## IPS9 in R: Two-way analysis of variance (Chapter 13)

Bonnie Lin and Nicholas Horton (nhorton@amherst.edu)

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## Introduction and background

These documents are intended to help describe how to undertake analyses introduced as examples in the Ninth Edition of *Introduction to the Practice of Statistics* (2017) by Moore, McCabe, and Craig.

More information about the book can be found here. The data used in these documents can be found under Data Sets in the Student Site. This file as well as the associated R Markdown reproducible analysis source file used to create it can be found at https://nhorton.people.amherst.edu/ips9/.

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 13: Two-way analysis of variance

This file replicates the analyses from Chapter 13: Two-way analysis of variance.

First, load the packages that will be needed for this document:

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

## Section 13.1: The two-way ANOVA model

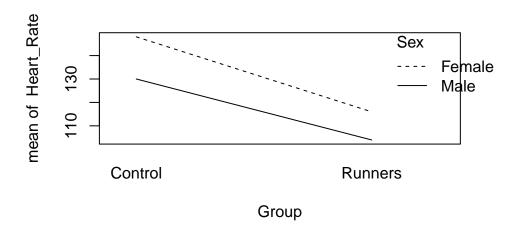
```
HRTRATE <- read_csv("https://nhorton.people.amherst.edu/ips9/data/chapter13/EG13-08HRTRATE.csv")
head(HRTRATE)</pre>
```

```
## # A tibble: 6 x 3
##
             Control Runners
     Sex
##
     <chr>>
               <int>
                        <int>
## 1 Female
                 159
                          119
## 2 Female
                 183
                           84
## 3 Female
                 140
                           89
## 4 Female
                 140
                          119
## 5 Female
                          127
                 125
## 6 Female
                          111
```

By default, the read\_csv() function will output the types of columns, as we see above. To improve readability for future coding, we will suppress the "Parsed with column specification" message by adding message = FALSE at the top of the code chunks.

We need to transform the data from wide to tall format using the gather() function.

```
HRTRATE_tidy <- HRTRATE %>%
  tidyr::gather(key = Group, value = Heart_Rate, Control, Runners)
head(HRTRATE_tidy)
## # A tibble: 6 x 3
##
     Sex
            Group
                    Heart_Rate
##
     <chr> <chr>
                         <int>
## 1 Female Control
                           159
## 2 Female Control
                           183
## 3 Female Control
                           140
## 4 Female Control
                           140
## 5 Female Control
                           125
## 6 Female Control
                           155
## Figure 13.4, age 710
favstats(Heart_Rate ~ Sex + Group, data = HRTRATE_tidy)
##
          Sex.Group min Q1 median Q3 max
                                               mean
                                                          sd
                                                               n missing
## 1 Female.Control 105 137
                               147 160 196 148.000 16.27095 200
       Male.Control 77 119
                               130 142 172 130.000 17.10035 200
                                                                        0
## 3 Female.Runners 78 106
                               116 126 164 115.985 15.97154 200
                                                                        0
       Male.Runners 69 96
                               103 112 146 103.975 12.49942 200
                                                                        Λ
## Figure 13.5, age 711
lm_HRTRATE <- lm(Heart_Rate ~ Group * Sex, data = HRTRATE_tidy)</pre>
msummary(lm_HRTRATE)
##
                        Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                         148.000
                                       1.100 134.511 < 2e-16 ***
## GroupRunners
                         -32.015
                                       1.556 -20.575 < 2e-16 ***
## SexMale
                         -18.000
                                       1.556 -11.568 < 2e-16 ***
## GroupRunners:SexMale
                           5.990
                                       2.201
                                               2.722 0.00663 **
##
## Residual standard error: 15.56 on 796 degrees of freedom
## Multiple R-squared: 0.5276, Adjusted R-squared: 0.5258
## F-statistic: 296.3 on 3 and 796 DF, p-value: < 2.2e-16
anova(lm_HRTRATE)
## Analysis of Variance Table
##
## Response: Heart_Rate
##
              Df Sum Sq Mean Sq F value Pr(>F)
## Group
               1 168432 168432 695.6470 < 2e-16 ***
                  45030
                          45030 185.9799 < 2e-16 ***
## Sex
               1
## Group:Sex
               1
                   1794
                           1794
                                  7.4095 0.00663 **
## Residuals 796 192730
                            242
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
We see that there is a significant interaction (p=0.007): the sex difference between the heart rates differs by
groups.
### Figure 13.6, page 712
with(HRTRATE_tidy, interaction.plot(Group, Sex, Heart_Rate))
```



Section 13.2: Inference for two-way ANOVA