### Workshop: Basic R and More, Part 3 of 3

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- The basic plotting commands in R are effective but the graphics are rudimentary and not easily extensible.
- ggplot2 is a state-of-the-art data visualization package in R (a member of the tidyverse).
- On balance, ggplot2 is every bit the equal of Tableau, Tableau Public, Data Studio, Power BI, etc., for data visualization. All have their pros and cons.
- The package does not fully anticipate what the user wants to do, but rather provides the mechanisms for combining different graphical components which the user chooses.

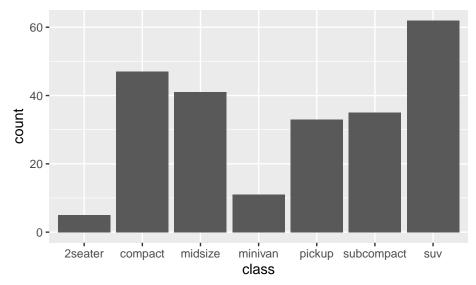
- The Grammar of Graphics (Wilkinson, 1999) provided a theoretical framework for the construction of graphics.
- ggplot2 is based upon a layered grammar of graphics, an extension of Wilkinson's work. This "grammar" gives a framework which enables one to easily update a graphic by modifying a single feature at a time, and suggests aspects of a graph that can be built up component by component.
  - <https://ggplot2.tidyverse.org/reference>
- Along with dplyr, ggplot2 has grown in use to become one of the most popular R packages. Companies now view the ability to work with these packages as a valuable technological skill.

- Let's create a *bar chart*. Here we consider a data set that gives the fuel efficiency of different classes of vehicles in two different years 1999 and 2008. This is a subset of data that the EPA makes publicly available and is included in the ggplot2 package as mpg.
- > data(mpg) ## Loads data set
- > ?mpg ## Read about the data set
  - There are three ways to determine what the columns are. First, we could submit str(mpg), which describes the 'structure' of an R object; in this case, our data frame. Second, we could submit glimpse(mpg), which is like str(), but tries to show you as much data as possible. Be aware that glimpse() requires the use of the tibble package. Third, we could use the Environment tab

```
> plot01 <- ggplot(mpg, aes(x = class))
> plot01 <- plot01 + geom_bar()</pre>
```

- aes stands for aesthetic, a visual property of the graph. It says the x-axis will be the car's class, which is indicated by the column named class.
- geom\_bar() creates a bar chart. geom stands for geometric object.
   They represent different ways to display variables and the relationship between them.
- Please note that the plus sign + serves as a pipe operator for ggplot2.
- When I create graphs using ggplot2, I prefer to code sequentially and save the graph.

> plot01



 Suppose we were interested in the distribution of highway miles per gallon (hwy).

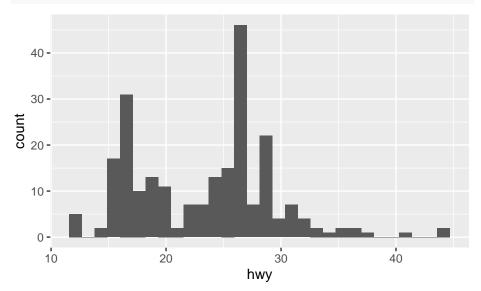
```
> plot02 <- ggplot(mpg, aes(x = hwy))
> plot02 <- plot02 + geom_histogram()</pre>
```

- geom histogram() creates a frequency histogram.
- You will get an ominous message about the default of 30 bins, which gives us too fine a partition of the data. To get a sensible number of classes for a histogram, use the Freedman-Diaconis rule:

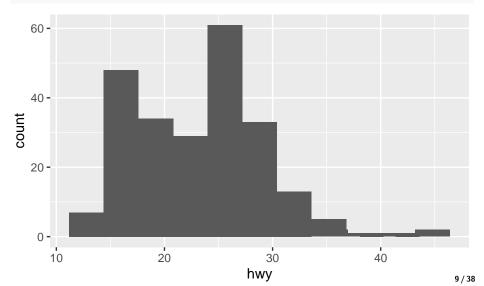
```
> nclass.FD(mpg$hwy)
```

[1] 11

> plot02



- > plot02 <- plot02 + geom\_histogram(bins = 11)</pre>
- > plot02

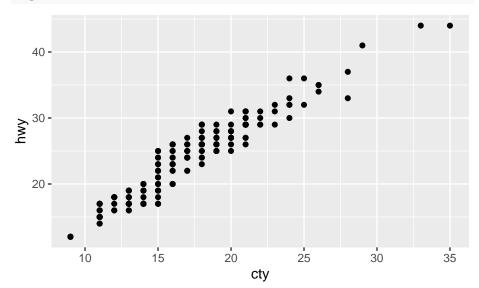


• Suppose we were also interested in the relationship between city miles per gallon (cty) and hwy.

