Data Carpentry

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https://nps.edu/web/core

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Welcome



Preface

 ${\rm init}$



Part I

Day 1



Chapter 1

R and RStudio

1.1 R

1.1.1 Installation

 $\rm https://cran.r-project.org/$

1.2 RStudio

1.2.1 Installation

https://rstudio.com/products/rstudio/download/

Chapter 2

The Basics

```
"Hello, World!"
#> [1] "Hello, World!"
```

2.1 R as a Calculator

```
1 + 1
                          # addition
#> [1] 2
1 - 1
                          # subtraction
#> [1] 0
2 * 3
                         # multiplication
#> [1] 6
1 + 1 * 3
                         # combining operations
#> [1] 4
(1 + 1) * 3
                         # operator precedence
#> [1] 6
3 / 2
                          # division
#> [1] 1.5
# ↓↓ pronounced "modulo"
3 %% 2
                          # division remainder
#> [1] 1
4 %/% 2
                          # integer division
#> [1] 2
3^2
                          # exponents
#> [1] 9
```





If your code doesn't form a complete *expression*, then R will look for more on the next line. Here's an example:

```
1 + isn't a complete expression, so R prompt for more code on subsequent lines. You'll see something like the following:
> 1 +
```

+ +

If this happens, press the **Esc**(scape) key (you may have to click on the Console pane first) and fix your code.

2.2 Fundamental Types

R has several basic data types that serve as the foundation upon which everything is built.

```
# double (short for double-precision floating-point number)
#> [1] 1
3.14
              # also double (we can just think of them as decimals)
#> [1] 3.14
              # integer (`L` for "Literal" or `long` integers)
1L
#> [1] 1
1111
              # character or string (kinda... we'll discuss later)
#> [1] "1"
              # logical (similar to `bool`s in other languages)
TRUE
#> [1] TRUE
FALSE
              # also logical
#> [1] FALSE
              # complex (we're never going to use these)
#> [1] 0+4i
```



2.3. VARIABLES CHAPTER 2. THE BASICS



Like most programming languages, R lets us mix *comments* into our code. Anything that follows # on the same line is ignored by R.

Comments enable us to annotate our work or temporarily (hopefully) disable lines of code.

```
-1 * -1000 # a negative number times a negative is positive
#> [1] 1000
# TRUE + FALSE # felt cute, might un-comment later
```

Leverage comments to communicate with humans! They're an opportunity for explaining what something does and (often more-importantly) why something works or is necessary.

Since comments are ubiquitous, it's worth pointing out two common conventions:

```
# TODO(CC): CC (initials) is going to implement some behavior
# FIXME(BK): BK is going to fix some problem
```

2.3 Variables

R's assignment operator is <-.

```
my_first_var <- "referring to data w/ names is handy!"
my_first_var
#> [1] "referring to data w/ names is handy!"
```

We can also use = like many other languages, but we *highly* discourage this (especially starting out) because we use = elsewhere. If you stick to <-, you'll never have to guess where you've assigned variables or rely on context clues to predict ='s intended purpose or behavior.

You should always prefer descriptive variable names so that others can more easily understand your code. Most of the time, the other person will just be you in the future.

```
length <- 2
width <- 4

area <- length * width
area
#> [1] 8
```

2.4 Multiple Values

We'll almost always need to deal with more than one value, so R let's us c() ombine values.

```
c(1, 2, 3, 4, 5)
#> [1] 1 2 3 4 5
```



2.5. FUNCTIONS CHAPTER 2. THE BASICS

We won't get into the nitty-gritty details just yet, but we typically call a collection of values of the same type (homogeneous) a vector.

R is special for a few reasons and having native vectors is definitely one of them. Understanding how they work is fundamental to writing good (and fast!) code.

2.5 Functions

```
# \( name of function
sqrt(x = 16)
#> [1] \( 4 \)
```

R comes with some handy variables built-in , such as letters and LETTERS.

```
letters
#> [1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l" "m" "n" "o" "p" "q" "r" "s" "t" "u" "v"
#> [23] "w" "x" "y" "z"

LETTERS
#> [1] "A" "B" "C" "D" "E" "F" "G" "H" "I" "J" "K" "L" "M" "N" "O" "P" "Q" "R" "S" "T" "U" "V"
#> [23] "W" "X" "Y" "Z"
```

We refer to x =letters as a *named* argument because we specify the parameter (x) to which we're passing our argument (letters), but we often don't specify the name of a parameter.

```
tolower(letters)
#> [1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l" "m" "n" "o" "p" "q" "r" "s" "t" "u" "v"
#> [23] "w" "x" "y" "z"
```



2.6. DOCUMENTATION CHAPTER 2. THE BASICS

We can't screw up too easily since tolower() and toupper() only have one paramter (x), but many functions can take multiple arguments.

Let's say we have a vector of unsorted_numbers:

```
unsorted_numbers <- c(3, 2, 10, 8, 1, 4, 9, 6, 5, 7)
unsorted_numbers
#> [1] 3 2 10 8 1 4 9 6 5 7
```

Like most languages, R has a built-in sort() function we can use, which works like so:

```
sort(x = unsorted_numbers)
#> [1] 1 2 3 4 5 6 7 8 9 10
```

By default, sort() sorts in ascending order, but we oftentimes will want to sort in descending (or decreasing) order.

Rather than having a separate function called sort_decreasing(), we pass an argument to sort()'s decreasing parameter.

```
sort(x = unsorted_numbers, decreasing = TRUE)
#> [1] 10 9 8 7 6 5 4 3 2 1
```

Even though sort() has multiple parameters, we can still skip the names if we pass our arguments by position.

```
sort(unsorted_numbers, TRUE)
#> [1] 10 9 8 7 6 5 4 3 2 1
```

Considering that x is sort()'s first parameter, and decreaing is sort()'s second parameter, we can pass our arguments (unsorted_numbers and TRUE) in the same order and R will know what we meant.

We can also mix positional and named argument (and often do), but we should always prioritize readable code.

```
sort(unsorted_numbers, decreasing = TRUE) # good
#> [1] 10 9 8 7 6 5 4 3 2 1

sort(x = unsorted_numbers, TRUE)  # avoid this
#> [1] 10 9 8 7 6 5 4 3 2 1

sort(decreasing = TRUE, unsorted_numbers) # just... no
#> [1] 10 9 8 7 6 5 4 3 2 1
```

2.6 Documentation

You're hopefully wondering "How could we know the order of sort()'s parameters?" which leads us to documentation.

If you want more information on a specific function, you should check out the documentation, which you can do with? or help().

Here's what that looks like for sort()



?sort

```
help(sort) # does the same thing as `?sort`
```

sort {base}

R Documentation

Sorting or Ordering Vectors

Description

Sort (or *order*) a vector or factor (partially) into ascending or descending order. For ordering along more than one variable, e.g., for sorting data frames, see <u>order</u>.

Usage

Arguments

Χ

for sort an R object with a class or a numeric, complex, character or logical vector. For sort.int, a numeric, complex, character or logical vector, or a factor.

decreasing

logical. Should the sort be increasing or decreasing? For the "radix" method, this can be a vector of length equal to the number of arguments in For the other methods, it must be length one. Not available for partial sorting.

