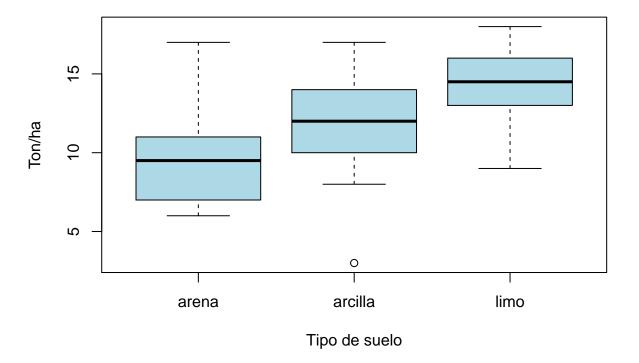
Clase 5.R

Usuario

2019-08-09

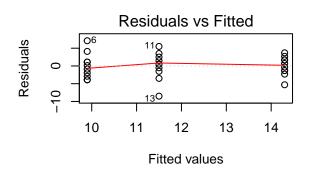
```
#Nallely Aguirre
#09/08/2019
#Clase_5
arena <- c(6, 10, 8, 6, 14, 17, 9, 11, 7, 11)
arcilla <- c(17, 15, 3, 11, 14, 12, 12, 8, 10, 13)
limo <- c(13, 16, 9,12, 15, 16, 17, 13, 18, 14)
y.ton <-c(arena, arcilla, limo)
suelo <- gl(3, 10, 30, labels=c("arena", "arcilla", "limo"))</pre>
prod <-data.frame(suelo, y.ton)</pre>
head(prod)
##
     suelo y.ton
## 1 arena
## 2 arena
             10
## 3 arena
              8
## 4 arena
              6
## 5 arena
              14
## 6 arena
              17
tapply(prod$y.ton, prod$suelo, mean)
##
     arena arcilla
                      limo
##
       9.9
              11.5
                      14.3
tapply(prod$y.ton, prod$suelo, var)
       arena
              arcilla
                            limo
## 12.544444 15.388889 7.122222
shapiro.test(prod$y.ton)
##
   Shapiro-Wilk normality test
##
##
## data: prod$y.ton
## W = 0.97214, p-value = 0.5993
bartlett.test(prod$y.ton, prod$suelo)
##
## Bartlett test of homogeneity of variances
##
## data: prod$y.ton and prod$suelo
## Bartlett's K-squared = 1.2764, df = 2, p-value = 0.5283
```

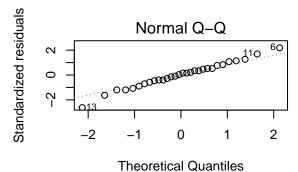


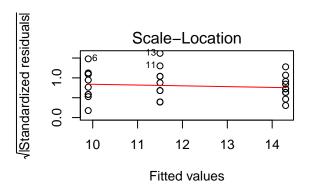
```
aov.suelo <- aov(prod$y.ton ~ prod$suelo)</pre>
aov.suelo
## Call:
      aov(formula = prod$y.ton ~ prod$suelo)
##
##
## Terms:
                    prod$suelo Residuals
##
## Sum of Squares
                          99.2
                                   315.5
## Deg. of Freedom
                             2
                                       27
##
## Residual standard error: 3.41836
## Estimated effects may be unbalanced
```

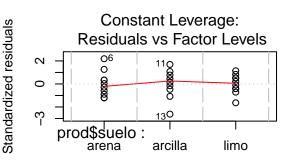
summary(aov.suelo)

```
## Df Sum Sq Mean Sq F value Pr(>F)
## prod$suelo 2 99.2 49.60 4.245 0.025 *
## Residuals 27 315.5 11.69
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
par(mfrow=c(2,2))
plot(aov(prod$y.ton ~ prod$suelo))
```









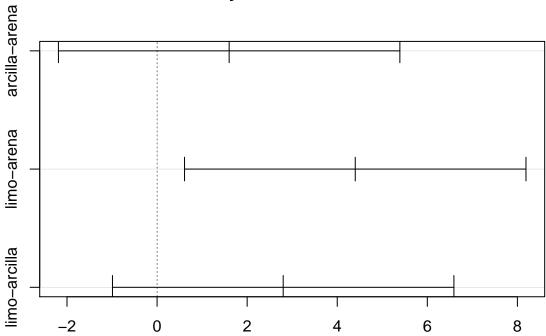
Factor Level Combinations

```
par(mfrow=c(1,1))
TukeyHSD(aov.suelo, conf.level = 0.95)
```

```
##
     Tukey multiple comparisons of means
##
       95% family-wise confidence level
##
## Fit: aov(formula = prod$y.ton ~ prod$suelo)
##
## $`prod$suelo`
                 diff
                             lwr
                                      upr
                                               p adj
## arcilla-arena 1.6 -2.1903777 5.390378 0.5546301
                  4.4 0.6096223 8.190378 0.0204414
## limo-arena
                  2.8 -0.9903777 6.590378 0.1785489
## limo-arcilla
```

plot(TukeyHSD(aov.suelo))

95% family-wise confidence level



Differences in mean levels of prod\$suelo

```
summary(aov.suelo)
              Df Sum Sq Mean Sq F value Pr(>F)
## prod$suelo
                                 4.245 0.025 *
               2
                    99.2
                           49.60
## Residuals
              27 315.5
                           11.69
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
summary.lm(aov.suelo)
##
## Call:
## aov(formula = prod$y.ton ~ prod$suelo)
## Residuals:
##
     Min
              1Q Median
                            3Q
                                  Max
     -8.5
            -1.8
                    0.3
                           1.7
                                  7.1
##
##
## Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                        9.900
                                   1.081
                                           9.158 9.04e-10 ***
## prod$sueloarcilla
                        1.600
                                   1.529
                                           1.047 0.30456
## prod$suelolimo
                        4.400
                                   1.529
                                           2.878 0.00773 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
##
```

Residual standard error: 3.418 on 27 degrees of freedom
Multiple R-squared: 0.2392, Adjusted R-squared: 0.1829
F-statistic: 4.245 on 2 and 27 DF, p-value: 0.02495

Ho, la media de la produccion de toneladas no es diferente en ninguno de los # tratamientos

H1, la media de la producion d elos tratamientos de al menos uno es diferente a # los demas