Lec 7: Relationships between two categorical variables (Two-way tables)

Instructor: Tomer Altman

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Learning objectives for today

- How to visualize and quantify relationships between two categorical variables
- $\bullet\,$ Two-way tables: marginal vs. conditional distributions
- Bar graphs: side by side vs. stacked
- Simpson's paradox

Readings

- Chapter 5 of Baldi & Moore
- Bar Plots
- Clustered Bar Charts

Two-way tables

- Two-way stands for 2×2 , as in a table with two columns and two rows
- Used to examine the relationship between two categorical variables, originally those with two levels
- Foundational to epidemiology, because of the types of variables we are often interested in

Classic 2×2 table format

Exposure group	Disease	No disease	Row total
Exposed	A	В	A+B
Not Exposed	\mathbf{C}	D	C+D
Column total	A+C	B+D	A+B+C+D

Example: Lung cancer and smoking

Group	Lung Cancer	No Lung Cancer	Row total
Smoker	12	238	250
Non-smoker	7	743	750
Column total	19	981	1000

Marginal distribution

- The **marginal distribution** of a variable is the one that is **in the margin** of the table (i.e., the Row total or the Column total are the two margins of a two-way table)
- The marginal distribution is the distribution for a single categorical variable
- We learned previously how to plot marginal distributions of categorical variables using geom_bar()

Marginal distribution

Group	Lung Cancer	No Lung Cancer	Row total
Smoker	12	238	250
Non-smoker	7	743	750
Column total	19	981	1000

- Overall, what % of the population has lung cancer?
 - Answer:
- Overall, what % of the population are smokers?
 - Answer:

Marginal distribution

Group	Lung Cancer	No Lung Cancer	Row total
Smoker	12	238	250
Non-smoker	7	743	750
Column total	19	981	1000

- Overall, what % of the population has lung cancer?
 - Answer: 19/1000 = 1.9%
- Overall, what % of the population are smokers?
 - Answer: 250/1000 25% smoking
- The marginal distribution of lung cancer is 1.9% lung cancer, 98.1% no lung cancer.

Conditional distribution

Group	Lung Cancer	No Lung Cancer	Row total
Smoker	12	238	250
Non-smoker	7	743	750
Column total	19	981	1000

- The **conditional distribution** is the distribution of one variable **within** or **conditional on** the level of a second variable
- What is the conditional distribution of lung cancer **given** smoking?
 - Answer:
- What is the conditional distribution of lung cancer **given** non-smoking?
 - Answer:

Conditional distribution

Group	Lung Cancer	No Lung Cancer	Row total
Smoker	12	238	250
Non-smoker	7	743	750
Column total	19	981	1000

- The **conditional distribution** is the distribution of one variable **within** or **conditional on** the level of a second variable
- What is the conditional distribution of lung cancer **given** smoking?
 - Answer: 12/250 = 4.8% lung cancer and 238/250 = 95.2% no lung cancer
- What is the conditional distribution of lung cancer **given** non-smoking?
 - Answer: 7/750 = 0.9% lung cancer and 743/750 = 99.1% no lung cancer

Marginal and conditional distributions in R

- We learned previously how to plot marginal distributions of categorical variables using geom bar()
- Can we generalize the use of geom_bar() to plot multiple conditional distributions? I.e., can we show the conditional distribution of lung cancer for smokers and non-smokers on the same plot?

First, we encode the data to read into R:

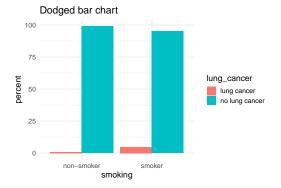
```
# students, you don't need to know how to do this
                                                        ~ percent, ~number,
two_way <- tribble(~ smoking,</pre>
                                    ~ lung_cancer,
                      "smoker",
                                       "lung cancer",
                                                          4.8,
                                                                     12,
                      "smoker",
                                       "no lung cancer", 95.2,
                                                                     238.
                      "non-smoker",
                                       "lung cancer",
                                                          0.9,
                                                                     7,
                      "non-smoker",
                                       "no lung cancer", 99.1,
                                                                     743
```

Visualization of conditional distributions

If there is an explanatory-response relationship, compare the conditional distribution of the response variable for the separate values of the explanatory variable.