. . .

arguments to be passed to or from methods or (for the default methods and objects without a class) to sort.int.

na.last

for controlling the treatment of NAs. If TRUE, missing values in the data are put last; if FALSE, they are put first; if NA, they are removed.

partial

NULL or a vector of indices for partial sorting.

method

character string specifying the algorithm used. Not available for partial sorting. Can be abbreviated.

index.return

logical indicating if the ordering index vector should be returned as well. Supported by method == "radix" for any na.last mode and data type, and the other methods when na.last = NA (the default) and fully sorting non-factors.

There's a ton of information here, but all we're interested in at the moment is the order in which we need to pass arguments to sort(), which we can find in the **Arguments** section.

2.7 Missingness and Nothingness

2.7.1 NA

You may have noticed the na.last argument in sort()'s documentation. R can represent nothingness with NULL (same as null or None in other languages), but it can also represent unknown or missing values with NA.



```
unsorted_numbers_with_nas <- c(3, 2, 10, 8, NA, 1, 4, 9, NA, 6, 5, 7)
unsorted_numbers_with_nas
#> [1] 3 2 10 8 NA 1 4 9 NA 6 5 7
```

sort()'s default behavior is na.last = NA, which simply removes any NAs.

```
sort(unsorted_numbers_with_nas)
#> [1] 1 2 3 4 5 6 7 8 9 10
```

If we want to keep NAs, we must specify whether sort() places them first or last.

```
sort(unsorted_numbers_with_nas, na.last = TRUE)
#> [1] 1 2 3 4 5 6 7 8 9 10 NA NA
sort(unsorted_numbers_with_nas, na.last = FALSE)
#> [1] NA NA 1 2 3 4 5 6 7 8 9 10
```

2.7.2 NULL

For the moment, think of the difference between NA and NULL as being that vectors (like unsorted_numbers_with_nas) can have NA values but they cannot have NULL values.

If we try to put NULL in a vector, it simply disappears.

```
c(3, 2, 10, 8, NULL, 1, 4, 9, NULL, 6, 5, 7)
#> [1] 3 2 10 8 1 4 9 6 5 7
```

But, how do we check if something is NA or NULL?

2.8 Predicate Functions

A predicate function is a function that returns either TRUE or FALSE based on some condition the function is checking.

Predicate functions *should* use a name that expresses this intent, such as is<some condition>, any<some condition>(), or all<some condition>().

If we want check if something is NULL, we use is.null().

```
is.null("this string isn't NULL!")
#> [1] FALSE
is.null(NULL)
#> [1] TRUE
```

R has many built-in predicate functions, including ones to check the basic data types that we've already seen.



```
is.double(1)
#> [1] TRUE
is.double(1L)
#> [1] FALSE
vec_dbl \leftarrow c(8, 6, 7, 5, 3, 0, 9)
is.double(vec_dbl)
#> [1] TRUE
is.integer(1)
#> [1] FALSE
is.integer(1L)
#> [1] TRUE
vec_int <- 1:10</pre>
is.integer(vec_int)
#> [1] TRUE
is.character(3.14)
#> [1] FALSE
is.character("is it though?")
#> [1] TRUE
is.character(letters)
#> [1] TRUE
is.logical("the year 2020")
#> [1] FALSE
is.logical(TRUE)
#> [1] TRUE
is.logical(FALSE)
#> [1] TRUE
vec_lgl <- c(TRUE, FALSE, TRUE)</pre>
is.logical(vec_lgl)
#> [1] TRUE
```

Similar to is.null(), there's is.na().

```
is.na("not NA!")
#> [1] FALSE
is.na(NA)
#> [1] TRUE
```

Recall our variable unsorted_numbers_with_nas.

```
unsorted_numbers_with_nas #> [1] 3 2 10 8 NA 1 4 9 NA 6 5 7
```

Consider the following:

- The predicate functions we've seen so far return either TRUE or FALSE.
- vectors can contain both NA and non-NA values.



Can you guess what is.na() returns?

is.na(unsorted_numbers_with_nas)
#> [1] FALSE FALSE FALSE FALSE TRUE FALSE FALS



We'll discuss accessing a vector's individual elements later, but is.na() is what we call a vectorized function: a function that takes vector argument and operates on every element simultaneously.

2.9 Vectorized Functions

As high-speed R coders, we should prefer vectorized solutions whenever possible as they're not only idiomatic (and thus easy for other R users to understand), but they're typically several orders of magnitude faster than other solutions.

While R isn't the fastest language out there, complaints about its speed usually come from poor code, including code that "speaks" R with a C or Python accent.

The simplest way to wrap our heads around vectorized operations is with math. let's first make a vector with five 0s in it.

We could do that like the following:

c(0, 0, 0, 0, 0) #> [1] 0 0 0 0 0

But, good coders are lazy and want to (correctly) automate everything they can. With that in mind, let's rep()eat 0.5 times.



```
zeros <- rep(0, length = 5)
zeros
#> [1] 0 0 0 0 0
```

For our purposes, the term *scalar* refers to an object that is a single value.

If we want to add 1 (a scalar) to every element of zeros, we can run zeros + 1 or 1 + zeros:

```
zeros + 1
#> [1] 1 1 1 1 1
```

R knows that 1 is a single value (and assumes we know what we're doing) and performs the operation (+) between it and every element of zeros. In R-speak, we refer to this behavior as *recycling*.

Let's see what happens when we add zeros and a vector containing two elements.

```
two_threes <- c(3, 3)

zeros + two_threes
#> Warning in zeros + two_threes: longer object length is not a multiple of shorter object length
#> [1] 3 3 3 3 3 3
```

That's probably not what you expected and R gives us a warning() to tell us something seems wrong.

R let's us get away with a lot of things it shouldn't, which includes

Part II

Day 2



Chapter 3

Tabular Data

- Aliases:
 - Tabular files
 - Flat
 - Delimited
- Includes:
 - Comma-Separated Value (.csv)
 - Tab-Separated Value (.tsv)

3.1 Basics

library(readr)

Here's some example data, modified from http://www.gapminder.org/data/

```
country,continent,year,lifeExp,pop,gdpPercap
Afghanistan,Asia,1952,28.801,8425333,779.4453145
Afghanistan,Asia,1957,30.332,9240934,820.8530296
Afghanistan,Asia,1962,31.997,10267083,853.10071
Afghanistan,Asia,1967,34.02,11537966,836.1971382
Afghanistan,Asia,1972,36.088,13079460,739.9811058
Afghanistan,Asia,1977,38.438,14880372,786.11336
Afghanistan,Asia,1982,39.854,12881816,978.0114388
Afghanistan,Asia,1987,40.822,13867957,852.3959448
```

- # header/column names, separated by commas
- # comma-separated values

```
csv_text <-
'country,continent,year,lifeExp,pop,gdpPercap
Afghanistan,Asia,1952,28.801,8425333,779.4453145
Afghanistan,Asia,1957,30.332,9240934,820.8530296
Afghanistan,Asia,1962,31.997,10267083,853.10071
Afghanistan,Asia,1967,34.02,11537966,836.1971382</pre>
```

