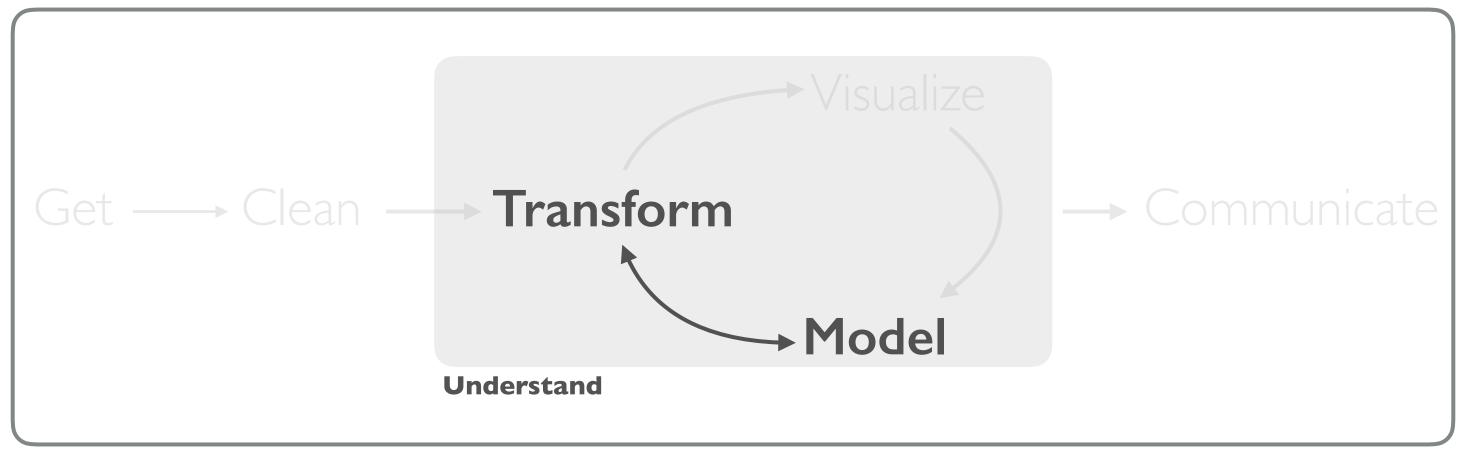
CONTROL STATEMENTS & ITERATION



Program

[†]A modified version of Hadley Wickham's analytic process

"Great design is iteration of good design."

– M. Cobanli

CONTROL STATEMENTS & ITERATION

- Reducing code duplication has three main benefits:
 - It's easier to see the intent of your code, because your eyes are drawn to what's different, not what stays the same.
 - It's easier to respond to changes in requirements. As your needs change, you only need to make changes in one place, rather than remembering to change every place that you copied-and-pasted the code.
 - You're likely to have fewer bugs because each line of code is used in more places.

Control statements and iteration functions provide two means to reduce code duplication

PREREQUISITES



PACKAGE PREREQUISITE

```
library(tidyverse)
Loading tidyverse: ggplot2
Loading tidyverse: tibble
Loading tidyverse: tidyr
Loading tidyverse: readr
Loading tidyverse: purrr
Loading tidyverse: dplyr
Conflicts with tidy packages
filter(): dplyr, stats
lag(): dplyr, stats
```

CONTROL STATEMENTS

```
if (test_expression) {
   statement
}
```

Evaluates if the test expression is TRUE

If TRUE: execute statement

If FALSE: do nothing

```
x <- c(8, 3, -2, 5)

if(x < 0) {
  print("x contains a negative number")
}</pre>
```

What do you expect to happen?

```
x <- c(8, 3, -2, 5)

if(x < 0) {
  print("x contains a negative number")
}
Warning message:
In if (x < 0) print("Negative") :
  the condition has length > 1 and only the first element will be used
```

Trick question - we actually get a warning

The if statement looks for a single conditional value. Here, x < 0 will return a vector of 4 conditional values.

```
x <- c(8, 3, -2, 5)

if(any(x < 0)) {
  print("x contains a negative number")
}</pre>
```

Instead, we can assess if any x values are less than 0

Now what do you expect to happen?

```
x <- c(8, 3, -2, 5)

if(any(x < 0)) {
  print("x contains a negative number")
}
[1] "x contains a negative number"</pre>
```

Instead, we can assess if any x values are less than 0

Now what do you expect to happen?

```
x <- c(8, 3, -2, 5)

if(any(x < 0)) {
  print("x contains a negative number")
}
[1] "x contains a negative number"</pre>
```

Instead, we can assess if any x values are less than 0

Now what do you expect to happen?

