# PRACTICAL DATA ANALYSIS FOR POLITICAL SCIENTISTS

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# About This Book

This book contains the course notes for Brenton Kenkel's course Statistics for Political Research II (PSCI 8357 at Vanderbilt University). It covers the basics of statistical modeling and programming with linear models, along with applications in R.

This book is written in R Markdown and published via Bookdown on GitHub Pages. You can find the R Markdown source files at https://github.com/brentonk/pdaps.

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# Principles of Programming

It may seem strange to begin a statistics class with two weeks on programming. It is strange. Here is why I have made this strange choice.

First, as a working social scientist, most of the time you spend on data analysis won't be on the *analysis* part. It'll be on obtaining and cleaning the data, to get it in a form that makes sense to analyze. Good programming skills will let you spend less time cleaning data and more time publishing papers.

Second, even if you don't want to develop good programming habits, journals are going to force you to. Every reputable political science journal requires that you provide replication scripts, and some of the best (e.g., American Journal of Political Science) have begun auditing the replication materials as a condition of publication. Better to learn The Right Way now when you have lots of time than to be forced to when you're writing a dissertation or on the market or teaching your own courses.

Third, while I feel embarrassed to invoke the cliché that is Big Data, that doesn't mean it's not a real thing. Political scientists have access to more data and more computing power than ever before. You can't collect, manage, clean, and analyze large quantities of data without understanding the basic principles of programming.

As Bowers (2011) puts it, "Data analysis is computer programming." By getting a PhD in political science, by necessity you're going to become a computer programmer. The choice before you is whether to be a good one or a bad one.

Wilson et al. (2014) list eight "best practices for scientific computing." The first two encapsulate most of what you need to know:

- 1. Write programs for people, not computers.
- 2. Let the computer do the work.

 $<sup>^{1}</sup>$  Or whatever other social science field.

#### 2.1 Write Programs for People, Not Computers

The first two words here—write programs—are crucial. When you are doing analysis for a research project, you should be writing and running scripts, not typing commands into the R (or Stata) console. The console is ephemeral, but scripts are forever, at least if you save them.

Like the manuscripts you will write to describe your findings, your analysis scripts are a form of scientific communication. You wouldn't write a paper that is disorganized, riddled with grammatical errors, or incomprehensible to anyone besides yourself. Don't write your analysis scripts that way either.

Each script should be self-contained, ideally accomplishing one major task. Using an omnibus script that runs every bit of analysis is like writing a paper without paragraph breaks. A typical breakdown of scripts for a project of mine looks like:

- 0-download.r: downloads the data
- 1-clean.r: cleans the data
- 2-run.r: runs the main analysis
- 3-figs.r: generates figures

The exact structure varies depending on the nature of the project. Notice that the scripts are numbered in the order they should be run.

Within each script, write the code to make it as easy as possible for your reader to follow what you're doing. You should indent your code according to style conventions such as http://adv-r.had.co. nz/Style.html. Even better, use the Code -> Reindent Lines menu option in R Studio to automatically indent according to a sane style.

```
my results <- c(mean(variable),
quantile(variable,
probs = 0.25),
max(variable))
# Better
my_results <- c(mean(variable),</pre>
                 quantile(variable,
```

max(variable))

Another way to make your code readable—one that, unfortunately, cannot be accomplished quite so algorithmically—is to add explanatory comments. The point of comments is not to document how the

probs = 0.25),

language works. The following comment is an extreme example of a useless comment.

```
# Take the square root of the errors and assign them to
# the output variable
output <- sqrt(errors)</pre>
```

A better use for the comment would be to explain why you're taking the square root of the errors, at least if your purpose in doing so would be unclear to a hypothetical reader of the code.

My basic heuristic for code readability is If I got hit by a bus tomorrow, could one of my coauthors figure out what the hell I was doing and finish the paper?

#### 2.2 Let the Computer Do the Work

Computers are really good at structured, repetitive tasks. If you ever find yourself entering the same thing into the computer over and over again, you are Doing It Wrong. Your job as the human directing the computer is to figure out the structure that underlies the repeated task and to program the computer to do the repetition.

For example, imagine you have just run a large experiment and you want to estimate effects by subgroups.<sup>2</sup> Your respondents differ across four variables—party ID (R or D), gender (male or female), race (white or nonwhite), and education (college degree or not)—giving you 16 subgroups. You could copy and paste your code to estimate the treatment effect 16 times. But this is a bad idea for a few reasons.

- Copy-paste doesn't scale. Copy-paste is managable (albeit misguided) for 16 iterations, but probably not for 50 and definitely not for more than 100.
- Making changes becomes painful. Suppose you decide to change how you calculate the estimate. Now you have to go back and individually edit 16 chunks of code.
- Copy-paste is error-prone, and insidiously so. If you do the calculation wrong all 16 times, you'll probably notice. But what if you screwed up for just one or two cases? Are you really going to go through and check that you did everything right in each individual case?

We're going to look at the most basic ways to get the computer to repeat structured tasks—functions and control flow statements. To illustrate these, we will use a result you discussed in Stat I: the central limit theorem.

<sup>&</sup>lt;sup>2</sup> There could be statistical problems with this kind of analysis, at least if the subgroups were specified post hoc. See https://xkcd.com/882/ ("Significant"). We're going to leave this issue aside for now, but we'll return to it later when we discuss the statistical crisis in science.

The central limit theorem concerns the sampling distribution of the sample mean,

$$\bar{X} = \frac{1}{N} \sum_{n=1}^{N} X_n,$$

where each  $X_n$  is independent and identically distributed with mean  $\mu$  and variance  $\sigma^2$ . Loosely speaking, the CLT says that as N grows large, the sampling distribution of  $\bar{X}$  becomes approximately normal with mean  $\mu$  and variance  $\sigma^2/N$ .

Here's what we would need to do to see the CLT in practice. We'd want to take a bunch of samples, each of size N, and calculate the sample mean of each. Then we'd have a sample of sample means, and we could check to verify that they are approximately normally distributed with mean  $\mu$  and variance  $\sigma^2/N$ . This is a structured, repetitive task—exactly the kind of thing that should be programmed. We'll try it out with a random variable from a Poisson distribution with  $\lambda = 3$ , which has mean  $\mu = 3$  and variance  $\sigma^2 = 3$ .

First things first. We can use the rpois function to draw a random sample of N numbers from the Poisson distribution.

```
samp <- rpois(10, lambda = 3)</pre>
samp
   [1] 2 3 8 3 5 4 3 4 2 2
```

To calculate the sample mean, we simply use the mean function.

```
mean(samp)
```

```
## [1] 3.6
```

We are interested in the distribution of the sample mean across many samples like this one. To begin, we will write a function that automates our core task—drawing a sample of N observations from Poisson(3) and calculating the sample mean. A function consists of a set of arguments (the inputs) and a body of code specifying which calculations to perform on the inputs to produce the output.

```
pois mean <- function(n obs) {</pre>
    samp <- rpois(n_obs, lambda = 3)</pre>
    ans <- mean(samp)</pre>
    return(ans)
}
```

This code creates a function called pois\_mean. It has a single argument, called n obs. It generates a random sample of n obs draws from Poisson(3) and calculates its sample mean. It then returns the sample mean as the output.

Let's try calling this function a few times, each with a sample size of N=30. Your output will differ slightly from what's printed here, since the function is generating random numbers.

```
pois_mean(n_obs = 30)
## [1] 3.0333
pois_mean(n_obs = 30)
## [1] 2.4667
pois mean(n obs = 30)
## [1] 2.9667
```

Remember that what we're interested in is the sampling distribution of the sample mean—the distribution of the sample mean across every possible sample of N observations. We can approximate this distribution by running pois\_mean many times (e.g., 1000 or more). This would be infeasible via copy-paste. Instead, we will use a **for loop**.

```
# Set up a vector to store the output
n_replicates <- 1000
sampling_dist <- rep(NA, n_replicates)</pre>
for (i in 1:n_replicates) {
    sampling_dist[i] <- pois_mean(n_obs = 30)</pre>
}
```

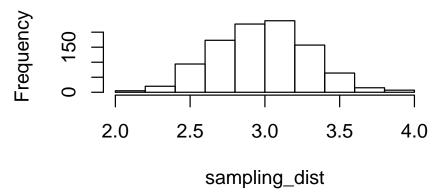
Here's how the for loop works. We specified i as the name of the index variable, with values 1:n\_replicates. The for loop takes each value in the sequence, assigns it to the variable i, runs the given expression (in this case, assigning the output of pois\_mean to the i'th element of sampling\_dist), and then moves on to the next value in sequence, until it reaches the end.

