An example of full path is:

```
my_file <- 'C:/My Research/data/price-data.csv'
```

Note the use of forwarding slashes (/) to designate the file directory. Relative references also work, as in:

```
my_file <- 'data/price-data.csv'
```

Here, it is assumed that, in the current working folder, there is a directory called data and, inside of it, exists a file called price-data.csv. If the file path is simply its name, such as in my_file <- 'price-data.csv', it is implicitly assumed that the file is located in the root of the working directory. From the previous chapter, recall that you can use setwd() to change the working folder to where the work is being done and simply create and use the "Projects" feature of RStudio, which automatically sets the working directory in the same location as the .Rproj file.



I again reinforce the use of tab and the autocomplete tool of RStudio. It is much easier and practical to find files on your computer's hard disk using tab navigation than to copy and paste the address from your file explorer. To use it, open double or quotes in RStudio, place the mouse cursor in between the quotes and press tab.

Going further, another very important point here is that the data from the file will be imported and exported as an object of type dataframe. That is, a table contained in an Excel or .csv file will become a dataframe object in R. When we export data, the most common format is this same type of object. Conveniently, dataframes are nothing more than tables, with rows and columns.

Each column in the dataframe will have its own class, the most common being numeric (numeric), text (character), factor (factor) and date (Date). When importing the data, it is imperative that each column is represented in the correct class. A vast amount of errors can be avoided by simply checking the column classes in the dataframe resulting from the import process. For now, we only need to understand this basic property of dataframes. We will study the details of this object in chapter 6.

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4.2 csv files

Consider a data file called CH04_SP500.csv, available from the book package. It contains daily closing prices of the SP500 index from 2010-01-04 until 2022-12-27. We will now use package {afedR3} for finding the file and copying it to your local folder. If you followed the instructions in the book preface chapter, you should have package {afedR3} already installed.

Once you install package {afedR3}, file CH04_SP500.csv and all other data files used in the book are downloaded from Github. The package also includes functions for facilitating the reproduction of code examples. Command data_path() will return the local path of a data file by looking up its name.

Let's copy CH04_SP500.csv to your "My Documents" folder with the following code using the tilde (~) shortcut:

```
my_f <- afedR3::data_path('CH04_SP500.csv')
fs::file_copy(my_f, '~')</pre>
```

Now, if it is your first time working with .csv files, use a file browser ("File Explorer" in Windows) and open CH04_SP500.csv in the "My Documents" folder with any text editor software such as Notepad. The first lines of CH04_SP500.csv, also called header lines, show the column names. Following international standards, rows are set using line breaks, and all columns are separated by commas (,). Next we show the textual content of the first ten lines of CH04_SP500.csv:

```
R> ref_date,price_close
R> 2010-01-04,1132.98999
R> 2010-01-05,1136.52002
R> 2010-01-06,1137.140015
R> 2010-01-07,1141.689941
R> 2010-01-08,1144.97998
R> 2010-01-11,1146.97998
R> 2010-01-12,1136.219971
R> 2010-01-13,1145.680054
R> 2010-01-14,1148.459961
```

The data in CH04_SP500.csv is organized in a standard way, and we should have no problem importing it in R. However, you should be aware this is not always the case. So, if you want to avoid the most common issues when importing data from csv files, I suggest you follow these steps:

1) Check the existence of text before the actual data. A standard .csv

file will only have the contents of a table but, sometimes, you will find a header text with metadata (extra information about the dataset). If necessary, you can control how many lines you skip in the *csv* reading function;

- 2) Verify the existence of names for all columns and if those names are readable;
- 3) Check the symbol for column separation. Normally it is a comma, but you never know for sure;
- 4) For the numerical data, verify the decimal symbol. R will expect it to be a dot. If necessary, you can adjust this information in the code, making sure R knows how to correctly parse numerical data.
- 5) Check the encoding of the text file. Normally it is one of UTF-8, Latin1 (ISO-8859) or Windows 1252. These are broad encoding formats and should suffice for most cases. Whenever you find strange symbols in the text columns of the resulting dataframe, the problem is probably due to a mismatch between the encoding of the file and R. Windows users can check the encoding of a text file by opening it in Notepad++1. The encoding format is available in the bottom right corner of the Notepad++ editor. Linux and Mac users can find the same information in any advanced text editor software such as Kate².



Whenever you find an unexpected text structure in a .csv file, use the arguments of the csv reading function to import the information correctly. As a rule of thumb, **never modify raw data manually**. Its far more efficient to use R code to deal with different structures of .csv files. It takes a bit of work, but such a policy will save you a lot of time in the future as, in a couple of months, you are unlikely to remember how you manually cleaned that csv file in order to import it more easily in your R session.

4.2.1 Importing Data

The {base} package includes a native function called **read.csv()** for importing data from .csv files. However, we will prefer the {**tidyverse**} alternative, **read_csv()**, as it is more efficient and easier to work with. In short, the

¹https://notepad-plus-plus.org/

²https://kate-editor.org/

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benefit of using **read_csv()** is that it reads the data very quickly, with clever internal rules for defining the classes of imported columns.

This is the first package from the {tidyverse} that we will use. Before doing so, it is necessary to install it in your R session. A simple way of installing all {tidyverse} packages as a bundle is as follows:

```
install.packages('tidyverse')
```

After running the previous code, all **{tidyverse}** packages will be installed on your computer. Once it finishes, let's load it.

```
# load library
library(tidyverse)
```

Back to importing data from .csv files, to load the contents of file CH04_SP500.csv in R, use the **read_csv()** function.

```
# set file to read
my_f <- afedR3::data_path('CH04_SP500.csv')

# read file
my_df_sp500 <- readr::read_csv(my_f)

# print it
print(head(my_df_sp500))</pre>
```

```
R> # A tibble: 6 x 2
R>
     ref_date price_close
R>
     <date>
                       <dbl>
R> 1 2010-01-04
                       1133.
R> 2 2010-01-05
                       1137.
R> 3 2010-01-06
                       1137.
R> 4 2010-01-07
                       1142.
R> 5 2010-01-08
                       1145.
R> 6 2010-01-11
                       1147.
```

As previously mentioned, the contents of the imported file becomes a dataframe object in R, and each column of a dataframe has a class. We can check the classes of object my_df_sp500 using function glimpse():

```
# Check the content of dataframe
dplyr::glimpse(my_df_sp500)
```

R> Rows: 3,269
R> Columns: 2

Note that the column of dates - ref_date - was imported as a Date vector and the closing prices - price_close - as numeric (dbl, double accuracy). This is exactly what we expected. Internally, function read_csv() identifies columns classes according to their content.

When calling read_csv() without extra arguments, the function will present in the prompt the classes of each column. Internally, the function sets the attributes of the columns by reading the first 1000 lines of the file. Column ref_date was imported with the *Date* class and column price_close was imported as *double* (*dbl*). We can use this information in our own code by copying the text and assigning it to a variable. Have a look:

```
# set cols from import message
my_cols <- readr::cols(
  ref_date = readr::col_date(),
  price_close = readr::col_double()
  )

# read file with readr::read_csv
my_df_sp500 <- readr::read_csv(my_f, col_types = my_cols)</pre>
```

This time, not message was set to the prompt. As an exercise, Let's import the same data, but use a character class for both columns:

```
# set cols from import message
# set cols from import message
my_cols <- readr::cols(
    ref_date = readr::col_character(),
    price_close = readr::col_character()
)

# read file with readr::read_csv
my_df_sp500 <- readr::read_csv(my_f, col_types = my_cols)
# glimpse the dataframe
dplyr::glimpse(my_df_sp500)</pre>
```

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As expected, both columns are now of class character.