Afghanistan, Asia, 1972, 36.088, 13079460, 739.9811058



```
Afghanistan, Asia, 1977, 38.438, 14880372, 786.11336
Afghanistan, Asia, 1982, 39.854, 12881816, 978.0114388
Afghanistan, Asia, 1987, 40.822, 13867957, 852.3959448'

csv_file <- tempfile(fileext = ".csv")
csv_file # a temporary file path
#> [1] "/tmp/RtmpkiXbVj/filec5e2405303c.csv"
writeLines(text = csv_text, con = csv_file) # write `csv_text` to `csv_file`
```

```
read_csv(file = csv_file)
#> Parsed with column specification:
#> cols(
#>
     country = col_character(),
#>
     continent = col_character(),
     year = col_double(),
#>
    lifeExp = col_double(),
#>
    pop = col_double(),
#>
     gdpPercap = col_double()
#> )
#> # A tibble: 8 x 6
#>
     country continent year lifeExp
                                               pop gdpPercap
                                            <db1>
#>
     <chr>
                 \langle chr \rangle \langle dbl \rangle \langle dbl \rangle
                                                        <d.b1.>
                                     28.8 8425333
#> 1 Afghanistan Asia
                            1952
                                                         779.
                            1957
#> 2 Afghanistan Asia
                                  30.3 9240934
                                                         821.
#> 3 Afghanistan Asia
                            1962 32.0 10267083
                                                         853.
#> 4 Afghanistan Asia
                            1967 34.0 11537966
                                                         836.
#> 5 Afghanistan Asia
                            1972
                                     36.1 13079460
                                                         740.
#> 6 Afghanistan Asia
                            1977
                                     38.4 14880372
                                                         786.
                            1982
                                                         978.
#> 7 Afghanistan Asia
                                     39.9 12881816
                             1987
                                                         852.
#> 8 Afghanistan Asia
                                     40.8 13867957
```

You may encounter Tab-Delimited data where values are separated by \t instead of ,. Instead of readr::read_csv(), we can use readr::read_tsv().

```
tsv_text <-
    'country\tcontinent\tyear\tlifeExp\tpop\tgdpPercap
Afghanistan\tAsia\t1952\t28.801\t8425333\t779.4453145
Afghanistan\tAsia\t1957\t30.332\t9240934\t820.8530296
Afghanistan\tAsia\t1962\t31.997\t10267083\t853.10071
Afghanistan\tAsia\t1967\t34.02\t11537966\t836.1971382
Afghanistan\tAsia\t1972\t36.088\t13079460\t739.9811058
Afghanistan\tAsia\t1977\t38.438\t14880372\t786.11336
Afghanistan\tAsia\t1982\t39.854\t12881816\t978.0114388
Afghanistan\tAsia\t1987\t40.822\t13867957\t852.3959448'

tsv_file <- tempfile(fileext = ".tsv")
writeLines(text = tsv_text, con = tsv_file)</pre>
```

```
read_tsv(file = tsv_file)
#> Parsed with column specification:
```



```
#> cols(
    country = col_character(),
#>
    continent = col_character(),
#>
    year = col_double(),
#>
    lifeExp = col_double(),
#>
    pop = col_double(),
#>
    qdpPercap = col_double()
#> )
#> # A tibble: 8 x 6
    country continent year lifeExp
                                          pop gdpPercap
     <chr>
               <chr> <dbl> <dbl>
                                        <db1>
                                                    <db1>
                         1952 28.8 8425333
                                                     779.
#> 1 Afghanistan Asia
                                 30.3 9240934
#> 2 Afghanistan Asia
                          1957
                                                     821.
#> 3 Afghanistan Asia
                         1962 32.0 10267083
                                                     853.
#> 4 Afghanistan Asia
                         1967
                                  34.0 11537966
                                                     836.
#> 5 Afghanistan Asia
                          1972
                                  36.1 13079460
                                                     740.
#> 6 Afghanistan Asia
                           1977
                                  38.4 14880372
                                                     786.
                          1982
                                                     978.
#> 7 Afghanistan Asia
                                  39.9 12881816
#> 8 Afghanistan Asia
                          1987
                                  40.8 13867957
                                                     852.
```

If we find ourselves reading delmited data that uses something other than \t or , to separate values, we can use readr::read_delim().

```
pipe_separated_values_text <-
    'country|continent|year|lifeExp|pop|gdpPercap
Afghanistan|Asia|1952|28.801|8425333|779.4453145
Afghanistan|Asia|1957|30.332|9240934|820.8530296
Afghanistan|Asia|1962|31.997|10267083|853.10071
Afghanistan|Asia|1967|34.02|11537966|836.1971382
Afghanistan|Asia|1972|36.088|13079460|739.9811058
Afghanistan|Asia|1977|38.438|14880372|786.11336
Afghanistan|Asia|1982|39.854|12881816|978.0114388
Afghanistan|Asia|1987|40.822|13867957|852.3959448'</pre>
psv_file <- tempfile(fileext = ".tsv")
writeLines(text = pipe_separated_values_text, con = psv_file)
```

```
read_delim(file = psv_file, delim = "|")
#> Parsed with column specification:
#> cols(
#>
     country = col_character(),
#>
     continent = col_character(),
#>
    year = col_double(),
#>
    lifeExp = col_double(),
#>
    pop = col_double(),
#>
                \dot{} = col\_double()
      qdpPercap
#> )
#> # A tibble: 8 x 6
#>
     country
                 continent year lifeExp
                                              pop `gdpPercap
#>
     <chr>
                <chr> <dbl> <dbl>
                                            <dbl>
                                                              <dbl>
#> 1 Afghanistan Asia
                           1952
                                 28.8 8425333
                                                              779.
```



```
1957
                                     30.3 9240934
                                                                821.
#> 2 Afghanistan Asia
#> 3 Afghanistan Asia
                             1962
                                     32.0 10267083
                                                                853.
#> 4 Afghanistan Asia
                             1967
                                     34.0 11537966
                                                                836.
#> 5 Afghanistan Asia
                             1972
                                     36.1 13079460
                                                                740.
#> 6 Afghanistan Asia
                             1977
                                     38.4 14880372
                                                                786.
#> 7 Afghanistan Asia
                             1982
                                     39.9 12881816
                                                                978.
#> 8 Afghanistan Asia
                             1987
                                                                852.
                                     40.8 13867957
```

```
country,continent,year,lifeExp,pop,gdpPercap Afghanistan,Asia,1952,28.801,8425333,779.4453145 Afghanistan,Asia,1957,30.332,9240934,820.8530296 Afghanistan,Asia,1962,31.997,10267083,853.10071 Afghanistan,Asia,1967,34.02,11537966,836.1971382 Afghanistan,Asia,1972,36.088,13079460,739.9811058 Afghanistan,Asia,1977,38.438,14880372,786.11336 Afghanistan,Asia,1982,39.854,12881816,978.0114388 Afghanistan,Asia,1987,40.822,13867957,852.3959448 Afghanistan,,,N/A,,
```

header/column names

notice that we're missing values

```
csv_text <-
'country,continent,year,lifeExp,pop,gdpPercap
Afghanistan,Asia,1952,28.801,8425333,779.4453145
Afghanistan,Asia,1957,30.332,9240934,820.8530296
Afghanistan,Asia,1962,31.997,10267083,853.10071
Afghanistan,Asia,1967,34.02,11537966,836.1971382
Afghanistan,Asia,1972,36.088,13079460,739.9811058
Afghanistan,Asia,1977,38.438,14880372,786.11336
Afghanistan,Asia,1982,39.854,12881816,978.0114388
Afghanistan,Asia,1987,40.822,13867957,852.3959448
Afghanistan,,N/A,,'</pre>
csv_file <- tempfile(fileext = ".csv")
writeLines(text = csv_text, con = csv_file)
```

