

Functional programming

July 2018

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Adapted from *Tidy Tools* by Hadley Wickham



Motivation

Copy and paste is a rich source of errors

```
# Fix missing values
df$a <- df$a[df$a == -99] <- NA
df$b <- df$b[df$b == -99] <- NA
df$c <- df$c[df$c == -99] <- NA
df$d <- df$d[df$d == -99] <- NA
df$e <- df$e[df$e == -99] <- NA
df$f <- df$f[df$f == -99] <- NA
df$g <- df$g[df$g == -98] <- NA
df$h <- df$h[df$h == -99] <- NA
df$i <- df$i[df$i == -99] <- NA
df$j <- df$i[df$j == -99] <- NA
df$k <- df$k[df$k == -99] <- NA
```

Copy and paste is a rich source of errors

```
# Fix missing values
```

```
df$a <- df$a[df$a == -99] <- NA
```

```
df$b <- df$b[df$b == -99] <- NA
```

```
df$c <- df$c[df$c == -99] <- NA
```

```
df$d <- df$d[df$d == -99] <- NA
```

```
df$e <- df$e[df$e == -99] <- NA
```

```
df$f <- df$f[df$f == -99] <- NA
```

```
df$g <- df$g[df$g == -98] <- NA
```

```
df$h <- df$h[df$h == -99] <- NA
```

```
df$i <- df$i[df$i == -99] <- NA
```

```
df$j <- df$i[df$j == -99] <- NA
```

```
df$k <- df$k[df$k == -99] <- NA
```

Functions can remove some sources of duplication

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df$a <- fix_missing(df$a)  
df$b <- fix_missing(df$b)  
df$c <- fix_missing(df$c)  
df$d <- fix_missing(df$d)  
df$e <- fix_missing(df$e)  
df$f <- fix_missing(df$f)  
df$g <- fix_missing(df$g)  
df$h <- fix_missing(df$h)  
df$h <- fix_missing(df$i)
```

Functions can remove some sources of duplication

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df$a <- fix_missing(df$a)  
df$b <- fix_missing(df$b)  
df$c <- fix_missing(df$c)  
df$d <- fix_missing(df$d)  
df$e <- fix_missing(df$e)  
df$f <- fix_missing(df$f)  
df$g <- fix_missing(df$g)  
df$h <- fix_missing(df$h)  
df$h <- fix_missing(df$i)
```

For loops can remove others

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}
```

```
for (i in seq_along(df)) {  
  df[[i]] <- fix_missing(df[[i]])  
}
```

Why for loops
are bad

Why for loops
~~are bad~~

suboptimal

What does this code do?

```
out1 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out1[[i]] <- mean(mtcars[[i]], na.rm = TRUE)
}
```

```
out2 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out2[[i]] <- median(mtcars[[i]], na.rm = TRUE)
}
```

For loops emphasise the objects

```
out1 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out1[[i]] <- mean(mtcars[[i]], na.rm = TRUE)
}
```

```
out2 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out2[[i]] <- median(mtcars[[i]], na.rm = TRUE)
}
```

Not the actions

```
out1 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out1[[i]] <- mean(mtcars[[i]], na.rm = TRUE)
}
```

```
out2 <- vector("double", ncol(mtcars))
for(i in seq_along(mtcars)) {
  out2[[i]] <- median(mtcars[[i]], na.rm = TRUE)
}
```

Functional programming emphasises the actions

```
library(purrr)
```

```
means <- map_dbl(mtcars, mean)  
medians <- map_dbl(mtcars, median)
```

Read: for each element of
mtcars, apply mean.

FP tools allow you to focus on what happens

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df$a <- fix_missing(df$a)  
df$b <- fix_missing(df$b)  
df$c <- fix_missing(df$c)  
df$d <- fix_missing(df$d)  
df$e <- fix_missing(df$e)  
df$f <- fix_missing(df$f)  
df$g <- fix_missing(df$g)  
df$h <- fix_missing(df$h)  
df$i <- fix_missing(df$i)
```

FP tools allow you to focus on what happens

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df <- modify(df, fix_missing)
```

And provide useful tools for **generalisation**

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df <- modify_if(df, fix_missing)
```


