

Reading in data with **readr** and **haven**

Data Wrangling: Session 6

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Statistical Horizons, April 2022

Load the packages, as always

```
library(here)      # manage file paths
```

```
## here() starts at /Users/kjhealy/Documents/courses/data_wrangling
```

```
library(socviz)    # data and some useful functions
```

```
##  
## Attaching package: 'socviz'  
## The following object is masked from 'package:kjhutils':  
##  
##      %nin%
```

```
library(tidyverse) # your friend and mine
```

```
## — Attaching packages ————— tidyverse 1.3.1 —
```

```
## ✓ ggplot2 3.3.5      ✓ purrr   0.3.4  
## ✓ tibble  3.1.6      ✓ dplyr  1.0.8  
## ✓ tidyr   1.2.0      ✓ stringr 1.4.0  
## ✓ readr   2.1.2      ✓ forcats 0.5.1
```

```
## — Conflicts ————— tidyverse_conflicts() —
```

```
## x readr::edition_get() masks testthat::edition_get()  
## x dplyr::filter()      masks stats::filter()  
## x purrr::is_null()     masks testthat::is_null()  
## x dplyr::lag()         masks stats::lag()  
## x readr::local_edition() masks testthat::local_edition()  
## x dplyr::matches()     masks tidyrr::matches(), testthat::matches()
```

```
library(haven)     # for Stata, SAS, and SPSS files
```

We've put a lot of pieces in place at this point

Including several things we haven't fully exploited yet

Data we want to get into R

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Nice, clean CSV files.

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Quite messy things like tables on web pages.

Data we want to get into R

Nice, clean CSV files.

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Other plain-text formats.

Foreign formats, like Stata.

Quite messy things like tables on web pages.

... and more besides.

Reading in CSV files

CSV is not really a proper format at all!

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Base R has `read.csv()`

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Base R has `read.csv()`

Corresponding tidyverse "underscored" version: `read_csv()`.

It is pickier and more talkative than the Base R version.

Where's the data? Using **here()**

If we're loading a file, it's coming from *somewhere*.

If it's on our local disk somewhere, we will need to interact with the file system. We should try to do this in a way that avoids *absolute* file paths.

```
# This is not portable  
df <- read_csv("/Users/kjhealy/Documents/data/misc/project/data/mydata.csv")
```

Where's the data? Using **here()**

If we're loading a file, it's coming from *somewhere*.

If it's on our local disk somewhere, we will need to interact with the file system. We should try to do this in a way that avoids *absolute* file paths.

```
# This is not portable  
df <- read_csv("/Users/kjhealy/Documents/data/misc/project/data/mydata.csv")
```

We should also do it in a way that is *platform independent*.

This makes it easier to share your work, move it around, etc. Projects should be self-contained.

Where's the data? Using **here()**

The `here` package, and **`here()`** function builds paths relative to the top level of your R project.

```
here() # this path will be different for you
```

```
## [1] "/Users/kjhealy/Documents/courses/data_wrangling"
```


Where's the data? Using **here()**

This seminar's files all live in an RStudio project. It looks like this:

```
## /Users/kjhealy/Documents/courses/data_wrangling
## |— LICENSE
## |— README.Rmd
## |— README.md
## |— code
## |— course_notes.Rmd
## |— data
## |— data_wrangling.Rproj
## |— docs
## |— office
## |— pdf_slides
## |— scratch.Rmd
## |— slides
## |— tests
```

I want to load files from the `data` folder, but I also want *you* to be able to load them. I'm writing this from somewhere deep in the `slides` folder, but you won't be there. Also, I'm on a Mac, but you may not be.

Where's the data? Using **here()**

So:

```
## Load the file relative to the path from the top of the project, without separators, etc  
organs <- read_csv(file = here("data", "organdonation.csv"))
```

Where's the data? Using **here()**

So:

```
## Load the file relative to the path from the top of the project, without separators, etc
organs <- read_csv(file = here("data", "organdonation.csv"))
```

organs

```
## # A tibble: 238 × 21
##   country year donors  pop pop.dens  gdp gdp.lag health health.lag pubhealth
##   <chr>   <dbl> <dbl> <dbl>   <dbl> <dbl> <dbl> <dbl>   <dbl>   <dbl>
## 1 Austra...  NA  NA   17065    0.220 16774 16591 1300    1224     4.8
## 2 Austra... 1991 12.1 17284    0.223 17171 16774 1379    1300     5.4
## 3 Austra... 1992 12.4 17495    0.226 17914 17171 1455    1379     5.4
## 4 Austra... 1993 12.5 17667    0.228 18883 17914 1540    1455     5.4
## 5 Austra... 1994 10.2 17855    0.231 19849 18883 1626    1540     5.4
## 6 Austra... 1995 10.2 18072    0.233 21079 19849 1737    1626     5.5
## 7 Austra... 1996 10.6 18311    0.237 21923 21079 1846    1737     5.6
## 8 Austra... 1997 10.3 18518    0.239 22961 21923 1948    1846     5.7
## 9 Austra... 1998 10.5 18711    0.242 24148 22961 2077    1948     5.9
## 10 Austra... 1999  8.67 18926    0.244 25445 24148 2231    2077     6.1
## # ... with 228 more rows, and 11 more variables: roads <dbl>, cerebvas <dbl>,
## #   assault <dbl>, external <dbl>, txp.pop <dbl>, world <chr>, opt <chr>,
## #   consent.law <chr>, consent.practice <chr>, consistent <chr>, ccode <chr>
```

And there it is.

Where's the data? Using **here()**

Get in the habit of putting this at the top of your files:

```
here::i_am("analysis.Rmd") # or whatever your Rmd or R file is called
```

See [the here project page](#) for more details.

`read_csv()` comes in different varieties

`read_csv()` Field separator is a comma: **,**

```
organs <- read_csv(file = here("data", "organdonation.csv"))
```

`read_csv2()` Field separator is a semicolon: **;**

```
# Example only  
my_data <- read_csv2(file = here("data", "my_euro_file.csv"))
```

Both are special cases of `read_delim()`

Other species are also catered to

`read_tsv()` Tab separated.

`read_fwf()` Fixed-width files.

`read_log()` Log files (i.e. computer log files).

`read_lines()` Just read in lines, without trying to parse them.

Also often useful ...

`read_table()`

Data that's separated by one (or more) columns of space.

You can read files remotely, too

You can give all of these functions local files, or they can point to URLs.

Compressed files will be automatically uncompressed.

(Be careful what you download from remote locations!)

```
organ_remote <- read_csv("http://kjhealy.co/organdonation.csv")
```

```
organ_remote
```

```
## # A tibble: 238 × 21
##   country year donors  pop pop.dens  gdp gdp.lag health health.lag pubhealth
##   <chr>   <dbl> <dbl> <dbl>    <dbl> <dbl>   <dbl>   <dbl>    <dbl>    <dbl>
## 1 Austr...   NA   NA   17065    0.220 16774  16591   1300    1224     4.8
## 2 Austr... 1991  12.1  17284    0.223 17171  16774   1379    1300     5.4
## 3 Austr... 1992  12.4  17495    0.226 17914  17171   1455    1379     5.4
## 4 Austr... 1993  12.5  17667    0.228 18883  17914   1540    1455     5.4
## 5 Austr... 1994  10.2  17855    0.231 19849  18883   1626    1540     5.4
## 6 Austr... 1995  10.2  18072    0.233 21079  19849   1737    1626     5.5
## 7 Austr... 1996  10.6  18311    0.237 21923  21079   1846    1737     5.6
## 8 Austr... 1997  10.3  18518    0.239 22961  21923   1948    1846     5.7
## 9 Austr... 1998  10.5  18711    0.242 24148  22961   2077    1948     5.9
## 10 Austr... 1999   8.67 18926    0.244 25445  24148   2231    2077     6.1
## # ... with 228 more rows, and 11 more variables: roads <dbl>, cerebvas <dbl>,
## #   assault <dbl>, external <dbl>, txp.pop <dbl>, world <chr>, opt <chr>,
## #   consent.law <chr>, consent.practice <chr>, consistent <chr>, ccode <chr>
```


An example: `read_table()`

England and Wales, Total Population, Death rates (period 1x1), Last modified: 02 Apr 2018; Methods Protocol: v6 (2017)

Year	Age	Female	Male	Total
1841	0	0.136067	0.169189	0.152777
1841	1	0.059577	0.063208	0.061386
1841	2	0.036406	0.036976	0.036689
1841	3	0.024913	0.026055	0.025480
1841	4	0.018457	0.019089	0.018772
1841	5	0.013967	0.014279	0.014123
1841	6	0.010870	0.011210	0.011040
1841	7	0.008591	0.008985	0.008788
1841	8	0.006860	0.007246	0.007053
1841	9	0.005772	0.006050	0.005911
1841	10	0.005303	0.005382	0.005343
1841	11	0.005114	0.005002	0.005057
1841	12	0.005145	0.004856	0.004999
1841	13	0.005455	0.004955	0.005202

1841	105	0.576987	1.727848	0.700373
1841	106	0.677711	6.000000	0.795287
1841	107	0.900000	.	0.900000
1841	108	1.388430	.	1.388430
1841	109	.	.	.
1841	110+	.	.	.
1842	0	0.148491	0.184007	0.166481
1842	1	0.063038	0.066596	0.064818
1842	2	0.035203	0.035854	0.035527

An example: `read_table()`

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Year	Age	Female	Male	Total
1841	0	0.136067	0.169189	0.152777
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1841	4	0.018457	0.019089	0.018772
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1841	6	0.010870	0.011210	0.011040
1841	7	0.008591	0.008985	0.008788
1841	8	0.006860	0.007246	0.007053
1841	9	0.005772	0.006050	0.005911
1841	10	0.005303	0.005382	0.005343
1841	11	0.005114	0.005002	0.005057
1841	12	0.005145	0.004856	0.004999
1841	13	0.005455	0.004955	0.005202

1841	105	0.576987	1.727848	0.700373
1841	106	0.677711	6.000000	0.795287
1841	107	0.900000	.	0.900000
1841	108	1.388430	.	1.388430
1841	109	.	.	.
1841	110+	.	.	.
1842	0	0.148491	0.184007	0.166481
1842	1	0.063038	0.066596	0.064818
1842	2	0.035203	0.035854	0.035527

```
engmort <- read_table(here("data", "mortality.txt"),  
                      skip = 2, na = ".")
```

```
engmort
```

```
## # A tibble: 222 × 5  
##   Year Age   Female   Male   Total  
##   <dbl> <chr>   <dbl>   <dbl> <dbl>  
## 1  1841 0     0.136   0.169   0.153  
## 2  1841 1     0.0596  0.0632  0.0614  
## 3  1841 2     0.0364  0.0370  0.0367  
## 4  1841 3     0.0249  0.0261  0.0255  
## 5  1841 4     0.0185  0.0191  0.0188  
## 6  1841 5     0.0140  0.0143  0.0141  
## 7  1841 6     0.0109  0.0112  0.0110  
## 8  1841 7     0.00859 0.00898 0.00879  
## 9  1841 8     0.00686 0.00725 0.00705  
## 10 1841 9     0.00577 0.00605 0.00591  
## # ... with 212 more rows
```

Attend to the **column specification**

```
engmort <- read_table(here("data", "mortality.txt"),  
                      skip = 2, na = ".")
```

```
##  
## — Column specification —————  
## cols(  
##   Year = col_double(),  
##   Age = col_character(),  
##   Female = col_double(),  
##   Male = col_double(),  
##   Total = col_double()  
## )
```

The column specification tells you what the read function did. That is, how it interpreted each of the columns. It will also report if things don't go as expected.

Attend to the **column specification**

```
engmort <- read_table(here("data", "mortality.txt"),  
                      skip = 2, na = ".")
```

```
##  
## — Column specification —————  
## cols(  
##   Year = col_double(),  
##   Age = col_character(),  
##   Female = col_double(),  
##   Male = col_double(),  
##   Total = col_double()  
## )
```

The column specification tells you what the read function did. That is, how it interpreted each of the columns. It will also report if things don't go as expected.

Why is age imported in character format?

Attend to the **column specification**

Absent you giving them a column specification, the `read_` functions try to *guess* what the type of each column is. They do this by looking at the first thousand rows of each column. They may guess incorrectly!

