Intro to R

Data Input/Output

Outline

- Part 1: reading CSV file, common new user mistakes in data reading, checking for problems in the read data
- Part 2: data input overview, working directories, relative vs. absolute paths, reading XLSX file (Excel file), other data inputs
- Part 3: writing CSV file
- Part 4: reading and saving R objects

Data We Use

- Everything we do in class will be using real publicly available data there are few 'toy' example datasets and 'simulated' data
- Baltimore Open Data and Data.gov will be sources for the first few days
- We have also added functionality to load these datasets directly in the jhur package

Data Input

- · 'Reading in' data is the first step of any real project/analysis
- R can read almost any file format, especially via add-on packages
- · We are going to focus on simple delimited files first
 - comma separated (e.g. '.csv')
 - tab delimited (e.g. '.txt')
 - Microsoft Excel (e.g. '.xlsx')

Data Input

Youth Tobacco Survey (YTS) dataset:

"The YTS was developed to provide states with comprehensive data on both middle school and high school students regarding tobacco use, exposure to environmental tobacco smoke, smoking cessation, school curriculum, minors' ability to purchase or otherwise obtain tobacco products, knowledge and attitudes about tobacco, and familiarity with pro-tobacco and anti-tobacco media messages."

Check out the data at: https://catalog.data.gov/dataset/youth-tobacco-survey-yts-data

Data Input: Dataset Location

Dataset is located at http://johnmuschelli.com/intro_to_r/data/Youth_Tobacco_Survey_YTS_Data.csv

- Download data by clicking the above link
 - Safari if a file loads in your browser, choose File -> Save As, select,
 Format "Page Source" and save

```
# load library `readr` that contains function `read csv`
library (readr)
dat = read csv("http://johnmuschelli.com/intro to r/data/Youth Tobacco Survey
# `head` displays first few rows of a data frame
head (dat, 5)
# A tibble: 5 x 31
  YEAR LocationAbbr LocationDesc TopicType TopicDesc MeasureDesc DataSour
 <dbl> <chr> <chr>
                                <chr> <chr> <chr>
                                                                   <chr>
 2015 AZ Arizona
                                Tobacco U... Cessation... Percent of C... YTS
2 2015 AZ Arizona Tobacco U... Cessation... Percent of C... YTS
  2015 AZ Arizona Tobacco U... Cessation... Percent of C... YTS
  2015 AZ Arizona Tobacco U... Cessation... Quit Attempt... YTS
  2015 AZ
                   Arizona
                                Tobacco U... Cessation... Quit Attempt... YTS
# ... with 24 more variables: Response <chr>, Data Value Unit <chr>,
   Data Value Type <chr>, Data Value <dbl>, Data Value Footnote Symbol <chr>,
   Data Value Footnote <chr>, Data Value Std Err <dbl>,
   Low Confidence Limit <dbl>, High Confidence Limit <dbl>, Sample Size <dbl>
   Gender <chr>, Race <chr>, Age <chr>, Education <chr>, GeoLocation <chr>,
   TopicTypeId <chr>, TopicId <chr>, MeasureId <chr>, StratificationID1 <chr>
#
   StratificationID2 <chr>, StratificationID3 <chr>, StratificationID4 <chr>,
   SubMeasureID <chr>, DisplayOrder <dbl>
```

So what is going on "behind the scenes"?

read_csv() parses a "flat" text file (.csv) and turns it into a **tibble** – a rectangular data frame, where data are split into rows and columns

- First, a flat file is parsed into a rectangular matrix of strings
- Second, the type of each column is determined (heuristic-based guess)

read csv() needs path to your file, will return a tibble

```
read_csv(file, col_names = TRUE, col_types = NULL,
  locale = default_locale(), na = c("", "NA"),
  quoted_na = TRUE, quote = "\"", comment = "", trim_ws = TRUE,
  skip = 0, n_max = Inf, guess_max = min(1000, n_max),
  progress = show_progress(), skip_empty_rows = TRUE
)
```

- file is the path to your file, in quotes
- · can be path in your local computer absolute file path or relative file path
- · can be path to a file on a website

```
## Examples

dat = read_csv("/Users/martakaras/Downloads/Youth_Tobacco_Survey_YTS_Data.csv"

dat = read_csv("Youth_Tobacco_Survey_YTS_Data.csv")

dat = read_csv("www.someurl.com/table1.csv")
```

read_csv() is a special case of read_delim() – a general function to read a delimited file into a data frame

read delim() needs path to your file and fileds delimiter, will return a tibble

```
read_delim(file, delim, quote = "\"", escape_backslash = FALSE,
  escape_double = TRUE, col_names = TRUE, col_types = NULL,
  locale = default_locale(), na = c("", "NA"), quoted_na = TRUE,
  comment = "", trim_ws = FALSE, skip = 0,
  n_max = Inf, guess_max = min(1000, n_max),
  progress = show_progress(), skip_empty_rows = TRUE
)
```