Dodged bar chart for the visualization of conditional distributions

```
ggplot(two_way, aes(x = smoking, y = percent)) +
geom_bar(aes(fill = lung_cancer), stat = "identity", position = "dodge") +
labs(title = "Dodged bar chart") + theme_minimal(base_size = 15)
```

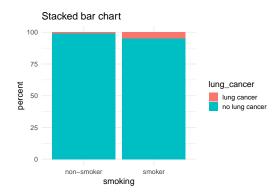


Syntax: Dodged bar chart for the visualization of conditional distributions

```
#students, remove eval=F if you copy this code chunk (or else the code won't compile)
ggplot(data, aes(x = exposure_variable, y = percent)) +
  geom_bar(aes(fill = outcome_variable), stat = "identity", position = "dodge") +
  labs(title = "Dodged bar chart") +
  theme_minimal(base_size = 15)
```

Stacked bar chart for the visualization of conditional distributions

```
ggplot(two_way, aes(x = smoking, y = percent)) +
geom_bar(aes(fill = lung_cancer), stat = "identity", position = "stack") +
labs(title = "Stacked bar chart") + theme_minimal(base_size = 15)
```



Syntax: Stacked bar chart for the visualization of conditional distributions

```
#students, remove eval=F if you copy this code chunk (or else the code won't compile)
ggplot(data, aes(x = exposure_variable, y = percent)) +
  geom_bar(aes(fill = outcome_variable), stat = "identity", position = "stack") +
  labs(title = "Stacked bar chart") +
  theme_minimal(base_size = 15)
```

Visualization of conditional distributions: three levels of response variable

- Stacked and dodged plots are less informative when there are only two levels of both variables
- This is because once you know the percent of lung cancer among smokers, you also know the percent of non-lung cancer among smokers. This makes some of the information redundant.
- The plots are more informative if there are 3 or more levels for at least one of the variables

Visualization of conditional distributions: three levels of response variable

• Example 2: Shoe support by gender (Data from Baldi & Moore page 124 of Ed.4):

Group	Men	Women	Row total
Good support	94	137	231
Average support	1348	581	1929
Poor support	30	1182	1212
Column total	1472	1900	3372

Check your understanding!

Visualization of conditional distributions: three levels of response variable

Group	Men	Women	Row total
Good support	94	137	231
Average support	1348	581	1929
Poor support	30	1182	1212
Column total	1472	1900	3372

- Example 2: Shoe support by gender (Data from Baldi & Moore page 124 of Ed.4):
- The question: How does the distribution of support of shoes worn vary between men and women?

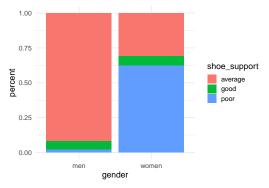
Visualization of conditional distributions: three levels of response variable

```
# students, you don't need to know how to do this
shoe_data <- tribble(~ shoe_support, ~ gender, ~ percent,</pre>
                       "good",
                                        "men",
                                                  94/1472,
                       "average",
                                        "men",
                                                  1348/1472,
                       "poor",
                                        "men",
                                                  30/1472,
                                        "women", 137/1900,
                       "good",
                       "average",
                                        "women", 581/1900,
                       "poor",
                                        "women", 1182/1900)
shoe_data
```

```
## # A tibble: 6 x 3
##
     shoe_support gender percent
##
                  <chr>
                            <dbl>
## 1 good
                  men
                           0.0639
## 2 average
                           0.916
                  men
## 3 poor
                           0.0204
                  men
## 4 good
                           0.0721
                  women
## 5 average
                           0.306
                  women
## 6 poor
                  women
                           0.622
```

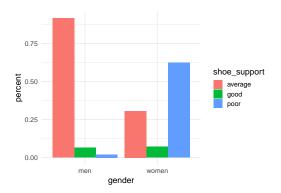
Stacked visualization when there are three levels of response

```
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "stack") +
  theme_minimal(base_size = 15)
```



