

### Some disclamers

My credentials

Reproducibility { rant time }

- Note: this talk
  - will be (mostly) **R**-based
  - inspired by a couple of talks from this lady

Advertisement time: join the <u>EMC coding café</u>

# POV: you look at code you wrote last year





## But enough about me...

You need to repeat an operation a few times (3-10). Chose a character:

- Copy-paste athlete
- □ For loop master
- Apply family mobster
- Vector-ciraptor

How confident are you writing a function:

- A what now?
- I can try if you really need me to
- definitely
- ☐ My functions write their own functions, b\*tch

code **7217 1669** 



# The holy trinity of code quality

stability

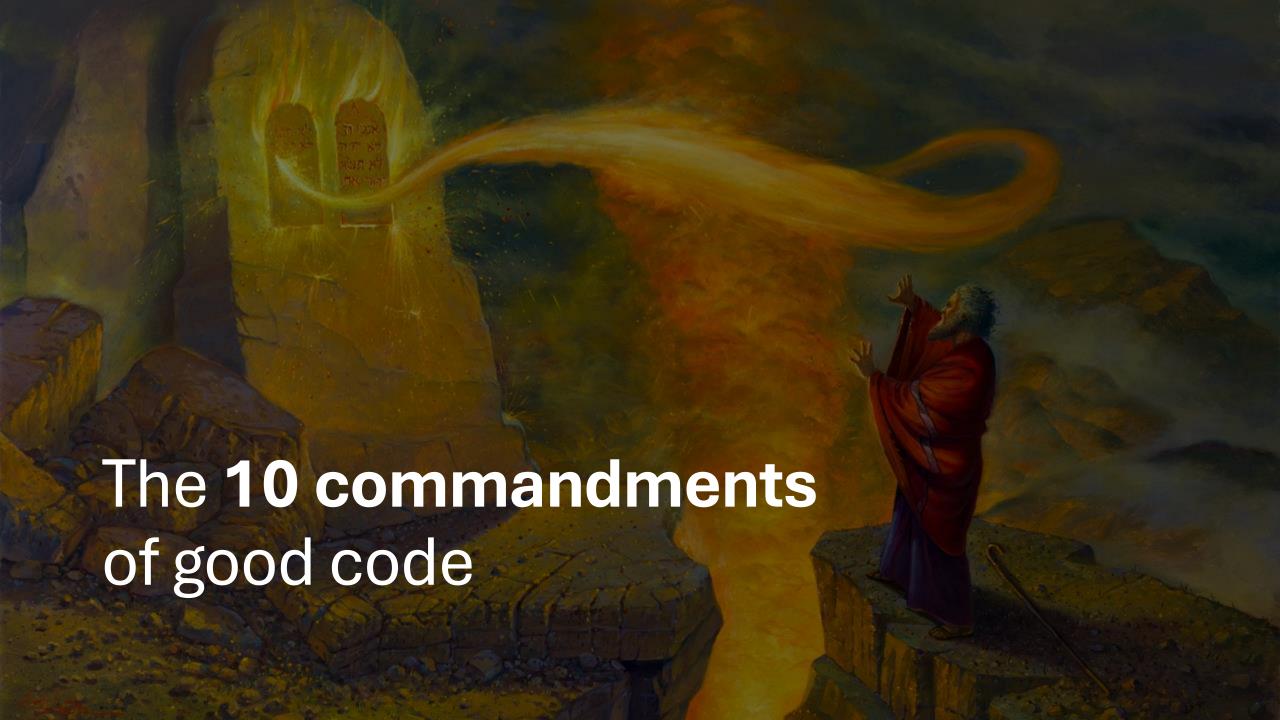
aka reproducible

flexibility

aka cheap to modify

readability

aka easy to understand



## **DO NOT REPEAT YOURSELF (DRY)**

#### Don't do this

```
# Welcome to my script

## [some code]

# Please input the path to save some output
write.csv(some_output, 'your/user/path/output1.csv')

## [some more code]

write.csv(some_other_output, 'your/user/path/output10.csv')
```

#### a. use **variables**

```
# Welcome to my script
# Please input the path where to save the output
user_path <- 'your/user/path'

## [some code]

write.csv(some_output, file.path(user_path, 'output1.csv'))

## [some more code]

write.csv(some_other_output, file.path(user_path, 'output10.csv'))</pre>
```

# 1 DRY level: beginner

#### Don't do this

```
# e.g. fitting linear models
fit1 <- lm(out1 ~ exp1 + cov1, data = data)
# [some more code]

fit2 <- lm(out1 ~ exp2 + cov1, data = data)
# [some more code]

fit3 <- lm(out1 ~ exp3 + cov1, data = data)
# [some more code]</pre>
```

