



WOMBAT 2024: Advanced R Tips & Tricks

Functional programming

workshop.nectric.com.au/advr-wombat24



Outline

1 Functional programming

2 Functional problem solving

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2 Functional problem solving

Functional programming

R is commonly considered a 'functional' programming language - and so far we have used functional programming.

Some programming paradigms in R

Functional programming

- Functions are created and used like any other object.
- Output should only depend on the function's inputs.

Some programming paradigms in R

Functional programming

- Functions are created and used like any other object.
- Output should only depend on the function's inputs.

Object-oriented programming

- Functions are associated with object types.
- Methods of the same 'function' produce object-specific output.

Some programming paradigms in R

Literate programming

- Natural language is interspersed with code.
- Aimed at prioritising documentation/comments.
- Now used to create reproducible reports/documents.

Some programming paradigms in R

Literate programming

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- Now used to create reproducible reports/documents.

Reactive programming

- Objects are expressed using code based on inputs.
- When inputs change, the object's value updates.

Functions are objects

```
square <- function(x) {  
  return(x^2)  
}  
square(8)
```

```
[1] 64
```

The square function is an object like any other in R.

Functions are objects

R functions can be printed,

```
print(square)
```

```
function(x) {  
  return(x^2)  
}
```

Functions are objects

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```
print(square)
```

```
function(x) {  
  return(x^2)  
}
```

inspected,

```
formals(square)
```

```
$x
```

Functions are objects

put in a list,

```
my_functions <- list(square, sum, min, max)
my_functions
```

```
[[1]]
```

```
function(x) {
  return(x^2)
}
```

```
[[2]]
```

```
function (... , na.rm = FALSE) .Primitive("sum")
```

```
[[3]]
```

```
function (... , na.rm = FALSE) .Primitive("min")
```

```
[[4]]
```

```
function (... , na.rm = FALSE) .Primitive("max")
```

Functions are objects

used within lists,

```
my_functions[[1]](8)
```

```
[1] 64
```

Functions are objects

used within lists,

```
my_functions[[1]](8)
```

```
[1] 64
```

but they can't be subsetted!

```
square$x
```

```
Error in square$x: object of type 'closure' is not subsettable
```

Handling input types

Functional programming handles different input types using control flow. The same code is ran regardless of object type.

```
square <- function(x) {  
  if(!is.numeric(x)) {  
    stop("`x` needs to be numeric")  
  }  
  return(x^2)  
}
```

What are functions?


A function is comprised of three components:

- The arguments/inputs (`formals()`)
- The body/code (`body()`)
- The environment (`environment()`)

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A function is comprised of three components:

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- The environment (`environment()`)


 Your turn!

Use these functions to take a closer look at `square()`.
Try modifying the function's `formals/body/env` with `<-`.

Functional programming

Since functions are like any other object, they can also be:

- **inputs** to functions


 Extensible design with function inputs

Using function inputs can improve your package's design! Rather than limiting users to a few specific methods, allow them to use and write any method with functions.

Function arguments

Consider a function which calculates accuracy measures:

```
accuracy <- function(e, measure, ...) {  
  if (measure == "mae") {  
    mean(abs(e), ...)  
  } else if (measure == "rmse") {  
    sqrt(mean(e^2, ...))  
  } else {  
    stop("Unknown accuracy measure")  
  }  
}
```


 Improving the design

This function is limited to only computing MAE and RMSE.

Function arguments

Using function operators allows any measure to be used.

```
MAE <- function(e, ...) mean(abs(e), ...)
RMSE <- function(e, ...) sqrt(mean(e^2, ...))
accuracy <- function(e, measure, ...) {
  ???
}
accuracy(rnorm(100), measure = RMSE)
```


 Your turn!

Complete the accuracy function to calculate accuracy statistics based on the function passed in to `measure`.

Functional programming

Since functions are like any other object, they can also be:

- **inputs** to functions
- **outputs** of functions

 Functions making functions?

These functions are known as *function factories*.

Where have you seen a function that creates a function?

Function factories


Let's generalise `square()` to raise numbers to any power.