- file is the path to your file, in quotes
- delim is what separates the fields within a record

```
## Examples
dat = read_delim("Youth_Tobacco_Survey_YTS_Data.csv", delim = ",")

dat = read_delim("www.someurl.com/table1.txt", delim = "\t")
```

Data Input: Read in Directly From File Path

```
dat = read_csv("../data/Youth_Tobacco_Survey_YTS_Data.csv")

— Column specification

cols(
    .default = col_character(),
    YEAR = col_double(),
    Data_Value = col_double(),
    Data_Value_Std_Err = col_double(),
    Low_Confidence_Limit = col_double(),
    High_Confidence_Limit = col_double(),
    Sample_Size = col_double(),
    DisplayOrder = col_double()
)
i Use `spec()` for the full column specifications.
```

The data is now successfully read into your R workspace. Colum specification of first few columns is printed to the console.

Common new user mistakes we have seen

- 1. Working directory problems: trying to read files that R "can't find"
 - Path misspecification
- 2. Typos (R is case sensitive, x and x are different)
 - RStudio helps with "tab completion"
- 3. Data type problems (is that a string or a number?)
- 4. Open ended quotes, parentheses, and brackets
- 5. Different versions of software

Data Input: Checking for problems

• The spec() and problems() functions show you the specification of how the data was read in.

```
problems (dat)
                       expected actual
[1] row
             col
<0 rows> (or 0-length row.names)
spec (dat)
cols(
  YEAR = col double(),
  LocationAbbr = col character(),
  LocationDesc = col character(),
  TopicType = col character(),
  TopicDesc = col character(),
  MeasureDesc = \overline{\text{col}} character(),
  DataSource = col character(),
  Response = col character(),
  Data Value Uni\overline{t} = col character(),
  Data Value Type = col character(),
  Data Value = col double(),
  Data Value Footnote Symbol = col character(),
  Data Value Footnote = col character(),
  Data Value Std Err = col double(),
  Low Confidence Limit = col double(),
  High Confidence Limit = col double(),
  Sample Size = col double(),
                                                                           13/42
  Gender = col character(),
```

Data Input: Checking for problems

The stop_for_problems() function will stop if your data had any problem when reading in (even if that problem did not cause the data reading to fail).

• Particularly useful to put after the data reading code e.g. in some automated R script that should not proceed in case some data "weirdness" occurred.

stop_for_problems(dat)

Help

For any function, you can write ?FUNCTION_NAME, or help("FUNCTION_NAME") to look at the help file:

```
?read_delim
help("read_delim")
```

Data Input: Read in From RStudio Toolbar

R Studio features some nice "drop-down" support, where you can run some tasks by selecting them from the toolbar.

For example, you can easily import text datasets using the File --> Import Dataset --> From Text (readr) command. Selecting this will bring up a new screen that lets you specify the formatting of your text file.

After importing a datatset, you get (printed in the R console) the corresponding R command that you can enter in the console if you want to re-import data.

Data Input: base R

There are also data importing functions provided in base R (rather than the readr package), like read.delim() and read.csv().

These functions have slightly different syntax for reading in data (e.g. header argument).

However, while many online resources use the base R tools, the latest version of RStudio switched to use these new readr data import tools, so we will use them in the class for slides. They are also up to two times faster for reading in large datasets, and have a progress bar which is nice.

But you can use whatever function you feel more comfortable with.

Lab Part 1

Lab

Website

Data Input: tbl_df (a tibble)

The read_csv(), read_delim() and related functions from readr all return tbl_df object (colloquially: a tibble) that is a data.frame object but with improved printing and subsetting properties.

We also get tol df (tibble) when using drop-down menu in RStudio.