Let's take a look at the results and compare them to our expectations.

```
mean(sampling_dist) # Expect 3
## [1] 2.9952
var(sampling_dist) # Expect 1/10
## [1] 0.096944
```

hist(sampling\_dist) # Expect roughly normal

# Histogram of sampling\_dist



For loops are fun, but don't overuse them. Many simple operations are vectorized and don't require a loop. For example, suppose you want to take the square of a sequence of numbers. You could use a for loop ...

```
input <-c(1, 3, 7, 29)
output <- rep(NA, length(input))</pre>
for (i in 1:length(input)) {
    output[i] <- input[i]^2</pre>
}
output
## [1]
              9
                49 841
```

... but it's faster (in terms of computational speed) and easier to just take advantage of vectorization:

```
input^2
## [1]
                 49 841
          1
```

Another useful piece of control flow is if/else statements. These check a logical condition—an expression whose value is TRUE or FALSE—and run different code depending on the value of the expression. (You may want to catch up on the comparison operators: ==, >, >=, <, <=, etc.)

Let's edit the pois\_mean function to allow us to calculate the median instead of the mean. We'll add a second argument to the function, and implement the option using an if/else statement.

```
pois_mean <- function(n_obs, use_median = FALSE) {</pre>
    samp <- rpois(n_obs, lambda = 3)</pre>
    if (use_median) {
        ans <- median(samp)
    } else {
        ans <- mean(samp)
    }
    return(ans)
}
```

A couple of things to notice about the structure of the function. We use a comma to separate multiple function arguments. Also, we've specified FALSE as the default for the use\_median argument. If we call the function without explicitly specifying a value for use\_median, the function sets it to FALSE.

```
pois_mean(n_obs = 9)
## [1] 3.7778
pois_mean(n_obs = 9, use_median = TRUE)
## [1] 2
pois_mean(n_obs = 9, use_median = FALSE)
## [1] 2.6667
```

There is a vectorized version of if/else statements called, naturally, the ifelse function. This function takes three arguments, each a vector of the same length: (1) a logical condition, (2) an output value if the condition is TRUE, (3) an output value if the condition is FALSE.

```
x <- 1:10
big_x <- x * 100
small_x <- x * -100
ifelse(x > 5, big_x, small_x)
   [1] -100 -200 -300 -400 -500 600 700
   [9]
        900 1000
```

Functions, for loops, and if/else statements are just a few of the useful tools for programming in R.<sup>3</sup> But even these simple tools are enough to allow you to do much more at scale than you could with a copy-paste philosophy.

<sup>&</sup>lt;sup>3</sup> Others include the replicate function, the apply family of functions (sapply, lapply, tapply, mapply, ...), the foreach package, the purrr package, just to name a few of the most useful off the top of my head.

# Working with Data

Some material in this chapter is adapted from notes Matt DiLorenzo wrote for the Spring 2016 session of PSCI 8357.

Let me repeat something I said last week. In your careers as social scientists, starting with your dissertation research—if not earlier—you will probably spend more time collecting, merging, and cleaning data than you will on statistical analysis. So it's worth taking some time to learn how to do this well.

Best practices for data management can be summarized in a single sentence: Record and document everything you do to the data.

The first corollary of this principle is that raw data is sacrosanct. You should never edit raw data "in place". Once you download the raw data file, that file should never change.<sup>1</sup>

In almost any non-trivial analysis, the "final" data—the format you plug into your analysis—will differ significantly from the raw data. It may consist of information merged from multiple sources. The variables may have been transformed, aggregated, or otherwise altered. The unit of observation may even differ from the original source. You must document every one of these changes, so that another researcher working from the exact same raw data will end up with the exact same final data.

The most sensible way to achieve this level of reproducibility is to do all of your data merging and cleaning in a script. In other words, no going into Excel and mucking around manually. Like any other piece of your analysis, your pipeline from raw data to final data should follow the principles of programming that we discussed last week.

Luckily for you,<sup>2</sup> the **tidyverse** suite of R packages (including **dplyr**, **tidyr**, and others) makes it easy to script your "data pipeline". We'll begin by loading the package.

<sup>1</sup> Even if it's data you collected yourself, that data should still have a "canonical" representation that never gets overwritten. See Leek (2015) for more on distributing your own data.

<sup>2</sup> But not for me, because these tools didn't exist when I was a PhD student. Also, get off my lawn!

library("tidyverse")

## 3.1 Loading

The first step in working with data is to acquire some data. Depending on the nature of your research, you will be getting some or all of your data from sources available online. When you download data from online repositories, you should keep track of where you got it from. The best way to do so is—you guessed it—to script your data acquisition.

The R function download.file() is the easiest way to download files from URLs from within R. Just specify where you're getting the file from and where you want it to go. For the examples today, we'll use an "untidied" version of the World Development Indicators data from the World Bank that I've posted to my website.

```
download.file(url = "http://bkenkel.com/data/untidy-data.csv",
   destfile = "my-untidy-data.csv")
```

Once you've got the file stored locally, use the utilities from the readr package (part of tidyverse) to read it into R as a data frame.<sup>3</sup> We have a CSV file, so we will use read\_csv. See help(package = "readr") for other possibilities.

```
untidy_data <- read_csv(file = "my-untidy-data.csv")</pre>
```

```
## Parsed with column specification:
## cols(
     country = col_character(),
##
##
     gdp.2005 = col_double(),
     gdp.2006 = col_double(),
##
##
     gdp.2007 = col_double(),
##
     gdp.2008 = col_double(),
     pop.2005 = col_double(),
##
##
     pop.2006 = col_double(),
     pop.2007 = col_double(),
##
##
     pop.2008 = col_double(),
##
     unemp.2005 = col_double(),
##
     unemp.2006 = col_double(),
##
     unemp.2007 = col_double(),
     unemp.2008 = col_double()
##
## )
```

Remember that each column of a data frame might be a different type, or more formally class, of object. read\_csv and its ilk try to guess the type of data each column contains: character, integer, decimal number ("double" in programming-speak), or something else.

<sup>3</sup> More precisely, the **readr** functions produce output of class "tbl\_df" (pronounced "tibble diff," I'm told), which are like data frames but better. See help(package = "tibble") for what can be done with tbl\_dfs.

The readout above tells you what guesses it made. If it gets something wrong—say, reading a column as numbers that ought to be characters—you can use the col\_types argument to set it straight.

FYI, you could also run read\_csv() directly on a URL, as in:

```
read csv("http://bkenkel.com/data/untidy-data.csv")
```

However, in analyses intended for publication, it's usually preferable to download and save the raw data. What's stored at a URL might change or disappear, and you'll need to have a hard copy of the raw data for replication purposes.

Now let's take a look at the data we've just loaded in.

#### head(untidy\_data)

```
## # A tibble: 6 × 13
##
     country gdp.2005 gdp.2006 gdp.2007
##
       <chr>
                 <dbl>
                          <dbl>
                                    <dbl>
## 1
          AD
               3.8423
                         4.0184
                                  4.0216
## 2
          AE 253.9655 278.9489 287.8318
                       10.3052 11.7212
## 3
          AF
               9.7630
## 4
          AG
               1.1190
                         1.2687
                                  1.3892
               9.2684
                                 10.3483
## 5
          AL
                         9.7718
## 6
          AM
               7.6678
                         8.6797
                                  9.8731
     ... with 9 more variables: gdp.2008 <dbl>,
## #
## #
       pop.2005 <dbl>, pop.2006 <dbl>,
## #
       pop.2007 <dbl>, pop.2008 <dbl>,
## #
       unemp.2005 <dbl>, unemp.2006 <dbl>,
       unemp.2007 <dbl>, unemp.2008 <dbl>
## #
```

We have a country variable giving country abbreviations. The other variables are numerical values: the country's GDP in 2005, 2006, 2007, and 2008; then the same for population and unemployment. Let's get this into a format we could use for analysis.