Going further, read_csv() has several other input options such as:

- change the format of the import data, including symbols for decimal places and encoding (locale option);
- change column names (argument col_names);
- skip *n* lines before importation (skip option);
- custom definition for NA values (na option)

As an example, let's study the case of a messy .csv file. In the book package we have a file called funky csv file.csv where:

- the header has textual information;
- the file will use the comma as a decimal;
- the file text will contain Latin characters.

The first 10 lines of the files contain the following content:

```
R> Example of funky file:
R> - columns separated by ";"
R> - decimal points as ","
R>
R> Data build in 2022-12-28
R> Origin: www.funkysite.com.br
R>
R> ID;Race;Age;Sex;Hour;IQ;Height;Died
R> 001;White;80;Male;00:00:00;92;68;FALSE
R> 002;Hispanic;25;Female;00:00:00;99;68;TRUE
```

We have a header text up to line number 7 and the columns being separated by a semicolon (";"). When importing the data with standard (and wrong) options, we will have the following output:

```
my_f <- afedR3::data_path('CH04_funky-csv-file.csv')
df_funky <- readr::read_csv(my_f)</pre>
```

The output shows that only one column of class *character* was read from the file. This happens because function **read_csv()** expects actual data (and not text) in the first line of the file. To solve it, we need to use several input arguments to handle the particularities of the file:

Note that the data has now been correctly imported, with the correct column classes. For that, we use the alternative function **read_delim()** with custom inputs. Package **{readr}** also provides several other functions for specific import situations.

4.2.2 Exporting Data

To write a .csv file, use the **write_csv()** function. First, we create a new dataframe with some random data:

```
R> y z
R> 1 0.29496694 a
```

```
R> 2 0.67641989 a
R> 3 0.69974341 a
R> 4 0.69661747 a
R> 5 0.93058643 a
R> 6 0.08149662 a
```

And now we use $write_csv()$ to save it in a new (and temporary) .csv file:

In the previous example, we save the object my_df into a temporary file with path file12c322795567.csv. We can read it back and check its contents using read_csv() once again:

```
# read it
my_df_imported <- readr::read_csv(f_out)

# print first five rows
print(head(my_df_imported))</pre>
```

```
R> # A tibble: 6 x 2
R> y z
R> <dbl> <chr>
R> 1 0.295 a
R> 2 0.676 a
R> 3 0.700 a
R> 4 0.697 a
R> 5 0.931 a
R> 6 0.0815 a
```

As we can see, the data imported from the file is identical to the one created in the other code chunk.

4.3 Excel Files (xlsx)

Although it is not an efficient or portable data storage format, Microsoft Excel is a popular software in Finance and Economics due to its spreadsheet-like capacities. It is not uncommon for data to be stored and distributed in this format.

The downside of using Excel files for storing data is its low portability and the longer reading and writing times. This may not be a problem for small tables, but when handling a large volume of data, using Excel files can be very frustrating. If you can, my advice is to avoid the use of Excel files in your work cycle.

4.3.1 Importing Data

R does not have a native function for importing Excel files. Therefore, we must install and use packages to perform this operation. There are several good options including $\{XLConnect\}$, $\{xlsx\}$, $\{readxl\}$ and $\{tidyxl\}$

Despite their similar goals, each package has its peculiarities. If reading Excel files is important to your work, I strongly advise the study of each package. For example, package {tidyxl} was specially designed to read unstructured Excel files, where the desired information is not contained in a tabular format. Alternatively, package {XLConnect} allows the user to open a live connection and, from R, control an Excel file, making it possible to export and send data, format cells, and so on.

In this section, we will give priority to package {readxl}, one of the most straightforward packages to interact with Excel files. Conveniently, it does not require the installation of external software.

Let's start with an example. Consider a file called CH04_SP500-Excel.xlsx that contains the same SP500 data from previous section. We can import the information from the file using function read_excel():

```
# set excel file
my_f <- afedR3::data_path('CH04_SP500-Excel.xlsx')

# read excel file
my_df <- readxl::read_excel(my_f, sheet = 'Sheet1')

# print with head (first five rows)
dplyr::glimpse(my_df)</pre>
```

One of the benefits of using Excel files is that the column's classes are directly inherited from the file. If the column classes are correct in the Excel file, then

they will automatically be correct in R. In our case, the date column of file CH04_SP500-Excel.xlsx was correctly set as a dttm object, a special type of DateTime class. Likewise, even if the Excel file used commas for decimals, the import process would still succeed as the conversion is handled internally.

4.3.2 Exporting Data

Exporting a dataframe to an Excel file is also easy. Again, no native function in R performs this procedure. We can, however, use package {writexl}:

4.4 RData and rds Files

R offers two native formats to write objects to a local file, *RData*, and *rds*. The benefit of using both is that the saved file is compact and its access is very fast. The downside is the low portability, i.e., it's difficult to use the files in other software.

The difference between *RData* and *rds* is that the first can save many R objects in a single file, while the latter only one. This, however, is not a hard restriction for the *rds* format as we can incorporate several objects into a single one using lists. In practice, a *rds* file can store as many objects as needed. We will learn more about lists in chapter 6.

4.4.1 Importing Data

To create a new .RData file, use the save() function. See the following example, where we create a .RData file with some content, clear R's memory, and then load the previously created file:

```
# set a object
my_x <- 1:100</pre>
```

```
# set name of RData file
my_file_rdata <- fs::file_temp(ext = 'RData')
my_file_rds <- fs::file_temp(ext = 'rds')

# save it in RData
save(list = c('my_x'), file = my_file)

# save it in rds
readr::write_rds(my_x, my_file_rds)</pre>
```

We can verify the existence of the file with the file_exists() function:

```
# check if file exists
fs::file_exists(my_file_rdata)
```

As expected, file price-data.csv is available.

Now, to import data from .rds files, we use function read rds():

```
# load content into workspace
my_x_2 <- readr::read_rds(file = my_file_rds)</pre>
```

Comparing the code between using .RData and .rds files, note that the .rds format allows the explicit definition of the output object. The contents of my_file_rds is saved in my_x_2. When we use the load() function for RData files, we cannot name the output directly. This is particularly inconvenient when you need to modify the name of the imported object.



As a suggestion, use the .rds format to write and read R data. Its use is more practical, resulting in cleaner code. The difference in speed between one and the other is minimal. The benefit of importing multiple objects into the same RData file becomes irrelevant when using list objects, which can incorporate other objects into its content.

4.4.2 Exporting Data

We can create a new RData file with command save():

```
# set vars
my_x <- 1:100
```

We can again check if the file exists:

```
fs::file_exists(my_file)

R> /tmp/Rtmp9FKiuz/file12c36c1f086a.RData
R> TRUE
```

The result is TRUE as expected.

As for .rds files, we save it with function **write_rds()**:

R> [1] TRUE

Command identical() tests if both objects are equal. Again, as expected, we find the result to be TRUE.

4.5 fst files

The fst format³, R package {fst}, is specially designed to enable quick writing and reading time from tabular data, with minimal disk space. Using this format is particularly beneficial when working with large databases in powerful computers. The trick here is the use of all computer cores to import

³http://www.fstpackage.org/

and export data, while all other formats only use one. If you have a computer with several cores, the gain in speed is impressive, as we will soon learn.

