Lecture 2: Investigating data patterns

EDUC 263: Managing and Manipulating Data Using R

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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
- 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

- o Generic syntax: install.packages("package_name")
- o 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:

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



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"
- o R Markdown documents have the extension .Rmd
 - ocan 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
 - b this is essentially a pdf document, where each page is a slide

Creating RMarkdown documents

Do this with a partner

Approach for creating a 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_simp

# - "knit" the entire .Rmd file

# - point-and-click OR shortcut: __Cmd/Ctrl + Shift + k__

#Don't worry about this. Ozan's notes
```

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
```

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 comples
- 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/MacTeX)
 - ▶ 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

- o 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 Definative Guide book [I prefer this book]

How you will learn R Markdown

- o Lectures written as .Rmd file
 - During class run "code chunks" and try to "knit" entire .Rmd file
- o 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 Definative Guide, about creating html_document
- o 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



3.1 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 visualizatoin (e.g., plot)

For example, sum() function calcualtes 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

- o function name (e.g., sum() , length() , seq())
- o 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

- o 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

- $\circ\,$ if you don't specify a value for that argument, the default value is inserted
- o 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 $\,^{\mathrm{NA}}$, output value is $\,^{\mathrm{NA}}$

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

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

- o na.rm argument asks whether to remove NA values prior to calculation
- o For most functions, default value is na.rm = FALSE
 - ▶ This means "do not remove NAs " 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
```

o 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 <code>?function_name</code> without parentheses

?sum ?seq

Contents of help files

- o Description. What the function does
- o **Usage**. Syntax, including default values for arguments
- o Arguments. Description of function arguments
- o **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
- o References. Additional reading
- o See Also. Related functions
- Examples. Examples of function in action
- o Bottom of help file identifies the package the function comes from

Practice!

- o when you encounter new function, spend two minutes reading help file
- o 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 $\ \dots$, referred to as the "dot-dot-dot" argument

```
?sum
?seq
```

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

- o 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

Working directory

(Current) Working directory

- the folder/directory in which you are currently working
- o 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] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
```

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

```
getwd()
#> [1] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG" "fp2.JPG"
#> [3] "lecture2.pdf" "lecture2.Rmd"
#> [5] "lecture2.tex" "sample_simple_rmarkdown.txt"
#> [7] "sample.Rmd" "text"
#> [9] "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] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG" "fp2.JPG"
#> [3] "lecture2.pdf" "lecture2.Rmd"
#> [5] "lecture2.tex" "sample_simple_rmarkdown.txt"
#> [7] "sample.Rmd" "text"
#> [9] "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] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
```

Absolute vs. relative filepath

Absolute file path: The absolute file path is the complete list of directories needed to locate a file or folder.

setwd("/Users/pm/Desktop/rclass/lectures/lecture2")

Relative file path: The relative file path is the path relative to your current location/directory.

Assuming your current working directory is in the "lecture2" folder and you want to change your directory to the data folder, your relative file path would look something like this:

```
setwd("../../data")
```

File path shortcuts

Key	Description
~	tilde is a shortcut for user's home directory (mine is my name pmartin)
/	moves up a level
//	moves up two level

Exercise

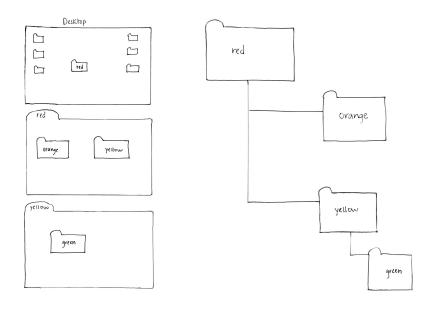
- 1. Let's create a folder on our desktop and name it red
- 2. Inside the red folder, create two subfolders named orange and yellow
- 3. Inside the yellow folder create another subfolder named green

Make sure to name these folders in lowercase.

You should have 1 folder on your desktop called red. Inside the red folder you have two folders called orange and yellow. Inside the yellow folder you have a folder called green.

Here is a visual of how it should look...

File path visual



Exercise continued

Let's say we want to get to the green folder using the absolute file path.

