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

Module I - Getting started with R/RStudio

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## 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. candidate 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.

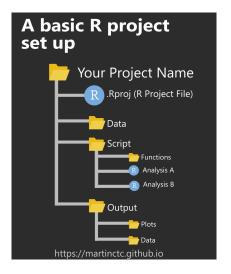
# Project management: working directory

► .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:

## 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)

# calculate the mean using the mean() function
mean(grades_M)</pre>
```

## [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)

# 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)</pre>
```

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

## [1] 81.25

- ► 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.

```
title: "Lab 1 - Intro to RMarkdown"
author: "Your name"
date: \today
output:
pdf_document:
latex_engine: pdflatex
fontsize: 12pt
editor_options:
chunk_output_type: console

11 ---
12
```

- ► 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.

```
78
79 ▼ ```{r name, error=TRUE, warning=FALSE} 

# brau brau, derp herp
head(data)

82 ▲
83
```

- ► 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

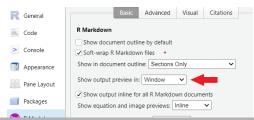
- In RMarkdown, rendering a document means converting the source RMarkdown file into its final output format (using pandoc).
- ► To render a document, we need to Knit, knitting is the process of taking the RMarkdown file and converting it into a single, cohesive document that can be rendered into different 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:

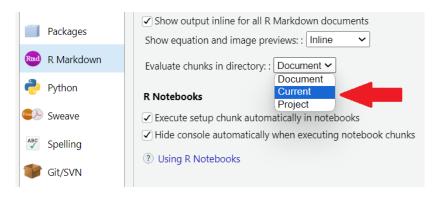


► Auxiliary window for output preview:



## 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 \leftarrow c(1:10)
```

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

```
x <- 10; y <- 20
mean(x y)
```

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).

```
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: reproducible code

- ▶ 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.
  - Include all code up to the error, or at least the necessary to reproduce it.
  - Comment your code for clarity.
  - **4.** If applicable, send the necessary **data** to reproduce the error.

#### **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 \leftarrow c(1, 2, 3)
print(x) # or simply, x
## [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 \leftarrow 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 \leftarrow 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</pre>
```

## Objects in R: atomic vectors

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

```
x \leftarrow c(1, 2, 3)
class(x)
## [1] "numeric"
y <- c(TRUE, FALSE, FALSE)
class(y)
## [1] "logical"
names <- c("Peter", "Paul", "Mary")</pre>
class(names)
## [1] "character"
```

## Objects in R: atomic vectors

► You can also coerce one type of vector into another:

```
x \leftarrow 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"</pre>
```