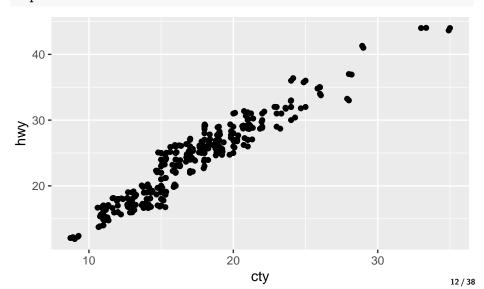
```
> plot03 <- ggplot(mpg, aes(x = cty, y = hwy))
> plot03 <- plot03 + geom_point()</pre>
```

- geom\_point() draws points for a scatter plot.
- There will be a minor overplotting problem. Counterintuitively, adding random noise to a plot can sometimes make it easier to read. Jittering is particularly useful for small data sets that are discretized. We will demonstrate jittering momentarily.

> plot03



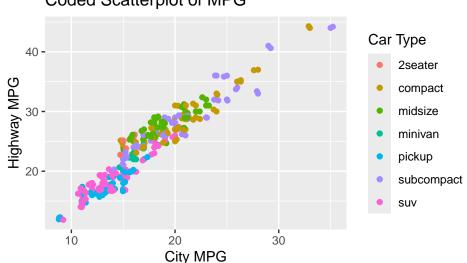
- > plot03 <- plot03 + geom\_jitter()</pre>
- > plot03



 We can also make a coded scatter plot. For example, suppose we were interested in the relationship between cty and hwy by class.

> plot03

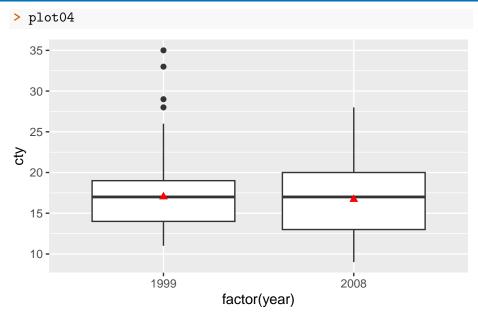




• Lastly, we can also make side-by-side box plots. For example, suppose we were interested in the relationship between cty by year, where year is the year of manufacture ({1998, 2008}).

- factor() tells R to treat year as a category, not an integer.
- geom\_boxplot() creates box plots.

- ggplot2 is a powerful, extensible way to obtain even the most complicated (but beautiful!) graphs in order to visualize your data.
- For example, let's suppose you work for the EPA and are interested in the mean city miles per gallon for each year. You then want to superimpose these values directly on each box plot. This is straightforward to do in ggplot2.



- Character strings can be an important class of data. Data being read into R often come in the form of character strings where different parts might mean different things.
  - For example a sample ID of "OBGYN06150902" might be a so-called "Patient Code" where OBGYN represents a department acronym, 061509 is the date in MMDDYY format, and 02 represents some designation for a location, doctor, etc.
- We need to have a set of utilities that allow us to split, combine, and examine character strings in a easy and consistent fashion. These will come in handy when managing and wrangling data and in more sophisticated use (e.g., text analytics).
- Here we will introduce a few of the more commonly used functions from the stringr package from the tidyverse.

 The str\_c() allows us to concatenate two strings or two vectors of strings. This is demonstrated through the following example.

```
> first_names <- c("Michael", "Sarah", "Mitch", "Jennifer")
> last_names <- c("Long", "Pearson", "Jones", "Smith")
> full_names <- str_c(first_names, last_names)
> cbind(first_names, last_names, full_names)
```

```
first_names last_names full_names
[1,] "Michael" "Long" "MichaelLong"
[2,] "Sarah" "Pearson" "SarahPearson"
[3,] "Mitch" "Jones" "MitchJones"
[4,] "Jennifer" "Smith" "JenniferSmith"
```

• The str\_length() function calculates the length of each string in the vector of strings passed to it. This could be important in text analytics.

```
> str_length(full_names)
```

#### [1] 11 12 10 13

- A powerful string operation is to take a string and match some pattern within it. For example, we might be interested in mining through full.names to detect if a pattern occurs.
- > str\_detect(full\_names, pattern = fixed("Mi"))
  - [1] TRUE FALSE TRUE FALSE

• To figure out positions where the first Mi patterns are in each string, we can use the str\_locate() function.