Dodged visualization when there are three levels of response

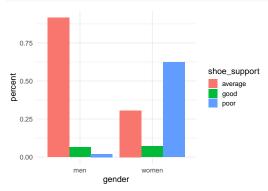
```
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "dodge") +
  theme_minimal(base_size = 15)
```



Dodged visualization when there are three levels of response

Question: what is misleading about the fill legend?

```
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "dodge") +
  theme_minimal(base_size = 15)
```



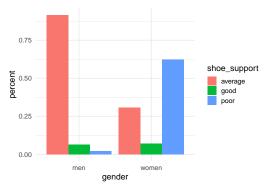
Dodged visualization when there are three levels of response

Question: what is misleading about the fill legend?

Answer: It is in alphabetic order, which is different from the natural order of this variable.

Question 2: How can we change the order in the legend?

```
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "dodge") +
  theme_minimal(base_size = 15)
```



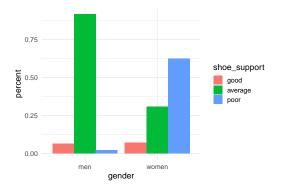
Dodged visualization when there are three levels of response

Question 2: How can we change the order in the legend?

Answer 2: Recall from the problem sets and lab how to reorder factor variables that affect the look of the plot:

```
shoe_data <- shoe_data %>%
  mutate(shoe_support = fct_relevel(shoe_support, "good", "average", "poor"))

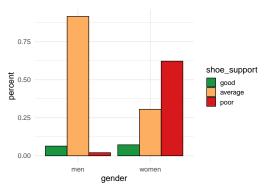
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "dodge") +
  theme_minimal(base_size = 15)
```



Dodged visualization when there are three levels of response

You might also want to specify the colors used to communicate that poor shoe support is painful!

```
ggplot(shoe_data, aes(x = gender, y = percent)) +
  geom_bar(stat = "identity", aes(fill = shoe_support), position = "dodge", col = "black") +
  theme_minimal(base_size = 15) +
  scale_fill_manual(values = c("#1a9641", "#fdae61", "#d7191c"))
```



Visualization of conditional distributions: three levels of response variable

In general, dodged plots are preferred over stacked plots. Why do you think that is?

Simpson's Paradox

Simpson's Paradox: Example from Baldi and Moore

• Let's load these data that examines mortality rates by community and age group across two communities

```
#this is the data from page 131 of edition 4 of baldi and moore
simp_data <- tribble(~ age_grp, ~ community, ~ deaths, ~ pop,</pre>
                        "0-34",
                                   "A",
                                                 20,
                                                           1000,
                                   "A",
                        "35-64",
                                                 120,
                                                           3000,
                        "65+",
                                   "A",
                                                 360,
                                                           6000,
                        "all",
                                   "A",
                                                 500,
                                                           10000,
                        "0-34",
                                   "B",
                                                180,
                                                           6000,
                                   "B",
                        "35-64",
                                                           3000,
                                                 150,
                        "65+",
                                   "B",
                                                 70,
                                                           1000,
                                                           10000)
                        "all",
                                   "B",
                                                 400,
simp_data <- simp_data %>%
  mutate(death_per_1000 = (deaths/pop) * 1000)
simp_data_no_all <- simp_data %>% filter(age_grp != "all")
```

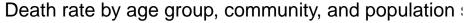
Simpson's Paradox: Example from Baldi and Moore

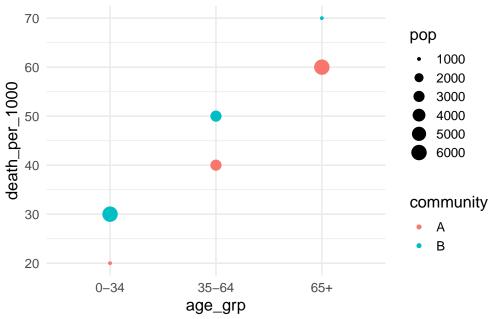
```
simp_data
## # A tibble: 8 x 5
##
     age_grp community deaths
                               pop death_per_1000
##
           <chr>
                       <dbl> <dbl>
     <chr>
## 1 0-34
            Α
                          20 1000
                                                20
## 2 35-64
                         120 3000
                                                40
            Α
## 3 65+
                         360 6000
                                                60
            Α
                         500 10000
## 4 all
            Α
                                                50
## 5 0-34
                         180 6000
                                                30
            В
                                                50
## 6 35-64
            В
                         150 3000
                                                70
## 7 65+
                          70 1000
            В
## 8 all
            В
                         400 10000
                                                40
```

