Week 2 Lab

Data Wrangling Using TidyVerse

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Goals for Today

- 1. Install and use packages
- 2. Use readr to load data
- 3. Use dplyr to manipulate data
- 4. Use tidyr to clean data
- 5. Use tibble to generate data

Next Week

• Use ggplot2 to visualize data

Install Packages

• Two ways to install and use packages

```
install.packages("tidyverse")
library(tidyverse)
```

install.packages("pacman")
p_load(tidyverse)

- 8 packages included in tidyverse
- dplyr
- readr
- tibble
- tidyr

- ggplot2
- forcats
- purrr
- stringr

Readr

• readr allows us to load data to R

```
cereal ← read_csv("cereal.csv")
```

• If you want to load excel data you will need to either save it as .csv file or use readxl package

Loading/Filtering Data

Usually when using dataframes, we need to get our hands dirty. We will evaluate our options with base functions before using functions from dplyr.

It may well be the case that there is far more data available than we will need. Three options;

- Cherry pick variables from the source,
- Trim variables from the file,
- Load entire file on R and trim down.

Loading/Filtering Data

```
p_load(gapminder)
head(gapminder)
```

```
#> # A tibble: 6 x 6
#>
    country
                continent year lifeExp
                                             pop gdpPercap
    <fct>
                <fct>
                                  <dbl>
                                           <int>
                                                     <dhl>
#>
                          <int>
#> 1 Afghanistan Asia
                           1952
                                   28.8 8425333
                                                      779.
#> 2 Afghanistan Asia
                           1957
                                   30.3 9240934
                                                      821.
  3 Afghanistan Asia
                           1962
                                   32.0 10267083
                                                      853.
#> 4 Afghanistan Asia
                           1967
                                                      836.
                                   34.0 11537966
#> 5 Afghanistan Asia
                           1972
                                                      740.
                                   36.1 13079460
#> 6 Afghanistan Asia
                                                      786.
                           1977
                                   38.4 14880372
```

Let's see some common dplyr functions using the gapminder dataframe.

Loading/Filtering Data

To generate a variable in your dataframe use %>% mutate()

To filter out particular rows from your dataframe use %>% filter()

```
EurAsia ← data_lnGDP %>% filter(continent %in% c("Asia", "Europe"))
# How many countries did I remove?
length(unique(gapminder$country)) - length(unique(EurAsia$country))
```

To summarize by groups, combine %>% group_by() and %>% summarize desc places arrange variables in descending order

```
sum_EurAsia ← EurAsia %>% group_by(country) %>% summarise(
    avg_pop = mean(pop),avg_gdp = mean(GDP)) %>% arrange(desc(avg_gdp))
sum_EurAsia
```

Binding

Binding vectors

Consider rbind and cbind: they treat the inputs as either rows or columns, and then binds them together.

```
name \(
\tau c("Pam", "George", "Sandy")
favorite \(
\tau c("Glazed Yams", "Leeks", "Daffodils")

# What are the dimensions of these?
rbind(name, favorite)
cbind(name, favorite)
```

rbind yields a 2x3

cbind yields a 3x2

Binding data frames

name work

#> [2,] "George" NA

#> [1,] "Pam" "Bus Driver"

#> [3,] "Sandy" "Shopkeeper"

#>

You can also use rbind and cbind to bind data frames.

```
# Create some data frames for us to work with
name_fav ← cbind(name, favorite)
name_work ← cbind(name, work = c("Bus Driver", NA, "Shopkeeper"))
name_fav

#> name favorite
#> [1,] "Pam" "Glazed Yams"
#> [2,] "George" "Leeks"
#> [3,] "Sandy" "Daffodils"
name_work
```

Binding data frames

cbind treats the objects as columns, so they're put side-by-side:

[A,B]

```
cbind(name_fav, name_work)
```

```
#> name favorite name work
#> [1,] "Pam" "Glazed Yams" "Pam" "Bus Driver"
#> [2,] "George" "Leeks" "George" NA
#> [3,] "Sandy" "Daffodils" "Sandy" "Shopkeeper"
```

Binding data frames

rbind treats the objects as rows, so they're stacked:

 $\begin{bmatrix} A \\ B \end{bmatrix}$

rbind(name_fav, name_work) #notice how rbind doesn't care about column names

```
#> name favorite
#> [1,] "Pam" "Glazed Yams"
#> [2,] "George" "Leeks"
#> [3,] "Sandy" "Daffodils"
#> [4,] "Pam" "Bus Driver"
#> [5,] "George" NA
#> [6,] "Sandy" "Shopkeeper"
```

