Managing and Manipulating Data Using R

Lecture 2

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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. Find homework groups
- 6. Investigating data frames, tidyverse approach
 - 6.1 Select variables
 - 6.2 Filter rows
 - 6.3 Arrange rows

Logistics

- 1. Homework expectations
 - bdue date: before class (except first homework due today)
 - > general instructions for homework
 - grading
- 2. If we don't get through all of lecture, finish on your own
 - > whenever we don't cover difficult concepts in class; I'll cover those concepts next week and slow down the pace of class

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")
- 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:

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

Approach for creating a RMarkdown document.

- 1. Point-and-click from within RStudio
 - ightharpoonup 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

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

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

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
- 2. "Run" code chunk or selected lines within code chunk
 - > Run selected line(s): Cmd/Ctrl + 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 https://document.com/https://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 Definative Guide book [I prefer this book]

How you will learn R Markdown

> prior to next week:

- 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
 - spend 10-15 minutes familiarizing yourself with Quick Reference

 - Read section 3.1 of R Markdown: The Definative Guide, about creating html_document
- Homework must be written in Rmd file
 - > vou 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 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())
- 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 ,

```
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

- o 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

#> \[17 \] 6

```
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
  o 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 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)
```

Help files for functions

To see help file on a function, type <code>?function_name</code> 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
- 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 ..., referred to as the "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) #> \[11 \] 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

getwd()

(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

```
#> [1] "/Users/Karina/rclass/lectures/lecture2"
Command list.files() lists all files located in working directory
getwd()
#> [1] "/Users/Karina/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG"
                                           "fp2.JPG"
                                          "lecture2.aux"
#> [3] "lecture2 ua.Rmd"
#> [5] "lecture2.knit.md"
                                          "lecture2.log"
#> [7] "lecture2.out"
                                          "lecture2.pdf"
#> [9] "lecture2.Rmd"
                                          "Lecture2, tex"
#> [11] "lecture2.utf8.md"
                                          "lecture2.vrb"
#> [13] "problemset2 solutions.html"
                                           "problemset2_solutions.html.zip"
#> [15] "problemset2_solutions.Rmd"
                                           "problemset2.html"
#> [17] "problemset2.Rmd"
                                           "sample\_simple\_rmarkdown.txt"
#> [19] "sample.html"
                                           "sample.Rmd"
#> [21] "text"
                                           "transform-logical.png"
```

Working directory, "Code chunks" vs. "console" and "R scripts"

When you run ${\bf code\ chunks}$ in RMarkdown files (.Rmd), the working directory is set to the filepath where the .Rmd file is stored

```
getwd()
#> [1] "/Users/Karina/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG"
                                          "fp2. JPG"
#> [3] "lecture2 ua.Rmd"
                                          "lecture2. aux"
#> [5] "lecture2.knit.md"
                                          "lecture2.log"
#> [7] "lecture2.out"
                                          "lecture2.pdf"
#> [9] "lecture2.Rmd"
                                          "Lecture2, tex"
#> [11] "lecture2.utf8.md"
                                          "lecture2.vrb"
#> [13] "problemset2 solutions.html"
                                          "problemset2 solutions.html.zip"
#> [15] "problemset2 solutions.Rmd"
                                          "problemset2.html"
#> [17] "problemset2.Rmd"
                                          "sample simple rmarkdown.txt"
#> [19] "sample.html"
                                          "sample.Rmd"
#> [21] "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] "/Users/Karina/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 pm) moves up a level
//	moves up two level

Exercise

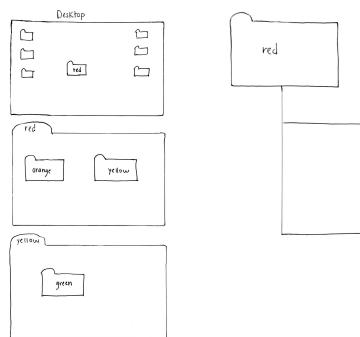
- 1. Let's create a folder on our desktop and name it red
- 2. Inside the red folder, create two subfolders named orange and yellow
- 3. Inside the yellow folder create another subfolder named green

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



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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: $\ldots/)$

Solution

```
getwd()
setwd("~/Desktop/red/yellow/green")
getwd()
setwd(".../orange")
getwd()
```

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/Karina/rclass/lectures/lecture2"

