

# Introduction to Programming with R

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# Agenda

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# Agenda

## Day 1

- RStudio setup
- Basic elements & data types of the R language
- Flow & conditional programming
- Loops & iteration
- Writing & using functions (part I)

## Day 2

- Writing & using functions (part II)
- Debugging
- Good programming practices

Open questions from day 1?

## Functions II

---

## Why write functions?

- They make code ...
  - shorter (less repetition)
  - easier to read and understand
- They help avoid copy-paste errors
- They make it easier to change your code
- They increase transferability to ...
  - other use cases
  - other projects
  - other persons
- They keep your work space clean

Writing a function:

```
RMSE <- get_RMSE(predictions, observations)
```

Not writing a function:

```
diff <- observations - predictions  
sq_diff <- diff^2  
m_sq_diff <- mean(diff)  
RMSE <- sqrt(m_sq_diff)
```



Writing a function:

```
summary(mtcars$mpg)
```

```
>      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
>   10.40   15.43   19.20   20.09   22.80   33.90
```

Not writing a function:

```
round(c("Min." = min(mtcars$mpg),  
      "1st Qu." = as.numeric(quantile(mtcars$mpg)[2]),  
      "Median" = median(mtcars$mpg),  
      "Mean" = mean(mtcars$mpg),  
      "3rd Qu." = as.numeric(quantile(mtcars$mpg)[4]),  
      "Max." = max(mtcars$mpg)), 2)
```

```
>      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.  
>    10.40   15.43   19.20   20.09   22.80   33.90
```

# Single return object

Pure functions return a single object.

- (Standard) The last evaluated object
- Object defined by return()



**Figure 1:** A pure function.

## Single return object

`return()` stops the computation, and returns the object.

```
return_early <- function(x = 1) {  
  x2 <- x*2  
  return(x2)  
  out <- x + x2    # not executed  
  out  
}  
return_early(2)  
  
> [1] 4
```

# Single return object

Multiple return objects can be combined in a list!

```
do_this <- function(vector, other_vector) {  
  # many computations  
  return(list(output1 = this,  
              output2 = that))  
}
```

# Name  
# Arguments  
# Body  
# Output

# Single Return Object

The return object is a list with multiple objects.

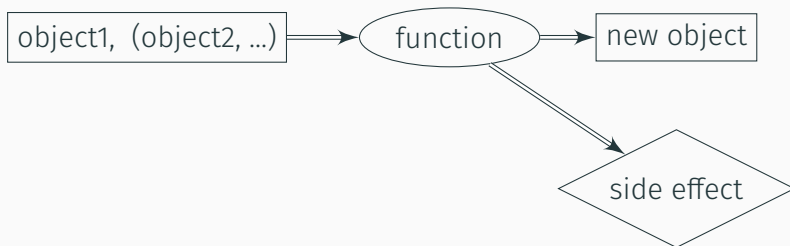
```
get_info <- function(x){  
  mean_x <- mean(x)  
  median_x <- median(x)  
  n_obs_x <- length(x)  
  range_x <- range(x)  
  return(list(mean = mean_x, median = median_x,  
              n_obs = n_obs_x, range = range_x))  
}  
str(get_info(airquality$Wind))
```

```
> List of 4  
> $ mean : num 9.96  
> $ median: num 9.7  
> $ n_obs : int 153  
> $ range : num [1:2] 1.7 20.7
```

# Side Effects

Functions can have “side effects”:

- console output
- plots
- write/save on drive
- ...



**Figure 2:** A function with side effect.

# Side Effects

Console output: ?cat and ?print

```
print_info <- function(x){  
  info <- get_info(x)  
  cat("There are ", info$n_obs,  
      " observed values. \nThe mean is ",  
      round(info$mean, 2), ". \nThe median is ",  
      round(info$median, 2), ". \n", sep = "")  
}  
print_info(airquality$Wind)
```

```
> There are 153 observed values.  
> The mean is 9.96.  
> The median is 9.7.
```



