

Lecture 2: Investigating data patterns

EDUC 263: Managing and Manipulating Data Using R

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1. Introduction
2. R Markdown
3. More R basics: functions and directories
4. Investigating objects, Base R approach
5. Investigating data frames, tidyverse approach

1 Introduction

What we will do today

1. Introduction
2. R Markdown
3. More R basics: functions and directories
 - 3.1 Introduction to using functions
 - 3.2 Directories and filepaths [PATRICIA - EDIT/COMPLETE THIS SECTION]
4. Investigating objects, Base R approach
 - 4.1 Variables names
 - 4.2 View and print data
 - 4.3 Missing values
5. Investigating data frames, tidyverse approach
 - 5.1 Select variables
 - 5.2 Filter rows
 - 5.3 Arrange rows

Libraries we will use today

“Load” the package we will use today (output omitted)

```
library(tidyverse)
```

If package not yet installed, then must install before you load. Install in “console” rather than .Rmd file

- Generic syntax: `install.packages("package_name")`
- Install “tidyverse”: `install.packages("tidyverse")`

Note: when we load package, name of package is not in quotes; but when we install package, name of package is in quotes:

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

2 R Markdown

What is R Markdown

Borrowing from Darin Christensen:

- R Markdown documents embed R code, the output associated with R code, and text into one document
- An R Markdown document is a “‘Living’ document that updates every time you compile [“knit”] it”
- R Markdown documents have the extension .Rmd
 - can think of them as text files with the extension .Rmd rather than .txt
- At top of .Rmd file you specify the “output” style, which dictates what kind of formatted document will be created
- When you compile [“knit”] a .Rmd file, the resulting formatted document can be an HTML document, a PDF document, an MS Word document, or many other types

How we will be using R Markdown files in this class:

- homework you submit will be .Rmd files, with “output” style will be `html_document` or `pdf_document`
- lectures we write are .Rmd files, where we the output style will usually be `beamer_presentation`
 - this is essentially a pdf document, where each page is a slide

Creating RMarkdown documents

Do this with a partner

Two approaches for creating an RMarkdown document.

1. Point-and-click from within RStudio

- ▶ Click on *File >> New File >> R Markdown >> Document >>* choose *HTML* >> click *OK*
- ▶ save the .Rmd file [any name, anywhere you can find it]
- ▶ “Knit” the entire .Rmd file
 - point-and-click OR shortcut: **Cmd/Ctrl + Shift + k**

2. From blank text file

- ▶ create blank text file
 - can give it any name, but change extension from .txt to .Rmd
- ▶ Open this blank .Rmd file in Rstudio
- ▶ copy this text into file [LINK HERE][PATRICIA CREATE LINK TO sample_simple_rmarkdown.txt FILE WHICH IS STORED IN LECTURE 2 DIRECTORY]
- ▶ “knit” the entire .Rmd file
 - point-and-click OR shortcut: **Cmd/Ctrl + Shift + k**

Components of a .Rmd file

An RMarkdown (.Rmd) file consists of several parts

1. **YAML header**

- ▷ YAML stands for “yet another markup language”
- ▷ controls settings that apply to the whole document (e.g., “output” should be `html_document` or `pdf_document`, whether to include table of contents, etc.)
- ▷ YAML header goes at very top of document
- ▷ starts with a line of three horizontal dashes `---`; ends with a line of three horizontal dashes `---`

2. **Text** in body of .Rmd file

- ▷ e.g., headings; description of results, etc.

3. **R code chunks** in body of .Rmd file

```
a <- c(2,4,6)
a
a-1
```

4. **R output** associated with code chunks

```
#> [1] 2 4 6
#> [1] 1 3 5
```

Comment: Running R code chunks vs. “knit” entire .Rmd file

Two ways to execute R commands in .Rmd file:

1. “Knit” entire .Rmd file

- ▶ shortcut: **Cmd/Ctrl + Shift + k**

2. “Run” code chunk or selected lines within code chunk

- ▶ Run selected line(s): **Cmd/Ctrl + Enter**

- ▶ Run current chunk: **Cmd/Ctrl + Shift + Enter**

Comment on default settings for RStudio:

- When you knit entire .Rmd file, “objects” created within .Rmd file will not be available after file completes
- When you run code chunk (or selected lines in chunk), objects created by lines you run will be in your “environment” until you remove them or quit R session

Output types of .Rmd file

Common/important output types:

- **html_document**: R Markdown originally designed to create HTML documents
 - ▷ Most features/code in .Rmd files were written for `html_document`
 - ▷ many of these features are available in other output types
 - ▷ When learning R Markdown, best to start by learning `html_document`
- **pdf_document**: Requires installation of LaTeX (MiKTeX)
 - ▷ How it works:
 - You write .Rmd code;
 - When you compile, this .Rmd code is transformed into LaTeX code
 - LaTeX “engine” creates the formatted .pdf file
 - ▷ Can include some of the same features available for *html_document*
 - ▷ Can insert LaTeX commands in .Rmd file with *pdf_document* output
- **beamer_presentation**: Requires installation of LaTeX
 - ▷ “beamer” is the name for presentations written in LaTeX
 - ▷ essentially creates PDF of presentation slides
 - ▷ Lectures for this class created with *beamer_presentation* output
 - ▷ note: YAML header includes `beamer_header.tex` file, which creates some formatting rules and additional commands