 #load dataset with one obs per recruiting event

 load(url("https://github.com/ozanj/rclass/raw/master/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_sc
#load("../../data/recruiting/recruit_school_somevars.Rdata")
```

Listing objects

Files in your working directory

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

 $\circ\,$ 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/Karina/rclass/lectures/lecture2"
list.files()
#> [1] "fp1.JPG"
                                         "fp2. JPG"
#> [3] "lecture2 ua.Rmd"
                                         "lecture2.aux"
#> [5] "lecture2.knit.md"
                                         "lecture2.log"
                                         "lecture2.pdf"
#> [7] "lecture2.out"
#> [9] "lecture2.Rmd"
                                         "lecture2 tex"
#> [11] "lecture2.utf8.md"
                                      "Lecture2.urb"
#> [13] "problemset2_solutions.html"
                                         "problemset2 solutions.html.zip"
#> [15] "problemset2 solutions.Rmd"
                                         "problemset2.html"
#> [17] "problemset2.Rmd"
                                         "sample\_simple\_rmarkdown.txt"
#> [19] "sample.html"
                                         "sample.Rmd"
#> [21] "text"
                                         "transform-logical.png"
```

Objects currently open in your R session

1s() function lists objects currently open in R

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

Removing objects

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

Command to remove all objects open in R (I don't run it)
rm(list = ls())
```

rm() function removes specified objects open in R

Describing objects, focus on data frames

type and length of a data frame object

- $\circ\,$ 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] 18680
dim(df_event) # shows number rows by columns
#> [1] 18680 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
- 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"
#> [16] "pct asian 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"
                            "total_students_pub" "total_students pri"
#> [28] "q12"
#> [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:

- 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] 18680
            #str(df_event)

    each element of the list (i.e., variable) has the same length
```

Variable names

Recall that object df_school has one obs per high school

- the variable visits_by_100751 shows number of visits by University of Alabama to each high school
- like all variables in a data frame, the var visits_by_100751 is just a vector

```
typeof(df_school$visits_by_100751)
#> [1] "integer"
length(df_school$visits_by_100751) # num elements in vector
#> [1] 21301
sum(df_school$visits_by_100751)
#> [1] 3338
```

Sp we perform calculations on a variable, just like we would any numeric vector

```
v <- c(2,4,6)
typeof(v)
#> [1] "double"
length(v)
#> [1] 3
sum(v)
#> [1] 12
```

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

```
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,]
```

obj_name[<rows>,<cols>] to print specific rows and columns of data frame
o particularly powerful when combined with sequences (e.g., 1:10)

Viewing and printing data

#> [1] "MA"

```
type obj_name$var_name to print specific elements (i.e., vars) in data frame
df event$zip

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

typeof(df event$zip)
#> [1] "character"
length(df event$zip)
#> [1] 18680

    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$event state[1:10]
df_event$event_type[6:10]
#> [1] "private hs" "private hs" "public hs" "private hs" "public hs"
Can also print multiple variables using combine() function
```

"MA"

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

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

"MA"

"MA "

Exercise

Create a printing exercise using the df_school data frame

- 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
- 3. 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"

 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
#> 1 AK public
                                 020000~ Beth~ 1006 R~ Beth~ 99559
                                                                                    11.8
                                 020000~ Ayaq~ 106 Vi~ Konq~ 99559
#> 2 AK
                public
                                                                                      0
#> 3 AK public
                                 020000~ Kwiq~ 108 Vi~ Kwiq~ 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>
```

3. 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

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 \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

#> 11423
```