4.5.1 Importing Data

Using fst file format is similar to the previous cases. We use function $read_fst()$ to read files:

```
# set file location
my_file <- afedR3::data_path('CHO4_example-fst.fst')</pre>
# read fst file
my_df <- fst::read_fst(my_file)</pre>
# check contents
dplyr::glimpse(my_df)
R> Rows: 100
R> Columns: 8
           <chr> "001", "002", "003", "004", "005", "006", "~
R> $ ID
R> $ Race <fct> Black, White, Hispanic, Black, White, White~
R> $ Age <int> 33, 35, 23, 87, 65, 51, 58, 67, 22, 52, 52,~
R> $ Sex
           <fct> Male, Female, Male, Female, Male, Fem~
R> $ Hour <dbl> 0.00000000, 0.00000000, 0.00000000, 0.000000~
R> $ IQ
            <dbl> 108, 108, 85, 106, 92, 92, 88, 100, 86, 80,~
R> $ Height <dbl> 72, 63, 77, 72, 71, 74, 64, 69, 63, 72, 70,~
R> $ Died
            <lgl> FALSE, TRUE, TRUE, FALSE, FALSE, TRUE, TRUE~
```

As with the other cases, the data from file CH04_example-fst.fst is available in the workspace.

4.5.2 Exporting Data

We use function write_fst() to save dataframes in the fst format:

```
# create dataframe
N <- 1000
my_file <- fs::file_temp(ext = 'fst')
my_df <- data.frame(x = runif(N))

# write to fst
fst::write_fst(x = my_df, path = my_file)</pre>
```

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4.5.3 Timing the *fst* format

As a test of the potential of the fst format, we will now time the read and write time between fst and rds for a large table: 5.000.000 rows and 2 columns. We will also report the size of the resulting file.

```
# set number of rows
N <- 500000
# create random dfs
my_df <- data.frame(y = seq(1,N),</pre>
                     z = rep('a',N))
# set files
my_file_1 <- fs::file_temp(ext = 'rds')</pre>
my_file_2 <- fs::file_temp(ext = 'fst')</pre>
# test write
time_write_rds <- system.time(readr::write_rds(my_df, my_file_1 ))</pre>
time_write_fst <- system.time(fst::write_fst(my_df, my_file_2 ))</pre>
# test read
time_read_rds <- system.time(readr::read_rds(my_file_1))</pre>
time_read_fst <- system.time(fst::read_fst(my_file_2))</pre>
# test file size (MB)
file_size_rds <- file.size(my_file_1)/1000000</pre>
file_size_fst <- file.size(my_file_2)/1000000
```

And now we check the results:

The difference in performance is impressive! The fst not only reads and writes faster but also results in smaller file sizes. Be aware, however, this result is found in a sixteen core computer in which the book was compiled. You may not be able to replicate the same result in a more modest machine.

R> Resulting FILE SIZE for .rds: 65.000177 MBs R> Resulting FILE SIZE for .fst: 14.791938 MBs



Due to the use of all the computer's cores, the *fst* format is highly recommended when working with large data on a powerful computer. Not only will the resulting files be smaller, but the writing and reading process will be considerably faster.

4.6 SQLite Files

The use of .csv or .rds files for storing objects has its limits as the volume of data increases. If you are waiting a long time to read a dataframe from a file or if you are only interested in a small portion of a large table, you should look for alternatives. Likewise, if you have access to the computer network in your institution, including databases, it is important to learn how to directly import the corporate data into your R session.

This brings us to the topic of **database software**. These specific programs usually work with a query language, called SQL (Structured Query Language). It allows the user to read portions of the data and even manipulate

it efficiently. Many options of database software integrate nicely with R. The list includes MySQL, SQLite and MariaDB. Here, we will provide a quick tutorial on this topic using SQLite, which is the easiest one to work with. Unlike other database software, SQLite stores data and configurations from a single local file, without the need of a formal server.

4.6.1 Importing Data

Before moving to the examples, we need to understand how to use database software. First, R will connect to the database and return a connection object. Based on this connection, we will send queries for manipulating data using the SQL language. The main advantage is we can have a large database of, let's say, 10 GB and only load a small portion of it in R. This operation is also very quick, allowing efficient access to the available tables.

Assuming the existence of an SQLite file in the computer, we can import its tables with package {RSQLite}:

It worked. The dataframe from table MyTable1 is exactly as expected.

Another example of using SQLite is with the actual SQL statements. Notice, in the previous code, we used function **dbReadTable()** to get the contents of all rows in table MyTable1. Now, let's use an SQL command to get from MyTable2 only the rows where the G column is equal to A.

It also worked, as expected.

In this simple example, we can see how easy it is to create a connection to a database, retrieve tables, and disconnect. If you have to work with large tables, which, in my opinion, is any database that occupies more than 4 GB of your computer memory, it is worth moving it to proper database software. You'll be able to retrieve data faster, without the need of loading the whole database in the computer's memory. If you have a server available in your workplace, I strongly advise learning how to connect to it and use the SQL language to your advantage.

4.6.2 Exporting Data

As an example of exporting data to an SQLite file, let's first create an SQLite database. For that, we will set two large dataframes with random data and save both in an SQLite file.

The TRUE output of **dbWriteTable()** indicates everything went well. A connection was opened using function **dbConnect()**, and both **dataframes** were written to an SQLite file, called file12c31d348fd0.SQLITE.

4.7 Unstructured Data and Other Formats

Another example of data importation is the case of reading and processing unstructured text files. If none of the previous packages can read the data, then it must be parsed line by line. Let's explore this problem.

4.7.1 Importing Data

You can read the contents of a text file with function read_lines():

```
# set file to read
my_f <- afedR3::data_path('CH04_price-and-prejudice.txt')
# read file line by line
my_txt <- readr::read_lines(my_f)</pre>
```

```
[1] "
R>
                                         [Illustration:"
    [2] ""
R>
    [3] "
R>
                                         GEORGE ALLEN"
    [4] "
                                            PUBLISHER"
R>
    [5] ""
R>
    [6] "
                                    156 CHARING CROSS ROAD"
R>
    [7] "
R>
                                             LONDON"
    [8] ""
R>
    [9]
R>
                                         RUSKIN HOUSE"
R> [10] "
                                                7"
R> [11] ""
R> [12] "
                                         [Illustration:"
R> [13] ""
                                                            _Cha"
R> [14] "
                          _Reading Jane's Letters._
R> [15] "
                                                ן יי
```

In this example, file CH04_price-and-prejudice.txt contains the whole content of the book *Pride and Prejudice* by Jane Austen, freely available in the Gutenberg⁴ project, and downloaded with package {gutenbergr}. We imported the entire content of the file as a character vector named my_txt. Each element of my_txt is a line from the raw text file. Based on it, we can check the number of lines in the book, and also the number of times that the name 'Bennet', one of the protagonists, appears in the text:

⁴http://www.gutenberg.org/

R> The number of lines found in the file is 14529. R> The word "Bennet" appears 696 in the book.

In the example, we used function **sapply()**. In this case, it allowed us to use a function for each element of my_txt. We searched and counted the number of times the word "Bennet" was found.

4.7.2 Exporting Data

A typical case of exporting unstructured text is saving the log record of a procedure. This is quite simple. Using function **write_lines()**, use the input **file** to set the name of the local file and **x** for the actual textual content.

In the previous example, we created a simple text object and saved it in data/temp.txt. We can check the result with the **read_lines()** function:

```
print(readr::read_lines(my_f))
R> [1] "Today is 2023-03-10" "Tomorrow is 2023-03-11"
As we can see, it worked as expected.
```

4.8 How to Select a Data File Format

The choice of file format is an important topic and might actually be a time-saver at your work. In that decision, my first and most important suggestion is **to avoid the Excel format at all costs**. As for alternatives, we must consider three points in the decision:

- **speed** of reading and writing operations;
- **size** of the resulting file;
- compatibility with other software and operating systems.

Usually, the use of csv files easily satisfies these requirements. A csv file is nothing more than a text file that can be opened, viewed, and imported into any other statistical software. This makes it easy to share it with other people. Also, the size of csv files is usually not restrictive and, if needed, the file can be compressed using the zip() function. For these reasons, the use of csv files for importing and exporting data is preferable in the vast majority of situations.