Learning more about R Markdown

Resources

- Cheat sheets and quick reference:
 - ▷ [Cheat Sheet](#)
 - ▷ [Quick Reference](#) [I prefer the quick reference]
- Chapters/books
 - ▷ [Chapter 27](#) of “R for Data Science” book
 - ▷ [R Markdown: The Definitive Guide](#) book [I prefer this book]

How you will learn R Markdown

- Lectures written as .Rmd file
 - ▷ During class run “code chunks” and try to “knit” entire .Rmd file
- I’ll assign **small** amount of reading on R Markdown
 - ▷ prior to next week:
 - spend 10-15 minutes familiarizing yourself with [Quick Reference](#)
 - Read section [3.1 of R Markdown: The Definitive Guide](#), about creating html_document
- Homework must be written in .Rmd file
 - ▷ you submit .Rmd file AND output of compiled file
 - ▷ for next week, you will submit homework as html_document output

3 More R basics: functions and directories

3.1 Introduction to using functions

What are functions

Functions are pre-written bits of code that accomplish some task.

Functions generally follow three sequential steps:

1. take in an **input** object(s)
2. **process** the input.
3. **return** (A) a new object or (B) a visualization (e.g., plot)

For example, `sum()` function calculates sum of elements in a vector

1. **input.** takes in a vector of elements (numeric or logical)
2. **processing.** Calculates the sum of elements
3. **return.** Returns numeric vector of length=1; value is sum of input vector

```
sum(c(1,2,3))
```

```
#> [1] 6
```

```
typeof(sum(c(1,2,3)))
```

```
#> [1] "double"
```

```
length(sum(c(1,2,3)))
```

```
#> [1] 1
```

```
sum(c(TRUE,TRUE,FALSE))
```

```
#> [1] 2
```

```
typeof(sum(c(TRUE,TRUE,FALSE))); length(sum(c(TRUE,TRUE,FALSE)))
```

```
#> [1] "integer"
```

```
#> [1] 1
```

Function syntax

Components of a function

- function name (e.g., `sum()` , `length()` , `seq()`)
- function arguments
 - Inputs that the function takes, which determine what function does
 - can be vectors, data frames, logical statements, etc.
 - In “function call” you specify values to assign to these function arguments
 - e.g., `sum(c(1,2,3))`
 - Separate arguments with a comma ,
 - e.g., `seq(10,15)` Example: the sequence function, `seq()`

```
seq(10,15)
```

```
#> [1] 10 11 12 13 14 15
```


Function syntax: More on function arguments

Usually, function arguments have names

- e.g., the `seq()` function includes the arguments `from`, `to`, `by`
- when you call the function, you need to assign values to these arguments; but you usually don't have to specify the name of the argument

```
seq(from=10, to=20, by=2)
#> [1] 10 12 14 16 18 20
seq(10,20,2)
#> [1] 10 12 14 16 18 20
```

Many function arguments have “default values”, set by whoever wrote function

- if you don't specify a value for that argument, the default value is inserted
- e.g., partial list of default values for `seq()`: `seq(from=1, to=1, by=1)`

```
seq()
#> [1] 1
seq(to=10)
#> [1] 1 2 3 4 5 6 7 8 9 10
seq(10) # R assigned value of 10 to "to" rather than "from" or "by"
#> [1] 1 2 3 4 5 6 7 8 9 10
```

Function arguments, the `na.rm` argument

When R performs calculation and an input has value `NA`, output value is `NA`

```
5+4+NA  
#> [1] NA
```

R functions that perform calculations often have argument named `na.rm`

- `na.rm` argument asks whether to remove `NA` values prior to calculation
- For most functions, default value is `na.rm = FALSE`
 - ▷ This means "do not remove `NA`s" prior to calculation
 - ▷ e.g., default values for `sum()` function: `sum(..., na.rm = FALSE)`

```
sum(c(1,2,3,NA), na.rm = FALSE) # default value  
#> [1] NA  
sum(c(1,2,3,NA))  
#> [1] NA
```

- if you specify, `na.rm = TRUE`, `NA` values removed prior to calculation

```
sum(c(1,2,3,NA), na.rm = TRUE)  
#> [1] 6
```

Help files for functions

To see help file on a function, type `?function_name` without parentheses

```
?sum
```

```
?seq
```

Contents of help files

- **Description.** What the function does
- **Usage.** Syntax, including default values for arguments
- **Arguments.** Description of function arguments
- **Details.** Details and idiosyncracies of about how the function works.
- **Value.** What (object) the function “returns”
 - ▷ e.g., `sum()` returns vector of length 1 whose value is sum of input vector
- **References.** Additional reading
- **See Also.** Related functions
- **Examples.** Examples of function in action
- Bottom of help file identifies the package the function comes from

Practice!