3.2 Common Pitfalls

3.2.1 Incorrect Column Types

```
data_frame_from_csv <- read_csv(file = csv_file)
#> Parsed with column specification:
#> cols(
#> country = col_character(),
#> continent = col_character(),
#> year = col_double(),
#> lifeExp = col_character(),
#> pop = col_double(),
#> gdpPercap = col_double()
```



```
#> )
data_frame_from_csv
#> # A tibble: 9 x 6
   country
              continent year lifeExp
                                            pop gdpPercap
                < chr >  < dbl > < chr >
#>
    <chr>
                                          <dbl>
                                                     <dbl>
                         1952 28.801
#> 1 Afghanistan Asia
                                         8425333
                                                     779.
#> 2 Afghanistan Asia
                         1957 30.332 9240934
                                                     821.
#> 3 Afghanistan Asia
                          1962 31.997 10267083
                                                     853.
#> 4 Afghanistan Asia
                          1967 34.02
                                       11537966
                                                     836.
#> 5 Afghanistan Asia
                          1972 36.088 13079460
                                                     740.
#> 6 Afghanistan Asia
                         1977 38.438 14880372
                                                     786.
#> 7 Afghanistan Asia
                          1982 39.854 12881816
                                                     978.
#> 8 Afghanistan Asia
                          1987 40.822 13867957
                                                     852.
#> 9 Afghanistan <NA>
                           NA N/A
                                             NA
                                                      NA
```

Notice that our year column says <dbl>, referring to it being of type double, yet all of our year values are whole numbers.

```
typeof(data_frame_from_csv$year)
#> [1] "double"
data_frame_from_csv$year
#> [1] 1952 1957 1962 1967 1972 1977 1982 1987 NA
```

We also have "N/A" in our lifeExp column, forcing R to interpret all lifeExp values as characters (<chr>).

```
typeof(data_frame_from_csv$lifeExp)
#> [1] "character"
data_frame_from_csv$lifeExp
#> [1] "28.801" "30.332" "31.997" "34.02" "36.088" "38.438" "39.854" "40.822" "N/A"
```

3.2.1.1 Solution

```
read_csv(
 file = csv_file,
  col_types = cols(
    country = col_character(),
    continent = col_character(),
                             # read `year` as `integer`
    year = col_integer(),
    lifeExp = col_double(),
                                 # read `lifeExp` as `double`
    pop = col_double(),
    gdpPercap = col_double()
 ),
 na = c("", "N/A")
                                 # be explicit about how `csv_file` represents missing values
)
#> # A tibble: 9 x 6
     country continent year lifeExp
                                               pop qdpPercap
              < chr >
                           \langle int \rangle \langle dbl \rangle
                                             <dbl>
                                                        <db1>
     <chr>
#> 1 Afghanistan Asia
                            1952
                                     28.8 8425333
                                                         779.
```



7	#> 2 Afghanistan Asia	1957	30.3 9240934	821.
7	#> 3 Afghanistan Asia	1962	32.0 10267083	<i>853</i> .
7	#> 4 Afghanistan Asia	1967	34.0 11537966	836.
7	#> 5 Afghanistan Asia	1972	36.1 13079460	740.
7	#> 6 Afghanistan Asia	1977	38.4 14880372	786.
7	#> 7 Afghanistan Asia	1982	39.9 12881816	978.
7	#> 8 Afghanistan Asia	1987	40.8 13867957	<i>852</i> .
7	#> 9 Afghanistan <na></na>	NA	NA NA	NA

Chapter 4

Manipulating Data Frames

```
library(tidyverse, warn.conflicts = FALSE)
#> -- Attaching packages -----
#> v qqplot2 3.3.2 v purrr 0.3.4
                   v dplyr 1.0.2
#> v tibble 3.0.3
#> v tidyr 1.1.2 v stringr 1.4.0
#> v readr 1.3.1
                   v forcats 0.5.0
#> -- Conflicts -----
                                                                 #> x dplyr::filter() masks stats::filter()
#> x dplyr::lag() masks stats::lag()
df <- tibble(</pre>
 group = c("a", "a", "b", "b", "b"),
 a = c(1, 4, NA, 3, 5),
 b = c(9, NA, 8, 10, 7),
 c = c(TRUE, FALSE, NA, FALSE, TRUE),
 d = c(LETTERS[1:3], NA, LETTERS[[5]]),
 e = factor(1:5, labels = c("tiny", "small", "medium", "big", "huge")),
 f_{col} = c(as.Date(NA), as.Date("2020-09-23") + c(3, 2, 1, 4)),
 g_{col} = c(as.POSIXct("2020-09-23 00:00:00") + 1:4 * 60 * 60 * 24 * 1.1, NA),
 col_h = list(c(1, 10), c(2, NA), c(3, 8), c(4, 7), c(5, 6)),
 col_i = list(NULL, pi, month.abb[6:10], iris, as.matrix(mtcars))
#> # A tibble: 5 x 10
    group a b c d e
                                      f\_col
                                                                   col\_h
                                                 g\_col
                                                                           col i
   <chr> <dbl> <dbl> <lql> <chr> <fct> <date>
                                                 \langle dttm \rangle
                                                                   t>
                 9 TRUE A tiny NA
                                                 2020-09-24 02:24:00 <dbl [2~ <NULL>
#> 1 a 1
#> 2 a
            4 NA FALSE B
                                small 2020-09-26 2020-09-25 04:48:00 <dbl [2~ <dbl [1]>
#> 3 b
           NA
                                medium 2020-09-25 2020-09-26 07:12:00 <dbl [2~ <chr [5]>
                 8 NA C
            3
                10 FALSE <NA> biq 2020-09-24 2020-09-27 09:36:00 <dbl [2~ <df[,5] [150 x ~
#> 5 b
                  7 TRUE E
                                huge 2020-09-27 NA
                                                                   <dbl [2~ <dbl[,11] [32 x~
glimpse(df)
#> Rows: 5
#> Columns: 10
#> $ group <chr> "a", "a", "b", "b", "b"
```

