

# ETC4500/ETC5450

## Advanced R programming

Week 5: Functional programming



# Programming paradigms

R code is typically structured using these paradigms:

- Functional programming
- Object-oriented programming
- Literate programming
- Reactive programming

Often several paradigms used together to solve a problem.

# Programming paradigms

## Functional programming (W5; today!)

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

# Programming paradigms

## Functional programming (W5; today!)

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

## Literate programming (W6)

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

# Programming paradigms

## Reactive programming (W7)

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

# Programming paradigms

## Reactive programming (W7)

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

## Object-oriented programming (W8 - W9)

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

# Functional programming

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

```
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

R functions can be printed,

```
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)  
}
```

# Handling input types

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

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

💡 Later in the semester...

We will see object-oriented programming, which handles different input types using different functions (methods)!

# 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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- 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?

# 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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- 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.

## `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

# 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.

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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)
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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)
```

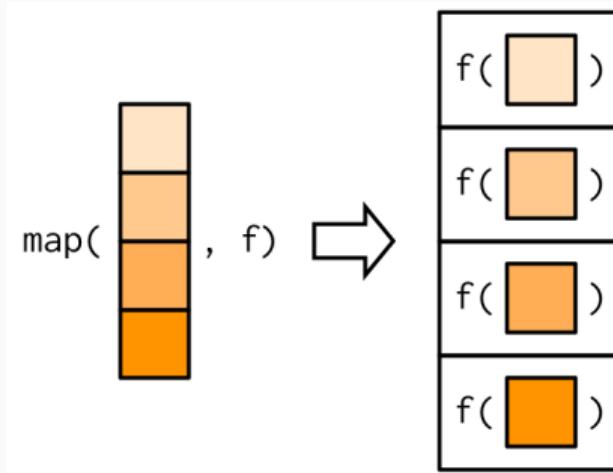
```
[[1]]
[1] 1
```

```
[[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
```

# for or map



## Advantages of map

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

# for or map

## 💡 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
52.5195	-0.0629	-0.0760	-1.4422	-3.1001

\$`6`

Call:

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

# 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
52.5195	-0.0629	-0.0760	-1.4422	-3.1001

\$`6`

Call:

```
lm(formula = 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 = .))
```

# 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

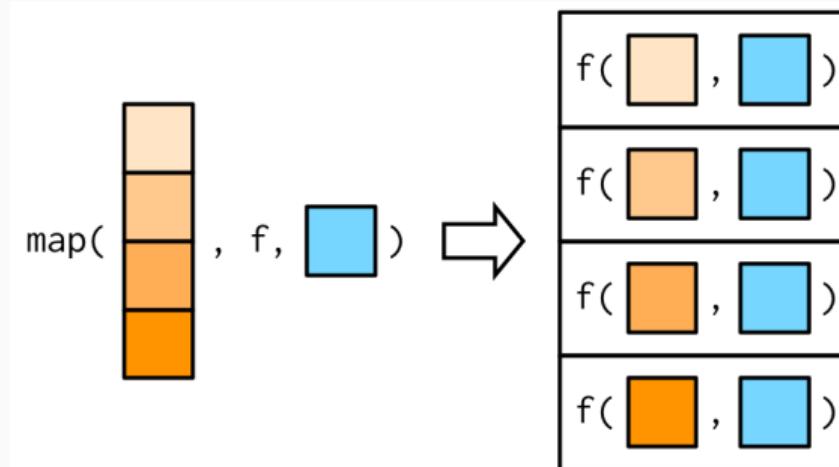
```
# lapply(mtcars_cyl, \(.) lm(mpg ~ disp + hp + drat + wt, data = .))  
mtcars_cyl |>  
  map(~ lm(mpg ~ disp + hp + drat + wt, data = .)) |>  
  map(coef)
```

```
$`4`  
(Intercept)          disp          hp          drat          wt  
      52.5195     -0.0629     -0.0760     -1.4422     -3.1001
```

```
$`6`
```

