L13 Vectors and Iteration

Data Science I (STAT 301-1)

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Overview

The goal of this lab is to improve our understanding of vectors (a basic structure in R) and iterative programming skills, using the purrr package from the tidyverse. This resource may help:

• purrr tidyverse homepage

Datasets

Datasets referenced in this lab are either from the tidyverse or from the familiar nycflights13 package.

Exercises

Please complete the following exercises. Be sure that your solutions are clearly indicated and that the document is neatly formatted.

```
# Loading package(s)
library(tidyverse)
library(nycflights13)
library(readr)
```

Load Packages

Exercise 1 (Website: 20.4.6 Ex. 1)

What does mean(is.na(x)) tell you about a vector x? What about sum(!is.finite(x))? mean(is.na(x)) would give you the proportion of NA in a vector. sum(!is.finite(x)) would give you a count of all the infinite elements in a vector.

Exercise 2 (Website: 20.4.6 Ex. 5)

```
Why is x[-which(x > 0)] not the same as x[x <= 0]?
```

```
x \leftarrow c(-3, -1, 0, 1, 4, Inf, -Inf, NA, NaN)
x[-which(x > 0)]
```

```
## [1] -3 -1 0 -Inf NA NaN x[x \le 0]
```

```
## [1] -3 -1 O -Inf NA NA
```

x[-which(x > 0)] leaves NA's and NaNs as is, whereas $x[x \le 0]$ replaces NaN's with NA's.

Exercise 3 (Website: 20.5.4 Ex. 2)

What happens if you subset a tibble as if you're subsetting a list? What are the key differences between a list and a tibble?

```
myTibble <- tibble('col1' = c(1, 2, 3, 4), 'col2' = c(5, 6, 7, 8))
myTibble[1]</pre>
```

```
## # A tibble: 4 x 1
## col1
## <dbl>
## 1 1
## 2 2
## 3 3
## 4 4
myTibble[[1]]
```

```
## [1] 1 2 3 4
```

Subsetting a tibble with one bracket returns a tibble, whereas doing the same for a list returns a list. The double bracket does the same for both tibbles and list: they return the individual elements.

Exercise 4 (Website: 20.7.4 Ex. 2)

Try to make a tibble containing columns with different lengths. What happens?

```
myTibble2 \leftarrow tibble('col1' = c(1, 2, 3), 'col2' = c(5, 6, 7, 8))
```

You can not create a tibble with different column lengths, you get an error.

Exercise 5 (Website: 20.7.4 Ex. 3)

Is it possible to have a list as a column of a tibble?

```
a <- list(4, 5, NA, 'a', 'b')
myTibble3 <- tibble('col1' = a, 'col2' = c(1, 2, 3, 4, 5))
myTibble3</pre>
```

```
## # A tibble: 5 x 2
## col1 col2
```

```
## t> <dbl>
## 1 <dbl [1]> 1
## 2 <dbl [1]> 2
## 3 <lgl [1]> 3
## 4 <chr [1]> 4
## 5 <chr [1]> 5
```

Theoretically you can, but it is not practically useful due to nesting issues.

Exercise 6 (Website: 21.2.1 Ex. 1)

Write for loops to:

[[1]]

- Compute the mean of every column in mtcars.
- Determine the type of each column in nycflights13::flights.
- Compute the number of unique values in each column of iris.
- Generate 10 random draws from a normal distribution with $\mu = -10, 0, 10,$ and 100.

Think about the output, sequence, and body of each loop before you start writing it.

```
means <- vector('double', ncol(mtcars))</pre>
for (i in seq_along(mtcars)) {
  means[i] = mean(mtcars[[i]])
}
means
##
    [1]
         20.090625
                      6.187500 230.721875 146.687500
                                                         3.596563
                                                                     3.217250
    [7]
         17.848750
                      0.437500
                                  0.406250
                                             3.687500
                                                         2.812500
col_type <- vector('character', ncol(flights))</pre>
for (i in seq_along(flights)) {
  col_type[i] = typeof(flights[[i]])
}
col_type
   [1] "integer"
                     "integer"
                                               "integer"
                                                            "integer"
                                                                         "double"
                                  "integer"
                     "integer"
                                  "double"
##
   [7] "integer"
                                               "character" "integer"
                                                                         "character"
## [13] "character" "character" "double"
                                               "double"
                                                            "double"
                                                                         "double"
## [19] "double"
view(flights)
num_unique <- vector('integer', ncol(iris))</pre>
for (i in seq_along(iris)) {
 num_unique[i] = length(unique(iris[[i]]))
}
num_unique
## [1] 35 23 43 22 3
norm_dist <- vector('list', 4)</pre>
mu \leftarrow c(-10, 0, 10, 100)
for (i in seq_along(norm_dist)) {
  norm_dist[[i]] = rnorm(10, mean = mu[i])
}
norm_dist
```

```
-8.570702 -9.868547 -10.782991 -9.135421 -10.327494 -9.640191
        -8.998848 -11.227847 -10.035393 -11.486441
##
##
## [[2]]
##
   [1] -3.1378973 2.1591393 -0.2440069 -0.1406911 1.1891850 0.1144654
   [7] 0.5604635 1.0042404 0.4802929 1.2633342
##
##
## [[3]]
##
   [1] 10.567096 10.890726 10.623337 9.660844 9.669222 11.338140 10.445854
##
   [8] 12.371605 10.050879 9.223419
##
## [[4]]
        99.92805 99.17429 100.20113 100.86725 100.21087 98.77748 101.37475
##
   [1]
   [8]
        99.41844 99.78901 101.97932
```