4.1 select() Columns

4.1.1 by Name



4.1.2 by Index



```
df %>%
    select(2:3, everything())
#> # A tibble: 5 x 10

#> a b group c d e f_col g_col col_h col_i
#> <dbl> <dbl> <chr> <lgl> <chr> <lgl> <chr> <lgl> <chr> <lgl> <chr> <ld>  1 9 a TRUE A tiny NA 2020-09-24 02:24:00 <dbl [2~ <NULL> 
#> 3 NA 8 b NA C medium 2020-09-25 2020-09-26 07:12:00 <dbl [2~ <chr [5]> 
#> 4 3 10 b FALSE 
#> 1 B Q20-09-24 2020-09-27 09:36:00 <dbl [2~ <df[,5] [150 x ~ dbl [2~ <dbl [,11] [32 x~ ]]
#> 5 5 7 b TRUE E huge 2020-09-27 NA
```

```
cols_to_select <- c(1, 3, 5)
df %>%
  select(all_of(cols_to_select))
#> # A tibble: 5 x 3
```



```
cols_{to}_{select} \leftarrow c(1, 3, 5, 1000)
df %>%
 select(any_of(cols_to_select))
#> # A tibble: 5 x 3
     group
              b d
     <chr> <dbl> <chr>
#> 1 a
             9 A
#> 2 a
             NAB
#> 3 b
              8 C
#> 4 b
             10 <NA>
#> 5 b
               7 E
```

4.1.3 by Name Pattern

contains() selects a column if any part of its name contains match=.

```
df %>%
 select(contains(match = "col"))
#> # A tibble: 5 x 4
#>
   f\_{\it col}
             g\_col
                                    col\_h
                                            col\_i
    <date>
               < dttm>
                                    < list>
                                              t>
#> 1 NA
              2020-09-24 02:24:00 <dbl [2]> <NULL>
#> 2 2020-09-26 2020-09-25 04:48:00 <dbl [2]> <dbl [1]>
#> 3 2020-09-25 2020-09-26 07:12:00 <dbl [2]> <chr [5]>
#> 4 2020-09-24 2020-09-27 09:36:00 <dbl [2]> <df[,5] [150 x 5]>
#> 5 2020-09-27 NA
                                    <dbl [2]> <dbl[,11] [32 x 11]>
```

starts_with() selects a column if its name starts with match=.

starts_with() selects a column if its name ends with match=.



matches()'s Selects a column if its name matches a regular expression pattern.

```
df %>%
 select(matches("(^\\w_)?col(_\\w)?"))
#> # A tibble: 5 x 4
             g\_col
    f\_{col}
                                     col\_h
                                              col\_i
                                              t>
#> <date>
                \langle dttm \rangle
                                     < list>
                2020-09-24 02:24:00 <dbl [2]> <NULL>
#> 1 NA
#> 2 2020-09-26 2020-09-25 04:48:00 <dbl [2]> <dbl [1]>
#> 3 2020-09-25 2020-09-26 07:12:00 <dbl [2]> <chr [5]>
#> 4 2020-09-24 2020-09-27 09:36:00 <dbl [2]> <df[,5] [150 x 5]>
#> 5 2020-09-27 NA
                                     <dbl [2]> <dbl[,11] [32 x 11]>
```

4.1.4 by Data Type

```
df %>%
    select(where(is.factor))
#> # A tibble: 5 x 1
#> e
#> <fct>
#> 1 tiny
#> 2 small
#> 3 medium
#> 4 big
#> 5 huge
```

```
df %>%
    select_if(is.factor)
#> # A tibble: 5 x 1
#>    e
#> <fct>
#> 1 tiny
#> 2 small
#> 3 medium
#> 4 big
#> 5 huge
```



```
df %>%
    select(where(is.factor), f_col)
#> # A tibble: 5 x 2
#> e    f_col
#> <fct> <date>
#> 1 tiny NA
#> 2 small 2020-09-26
#> 3 medium 2020-09-25
#> 4 big 2020-09-24
#> 5 huge 2020-09-27
```

```
df %>%
  select_if(~ is.character(.x) | is.factor(.x))
#> # A tibble: 5 x 3
#> group d e
```



4.2 filter() Rows

4.2.1 by row_number()

```
df %>%
    filter(row_number() == 1)
#> # A tibble: 1 x 10
#> group a b c d e f_col g_col col_h col_i
#> <chr> <dbl> <dbl> <lgl> <chr> <dbl> <dbl> <lgl> <chr> <dbl> <dbl> <lgl> <chr> < dbl> <dbl > (lgl> <chr> <fct) <date> dtm> (list> tist> #> 1 a 1 9 TRUE A tiny NA 2020-09-24 02:24:00 <dbl [2]> <NULL>
```

4.2.2 by Name

```
df %>%
    filter(a == 2)
#> # A tibble: 0 x 10
#> # ... with 10 variables: group <chr>, a <dbl>, b <dbl>, c <lgl>, d <chr>, e <fct>,
#> # f_col <date>, g_col <dttm>, col_h list>, col_i list>
```



```
df %>%
filter(c)
#> # A tibble: 2 x 10
df %>%
filter(!c)
#> # A tibble: 2 x 10
#> group a b c d e f\_col g\_col col\_h col\_i #> <chr> <math><dbl> <dbl> <dgl> <chr> <math><fct> <date> <dttm> <math>(list> )
3 10 FALSE <NA> big 2020-09-24 2020-09-27 09:36:00 <dbl [2]> <df[,5] [150 x ~
#> 2 b
df %>%
filter(a == 5, d == "E")
#> # A tibble: 1 x 10
#> group a b c d e f_col g_col

#> <chr> <dbl> <dbl> <lgl> <chr> <fct> <date> <dttm>

#> 1 b 5 7 TRUE E huge 2020-09-27 NA
                                               <dbl [2]> <dbl[,11] [32 x~
df %>%
filter(a >= 3 | f_col == "2020-09-24")
#> # A tibble: 3 x 10
#> group a b c d e f\_col g\_col col\_h col\_i #> <chr> <math><dbl> <dbl> <lgl> <math><chr> <fct> <date> <dttm>
3 10 FALSE <NA> big 2020-09-24 2020-09-27 09:36:00 <dbl [2]> <df[,5] [150 x \sim
#> 2 b
         5 7 TRUE E huge 2020-09-27 NA  <dbl [2] > <dbl [,11] [32 x^{2}] 
df %>%
filter(a < 2 | c)
#> # A tibble: 2 x 10
<dbl [2]> <dbl[,11] [32 x~
filter(!is.na(a), !is.na(b), !is.na(d))
#> # A tibble: 2 x 10
```



4.2.3 by Type

```
df %>%
    filter(across(where(is.numeric), ~ .x >= 5))
#> # A tibble: 1 x 10
#> group a b c d e f_col g_col col_h col_i
#> <chr> <dbl> <dbl> <dbl> <dbl> <dgl> <chr> <dbl> <dbl> <dbl> description of the col of t
```

```
df %>%
    filter_if(is.numeric, ~ .x >= 5)
#> # A tibble: 1 x 10
#> group a b c d e f_col g_col col_h col_i
#> <chr> <dbl> <dbl> <dbl> <dp> <chr> <dbl> <dbl> <dbl> <dfc> <date> <dttm> <dbl (2]> <dbl (11] [32 x~</p>
```