- when you encounter new function, spend two minutes reading help file
- over time, help files will feel less cryptic and will start to feel helpful

Function arguments, the dot-dot-dot (`...`) argument

On help file for many functions, you will see an argument called `...`, referred to as the “dot-dot-dot” argument

```
?sum  
?seq
```

“Dot-dot-dot” arguments have several uses. What you should know for now:

- `...` refers to arguments that are “un-named”; but user can specify values
 - ▷ e.g., default syntax for `sum()` : `sum(..., na.rm = FALSE)`
 - argument `na.rm` is “named” (name is `na.rm`); argument `...` un-named
- `...` used to allow a function to take an arbitrary number of arguments:

```
sum(c(10,5,NA),na.rm=TRUE)  
#> [1] 15
```

#Here the sum function takes 3 un-named arguments

```
sum(10,5,NA,na.rm=TRUE)  
#> [1] 15
```

#Here the sum function takes 5 un-named arguments

```
sum(10,5,10,20,NA,na.rm=TRUE)  
#> [1] 45
```

3.2 Directories and filepaths [PATRICIA - EDIT/COMPLETE THIS SECTION]

Working directory

(Current) Working directory

- the folder/directory in which you are currently working
- this is where R looks for files
- Files located in your current working directory can be accessed without specifying a filepath because R automatically looks in this folder

Function `getwd()` shows current working directory

```
getwd()
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
```

Command `list.files()` lists all files located in working directory

```
getwd()
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
list.files()
#> [1] "lecture2.pdf"           "lecture2.Rmd"
#> [3] "lecture2.tex"          "sample.html"
#> [5] "sample.Rmd"            "sample_simple_rmarkdown.txt"
#> [7] "text"                  "transform-logical.png"
```

Working directory, “Code chunks” vs. “console” and “R scripts”

When you run **code chunks** in RMarkdown files (.Rmd), the working directory is set to the filepath where the .Rmd file is stored

```
getwd()
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
list.files()
#> [1] "lecture2.pdf"           "lecture2.Rmd"
#> [3] "lecture2.tex"           "sample.html"
#> [5] "sample.Rmd"             "sample_simple_rmarkdown.txt"
#> [7] "text"                   "transform-logical.png"
```

When you run code from the **R Console** or an **R Script**, the working directory is...

Command `getwd()` shows current working directory

```
getwd()
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
```

Absolute vs. relative filepath

4 Investigating objects, Base R approach

Load .Rdata data frames we will use today

Data on off-campus recruiting events by public universities

- Data frame object `df_event`
 - One observation per university, recruiting event
- Data frame object `df_event`
 - One observation per high school (visited and non-visited)

PATRICIA; REPLACE FILE PATH W/ URL; KEEP EXISTING LOAD COMMANDS TOO BUT COMMENT THEM OUT

```
rm(list = ls()) # remove all objects in current environment

getwd()
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
#load dataset with one obs per recruiting event
load("../..data/recruiting/recruit_event_somevars.Rdata")

#load dataset with one obs per high school
load("../..data/recruiting/recruit_school_somevars.Rdata")
```

Listing objects

Files in your working directory

`list.files()` function lists files in your current working directory

- if you run this code from .Rmd file, working directory is location .Rmd file is stored

```
getwd() # what is your current working directory
#> [1] "C:/Users/ozanj/Documents/rclass/lectures/lecture2"
list.files()
#> [1] "lecture2.pdf"           "lecture2.Rmd"
#> [3] "lecture2.tex"          "sample.html"
#> [5] "sample.Rmd"            "sample_simple_rmarkdown.txt"
#> [7] "text"                  "transform-logical.png"
```

Objects currently open in your R session

`ls()` function lists objects currently open in R

```
x <- "hello!"
ls() # Objects open in R
#> [1] "df_event" "df_school" "x"
```

Removing objects

`rm()` function removes specified objects open in R

```
rm(x)  
ls()  
#> [1] "df_event" "df_school"
```

Command to remove all objects open in R (I don't run it)

```
rm(list = ls())
```

Describing objects, focus on **data frames**

type and **length** of a data frame object

- Recall that a data frame is an object where **type** is a list
- **Length** of an object is the number of elements
 - ▷ When object is a data frame, number of elements = number of variables

```
typeof(df_event)
#> [1] "list"
length(df_event) # = num elements = num columns
#> [1] 33
```

Number of **columns** and **rows** of data frame object

- number of columns = number of elements = number of variables
- number of rows = number of observations

```
ncol(df_event) # num columns = num variables
#> [1] 33
nrow(df_event) # num rows = num observations
#> [1] 17976
dim(df_event) # shows number rows by columns
#> [1] 17976    33
```

`str()` provides compact information on structure any object (output omitted)

```
str(df_event)
```