- One of my favorite stringr functions is str\_count(). It counts the number of designated matches in a string (e.g., the number of vowels/full name). Note that '|' is a logical operator that stands for 'or'.
- > str\_count(full\_names, "a|e|i|o|u|A|E|I|O|U")

```
[1] 4 5 3 4
```

[4,] NA NA

- Using dates in any software package can be surprisingly difficult.
- They require some special organization because there are several conventions for how to write them and because the sort order should be in the order that the dates occur in time.
- We are going to use the package lubridate (which belongs to the tidyverse) to work with dates and times. lubridate gets installed as part of the tidyverse installation. However, because it's so specialized, it does not belong to the *core* tidyverse, so you have to load it explicitly with library(lubridate).

- To create a Date object, we need to take a string or number that represents a date and give R a consistent order for the date so that it can uniquely identify which piece is the year, the month, and the day.
- We demonstrate the use of the mdy() function. Many other functions exist depending on how the date is specified.
- > library(lubridate)
  > mdy("April 16, 2019", "4-16-19", "4-16-2019", "4/16/19")

```
[1] "2019-04-16" "2019-04-16" "2019-04-16" "2019-04-16"
```

• Suppose we also had the hours and minutes. Then we can add this time to the date if we wanted to using the mdy\_hm() function.

```
> mdy_hm("April 16, 2019 1:00 PM", "4-16-2019 13:00")
```

```
[1] "2019-04-16 13:00:00 UTC" "2019-04-16 13:00:00 UTC"
```

• UTC stands for Coordinated Universal Time. However, there are options for specifying a different time zone (e.g., EST, EDT, etc.). See OlsonNames() for your personal favorite. Notice the tz option.

```
> mdy_hm("4-16-2019 13:00", tz = "US/Eastern")
```

```
[1] "2019-04-16 13:00:00 EDT"
```

 The lubridate package provides many functions for extracting information from the date and time. We demonstrate just a few of them.

```
> x <- mdy_hm("4-16-2019 13:00", tz = "US/Eastern")
> year(x) ## Return year
```

```
[1] 2019
```

```
> hour(x) ## Return hour of the day
```

```
[1] 13
```

```
> wday(x) ## Return day of the week (Sunday = 1)
```

```
[1] 3
```

```
> yday(x) ## Return day of the year
```

```
[1] 106
```

- Once we have two or more Date objects defined, we can perform appropriate mathematical operations. For example, we might want to the know the number of days there are between two dates, aka 'tenure time'.
- For example, for some patient, suppose s/he was admitted into the hospital on 2019-6-23 and discharged on 2019-9-15.

```
> Admission <- ymd("2019-6-23") ## "time when"
> Discharge <- ymd("2019-9-15") ## "time when"
> LOS <- Discharge - Admission
> LOS ## "time duration"
```

Time difference of 84 days

- Let's pause for a moment. This is a great demonstration of the power of R as a tool in data science. Why?
- We have demonstrated proof of concept for one patient. But what if a
  hospital had such data for thousands of people? Scaling up to get
  everyone's time difference would present no challenge at all to R. The
  creation of a 'time difference' variable is sometimes called feature
  extraction among machine learning modelers.
- This feature could then be used in a machine learning model to help us understand and predict some behavior that might be of interest to the hospital. Of course, the modeling would be done in R in a seamless workflow.

- I always work with script (.R) files instead of typing directly into the
  console. If you have a data analysis of any substance, it is virtually
  impossible to write all your R code correctly the first time and
  remember all that you did later on if the need arises.
- Writing a script file fully documents how you did your analysis. Hence, having a script makes it easy to re-run an analysis after a change in the data (additional data values, transformed data, or removal of outliers) or if a journal referee wants revisions.
- Let's see an example. Open your browser, go to <a href="https://github.com/philturk/R\_Wshop">https://github.com/philturk/R\_Wshop</a>> and click on Ex01.R. Keep this window open.

