# CS&SS 321 - Data Science and Statistics for Social Sciences

Module I - Getting started with R/RStudio

Ramses Llobet

# Welcome!



# (But Seriously) Welcome!

- Welcome to the first quiz section of CS&SS / SOC / STAT 321!
- I am Ramses Llobet (rllobet@uw.edu), I am a Ph.D. student in Political Science.
- My research interest are in political economy and applied statistics.
- Please DO NOT hesitate to stop me if you don't hear or understand me properly.
- ▶ DO NOT hesitate to ask questions. No question is silly. :)

Now it's your turn

Name and major/year (or intended major)?

• Why are you take this course?

► What is your experience with R (zero shame)?

# R setup

- ► How to install R and R-studio.
  - R-4.3.2 for Windows
  - ► R-4.3.2 for macOS
- R-studio can be downloaded from posit's repository.
- I recommend this tutorial from Casey Bates for an overview of R and RStudio.
- Live coding: how to install packages and start tutorials (setting\_up.R).
- For Mac users, installation of the **qss** package may sometimes fail if **pandoc** or **curl** is not installed or updated on your Mac. To resolve this, you can:
  - 1. Install the package manager Homebrew package.
  - Then use the macOS terminal to install pandoc or curl using the commands brew install pandoc or brew install curl.

## Useful free online R resources

Introductory:

► Grolemund (2014) Hands-On Programming with R.

Intermediary:

▶ Wickham et al. (2023) *R* for Data Science. 2nd Edition.

R Markdown

► Xie et al. (2022) R Markdown: The Definitive Guide

Others

Stack Overflow.

ChatGPT

# Project management and working directory

- A good practice is to keep your projects and files organized and tidy.
  - Avoid accumulating data and R files in your downloads folder.
- I recommend creating an R project file in your course folder materials. R projects have several advantages:
  - Centralized and efficient *workflow*.
  - Sets the **current** (*root*) working directory.
  - See more in Martin Chan's beginner's guide.

# What are working directories?

- A directory is a folder in a file system that stores files and other sub-directories.
- A path is a string that specifies the location of a directory in a file system.
- ► For example:
  - ▶ D:\Google Drive
  - D:\Google Drive\Phd UW\Courses\Third Year\CSSS 594
    - Text as Data
- When you run a command or script, R looks for files and sub-directories based on relative paths to your current working directory.

# Absolute and relative paths

- Absolute Path: Specifies the full path from the root directory to the file or directory.
  - For example: D:\Google Drive\Phd UW\Courses\Third Year\CSSS 321\scripts\setting\_up.R is an absolute path.
- Relative Path: Specifies the path relative to the current working directory.
  - For example, if the working directory is D:\Google Drive\Phd UW\Courses\Third Year\CSSS 321, then
    - scripts\setting\_up.R is a relative path.

# Project management: working directory

 .Rproj (R Project File) in your project folder establishes the working directory as its absolute path upon opening R.



- Employing .Rproj and relative paths in R streamlines project management and collaboration by overseeing files, inputs, and outputs.
  - Live demostration of how to create and manage an R Project File.

Project management: workflow



# Working directories: obsolete practices

- ► Workflow with .*Rproj* is relatively **new**.
- Until recently, users had to manually set working directories using functions or specialized packages. See example:

#### setwd() # function to set directory

setwd("D:/Google Drive/Phd UW/Courses
 /Third Year/CSSS 594 - Text as Data
 /presentation") # remember to put quotes

# What are functions?

- They are a set of instructions that performs a specific task in R.
- Functions often take one or more arguments, which are inputs that are used to customize the behavior of the function.
- The mean() function takes one required argument, which is the vector of numbers to calculate the mean of.

```
# create a vector consisting of midterm scores.
grades_M <- c(76, 82, 94, 45, 75)</pre>
```

# calculate the mean using the mean() function mean(grades\_M)

## [1] 74.4

# What are functions?

the mean() function also has additional optional arguments, which can be used to further customize the behavior of the function.

```
# create a vector consisting of final scores.
grades F <- c(82, 90, 89, NA, 64)</pre>
```

```
# calculate the mean using the mean() function
mean(grades_F)
```

## [1] NA

# use the argument `na.rm` to evaluate the removal of NAs
mean(grades\_F, na.rm= TRUE)

```
## [1] 81.25
```

Remember: use ? or help() to see the documentation of a function.