Change -2 to 2. Now what happens when you run this code?

SYNTAX OF AN if...else STATEMENT

```
if (test_expression) {
   statement 1
} else {
   statement 2
}
```

We can extend an if statement with an else statement.

SYNTAX OF AN if...else STATEMENT

```
if(any(x < 0)) {
  print("x contains negative number(s)")
} else{
  print("x contains all positive numbers")
}
[1] "x contains negative number(s)"</pre>
```

Now, if the test expression is TRUE we print statement 1 and if it is FALSE we print statement 2.

SYNTAX OF AN if...else STATEMENT

```
x <- 7
if(x >= 10) {
  print("x exceeds acceptable tolerance levels")
} else if(x >= 0 & x < 10) {
  print("x is within acceptable tolerance
levels")
} else {
  print("x is negative")
   "x is within acceptable tolerance levels"
```

And we can expand this further to include multiple test expressions.

YOURTURN - PART I

In your data folder you have many data sets. You want to create an if else control statement that will look to see if a particular "Month-XX.csv" file exists in this folder. For Part 1, build your your if-else statement so that you create the appropriate file name based on a month input:

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 13
                                           If month equals 1-9 the output
                                           should look like "Month-08.csv"
if (month %in% 1:9) {
                                           If month equals 10-12 the
} else if (month %in% 10:12) {
                                           output should look like
                                           "Month-12.csv"
} else {
                                           If month is not 1-12 the output
                                           should look like "Invalid month"
```

SOLUTION

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 1
if (month %in% 1:9) {
  paste0("data/", file_1, 0, month, file_2)
} else if (month %in% 10:12) {
  paste0("data/", file_1, month, file_2)
} else {
  print("Invalid month")
[1] "data/Month-01.csv"
```

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 13
if (month %in% 1:9) {
  paste0("data/", file_1, 0, month, file_2)
} else if (month %in% 10:12) {
  paste0("data/", file_1, month, file_2)
} else {
  print("Invalid month")
[1] "Invalid month"
```

YOURTURN - PART 2

The **file.exists** function will check to see if a file exists at the given path. Incorporate **file.exists** in your if-else statement so now the output provides a TRUE, FALSE, or "Invalid month" response. Is the December data ("Month-I 2.csv") present?

SOLUTION

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 12
if(month %in% 1:9) {
  file_name <- paste0("data/", file_1, 0, month, file_2)
  file.exists(file_name)
} else if(month %in% 10:12) {
  file_name <- paste0("data/", file_1, month, file_2)
  file.exists(file_name)
} else {
 print("Invalid month")
[1] FALSE
```

ITERATION

for loops



```
for(i in 1:100) {
     <do stuff here with i>
}
```

For each element in a sequence performs defined tasks

For each element in a sequence performs defined tasks

Alternative - why use seq_along?

```
years <- 2010:2017

for (i in seq_along(years)) {
  output <- paste("The year is", years[i])
  print(output)
}</pre>
```

What does this for loop do?

```
years <- 2010:2017
for (i in seq_along(years)) {
  output <- paste("The year is", years[i])
  print(output)
[1] "The year is 2010"
[1] "The year is 2011"
   "The year is 2012"
   "The year is 2013"
[1] "The year is 2014"
[1] "The year is 2015"
[1] "The year is 2016"
[1] "The year is 2017"
```

What does this for loop do?

```
result <- vector(mode = "character",
                 length = length(years))
for (i in seq_along(years)) {
  output <- paste("The year is", years[i])
  result[i] <- output
result
[1] "The year is 2010" "The year is 2011"
[3] "The year is 2012" "The year is 2013"
[5] "The year is 2014" "The year is 2015"
[7] "The year is 2016" "The year is 2017"
```

When saving results from a for loop we want to:

- 1. initiate the output outside of the for loop
- 2. save the results from the for loop to the output

```
x <- 7
if(x >= 10) {
  print("x exceeds acceptable tolerance levels")
} else if(x >= 0 & x < 10) {
  print("x is within acceptable tolerance
levels")
} else {
  print("x is negative")
[1] "x is within acceptable tolerance levels"
```

Remember this extended if...else statement?