However, there are cases where the speed of import and export operations matter. If you don't mind giving up portability, the *rds* format is a great choice for most projects. If you have good hardware and execution speed with *rds* is still not great, then the best alternative is the *fst* format, which uses all cores to import and export data.

4.9 Exercises

Export the resulting dataframe to each of the five formats: csv, rds, xlsx, fst. Which of the formats took up the most space in the computer's memory? Tip: file.size calculates the size of files within R.

02 - Improve the previous code by measuring the execution time for saving the data in different formats. Which file format resulted in the fastest execution for exporting data? Tip: use the system.time function or thetictoc package to calculate the execution times.

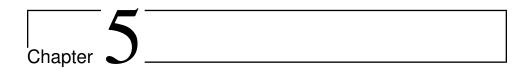
03 - For the previous code, reset the value of my_N to 1000000. Does it change the answers to the last two questions?

04 - Use afedR3::data_path function to access the CH04_SP500.csv file in the book's data repository. Import the contents of the file into R with the function readr::read_csv. How many lines are there in the resulting dataframe?

05 - At link https://eeecon.uibk.ac.at/~zeileis/grunfeld/Grunfeld.csv/ you'll find a .csv file for the Grunfeld data. This is a particularly famous table due to its use as reference data in econometric models. Using readr::read_csv function, read this file using the direct link as input read_csv. How many columns do you find in the resulting dataframe?

06 - Use function afedR3::data_path function to access the CH04_example-tsv.tsv file in the book's data repository. Note that the columns of the data are separated by the tab symbol ('\t'), and not the usual comma. After reading the readr::read_delim manual, import the information from this file to your R session. How many rows does the resulting dataframe contain?

07 - In the book package you'll find data file called CH04_another-funky-csv-file.csv, with a particularly bizarre format. Open it in a text editor and try to understand how the columns are separated and what is symbol for decimals. After that, study the inputs of function utils::read.table and import the table into your R session. If we add the number of rows to the number of columns in the imported table, what is the result?



Importing Data from the Internet

One of the great advantages of using R in Finance and Economics is the large amount of data that can be imported using the internet, substituting the tedious, unreliable and soul-crushing work of manual data collection. It also becomes easier to share reproducible code, as anyone can feasibly download the same tables with a single line of code.

In this chapter, we will study the most important and reliable packages for data importation in the fields of Finance and Economics. It is a small, but comprehensive list of packages that cover a large range of research topics. The list includes:

- **{GetQuandlData}** Imports economical and financial data from the Quandl platform.
- **{yfR}** Imports adjusted and unadjusted stock price data from Yahoo Finance.
- **{simfinapi}** Imports financial statements and adjusted stock prices from the SimFin project¹.
- **{tidyquant}** Imports several financial information about stock prices and fundamental data.

5.1 Package {GetQuandlData}

Quandl is an established and comprehensive platform that provides access to a series of free and paid data. Several central banks and research institutions

¹https://simfin.com/

provide free economic and financial information on this platform. I strongly recommend browsing the available tables from the Quandl website². It is likely that you'll find datasets that you're familiar with.

In R, package {Quandl} is the official extension offered by the company and available in CRAN. However, the package has some issues (see blog post here³), which are fixed with the alternative package {GetQuandlData}.

The first and mandatory step in using {GetQuandlData} is to register a user at the Quandl website⁴. Soon after, go to account settings and click $API \ KEY$. This page should show a code, such as Asv8Ac7zuZzJSCGxynfG. Copy it to the clipboard (control + c) and, in R, define a character object containing the copied content as follows:

```
# set FAKE api key to quandl
my_api_key <- 'Asv8Ac7zuZzJSCGxynfG'</pre>
```

The API key is unique to each user, and the one presented here will not work on your computer. You'll need to get your own API key to run the examples of the book. After finding and setting your key, go to Quandl's website and use the search box to look for the symbol of the time series of interest. As an example, we will use data for gold prices in the London Market, with a Quandl code equivalent to 'LBMA/GOLD'. Do notice that the structure of a Quandl code is always the same, with the name of the main database at first, and the name of table second, separated by a forward slash (/).

Now, with the API key and the Quandl symbol, we use function **get_Quandl_series()** to download the data from 1980-01-01 to 2023-01-01:

```
# set symbol and dates
my_symbol <- c('GOLD' = 'LBMA/GOLD')
first_date <- '1980-01-01'
last_date <- '2023-01-01'

# get data!
df_quandl <- GetQuandlData::get_Quandl_series(
   id_in = my_symbol,
        api_key = my_api_key,
        first_date = first_date,
        last_date = last_date,</pre>
```

²https://data.nasdaq.com/

³https://www.msperlin.com/post/2019-10-01-new-package-getquandldata/

⁴https://data.nasdag.com/

```
do_cache = FALSE)

# check it
dplyr::glimpse(df_quandl)
```

```
R> Rows: 10,866
R> Columns: 9
             <chr> "559", "632", "596", "634", "615.75", ~
R> $ `USD (AM)`
R> $ `USD (PM)`
             <chr> "559.5", "634", "588", "633.5", "610",~
             <chr> "251.123", "281.577", "266.285", "281.~
R> $ `GBP (AM)`
             <chr> "250.841", "282.468", "262.77", "280.0~
R> $ `GBP (PM)`
R> $ series_name <chr> "GOLD", "GOLD", "GOLD", "GOLD", "GOLD"~
R> $ ref_date
             <date> 1980-01-02, 1980-01-03, 1980-01-04, 1~
R> $ id_quandl
             <chr> "LBMA/GOLD", "LBMA/GOLD", "LBMA/GOLD",~
```

Notice how we set the name of the time series in line id_in = c('GOLD' = 'LBMA/GOLD'). The name of the element becomes the value of column series_name in df_quandl. If we had more time series, they would be stacked in the same table, but with different series name value.

There are other Quand1 API options available with inputs order, collapse and transform. If using Quand1 is important to your work, I strongly recommend reading the available parameters for querying data⁵. Several choices for data transformations can be passed to function get_Quandl_series()

As an inspection check, let's plot the prices of Gold in USD over time.

Overall, gold prices were fairly stable between 1980 and 2000, reaching a spike after 2010. One possible explanation is the higher demand for safer assets, such as gold, after the 2009 financial crisis. However, gold was never an efficient long term investment. To show that, let's calculate its compound annual return from 1980-01-02 to 2022-12-30:

```
library(dplyr)

# sort the rows

df_quandl <- df_quandl |>
  mutate(USD = as.numeric(`USD (AM)`)) |>
  arrange(ref_date)
```

⁵https://docs.data.nasdaq.com/docs/parameters-2

```
R> [1] 0.02771743
```

We find the result that Gold prices in USD compounded in a rate equal to 2.77% per year. This is not an impressive investment result by any means. As a comparison, the annual inflation for the US in the same period is 3.21%. In other words, the investor that bought gold in 1980 and held to it until 2023, perceived a lower nominal return than the inflation.