Normalizing names and recoding

```
read_table(here("data", "mortality.txt"),  
           skip = 2, na = ".")
```

```
## # A tibble: 222 × 5  
##   Year Age   Female   Male   Total  
##   <dbl> <chr>   <dbl>   <dbl>   <dbl>  
## 1  1841 0     0.136   0.169   0.153  
## 2  1841 1     0.0596  0.0632  0.0614  
## 3  1841 2     0.0364  0.0370  0.0367  
## 4  1841 3     0.0249  0.0261  0.0255  
## 5  1841 4     0.0185  0.0191  0.0188  
## 6  1841 5     0.0140  0.0143  0.0141  
## 7  1841 6     0.0109  0.0112  0.0110  
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## 10 1841 9     0.00577 0.00605 0.00591  
## # ... with 212 more rows
```

Normalizing names and recoding

```
read_table(here("data", "mortality.txt"),  
           skip = 2, na = ".") %>%  
  janitor::clean_names()
```

```
## # A tibble: 222 × 5  
##   year age   female   male   total  
##   <dbl> <chr>   <dbl>   <dbl>   <dbl>  
## 1  1841 0     0.136   0.169   0.153  
## 2  1841 1     0.0596  0.0632  0.0614  
## 3  1841 2     0.0364  0.0370  0.0367  
## 4  1841 3     0.0249  0.0261  0.0255  
## 5  1841 4     0.0185  0.0191  0.0188  
## 6  1841 5     0.0140  0.0143  0.0141  
## 7  1841 6     0.0109  0.0112  0.0110  
## 8  1841 7     0.00859 0.00898 0.00879  
## 9  1841 8     0.00686 0.00725 0.00705  
## 10 1841 9     0.00577 0.00605 0.00591  
## # ... with 212 more rows
```

Normalizing names and recoding

```
read_table(here("data", "mortality.txt"),  
           skip = 2, na = ".") %>%  
  janitor::clean_names() %>%  
  mutate(age = as.integer(recode(age, "110+" = "110")))
```

```
## # A tibble: 222 × 5  
##   year   age female   male  total  
##   <dbl> <int>   <dbl>   <dbl> <dbl>  
## 1  1841     0  0.136  0.169  0.153  
## 2  1841     1  0.0596  0.0632  0.0614  
## 3  1841     2  0.0364  0.0370  0.0367  
## 4  1841     3  0.0249  0.0261  0.0255  
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## 6  1841     5  0.0140  0.0143  0.0141  
## 7  1841     6  0.0109  0.0112  0.0110  
## 8  1841     7  0.00859 0.00898 0.00879  
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## # ... with 212 more rows
```


Normalizing names and recoding

```
read_table(here("data", "mortality.txt"),
           skip = 2, na = ".") %>%
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```

```
## # A tibble: 222 × 5
##   year   age female   male  total
##   <dbl> <int>   <dbl>   <dbl> <dbl>
## 1  1841     0  0.136  0.169  0.153
## 2  1841     1  0.0596  0.0632  0.0614
## 3  1841     2  0.0364  0.0370  0.0367
## 4  1841     3  0.0249  0.0261  0.0255
## 5  1841     4  0.0185  0.0191  0.0188
## 6  1841     5  0.0140  0.0143  0.0141
## 7  1841     6  0.0109  0.0112  0.0110
## 8  1841     7  0.00859  0.00898  0.00879
## 9  1841     8  0.00686  0.00725  0.00705
## 10 1841     9  0.00577  0.00605  0.00591
## # ... with 212 more rows
```

The `janitor` package is very handy!

The main cost of normalizing names comes with, e.g., data where there is a codebook you need to consult. But in general it's worth it.

More on column specifications

CDC/NCHS data: **Provisional COVID-19 Death Counts by Sex, Age, and State**



Centers for Disease Control and Prevention
CDC 24/7: Saving Lives. Protecting People.™

Data.CDC.gov

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Sign In

Provisional COVID-19 Death Counts by Sex, Age, and State

NCHS

View Data

Visualize ▾

Export

API

...

Deaths involving coronavirus disease 2019 (COVID-19), pneumonia, and influenza reported to NCHS by sex and age group and state.

NOTICE TO USERS: As of September 2, 2020, this data file includes the following age groups

[More](#)

Updated

April 14, 2021

Data Provided by

National Center for Health Statistics

More on column specifications

What's in this Dataset?

Rows	Columns
52.3K	16

Columns in this Dataset

Column Name	Description	Type	
Data As Of	Date of analysis	Date & Time 𐀀	▼
Start Date	First date of data period	Date & Time 𐀀	▼
End Date	Last date of data period	Date & Time 𐀀	▼
Group	Indicator of whether data measured by Month, by Year, or ...	Plain Text T	▼
Year	Year in which death occurred	Number #	▼
Month	Month in which death occurred	Number #	▼

Let's try to load it

```
nchs <- with_edition(1, read_csv(here("data", "SAS_on_2021-04-13.csv")))
```

```
## Warning: 88128 parsing failures.
```

```
##   row  col                expected actual                                     file
## 2755 Year 1/0/T/F/TRUE/FALSE    2020  '/Users/kjhealy/Documents/courses/data_wrangling/data/SAS_on_2021-04-13.csv'
## 2756 Year 1/0/T/F/TRUE/FALSE    2020  '/Users/kjhealy/Documents/courses/data_wrangling/data/SAS_on_2021-04-13.csv'
## 2757 Year 1/0/T/F/TRUE/FALSE    2020  '/Users/kjhealy/Documents/courses/data_wrangling/data/SAS_on_2021-04-13.csv'
## 2758 Year 1/0/T/F/TRUE/FALSE    2020  '/Users/kjhealy/Documents/courses/data_wrangling/data/SAS_on_2021-04-13.csv'
## 2759 Year 1/0/T/F/TRUE/FALSE    2020  '/Users/kjhealy/Documents/courses/data_wrangling/data/SAS_on_2021-04-13.csv'
## .....
## See problems(...) for more details.
```

Let's try to load it

```
problems(nchs)
```

```
## # A tibble: 88,128 × 5
##   row col   expected      actual file
##   <int> <chr> <chr>      <chr> <chr>
## 1  2755 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 2  2756 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 3  2757 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 4  2758 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 5  2759 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 6  2760 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 7  2761 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 8  2762 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 9  2763 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 10 2764 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## # ... with 88,118 more rows
```

Let's try to load it

```
problems(nchs)
```

```
## # A tibble: 88,128 × 5
##   row col   expected      actual file
##   <int> <chr> <chr>      <chr> <chr>
## 1  2755 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 2  2756 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 3  2757 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 4  2758 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 5  2759 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 6  2760 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 7  2761 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 8  2762 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 9  2763 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 10 2764 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## # ... with 88,118 more rows
```

Problems are stored as an attribute of the nchs object, so we can revisit them.

Let's try to load it

```
problems(nchs)
```

```
## # A tibble: 88,128 × 5
##   row col expected actual file
##   <int> <chr> <chr>      <chr> <chr>
## 1  2755 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 2  2756 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 3  2757 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 4  2758 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 5  2759 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 6  2760 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 7  2761 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 8  2762 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 9  2763 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## 10 2764 Year 1/0/T/F/TRUE/FALSE 2020 '/Users/kjhealy/Documents/courses/data...'
## # ... with 88,118 more rows
```

Problems are stored as an attribute of the `nchs` object, so we can revisit them.

Parsing failures tend to cascade. Our data only has 56k rows but we got 88k failures.

Take a look with **head()**

```
head(nchs)
```

```
## # A tibble: 6 × 16
##   `Data As Of` `Start Date` `End Date` Group Year  Month State Sex   `Age Group`
##   <chr>        <chr>        <chr>    <chr> <lgl> <lgl> <chr> <chr> <chr>
## 1 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... All Ages
## 2 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... Under 1 ye...
## 3 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... 0-17 years
## 4 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... 1-4 years
## 5 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... 5-14 years
## 6 04/07/2021   01/01/2020   04/03/2021 By T... NA    NA    Unit... All ... 15-24 years
## # ... with 7 more variables: `COVID-19 Deaths` <dbl>, `Total Deaths` <dbl>,
## #   `Pneumonia Deaths` <dbl>, `Pneumonia and COVID-19 Deaths` <dbl>,
## #   `Influenza Deaths` <dbl>, `Pneumonia, Influenza, or COVID-19 Deaths` <dbl>,
## #   Footnote <chr>
```


Take a look with `tail()`

```
tail(nchs)
```

```
## # A tibble: 6 × 16
##   `Data As Of` `Start Date` `End Date` Group Year  Month State Sex   `Age Group`
##   <chr>        <chr>        <chr>    <chr> <lgl> <lgl> <chr> <chr> <chr>
## 1 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 45-54 years
## 2 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 50-64 years
## 3 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 55-64 years
## 4 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 65-74 years
## 5 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 75-84 years
## 6 04/07/2021   04/01/2021   04/03/2021 By M... NA    NA    Puer... Fema... 85 years a...
## # ... with 7 more variables: `COVID-19 Deaths` <dbl>, `Total Deaths` <dbl>,
## #   `Pneumonia Deaths` <dbl>, `Pneumonia and COVID-19 Deaths` <dbl>,
## #   `Influenza Deaths` <dbl>, `Pneumonia, Influenza, or COVID-19 Deaths` <dbl>,
## #   Footnote <chr>
```

Take a look with `slice_sample()`

```
nchs %>%  
  slice_sample(n = 10)
```

```
## # A tibble: 10 × 16  
##   `Data As Of` `Start Date` `End Date` Group   Year Month State      Sex  
##   <chr>        <chr>        <chr>    <chr>   <lgl> <lgl> <chr>    <chr>  
## 1 04/07/2021  04/01/2020  04/30/2020 By Month NA    NA    Tennessee All ...  
## 2 04/07/2021  11/01/2020  11/30/2020 By Month NA    NA    Arizona    All ...  
## 3 04/07/2021  10/01/2020  10/31/2020 By Month NA    NA    Connecticut Male  
## 4 04/07/2021  01/01/2021  01/31/2021 By Month NA    TRUE  District of ... All ...  
## 5 04/07/2021  08/01/2020  08/31/2020 By Month NA    NA    Maine       Male  
## 6 04/07/2021  01/01/2020  01/31/2020 By Month NA    TRUE  Connecticut Fema...  
## 7 04/07/2021  04/01/2021  04/03/2021 By Month NA    NA    Puerto Rico Male  
## 8 04/07/2021  03/01/2021  03/31/2021 By Month NA    NA    Idaho       Male  
## 9 04/07/2021  12/01/2020  12/31/2020 By Month NA    NA    Ohio        Fema...  
## 10 04/07/2021  04/01/2021  04/03/2021 By Month NA    NA    Oregon      Fema...  
## # ... with 8 more variables: `Age Group` <chr>, `COVID-19 Deaths` <dbl>,  
## #   `Total Deaths` <dbl>, `Pneumonia Deaths` <dbl>,  
## #   `Pneumonia and COVID-19 Deaths` <dbl>, `Influenza Deaths` <dbl>,  
## #   `Pneumonia, Influenza, or COVID-19 Deaths` <dbl>, Footnote <chr>
```

Aside: one that happened earlier ...

```
nchs %>%  
  slice_sample(n = 10)
```

```
## # A tibble: 10 x 16
```

##		`Data As Of`	`Start Date`	`End Date`	Group	Year	Month	State	Sex
##		<chr>	<chr>	<chr>	<chr>	<lgl>	<lgl>	<chr>	<chr>
##	1	04/07/2021	01/01/2020	04/03/2021	By Tot...	NA	NA	Minnesota	Male
##	2	04/07/2021	02/01/2020	02/29/2020	By Mon...	NA	NA	Georgia	Male
##	3	04/07/2021	02/01/2021	02/28/2021	By Mon...	NA	NA	Maine	Male
##	4	04/07/2021	11/01/2020	11/30/2020	By Mon...	NA	NA	New Jersey	Female
##	5	04/07/2021	01/01/2020	12/31/2020	By Year	NA	NA	Rhode Island	All Se...
##	6	04/07/2021	01/01/2020	01/31/2020	By Mon...	NA	TRUE	New York	All Se...
##	7	04/07/2021	05/01/2020	05/31/2020	By Mon...	NA	NA	District of...	Male
##	8	04/07/2021	04/01/2021	04/03/2021	By Mon...	NA	NA	North Carol...	Female
##	9	04/07/2021	03/01/2021	03/31/2021	By Mon...	NA	NA	Kentucky	Male
##	10	04/07/2021	04/01/2021	04/03/2021	By Mon...	NA	NA	New Mexico	Female

```
## # ... with 8 more variables: Age Group <chr>, COVID-19 Deaths <dbl>,  
## #   Total Deaths <dbl>, Pneumonia Deaths <dbl>,  
## #   Pneumonia and COVID-19 Deaths <dbl>, Influenza Deaths <dbl>,  
## #   Pneumonia, Influenza, or COVID-19 Deaths <dbl>, Footnote <chr>
```

Take a look with `slice()`

Let's look at the rows `read_csv()` complained about.

```
nchs %>%  
  slice(2750:2760)
```

```
## # A tibble: 11 × 16  
##   `Data As Of` `Start Date` `End Date` Group   Year Month State      Sex  
##   <chr>        <chr>        <chr>    <chr>   <lgl> <lgl> <chr>    <chr>  
## 1 04/07/2021  01/01/2020  04/03/2021 By Total NA    NA    Puerto Rico Fema...  
## 2 04/07/2021  01/01/2020  04/03/2021 By Total NA    NA    Puerto Rico Fema...  
## 3 04/07/2021  01/01/2020  04/03/2021 By Total NA    NA    Puerto Rico Fema...  
## 4 04/07/2021  01/01/2020  04/03/2021 By Total NA    NA    Puerto Rico Fema...  
## 5 04/07/2021  01/01/2020  04/03/2021 By Total NA    NA    Puerto Rico Fema...  
## 6 04/07/2021  01/01/2020  12/31/2020 By Year  NA    NA    United States All ...  
## 7 04/07/2021  01/01/2020  12/31/2020 By Year  NA    NA    United States All ...  
## 8 04/07/2021  01/01/2020  12/31/2020 By Year  NA    NA    United States All ...  
## 9 04/07/2021  01/01/2020  12/31/2020 By Year  NA    NA    United States All ...  
## 10 04/07/2021 01/01/2020 12/31/2020 By Year  NA    NA    United States All ...  
## 11 04/07/2021 01/01/2020 12/31/2020 By Year  NA    NA    United States All ...  
## # ... with 8 more variables: `Age Group` <chr>, `COVID-19 Deaths` <dbl>,  
## #   `Total Deaths` <dbl>, `Pneumonia Deaths` <dbl>,  
## #   `Pneumonia and COVID-19 Deaths` <dbl>, `Influenza Deaths` <dbl>,  
## #   `Pneumonia, Influenza, or COVID-19 Deaths` <dbl>, Footnote <chr>
```

Take a look with `slice()`

```
nchs %>%  
  slice(2750:2760) %>%  
  select(Year, Month, State)
```

```
## # A tibble: 11 × 3  
##   Year  Month State  
##   <lgl> <lgl> <chr>  
## 1 NA    NA    Puerto Rico  
## 2 NA    NA    Puerto Rico  
## 3 NA    NA    Puerto Rico  
## 4 NA    NA    Puerto Rico  
## 5 NA    NA    Puerto Rico  
## 6 NA    NA    United States  
## 7 NA    NA    United States  
## 8 NA    NA    United States  
## 9 NA    NA    United States  
## 10 NA   NA    United States  
## 11 NA   NA    United States
```

Hm, something to do with the transition to national numbers maybe?

Take a look with `select()` and `filter()`

```
nchs %>%  
  select(Year, Month, State) %>%  
  filter(State == "New York")
```

```
## # A tibble: 969 × 3  
##   Year Month State  
##   <lgl> <lgl> <chr>  
## 1 NA     NA     New York  
## 2 NA     NA     New York  
## 3 NA     NA     New York  
## 4 NA     NA     New York  
## 5 NA     NA     New York  
## 6 NA     NA     New York  
## 7 NA     NA     New York  
## 8 NA     NA     New York  
## 9 NA     NA     New York  
## 10 NA    NA     New York  
## # ... with 959 more rows
```

Take a look with `is.na()`

```
nchs %>%  
  select(Year, Month, State) %>%  
  filter(!is.na(Year))
```

```
## # A tibble: 0 × 3  
## # ... with 3 variables: Year <lgl>, Month <lgl>, State <chr>
```

It really has been read in as a completely empty column.

That doesn't seem like it can be right.

Take a look with `distinct()`

```
nchs %>%  
  select(Year) %>%  
  distinct(Year)
```

```
## # A tibble: 1 × 1  
##   Year  
##   <lgl>  
## 1 NA
```

Again, it's been read in as a completely empty column.

Take a look with `read_lines()`

Time to reach for a different kitchen knife.

```
read_lines(here("data", "SAS_on_2021-04-13.csv"), n_max = 10)
```

```
## [1] "Data As Of,Start Date,End Date,Group,Year,Month,State,Sex,Age Group,COVID-19 Deaths,Total Deaths,Pneumonia Deaths,Pneumoni
## [2] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,All Ages,539723,4161167,466437,263147,9037,750804,"
## [3] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,Under 1 year,59,22626,246,10,21,316,"
## [4] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,0-17 years,251,39620,667,46,179,1051,"
## [5] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,1-4 years,31,4069,137,5,61,224,"
## [6] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,5-14 years,89,6578,195,19,76,341,"
## [7] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,15-24 years,804,42596,930,317,81,1493,"
## [8] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,18-29 years,1996,75339,2184,884,150,3434,"
## [9] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,25-34 years,3543,88196,3493,1617,237,5638,"
## [10] "04/07/2021,01/01/2020,04/03/2021,By Total,,,United States,All Sexes,30-39 years,5792,107348,5276,2658,318,8706,"
```

We can get the whole thing this way

```
raw_file <- read_lines(here("data", "SAS_on_2021-04-13.csv"))
```

This imports the data as a long, long character vector, with each element being a line.

```
# reminder: indexing 1D vectors  
letters[5:6]
```

```
## [1] "e" "f"
```

Now we're just looking at lines in a file

```
# This is not a tibble; we have to index it the basic way  
raw_file[2753:2758]
```

```
## [1] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,65-74 years,203,2650,410,151,,466,One or more data cells hav  
## [2] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,75-84 years,234,4274,656,154,16,751,"  
## [3] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,85 years and over,222,6164,795,136,29,909,"  
## [4] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,All Ages,380949,3372967,349667,178222,8779,560025,"  
## [5] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,Under 1 year,48,19356,224,9,21,284,"  
## [6] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,0-17 years,189,33808,598,35,178,930,"
```

Now we're just looking at lines in a file

```
# This is not a tibble; we have to index it the basic way  
raw_file[2753:2758]
```

```
## [1] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,65-74 years,203,2650,410,151,,466,One or more data cells hav  
## [2] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,75-84 years,234,4274,656,154,16,751,"  
## [3] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,85 years and over,222,6164,795,136,29,909,"  
## [4] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,All Ages,380949,3372967,349667,178222,8779,560025,"  
## [5] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,Under 1 year,48,19356,224,9,21,284,"  
## [6] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,0-17 years,189,33808,598,35,178,930,"
```

There you are, you bastard.

Now we're just looking at lines in a file

```
# This is not a tibble; we have to index it the basic way
raw_file[2753:2758]
```

```
## [1] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,65-74 years,203,2650,410,151,,466,One or more data cells hav
## [2] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,75-84 years,234,4274,656,154,16,751,"
## [3] "04/07/2021,01/01/2020,04/03/2021,By Total,,,Puerto Rico,Female,85 years and over,222,6164,795,136,29,909,"
## [4] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,All Ages,380949,3372967,349667,178222,8779,560025,"
## [5] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,Under 1 year,48,19356,224,9,21,284,"
## [6] "04/07/2021,01/01/2020,12/31/2020,By Year,2020,,United States,All Sexes,0-17 years,189,33808,598,35,178,930,"
```

There you are, you bastard.

In this case, this is due to the kind of data this is, mixing multiple reporting levels and totals. That is, it's not a mistake in the *data*, but rather in the *parsing*.

OK, let's go back to the colspec!

```
nchs <- with_edition(1, read_csv(here("data", "SAS_on_2021-04-13.csv")))
```

```
##  
## — Column specification —————  
## cols(  
##   `Data As Of` = col_character(),  
##   `Start Date` = col_character(),  
##   `End Date` = col_character(),  
##   Group = col_character(),  
##   Year = col_logical(),  
##   Month = col_logical(),  
##   State = col_character(),  
##   Sex = col_character(),  
##   `Age Group` = col_character(),  
##   `COVID-19 Deaths` = col_double(),  
##   `Total Deaths` = col_double(),  
##   `Pneumonia Deaths` = col_double(),  
##   `Pneumonia and COVID-19 Deaths` = col_double(),  
##   `Influenza Deaths` = col_double(),  
##   `Pneumonia, Influenza, or COVID-19 Deaths` = col_double(),  
##   Footnote = col_character()  
## )
```

We can just copy it from the console output! It's valid code.

We use it with **col_types**

```
nchs <- with_edition(1, read_csv(here("data", "SAS_on_2021-04-13.csv"),
  col_types = cols(
    `Data As Of` = col_character(),
    `Start Date` = col_character(),
    `End Date` = col_character(),
    Group = col_character(),
    Year = col_logical(),
    Month = col_logical(),
    State = col_character(),
    Sex = col_character(),
    `Age Group` = col_character(),
    `COVID-19 Deaths` = col_double(),
    `Total Deaths` = col_double(),
    `Pneumonia Deaths` = col_double(),
    `Pneumonia and COVID-19 Deaths` = col_double(),
    `Influenza Deaths` = col_double(),
    `Pneumonia, Influenza, or COVID-19 Deaths` = col_double(),
    Footnote = col_character()
  )))
```

But we know we need to make some adjustments.

Fixes

```
# Date format
us_style <- "%m/%d/%Y"

nchs <- with_edition(1, read_csv(
  here("data", "SAS_on_2021-04-13.csv"),
  col_types = cols(
    `Data As Of` = col_date(format = us_style),
    `Start Date` = col_date(format = us_style),
    `End Date` = col_date(format = us_style),
    Group = col_character(),
    Year = col_character(),
    Month = col_character(),
    State = col_character(),
    Sex = col_character(),
    `Age Group` = col_character(),
    `COVID-19 Deaths` = col_integer(),
    `Total Deaths` = col_integer(),
    `Pneumonia Deaths` = col_integer(),
    `Pneumonia and COVID-19 Deaths` = col_integer(),
    `Influenza Deaths` = col_integer(),
    `Pneumonia, Influenza, or COVID-19 Deaths` = col_integer(),
    Footnote = col_character()
  )) %>%
janitor::clean_names() %>%
select(-footnote) %>%
mutate(age_group = stringr::str_to_sentence(age_group)) %>%
filter(!stringr::str_detect(state, "Total"))
)
```


Now let's look again

```
dim(nchs)
```

```
## [1] 52326    15
```

```
nchs %>%  
  select(year, month, state) %>%  
  filter(!is.na(year))
```

```
## # A tibble: 49,572 × 3  
##   year month state  
##   <chr> <chr> <chr>  
## 1 2020 <NA> United States  
## 2 2020 <NA> United States  
## 3 2020 <NA> United States  
## 4 2020 <NA> United States  
## 5 2020 <NA> United States  
## 6 2020 <NA> United States  
## 7 2020 <NA> United States  
## 8 2020 <NA> United States  
## 9 2020 <NA> United States  
## 10 2020 <NA> United States  
## # ... with 49,562 more rows
```

Now let's look again

```
nchs %>%  
  distinct(year)
```

```
## # A tibble: 3 × 1  
##   year  
##   <chr>  
## 1 <NA>  
## 2 2020  
## 3 2021
```

Lessons learned

Lessons learned

I said at the start that it was no fun, but also weirdly satisfying.

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When `read_csv()` warns you of a parsing failure, **don't ignore it**.

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When `read_csv()` warns you of a parsing failure, **don't ignore it**.

`read_lines()` lets you get the file in a nearly unprocessed form.

Lessons learned

I said at the start that it was no fun, but also weirdly satisfying.

When `read_csv()` warns you of a parsing failure, **don't ignore it**.

`read_lines()` lets you get the file in a nearly unprocessed form.

The `colspec` output is your friend.

If we wanted to ...

```
library(stringr) # it's back!
```


If we wanted to ...

```
library(stringr) # it's back!
```

```
nchs
```

```
## # A tibble: 52,326 × 15
##   data_as_of start_date end_date   group   year month state
##   <date>      <date>    <date>   <chr>   <chr> <chr> <chr>
## 1 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 2 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 3 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 4 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 5 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 6 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 7 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 8 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 9 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## 10 2021-04-07 2020-01-01 2021-04-03 By Total <NA> <NA> United
## # ... with 52,316 more rows, and 6 more variables: covid_19_deaths
## #   total_deaths <int>, pneumonia_deaths <int>,
## #   pneumonia_and_covid_19_deaths <int>, influenza_deaths <int>
## #   pneumonia_influenza_or_covid_19_deaths <int>
```

If we wanted to ...

```
library(stringr) # it's back!
```

```
nchs %>%
```

```
  select(!c(data_as_of:end_date, year, month))
```

```
## # A tibble: 52,326 × 10
```

```
##   group      state sex  age_group covid_19_deaths total_deaths
```

```
##   <chr>      <chr> <chr> <chr>          <int>         <int>
```

```
## 1 By Total Unite... All ... All ages          539723        4161167
```

```
## 2 By Total Unite... All ... Under 1 ...           59          22626
```

```
## 3 By Total Unite... All ... 0-17 yea...          251         39620
```

```
## 4 By Total Unite... All ... 1-4 years           31          4069
```

```
## 5 By Total Unite... All ... 5-14 yea...           89          6578
```

```
## 6 By Total Unite... All ... 15-24 ye...          804         42596
```

```
## 7 By Total Unite... All ... 18-29 ye...         1996         75339
```

```
## 8 By Total Unite... All ... 25-34 ye...         3543         88196
```

```
## 9 By Total Unite... All ... 30-39 ye...         5792        107348
```

```
## 10 By Total Unite... All ... 35-44 ye...         9259        126848
```

```
## # ... with 52,316 more rows, and 3 more variables:
```

```
## #   pneumonia_and_covid_19_deaths <int>, influenza_deaths <int>
```

```
## #   pneumonia_influenza_or_covid_19_deaths <int>
```

If we wanted to ...

```
library(stringr) # it's back!
```

```
nchs %>%
```

```
  select(!(c(data_as_of:end_date, year, month))) %>%
```

```
  pivot_longer(covid_19_deaths:pneumonia_influenza_or_covid_19  
               names_to = "outcome",  
               values_to = "n")
```

```
## # A tibble: 313,956 × 6
```

```
##   group      state      sex      age_group      outcome
```

```
##   <chr>      <chr>      <chr>      <chr>      <chr>
```

```
## 1 By Total United States All Sexes All ages covid_19_death
```

```
## 2 By Total United States All Sexes All ages total_deaths
```

```
## 3 By Total United States All Sexes All ages pneumonia_deat
```

```
## 4 By Total United States All Sexes All ages pneumonia_and_
```

```
## 5 By Total United States All Sexes All ages influenza_deat
```

```
## 6 By Total United States All Sexes All ages pneumonia_infl
```

```
## 7 By Total United States All Sexes Under 1 year covid_19_death
```

```
## 8 By Total United States All Sexes Under 1 year total_deaths
```

```
## 9 By Total United States All Sexes Under 1 year pneumonia_deat
```

```
## 10 By Total United States All Sexes Under 1 year pneumonia_and_
```

```
## # ... with 313,946 more rows
```

If we wanted to ...

```
library(stringr) # it's back!
```

```
nchs %>%
```

```
  select(!c(data_as_of:end_date, year, month)) %>%
```

```
  pivot_longer(covid_19_deaths:pneumonia_influenza_or_covid_19  
               names_to = "outcome",  
               values_to = "n") %>%
```

```
  mutate(outcome = str_to_sentence(outcome),  
         outcome = str_replace_all(outcome, "_", " "),  
         outcome = str_replace(outcome, "(C|c)ovid 19", "COVID
```

```
## # A tibble: 313,956 × 6
```

```
##   group      state      sex      age_group      outcome
```

```
##   <chr>      <chr>      <chr>      <chr>      <chr>
```

```
## 1 By Total United States All Sexes All ages COVID-19 death
```

```
## 2 By Total United States All Sexes All ages Total deaths
```

```
## 3 By Total United States All Sexes All ages Pneumonia death
```

```
## 4 By Total United States All Sexes All ages Pneumonia and
```

```
## 5 By Total United States All Sexes All ages Influenza death
```

```
## 6 By Total United States All Sexes All ages Pneumonia infl
```

```
## 7 By Total United States All Sexes Under 1 year COVID-19 death
```

```
## 8 By Total United States All Sexes Under 1 year Total deaths
```

```
## 9 By Total United States All Sexes Under 1 year Pneumonia death
```

```
## 10 By Total United States All Sexes Under 1 year Pneumonia and
```

```
## # ... with 313,946 more rows
```

If we wanted to ...

```
library(stringr) # it's back!
```

```
nchs %>%  
  select(!c(data_as_of:end_date, year, month)) %>%  
  pivot_longer(covid_19_deaths:pneumonia_influenza_or_covid_19  
    names_to = "outcome",  
    values_to = "n") %>%  
  mutate(outcome = str_to_sentence(outcome),  
    outcome = str_replace_all(outcome, "_", " "),  
    outcome = str_replace(outcome, "(C|c)ovid 19", "COVID
```

```
## # A tibble: 313,956 × 6  
##   group      state      sex      age_group outcome  
##   <chr>    <chr>    <chr>    <chr>    <chr>  
## 1 By Total United States All Sexes All ages COVID-19 death  
## 2 By Total United States All Sexes All ages Total deaths  
## 3 By Total United States All Sexes All ages Pneumonia death  
## 4 By Total United States All Sexes All ages Pneumonia and  
## 5 By Total United States All Sexes All ages Influenza death  
## 6 By Total United States All Sexes All ages Pneumonia infl  
## 7 By Total United States All Sexes Under 1 year COVID-19 death  
## 8 By Total United States All Sexes Under 1 year Total deaths  
## 9 By Total United States All Sexes Under 1 year Pneumonia death  
## 10 By Total United States All Sexes Under 1 year Pneumonia and  
## # ... with 313,946 more rows
```

Put this in `nchs_fmt`

... we could make a table or graph

```
nchs_fmt %>%  
  select(state, age_group, outcome, n)
```

```
## # A tibble: 313,956 × 4  
##   state      age_group outcome      n  
##   <chr>      <chr>    <chr>    <int>  
## 1 United States All ages COVID-19 deaths 539723  
## 2 United States All ages Total deaths 4161167  
## 3 United States All ages Pneumonia deaths 466437  
## 4 United States All ages Pneumonia and COVID-19 deaths 263147  
## 5 United States All ages Influenza deaths 9037  
## 6 United States All ages Pneumonia influenza or COVID-19 deaths 750804  
## 7 United States Under 1 year COVID-19 deaths 59  
## 8 United States Under 1 year Total deaths 22626  
## 9 United States Under 1 year Pneumonia deaths 246  
## 10 United States Under 1 year Pneumonia and COVID-19 deaths 10  
## # ... with 313,946 more rows
```

Cleaned up (but not tidy)

```
nchs_fmt %>%  
  distinct(group)
```

```
## # A tibble: 3 × 1  
##   group  
##   <chr>  
## 1 By Total  
## 2 By Year  
## 3 By Month
```

Cleaned up (but not tidy)

```
nchs_fmt %>%  
  distinct(group)
```

```
## # A tibble: 3 × 1  
##   group  
##   <chr>  
## 1 By Total  
## 2 By Year  
## 3 By Month
```

```
nchs_fmt %>%  
  distinct(age_group)
```

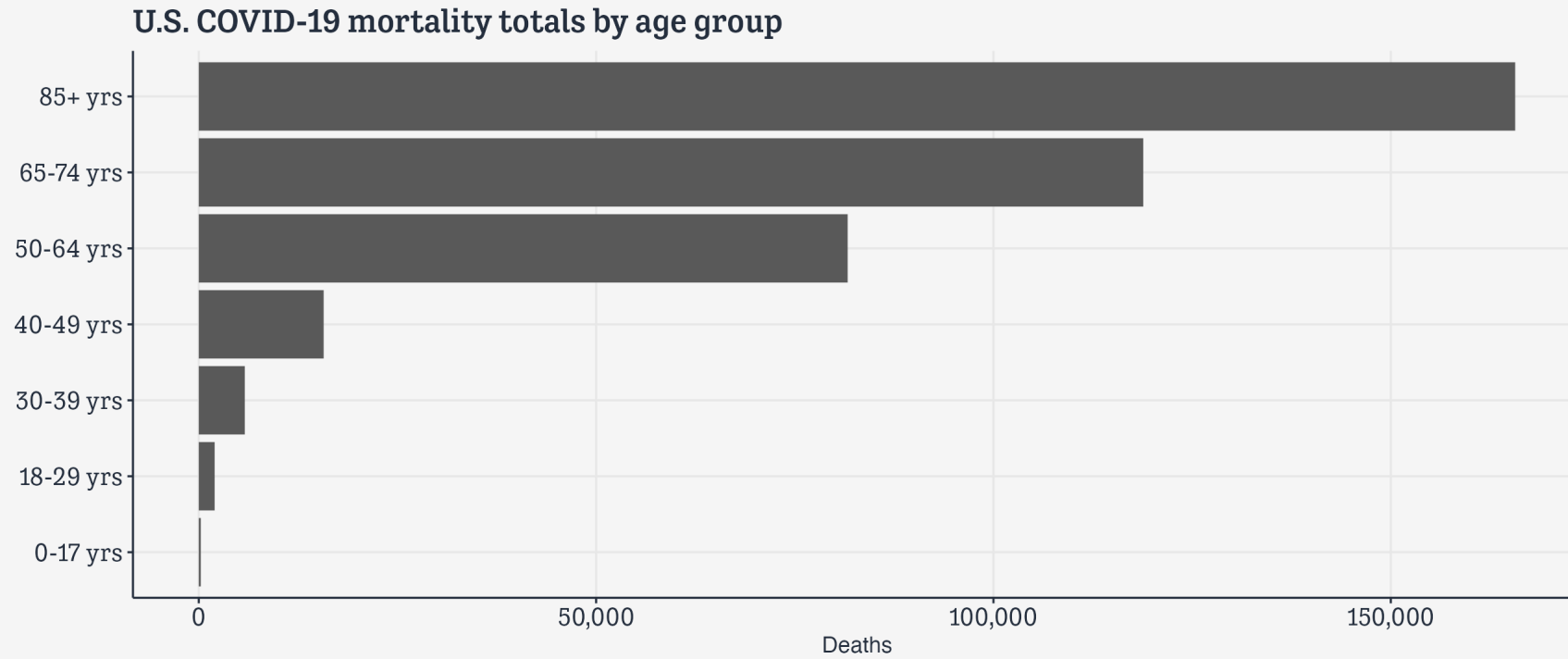
```
## # A tibble: 17 × 1  
##   age_group  
##   <chr>  
## 1 All ages  
## 2 Under 1 year  
## 3 0-17 years  
## 4 1-4 years  
## 5 5-14 years  
## 6 15-24 years  
## 7 18-29 years  
## 8 25-34 years  
## 9 30-39 years  
## 10 35-44 years  
## 11 40-49 years  
## 12 45-54 years  
## 13 50-64 years  
## 14 55-64 years  
## 15 65-74 years  
## 16 75-84 years  
## 17 85 years and over
```


Make our plot

```
p_out <- nchs_fmt %>%  
  filter(group %in% "By Total",  
         sex %in% "All Sexes",  
         state %in% "United States",  
         age_group %in% c("0-17 years",  
                           "18-29 years",  
                           "30-39 years",  
                           "40-49 years",  
                           "50-64 years",  
                           "65-74 years",  
                           "85 years and over"),  
         outcome %in% "COVID-19 deaths") %>%  
  mutate(age_group = str_replace(age_group, "years", "yrs"),  
         age_group = str_replace(age_group, " and over", ""),  
         age_group = str_replace(age_group, "85", "85+")) %>%  
  ggplot(mapping = aes(x = n, y = age_group)) +  
  geom_col() + scale_x_continuous(labels = scales::comma) +  
  labs(x = "Deaths", y = NULL, title = "U.S. COVID-19 mortality totals by age group")
```

Result

p_out



**Every dataset is
different**

Dropping missing values: a quick demo

```
df <- tribble(  
  ~a, ~b, ~c,  
  1, NA, 2,  
  NA, NA, NA,  
  2, 2, 2  
)
```

df

```
## # A tibble: 3 × 3  
##       a     b     c  
##   <dbl> <dbl> <dbl>  
## 1     1    NA     2  
## 2    NA    NA    NA  
## 3     2     2     2
```

Dropping **missing values**: a quick demo

```
# 1 Write it yourself
df %>%
  filter(complete.cases(.))
```

```
## # A tibble: 1 × 3
##       a       b       c
##   <dbl> <dbl> <dbl>
## 1     2     2     2
```

```
# 2 Convenience function
df %>%
  drop_na()
```

```
## # A tibble: 1 × 3
##       a       b       c
##   <dbl> <dbl> <dbl>
## 1     2     2     2
```

Both drop all rows with *any* missing cases.

Dropping **missing values**: a quick demo

What if we only want to drop all rows with *all* missing cases?

```
# 3
df %>%
  filter(!if_all(everything(), ~ is.na(.x))) # Pronoun
```

```
## # A tibble: 2 × 3
##       a         b         c
##   <dbl> <dbl> <dbl>
## 1     1     NA     2
## 2     2     2     2
```

```
# 4 Convenience function from janitor
df %>%
  janitor::remove_empty("rows")
```

```
## # A tibble: 2 × 3
##       a         b         c
##   <dbl> <dbl> <dbl>
## 1     1     NA     2
## 2     2     2     2
```

Read `. x` as "the column we're looking at" or "the variable we're computing on right now".

Cleaning a table

With that in mind ... Some marketing data

SEGMENT	DESCRIPTION	R	F	M
Champions	Bought recently, buy often and spend the most	4- 5	4- 5	4- 5
Loyal Customers	Spend good money. Responsive to promotions	2- 5	3- 5	3- 5
Potential Loyalist	Recent customers, spent good amount, bought more than once	3- 5	1- 3	1- 3
New Customers	Bought more recently, but not often	4- 5	<= 1	<= 1
Promising	Recent shoppers, but haven't spent much	3- 4	<= 1	<= 1
Need Attention	Above average recency, frequency & monetary values	2- 3	2- 3	2- 3
About To Sleep	Below average recency, frequency & monetary values	2- 3	<= 2	<= 2
At Risk	Spent big money, purchased often but long time ago	<= 2	2- 5	2- 5
Can't Lose Them	Made big purchases and often, but long time ago	<= 1	4- 5	4- 5
Hibernating	Low spenders, low frequency, purchased long time ago	1- 2	1- 2	1- 2
Lost	Lowest recency, frequency & monetary scores	<= 2	<= 2	<= 2

Cleaning a table

```
read_csv(here("data", "rfm_table.csv"))
```

```
## # A tibble: 23 × 5
##   SEGMENT      DESCRIPTION      R      F      M
##   <chr>      <chr>      <chr> <chr> <chr>
## 1 <NA>      <NA>      <NA> <NA> <NA>
## 2 Champions Bought recently, buy often and spend th... 4- 5 4- 5 4- 5
## 3 <NA>      <NA>      <NA> <NA> <NA>
## 4 Loyal Customers Spend good money. Responsive to promoti... 2- 5 3- 5 3- 5
## 5 <NA>      <NA>      <NA> <NA> <NA>
## 6 Potential Loyalist Recent customers, spent good amount, bo... 3- 5 1- 3 1- 3
## 7 <NA>      <NA>      <NA> <NA> <NA>
## 8 New Customers Bought more recently, but not often      4- 5 <= 1 <= 1
## 9 <NA>      <NA>      <NA> <NA> <NA>
## 10 Promising Recent shoppers, but haven't spent much 3- 4 <= 1 <= 1
## # ... with 13 more rows
```


Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%  
  janitor::clean_names()
```

```
## # A tibble: 23 × 5  
##   segment      description      r      f  
##   <chr>      <chr>      <chr> <chr>  
## 1 <NA>      <NA>      <NA> <NA>  
## 2 Champions Bought recently, buy often and spend th... 4- 5 4- 5  
## 3 <NA>      <NA>      <NA> <NA>  
## 4 Loyal Customers Spend good money. Responsive to promoti... 2- 5 3- 5  
## 5 <NA>      <NA>      <NA> <NA>  
## 6 Potential Loyalist Recent customers, spent good amount, bo... 3- 5 1- 3  
## 7 <NA>      <NA>      <NA> <NA>  
## 8 New Customers Bought more recently, but not often      4- 5 <= 1  
## 9 <NA>      <NA>      <NA> <NA>  
## 10 Promising Recent shoppers, but haven't spent much 3- 4 <= 1  
## # ... with 13 more rows
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%  
  janitor::clean_names() %>%  
  janitor::remove_empty("rows")
```

```
## # A tibble: 11 × 5  
##   segment      description      r      f  
##   <chr>      <chr>      <chr> <chr>  
## 1 Champions Bought recently, buy often and spend th... 4- 5 4- 5  
## 2 Loyal Customers Spend good money. Responsive to promoti... 2- 5 3- 5  
## 3 Potential Loyalist Recent customers, spent good amount, bo... 3- 5 1- 3  
## 4 New Customers Bought more recently, but not often      4- 5 <= 1  
## 5 Promising Recent shoppers, but haven't spent much    3- 4 <= 1  
## 6 Need Attention Above average recency, frequency & mone... 2- 3 2- 3  
## 7 About To Sleep Below average recency, frequency & mone... 2- 3 <= 2  
## 8 At Risk Spent big money, purchased often but lo... <= 2 2- 5  
## 9 Can't Lose Them Made big purchases and often, but long ... <= 1 4- 5  
## 10 Hibernating Low spenders, low frequency, purchased ... 1- 2 1- 2  
## 11 Lost Lowest recency, frequency & monetary sc... <= 2 <= 2
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%  
  janitor::clean_names() %>%  
  janitor::remove_empty("rows") %>%  
  pivot_longer(cols = r:m)
```

```
## # A tibble: 33 × 4  
##   segment      description name  
##   <chr>      <chr>      <chr>  
## 1 Champions Bought recently, buy often and spend the most r  
## 2 Champions Bought recently, buy often and spend the most f  
## 3 Champions Bought recently, buy often and spend the most m  
## 4 Loyal Customers Spend good money. Responsive to promotions r  
## 5 Loyal Customers Spend good money. Responsive to promotions f  
## 6 Loyal Customers Spend good money. Responsive to promotions m  
## 7 Potential Loyalist Recent customers, spent good amount, bought m... r  
## 8 Potential Loyalist Recent customers, spent good amount, bought m... f  
## 9 Potential Loyalist Recent customers, spent good amount, bought m... m  
## 10 New Customers Bought more recently, but not often r  
## # ... with 23 more rows
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%
  janitor::clean_names() %>%
  janitor::remove_empty("rows") %>%
  pivot_longer(cols = r:m) %>%
  separate(col = value, into = c("lo", "hi"),
           remove = FALSE, convert = TRUE,
           fill = "left")
```

```
## # A tibble: 33 × 6
##   segment      description      name  value  lo
##   <chr>      <chr>      <chr> <chr> <int> <
## 1 Champions Bought recently, buy often and sp... r    4- 5    4
## 2 Champions Bought recently, buy often and sp... f    4- 5    4
## 3 Champions Bought recently, buy often and sp... m    4- 5    4
## 4 Loyal Customers Spend good money. Responsive to p... r    2- 5    2
## 5 Loyal Customers Spend good money. Responsive to p... f    3- 5    3
## 6 Loyal Customers Spend good money. Responsive to p... m    3- 5    3
## 7 Potential Loyalist Recent customers, spent good amou... r    3- 5    3
## 8 Potential Loyalist Recent customers, spent good amou... f    1- 3    1
## 9 Potential Loyalist Recent customers, spent good amou... m    1- 3    1
## 10 New Customers Bought more recently, but not oft... r    4- 5    4
## # ... with 23 more rows
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%
  janitor::clean_names() %>%
  janitor::remove_empty("rows") %>%
  pivot_longer(cols = r:m) %>%
  separate(col = value, into = c("lo", "hi"),
           remove = FALSE, convert = TRUE,
           fill = "left") %>%
  select(-value)
```

```
## # A tibble: 33 × 5
##   segment      description      name      lo
##   <chr>      <chr>      <chr> <int> <chr>
## 1 Champions Bought recently, buy often and spend th... r      4
## 2 Champions Bought recently, buy often and spend th... f      4
## 3 Champions Bought recently, buy often and spend th... m      4
## 4 Loyal Customers Spend good money. Responsive to promoti... r      2
## 5 Loyal Customers Spend good money. Responsive to promoti... f      3
## 6 Loyal Customers Spend good money. Responsive to promoti... m      3
## 7 Potential Loyalist Recent customers, spent good amount, bo... r      3
## 8 Potential Loyalist Recent customers, spent good amount, bo... f      1
## 9 Potential Loyalist Recent customers, spent good amount, bo... m      1
## 10 New Customers Bought more recently, but not often      r      4
## # ... with 23 more rows
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%
  janitor::clean_names() %>%
  janitor::remove_empty("rows") %>%
  pivot_longer(cols = r:m) %>%
  separate(col = value, into = c("lo", "hi"),
           remove = FALSE, convert = TRUE,
           fill = "left") %>%
  select(-value) %>%
  pivot_wider(names_from = name,
             values_from = lo:hi)
```

```
## # A tibble: 11 × 8
##   segment      description lo_r lo_f lo_m hi_r hi_f
##   <chr>      <chr>      <int> <int> <int> <int> <int>
## 1 Champions Bought recently, buy ...     4     4     4     5     5
## 2 Loyal Customers Spend good money. Res...     2     3     3     5     5
## 3 Potential Loyalist Recent customers, spe...     3     1     1     5     3
## 4 New Customers Bought more recently,...     4    NA    NA     5     1
## 5 Promising Recent shoppers, but ...     3    NA    NA     4     1
## 6 Need Attention Above average recency...     2     2     2     3     3
## 7 About To Sleep Below average recency...     2    NA    NA     3     2
## 8 At Risk Spent big money, purc...    NA     2     2     2     5
## 9 Can't Lose Them Made big purchases an...    NA     4     4     1     5
## 10 Hibernating Low spenders, low fre...     1     1     1     2     2
## 11 Lost Lowest recency, frequ...    NA    NA    NA     2     2
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%
  janitor::clean_names() %>%
  janitor::remove_empty("rows") %>%
  pivot_longer(cols = r:m) %>%
  separate(col = value, into = c("lo", "hi"),
    remove = FALSE, convert = TRUE,
    fill = "left") %>%
  select(-value) %>%
  pivot_wider(names_from = name,
    values_from = lo:hi) %>%
  mutate(across(where(is.integer), replace_na, 0))
```

```
## # A tibble: 11 × 8
##   segment      description    lo_r  lo_f  lo_m  hi_r  hi_f
##   <chr>      <chr>      <int> <int> <int> <int> <int>
## 1 Champions Bought recently, buy ...     4     4     4     5     5
## 2 Loyal Customers Spend good money. Res...     2     3     3     5     5
## 3 Potential Loyalist Recent customers, spe...     3     1     1     5     3
## 4 New Customers Bought more recently,...     4     0     0     5     1
## 5 Promising Recent shoppers, but ...     3     0     0     4     1
## 6 Need Attention Above average recency...     2     2     2     3     3
## 7 About To Sleep Below average recency...     2     0     0     3     2
## 8 At Risk Spent big money, purc...     0     2     2     2     5
## 9 Can't Lose Them Made big purchases an...     0     4     4     1     5
## 10 Hibernating Low spenders, low fre...     1     1     1     2     2
## 11 Lost Lowest recency, frequ...     0     0     0     2     2
```

Cleaning a table

```
read_csv(here("data", "rfm_table.csv")) %>%
  janitor::clean_names() %>%
  janitor::remove_empty("rows") %>%
  pivot_longer(cols = r:m) %>%
  separate(col = value, into = c("lo", "hi"),
    remove = FALSE, convert = TRUE,
    fill = "left") %>%
  select(-value) %>%
  pivot_wider(names_from = name,
    values_from = lo:hi) %>%
  mutate(across(where(is.integer), replace_na, 0))
select(segment,
  lo_r, hi_r,
  lo_f, hi_f,
  lo_m, hi_m,
  description)
```

```
## # A tibble: 11 × 8
##   segment lo_r hi_r lo_f hi_f lo_m hi_m description
##   <chr>   <int> <int> <int> <int> <int> <int> <chr>
## 1 Champions      4     5     4     5     4     5 Bought recently, h
## 2 Loyal Customers  2     5     3     5     3     5 Spend good money.
## 3 Potential Loyalist 3     5     1     3     1     3 Recent customers,
## 4 New Customers   4     5     0     1     0     1 Bought more recent
## 5 Promising       3     4     0     1     0     1 Recent shoppers, h
## 6 Need Attention  2     3     2     3     2     3 Above average rece
## 7 About To Sleep  2     3     0     2     0     2 Below average rece
## 8 At Risk         0     2     2     5     2     5 Spent big money, p
## 9 Can't Lose Them  0     1     4     5     4     5 Made big purchases
## 10 Hibernating    1     2     1     2     1     2 Low spenders, low
## 11 Lost          0     2     0     2     0     2 Lowest recency, fr
```


Maybe a candidate for `rowwise()`?

```
rfm_table
```

```
## # A tibble: 11 × 8
##   segment      lo_r hi_r lo_f hi_f lo_m hi_m description
##   <chr>    <int> <int> <int> <int> <int> <int> <chr>
## 1 Champions      4     5     4     5     4     5 Bought recently, buy ...
## 2 Loyal Customers  2     5     3     5     3     5 Spend good money. Res...
## 3 Potential Loyalist 3     5     1     3     1     3 Recent customers, spe...
## 4 New Customers   4     5     0     1     0     1 Bought more recently,...
## 5 Promising      3     4     0     1     0     1 Recent shoppers, but ...
## 6 Need Attention  2     3     2     3     2     3 Above average recency...
## 7 About To Sleep  2     3     0     2     0     2 Below average recency...
## 8 At Risk        0     2     2     5     2     5 Spent big money, purc...
## 9 Can't Lose Them  0     1     4     5     4     5 Made big purchases an...
## 10 Hibernating    1     2     1     2     1     2 Low spenders, low fre...
## 11 Lost          0     2     0     2     0     2 Lowest recency, frequ...
```

Maybe a candidate for `rowwise()`?

This does what we expect:

```
rfm_table %>%  
  mutate(sum_lo = lo_r + lo_f + lo_m,  
         sum_hi = hi_r + hi_f + hi_m) %>%  
  select(segment, sum_lo, sum_hi, everything())
```

```
## # A tibble: 11 × 10  
##   segment      sum_lo sum_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <int>  <int> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions      12     15     4     5     4     5     4     5 Bought rec...  
## 2 Loyal Customers    8     15     2     5     3     5     3     5 Spend good...  
## 3 Potential Loya...    5     11     3     5     1     3     1     3 Recent cus...  
## 4 New Customers     4      7     4     5     0     1     0     1 Bought mor...  
## 5 Promising        3      6     3     4     0     1     0     1 Recent sho...  
## 6 Need Attention    6      9     2     3     2     3     2     3 Above aver...  
## 7 About To Sleep    2      7     2     3     0     2     0     2 Below aver...  
## 8 At Risk          4     12     0     2     2     5     2     5 Spent big ...  
## 9 Can't Lose Them   8     11     0     1     4     5     4     5 Made big p...  
## 10 Hibernating      3      6     1     2     1     2     1     2 Low spende...  
## 11 Lost            0      6     0     2     0     2     0     2 Lowest rec...
```

This adds each column, elementwise.

Maybe a candidate for `rowwise()`?

But this does not:

```
rfm_table %>%  
  mutate(sum_lo = sum(lo_r, lo_f, lo_m),  
         sum_hi = sum(hi_r, hi_f, hi_m)) %>%  
  select(segment, sum_lo, sum_hi, everything())
```

```
## # A tibble: 11 × 10  
##   segment      sum_lo sum_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <int>  <int> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions      55    105   4    5   4    5   4    5 Bought rec...  
## 2 Loyal Customers  55    105   2    5   3    5   3    5 Spend good...  
## 3 Potential Loya...  55    105   3    5   1    3   1    3 Recent cus...  
## 4 New Customers   55    105   4    5   0    1   0    1 Bought mor...  
## 5 Promising       55    105   3    4   0    1   0    1 Recent sho...  
## 6 Need Attention   55    105   2    3   2    3   2    3 Above aver...  
## 7 About To Sleep   55    105   2    3   0    2   0    2 Below aver...  
## 8 At Risk         55    105   0    2   2    5   2    5 Spent big ...  
## 9 Can't Lose Them  55    105   0    1   4    5   4    5 Made big p...  
## 10 Hibernating     55    105   1    2   1    2   1    2 Low spende...  
## 11 Lost           55    105   0    2   0    2   0    2 Lowest rec...
```

Sum is taking all the columns, adding them up (into a single number), and putting that result in each row.

Maybe a candidate for `rowwise()`?

Similarly, this will not give the answer we probably expect:

```
rfm_table %>%  
  mutate(mean_lo = mean(c(lo_r, lo_f, lo_m)),  
         mean_hi = mean(c(hi_r, hi_f, hi_m))) %>%  
  select(segment, mean_lo, mean_hi, everything())
```

```
## # A tibble: 11 × 10  
##   segment mean_lo mean_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <dbl>  <dbl> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions      1.67    3.18     4     5     4     5     4     5 Bought rec...  
## 2 Loyal Custom...  1.67    3.18     2     5     3     5     3     5 Spend good...  
## 3 Potential Lo...  1.67    3.18     3     5     1     3     1     3 Recent cus...  
## 4 New Customers  1.67    3.18     4     5     0     1     0     1 Bought mor...  
## 5 Promising      1.67    3.18     3     4     0     1     0     1 Recent sho...  
## 6 Need Attenti...  1.67    3.18     2     3     2     3     2     3 Above aver...  
## 7 About To Sle...  1.67    3.18     2     3     0     2     0     2 Below aver...  
## 8 At Risk        1.67    3.18     0     2     2     5     2     5 Spent big ...  
## 9 Can't Lose T...  1.67    3.18     0     1     4     5     4     5 Made big p...  
## 10 Hibernating    1.67    3.18     1     2     1     2     1     2 Low spende...  
## 11 Lost          1.67    3.18     0     2     0     2     0     2 Lowest rec...
```

Maybe a candidate for `rowwise()`?

But this will:

```
rfm_table %>%  
  rowwise() %>%  
  mutate(mean_lo = mean(c(lo_r, lo_f, lo_m)),  
         mean_hi = mean(c(hi_r, hi_f, hi_m))) %>%  
  select(segment, mean_lo, mean_hi, everything())
```

```
## # A tibble: 11 × 10
```

```
## # Rowwise:
```

```
##   segment    mean_lo mean_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <dbl>   <dbl> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions      4       5      4     5     4     5     4     5 Bought rec...  
## 2 Loyal Custom... 2.67     5      2     5     3     5     3     5 Spend good...  
## 3 Potential Lo... 1.67     3.67    3     5     1     3     1     3 Recent cus...  
## 4 New Customers  1.33     2.33    4     5     0     1     0     1 Bought mor...  
## 5 Promising      1       2      3     4     0     1     0     1 Recent sho...  
## 6 Need Attenti... 2       3      2     3     2     3     2     3 Above aver...  
## 7 About To Sle... 0.667    2.33    2     3     0     2     0     2 Below aver...  
## 8 At Risk        1.33     4      0     2     2     5     2     5 Spent big ...  
## 9 Can't Lose T... 2.67     3.67    0     1     4     5     4     5 Made big p...  
## 10 Hibernating   1       2      1     2     1     2     1     2 Low spende...  
## 11 Lost          0       2      0     2     0     2     0     2 Lowest rec...
```

Rowwise isn't very efficient

In general, you'll want to see if some vectorized ("operating on columns, but elementwise") function exists, as it'll be faster.

And most of the time, R and the tidyverse "wants" you to work in vectorized, columnar terms ... hence your first move will often be to pivot the data into long format.

So, **rowwise()** is not likely to see a whole lot of further development.

You probably want `group_by()` instead

```
rfm_table %>%  
  group_by(segment) %>%  
  mutate(mean_lo = mean(c(lo_r, lo_f, lo_m)),  
         mean_hi = mean(c(hi_r, hi_f, hi_m))) %>%  
  select(segment, mean_lo, mean_hi, everything())
```

```
## # A tibble: 11 × 10  
## # Groups:   segment [11]  
##   segment    mean_lo mean_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <dbl>   <dbl> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions      4       5      4     5     4     5     4     5 Bought rec...  
## 2 Loyal Custom... 2.67     5      2     5     3     5     3     5 Spend good...  
## 3 Potential Lo... 1.67     3.67    3     5     1     3     1     3 Recent cus...  
## 4 New Customers  1.33     2.33    4     5     0     1     0     1 Bought mor...  
## 5 Promising      1       2      3     4     0     1     0     1 Recent sho...  
## 6 Need Attenti... 2       3      2     3     2     3     2     3 Above aver...  
## 7 About To Sle... 0.667    2.33    2     3     0     2     0     2 Below aver...  
## 8 At Risk        1.33     4      0     2     2     5     2     5 Spent big ...  
## 9 Can't Lose T... 2.67     3.67    0     1     4     5     4     5 Made big p...  
## 10 Hibernating   1       2      1     2     1     2     1     2 Low spende...  
## 11 Lost          0       2      0     2     0     2     0     2 Lowest rec...
```

You probably want `group_by()` instead

```
rfm_table %>%  
  group_by(segment) %>%  
  mutate(sum_lo = sum(lo_r, lo_f, lo_m),  
         sum_hi = sum(hi_r, hi_f, hi_m)) %>%  
  select(segment, sum_lo, sum_hi, everything())
```

```
## # A tibble: 11 × 10  
## # Groups:   segment [11]  
##   segment      sum_lo sum_hi lo_r hi_r lo_f hi_f lo_m hi_m description  
##   <chr>      <int> <int> <int> <int> <int> <int> <int> <int> <chr>  
## 1 Champions        12     15     4     5     4     5     4     5 Bought rec...  
## 2 Loyal Customers     8     15     2     5     3     5     3     5 Spend good...  
## 3 Potential Loya...    5     11     3     5     1     3     1     3 Recent cus...  
## 4 New Customers       4      7     4     5     0     1     0     1 Bought mor...  
## 5 Promising          3      6     3     4     0     1     0     1 Recent sho...  
## 6 Need Attention      6      9     2     3     2     3     2     3 Above aver...  
## 7 About To Sleep      2      7     2     3     0     2     0     2 Below aver...  
## 8 At Risk            4     12     0     2     2     5     2     5 Spent big ...  
## 9 Can't Lose Them     8     11     0     1     4     5     4     5 Made big p...  
## 10 Hibernating        3      6     1     2     1     2     1     2 Low spende...  
## 11 Lost              0      6     0     2     0     2     0     2 Lowest rec...
```


What about Stata?

Using **haven**

Haven is the Tidyverse's package for reading and managing files from Stata, SPSS, and SAS. You should prefer it to the older Base R package `foreign`, which has similar functionality.

We're going to import a General Social Survey dataset that's in Stata's `.dta` format.

```
library(haven)

# This will take a moment
gss_panel <- read_stata(here("data", "gss_panel_long.dta"))
```

We'll do some of the common recoding and reorganizing tasks that accompany this.

The GSS panel

The data:

```
gss_panel
```

```
## # A tibble: 14,610 × 2,757
##   firstyear firsttid year   id   vpsu vstrat adults ballot dateintv famgen
##   <dbl> <dbl+lbl> <dbl> <dbl> <dbl+> <dbl+> <dbl+> <dbl+l> <dbl+lb> <dbl+l>
## 1     2006      9 2006     9     2  1957     1 3 [BAL...    709 1 [1 G...
## 2     2006      9 2008   3001    NA    NA     2 3 [BAL...    503 1 [1 G...
## 3     2006      9 2010   6001 NA(i)    NA     2 3 [BAL...    508 1 [1 G...
## 4     2006     10 2010   6002 NA(i)    NA     1 1 [BAL...    408 1 [1 G...
## 5     2006     10 2006     10     2  1957     2 1 [BAL...    630 2 [2 G...
## 6     2006     10 2008   3002    NA    NA     2 1 [BAL...    426 2 [2 G...
## 7     2006     11 2008   3003    NA    NA     2 3 [BAL...    718 4 [2 G...
## 8     2006     11 2010   6003 NA(i)    NA  NA(n) 3 [BAL...    518 2 [2 G...
## 9     2006     11 2006     11     2  1957     2 3 [BAL...    630 4 [2 G...
## 10    2006     12 2010   6004 NA(i)    NA     4 1 [BAL...    324 2 [2 G...
## # ... with 14,600 more rows, and 2,747 more variables: form <dbl+lbl>,
## #   formwt <dbl>, gender1 <dbl+lbl>, hompop <dbl+lbl>, intage <dbl+lbl>,
## #   intid <dbl+lbl>, intyrs <dbl+lbl>, mode <dbl+lbl>, oversamp <dbl>,
## #   phase <dbl+lbl>, race <dbl+lbl>, reg16 <dbl+lbl>, region <dbl+lbl>,
## #   relate1 <dbl+lbl>, relhh1 <dbl+lbl>, relhhd1 <dbl+lbl>, respnum <dbl+lbl>,
## #   rvisitor <dbl+lbl>, sampcode <dbl+lbl>, sample <dbl+lbl>, sex <dbl+lbl>,
## #   size <dbl+lbl>, spaneng <dbl+lbl>, srcbelt <dbl+lbl>, version <dbl>, ...
```

The GSS panel

Many variables.

Stata's missing value types are preserved

Data types are things like `dbl+lbl` indicating that Stata's numeric values and variable labels have been preserved.

The GSS panel

You can see the labeling system at work:

```
gss_panel %>%  
  select(degree) %>%  
  group_by(degree) %>%  
  tally()
```

```
## # A tibble: 6 × 2  
##           degree      n  
##           <dbl+lbl> <int>  
## 1      0 [LT HIGH SCHOOL] 1850  
## 2      1 [HIGH SCHOOL]    7274  
## 3      2 [JUNIOR COLLEGE] 1161  
## 4      3 [bachelor]       2767  
## 5      4 [graduate]       1556  
## 6 NA(d)                  2
```

The GSS panel

Values get pivoted, not labels, though.

```
gss_panel %>%  
  select(sex, degree) %>%  
  group_by(sex, degree) %>%  
  tally() %>%  
  pivot_wider(names_from = sex, values_from = n)
```

```
## # A tibble: 6 × 3  
##           degree  `1`  `2`  
##      <dbl+lbl> <int> <int>  
## 1  0 [LT HIGH SCHOOL]   814  1036  
## 2  1 [HIGH SCHOOL]     3131  4143  
## 3  2 [JUNIOR COLLEGE]   440   721  
## 4  3 [bachelor]        1293  1474  
## 5  4 [graduate]         696   860  
## 6 NA(d)                NA     2
```

The GSS panel

Option 1: Just drop all the labels.

```
gss_panel %>%  
  zap_missing() %>%  
  zap_labels()
```

```
## # A tibble: 14,610 × 2,757  
##   firstyear firstid year   id vpsu vstrat adults ballot dateintv famgen  
##   <dbl>   <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>   <dbl> <dbl>  
## 1     2006     9  2006     9     2  1957     1     3     709     1  
## 2     2006     9  2008  3001    NA    NA     2     3     503     1  
## 3     2006     9  2010  6001    NA    NA     2     3     508     1  
## 4     2006    10  2010  6002    NA    NA     1     1     408     1  
## 5     2006    10  2006    10     2  1957     2     1     630     2  
## 6     2006    10  2008  3002    NA    NA     2     1     426     2  
## 7     2006    11  2008  3003    NA    NA     2     3     718     4  
## 8     2006    11  2010  6003    NA    NA    NA     3     518     2  
## 9     2006    11  2006    11     2  1957     2     3     630     4  
## 10    2006    12  2010  6004    NA    NA     4     1     324     2  
## # ... with 14,600 more rows, and 2,747 more variables: form <dbl>, formwt <dbl>,  
## #   gender1 <dbl>, hompop <dbl>, intage <dbl>, intid <dbl>, intyrs <dbl>,  
## #   mode <dbl>, oversamp <dbl>, phase <dbl>, race <dbl>, reg16 <dbl>,  
## #   region <dbl>, relate1 <dbl>, relhh1 <dbl>, relhhd1 <dbl>, respnum <dbl>,  
## #   rvisitor <dbl>, sampcode <dbl>, sample <dbl>, sex <dbl>, size <dbl>,  
## #   spaneng <dbl>, srcbelt <dbl>, version <dbl>, visitors <dbl>, wtss <dbl>,  
## #   wtssall <dbl>, wtssnr <dbl>, xnorcsiz <dbl>, hispanic <dbl>, ...
```

The GSS panel

Option 2: Convert the labels

Let's focus on a few measures of interest, and do some recoding.

```
## Categorical vars
cat_vars <- c("race", "sex", "degree", "relig", "income", "polviews", "fefam")

## Integer vars
int_vars <- c("year", "id", "ballot", "age", "tvhours")

## Survey design
wt_vars <- c("vpsu",
            "vstrat",
            "oversamp",
            "formwt",      # weight to deal with experimental randomization
            "wtssall",     # weight variable
            "sampcode",    # sampling error code
            "sample")      # sampling frame and method

my_gss_vars <- c(int_vars, cat_vars, wt_vars)
```

Now we're ready to go ...

Cut down the dataset

```
gss_sub <- gss_panel %>%  
  select(all_of(my_gss_vars))
```

```
gss_sub
```

```
## # A tibble: 14,610 × 19  
##   year   id ballot age tvhours race sex degree relig  
##   <dbl> <dbl> <dbl+lbl> <dbl+lb> <dbl+lbl> <dbl+lb> <dbl+lb> <dbl+lb> <dbl+lb>  
## 1  2006     9 3 [BALLOT C] 23 NA(a) [iap] 2 [bla... 2 [fem... 3 [bac... 4 [non...  
## 2  2008  3001 3 [BALLOT C] 25 NA(i)      3 [oth... 2 [fem... 3 [bac... 4 [non...  
## 3  2010  6001 3 [BALLOT C] 27 NA(i)      2 [bla... 2 [fem... 3 [bac... 4 [non...  
## 4  2010  6002 1 [BALLOT A] 36      3      1 [whi... 2 [fem... 4 [gra... 4 [non...  
## 5  2006    10 1 [BALLOT A] 32      3      3 [oth... 2 [fem... 4 [gra... 4 [non...  
## 6  2008  3002 1 [BALLOT A] 34      3      3 [oth... 2 [fem... 4 [gra... 4 [non...  
## 7  2008  3003 3 [BALLOT C] 83 NA(i)      2 [bla... 2 [fem... 0 [LT ... 1 [pro...  
## 8  2010  6003 3 [BALLOT C] 85 NA(i)      2 [bla... 2 [fem... 0 [LT ... 1 [pro...  
## 9  2006    11 3 [BALLOT C] 81 NA(a) [iap] 2 [bla... 2 [fem... 0 [LT ... 1 [pro...  
## 10 2010  6004 1 [BALLOT A] 51    10      3 [oth... 1 [mal... 1 [HIG... 2 [cat...  
## # ... with 14,600 more rows, and 10 more variables: income <dbl+lbl>,  
## #   polviews <dbl+lbl>, fefam <dbl+lbl>, vpsu <dbl+lbl>, vstrat <dbl+lbl>,  
## #   oversamp <dbl>, formwt <dbl>, wtssall <dbl+lbl>, sampcode <dbl+lbl>,  
## #   sample <dbl+lbl>
```

The GSS Panel: Recoding

```
gss_sub %>%
  mutate(across(everything(), zap_missing)) %>%
  mutate(across(all_of(wt_vars), as.numeric)) %>%
  mutate(across(all_of(int_vars), as.integer)) %>%
  mutate(across(all_of(cat_vars), as_factor)) %>%
  mutate(across(all_of(cat_vars), fct_relabel, tolower)) %>%
  mutate(across(all_of(cat_vars), fct_relabel, tools::toTitleCase)) %>%
  mutate(income = stringr::str_replace(income, " - ", "-"))
```

```
## # A tibble: 14,610 × 19
##   year   id ballot  age tvhours race  sex  degree  relig income polviews
##   <int> <int> <int> <int>   <int> <fct> <fct> <fct>   <fct> <chr>   <fct>
## 1  2006     9     3    23     NA Black Female Bachelor None  $2500... Conserv...
## 2  2008   3001     3    25     NA Other Female Bachelor None  $2500... Extreme...
## 3  2010   6001     3    27     NA Black Female Bachelor None  $2500... Extreme...
## 4  2010   6002     1    36      3 White Female Graduate None  $2500... Liberal
## 5  2006    10     1    32      3 Other Female Graduate None  <NA>    Slightl...
## 6  2008   3002     1    34      3 Other Female Graduate None  $2500... Moderate
## 7  2008   3003     3    83     NA Black Female Lt High ... Prot... $2000... Liberal
## 8  2010   6003     3    85     NA Black Female Lt High ... Prot... <NA>    Moderate
## 9  2006    11     3    81     NA Black Female Lt High ... Prot... <NA>    Moderate
## 10 2010   6004     1    51     10 Other Male  High Sch... Cath... Lt $1... Liberal
## # ... with 14,600 more rows, and 8 more variables: fefam <fct>, vpsu <dbl>,
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampcode <dbl>,
## #   sample <dbl>
```

How we'd actually write this

```
gss_sub <- gss_sub %>%  
  mutate(across(everything(), zap_missing),  
    across(all_of(wt_vars), as.numeric),  
    across(all_of(int_vars), as.integer),  
    across(all_of(cat_vars), as_factor),  
    across(all_of(cat_vars), fct_relabel, tolower),  
    across(all_of(cat_vars), fct_relabel, tools::toTitleCase),  
    income = stringr::str_replace(income, " - ", "-"))
```

The GSS panel: more recoding

Age quintiles: find the cutpoints

```
# seq can make all kinds of sequences  
seq(from = 0, to = 1, by = 0.2)
```

```
## [1] 0.0 0.2 0.4 0.6 0.8 1.0
```

```
age_quintiles <- quantile(as.numeric(gss_panel$age),  
                          probs = seq(0, 1, 0.2),  
                          na.rm = TRUE)
```

```
## These are the quintile cutpoints  
age_quintiles
```

```
##    0%   20%   40%   60%   80%  100%  
##    18    33    43    53    65    89
```

The GSS panel: more recoding

Age quintiles: create the quintile variable

```
## Apply the cut
gss_sub %>%
  mutate(agequint = cut(x = age,
                        breaks = unique(age_quintiles),
                        include.lowest = TRUE)) %>%
  pull(agequint) %>% # grab a column and make it an ordinary vector
  table()
```

```
## .
## [18,33] [33,43] [43,53] [53,65] [65,89]
##      3157      2680      2851      3057      2720
```

We'll need to clean up those labels.

The GSS panel: more recoding

I told you that regexp stuff would pay off.

```
convert_agegrp <- function(x){  
  x <- stringr::str_remove(x, "\\(") # Remove open paren  
  x <- stringr::str_remove(x, "\\[") # Remove open bracket  
  x <- stringr::str_remove(x, "\\]") # Remove close bracket  
  x <- stringr::str_replace(x, ",", "-") # Replace comma with dash  
  x <- stringr::str_replace(x, "-89", "+") # Replace -89 with +  
  regex <- "^(.*)$" # Matches everything in string to end of line  
  x <- stringr::str_replace(x, regex, "Age \\1") # Preface string with "Age"  
  x  
}
```

The GSS panel: more recoding

gss_sub

```
## # A tibble: 14,610 × 19
##   year      id ballot  age tvhours race  sex  degree  relig  inco
##   <int> <int>  <int> <int>   <int> <fct> <fct>  <fct>  <fct> <chr
## 1  2006      9      3   23     NA Black Female Bachelor None $250
## 2  2008   3001      3   25     NA Other Female Bachelor None $250
## 3  2010   6001      3   27     NA Black Female Bachelor None $250
## 4  2010   6002      1   36      3 White Female Graduate None $250
## 5  2006     10      1   32      3 Other Female Graduate None <NA>
## 6  2008   3002      1   34      3 Other Female Graduate None $250
## 7  2008   3003      3   83     NA Black Female Lt High ... Prot... $200
## 8  2010   6003      3   85     NA Black Female Lt High ... Prot... <NA>
## 9  2006     11      3   81     NA Black Female Lt High ... Prot... <NA>
## 10 2010   6004      1   51     10 Other Male  High Sch... Cath... Lt $
## # ... with 14,600 more rows, and 8 more variables: fefam <fct>, vpsu <d
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco
## #   sample <dbl>
```

The GSS panel: more recoding

```
gss_sub %>%  
  mutate(agequint = cut(x = age,  
                        breaks = unique(age_quintiles)  
                        include.lowest = TRUE))
```

```
## # A tibble: 14,610 × 20  
##   year   id ballot  age tvhours race  sex  degree  relig  inco  
##   <int> <int>   <int> <int>   <int> <fct> <fct>   <fct>   <fct> <chr>  
## 1  2006     9     3    23      NA Black Female Bachelor None $250  
## 2  2008   3001     3    25      NA Other Female Bachelor None $250  
## 3  2010   6001     3    27      NA Black Female Bachelor None $250  
## 4  2010   6002     1    36       3 White Female Graduate None $250  
## 5  2006    10     1    32       3 Other Female Graduate None <NA>  
## 6  2008   3002     1    34       3 Other Female Graduate None $250  
## 7  2008   3003     3    83      NA Black Female Lt High ... Prot... $200  
## 8  2010   6003     3    85      NA Black Female Lt High ... Prot... <NA>  
## 9  2006    11     3    81      NA Black Female Lt High ... Prot... <NA>  
## 10 2010   6004     1    51     10 Other Male  High Sch... Cath... Lt $  
## # ... with 14,600 more rows, and 9 more variables: fefam <fct>, vpsu <d  
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco  
## #   sample <dbl>, agequint <fct>
```


The GSS panel: more recoding

```
gss_sub %>%  
  mutate(agequint = cut(x = age,  
    breaks = unique(age_quintiles)  
    include.lowest = TRUE)) %>%  
  mutate(agequint = fct_relabel(agequint, convert_agec
```

```
## # A tibble: 14,610 × 20  
##   year   id ballot  age tvhours race  sex  degree  relig  inco  
##   <int> <int> <int> <int>   <int> <fct> <fct> <fct>   <fct> <chr>  
## 1  2006     9      3   23      NA Black Female Bachelor None $250  
## 2  2008  3001      3   25      NA Other Female Bachelor None $250  
## 3  2010  6001      3   27      NA Black Female Bachelor None $250  
## 4  2010  6002      1   36       3 White Female Graduate None $250  
## 5  2006    10      1   32       3 Other Female Graduate None <NA>  
## 6  2008  3002      1   34       3 Other Female Graduate None $250  
## 7  2008  3003      3   83      NA Black Female Lt High ... Prot... $200  
## 8  2010  6003      3   85      NA Black Female Lt High ... Prot... <NA>  
## 9  2006    11      3   81      NA Black Female Lt High ... Prot... <NA>  
## 10 2010  6004      1   51     10 Other Male  High Sch... Cath... Lt $  
## # ... with 14,600 more rows, and 9 more variables: fefam <fct>, vpsu <d  
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco  
## #   sample <dbl>, agequint <fct>
```

The GSS panel: more recoding

```
gss_sub %>%  
  mutate(agequint = cut(x = age,  
    breaks = unique(age_quintiles)  
    include.lowest = TRUE)) %>%  
  mutate(agequint = fct_relabel(agequint, convert_ageq  
  mutate(year_f = droplevels(factor(year)))
```

```
## # A tibble: 14,610 × 21  
##   year   id ballot  age tvhours race  sex  degree  relig inco  
##   <int> <int> <int> <int>   <int> <fct> <fct> <fct>   <fct> <chr>  
## 1  2006     9     3   23     NA Black Female Bachelor None $250  
## 2  2008   3001     3   25     NA Other Female Bachelor None $250  
## 3  2010   6001     3   27     NA Black Female Bachelor None $250  
## 4  2010   6002     1   36      3 White Female Graduate None $250  
## 5  2006    10     1   32      3 Other Female Graduate None <NA>  
## 6  2008   3002     1   34      3 Other Female Graduate None $250  
## 7  2008   3003     3   83     NA Black Female Lt High ... Prot... $200  
## 8  2010   6003     3   85     NA Black Female Lt High ... Prot... <NA>  
## 9  2006    11     3   81     NA Black Female Lt High ... Prot... <NA>  
## 10 2010   6004     1   51     10 Other Male  High Sch... Cath... Lt $  
## # ... with 14,600 more rows, and 10 more variables: fefam <fct>, vpsu <  
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco  
## #   sample <dbl>, agequint <fct>, year_f <fct>
```

The GSS panel: more recoding

```
gss_sub %>%
  mutate(agequint = cut(x = age,
                        breaks = unique(age_quintiles)
                        include.lowest = TRUE)) %>%
  mutate(agequint = fct_relabel(agequint, convert_ageq
  mutate(year_f = droplevels(factor(year))) %>%
  mutate(young = ifelse(age < 26, "Yes", "No"))
```

```
## # A tibble: 14,610 × 22
##   year   id ballot  age tvhours race  sex  degree  relig inco
##   <int> <int> <int> <int>   <int> <fct> <fct> <fct>   <fct> <chr>
## 1  2006     9     3    23     NA Black Female Bachelor None $250
## 2  2008   3001     3    25     NA Other Female Bachelor None $250
## 3  2010   6001     3    27     NA Black Female Bachelor None $250
## 4  2010   6002     1    36      3 White Female Graduate None $250
## 5  2006    10     1    32      3 Other Female Graduate None <NA>
## 6  2008   3002     1    34      3 Other Female Graduate None $250
## 7  2008   3003     3    83     NA Black Female Lt High ... Prot... $200
## 8  2010   6003     3    85     NA Black Female Lt High ... Prot... <NA>
## 9  2006    11     3    81     NA Black Female Lt High ... Prot... <NA>
## 10 2010   6004     1    51     10 Other Male  High Sch... Cath... Lt $
## # ... with 14,600 more rows, and 11 more variables: fefam <fct>, vpsu <
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco
## #   sample <dbl>, agequint <fct>, year_f <fct>, young <chr>
```