Let's implement it into a for loop

```
x \leftarrow c(-1, 7, 8, 11)
tolerance <- vector(mode = "character",
                      length = length(x)
for (i in seq_along(x)) {
  if(x[i] >= 10) {
    value <- "x exceeds acceptable tolerance levels"
  } else if(x[i] >= 0 & x[i] < 10) {</pre>
    value <- "x is within acceptable tolerance levels"
  } else {
    value <- "x is negative"</pre>
  tolerance[i] <- value</pre>
```

- 1. Vector to analyze
- 2. Initiate output shell
- 3. For each element in our x
- 4. Assess test expressions, execute relevant statement, and save as value

5. For that respective element add the value in the corresponding element in out output shell

YOURTURN

Remember this if-else statement we made?

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 12
if(month %in% 1:9) {
  file_name <- paste0("data/", file_1, 0, month, file_2)
  file.exists(file_name)
} else if(month %in% 10:12) {
  file_name <- paste0("data/", file_1, month, file_2)
  file.exists(file_name)
} else {
  print("Invalid month")
```

YOURTURN

Develop a for loop that will loop through the months provided and import those .csv files that are present.

YOURTURN

Develop a for loop that will loop through the months provided and import those .csv files that are present.

Run your for loop for months 1-13:

- If a particular month is available import it as "df.month. I", "df.month.2", ...
- I a particular month is not available, provide the response: "There is no data for month x"
- If a particular month is invalid (i.e. 13), provide the response: "x is an invalid month"

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 1:13
for(i in month) {
  # create file name
  if(i %in% 1:9) {
    file_name <- paste0("data/", file_1, 0, i, file_2)
  } else if(i %in% 10:12) {
    file_name <- paste0("data/", file_1, i, file_2)
  } else {
    response <- paste(i, "is an invalid month")</pre>
    print(response)
    next
  # import data
  if(file.exists(file_name)) {
    df <- read_csv(file_name)</pre>
    assign(paste0("df.month.", i), df)
    rm(df)
  } else {
    response <- paste("There is no available data for month", i)
    print(response)
```

SOLUTION

1. Create file names.

2. If invalid month, provide response and skip to next iteration.

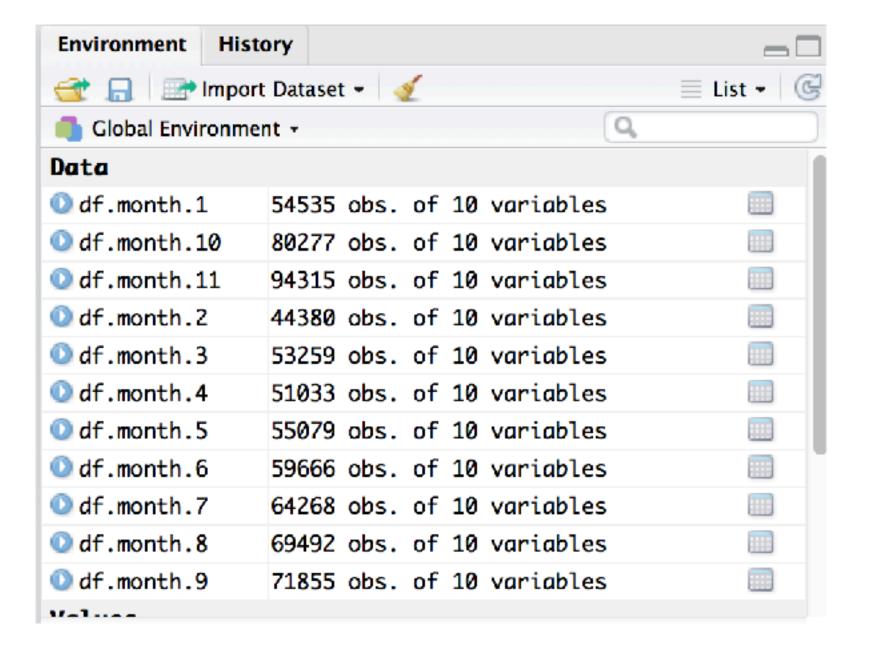
3. If the file exists, import and rename.