#### 3.2 *Tidying*

Wickham (2014) outlines three qualities that make data "tidy":

- 1. Each variable forms a column.
- 2. Each observation forms a row.
- 3. Each type of observational unit forms a table.

For one thing, this means that whether a dataset is tidy or not depends—at least in part (some data collections are messy from any angle)—on the purpose it's being analyzed for.

Each row of untidy\_data is a country. In observational studies in comparative politics and international relations, more commonly the unit of observation is the country-year.<sup>4</sup> How can we take untidy\_data and easily make it into country-year data?

We'll use the **tidyr** package (again, part of **tidyverse**) to clean up this data. The biggest problem right now is that each column, besides the country identifier, really encodes two pieces of information: the year of observation and the variable being observed. To deal with this, we'll have to first transform the data from one untidy format to another. We're going to use the gather() function to make each row a country-year-variable.

What gather() does is make a row for each entry from a set of columns. It's probably easiest to understand it by seeing it in practice:

```
long_data <- gather(untidy_data, key = variable,</pre>
    value = number, gdp.2005:unemp.2008)
head(long_data)
## # A tibble: 6 × 3
##
     country variable
                         number
##
       <chr>
                 <chr>
                           <dbl>
## 1
          AD gdp.2005
                         3.8423
## 2
          AE gdp.2005 253.9655
          AF gdp.2005
## 3
                         9.7630
## 4
          AG gdp.2005
                         1.1190
## 5
          AL gdp.2005
                         9.2684
          AM gdp.2005
## 6
                         7.6678
```

With the first argument, we told gather() to use the untidy\_data data frame. With the last argument, we told it the set of columns to "gather" together into a single column. The key column specifies the name of the variable to store the "key" (original column name) in, and the value column specifies the name of the variable to store the associated value. For example, the second row of long data encodes what we previously saw as the gdp.2005 column of untidy\_data.

Now we have a new problem, which is that variable encodes two pieces of information: the variable and the year of its observation. tidyr provides the separate() function to solve that, splitting a single variable into two.

```
long_data <- separate(long_data, col = variable,</pre>
    into = c("var", "year"))
head(long_data)
## # A tibble: 6 × 4
##
     country
                var year
                             number
```

<sup>4</sup> Insert lame joke about how Americanists haven't heard of other countries. But, seriously, if you're confused because you haven't heard of other countries, just think of "state-years".

```
##
       <chr> <chr> <chr>
                               <dbl>
## 1
           AD
                gdp
                      2005
                              3.8423
## 2
           ΑE
                      2005 253.9655
                gdp
                              9.7630
## 3
           AF
                gdp
                      2005
## 4
           AG
                      2005
                              1.1190
                gdp
## 5
                      2005
                              9.2684
           ΑL
                gdp
## 6
           AM
                gdp
                      2005
                              7.6678
```

So now we have country-year-variable data, with the year and variable conveniently stored in different columns. To turn this into country-year data, we can use the spread() function, which is like the inverse of gather(). spread() takes a key column and a value column, and turns each different key into a column of its own.

```
clean_data <- spread(long_data, key = var, value = number)</pre>
head(clean_data)
```

```
## # A tibble: 6 × 5
##
     country
              year
                          gdp
                                   pop unemp
##
       <chr> <chr>
                       <dbl>
                                 <dbl> <dbl>
## 1
          AD
               2005
                      3.8423 0.081223
                                           NA
## 2
               2006
                      4.0184 0.083373
          AD
                                           NA
## 3
          AD
               2007
                      4.0216 0.084878
                                           NA
## 4
          AD
               2008
                      3.6759 0.085616
                                           NA
## 5
          ΑE
               2005 253.9655 4.481976
                                          3.1
               2006 278.9489 5.171255
## 6
          ΑE
                                          3.3
```

When using spread() on data that you didn't previously gather(), be sure to set the fill argument to tell it how to fill in empty cells. A simple example:

### test\_data

```
## # A tibble: 3 × 3
##
           id
                  k
       <chr> <chr> <dbl>
##
## 1 brenton
                       10
                  а
                       20
## 2 brenton
                  b
## 3 patrick
                  b
                        5
spread(test_data, key = k, value = v)
## # A tibble: 2 × 3
##
           id
                  a
                        b
## *
       <chr> <dbl> <dbl>
## 1 brenton
                 10
                       20
## 2 patrick
                 NA
                        5
```

```
spread(test_data, key = k, value = v, fill = 100)
## # A tibble: 2 × 3
##
          id
                        b
                  a
       <chr> <dbl> <dbl>
## *
## 1 brenton
                 10
                       20
## 2 patrick
                100
                        5
```

One more important note on **tidyverse** semantics. It includes a fabulous feature called the pipe, %>%, which makes it easy to string together a truly mind-boggling number of commands.

In pipe syntax, x %>% f() is equivalent to f(x). That seems like a wasteful and confusing way to write f(x), and it is. But if you want to string together a bunch of commands, it's much easier to comprehend

```
x %>% f() %>% g() %>% h() %>% i()
  than i(h(g(f(x)))).
```

You can pass function arguments using the pipe too. For example, f(x, bear = "moose") is equivalent to x %% f(bear = "moose").

The key thing about the **tidyverse** functions is that each of them takes a data frame as its first argument, and returns a data frame as its output. This makes them highly amenable to piping. For example, we can combine all three steps of our tidying above with a single command, thanks to the pipe:<sup>5</sup>

```
untidy_data %>% gather(key = variable, value = number,
    gdp.2005:unemp.2008) %>% separate(col = variable,
    into = c("var", "year")) %>% spread(key = var,
    value = number)
## # A tibble: 860 × 5
##
     country year
                         gdp
                                  pop unemp
## *
       <chr> <chr>
                       <dbl>
                                <dbl> <dbl>
## 1
          AD
              2005
                      3.8423 0.081223
                                          NA
## 2
          AD
              2006
                      4.0184 0.083373
                                          NA
## 3
          AD
              2007
                      4.0216 0.084878
                                          NA
## 4
              2008
                      3.6759 0.085616
          AD
                                         NΑ
              2005 253.9655 4.481976
## 5
          ΑE
                                         3.1
## # ... with 855 more rows
```

Without the pipe, if we wanted to run all those commands together, we would have to write:

```
spread(separate(gather(untidy_data, key = variable,
   value = number, gdp.2005:unemp.2008), col = variable,
   into = c("var", "year")), key = var, value = number)
  Sad!
```

<sup>5</sup> If you are reading the PDF copy of these notes (i.e., the ones I hand out in class), the line breaks are eliminated, making the piped commands rather hard to read. I am working on fixing this. For now, you may find the online notes at http://bkenkel.com/pdaps easier to follow.

#### 3.3 Transforming and Aggregating

Tidying the data usually isn't the end of the process. If you want to perform further calculations on the raw, that's where the tools in dplyr (part of, you guessed it, the tidyverse) come in.

Perhaps the simplest **dplyr** function (or "verb", as the R hipsters would say) is rename(), which lets you rename columns.

```
clean_data %>% rename(gross_domestic_product = gdp)
```

```
## # A tibble: 860 × 5
     country year gross_domestic_product
## *
       <chr> <chr>
                                      <dbl>
              2005
                                     3.8423
## 1
          AD
## 2
              2006
                                     4.0184
          AD
## 3
          AD
              2007
                                     4.0216
## 4
          AD
              2008
                                     3.6759
                                   253.9655
              2005
## 5
          ΑE
     ... with 855 more rows, and 2 more
## #
       variables: pop <dbl>, unemp <dbl>
```

The **dplyr** functions, like the vast majority of R functions, do not modify their inputs. In other words, running rename() on clean\_data will return a renamed copy of clean\_data, but won't overwrite the original.

