R: Tibbles and Dataframes

Sakol Suethanapornkul

28 October 2020

Quiz

Use the following vectors to answer the questions below:

```
scores <- c(3, 12, 8, 2, 4, 11, 15, 19, 3, 7, 6, 9)
answer <- c(TRUE, FALSE, FALSE, TRUE, TRUE, FALSE)
```

- [1] What kind of vector is scores? How many elements does it have?
- [2] Check whether values of scores are either below 5 or above 10.
- [3] Check whether values of scores are 11, 12, and 15.
- [4] What kind of vector is answer? Find the average of answer.
- [5] Bonus: Extract scores below 7?

Answers

Here are the answers to the first four questions:

- [1] typeof(scores) and length(scores)
- [2] scores < 5 | scores > 10
- [3] scores == 11 | scores == 12 | scores == 15
- [4] typeof(answer) and mean(answer)

Answers

But our answer to [3] is cumbersome and prone to error.

- If we want to check if x is one of the "things" in y, we can use %in%.
- So, a better answer is: scores %in% c(11, 12, 15)

Answers

So far, we have learned: == != < = >= and x %in% c() Answer to [5] needs a bit more explanation about indices.

Index & Subsetting

Say, we write:

```
scores <- c(3, 12, 8, 2, 4, 11, 15, 19, 3, 7, 6, 9)
scores < 7 #or check <- scores < 7</pre>
```

What do we get with this code?

Index & Subsetting

We'd like to extract scores below 7 from the vector (i.e., 3, 2, 4, 3, 6). We do not want T or F as an answer.

```
scores <- c(3, 12, 8, 2, 4, 11, 15, 19, 3, 7, 6, 9)
```

So, we'll need to talk about subsetting. Let's begin with subsetting with positive integers.

```
scores[1:2]
scores[c(1, 3, 5)]
scores[[2]]
```

Index & Subsetting

We can subset with a comparison function, which will keep values that are TRUE:

```
scores[scores < 7]
```

And if we need position numbers instead:

```
which(scores < 7) #then
scores[which(scores < 7)]</pre>
```

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

- numbers <- c(1, 2, 5, 6); numbers <- 1:10
- typeof(numbers); length(numbers)
- numbers[1:2]; numbers[[2]]
- numbers >= 5; (numbers / 2) == 0 | numbers >= 3
- numbers %in% c(3, 5, 7)
- numbers * 3

R contains numerous packages to perform various functions, and one of the first things we need to know is how to install external packages from CRAN:

```
install.packages("<package_name>")
```

For instance, to do data science, we'll need a package called tidyverse. We can install that on our laptop with:

```
install.packages("tidyverse")
```

Once we have a package installed, we will need to load it with library():

library(tidyverse)

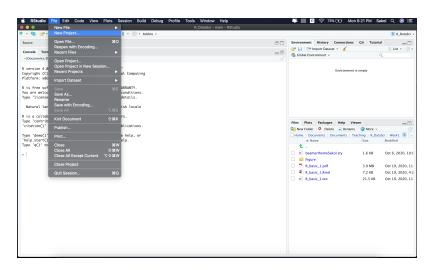
EXERCISE: Try calling a package lme4 with library(lme4). What happened?

Now that we have discussed library(), it's time we talk about our workflow!

- Ideally, we want to call all the packages we need for our analysis once;
- We then want to run some analysis on our data and save all the results in one place;
- We also want to keep a record of what we have done in R, so we can recreate all these things later.

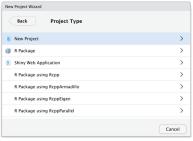
That can be achieved with an R project!

R project can be created right from RStudio:



R project can be created right from RStudio:





You can choose to place this newly-created R project in any folder you want. You can then check where your project lives with:

getwd()

Alternatively, you can see your current directory under Console.

EXERCISE: Why does this matter? What's the point of creating a project?

Run the following code (adapted from R for Data Science, p. 116):

```
library(tidyverse)
ggplot(diamonds, aes(carat, price)) +
  geom_point(alpha = 0.2) +
  theme bw()
ggsave("diamonds.png")
write csv(diamonds, "diamonds.csv")
```

How can we foster this habit? A short answer is: create an R script!

An R Script, not the console, should be where your codes live.

- Begin with general comment about your script;
- Then, load all the packages you need in the preamble;
- Use comments to provide context for your analysis;
- Indent your codes and use new lines to your advantage.

In most data analysis projects, we deal with data frames. Each data frame is a named list of vectors.

In data frames, the length of each element must be the same.

We can inspect the structure of a data frame with:

```
str(dat)
```

which lists on each line an object with a column name.

Another useful function, which provides a descriptive summary, is:

```
summary(dat)
```

And, finally, we can peak inside data frames:

```
head(dat, n = 3)

tail(dat, n = 3)
```

We will work with data frames that come with the tidyverse package. Let's load two of the data sets:

```
data(diamonds)
data(mpg)
```

To find out more about these data sets, you can run help() such as help(diamonds).

EXERCISE: Answer the following questions

- [1] How many columns does diamonds have? How about mpg?
- [2] In diamonds, what vector are cut and color?
- [3] Figure out what mpg is about by running help(mpg).

If we create a data frame with data.frame() or read in external data with base R functions (e.g., read.__), we get a data frame.

Data frames have some undesirable behaviors. As shown in diamonds, some character vectors are converted to **factors**.

```
str(diamonds, vec.len = 0.5)
## tibble [53,940 x 10] (S3: tbl df/tbl/data.frame)
##
   $ carat : num [1:53940] 0.23 ...
##
   $ cut : Ord.factor w/ 5 levels "Fair"<"Good"<...: 5
   $ color : Ord.factor w/ 7 levels "D"<"E"<"F"<"G"<...: 2</pre>
##
   $ clarity: Ord.factor w/ 8 levels "I1"<"SI2"<"SI1"<...</pre>
##
   $ depth : num [1:53940] 61.5 ...
##
## $ table : num [1:53940] 55 ...
   $ price : int [1:53940] 326 ...
##
## $ x : num [1:53940] 3.95 ...
##
   $ y : num [1:53940] 3.98 ...
   $ 7.
             : num [1:53940] 2.43 ...
##
```

In this class, we will work with tibbles, which can be created with a function called tibble():

Right now, we will not dwell on why we should work with tibbles rather than data frames. We will instead focus on some helpful functions. Previously:

This is tedious and error-prone. Instead:

```
class <- rep(c("A", "B", "C"), times = 4)
class <- rep(c("A", "B", "C"), each = 4)</pre>
```

You can rep() with various types of vectors.

```
a <- 1:5
b <- letters[1:5]

rep(a, each = 3)
rep(b, each = 2, times = 3)</pre>
```

Alternatively, you can generate a sequence of numbers with:

```
seq(from = 0, to = 1, by = 0.1)
seq(from = 0, to = 1, length.out = 100)
```

Wrap-up

We have discussed the following functions for data frames:

```
data.frame(a = , b = , ...)
 str()
 summary()
 head()
 tail()
And helpful functions to generate sequences for vectors:
 seq()
 rep()
Sandwiched in between are the basics:
 install.packages()
 library()
```

Homework

- First, create your own R project;
- Then, move some data file into that folder; and
- Last, create an R script for future analyses.