4. If the file does not exist for a given month, provide a response

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 1:13
for(i in month) {
  # create file name
  if(i %in% 1:9) {
    file_name <- paste0("data/", file_1, 0, i, file_2)
  } else if(i %in% 10:12) {
    file_name <- paste0("data/", file_1, i, file_2)
  } else {
    response <- paste(i, "is an invalid month")</pre>
    print(response)
    next
  # import data
  if(file.exists(file_name)) {
    df <- read_csv(file_name)</pre>
    assign(paste0("df.month.", i), df)
    rm(df)
  } else {
    response <- paste("There is no available data for month", i)
    print(response)
```

SOLUTION

When you run this for months 1:13



and for months 12 & 13 the response is:

- [1] "There is no available data for month 12"
- [1] "13 is an invalid month"

```
file_1 <- "Month-"
  file_2 <- ".csv"
 month <- 1:13
  for(i in month) {
   # create file name
   if(i %in% 1:9) {
     file_name <- paste0("data/", file_1, 0, i, file_2)
   } else if(i %in% 10:12) {
 Since all our data frames have similar variables (we just get
response <- paste(i, "is an invalid month")

updated data each month), how could we change this so you
                      just create one single data frame?
   # import data
   if(file.exists(file_name)) {
     df <- read_csv(file_name)</pre>
     assign(paste0("df.month.", i), df)
     rm(df)
   } else {
     response <- paste("There is no available data for month", i)
     print(response)
```

SOLUIION

When you run this for months 1:13



and for months 12 & 13 the response is:

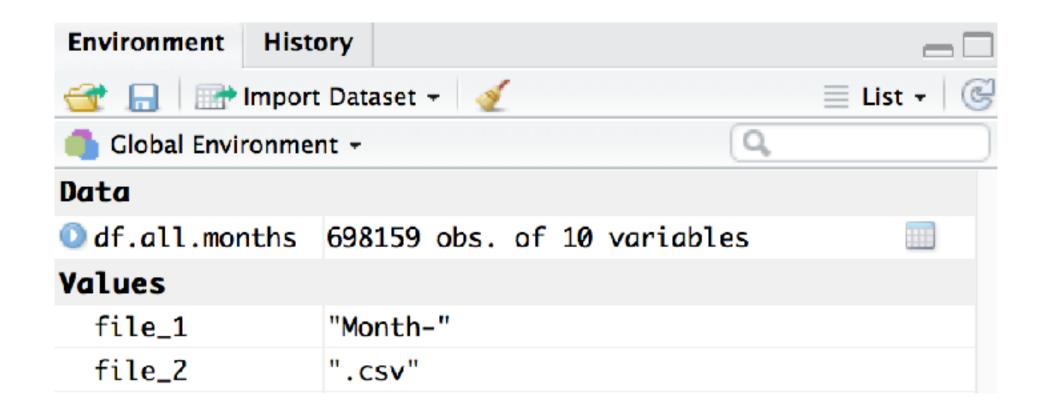
- [1] "There is no available data for month 12"
- [1] "13 is an invalid month"

