## 36-463/663: Multilevel & Hierarchical Models Fall 2016 HW02 – Due Tue, 13 Sept 2016

## Announcements

- Homework is due Tues 13 Sept on Blackboard. Submit a single file only. pdf preferred, doc or docx acceptable.
- You should have read or be finishing Gelman & Hill (G&H), Chapters 1 and 2. Please get started on reading G&H Chapters 3–4.
- This assignment involves both
  - Material from Ch 4 of Using R for Data Analysis and Graphics: Introduction, Code and Commentary, by J.
     H. Maindonald, Centre for Mathematics and Its Applications, Australian National University (usingR.pdf), and the accompanying data file (usingR.RData). You can get these at

http://www.stat.cmu.edu/~brian/463/hw01

just like last week.

You will also need the package "ggplot2"<sup>1</sup>, which you can install from the "Packages" menu in the R console window. After you install the lattice package, use the command library(ggplot2) to make ggplot functions like ggplot() or qplot() available for use. You can get a list of functions in the lattice library with library(help=ggplot2), and help on specific functions like qplot() with help(qplot).

- Examples and data sets from G&H, Chapters 1 and 2. You can get these from

http://www.stat.columbia.edu/~gelman/arm/examples/

## Exercises

- 1. [5 parts] Read all of Chapter 4 of Maindonald. Since the examples are in lattice rather than ggplot2 you do not have to try the things in the chapter, unless you are curious. You will need some trial-and-error to make sense of things. Then please do and turn in the following exercises from Chapter 4, using basic R graphics and ggplot2 as needed<sup>2</sup> (instead of lattice):
  - (a) Chapter 4, #1. Hint: To enter the data try something like this (see also additional hint on next page):

<sup>&</sup>lt;sup>1</sup>Recall, we are using ggplot2 instead of the lattice package discussed in the Maindonald pdf.

<sup>&</sup>lt;sup>2</sup>Please consult 02a-intro-ggplot-graphs.r and 02b-intro-ggplot-london.r from last week's lecture for examples of the kinds of plotting functions you'll need.

You will need commands like

- ggplot(...) + geom\_boxplot(...) + xlab(" ") + coord\_flip()
- ggplot(...) + geom\_dotplot(...)
- (b) Chapter 4, #2. Despite the use of the plural ("histograms", "normal probability plots", "density plots"), make one plot for each of the three parts of #2. Your plotting commands will be something like this (using ggplot2):
  - ggplot(...) + geom\_histogram(...)
  - ggplot(...) + stat\_qq(...)
  - ggplot(...) + geom\_density(...)
- (c) Re-do part (1b) but now using the built-in plotting functions:
  - hist()
  - qqnorm()
  - density()

Comment on any differences between (1b) and (1c).

(d) Chapter 4, #12. Hint: First, try something like

```
ggplot(possum,aes(x=hdlngth)) + geom_histogram() + facet_grid(sex ~ Pop)
This makes a set of histograms of hdlngth, taking into account sex and Pop. You want to do something
similar for each part of this problem.
```

- (e) Chapter 4, #13. Use ggplot2 commands, of course.
- 2. [4 parts] Please read and try everything in G&H Chapter 2. Note that some of the code is written out for you in the file book\_codes.zip at http://www.stat.columbia.edu/~gelman/arm/examples/. Then please do and turn in:
  - (a) G&H Chapter 2, #2. Note that part (b) asks you to conduct a hypothesis test, similar to the discussion in sections 2.3 and 2.4, but now comparing to a chi-squared distribution (qchisq()) under the null hypothesis.
  - (b) G&H Chapter 2, #3. Hint: Here are two possible correct solutions for this problem:

```
sums <- NULL
sims <- matrix(runif(20*1000),nrow=1000)</pre>
sums <- apply(sims,1,sum)</pre>
                                                     for (i in 1:1000) {
hist(sums,probability=TRUE)
                                                       data <- runif(20)</pre>
                                                       newsum <- sum(data)</pre>
m <- mean(sums)</pre>
                                                        sums <- c(sums,newsum)</pre>
s <- sd(sums)</pre>
x <- seq(min(sums),max(sums),length=100)</pre>
                                                       }
lines(x,dnorm(x,m,s))
                                                     hist(sums,probability=TRUE)
                                                     m <- mean(sums)</pre>
                                                     s <- sd(sums)</pre>
                                                     x <- seq(min(sums),max(sums),length=100)</pre>
                                                     lines(x,dnorm(x,m,s))
```

Whether you copy mine or create your own, write in detail about what each line of the code you use does, and how it helps move toward a solution to the problem. Use the help() function liberally, experiment with the functions to see what they do, and use any other tools you need, to give a good explanation of each line.

(c) G&H Chapter 2, #4. *Hints:* The first part of this problem is basically a more elaborate version of problem 2b: 1000 times, you need to do something like this:

```
mens_heights <- rnorm(n=100,mean=69.1,sd=2.9)
womens_heights <- rnorm(n=100,mean=63.7,sd=2.7)
x <- mean(mens_heights)
y <- mean(womens_heights)
height_diff <- x - y</pre>
```

Then make a histogram of your 1000 values of  $height_diff$ . Is the exact mean of x - y roughly in the middle of the histogram? Is it roughly equal to the mean of the 1000 values of  $height_diff$ ? Is the exact standard deviation (SD) roughly equal to the sample SD of the 1000 values of  $height_diff$ ? Is the histogram approximately six SD's wide [why is this an appropriate question?]?

(d) G&H Chapter 2, #5. Use the following information to complete this problem: The heights of men are approximately normally distributed with mean 69.1 inches and SD 2.9 inches. The heights of women are approximately normally distributed with mean 63.7 inches and SD 2.7 inches.

Hints:

• You should first find Cov(x, y) from the fact that Cor(x, y) = 0.3, using the well-known formula

$$\operatorname{Cor}(x, y) = \frac{\operatorname{Cov}(x, y)}{\operatorname{SD}(x)\operatorname{SD}(y)}$$

• Then you will need to apply the well-known formulas

$$E[ax + by + c] = aE[x] + bE[y] + c$$
  
Var (ax + by + c) = a<sup>2</sup>Var (x) + 2abCov (x, y) + b<sup>2</sup>Var (y)

No R is required for this problem. It is a straight calculation, like in 36-226 or a similar math stat course, using a little algebra and a calculator.