# Reports and manuscripts with Quarto

# What is quarto?

- Open-source document format and computational notebook system
- Integrates text, code, and outout
- Can create multiple different types of products (documents, slides, websites, books)



# Why not R Markdown?

Only because quarto is newer and more featured!

- Anything you already know how to do in R Markdown you can do in quarto, and more!
- All of these slides, website, etc. are all made in quarto.
- If you know and love R Markdown, by all means keep using it!

## Quarto workflow

- 1. Create a Quarto document
- 2. Write code
- 3. Write text
- 4. Repeat 2-3 in whatever order you want
- 5. Render repeatedly as you go

### How does it work?

- You text in markdown and code in R
- knitr processes the code chunks, executes the R code, and inserts the code outputs (e.g., plots, tables) back into the markdown document
- pandoc transforms the markdown document into various output formats



### Text and code...

```
1 # My header
 3 Some text
 5 Some *italic text*
 6
 7 Some **bold text**
 8
 9 - Eggs
10 - Milk
11
12 \\ \rangle \{r\}
13 x <- 3
```

### ... becomes ...

### Mytitle

Some text

Some italic text

Some bold text

- Eggs
- Milk

```
1 x <- 3
2 x
```

# If you prefer, you can use the visual editor

### My title

Some text

Some *italic text* 

#### Some **bold text**

```
{r}
x <- 3
x
```

### Rchunks

Everything within the chunks has to be valid R<sup>1</sup>

```
1 ```{r}
2 x <- 3
3 ```

1 ```{r}
2 x + 4
3 ```
```

Chunks run in order, continuously, like a single script

### YAML

At the top of your Quarto document, a header written in yaml describes options for the document

```
1 ---
2 title: "My document"
3 author: Louisa Smith
4 format: html
5 ---
```

There are a *ton* of possible options, but importantly, this determines the document output

## Output



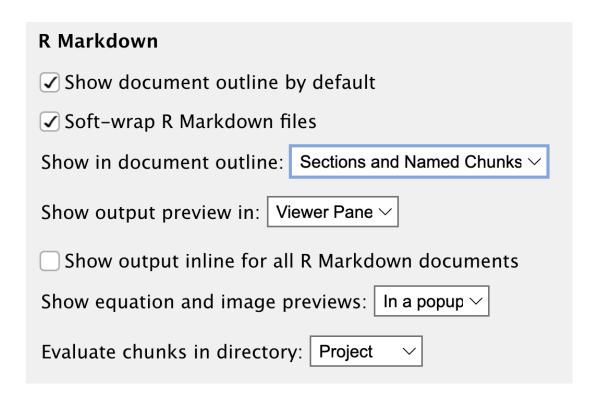
https://quarto.org/docs/output-formats/all-formats.html

We're going to focus on html output

- It's easy to transition to Word (format: docx) but it's not as good for constant re-rendering
- You need a LaTeX installation for pdf (format: pdf)
  - I recommend {tinytex}

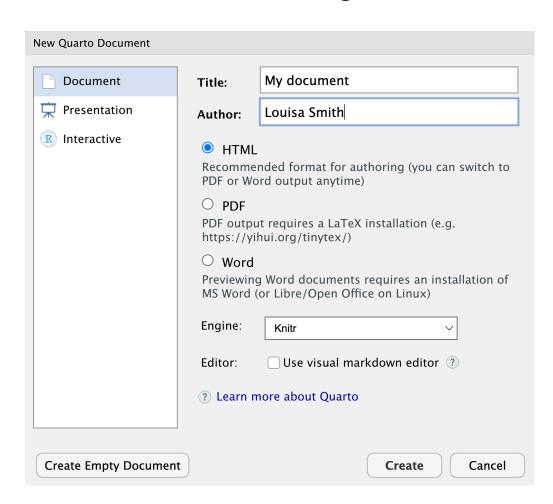
You can choose whether you want to have chunk output show up within the document (vs. just the console) when you are running Quarto (and RMarkdown) documents interactively

### Tools > Global options

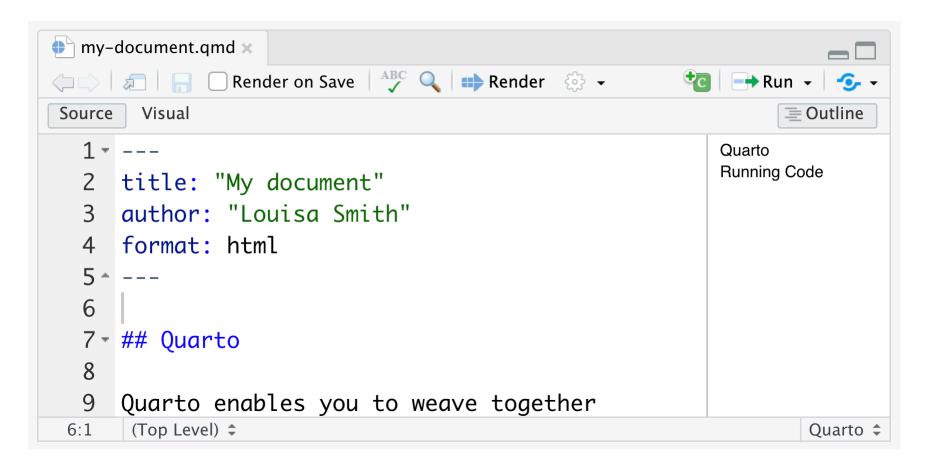


Open up your epi590r-2023-in-class R project!

File > New File > Quarto Document



- Try toggling between Source and Visual views
- Toggle on and off the Outline
- Click Render and look at the output



# Quartooptions

# Chunk options

In your Quarto document, you had a chunk:

```
1 ```{r}
2 #| echo: false
3 2 * 2
4 ```
```

#| echo: false tells knitr not to show the code within that chunk

In RMarkdown, you would have written this  $\{r, echo = FALSE\}$ . You can still do that with Quarto, but it's generally easier to read, particularly for long options (like

# Chunk options

Some of the ones I find myself using most often:

- #| eval: false: Don't evaluate this chunk! Really helpful if you're trying to isolate an error, or have a chunk that takes a long time
- #| error: true: Render this *even if* the chunk causes an error
- #| cache: true: Store the results of this chunk so that it doesn't need to re-run every time, as long as there are no changes
- #| warning: false: Don't print warnings
- #| message: false: Don't print messages

# Document options

You can tell the *entire* document not to evaluate or print code (so just include the text!) at the top:

```
title: "My document"
author: Louisa Smith
format: html
execute:
eval: false
ceho: false
```

Careful! YAML is *really* picky about spacing.

# Document options

There are lots of different options for the document

• For example, you can choose a theme:

```
1 ---
2 format:
3 html:
4 theme: yeti
5 ---
```

 Remember the pickiness: when you have a format option, <a href="https://html:">html:</a> moves to a new line and the options are indented 2 spaces

Download the quarto document with some {gtsummary} tables from yesterday

- Change some options to hide code and output
- Add another code chunk
- Play with themes
- Deal with errors
- Add some text

# Quarto tables, figures, and stats

# Chunks can produce figures and tables

```
1 ```{r}
2 #| label: tbl-one
3 #| tbl-cap: "This is a great table"
4 knitr::kable(mtcars)
5 ```
```

Table 1: This is a great table

	mpg	cyl	disp	hp	drat	wt	qse
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.40
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.07
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.6
Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44

	mpg	cyl	disp	hp	drat	wt	qse
Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02
Valiant	18.1	6	225.0	105	2.76	3.460	20.2
Duster 360	14.3	8	360.0	245	3.21	3.570	15.84
Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00
Merc 230	22.8	4	140.8	95	3.92	3.150	22.90
Merc 280	19.2	6	167.6	123	3.92	3.440	18.30
Merc 280C	17.8	6	167.6	123	3.92	3.440	18.90
Merc 450SE	16.4	8	275.8	180	3.07	4.070	17.4(
Merc 450SL	17.3	8	275.8	180	3.07	3.730	17.60

	mpg	cyl	disp	hp	drat	wt	qse
Merc 450SLC	15.2	8	275.8	180	3.07	3.780	18.00
Cadillac Fleetwood	10.4	8	472.0	205	2.93	5.250	17.98
Lincoln Continental	10.4	8	460.0	215	3.00	5.424	17.8%
Chrysler Imperial	14.7	8	440.0	230	3.23	5.345	17.47
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.4'
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.5%

	mpg	cyl	disp	hp	drat	wt	qse
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90
Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.0
Dodge Challenger	15.5	8	318.0	150	2.76	3.520	16.8′
AMC Javelin	15.2	8	304.0	150	3.15	3.435	17.3(
Camaro Z28	13.3	8	350.0	245	3.73	3.840	15.4
Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.0

	mpg	cyl	disp	hp	drat	wt	qse
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.5(
Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50
Maserati Bora	15.0	8	301.0	335	3.54	3.570	14.60
Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60

# Chunks can produce figures or tables

```
1 ```{r}
2 #| label: fig-hist
3 #| fig-cap: "This is a histogram"
4 hist(rnorm(100))
5 ```
```

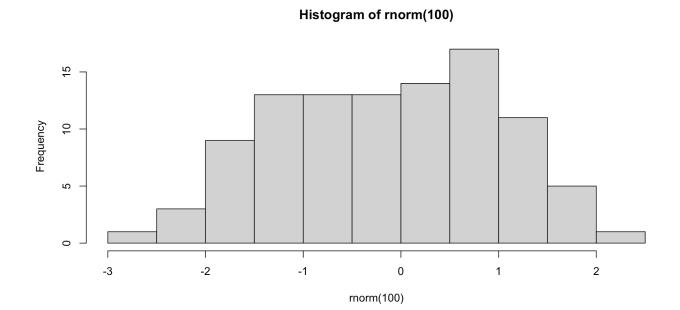


Figure 1: This is a histogram

# Cross-referencing

You can then refer to those with <a href="etable-one">@tbl-one</a> and <a href="etage-hist">@fig-hist</a> and the Table and Figure ordering will be correct (and linked)

@fig-hist contains a histogram and @tbl-one a table.

gets printed as:

Figure 1 contains a histogram and Table 1 a table.

### Inline R

Along with just regular text, you can also run R code within the text:

```
There were `r 3 + 4` participants
```

becomes:

There were 7 participants

### Inline R

This is helpful for reporting statistics, e.g. the sample size:

```
There were `r nrow(nlsy)` participants
```

becomes:

There were 1.2686^{4} participants

### Inline stats

You can also create an object in a chunk and then reference it later in the text

```
1 ```{r}
2 total_sample <- nrow(nlsy)
3 ```
There were `r total_sample` participants</pre>
```

# Inline stats (aside)

I often create a list of stats that I want to report in a manuscript:

```
1 stats <- list(n = nrow(data),
2 mean_age = mean(data$age))
```

I can then print these numbers in the text with:

There were `r stats\$n` participants with a mean age of `r stats\$mean\_age`.

which turns into:

There were 1123 participants with a mean age of 43.5.