5.1.0.1 Fetching many time series

When asking for multiple time series from Quandl, package {GetQuandl-Data} stacks all the data in a single dataframe, making it easier to work with the {tidyverse} tools. As an example, let's look at Quandl database RATEINF, which contains a time series of inflation rates around the world. First, we need to see what are the available datasets:

```
# databse to get info
db_id <- 'RATEINF'</pre>
# get info
df_db <- GetQuandlData::get_database_info(db_id, my_api_key)</pre>
dplyr::glimpse(df_db)
R> Rows: 26
R> Columns: 8
R> $ code
                  <chr> "CPI_ARG", "CPI_AUS", "CPI_CAN", "CPI~
R> $ name
                  <chr> "Consumer Price Index - Argentina", "~
R> $ description <chr> "Please visit <a href=http://www.rate~
R> $ refreshed_at <dttm> 2020-10-10 02:03:32, 2023-03-04 02:0~
                  <date> 1988-01-31, 1948-09-30, 1989-01-31, ~
R> $ from_date
R> $ to_date
                  <date> 2013-12-31, 2022-12-31, 2023-01-31, ~
                  <chr> "RATEINF/CPI_ARG", "RATEINF/CPI_AUS",~
R> $ quandl_code
                  <chr> "RATEINF", "RATEINF", "RATEINF", "RAT~
R> $ quandl_db
```

Column name contains the description of tables. If we dig deeper, we'll find

the following names:

print(unique(df_db\$name))

```
[1] "Consumer Price Index - Argentina"
R>
R>
    [2] "Consumer Price Index - Australia"
    [3] "Consumer Price Index - Canada"
R>
   [4] "Consumer Price Index - Switzerland"
R>
R>
    [5] "Consumer Price Index - Germany"
   [6] "Consumer Price Index - Euro Area"
R>
   [7] "Consumer Price Index - France"
R.>
    [8] "Consumer Price Index - UK"
R>
R> [9] "Consumer Price Index - Italy"
R> [10] "Consumer Price Index - Japan"
R> [11] "Consumer Price Index - New Zealand"
R> [12] "Consumer Price Index - Russia"
R> [13] "Consumer Price Index - USA"
R> [14] "Inflation YOY - Argentina"
R> [15] "Inflation YOY - Australia"
R> [16] "Inflation YOY - Canada"
R> [17] "Inflation YOY - Switzerland"
R> [18] "Inflation YOY - Germany"
R> [19] "Inflation YOY - Euro Area"
R> [20] "Inflation YOY - France"
R> [21] "Inflation YOY - UK"
R> [22] "Inflation YOY - Italy"
R> [23] "Inflation YOY - Japan"
R> [24] "Inflation YOY - New Zealand"
R> [25] "Inflation YOY - Russia"
```

R> [26] "Inflation YOY - USA"

What we want is the 'Inflation YOY - *' datasets, which contain the year-on-year inflation rates for different countries. Let's filter the dataframe to keep the series with the yearly inflation, and select four countries:

Now we grab the data using get_Quandl_series:

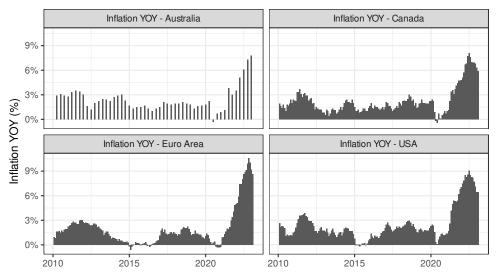
```
my_id <- df_db$quandl_code
names(my_id) <- df_db$name
first_date <- '2010-01-01'
last_date <- Sys.Date()

df_inflation <- GetQuandlData::get_Quandl_series(
   id_in = my_id,
   api_key = my_api_key,
   first_date = first_date,
   last_date = last_date)

dplyr::glimpse(df_inflation)</pre>
```

And, finally, we create an elegant plot to see the behavior of the inflation rates in the selected countries:

Inflation in the World 2010-01-31 to 2023-01-31



As you can see, the {GetQuandlData} output is formatted to work well with the {tidyverse} tools. The resulting data shows some correlation between the inflation rates, specially between the Euro Area and USA. We can also see the impact of covid in 2020, with large increases of inflation rates across the world.

5.2 Package {yfR}

Package {yfR} is all about downloading stock price data from Yahoo Finance. Unlike other packages, {yfR} focuses on large batch downloads of structured and clean/tidy data. Its main features are:

- Fetches daily/weekly/monthly/annual stock prices/returns from yahoo finance and outputs a dataframe (tibble) in the long format (stacked data);
- A feature called collections facilitates download of multiple tickers from a particular market/index. You can, for example, download data for all stocks in the SP500 index with a simple call to yf_collection_get("SP500");
- A session-persistent smart cache system is available by default. This
 means that the data is saved locally and only missing portions are
 downloaded, if needed.
- All dates are compared to a benchmark ticker such as SP500 and, whenever an individual asset does not have a sufficient number of dates, the software drops it from the output. This means you can choose to ignore tickers with a high proportion of missing dates.
- A customized function called **yf_convert_to_wide()** can transform a long **dataframe** into a wide format (tickers as columns), much used in portfolio optimization. The output is a list where each element is a different target variable (prices, returns, volumes).
- Parallel computing with package furrr is available, speeding up the data importation process.

As an example of usage, let's download the prices of four stocks in the previous five years using function **yf_get()**. We choose these companies: Microsoft (MSFT), Google (GOOGL), JP Morgan (JPM) and General Electric (GE).

In the call to function $yf_get()$, we set arguments thresh_bad_data = 0.95 and bench_ticker = '^GSPC'. These choices make sure that all returned data have at least 95% of valid prices when compared to data from the SP500 index (ticker '^GSPC').

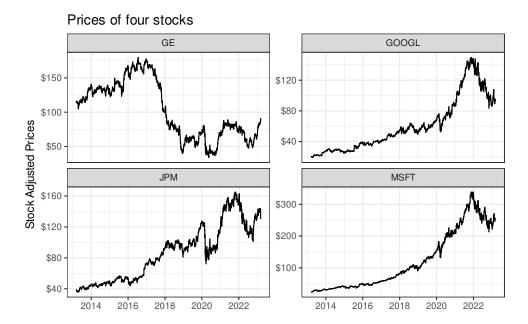
The output of **yf_get()** is an object of type **dataframe**, a table with prices and returns:

```
# print df.tickers
dplyr::glimpse(df_yf)
```

```
R> Rows: 10,068
R> Columns: 11
R> $ ticker
                            <chr> "GE", "GE", "GE", "GE", "GE~
R> $ ref_date
                            <date> 2013-03-12, 2013-03-13, 20~
R> $ price_open
                            <dbl> 141.6562, 140.7554, 141.416~
                            <dbl> 141.7162, 141.2358, 142.677~
R> $ price_high
R> $ price_low
                            <dbl> 140.0949, 140.5152, 141.175~
                            <dbl> 140.5753, 141.0557, 142.256~
R> $ price_close
R> $ volume
                            <dbl> 6093832, 4022099, 5766901, ~
R> $ price_adjusted
                            <dbl> 114.8972, 115.2898, 116.271~
R> $ ret_adjusted_prices
                            <dbl> NA, 0.0034171676, 0.0085141~
R> $ ret_closing_prices
                            <dbl> NA, 0.0034173293, 0.0085142~
R> $ cumret_adjusted_prices <dbl> 1.0000000, 1.0034172, 1.011~
```

As expected, we find information about stock prices, daily returns and traded volumes. Notice it also includes column ticker, which contains the symbols of the stocks. In the tidy format, each stock has a chunk of data that is pilled in top of each other. Later, in chapter 8, we will use this column to

split the data and build summary tables. To inspect the data, create a figure with the prices:



Data from Yahoo Finance

We see that General Eletric (GE) stock was not kind to its investors. Someone that bought the stock at its peak in mid-2016 has found its current value at less than half. Now, when it comes to the GOOGL, JPM and MSFT, we see an overall upward increase in stock prices, and a recent drop. These are profitable and competitive companies in their sectors and not surprisingly, the stock prices surged over the long period of time.