# 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))
```

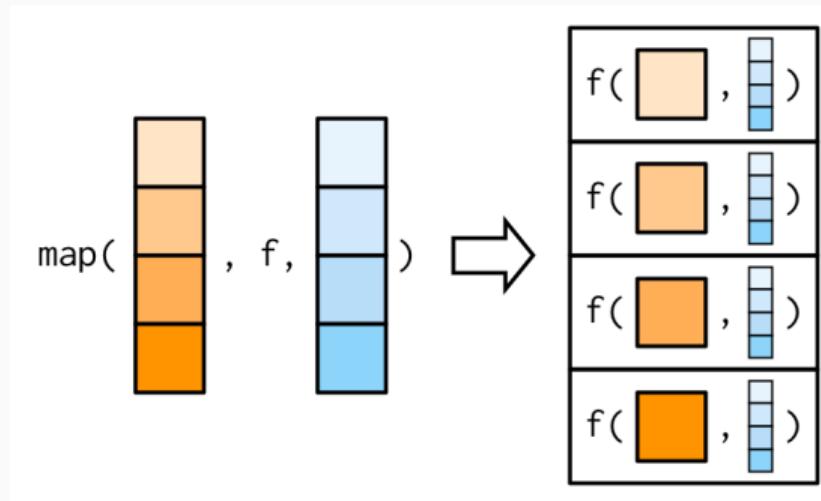
```
[1] 3.0 5.5
```

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

```
[1] 3.0 5.5
```

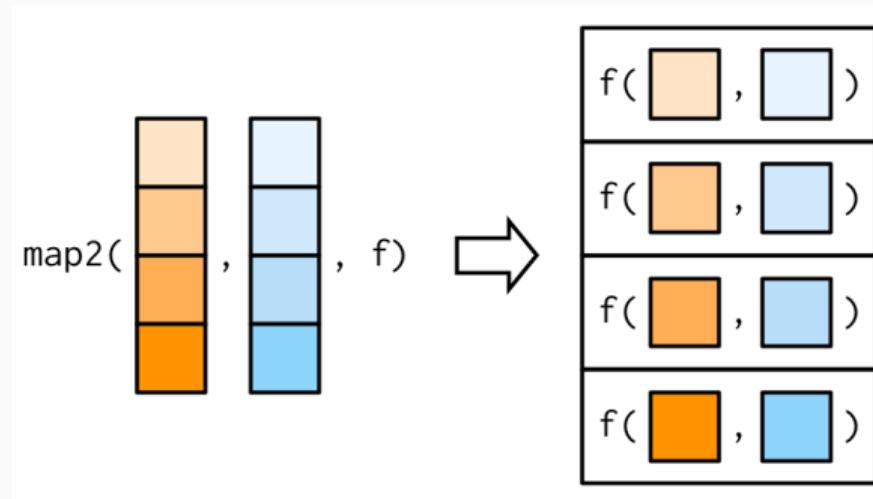
# Mapping arguments

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] 0.598 0.427 0.444 0.411 0.488 0.505 0.313 0.492
```

# Mapping multiple arguments

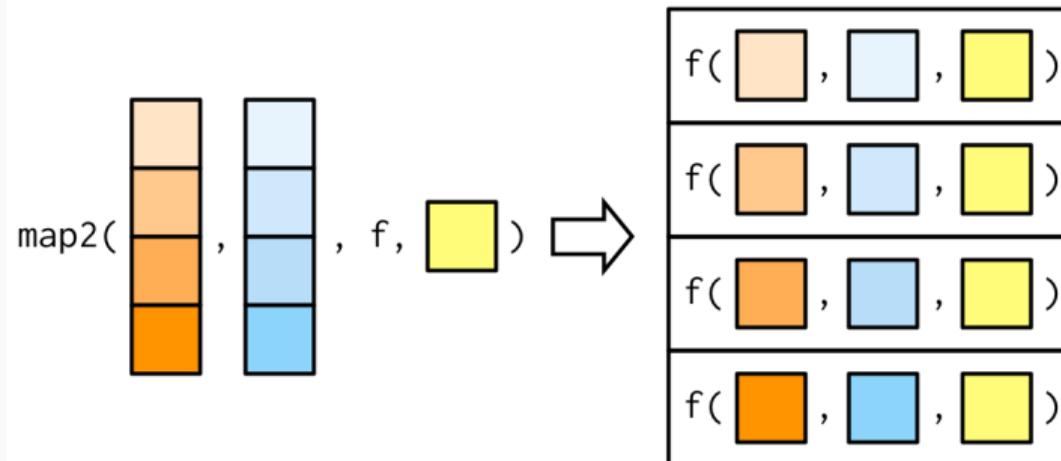
```
xs <- map(1:8, ~ ifelse(runif(10) > 0.8, NA, runif(10)))
map_vec(xs, mean, na.rm = TRUE)
```

```
[1] 0.598 0.427 0.444 0.411 0.488 0.505 0.313 0.492
```