Binding data frames

dplyr has very similar functions bind_rows and bind_cols. They work best with tibbles, so we'll go ahead and create tibble versions of our data.

```
name_fav_tib ← as_tibble(name_fav)
name_work_tib ← as_tibble(name_work)
```

```
#> # A tibble: 3 x 2
                                         #> # A tibble: 3 x 2
         favorite
#>
                                                     work
    name
                                         #>
                                              name
    <chr> <chr>
                                              <chr> <chr>
#>
                                         #>
#> 1 Pam Glazed Yams
                                         #> 1 Pam Bus Driver
#> 2 George Leeks
                                         #> 2 George <NA>
#> 3 Sandy Daffodils
                                         #> 3 Sandy Shopkeeper
```

Binding data frames

```
bind cols(name fav tib, name work tib
                                            bind rows(name fav tib, name work tib
#> # A tibble: 3 x 4
                                           #> # A tibble: 6 x 3
#>
     name ... 1 favorite
                          name ... 3 work
                                           #>
                                                name
                                                       favorite
                                                                   work
                                                                   <chr>
#>
     <chr>
              <chr>
                          <chr>
                                   <chr>
                                           #>
                                                <chr> <chr>
          Glazed Yams Pam
                                   Bus Driv#x 1 Pam
                                                       Glazed Yams <NA>
#> 1 Pam
            Leeks
                                           #> 2 George Leeks
#> 2 George
                          George
                                   <NA>
                                                                   <NA>
#> 3 Sandv
              Daffodils
                          Sandv
                                   Shopkeepter 3 Sandy
                                                       Daffodils
                                                                   <NA>
                                           #> 4 Pam
                                                       <NA>
                                                                   Bus Driver
                                           #> 5 George <NA>
                                                                   <NA>
                                           #> 6 Sandy
                                                                   Shopkeeper
                                                       <NA>
```

Set Operations

The dplyr set operation functions are union, intersect, and setdiff.

These set operations treat observations (rows) as if they were set elements.

```
table 1 ← tribble(
 ~"name", ~"favorites",
 #----
 "Pam", "Glazed Yams",
 "George", "Leeks",
  "Sandy", "Daffodils"
table 2 ← tribble(
 ~"name", ~"favorites",
 #----
 "Pam", "Glazed Yams",
 "Gus", "Fish Tacos"
```

Set Operations

Create tibbles using an easier to read row-by-row layout. This is useful for small tables of data where readability is important

```
table 2
table 1
#> # A tibble: 3 x 2
                                          #> # A tibble: 2 x 2
          favorites
                                               name favorites
#>
    name
                                          #>
#>
    <chr> <chr>
                                               <chr> <chr>
                                          #>
           Glazed Yams
                                          #> 1 Pam Glazed Yams
#> 1 Pam
                                          #> 2 Gus Fish Tacos
#> 2 George Leeks
           Daffodils
#> 3 Sandy
```

Set Operations

union will give you all the observations (rows) that appear in either or both tables. This is similar to bind_rows, but union will remove duplicates.

```
union(table_1, table_2)
#> # A tibble: 4 x 2
```

```
#> # A CIDDLE: 4 x 2
#> name favorites
#> <chr> <chr>
#> 1 Pam Glazed Yams
#> 2 George Leeks
#> 3 Sandy Daffodils
#> 4 Gus Fish Tacos
```

Set Operations

intersect will give you only the observations that appear both in table_1 and in table_2: in the intersection of the two tables.

```
intersect(table_1, table_2)
#> # A tibble: 1 x 2
```

```
#> # A tibble: 1 x 2
#> name favorites
#> <chr> <chr> #> 1 Pam Glazed Yams
```

#> 1 George Leeks

#> 2 Sandy Daffodils

Set Operations

setdiff(table_1, table_2) gives you all the observations in table_1 that are not in table_2.

```
setdiff(table_1, table_2)

#> # A tibble: 2 x 2

#> name favorites

#> <chr> <chr>
```

Mutating joins

Mutating joins take the first table and add columns from the second table. There are 3 mutating joins: left_join, inner_join, and full_join.

```
# We'll create 2 new data frames to learn mutating joins:
favorites ← tribble(
 ~"name", ~"fav",
 #----
 "Pam", "Glazed Yams",
 "George", "Leeks",
 "Sandy", "Daffodils"
jobs ← tribble(
 ~"name", ~"work",
 #----
 "Pam", "Bus Driver",
 "Gus", "Bartender",
 "Sandy", "Shopkeeper"
                                                                           19 / 33
```