- It often makes your script more readable if you break a single command up into multiple lines. R will disregard all whitespace (including line breaks) so you can safely spread your command over multiple lines.
- It's also useful to leave comments in the script for things such as
  explaining a tricky step, who wrote the code and when, or why you
  chose a particular name for a variable. So 'comment' your R code and
  be kind to your future self!

- One way to create a new .R script in RStudio is to go to File, New File, R Script. This opens a blank script in the Source window where you can type commands and functions.
- Copy-and-paste or type the R code from Ex01.R in the Source window and then you can execute the code as follows:
  - $\bullet$  Click the line of code you want to run, and then press Cmd + Return or click the Run button (Windows users, try Ctrl + Enter).
  - Highlight the block of code you want to run, and then again press Cmd
     Return or click the Run button.
  - Run the entire script by clicking anywhere in the source editor and then
    pressing Cmd + Shift + Return (Windows users, try Ctrl + Alt + R)

- Working with R scripts is an acceptable way of documenting what you
  did, but the script file itself doesn't contain the actual results of
  commands that were run, nor does it show you the plots.
- Also, anytime I want to comment on some output, it needs to be offset with the commenting character #.
- It would be nice to have all the commands, the results, and comments merged into one document. This is what an R Markdown file does for us. Alas, due to time, this is where our journey ends:(
- However, you now have enough knowledge under your belt to check out R Markdown on your own at <a href="https://rmarkdown.rstudio.com">https://rmarkdown.rstudio.com</a>>.
- A newer open-source scientific and technical publishing system called Quarto is gaining traction (<a href="https://quarto.org">https://quarto.org</a>).

# **Appendix**

Optional material on basic programming follows

- It is often necessary to write R code that can perform different actions depending on the data or to automate a task that must be repeated many times. We now cover just a few of these tools.
- Here is an 'if' statement. (The 'else' part is sometimes optional.)

```
> flip <- rbinom(n = 1, size = 1, prob = 0.5)
> flip ## Flip a coin, and get a 0 or 1

[1] 0
> if(flip == 0){ ## Convert the 0/1 to Tail/Head
+ flip <- "Tail"
+ }else{ ## Keep else on same line as closing } for if
+ flip <- "Head"
+ }
> flip
```

 In my experience, the ifelse() function is very useful because it operates on vectors.

```
> x <- 1:5
> x
```

```
[1] 1 2 3 4 5
```

```
> ifelse(x <= 2, "Small", "Large")</pre>
```

```
[1] "Small" "Small" "Large" "Large" "Large"
```

 Be aware that there is also an if\_else() function in dplyr that is faster (but a bit more temperamental).

 A while loop will repeat a section of R code over and over until a stopping condition is met.

```
> x <- pi ## Initialize x
> while(x < 50){ ## Multiple lines of R code
+ x <- 2*x ## Important to update x
+ print(x)
+ }</pre>
```

- [1] 6.283185
- [1] 12.56637
- [1] 25.13274
- [1] 50.26548

 Aside: When "programming" at the command line for this workshop, do not close with a } brace until you are done typing. (We won't program this way in practice.)

 When we know in advance how many times we should go through a loop of R code, a 'for' loop can be quite useful.

```
> stuff <- NULL ## Create an empty object
> for(i in 1:4){ ## Append elements to stuff
+ stuff[i] <- i*i
+ print(stuff)
+ }</pre>
```

```
[1] 1
[1] 1 4
[1] 1 4 9
[1] 1 4 9 16
```

• In the for loop, 'i' can be called anything. It's a variable that you choose that is a placeholder.

- Sometimes it can be useful to define your own functions. A rigorous treatment of R programming would require its own short course :)
- Let's start by defining a function F\_to\_C that converts temperatures from Fahrenheit to Celsius.

```
> F_to_C <- function(temp_F){ ## temp_F is placeholder
+ temp_C <- ((temp_F - 32) * (5 / 9)) ## Do conversion
+ return(temp_C) ## Show result temp_C
+ }</pre>
```

> F\_to\_C(32) ## Check it!

[1] -17.77778 0.00000 20.00000

• Calling our own function is no different from calling any other function already in R. We can do it for one temperature ( $^{\circ}F$ ) or a vector of temperatures ( $^{\circ}F$ ).

```
[1] 0
> lots_temps <- c(0, 32, 68)
> F_to_C(lots_temps)

[1] -17.77778   0.00000   20.00000
> my_Cs <- F_to_C(lots_temps)
> my_Cs ## Can assign function output to object
```