IPS9 in R: One-way analysis of variance (Chapter 12)

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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 12: One-way analysis of variance

This file replicates the analyses from Chapter 12: One-way analysis of variance.

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

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

Complicated computations do not guarantee a valid statistical analysis. (648)

Section 12.1: Inference for one-way analysis of variance

We begin with Example 12.3 in page 648. Let's read in our data

```
Friends <- read_csv("https://nhorton.people.amherst.edu/ips9/data/chapter12/EG12-03FRIENDS.csv") head(Friends)
```

```
## # A tibble: 6 x 3
##
     Friends Participant Score
##
       <int>
                     <int> <dbl>
## 1
          102
                         1
                              3.8
                         2
## 2
          102
                              3.6
## 3
          102
                         3
                              3.2
## 4
          102
                         4
                              2.4
                         5
                              4.8
## 5
          102
## 6
```

We want to get a nice summary table like the one in example 12.3. To do this we have to group the dataset by the number of friends. Then we can use some aggregating functions to get the desired table structure.

```
SummaryFriends <- Friends %>%
  group_by(Friends) %>%
  summarize(n = n(), x_hat = mean(Score), s = sd(Score))
SummaryFriends
```

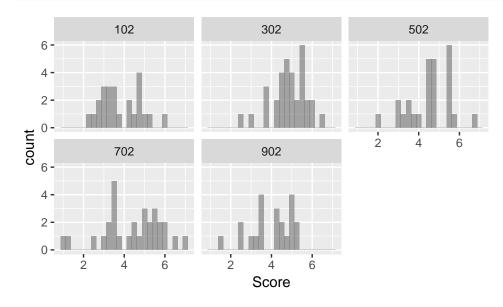
```
## # A tibble: 5 x 4
##
     Friends
                 n x_hat
##
       <int> <int> <dbl> <dbl>
## 1
         102
                24 3.82 0.999
                    4.88 0.851
## 2
         302
                33
## 3
         502
                26
                    4.56 1.07
## 4
         702
                30
                    4.41 1.43
## 5
         902
                21
                    3.99 1.02
```

In R the combination of group_by() and summary() is a powerful one. With these two data verbs we can create summary outputs like the one above with very little code.

There are some helpful visualizations starting in page 649 that will help us in our analysis of our Friends data.

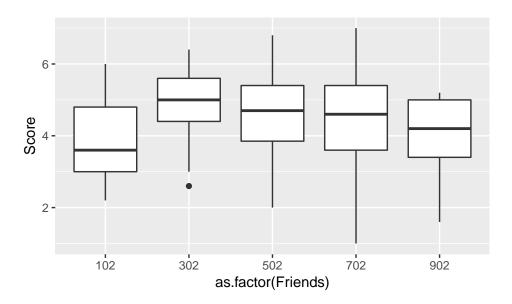
To recreate the histogram (note the | used to facet based on the number of friends):

gf_histogram(~ Score | Friends, data = Friends) #XX need to make this match the figure



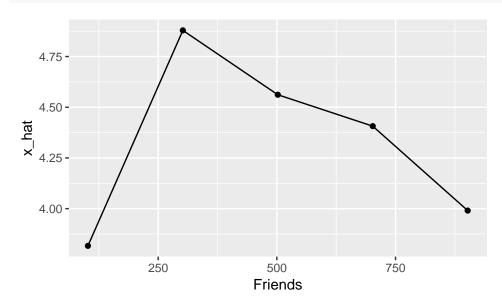
To recreate the boxplot (note: we have to convert Friends to a factor for the boxplots to work):

```
gf_boxplot(Score ~ as.factor(Friends), data = Friends)
```



And to recreate the linegraph (note the piping through gf_point to add the points to the line):

gf_line(x_hat ~ Friends, data = SummaryFriends) %>% gf_point()



These visualizations are a quick way to know what is going on with your data. Just a few lines of codes can save much time in your analysis.

XX EXample 12.7 There is a table to recreate here.