Alternatives:

- read.csv() "base" R function, still largely used; returns data.frame object
- fread() from data.table R package; returns data.table object

Data Input: tbl_df (a tibble)

```
head (dat, 3)
# A tibble: 3 x 31
  YEAR LocationAbbr LocationDesc TopicType TopicDesc MeasureDesc DataSour
 Arizona Tobacco Us... Cessatio... Percent of C... YTS
1 2015 AZ
2 2015 AZ
               Arizona Tobacco Us... Cessatio... Percent of C... YTS
  2015 AZ
                  Arizona Tobacco Us... Cessatio... Percent of C... YTS
 ... with 24 more variables: Response <chr>, Data Value Unit <chr>,
   Data Value Type <chr>, Data Value <dbl>, Data Value Footnote Symbol <chr>,
   Data Value Footnote <chr>, Data Value Std Err <dbl>,
   Low Confidence Limit <dbl>, High Confidence Limit <dbl>, Sample Size <dbl>
   Gender <chr>, Race <chr>, Age <chr>, Education <chr>, GeoLocation <chr>,
   TopicTypeId <chr>, TopicId <chr>, MeasureId <chr>, StratificationID1 <chr>
   StratificationID2 <chr>, StratificationID3 <chr>, StratificationID4 <chr>,
   SubMeasureID <chr>, DisplayOrder <dbl>
class (dat)
[1] "spec tbl df" "tbl df"
                             "tbl"
                                          "data.frame"
```

Data Input

- nrow() displays the number of rows of a data frame
- ncol() displays the number of columns
- dim() displays a vector of length 2: # rows, # columns
- colnames() displays the column names (if any) and rownames() displays the row names (if any)

```
dim(dat)
[1] 9794 31
nrow(dat)
[1] 9794
ncol(dat)
[1] 31
```

Data Input

- nrow() displays the number of rows of a data frame
- ncol () displays the number of columns
- dim() displays a vector of length 2: # rows, # columns
- colnames() displays the column names (if any) and rownames() displays the row names (if any)

colnames (dat)

```
"LocationAbbr"
 [1] "YEAR"
 [3] "LocationDesc"
                                   "TopicType"
                                   "MeasureDesc"
 [5] "TopicDesc"
 [7] "DataSource"
                                   "Response"
 [9] "Data Value Unit"
                                   "Data Value Type"
                                   "Data Value Footnote Symbol"
[11] "Data Value"
                                   "Data Value Std Err"
[13] "Data Value Footnote"
[15] "Low Confidence Limit"
                                   "High Confidence Limit"
[17] "Sample Size"
                                   "Gender"
[19] "Race"
                                   "Age"
[21] "Education"
                                   "GeoLocation"
[23] "TopicTypeId"
                                   "TopicId"
                                   "StratificationID1"
[25] "MeasureId"
[27] "StratificationID2"
                                   "StratificationID3"
[29] "StratificationID4"
                                   "SubMeasureID"
[31] "DisplayOrder"
```

Working Directories

- R "looks" for files on your computer relative to the working directory
- Many people recommend not setting a working directory in the scripts
 - If you open an R file with a new RStudio session, it assumes the working directory is where the script is in
- If you do set a working directory, do it at the beginning of your script
- Example of getting and setting the working directory:

```
# get the working directory
getwd()
setwd("~/Lectures")
```

Setting a Working Directory

- Setting the directory can sometimes be finicky
 - **Windows**: Default directory structure involves single backslashes ("\"), but R interprets these as "escape" characters. So you must replace the backslash with forward slashes ("/") or two backslashes ("\\")
 - Mac/Linux: Default is forward slashes, so you are okay
- Typical directory structure syntax applies
 - ".." goes up one level
 - "./" is the current directory
 - "~" is your "home" directory

Working Directory

Note that the dir() function interfaces with your operating system and can show you which files are in your current working directory.

You can try some directory navigation:

```
dir("./") # shows directory contents
 [1] "Data IO.html"
                                   "Data IO.pdf"
 [3] "Data IO.R"
                                   "index.html"
 [5] "index.pdf"
                                   "index.R"
 [7] "index.Rmd"
                                   "lab"
 [9] "makefile"
                                   "YouthTobacco newNames.csv"
[11] "yts dataset.rds"
dir("...")
 [1] "all functions.xlsx"
 [2] "all_the functions.csv"
 [3] "all the packages.txt"
 [4] "Arrays Split"
 [5] "Basic \overline{R}"
 [6] "Best Model Coefficients.csv"
 [7] "Best Model Coefficients.xlsx"
 [8] "bibliography.bib"
 [9] "black and white theme.pdf"
[10] "bloomberg.logo.small.horizontal.blue.png"
[11] "data"
                                                                           25/42
    "Data Classes"
```

Relative vs. absolute paths (From Wiki)

An **absolute or full path** points to the same location in a file system, regardless of the current working directory. To do that, it must include the root directory.

