Gov 50: 21. More Hypothesis testing

Matthew Blackwell

Harvard University

Roadmap

- 1. Hypothesis testing using infer
- 2. Two-sample tests
- 3. Two-sample permutation tests with infer

1/ Hypothesis testing using infer

Statistical hypothesis testing

- · Statistical hypothesis testing is a thought experiment.
- What would the world look like if we knew the truth?
- Conducted with several steps:
 - 1. Specify your null and alternative hypotheses
 - 2. Choose an appropriate **test statistic** and level of test α
 - 3. Derive the **reference distribution** of the test statistic under the null.
 - 4. Use this distribution to calculate the **p-value**.
 - 5. Use p-value to decide whether to reject the null hypothesis or not

GSS data from infer

library(infer) gss

```
A tibble: 500 x 11
##
                        college partyid hompop hours income
      vear
             age sex
     <dhl> <dhl> <fct>
                        <fct> <fct>
                                           <dhl> <dhl> <ord>
##
   1 2014
                                                    50 $25000~
##
              36 male
                        degree
                                 ind
                                               3
                                                    31 $20000~
##
   2 1994
             34 female no degree rep
   3 1998
              24 male
                                                    40 $25000~
##
                        degree
                                  ind
                                                    40 $25000~
##
      1996
              42 male
                        no degree ind
                                                    40 $25000~
##
   5 1994
              31 male
                        degree
                                  rep
##
   6 1996
              32 female no degree rep
                                               4
                                                    53 $25000~
##
   7 1990
              48 female no degree dem
                                                    32 $25000~
##
   8 2016
              36 female degree
                                  ind
                                                    20 $25000~
##
      2000
              30 female degree
                                rep
                                                    40 $25000~
                                                    40 $15000~
##
  10
      1998
              33 female no degree dem
  # i 490 more rows
    i 3 more variables: class <fct>, finrela <fct>,
## #
      weight <dbl>
```

What is the average hours worked?

```
dplyr way:
```

```
gss |>
  summarize(mean(hours))
```

```
## # A tibble: 1 x 1
## `mean(hours)`
## <dbl>
## 1 41.4
```

infer way:

```
observed_mean <- gss |>
  specify(response = hours) |>
  calculate(stat = "mean")
observed_mean
```

```
## Response: hours (numeric)
## # A tibble: 1 x 1
## stat
## <dbl>
## 1 41.4
```

Hypothesis test

Could we get a mean this different from 40 hours if that was the true population average of hours worked?

Null and alternative:

$$H_0: \mu_{\text{hours}} = 40$$

$$H_1: \mu_{\text{hours}} \neq 40$$

How do we perform this test using infer? The **bootstrap!**

Specifying the hypotheses

```
specify(response = hours) |>
 hypothesize(null = "point", mu = 40)
  Response: hours (numeric)
  Null Hypothesis: point
  # A tibble: 500 x 1
##
     hours
##
     <dh1>
##
   1
        50
## 2 31
   3 40
##
## 4 40
##
   5 40
##
   6
     53
##
   7 32
##
        20
##
       40
## 10
        40
  # i 490 more rows
```

Generating the null distribution

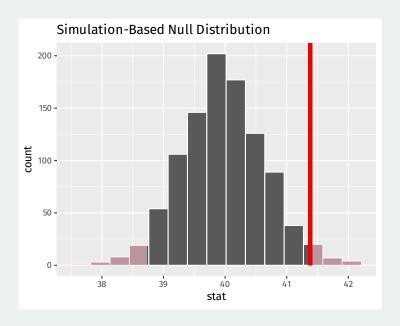
We can use the bootstrap to determine how much variation there will be around 40 in the null distribution.

```
null_dist <- gss |>
   specify(response = hours) |>
   hypothesize(null = "point", mu = 40) |>
   generate(reps = 1000, type = "bootstrap") |>
   calculate(stat = "mean")
null_dist
```

```
## Response: hours (numeric)
## Null Hypothesis: point
  # A tibble: 1,000 x 2
## replicate stat
        <int> <dhl>
##
## 1
           1 40.7
## 2
           2 40.5
## 3
           3 39.9
## 4
          4 41.1
## 5 5 41.3
       6 40.5
## 6
##
           7 40.8
```