## Side effects

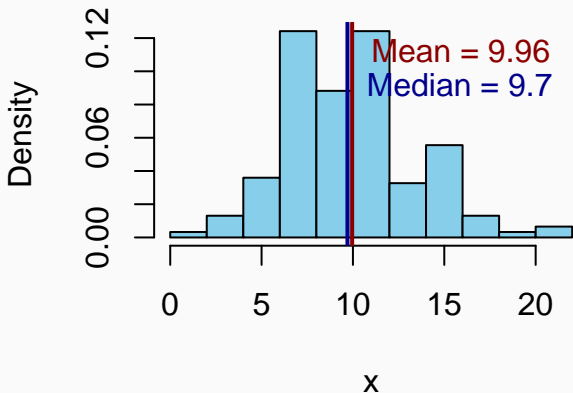
Graphics output: Standard plot, ggplot2, lattice

```
hist2 <- function(x, title){  
  info <- get_info(x)  
  mean_median <- as.numeric(info[c("mean", "median")])  
  hist(x, col = "skyblue", freq = FALSE,  
       main = paste0(title, " (n = ", info$n_obs, ")"))  
  abline(v = mean_median, lwd = 2,  
         col = c("darkred", "darkblue"))  
  text(mean_median, y = c(.11, .09),  
       labels = paste(c("Mean", "Median"),  
                     round(mean_median, 2),  
                     sep = " = "),  
       col = c("darkred", "darkblue"), pos = 4)  
}  
hist2(airquality$Wind, "Wind")
```

# Side effects

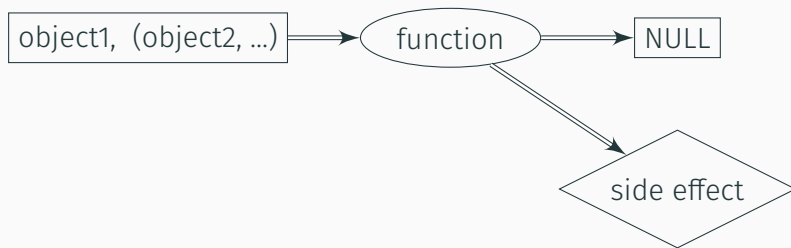
Graphics output

## Wind (n = 153)



## Programming advice

- Write pure functions (no-side effects)
- Write separate functions for side effects
- Plotting functions should return **NULL** or the plot as an object



**Figure 3:** A side effect function.

## Error, warning, & message

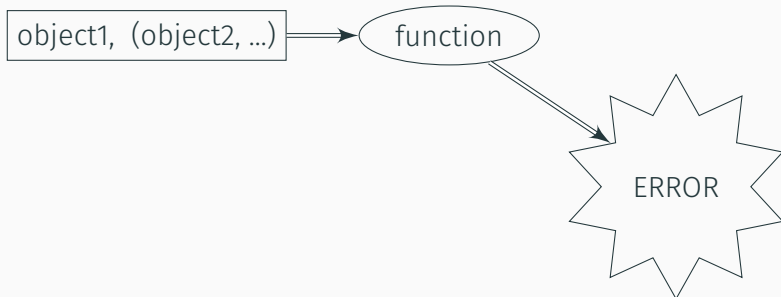
Error: computation is interrupted without return object!

?stop

```
get_log_xtox <- function(x) {  
  if(!is.numeric(x)) stop("This does not work!")  
  x_x <- x^x  
  return(log(x_x))  
}  
get_log_xtox("a")  
  
> Error in get_log_xtox("a"): This does not work!
```

## Error, warning, & message

Error: computation is interrupted without return object!



**Figure 4:** Computation with Error.

## Error, warning, & message

?stopifnot is an abbreviation for `if(!test) stop()`:

```
get_log_xtox <- function(x) {  
  stopifnot(is.numeric(x))  
  x_x <- x^x  
  return(log(x_x))  
}  
get_log_xtox("a")
```

```
> Error in get_log_xtox("a"): is.numeric(x) is not TRUE
```

## Error, warning, & message

Message: To inform the user about something.

?message

```
get_log_xtox <- function(x) {  
  x_x <- x^x  
  message("Thank you for using this function!")  
  return(log(x_x))  
}  
get_log_xtox(2)
```

```
> Thank you for using this function!
```

```
> [1] 1.386294
```

## Error, warning, & message

Warning: Warn the user that something may be fishy.

?warning

```
get_log_xtox <- function(x) {  
  if(x < 0 && (x %% 2 == 0))  
    warning("Not sure you can trust the result.",  
           call. = FALSE)  
  x_x <- x^x  
  return(log(x_x))  
}  
get_log_xtox(-2)
```

```
> Warning: Not sure you can trust the result.
```

```
> [1] -1.386294
```



## Error, warning, & message

Message & warning: computation is NOT interrupted!

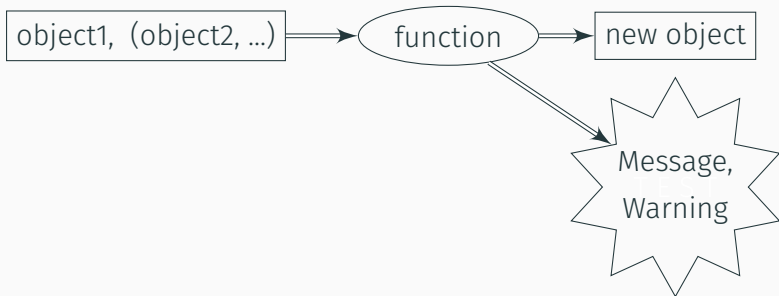


Figure 5: A message or warning.

## Programming advice

- Choose carefully when something warrants a message, warning or error
- Write clear and helpful warnings, errors, messages

## Default arguments

What happens if the user omits an argument?

```
add_ten <- function(x) {  
  return(x + 10)  
}  
add_ten()
```

```
> Error in add_ten(): argument "x" is missing, with no  
default
```

## Default arguments

Default arguments are made for such instances!

```
add_ten_default <- function(x = 0) {  
  return(x + 10)  
}  
add_ten_default()  
  
> [1] 10
```

# Default arguments

Additional arguments give (the user) flexibility. Default arguments keep the function easy to use.

Try ?lm

## Programming advice

- Think which arguments to include, and which should (not) have defaults
- Choose sensible defaults

# Lazy Evaluation

R only considers (evaluates) an argument when it is used.

```
add_ten_lazy <- function(x, y) {  
  return(x + 10)  
}  
add_ten_lazy(2, y = stop("This is not evaluated"))  
  
> [1] 12
```

# Lazy Evaluation

R only considers (evaluates) an argument when it is used. But, you can **force** the evaluation:

```
add_ten_force <- function(x, y) {  
  force(y)  
  return(x + 10)  
}  
add_ten_force(2, y = stop("Evaluation was forced"))  
  
> Error in force(y): Evaluation was forced
```

?force

# Exercises





## Functions III

---

## Where does a function find objects?

R uses specific rules to find objects, which lead to the following:

```
a <- 55
add_a <- function(x){
  return(x + a)
}
add_a(5)

> [1] 60
```

When a function is called, the computations in the body are run line by line. When R cannot find an object inside the function, it looks outside the function.

# Where does a function find objects?

## Name masking!

Objects inside the function mask objects outside the function with the same name.

```
a <- 55
add_a <- function(x){
  a <- 5
  return(x + a)
}
add_a(5)

> [1] 10
```

# Where does a function find objects?

R uses specific rules to find objects.

```
a <- b <- c <- d <- "fourth"
find_object <- function(a, b = "third", c = "third"){
  a <- "first"
  return(c(a = a, b = b, c = c, d = d))
}
find_object(b = "second")

>      a      b      c      d
> "first" "second" "third" "fourth"
```

## Where does a function find objects?

R uses specific rules to find objects.

1. in the function body
2. in the function call
3. in the function definition
4. outside the function

Watch out with number 4! Frequently restart R: **Ctrl + shift + F4**

# Functional programming

The return object should only depend on the arguments of the function, ***not*** on the context!

**BAD:**

```
a <- 55
add_a <- function(x){
  return(x + a)
}
add_a(5)

> [1] 60
```

# Functional programming

The return object should only depend on the arguments of the function, *not* on the context!

GOOD:

```
add_a <- function(x, a = 55){  
  return(x + a)  
}  
add_a(5)  
  
> [1] 60
```

# Functional programming

The function should not change the context.

**BAD**

```
a <- 55
change_a <- function(new_a){
  a <- new_a
  return(invisible(NULL))
}
change_a(5)
a

> [1] 5
```



## ... dot-dot-dot

R has a special argument (in the definition of the function):

... (dot-dot-dot)

Examples:

- ?sum
- ?save
- ...

## ... dot-dot-dot

... can take *any* number of additional arguments

Useful for passing arguments to other functions like:

- ?apply
- ?plot
- ...

Useful when you don't know how many arguments there will be.