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

```
[1] 0.595 0.408 0.399 0.460 0.445 0.531 0.318 0.452
```

# Mapping multiple arguments



# Mapping many arguments

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

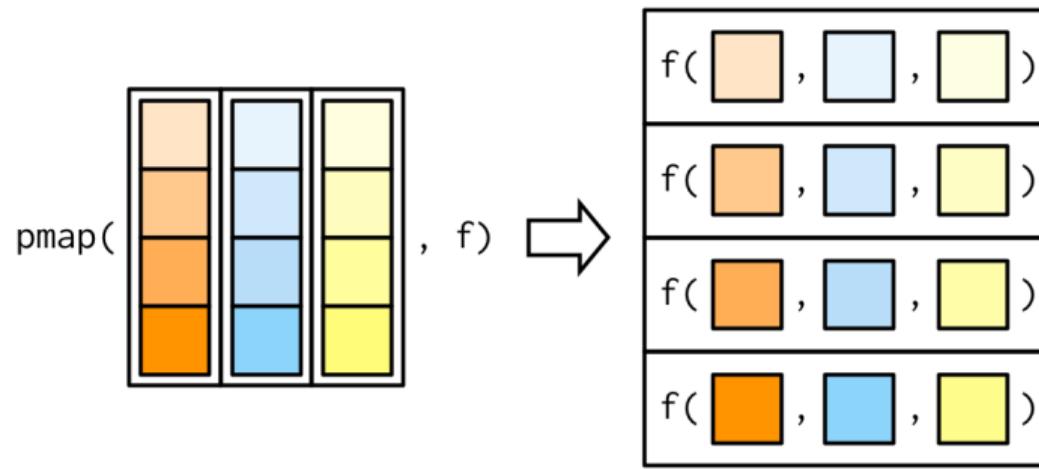
```
n <- 1:3
min <- c(0, 10, 100)
max <- c(1, 100, 1000)
pmap(list(n, min, max), runif) # .mapply(runif, list(n, min, max), list())
```

```
[[1]]
[1] 0.619
```

```
[[2]]
[1] 33.0 40.2
```

```
[[3]]
[1] 140 101 794
```

# 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] 0.598 0.427 0.444 0.411 0.488 0.505 0.313 0.492
```

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

```
[1] 0.595 0.408 0.399 0.460 0.445 0.531 0.318 0.452
```

# 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] 31 16 76 33 2 20 69 24 53 80
```

```
sum(x)
```

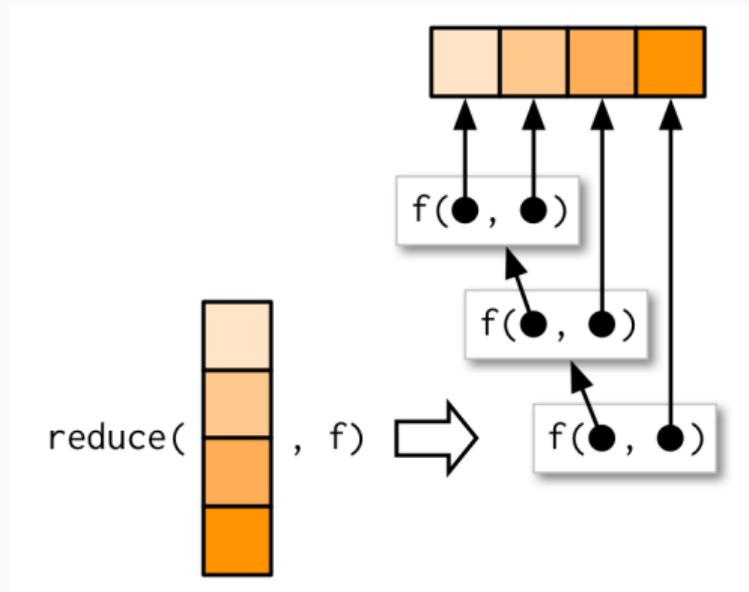
```
[1] 404
```

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

```
[1] 404
```

# 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] "j" "v" "o" "l" "t" "g" "u" "z" "d" "f"
```

```
[[2]]  
[1] "v" "t" "s" "j" "a" "b" "c" "g" "j" "v" "h" "f" "p" "a" "q" "r" "u"  
[18] "v" "h" "v" "r" "h" "w" "f"
```

```
[[3]]
```

# 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.

## Changing results

- `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*.