4.3 arrange() Rows





Part III Spatial Data



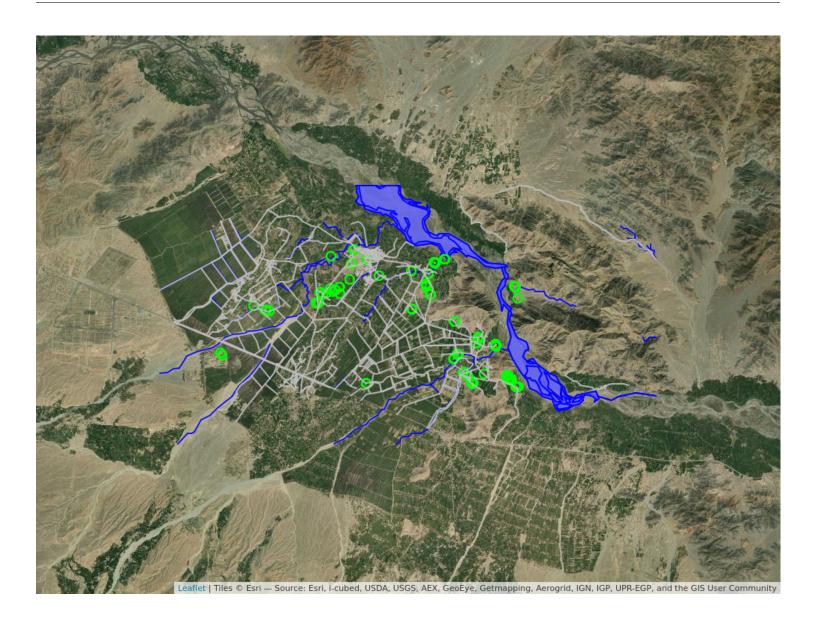
35

Chapter 5

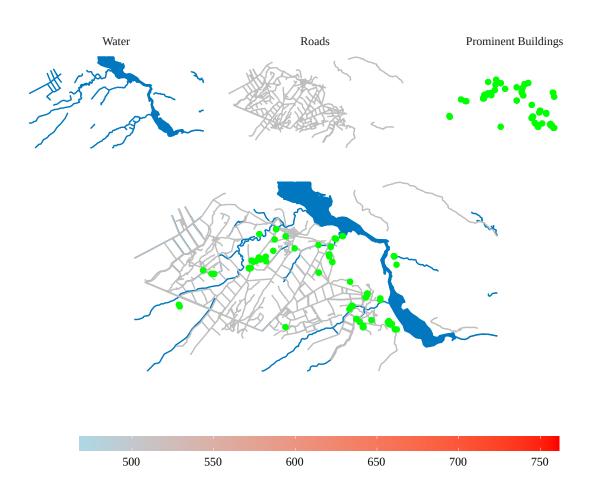
Types of Spatial Data

```
packages <- c(
 "leaflet", # interactive web mapping
  "osmdata", # Open Street Maps API data
  \hbox{"raster", } \hbox{\it\# obtaining administrative boundary data and spatial raster data handling}
           # spatial vector data handling
 "sf",
 "stars", # {sf}'s spatio-temporal raster counterpart
 "tidyverse" # data manipulation
install.packages(
 packages[!sapply(packages, requireNamespace, quietly = TRUE)]
library(leaflet)
library(sf)
#> Linking to GEOS 3.7.1, GDAL 2.4.2, PROJ 5.2.0
library(stars)
#> Loading required package: abind
library(tidyverse)
#> -- Attaching packages -----
#> v ggplot2 3.3.2 v purrr 0.3.4
#> v tibble 3.0.3 v dplyr 1.0.2
#> v tidyr 1.1.2 v stringr 1.4.0
#> v readr 1.3.1 v forcats 0.5.0
#> -- Conflicts -----
                                                                    #> x dplyr::filter() masks stats::filter()
#> x dplyr::lag() masks stats::lag()
```







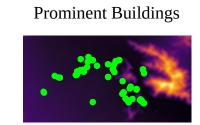


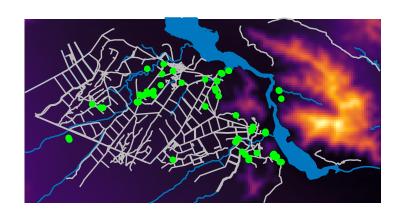
489	487	485	499	543	584	585	551	518	501	499	503	502	508	532	549	535
527	525	521	530	565	598	595	564	535	519	515	518	516	527	558	575	550
569	582	588	597	618	639	639	618	591	570	557	556	561	574	593	590	549
576	619	641	649	661	678	691	690	673	648	627	622	633	643	629	589	535
553	608	632	630	635	656	685	709	716	710	705	707	711	694	638	567	513
520	559	569	558	559	583	621	665	705	736	757	761	747	699	617	539	498
492	509	511	504	507	530	565	617	682	735	755	741	711	662	591	529	495
478	482	483	484	492	514	552	608	677	720	709	664	626	598	562	527	500
475	476	477	481	489	516	564	625	677	688	645	582	544	532	524	514	501
471	474	478	482	487	510	561	615	641	622	573	522	495	492	495	497	497
471	473	478	486	491	508	547	581	579	548	514	492	482	482	483	486	492
485	480	483	495	509	526	549	556	534	500	482	480	480	480	481	483	488
525	521	515	516	525	536	543	533	508	482	474	478	481	481	482	484	485
557	565	551	530	521	522	522	513	494	478	473	477	480	481	482	483	485
534	549	537	512	502	509	517	515	497	477	471	476	479	480	482	482	483
486	498	492	479	481	501	523	529	507	478	468	475	480	481	482	482	482





Water Roads





Chapter 6

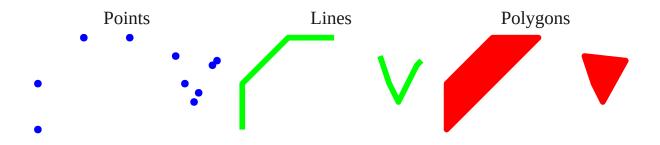
Vector Geometries

Spatial vector data represent the world as a collection of points which, for two-dimensional data, are stored as x and y coordinates.

```
suppressPackageStartupMessages(library(tidyverse))
library(sf)
#> Linking to GEOS 3.7.1, GDAL 2.4.2, PROJ 5.2.0
```

Points can be joined in order to make lines, which themselves can be joined to make polygons.





```
practice_coords <- tibble(</pre>
  lng = c(-20, -20, -10, -10, 20, 20, 10, 10),
  lat = c(-20, 10, 20, -10, -10, 10, 20, -20),
  lab = c("A", "B", "C", "D", "E", "F", "G", "H"),
  grp = c("a", "a", "a", "a", "b", "b", "b", "b")
practice_coords
#> # A tibble: 8 x 4
       lng lat lab
                        grp
#>
     <dbl> <dbl> <chr> <chr>
#> 1
       -20
             -20 A
#> 2
       -20
              10 B
#> 3
       -10
              20 C
#> 4
       -10
             -10 D
                        \boldsymbol{a}
#> 5
        20
             -10 E
                        b
#> 6
        20
             10 F
                        b
#> 7
        10
              20 G
                        b
#> 8
        10
             -20 H
```

6.1 POINT

POINT refers to the location of a single point in space.

