Data Summarization

Recap

- select(): subset and/or reorder columns
- filter(): remove rows
- arrange(): reorder rows
- mutate(): create new columns or modify them
- select() and filter() can be combined together
- remove a column: select() with ! mark(!col_name)
- you can do sequential steps: especially using pipes %>%

Day 3 Cheatsheet

The Data

We can use the CO heat-related ER visits dataset to explore different ways of summarizing data.

Reminder: This dataset contains information about the number and rate of visits for heat-related illness to ERs in Colorado from 2011-2022, adjusted for age.

The Data

We can use the CO heat-related ER visits dataset to explore different ways of summarizing data.

The head function displays the first rows of an object:

```
er <-
 read_csv("../../data/CO_ER_heat_visits.csv")
head(er)
# A tibble: 6 × 6
 county rate lower95cl upper95cl visits year
 <chr> <dbl>
                 <dbl>
                           <dbl> <dbl> <dbl>
1 Adams
        6.73
                            9.24
                                    29 2011
                 NA
2 Adams
        4.84
                2.85
                                    23 2012
                           NA
3 Adams
       6.84
                         9.31
                 4.36
                                    31 2013
4 Adams
                  1.71
                         4.85
                                    15 2014
        3.08
5 Adams
        3.36
                  1.89
                            5.23
                                    16 2015
6 Adams
         8.85
                                    42 2016
                  6.12
                           11.6
```

Behavior of pull() function

pull() converts a single data column into a vector.

er %>% pull(visits)

Data Summarization

Now that we have a vector of numbers.. what can we do with it?

- Basic statistical summarization
 - mean(x): takes the mean of x
 - sd(x): takes the standard deviation of x
 - median(x): takes the median of x
 - quantile(x): displays sample quantiles of x. Default is min, IQR, max
 - range(x): displays the range. Same as c(min(x), max(x))
 - sum(x): sum of x
 - max(x): maximum value in x
 - min(x): minimum value in x

Pipe (%>%) vectors into summarizing functions

A vector can be summarized:

```
er %>% pull(visits) %>% mean()
[1] NA
```

Add the na.rm = argument for missing data

```
er %>% pull(visits) %>% mean(na.rm=T)
[1] 7.189247
```

GUT CHECK!

What kind of object do we need to run summary operators like mean()?

- A. A vector of numbers
- B. A vector of characters
- C. A dataset

Summarization on tibbles (data frames)

summarize works on datasets without pull().

Multiple summary statistics can be calculated at once!

summarize() can do multiple operations at once. Just separate by a comma.

Note that summarize() creates a separate tibble from the original data.

If you want to save a summary statistic in the original data, use mutate() instead to create a new column for the summary statistic.

summary() Function

Using summary() can give you rough snapshots of each numeric column (character columns are skipped):

summary(er)

| county | rate | lower95cl | upper95cl |
|------------------|----------------|----------------|----------------|
| Length: 768 | Min. : 0.000 | Min. : 0.000 | Min. : 0.000 |
| Class :character | 1st Qu.: 0.000 | 1st Qu.: 0.000 | 1st Qu.: 0.000 |
| Mode :character | Median : 0.000 | Median : 0.000 | Median : 0.000 |
| | Mean : 2.431 | Mean : 1.449 | Mean : 3.526 |
| | 3rd Qu.: 3.509 | 3rd Qu.: 1.895 | 3rd Qu.: 5.173 |
| | Max. :89.275 | Max. :43.398 | Max. :151.420 |
| | NA's :303 | NA's :304 | NA's :304 |
| visits | year | | |
| Min. : 0.000 | Min. :2011 | | |
| 1st Qu.: 0.000 | 1st Qu.:2014 | | |
| Median : 0.000 | Median :2016 | | |
| Mean : 7.189 | Mean :2016 | | |
| 3rd Qu.:13.000 | 3rd Qu.:2019 | | |
| Max. :48.000 | Max. :2022 | | |
| NA's :303 | | | |

Summary & Lab Part 1

- pull() creates a vector
- don't forget the na.rm = TRUE argument!
- summary(x): quantile information
- summarize: creates a summary table of columns of interest
- summary stats (mean()) work with vectors or with summarize()
- Class Website
- Lab

distinct() values

distinct(x) will return the unique elements of column x.

```
er %>%
  distinct(county)
# A tibble: 64 \times 1
   county
   <chr>
 1 Adams
 2 Alamosa
 3 Arapahoe
 4 Archuleta
 5 Baca
 6 Bent
 7 Boulder
 8 Broomfield
 9 Chaffee
10 Cheyenne
# 🛮 54 more rows
```

How many distinct() values?

n_distinct() tells you the number of unique elements.