The ANOVA analysis can be done in a straightforward manner in R. We will create a linear model out of the Scores given their Friends group and pipe it into the anova function. (Note: the anova function takes an lm object. This is useful to us since we would maybe want to use the lm object for other purposes too.)

```
modFriends <- lm(Score ~ Friends, data = Friends)
modFriends %>%
anova()
```

Analysis of Variance Table
##

Eyes <- read_csv("https://nhorton.people.amherst.edu/ips9/data/chapter12/EG12-15EYES.csv")</pre>

We will recreate the ouput from the Excel spreadsheet. With the help of the group_by() and summarize() idiom we can use the aggregating functions to summarize our dataset. To recreate the anova part of the output we use again the anova() function that takes an lm object.

```
Eyes %>%
  group_by(Group) %>%
  summarize(n = n(),
            Sum = sum(Score),
            Average = Sum / n,
            Variance = sd(Score)^2)
## # A tibble: 4 x 5
     Group
                   Sum Average Variance
##
               n
     <chr> <int> <dbl>
##
                          <dbl>
                                   <dbl>
                                    3.08
## 1 Blue
              67
                  214
                           3.19
## 2 Brown
              37
                  138.
                           3.72
                                    2.94
## 3 Down
                  127.
                           3.11
                                    2.33
              41
                           3.86
## 4 Green
              77 297.
                                    2.78
modEyes <- lm(Score ~ Group, data = Eyes)</pre>
modEves %>%
  anova()
## Analysis of Variance Table
##
## Response: Score
              Df Sum Sq Mean Sq F value Pr(>F)
##
               3 24.42 8.1399 2.8941 0.03618 *
## Group
## Residuals 218 613.14 2.8126
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
XXExample 12.16 recreate table equal variance test
```

Section 12.2: Comparing the means

Response: Score

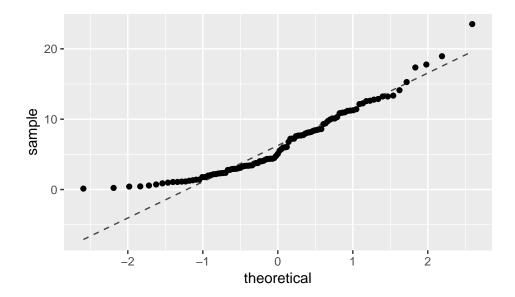
Anova gives us the answer to the question "are the differences between the means of the groups statistically significant?" However, it gives no information on what these differences are. This section covers those PostHocTests.

Let's read in the dataset for the times for times people spend on Facebook.

```
Facetym <- read_csv("https://nhorton.people.amherst.edu/ips9/data/chapter12/EG12-17FACETYM.csv")
```

The analysis starts by checking the data. We will check the distribution of our data with a qqplot.

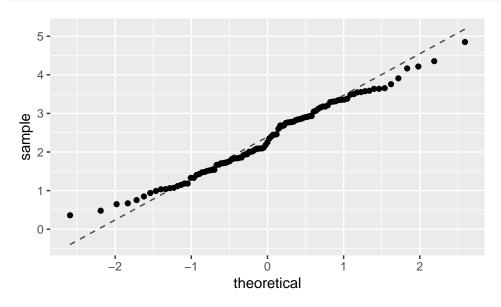
```
gf_qq(~ Time, data = Facetym) %>%
gf_qqline()
```



The gf_qq() is piped to gf_qqline() in order to get the reference line through the middle of the plot. Note the skewdness of the data.

We now visualize the transformed data.

```
gf_qq(~ SqrtTime, data = Facetym) %>%
gf_qqline()
```



Much better. Next step in our analysis is to get some descriptive statistics about the data. After that we perform the Anova test with anova().

```
Facetym %>%
  group_by(Grp) %>%
  summarize(n = n(),
     Sum = sum(SqrtTime),
     Average = mean(SqrtTime),
     SD = sd(SqrtTime),
     min = min(SqrtTime),
```

```
max = max(SqrtTime)
## # A tibble: 5 x 7
##
       Grp
                n
                    Sum Average
                                     SD
                                          min
                                                 max
##
     <int> <int> <dbl>
                           <dbl> <dbl> <dbl> <dbl>
## 1
         1
               21
                   52.9
                            2.52 0.850 0.849
                                               3.76
## 2
         2
               21
                   54.3
                            2.58 0.892 1.04
                                                4.35
         3
## 3
               21
                   50.5
                            2.40 0.921 1.06
                                                4.17
## 4
         4
               21
                   54.9
                            2.61 1.04 0.480 4.85
                            1.60 0.834 0.361 3.65
## 5
         5
               21
                   33.6
modFacetym <- lm(SqrtTime ~ factor(Grp), data = Facetym)</pre>
modFacetym %>%
  anova()
## Analysis of Variance Table
##
## Response: SqrtTime
##
                 Df Sum Sq Mean Sq F value
                                       4.545 0.002051 **
                  4 15.08 3.7701
## factor(Grp)
## Residuals
                100 82.95 0.8295
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Since the result is statistically significant, there is enough evidence to reject the null hypothesis that the mean
differences are all 0. However, we now need PostHocTests in order to find out what those differences are.
We will use a package called DescTools which has many multiple comparisons tests for after you reject the
null hypothesis based on your anova() output.
To use the PostHocTest() function we have to pass it an object returned by the aov() function, which
does the ANOVA test for a formula (and not an 1m object). After that we only have to specify the method
parameter, which is equal to the name of the multiple comparisons test that you want to perform.
library(DescTools)
## Attaching package: 'DescTools'
## The following object is masked from 'package:mosaic':
##
##
       MAD
PostHocTest(aov(SqrtTime ~ factor(Grp), data = Facetym), method = "bonferroni")
##
##
     Posthoc multiple comparisons of means : Bonferroni
##
       95% family-wise confidence level
##
## $`factor(Grp)`
##
               diff
                         lwr.ci
                                       upr.ci
                                                 pval
## 2-1 0.06723591 -0.7396166
                                0.874088419 1.0000
## 3-1 -0.11275242 -0.9196049
                                 0.694100090 1.0000
## 4-1 0.09736089 -0.7094916
                                 0.904213396 1.0000
```

5-1 -0.91714318 -1.7239957 -0.110290666 0.0151 *
3-2 -0.17998833 -0.9868408 0.626864181 1.0000
4-2 0.03012498 -0.7767275 0.836977488 1.0000

```
## 5-2 -0.98437909 -1.7912316 -0.177526574 0.0069 **
## 4-3 0.21011331 -0.5967392 1.016965818 1.0000
## 5-3 -0.80439076 -1.6112433 0.002461756 0.0513 .
## 5-4 -1.01450406 -1.8213566 -0.207651551 0.0048 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
PostHocTest(aov(SqrtTime ~ factor(Grp), data = Facetym), method = "lsd")
##
##
    Posthoc multiple comparisons of means : Fisher LSD
##
      95% family-wise confidence level
##
## $`factor(Grp)`
##
             diff
                      lwr.ci
                                 upr.ci
                                           pval
## 2-1 0.06723591 -0.4903979 0.6248697 0.81143
## 3-1 -0.11275242 -0.6703863 0.4448814 0.68916
## 4-1 0.09736089 -0.4602729 0.6549947 0.72977
## 5-1 -0.91714318 -1.4747770 -0.3595093 0.00151 **
## 3-2 -0.17998833 -0.7376222 0.3776455 0.52340
## 4-2 0.03012498 -0.5275089 0.5877588 0.91486
## 5-2 -0.98437909 -1.5420129 -0.4267453 0.00069 ***
## 4-3 0.21011331 -0.3475205 0.7677471 0.45649
## 5-3 -0.80439076 -1.3620246 -0.2467569 0.00513 **
## 5-4 -1.01450406 -1.5721379 -0.4568702 0.00048 ***
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
XX Anova power?
```