Mutating joins

left_join(x, y) takes x and adds the columns of y where the **key** matches. The **key** is a variable that shows up in both tables and you'll specify it with by = "key_variable".

```
left_join(favorites, jobs, by = "name

#> # A tibble: 3 x 3

#> name fav work

#> <chr> <chr> <chr>
#> 1 Pam Glazed Yams Bus Driver

#> 2 George Leeks <NA>

#> 3 Sandy Daffodils Shopkeeper
```

```
# What will be the output?
left_join(jobs, favorites, by = "name")
```

Mutating joins

left_join(x, y) takes x and adds the columns of y where the key
matches. The key is a variable that shows up in both tables and you'll
specify it with by = "key_variable".

```
left_join(favorites, jobs, by = "name")
#> # A tibble: 3 x 3
#>
    name
           fav
                       work
    <chr> <chr>
                       <chr>
#>
           Glazed Yams Bus Driver
#> 1 Pam
#> 2 George Leeks
                       <NA>
#> 3 Sandy
           Daffodils
                       Shopkeeper
```

```
# What will be the output?
left_join(jobs, favorites, by = "name

#> # A tibble: 3 x 3
#> name work fav
#> <chr> <chr> <chr> #> 1 Pam Bus Driver Glazed Yams
#> 2 Gus Bartender <NA>
#> 3 Sandy Shopkeeper Daffodils
```

Mutating joins

inner_join(x, y) takes the **intersect** of the key variable and adds columns from both tables.

Mutating joins

full_join(x, y) takes the **union** of the key variable and adds columns from both tables.

```
full_join(favorites, jobs, by = "name")

#> # A tibble: 4 x 3

#> name fav work

#> <chr> <chr> <chr>
#> 1 Pam Glazed Yams Bus Driver

#> 2 George Leeks <NA>

#> 3 Sandy Daffodils Shopkeeper

#> 4 Gus <NA> Bartender
```

Filtering joins

Unlike mutating joins, filtering joins will only preserve data from the first table. The observations that are kept depends on the second table. dplyr has 2 types of filtering joins: semi_join and anti_join.

```
semi_join(x, y) keeps all rows in x where the key matches in y.
```

```
semi_join(favorites, jobs, by = "name")
```

```
#> # A tibble: 2 x 2
#> name fav
#> <chr> <chr>
#> 1 Pam Glazed Yams
#> 2 Sandy Daffodils
```

Filtering joins

anti_join(x, y) keeps rows in x as long as the key **doesn't** have a match in y.

```
anti_join(favorites, jobs, by = "name")

#> # A tibble: 1 x 2

#> name fav

#> <chr> <chr>
#> 1 George Leeks
```

Pivoting

pivot_wider() and pivot_longer() aren't two-table topics, but they are useful data manipulation tools in the tidyverse.

```
prefs ← tribble(
    ~"name", ~"preference", ~"item",
    #/----/-----/
    "Pam", "loves", "Glazed Yams",
    "Pam", "likes", "Daffodils",
    "Pam", "hates", "Horseradish",
    "George", "loves", "Leeks",
    "George", "likes", "Hazelnuts",
    "George", "hates", "Dandelions"
)
```

Take a look at the data. There are 2 people (Pam and George). Each person has one "love", one "like", and one "hate" item.

Pivoting

Suppose instead we wanted our data in a different format. What if we had 4 columns instead of 3: name, the thing that person loves, the thing that person likes, and the thing that person hates. We'd only need 2 rows (Pam and George).

We want our data to go from having 3 columns to having 4, so we know we can use tidyr::pivot_wider.

```
pivot_wider(prefs, names_from = preference, values_from = item)
```

```
#> # A tibble: 2 x 4

#> name loves likes hates

#> <chr> <chr> <chr> #> 1 Pam Glazed Yams Daffodils Horseradish
#> 2 George Leeks Hazelnuts Dandelions
```

Pivoting

Now suppose we want to reverse that operation! We'll start with prefs_wide and pivot in to get prefs again.

pivot_longer() has these arguments:

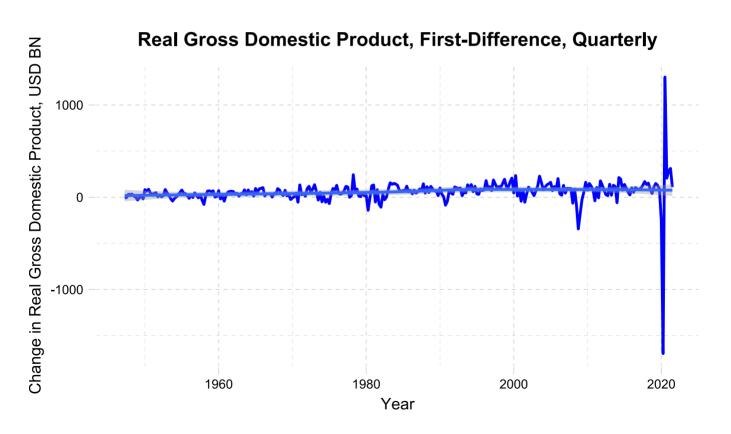
- **cols**: columns to pivot into the longer format. For us, that will be the columns loves, likes, and hates. We can also say columns 2 through 4: cols = 2:4.
- names_to: A string. What we should call the new column that holds
 those old column names: loves, likes, hates: names_to =
 "preferences"
- values_to: A string. what we should call the values that are now being pivoted in? Glazed Yams, Daffodils, etc. So we want values_to = "items"

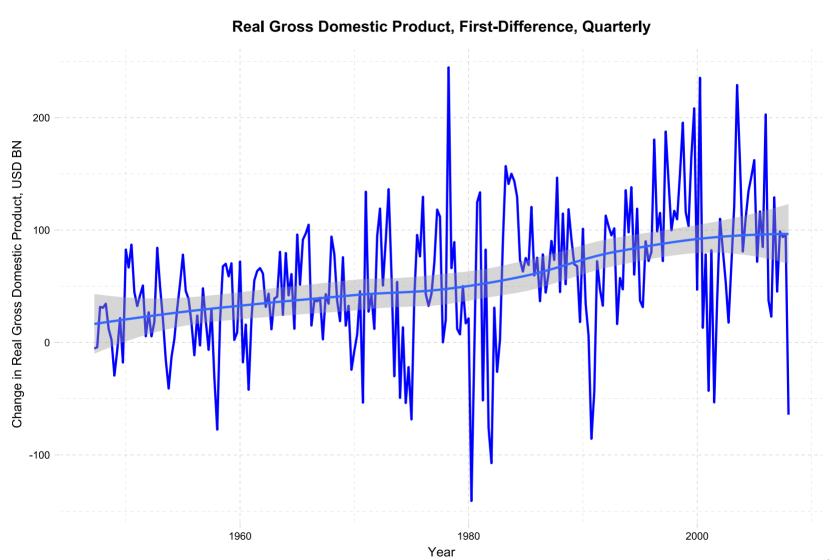
Pivoting

```
prefs_wide %>% pivot_longer(cols = 2:4, names_to = "preferences", values_to = "i
```

```
#> # A tibble: 6 x 3
           preferences items
#>
    name
   <chr> <chr>
                      <chr>
#>
#> 1 Pam
        loves
                      Glazed Yams
#> 2 Pam
        likes
                      Daffodils
#> 3 Pam
         hates
                      Horseradish
#> 4 George loves
                  Leeks
#> 5 George likes
                      Hazelnuts
                      Dandelions
#> 6 George hates
```

Outliers may also be present in data. In macroeconomics, one may be trying to assess mean and variance of real gross domestic product in the United States.





Consider the 1.5 IQR rule of thumb. This is used to identify mild outliers. For extreme outliers only, shift to a 3 IQR.

- IQR = 75th Percentile Value 25th Percentile Value
- Lower Outlier Boundary = 25th 1.5*IQR
- Upper Outlier Boundary = 75th + 1.5*IQR

```
#Extreme Outliers identified and cleaned
Phase1 ← summary(data$columnX)
OutLower ← Phase1[2]-3*(Phase1[5]-Phase1[2])
OutHigher ← Phase1[5]+3*(Phase1[5]-Phase1[2])
house_w ← filter(house, columnX > OutLower)
house_w ← filter(house, columnX < OutHigher)</pre>
```

See example of extreme outlier cleaning in Davies, R., & T., Jeppesen, 2015 "Export mode, firm heterogeneity, and source country characteristics, Review of World Economics, Vol. 151(2), pp 169-195.

Resources

- Datacamp: Joining data with dplyr
- RStudio dplyr Cheat Sheet
- Tidyverse