Now, let's look at an example of a large download of stock price data. Here, we will download the current composition of the SP500 index using function yf_collection_get().

```
# set dates
first_date <- '2020-01-01'
last_date <- '2023-01-01'

df_dow <- yfR::yf_collection_get(
   "DOW",
   first_date,
   last_date
)</pre>
```

And now we check the resulting data:

```
dplyr::glimpse(df_dow)
```

```
R> Rows: 22,680
R> Columns: 11
                            <chr> "AAPL", "AAPL", "AAPL", "AA~
R> $ ticker
R> $ ref_date
                            <date> 2020-01-02, 2020-01-03, 20~
                            <dbl> 74.0600, 74.2875, 73.4475, ~
R> $ price_open
R> $ price_high
                            <dbl> 75.1500, 75.1450, 74.9900, ~
                            <dbl> 73.7975, 74.1250, 73.1875, ~
R> $ price_low
R> $ price_close
                            <dbl> 75.0875, 74.3575, 74.9500, ~
R> $ volume
                            <dbl> 135480400, 146322800, 11838~
R> $ price_adjusted
                            <dbl> 73.44937, 72.73532, 73.3148~
R> $ ret_adjusted_prices
                            <dbl> NA, -0.009721663, 0.0079681~
                            <dbl> NA, -0.009722044, 0.0079682~
R> $ ret_closing_prices
R> $ cumret adjusted prices <dbl> 1.0000000, 0.9902783, 0.998~
```

We get a fairly sized table with 22680 rows and 11 columns in object df_dow. Notice how easy it was to get that large volume of data from Yahoo Finance with a simple call to yf_collection_get().



Be aware that Yahoo Finance (YF) data for adjusted prices of single stocks over long periods of time is not trustworthy. If you compare it to other data vendors, you'll easily find large differences. The issue is that Yahoo Finance does not adjust for dividends, only for stock splits. This means that, when looking at a price series over a long period of time, there is a downward bias in overall return. As a rule of thumb, in a formal research, never use individual stock data from Yahoo Finance, specially if the stock return is important to the research. The exception is for financial indexes, such as the SP500, where Yahoo Finance data is quite reliable since indexes do not undergo the same adjustments as individual stocks.

5.3 Package {simfinapi}

SimFin⁶ is a reliable repository of financial data from around the world. It works by gathering, cleaning and organizing data from different stock exchanges and financial reports. From its own website⁷:

⁶https://simfin.com/

⁷https://simfin.com/

Our core goal is to make financial data as freely available as possible because we believe that having the right tools for investing/research shouldn't be the privilege of those that can afford to spend thousands of dollars per year on data.

As of february 2023, the platform offers a generous free plan, with a daily limit of 2000 api calls. This is enough calls for most projects. If you need more calls, the premium version⁸ is a fraction of what other data vendors usually request.

Package {simfinapi} facilitates importing data from the SimFin API. First, it makes sure the requested data exists and only then calls the api. As usual, all api queries are saved locally using package memoise. This means that the second time you ask for a particular data about a company/year, the function will load a local copy, and will not call the web api, helping you stay below the API limits.

5.3.1 Example 01 - Apple Inc Annual Profit

The first step in using simfinR is registering at the SimFin website. Once done, click on Data Access⁹. It should now show an API key such as 'rluwSlN304NpyJeBjlxZPspfBBhfJR4o'. Save it in an R object for later use.

```
my_api_key <- 'rluwS1N3O4NpyJeBjlxZPspfBBhfJR4o'</pre>
```

Be aware that the **API key in my_api_key is fake** and will not work for you. You need to get your own to execute the examples.

With the API key in hand, the second step is to find the numerical id of the company of interest. For that, we can find all available companies and their respective ids and ticker with **sfa_get_entities()**.

```
cache_dir <- fs::path_temp("cache-simfin")
fs::dir_create(cache_dir)

simfinapi::sfa_set_api_key(my_api_key)
simfinapi::sfa_set_cache_dir(cache_dir)

# get info
df_info_companies <- simfinapi::sfa_get_entities()</pre>
```

⁸https://simfin.com/simfin-plus

⁹https://simfin.com/data/access/api

```
# check it
glimpse(df_info_companies)
R> Rows: 3,282
R> Columns: 2
R> $ simfin_id <int> 854465, 45846, 1253413, 1333027, 367153,~
               <chr> "1COV.DE", "A", "A18", "A21", "AA", "AAC~
R> $ ticker
Now, based on ticker "AAPL", lets download the download the profit and
loss (PL) statement for Apple INC in 2022:
ticker <- "AAPL" # ticker of APPLE INC
type_statements <- 'pl' # profit/loss statement
period <- 'fy' # final year</pre>
fiscal_year <- 2022
PL_aapl <- simfinapi::sfa_get_statement(
 ticker = ticker,
  statement = type_statements,
 period = period,
 fyear = fiscal_year)
# select columns
PL_aapl <- PL_aapl |>
  dplyr::select(ticker, simfin_id, revenue, net_income)
dplyr::glimpse(PL_aapl)
R> Rows: 1
R> Columns: 4
R> $ ticker
             <chr> "AAPL"
R> $ simfin_id <int> 111052
R> $ revenue <dbl> 3.94328e+11
```

Not bad! Financially speaking, the year 2022 was very good for Apple.

R> \$ net_income <dbl> 9.9803e+10

5.3.2 Example 02 - Annual Net Profit of Many Companies

Package simfinapi can also fetch information for many companies in a single call. Let's run another example by selecting three companies: Apple, Google and Amazon, and downloading end of year information:

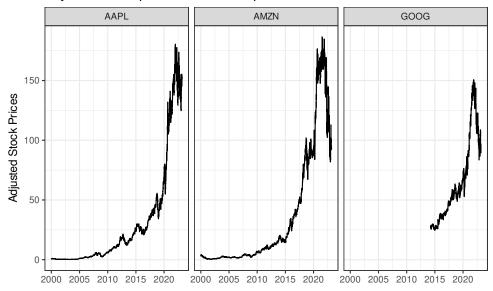
```
tickers <- c("AAPL", 'GOOG', "AMZN") # ticker of APPLE INC
type_statements <- 'pl' # profit/loss statement</pre>
period <- 'fy' # final year
fiscal_year <- 2022
PL <- simfinapi::sfa_get_statement(
  ticker = tickers,
  statement = type_statements,
  period = period,
  fyear = fiscal_year)
# select columns
PL <- PL |>
  dplyr::select(ticker, simfin_id, revenue, net_income)
dplyr::glimpse(PL)
R> Rows: 3
R> Columns: 4
             <chr> "AAPL", "AMZN", "GOOG"
R> $ ticker
R> $ simfin id <int> 111052, 62747, 18
R> $ revenue <dbl> 3.94328e+11, 5.13983e+11, 2.82836e+11
R> $ net_income <dbl> 99803000000, -2722000000, 59972000000
```

As you can see, it is fairly straightforward to download financial data for multiple companies using package {simfinapi}.

5.3.3 Example 03 - Fetching price data

The simfin project also provides prices of stocks, adjusted for dividends, splits and other corporate events. Have a look at the next example, where we download adjusted stock prices for the previous three companies:

Adjusted stock prices for four companies



Price data from simfin

As you can see, the data is comprehensive and should suffice for many different corporate finance research topics.

5.4 Package {tidyquant}

Package {tidyquant} provides functions related to financial data acquisition and analysis. It is an ambitious project that offers many solutions in the field of finance. The package includes functions for obtaining financial data from the web, manipulation of such data, and the calculation of performance measures of portfolios.

First, we will obtain price data for Apple stocks (AAPL) using function $\mathbf{tq}_{\mathbf{get}}()$.

As we can see, except for column names, the price data has a similar format to the one we got with $\{yfR\}$. This is not surprising as both share the same origin, Yahoo Finance.