## clean\_data

```
## # A tibble: 860 × 5
##
     country year
                         gdp
                                   pop unemp
## *
       <chr> <chr>
                       <dbl>
                                 <dbl> <dbl>
## 1
          AD
               2005
                      3.8423 0.081223
                                          NA
## 2
          AD
              2006
                      4.0184 0.083373
                                          NA
## 3
                      4.0216 0.084878
          AD
              2007
                                          NA
## 4
          AD
              2008
                      3.6759 0.085616
                                          NA
              2005 253.9655 4.481976
## 5
          ΑE
                                         3.1
## # ... with 855 more rows
```

If you wanted to make the change stick, you would have to run:

```
clean_data <- clean_data %>% rename(gross_domestic_product = gdp)
```

select() lets you keep a couple of columns and drop all the others. Or vice versa if you use minus signs.

```
clean_data %>% select(country, gdp)
```

```
## # A tibble: 860 × 2
##
     country
                  gdp
## *
       <chr>
                <dbl>
## 1
          AD
               3.8423
## 2
          AD
               4.0184
## 3
          AD
               4.0216
## 4
          AD
               3.6759
## 5
         AE 253.9655
## # ... with 855 more rows
clean_data %>% select(-pop)
## # A tibble: 860 \times 4
     country year
                        gdp unemp
## *
       <chr> <chr>
                      <dbl> <dbl>
## 1
          AD 2005
                     3.8423
                               NA
## 2
          AD 2006
                     4.0184
                               NA
## 3
          AD 2007
                     4.0216
                               NA
## 4
          AD 2008
                     3.6759
                              NA
## 5
          AE 2005 253.9655
                               3.1
## # ... with 855 more rows
  mutate() lets you create new variables that are transformations of
old ones.
clean_data %>% mutate(gdppc = gdp/pop, log_gdppc = log(gdppc))
## # A tibble: 860 × 7
     country year
                      gdp
                                 pop unemp
##
       <chr> <chr>
                      <dbl>
                               <dbl> <dbl>
## 1
          AD 2005 3.8423 0.081223
                                         NΑ
## 2
          AD 2006 4.0184 0.083373
                                         NA
## 3
          AD 2007
                     4.0216 0.084878
                                         NA
                     3.6759 0.085616
## 4
          AD 2008
                                         NA
          AE 2005 253.9655 4.481976
                                        3.1
## # ... with 855 more rows, and 2 more
       variables: gdppc <dbl>, log_gdppc <dbl>
  filter() cuts down the data according to the logical condition(s)
you specify.
clean_data %>% filter(year == 2006)
## # A tibble: 215 × 5
##
     country year
                        gdp
                                   pop unemp
##
       <chr> <chr>
                      <dbl>
                                 <dbl> <dbl>
## 1
          AD 2006
                     4.0184 0.083373
                                          NA
```

```
## 2
          ΑE
              2006 278.9489 5.171255
                                          3.3
## 3
          AF
              2006
                    10.3052 25.183615
                                          8.8
## 4
          AG
              2006
                      1.2687
                              0.083467
                                          NΑ
## 5
          ΑL
              2006
                      9.7718
                              2.992547
                                       12.4
## # ... with 210 more rows
```

summarise() calculates summaries of the data. For example, let's find the maximum unemployment rate.

```
clean_data %>% summarise(max_unemp = max(unemp,
    na.rm = TRUE))
## # A tibble: 1 × 1
     max_unemp
##
##
         <dbl>
## 1
          37.6
```

This seems sort of useless, until you combine it with the group\_by() function. If you group the data before summarise-ing it, you'll calculate a separate summary for each group. For example, let's calculate the maximum unemployment rate for each year in the data.

```
clean_data %>% group_by(year) %>% summarise(max_unemp = max(unemp,
    na.rm = TRUE))
## # A tibble: 4 × 2
##
      year max_unemp
##
     <chr>>
                <dbl>
## 1
      2005
                 37.3
## 2
      2006
                 36.0
## 3
      2007
                 34.9
## 4
      2008
                 37.6
```

summarise() produces a "smaller" data frame than the input—one row per group. If you want to do something similar, but preserving the structure of the original data, use mutate in combination with group\_by.

```
clean_data %>% group_by(year) %>% mutate(max_unemp = max(unemp,
   na.rm = TRUE), unemp_over_max = unemp/max_unemp) %>%
    select(country, year, contains("unemp"))
## Source: local data frame [860 x 5]
## Groups: year [4]
##
##
     country year unemp max_unemp
##
       <chr> <chr> <dbl>
                             <dbl>
```

```
## 1
           AD
               2005
                        NA
                                 37.3
## 2
               2006
           AD
                        NA
                                 36.0
               2007
## 3
           AD
                        NA
                                 34.9
## 4
           AD
               2008
                        NA
                                 37.6
## 5
           ΑE
               2005
                       3.1
                                 37.3
     ... with 855 more rows, and 1 more
## #
       variables: unemp_over_max <dbl>
## #
```

This gives us back the original data, but with a max\_unemp variable recording the highest unemployment level that year. We can then calculate each individual country's unemployment as a percentage of the maximum. Whether grouped mutate or summarise is better depends, of course, on the purpose and structure of your analysis.

Notice how I selected all of the unemployment-related columns with contains("unemp"). See ?select\_helpers for a full list of helpful functions like this for select-ing variables.

## 3.4 Merging

Only rarely will you be lucky enough to draw all your data from a single source. More often, you'll be merging together data from multiple sources.

The key to merging data from separate tables is to have consistent identifiers across tables. For example, if you run an experiment, you might have demographic data on each subject in one table, and each subject's response to each treatment in another table. Naturally, you'll want to have a subject identifier that "links" the records across tables, as in the following hypothetical example.

### subject\_data

```
## # A tibble: 3 × 4
##
        id gender loves_bernie does_yoga
     <dbl>
             <chr>>
                           <chr>>
                                      <chr>
##
## 1
      1001
              male
                             yes
                                         no
## 2
      1002 female
                              no
                                        yes
## 3
      1003
              male
                              no
                                         no
```

subject\_response\_data

```
## # A tibble: 6 × 3
##
        id treatment response
     <dbl>
##
                <chr>>
                         <chr>>
      1001 read book
## 1
                           sad
## 2
      1001 watch_tv
                           sad
## 3
     1002 read_book
                         happy
```

```
1002 watch_tv
                          sad
## 5
     1003 read_book
                          sad
## 6
     1003 watch_tv
                        happy
```

## # ... with 1,065 more rows

Let's practice merging data with our cleaned-up country-year data. We'll take two datasets from my website: a country-level dataset with latitudes and longitudes, and a country-year-level dataset with inflation over time.

```
latlong_data <- read_csv("http://bkenkel.com/data/latlong.csv")</pre>
latlong_data
## # A tibble: 245 × 3
     country latitude longitude
##
##
       <chr>>
                 <dbl>
                            <dbl>
## 1
          AD
                42.546
                           1.6016
## 2
          ΑE
                23.424
                          53.8478
## 3
          AF
                33.939
                          67.7100
## 4
          AG
                17.061
                        -61.7964
## 5
          ΑI
                18.221
                        -63.0686
## # ... with 240 more rows
inflation data <- read csv("http://bkenkel.com/data/inflation.csv")
inflation_data
## # A tibble: 1,070 \times 3
##
     country year inflation
##
       <chr> <int>
                         <dbl>
## 1
          AD
               2004
                            NA
## 2
               2005
          AD
                            NA
## 3
          AD
               2006
                            NA
## 4
          AD
               2007
                            NA
               2008
## 5
          AD
                            NA
```

For your convenience, both of these datasets use the same two-letter country naming scheme as the original data. Unfortunately, out in the real world, data from different sources often use incommensurate naming schemes. Converting from one naming scheme to another is part of the data cleaning process, and it requires careful attention.

dplyr contains various \_join() functions for merging. Each of these take as arguments the two data frames to merge, plus the names of the identifier variables to merge them on. The one I use most often is left\_join(), which keeps every row from the first ("left") data frame and merges in the columns from the second ("right") data frame.