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] 18680
table(df_event$g12offered, useNA="always")
#>
#> 1 <NA>
#> 11423 7257
```

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:

command str(df_event) shows which variables have missing values

5 Find homework groups

What we'll do to choose homework groups

- Meet new people (10-15 minutes of speed-dating!)
 - find someone in class you don't know and talk to them for three minutes about whatever!
 - e.g., where you from, what program, what are research interests, what you like doing outside of school, work
- We'll spend 10-15 minutes where enrolled students choose homework groups
 must be groups of 3
 - > cannot have more than 2 people from same academic program (e.g., HEOC, HDP)
- Auditors not part of "official" homework groups of 3, but they are welcome to join any homework group or form their own homework group

Use Zoom for group meetings! - https://ucla.zoom.us/

6 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	- mutate() creates 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

6.1 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: 18,680 x 5
#> instnm event_date event_type event_state med_inc
\#> < chr> < date> < chr>
                                              <d.b1.>
#> 1 UM Amherst 2017-10-12 public hs MA
                                            71714.
#> 2 UM Amherst 2017-10-04 public hs MA
                                             89122.
#> 3 UM Amherst 2017-10-25 public hs MA 70136.
#> 4 UM Amherst 2017-10-26 public hs MA
                                           70136.
#> 5 Stony Brook 2017-10-02 public hs MA
                                          71024.
#> 6 USCC 2017-09-18 private hs MA
                                          71024.
#> 7 UM Amherst 2017-09-18 private hs MA
                                          71024.
#> 8 UM Amherst 2017-09-26 public hs MA
                                         97225
#> 9 UM Amherst 2017-09-26 private hs MA
                                           97225
#> 10 UM Amherst 2017-10-12 public hs MA
                                             77800.
#> # ... with 18.670 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
#?qlimpse
glimpse(select(df_event,instnm,event_date,event_type,event_state,med_inc))
#> Observations: 18,680
#> Variables: 5
#> $ instnm <chr> "UM Amherst", "UM Amherst", "UM Amherst", "UM Amhe...
#> $ event date <date> 2017-10-12, 2017-10-04, 2017-10-25, 2017-10-26, 2...
#> $ event type <chr> "public hs", "public hs", "public hs", "public hs"...
#> $ event_state <chr> "MA", "
#> $ med inc <dbl> 71713.5, 89121.5, 70136.5, 70136.5, 71023.5, 71023...
```

Select variables using select() function

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

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: 18,680 x 8
#> instnm event date event type event state event inst event name
#> <chr> <date> <chr> <chr> <chr>
                                        In-State
                                                  Amherst-P~
#> 1 UM Am~ 2017-10-12 public hs MA
\# 2 UM Am~ 2017-10-04 public hs MA In-State Hampshire~
#> 3 UM Am~ 2017-10-25 public hs MA
                                        In-State Chicopee ~
#> 4 UM Am~ 2017-10-26 public hs MA
                                        In-State
                                                  Chicopee ~
#> 5 Stony~ 2017-10-02 public hs MA Out-State Easthampt~
                                       Out-State Williston~
#> 6 USCC 2017-09-18 private hs MA
#> 7 UM Am~ 2017-09-18 private hs MA
                                       In-State
                                                  Wi.1.1.i.st.on~
#> 8 UM Am~ 2017-09-26 public hs MA In-State Granby Jr~
#> 9 UM Am~ 2017-09-26 private hs MA
                                       In\text{-}State
                                                  MacDuffie~
#> 10 UM Am~ 2017-10-12 public hs MA
                                        In\text{-}State
                                                  Smith Aca-
#> # ... with 18,670 more rows, and 2 more variables:
#> # event location name <chr>, event datetime start <dttm>
```

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
- 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_stat
names(rename_event)
rm(rename_event)</pre>
```

6.2 Filter rows

The filter() function

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

- Arguments
 - b 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 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))

• 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

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

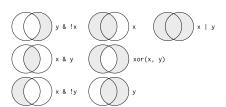


Figure 1: "Boolean" operations, x=left circle, y=right circle, from Wichkam (2018)

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.
\#> \langle i,n,t,>
#> 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 2.1
#at least 50% white
count(filter(df school, school type == "public", pct white >= 50,
             state code == "AL"))
#> # A tibble: 1 x 1
#>
         n
#>
   \langle i, n, t, \rangle
#> 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
#>
#> <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.
\#> \langle i,n,t,>
#> 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 shows data type of a variable

table() to show potential values of categorical variables

```
class(df event$event type)
#> [1] "character"
str(df event$event type)
#> chr [1:18680] "public hs" "public hs" "public hs" "public hs" ...
table(df_event$event_type)
#>
#> 2yr college 4yr college other private hs public hs
    951
                  531 2001 3774 11423
#>
class(df_event$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 [1:18680] 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"))
#> # A tibble: 1 x 1
#> n
#> cont
```

Exercises

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
- 2. Count the number of recruiting events by Pitt at public or private high schools
- Count the number of recruiting events by Pitt at public or private high schools located in the state of PA
- 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

 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

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

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:"

```
#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
#>
\#> \langle i,n,t,>
#> 1 910
#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
#>
#> <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
#>
#> <int>
#> 1
       936
```