- 1. View your current working directory getwd()
- 2. Set your working directory to the green folder using the absolute file path
- 3. Now set your working directory to the orange folder using the relative file path (hint: ./)

Solution

```
getwd()
#> [1] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
setwd("~/Desktop/red/yellow/green")
getwd()
#> [1] "/Users/patriciamartin/Desktop/red/yellow/green"
setwd("../../orange")
getwd()
#> [1] "/Users/patriciamartin/Desktop/red/orange"
```

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_school
 - One observation per high school (visited and non-visited)

```
rm(list = ls()) # remove all objects in current environment
getwd()
#> [1] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
#load dataset with one obs per recruiting event
load(url("https://github.com/ozanj/rclass/raw/master/data/recruiting/recruit_eve
#load("../../data/recruiting/recruit_event_somevars.Rdata")

#load dataset with one obs per high school
load(url("https://github.com/ozanj/rclass/raw/master/data/recruiting/recruit_sch
#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] "/Users/patriciamartin/Desktop/GitHub/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG" "fp2.JPG"
#> [3] "lecture2.pdf" "lecture2.Rmd"
#> [5] "lecture2.tex" "sample_simple_rmarkdown.txt"
#> [7] "sample.Rmd" "text"
#> [9] "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

- o number of columns = number of elements = number of variables
- o 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
```

```
\begin{tabular}{ll} \tt str() & provides compact information on structure any object (output omitted) \\ \tt str(df\_event) & \end{tabular}
```

4.1 Variables names

Variable names

names() function lists names of elements in an object

?names

When object is a data frame:

- o each element is a variable
- o 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"
                                                       "pct amerindian_zip"
#> [16] "pct_asian_zip"
                               "pct_hispanic_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:

```
o obj_name$element_name
```

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

- o data_fram_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)
```

- o 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

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

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

Examples:

Print first five rows

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

o Print first five rows and first three columns

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

o Print first three columns of the 100th observation

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

o 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

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"
#> [6] "public hs" "public hs" "public hs" "public hs" "private hs"
```

Exercise

Create a printing exercise using the df_school

- 1. Use obj_name(<rows>,<cols>) to print the first 5 rows and 3 columns of data frame
- 2. Use head() to print first 4 observations
- Use obj_name\$var_name[1:10] to print the first 10 observations of a variable
- Use combine() to print the first 3 observations of variables "school_type" & "name"

1. Use obj_name[<rows>,<cols>] to print the first 5 rows and 3 columns of data frame

2. Use head() to print first 4 observations

```
head(df school, n=4)
#> # A tibble: 4 x 26
#> state code school type ncessch name address city zip code pct white
#> <chr> <chr
                                                                               <db1>
#> 1 AK public 020000~ Beth~ 1006 R~ Beth~ 99559
                                                                                11.8
#> 2 AK public 020000~ Ayag~ 106 Vi~ Kong~ 99559
#> 3 AK
                 public
                                020000~ Kwig~ 108 Vi~ Kwig~ 99622
#> 4 AK
                 public
                                020000~ Nels~ 118 Vi~ Toks~ 99637
#> # ... with 18 more variables: pct black <dbl>, pct hispanic <dbl>,
#> # pct asian <dbl>, pct_amerindian <dbl>, pct_other <dbl>,
#> #
        num_fr_lunch <dbl>, total_students <dbl>, num_took_math <dbl>,
#> #
        num prof math <dbl>, num took rla <dbl>, num prof rla <dbl>,
#> #
        avgmedian inc 2564 <dbl>, visits by 110635 <int>,
#> # visits by 126614 <int>, visits by 100751 <int>, inst 110635 <chr>,
        inst 126614 <chr>, inst 100751 <chr>
#> #
```

 Use obj_name\$var_name[1:10] to print the first 10 observations of a variable

```
df_school$name[1:10]
#> [1] "Bethel Regional High School" "Ayagina'ar Elitnaurvik"
#> [3] "Kwigillingok School" "Nelson Island Area School"
#> [5] "Alakanuk School" "Emmonak School"
#> [7] "Hooper Bay School" "Ignatius Beans School"
#> [9] "Pilot Station School" "Kotlik School"
```