Here, we use st_as_sf() to convert a regular data.frame into an sf object.

• Steps:

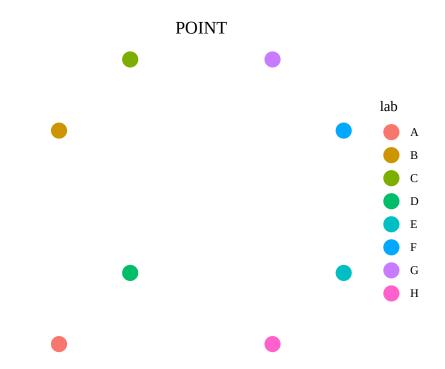


- 1. take practice_coords
- 2. convert to sf object with $st_as_sf()$, providing a character vector indicating the names of practice_coords in (x,y) / (longitude, latitude) order
- 3. mutate() to a add a column named shape, which we obtain using st_geometry_type().

```
point_sf <- practice_coords %>%
                                                     # Step 1.
  st_as_sf(coords = c(x = "lng", y ="lat")) %>%
                                                     # 2.
                                                     # 3.
  mutate(shape = st_geometry_type(geometry))
point_sf
#> Simple feature collection with 8 features and 3 fields
#> geometry type: POINT
#> dimension:
                   XY
                   xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> bbox:
#> CRS:
#> # A tibble: 8 x 4
     lab grp geometry shape
#> * <chr> <chr> <POINT> <fct>
                (-20 -20) POINT
#> 1 A
        \boldsymbol{a}
#> 2 B
                (-20 10) POINT
#> 3 C
                 (-10 20) POINT
         \boldsymbol{a}
#> 4 D
          \boldsymbol{a}
                (-10 -10) POINT
         b
#> 5 E
                (20 -10) POINT
#> 6 F
         b
                  (20 10) POINT
#> 7 G
           b
                   (10 20) POINT
#> 8 H
                  (10 -20) POINT
```

The data in our lng and lat columns are moved to a new geometry column.

```
ggplot(data = point_sf) +
  geom_sf(aes(color = lab), size = 5, show.legend = "point") +
  labs(title = "POINT")
```



6.2 MULTIPOINT

MULTIPOINT refers to a collection of POINTs.

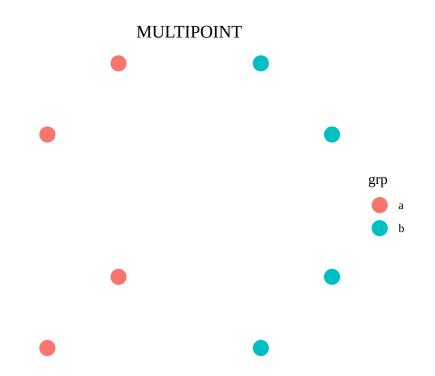
- Steps:
 - 1. take point_sf
 - 2. using group_by(), group the rows together based on the values in their grp column
 - 3. summarise() each group, which combines the points of each group into a MULTIPOINT
 - 4. mutate() the shape column to change it to the new st_geometry_type()

```
multi_point_sf <- point_sf %>%
                                              # Step 1.
                                              # 2.
 group_by(grp) %>%
  summarise() %>%
 mutate(shape = st_geometry_type(geometry)) # 4.
   `summarise()` ungrouping output (override with `.groups` argument)
multi_point_sf
#> Simple feature collection with 2 features and 2 fields
#> geometry type: MULTIPOINT
#> dimension:
                   XY
#> bbox:
                   xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> CRS:
#> # A tibble: 2 x 3
                                              geometry shape
     grp
                                          <MULTIPOINT> <fct>
#> * <chr>
#> 1 a
           ((-20 -20), (-20 10), (-10 -10), (-10 20)) MULTIPOINT
               ((10 -20), (10 20), (20 -10), (20 10)) MULTIPOINT
#> 2 b
```



Instead of the 8 separate POINTs with which we started, we now have 2 rows of MULTIPOINTs, each of which contain 4 points.

```
ggplot(data = multi_point_sf) +
geom_sf(aes(color = grp), size = 5, show.legend = "point") +
labs(title = "MULTIPOINT")
```



6.3 LINESTRING

LINESTRING is how we represent individual lines.

- Steps:
 - 1. take multi_point_sf
 - 2. cast the geometry to= LINESTRING using st_cast() 3 mutate() the shape column to change it to the new st_geometry_type()



```
#> grp shape geometry

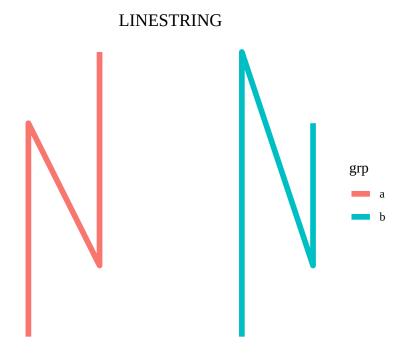
#> * <chr> <fct> <LINESTRING>

#> 1 a LINESTRING (-20 -20, -20 10, -10 -10, -10 20)

#> 2 b LINESTRING (10 -20, 10 20, 20 -10, 20 10)
```

Now we have 2 rows that each contain a LINESTRING, which was built by connecting each point to the next.

```
ggplot(data = linestring_sf) +
  geom_sf(aes(color = grp), size = 2, show.legend = "line") +
  labs(title = "LINESTRING")
```



6.4 MULTILINESTRING

Similar to MULTIPOINTs that contain multiple points, we also have MULTILINESTRINGS.

- Steps:
 - 1. take linestring_sf
 - 2. summarise() the rows, combining them all into a single MULTILINESTRING
 - 3. mutate() the shape column to change it to the new st_geometry_type() and replace the grp column that is dropped when we summarise() without using group_by()

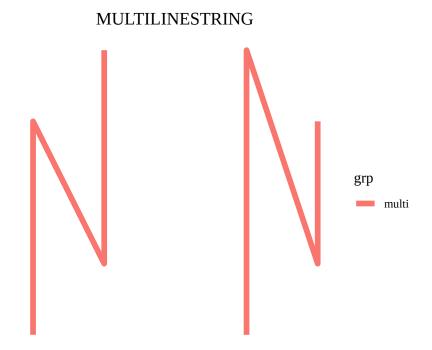
```
multi_linestring_sf <- linestring_sf %>%  # Step 1.
summarise() %>%  # 2.
mutate(shape = st_geometry_type(geometry), # 3.
```



```
grp = "multi")
                                              # 4.
multi_linestring_sf
#> Simple feature collection with 1 feature and 2 fields
#> geometry type: MULTILINESTRING
#> dimension:
                   XY
#> bbox:
                   xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> CRS:
#> # A tibble: 1 x 3
                                                                  geometry shape
                                                                                            grp
#> *
                                                         <MULTILINESTRING> <fct>
                                                                                            <chr>
#> 1 ((-20 -20, -20 10, -10 -10, -10 20), (10 -20, 10 20, 20 -10, 20 10)) MULTILINESTRING multi
```

Now we have 2 lines embedded inside a single MULTILINESTRING row.

```
ggplot(data = multi_linestring_sf) +
  geom_sf(aes(color = grp), size = 2, show.legend = "line") +
  labs(title = "MULTILINESTRING")
```



6.5 POLYGON

POLYGONs are essentially sets of lines that close to form a ring, although POLGYONs can also contain holes. We can easily wrap a shape around any geometry using st_convex_hull() to form a convex hull polygon.

- Steps:
 - 1. take point_sf

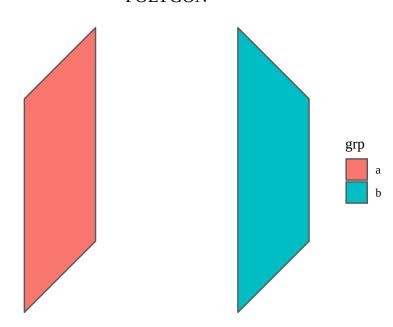


- 2. using group_by(), group the rows together based on the values in their grp column
- 3. summarise() each group, combining them into MULTIPOINTs
- 4. wrap the MULTIPOINTs in a polygon using st_convex_hull()
- 5. mutate() the shape column to change it to the new st_geometry_type()

```
polygon_sf <- point_sf %>%
                                              # Step 1.
  group_by(grp) %>%
                                              # 2.
  summarise() %>%
                                              # 3.
  st_convex_hull() %>%
                                              # 4.
  mutate(shape = st_geometry_type(geometry)) # 5.
  `summarise()` ungrouping output (override with `.groups` argument)
polygon_sf
#> Simple feature collection with 2 features and 2 fields
#> geometry type: POLYGON
#> dimension:
                   XY
#> bbox:
                   xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> CRS:
#> # A tibble: 2 x 3
   grp
                                                geometry shape
#> * <chr>
                                                <POLYGON> <fct>
          ((-20 -20, -20 10, -10 20, -10 -10, -20 -20)) POLYGON
                ((10 -20, 10 20, 20 10, 20 -10, 10 -20)) POLYGON
#> 2 b
```

```
ggplot(data = polygon_sf) +
  geom_sf(aes(fill = grp), show.legend = "polygon") +
  labs(title = "POLYGON")
```

POLYGON



6.6 MULTIPOLYGON

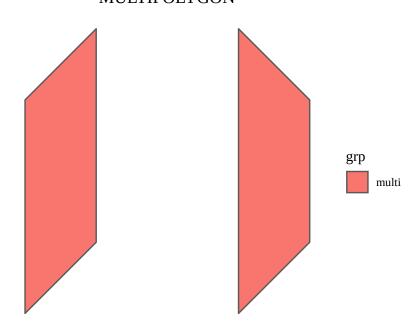
POLYGONs can also be grouped together to form MULTIPOLYGONs.

- Steps:
 - 1. take polygon_sf
 - 2. summarise() the rows, combining them all into a single MULTILIPOLYGON
 - 3. mutate() the shape column to change it to the new st_geometry_type() and replace the grp column that is dropped when we summarise() without using group_by()

```
multi_polygon_sf <- polygon_sf %>%
                                                 # Step 1.
  summarise() %>%
                                                 # 2.
  mutate(shape = st_geometry_type(geometry), # 3.
         grp = "multi")
multi_polygon_sf
#> Simple feature collection with 1 feature and 2 fields
#> geometry type: MULTIPOLYGON
#> dimension:
                   XY
#> bbox:
                    xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> CRS:
#> # A tibble: 1 x 3
#>
                                                                               geometry shape
#>
                                                                         <\!\!\mathit{MULTIPOLYGON}\!\!> <\!\!\mathit{fct}\!\!>
                                                                                                     <chr>
#> 1 (((-20 -20, -20 10, -10 20, -10 -10, -20 -20)), ((10 -20, 10 20, 20 10, 20 ~ MULTIPOLY~ multi
```

```
ggplot(data = multi_polygon_sf) +
  geom_sf(aes(fill = grp), show.legend = "polygon") +
  labs(title = "MULTIPOLYGON")
```

MULTIPOLYGON





6.7 GEOMETRY

GEOMETRY is a special geometry type. It refers to a column of mixed geometries, i.e. we have multiple geometry types in our geometry column.

```
geometry_sf <- list(point_sf, multi_point_sf, linestring_sf,</pre>
                    multi_linestring_sf, polygon_sf, multi_polygon_sf) %>%
 map_if(~ "lab" %in% names(.x), select, -lab) %>%
  do.call(what = rbind) %>%
  mutate(grp = if_else(shape == "POINT", as.character(row_number()), grp))
geometry_sf
#> Simple feature collection with 16 features and 2 fields
#> geometry type: GEOMETRY
#> dimension:
                   XY
#> bbox:
                   xmin: -20 ymin: -20 xmax: 20 ymax: 20
#> CRS:
#> # A tibble: 16 x 3
     qrp
                                                                              geometry shape
#>
   * <chr>
                                                                            <GEOMETRY> <fct>
   1 1
                                                                      POINT (-20 -20) POINT
                                                                       POINT (-20 10) POINT
#> 22
                                                                       POINT (-10 20) POINT
#> 33
#> 4 4
                                                                      POINT (-10 -10) POINT
#> 5 5
                                                                       POINT (20 -10) POINT
#> 6 6
                                                                        POINT (20 10) POINT
#> 77
                                                                        POINT (10 20) POINT
                                                                       POINT (10 -20) POINT
#> 88
#>
  9 a
                                MULTIPOINT ((-20 -20), (-20 10), (-10 -10), (-10 20)) MULTIPOINT
#> 10 b
                                    MULTIPOINT ((10 -20), (10 20), (20 -10), (20 10)) MULTIPOINT
#> 11 a
                                        LINESTRING (-20 -20, -20 10, -10 -10, -10 20) LINESTRING
                                            LINESTRING (10 -20, 10 20, 20 -10, 20 10) LINESTRING
#> 13 multi MULTILINESTRING ((-20 -20, -20 10, -10 -10, -10 20), (10 -20, 10 20, 20 ~ MULTILINEST~
#> 14 a
                                POLYGON ((-20 -20, -20 10, -10 20, -10 -10, -20 -20)) POLYGON
#> 15 b
                                     POLYGON ((10 -20, 10 20, 20 10, 20 -10, 10 -20)) POLYGON
#> 16 multi MULTIPOLYGON (((-20 -20, -20 10, -10 20, -10 -10, -20 -20)), ((10 -20, 1~ MULTIPOLYGON
```

```
ggplot(data = geometry_sf) +
geom_sf(aes(color = grp, fill = grp), size = 2, show.legend = FALSE) +
facet_wrap(~ shape, nrow = 2) +
labs(title = "GEOMETRY")
```