6.3 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

```
o 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 sell
```

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: 18,680 x 4
#> instnm event date event type school id
\#> < chr> < date> < 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>

Exercise, arranging

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

- 1. Sort ascending by "univ_id" and descending by "event_date"
- Select four variables in total and sort ascending by "univ_id" and descending by "event_date"
- Now using the same variables from above, sort by is.na to put missing values in "school_id" first

1. Sort ascending by "univ_id" and descending by "event_date"

```
arrange(df_event, univ_id, desc(event_date))
#> # A tibble: 18,680 x 33
    instrm univ id instst pid event date event type zip school id
#>
#> <chr> <int> <chr> <int> <chr> <chr> <chr> <chr> <chr>
#> 1 Bama 100751 AL
                          7115 2017-12-18 private hs 77089 A9106483
   2 Bama 100751 AL
                          7121 2017-12-18 other <NA> <NA>
#>
#>
  3 Bama 100751 AL
                          7114 2017-12-15 public hs 75165 48447300~
#> 4 Bama 100751 AL
                          7100 2017-12-15 public hs 93012 06292700~
   5 Bama 100751 AL
                       7073 2017-12-15 other
                                                   98027 <NA>
#>
#> 6 Bama 100751 AL
                       7072 2017-12-14 other 98007 <NA>
#> 7 Bama 100751 AL
                          7118 2017-12-13 public hs 31906 13038700~
#> 8 Bama 100751 AL 7099 2017-12-13 private hs 90293 00071151
#>
   9 Bama 100751 AL 7109 2017-12-13 public hs 92630 06338600~
            100751 AL. 7071 2017-12-13 other
#> 10 Ba.ma.
                                                   98032 <NA>
#> # ... with 18,670 more rows, and 25 more variables: ipeds id <int>,
#> #
      event state <chr>, event inst <chr>, med inc <dbl>, pop total <dbl>,
#> #
      pct white zip <dbl>, pct black zip <dbl>, pct asian zip <dbl>,
#> #
      pct hispanic zip <dbl>, pct amerindian zip <dbl>,
#> #
      pct nativehawaii zip <dbl>, pct tworaces zip <dbl>,
#> #
      pct_otherrace_zip <dbl>, fr_lunch <dbl>, titlei_status_pub <fct>,
#> #
      total 12 <dbl>, school type pri <int>, school type pub <int>,
      q12offered <dbl>, q12 <dbl>, total students pub <dbl>,
#> #
#> #
      total students pri <dbl>, event name <chr>, event location name <chr>,
      event datetime start <dttm>
#> #
```

Select four variables in total and sort ascending by "univ_id" and descending by "event_date"

```
select(arrange(df_event, univ_id, desc(event_date)), univ_id, event_date,
      instnm, event type)
#> # A tibble: 18.680 x 4
#> univ id event date instnm event type
\#> <int><date> <chr><
#> 1 100751 2017-12-18 Bama private hs
#> 2 100751 2017-12-18 Bama other
   3 100751 2017-12-15 Bama public hs
#>
#>
  4 100751 2017-12-15 Bama public hs
#>
   5 100751 2017-12-15 Bama other
#> 6 100751 2017-12-14 Bama other
#> 7 100751 2017-12-13 Bama public hs
#> 8 100751 2017-12-13 Bama private hs
#> 9 100751 2017-12-13 Bama public hs
#> 10 100751 2017-12-13 Bama other
#> # ... with 18,670 more rows
```

 Select the variables "univ_id", "event_date", and "school_id" and sort by is.na to put missing values in "school_id" first.

```
select(arrange(df_event, univ_id, desc(event_date), desc(is.na(school_id))),
       univ id, event date, school id)
#> # A tibble: 18,680 x 3
#> univ id event date school id
#>
     \langle int \rangle \langle date \rangle \langle chr \rangle
#> 1 100751 2017-12-18 <NA>
#> 2 100751 2017-12-18 A9106483
#> 3 100751 2017-12-15 <NA>
   4 100751 2017-12-15 484473005095
#>
#> 5 100751 2017-12-15 062927004516
#> 6 100751 2017-12-14 <NA>
#> 7 100751 2017-12-13 <NA>
#> 8 100751 2017-12-13 130387001439
#> 9 100751 2017-12-13 00071151
#> 10 100751 2017-12-13 063386005296
#> # ... with 18,670 more rows
```