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

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))

```
head(mtcars, 2)
##
               mpg cyl disp hp drat wt qsec vs am gear carb
                     6 160 110 3.9 2.620 16.46 0 1
## Mazda RX4
                21
## Mazda RX4 Wag 21 6 160 110 3.9 2.875 17.02 0 1
mtcars %>%
 mutate(across(.cols = c(disp, drat, wt, qsec), round)) %>%
 head(2)
               mpg cyl disp hp drat wt qsec vs am gear carb
##
## Mazda RX4
                21
                     6 160 110
                                         16 0 1
## Mazda RX4 Wag 21
                     6
                        160 110
```

functions in R

```
my_function <- function(x){x + 1} # alt: \(x){x + 1}
my_function

## function(x){x + 1}

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 x ... 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:

```
mtcars %>%
 mutate(across(.cols = c(disp, drat, wt, qsec), ~ round(.x))) >>>
 head(2)
##
               mpg cyl disp hp drat wt qsec vs am gear carb
## Mazda RX4
               21 6 160 110
                                4 3 16 0 1
## Mazda RX4 Wag 21 6 160 110
                                4 3 17 0 1 4 4
mtcars %>%
 mutate(across(.cols = c(disp, drat, wt, qsec), round)) %%
 head(2)
               mpg cyl disp hp drat wt qsec vs am gear carb
## Mazda RX4
            21 6 160 110
                                4 3 16 0 1
                                4 3 17 0 1 4 4
## Mazda RX4 Wag 21 6 160 110
```

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 $_{\bullet}x$ (or $_{\bullet}$) 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

```
mtcars %>%
 mutate(across(.cols = disp:wt, round)) %>%
 head(2)
##
                mpg cyl disp hp drat wt qsec vs am gear carb
                 21
## Mazda RX4
                     6 160 110
                                   4 3 16.46 0 1
## Mazda RX4 Wag 21 6 160 110 4 3 17.02
mtcars %>%
 mutate(across(.cols = everything(), round))%>%
 head(2)
                mpg cyl disp hp drat wt qsec vs am gear carb
##
                      6 160 110
                                   4 3
4 3
## Mazda RX4
                 21
                                          16 0 1
## Mazda RX4 Wag 21
                     6
                        160 110
```

Previously we filtered for patterns or conditions...

Dilemma: Seems a bit repetitive!

Now we can filter multiple columns!

if_all(): helps us filter on multiple similar conditions simultaneously!

if_any() and if_all() are 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
                                6.06.15.72.5 virginica2.5 virginica2.5 virginica
## 1
               6.3
                           3.3
                           3.6
## 2
               7.2
## 3
               6.7
                            3.3
                                                      2.5 virginica
iris %>% filter(if_all(Sepal.Length:Petal.Width, ~ . > 2.4))
     Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
                                         6.06.15.72.5 virginica2.5 virginica2.5 virginica
                           3.3
3.6
3.3
               6.3
7.2
## 1
## 2
                                                       2.5 virginica
## 3
```

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)

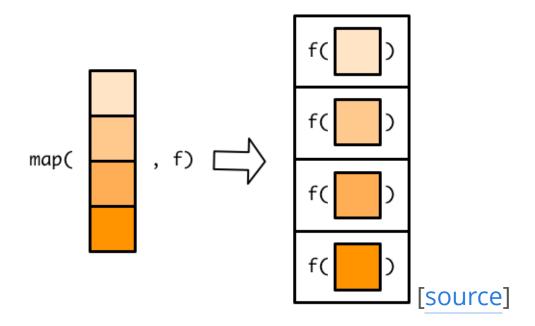
map2 and map2*

- 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

Using map()



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

[1] 1 2 4 5

Using map_df()

Using modify()

purrr:map_df or modify

21

2

160

110

map_df library(purrr) head(mtcars, 2) ## mpg cyl disp hp drat wt qsec vs am gear carb 6 160 110 3.9 2.620 16.46 0 ## Mazda RX4 21 1 ## Mazda RX4 Wag 21 6 160 110 3.9 2.875 17.02 0 1 mtcars %>% map_df(round) %>% head(2)## # A tibble: 2 × 11 ## mpg cyl disp hp drat wt qsec vs am gear carb <dbl> <dbl > <d ## 3 21 110 6 160 4 16 ## 1

4

17

purrr:map_df or modify

modify

```
mtcars %>%
    modify(round) %>%
head(2)

## mpg cyl disp hp drat wt qsec vs am gear carb
## 1 21 6 160 110 4 3 16 0 1 4 4
## 2 21 6 160 110 4 3 17 0 1 4 4
```

It's a bit simpler than across...

