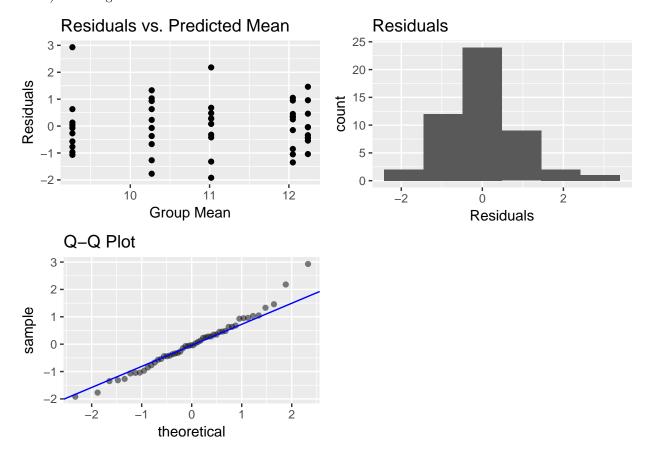
Chapter 8 And 9

Kyle Ligon

9.13 a) Checking the results from Proc Mixed in order to do ANOVA

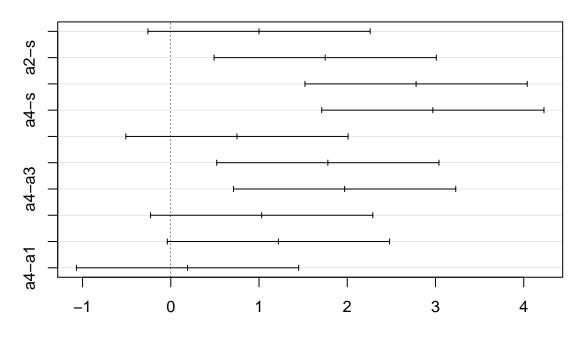


9.13b) Perform ANOVA test on the data: Show ANOVA Table First, then Run the Test
anova_mod

```
## Call:
##
      aov(formula = wt_loss ~ treatment, data = gather_frame)
##
## Terms:
##
                   treatment Residuals
## Sum of Squares
                       61.618
                                 44.207
## Deg. of Freedom
                                     45
## Residual standard error: 0.9911497
## Estimated effects may be unbalanced
summary(anova_mod)
##
               Df Sum Sq Mean Sq F value
                                    15.68 4.16e-08 ***
## treatment
                   61.62 15.404
                   44.21
                            0.982
## Residuals
               45
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
9.13 c) Perform Tukey's W on the significant pairs
real_w <- TukeyHSD(anova_mod, ordered = TRUE)</pre>
real_w$treatment
         diff
                     lwr
                              upr
                                         p adj
## a3-s
         1.00 -0.2594887 2.259489 1.784060e-01
         1.75
              0.4905113 3.009489 2.428628e-03
## a1-s
        2.78
              1.5205113 4.039489 1.200843e-06
        2.97
               1.7105113 4.229489 2.780828e-07
## a2-a3 0.75 -0.5094887 2.009489 4.490082e-01
## a1-a3 1.78 0.5205113 3.039489 1.980323e-03
## a4-a3 1.97
              0.7105113 3.229489 5.243121e-04
## a1-a2 1.03 -0.2294887 2.289489 1.563263e-01
## a4-a2 1.22 -0.0394887 2.479489 6.176067e-02
## a4-a1 0.19 -1.0694887 1.449489 9.927171e-01
plot(real_w)
```

95% family-wise confidence level



Differences in mean levels of treatment

9.13 d) Use Dunnett's to see if any of the new agents have significantly larger mean weights loss as compared to the standard agent. alpha = 0.05

```
##
## Simultaneous Tests for General Linear Hypotheses
##
## Multiple Comparisons of Means: User-defined Contrasts
##
```

```
##
## Fit: aov(formula = wt_loss ~ treatment, data = gather_frame)
##
## Linear Hypotheses:
##
              Estimate Std. Error t value Pr(>|t|)
## a1 - s == 0
                 2.7800
                            0.4433
                                     6.272
                                             <0.001 ***
## a2 - s
                 1.7500
                            0.4433
                                     3.948
                                             <0.001 ***
## a3 - s == 0
                 1.0000
                            0.4433
                                     2.256
                                              0.093 .
## a4 - s == 0
                 2.9700
                            0.4433
                                     6.700
                                             <0.001 ***
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Adjusted p values reported -- single-step method)
```