# Inline stats from {gtsummary}

We saw very, very briefly yesterday:

```
library(gtsummary)
          2 income table <- tbl uvregression(</pre>
              nlsy,
          4 	 y = income_{r}
          5 include = c(
           sex cat, race eth cat,
                eyesight_cat, income, age_bir
              method = lm
         10)
            inline text(income table, variable = "age bir")
[1] "595 (95% CI 538, 652; p<0.001)"
```

We pulled a statistic from our univariate table

# If we're making a table, we probably want to report numbers from it

```
1 \``{r}
2 # | label: tbl-descr
3 # | tbl-cap: "Descriptive statistics"
4 # output-location: slide
5 table1 <- tbl summary(</pre>
   nlsy,
 7 	 by = sex cat,
   include = c(sex_cat, race_eth_cat, region_cat,
                  eyesight cat, glasses, age bir)) |>
     add overall(last = TRUE)
   table1
12
```

# If we're making a table, we probably want to report numbers from it

Table 2: Descriptive statistics

	Male	Female	Overall
Characteristic	N = 6,403	$N = \frac{6}{1},283$	N = 12,686
race_eth_cat			
Hispanic	1,000 (16%)	1,002 (16%)	2,002 (16%)
Black	1,613 (25%)	1,561 (25%)	3,174 (25%)
Non-Black, Non-Hispanic	3,790 (59%)	3,720 (59%)	7,510 (59%)
region_cat			
Northeast	1,296 (21%)	1,254 (20%)	2,550 (20%)
North Central	1,488 (24%)	1,446 (23%)	2,934 (24%)
South	2,251 (36%)	2,317 (38%)	4,568 (37%)
West	1,253 (20%)	1,142 (19%)	2,395 (19%)
Unknown	115	124	239
eyesight_cat			
1			

	Male	Female	Overall
Characteristic	$N = \frac{6}{1},403$	N = 6,283	N = 12,686
Excellent	1,582 (38%)	1,334 (31%)	2,916 (35%)
Very good	1,470 (35%)	1,500 (35%)	2,970 (35%)
Good	792 (19%)	1,002 (23%)	1,794 (21%)
Fair	267 (6.4%)	365 (8.5%)	632 (7.5%)
Poor	47 (1.1%)	85 (2.0%)	132 (1.6%)
Unknown	2,245	1,997	4,242
glasses	1,566 (38%)	2,328 (54%)	3,894 (46%)
Unknown	2,241	1,995	4,236
age_bir	25 (21, 29)	22 (19, 27)	23 (20, 28)
Unknown	3,652	3,091	6,743
<sup>1</sup> n (%); Median (Q1, Q3)			

#### I want to report some stats!

The help file for <a href="inline\_text">inline\_text</a>() is helpful and tells us that we can look at <a href="table\_body">table1\$table\_body</a> to help figure out what data to extract.

How about the median (IQR) age of the male participants at the birth of their first child?

```
1 inline_text(table1, variable = "age_bir", column = "stat_]
[1] "25 (21, 29)"
```

## Formatting

We can add sample sizes for the overall stats on people who wear glasses using the <a href="mailto:pattern">pattern</a> = argument:

## Formatting for regression statistics

Remove some details:

#### Better yet...

We can integrate these into the text of our manuscript:

```
A greater proportion of female (`r inline_text(table1, variable = "glasses", column = "stat_2")`) than male

(`r inline_text(table1, variable = "glasses", column = "stat_1")`) participants wore glasses.
```

#### Which becomes:

A greater proportion of female (2,328 (54%)) than male (1,566 (38%)) participants wore glasses.

### Readability

Because this can be hard to read, I'd suggest storing those stats in a chunk before the text:

#### Exercises

Return to the quarto document with the tables.

- Choose a table to label and caption, and then write a sentence that cross-references it (e.g., Table 1 shows the descriptive statistics)
- Edit or create a new table that has the median hours of sleep on weekends in it (in the overall sample).
- Pull that value from a table.

## Functions

#### Functions in R

I've been denoting functions with parentheses: func()
We've seen functions such as:

- mean()
- tbl\_summary()
- init()
- create\_github\_token

Functions take **arguments** and return **values** 

## Looking inside a function

If you want to see the code within a function, you can just type its name without the parentheses:

```
1 usethis::create_github_token
```

```
function (scopes = c("repo", "user", "gist", "workflow"), description = "DESCRIBE THE TOKEN'S USE
CASE",
   host = NULL)
   scopes <- glue collapse(scopes, ",")</pre>
   host <- get hosturl(host % | | % default api url())
   url <- glue("{host}/settings/tokens/new?scopes={scopes}&description={description}")</pre>
   withr::defer(view url(url))
   hint <- code hint with host("gitcreds::gitcreds set", host)</pre>
   message <-c() = "Call {.run {hint}} to register this token in the local Git\n
credential store.")
   if (is linux()) {
       message <- c(message, `!` = "On Linux, it can be tricky to store credentials
persistently.",
            i = "Read more in the {.href ['Managing Git(Hub) Credentials' article]
(https://usethis.r-lib.org/articles/articles/git-credentials.html)}.")
   message <- c(message, i = "It is also a great idea to store this token in any\n
password-management software that you use.")
```

```
ui_bullets(message)
   invisible()
}
<bytecode: 0x141ac26a0>
<environment: namespace:usethis>
```

```
1 func <- function()</pre>
```

You can name your function like you do any other object

Just avoid names of existing functions

What objects/values do you need to make your function work?

 You can give them default values to use if the user doesn't specify others

Everything else goes within curly braces

 Code in here will essentially look like any other R code, using any inputs to your functions

- One thing you'll likely want to do is make new objects along the way
- These aren't saved to your environment (i.e., you won't see them in the upper-right window) when you run the function
- You can think of them as being stored in a temporary environment within the function

Return something new that the code has produced

• The return() statement is actually optional. If you don't put it, it will return the last object in the code. When you're starting out, it's safer to always explicitly write out what you want to return.

### Example: a new function for the mean

Let's say we are not satisfied with the mean() function and want to write our own.

Here's the general structure we'll start with.

```
1 new_mean <- function() {
2
3 }</pre>
```

#### New mean: arguments

We'll want to take the mean of a vector of numbers.

It will help to make an example of such a vector to think about what the input might look like, and to test the function. We'll call it  $\mathbf{x}$ :

```
1 \times < -c(1, 3, 5, 7, 9)
```

We can add x as an argument to our function:

```
1 new_mean <- function(x) {
2
3 }</pre>
```

### New mean: function body

Let's think about how we calculate a mean in math, and then translate it into code:

$$\overline{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_{i}$$

So we need to sum the elements of x together, and then divide by the number of elements.

We can use the functions sum() and length() to help us.

We'll write the code with our test vector first, before inserting it into the function:

```
1 n <- length(x)
2 sum(x) / n
```

### New mean: function body

Our code seems to be doing what we want, so let's insert it. To be explicit, I've stored the answer (within the function) as mean\_val, then returned that value.

```
1 new_mean <- function(x) {
2    n <- length(x)
3    mean_val <- sum(x) / n
4    return(mean_val)
5 }</pre>
```

## Testing a function

Let's plug in the vector that we created to test it:

```
1 new_mean(x = x)
[1] 5
```

And then try another one we create on the spot:

```
1 new_mean(x = c(100, 200, 300))
```

#### Exercises

Create some functions!

Create an R script in your project called **functions.** R to save your work!

## Functions, continued

## Adding another argument

Let's say we plan to be using our <a href="new\_mean">new\_mean</a>() function to calculate proportions (i.e., the mean of a binary variable). Sometimes we'll want to report them as as percentage by multiplying the proportion by 100.

Let's name our new function prop(). We'll use the same structure as we did with new\_mean().

```
1 prop <- function(x) {
2    n <- length(x)
3    mean_val <- sum(x) / n</pre>
```

## Testing the code

Now we'll want to test on a vector of 1's and 0's.

```
1 \times < -c(0, 1, 1)
```

To calculate the proportion and turn it into a percentage, we'll just multiply the mean by 100.

```
1 multiplier <- 100
2 multiplier * sum(x) / length</pre>
```

[1] 66.66667

## Testing the code

We want to give users the option to choose between a proportion and a percentage. So we'll add an argument multiplier. When we want to just return the proportion, we can just set multiplier to be 1.

```
1 multiplier <- 1
         2 multiplier * sum(x) / length
11 0.6666667
         1 multiplier <- 100
         2 multiplier * sum(x) / length
```

### Adding another argument

If we add multiplier as an argument, we can refer to it in the function body.

```
1 prop <- function(x, multipli
2    n <- length(x)
3    mean_val <- multiplier * s
4    return(mean_val)
5 }</pre>
```

### Adding another argument

Now we can test:

```
1 prop(x = c(1, 0, 1, 0), multiplier = 1)

[1] 0.5

1 prop(x = c(1, 0, 1, 0), multiplier = 100)
[1] 50
```

## Making a default argument

Since we don't want users to have to specify multiplier = 1 every time they just want a proportion, we can set it as a default.

```
prop <- function(x, multiplier = 1) {
    n <- length(x)
    mean_val <- multiplier * sum(x) / n
    return(mean_val)
}</pre>
```

Now we only need to specify that argument if we want a percentage.

```
1 \text{ prop}(x = c(0, 1, 1, 1))
1 \text{ prop}(x = c(0, 1, 1, 1), \text{ multiplier} = 100)
```

#### Caveats

- This is obviously not the best way to write this function!
- For example, it will still work if x = c(123, 593, -192).... but it certainly won't give you a proportion or a percentage!
- We could also put multiplier = any number, and we'll just be multiplying the answer by that number this is essentially meaningless.
- We also haven't done any checking to see whether the user is even entering numbers! We could put in better error messages so users don't just get an R default error message if they do something wrong.

```
1 prop(x = c("blah", "blah", "blah"))
```

## Functions, continued

# Writing functions to reduce copy and paste

If you find yourself doing something over and over, you can probably write a function for it.

One is example is the regression tables we created yesterday.

# Let's imagine we are fitting multiple similar models

```
1 logistic_model <- glm(glasses ~ eyesight_cat + sex_cat + i
2    data = nlsy, family = binomial()
3 )
4 poisson_model <- glm(nsibs ~ eyesight_cat + sex_cat + incomplete to the sex_cat + incomp
```

# We have a model for a table we want to create

Even though this is already a function, we can *wrap* it in a new function!