Foundations

Warmup: Guess the output, then run to check.

```
# Guess the output  
fun1 <- mean  
fun1(c(1, 3))
```

```
# Guess the output  
fun2 <- `+`  
fun2(1, 3)
```

How can we remove the duplication here?

```
df <- tibble::tibble(  
  a = c(rnorm(9), -99),  
  b = c(-999, -99, rnorm(8)),  
  c = c(0, rnorm(9)),  
  d = rnorm(10)  
)  
  
# Center columns  
df$a <- df$a - mean(df$a)  
df$b <- df$b - median(df$b)  
df$c <- df$c - dplyr::first(df$c)
```

Your Turn

Write the function `recenter()` to remove the repetition:

```
df$a <- recenter(df$a, mean)
df$b <- recenter(df$b, median)
df$c <- recenter(df$c, dplyr::first)
```

Identify what might change

```
df$a - mean(df$a)
df$b - median(df$b)
df$c - dplyr::first(df$c)
```

Give them names,

x

fun

x

```
df$a - mean(df$a)
df$b - median(df$b)
df$c - dplyr::first(df$c)
```

Make the template

```
recenter <- function(x, fun){  
  
}
```

Copy in an example

```
recenter <- function(x, fun){  
  df$a - mean(df$a)  
}
```


And use the argument names

```
recenter <- function(x, fun){  
  x - mean(x)  
}
```

And use the argument names

```
recenter <- function(x, fun){  
  x - fun(x)  
}
```

And use the argument names

```
recenter <- function(x, fun){  
  x - fun(x)  
}
```

Passing in a function as an argument

```
df$a <- recenter(df$a, mean)  
df$b <- recenter(df$b, median)  
df$c <- recenter(df$c, dplyr::first)
```

Functions are **first class** citizens in R

Anything you can do with a vector you can do with a function, e.g.:

1. Functions can be assigned to variables.
2. Functions can be arguments to functions.
3. Functions can be returned from a function.
4. Functions can be stored in lists.

Functions are **first class** citizens in R

Anything you can do with a vector you can do with a function, e.g.:

1. Functions can be assigned to variables.
- 2. Functions can be arguments to functions.**
3. Functions can be returned from a function.
4. Functions can be stored in lists.

Solving iteration problems with the `map()` family

A family of functionals

Map strategy

For an iteration task:

1. Solve for single x
2. Generalise solution with appropriate `map()` function
3. Simplify (if possible)

Find first element of compound string

```
x1 <- c("a|b", "a|b|c", "d|e", "b|c|d")
```

```
# We want:
```

```
# "a" "a" "d" "b"
```

```
# A useful intermediate object
```

```
x2 <- strsplit(x1, "|", fixed = TRUE)
```

```
# For each element of x2
```

```
# pull out the first element
```

```
# [[1]]  
# [1] "a" "b"  
#  
# [[2]]  
# [1] "a" "b" "c"  
#  
# [[3]]  
# [1] "d" "e"  
#  
# [[4]]  
# [1] "b" "c" "d"
```


1. Solve for single .x

Pull out one element

```
.x <- x2[[1]]
```

Specially named pronoun that map understands

```
.x
```

[1] "a" "b"

Get first element

```
.x[[1]]
```

Solved!

2. Generalise solution with map()

```
# Solution for one element  
  .x[[1]]
```

```
# Turn into a recipe with ~ and pass to map  
map(x2, ~ .x[[1]])
```

For each
element of
x2,

take it, and extract the
first element

Your Turn

Compute the mean of every column in mtcars.

Generate a sample of size 10 from Normals with the following means: -10, 0, 10, 100

Compute the number of unique values in each column of iris

Compute the mean of every column in mtcars

```
# Solve for one
```

```
.x <- mtcars[[1]]
```

```
mean(.x)
```

```
# Generalise
```

```
map(mtcars, ~ mean(.x))
```

Generate 10 random normals

```
mu <- c(-10, 0, 10, 100)
```

```
# Solve for one
```

```
.x <- mu[[1]]
```

```
rnorm(10, mean = .x)
```

```
# Generalise
```

```
map(mu, ~ rnorm(10, mean = .x))
```

Compute the number of unique values in each column

```
# Solve for one
```

```
.x <- iris[[1]]
```

```
length(unique(.x))
```

```
# Generalise
```

```
map(iris, ~ length(unique(.x)))
```

Map strategy

For an iteration task:

1. Solve for single x
2. Generalise solution with
appropriate `map()` function
3. Simplify (if possible)

Each variant always produces the same type

Function	Output
<code>map_lgl()</code>	Logical vector
<code>map_int()</code>	Integer vector
<code>map_dbl()</code>	Double vector
<code>map_chr()</code>	Character vector
<code>map()</code>	List
<code>map_dfc()</code>	Data frame (by col)
<code>map_dfr()</code>	Data frame (by row)

Map strategy

For an iteration task:

1. Solve for single x
2. Generalise solution with appropriate `map()` function
3. **Simplify** (if possible)

Simplify extraction

```
map(z, ~ .x[[1]])
```

```
map(z, 1)
```

```
map(z, ~ .x[["string"]])
```

```
map(z, "string")
```

```
map(z, ~ .x[["string"]][[1]] %||% NA)
```

```
map(z, list("string", 1), .default = NA))
```

Simplify function calls

`map(z, ~ f(.x))`

`map(z, f)`

`map(z, ~ f(.x, a = 1, b = 2))`

`map(z, f, a = 1, b = 2)`

`map(z, ~ f(1, .x))`

`map(z, f, first_arg = 1)`

Your Turn

For the three examples earlier:

```
map(mu, ~ rnorm(10, mean = .x))
```

```
map(mtcars, ~ mean(.x))
```

```
map(iris, ~ length(unique(.x)))
```

Use an appropriate map function, and simplify if possible.

Compute the mean of every column in mtcars

```
# Appropriate function
```

```
map_dbl(mtcars, ~ mean(.x))
```

```
# Simplify (optional)
```

```
map_dbl(mtcars, mean)
```

Generate 10 random normals

```
mu <- c(-10, 0, 10, 100)
```

```
# Appropriate function?
```

```
map(mu, ~ rnorm(10, mean = .x))
```

```
map_dfc(mu, ~ rnorm(10, mean = .x)) #?
```

```
# Simplify (optional)
```

```
map(mu, rnorm, n = 10)
```

Compute the number of unique values in each column

Appropriate function

```
map_int(iris, ~ length(unique(.x)))
```

Simplify ?

```
nunique <- function(x) length(unique(x))
```

```
map_int(iris, ~ nunique(.x))
```

```
map_int(iris, nunique)
```

← **Typo**

FP tools allow you to focus on what happens

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}  
df$a <- fix_missing(df$a)  
df$b <- fix_missing(df$b)  
df$c <- fix_missing(df$c)  
df$d <- fix_missing(df$d)  
df$e <- fix_missing(df$e)  
df$f <- fix_missing(df$f)  
df$g <- fix_missing(df$g)  
df$h <- fix_missing(df$h)  
df$i <- fix_missing(df$i)
```


Why isn't this quite what we want?

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}
```

```
df2 <- map(df, fix_missing)
```

```
# Hint: look at the structure of df2
```

Why isn't this quite what we want?

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}
```

```
df2 <- map(df, fix_missing)
```

```
str(df2)
```

```
# List of 4
```

```
# $ a: num [1:10] 10.53 10.32 9.87 10.5 8.07 ...
```

```
# $ b: num [1:10] -998.33 -98.33 0.419 -1.084 -0.258 ...
```

```
# $ c: num [1:10] 0 0.121 0.863 0.554 0.677 ...
```

```
# $ d: num [1:10] -0.769 1.325 1.972 -0.185 -0.289 ...
```

Why isn't this quite what we want?

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}
```

```
df[] <- map(df, fix_missing)  
str(df)
```

```
# Classes 'tbl_df', 'tbl' and 'data.frame': 10 obs. of  4 variables:  
# $ a: num  10.53 10.32 9.87 10.5 8.07 ...  
# $ b: num -998.33 -98.33 0.419 -1.084 -0.258 ...  
# $ c: num  0 0.121 0.863 0.554 0.677 ...  
# $ d: num -0.769 1.325 1.972 -0.185 -0.289 ...
```

Why isn't this quite what we want?

```
fix_missing <- function(x) {  
  x[x == -99] <- NA  
  x  
}
```

```
df <- modify(df, fix_missing)  
str(df)
```

```
# Classes 'tbl_df', 'tbl' and 'data.frame': 10 obs. of  4 variables:  
# $ a: num  10.53 10.32 9.87 10.5 8.07 ...  
# $ b: num -998.33 -98.33 0.419 -1.084 -0.258 ...  
# $ c: num  0 0.121 0.863 0.554 0.677 ...  
# $ d: num -0.769 1.325 1.972 -0.185 -0.289 ...
```

Why not base R?

Compared to purrr, base R functions:

Have inconsistent names (`lapply()` vs. `Map()`)

Have inconsistent argument order (`lapply()` vs. `mapply()`)

Require functions (no `1`, or extract helpers)

Are either type-unstable (`sapply()`) or verbose (`vapply()`)

Lack side-effect form (no `walk()`)

Lack paired maps (no `map2()`)

Lack data frame output (no `_dfc()`, `_dfr()`)

purrr provides a full set of functions

Number of inputs		Output is a scalar	Output is anything	Output is nothing
	1	map_lgl(), map_int(), map_dbl(), map_chr()	map()	walk()
	2	map2_lgl(), map2_int(), map2_dbl(), map2_chr()	map2()	walk2()
	n	pmap_lgl(), pmap_int(), pmap_dbl(), pmap_chr()	pmap()	pwalk()

Base R only provides a partial set of functions

Number of inputs		Output is a scalar	Output is anything	Output is nothing
	1	<code>sapply()</code> / <code>vapply()</code>	<code>lapply()</code>	
	2			
	n	<code>mapply()</code>	<code>Map()</code>	

Paired map

stringr application

How do we go from locations to words?

Easy if we have a single location

```
pos <- str_locate(sentences, "\\b\\w{5,}\\b")
```

```
str_sub(sentences, pos)
```

NB: str_sub can take one

2 column matrix, or two vectors



A word with 5 or more
letters.

What if we have multiple locations?

```
pos <- str_locate_all(  
  sentences, "\\b\\w{5,}\\b"  
)
```

Solve for one instance: now have two inputs!

```
.x <- sentences[[1]]
```

```
.y <- pos[[1]]
```



Another special pronoun

```
str_sub(.x, .y)
```

Generalise & simplify

Generalise

```
map2(sentences, pos, ~ str_sub(.x, .y))
```



1st input



2nd input

Simplify

```
map2(sentences, pos, str_sub)
```

walk2() is often useful when writing files

```
diamonds <- ggplot2::diamonds  
by_color <- split(diamonds, diamonds$color)  
paths <- paste0(names(by_color), ".csv")
```

```
# Solve for one  
.x <- by_color[[1]]  
.y <- paths[[1]]  
write.csv(.x, .y)
```

```
# Solve for all  
walk2(by_color, paths, ~ write.csv(.x, .y))
```

```
# Simplify  
walk2(by_color, paths, write.csv)
```

Principle:

Compose value functions with `map()`;
compose effect functions with `walk()`

Your Turn

```
diamonds <- ggplot2::diamonds
by_color <- split(diamonds, diamonds$color)
plots <- map(by_color, ~ ggplot(.x, aes(clarity, price)) +
              geom_point())
plot_paths <- paste0(names(by_color), ".pdf")
```

Save the plots to the corresponding paths.

Your Turn

```
# Solve for one
```

```
.x <- plots[[1]]
```

```
.y <- plot_paths[[1]]
```

```
ggsave(.y, .x)
```

```
# Generalise
```

```
walk2(plots, plot_paths, ~ ggsave(.y, .x))
```

```
# Simplify
```

```
walk2(plot_paths, plots, ggsave)
```

```
# To clean up
```

```
file.remove(paths)
```

```
file.remove(plot_paths)
```


Functions are **first class** citizens in R

Anything you can do with a vector you can do with a function, e.g.:

1. Functions can be assigned to variables.
2. Functions can be arguments to functions.
- 3. Functions can be returned from a function.**
4. Functions can be stored in lists.

Handling errors with `safely()`

A functional operator

What happens when there is an error?

```
input <- list(1:10, sqrt(4), 5, "n")  
map(input, log)
```

What does safely() do?

```
# safely() modifies a function so it never fails
```

```
safe_log <- safely(log)
```

```
# The output of safely() is a function
```

```
safe_log
```

```
# function (...)
```

```
# capture_error(.f(...), otherwise, quiet)
```

```
# <bytecode: 0x7fcb996c2398>
```

```
# <environment: 0x7fcb99679058>
```

What does safely() do?

What does it return when the function succeeds?

```
safe_log(1:10)
```

What does it return when the function fails?

```
safe_log("n")
```

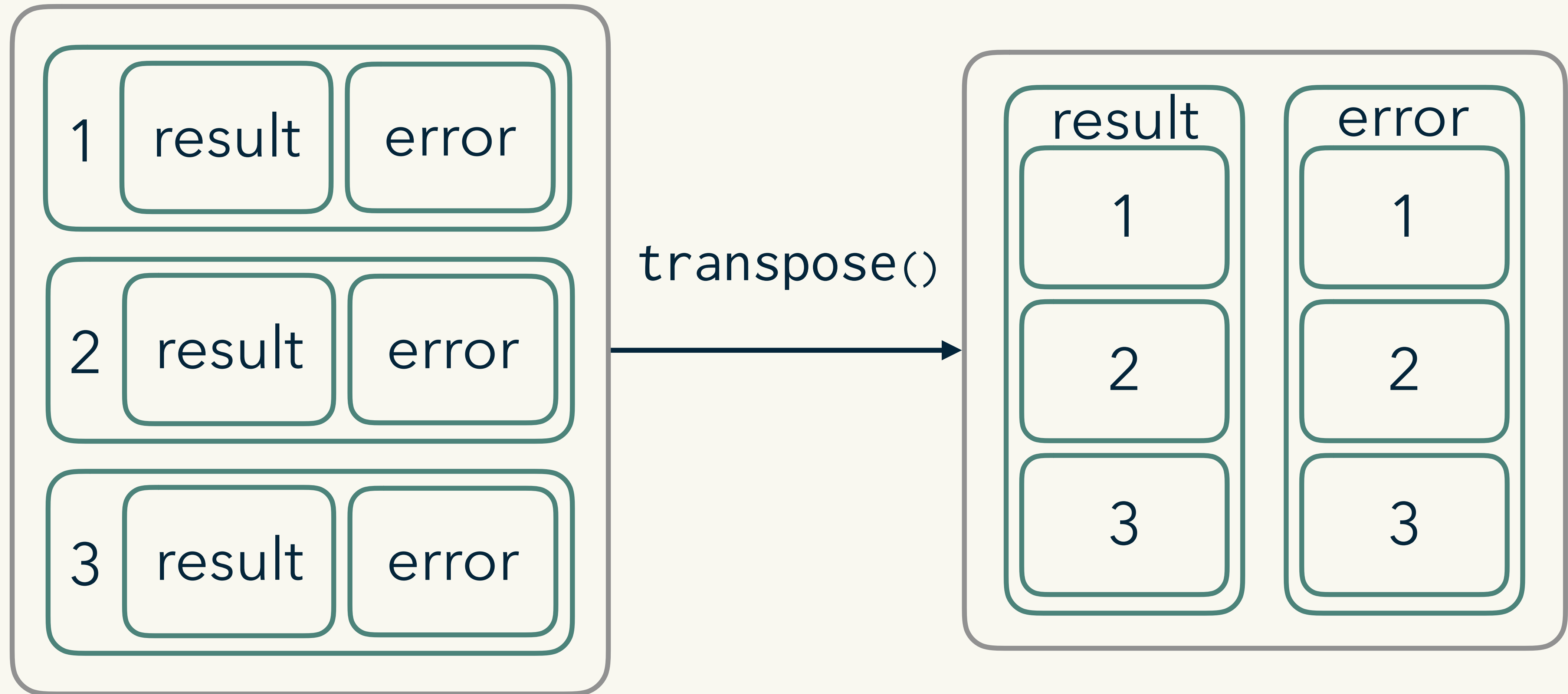
A more useful example

```
urls <- c(
  "https://google.com",
  "https://en.wikipedia.org",
  "asdfasdaskfjlda"
)

# Fails
contents <- map(urls, readLines, warn = FALSE)

# Always succeeds
contents <- urls %>%
  map(safely(readLines), warn = FALSE)
str(contents)
```

But `map()` + `safely()` gives awkward output



Your turn

```
urls <- c(
  "https://google.com",
  "https://en.wikipedia.org",
  "asdfasdasdkfjlda"
)
contents <- urls %>%
  map(safely(readLines), warn = FALSE)
```

Apply `transpose()` to `contents` then:

1. Make logical vector that is TRUE if download succeeded. (Hint: use `map_lgl()`)
2. List failed urls
3. Extract successfully retrieved text

Common pattern with safely()

```
contents <- urls %>%  
  map(safely(readLines)) %>%  
  transpose()
```

```
ok <- map_lgl(contents$error, is.null)  
# This is suboptimal:  
ok <- !map_lgl(contents$result, is.null)
```

```
urls[!ok]  
contents$result[ok]
```

Case Study

Switch to project
case_study

Your Turn

Open 02-report.R.

Pay attention to contents of files, file_paths and states.

Remove the duplication in the data import step:

```
# Import data
```

```
-----
```

```
or <- read_csv(or_file_path)
```

```
bc <- read_csv(bc_file_path)
```

```
wa <- read_csv(wa_file_path)
```

One Solution

```
# Import data
```

```
-----
```

```
all_states <- map(file_paths, read_csv)
```

Requires downstream edits too...

```
# Instead of this
```

```
or_weekly <- summarise_weekly(or)
```

```
bc_weekly <- summarise_weekly(bc)
```

```
wa_weekly <- summarise_weekly(wa)
```

```
# Will now need
```

```
all_states_weekly <- map(all_states, summarise_weekly)
```

Requires downstream edits too...

```
# Instead of this
```

```
or_plot <- plot_weekly(or_weekly, "Oregon")
```

```
bc_plot <- plot_weekly(bc_weekly, "British Columbia")
```

```
wa_plot <- plot_weekly(wa_weekly, "Washington")
```

```
# Will now need
```

```
all_states_plots <- map2(all_states_weekly,  
  states_long_names,  
  plot_weekly)
```

Your Turn

```
or_plot <- plot_weekly(or_weekly, "Oregon")  
or_image_path <- path("images",  
  paste(or_file, "n_sales", sep = "_"))  
  
ggsave(paste0(or_image_path, ".pdf"), or_plot, height = 3, width = 8)  
ggsave(paste0(or_image_path, ".png"), or_plot, height = 3, width = 8)
```

Use an appropriate map function to save or_plot to all extensions.

1. Solve for one extension

```
exts <- c(".pdf", ".png")
```

```
.x <- exts[[1]]
```

```
ggsave(paste0(or_image_path, .x),  
        or_plot, height = 3, width = 8)
```

2. Generalize & Simplify

Generalize

```
walk(exts, ~ ggsave(paste0(or_image_path, .x),  
  or_plot, height = 3, width = 8))
```

Simplify

```
walk(exts, ~ ggsave(paste0(or_image_path, .x)),  
  plot = or_plot, height = 3, width = 8)
```

Even better

```
image_paths <- paste0(or_image_path, exts)  
walk(image_paths, ggsave,  
  plot = or_plot, height = 3, width = 8)
```

Challenge

Try writing the `ggsave_multiple()` function:

```
exts <- c(".pdf", ".png")  
image_paths <- paste0(or_image_path, exts)  
  
walk(image_paths, ggsave,  
      plot = or_plot, height = 3, width = 8)
```

1. What might vary? What are our argument names?

```
exts <- c(".pdf", ".png")
```

```
image_paths <- paste0(or_image_path, exts)
```

```
walk(image_paths, ggsave,  
      plot = or_plot, height = 3, width = 8)
```

1. What might vary? What are our argument names?

```
exts <- c(".pdf", ".png")
```

exts

```
image_paths <- paste0(or_image_path, exts)
```

filename

```
walk(image_paths, ggsave,
```

```
  plot = or_plot, height = 3, width = 8)
```

plot

any other args to ggsave

Put in function template, matching ggsave() arg order

```
ggsave_multiple <- function(filename, plot, exts, ...){  
  paths <- paste0(filename, exts)  
  
  walk(paths, ggsave,  
    plot = plot, ...)  
}  
ggsave_multiple(or_image_path, or_plot, c(".pdf", ".png"),  
  height = 3, width = 8)
```

