R for Psych Handbook

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Preface

This book serves as a collection of resources used in the LSU psychological statistics courses. It contains some lecture notes, as well as R code and data to run each of the examples in R.

The book is still under construction.

6

Introduction to R

You can label chapter and section titles using {#label} after them, e.g., we can reference Chapter 2. If you do not manually label them, there will be automatic labels anyway, e.g., Chapter ??.

Figures and tables with captions will be placed in figure and table environments, respectively.

```
par(mar = c(4, 4, .1, .1))
plot(pressure, type = 'b', pch = 19)
```

Reference a figure by its code chunk label with the fig: prefix, e.g., see Figure 2.1. Similarly, you can reference tables generated from knitr::kable(), e.g., see Table 2.1.

```
knitr::kable(
  head(iris, 20), caption = 'Here is a nice table!',
  booktabs = TRUE
)
```

You can write citations, too. For example, we are using the **bookdown** package (Xie, 2017) in this sample book, which was built on top of R Markdown and **knitr** (Xie, 2015).

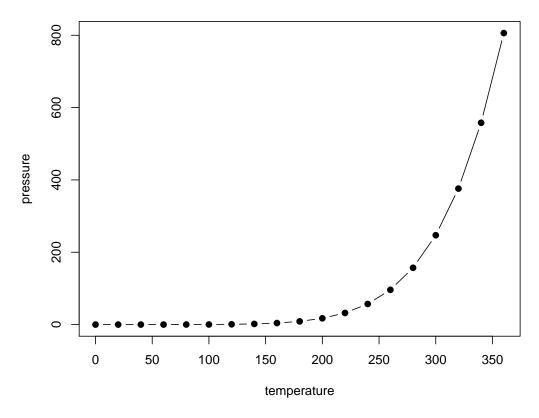


Figure 2.1: Here is a nice figure!

Table 2.1: Here is a nice table!

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa
4.6	3.4	1.4	0.3	setosa
5.0	3.4	1.5	0.2	setosa
4.4	2.9	1.4	0.2	setosa
4.9	3.1	1.5	0.1	setosa
5.4	3.7	1.5	0.2	setosa
4.8	3.4	1.6	0.2	setosa
4.8	3.0	1.4	0.1	setosa
4.3	3.0	1.1	0.1	setosa
5.8	4.0	1.2	0.2	setosa
5.7	4.4	1.5	0.4	setosa
5.4	3.9	1.3	0.4	setosa
5.1	3.5	1.4	0.3	setosa
5.7	3.8	1.7	0.3	setosa
5.1	3.8	1.5	0.3	setosa

Data Manipulation in R

Here is a review of existing methods.

Episotomology of Statistics

We describe our methods in this chapter.

Descriptive Statistics, z Scores, Central Limit

Some *significant* applications are demonstrated in this chapter.

- 5.1 Example one
- 5.2 Example two

Data Transformations and Confidence Intervals

We have finished a nice book.

Power and Effect Size Measures

attempt here

Correlation and Regression

attempt here

Matched T Test

One Way ANOVA

Multiple Comparisons

Simple Effects, Interactions, Mixed Designs

Repeated Measures ANOVA:

Mixed Two Way ANOVA

Multiple Regression

Chi-Square

Non-Parametric Data

Advanced Data Cleaning

Advanced Multiple Regression

Logistic Regression

Mediation and Moderation

ANCOVA

22.1 Theory

22.1.1 Reducing Noise

Just like the ANOVA (Analysis of Variance), the ANCOVA (Analysis of Covariance) is used to analyze experiments by calculating an F ratio for more than 2 groups. As with any F calculation, differences in the dependent variable are caused by two things:

- 1. The independent variable (signal, systematic variation)
- 2. Error (noise, unsysematic variation)

The F statistic captures this

$$F = \frac{Variation \ Due \ to \ IV}{Variation \ Due \ to \ Error}$$

When there is small variation within groups and large variation between groups, we get a large F.

CHART HERE

When there is large variatin within groups and small variation between groups, we get a small F.

