Introduction to probability

Kyndall Payton

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Upcoming Deadline

- Exam due by 11:59 PM on February 22nd.
- Lab in groups on Thursday.

Main ideas

- Use formulas to compute probabilities from tabular data
- Compute empirical probabilities in R via simulation

Packages

```
library(tidyverse)
library(vcd) # used for Arthritis data
library(usethis)
use_git_config(user.name="kyndallpayton", user.email="kp256@duke.edu")
```

Computing probabilities

Take a look at the help for Arthritis to understand where this data comes from and the variable meanings.

Let's look at the data in a tabular view. Don't worry about understanding these functions, we're only using it to better visualize our data via a table.

```
xtabs(~ Treatment + Improved, data = Arthritis) %>%
addmargins()
```

```
#>
            Improved
#> Treatment None Some Marked Sum
               29
                     7
#>
    Placebo
                            7
                               43
     Treated
               13
                     7
                            21
                               41
#>
               42
                            28 84
#>
     Sum
                    14
```

- How many patients were enrolled in the clinical trial? 84
- What is the probability a randomly selected patient received the placebo? 43/84
- \bullet What is the probability a randomly selected patient received the placebo and had a marked improvement? 7/84
- What is the probability a randomly selected patient received the placebo and the treatment? 0
- What is the probability a randomly selected patient had some improvement or was on the treatment?
 48/84

Using computer simulations to calculate probabilities

Example Recall that a **vector** is the basic building block in R. Let's create a vector called marbles.

```
marbles <- c("red", "red", "white", "red", "blue", "blue", "red", "blue")
```

Suppose we draw a single marble from our imaginary box, where all the marbles are equally likely to be selected. What is the probability the marble is blue? How about white? 3/8; 1/8

We can simulate this "drawing" with the sample() function.

```
sample(marbles, size = 1)
```

```
#> [1] "blue"
```

We produced one random outcome from this experiment. To estimate the probability of say getting a white marble, we need to repeat this experiment many many times.

In the sample() function we can change the size argument and set replace = TRUE. Setting replace = TRUE allows to draw from our population of eight marbles each time. This way we can easily simulate our marble-drawing experiment.

```
draw_results <- sample(marbles, size = 10000, replace = TRUE)
counts <- table(draw_results)
prop.table(counts)</pre>
```

```
#> draw_results
#> blue red white
#> 0.3762 0.5012 0.1226
```

How close is this value to the "true" probability? Quite close as red marbles make up half the dataset and its probability for this number of draws is 49.87%. Blue makes up 37.5% of the dataset and its probability for this size is 37.66%.

To summarize our process:

- 1. We defined the sample space for our experiment ${\tt marbles}$
- 2. We simulated this experiment many many times and recorded the outcomes from each of the simulations.
- 3. We computed the relative frequency of the observed outcomes from our many simulations.

Another example What if we want to compute the probability of getting two marbles of the same color if we make two draws with replacement? We haven't discussed how to compute this theoretically yet, but this is what computers are good at.

Before we do this, what is your guess as to what the probability will be? $\sim 1/2$, fairly likely

We'll still use sample() to run our simulation many times, but we'll use dplyr functions to compute the relative frequencies.

```
two_draw_results <- tibble(</pre>
  draw_1 = sample(marbles, size = 10000, replace = TRUE),
  draw_2 = sample(marbles, size = 10000, replace = TRUE)
)
two_draw_results
#> # A tibble: 10,000 x 2
#>
      draw_1 draw_2
#>
      <chr> <chr>
#>
   1 white red
#>
    2 red
             blue
#>
   3 red
             blue
#>
  4 red
             blue
#> 5 white blue
#>
   6 blue
            red
#> 7 blue
             blue
#> 8 blue
             red
#> 9 white blue
#> 10 blue
             red
#> # ... with 9,990 more rows
How can we add a variable to two_draw_results to see if draw_1 and draw_2 match?
two_draw_results <- two_draw_results %>%
  mutate(color_match = draw_1 == draw_2)
two_draw_results
#> # A tibble: 10,000 x 3
#>
      draw_1 draw_2 color_match
#>
      <chr> <chr> <lgl>
#>
   1 white red
                    FALSE
#>
    2 red
             blue
                    FALSE
#>
    3 red
             blue
                    FALSE
#>
   4 red
             blue FALSE
#>
   5 white blue
                  FALSE
#> 6 blue
             red
                    FALSE
    7 blue
             blue
                    TRUE
#>
#>
  8 blue
            red
                    FALSE
#> 9 white blue
                  FALSE
#> 10 blue
             red
                    FALSE
#> # ... with 9,990 more rows
All that remains is to compute the relative frequency of the observed outcomes from our many simulations.
two_draw_results %>%
  count(color_match) %>%
  mutate(proportion = n / sum(n))
#> # A tibble: 2 x 3
     color_match
                     n proportion
#> * <lgl>
                 <int>
                            <dbl>
#> 1 FALSE
                  5903
                            0.590
#> 2 TRUE
                  4097
                            0.410
```

Practice

Suppose you roll two fair six-sided dice. Which has a higher probability: the square of dice roll 1 is equal to dice roll 2; or the absolute value of the difference between dice roll 1 and dice roll 2 is equal to 4.

Perform a simulation to compute this empirical probability.

Write down your guess to the answer before you calculate it. The probability of scenario 2 is about twice as large as the probability of scenario 1.

```
dice = c(1:6)
print(dice)
#> [1] 1 2 3 4 5 6
roll_results<- tibble(</pre>
die_1= replicate(n= 10000, expr= sample(dice, size= 1)),
die_2= replicate(n=10000, expr= sample(dice, size = 1))
)
roll_results<-
  roll_results %>%
  mutate(die1_squared = die_2== die_1*die_1,
         absolute_val_four= abs(die_1-die_2)==4)
roll_results %>%
  summarize(die1_squared_prob = mean(die1_squared),
            absolute_val_four_prob = mean(absolute_val_four))
#> # A tibble: 1 x 2
     die1_squared_prob absolute_val_four_prob
#>
#>
                 <dbl>
                                         <dbl>
                0.0553
                                         0.112
#> 1
```

Additional Resources-please look at before Weds.

• Open Intro Stats Sections 3.1 and 3.2