```
power <- function(x, exp) {  
  x^exp  
}  
power(8, exp = 2)
```

```
[1] 64
```

```
power(8, exp = 3)
```

```
[1] 512
```

 Starting a factory

What if the function returned a function instead?

Function factories

```
power_factory <- function(exp) {  
  # R is lazy and won't look at exp unless we ask it to  
  force(exp)  
  # Return a function, which finds exp from this environment  
  function(x) {  
    x^exp  
  }  
}  
square <- power_factory(exp = 2)  
square(8)
```

```
[1] 64
```

Function factories

```
power_factory <- function(exp) {  
  # R is lazy and won't look at exp unless we ask it to  
  force(exp)  
  # Return a function, which finds exp from this environment  
  function(x) {  
    x^exp  
  }  
}  
square <- power_factory(exp = 2)  
square(8)
```

```
[1] 64
```


```
cube <- power_factory(exp = 3)  
cube(8)
```

```
[1] 512
```


Function factories

Consider this function to calculate plot breakpoints of vectors.

```
breakpoints <- function(x, n.breaks) {  
  seq(min(x), max(x), length.out = n.breaks)  
}
```

 Your turn!

Convert this function into a function factory.

Is it better to create functions via `x` or `n.breaks`?

Outline

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Split, apply, combine

Many problems can be simplified/solved using this process:

- split (break the problem into smaller parts)
- apply (solve the smaller problems)
- combine (join solved parts to solve original problem)

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Many problems can be simplified/solved using this process:

- split (break the problem into smaller parts)
- apply (solve the smaller problems)
- combine (join solved parts to solve original problem)

This technique applies to both

- writing functions (rewriting a function into sub-functions)
- working with data (same function across groups or files)

`data |> group_by() |> summarise()`

An example of split-apply-combine being used to work with data is when `group_by()` and `summarise()` are used together.

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- **split:** `group_by()` splits up the data into groups
- **apply:** your `summarise()` code calculates a single value
- **combine:** `summarise()` combines the results into a vector

data |> group_by() |> summarise()

An example of split-apply-combine being used to work with data is when `group_by()` and `summarise()` are used together.

- **split:** `group_by()` splits up the data into groups
- **apply:** your `summarise()` code calculates a single value
- **combine:** `summarise()` combines the results into a vector

```
library(dplyr)
mtcars |>
  group_by(cyl) |>
  summarise(mean(mpg))
```

```
# A tibble: 3 x 2
  cyl `mean(mpg)`
<dbl> <dbl>
1     4     26.7
2     6     19.7
3     8     15.1
```

Split-apply-combine for vectors and lists

The same idea can be used for calculations on vectors.

Split-apply-combine for vectors and lists

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There are two main implementations we consider:

- base R: The `*apply()` functions
- purrr: The `map*()` functions

Split-apply-combine for vectors and lists

The same idea can be used for calculations on vectors.

There are two main implementations we consider:

- base R: The `*apply()` functions
- purrr: The `map*()` functions

We will use purrr and but I'll also share the base R equivalent.

for or map?

Let's square() a vector of numbers with a for loop.

```
x <- c(1, 3, 8)
x2 <- numeric(length(x))
for (i in seq_along(x)) {
  x2[i] <- square(x[i])
}
x2
```

```
[1] 1 9 64
```

for or map?

Let's `square()` a vector of numbers with a for loop.

```
x <- c(1, 3, 8)
x2 <- numeric(length(x))
for (i in seq_along(x)) {
  x2[i] <- square(x[i])
}
x2
```

```
[1] 1 9 64
```

Vectorisation?

Of course `square()` is vectorised, so we should use `square(x)`.
Other functions like `lm()` or `read.csv()` are not!

for or map?

Instead using `map()` we get...

```
library(purrr)
x <- c(1, 3, 8)
map(x, square) # lapply(x, square)
```

```
[[1]]
```

```
[1] 1
```

```
[[2]]
```

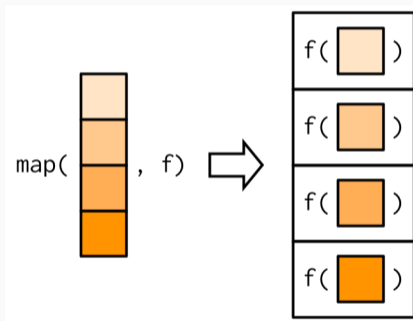
```
[1] 9
```

```
[[3]]
```

```
[1] 64
```

Mapping vectors

The same result, but it has been combined differently!



Mapping vectors

To combine the results into a vector rather than a list, we instead use `map_vec()` to combine results into a vector.

