Functional Programming

Data Wrangling in R

Functional Programming

"R, at its heart, is a functional programming (FP) language. This means that it provides many tools for the creation and manipulation of functions. In particular, R has what's known as first class functions. You can do anything with functions that you can do with vectors: you can assign them to variables, store them in lists, pass them as arguments to other functions, create them inside functions, and even return them as the result of a function." - Hadley Wickham

Don't need to write for-loops! - check this video.

Allows you to flexibly iterate functions to multiple elements of a data object!

Useful when you want to apply a function to:

- * lots of columns in a tibble
- * multiple tibbles
- * multiple data files
- * or perform fancy functions with vectors (or tibble columns)

Working across multiple columns

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Mazda RX4

Say we wanted to round multiple columns of the mtcars data. We could do so one column at a time, or we could use the across function from the dplyr package. Needs to be used within other dplyr functions such as mutate.

mutate (across (which columns, which function or operation))

6 160 110

Mazda RX4 Wag 21 6 160 110 4 3 17 0 1

4 3 16 0 1

functions in R

```
my_function <- function(x) {x + 1}
my_function

## function(x) {x + 1}
## <environment: 0x10b670f90>

my_data <- c(2,3,4)
my_function(x = my_data)

## [1] 3 4 5

my_function(my_data)

## [1] 3 4 5</pre>
```

Special tilda use

If you see $\sim .x$ or ... this means function (x) $\{x\}$. In other words, we are passing x to a function. See purrr-shortcuts

For example - this is not necessary but you could use it here:

Using across with arguments

If you wish to also pass arguments to the function that you are applying to the various columns, then you need to use the \sim and .x (or .) as a place holder for what you the values you will be passing into the function.

Using across with helpers to apply function to multiple columns

?tidyr tidy select

if_any() and if_all() are also helpful!

Work inside other tidyverse functions, like across ()

```
iris %>% filter(Sepal.Length > 2.4 & Sepal.Width > 2.4 &
              Petal.Length > 2.4 & Petal.Width > 2.4)
##
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
          6.3 3.3 6.0 2.5 virginica
7.2 3.6 6.1 2.5 virginica
6.7 3.3 5.7 2.5 virginica
## 1
## 2
## 3
iris %>% filter(if all(Sepal.Length:Petal.Width, ~ . > 2.4))
##
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
     6.3 3.3 6.0 2.5 virginica
## 1
         7.2 3.6 6.1 2.5 virginica
## 2
## 3 6.7 3.3 5.7 2.5 virginica
iris %>% filter(if all(where(is.numeric), ~ . > 2.4))
##
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1 6.3 3.3 6.0 2.5 virginica
## 2 7.2 3.6 6.1 2.5 virginica
## 3 6.7 3.3 5.7 2.5 virginica
```

Previously we filtered for patterns or conditions..

Seems a bit repetitive!

Now we can filter multiple columns!

Multiple conditions simultaneously!

purrr is also a super helpful package!

"Designed to make your functions purrr."

dplyr is designed for data frames purrr is designed for vectors

The purrr package can be very helpful!

- https://purrr.tidyverse.org/
- https://github.com/rstudio/cheatsheets/raw/master/purrr.pdf
- https://jennybc.github.io/purrr-tutorial/

purrr main functions

map and map * and modify

- **applies function** to each element of an vector or object (map returns a list, modify returns the same object type)

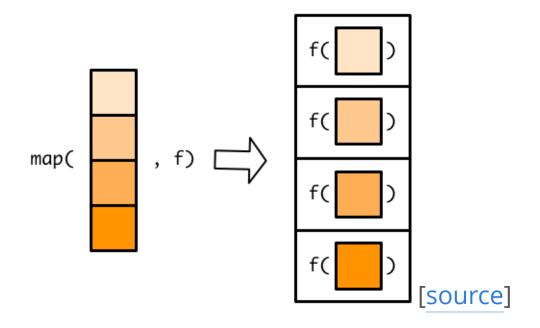
map2 and map2*

- applies function to each element of **two** vectors or objects

pmap and pmap_* - applies function to each element of **3+** vector or objects (requires a list for input)

