IS5 in R: Comparing Groups (Chapter 17)

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Introduction and background

This document is intended to help describe how to undertake analyses introduced as examples in the Fifth Edition of *Intro Stats* (2018) by De Veaux, Velleman, and Bock. More information about the book can be found at http://wps.aw.com/aw_deveaux_stats_series. This file as well as the associated R Markdown reproducible analysis source file used to create it can be found at http://nhorton.people.amherst.edu/is5.

This work leverages initiatives undertaken by Project MOSAIC (http://www.mosaic-web.org), an NSF-funded effort to improve the teaching of statistics, calculus, science and computing in the undergraduate curriculum. In particular, we utilize the mosaic package, which was written to simplify the use of R for introductory statistics courses. A short summary of the R needed to teach introductory statistics can be found in the mosaic package vignettes (http://cran.r-project.org/web/packages/mosaic). A paper describing the mosaic approach was published in the R Journal: https://journal.r-project.org/archive/2017/RJ-2017-024.

Chapter 17: Comparing Groups

```
library(mosaic)
library(readr)
library(janitor)
```

Section 17.1: A Confidence Interval for the Difference Between Two Proportions

Here, the do() function creates a number of rows for the data frame.

```
set.seed(234)
numsim <- 10000

# What does do() do?
resample(Seatbelts) %>%
    group_by(passenger) %>%
    summarise(proportion = sum(belted)/n()) %>%
    summarise(diffprop = abs(diff(proportion))) # Difference of proportions from one random resample

## # A tibble: 1 x 1
## diffprop
## <dbl>
```

```
## 1
        0.170
resample(Seatbelts) %>%
  group_by(passenger) %>%
  summarise(proportion = sum(belted)/n()) %>%
  summarise(diffprop = abs(diff(proportion))) # Difference of proportions from another random resample
## # A tibble: 1 x 1
     diffprop
##
##
        <dbl>
## 1
        0.178
do(2) * resample(Seatbelts) %>%
  group_by(passenger) %>%
  summarise(proportion = sum(belted)/n()) %>%
  summarise(diffprop = abs(diff(proportion))) # Calculates two differences
##
      diffprop
## 1 0.1573754
## 2 0.1548267
# We need 10000 differences of proportions
seatbeltresamples <- do(numsim) * resample(Seatbelts) %>%
  group_by(passenger) %>%
  summarise(proportion = sum(belted)/n()) %>%
  summarise(diffprop = abs(diff(proportion)))
# Figure 17.1, page 542
gf_histogram(~ diffprop, data = seatbeltresamples) %>%
 gf_labs(x = "Difference of Proportions", y = "# of Resamples")
   1200 -
    900 -
of Resamples
    600 -
    300 -
      0 -
                   0.14
                                0.16
                                            0.18
                                                         0.20
       0.12
                           Difference of Proportions
```

Example 17.1: Finding the Standard Error of a Difference in Proportions

```
do(256 * .30 + 1) * data.frame(gender = "F", profile = FALSE)
tally(~ gender, data = OnlineProf)
## gender
## M F
## 248 256
OnlineProfM <- OnlineProf %>%
 filter(gender == "M")
sepboys <- ((mean(~ profile, data = OnlineProfM)) * (1 - mean(~ profile, data = OnlineProfM)))/nrow(OnlineProfM)</pre>
sepboys
## [1] 0.03145024
OnlineProfF <- OnlineProf %>%
 filter(gender == "F")
sepgirls <- ((mean(~ profile, data = OnlineProfF) * (1 - mean(~ profile, data = OnlineProfF)))/nrow(Onl</pre>
sepgirls
## [1] 0.02866236
sep <- (sepboys^2 + sepgirls^2)^.5</pre>
## [1] 0.04255171
Example 17.2: Finding a Two-Proportion z-Interval
zstats <-qnorm(p = c(.025, .975))
(mean(~ profile, data = OnlineProfF) - mean(~ profile, data = OnlineProfM)) + zstats * sep
## [1] 0.04727054 0.21407019
Section 17.2: Assumptions and Conditions for Comparing Proportions
Section 17.3: The Two-Sample z-Test: Testing for the Difference Between Proportions
Step-By-Step Example: A Two-Proportion z-Test
```

```
# Create the data set
SleepHabits <- rbind(</pre>
               * data.frame(gen = "GenY", internet = TRUE),
  do(293 - 205) * data.frame(gen = "GenY", internet = FALSE),
              * data.frame(gen = "GenX", internet = TRUE),
  do(469 - 235) * data.frame(gen = "GenX", internet = FALSE)
# Mechanics
ngeny <- nrow(filter(SleepHabits, gen == "GenY"))</pre>
ngeny
ygeny <- nrow(filter(SleepHabits, gen == "GenY" & internet == TRUE))
ygeny
```

```
## [1] 205
pgeny <- mean(~ internet, data = filter(SleepHabits, gen == "GenY"))</pre>
pgeny
## [1] 0.6996587
ngenx <- nrow(filter(SleepHabits, gen == "GenX"))</pre>
ngenx
## [1] 469
ygenx <- nrow(filter(SleepHabits, gen == "GenX" & internet == TRUE))</pre>
ygenx
## [1] 235
pgenx <- mean(~ internet, data = filter(SleepHabits, gen == "GenX"))</pre>
pgenx
## [1] 0.5010661
sepgen <- ((pgeny * (1 - pgeny))/ngeny + (pgenx * (1 - pgenx))/ngenx)^.5</pre>
sepgen
## [1] 0.03535867
pdiff <- pgeny - pgenx
pdiff
## [1] 0.1985926
z <- (pdiff - 0)/sepgen
## [1] 5.616518
2 * pnorm(q = z, lower.tail = FALSE)
## [1] 1.948444e-08
```

Section 17.4: A Confidence Interval for the Difference Between Two Means

Example 17.7: Finding a Confidence Interval for the Difference in Sample Means

```
# Not sure if creating data set is really necessary
# page 555
nord <- 27
nref <- 27
yord <- 8.5
yref <- 14.7
sord <- 6.1
sref <- 8.4

seys <- 2.0
diffy <- yref - yord # 6.2
tstats <- qt(p = c(.025, .975), df = 47.46)
tstats</pre>
```

[1] -2.011226 2.011226

```
me <- tstats * seys
me
## [1] -4.022452 4.022452
diffy + me
## [1] 2.177548 10.222452
```

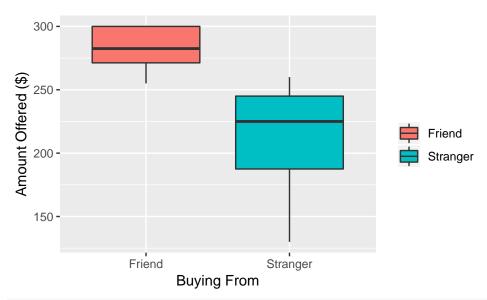
Section 17.5: The Two-Sample t-Test: Testing for the Difference Between Two Means

```
Step-By-Step Example: A Two-Sample t-Test for the Difference Between the Two Means
# page 556
BuyingCam <- read_csv("http://nhorton.people.amherst.edu/is5/data/Buy_from_a_friend.csv")</pre>
## Parsed with column specification:
## cols(
    Friend = col_integer(),
     Stranger = col_integer()
##
## )
By default, read_csv() prints the variable names. These messages can be suppressed using the
message=FALSE code chunk option to save space and improve readability.
library(tidyr) # for gather() function
## Attaching package: 'tidyr'
## The following object is masked from 'package:Matrix':
##
##
       expand
BuyingCam <- BuyingCam %>%
  gather(key = buying_type, value = amount_offered, Friend, Stranger)
# Model
```

Warning: Removed 1 rows containing non-finite values (stat_boxplot).

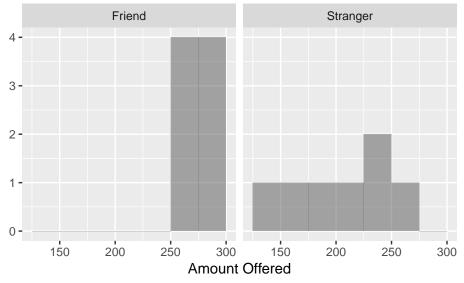
gf_labs(x = "Buying From", y = "Amount Offered (\$)", fill = "")

gf_boxplot(amount_offered ~ buying_type, fill = ~ buying_type, data = BuyingCam) %>%



```
gf_histogram(~ amount_offered, binwidth = 25, center = 12.5, data = BuyingCam) %>% # doesn't exactly ma
gf_facet_wrap(buying_type ~ .) %>%
gf_labs(x = "Amount Offered", y = "")
```

Warning: Removed 1 rows containing non-finite values (stat_bin).



Section 17.6: Randomization Tests and Confidence Intervals for Two Means

```
Cars <- read_csv("http://nhorton.people.amherst.edu/is5/data/Car_speeds.csv")</pre>
```

Parsed with column specification:

```
## cols(
## direction = col_character(),
## speed = col_double()
## )
# Figure 17.2 (page 560) is the same as Figure 4.4 (page 102)
favstats(~ speed | direction, data = Cars)
                         Q1 median
##
   direction min
                                       Q3 max
         Down 10.27 20.4675 22.885 25.3525 32.95 22.71708 3.622006 250
           Up 15.08 22.4975 25.155 28.1600 34.97 25.25172 3.856331 250
## 2
## missing
## 1
## 2
          0
```

Section 17.7: Pooling

Section 17.8: The Standard Deviation of a Difference