```
library(purrr)
x <- c(1, 3, 8)
map_vec(x, square) # vapply(x, square, numeric(1L))
```

```
[1] 1 9 64
```

Advantages of map

- Less coding (less bugs!)
- Easier to read and understand.

Advantages of map

- Less coding (less bugs!)
- Easier to read and understand.

Disadvantages of map

- Less control over loop
- Cannot solve sequential problems

Functional mapping

Recall `group_by()` and `summarise()` from `dplyr`:

```
mtcars |>  
  group_by(cyl) |>  
  summarise(mean(mpg))
```

Your turn!

Use `split()` and `map_vec()` to achieve a similar result.

Hint: `split(mtcars$mpg, mtcars$cyl)` creates a list that splits `mtcars$mpg` by each value of `mtcars$cyl`.

Anonymous mapper functions

Suppose we want to separately model `mpg` for each `cyl`.

```
lm(mpg ~ disp + hp + drat + wt, mtcars[mtcars$cyl == 4,])  
lm(mpg ~ disp + hp + drat + wt, mtcars[mtcars$cyl == 6,])  
lm(mpg ~ disp + hp + drat + wt, mtcars[mtcars$cyl == 8,])
```

Anonymous mapper functions

We can split the data by `cyl` with `split()`,

```
mtcars_cyl <- split(mtcars, mtcars$cyl)
```

but `map(mtcars_cyl, lm, mpg ~ disp + hp + drat + wt)`
won't work - why?

Anonymous mapper functions

We can split the data by `cyl` with `split()`,

```
mtcars_cyl <- split(mtcars, mtcars$cyl)
```

but `map(mtcars_cyl, lm, mpg ~ disp + hp + drat + wt)`
won't work - why?

! Difficult to map

Using `map(mtcars_cyl, lm)` will apply `lm(mtcars_cyl[i])`.
The mapped vector is always used as the first argument!

Anonymous mapper functions

We can write our own functions!

```
mtcars_lm <- function(.) lm(mpg ~ disp + hp + drat + wt, data = .)
map(mtcars_cyl, mtcars_lm)
```

```
$`4`
```

Call:

```
lm(formula = mpg ~ disp + hp + drat + wt, data = .)
```

Coefficients:

(Intercept)	disp	hp	drat	wt
52.51953	-0.06294	-0.07602	-1.44216	-3.10007

```
$`6`
```

Anonymous mapper functions

Or use `~ body` to create anonymous functions.

```
# lapply(mtcars_cyl, \(.) lm(mpg ~ disp + hp + drat + wt, data = .))  
map(mtcars_cyl, ~ lm(mpg ~ disp + hp + drat + wt, data = .))
```

```
$`4`
```

Call:

```
lm(formula = mpg ~ disp + hp + drat + wt, data = .)
```

Coefficients:

(Intercept)	disp	hp	drat	wt
52.51953	-0.06294	-0.07602	-1.44216	-3.10007

```
$`6`
```

Mapping mapping mapping

How would you then get the coefficients from all 3 models?

```
# mtcars_cyl |> lapply(\(.) lm(mpg ~ disp + hp + drat + wt, data = .))  
mtcars_cyl |>  
  map(~ lm(mpg ~ disp + hp + drat + wt, data = .))
```


Mapping mapping mapping

How would you then get the coefficients from all 3 models?

```
# mtcars_cyl |> lapply(\(.) lm(mpg ~ disp + hp + drat + wt, data = .))
mtcars_cyl |>
  map(~ lm(mpg ~ disp + hp + drat + wt, data = .))
```

Solution

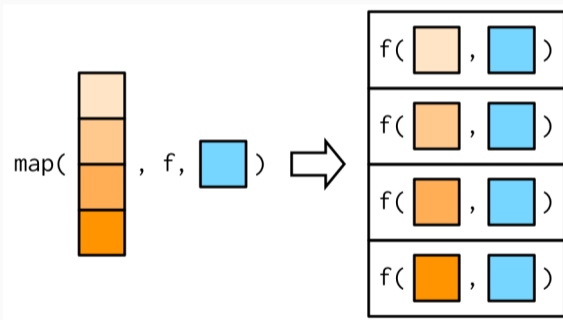
```
# lapply(mtcars_cyl, \(.) lm(mpg ~ disp + hp + drat + wt, data = .))
mtcars_cyl |>
  map(~ lm(mpg ~ disp + hp + drat + wt, data = .)) |>
  map(coef)
```

```
$`4`
```

```
(Intercept)          disp          hp          drat          wt
52.51952502 -0.06293845 -0.07601929 -1.44215918 -3.10006904
```

Mapping arguments

Any arguments after your function are passed to all functions.



Mapping arguments

This works by passing through ... to the function.