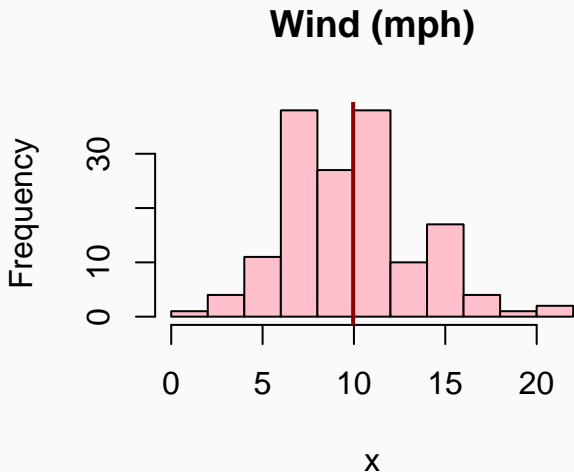
## ... dot-dot-dot

### plot example

```
hist3 <- function(x, ...){  
  hist(x, ...)  
  abline(v = mean(x, ...),  
         col = "darkred",  
         lwd = 2)  
}  
hist3(airquality$Wind, col = "pink",  
      main = "Wind (mph)")
```

... dot-dot-dot

plot example



## ... dot-dot-dot

apply example.

```
get_quantiles <- function(x, ...){  
  out <- lapply(x, quantile, ...)  
  return(do.call(rbind, out))  
}  
get_quantiles(airquality, na.rm = TRUE,  
              probs = c(.25, .5, .27))
```

```
>           25%    50%    27%  
> Ozone      18.00   31.5   18.05  
> Solar.R  115.75  205.0  127.00  
> Wind        7.40    9.7    7.40  
> Temp       72.00   79.0   73.00  
> Month        6.00    7.0    6.00  
> Day         8.00   16.0    9.00
```

## ... dot-dot-dot

**WARNING!** Watch out with spelling mistakes, arguments can get lost!

```
get_quantiles <- function(x, ...){  
  out <- lapply(x, quantile, ...)  
  return(do.call(rbind, out))  
}  
get_quantiles(airquality, na.rm = TRUE,  
              probs = c(.25, .5, .27))
```

```
>           0%    25%   50%   75%  100%  
> Ozone      1.0  18.00  31.5  63.25 168.0  
> Solar.R    7.0 115.75 205.0 258.75 334.0  
> Wind       1.7   7.40   9.7  11.50  20.7  
> Temp      56.0  72.00  79.0  85.00  97.0  
> Month      5.0   6.00   7.0   8.00   9.0  
> Day        1.0   8.00  16.0  23.00  31.0
```

## on.exit()

Performing an action when the function terminates.

```
add_ten_on_exit <- function(x) {  
  on.exit(cat("Finished 'add_ten_on_exit', with input '",  
             x, "'. \n", sep = ""))  
  return(x + 10)  
}  
add_ten_on_exit(1)  
  
> Finished 'add_ten_on_exit', with input '1'.  
> [1] 11
```

## on.exit()

Performing an action when the function terminates.

```
add_ten_on_exit <- function(x) {  
  on.exit(cat("Finished 'add_ten_on_exit', with input '",  
            x, "'. \n", sep = ""))  
  return(x + 10)  
}  
add_ten_on_exit("one")
```

```
> Error in x + 10: non-numeric argument to binary  
operator
```

```
> Finished 'add_ten_on_exit', with input 'one'.
```