Exercise 7 (Website: 21.3.5 Ex. 1)

Imagine that you have a directory full of CSV files that you want to read in to R. You have their paths in a vector, files <- dir("data/", pattern = "\\.csv\$", full.names = TRUE), and want to read each one with read_csv(). Write a for loop that will load them into a single data frame.

```
files <- dir("data/", pattern = "\\.csv$", full.names = TRUE)
num_files <- length(files)
file_df <- vector('list', num_files)
for (i in seq_along(files)) {
  file_df[[i]] = read_csv(files[[i]])
}
file_df <- bind_rows(file_df)</pre>
```

Exercise 8 (Website: 21.3.5 Ex. 3)

Write a function that prints the mean of each numeric column in a data frame along with its name. For example, show_mean(iris) should print:

```
show_mean(iris)
#> Sepal.Length: 5.84
#> Sepal.Width: 3.06
#> Petal.Length: 3.76
#> Petal.Width: 1.20

show_mean <- function(df) {
   for (name in seq_along(df)) {
      if (is.numeric(df[[name]])) {
        print(c(names(df)[name], ": ", mean(df[[name]])))
      }
   }
}
show_mean(iris)</pre>
```

```
## [1] "Sepal.Length" ": " "5.8433333333333333"
## [1] "Sepal.Width" ": " "3.057333333333333"
## [1] "Petal.Length" ": " "3.758"
## [1] "Petal.Width" ": " "1.199333333333333"
```

Note: it's not pretty but it gets the job done, and I'm not very familiar with from atting print lines.

Exercise 9 (Website: 21.5.3 Ex. 1)

\$origin

[1] "character"

Write code that uses one of the map functions to:

- Compute the mean of every column in mtcars.
- Determine the type of each column in nycflights13::flights.
- Compute the number of unique values in each column of iris.
- Generate 10 random normals for each of $\mu = -10$, 0, 10, and 100.

```
map_dbl(mtcars, mean)
##
                       \operatorname{cyl}
                                  disp
                                                          drat
                                                                        wt
                                                                                  qsec
          mpg
                                                hp
    20.090625
                 6.187500 230.721875 146.687500
                                                     3.596563
                                                                 3.217250
                                                                           17.848750
##
##
            ٧s
                       am
                                  gear
                                              carb
     0.437500
                 0.406250
                             3.687500
                                         2.812500
map(flights, typeof)
## $year
## [1] "integer"
##
## $month
## [1] "integer"
##
## $day
## [1] "integer"
##
## $dep_time
   [1] "integer"
##
##
## $sched_dep_time
## [1] "integer"
##
## $dep_delay
   [1] "double"
##
## $arr_time
   [1] "integer"
##
##
## $sched_arr_time
## [1] "integer"
##
## $arr_delay
   [1] "double"
##
##
## $carrier
## [1] "character"
##
## $flight
## [1] "integer"
##
## $tailnum
## [1] "character"
##
```

```
##
## $dest
## [1] "character"
##
## $air_time
## [1] "double"
##
## $distance
## [1] "double"
##
## $hour
## [1] "double"
## $minute
## [1] "double"
##
## $time_hour
## [1] "double"
map_int(iris, n_distinct)
## Sepal.Length Sepal.Width Petal.Length Petal.Width
                                                             Species
map(c(-10, 0, 10, 100), \sim rnorm(n = 10, mean = .))
## [[1]]
  [1] -11.516111 -11.655894 -9.992923 -10.101011 -9.701573 -10.656301
##
   [7] -10.542919 -8.254765 -9.144136 -9.796250
##
## [[2]]
   [1] 0.12626639 -1.22695731 -0.49251532 -0.77321671 -0.72359170 -1.32026626
##
##
   [7] 1.31965260 -0.05420319 -0.19098626 -0.59998502
##
## [[3]]
##
  [1] 9.447817 10.882168 10.779855 9.281482 9.876865 10.118011 11.274408
  [8] 9.831129 10.248930 10.646574
##
##
## [[4]]
## [1] 99.81377 100.54563 99.21948 99.40902 100.63160 96.72199 99.97632
  [8] 100.29100 100.96141 99.64610
Exercise 10 (Website: 21.5.3 Ex. 5)
Rewrite map(x, function(df) lm(mpg ~ wt, data = df)) to eliminate the anonymous function.
x <- split(mtcars, mtcars$cyl)</pre>
map(x, ~lm(mpg ~ wt, data = .))
## $`4`
##
## Call:
## lm(formula = mpg ~ wt, data = .)
## Coefficients:
```

```
ercept) wt
39.571 -5.647
## (Intercept)
##
##
##
## $`6`
##
## lm(formula = mpg ~ wt, data = .)
## Coefficients:
                       wt
        ercept) wt
28.41 -2.78
## (Intercept)
##
##
##
## $`8`
##
## Call:
## lm(formula = mpg ~ wt, data = .)
## Coefficients:
## (Intercept) wt
## 23.868 -2.192
```