```
x <- list(1:5, c(1:10, NA))  
map_dbl(x, ~ mean(.x, na.rm = TRUE))
```

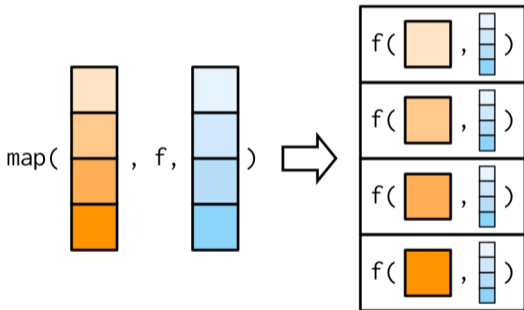
```
[1] 3.0 5.5
```

```
map_dbl(x, mean, na.rm = TRUE)
```

```
[1] 3.0 5.5
```

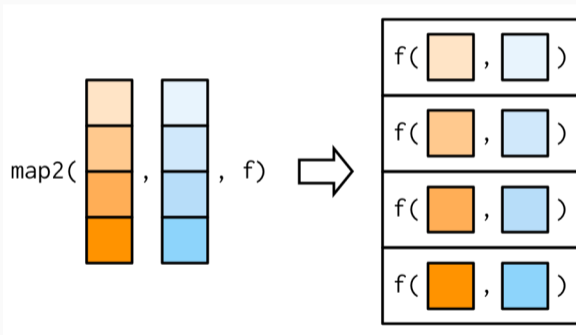
Mapping arguments

These additional arguments are not decomposed / mapped.



Mapping multiple arguments

It is often useful to map multiple arguments.



Mapping multiple arguments

```
xs <- map(1:8, ~ ifelse(runif(10) > 0.8, NA, runif(10)))  
map_vec(xs, mean, na.rm = TRUE)
```

```
[1] 0.5298053 0.4535120 0.4972954 0.3761379 0.5603879 0.42  
[8] 0.4608077
```

Mapping multiple arguments

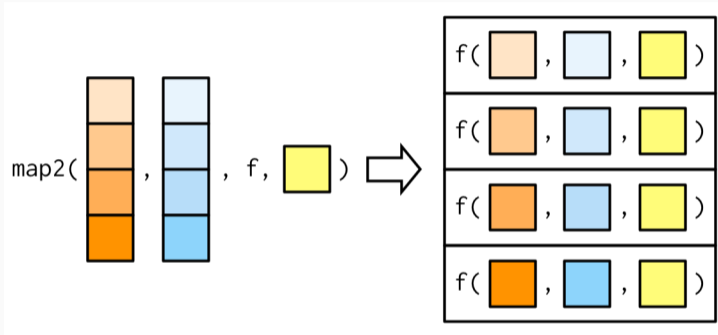
```
xs <- map(1:8, ~ ifelse(runif(10) > 0.8, NA, runif(10)))  
map_vec(xs, mean, na.rm = TRUE)
```

```
[1] 0.5298053 0.4535120 0.4972954 0.3761379 0.5603879 0.42  
[8] 0.4608077
```

```
ws <- map(1:8, ~ rpois(10, 5) + 1)  
map2_vec(xs, ws, weighted.mean, na.rm = TRUE)
```

```
[1] 0.5199651 0.4452852 0.4631680 0.3489870 0.5464348 0.49  
[8] 0.4804868
```

Mapping multiple arguments



Mapping many arguments

It is also possible to map any number of inputs with `pmap`.