It needs a vector so you must pull the column first!

```
er %>%
  pull(county) %>%
  n_distinct()

[1] 64
```

Use count () to return row count per category.

```
er %>% count(county)
# A tibble: 64 \times 2
   county
                  n
   <chr>
              <int>
 1 Adams
                 12
 2 Alamosa
                 12
 3 Arapahoe
                 12
 4 Archuleta
                 12
 5 Baca
                 12
 6 Bent
                12
 7 Boulder
                 12
8 Broomfield
                12
 9 Chaffee
                 12
10 Cheyenne
                 12
# 0 54 more rows
```

Looks like 12 rows/observations per county!

Multiple columns listed further subdivides the count ()

```
er %>% count(county, year)
# A tibble: 768 × 3
   county year
                    n
   <chr> <dbl> <int>
 1 Adams
           2011
                    1
 2 Adams
           2012
 3 Adams
           2013
 4 Adams
           2014
                    1
 5 Adams
           2015
 6 Adams
           2016
 7 Adams
           2017
 8 Adams
           2018
 9 Adams
           2019
10 Adams
           2020
# 0 758 more rows
```

Looks like 1 row/observation per county and year!

GUT CHECK!

The count () function can help us tally:

- A. Sample size
- B. Rows per each category
- C. How many categories

Grouping

Goal

We want to find the mean number of ER visits per year in the dataset.

How do we do this?

Perform Operations By Groups: dplyr

First, let's group the data.

group_by allows you group the data set by variables/columns you specify:

```
er grouped <- er %>% group by(year)
er grouped
# A tibble: 768 × 6
          year [12]
# Groups:
  county rate lower95cl upper95cl visits year
  <chr> <dbl>
                   <dbl>
                             <dbl> <dbl> <dbl>
 1 Adams
          6.73
                   NA
                              9.24
                                       29
                                          2011
 2 Adams
          4.84
                    2.85
                                       23 2012
                             NA
 3 Adams
          6.84
                    4.36
                                       31 2013
                              9.31
 4 Adams
          3.08
                    1.71
                              4.85
                                       15 2014
 5 Adams
          3.36
                                       16 2015
                    1.89
                              5.23
 6 Adams
          8.85
                    6.12
                             11.6
                                       42
                                           2016
 7 Adams
          6,63
                    4.29
                                       32
                                           2017
                          8.98
 8 Adams
          7.11
                    4.77
                              9.44
                                       37 2018
 9 Adams
          6.76
                    4.53
                              8.99
                                       36 2019
10 Adams
          4.76
                    2.82
                              6.70
                                       24
                                           2020
# 0 758 more rows
```

Summarize the grouped data

It's grouped! Grouping doesn't change the data in any way, but how **functions operate on it**. Now we can summarize **visits** by group:

```
er_grouped %>%
 summarize(avg_visits = mean(visits, na.rm = TRUE))
# A tibble: 12 × 2
   year avg_visits
  <dbl>
            <dbl>
1 2011
             5.20
2 2012
             5.89
3 2013
             5.63
4 2014
             4.12
5 2015
             6.4
6 2016
             10.1
7 2017
            7.24
  2018
            11.7
  2019
          9.12
10 2020
             6.26
11 2021
             8.06
12 2022
             9.29
```

Do it in one step: use %>% to string these together!

Pipe CO_heat_ER into group_by, then pipe that into summarize:

```
er %>%
 group_by(year) %>%
 summarize(avg_visits = mean(visits, na.rm = TRUE))
# A tibble: 12 × 2
   year avg_visits
  <dbl>
            <dbl>
1 2011
             5.20
2 2012 5.89
 3 2013
             5.63
4 2014
             4.12
  2015
           6.4
6 2016
            10.1
7 2017
            7.24
  2018
            11.7
  2019
          9.12
10 2020
         6.26
11 2021
             8.06
12 2022
             9.29
```

Group by as many variables as you want

group_by county and year:

```
er %>%
 group_by(year, county) %>%
  summarize(avg_visits = mean(visits, na.rm = TRUE))
# A tibble: 768 × 3
# Groups: year [12]
   year county avg_visits
  <dbl> <chr>
                        <dbl>
1 2011 Adams
                           29
 2 2011 Alamosa
                            0
 3 2011 Arapahoe
                           33
 4 2011 Archuleta
                            0
   2011 Baca
                            0
   2011 Bent
                            0
   2011 Boulder
                           12
   2011 Broomfield
                          NaN
 9 2011 Chaffee
                            0
10 2011 Cheyenne
                            0
# 0 758 more rows
```

