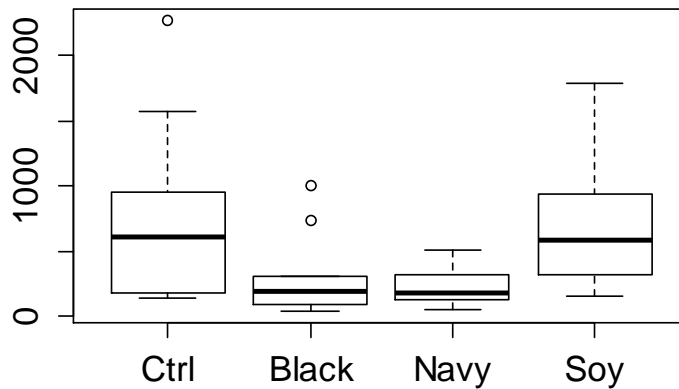


## STAT511 HW#7 KEY

32 points total, 2 points per problem unless otherwise noted.

### #1 Dog Diets ANOVA

#### A. Boxplots



#### B. ANOVA Table (Original Scale)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Diet	3	2623607	874536	3.723	0.0198 *
Residuals	36	8457190	234922		

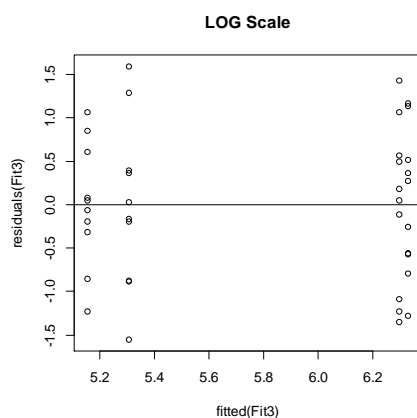
#### C. ANOVA Assumptions (4 pts)

From the plot of residuals vs fitted values, we see a clear megaphone shape indicating that the assumption of equal variances is NOT satisfied.

From the qqplot of residuals, we see curvature indicating that the assumption of normality is NOT satisfied.

D. Levene's test p-value = 0.0484. Reject  $H_0$ ; we cannot conclude equal variances.

#### E. Log Transformation



F. ANOVA (after log transformation)

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Diet	3	11.80	3.934	5.137	0.00464 **
Residuals	36	27.57	0.766		

p-value < 0.05

Reject H0 and conclude not all the means are the same

G. Unadjusted pairwise comparisons

	Estimate	Std. Error	t value	Pr(> t )
Black - Ctrl == 0	-0.98818	0.39133	-2.525	0.01611 *
Navy - Ctrl == 0	-1.14118	0.39133	-2.916	0.00607 **
Soy - Ctrl == 0	0.03198	0.39133	0.082	0.93532
Navy - Black == 0	-0.15300	0.39133	-0.391	0.69813
Soy - Black == 0	1.02016	0.39133	2.607	0.01321 *
Soy - Navy == 0	1.17316	0.39133	2.998	0.00490 **

H. Tukey Comparisons

	Estimate	Std. Error	t value	Pr(> t )
Black - Ctrl == 0	-0.98818	0.39133	-2.525	0.0731 .
Navy - Ctrl == 0	-1.14118	0.39133	-2.916	0.0295 *
Soy - Ctrl == 0	0.03198	0.39133	0.082	0.9998
Navy - Black == 0	-0.15300	0.39133	-0.391	0.9794
Soy - Black == 0	1.02016	0.39133	2.607	0.0606 .
Soy - Navy == 0	1.17316	0.39133	2.998	0.0242 *

I. HSD= 1.05415

J. "Lines" display

Navy	Black	Ctrl	Soy
5.155598	5.308594	6.296778	6.328757

K. Dunnetts' Comparisons

	Estimate	Std. Error	t value	Pr(> t )
Black - Ctrl == 0	-0.98818	0.39133	-2.525	0.0421 *
Navy - Ctrl == 0	-1.14118	0.39133	-2.916	0.0166 *
Soy - Ctrl == 0	0.03198	0.39133	0.082	0.9996

L. Contrasts (8 pts)

DietCtrl	DietBlack	DietNavy	DietSoy
6.297	5.309	5.156	6.329

Estimate	Std. Error	t value	Pr(> t )
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C1 == 0	0.1530	0.3913	0.391	0.698129	
C2 == 0	-1.0647	0.3389	-3.142	0.003354	**
C3 == 0	-1.0967	0.3389	-3.236	0.002602	**
C4 == 0	-1.0807	0.2767	-3.905	0.000397	***

## #R Code

### #Q1

```
Diets <-
read.csv("C:/hess/STAT511_FA11/HW_2015/HW7/DogDiets.csv")
Diets <- data.frame(Diets, sqrtTrig = sqrt(Diets$Trig), logTrig
= log(Diets$Trig))
str(Diets)
levels(Diets$Diet)
Diets$Diet <- factor(Diets$Diet, levels(Diets$Diet)[c(2, 1, 3,
4)])
```

### #Original Scale

```
boxplot(Trig ~ Diet, data = Diets)
Fit1 <- aov(Trig ~ Diet, data = Diets)
summary(Fit1)
plot(Fit1)
plot(residuals(Fit1) ~ fitted(Fit21), main = "Original
Scale");abline(h=0)
library(car)
leveneTest(Trig ~ Diet, data = Diets)
```

### #SQRT Scale

```
Fit2 <- aov(sqrtTrig ~ Diet, data = Diets)
plot(residuals(Fit2) ~ fitted(Fit2), main = "SQRT
Scale");abline(h=0)
leveneTest(sqrtTrig ~ Diet, data = Diets)
```

### #LOG Scale

```
Fit3 <- aov(logTrig ~ Diet, data = Diets)
plot(residuals(Fit3) ~ fitted(Fit3), main = "LOG
Scale");abline(h=0)
leveneTest(logTrig ~ Diet, data = Diets)
summary(Fit3)
```

### #Pairwise Comparisons (G-K)

```
library(multcomp)
PairComps <- glht(Fit3, linfct = mcp(Diet = "Tukey"))
summary(PairComps, test = adjusted(type = "none"))
summary(PairComp)
HSD <- qtkey(0.95, 4, 36)*sqrt(0.766/10)
cld(PairComps)
DunnettComps <- glht(Fit3, linfct = mcp(Diet = "Dunnett"))
summary(DunnettComps)
```

### #Contrasts (L)

```
C1 <- c(0, 1, -1, 0)
```

```
C2 <- c(-1, 0.5, 0.5, 0)
C3 <- c(0, 0.5, 0.5, -1)
C4 <- c(-0.5, 0.5, 0.5, -0.5)
Cmat <- t(cbind(C1, C2, C3, C4))
colnames(Cmat) <- c("Ctrl", "Black", "Navy", "Soy")
Fit4 <- lm(logTrig ~ Diet - 1, data = Diets)
Fit4
contrresults <- glht(Fit4, linfct = Cmat)
contrresults
summary(contrresults, test = adjusted(type= "none"))
```