```
file_1 <- "Month-"
file_2 <- ".csv"
month <- 1:13
# create empty data frame
df.all.months <- data.frame(NULL)</pre>
for(i in month) {
  # create file name
  if(i %in% 1:9) {
    file_name <- paste0("data/", file_1, 0, i, file_2)</pre>
  } else if(i %in% 10:12) {
    file_name <- paste0("data/", file_1, i, file_2)</pre>
  } else {
    response <- paste(i, "is an invalid month")</pre>
    print(response)
    next
  # import data
  if(file.exists(file_name)) {
    df <- read_csv(file_name)</pre>
    df.all.months <- rbind(df.all.months, df)</pre>
    rm(df)
  } else {
    response <- paste("There is no available data for month", i)</pre>
    print(response)
```

SOLUTION

1. Create an empty data frame.

2. Then as you import the files, you can rbind the new data to our empty data frame.



ITERATION

map functions



INTRODUCING purrr

The purrr package provides an alternative means to iterate over elements and perform a set function

Each map function works the same $map_*(x, f)$:

- 1. loop over x
- 2. apply function f to each element
- 3. return the results

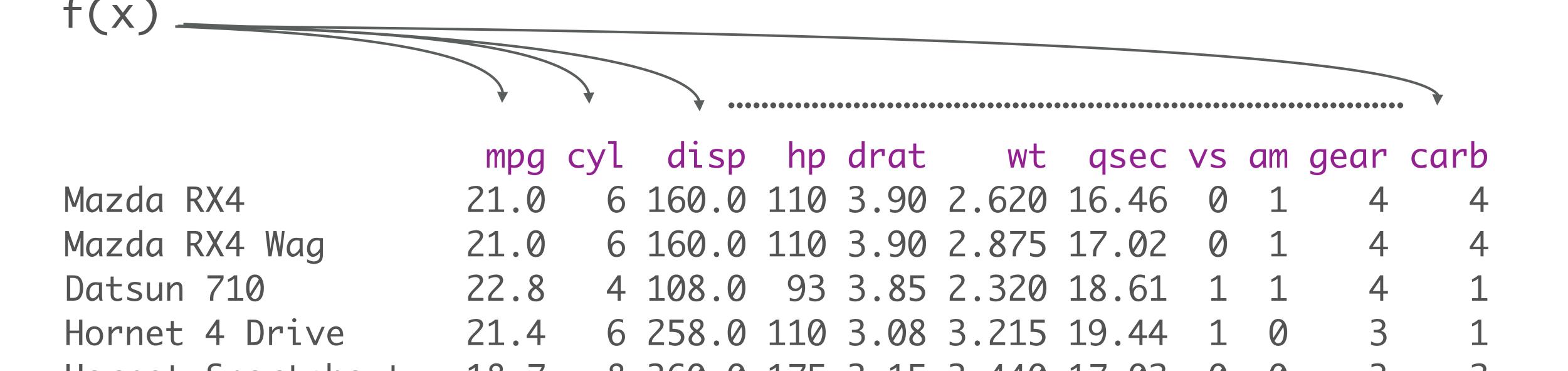




map_*(vector, function)



map_*(data.frame, function)



map_*(list, function)

```
f(x) $item1

[,1] [,2] [,3] [,4] [,5]

[1,] 0.1843049 0.05852987 0.2002625 0.2860744 0.7361251
[2,] 0.3005814 0.77366238 0.3798870 0.5418013 0.1937544

$item2

[,1] [,2] [,3] [,4] [,5]

[1,] 0.9181339 0.6694026 0.3084275 0.4940026 0.08608652
[2,] 0.5877923 0.5218446 0.9769865 0.6842613 0.04770349
```

The map_* function we use is determined by the output of the function

- map() returns a list
- map_dbl() returns a double vector
- map_lgl() returns a logical vector
- map_int() returns a integer vector
- map_chr() returns a character vector

```
map_dbl(mtcars, mean)
mtcars %>% map_dbl(mean)
                          disp
                                       hp
          cyl
      mpg
 20.090625 6.187500 230.721875 146.687500
     drat
                 wt
                          qsec
                                       VS
  3.596563 3.217250 17.848750 0.437500
                          carb
                gear
       am
           3.687500 2.812500
  0.406250
mtcars %>% map(mean)
$mpg
[1] 20.09062
$cyl
```

The map_* functions incorporate the %>% operator

YOURTURN!

With the iris data set, use the map functions to answer these three questions:

- 1. what is the class of each variable?
- 2. what is the mean value for each variable?
- 3. which variables have a mean value greater than 5

SOLUTION