4.1 Variables names

Variable names

`names()` function lists names of elements in an object

?names

When object is a data frame:

- each element is a variable
- each element name is a variable name

```
names(df_event)
```

```
#> [1] "instnm" "univ_id" "instst"
#> [4] "pid" "event_date" "event_type"
#> [7] "zip" "school_id" "ipeds_id"
#> [10] "event_state" "event_inst" "med_inc"
#> [13] "pop_total" "pct_white_zip" "pct_black_zip"
#> [16] "pct_asian_zip" "pct_hispanic_zip" "pct_amerindian_zip"
#> [19] "pct_nativehawaii_zip" "pct_tworaces_zip" "pct_otherrace_zip"
#> [22] "fr_lunch" "titlei_status_pub" "total_12"
#> [25] "school_type_pri" "school_type_pub" "g12offered"
#> [28] "g12" "total_students_pub" "total_students_pri"
#> [31] "event_name" "event_location_name" "event_datetime_start"
```

Variable names

Refer to specific named elements of an object using this syntax:

- `obj_name$element_name`

When object is data frame, refer to specific variables using this syntax:

- `data_frame_name$varname`
- **This approach to isolating variables very useful for investigating data**

```
typeof(df_event$instnm)
#> [1] "character"
typeof(df_event$med_inc)
#> [1] "double"
```


Variable names

Recall that data frames are lists with following criteria:

- each element of the list is a vector
 - ▷ each element of list is a variable; length of data frame = number of variables

```
length(df_event)
```

```
#> [1] 33
```

```
nrow(df_event)
```

```
#> [1] 17976
```

```
#str(df_event)
```

- each element of the list (i.e., variable) has the same length
 - ▷ Length of each variable is equal to number of observations in data frame

```
typeof(df_event$event_state)
```

```
#> [1] "character"
```

```
length(df_event$event_state)
```

```
#> [1] 17976
```

```
str(df_event$event_state)
```

```
#> chr [1:17976] "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" ...
```

```
typeof(df_event$med_inc)
```

```
#> [1] "double"
```

```
length(df_event$med_inc)
```

```
#> [1] 17976
```

```
str(df_event$med_inc)
```

```
#> num [1:17976] 71714 89122 70137 70137 71024 ...
```

4.2 View and print data

Viewing and printing data frames

Three ways to view/print a data frame object

1. Simply type the object name (output omitted)

- ▶ number of observations and rows printed depend on YAML header settings and on attributes (discussed next week) of the object

```
df_event
```

2. Use the `View()` function to view data in a browser

```
View(df_event)
```

3. `head()` to show the first n rows

```
##?head  
head(df_event, n=5)
```

Viewing and printing data frames

`{obj_name[<rows>,<cols>]}` to print specific rows and columns of data frame

- particularly powerful when combined with sequences (e.g., `1:10`)

Examples:

- Print first five rows

```
df_event[1:5, ]
```

- Print first five rows and first three columns

```
df_event[1:5, 1:3]
```

- Print first three columns of the 100th observation

```
df_event[100, 1:3]
```

- Print the 50th observation, all variables

```
df_event[50,]
```

Viewing and printing data

type `obj_name$var_name` to print specific elements (i.e., variables) in a data frame

```
df_event$zip
```

- o recall that these elements are vectors, with length = number of obs

```
typeof(df_event$zip)
```

```
#> [1] "character"
```

```
length(df_event$zip)
```

```
#> [1] 17976
```

- o `obj_name$var_name` syntax can be combined with sequences
 - ▷ vectors don't have "rows" or "columns"; they just have elements
 - ▷ so use sequence to identify which elements you want to print

```
df_event$event_state[1:10]
```

```
#> [1] "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA"
```

```
df_event$event_type[6:10]
```

```
#> [1] private hs public hs private hs public hs public hs
```

```
#> Levels: public hs private hs 2yr college 4yr college other
```

- o can also print multiple variables using `combine()` function

```
c(df_event$event_state[1:5],df_event$event_type[1:5])
```

```
#> [1] "MA" "MA" "MA" "MA" "MA" "1" "1" "1" "1" "2"
```

Exercise

PATRICIA: PLEASE ADD TO THIS/MODIFY

Create a printing exercise using the df_school

1. Use `head()` to print first 5 observations
2. Use `obj_name$var_name[1:10]` to print the first 10 observations of a variable
3. use `combine()`

4.3 Missing values

Missing values

Missing values have the value `NA`

- `NA` is a special keyword, not the same as the character string `"NA"`

use `is.na()` function to determine if a value is missing

- `is.na()` returns a logical vector

```
is.na(5)
#> [1] FALSE
is.na(NA)
#> [1] TRUE
is.na("NA")
#> [1] FALSE
typeof(is.na("NA")) # example of a logical vector
#> [1] "logical"

nvector <- c(10,5,NA)
is.na(nvector)
#> [1] FALSE FALSE TRUE
typeof(is.na(nvector)) # example of a logical vector
#> [1] "logical"

svector <- c("e","f",NA,"NA")
is.na(svector)
#> [1] FALSE FALSE TRUE FALSE
```


Missing values are “contagious”

What does “contagious” mean?

