Statistics 312/612, fall 2016 Homework # 4 Due: Monday 3 October

[1] The catheter data set is taken from a well known text book. If you happen to know the book please do not just repeat the analysis it presents.

```
cath <- read.table("catheter.txt",header=T)
outHW <- lm(distance ~ height + weight,cath)
outH <- lm(distance ~ height, cath)
outW <-lm(distance ~ weight, cath)
cath[1:3,]
## height weight distance
## 1 42.8 40.0 37.0
## 2 63.5 93.5 49.5
## 3 37.5 35.5 34.5</pre>
```

The summary information for each fit (outHW, outH, outW) seems to suggest that height by itself is a good predictor of distance, that weight by itself is a good predictor of distance, but when both predictors are used then neither is particularly useful. (The stars in the summary table suggest 'significance'.)

(i) (15 points) Add more variables to the cath data.frame: height and weight centered to zero means (call them heen and ween); and things like wres <-lm(wcen~ heen, cath)\$res. Explain why some of the coeffs and std. errors are the same and some are different for the models:

```
distance ~ height
distance ~ hcen
distance ~ hcen + wres
distance ~ height+wres
distance ~ hcen+wcen
distance ~ height+weight
```

- (ii) (5 points) Explains why summary(outHW) is misleading regarding the value of height and weight as predictors.
- (iii) (5 points) Explain how the output from cor(cath) is relevant to the problem.
- [2] The handout two_factors.pdf showed how to calculate several least squares fits using the Box-Cox data:

```
BC <- read.table("../Handouts/boxcox.data", header=T,sep="\t")
BC$rate <- 1/BC$time # transformation suggested by BHH page 235
BC$Htreatment <- C(BC$treatment,helmert)
out5 <- lm(rate ~ -1 + treatment,BC)
out7 <- lm(rate ~ Htreatment,BC)</pre>
```

In class I showed (page 9 of the handout) how to transform results from one parametrization into results for a different parametrization, using out5 and out6 as an example. For this homework problem I want you to recreate the shortened summary

```
## lm(formula = rate ~ -1 + treatment, data = BC)
## tA tB tC tD
## Est 3.519 1.862 2.947 2.161
## StdErr 0.292 0.292 0.292 0.292
```

using only the information contained in out7, which is essentially the same as the out9 generated by:

```
C7 <- contrasts(BC$Htreatment)

dummyT <- outer(BC$treat,levels(BC$treat),"==")+0

X7 <- cbind(1, dummyT ½*% C7)

out9 <- lm(BC$rate ~ -1+X7)
```

Display all the \mathbf{R} code that you use.

- (i) (5 points) Show that X7 is equal to dummyT %*% K7 where K7 <- cbind(1,C7).
- (ii) (10 points) If \hat{g} is the vector of coefficients from out7 and \hat{b} is the vector of coefficients from out5, show that $\hat{b} = K_7 \hat{g}$.
- (iii) (10 points) Use (ii) and $\tt out7$ to recreate the shortened summary for $\tt out5.$