For example, let's merge the latitude and longitude data for each country into clean\_data.

left\_join(clean\_data, latlong\_data, by = "country")

```
## # A tibble: 860 × 7
##
     country year
                         gdp
                                   pop unemp
##
       <chr> <chr>
                       <dbl>
                                 <dbl> <dbl>
## 1
          AD
              2005
                      3.8423 0.081223
## 2
          AD
              2006
                      4.0184 0.083373
                                          NA
## 3
              2007
                      4.0216 0.084878
          AD
                                          NA
## 4
          AD
              2008
                      3.6759 0.085616
## 5
          ΑE
              2005 253.9655 4.481976
                                         3.1
     ... with 855 more rows, and 2 more
       variables: latitude <dbl>,
## #
       longitude <dbl>
```

Since latlong\_data is country-level, the value is the same for each year. So the merged data contains redundant information. This is one reason to store data observed at different levels in different tables—with redundant observations, it is easier to make errors yet harder to catch them and fix them.

We can also merge data when the identifier is stored across multiple columns, as in the case of our country-year data. But first, a technical note.<sup>6</sup> You might notice that the year column of clean\_data is labeled <chr>, as in character data. Yet the year column of inflation\_data is labeled <int>, as in integer data. We can check that by running class() on each respective column.

<sup>6</sup> This won't be the case if you got clean\_data by loading it in directly from clean-data.csv on my website, since read\_csv() will have correctly encoded year as an integer.

```
class(clean_data$year)
## [1] "character"
class(inflation_data$year)
## [1] "integer"
```

From R's perspective, the character string "1999" is a very different thing than the integer number 1999. Therefore, if we try to merge clean\_data and inflation\_data on the year variable, it will throw an error.

## Error in eval(substitute(expr), envir, enclos): Can't join on 'year' x 'year' because of incompatible

To fix this, let's use mutate() to convert the year column of clean\_data to an integer. We probably should have done this in the first place—after all, having the year encoded as a character string would have thrown off plotting functions, statistical functions, or anything else where it would be more natural to treat the year like a number.

```
clean_data <- mutate(clean_data, year = as.integer(year))
clean_data</pre>
```

```
## # A tibble: 860 × 5
##
     country year
                         gdp
                                  pop unemp
##
       <chr> <int>
                       <dbl>
                                <dbl> <dbl>
## 1
          AD
              2005
                      3.8423 0.081223
## 2
          AD
              2006
                      4.0184 0.083373
                                          NA
## 3
          AD
              2007
                      4.0216 0.084878
                                          NA
## 4
          AD
              2008
                      3.6759 0.085616
                                          NA
## 5
          ΑE
              2005 253.9655 4.481976
                                         3.1
## # ... with 855 more rows
```

Looks the same as before, except with an important difference: year is now labeled <int>.

Now we can merge the two datasets together without issue. Notice how we use a vector in the by argument to specify multiple columns to merge on.

```
## # A tibble: 860 × 6
##
     country year
                         gdp
                                  pop unemp
                                <dbl> <dbl>
##
       <chr> <int>
                       <dbl>
## 1
          AD
              2005
                      3.8423 0.081223
## 2
          AD
              2006
                      4.0184 0.083373
                                         NA
## 3
          AD
              2007
                      4.0216 0.084878
                                         NΑ
## 4
          AD
              2008
                      3.6759 0.085616
                                          NA
## 5
          ΑE
              2005 253.9655 4.481976
    ... with 855 more rows, and 1 more
## #
       variables: inflation <dbl>
```

You might remember that inflation\_data contained some countryyears not included in the original data (namely, observations from 2004). If we want the merged data to use the observations from inflation\_data rather than clean\_data, we can use the right\_join() function.

```
## # A tibble: 1,070 \times 6
##
     country year
                       gdp
                                pop unemp
##
       <chr> <int>
                    <dbl>
                              <dbl> <dbl>
## 1
          AD
              2004
                                 NA
                        NA
## 2
          AD
              2005 3.8423 0.081223
                                        NA
## 3
          AD
              2006 4.0184 0.083373
                                        NA
## 4
          AD
              2007 4.0216 0.084878
                                        NA
              2008 3.6759 0.085616
## 5
          AD
                                        NA
## # ... with 1,065 more rows, and 1 more
## #
       variables: inflation <dbl>
```

One last common issue in merging is that the identifier variables have different names in the two datasets. If it's inconvenient or infeasible to correct this by renaming the columns in one or the other, you can specify the by argument as in the following example.

```
inflation_data <- rename(inflation_data, the_country = country,
    the_year = year)
inflation_data
## # A tibble: 1,070 \times 3
##
     the_country the_year inflation
           <chr>>
                     <int>
##
                               <dbl>
## 1
              AD
                      2004
                                  NA
## 2
              AD
                      2005
                                   NA
## 3
              AD
                      2006
                                  NA
## 4
              AD
                      2007
                                  NA
## 5
              AD
                      2008
                                   NA
## # ... with 1,065 more rows
left_join(clean_data, inflation_data, by = c(country = "the_country",
    year = "the_year"))
## # A tibble: 860 × 6
##
     country year
                         gdp
                                  pop unemp
##
       <chr> <int>
                       <dbl>
                                <dbl> <dbl>
## 1
          AD
              2005
                      3.8423 0.081223
                                          NΑ
## 2
                      4.0184 0.083373
          AD
              2006
                                          NA
## 3
              2007
                      4.0216 0.084878
          AD
                                          NA
## 4
          AD
              2008
                      3.6759 0.085616
                                          NA
              2005 253.9655 4.481976
## 5
          ΑE
                                         3.1
## # ... with 855 more rows, and 1 more
       variables: inflation <dbl>
```

#### 3.5Appendix: Creating the Example Data

I used the same tools this chapter introduces to create the untidy data. I may as well include the code to do it, in case it helps further illustrate how to use the tidyverse tools (and, as a bonus, the WDI package for downloading World Development Indicators data).

First I load the necessary packages.

```
library("tidyverse")
library("WDI")
library("countrycode")
library("stringr")
```

Next, I download the relevant WDI data. I used the WDIsearch() function to locate the appropriate indicator names.

```
dat_raw <- WDI(country = "all",</pre>
               indicator = c("NY.GDP.MKTP.KD",
                                                  # GDP in 2000 USD
                              "SP.POP.TOTL",
                                                  # Total population
                              "SL.UEM.TOTL.ZS"), # Unemployment rate
               start = 2005,
               end = 2008)
```

#### head(dat\_raw)

```
country year NY.GDP.MKTP.KD
##
     iso2c
                                1.6428e+12
## 1
        1A Arab World 2005
## 2
        1A Arab World 2006
                                 1.7629e+12
## 3
        1A Arab World 2007
                                 1.8625e+12
## 4
        1A Arab World 2008
                                 1.9799e+12
## 5
        1W
                 World 2005
                                 5.7703e+13
                 World 2006
                                 6.0229e+13
## 6
        1W
     SP.POP.TOTL SL.UEM.TOTL.ZS
##
## 1
       313430911
                         12.1402
## 2
       320906736
                         11.3296
## 3
       328766559
                         10.8961
## 4
       336886468
                         10.5060
      6513959904
## 5
                          6.1593
                          5.9000
## 6
      6594722462
```

I want to get rid of the aggregates, like the "Arab World" and "World" we see here. As a rough tack at that, I'm going to exclude those so-called countries whose ISO codes don't appear in the coun**trycode** package data.<sup>7</sup>

```
dat_countries <- dat_raw %>% filter(iso2c %in%
    countrycode_data$iso2c)
```

<sup>&</sup>lt;sup>7</sup> countrycode is a very useful, albeit imperfect, package for converting between different country naming/coding schemes.