```
n <- 1:3
min <- c(0, 10, 100)
max <- c(1, 100, 1000)
pmap(list(n, min, max), runif) # .mapply(runif, list(n, min, max), list())
```

```
[[1]]
```

```
[1] 0.8066672
```

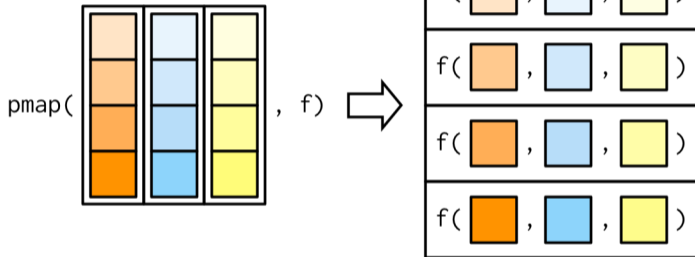
```
[[2]]
```

```
[1] 35.75897 52.32907
```

```
[[3]]
```

```
[1] 751.5277 596.4991 941.6216
```

Mapping many arguments



Parallel mapping

Split-apply-combine problems are *embarrassingly parallel*.

Parallel mapping

Split-apply-combine problems are *embarrassingly parallel*.

The `furrr` package (`future` + `purrr`) makes it easy to use `map()` in parallel, providing `future_map()` variants.

```
library(furrr)
plan(multisession, workers = 4)
future_map_dbl(xs, mean, na.rm = TRUE)
```

```
[1] 0.5298053 0.4535120 0.4972954 0.3761379 0.5603879 0.42
[8] 0.4608077
```

```
future_map2_dbl(xs, ws, weighted.mean, na.rm = TRUE)
```

```
[1] 0.5199651 0.4452852 0.4631680 0.3489870 0.5464348 0.49
```

Reduce vectors to single values

Sometimes you want to collapse a vector, reducing it to a single value. `reduce()` always returns a vector of length 1.

```
x <- sample(1:100, 10)
x
```

```
[1] 70 42 35 61 85 81 77 65 68 40
```

```
sum(x)
```

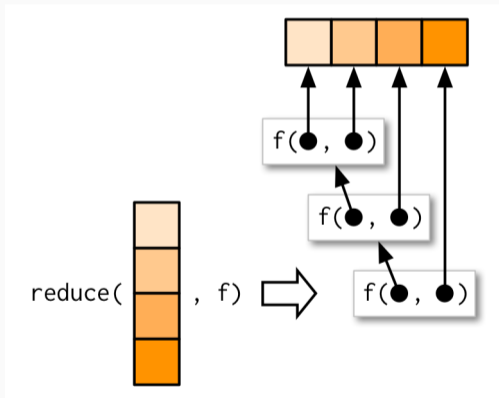
```
[1] 624
```

```
# Alternative to sum()
reduce(x, `+`) # Reduce(`+`, x)
```

```
[1] 624
```

Reduce vectors to single values

The result from the function is re-used as the first argument.



Reduce vectors to single values

🔥 Your turn!

We're studying the letters in 3 bowls of alphabet soup.



Reduce vectors to single values

Your turn!

We're studying the letters in 3 bowls of alphabet soup.
Use `reduce()` to find the letters were in all bowls of soup!
Are all letters found in the soups?

```
alphabet_soup <- map(c(10,24,13), sample, x=letters, replace=TRUE)
alphabet_soup
```

```
[[1]]
```

```
[1] "k" "h" "a" "h" "b" "e" "k" "x" "c" "y"
```

```
[[2]]
```

```
[1] "k" "e" "d" "m" "k" "r" "w" "e" "d" "o" "k" "y" "p" "u" "u" "n" "r" "u" "f"
[20] "a" "m" "k" "q" "d"
```


Functional adverbs

purrr also offers many *adverbs*, which modify a function.

Capturing conditions

- `possibly(.f, otherwise)`: If the function errors, it will return `otherwise` instead.
- `safely(.f)`: The function now returns a list with 'result' and 'error', preventing errors.
- `quietly(.f)`: Any conditions (messages, warnings, printed output) are now captured into a list.

Functional adverbs

purrr also offers many *adverbs*, which modify a function.

Capturing conditions

- `negate(.f)` will return `!result`.

Chaining functions

- `compose(...)` will chain functions together like a chain of piped functions.

Functional adverbs

purrr also offers many *adverbs*, which modify a function.

💡 Functions modifying functions?

These functions are all *function factories*!

More specifically they are known as *function operators* since both the input and output is a function.

`memoise::memoise()` is also a *function operator*.