Permuted Data and Graphics

Unit 2 - Lab 6

Directions: Follow along with the slides and answer the questions in **BOLDED** font in your journal.

Why permute data?

- Permuting data is a really simple technique to shuffle or randomize our data.
 - You actually permuted data when you completed the *Horror Movie Shuffle* lab.
- Why do we bother?
 - So we can compare our data's distributions with examples that we know for a fact (100% certain) are random.
- And why is this helpful?
 - It lets us see what Chance World looks like so that we can compare our real world to Chance World

The Titanic

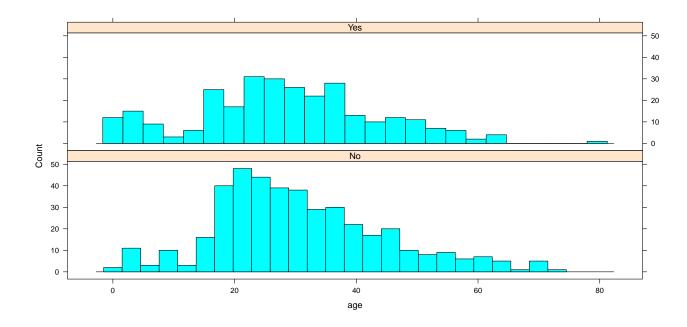
- The Titanic was a ship that sank en route to the U.S.A. from England after hitting an Iceberg in 1912.
- One common thought is that younger people were given priority when boarding the lifeboats.
- Use the following to load our titanic passenger and survival data.

data(titanic)

• In this lab, we'll take a brief look at missing data and practice comparing shuffled data to real data.

What do you think?

- Take a look at the following plot about the age of survivors and non-survivors.
- Was the *typical* survivor younger than the *typical* non-survivor? Give an argument as to why.



Let's find an answer!

• Let's start by calculating how much younger the *typical* survivor was than the *typical* non-survivor in our data.

```
mean(~age | survived, data = titanic)
```

- Notice that we get a warning message!
 - It appears we're missing 125 non-survivor's ages and 52 survivors ages.
 - Let's look at our favstats to see if this will be a problem.

```
favstats(~age | survived, data = titanic)
```

Missing values

```
favstats(~age | survived, data = titanic)
```

- Even though we're missing people in our data, we still have quite a few data points to use for calculating the *mean*.
- Write down the number of people who are NOT missing for each group
- What do you think we mean when we say missing data? Why might it be missing?

Back to our task

- Recalculate the mean age of survivors and non-survivors in our data.
- How much younger is the average suvivor than the average non-survivor in our *actual* titanic data?
 - Write down this number in your data science journal.

Randomizing our data

- Now that we've calculated the difference of our passenger's average ages, we'd like to compare it to data that we've randomized.
- Run the following to completely randomize the survived status for each person's age:

Resampling (a.k.a. permuting)

- When we write, data = resample(titanic, shuffled="survived"), we're telling R to:
 - Start with our original titanic data and ...
 - Take the Yes and No values in our survived variable and mix them up...
- Then! With this randomized data:
 - Find the average age of our *randomized* survivors and non-survivors.

Why shuffle (or resample)?

- By shuffling the values of Yes and No in our survived variable:
 - We keep the same number of total survivors.
 - But! We randomly choose which passengers survived (or didn't ...)
- This means that, any relationship between age and those who survived is lost!

Answer the following:

- How much younger is the average survivor than the average non-survivor in our *random-ized* titanic data?
- Is this number very different than the difference we computed for our *actual* titanic data?

But how "different" is 'different'?

• Comparing our actual data to the randomized data is not quite fair. -The randomized data is different every time.

- Instead of comparing to a single randomized data set, we want to compare to the *typical outcomes* we'd get from randomizing.
 - To do this, we need to randomize many, many times."
 - Then we can get a sense of how likely our *actual* values randomly appear.

do-ing things many times

- Just like how we computed our randomized data once, we can use the do() function to repeat our calculations many times.
- Run the following example:

- How many times did we randomize our data and then compute the average?
- Write down the code you would run to do our calculations 10 times.
 - Test your code to ensure it works.

Calculate many means

- When we do a calculation many times, the results are stored as a data.frame type object.
 - This means we can save and manipulate our many calculations.
- Re-run your code from the previous slide BUT \dots
 - do the calculation 300 times AND ...*
 - Assign the object the name shuffled_means
- Note: Doing something 300 times can take a while. Be patient while your code runs.
 - You'll know it's done when the > symbol re-appears in your console.

Find many differences

we want to calculate how much younger our average chance-world survivor was than our average chance world non-survivor

- Now we want to calculate how much younger our average *Chance-World* survivor was than our average *Chance World* non-survivor.
- To do this, run:

Looking at our differences

- After calculating our 300 randomized differences, make an appropriate plot to visualize them.
 - Write down the code you used to make the plot of differences.
- The values on the x-axis represent how much *younger* the average survivor was than the average non-survivor.
 - What does it mean for our difference in average age to be negative?

Making the call

- Compare the difference from our actual data, -2.2825, to the histogram of our randomized data.
 - Is our *actual* difference close to the values in the center of our histogram? Or are they far away?
 - What does it mean for our *actual* value to be far away from the center of our randomized values?

Making the call

- Read this part carefully:
 - If a real-life event is very common in Chance-World, then we might suspect that our real-life outcome was just due to chance, and not very meaningful.
 - But! if the *real-life* outcome is rare or unusual (i.e. Doesn't occur very often) in chance world, then we have evidence that the outcome is *meaningful*.
- Based on your shufflings, do you think the difference in age of survivors is meaningful? Explain.