Visualizing the p-value

We can visualize our bootstrapped null distribution and the p-value as a shaded region:



2/ Two-sample tests

Social pressure experiment

- Experimental study where each household for 2006 MI primary was randomly assigned to one of 4 conditions:
 - · Control: no mailer
 - Civic Duty: mailer saying voting is your civic duty.
 - · Hawthorne: a "we're watching you" message.
 - Neighbors: naming-and-shaming social pressure mailer.
- · Outcome: whether household members voted or not.
- · We'll focus on Neighbors vs Control
- Randomized implies samples are independent

Neighbors mailer

Dear Registered Voter:

WHAT IF YOUR NEIGHBORS KNEW WHETHER YOU VOTED?

Why do so many people fail to vote? We've been talking about the problem for years, but it only seems to get worse. This year, we're taking a new approach. We're sending this mailing to you and your neighbors to publicize who does and does not vote.

The chart shows the names of some of your neighbors, showing which have voted in the past. After the August 8 election, we intend to mail an updated chart. You and your neighbors will all know who voted and who did not.

DO YOUR CIVIC DUTY - VOTE!

MAPLE DR	Aug 04	Nov 04	Aug 06
9995 JOSEPH JAMES SMITH	Voted	Voted	
9995 JENNIFER KAY SMITH		Voted	
9997 RICHARD B JACKSON		Voted	
9999 KATHY MARIE JACKSON		Voted	

Social pressure data

```
data(social, package = "qss")
social <- as_tibble(social)
social</pre>
```

```
## # A tibble: 305,866 x 6
##
     sex
           yearofbirth primary2004 messages primary2006 hhsize
##
     <chr>>
                 <int>
                            <int> <chr>
                                                 <int> <int>
##
   1 male
                                0 Civic D~
                 1941
                                                     0
                                O Civic D~
##
   2 fema~
                  1947
                                                     0
##
   3 male
                  1951
                                0 Hawthor~
   4 fema∼
                  1950
                                0 Hawthor~
##
                                0 Hawthor~
##
   5 fema∼
                  1982
   6 male
                                0 Control
                                                     0
##
                  1981
##
   7 fema∼
                  1959
                                0 Control
                                                     1
                                                            3
##
   8 male
                  1956
                                0 Control
   9 fema~
                  1968
                                0 Control
                                                     0
##
  10 male
                                0 Control
                                                     0
                  1967
  # i 305,856 more rows
```

Two-sample hypotheses

- Parameter: **population ATE** $\mu_T \mu_C$
 - μ_T : Turnout rate in the population if everyone received treatment.
 - μ_C : Turnout rate in the population if everyone received control.
- · Goal: learn about the population difference in means
- Usual null hypothesis: no difference in population means (ATE = 0)
 - Null: $H_0: \mu_T \mu_C = 0$
 - Two-sided alternative: $H_1: \mu_T \mu_C \neq 0$
- In words: are the differences in sample means just due to chance?

Permutation test

How do we generate draws of the difference in means under the null?

$$H_0: \mu_T - \mu_C = 0$$

If the voting distribution is the same in the treatment and control groups, we could randomly swap who is labelled as treated and who is labelled as control and it shouldn't matter.

Permutation test: generate the null distribution by permuting the group labels and see the resulting distribution of differences in proportions

Permuting the labels

```
social <- social |>
  filter(messages %in% c("Neighbors", "Control"))

social |>
  mutate(messages_permute = sample(messages)) |>
  select(primary2006, messages, messages_permute)
```

```
## # A tibble: 229,444 x 3
##
     primary2006 messages messages permute
##
           <int> <chr> <chr>
##
   1
               0 Control Control
               1 Control Control
## 2
## 3
               1 Control
                         Neighbors
## 4
              0 Control
                         Control
## 5
              0 Control
                         Control
##
   6
              1 Control
                         Neighbors
                         Control
##
              0 Control
                         Control
## 8
              1 Control
##
   9
               1 Control
                         Control
## 10
              1 Control
                         Control
## # i 229,434 more rows
```

3/ Two-sample permutation tests with infer

Calculating the difference in proportion

infer functions with binary outcomes work best with factor variables:

```
social <- social |>
 mutate(turnout = if else(primary2006 == 1, "Voted", "Didn't Vote"))
est ate <- social |>
 specify(turnout ~ messages, success = "Voted") |>
 calculate(stat = "diff in props", order = c("Neighbors", "Control"))
est ate
## Response: turnout (factor)
## Explanatory: messages (factor)
## # A tibble: 1 x 1
## stat
## <dbl>
## 1 0.0813
```

Specifying the relationship of interest

infer functions with binary outcomes work best with factor variables:

```
social |>
  specify(turnout ~ messages, success = "Voted")
```

```
Response: turnout (factor)
  Explanatory: messages (factor)
## # A tibble: 229,444 x 2
## turnout messages
## <fct> <fct>
##
   1 Didn't Vote Control
##
   2 Voted Control
##
   3 Voted Control
   4 Didn't Vote Control
##
##
   5 Didn't Vote Control
##
   6 Voted Control
  7 Didn't Vote Control
##
##
   8 Voted Control
##
   9 Voted Control
## 10 Voted Control
  # i 229,434 more rows
```

Setting the hypotheses

The null for these two-sample tests is called "independence" for the infer package because the assumption is that the two variables are statistically independent.

```
social |>
  specify(turnout ~ messages, success = "Voted") |>
  hypothesize(null = "independence")
```

```
## Response: turnout (factor)
  Explanatory: messages (factor)
  Null Hypothesis: independence
  # A tibble: 229,444 x 2
##
     turnout messages
##
  <fct> <fct>
##
   1 Didn't Vote Control
   2 Voted Control
##
   3 Voted Control
##
   4 Didn't Vote Control
##
   5 Didn't Vote Control
##
##
   6 Voted Control
##
   7 Didn't Vote Control
##
   8 Voted Control
```

Generating the permutations

We can tell infer to do our permutation test by using the argument type = "permute" to generate():

```
social |>
  specify(turnout ~ messages, success = "Voted") |>
  hypothesize(null = "independence") |>
  generate(reps = 1000, type = "permute")
```

20 / 25

```
## Response: turnout (factor)
  Explanatory: messages (factor)
## Null Hypothesis: independence
  # A tibble: 229,444,000 x 3
  # Groups: replicate [1,000]
  turnout messages replicate
##
## <fct> <fct> <int>
##
   1 Voted Control
##
   2 Didn't Vote Control
##
   3 Voted Control
   4 Didn't Vote Control
##
##
   5 Didn't Vote Control
##
   6 Voted Control
##
   7 Voted Control
```

Calculating the diff in proportions in each sample

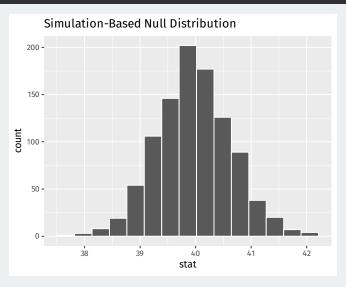
```
null_dist <- social |>
   specify(turnout ~ messages, success = "Voted") |>
   hypothesize(null = "independence") |>
   generate(reps = 1000, type = "permute") |>
   calculate(stat = "diff in props", order = c("Neighbors", "Control"))
```

null_dist

```
## Response: hours (numeric)
  Null Hypothesis: point
  # A tibble: 1,000 x 2
##
     replicate stat
##
##
         <int> <dbl>
##
   1
             1 40.7
##
   2
             2 40.5
##
             3 39.9
             4 41.1
##
   4
##
             5 41.3
##
   6
             6 40.5
               40.8
##
##
   8
             8
               39.8
                39.5
##
## 10
            10 39.9
## # i 990 more rows
```

Visualizing

null_dist |>
 visualize()



Calculating p-values

```
ate_pval <- null_dist |>
  get_p_value(obs_stat = est_ate, direction = "both")
ate_pval
```

```
## # A tibble: 1 x 1
## p_value
## <dbl>
## 1 0
```

Visualizing p-values

```
null_dist |>
  visualize() +
  shade_p_value(obs_stat = est_ate, direction = "both")
```