the _* options specify the type of data output

map (and modify)



```
x <-c(1.2, 2.3, 3.5, 4.6)
map(x, round) %>% unlist()
```

[1] 1 2 4 5

map (and modify)

```
x < -tibble(values = c(1.2, 2.3, 3.5, 4.6))
map df(x, round)
## # A tibble: 4 × 1
## values
## <dbl>
## 1
## 2
## 3 4
## 4
modify(x, round)
## # A tibble: 4 × 1
## values
## <dbl>
## 1
## 2
## 3
## 4
```

purrr - apply function to all columns

two options map_df or modify

Lots of variations of map based on output

```
library (purrr)
head (mtcars, 2)
                                                                mpg cyl disp hp drat wt qsec vs am qear carb
##
## Mazda RX4 21 6 160 110 3.9 2.620 16.46 0 1
## Mazda RX4 Wag 21 6 160 110 3.9 2.875 17.02 0 1 4
   mtcars %>%
           map df(round) %>% # will be a tibble now - will remove rownames
          head (2)
## # A tibble: 2 × 11
 ##
                          mpg cyl disp hp drat wt gsec
                                                                                                                                                                                                 VS
                                                                                                                                                                                                                          am gear carb
## <dbl> <db
                                                          6 160
## 1 21
                                                                                                110
                                                                                                                               4 3
                                                                                                                                                                         16
                                                                                                                                                                                                                                                                             4
                                                                                                                              4 3
## 2 21 6 160 110
                                                                                                                                                                          17
   mtcars %>%
           modify(round) %>% # modify keeps original data type
           head (2)
                                                                                                                                                                                                                                                                                15/34
##
                   mpg cyl disp hp drat wt gsec vs am gear carb
```

6 160 110 4 3 16 0 1 4

It's a bit simpler than across...

purrr apply function to some columns like accross

Using modify_if() or map_if(), we can specify what columns to modify

```
head(as tibble(iris), 3)
## # A tibble: 3 \times 5
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
    <dbl> <dbl> <dbl> <dbl> <fct>
         5.1 3.5 1.4 0.2 setosa
4.9 3 1.4 0.2 setosa
4.7 3.2 1.3 0.2 setosa
## 1
## 2
## 3
as tibble(iris) %>%
 modify if (is.numeric, as.character) %>%
 head(3)
## # A tibble: 3 \times 5
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
as tibble(iris) %>%
 map if(is.numeric, as.character) %>%
 class()
```

modify_if vs mutate/across/where

What is a 'list'?

- Lists are the most flexible/"generic" data class in R
- Can be created using list()
- · Can hold vectors, strings, matrices, models, list of other lists, lists upon lists!
- · Can reference data using \$ (if the elements are named), or using [], or [[]]

```
> mylist <- list(
+ letters = c("A", "b", "c"),
+ numbers = 1:3,
+ matrix(1:25, ncol = 5),
+ matrix(1:25, ncol = 5)
+ )</pre>
```

List Structure

```
> head(mylist)
$letters
[1] "A" "b" "c"
$numbers
[1] 1 2 3
[[3]]
     [,1] [,2] [,3] [,4] [,5]
[1,]
                 11
                            21
                           22
               12
[2,]
                      17
           8 13
[3,]
                      18
                           23
               14
                      19
                           24
[4,]
            10
                 15
                       20
                            25
[5,]
[[4]]
     [,1] [,2] [,3] [,4] [,5]
[1,]
                 11
                            21
               12
                           22
                       17
[2,]
[3,]
               13
                      18
                           23
                14
                      19
                           24
[4,]
            10
                 15
                            25
                       20
[5,]
```

List referencing

```
> mylist[1] # returns a list

$letters
[1] "A" "b" "c"

> mylist["letters"] # returns a list

$letters
[1] "A" "b" "c"
```

List referencing

```
> mylist[[1]] # returns the vector 'letters'

[1] "A" "b" "c"
> mylist$letters # returns vector

[1] "A" "b" "c"
> mylist[["letters"]] # returns the vector 'letters'

[1] "A" "b" "c"
->
```

List referencing

You can also select multiple lists with the single brackets.

```
> mylist[1:2] # returns a list
$letters
[1] "A" "b" "c"
$numbers
[1] 1 2 3
```

... Why lists?

great example with split()

```
head (mtcars)
##
                     mpg cyl disp hp drat wt qsec vs am gear carb
                    21.0
## Mazda RX4
                              160 110 3.90 2.620 16.46
## Mazda RX4 Wag 21.0 6
                              160 110 3.90 2.875 17.02
                     22.8 4
                              108
                                    93 3.85 2.320 18.61
## Datsun 710
                     21.4 6 258 110 3.08 3.215 19.44
## Hornet 4 Drive
## Hornet Sportabout 18.7 8 360 175 3.15
                                           3.440 17.02
                         6 225 105 2.76 3.460 20.22 1
## Valiant
                    18.1
str(mtcars %>% split(.$cyl))
## List of 3
##
    $ 4:'data.frame': 11 obs. of 11 variables:
     ..$ mpg : num [1:11] 22.8 24.4 22.8 32.4 30.4 33.9 21.5 27.3 26 30.4 ...
##
##
     ..$ cyl : num [1:11] 4 4 4 4 4 4 4 4 4 4
     ..$ disp: num [1:11] 108 146.7 140.8 78.7 75.7 ...
##
##
     ..$ hp : num [1:11] 93 62 95 66 52 65 97 66 91 113 ...
     ..$ drat: num [1:11] 3.85 3.69 3.92 4.08 4.93 4.22 3.7 4.08 4.43 3.77 ...
##
##
                          2.32 3.19 3.15 2.2 1.61 ...
     ..$ wt : num [1:11]
##
     ..$ qsec: num [1:11]
                          18.6
                                 22.9 19.5
##
     ..$ vs : num [1:11]
##
     ..$ am : num [1:11]
##
     ..$ gear: num [1:11]
                              4 4
     ..$ carb: num [1:11] 1 2 2 1
##
    $ 6:'data.frame':
                       7 obs. of
                                 11 variables:
##
     ..$ mpg : num [1:7] 21 21 21.4 18.1 19.2 17.8 19.7
                                                                      25/34
##
     ..$ cyl : num [1:7]
                         6 6 6 6 6 6
```

great example with split()

```
mtcars %>%
   split(.$cyl) %>% # creates split of data for each unique cyl value
   map(~lm(mpg ~ wt, data = .)) %>% # apply linear model to each
   map(summary) %>%
   map_dbl("r.squared")

## 4 6 8
## 0.5086326 0.4645102 0.4229655
```

Lists from multiple files!

This comes up a lot in data cleaning and also when reading in multiple files!

```
library(here)
library(readr)
list.files(here::here("data", "iris"), pattern = "*.csv")
## [1] "iris q1.csv" "iris q4.csv" "iris q5.csv"
file list <- paste0(here::here(), "/data/iris/", list.files(here::here("data", "iris"), pattern = "*.csv"))
file list
## [1] "/Users/avahoffman/Dropbox/JHSPH/Data-Wrangling SISBID/data/iris/iris q1.csv"
## [2] "/Users/avahoffman/Dropbox/JHSPH/Data-Wrangling SISBID/data/iris/iris q4.csv"
## [3] "/Users/avahoffman/Dropbox/JHSPH/Data-Wrangling SISBID/data/iris/iris q5.csv"
multifile data <- file list %>%
 map(read csv)
class(multifile_data)
## [1] "list"
```

multifile_data[[1]]

```
## # A tibble: 150 × 5
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
            <dbl>
                        <dbl>
                                                 <dbl> <chr>
                                     <dbl>
## 1
              5.1
                          3.5
                                       1.4
                                                   0.2 setosa
## 2
              4.9
                          3
                                       1.4
                                                  0.2 setosa
              4.7
                          3.2
                                       1.3
                                                  0.2 setosa
## 3
              4.6
                          3.1
                                       1.5
                                                   0.2 setosa
                          3.6
## 5
                                       1.4
                                                   0.2 setosa
## 6
                          3.9
                                       1.7
                                                   0.4 setosa
              5.4
## 7
              4.6
                          3.4
                                       1.4
                                                  0.3 setosa
                                                   0.2 setosa
## 8
                          3.4
                                       1.5
              5
                          2.9
                                       1.4
             4.4
                                                   0.2 setosa
## 10
              4.9
                          3.1
                                                   0.1 setosa
                                       1.5
## # i 140 more rows
```

multifile_data[[2]]

multifile_data[[3]]

```
## # A tibble: 150 × 5
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
                                                <dbl> <chr>
##
           <dbl>
                        <dbl>
                                     <dbl>
                                                  0.2 setosa
## 1
          -999
                          3.5
                                      1.4
## 2
                          3
                                      1.4
                                                  0.2 setosa
          -999
## 3
          -999
                          3.2
                                      1.3
                                                  0.2 setosa
             4.6
                          3.1
                                      1.5
                                                  0.2 setosa
                          3.6
                                      1.4
                                                  0.2 setosa
## 6
                          3.9
                                      1.7
                                                  0.4 setosa
             5.4
             4.6
                          3.4
                                      1.4
                                                  0.3 setosa
                          3.4
                                      1.5
## 8
              5
                                                  0.2 setosa
                          2.9
                                      1.4
             4.4
                                                  0.2 setosa
## 10
             4.9
                          3.1
                                                  0.1 setosa
                                      1.5
## # i 140 more rows
```

Fixing the second file

First, separating by the :.

2 4.9

3 4.7

3.2

```
multifile data[[2]] <-</pre>
  separate(
   multifile data[[2]],
   col = `Sepal.Length:Sepal.Width:Petal.Length:Petal.Width:Species`,
   into = c(
     "Sepal.Length",
     "Sepal.Width",
     'Petal.Length',
     "Petal.Width",
     "Species"
   sep = ":"
head(multifile data[[2]], 3)
## # A tibble: 3 × 5
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## <chr> <chr>
                            <chr>
                                        <chr>
                                                    <chr>
              3.5
## 1 5.1
                                        0.2
                            1.4
                                                    setosa
```

setosa

setosa

0.2

0.2

1.4

1.3

Fixing the second file

Second, making sure values are numeric.

```
multifile data[[2]] <-
 multifile data[[2]] %>%
 mutate (across (!Species, as.numeric))
head(multifile data[[2]], 3)
## # A tibble: 3 \times 5
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
   <dbl> <dbl> <chr>
## 1 5.1 3.5 1.4 0.2 setosa
       4.9
## 2
                           1.4 0.2 setosa
## 3
       4.7
                           1.3
                                    0.2 setosa
```

The bind rows () function can be great for simply combining data.

```
# bind rows(multifile data[[1]], multifile data[[3]], multifile_data[[2]])
bindrows data <- multifile data %>%
   map df(bind rows, .id = "experiment") # recall that modify keeps the same data type
# so that will not do what we want here because we want a data frame instead of a list!
dim(bindrows data)
## [1] 450
tail(bindrows data, 2)
## # A tibble: 2 × 6
    experiment Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## <chr>
                      <dbl>
                                  <dbl>
                                               <dbl>
                                                         <dbl> <chr>
                                                          2.3 virginica
## 1 3
                       6.2
                                  3.4
                                                 5.4
## 2 3
                                                            1.8 virginica
                       5.9
                                    3
                                                 5.1
```

See https://www.opencasestudies.org/ocs-bp-vaping-case-study for more information!

Lab

https://sisbid.github.io/Data-Wrangling/labs/functional-program-lab.Rmd