Let's check on which countries are left. (I cut it down to max six characters per country name for printing purposes.)

```
dat_countries$country %>% unique() %>% str_sub(start = 1,
    end = 6
##
     [1] "Andorr" "United" "Afghan" "Antigu"
##
     [5] "Albani" "Armeni" "Angola" "Argent"
     [9] "Americ" "Austri" "Austra" "Aruba"
##
    [13] "Azerba" "Bosnia" "Barbad" "Bangla"
##
    [17] "Belgiu" "Burkin" "Bulgar" "Bahrai"
##
##
    [21] "Burund" "Benin" "Bermud" "Brunei"
    [25] "Bolivi" "Brazil" "Bahama" "Bhutan"
##
##
    [29] "Botswa" "Belaru" "Belize" "Canada"
    [33] "Congo," "Centra" "Congo," "Switze"
##
    [37] "Cote d" "Chile" "Camero" "China"
##
    [41] "Colomb" "Costa " "Cuba"
                                     "Cabo V"
##
##
    [45] "Curaca" "Cyprus" "Czech " "German"
    [49] "Djibou" "Denmar" "Domini" "Domini"
##
##
    [53] "Algeri" "Ecuado" "Estoni" "Egypt,"
##
    [57] "Eritre" "Spain"
                           "Ethiop" "Finlan"
    [61] "Fiji"
                  "Micron" "Faroe " "France"
##
    [65] "Gabon"
                  "United" "Grenad" "Georgi"
##
    [69] "Ghana"
                  "Gibral" "Greenl" "Gambia"
##
##
    [73] "Guinea" "Equato" "Greece" "Guatem"
##
    [77] "Guam"
                  "Guinea" "Guyana" "Hong K"
    [81] "Hondur" "Croati" "Haiti"
                                     "Hungar"
##
##
    [85] "Indone" "Irelan" "Israel" "Isle o"
    [89] "India"
                  "Iraq"
                            "Iran, " "Icelan"
##
    [93] "Italy"
                  "Jamaic" "Jordan" "Japan"
##
                  "Kyrgyz" "Cambod" "Kiriba"
##
    [97] "Kenya"
   [101] "Comoro" "St. Ki" "Korea," "Korea,"
   [105] "Kuwait" "Cayman" "Kazakh" "Lao PD"
   [109] "Lebano" "St. Lu" "Liecht" "Sri La"
   [113] "Liberi" "Lesoth" "Lithua" "Luxemb"
   [117] "Latvia" "Libya"
                           "Morocc" "Monaco"
  [121] "Moldov" "Monten" "St. Ma" "Madaga"
##
## [125] "Marsha" "Macedo" "Mali"
                                     "Myanma"
  [129] "Mongol" "Macao " "Northe" "Maurit"
## [133] "Malta"
                  "Maurit" "Maldiv" "Malawi"
## [137] "Mexico" "Malays" "Mozamb" "Namibi"
## [141] "New Ca" "Niger"
                           "Nigeri" "Nicara"
## [145] "Nether" "Norway" "Nepal"
                                     "Nauru"
                            "Panama" "Peru"
## [149] "New Ze" "Oman"
```

## [153] "French" "Papua " "Philip" "Pakist"

```
## [157] "Poland" "Puerto" "West B" "Portug"
## [161] "Palau" "Paragu" "Qatar"
                                    "Romani"
## [165] "Serbia" "Russia" "Rwanda" "Saudi "
## [169] "Solomo" "Seyche" "Sudan"
                                    "Sweden"
## [173] "Singap" "Sloven" "Slovak" "Sierra"
## [177] "San Ma" "Senega" "Somali" "Surina"
## [181] "South " "Sao To" "El Sal" "Sint M"
## [185] "Syrian" "Swazil" "Turks " "Chad"
## [189] "Togo"
                  "Thaila" "Tajiki" "Timor-"
## [193] "Turkme" "Tunisi" "Tonga"
                                    "Turkey"
## [197] "Trinid" "Tuvalu" "Tanzan" "Ukrain"
## [201] "Uganda" "United" "Urugua" "Uzbeki"
## [205] "St. Vi" "Venezu" "Britis" "Virgin"
## [209] "Vietna" "Vanuat" "Samoa"
                                    "Yemen,"
## [213] "South " "Zambia" "Zimbab"
```

With that out of the way, there's still some cleaning up to do. The magnitudes of GDP and population are too large, and the variable names are impenetrable. Also, the country variable, while helpful, is redundant now that we're satisfied with the list of countries remaining.

```
dat_countries <- dat_countries %>% select(-country) %>%
   rename(gdp = NY.GDP.MKTP.KD, pop = SP.POP.TOTL,
       unemp = SL.UEM.TOTL.ZS, country = iso2c) %>%
   mutate(gdp = gdp/1e+09, pop = pop/1e+06)
```

#### head(dat\_countries)

```
##
     country year
                                 pop unemp
                        gdp
          AD 2005
## 1
                     3.8423 0.081223
                                         NA
## 2
          AD 2006
                     4.0184 0.083373
                                         NA
## 3
          AD 2007
                     4.0216 0.084878
                                         NA
## 4
          AD 2008
                     3.6759 0.085616
                                         NA
          AE 2005 253.9655 4.481976
## 5
                                        3.1
## 6
          AE 2006 278.9489 5.171255
                                        3.3
```

Now I convert the data to "long" format.

```
dat_countries_long <- dat_countries %>% gather(key = variable,
   value = value, gdp:unemp)
```

#### head(dat\_countries\_long)

```
##
     country year variable
                                value
## 1
          AD 2005
                               3.8423
                         gdp
## 2
          AD 2006
                               4.0184
                         gdp
```

```
## 3
          AD 2007
                        gdp
                               4.0216
## 4
          AD 2008
                        gdp
                               3.6759
                        gdp 253.9655
## 5
          AE 2005
## 6
          AE 2006
                        gdp 278.9489
```

I then smush variable and year into a single column, and drop the individual components.

```
dat_countries_long <- dat_countries_long %>% mutate(var_year = paste(variable,
    year, sep = ".")) %>% select(-variable, -year)
```

#### head(dat\_countries\_long)

```
value var_year
##
     country
## 1
          AD
               3.8423 gdp.2005
## 2
          AD
               4.0184 gdp.2006
## 3
          AD
               4.0216 gdp.2007
## 4
          AD
               3.6759 gdp.2008
## 5
          AE 253.9655 gdp.2005
          AE 278.9489 gdp.2006
## 6
```

Finally, I "widen" the data, so that each var\_year is a column of

```
dat_countries_wide <- dat_countries_long %>% spread(key = var_year,
    value = value)
```

#### head(dat countries wide)

```
##
    country gdp.2005 gdp.2006 gdp.2007
## 1
         AD
              3.8423
                       4.0184
                                4.0216
## 2
         AE 253.9655 278.9489 287.8318
## 3
         AF
              9.7630 10.3052 11.7212
## 4
         AG
              1.1190
                       1.2687
                                1.3892
## 5
              9.2684
                       9.7718 10.3483
         AL
## 6
         MA
              7.6678
                       8.6797
                                9.8731
    gdp.2008 pop.2005 pop.2006 pop.2007
##
## 1
      3.6759 0.081223 0.083373 0.084878
## 2 297.0189 4.481976 5.171255 6.010100
## 3 12.1445 24.399948 25.183615 25.877544
## 4
      1.3902 0.082565 0.083467 0.084397
    11.1275 3.011487 2.992547 2.970017
## 5
## 6 10.5544 3.014917 3.002161 2.988117
     pop.2008 unemp.2005 unemp.2006 unemp.2007
##
## 1 0.085616
                      NA
                                 NA
                                            NA
## 2 6.900142
                     3.1
                                3.3
                                           3.4
```

```
8.4
## 3 26.528741
                       8.5
                                   8.8
## 4 0.085350
                        NA
                                               NA
                                    NA
## 5 2.947314
                      12.5
                                             13.5
                                  12.4
## 6 2.975029
                      27.8
                                             28.4
                                  28.6
##
     unemp.2008
## 1
             NA
## 2
            4.0
## 3
            8.9
## 4
             NA
## 5
           13.0
## 6
           16.4
```

Now we have some ugly data. I save the output to upload to my website.

```
write_csv(dat_countries_wide, path = "untidy-data.csv")
```

And here's how I made the second country-year dataset used in the merging section. The country dataset with latitudes and longitudes is from https://developers.google.com/public-data/docs/ canonical/countries\_csv.

```
dat_2 <- WDI(country = "all", indicator = "FP.CPI.TOTL.ZG",</pre>
   start = 2004, end = 2008) %>% as_data_frame() %>%
   select(country = iso2c, year, inflation = FP.CPI.TOTL.ZG) %>%
   mutate(year = as.integer(year)) %>% filter(country %in%
    clean_data$country) %>% arrange(country, year)
write_csv(dat_2, path = "inflation.csv")
```

# Data Visualization

Visualization is most important at the very beginning and the very end of the data analysis process. In the beginning, when you've just gotten your data together, visualization is perhaps the easiest tool to explore each variable and learn about the relationships among them. And when your analysis is almost complete, you will (usually) use visualizations to communicate your findings to your audience.