Sesignals 0 -10-

150

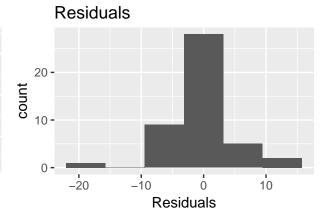
130

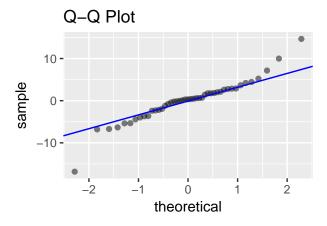
170

Group Mean

190

Residuals vs. Predicted Mean





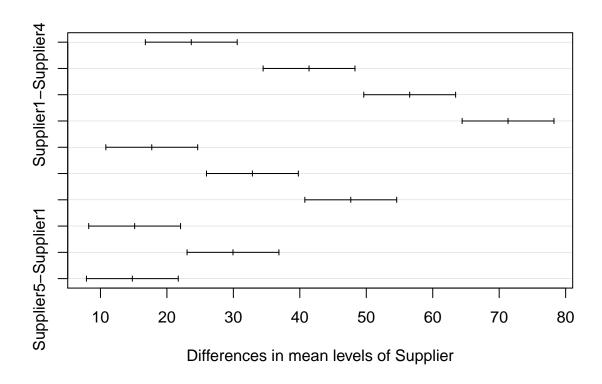
b) Perform an Anova

anova_lenses

```
## Call:
## aov(formula = lov ~ Supplier, data = gather_lenses)
##
## Terms:
## Supplier Residuals
## Sum of Squares 28024.350 1053.789
## Deg. of Freedom 4 40
##
## Residual standard error: 5.132711
## Estimated effects may be unbalanced
```

```
summary(anova_lenses)
               Df Sum Sq Mean Sq F value Pr(>F)
##
               4 28024 7006 265.9 <2e-16 ***
## Supplier
## Residuals
               40
                  1054
                              26
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
  c) Run a Kruskal-Wallis
kw_lenses <- kruskal.test(lov ~ Supplier, data = gather_lenses)</pre>
kw_lenses
##
## Kruskal-Wallis rank sum test
## data: lov by Supplier
## Kruskal-Wallis chi-squared = 41.596, df = 4, p-value = 2.023e-08
kw_lenses$statistic
## Kruskal-Wallis chi-squared
##
                      41.5963
 d) Use Tukey's W to find out which pairs are significantly different.
real_lenses <- TukeyHSD(anova_lenses, ordered = TRUE)</pre>
real_lenses$treatment
## NULL
plot(real_lenses)
```

95% family-wise confidence level



real_lenses\$Supplier

```
## Supplier3-Supplier4 23.66667 16.756116 30.57722 3.656697e-11
## Supplier2-Supplier4 41.38889 34.478338 48.29944 4.636291e-13
## Supplier1-Supplier4 56.53333 49.622783 63.44388 4.636291e-13
## Supplier5-Supplier4 71.33333 64.422783 78.24388 4.636291e-13
## Supplier2-Supplier3 17.72222 10.811672 24.63277 6.520741e-08
## Supplier1-Supplier3 32.86667 25.956116 39.77722 4.682921e-13
## Supplier5-Supplier3 47.66667 40.756116 54.57722 4.636291e-13
## Supplier1-Supplier2 15.14444 8.233894 22.05499 1.978429e-06
## Supplier5-Supplier2 29.94444 23.033894 36.85499 5.020429e-13
## Supplier5-Supplier1 14.80000 7.889450 21.71055 3.129018e-06
```

f) Run a KW pairwise comparison on the different suppliers to see if there's a difference in variability of power

```
ranks_1 <- c(39,32, 33, 28, 29, 34, 35, 31, 36)
ranks_2 <- c(19, 20, 21, 22, 23, 24, 25, 26, 27)
ranks_3 <- c(10, 11, 12, 13, 14, 15, 16, 17, 18)
ranks_4 <- c(1, 2, 3, 4, 5, 6, 7, 8, 9)
ranks_5 <- c(38, 39, 30, 40, 41, 42, 43, 44, 45)
ranks_table = data.frame(cbind(ranks_1, ranks_2, ranks_3, ranks_4, ranks_5))
colnames(ranks_table) <- c("Ranks_Group1", "Ranks_Group2", "Ranks_Group3", "Ranks_Group4", "Ranks
```