But across keeps rownames...

purrr apply function to some columns like across

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

```
head(as_tibble(iris), 3)
## # A tibble: 3 × 5
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
          <dbl>
                    <dbl>
                               <dbl>
                                         <dbl> <fct>
## 1
           5.1
                     3.5
                                1.4
                                           0.2 setosa
## 2
           4.9 3
                                1.4
                                           0.2 setosa
## 3
           4.7
                     3.2
                                1.3
                                           0.2 setosa
as tibble(iris) >>%
 modify_if(is.numeric, as.character) >>>
 head(3)
## # A tibble: 3 × 5
   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
  <chr>
               <chr>
                         <chr>
                                    <chr>
                                              <fct>
## 1 5.1
        3.5
                         1.4
                                    0.2
                                              setosa
## 2 4.9
        3
                         1.4
                                   0.2
                                              setosa
               3.2
                         1.3
                                    0.2
## 3 4.7
                                              setosa
```

Speed test! 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]
                                     21
##
    [1,]
                              16
             1
2
3
4
5
                        11
   [2, ]
[3, ]
                                     22
23
                        12
                              17
##
##
                        13
                              18
                              19
                                     24
##
    [4,]
                        14
##
                         15
                                     25
                  10
                               20
    [5,]
##
   [[4]]
##
                                  [,5]
21
##
          [,1]
               [,2] [,3] [,4]
    [1,]
                        11
##
                               16
                                     22
23
    [2,]
                        12
                              17
##
   [3, ]
                        13
                              18
##
                        14
                              19
                                     24
##
   [4,]
                                     25
                  10
                         15
                               20
```

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?

List: split() the dataset by cyl

```
head(mtcars)
                        mpg cyl disp hp drat
                                                    wt qsec vs am gear carb
##
                                  160 110 3.90 2.620
                       21.0
                                                       16.46
## Mazda RX4
                       21.0
## Mazda RX4 Wag
                                  160 110 3.90
## Datsun 710 22.8 4 108 93 3.85 ## Hornet 4 Drive 21.4 6 258 110 3.08 ## Hornet Sportabout 18.7 8 360 175 3.15
                                  225 105 2.76 3.460 20.22
## Valiant
                       18.1
str(mtcars %>% split(.$cyl))
## List of 3
    $ 4:'data.frame': 11 obs. of 11 variables:
                             22.8 24.4 22.8 32.4 30.4 33.9 21.5 27.3 26 30.4 ...
      ..$ mpg : num [1:11]
##
                     [1:11]
##
                             4 4 4 4 4 4 4 4 4 4
      ..$ cyl : num
##
      ..$ disp: num
                             108 146.7 140.8 78.7 75.7
##
                             93 62 95 66 52 65 97 66 91 113 ...
      ..$ hp : num
                                  3.69 3.92 4.08 4.93 4.22 3.7 4.08 4.43 3.77 ...
##
      ..$ drat: num
##
      ..$ wt : num
##
      ..$ qsec: num
##
      ..$ vs : num
##
      ..$ am : num
##
      ..$ gear: num
##
      ..$ carb: num
##
    $ 6:'data.frame':
                             obs.
##
                                  21.4 18.1 19.2 17.8 19.7
      ..$ mpg : num
                                                                             27/36
                     [1:7]
                            6 6 6
##
      ..$ cyl : num
                                  6 6 6 6
```

List: model on each

```
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

Use lists to iterate through 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)
```

multifile_data[[1]]

```
## # A tibble: 150 × 5
      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
             <dbl>
                         <dbl>
                                      <dbl>
                                                  <dbl> <chr>
##
## 1
               5.1
                           3.5
                                        1.4
                                                    0.2 setosa
## 2
               4.9
                           3
                                        1.4
                                                    0.2 setosa
## 3
               4.7
                           3.2
                                        1.3
                                                    0.2 setosa
## 4
               4.6
                           3.1
                                        1.5
                                                    0.2 setosa
## 5
               5
                           3.6
                                        1.4
                                                    0.2 setosa
## 6
               5.4
                           3.9
                                        1.7
                                                    0.4 setosa
## 7
               4.6
                           3.4
                                        1.4
                                                    0.3 setosa
               5
## 8
                           3.4
                                        1.5
                                                    0.2 setosa
               4.4
                          2.9
                                        1.4
                                                    0.2 setosa
## 9
               4.9
                           3.1
## 10
                                        1.5
                                                    0.1 setosa
## # i 140 more rows
```

multifile_data[[2]]

multifile_data[[3]]

```
## # A tibble: 150 × 5
      Sepal.Length Sepal.Width Petal.Length Petal.Width Species
             <dbl>
                         <dbl>
                                      <dbl>
                                                  <dbl> <chr>
##
## 1
            -999
                           3.5
                                        1.4
                                                    0.2 setosa
## 2
           -999
                           3
                                        1.4
                                                    0.2 setosa
## 3
           -999
                           3.2
                                        1.3
                                                    0.2 setosa
## 4
               4.6
                           3.1
                                        1.5
                                                    0.2 setosa
## 5
               5
                           3.6
                                        1.4
                                                    0.2 setosa
## 6
               5.4
                           3.9
                                        1.7
                                                    0.4 setosa
## 7
               4.6
                           3.4
                                        1.4
                                                    0.3 setosa
               5
## 8
                           3.4
                                        1.5
                                                    0.2 setosa
               4.4
                           2.9
                                        1.4
                                                    0.2 setosa
## 9
               4.9
                           3.1
## 10
                                        1.5
                                                    0.1 setosa
## # i 140 more rows
```

Fixing the second file

3

3.2

1.4

1.3

First, separating by the :.

2 4.9

3 4.7

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

setosa

setosa

0.2

0.2

Fixing the second file

Second, making sure values are numeric.

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

The bind_rows() function can be great for combining data.

recall that modify keeps the same data type (here, a list). We want a data frame instead.

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

Summary

- function(x){ } or (x){ } denotes a function. You also commonly see ~ x inside across
- map_df and modify apply functions to each element of an object. map returns
 a list, modify returns the same object type.
- The purrr package has other useful functional programming features
- lists can be great for storing iterative work.

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