#### New table function

We can refer generically to model and then put the model we want a table for as an argument

```
new table function <- function(model) {</pre>
     tbl regression(
       model,
 4 exponentiate = TRUE,
 5 label = list(
         sex cat ~ "Sex",
         eyesight cat ~ "Eyesight",
         income ~ "Income"
10
11 }
```

- Copy and paste this function into your script
- Create each of the models and run
- Try to figure out how you could allow someone to pass the tidy\_fun argument (from the regression exercises yesterday)

# {renv}

Package management for R

# What is {renv}?

{renv} is an R package for managing project dependencies and creating reproducible environments

# Benefits of using {renv}

- 1. **Isolation:** Creates project-specific environments separate from the global R library.
- 2. **Reproducibility:** Ensures consistent package versions for code reproducibility.
- 3. **Collaboration:** Facilitates sharing and collaborating on projects with others.

# Getting Started with {renv}

1. Install {renv}

```
1 install.packages("renv")
```

2. Initialize a project

```
1 renv::init()
```

3. Install packages

```
install.packages("other_package")

# only an option when using renv!

install.packages("github_user/github_package")
```

4. Track dependencies via a lockfile

```
1 renv::snapshot()
```

#### Behind the scenes

• Your project .Rprofile is updated to include:

```
1 source("renv/activate.R")
```

- This is run every time R starts, and does some management of the library paths to make sure when you call install.packges("package") or library(package) it does to the right place (renv/library/R-{version}/{computer-specifics})
- A renv. lock file (really just a text file) is created to store the names and versions of the packages.

## renv.lock

```
"R": {
  "Version": "4.3.0",
  "Repositories": [
      "Name": "CRAN",
      "URL": "https://cran.rstudio.com"
"Packages": {
  "R6": {
    "Package": "R6",
    "Version": "2.5.1",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
      "R"
    "Hash": "470851b6d5d0ac559e9d01bb352b4021"
  base64enc": {
    "Package": "base64enc",
    "Version": "0.1-3",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
```

```
"R"
],
"Hash": "543776ae6848fde2f48ff3816d0628bc"
},
```

# Using {renv} later

Restore an environment

```
1 renv::restore()
```

Install new packages

```
1 install.packages("other_package")
```

Update the lockfile

```
1 renv::snapshot()
```

# Collaboration with {renv}

- Share the project's renv. lock file with collaborators to ensure consistent environments
- When they run <a href="renv::restore">renv::restore</a>(), the correct versions of the packages will be installed on their computer

```
1 renv::restore()
```

# Other helpful functions

Remove packages that are no longer used:

```
1 renv::clean()
```

Check the status of the project library with respect to the lockfile:

```
1 renv::status()
```

This will tell you to <a href="renv::snapshot">renv::snapshot</a>() to add packages you've installed but haven't snapshotted, or <a href="renv::restore">renv::restore</a>() if you're missing packages you need but which aren't installed

#### Conclusion

{renv} benefits:

Isolation, reproducibility, and collaboration

Getting started with {renv}

- 1. Initialize a project using renv::init()
- 2. Install packages and store with renv::snapshot()
- 3. Restore later or elsewhere with renv::restore()

- 3. Install a new R package of your choice. (Not sure what to choose? Try one of these fun packages. For example, I did install.packages("hadley/emo").)
- 4. Create an R script and save it in your R project. Include some code that requires the package. For example:

```
1 emo::ji("banana")
```

- 4. Run renv::status() to make sure your project picked up the package as a dependency.
- 5. Run renv::snapshot() to record that package in your lockfile.
- 6. Open your lockfile and look for your new package. For example, mine now has:

```
"emo": {
      "Package": "emo",
      "Version": "0.0.0.9000",
      "Source": "git",
      "RemoteType": "git",
      "RemoteUrl": "https://github.com/hadley/emo",
      "RemoteHost": "api.github.com",
      "RemoteUsername": "hadley",
      "RemoteRepo": "emo",
      "RemoteRef": "master",
      "RemoteSha": "3f03b11491ce3d6fc5601e210927eff73bf8e350",
      "Requirements": [
        "R",
        "assertthat",
        "crayon",
        "glue",
        "lubridate",
        "magrittr",
```

# Reports and manuscripts with Quarto

# What is quarto?

- Open-source document format and computational notebook system
- Integrates text, code, and outout
- Can create multiple different types of products (documents, slides, websites, books)



# Why not R Markdown?

Only because quarto is newer and more featured!

- Anything you already know how to do in R Markdown you can do in quarto, and more!
- All of these slides, website, etc. are all made in quarto.
- If you know and love R Markdown, by all means keep using it!

## Quarto workflow

- 1. Create a Quarto document
- 2. Write code
- 3. Write text
- 4. Repeat 2-3 in whatever order you want
- 5. Render repeatedly as you go

#### How does it work?

- You text in markdown and code in R
- knitr processes the code chunks, executes the R code, and inserts the code outputs (e.g., plots, tables) back into the markdown document
- pandoc transforms the markdown document into various output formats



#### Text and code...

```
1 # My header
 3 Some text
 5 Some *italic text*
 6
 7 Some **bold text**
 8
 9 - Eggs
10 - Milk
11
12 \\ \rangle \{r\}
13 x <- 3
```

#### ... becomes ...

#### Mytitle

Some text

Some italic text

Some bold text

- Eggs
- Milk

```
1 x <- 3
2 x
```

# If you prefer, you can use the visual editor

#### My title

Some text

Some *italic text* 

#### Some **bold text**

```
{r}
x <- 3
x
```

#### Rchunks

Everything within the chunks has to be valid R<sup>1</sup>

```
1 ```{r}
2 x <- 3
3 ```

1 ```{r}
2 x + 4
3 ```
```

Chunks run in order, continuously, like a single script

#### YAML

At the top of your Quarto document, a header written in yaml describes options for the document

```
1 ---
2 title: "My document"
3 author: Louisa Smith
4 format: html
5 ---
```

There are a *ton* of possible options, but importantly, this determines the document output

## Output



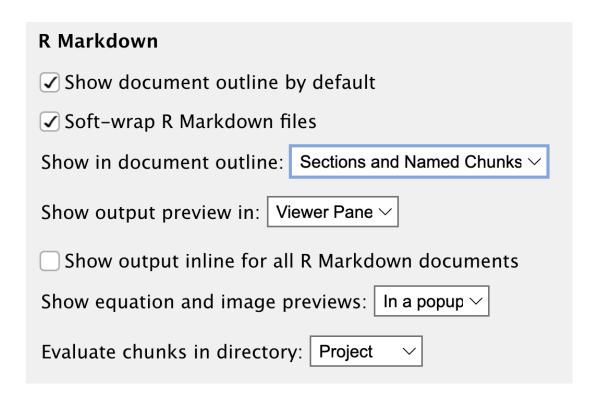
https://quarto.org/docs/output-formats/all-formats.html

We're going to focus on html output

- It's easy to transition to Word (format: docx) but it's not as good for constant re-rendering
- You need a LaTeX installation for pdf (format: pdf)
  - I recommend {tinytex}

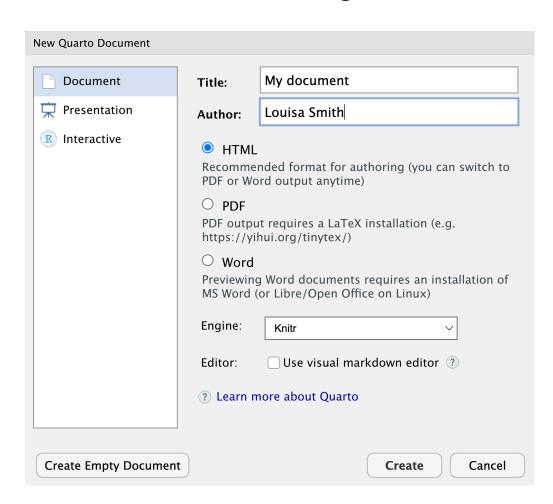
You can choose whether you want to have chunk output show up within the document (vs. just the console) when you are running Quarto (and RMarkdown) documents interactively

#### Tools > Global options

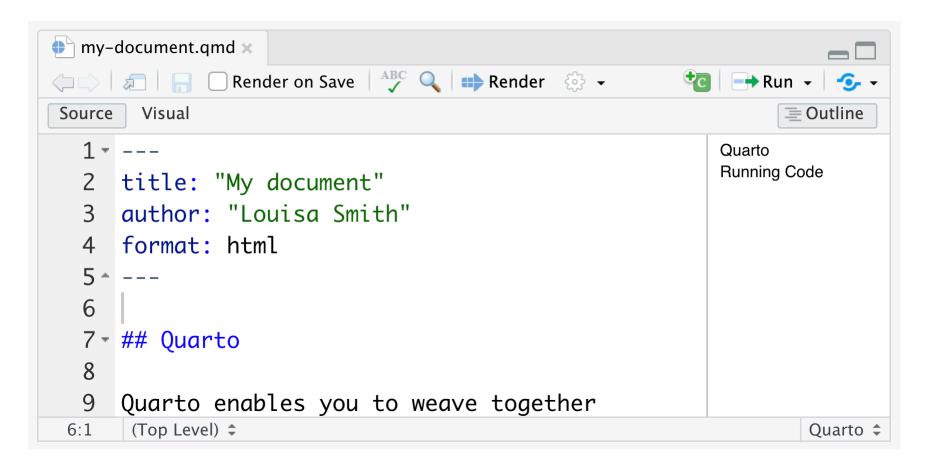


Open up your epi590r-2023-in-class R project!

File > New File > Quarto Document



- Try toggling between Source and Visual views
- Toggle on and off the Outline
- Click Render and look at the output



# Quartooptions

# Chunk options

In your Quarto document, you had a chunk:

```
1 ```{r}
2 #| echo: false
3 2 * 2
4 ```
```

#| echo: false tells knitr not to show the code within that chunk

In RMarkdown, you would have written this  $\{r, echo = FALSE\}$ . You can still do that with Quarto, but it's generally easier to read, particularly for long options (like

# Chunk options

Some of the ones I find myself using most often:

- #| eval: false: Don't evaluate this chunk! Really helpful if you're trying to isolate an error, or have a chunk that takes a long time
- #| error: true: Render this *even if* the chunk causes an error
- #| cache: true: Store the results of this chunk so that it doesn't need to re-run every time, as long as there are no changes
- #| warning: false: Don't print warnings
- #| message: false: Don't print messages

# Document options

You can tell the *entire* document not to evaluate or print code (so just include the text!) at the top:

```
title: "My document"
author: Louisa Smith
format: html
execute:
eval: false
ceho: false
```

Careful! YAML is *really* picky about spacing.