### b. use a for loop

```
fits <- list()

for (exp in c('exp1','exp2','exp3')) {
    # Construct formula as a string
    my_formula <- paste('out1 ~', exp, '+ cov1')
    # Save linear model output inside a list
    fits[[paste0('fit_', exp)]] <- lm(as.formula(my_formula), data=data)
}

# Access model information (e.g. coefficients)
coef(fits$fit_exp1) # or fits[['fit_exp1']] or fits[1]]</pre>
```

# 1 DRY level: intermediate

#### Don't do this

```
# e.g. fitting linear models
fit1 <- lm(out1 ~ exp1 + cov1, data = data)
# [some more code]

fit2 <- lm(out1 ~ exp2 + cov1, data = data)
# [some more code]

fit3 <- lm(out1 ~ exp3 + cov1, data = data)
# [some more code]</pre>
```

Unlock **vectorization**! →

#### c. use a **function**

```
fit_model <- function(exp) {</pre>
  # Construct formula as a string
  my_formula <- paste('out1 ~', exp, '+ cov1')</pre>
  # Fit the linear model
  fit <- lm(as.formula(my_formula), data=data)</pre>
  # [some more code]
  return(fit)
# Fit the model with each exposure
fits <- lapply(c('exp1','exp2','exp3'), fit_model)</pre>
fits <- list()
for (exp in c('exp1', 'exp2', 'exp3')) {
  fits[[paste0('fit_', exp)]] <- fit_model(exp)
```

readability

flexibility

# 1 DRY level: advanced

#### Don't do this

```
# e.g. fitting linear models
fit1 \leftarrow lm(out1 \sim exp1 + cov1, data = data)
# [some more code]
fit2 \leftarrow lm(out1 \sim exp2 + cov1, data = data)
# [some more code]
fit3 \leftarrow lm(out1 \sim exp3 + cov1, data = data)
# [some more code]
fit4 \leftarrow lm(out2 \sim exp1 + cov1, data = data)
# [some more code]
fit5 < lm(out2 \sim exp2 + cov1, data = data)
# [some more code]
```

### d. add *arguments* to your **function**

```
fit_model <- function(exp, out) {
   my_formula <- paste(out, '~', exp, '+ cov1')
   fit <- lm(as.formula(my_formula), data=data)
}

fits <- list()
for (out in c('out1','out2')) {
   for (exp in c('exp1','exp2','exp3')) {
     fits[[paste0('fit_', out, exp)]] <- fit_model(exp, out)
   }
}</pre>
```

# 1 DRY level: brat

#### Don't do this

```
# e.g. fitting linear models
fit1 \leftarrow lm(out1 \sim exp1 + cov1, data = data)
# [some more code]
fit2 \leftarrow lm(out1 \sim exp2 + cov1, data = data)
# [some more code]
fit3 \leftarrow lm(out1 \sim exp3 + cov1, data = data)
# [some more code]
fit4 \leftarrow lm(out2 \sim exp1 + cov1, data = data)
# [some more code]
fit5 < lm(out2 \sim exp2 + cov1, data = data)
# [some more code]
```

```
fit_model <- function(exp, out, cov = '+ cov1') {</pre>
  my_formula <- paste(out, '~', exp, cov)</pre>
 fit <- lm(as.formula(my_formula), data=data)</pre>
  print(summary(fit))
 NULL
# Define all possible combinations of arguments
exps <- c('exp1','exp2','exp3')
outs <- c('out1','out2')</pre>
covs <- c('+ cov1', '+ cov1 + cov2')
all_models <- expand.grid(exp = exps, out = outs, cov = covs)
# Apply the model
do.call(mapply, c(fit_model, all_models))
# OR
# use purrr::pmap()
```

readability

flexibility

# comment. your. code.

Explain yourself, what seems obvious now may become confusing once you forgot the context (...for me: pretty darn soon)

Don't expect anyone to read the comments

Good code should "read like English"

---> Pick the right **names** your functions and variables

--- (in R) use **piping** ( %>% , > )

"there are only two hard things in Computer Science: cache invalidation and naming things."

The code should explain what, the comment should explain why

### 4

## Use "chapters" and "paragraphs"

Avoid very long scripts (with 500+ lines)

- ---> Divide your pipeline into "chapter" **files**
- ... number them (when it makes sense)
- ---> *Tip*: access variables or functions defined inside other files:

```
source("0-functions.R")
```

from definitions.backend import \*

<u>Divide</u> your script into **sections** using

```
# [section title] ------
```

- 0-Data\_prep.R
  1-Descriptives.R
  - 2-Main\_analyses.R
  - 🔓 3-Regr-diagnostics.R
  - 4-Figures.ipynb
  - README.md
  - run.slurm.0

A few small functions > one monster function

readability

flexibility

### You have to break it (before you can fix it)

Repeat after me: errors should never pass silently.

In the face of ambiguity (which will probably be there), don't guess

---> Check your **user input** (...often: classes and types)

```
if (is.numeric(variable)) {
    mean(variable)
} else if (is.factor(variable)) {
    warning("Factor means are not meaningful, how about frequencies.")
    table(variable) |
} else {
    stop("You need to input either a numeric variable or a factor.")
}
```

Don't comment & uncomment code, especially if you need to do it in multiple places!