- Save the following Cheat Sheet for RMarkdown.
- If any of you is looking for an general introduction for RMarkdown, I suggest you to check Chapter 27 from Wickham and Grolemund (2017) - R for Data Science.
- If you want a more comprehensive guide, then check Xie et al. (2021) R Markdown: The Definitive Guide.
- Another, more applied, resource is Xie et al. (2022) R Markdown Cookbook.

- RMarkdown is a document format that allows you to integrate R code and output into a single document.
- Besides R code and output, it can also include text, images, and other multimedia elements, allowing for rich and informative documents.
- Pandoc is a free and open-source document converter that can convert documents from one markup language to another.
  - In the context of Rmarkdown, pandoc is the underlying document converter (sfotware) that converts the R-markdown file into a final output format, such as HTML, PDF, or Word.

The output format of the final document can be customized using options in the YAML header or external templates.



- The YAML header in RMarkdown is a block of configuration settings at the beginning of the document enclosed by three hyphens (---).
- It is used to specify document metadata and other settings such as the document title, author, output format, and more.

► Code chunks are sections of R code that can be executed and embedded within an RMarkdown document.



- Code chunks can be inserted using the syntax {r} and closed with "'.
  - ► Short cut in Windows: Ctrl + Alt + I
  - Short cut in macOS: Cmd + Option + I
- Code chunks can be customized with various **chunk options**.
- Note: set the function knitr::opts\_chunk\$set() with any general setting without repeating it in every code chunk.

#### Frequently used chunk options

Option	Description
include	If FALSE, knitr will run the chunk but <b>not</b> include the chunk in the final document
echo	If FALSE, knitr will <b>not</b> display the code in the code chunk above it's results in the final document.
error	If FALSE, knitr will <b>not</b> display any error messages generated by the code.
message	If FALSE, knitr will <b>not</b> display any messages generated by the code.
warning	If FALSE, knitr will <b>not</b> display any warning messages generated by the code.

# **Recommendation for Homework**

Option	HW setting
include	TRUE
echo	TRUE
error	FALSE
message	FALSE
warning	FALSE



- In RMarkdown, rendering a document means converting the source RMarkdown file into its final output format (using pandoc).
- To render a document, we need to Knit, knitting is the process of taking the RMarkdown file and converting it into a single, cohesive document that can be rendered into different formats (HTML, PDF, etc).
  - ► To produce **PDF file**, you need TeX files.
- Easy way: Install the tinytex package: install.packages("tinytex"). Then run tinytex::install\_tinytex().

# Knitting

► To knit:

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► Auxiliary window for output preview:



# Working directories and R-Markdown

- When opening an RMarkdown file, this will set the file location as the working directory.
- Change the following option in the global options to avoid this behavior:



Live demonstration and in-class exercise:

Open the file RMarkdown\_sample.Rmd

Reference: David Robinson

x <- 10 + foo

Error: object 'foo' not found: You tried to access a variable that doesn't exist.

You might have:

- misspelled the variable name
- ▶ incorrectly **capitalized** the variable name (R is case sensitive!)
- forgotten to run the line that defines the variable in the first place, or run into an error on that line.

x <- foo(...)

Error: could not find function "foo": You tried to use a function that doesn't exist. You might have:

- misspelled the function name
- incorrectly capitalized the function name
- ▶ forgotten to **load the library** that provides this function.

x <- c(1:10))

Error: unexpected ')' in ...: There is an extra end parenthesis in your line

Error: unexpected symbol in ...: The most common cause of this is forgetting a punctuation mark such as a comma: for example, foo(bar1 bar2) instead of foo(bar1, bar2).

Error: unexpected numeric constant is similar: it just means the value after the missing punctuation is a number (for example, x 2 instead of x = 2).

paste("welcome to CSSS, 321)

- You might see a + sign in the interpreter after you hit return. This means the previous statement is unfinished:
- it might have an open parenthesis that never closes,
  - ▶ an open " or ' that is unmatched, or
  - it could end with an operator like + or that expects the line to continue afterwards.
- Find the problem in your previous lines (count parentheses, and check your quotes) and fix it.

# Getting help: Using the Internet to Your Advantage

- When encountering coding error messages, use Google or post on Stack Overflow for solutions.
- Both beginners and experts often rely on online searches for coding assistance.
- For example, let's say that I want to know how to rename a column in my dataset. I could Google:
  - ► "How to rename a column in R" ... and look to the answer.
  - Make sure that you understand the terminology.

# Getting help: minimal reproducible example

- If you feel stuck with an error, seek help but remember to provide reproducible code in an R-script file:
  - 1. Load necessary packages at the beginning.
  - 2. Include all code up to the error, or at least the **necessary** to reproduce it.
  - 3. Comment your code for clarity.
  - 4. If applicable, send the necessary data to reproduce the error.

Getting help: practice

- Open the file common\_errors.Rmd, and try to solve the problems.
- You will need to submit your work to me via Slack.

# **Review of basics**

The subsequent slides show some R basics that we have already covered during the live demonstrations but feel free to review them on your own.

# Running R code and operators

# Arithmetic Operators 1 + 1
## [1] 2
2 * 8
## [1] 16
9 / 3
## [1] 3
2^3
## [1] 8

## Running R code and operators

# Relational Operators
10 > 8 # is 10 bigger than 8?

## [1] TRUE

7 <= 6 # is 7 less or equal to 6?

## [1] FALSE

(2 \* 5) == 10 # is 2\*5 equal to 10?

## [1] TRUE

1 != 2 # is 1 unequal to 2?

#### ## [1] TRUE

## **Objects in R: vectors and assignment**

```
# Concatenate vectors into a new vector
c(1, 2, 3)
```

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

```
# Assign them to a new object for manipulation
x <- c(1, 2, 3)
print(x) # or simply, x</pre>
```

## [1] 1 2 3

# Operators on vector
x + 1

## [1] 2 3 4

# Logical test on vector
x == 1

# **Objects in R: vectors and functions**

# Use an object as input to a function  $x \le c(1, 2, 3)$ 

# Functions take input(s) and produce output(s)
class(x)

## [1] "numeric"

length(x)

## [1] 3

mean(x)

## [1] 2

## **Objects in R: introductory tips**

 Unless you assign (<- ) some operations or transformations to an object, those values will not be registered

```
x < -c(1, 2, 3)
print(x + 1)
## [1] 2 3 4
print(x)
## [1] 1 2 3
x < -x + 1
print(x)
```

## [1] 2 3 4

# **Objects in R: introductory tips**

New assignment will overwrite the original values if you assign some values to an existing object. It is a major source of errors. One advise is to keep distinct object names

```
x <- c(1, 2, 3)
length(x)
## [1] 3
x <- c(1, 2, 3, 4, 5)
length(x)
```

## [1] 5

## **Objects in R: atomic vectors**

 Most common types of atomic vectors: numeric (integer, double), logical, character

```
x <- c(1, 2, 3)
class(x)</pre>
```

```
## [1] "numeric"
```

```
y <- c(TRUE, FALSE, FALSE)
class(y)</pre>
```

```
## [1] "logical"
```

```
names <- c("Peter", "Paul", "Mary")
class(names)</pre>
```

```
## [1] "character"
```

## **Objects in R: atomic vectors**

You can also coerce one type of vector into another:

```
x <- c(1, 2, 3)
x <- as.character(x)
print(x)
## [1] "1" "2" "3"
```

class(x)

## [1] "character"

# **Objects in R: reading data**

▶ You can import any data file and assign it into an object

```
x <- c(1, 2, 3)
x <- as.character(x)
print(x)
## [1] "1" "2" "3"
```

class(x)

```
## [1] "character"
```