# Document options

There are lots of different options for the document

• For example, you can choose a theme:

```
1 ---
2 format:
3 html:
4 theme: yeti
5 ---
```

 Remember the pickiness: when you have a format option, <a href="https://html:">html:</a> moves to a new line and the options are indented 2 spaces

Download the quarto document with some {gtsummary} tables from yesterday

- Change some options to hide code and output
- Add another code chunk
- Play with themes
- Deal with errors
- Add some text

# Quarto tables, figures, and stats

# Chunks can produce figures and tables

```
1 ```{r}
2 #| label: tbl-one
3 #| tbl-cap: "This is a great table"
4 knitr::kable(mtcars)
5 ```
```

Table 1: This is a great table

	mpg	cyl	disp	hp	drat	wt	qse
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.40
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.07
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.6
Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44

	mpg	cyl	disp	hp	drat	wt	qse
Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02
Valiant	18.1	6	225.0	105	2.76	3.460	20.2
Duster 360	14.3	8	360.0	245	3.21	3.570	15.84
Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00
Merc 230	22.8	4	140.8	95	3.92	3.150	22.90
Merc 280	19.2	6	167.6	123	3.92	3.440	18.30
Merc 280C	17.8	6	167.6	123	3.92	3.440	18.90
Merc 450SE	16.4	8	275.8	180	3.07	4.070	17.4(
Merc 450SL	17.3	8	275.8	180	3.07	3.730	17.60

	mpg	cyl	disp	hp	drat	wt	qse
Merc 450SLC	15.2	8	275.8	180	3.07	3.780	18.00
Cadillac Fleetwood	10.4	8	472.0	205	2.93	5.250	17.98
Lincoln Continental	10.4	8	460.0	215	3.00	5.424	17.8%
Chrysler Imperial	14.7	8	440.0	230	3.23	5.345	17.47
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.4'
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.5%

	mpg	cyl	disp	hp	drat	wt	qse
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.90
Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.0
Dodge Challenger	15.5	8	318.0	150	2.76	3.520	16.8′
AMC Javelin	15.2	8	304.0	150	3.15	3.435	17.3(
Camaro Z28	13.3	8	350.0	245	3.73	3.840	15.4
Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.0

	mpg	cyl	disp	hp	drat	wt	qse
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.5(
Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50
Maserati Bora	15.0	8	301.0	335	3.54	3.570	14.60
Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60

# Chunks can produce figures or tables

```
1 ```{r}
2 #| label: fig-hist
3 #| fig-cap: "This is a histogram"
4 hist(rnorm(100))
5 ```
```

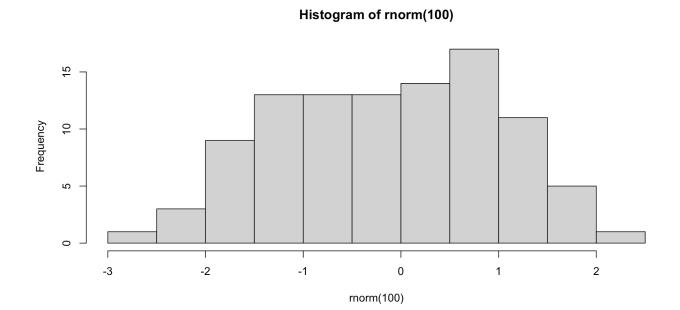


Figure 1: This is a histogram

### Cross-referencing

You can then refer to those with <a href="etable-one">@tbl-one</a> and <a href="etage-hist">@fig-hist</a> and the Table and Figure ordering will be correct (and linked)

@fig-hist contains a histogram and @tbl-one a table.

gets printed as:

Figure 1 contains a histogram and Table 1 a table.

#### Inline R

Along with just regular text, you can also run R code within the text:

```
There were `r 3 + 4` participants
```

becomes:

There were 7 participants

#### Inline R

This is helpful for reporting statistics, e.g. the sample size:

```
There were `r nrow(nlsy)` participants
```

becomes:

There were 1.2686^{4} participants

#### Inline stats

You can also create an object in a chunk and then reference it later in the text

```
1 ```{r}
2 total_sample <- nrow(nlsy)
3 ```
There were `r total_sample` participants</pre>
```

# Inline stats (aside)

I often create a list of stats that I want to report in a manuscript:

```
1 stats <- list(n = nrow(data),
2 mean_age = mean(data$age))
```

I can then print these numbers in the text with:

There were `r stats\$n` participants with a mean age of `r stats\$mean\_age`.

which turns into:

There were 1123 participants with a mean age of 43.5.

# Inline stats from {gtsummary}

We saw very, very briefly yesterday:

```
library(gtsummary)
          2 income table <- tbl uvregression(</pre>
              nlsy,
          4 	 y = income_{r}
          5 include = c(
           sex cat, race eth cat,
                eyesight_cat, income, age_bir
              method = lm
         10)
            inline text(income table, variable = "age bir")
[1] "595 (95% CI 538, 652; p<0.001)"
```

We pulled a statistic from our univariate table

# If we're making a table, we probably want to report numbers from it

```
1 \``{r}
2 # | label: tbl-descr
3 # | tbl-cap: "Descriptive statistics"
4 # output-location: slide
5 table1 <- tbl summary(</pre>
   nlsy,
 7 	 by = sex cat,
   include = c(sex_cat, race_eth_cat, region_cat,
                  eyesight cat, glasses, age bir)) |>
     add overall(last = TRUE)
   table1
12
```

# If we're making a table, we probably want to report numbers from it

Table 2: Descriptive statistics

	Male	Female	Overall
Characteristic	N = 6,403	$N = \frac{6}{1},283$	N = 12,686
race_eth_cat			
Hispanic	1,000 (16%)	1,002 (16%)	2,002 (16%)
Black	1,613 (25%)	1,561 (25%)	3,174 (25%)
Non-Black, Non-Hispanic	3,790 (59%)	3,720 (59%)	7,510 (59%)
region_cat			
Northeast	1,296 (21%)	1,254 (20%)	2,550 (20%)
North Central	1,488 (24%)	1,446 (23%)	2,934 (24%)
South	2,251 (36%)	2,317 (38%)	4,568 (37%)
West	1,253 (20%)	1,142 (19%)	2,395 (19%)
Unknown	115	124	239
eyesight_cat			
1			

	Male	Female	Overall
Characteristic	N = 6,403	N = 6,283	N = 12,686
Excellent	1,582 (38%)	1,334 (31%)	2,916 (35%)
Very good	1,470 (35%)	1,500 (35%)	2,970 (35%)
Good	792 (19%)	1,002 (23%)	1,794 (21%)
Fair	267 (6.4%)	365 (8.5%)	632 (7.5%)
Poor	47 (1.1%)	85 (2.0%)	132 (1.6%)
Unknown	2,245	1,997	4,242
glasses	1,566 (38%)	2,328 (54%)	3,894 (46%)
Unknown	2,241	1,995	4,236
age_bir	25 (21, 29)	22 (19, 27)	23 (20, 28)
Unknown	3,652	3,091	6,743
<sup>1</sup> n (%); Median (Q1, Q3)			

#### I want to report some stats!

The help file for <a href="inline\_text">inline\_text</a>() is helpful and tells us that we can look at <a href="table\_body">table1\$table\_body</a> to help figure out what data to extract.

How about the median (IQR) age of the male participants at the birth of their first child?

```
1 inline_text(table1, variable = "age_bir", column = "stat_]
[1] "25 (21, 29)"
```

# Formatting

We can add sample sizes for the overall stats on people who wear glasses using the <a href="mailto:pattern">pattern</a> = argument:

# Formatting for regression statistics

Remove some details:

```
1 inline_text(income_table, variable = "age_bir")

[1] "595 (95% CI 538, 652; p<0.001)"

1 inline_text(income_table, variable = "age_bir",
2 pattern = "{estimate} ({conf.low}, {conf.high})

[1] "595 (538, 652)"</pre>
```

#### Better yet...

We can integrate these into the text of our manuscript:

```
A greater proportion of female (`r inline_text(table1, variable = "glasses", column = "stat_2")`) than male

(`r inline_text(table1, variable = "glasses", column = "stat_1")`) participants wore glasses.
```

#### Which becomes:

A greater proportion of female (2,328 (54%)) than male (1,566 (38%)) participants wore glasses.

### Readability

Because this can be hard to read, I'd suggest storing those stats in a chunk before the text:

#### Exercises

Return to the quarto document with the tables.

- Choose a table to label and caption, and then write a sentence that cross-references it (e.g., Table 1 shows the descriptive statistics)
- Edit or create a new table that has the median hours of sleep on weekends in it (in the overall sample).
- Pull that value from a table.

# Functions

#### Functions in R

I've been denoting functions with parentheses: func()
We've seen functions such as:

- mean()
- tbl\_summary()
- init()
- create\_github\_token

Functions take **arguments** and return **values** 

# Looking inside a function

If you want to see the code within a function, you can just type its name without the parentheses:

```
1 usethis::create_github_token
```

```
function (scopes = c("repo", "user", "gist", "workflow"), description = "DESCRIBE THE TOKEN'S USE
CASE",
   host = NULL)
   scopes <- glue collapse(scopes, ",")</pre>
   host <- get hosturl(host % | | % default api url())
   url <- glue("{host}/settings/tokens/new?scopes={scopes}&description={description}")</pre>
   withr::defer(view url(url))
   hint <- code hint with host("gitcreds::gitcreds set", host)</pre>
   message <-c() = "Call {.run {hint}} to register this token in the local Git\n
credential store.")
   if (is linux()) {
       message <- c(message, `!` = "On Linux, it can be tricky to store credentials
persistently.",
            i = "Read more in the {.href ['Managing Git(Hub) Credentials' article]
(https://usethis.r-lib.org/articles/articles/git-credentials.html)}.")
   message <- c(message, i = "It is also a great idea to store this token in any\n
password-management software that you use.")
```

```
ui_bullets(message)
   invisible()
}
<bytecode: 0x141ac26a0>
<environment: namespace:usethis>
```

```
1 func <- function()</pre>
```

You can name your function like you do any other object

Just avoid names of existing functions

What objects/values do you need to make your function work?

 You can give them default values to use if the user doesn't specify others

Everything else goes within curly braces

 Code in here will essentially look like any other R code, using any inputs to your functions

- One thing you'll likely want to do is make new objects along the way
- These aren't saved to your environment (i.e., you won't see them in the upper-right window) when you run the function
- You can think of them as being stored in a temporary environment within the function

Return something new that the code has produced

• The return() statement is actually optional. If you don't put it, it will return the last object in the code. When you're starting out, it's safer to always explicitly write out what you want to return.

### Example: a new function for the mean

Let's say we are not satisfied with the mean() function and want to write our own.

Here's the general structure we'll start with.

```
1 new_mean <- function() {
2
3 }</pre>
```

#### New mean: arguments

We'll want to take the mean of a vector of numbers.

It will help to make an example of such a vector to think about what the input might look like, and to test the function. We'll call it  $\mathbf{x}$ :

```
1 \times < -c(1, 3, 5, 7, 9)
```

We can add x as an argument to our function:

```
1 new_mean <- function(x) {
2
3 }</pre>
```

#### New mean: function body

Let's think about how we calculate a mean in math, and then translate it into code:

$$x = \frac{1}{n} \sum_{i=1}^{n} x_i$$

So we need to sum the elements of x together, and then divide by the number of elements.

We can use the functions sum() and length() to help us.

We'll write the code with our test vector first, before inserting it into the function:

```
1 n <- length(x)
2 sum(x) / n
```

### New mean: function body

Our code seems to be doing what we want, so let's insert it. To be explicit, I've stored the answer (within the function) as mean\_val, then returned that value.

```
1 new_mean <- function(x) {
2    n <- length(x)
3    mean_val <- sum(x) / n
4    return(mean_val)
5 }</pre>
```

## Testing a function

Let's plug in the vector that we created to test it:

```
1 new_mean(x = x)
[1] 5
```

And then try another one we create on the spot:

```
1 new_mean(x = c(100, 200, 300))
```

#### Exercises

Create some functions!

Create an R script in your project called **functions.** R to save your work!

# Functions, continued

### Adding another argument

Let's say we plan to be using our <a href="new\_mean">new\_mean</a>() function to calculate proportions (i.e., the mean of a binary variable). Sometimes we'll want to report them as as percentage by multiplying the proportion by 100.

Let's name our new function prop(). We'll use the same structure as we did with new\_mean().

```
1 prop <- function(x) {
2    n <- length(x)
3    mean_val <- sum(x) / n</pre>
```

## Testing the code

Now we'll want to test on a vector of 1's and 0's.

```
1 \times < -c(0, 1, 1)
```

To calculate the proportion and turn it into a percentage, we'll just multiply the mean by 100.

```
1 multiplier <- 100
2 multiplier * sum(x) / length</pre>
```

[1] 66.66667

## Testing the code

We want to give users the option to choose between a proportion and a percentage. So we'll add an argument multiplier. When we want to just return the proportion, we can just set multiplier to be 1.

```
1 multiplier <- 1
         2 multiplier * sum(x) / length
11 0.6666667
         1 multiplier <- 100
         2 multiplier * sum(x) / length
```

#### Adding another argument

If we add multiplier as an argument, we can refer to it in the function body.

```
1 prop <- function(x, multipli
2    n <- length(x)
3    mean_val <- multiplier * s
4    return(mean_val)
5 }</pre>
```

#### Adding another argument

Now we can test:

```
1 prop(x = c(1, 0, 1, 0), multiplier = 1)

[1] 0.5

1 prop(x = c(1, 0, 1, 0), multiplier = 100)
[1] 50
```

## Making a default argument

Since we don't want users to have to specify multiplier = 1 every time they just want a proportion, we can set it as a default.

```
prop <- function(x, multiplier = 1) {
    n <- length(x)
    mean_val <- multiplier * sum(x) / n
    return(mean_val)
}</pre>
```

Now we only need to specify that argument if we want a percentage.

```
1 \text{ prop}(x = c(0, 1, 1, 1))
1 \text{ prop}(x = c(0, 1, 1, 1), \text{ multiplier} = 100)
```

#### Caveats

- This is obviously not the best way to write this function!
- For example, it will still work if x = c(123, 593, -192).... but it certainly won't give you a proportion or a percentage!
- We could also put multiplier = any number, and we'll just be multiplying the answer by that number this is essentially meaningless.
- We also haven't done any checking to see whether the user is even entering numbers! We could put in better error messages so users don't just get an R default error message if they do something wrong.

```
1 prop(x = c("blah", "blah", "blah"))
```

# Functions, continued

# Writing functions to reduce copy and paste

If you find yourself doing something over and over, you can probably write a function for it.

One is example is the regression tables we created yesterday.

# Let's imagine we are fitting multiple similar models

```
1 logistic_model <- glm(glasses ~ eyesight_cat + sex_cat + i
2    data = nlsy, family = binomial()
3 )
4 poisson_model <- glm(nsibs ~ eyesight_cat + sex_cat + incomplete to the sex_cat + incomp
```

# We have a model for a table we want to create

Even though this is already a function, we can *wrap* it in a new function!

#### New table function

We can refer generically to model and then put the model we want a table for as an argument

```
new table function <- function(model) {</pre>
     tbl regression(
       model,
 4 exponentiate = TRUE,
 5 label = list(
         sex cat ~ "Sex",
         eyesight cat ~ "Eyesight",
         income ~ "Income"
10
11 }
```

#### Exercise

- Copy and paste this function into your script
- Create each of the models and run
- Try to figure out how you could allow someone to pass the tidy\_fun argument (from the regression exercises yesterday)

# {renv}

Package management for R

## What is {renv}?

{renv} is an R package for managing project dependencies and creating reproducible environments

# Benefits of using {renv}

- 1. **Isolation:** Creates project-specific environments separate from the global R library.
- 2. **Reproducibility:** Ensures consistent package versions for code reproducibility.
- 3. **Collaboration:** Facilitates sharing and collaborating on projects with others.

## Getting Started with {renv}

1. Install {renv}

```
1 install.packages("renv")
```

2. Initialize a project

```
1 renv::init()
```

3. Install packages

```
install.packages("other_package")

# only an option when using renv!

install.packages("github_user/github_package")
```

4. Track dependencies via a lockfile

```
1 renv::snapshot()
```

#### Behind the scenes

• Your project .Rprofile is updated to include:

```
1 source("renv/activate.R")
```

- This is run every time R starts, and does some management of the library paths to make sure when you call install.packges("package") or library(package) it does to the right place (renv/library/R-{version}/{computer-specifics})
- A renv. lock file (really just a text file) is created to store the names and versions of the packages.

#### renv.lock

```
"R": {
  "Version": "4.3.0",
  "Repositories": [
      "Name": "CRAN",
      "URL": "https://cran.rstudio.com"
"Packages": {
  "R6": {
    "Package": "R6",
    "Version": "2.5.1",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
      "R"
    "Hash": "470851b6d5d0ac559e9d01bb352b4021"
  base64enc": {
    "Package": "base64enc",
    "Version": "0.1-3",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
```

```
"R"
],
"Hash": "543776ae6848fde2f48ff3816d0628bc"
},
```

# Using {renv} later

Restore an environment

```
1 renv::restore()
```

Install new packages

```
1 install.packages("other_package")
```

Update the lockfile

```
1 renv::snapshot()
```

#### Collaboration with {renv}

- Share the project's renv. lock file with collaborators to ensure consistent environments
- When they run <a href="renv::restore">renv::restore</a>(), the correct versions of the packages will be installed on their computer

```
1 renv::restore()
```

#### Other helpful functions

Remove packages that are no longer used:

```
1 renv::clean()
```

Check the status of the project library with respect to the lockfile:

```
1 renv::status()
```

This will tell you to <a href="renv::snapshot">renv::snapshot</a>() to add packages you've installed but haven't snapshotted, or <a href="renv::restore">renv::restore</a>() if you're missing packages you need but which aren't installed

#### Conclusion

{renv} benefits:

Isolation, reproducibility, and collaboration

Getting started with {renv}

- 1. Initialize a project using renv::init()
- 2. Install packages and store with renv::snapshot()
- 3. Restore later or elsewhere with renv::restore()

#### Exercises

- 3. Install a new R package of your choice. (Not sure what to choose? Try one of these fun packages. For example, I did install.packages("hadley/emo").)
- 4. Create an R script and save it in your R project. Include some code that requires the package. For example:

```
1 emo::ji("banana")
```

- 4. Run renv::status() to make sure your project picked up the package as a dependency.
- 5. Run renv::snapshot() to record that package in your lockfile.
- 6. Open your lockfile and look for your new package. For example, mine now has:

```
"emo": {
      "Package": "emo",
      "Version": "0.0.0.9000",
      "Source": "git",
      "RemoteType": "git",
      "RemoteUrl": "https://github.com/hadley/emo",
      "RemoteHost": "api.github.com",
      "RemoteUsername": "hadley",
      "RemoteRepo": "emo",
      "RemoteRef": "master",
      "RemoteSha": "3f03b11491ce3d6fc5601e210927eff73bf8e350",
      "Requirements": [
        "R",
        "assertthat",
        "crayon",
        "glue",
        "lubridate",
        "magrittr",
```

# Reports and manuscripts with Quarto

### What is quarto?

- Open-source document format and computational notebook system
- Integrates text, code, and outout
- Can create multiple different types of products (documents, slides, websites, books)



#### Why not R Markdown?

Only because quarto is newer and more featured!

- Anything you already know how to do in R Markdown you can do in quarto, and more!<sup>1</sup>
- All of these slides, website, etc. are all made in quarto.
- If you know and love R Markdown, by all means keep using it!

3

#### Quarto workflow

- 1. Create a Quarto document
- 2. Write code
- 3. Write text
- 4. Repeat 2-3 in whatever order you want
- 5. Render

#### How does it work?

- You text in markdown and code in R
- knitr processes the code chunks, executes the R code, and inserts the code outputs (e.g., plots, tables) back into the markdown document
- pandoc transforms the markdown document into various output formats



#### Text and code...

```
1 # My header
 3 Some text
 5 Some *italic text*
 6
 7 Some **bold text**
 8
 9 – Eggs
10 - Milk
11
12 ```{r}
13 x <- 3
```

### ... becomes ...

#### Mytitle

Some text

Some italic text

Some bold text

- Eggs
- Milk

```
1 x <- 3
2 x

[1] 3
```

## If you prefer, you can use the visual editor

#### My title

Some text

Some *italic text* 

#### Some **bold text**

```
{r}
x <- 3
x
```

#### Rchunks

Everything within the chunks has to be valid R<sup>1</sup>

```
1 ```{r}
2 x <- 3
3 ```

1 ```{r}
2 x + 4
3 ```
```

Chunks run in order, continuously, like a single script

#### YAML

At the top of your Quarto document, a header written in yaml describes options for the document

```
1 ---
2 title: "My document"
3 author: Louisa Smith
4 format: html
5 ---
```

There are a *ton* of possible options, but importantly, this determines the document output

## Output



https://quarto.org/docs/output-formats/all-formats.html

We're going to focus on html output

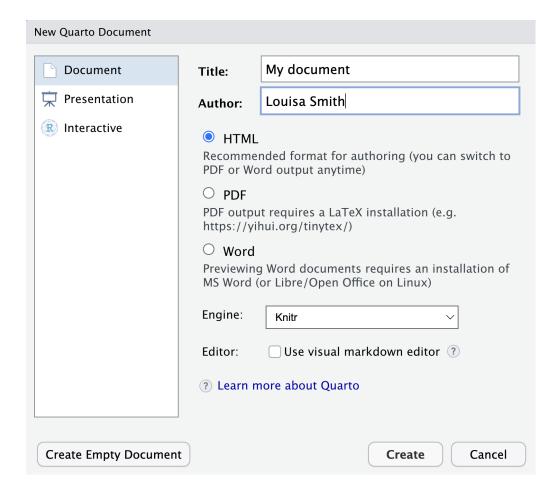
- It's easy to transition to Word (format: docx) but it's not as good for constant re-rendering
- You need a LaTeX installation for pdf
  - I recommend {tinytex}

You can choose whether you want to have chunk output show up within the document (vs. just the console) when you are running Quarto (and RMarkdown) documents interactively