- operations involving a missing value will yield a missing value

```
7>5
#> [1] TRUE
7>NA
#> [1] NA
0==NA
#> [1] NA
2*c(0,1,2,NA)
#> [1] 0 2 4 NA
NA*c(0,1,2,NA)
#> [1] NA NA NA NA
```

Function and missing values, the `table()` function

`table()` function useful for investigating categorical variables

```
table(df_event$g12offered)
#>
#>      1
#> 11025
```

By default `table()` ignores `NA` values

- `useNA` argument determines whether to include `NA` values
 - ▷ “allowed values correspond to never (“no”); only if count is positive (“ifany”); and even for zero counts (“always”)”

```
nrow(df_event)
#> [1] 17976
table(df_event$g12offered, useNA="always")
#>
#>      1  <NA>
#> 11025 6951
```

Broader point:

- Most functions that create descriptive statistics have options about how to treat missing values
- When investigating data, good practice to *a/ways* show missing values

Tip:

- command `str(df_event)` shows which variables have missing values

5 Investigating data frames, tidyverse approach

Introduction to the `dplyr` library

`dplyr`, a package within the `tidyverse` suite of packages, provide tools for manipulating data frames

- Wickham describes functions within `dplyr` as a set of “verbs” that fall in the broader categories of **subsetting**, **sorting**, and **transforming**

Today	Next two weeks
Subsetting data <ul style="list-style-type: none">- <code>select()</code> variables- <code>filter()</code> observations Sorting data <ul style="list-style-type: none">- <code>arrange()</code>	Transforming data <ul style="list-style-type: none">- <code>mutate()</code> creates new variables- <code>summarize()</code> calculates across rows- <code>group_by()</code> to calculate across rows within groups

All `dplyr` verbs (i.e., functions) work as follows

1. first argument is a data frame
2. subsequent arguments describe what to do with variables and observations in data frame
 - ▷ refer to variable names without quotes
3. result of the function is a new data frame

5.1 Select variables

Select variables using `select()` function

Printing observations is key to investigating data, but datasets often have hundreds, thousands of variables

`select()` function selects **columns** of data (i.e., variables) you specify

- first argument is the name of data frame object
- remaining arguments are variable names, which are separated by commas and without quotes

Without **assignment**, `select()` function by itself simply prints selected vars

```
select(df_event, instnm, event_date, event_type, event_state, med_inc)
#> # A tibble: 17,976 x 5
#>   instnm      event_date event_type event_state med_inc
#>   <chr>      <date>      <fct>      <chr>      <dbl>
#> 1 UM Amherst 2017-10-12 public hs    MA          71714.
#> 2 UM Amherst 2017-10-04 public hs    MA          89122.
#> 3 UM Amherst 2017-10-26 public hs    MA          70136.
#> 4 UM Amherst 2017-10-25 public hs    MA          70136.
#> 5 USCC       2017-09-18 private hs  MA          71024.
#> 6 UM Amherst 2017-09-18 private hs  MA          71024.
#> 7 Stony Brook 2017-10-02 public hs    MA          71024.
#> 8 UM Amherst 2017-09-26 private hs  MA          97225
#> 9 UM Amherst 2017-09-26 public hs    MA          97225
#> 10 UM Amherst 2017-10-12 public hs    MA          77800.
#> # ... with 17,966 more rows
```

Select variables using `select()` function

Recall that all `dplyr` functions (e.g., `select()`) return a new data frame object

- **type** equals "list"
- **length** equals number of vars you select

```
typeof(select(df_event, instnm, event_date, event_type, event_state, med_inc))  
#> [1] "list"
```

```
length(select(df_event, instnm, event_date, event_type, event_state, med_inc))  
#> [1] 5
```

`glimpse()` function – a tidyverse function for viewing data frames – is a cross between `str()` and simply printing data

```
##?glimpse  
glimpse(select(df_event, instnm, event_date, event_type, event_state, med_inc))  
#> Observations: 17,976  
#> Variables: 5  
#> $ instnm      <chr> "UM Amherst", "UM Amherst", "UM Amherst", "UM Amhe...  
#> $ event_date  <date> 2017-10-12, 2017-10-04, 2017-10-26, 2017-10-25, 2...  
#> $ event_type  <fct> public hs, public hs, public hs, public hs, privat...  
#> $ event_state <chr> "MA", "MA", "MA", "MA", "MA", "MA", "MA", "MA", "M...  
#> $ med_inc     <dbl> 71713.5, 89121.5, 70136.5, 70136.5, 71023.5, 71023.5...
```

Select variables using `select()` function

With **assignment**, `select()` creates a new object containing only the variables you specify

```
event_small <- select(df_event, instnm, event_date, event_type, event_state, med_inc)
glimpse(event_small)
#> Observations: 17,976
#> Variables: 5
#> $ instnm      <chr> "UM Amherst", "UM Amherst", "UM Amherst", "UM Amhe...
#> $ event_date  <date> 2017-10-12, 2017-10-04, 2017-10-26, 2017-10-25, 2...
#> $ event_type  <fct> public hs, public hs, public hs, public hs, privat...
#> $ event_state <chr> "MA", "MA", "MA", "MA", "MA", "MA", "MA", "MA", "M...
#> $ med_inc     <dbl> 71713.5, 89121.5, 70136.5, 70136.5, 71023.5, 71023...
```


Select

`select()` can use “helper functions” `starts_with()`, `contains()`, and `ends_with()` to choose columns

Example:

```
#names(df_event)
select(df_event, instnm, starts_with("event"))
#> # A tibble: 17,976 x 8
#>   instnm event_date event_type event_state event_inst event_name
#>   <chr>   <date>      <fct>      <chr>      <chr>      <chr>
#> 1 UM Am~ 2017-10-12 public hs MA          In-State Amherst-P~
#> 2 UM Am~ 2017-10-04 public hs MA          In-State Hampshire~
#> 3 UM Am~ 2017-10-26 public hs MA          In-State Chicopee ~
#> 4 UM Am~ 2017-10-25 public hs MA          In-State Chicopee ~
#> 5 USCC   2017-09-18 private hs MA          Out-State Williston~
#> 6 UM Am~ 2017-09-18 private hs MA          In-State Williston~
#> 7 Stony~ 2017-10-02 public hs MA          Out-State Easthampt~
#> 8 UM Am~ 2017-09-26 private hs MA          In-State MacDuffie~
#> 9 UM Am~ 2017-09-26 public hs MA          In-State Granby Jr~
#> 10 UM Am~ 2017-10-12 public hs MA          In-State Smith Aca~
#> # ... with 17,966 more rows, and 2 more variables:
#> #   event_location_name <chr>, event_datetime_start <dtm>
```

Exercise

The data frame `df_school` has one observation for each high school and indicators for whether the high school received a recruiting visit.

```
names(df_school)
```

1. Use `select()` to familiarize yourself with variables in the data frame
2. Practice using the `contains()` and `ends_with()` helper functions to choose variables

Rename variables

`rename()` function renames variables within a data frame object

Syntax:

- `rename(obj_name, new_name = old_name,...)`

```
rename(df_event, g12_offered = g12offered, titlei = titlei_status_pub)
names(df_event)
```

Variable names do not change permanently unless we combine rename with assignment

```
rename_event <- rename(df_event, g12_offered = g12offered, titlei = titlei_status_pub)
names(rename_event)
rm(rename_event)
```

5.2 Filter rows

The `filter()` function

`filter()` allows you to **select observations** based on values of variables

- Arguments
 - first argument is name of data frame
 - subsequent arguments are *logical expressions* to filter the data frame
 - Multiple expressions separated by commas work as **AND** operators (e.g., condition 1 `TRUE` AND condition 2 `TRUE`)
- What is the result of a `filter()` command?
 - `filter()` returns a data frame consisting of rows where the condition is `TRUE`

Example using data frame object `df_school`, where each observation is a high school

- Show all obs where the high school received 1 visit from UC Berkeley (110635) [output omitted]

```
filter(df_school,visits_by_110635 == 1)
```

Note that resulting object is list, consisting of obs where condition `TRUE`

```
nrow(df_school)
#> [1] 21301
nrow(filter(df_school,visits_by_110635 == 1))
#> [1] 528
```

The `filter()` function

PATRICIA - CONVERT THIS SLIDE INTO A STUDENT TASK PLEASE. THX! CAN PUT SOLUTION ON SUBSEQUENT SLIDE

Task

- Create a filter to identify all the high schools that recieved 1 visit from UC Berkeley (110635) AND 1 visit from CU Boulder (126614)[output omitted]

```
filter(df_school,visits_by_110635 == 1, visits_by_126614==1)

nrow(filter(df_school,visits_by_110635 == 1, visits_by_126614==1))
count(filter(df_school,visits_by_110635 == 1, visits_by_126614==1))
```

- Must **assign** to create new object based on filter

```
berk_boulder <- filter(df_school,visits_by_110635 == 1, visits_by_126614==1)
count(berk_boulder)
```

Filter, character variables

Use single quotes `' '` or double quotes `" "` to refer to values of character variables

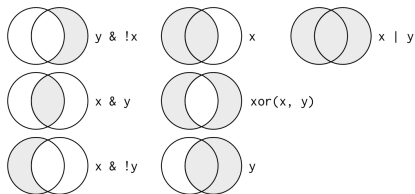
Below, we identify all private high schools in CA that got visit by particular universities

```
glimpse(df_school)
#Berkeley
filter(df_school,visits_by_110635 == 1, school_type == "private",
       state_code == "CA")
#Bama
filter(df_school,visits_by_100751 == 1, school_type == "private",
       state_code == "CA")

#Berkeley and Bama
filter(df_school,visits_by_100751 == 1, visits_by_110635 == 1,
       school_type == "private", state_code == "CA")
```