The idea with the ANCOVA is that you can reduce your error term (the denominator from above) by choosing a covariate (CV) that is related to the DV and will soak up some random variation in your F test to give you a clearer picture of what is going on. Usually this means controlling for some sort of variable. The classic example is if you wanted to give some kids a test of some mental ability in a between subjects design but you don't want their age to skew your results. You might have had a control condition, an intervention, and some sort of alternate intervention. Typically you would run an ANOVA on the three groups, but since you know age has been accounted for and you want to remove that effect from the model, you enter age as a covariate in this calculation. Outside of this there are **three** major applications for ANCOVA.

- 1. Increase test sensitivity by using the CV(s) to account for more of the error variance
- 2. Adjust DV scores to what they would be if everyone scored the same on the CV(s)
- 3. Adjustment of a DV for other DVs taken as CVs

Note that use of a CV can adjust DV scores and show a larger effect or the CV can even eliminate the effect. Looking at this another way:

- 1. Reduces random error by increasing the size of F
- 2. Reduces systematic error by adjusting for differences in means
- 3. May increase differences by soaking up error

CHAPTER 22. ANCOVA

It's a **good** idea to use ANCOVA when you are removing variance in the DV related to covariate, but not related to the grouping variable. This decreases the error term and increases power.

It's a bad idea to use ANCOVA when groups differ on their mean level of the covariate. Usually here the covariate and the grouping variable are not independent. An example of this might be when "controlling" for anxiety when studying people with and without depression. Clearly people with depression will have higher levels of anxiety than their controls by nature of having depression!

22.1.2 Assumptions of ANOVA

- 1. All ANOVA assumptions apply (errors are RIND)
- Random
- Independent
- Normally distributed
- 2. The Covariate z is unrelated to x, and that z is related to, and in a sense, acts as a suppressor of y
- 3. The covariate has a linear relationship with the dependent variable
- 4. Variance of groups are equal.
- 5. Correlation between y and z is equal for all levels of x (Homogeneity of Regression)

You test for homogeneity of regression slopes by including a covariate by group interaction. If it's significant, you violated an assumption and can't to ANCOVA!

CHART

CHART

22.1.3 Best Practice

Things you should control for: 1. Broad Descriptors (Age, Social Class) 2. General Ability (intelligence, memory, speed of movement) 3. Personality measures (extraversion, neuroticism) 4. Things you think are relevant 5. Pick CVs that are correlated with DV * Height good for basketball * Height poor for test of math skills 6. CV should not be influenced by treatment

22.2 Practice

In order to get our hands dirty with ANCOVA we're going to look at a dataset where a bunch of sheep were slaughtered. We have three types of sheep (ewe, wether, ram) and a measure of fatness and weight.

Let's first run an ANOVA to see if fatness differs between animal type.

```
library(data.table)
## Warning: package 'data.table' was built under R version 3.4.2
deadsheep <- fread("datasets/deadsheep.csv")</pre>
deadsheep[, Animal := as.factor(animal)]
sheep.model.1 <- aov(fatness ~ Animal, data=deadsheep)</pre>
summary(sheep.model.1)
##
               Df Sum Sq Mean Sq F value Pr(>F)
## Animal
                  124.4
                           62.21
                                   3.165 0.0566 .
## Residuals
               30 589.6
                           19.65
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

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Looking at the output, we note that There was no significant effect of fatness on levels of animal,

$$F(2,30) = 3.165, p > .05$$

.

Now since we know that the whole weight of the animal might be confounding our answer (it's related to the DV of fatness, but not nessecarily related to the type of sheep), let's run an ANCOVA with carcass weight as a covariate and see if that changes our answer.

Before we run the ANCOVA, we need to first check that our IV is not significantly related to our CV. We do this by running an ANOVA predicting weight by animal.

```
#Test IND of IV and CV ind.sheep <- aov(weight ~ Animal, data=deadsheep) ## Check CV for IV indep, want non-sig summary(ind.sheep) # p=.104
```

```
## Df Sum Sq Mean Sq F value Pr(>F)
## Animal 2 14.20 7.102 2.438 0.104
## Residuals 30 87.41 2.914
```

Here we get

$$F(2,30) = 2.44, p > .05$$

and can move forward. Let's now run our ANCOVA. Note that R's default is Type I Sum of Squares, we need to load the car package to give us the correct sum of squares. For further reading on this, check out the Andy Field book on this.

```
sheepANCOVA <- aov(fatness ~ Animal + weight, data=deadsheep)
library(car)
Anova(sheepANCOVA, type="III")</pre>
```

We can then report

The covariate, weight, was not significantly related to the fatness of the animal,

$$F(1,29) =, p = .0565$$

and that there was a significant effect of the animal after controlling for weight of the animal,

$$F(2,28) = 37.62, p < .05.$$

.

Let's now check on our marginal means. These are our group means after we have controlled for our covariate. For this we need the effects package.

```
library(effects)
```

```
## Loading required package: carData
```

```
##
## Attaching package: 'carData'
## The following objects are masked from 'package:car':
##
##
       Guyer, UN, Vocab
## lattice theme set by effectsTheme()
## See ?effectsTheme for details.
mean.sheep <- effect("Animal", sheepANCOVA, se=TRUE)</pre>
summary(mean.sheep)
##
##
    Animal effect
  Animal
##
        Ewe
                 Ram
                        Wether
  18.90219 10.53996 14.83058
##
##
   Lower 95 Percent Confidence Limits
##
## Animal
##
         Ewe
                    Ram
                           Wether
## 17.565871 9.183322 13.532447
##
##
    Upper 95 Percent Confidence Limits
## Animal
##
        F.we
                  R.am
                        Wether
## 20.23851 11.89660 16.12870
```

We can then note the estimated marginal means here (18.9 for Ewe, 10.53 for Ram, and 14.83 for Wether) are mean values for fatness once the CV of weight has been controlled for.

Lastly, let's test the homogenaity of the regression slopes assumption (though we should have done this first!). We do this by running the model with an interaction term included.

```
hoRS.sheep <- aov(fatness ~ weight + Animal + animal:weight, data=deadsheep)
Anova(hoRS.sheep, type="III")</pre>
```

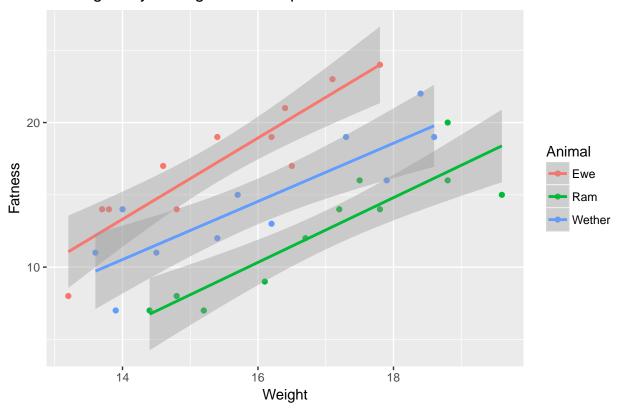
```
## Anova Table (Type III tests)
##
## Response: fatness
##
               Sum Sq Df F value
                                  Pr(>F)
## (Intercept)
               64.571 1 14.6355 0.0007007 ***
## weight
               5.293
                      2 0.5998 0.5560672
## Animal
## weight:animal
                8.957 2
                         1.0151 0.3757923
## Residuals
              119.123 27
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

The interaction term is not significant, thus no violation. We can also plot this to see it. All regression lines should be pointing in the same direction when we plot the groups separately.

```
library(ggplot2)
ggplot(deadsheep, aes(x = weight, y = fatness, color = Animal)) +
  geom_point() +
  geom_smooth(method = "lm") +
  labs(title = "Homogeneity of Regression Slopes", x = "Weight", y = "Fatness")
```

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Homogeneity of Regression Slopes



If one of these lines would be pointed downwards, we would have an interaction.

MANOVA

Repeated Measures ANOVA

Factor Analysis

Mixed Effects Models

Confirmatory Factor Analysis

This is a template file

Bibliography

Xie, Y. (2015). Dynamic Documents with R and knitr. Chapman and Hall/CRC, Boca Raton, Florida, 2nd edition. ISBN 978-1498716963.

Xie, Y. (2017). bookdown: Authoring Books and Technical Documents with R Markdown. R package version 0.5.