Use combine() to print the first 3 observations of variables "school_type" & "name"

4.3 Missing values

Missing values

Missing values have the value NA

 \circ NA is a special keyword, not the same as the character string "NA"

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

o 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 \leftarrow 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")</pre>
is.na(svector)
#> [1] FALSE FALSE TRUE FALSE
```

Missing values are "contageous"

What does "contageous" mean?

o 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

- o 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 always show missing values

Tip:

o 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			Transforming data			
-	select()	variables	-	<pre>mutate()</pre>	cr	eates new variables
-	filter()	observations	-	summarize	()	calculates across rows
Sorting data		-	group_by()	to calculate across rows within groups	
-	arrange())				

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

- 1. first argument is a data frame
- 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

- o 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> <chr> <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

- o type equals "list"
- o **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
```

 $\begin{tabular}{ll} glimpse() & function - a tidyverse function for viewing data frames - is a cross \\ between & str() & and simply printing data \\ \end{tabular}$

Select variables using select() function

With **assignment**, <code>select()</code> creates a new object containing only the variables you specify

Select

```
{\tt select()} \ \ {\tt can} \ \ {\tt use} \ \ "{\tt helper} \ \ {\tt functions}" \ \ {\tt starts\_with()} \ , \ \ {\tt contains()} \ , \ {\tt and} \ \\ {\tt ends\_with()} \ \ \ {\tt to} \ \ {\tt choose} \ \ {\tt 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> <chr>
                              <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
                                                  Williston~
                                        Out-State
#> 6 UM Am~ 2017-09-18 private hs MA
                                                  Williston~
                                        In-State
#> 7 Stony~ 2017-10-02 public hs MA
                                        Out-State
                                                  Easthampt~
#> 8 UM Am~ 2017-09-26 private hs MA
                                                  MacDuffie~
                                        In-State
#> 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 <dttm>
```

Exercise

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

```
names(df_school)
```

- 1. Use <code>select()</code> to familiarize yourself with variables in the data frame
- Practice using the contains() and ends_with() helper functions to to choose variables

Rename variables

rename() function renames variables within a data frame object

Syntax:

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

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

```
rename_event <- rename(df_event, g12_offered = g12offered, titlei = titlei_statu
names(rename_event)
rm(rename_event)</pre>
```

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., condtion 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 ${\tt df_school}$, 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
```

Exercise

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))
```

o 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)</pre>
```

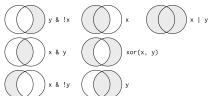
Filter, character variables

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

Logical operators for comparisons

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



WEBSITE AND GIVE LINK TO URL

PATRICIA - TRY TO PUT ON IMAGE ON

Filters and comparisons, Demonstration

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

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
#>
```

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?

Easier way to do this is with %in% operator

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

Identifying data type and possible values of variable is helpful for filtering of class() and str() shows data type of a variable

o table() to show potential values of categorical variables

```
class(df_event$event_type)
#> [1] "character"
str(df_event$event_type)
#> chr [1:17976] "public hs" "public hs" "public hs" "public hs" ...
table(df event$event type)
#>
#> 2yr college 4yr college other private hs public hs
      769
             431
                        2107 3644
                                              11025
#>
class(df event$event state)
#> [1] "character"
str(df event$event state) # double quotes indicate character
class(df event$med inc)
#> [1] "numeric"
str(df_event$med_inc)
#> num [1:17976] 71714 89122 70136 70136 71024 ...
```

Now that we know event_type is a character, we can filter values

```
count(filter(df_event, event_type == "public hs", event_state =="CA"))
```

Exercises

[Patricia]

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

- Count the number of events attended by the University of Pittsburgh (Pitt) univ_id == 215293
- 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
- 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
- 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

1. Count the number of events attended by the University of Pittsburgh (Pitt) univ_id == 215293

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

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

o 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))</pre>
#> # 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)))</pre>
#> # 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,...)

- o 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</pre>
```

arrange() function

Ascending and descending order

- o arrange() sorts in **ascending** order by default
- o use <code>desc()</code> 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 $% \left(1\right) =\left(1\right) \left(1\right)$

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

2017-12-12 public bg 120297001/20

#\ Q Dama