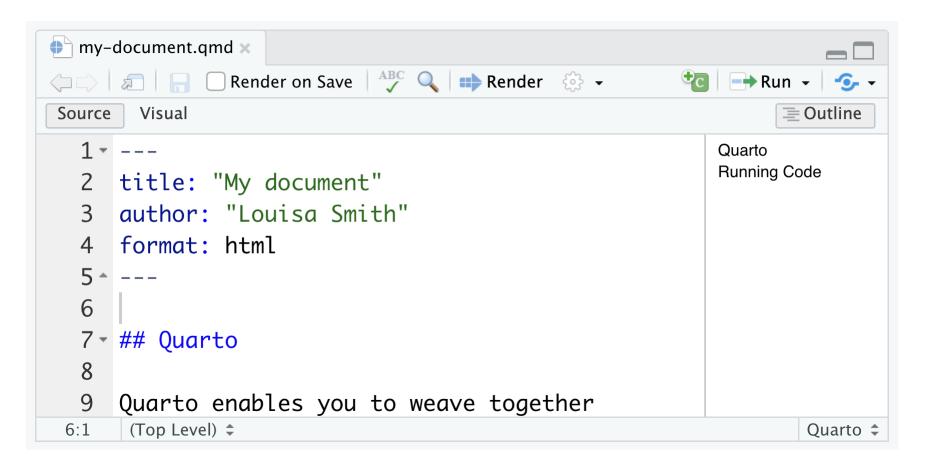
#### Tools > Global options

R Markdown					
✓ Show document outline by default					
✓ Soft-wrap R Markdown files					
Show in document outline: Sections and Named Chunks V					
Show output preview in: Viewer Pane ~					
Show output inline for all R Markdown documents					
Show equation and image previews: $oxed{ In a popup} ee$					
Evaluate chunks in directory: Project					

Open up your epi590r-2023-in-class R project! File > New File > Quarto Document



- Try toggling between Source and Visual views
- Toggle on and off the Outline
- Click Render and look at the output



## Quartooptions

## Chunk options

In your Quarto document, you had a chunk:

```
1 ```{r}
2 #| echo: false
3 2 * 2
4 ```
```

#| echo: false tells knitr not to show the code within that chunk

In RMarkdown, you would have written this  $\{r, echo = FALSE\}$ . You can still do that with Quarto, but it's generally easier to read, particularly for long options (like

## Chunk options

Some of the ones I find myself using most often:

- #| eval: false: Don't evaluate this chunk! Really helpful if you're trying to isolate an error, or have a chunk that takes a long time
- #| error: true: Render this even if the chunk causes an error
- #| cache: true: Store the results of this chunk so that it doesn't need to re-run every time, as long as there are no changes
- #| warning: false: Don't print warnings
- #| message: false: Don't print messages

## Document options

You can tell the *entire* document not to evaluate or print code (so just include the text!) at the top:

```
1 ---
2 title: "My document"
3 author: Louisa Smith
4 format: html
5 execute:
6 eval: false
7 echo: false
8 ---
```

Careful! YAML is *really* picky about spacing.

## Document options

There are lots of different options for the document

• For example, you can choose a theme:

```
1 ---
2 format:
3 html:
4 theme: yeti
5 ---
```

 Remember the pickiness: when you have a format option, <a href="https://html:">html:</a> moves to a new line and the options are indented 2 spaces

Download the quarto document with some {gtsummary} tables from yesterday

- There's an error in the code! Try to render it. Play around with eval: and error: chunk and document options to help you a) find the error and b) render the document despite the error. Then fix the error.
- I don't like the output from the first chunk, where the passages are loaded. Make it so that we don't see this chunk's code or output.
- Play around with themes!

# Quarto tables, figures, and stats

## Chunks can produce figures and tables

```
1 ```{r}
2 #| label: tbl-one
3 #| tbl-cap: "This is a great table"
4 knitr::kable(mtcars)
5 ```
```

Table 1: This is a great table

	mpg	cyl	disp	hp	drat	wt	qse
Mazda RX4	21.0	6	160.0	110	3.90	2.620	16.40
Mazda RX4 Wag	21.0	6	160.0	110	3.90	2.875	17.02
Datsun 710	22.8	4	108.0	93	3.85	2.320	18.6

	mpg	cyl	disp	hp	drat	wt	qse
Hornet 4 Drive	21.4	6	258.0	110	3.08	3.215	19.44
Hornet Sportabout	18.7	8	360.0	175	3.15	3.440	17.02
Valiant	18.1	6	225.0	105	2.76	3.460	20.2
Duster 360	14.3	8	360.0	245	3.21	3.570	15.84
Merc 240D	24.4	4	146.7	62	3.69	3.190	20.00
Merc 230	22.8	4	140.8	95	3.92	3.150	22.90
Merc 280	19.2	6	167.6	123	3.92	3.440	18.30
Merc 280C	17.8	6	167.6	123	3.92	3.440	18.90
Merc 450SE	16.4	8	275.8	180	3.07	4.070	17.4(

	mpg	cyl	disp	hp	drat	wt	qse
Merc 450SL	17.3	8	275.8	180	3.07	3.730	17.60
Merc 450SLC	15.2	8	275.8	180	3.07	3.780	18.00
Cadillac Fleetwood	10.4	8	472.0	205	2.93	5.250	17.98
Lincoln Continental	10.4	8	460.0	215	3.00	5.424	17.8%
Chrysler Imperial	14.7	8	440.0	230	3.23	5.345	17.47
Fiat 128	32.4	4	78.7	66	4.08	2.200	19.4'
Honda Civic	30.4	4	75.7	52	4.93	1.615	18.5%

	mpg	cyl	disp	hp	drat	wt	qse
Toyota Corolla	33.9	4	71.1	65	4.22	1.835	19.9(
Toyota Corona	21.5	4	120.1	97	3.70	2.465	20.0
Dodge Challenger	15.5	8	318.0	150	2.76	3.520	16.8'
AMC Javelin	15.2	8	304.0	150	3.15	3.435	17.3(
Camaro Z28	13.3	8	350.0	245	3.73	3.840	15.4
Pontiac Firebird	19.2	8	400.0	175	3.08	3.845	17.0

	mpg	cyl	disp	hp	drat	wt	qse
Fiat X1-9	27.3	4	79.0	66	4.08	1.935	18.90
Porsche 914-2	26.0	4	120.3	91	4.43	2.140	16.70
Lotus Europa	30.4	4	95.1	113	3.77	1.513	16.90
Ford Pantera L	15.8	8	351.0	264	4.22	3.170	14.5(
Ferrari Dino	19.7	6	145.0	175	3.62	2.770	15.50
Maserati Bora	15.0	8	301.0	335	3.54	3.570	14.60
Volvo 142E	21.4	4	121.0	109	4.11	2.780	18.60

## Chunks can produce figures or tables

```
1 ```{r}
2 #| label: fig-hist
3 #| fig-cap: "This is a histogram"
4 hist(rnorm(100))
5 ```
```

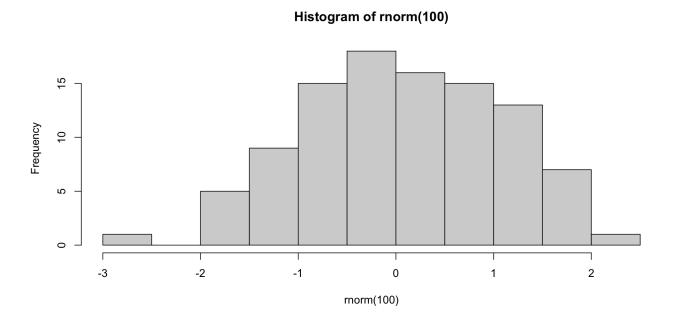


Figure 1: This is a histogram

## Cross-referencing

You can then refer to those with <a href="etable-one">@tbl-one</a> and <a href="etage-hist">@fig-hist</a> and the Table and Figure ordering will be correct (and linked)

@fig-hist contains a histogram and @tbl-one a table.

gets printed as:

Figure 1 contains a histogram and Table 1 a table.

#### Inline R

Along with just regular text, you can also run R code within the text:

```
There were `r 3 + 4` participants
```

becomes:

There were 7 participants

#### Inline stats

I often create a list of stats that I want to report in a manuscript:

I can then print these numbers in the text with:

There were `r stats\$n` participants with a mean age of `r stats\$mean\_age`.

which turns into:

There were 1123 participants with a mean age of 43.5.

## Inline stats from {gtsummary}

We saw very, very briefly yesterday:

```
1 inline_text(income_table, variable = "age_bir")
[1] "595 (95% CI 538, 652; p<0.001)"</pre>
```

We pulled a statistic from our univariate table

## If we're making a table, we probably want to report numbers from it

```
1 \\ (r)
   # | label: tbl-descr
   # tbl-cap: "Descriptive statistics"
   #| output-location: slide
 5 table1 <- tbl summary(</pre>
     nlsy,
     by = sex cat,
    include = c(sex cat, race eth cat, region cat,
                  eyesight cat, glasses, age bir)) |>
     add_overall(last = TRUE)
10
   table1
```

## If we're making a table, we probably want to report numbers from it

Table 2: Descriptive statistics

Characteristic	<b>Male</b> , $N = 6,403^{1}$	<b>Female,</b> $N = 6,283^{\circ}$	<b>Overall</b> , N = 12,686 <sup>1</sup>
race_eth_cat			
Hispanic	1,000 (16%)	1,002 (16%)	2,002 (16%)
Black	1,613 (25%)	1,561 (25%)	3,174 (25%)
Non-Black, Non-Hispanic	3,790 (59%)	3,720 (59%)	7,510 (59%)
region_cat			
Northeast	1,296 (21%)	1,254 (20%)	2,550 (20%)
North Central	1,488 (24%)	1,446 (23%)	2,934 (24%)
South	2,251 (36%)	2,317 (38%)	4,568 (37%)
West	1,253 (20%)	1,142 (19%)	2,395 (19%)
Unknown	115	124	239
eyesight_cat			
Excellent	1,582 (38%)	1,334 (31%)	2,916 (35%)
Very good	1,470 (35%)	1,500 (35%)	2,970 (35%)
<sup>1</sup> n (%); Median (IQR)			

Characteristic	<b>Male</b> , N = $6,403^{1}$	<b>Female,</b> $N = 6,283^{1}$	<b>Overall</b> , N = 12,686 <sup>1</sup>
Good	792 (19%)	1,002 (23%)	1,794 (21%)
Fair	267 (6.4%)	365 (8.5%)	632 (7.5%)
Poor	47 (1.1%)	85 (2.0%)	132 (1.6%)
Unknown	2,245	1,997	4,242
glasses	1,566 (38%)	2,328 (54%)	3,894 (46%)
Unknown	2,241	1,995	4,236
age_bir	25 (21, 29)	22 (19, 27)	23 (20, 28)
Unknown	3,652	3,091	6,743
<sup>1</sup> n (%); Median (IQR)			

## I want to report some stats!

How about the median (IQR) age of the male participants at the birth of their first child?

```
1 inline_text(table1, variable = "age_bir", column = "Male")
[1] "25 (21, 29)"
```

Or the frequency and percentage of women from the South?

```
1 inline_text(table1, variable = "region_cat", level = "South", colu
[1] "2,317 (38%)"
```

And the overall stats on people who wear glasses?

## Better yet...

We can integrate these into the text of our manuscript:

```
A greater proportion of female (`r inline_text(table1, variable = "glasses", column = "Female")`) than male

(`r inline_text(table1, variable = "glasses", column = "Male")`) participants wore glasses.
```

#### Which becomes:

A greater proportion of female (2,328 (54%)) than male (1,566 (38%)) participants wore glasses.

## Readability

Because this can be hard to read, I'd suggest storing those stats in a chunk before the text:

Return to the quarto document with the tables.

- Choose a table to label and caption, and then write a sentence that cross-references it (e.g., Table 1 shows the descriptive statistics)
- From that table, choose at least two statistics to pull out of the table and include in the text using inline\_text().
- Add another statistic to the text that you calculate yourself using the nlsy data, e.g., the mean number of hours of sleep on weekends.

# {renv}

Package management for R

### What is {renv}?

{renv} is an R package for managing project
dependencies and creating reproducible environments

### Benefits of using {renv}

- 1. **Isolation:** Creates project-specific environments separate from the global R library.
- 2. **Reproducibility:** Ensures consistent package versions for code reproducibility.
- 3. **Collaboration:** Facilitates sharing and collaborating on projects with others.

### Getting Started with {renv}

1. Install {renv}

```
1 install.packages("renv")
```

2. Initialize a project

```
1 renv::init()
```

3. Install packages

```
install.packages("other_package")

# only an option when using renv!

install.packages("github_user/github_package")
```

4. Track dependencies via a lockfile

```
1 renv::snapshot()
```

#### Behind the scenes

• Your project .Rprofile is updated to include:

```
1 source("renv/activate.R")
```

- This is run every time R starts, and does some management of the library paths to make sure when you call install.packges("package") or library(package) it does to the right place (renv/library/R-{version}/{computer-specifics})
- A renv. lock file (really just a text file) is created to store the names and versions of the packages.

### renv.lock

```
"R": {
  "Version": "4.3.0",
  "Repositories": [
      "Name": "CRAN",
      "URL": "https://cran.rstudio.com"
"Packages": {
  "R6": {
    "Package": "R6",
    "Version": "2.5.1",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
      "R"
    "Hash": "470851b6d5d0ac559e9d01bb352b4021"
  base64enc": {
    "Package": "base64enc",
    "Version": "0.1-3",
    "Source": "Repository",
    "Repository": "CRAN",
    "Requirements": [
```

```
"R"
],
"Hash": "543776ae6848fde2f48ff3816d0628bc"
},
```

### Using {renv} later

Restore an environment

```
1 renv::restore()
Install new packages
1 install.packages("other_package")
Update the lockfile
1 renv::snapshot()
```

### Collaboration with {renv}

- Share the project's renv. lock file with collaborators to ensure consistent environments
- When they run <a href="renv::restore">renv::restore</a>(), the correct versions of the packages will be installed on their computer

```
1 renv::restore()
```

### Other helpful functions

Remove packages that are no longer used:

```
1 renv::clean()
```

Check the status of the project library with respect to the lockfile:

```
1 renv::status()
```

This will tell you to <a href="renv::snapshot">renv::snapshot</a>() to add packages you've installed but haven't snapshotted, or <a href="renv::restore">renv::restore</a>() if you're missing packages you need but which aren't installed

#### Conclusion

{renv} benefits:

Isolation, reproducibility, and collaboration

Getting started with {renv}

- 1. Initialize a project using renv::init()
- 2. Install packages and store with renv::snapshot()
- 3. Restore later or elsewhere with renv::restore()

#### Exercises

- 3. Install a new R package of your choice. (Not sure what to choose? Try one of these fun packages. For example, I did install.packages("hadley/emo").)
- 4. Create an R script and save it in your R project. Include some code that requires the package. For example:

```
1 emo::ji("banana")
```

- 4. Run renv::status() to make sure your project picked up the package as a dependency.
- 5. Run renv::snapshot() to record that package in your lockfile.
- 6. Open your lockfile and look for your new package. For example, mine now has:

```
"emo": {
      "Package": "emo",
      "Version": "0.0.0.9000",
      "Source": "git",
      "RemoteType": "git",
      "RemoteUrl": "https://github.com/hadley/emo",
      "RemoteHost": "api.github.com",
      "RemoteUsername": "hadley",
      "RemoteRepo": "emo",
      "RemoteRef": "master",
      "RemoteSha": "3f03b11491ce3d6fc5601e210927eff73bf8e350",
      "Requirements": [
        "R",
        "assertthat",
        "crayon",
        "glue",
        "lubridate",
        "magrittr",
```

## Functions

#### Functions in R

I've been denoting functions with parentheses: func()
We've seen functions such as:

- mean()
- tbl\_summary()
- init()
- create\_github\_token

Functions take **arguments** and return **values** 

### Looking inside a function

If you want to see the code within a function, you can just type its name without the parentheses:

```
1 usethis::create github token
function (scopes = c("repo", "user", "gist", "workflow"), description = "DESCRIBE THE TOKEN'S USE
CASE",
   host = NULL)
   scopes <- glue collapse(scopes, ",")</pre>
   host <- get hosturl(host %||% default api url())</pre>
   url <- glue("{host}/settings/tokens/new?scopes={scopes}&description={description}")</pre>
   withr::defer(view url(url))
   hint <- code hint with host("gitcreds::gitcreds set", host)</pre>
   ui todo("\n Call {ui code(hint)} to register this token in the \\\n local Git
software that you use")
   invisible()
<br/>
<br/>
bytecode: 0x104c98aa0>
<environment: namespace:usethis>
```

```
1 func <- function()</pre>
```

You can name your function like you do any other object

Just avoid names of existing functions

What objects/values do you need to make your function work?

 You can give them default values to use if the user doesn't specify others

Everything else goes within curly braces

 Code in here will essentially look like any other R code, using any inputs to your functions

- One thing you'll likely want to do is make new objects along the way
- These aren't saved to your environment (i.e., you won't see them in the upper-right window) when you run the function
- You can think of them as being stored in a temporary environment within the function

Return something new that the code has produced

• The return() statement is actually optional. If you don't put it, it will return the last object in the code. When you're starting out, it's safer to always explicitly write out what you want to return.

### Example: a new function for the mean

Let's say we are not satisfied with the mean() function and want to write our own.

Here's the general structure we'll start with.

```
1 new_mean <- function() {
2
3 }</pre>
```

### New mean: arguments

We'll want to take the mean of a vector of numbers.

It will help to make an example of such a vector to think about what the input might look like, and to test the function. We'll call it  $\mathbf{x}$ :

```
1 \times < -c(1, 3, 5, 7, 9)
```

We can add x as an argument to our function:

```
1 new_mean <- function(x) {
2
3 }</pre>
```

### New mean: function body

Let's think about how we calculate a mean in math, and then translate it into code:

$$x = \frac{1}{n} \sum_{i=1}^{n} x_i$$

So we need to sum the elements of x together, and then divide by the number of elements.

We can use the functions sum() and length() to help us.

We'll write the code with our test vector first, before inserting it into the function:

```
1 n <- length(x)
2 sum(x) / n</pre>
```

### New mean: function body

Our code seems to be doing what we want, so let's insert it. To be explicit, I've stored the answer (within the function) as mean\_val, then returned that value.

```
1 new_mean <- function(x) {
2   n <- length(x)
3   mean_val <- sum(x) / n
4   return(mean_val)
5 }</pre>
```

### Testing a function

Let's plug in the vector that we created to test it:

```
1 \quad \text{new\_mean}(x = x)
[1] 5
```

And then try another one we create on the spot:

```
1 new_mean(x = c(100, 200, 300))
[1] 200
```

### Adding another argument

Let's say we plan to be using our <a href="new\_mean">new\_mean</a>() function to calculate proportions (i.e., the mean of a binary variable). Sometimes we'll want to report them as as percentage by multiplying the proportion by 100.

Let's name our new function prop(). We'll use the same structure as we did with new\_mean().

```
1 prop <- function(x) {
2   n <- length(x)
3  mean_val <- sum(x) / n</pre>
```

### Testing the code

Now we'll want to test on a vector of 1's and 0's.

```
1 \times < -c(0, 1, 1)
```

To calculate the proportion and turn it into a percentage, we'll just multiply the mean by 100.

```
1 multiplier <- 100
2 multiplier * sum(x) / length(x)
[1] 66.66667</pre>
```

### Testing the code

We want to give users the option to choose between a proportion and a percentage. So we'll add an argument multiplier. When we want to just return the proportion, we can just set multiplier to be 1.

```
multiplier <- 1
  multiplier * sum(x) / length(x)
11 0.6666667
  multiplier <- 100
  multiplier * sum(x) / length(x)
```

### Adding another argument

If we add multiplier as an argument, we can refer to it in the function body.

```
1 prop <- function(x, multiplier) {
2   n <- length(x)
3   mean_val <- multiplier * sum(x) / :
4   return(mean_val)
5 }</pre>
```

### Adding another argument

Now we can test:

```
1 prop(x = c(1, 0, 1, 0), multiplier = 1)

[1] 0.5

1 prop(x = c(1, 0, 1, 0), multiplier = 100)

[1] 50
```

### Making a default argument

Since we don't want users to have to specify multiplier = 1 every time they just want a proportion, we can set it as a default.

```
1 prop <- function(x, multiplier = 1) {
2   n <- length(x)
3   mean_val <- multiplier * sum(x) / n
4   return(mean_val)
5 }</pre>
```

Now we only need to specify that argument if we want a percentage.

```
1 prop(x = c(0, 1, 1, 1))

[1] 0.75

1 prop(x = c(0, 1, 1, 1), multiplier = 100)
```

#### Caveats

- This is obviously not the best way to write this function!
- For example, it will still work if x = c(123, 593, -192).... but it certainly won't give you a proportion or a percentage!
- We could also put multiplier = any number, and we'll just be multiplying the answer by that number this is essentially meaningless.
- We also haven't done any checking to see whether the user is even entering numbers! We could put in better error messages so users don't just get an R default error message if they do something wrong.

```
1 prop(x = c("blah", "blah", "blah"))
```

#### Exercises

Create some functions!

Create an R script in your project called **functions.** R to save your work!

# {targets}

### What is {targets}?



"a Make-like pipeline tool for statistics and data science in R"

- manage a sequence of computational steps
- only update what needs updating
- ensure that the results at the end of the pipeline are still valid

#### Script-based workflow

#### 01-data.R

```
1 library(tidyverse)
2 data <- read_csv("data.csv", col_types = cols()) %>%
3     filter(!is.na(Ozone))
4 write_rds(data, "data.rds")
```

#### 02-model.R

```
1 library(tidyverse)
2 data <- read_rds("data.rds")
3 model <- lm(Ozone ~ Temp, data) %>%
4    coefficients()
5 write_rds(model, "model.rds")
```

#### 03-plot.R

```
1 library(tidyverse)
2 model <- read_rds("model.rds")
3 data <- read_rds("data.rds")
4 ggplot(data) +
5     geom_point(aes(x = Temp, y = Ozone)) +
6     geom_abline(intercept = model[1], slope = model[2])
7 ggsave("plot.png", plot)</pre>
```

#### Problems with script-based workflow

- Reproducibility: if you change something in one script, you have to remember to re-run the scripts that depend on it
- Efficiency: that means you'll usually rerun all the scripts even if they don't depend on the change
- Scalability: if you have a lot of scripts, it's hard to keep track of which ones depend on which
- File management: you have to keep track of which files are inputs and which are outputs and where they're saved

### {targets} workflow

#### R/functions.R

```
get data <- function(file) {</pre>
      read csv(file, col types = cols()) %>%
 3
        filter(!is.na(Ozone))
 5
   fit model <- function(data) {</pre>
      lm(Ozone ~ Temp, data) %>%
        coefficients()
10
   plot model <- function(model, data) {</pre>
12
      ggplot(data) +
13
        geom point(aes(x = Temp, y = Ozone)) +
```

### {targets} workflow

#### \_targets.R

```
library(targets)
 3 tar source()
   tar option set(packages = c("tidyverse"))
 5
   list(
     tar target(file, "data.csv", format = "file"),
     tar target(data, get data(file)),
     tar target(model, fit model(data)),
10
     tar target(plot, plot model(model, data))
|11|
```

Run tar\_make() to run pipeline

#### {targets} workflow

Targets are "hidden" away where you don't need to manage them

```
— _targets.R

— data.csv

— R/

| — functions.R

— _targets/

| — objects

| — data

| — model

| — plot
```

## My typical workflow with {targets}

- 1. Read in some data and do some cleaning until it's in the form I want to work with.
- 2. Wrap that in a function and save the file in R/.
- 3. Run use\_targets() and edit \_targets.R accordingly, so that I list the data file as a target and clean\_data as the output of the cleaning function.
- 4. Run tar\_make().
- 5. Run tar\_load(clean\_data) so that I can work on the next step of my workflow.
- 6. Add the next function and corresponding target when I've solidified that step.

#### \_targets.R tips and tricks

```
list(
      tar_target(
        data file,
 3
        "data/raw data.csv",
        format = "file"
 5
 6
      ),
      tar target(
        raw_data,
 8
        read.csv(data file)
10
11
     tar_target(
12
        clean_data,
13
        clean_data_function(raw_data)
```

#### targets.R tips and tricks

```
preparation <- list(</pre>
      tar_target(
 3
        clean_data,
 5
        clean data function(raw data)
 6
   modeling <- list(</pre>
      tar target(
        linear model,
10
11
        linear_model_function(clean_data)
12
13
```

By grouping the targets into lists, I can easily comment out chunks of the pipeline

#### \_targets.R tips and tricks

```
1 ## prepare ----
   prepare <- list(</pre>
 3
      ### cleanData.csv ----
      tar target(
 5
        cleanData.csv,
        file.path(path to data,
 6
                   "cleanData.csv"),
        format = "file"
 8
10
      ### newdat ----
11
      tar target(
12
        newdat,
13
        read csv(cleanData.csv,
```

```
prepare
                          raw
cleanData.csv
                           log_res_raw
newdat
                           mult res raw
fulldat
                           ctc res raw
flow
                           tte res raw
loglinear
                           tte res raw very
                           all res raw
log_dat
log_dat_covid_mild...
                          imputation
log_dat_covid
                           mice params
                           cols_to_impute
log_est
log est severity
                           fulldat_imputed
loglinear_spon
                           log_dat_imputed
                           mice_comparison
log est spon
log_est_severity_spon
                          tabs figs
log est ind
                           descriptive table
log est severity ind
                           descriptive_table_o...
                           outcomes table
mult est
mult_est_severity
                           risk_plot
ctc
                           tte_plot
                           followup_tab
ctc dat
cases
                           testing_plot
                          all targets
controls
```

### Key {targets} functions

- use\_targets() gets you started with a \_targets.R script to fill in
- tar\_make() runs the pipeline and saves the results in \_targets/objects/
- tar\_make\_future() runs the pipeline in parallel<sup>1</sup>
- tar\_load() loads the results of a target into the global environment (e.g., tar\_load(clean\_data))
- tar\_read() reads the results of a target into the global environment (e.g., dat <- tar\_read(clean\_data))</li>
- tar\_visnetwork() creates a network diagram of the pipeline
- tar\_outdated() checks which targets need to be updated
- tar\_prune() deletes targets that are no longer in \_targets.R
- tar\_destroy() deletes the .targets/ directory if you need to burn everything down and start again

# Advanced {targets}

#### "target factories"



repo status Active <u>tarchetypes</u> makes it easy to add certain kinds of common tasks to reproducible pipelines. Most of its <u>functions</u> create families of targets for <u>parameterized R Markdown</u>, <u>simulation studies</u>, and other general-purpose scenarios.



repo status Active stantargets is a workflow framework for Bayesian data analysis with cmdstanr. With concise, easy-to-use syntax, it defines versatile families of targets tailored to Bayesian statistics, from a single MCMC run with postprocessing to large simulation studies.



repo status Active Like stantargets, jagstargets is a workflow framework for Bayesian data analysis, with support for both single MCMC runs and large-scale simulation studies. It invokes <u>JAGS</u> through the <u>R2jags</u> package, which has nice features such as the ability to parallelize chains across local R processes.

#### {tarchetypes}: reports

Render documents that depend on targets loaded with tar\_load() or tar\_read().

- tar\_render() renders an R Markdown document
- tar\_quarto() renders a Quarto document (or project)

#### What does report. qmd look like?

```
2 title: "My report"
   ```{r}
 5 library(targets)
 6 tar load(results)
 7 tar load(plots)
   There were `r results$n` observations with a mean age of `r result
   ```{r}
|11 library(ggplot2)
12 plots$age plot
13
```

Because report qmd depends on results and plots, it will only be re-rendered if either of those targets change.

## {tarchetypes}: branching

Using data from the National Longitudinal Survey of Youth,

#### \_targets.R

```
1 library(targets)
 2 library(tarchetypes)
   tar source()
   targets setup <- list(</pre>
      tar target(
        CSV,
        "data/nlsy.csv",
        format = "file"
10
11
      tar target(
12
        dat,
13
        readr::read csv(csv,
14
          show col types = FALSE)
15
16)
```

#### R/functions.R

we want to investigate the relationship between age at first birth and hours of sleep on weekdays and weekends among moms and dads separately

#### Option 1

Create (and name) a separate target for each combination of sleep variable ("sleep\_wkdy", "sleep\_wknd") and sex (male: 1, female: 2):

```
targets 1 <- list(</pre>
     tar target(
 3
       model 1,
       model function(outcome var = "sleep wkdy", sex val = 1, dat =
     tar target(
       coef 1,
 8
       coef function(model 1)
10
```

... and so on...

#### Option 2

Use tarchetypes::tar\_map() to map over the combinations for you (static branching):

```
targets 2 <- tar map(</pre>
     values = tidyr::crossing(
        outcome = c("sleep wkdy", "sleep wknd"),
 3
       sex = 1:2
     tar target(
       model 2,
 8
       model function(outcome var = outcome, sex val = sex, dat = dat
     tar_target(
10
111
       coef 2,
12
       coef function(model 2)
13
```

#### Option 2, cont.

Use tarchetypes::tar\_combine() to combine the results of a call to tar\_map():

```
combined <- tar_combine(
combined_coefs_2,
  targets_2[["coef_2"]],
  command = vctrs::vec_c(!!!.x),

tar_read(combined_coefs_2)</pre>
```

command = vctrs::vec\_c(!!!.x) is the default, but you can supply your own function to combine the results

#### Option 3

Use the pattern = argument of tar\_target() (dynamic branching):

```
1 targets 3 <- list(</pre>
 2
     tar target(
       outcome_target,
 3
       c("sleep wkdy", "sleep wknd")
 5
     tar_target(
 6
       sex target,
      1:2
 8
     tar_target(
10
       model 3,
11
       model_function(outcome_var = outcome_target, sex_val = sex_tar
12
13
       pattern = cross(outcome target, sex target)
```

# Branching

Dynamic	Static
Pipeline creates new targets at runtime.	All targets defined in advance.
Cryptic target names.	Friendly target names.
Scales to hundreds of branches.	Does not scale as easily for tar_visnetwork() etc.
No metaprogramming required.	Familiarity with metaprogramming is helpful.

#### Branching

- The book also has an example of using metaprogramming to map over different functions
  - i.e. fit multiple models with the same arguments
- Static and dynamic branching can be combined
  - e.g. tar\_map(values = ..., tar\_target(..., pattern
    = map(...)))
- Branching can lead to slowdowns in the pipeline (see book for suggestions)

#### {tarchetypes}: repetition

tar\_rep() repeats a target multiple times with the same arguments

```
targets 4 <- list(</pre>
     tar_rep(
       bootstrap coefs,
       dat >
          dplyr::slice sample(prop = 1, replace = TRUE) |>
         model_function(outcome_var = "sleep_wkdy", sex_val = 1, dat
 6
         coef function(),
      batches = 10,
       reps = 10
10
11 )
```

The pipeline gets split into batches x reps chunks, each with its own random seed

# {tarchetypes}: mapping over iterations

```
1 sensitivity_scenarios <- tibble::tibble(
2 error = c("small", "medium", "large"),
3 mean = c(1, 2, 3),
4 sd = c(0.5, 0.75, 1)
5 )</pre>
```

tar\_map\_rep() repeats a target multiple times with different arguments

```
1 targets_5 <- tar_map_rep(
2    sensitivity_analysis,
3    dat |>
4         dplyr::mutate(sleep_wkdy = sleep_wkdy + rnorm(nrow(dat), mean,
5         model_function(outcome_var = "sleep_wkdy", sex_val = 1, dat =
6         coef_function() |>
7         data.frame(coef = _),
8         values = sensitivity_scenarios,
```

```
9 batches = 10,
10 reps = 10
11 )
```

# {tarchetypes}: mapping over iterations

```
1 tar_read(sensitivity_analysis) |> head()
```

Ideal for sensitivity analyses that require multiple iterations of the same pipeline with different parameters

#### Summary

- {targets} is a great tool for managing complex workflows
- {tarchetypes} makes it even more powerful
- The user manual is a great resource for learning more

#### Exercises

We'll clone a repo with {targets} already set up and add some additional steps to the analysis.