Simpson's Paradox Example: Plot only the conditional data

• Plot the mortality rates according to age group and community and link the point size to population size

```
ggplot(simp_data_no_all, aes(x = age_grp, y = death_per_1000)) +
geom_point(aes(col = community, size = pop)) +
labs(title = "Death rate by age group, community, and population size") +
theme_minimal(base_size = 15)
```





Observations from this visualization:

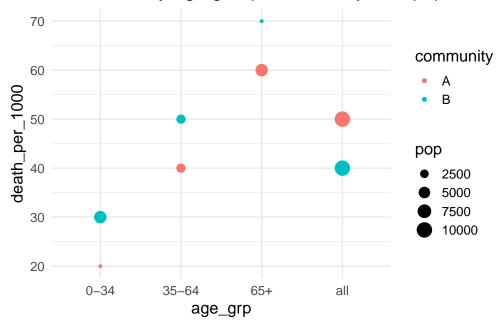
- 1.
- 2.
- 3.

If someone ask you which community has higher mortality, what would you say?

Simpson's Paradox Example: Add the marginal data

- Add in the **marginal** data (not conditional on age)
- Notice that the mortality rates for the communities overall show community A having a higher rate than community B. Why?

Death rate by age group, community, and population :

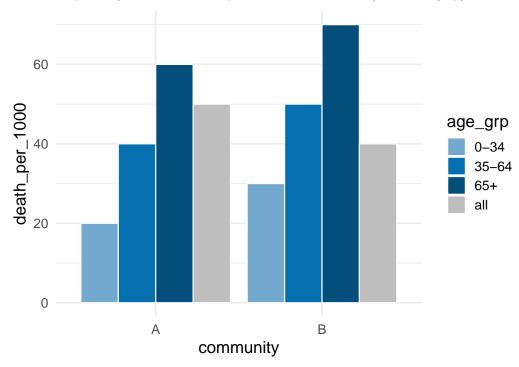


Simpson's Paradox

"An association or comparison that holds for all of several groups can **reverse direction** when the data are combined to form a single group. This reversal is called **Simpson's Paradox**"

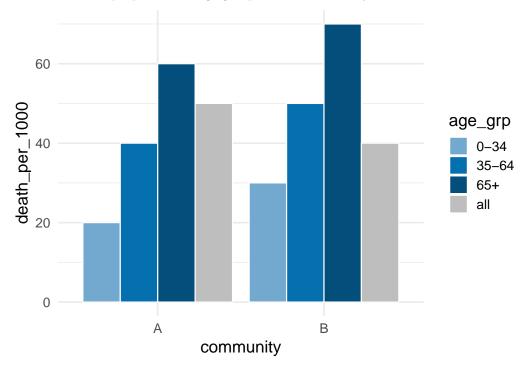
Simpson's Paradox

- Here are the same data shown using a bar chart
- Notice that the mortality rate for each of the blue-shaded bars in community B is higher than the correponding bar for community A, but the overall bar (shaded in gray) shows a reversal.



Simpson's Paradox

- With a bar chart we can't use aes(size = pop), so it is harder to see why the paradox is occurring.
- It is because we are taking a weighted average of each age-specific bar with weights proportional to the number of people of each age group in each community



Check your understanding

Simpson's Paradox in Berkeley Admissions

- There is a famous example of Simpson's paradox related to admissions to Berkeley by gender
- Watch it here!

Recap: What new code and statistical concepts did we learn?

- 1. geom_bar(aes(col = var), stat = "identity", position = "dodge")
- 2. geom_bar(aes(col = var), stat = "identity", position = "stack")
- 3. Marginal distribution vs. conditional distribution
- 4. Simpson's Paradox