```
# what is the class of each variable?
iris %>% map_chr(class)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
   "numeric" "numeric" "numeric" "numeric" "factor"
# what is the mean value for each variable?
iris %>% map_dbl(mean)
Sepal.Length Sepal.Width Petal.Length Petal.Width
                                               Species
   5.843333 3.057333 3.758000 1.199333
                                                        NA
```

SOLUTION

```
# which variables have a mean value greater than 5?
# option 1
iris %>%
  map_dbl(mean) %>%
  map_lgl(\sim . > 5)
# option 2
iris %>% map_lgl(~ mean(.) > 5)
                                                        Species
Sepal.Length Sepal.Width Petal.Length Petal.Width
                          FALSE
                   FALSE
        TRUE
                                             FALSE
                                                             NA
```

ADVANTAGES OF THE MAP FUNCTIONS

- Handy shortcuts for specifying . f
- More consistent than sapply(), lapply(), which makes them better for programming
- Takes much less time to solve iteration problems (because they are written in C)

SHORTCUTS FOR SPECIFYING. f

map(df, summary)

An existing function

map(df, myfunction)

An existing function you created

map(df, function(x) sum(is.na(x))

An anonymous function defined on the fly

map(df, ~ sum(is.na(.)))

An anonymous function defined using a formula shortcut

SHORTCUTS FOR SPECIFYING. f

Can you find what variables in the nycflights13::flights data have missing values and how many missing values they have?

SHORTCUTS FOR SPECIFYING. f

Can you find what variables in the nycflights::flights data have missing values and how many missing values they have?

```
nycflights13::flights %>% map_dbl(~ sum(is.na(.)))
                                                    dep_time sched_dep_time
                        month
                                          day
          year
                                                        8255
                     arr_time sched_arr_time
                                                   arr_delay
     dep_delay
                                                                    carrier
          8255
                                                        9430
                         8713
        flight
                      tailnum
                                      origin
                                                        dest
                                                                   air_time
                                                                       9430
                         2512
      distance
                                      minute
                                                   time_hour
                         hour
```

```
cyl <- split(mtcars, mtcars$cyl)</pre>
str(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 ...
  ..$ 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 ...
  ..$ wt : num [1:11] 2.32 3.19 3.15 2.2 1.61 ...
  ..$ qsec: num [1:11] 18.6 20 22.9 19.5 18.5 ...
  ..$ vs : num [1:11] 1 1 1 1 1 1 1 0 1 ...
```

```
cyl[[1]]
             mpg cyl disp hp drat wt qsec vs am gear carb
Datsun 710
                  4 108.0 93 3.85 2.320 18.61 1 1
            22.8
            24.4
                  4 146.7 62 3.69 3.190 20.00 1 0 4 2
Merc 240D
                  4 140.8 95 3.92 3.150 22.90 1 0
            22.8
Merc 230
                  4 78.7 66 4.08 2.200 19.47 1 1
       32.4
Fiat 128
Honda Civic 30.4
                  4 75.7 52 4.93 1.615 18.52 1 1
Toyota Corolla 33.9
                  4 71.1 65 4.22 1.835 19.90 1 1
Toyota Corona 21.5 4 120.1 97 3.70 2.465 20.01 1
            27.3
Fiat X1-9
                  4 79.0 66 4.08 1.935 18.90 1 1
Porsche 914-2 26.0
                  4 120.3 91 4.43 2.140 16.70 0 1
            30.4
                  4 95.1 113 3.77 1.513 16.90 1 1
Lotus Europa
```

Say we want to apply a model over a list of data frames

```
mtcars %>%
  split(.$cyl) %>%
  map(\sim lm(mpg \sim wt, data = .))
$`4`
Call:
lm(formula = mpg \sim wt, data = .)
Coefficients:
(Intercept)
     39.571
                   -5.647
$`6`
```

I. Apply regression model over each data frame

```
mtcars %>%
  split(.$cyl) %>%
  map(\sim lm(mpg \sim wt, data = .)) \%>\%
  map(summary)
$`4`
Call:
lm(formula = mpg \sim wt, data = .)
Residuals:
    Min
            10 Median
                         30
                                   Max
-4.1513 -1.9795 -0.6272 1.9299 5.2523
```

- 1. Apply regression model over each data frame
- 2. Access results with the summary function

- 1. Apply regression model over each data frame
- 2. Access results with the summary function
- 3. Get specific results by subsetting each list item

- 1. Apply regression model over each data frame
- 2. Access results with the summary function
- 3. Get specific results by subsetting each list item

YOURTURN!

- 1. split the ggplot2::diamonds data set by cut
- 2. run a regression on each list item $lm(price \sim carat, data = .)$
- 3. get the summary of the regression
- 4. how does the "r.squared" compare across models?
- 5. how do the model slopes (aka "coefficients") compare?

SOLUTION

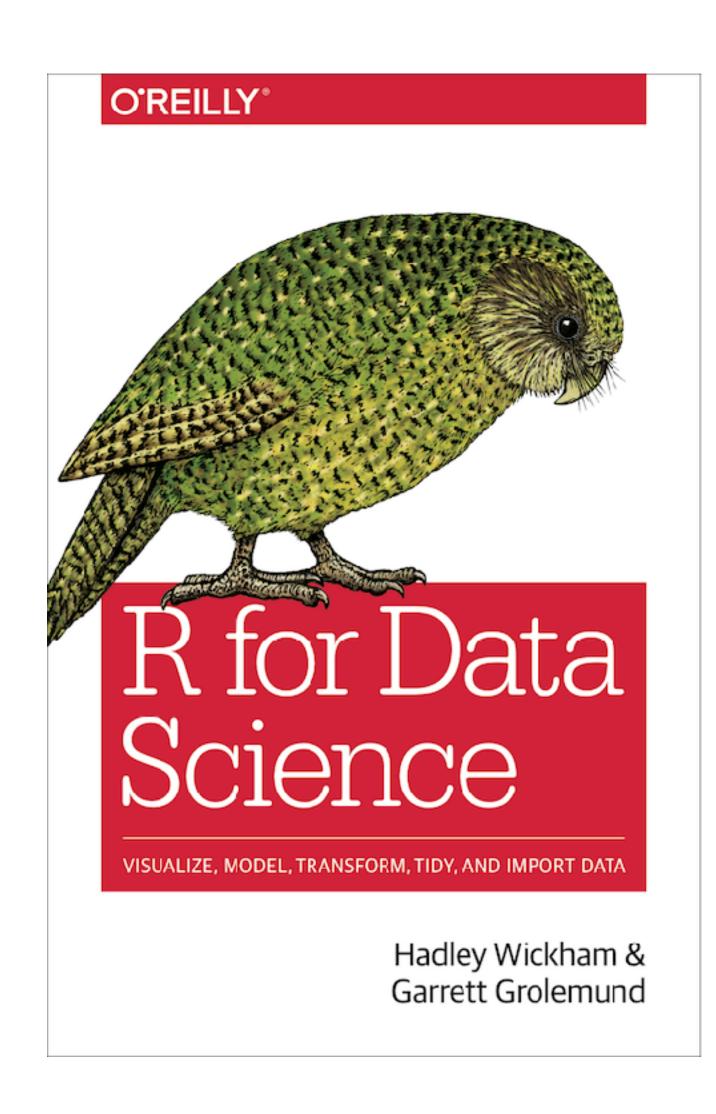
Compare R² values

SOLUTION

Compare slopes values



LEARN MORE



Use R! **Bradley Boehmke** Data Wrangling with R 2 Springer

WHATTO REMEMBER

FUNCTIONS TO REMEMBER

| Operator/Function | Description |
|--|---|
| if, ifelse, ifelse | Conditional control statements |
| for, while, repeat | Looping control statements |
| seq_along | Sequencing argument |
| break, next | Arguments to exit or skip a loop iteration |
| <pre>map, map_dbl, map_int, map_chr, map_lgl</pre> | Family of iteration functions |
| split | Split a data frame based on categorical variable levels |