Counting rows/observations

There are other summarizing functions, such as n() count the number of rows/observations (NAs included).

```
er %>%
 group_by(year) %>%
 summarize(n = n(),
          mean = mean(visits, na.rm = TRUE))
# A tibble: 12 × 3
   vear
           n mean
  <dbl> <int> <dbl>
1 2011
          64 5.20
2 2012 64 5.89
3 2013
         64 5.63
         64 4.12
4 2014
 5 2015
         64 6.4
6 2016
         64 10.1
  2017
        64 7.24
8
   2018
         64 11.7
   2019 64 9.12
10 2020
         64 6.26
11 2021
         64 8.06
12 2022
          64 9.29
```

Counting: count() and n()

count() and n() can give very similar information.

```
# Here we use count()
er %>% count(year)
# A tibble: 12 × 2
   year
            n
   <dbl> <int>
 1 2011
            64
 2 2012
            64
   2013
            64
   2014
            64
   2015
            64
 6
   2016
            64
   2017
            64
   2018
            64
 8
   2019
            64
 9
10 2020
            64
   2021
11
            64
12 2022
            64
```

Counting: count() and n()

count() and n() can give very similar information.

```
# n() with summarize
er %>% group_by(year) %>% summarize(n())
# A tibble: 12 × 2
   year `n()`
  <dbl> <int>
 1 2011
           64
 2 2012
           64
 3 2013
          64
 4 2014
          64
 5 2015
           64
   2016
           64
   2017
           64
   2018
           64
   2019
           64
 9
10 2020
           64
11 2021
           64
12 2022
           64
```

A few miscellaneous topics ..

Base R functions you might see: length and unique

These functions require a column as a vector using pull().

```
er_year <- er %>% pull(year) # pull() to make a vector
er_year %>% unique() # similar to distinct()
[1] 2011 2012 2013 2014 2015 2016 2017 2018 2019 2020 2021 2022
```

Base R functions you might see: length and unique

These functions require a column as a vector using pull().

```
er_year %>% unique() %>% length() # similar to n_distinct()
[1] 12
```

summary() vs. summarize()

- summary() (base R) gives statistics table on a dataset.
- summarize() (dplyr) creates a more customized summary tibble/dataframe.

Summary

- count(x): what unique values do you have?
 - distinct(): what are the distinct values?
 - n_distinct() with pull(): how many distinct values?
- group_by(): changes all subsequent functions
 - combine with summarize() to get statistics per group
- summarize() with n() gives the count (NAs included)

Lab Part 2

- Class Website
- Lab
- Day 4 Cheatsheet
- Posit's data transformation Cheatsheet

For more advanced learning:

- https://www.danieldsjoberg.com/gtsummary/ for tables
- · extra slides in this file.



Image by Gerd Altmann from Pixabay

Extra Slides: More advanced summarization

Data Summarization on data frames

- Statistical summarization across the data frame
 - rowMeans(x): takes the means of each row of x
 - colMeans(x): takes the means of each column of x
 - rowSums(x): takes the sum of each row of x
 - colSums(x): takes the sum of each column of x

```
yearly_co2 <- read_csv("../../data/Yearly_C02_Emissions_1000_tonnes.csv")</pre>
```

rowMeans() example

Get means for each row.

Let's see what the mean CO2 emissions is across the years 2010-2014 for each row (country):

```
yearly_co2 %>%
  select(starts_with("201")) %>%
  rowMeans(na.rm = TRUE) %>%
  head(n = 5)
             5106 129800
                           487 32040
[1]
     10254
yearly_co2 %>%
  group_by(country) %>%
  summarize(mean = rowMeans(across(starts_with("201")), na.rm = TRUE)) %>%
  head(n = 5)
# A tibble: 5 \times 2
  country
                mean
  <chr>
               <dbl>
1 Afghanistan 10254
2 Albania
                5106
3 Algeria
              129800
4 Andorra
                 487
5 Angola
               32040
```

colMeans() example

Get means for each column.

Let's see what the mean is across each column (year):

```
yearly_co2 %>%
  select(starts_with("201")) %>%
 colMeans(na.rm = TRUE) %>%
 head(n = 5)
    2010
                     2012
            2011
                              2013
                                        2014
165334.1 171764.9 174033.4 174856.2 175992.5
yearly_co2 %>%
  summarize(across(starts_with("201"), ~mean(.x, na.rm = TRUE)))
# A tibble: 1 \times 5
   `2010` `2011` `2012` `2013` `2014`
    <dbl> <dbl> <dbl> <dbl>
                                   <dbl>
1 165334, 171765, 174033, 174856, 175993,
```

* New! * Many dplyr functions now have a .by= argument

Pipe er into group_by, then pipe that into summarize: