IS5 in R: Testing Hypotheses (Chapter 15)

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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 15: Testing Hypotheses

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

Section 15.1: Hypotheses

Section 15.2: P-Values

Section 15.3: The Reasoning of Hypothesis Testing

Example 15.5: Finding A P-Value

```
n <- 90
x <- 61
p <- .8
phat <- x/n
sdphat <- ((p * (1 - p))/n)^.5
z <- (phat - p)/sdphat
pnorm(z)
```

```
## [1] 0.00187324
```

```
# Or, without calculating the z-score:
pnorm(q = phat, mean = p, sd = sdphat)
```

```
## [1] 0.00187324
```

Section 15.4: A Hypothesis Test for the Mean

```
GestationTime <- read_csv("http://nhorton.people.amherst.edu/is5/data/Nashville.csv")
```

```
## Parsed with column specification:
## cols(
##   Gestation = col_integer(),
##   Time = col_character()
## )
```

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.

```
# 2. Model (page 482)
gf_histogram(~ Gestation, data = GestationTime, binwidth = 7.5, center = 3.75) %>%
  gf_labs(x = "Gestation Time (days)", y = "# of Births")
```
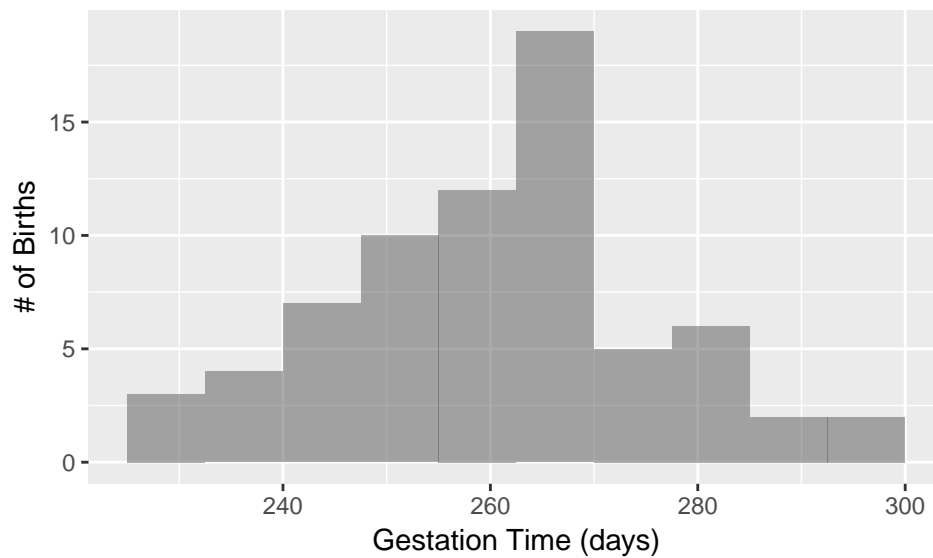


```
# 3. Mechanics
gf_dist(dist = "t", df = 69) %>%
  gf_vline(xintercept = -3.118) %>%
    gf_vline(xintercept = 3.118) %>%
  gf_labs(x = "", y = "") +
  xlim(-3.347, 3.347)
```
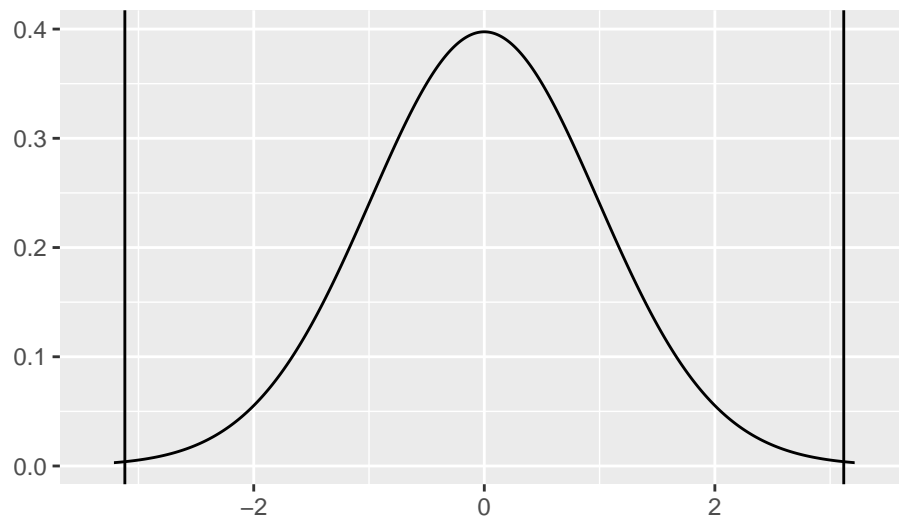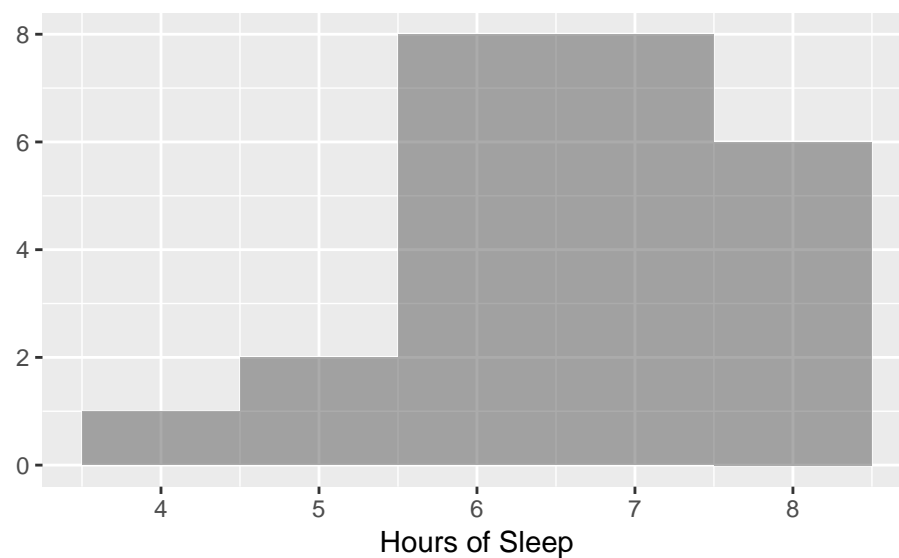
Step-By-Step Example: A One-Sample *t*-Test for the Mean

```
Sleep <- read_csv("http://nhorton.people.amherst.edu/is5/data/Sleep.csv")
```
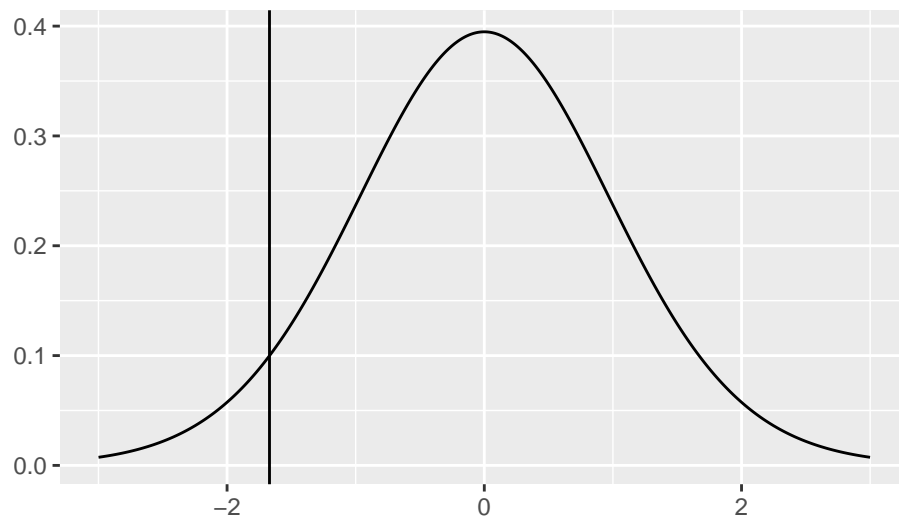
```
## Parsed with column specification:
## cols(
##   Sleep = col_integer()
## )
```

```
# Plan
gf_histogram(~ Sleep, data = Sleep, binwidth = 1) %>%
  gf_labs(x = "Hours of Sleep", y = "")
```



```
gf_dist(dist = "t", df = 24) %>%
  gf_vline(xintercept = -1.67) %>%
  gf_labs(x = "", y = "") +
  xlim(-3, 3)
```

```
## Warning: Removed 674 rows containing missing values (geom_path).
```

```
# Mechanics
n <- 25
mean <- 7.0
df <- 24
y <- 6.64
s <- 1.075
sey <- s/(n^.5)
t <- (y - mean)/sey # t-statistic
pt(q = t, df = df) # p-value
```

```
## [1] 0.05351625
```
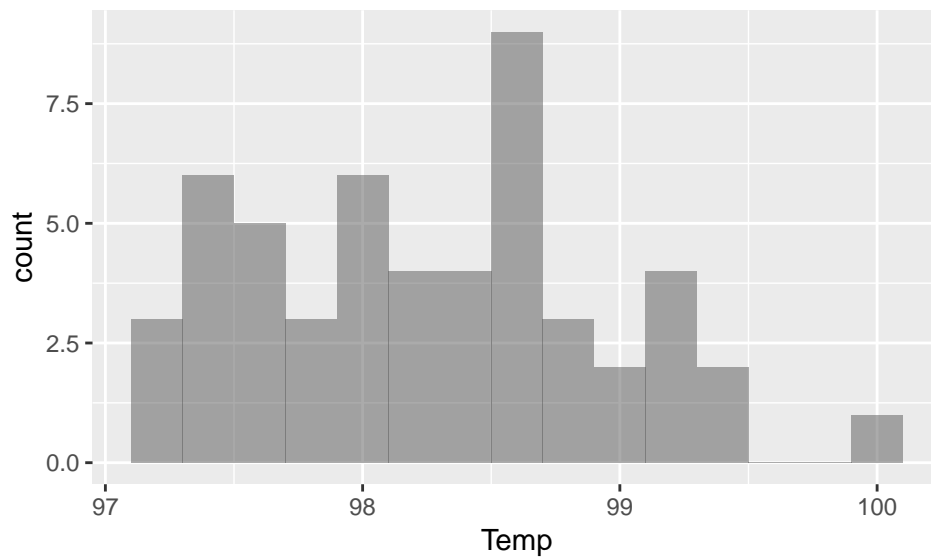
Section 15.5: Intervals and Tests

```
# page 487
Temperatures <- read_csv("http://nhorton.people.amherst.edu/is5/data/Normal_temperature.csv")
```

```
## Parsed with column specification:
## cols(
##   Temp = col_double()
## )
```

```
favstats(~ Temp, data = Temperatures)
```

```
##    min    Q1 median   Q3 max    mean        sd  n missing
##   97.2 97.675   98.2 98.7 100 98.28462 0.6823789 52       0
```

```
gf_histogram(~ Temp, data = Temperatures, binwidth = .2)
```

4

```r
# Confidence interval
y <- mean(~ Temp, data = Temperatures)
y
```

```
## [1] 98.28462
```

```r
s <- sd(~ Temp, data = Temperatures)
s
```

```
## [1] 0.6823789
```

```r
n <- nrow(Temperatures)
n
```

```
## [1] 52
```

```r
tstats <- qt(df = n - 1, p = c(.005, .995))
tstats
```

```
## [1] -2.675722  2.675722
```

```r
y + (tstats * (s/(n^.5)))
```

```
## [1] 98.03141 98.53782
```

```r
# Hypothesis test
mu <- 98.6
t <- (y - mu)/(s/(n^.5))
t
```

```
## [1] -3.332856
```

```r
2 * pt(q = t, df = n - 1) # two sided test
```

```
## [1] 0.001605849
```

Step-By-Step Example: Tests and Intervals

```r
# Creating the data set
Baseball <- rbind(
  do(1308) * (winner = "HOME"),
```

```
  do(2431 - 1308) * (winner = "AWAY")
) %>%
  rename(winner = result)
# Mechanics (page 490)
n <- nrow(Baseball)
p <- .5
phat <- Baseball %>%
  filter(winner == "HOME") %>%
  nrow()/n
phat
```

[1] 0.5380502

```
sdphat <- ((p * (1 - p))/n)^.5
sdphat
```

[1] 0.01014092

```
z <- (phat - p)/sdphat # z-value
z
```

[1] 3.752142

```
1 - pnorm(z) # p-value
```

[1] 8.76651e-05

```
# Or, without calculating the z-score:
1 - pnorm(q = phat, mean = p, sd = sdphat)
```

[1] 8.76651e-05

```
# Mechanics (page 491)
sep <- ((phat * (1 - phat))/n)^.5
sep
```

[1] 0.01011152

```
me <- 1.96 * sep
phat - me # lower bound of 95% confidence
```

[1] 0.5182316

```
phat + me # upper bound of 95% confidence
```

[1] 0.5578688

Section 15.6: P-Values and Decisions: What to Tell About a Hypothesis Test