Logical operators for comparisons

Symbol	Meaning
<code>==</code>	Equal to
<code>!=</code>	Not equal to
<code>></code>	greater than
<code>>=</code>	greater than or equal to
<code><</code>	less than
<code><=</code>	less than or equal to
<code>&</code>	AND
<code> </code>	OR
<code>%in</code>	includes



PATRICIA - TRY TO PUT ON IMAGE ON

WEBSITE AND GIVE LINK TO URL

Filters and comparisons

PATRICIA - MAKE A CALL AS TO WHETHER THE NEXT FEW SLIDES SHOULD BE STUDENT EXERCISES OR DEMONSTRATIONS. THX!

Schools visited by Bama (100751) and/or Berkeley (110635)

```
#berkeley and bama
filter(df_school,visits_by_100751 >= 1, visits_by_110635 >= 1)
filter(df_school,visits_by_100751 >= 1 & visits_by_110635 >= 1) # same same
#berkeley or bama
filter(df_school,visits_by_100751 >= 1 | visits_by_110635 >= 1)
```

Apply `count()` function on top of `filter()` function to count the number of observations that satisfy criteria

- Avoids printing individual observations

```
#Number of schools that git visit by Berkeley AND Bama
count(filter(df_school,visits_by_100751 >= 1 & visits_by_110635 >= 1))
#> # A tibble: 1 x 1
#>       n
#>   <int>
#> 1   247
#Number of schools that git visit by Berkeley OR Bama
count(filter(df_school,visits_by_100751 >= 1 | visits_by_110635 >= 1))
#> # A tibble: 1 x 1
#>       n
#>   <int>
#> 1  2763
```

Filters and comparisons, >=

Number of public high schools that are at least 50% Black in Alabama compared to number of schools that received visit by Bama

```
#at least 50% black
count(filter(df_school, school_type == "public", pct_black >= 50,
             state_code == "AL"))
#> # A tibble: 1 x 1
#>       n
#>   <int>
#> 1     86
count(filter(df_school, school_type == "public", pct_black >= 50,
             state_code == "AL", visits_by_100751 >= 1))
#> # A tibble: 1 x 1
#>       n
#>   <int>
#> 1     21
#at least 50% white
count(filter(df_school, school_type == "public", pct_white >= 50,
             state_code == "AL"))
#> # A tibble: 1 x 1
#>       n
#>   <int>
#> 1    238
count(filter(df_school, school_type == "public", pct_white >= 50,
             state_code == "AL", visits_by_100751 >= 1))
#> # A tibble: 1 x 1
#>       n
```

Filters and comparisons, not equals (!=)

Count the number of high schools visited by University of Colorado (126614) that are not located in CO

```
#number of high schools visited by U Colorado  
count(filter(df_school, visits_by_126614 >= 1))  
#> # A tibble: 1 x 1  
#>       n  
#>   <int>  
#> 1  1056
```

```
#number of high schools visited by U Colorado not located in CO  
count(filter(df_school, visits_by_126614 >= 1, state_code != "CO"))  
#> # A tibble: 1 x 1  
#>       n  
#>   <int>  
#> 1   873
```

```
#number of high schools visited by U Colorado located in CO  
#count(filter(df_school, visits_by_126614 >= 1, state_code == "CO"))
```

Filters and comparisons, %in% operator

What if you wanted to count the number of schools visited by Bama (100751) in a group of states?

```
count(filter(df_school,visits_by_100751 >= 1, state_code == "MA" | state_code ==  
#> # A tibble: 1 x 1  
#>       n  
#>   <int>  
#> 1    108
```

Easier way to do this is with %in% operator

```
count(filter(df_school,visits_by_100751 >= 1, state_code %in% c("MA","ME","VT"))  
#> # A tibble: 1 x 1  
#>       n  
#>   <int>  
#> 1    108
```

Select the private high schools that got either 2 or 3 visits from Bama

```
count(filter(df_school, visits_by_100751 %in% 2:3, school_type == "private"))  
#> # A tibble: 1 x 1  
#>       n  
#>   <int>  
#> 1    183
```

Identifying data type and possible values of variable is helpful for filtering

PATRICIA - event_type IS CURRENTLY FACTOR VARIABLE, NOT CHARACTER [BUT YOU CAN REFER TO IT USING CHARACTER]; SHOULD WE CHANGE THIS VAR TO CHARACTER IN READ_CSV OR LEAVE IT AS FACTOR?