## Error, warning, & message

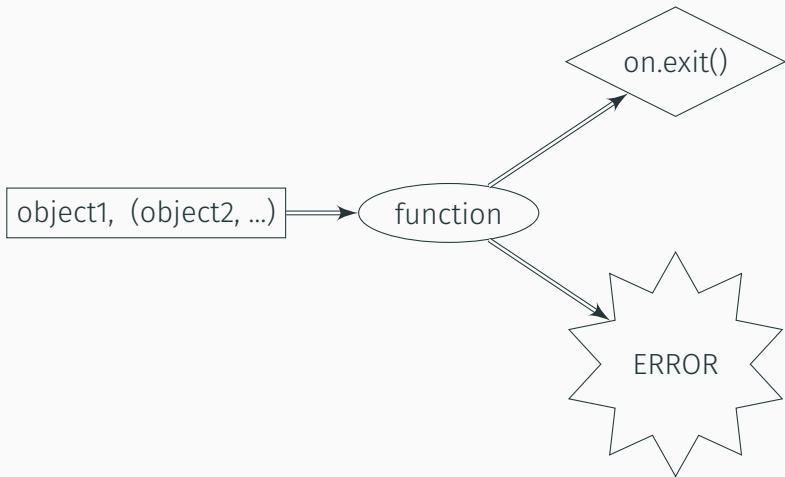


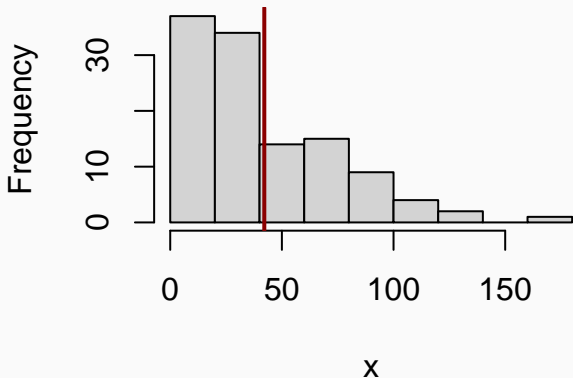
Figure 6: on.exit() with error.

Useful when your function has side effects:

```
hist3 <- function(x, ...){  
  old_options <- getOption("warn")  
  on.exit(options(warn = old_options))  
  options(warn = -1)  
  hist(x, ...)  
  abline(v = mean(x, ...),  
         col = "darkred", lwd = 2)  
}  
hist3(airquality$Ozon, na.rm = TRUE)
```

Useful when your function has side effects:

## Histogram of x



*“To understand computations in R, two slogans are helpful: Everything that exists is an object. Everything that happens is a function call.”*

— John Chambers

# Functions are objects

Functions are also objects. They can be arguments.

- `apply`-family
- ...

```
do_this_that <- function(function1, function2, x){  
  function2(function1(x))  
}  
do_this_that(sum, log, 0:3)  
  
> [1] 1.791759
```

# Functions are objects

Anonymous functions = functions without a name

```
do_this_that(function(x) x^2,  
              function2 = function(y) sum(y) / (length(y) - 1)),  
              -1:5)
```

```
> [1] 8
```

# Functions are objects

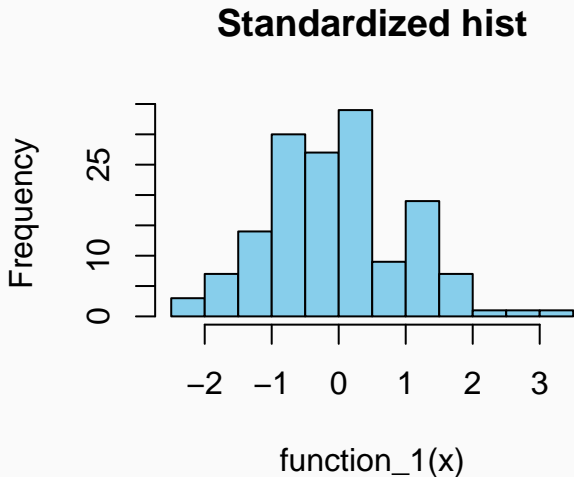
The return objects can also be functions:

```
combine_2fun <- function(function_1, function_2){  
  out_function <- function(x, ...) {  
    function_2(function_1(x), ...)  
  }  
  return(out_function)  
}
```

```
standardized_hist <- combine_2fun(scale, hist)  
standardized_hist(airquality$Wind,  
                  col = "skyblue",  
                  main = "Standardized hist")
```

# Functions are objects

The return objects can also be functions:





# Functions are objects

The return objects can also be functions:

```
combine_2fun <- function(function_1, function_2){  
  out_function <- function(x, ...) {  
    function_2(function_1(x), ...)  
  }  
  return(out_function)  
}  
mean_abs_deviation <- combine_2fun(abs, mean)  
mean_abs_deviation(airquality$Ozone, na.rm = TRUE)  
  
> [1] 42.12931
```

# Functions are objects

The return objects can also be functions:

```
normalize <- combine_2fun(  
  function(x) {x - min(x, na.rm = TRUE)},  
  function(x) {x / max(x, na.rm = TRUE)})  
normalize(airquality$Ozone)[1:4]  
  
> [1] 0.23952096 0.20958084 0.06586826 0.10179641
```

# Writing Functions

Before creating the function

- What should my function do?
- Which input objects (Arguments)?
- which additional options (Arguments)?
- What should the output object be?

After creating the function

- Test it
- Add input validation
- Document

# What makes a good function?

## Pure functions!

- no side effects
- no dependency on global environment
- only input via arguments (functional programming)

Results in easier understanding and higher portability.

# Exercises



# Debugging

---

- `browser()`
- `traceback()`
- `options(error = recover)`
- `options(warn = 2)`

# browser()

Inspecting a function interactively

```
some_function <- function(x, y) {  
  z <- x + y  
  browser()  
  z  
}  
some_function(x = 1, y = 5)
```



# browser()

```
> some_function <- function(x, y) {  
+   z <- x + y  
+   browser()  
+   z  
+ }  
> some_function(x = 1, y = 5)  
Called from: some_function(x = 1, y = 5)  
Browse[1]> |
```

# browser()

Navigating within a browser:

ls() Show existing objects in the current environment

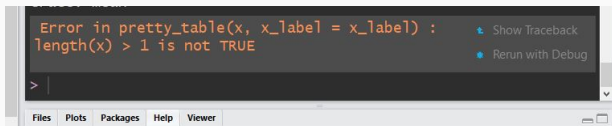
c Exit the browser and continue execution

Q Exit the browser, return to top level

where Show call stack

# traceback()

Understanding the call stack:



# traceback()

## Understanding the call stack:

```
Error in pretty_table(x, x_label = x_label) :  
length(x) > 1 is not TRUE  
13. stop(simpleError(msg, call = if (p <- sys.parent(1L)) sys.c  
all(p)))  
12. stopifnot(length(x) > 1)  
11. pretty_table(x, x_label = x_label)  
10. pretty_statistics(sub_dat$cy1, x_label = "cy1")  
9. FUN(data[x, , drop = FALSE], ...)  
8. FUN(X[[i]], ...)  
7. lapply(X = ans[index], FUN = FUN, ...)  
6. tapply(seq_len(32L), list('mtcars$carb' = c(4, 4, 1, 1, 2,  
1,  
4, 2, 2, 4, 4, 3, 3, 3, 4, 4, 4, 1, 2, 1, 1, 2, 2, 4, 2, 1,  
2,  
2, 4, 6, 8, 2)), function (x)  
FUN(data[x, , drop = FALSE], ...), simplify = TRUE)  
5. eval(substitute(tapply(seq_len(nd), IND, FUNx, simplify = s  
implify)),  
data)  
4. eval(substitute(tapply(seq_len(nd), IND, FUNx, simplify = s  
implify)),  
data)  
3. structure(eval(substitute(tapply(seq_len(nd), IND, FUNx, si  
mplify = simplify)),  
data), call = match.call(), class = "by")  
2. by.data.frame(mtcars, mtcars$carb, function(sub_dat) {  
pretty_statistics(sub_dat$cy1, x_label = "cy1")  
})  
1. by(mtcars, mtcars$carb, function(sub_dat) {  
pretty_statistics(sub_dat$cy1, x_label = "cy1")  
})  
> |
```

Being able to chose an environment from the call stack:

```
# on
options(error = recover)

# off
options(error = NULL)
```

Being able to choose an environment from a call stack:

```
Error in pretty_table(x, x_label = x_label) : length(x) > 1 is not TRUE
Enter a frame number, or 0 to exit

1: by(mtcars, mtcars$carb, function(sub_dat) {
  pretty_statistics(sub_dat$cyl, x_label = "Cyl")
2: by.data.frame(mtcars, mtcars$carb, function(sub_dat) {
  pretty_statistics(sub_dat$cyl, x_label = "Cyl")
3: structure(eval(substitute(tapply(seq_len(nd), IND, FUNx, simplify = simplify))), data), call =
4: eval(substitute(tapply(seq_len(nd), IND, FUNx, simplify = simplify))), data)
5: eval(substitute(tapply(seq_len(nd), IND, FUNx, simplify = simplify))), data)
6: tapply(seq_len(32), list(mtcars$carb = c(4, 4, 1, 1, 2, 1, 4, 2, 2, 4, 4, 3, 3, 3, 4, 4, 4,
7: tapply(X = ans[index], FUN = FUN, ...)
8: FUN(X[[i]], ...)
9: FUN(data[x, , drop = FALSE], ...)
10: #2: pretty_statistics(sub_dat$cyl, x_label = "Cyl")
11: #3: pretty_table(x, x_label = x_label)
12: #2: stopifnot(length(x) > 1)

Selection: |
```

# Warnings

Turning warnings into errors

```
# on  
options(warn = 2)  
  
# off  
options(warn = 1)
```

# Exercises





# Good programming practices

---

“Write code for humans, not for machines!”

Invest time in writing readable R-code.

- It will make collaborations easier
- It will make debugging easier
- It will help make your analyses reproducible

There is a complete *tidyverse* style-guide

<https://style.tidyverse.org/>.

## Go easy on your eyes

- with spaces before and after: `- + / * = <- < == >`
- always use `<-` for assignments
- only use `=` in function calls
- use indentation (largely automatic in RStudio)
- **`CamelCaseNames`** vs **`snake_case_names`**
- be consistent!
- wrap long lines at column 70-80 (Rstudio)

## White space

```
new_var=(var1*var2/2)-5/(var3+var4)
```

```
# versus
```

```
new_var <- (var1 * var2 / 2) - 5 / (var3 + var4)
```

# Indentation

```
for(name in names){formula=as.formula(paste0("y~.-",name))
fit<-lm(formula,data=my_data)
coefs[["name"]]=coef(fit)
print(name)
print(summary(fit))}
```

# versus

```
for(name in names){
  formula <- as.formula(paste0("y~.-", name))
  fit <- lm(formula, data = my_data)
  coefs[["name"]] <- coef(fit)
  print(name)
  print(summary(fit))
}
```

## Wrap long lines

```
final_results <- data.frame(first_variable =  
  sqrt(results$mean_squared_error), second_variable =  
  paste0(results$condition, results$class, sep = ":"),  
  third_variable = results$bias)
```

# versus

```
final_results <- data.frame(  
  first_variable = sqrt(results$mean_squared_error),  
  second_variable = paste0(results$condition,  
                           results$class, sep = ":"),  
  third_variable = results$bias)
```

# Go easy on your mind

- use meaningful names: “self-explainable”
- always write the formal arguments in function calls (except the first)
- benefit from autocompletion (`<tab>`) => embrace longer names
- use **TRUE** and **FALSE** not **T** and **F**
- comment, comment, comment
  - NOT what (should be clear from the code)
  - but why
  - explain the reasoning, not the code



## Use meaningful names

```
V <- myFun(m1_B)
```

```
# versus
```

```
RMSE_age_gender <- get_RMSE(lm_age_gender)
```

### Programming advice

Use **verbs** for functions and **nouns** for other objects.

# Write formal arguments

Benefit from auto completion using `tab`

```
m1_B <- lm(outcome ~ age*gender,  
            exp1, condition_1, freq)
```

# versus

```
lm_age_gender <- lm(outcome ~ age * gender,  
                    data = exp1,  
                    subset = condition_1,  
                    weights = freq)
```

## Comment, comment, comment

```
## Start every Rscript with a comment that explains
## what the code in the script does, why it does
## this, and to which project it belongs.
## Your future self will be very thankful!
##
## Mention which packages you are using in this Rscript.

## Use sections to separate chunks -----

## Maybe even subsections =====

## Recode variables so that missings are coded as "NA"
dat[dat %in% c(99, 999)] <- NA # missings coded 99 or 999
```

## Keep your code slim

Try to limit your *package-dependencies*.

Only load `library()` the packages that you absolutely need. If you are only using `dplyr`, it does not make sense to load the complete `tidyverse`.

**Controversial:** when possible, use the `::` operator (and consider not loading the package).

`<package>::<function>`

- explicit dependencies
- less name conflicts

# Never Attach

Forget about `attach()`!

Don't use it, unless you completely understand what happens (see `?attach`).

Use `with(data.frame, expression)` instead.

```
# using with()
n <- 2e+4
data <- data.frame(x = runif(n),
                  y = runif(n),
                  z = seq_len(n))
result <- with(data, exp(x) / log(z) + 5 * sqrt(y))
```

Writing code is error prone. Incorporate tests and checks in your workflow.

- minimal examples
- write tests and checks
- helpful packages: `testthat`, `RUnit`, `testit`, ...

Computing speed can become an issue. Avoid common pitfalls:

- don't grow, but replace
- vectorize where possible
- check the computing speed

?`system.time`, microbenchmark or profiling tools

```
n <- 2e+4
data <- data.frame(x = runif(n),
                  y = runif(n),
                  z = seq_len(n))
```

# Speed

Don't grow!

```
system.time({  
  new_data <- NULL  
  
  for(row_nr in seq_len(NROW(data))){  
    new_data <- cbind(  
      data[row_nr,],  
      result = exp(data$x[row_nr]) /  
        log(data$z[row_nr]) +  
        5 * sqrt(data$y[row_nr]))  
  }  
})
```

```
>   user  system elapsed  
> 3.77    0.00    3.77
```



Replace!

```
system.time({  
  n_rows <- dim(data)[1]  
  data$result <- rep(NA, n_rows)  
  
  for(row_nr in seq_len(n_rows)){  
    data$result[row_nr] <- exp(data$x[row_nr]) /  
      log(data$z[row_nr]) +  
      5 * sqrt(data$y[row_nr])  
  }  
})
```

```
>   user  system elapsed  
> 0.51    0.04    0.54
```

Vectorize!

```
system.time({  
  data$result <- exp(data$x) / log(data$z) +  
    5 * sqrt(data$y)  
})
```

```
>      user  system elapsed  
>         0         0         0
```

# Speed

Compare the speed of different implementations using:

`microbenchmark::microbenchmark`

```
get_mean1 <- function(x){  
  weight <- 1/length(x)  
  out <- 0  
  for(i in seq_along(x)){  
    out <- out + x[i] * weight  
  }  
  return(out)  
}  
  
get_mean2 <- function(x){  
  sum(x)/length(x)  
}
```

# Speed

Compare the speed of different implementations using:

`microbenchmark::microbenchmark`

```
x <- rnorm(500)
microbenchmark::microbenchmark(
  mean(x), get_mean1(x), get_mean2(x))
```

```
> Unit: microseconds
```

>	expr	min	lq	mean	median	uq	max	neval	cld
>	mean(x)	4.1	4.2	4.793	4.3	4.50	27.0	100	a
>	get_mean1(x)	29.3	30.3	81.544	30.3	30.45	5128.3	100	a
>	get_mean2(x)	1.5	1.5	20.609	1.6	1.70	1877.8	100	a

## Programming advice

Don't worry about speed before it becomes an issue.

“Every project should get an RStudio Project!”

Don't use `setwd(``path to my local_folder'')`

Issues when:

- folders names are changed
- folders are moved
- a shared drive is used
- you ZIP and send the folder

Don't save work space to `.RData`.

- Tools < Global Options < Workspace < Save workspace ....
- Save the code instead!
- Use `saveRDS()` and `readRDS()` for objects that require long computations

Don't use `rm(list = ls())` at the start of an Rscript.

- Start clean, every time.
- Keep it clean. No outside code, no outside computing.
- Regularly completely clean the work space (or restart the session).

```
.rs.restartR()
```



Keep it clean

- one folder per project!
- work on different projects in different RStudio instances!
- each with own R console, working directory, ...

# Working with RStudio

## Organize your project folder

- R-folder with R scripts
- Data-folder with data
- split long scripts in meaningful chunks
- use relative paths (alternative: here-package)

```
# read data
this_data <- read.csv("Data\the-correct-file.csv")

# source Rscript
source("R\0_first-script-to-source.R")
```

## Use keyboard shortcuts

- Can make working in RStudio more efficient
- Completely tunable: Tools < Modify Keyboard Shortcuts...
- Useful shortcuts (defaults):
  - jump to editor: `ctrl + 1`
  - jump to console: `ctrl + 2`
  - jump to ...: `ctrl + 3-9`
  - jump to next tab: `ctrl + tab`
  - jump to previous tab: `ctrl + shift + tab`

More useful shortcuts (defaults):

- run selection/selected line: `ctrl + enter`
- save current file: `ctrl + s`
- close current file: `ctrl + w`
- restart R: `ctrl + shift + F10`
- Show help (for function at cursor) `F1`
- Show source code (for function at cursor) `F2`

More on this [HERE](#).

# Exercises



## Wrap Up

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## General Advice

- Investing time in learning R pays off
- It's a steady learning curve
- Learn from masters
- Rewrite important code - the first attempt is usually not the best approach

# General R Advice

- Document well
- Use a consistent style
- Write functions
- Split long functions in smaller ones
- Write wrappers
- Use Iteration (don't copy paste)
- Use matrix operations and vectorized functions instead of loops
- Use git



## R Resources

- Advanced R Ed. 1 (<http://adv-r.had.co.nz/>)
- Advanced R Ed. 2 (<https://adv-r.hadley.nz/>)
- R Inferno ([https://www.burns-stat.com/pages/Tutor/R\\_inferno.pdf](https://www.burns-stat.com/pages/Tutor/R_inferno.pdf))
- R Packages (<https://r-pkgs.org/>)
- Clean Code (<https://enos.itcollege.ee/~jpoial/oop/naited/Clean%20Code.pdf>)

Thank you for your attention!

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Questions? Remarks?