---> *Throw* **errors** and/or *catch* them using tryCatch()

readability flexibility

### **BUT...** beware of **onion functions**

```
get_some_data <- function(config, outfile) {</pre>
 if (config_ok(config)) {
    if (can_write(outfile)) {
      if (can_open_network_connection(config)) {
        data <- parse_something_from_network()</pre>
        if(makes_sense(data)) {
          data <- beautify(data)
          write_it(data, outfile)
          return(TRUE)
        } else {
          return(FALSE)
      } else {
        stop("Can't access network")
    } else {
  } else {
```

### 6

### Flat is better than nested

Not all **if**s need an else...

```
Use quick stop() and return() ("guard clauses")
e.g. stopifnot(is.numeric(x) || is.logical(x))
```

Less indentation > more indentation.

Use control-flow alternatives:

```
library(dplyr)

tibble(
    age_yrs = c(0, 4, 15, 24, 55),
    age_cat = case_when(
        age_yrs < 2 ~ "baby",
        age_yrs < 13 ~ "kid",
        age_yrs < 20 ~ "teen",
        TRUE ~ "adult"
)
)</pre>
```

### **Code independence**

<u>Avoid</u> adding dependencies

(unless you *really* need them)

Packages:

are living things, they **change** all the time **depend** on other packages, which

**Good practice::contextualize functions!** 

```
# Read file
foreign::read.spss(my_spss_file)
```

#### Don't do this

```
# Upload packages
library('foreign')
library('table1')
library('tidyverse')
library('dplyr')
library('gt')
library('plyr')
library('car')
library('nnet')
library('pscl')
library('broom')
library('xtable')
library('car')
library('MASS')
library('emmeans')
library('mice')
```

## Version control (a): use environments

A programming environment = "the infrastructure (including the compiler and the tools) where the code can run" ...and, sometimes, the OS

Basically: the **version** of R and the packages

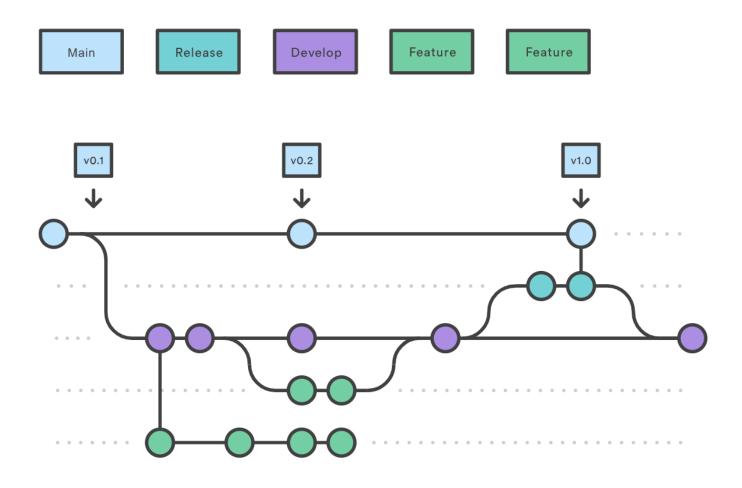
Environment managers: conda, venv, pyenv

From R:

```
# 1. Activate the environment
renv::activate()
# 2. Create a "lockfile": a "snapshot" of the current project
renv::snapshot()
# 3. Reproduce it any place else
renv::restore()
```

flexibility readability stability 9

## Version control (b): use git



[note] Upcoming talk:

# Introduction to git and github

PhD meeting 5 Feb 2024,14:00

readability

flexibility

## Version control (a & b): use projects

9 I often see scripts starting like this:

setwd('path/to/project/folder/')

Use a **project** instead...

**RStudio** makes this very easy! [demo]



Once you create a project, it's <u>much easier</u> to manage your files, your dependencies (with renv), track code versions (with git) and (eventually) give it somebody else.

# 10 Peer review, baby 😇



- As an author, ask people to review your code!
  - ---> Best medicine against *code blindness*
  - ---> If you can (and you can) have at least one co-author review the code

As a reviewer, ask to see the code: that's the real scientific product

### In summary...

- 1 Don't repeat yourself: use variables, functions and loops
- **2** #comment your code
- Pick names so you can read\_code\_like("English")
- 4 Keep your files (and functions) short & organised
- 5 Throw and catch errors
- <u>Avoid</u> (a lot of) **indentation**
- 7 Avoid package dependencies
- 8 Use **environments**
- 9 Use git

10 Peer review the code

control those versions

### **Bonus:** some extra **don't**s

- X forget to set.seed()
- X refer to columns in a data frame using their numbers
- X make functions dependent on global enviroment variables

# Thanks for listening!

(happy\_coding\_everyone;)