This means if I try your code, and you use absolute paths, it won't work unless we have the exact same folder structure where R is looking (bad).

By contrast, a **relative path starts from some given working directory**, avoiding the need to provide the full absolute path. A filename can be considered as a relative path based at the current working directory.

Setting the Working Directory From RStudio Toolbar

In RStudio, go to Session -> Set Working Directory -> To Source File
Location

RStudio should put code in the Console, similar to this:

```
setwd("~/Lectures/Data_IO/lecture")
```

Again, if you open an R file with a new RStudio session, RStudio sets working directory to the file's location for you.

Setting the Working Directory

You may need to make sure that RStudio is the default application to open .R files

Mac - right click -> Get Info --> Open With: RStudio --> Change All

Data Input - Excel

Many data analysts collaborate with researchers who use Excel to enter and curate their data. Often times, this is the input data for an analysis. You therefore have two options for getting this data into R:

- Saving the Excel sheet as a .csv file, and using read_csv()
- Using an add-on package, like xlsx, readxl, or openxlsx

For single worksheet .xlsx files, I often just save the spreadsheet as a .csv file (because I often have to strip off additional summary data from the columns)

For an .xlsx file with multiple well-formated worksheets, I use the readx1 package for reading in the data.

Data Input - Other Software

- readr package has read_csv()/write_csv() and read_table() functions similar to read.csv()/write.csv() and read.table() from base R; has different defaults, but can read much faster for very large data sets
- readxl package the read_excel() function can read Excel sheets easily
- haven package reads in SAS, SPSS, Stata formats
- sas7bdat reads .sas7bdat files (SAS files)
- foreign package can read all the formats as haven. Around longer (aka more testing), but not as maintained (bad for future)

Some of these are now available in the RStudio drop-down list.

Lab Part 2

Lab

Website

While its nice to be able to read in a variety of data formats, it's equally important to be able to output data somewhere.

There are also data exporting functions in the readr package, which have the pattern write * like write csv() and write delim()

```
write_csv(x, file,
  na = "NA", append = FALSE,
  col_names = !append, quote_escape = "double",
  eol = "\n", path = deprecated()
)

write_delim(x, file, delim = " ",
  na = "NA", append = FALSE,
  col_names = !append, quote_escape = "double",
  eol = "\n", path = deprecated()
)
```

x: data frame or tibble you want to write

file: the file name where you want to R object written

• It can be an absolute path, a relative path (relative to your working directory), or a filename (which writes the file to your working directory)

delim: what character separates the columns?

- · ", " = comma delimted
- . "\t" = tab delimited

There other functions in base R to do the writing, like write.table() and write.csv()

- Some of their arguments may be named differently than in write_csv() look up teir documentation (e.g. run ?write.table)
- Note base R writing functions by default do write out row names, which you
 can change by setting e.g. row.names = TRUE. I do this a lot since I often
 email these to collaborators who open them in Excel

For example, we can write back out the Youth Tobacco dataset with the new column name:

```
# load `dplyr` package that has `rename` function
library(dplyr)
dat = rename(dat, Location_Abbr = LocationAbbr)
write_csv(dat, path = "YouthTobacco_newNames.csv")
```

Lab Part 3

Lab

Website

Write a single R object to a file

If you want to save **one** R object, you can use write_rds() from readr package to save an R object (e.g., a tibble) to an R binary file with rds extension

```
write_rds(dat, file = "yts_dataset.rds")
```

rds is R's native format for storing single objects

Read a single R object from a file

To read this back in to R, you need to use read_rds(). Note you need to assign it:

```
dat2 = read_rds(file = "yts_dataset.rds")
identical(dat, dat2) # test if they are the same
```

[1] TRUE

Lab Part 4

Lab

Website

Write multiple R objects to a file

The save command can save a set of R objects into an "R data file", with the extension .rda or .RData.

Load multiple R objects from a file

The opposite of save is load.

The ls() command lists the items in the workspace/environment and rm() removes them:

```
ls() # list things in the workspace

[1] "dat" "dat2" "x" "yts"

rm(list = c("x", "yts"))
ls()

[1] "dat" "dat2"

z = load("yts_data.rda")
ls()

[1] "dat" "dat2" "x" "yts" "z"
```

Load multiple R objects from a file

```
z = load("yts_data.rda")
print(z)

[1] "yts" "x"
```

Note, z is a **character vector** of the **names** of the objects loaded, **not** the objects themselves.