We only have time to scratch the surface of data visualization. This chapter will cover the plotting techniques I find most useful for exploratory and descriptive data analysis. We will talk about graphical techniques for presenting the results of regression analyses later in the class—once we've, you know, learned something about regression.

#### 4.1 Basic Plots

We will use the **ggplot2** package, which is part of—I'm as tired of it as you are—the **tidyverse**.

## library("tidyverse")

For the examples today, we'll be using a dataset with statistics about the fifty U.S. states in 1977, which is posted on my website.

<sup>1</sup> Why 1977? Because it was easily available. See the appendix to this schapter

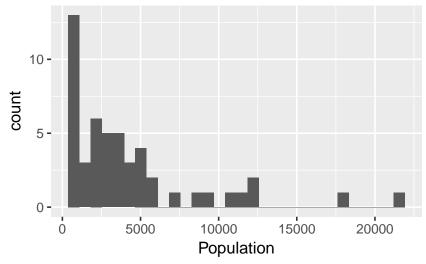
state\_data <- read\_csv("http://bkenkel.com/data/state-data.csv")chapter.
state\_data</pre>

```
## # A tibble: 50 × 12
##
          State Abbrev Region Population Income
##
           <chr>
                  <chr>
                          <chr>
                                      <dbl>
                                              <dbl>
## 1
        Alabama
                     AL
                          South
                                       3615
                                              3624
## 2
         Alaska
                     AK
                           West
                                        365
                                              6315
## 3
                                       2212
                                              4530
        Arizona
                     AZ
                           West
                                       2110
## 4
       Arkansas
                     AR
                          South
                                              3378
## 5 California
                     CA
                                      21198
                                              5114
                           West
```

```
... with 45 more rows, and 7 more
## #
       variables: Illiteracy <dbl>,
## #
       LifeExp <dbl>, Murder <dbl>,
       HSGrad <dbl>, Frost <dbl>, Area <dbl>,
## #
## #
       IncomeGroup <chr>
```

When I obtain data, I start by looking at the univariate distribution of each variable via a histogram. The following code creates a histogram in ggplot.

```
ggplot(state_data, aes(x = Population)) + geom_histogram()
## `stat_bin()` using `bins = 30`. Pick
## better value with `binwidth`.
```



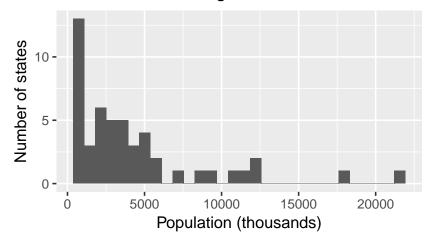
Let's walk through the syntax there. In the first line, we call ggplot(), specifying the data frame to draw from, then in the aes() command (which stands for "aesthetic") we specify the variable to plot. If this were a bivariate analysis, here we would have also specified a y variable to put on the y-axis. If we had just stopped there, we would have a sad, empty plot. The + symbol indicates that we'll be adding something to the plot. geom\_histogram() is the command to overlay a histogram.

We'll only be looking at a few of the ggplot commands today. I recommend taking a look at the online package documentation at http://docs.ggplot2.org to see all of the many features available.

When you're just making graphs for yourself to explore the data, you don't need to worry about things like axis labels as long as you can comprehend what's going on. But when you prepare graphs for others to read (including those of us grading your problem sets!) you need to include an informative title and axis labels. To that end, use the xlab(), ylab(), and ggtitle() commands.

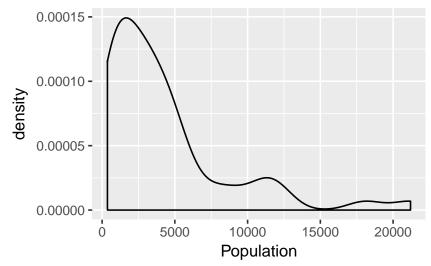
```
ggplot(state_data, aes(x = Population)) + geom_histogram() +
   xlab("Population (thousands)") + ylab("Number of states") +
    ggtitle("Some states are big, but most are small")
## `stat_bin()` using `bins = 30`. Pick
## better value with `binwidth`.
```

## Some states are big, but most are small



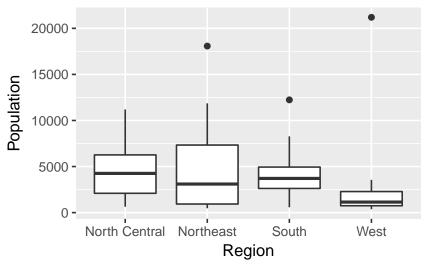
The density plot is a close relative of the histogram. It provides a smooth estimate of the probability density function of the data. Accordingly, the area under the density plot integrates to one. Depending on your purposes, this can make the y-axis of a density plot easier or (usually) harder to interpret than the count given by a histogram.





The box plot is a common way to look at the distribution of a continuous variable across different levels of a categorical variable.

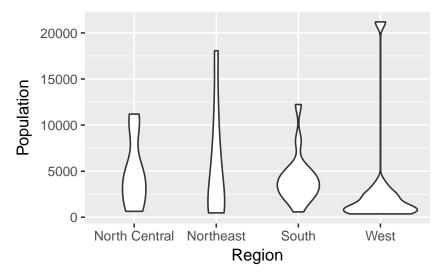
```
ggplot(state_data, aes(x = Region, y = Population)) +
    geom_boxplot()
```



A box plot consists of the following components:

- Center line: median of the data
- Bottom of box: 25th percentile
- Top of box: 75th percentile
- Lower "whisker": range of observations no more than 1.5 IQR (height of box) below the 25th percentile
- $\bullet~$  Upper "whisker": range of observations no more than 1.5 IQR above the 75th percentile
- Plotted points: any data lying outside the whiskers

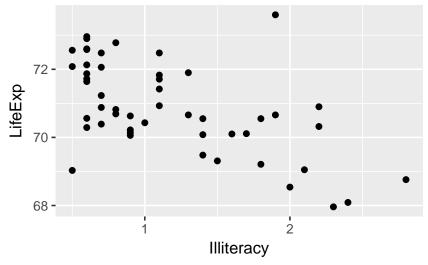
If you want to skip the summary and plot the full distribution of a variable across categories, you can use a violin plot.



Technically, violin plots convey more information than box plots since they show the full distribution. However, readers aren't as likely to be familiar with a violin plot. It's harder to spot immediately where the median is (though you could add that to the plot if you wanted). Plus, violin plots look goofy with outliers—see the "West" column above—whereas box plots handle them easily.

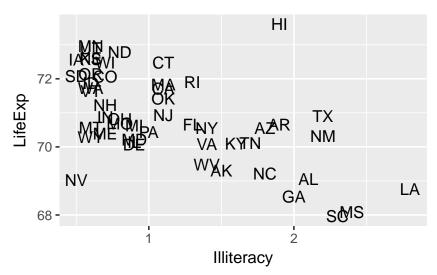
For visualizing relationships between continuous variables, nothing beats the scatterplot.

```
ggplot(state_data, aes(x = Illiteracy, y = LifeExp)) +
   geom_point()
```



When you're plotting states or countries, a hip thing to do is plot abbreviated names instead of points. To do that, you can use geom\_text() instead of geom\_point(), supplying an additional aesthetic argument telling ggplot where to draw the labels from.

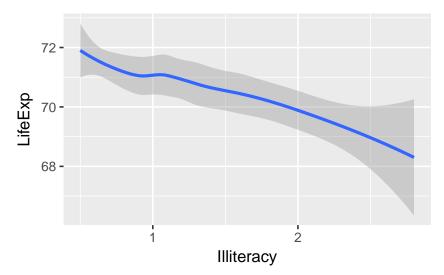
```
ggplot(state_data, aes(x = Illiteracy, y = LifeExp)) +
   geom_text(aes(label = Abbrev))
```



Maybe it's overwhelming to look at all that raw data and you just want a summary. For example, maybe you want an estimate of expected LifeExp for each value of Illiteracy. This is called the conditional expectation and will be the subject of much of the rest of the course. For now, just now that you can calculate a smoothed conditional expectation via geom\_smooth().

