

# ETC4500/ETC5450

## Advanced R programming

Week 5: Functional programming



# Outline

- 1 Programming paradigms
- 2 Functional programming
- 3 Functional problem solving

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# 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.

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# 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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A function is comprised of three components:

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

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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:

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- 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.



## data |> group\_by() |> summarise()

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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.

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- 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

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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.

# 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] 0.533 0.492 0.613 0.446 0.414 0.437 0.516 0.316
```

# Mapping multiple arguments

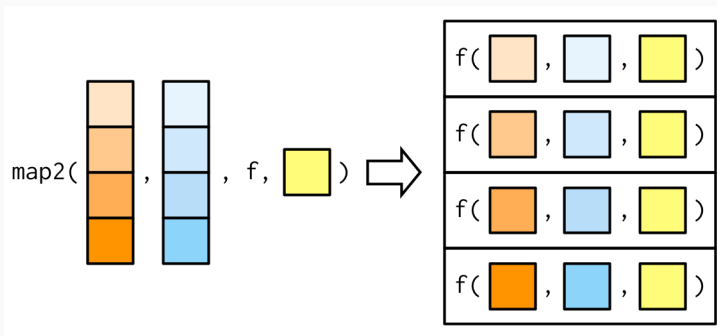
```
xs <- map(1:8, ~ ifelse(runif(10) > 0.8, NA, runif(10)))  
map_vec(xs, mean, na.rm = TRUE)
```

```
[1] 0.533 0.492 0.613 0.446 0.414 0.437 0.516 0.316
```

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

```
[1] 0.511 0.513 0.585 0.460 0.415 0.441 0.524 0.271
```

# 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] 0.558
```

```
[[2]]
```

```
[1] 14.9 87.8
```

```
[[3]]
```

```
[1] 934 407 181
```

# 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] 0.533 0.492 0.613 0.446 0.414 0.437 0.516 0.316
```

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

```
[1] 0.511 0.513 0.585 0.460 0.415 0.441 0.524 0.271
```

# 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] 46 43  8 39 95 85 29 15 16 32
```

```
sum(x)
```

```
[1] 408
```

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

```
[1] 408
```



# 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] "b" "d" "l" "m" "k" "f" "a" "e" "o" "r"
```

```
[[2]]
```

```
[1] "s" "t" "t" "a" "e" "g" "y" "w" "x" "z" "x" "o" "x" "t" "n" "c" "s"  
[18] "c" "t" "q" "o" "b" "f" "i"
```

```
[[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*.