Downstream changes

This

```
ggsave_multiple(or_image_path, or_plot, c(".pdf", ".png"),  
  height = 3, width = 8)  
ggsave_multiple(bc_image_path, bc_plot, c(".pdf", ".png"),  
  height = 3, width = 8)  
ggsave_multiple(wa_image_path, wa_plot, c(".pdf", ".png"),  
  height = 3, width = 8)
```

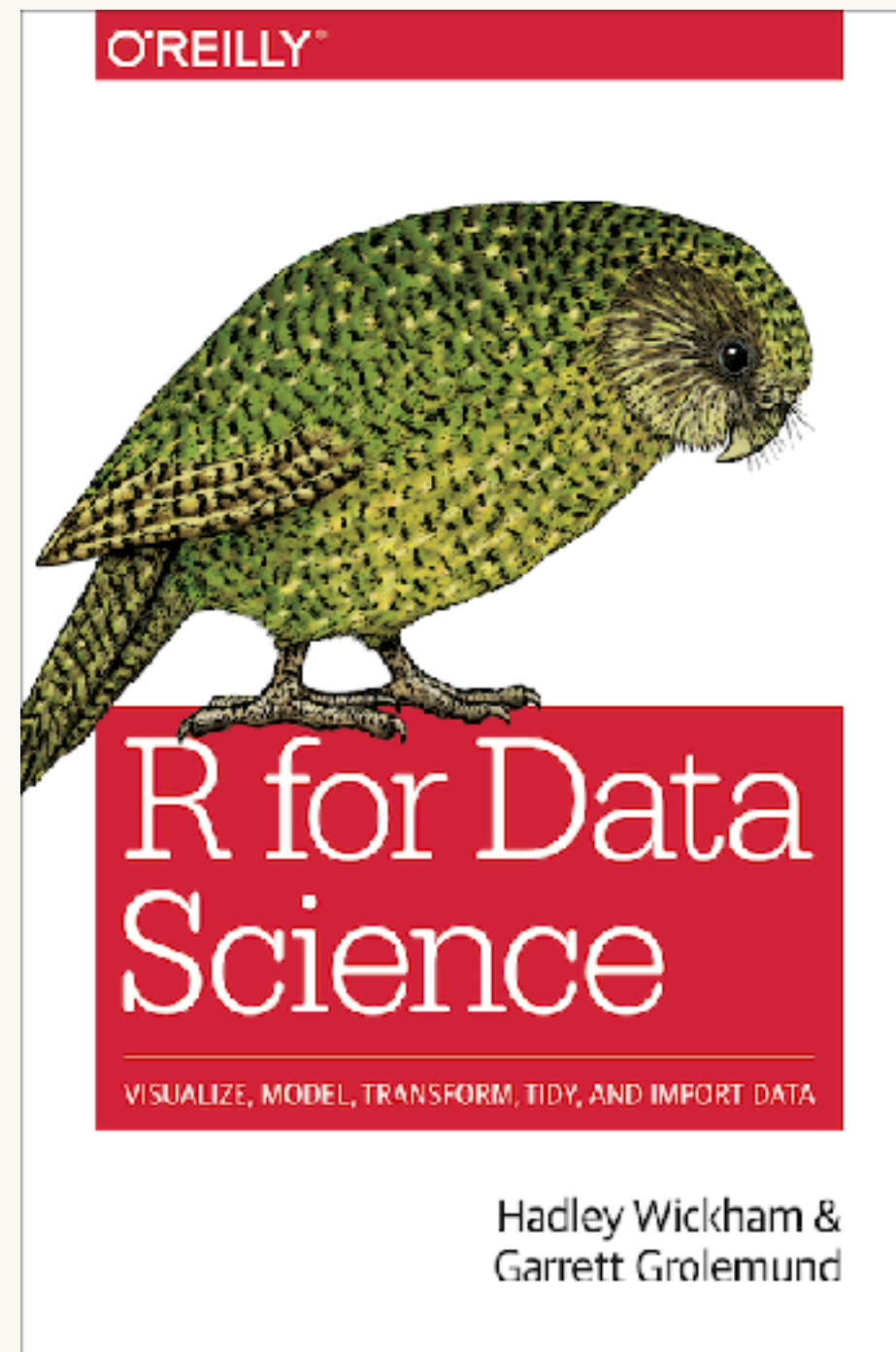
Becomes

```
image_paths <- path("images",  
  paste(files, "n_sales", sep = "_"))
```

```
walk2(image_paths, all_states_plots, ggsave_multiple,  
  exts = c(".pdf", ".png"), height = 3, width = 8)
```