- `class()` and `str()` shows data type of a variable
- `table()` to show potential values of categorical variables

```
class(df_event$event_type)
#> [1] "factor"
str(df_event$event_type)
#> Factor w/ 5 levels "public hs","private hs",...: 1 1 1 1 2 2 1 2 1 1 ...
table(df_event$event_type)
#>
#>   public hs   private hs 2yr college 4yr college      other
#>         11025         3644         769         431         2107

class(df_event$event_state)
#> [1] "character"
str(df_event$event_state) # double quotes indicate character
#> chr [1:17976] "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" "MA" ...

class(df_event$med_inc)
#> [1] "numeric"
str(df_event$med_inc)
#> num [1:17976] 71714 89122 70137 70137 71024 ...
```

Exercises

PATRICIA: CREATE SEPARATE SLIDE W/ JUST THE QUESTION AND THEN SEPARATE SLIDE OF JUST THE ANSWERS; ALSO LOOK AT BELOW QUESTIONS AND MODIFY/DELETE/ADD IF YOU THINK OF BETTER OPTIONS. I WROTE THESE REAL QUICK.

Use the data from `df_event`, which has one observation for each off-campus recruiting event a university attends

1. Count the number of events attended by the University of Pittsburgh (Pitt)
`univ_id == 215293`
2. Count the number of recruiting events by Pitt at public or private high schools
3. Count the number of recruiting events by Pitt at public or private high schools located in the state of PA
4. Count the number of recruiting events by Pitt at public high schools not located in PA where median income is less than 100,000
5. Count the number of recruiting events by Pitt at public high schools not located in PA where median income is greater than or equal to 100,000
6. Count the number of out-of-state recruiting events by Pitt at private high schools or public high schools with median income of at least 100,000

Filtering and missing values

Wickham (2018) states:

- `filter()` only includes rows where condition is TRUE; it excludes both `FALSE` and `NA` values. To preserve missing values, ask for them explicitly:

Investigate var `df_event$fr_lunch`, number of free/reduced lunch students

- only available for visits to public high schools

```
#visits to public HS with less than 50 students on free/reduced lunch  
count(filter(df_event,event_type == "public hs", fr_lunch<50))
```

```
#> # A tibble: 1 x 1
```

```
#>       n
```

```
#>   <int>
```

```
#> 1     890
```

```
#visits to public HS, where free/reduced lunch missing
```

```
count(filter(df_event,event_type == "public hs", is.na(fr_lunch)))
```

```
#> # A tibble: 1 x 1
```

```
#>       n
```

```
#>   <int>
```

```
#> 1      26
```

```
#visits to public HS, where free/reduced is less than 50 OR is missing
```

```
count(filter(df_event,event_type == "public hs", fr_lunch<50 | is.na(fr_lunch)))
```

```
#> # A tibble: 1 x 1
```

```
#>       n
```

```
#>   <int>
```

```
#> 1     916
```

5.3 Arrange rows

arrange() function

`arrange()` function “arranges” rows in a data frame; said different, it sorts observations

Syntax: `arrange(x, ...)`

- First argument, `x`, is a data frame
- Subsequent arguments are a “comma separated list of unquoted variable names”

```
arrange(df_event, event_date)
```

Data frame goes back to previous order unless you **assign** the new order

```
df_event  
df_event <- arrange(df_event, event_date)  
df_event
```

arrange() function

Ascending and descending order

- `arrange()` sorts in **ascending** order by default
- use `desc()` to sort a column by descending order

```
arrange(df_event, desc(event_date))
```

Can sort by multiple variables

```
arrange(df_event, univ_id, desc(event_date), desc(med_inc))
```

```
#sort by university and descending by size of 12th grade class; combine with sel  
select(arrange(df_event, univ_id, desc(g12)), instnm, event_type, event_date, g12)
```

arrange() , missing values sorted at the end

Missing values automatically sorted at the end, regardless of whether you sort ascending or descending

Below, we sort by university, then by date of event, then by ID of high school

```
#by university, date, ascending school id
select(arrange(df_event, univ_id, desc(event_date), school_id),
        instnm,event_date,event_type,school_id)
```

```
#by university, date, descending school id
select(arrange(df_event, univ_id, desc(event_date), desc(school_id)),
        instnm,event_date,event_type,school_id)
```

Can sort by `is.na` to put missing values first

```
select(arrange(df_event, univ_id, desc(event_date), desc(is.na(school_id))),
        instnm,event_date,event_type,school_id)
#> # A tibble: 17,976 x 4
#>   instnm event_date event_type school_id
#>   <chr>   <date>     <fct>    <chr>
#> 1 Bama   2017-12-18 other     <NA>
#> 2 Bama   2017-12-18 private hs A9106483
#> 3 Bama   2017-12-15 other     <NA>
#> 4 Bama   2017-12-15 public hs 484473005095
#> 5 Bama   2017-12-15 public hs 062927004516
#> 6 Bama   2017-12-14 other     <NA>
#> 7 Bama   2017-12-13 other     <NA>
#> 8 Bama   2017-12-13 public hs 130387001439
```

Exercises, arranging

PATRICIA - CAN YOU CREATE A COUPLE OF EXERCISES FOR ARRANGE?