We can also get information about components of an index using function $tq_index()$. The available market indices are:

R> # A tibble: 157 x 8

```
R>
                            ident~1 sedol
                                           weight sector share~2
      symbol
              company
R>
      <chr>>
              <chr>
                            <chr>
                                    <chr>
                                            <dbl> <chr>
                                                            <dbl>
R>
    1 UCG-IT
              UniCredit S~ BYMXPS
                                    BYMX~ 0.0119 Finan~
                                                            65121
R>
    2 NVDA
              NVIDIA Corp~ 67066G~ 2379~ 0.00999 Infor~
                                                             4726
R>
    3 GE
              General Ele~ 369604~ BL59~ 0.00975 Indus~
                                                            11807
R>
    4 BBVA-ES Banco Bilba~ 550190
                                    5501~ 0.00973 Finan~
                                                           139961
    5 SAN-ES Banco Santa~ 570594
                                   5705~ 0.00962 Finan~
R>
                                                           267887
                                                             6468
R>
    6 SIE-DE
              Siemens Akt~ 572797
                                    5727~ 0.00932 Indus~
    7 5401-JP NIPPON STEE~ 664256
                                   6642~ 0.00925 Mater~
R>
                                                            42618
R>
    8 SGO-FR Compagnie d~ 738048
                                   7380~ 0.00879 Indus~
                                                            16367
    9 8306-JP Mitsubishi ~ 633517
                                    6335~ 0.00877 Finan~
                                                           133448
R> 10 8411-JP Mizuho Fina~ 659101
                                    6591~ 0.00871 Finan~
                                                            59222
R> # ... with 147 more rows, 1 more variable:
       local_currency <chr>, and abbreviated variable names
       1: identifier, 2: shares held
R> #
```

We only looked into a few functions from the package. **{tidyquant}** also offers solutions for the usual financial manipulations, such as calculating returns and functions for portfolio analytics. You can find more details about this package in its website¹⁰.

5.5 Other Packages

In CRAN, you'll find many more packages for importing financial datasets in R. In this section, we focused on packages, which are free and easy to use. Interface with commercial data sources is also possible. Several companies provide APIs for serving data to their clients. Packages such as {Rblpapi} for Bloomberg, {IBrokers} for Interactive Brokers can make R communicate with commercial platforms. If the company you use is not available, check the list of packages in CRAN¹¹. It is very likely you'll find what you need.

5.6 Accessing Data from Web Pages (web-scraping)

Packages from previous section facilitates data importation over the internet. However, in many cases, the information of interest is not available through a package, but on a web page. Fortunately, we can use R to read the data

¹⁰https://business-science.github.io/tidyquant/

¹¹https://cran.r-project.org/

and import the desired information into an R session. The main advantage is that, every time we execute the code, we get the same content available in the website.

The process of extracting information from web pages is called *webscraping*. Depending on the structure and technology used on the internet page, importing its content can be as trivial as a single line in R or a complex process, taking hundreds of lines of code.

5.6.1 Scraping the Components of the SP500 Index from Wikipedia

As an example of webscraping, let's retrieve tabular information about the SP500 index from Wikipedia. In its website, Wikipedia offers a section¹² about the components of the SP500 index. This information is presented in a tabular format, Figure 5.1.

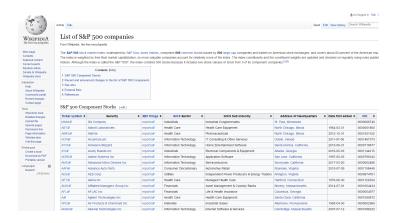


Figure 5.1: Mirror of Wikipedia page on SP500 components

The information on this web page is constantly updated, and we can use it to import information about the stocks belonging to the SP500 index. Before delving into the R code, we need to understand how a webpage works. Briefly, a webpage is nothing more than a lengthy code interpreted by your browser. A numerical value or text presented on the website can usually be found within the code. This code has a particular tree-like structure with branches and classes. Moreover, every element of a webpage has an address, called *xpath*. In chrome and firefox browsers, you can see the actual code of a webpage by using the mouse to right-click any part of the webpage and selecting *View page source*.

 $^{^{12} \}rm https://en.wikipedia.org/wiki/List_of_S\%26P_500_companies$

The first step in webscraping is finding out the location of the information you need. In Chrome, you can do that by right-clicking in the specific location of the number/text on the website and selecting *inspect*. This will open an extra window in the browser. Once you do that, right-click in the selection and click in *copy* and *copy xpath*. In Figure 5.2, we see a mirror of what you should be seeing in your browser.

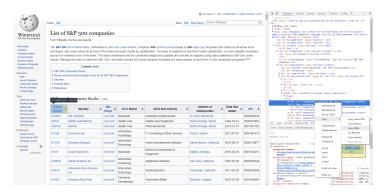


Figure 5.2: Finding xpath from website

Here, the copied *xpath* is:

```
'//*[@id="mw-content-text"]/table[1]/thead/tr/th[2]'
```

This is the address of the header of the table. For the whole content of the table, including header, rows, and columns, we need to set an upper level of the HTML tree. This is equivalent to address //*[@id="MW-content-text"]/table[1].

Now that we have the location of what we want, let's load package {rvest} and use functions read_html(), html_nodes() and html_table() to import the desired table into R:

```
# isolate it
df_SP500_comp <- df_SP500_comp[[1]]</pre>
# change column names (remove space)
names(df_SP500_comp) <- make.names(names(df_SP500_comp))</pre>
# print it
dplyr::glimpse(df_SP500_comp)
R> Rows: 503
R> Columns: 8
R> $ Symbol
                            <chr> "MMM", "AOS", "ABT", "ABBV",~
R> $ Security
                            <chr> "3M", "A. O. Smith", "Abbott~
R> $ GICS.Sector
                            <chr> "Industrials", "Industrials"~
R> $ GICS.Sub.Industry
                           <chr> "Industrial Conglomerates", ~
R> $ Headquarters.Location <chr> "Saint Paul, Minnesota", "Mi~
R> $ Date.added
                            <chr> "1957-03-04", "2017-07-26", ~
R> $ CIK
                            <int> 66740, 91142, 1800, 1551152,~
R> $ Founded
                            <chr> "1902", "1916", "1888", "201~
```

Object df_SP500_comp contains a mirror of the data from the Wikipedia website. The names of the columns require some work, but the raw data is intact and could be further used in a script.

Learning webscraping techniques can give you access to an immense amount of information available on the web. However, each scenario of webscraping is particular. It is not always the case you can import data directly and easily as in previous example.

Another problem is that the webscrapping code depends on the structure of the website. Any simple change in the *html* structure and your code will fail. You should be aware that maintaining a *webscraping* code can demand significant time and effort from the developer. If possible, you should always check for alternative sources of the same information.

Readers interested in learning more about this topic should study the functionalities of packages {XML} and {RSelenium}

5.7 Exercises

01 - Using the yfR package, download daily data of the Facebook stock

(META) from Yahoo Finance for the period between 2019 and 2023. What is the lowest unadjusted closing price (column price.close) in the analyzed period?

02 - If you have not already done so, create a profile on the Quandl website¹³ and download the arabica coffee price data in the CEPEA database (Center for Advanced Studies in Applied Economics)) between 2010-01-01 and 2020-12-31. What is the value of the most recent price?

03 - Use function simfinapi::sfa_get_entities() to import data about all available companies in Simfin. How many companies do you find? (see function dplyr::n_distinct()).

04 - With package simfinapi, download the PL (profit/loss) statement for FY (final year) data for TESLA (ticker = "TSLA") for year 2022. What is the latest Profit/Loss of the company for that particular year?

05 - Using function tidyquant::tq_index, download the current composition of index SP500. What is the company with the highest percentage in the composition of the index?

Be aware that the answer is time-dependent and the reported result might be different from what you actually got in your R session.

06 - Using again the yfR package, download financial data between 2019-01-01 and 2020-01-01 for the following tickers:

• AAPL: Apple Inc

• BAC: Bank of America Corporation

• GE: General Electric Company

TSLA: Tesla, Inc.SNAP: Snap Inc.

Using the **adjusted closing price** column, what company provided higher return to the stock holder during the analyzed period?

 $^{^{13} \}rm https://www.quandl.com/$

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Tip: this is an advanced exercise that will require some coding. To solve it, check out function split to split the dataframe of price data and lapply to map a function to each dataframe.

Chapter 6

Dataframes and Other Objects

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6.1 Dataframes

6.1.1 Creating dataframes



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6.1.2 Inspecting a Dataframe



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6.1.3 The *pipeline* Operators (|> and |>)



6.1. DATAFRAMES 133

6.1.4 Accessing Columns



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6.1.5 Modifying a dataframe

•

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6.1.6 Filtering rows of a dataframe



6.1.7 Sorting a dataframe



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6.1.8 Combining and Aggregating dataframes



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6.1.9 Extensions of the dataframe Class



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6.1.10 Other Useful Functions for Handling dataframes



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6.2 Lists



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6.2.1 Creating lists



6.2.2 Accessing the Elements of a list



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6.2.3 Adding and Removing Elements from a list



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6.2.4 Processing the Elements of a list



6.3. MATRICES 137

6.2.5 Other Useful Functions



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6.3 Matrices



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6.3.1 Selecting Elements from a matrix



6.3.2 Other Useful Functions



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6.4 Exercises



Chapter 7

Basic Object Classes

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7.1 Numeric Objects

7.1.1 Creating and Manipulating numeric Objects



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7.1.2 Creating a numeric Sequence



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7.1.3 Creating Vectors with Repeated Elements



7.1.4 Creating Vectors with Random Numbers



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7.1.5 Accessing the Elements of a numeric Vector



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7.1.6 Modifying and Removing Elements of a numeric Vector



7.1.7 Creating Groups



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7.1.8 Other Useful Functions

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7.2 Character Objects



7.2.1 Creating a Simple character Object



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7.2.2 Creating Structured character Objects



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7.2.3 character Constants



7.2.4 Selecting Pieces of a Text Object



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7.2.5 Finding and Replacing Characters of a Text



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7.2.6 Splitting Text



7.2.7 Finding the Number of Characters in a Text



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7.2.8 Generating Combinations of Text



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7.2.9 Encoding of character Objects



7.2.10 Other Useful Functions



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7.3 Factor Objects



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7.3.1 Creating factors



7.3.2 Modifying factors



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7.3.3 Converting factors to Other Classes



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7.3.4 Creating Contingency Tables



7.3.5 Other Useful Functions



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7.4 Logical Objects



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7.4.1 Creating logical Objects



7.5 Date and Time



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7.5.1 Creating Simple Dates



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7.5.2 Creating a Sequence of Dates



7.5.3 Operations with Dates



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7.5.4 Dealing with Time



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7.5.5 Customizing the Format of Dates and Times



7.5.6 Extracting Elements of a Date



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7.5.7 Find the Current Date and Time



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7.5.8 Other Useful Functions



7.6 Missing Data - NA (Not available)



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7.6.1 Defining NA Values



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7.6.2 Finding and Replacing NA



7.6.3 Other Useful Functions



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7.7 Exercises



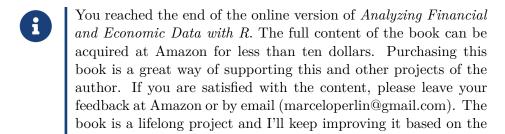
Chapter 8

Programming and Data Analysis

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8.1 R Functions

8.2 Using for Loops



8.3 Conditional Statements (if, else, switch)

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8.4 Functional Programming

8.4.1 Using lapply()



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8.4.2 Using sapply()



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8.4.3 Using tapply()



8.4.4 Using mapply()



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8.4.5 Using apply()



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8.4.6 Using by()



8.5 Using package {purrr}



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8.5.1 Function map()



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8.5.2 Function safely()



8.5.3 Function pmap()



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8.6 Data Manipulation with Package {dplyr}



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8.6.1 Group Operations



8.6.2 Complex Group Operations



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8.7 Exercises



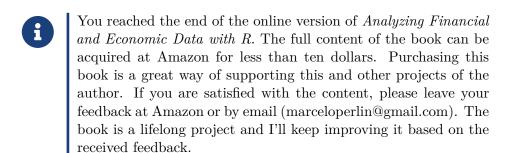
Chapter 9

Cleaning and Structuring Data

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9.1 The Format of a dataframe

9.1.1 Converting a dataframe Structure (long and wide)



9.2 Converting lists into dataframes

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9.3 Removing Outliers

9.3.1 Treating Outliers in dataframes



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9.4 Inflation and Price Data



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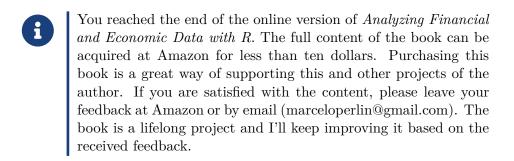
9.5 Modifying Time Frequency and Aggregating Data



9.6 Exercises



Data Visualization with {ggplot2}



10.1 Principles for Data visualization

10.2 The {ggplot2} Package



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10.3 Using Graphics Windows



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10.4 Creating Figures with Function ggplot()



10.5 Data Visualization for Groups



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10.5.1 The US Yield Curve



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10.6 Using Themes



10.7 Creating Panels with facet_wrap



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10.8 Using the Pipeline



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10.9 Creating Statistical Graphics



10.9.1 Creating Histograms



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10.9.2 Creating boxplot Figures



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10.9.3 Creating QQ Plots



10.10 Saving Graphics to a File



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10.11 Exercises



Chapter 11

Financial Econometrics with R

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11.1 Linear Models (OLS)

11.1.1 Simulating a Linear Model



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11.1.2 Estimating a Linear Model



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11.1.3 Statistical Inference in Linear Models



11.2 Generalized Linear Models (GLM)



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11.2.1 Simulating a GLM Model



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11.2.2 Estimating a GLM Model



11.3 Panel Data Models



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11.3.1 Simulating Panel Data Models



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11.3.2 Estimating Panel Data Models



11.4 Arima Models



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11.4.1 Simulating Arima Models



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11.4.2 Estimating Arima Models



11.4.3 Forecasting Arima Models



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11.5 GARCH Models



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11.5.1 Simulating Garch Models



11.5.2 Estimating Garch Models



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11.5.3 Forecasting Garch Models



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11.6 Dealing with Several Models



11.6.1 Using tapply() and sapply()



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11.6.2 Using by()



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11.6.3 Using dplyr::group_by()



11.7 Exercises



 $\boxed{12}$

Reporting Results

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12.1 Reporting Tables

12.2 Reporting Models



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12.3 Creating Reports with *RMarkdown*



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12.4 Exercises



Optimizing Code

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13.1 Optimizing your Programming Time

13.2 Optimizing Code Speed



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13.2.1 Profiling Code



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13.2.2 Simple Strategies to Improve Code Speed



13.2.2.1 Use Vector Operations



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13.2.2.2 Repetitive binding of dataframes



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13.2.3 Using C++ code (package {Rcpp})



13.2.4 Using cache (package {memoise})



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13.2.4.1 Using parallel processing (package {furrr})



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13.3 Exercises



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