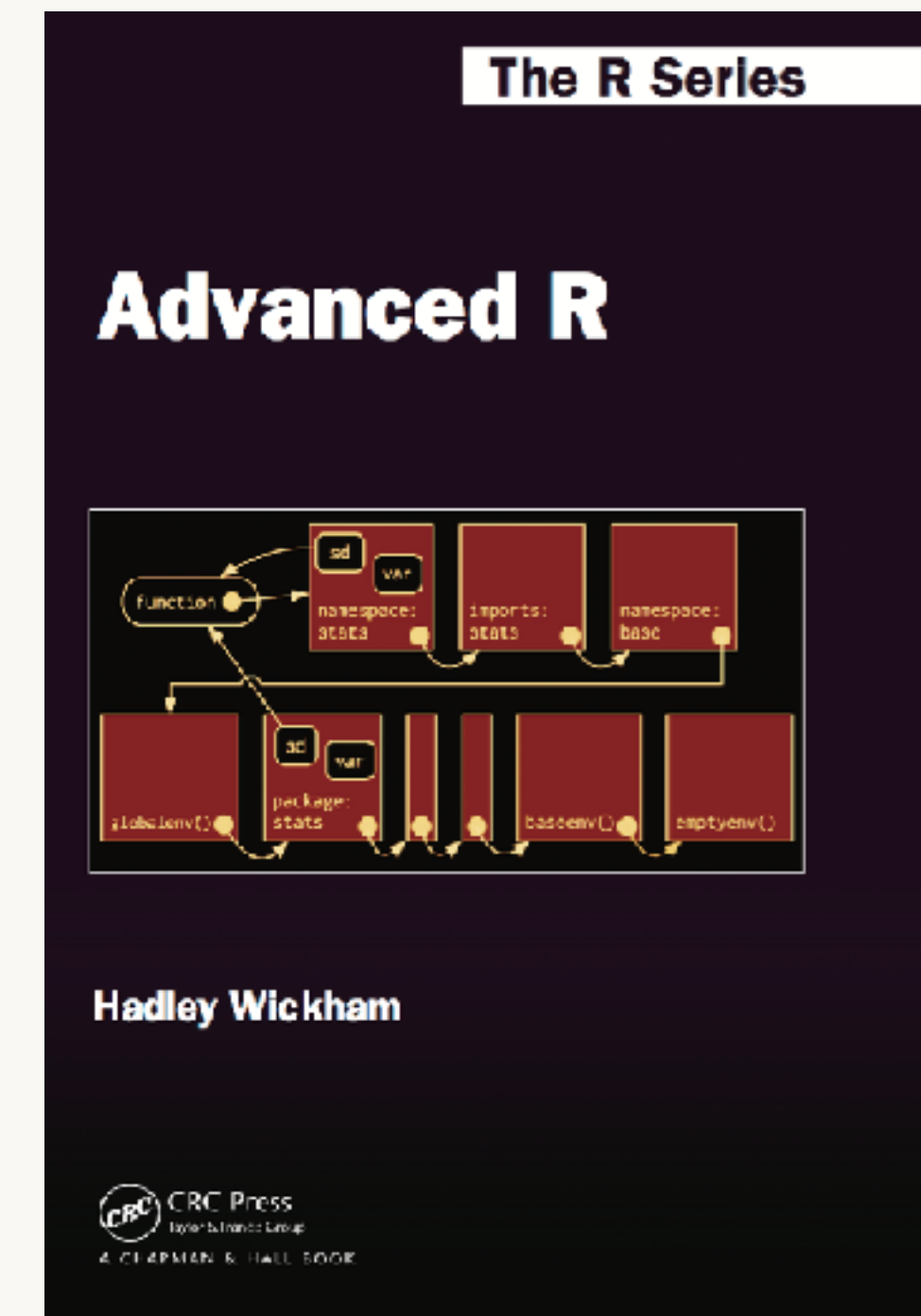
Learning more

Learning more



Iteration

<http://r4ds.had.co.nz/iteration.html>



Functional Programming

<https://adv-r.hadley.nz/functional-programming.html>

Adapted from *Tidy Tools* by Hadley Wickham

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