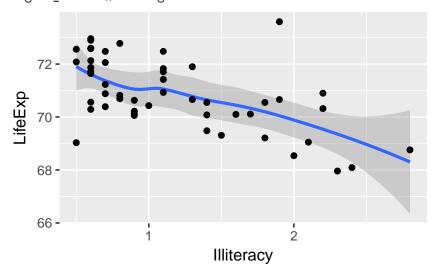
```
ggplot(state_data, aes(x = Illiteracy, y = LifeExp)) +
   geom_smooth()
```

## `geom\_smooth()` using method = 'loess'



And if you're the kind of overachiever who likes to have the raw data and the summary, you can do it. Just add them both to the ggplot() call.

```
ggplot(state_data, aes(x = Illiteracy, y = LifeExp)) +
    geom_smooth() + geom_point()
```



## ## 'geom\_smooth()' using method = 'loess'

#### 4.2 Saving Plots

When you're writing in R Markdown, the plots go straight into your document without much fuss. Odds are, your dissertation will contain plots but won't be written in R Markdown, which means you'll need to learn how to save them.

It's pretty simple:

- 1. Assign your ggplot() call to a variable.
- 2. Pass that variable to the ggsave() function.

```
pop_hist <- ggplot(state_data, aes(x = Population)) +</pre>
    geom_histogram()
```

```
ggsave(filename = "pop-hist.pdf", plot = pop_hist,
   width = 6, height = 3)
```

If you want plot types other than PDF, just set a different extension. See ?ggsave for the possibilities.

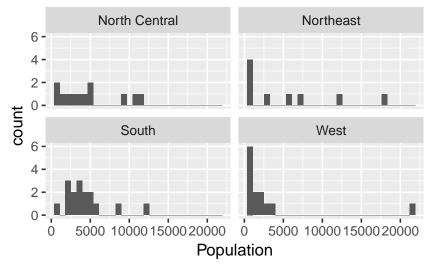
#### 4.3 Faceting

Suppose you want to split the data into subgroups, as defined by some variable in the data (e.g., the region states are in), and make the same plot for each subgroup. ggplot's faceting functions, facet\_wrap() and facet\_grid(), make this easy.

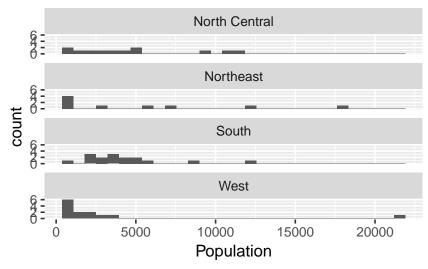
To split up plots according to a single grouping variable, use facet\_wrap(). This uses R's formula syntax, defined by the tilde ~, which you'll become well acquainted with once we start running regressions.

```
ggplot(state_data, aes(x = Population)) + geom_histogram() +
   facet_wrap(~Region)
```

```
## `stat_bin()` using `bins = 30`. Pick
## better value with `binwidth`.
```

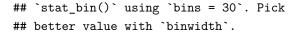


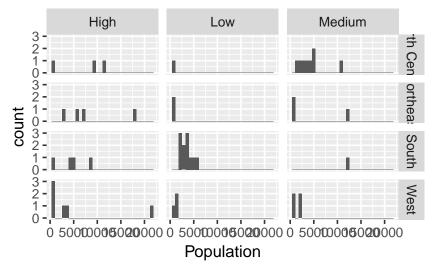
If you don't like the default arrangement, use the ncol argument.



For two grouping variables, use facet\_grid(), putting variables on both sides of the formula.

```
ggplot(state_data, aes(x = Population)) + geom_histogram() +
    facet_grid(Region ~ IncomeGroup)
```



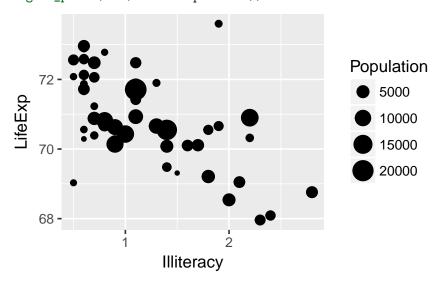


#### Aesthetics 4.4

Faceting is one way to incorporate information about additional variables into what would otherwise be a plot of just one or two variables. Aesthetics—which alter the appearance of particular plot features depending on the value of a variable—provide another way to do that.

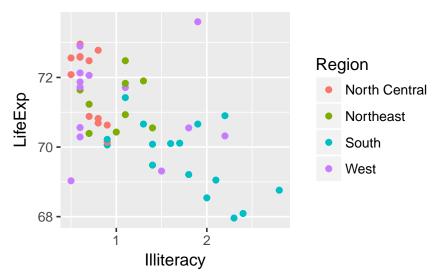
For example, when visualizing the relationship between statewide illiteracy and life expectancy, you might want larger states to get more visual weight. You can set the size aesthetic of the point geometry to vary according to the state's population.

ggplot(state\_data, aes(x = Illiteracy, y = LifeExp)) + geom\_point(aes(size = Population))



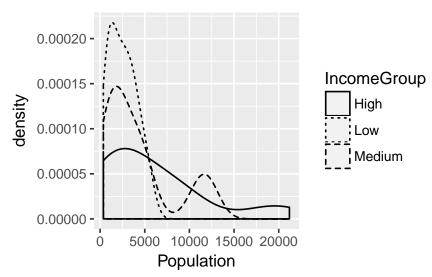
The **ggplot2** documentation lists the available aesthetics for each function. Another popular one is colour, which is great for on-screen display but not so much for the printed page. (And terrible for the colorblind!)

ggplot(state\_data, aes(x = Illiteracy, y = LifeExp)) + geom\_point(aes(colour = Region))



For line graphs or density plots, you can set the linetype to vary by category.

ggplot(state\_data, aes(x = Population)) + geom\_density(aes(linetype = IncomeGroup))



(I always find these incomprehensible with more than two lines, but maybe that's just me.) You can use multiple aesthetics together, and you can even combine aesthetics with faceting, as in the following example.

```
ggplot(state_data, aes(x = Illiteracy, y = LifeExp)) +
    geom_smooth() + geom_text(aes(label = Abbrev,
    colour = Region, size = Population)) + facet_wrap(~IncomeGroup)
## `geom_smooth()` using method = 'loess'
                                                   North Central
            High
                        Low
                                  Medium
                                                   Northeast
      74 .
                                                   South
                                                   West
                                               Population
                                                a 5000
      68
                                                a 10000
                                                   15000
               2
                           2
                                       2
                                   1
                     Illiteracy
                                                a 20000
```

But the fact that you can do something doesn't mean you should. That plot is so cluttered that it's hard to extract the relevant information from it. Data visualizations should communicate a clear message to viewers without overwhelming them. To do this well takes practice, patience, and maybe even a bit of taste.

## 4.5 Appendix: Creating the Example Data

The example data comes from data on U.S. states in 1977 that are included with base R. See ?state.

```
library("tidyverse")
```

```
state_data <- state.x77 %>% as_tibble() %>% add_column(State = rownames(state.x77),
    Abbrev = state.abb, Region = state.region,
    .before = 1) %>% rename(LifeExp = `Life Exp`,
   HSGrad = `HS Grad`) %>% mutate(IncomeGroup = cut(Income,
   breaks = quantile(Income, probs = seq(0, 1,
        by = 1/3), labels = c("Low", "Medium",
        "High"), include.lowest = TRUE))
write_csv(state_data, path = "state-data.csv")
```

# Bibliography

Bowers, J. (2011). Six Steps to a Better Relationship with Your Future Self. *The Political Methodologist*, 18(2):2–8.

Leek, J. (2015). The Elements of Data Analytic Style. Leanpub.

Wickham, H. (2014). Tidy Data. Journal of Statistical Software, 59(10).

Wilson, G., Aruliah, D. A., Brown, C. T., Hong, N. P. C., Davis, M., Guy, R. T., Haddock, S. H. D., Huff, K. D., Mitchell, I. M., Plumbley, M. D., Waugh, B., White, E. P., and Wilson, P. (2014). Best Practices for Scientific Computing. *PLOS Biology*, 12(1):e1001745.