The GSS panel: more recoding

```
gss_sub %>%
  mutate(agequint = cut(x = age,
                        breaks = unique(age_quintiles)
                        include.lowest = TRUE)) %>%
  mutate(agequint = fct_relabel(agequint, convert_ageq
  mutate(year_f = droplevels(factor(year))) %>%
  mutate(young = ifelse(age < 26, "Yes", "No")) %>%
  mutate(fefam_d = fct_recode(fefam,
                             Agree = "Strongly Agree"
                             Disagree = "Strongly Dis
```

```
## # A tibble: 14,610 × 23
##   year   id ballot  age tvhours race  sex  degree  relig inco
##   <int> <int> <int> <int>   <int> <fct> <fct> <fct>   <fct> <chr>
## 1  2006     9     3   23     NA Black Female Bachelor None $250
## 2  2008   3001     3   25     NA Other Female Bachelor None $250
## 3  2010   6001     3   27     NA Black Female Bachelor None $250
## 4  2010   6002     1   36      3 White Female Graduate None $250
## 5  2006    10     1   32      3 Other Female Graduate None <NA>
## 6  2008   3002     1   34      3 Other Female Graduate None $250
## 7  2008   3003     3   83     NA Black Female Lt High ... Prot... $200
## 8  2010   6003     3   85     NA Black Female Lt High ... Prot... <NA>
## 9  2006    11     3   81     NA Black Female Lt High ... Prot... <NA>
## 10 2010   6004     1   51     10 Other Male  High Sch... Cath... Lt $
## # ... with 14,600 more rows, and 12 more variables: fefam <fct>, vpsu <
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco
## #   sample <dbl>, agequint <fct>, year_f <fct>, young <chr>, fefam_d
```

The GSS panel: more recoding

```
gss_sub %>%
  mutate(agequint = cut(x = age,
                        breaks = unique(age_quintiles)
                        include.lowest = TRUE)) %>%
  mutate(agequint = fct_relabel(agequint, convert_ageq
  mutate(year_f = droplevels(factor(year))) %>%
  mutate(young = ifelse(age < 26, "Yes", "No")) %>%
  mutate(fefam_d = fct_recode(fefam,
                             Agree = "Strongly Agree"
                             Disagree = "Strongly Dis
  mutate(degree = factor(degree,
                          levels = levels(gss_sub$degree
                          ordered = TRUE)) #<<

## # A tibble: 14,610 × 23
##   year   id ballot  age tvhours race  sex  degree  relig inco
##   <int> <int> <int> <int>   <int> <fct> <fct> <ord>   <fct> <chr>
## 1  2006     9     3   23     NA Black Female Bachelor None $250
## 2  2008   3001     3   25     NA Other Female Bachelor None $250
## 3  2010   6001     3   27     NA Black Female Bachelor None $250
## 4  2010   6002     1   36      3 White Female Graduate None $250
## 5  2006    10     1   32      3 Other Female Graduate None <NA>
## 6  2008   3002     1   34      3 Other Female Graduate None $250
## 7  2008   3003     3   83     NA Black Female Lt High ... Prot... $200
## 8  2010   6003     3   85     NA Black Female Lt High ... Prot... <NA>
## 9  2006    11     3   81     NA Black Female Lt High ... Prot... <NA>
## 10 2010   6004     1   51     10 Other Male  High Sch... Cath... Lt $
## # ... with 14,600 more rows, and 12 more variables: fefam <fct>, vpsu <
## #   vstrat <dbl>, oversamp <dbl>, formwt <dbl>, wtssall <dbl>, sampco
## #   sample <dbl>, agequint <fct>, year_f <fct>, young <chr>, fefam_d
```

How we'd actually write this

```
gss_sub <- gss_sub %>%  
  mutate(agequint = cut(x = age,  
                        breaks = unique(age_quintiles),  
                        include.lowest = TRUE),  
         agequint = fct_relabel(agequint, convert_agegrp),  
         year_f = droplevels(factor(year)),  
         young = ifelse(age < 26, "Yes", "No"),  
         fefam_d = fct_recode(fefam,  
                              Agree = "Strongly Agree",  
                              Disagree = "Strongly Disagree"),  
         degree = factor(degree,  
                          levels = levels(gss_sub$degree),  
                          ordered = TRUE))
```

How we'd actually write this

```
gss_sub <- gss_sub %>%  
  mutate(agequint = cut(x = age,  
                        breaks = unique(age_quintiles),  
                        include.lowest = TRUE),  
         agequint = fct_relabel(agequint, convert_agegrp),  
         year_f = factor(year),  
         young = ifelse(age < 26, "Yes", "No"),  
         fefam_d = fct_recode(fefam,  
                              Agree = "Strongly Agree",  
                              Disagree = "Strongly Disagree"),  
         degree = factor(degree,  
                          levels = levels(gss_sub$degree),  
                          ordered = TRUE))
```

How we'd actually write this

```
gss_sub <- gss_sub %>%  
  mutate(agequint = cut(x = age,  
                        breaks = unique(age_quintiles),  
                        include.lowest = TRUE),  
    agequint = fct_relabel(agequint, convert_agegrp),  
    year_f = droplevels(factor(year)),  
    young = ifelse(age < 26, "Yes", "No"),  
    fefam_d = fct_recode(fefam,  
                        Agree = "Strongly Agree",  
                        Disagree = "Strongly Disagree"),  
    degree = factor(degree,  
                    levels = levels(gss_sub$degree),  
                    ordered = TRUE))
```


GSS Panel

```
gss_sub %>%  
  select(sex, year, year_f, age, young, fefam, fefam_d) %>%  
  sample_n(15)
```

```
## # A tibble: 15 × 7
```

##	sex	year	year_f	age	young	fefam	fefam_d
##	<fct>	<int>	<fct>	<int>	<chr>	<fct>	<fct>
##	1 Female	2012	2012	89	No	<NA>	<NA>
##	2 Male	2014	2014	42	No	<NA>	<NA>
##	3 Female	2012	2012	56	No	Strongly Agree	Agree
##	4 Male	2012	2012	29	No	Agree	Agree
##	5 Female	2012	2012	28	No	<NA>	<NA>
##	6 Female	2012	2012	35	No	Disagree	Disagree
##	7 Male	2014	2014	NA	<NA>	Disagree	Disagree
##	8 Female	2008	2008	28	No	Disagree	Disagree
##	9 Female	2012	2012	51	No	<NA>	<NA>
##	10 Female	2008	2008	31	No	Agree	Agree
##	11 Female	2010	2010	44	No	Strongly Agree	Agree
##	12 Female	2010	2010	60	No	Disagree	Disagree
##	13 Female	2010	2010	38	No	Disagree	Disagree
##	14 Female	2010	2010	62	No	Agree	Agree
##	15 Male	2012	2012	73	No	Agree	Agree

GSS Panel

```
gss_sub %>%  
  select(sex, degree) %>%  
  group_by(sex, degree) %>%  
  tally() %>%  
  pivot_wider(names_from = sex, values_from = n)
```

```
## # A tibble: 6 × 3  
##   degree      Male Female  
##   <ord>      <int> <int>  
## 1 Lt High School    814   1036  
## 2 High School    3131   4143  
## 3 Junior College   440    721  
## 4 Bachelor    1293   1474  
## 5 Graduate     696    860  
## 6 <NA>         NA      2
```

More on **factors**

We've already seen `fct_relabel()` and `fct_recode()` from `forcats`.

There are numerous other convenience functions for factors.

More on factors

We've already seen `fct_relabel()` and `fct_recode()` from `forcats`.

There are numerous other convenience functions for factors.

```
gss_sub %>%  
  count(degree)
```

```
## # A tibble: 6 × 2  
##   degree      n  
##   <ord>    <int>  
## 1 Lt High School 1850  
## 2 High School   7274  
## 3 Junior College 1161  
## 4 Bachelor     2767  
## 5 Graduate     1556  
## 6 <NA>         2
```

```
levels(gss_sub$degree)
```

```
## [1] "Lt High School" "High School"    "Junior College" "Bachelor"  
## [5] "Graduate"
```

More on factors

Make the **NA** values an explicit level

```
gss_sub %>%  
  mutate(degree_na = fct_explicit_na(degree)) %>%  
  count(degree_na)
```

```
## # A tibble: 6 × 2  
##   degree_na      n  
##   <ord>      <int>  
## 1 Lt High School 1850  
## 2 High School   7274  
## 3 Junior College 1161  
## 4 Bachelor      2767  
## 5 Graduate      1556  
## 6 (Missing)      2
```

More on factors

Relevel by frequency

```
gss_sub %>%  
  mutate(degree_freq = fct_infreq(degree)) %>%  
  count(degree_freq)
```

```
## # A tibble: 6 × 2  
##   degree_freq      n  
##   <ord>         <int>  
## 1 High School    7274  
## 2 Bachelor       2767  
## 3 Lt High School 1850  
## 4 Graduate       1556  
## 5 Junior College 1161  
## 6 <NA>           2
```

More on **factors**

Relevel manually

```
is.ordered(gss_sub$sex)
```

```
## [1] FALSE
```

```
levels(gss_sub$sex)
```

```
## [1] "Male"  "Female"
```

More on factors

```
summary(lm(age ~ sex, data = gss_sub))
```

```
##
## Call:
## lm(formula = age ~ sex, data = gss_sub)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -31.431 -13.972  -0.431  12.569  40.028
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  48.9720     0.2149  227.846  <2e-16 ***
## sexFemale    0.4594     0.2864   1.604    0.109
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 17.08 on 14463 degrees of freedom
## (145 observations deleted due to missingness)
## Multiple R-squared:  0.0001779,    Adjusted R-squared:  0.0001088
## F-statistic: 2.573 on 1 and 14463 DF,  p-value: 0.1087
```


More on factors

Relevel manually

```
gss_sub <- gss_sub %>%  
  mutate(sex = fct_relevel(sex, "Female"))  
  
levels(gss_sub$sex)
```

```
## [1] "Female" "Male"
```

More on factors

Relevel manually

```
gss_sub <- gss_sub %>%  
  mutate(sex = fct_relevel(sex, "Female"))  
  
levels(gss_sub$sex)
```

```
## [1] "Female" "Male"
```

```
summary(lm(age ~ sex, data = gss_sub))
```

```
##  
## Call:  
## lm(formula = age ~ sex, data = gss_sub)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max   
## -31.431 -13.972  -0.431  12.569  40.028   
##  
## Coefficients:  
##              Estimate Std. Error t value Pr(>|t|)      
## (Intercept)  49.4313     0.1892  261.233  <2e-16 ***  
## sexMale      -0.4594     0.2864   -1.604    0.109      
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 17.08 on 14463 degrees of freedom  
## (145 observations deleted due to missingness)  
## Multiple R-squared:  0.0001779,    Adjusted R-squared:  0.0001088   
## F-statistic: 2.573 on 1 and 14463 DF,  p-value: 0.1087
```

More on factors

Interact or cross factors

```
gss_sub <- gss_sub %>%  
  mutate(degree_by_race = fct_cross(race, degree))  
  
gss_sub %>%  
  count(degree_by_race)
```

```
## # A tibble: 16 × 2  
##   degree_by_race      n  
##   <fct>          <int>  
## 1 White:Lt High School 1188  
## 2 Black:Lt High School  379  
## 3 Other:Lt High School  283  
## 4 White:High School    5548  
## 5 Black:High School    1180  
## 6 Other:High School     546  
## 7 White:Junior College  885  
## 8 Black:Junior College  206  
## 9 Other:Junior College   70  
## 10 White:Bachelor      2334  
## 11 Black:Bachelor       233  
## 12 Other:Bachelor       200  
## 13 White:Graduate      1293  
## 14 Black:Graduate       116  
## 15 Other:Graduate       147  
## 16 <NA>                 2
```

More on factors

Relevel manually by lumping ... the least frequent n

```
gss_sub %>%  
  mutate(degree_n = fct_lump_n(degree, n = 3)) %>%  
  count(degree_n)
```

```
## # A tibble: 5 × 2  
##   degree_n      n  
##   <ord>      <int>  
## 1 Lt High School 1850  
## 2 High School    7274  
## 3 Bachelor       2767  
## 4 Other          2717  
## 5 <NA>           2
```

More on factors

Relevel manually by lumping ...to other, manually

```
gss_sub %>%  
  mutate(degree_o = fct_other(degree,  
                              keep = c("Lt High School",  
                                       "High School")) %>%  
  count(degree_o)
```

```
## # A tibble: 4 × 2  
##   degree_o      n  
##   <ord>      <int>  
## 1 Lt High School 1850  
## 2 High School   7274  
## 3 Other        5484  
## 4 <NA>          2
```