

# Clase6.R

Usuario

2019-08-12

```
#Emmanuel Ferrer
```

```
#Clase 6
```

```
#09/08/2019
```

```
#Analisis co-varianza
```

```
rascon <-read.csv("C:/MCF202-2019/rascon.csv", header= T)
head(rascon)
```

```
##   arbol  DAP EDAD      SP
## 1     1 27.4   59 arizonica
## 2     2 19.5   29 arizonica
## 3     3 20.0   24 arizonica
## 4     4 22.0   40 arizonica
## 5     5 34.0   50 arizonica
## 6     6 33.1   44 arizonica
```

```
summary(rascon)
```

```
##      arbol      DAP      EDAD      SP
## Min.   : 1.00   Min.   : 7.70   Min.   : 14.00   arizonica :30
## 1st Qu.:15.75   1st Qu.:16.62   1st Qu.: 28.75   durangensis:30
## Median :30.50   Median :22.05   Median : 41.50
## Mean   :30.50   Mean   :23.24   Mean   : 48.03
## 3rd Qu.:45.25   3rd Qu.:28.98   3rd Qu.: 58.25
## Max.   :60.00   Max.   :45.60   Max.   :128.00
```

```
# Actividad a realizar -----
```

```
#Estadística descriptiva
```

```
mean(rascon$DAP)
```

```
## [1] 23.24167
```

```
mean(rascon$EDAD)
```

```
## [1] 48.03333
```

```
sd(rascon$DAP)
```

```
## [1] 9.244921
```

```
sd(rascon$EDAD)
```

```
## [1] 28.26507
```

```
var(rascon$DAP)
```

```
## [1] 85.46857
```

```
var(rascon$EDAD)
```

```
## [1] 798.9141
```

```

# Correlacion -----
shapiro.test(rascon$EDAD)

##
##  Shapiro-Wilk normality test
##
## data:  rascon$EDAD
## W = 0.87401, p-value = 1.684e-05

shapiro.test(rascon$DAP)

##
##  Shapiro-Wilk normality test
##
## data:  rascon$DAP
## W = 0.97282, p-value = 0.2

cor.test(rascon$EDAD, rascon$DAP)

##
##  Pearson's product-moment correlation
##
## data:  rascon$EDAD and rascon$DAP
## t = 9.9917, df = 58, p-value = 3.201e-14
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6783908 0.8729475
## sample estimates:
##          cor
## 0.7953145

# Conclusion de correlacion -----
#la correlacion r=0.7953145

#correlacion:
#la correlacion entre las variables es siginificativa porque se
#observa que el valor de p-value < 3.201e-14.
#cuales el coeficiente de correlacion (r)?

# Analisis de covarianza -----
lm.rasco <- lm(rascon$EDAD ~ rascon$DAP)
lm.rasco

##
## Call:
## lm(formula = rascon$EDAD ~ rascon$DAP)
##
## Coefficients:
## (Intercept)  rascon$DAP
##      -8.480      2.432

summary(lm.rasco)

##
## Call:
## lm(formula = rascon$EDAD ~ rascon$DAP)

```

```
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -28.005 -12.539   0.270   7.457  49.630
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  -8.4803     6.0802  -1.395   0.168
## rascon$DAP    2.4316     0.2434   9.992 3.2e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 17.28 on 58 degrees of freedom
## Multiple R-squared:  0.6325, Adjusted R-squared:  0.6262
## F-statistic: 99.83 on 1 and 58 DF,  p-value: 3.201e-14
```

```
length(rascon$EDAD)
```

```
## [1] 60
```

```
# identificar columna SP como factor
```

```
arizedad <- factor(rascon$SP)
arizedad
```

```
## [1] arizonica arizonica arizonica arizonica arizonica
## [6] arizonica arizonica arizonica arizonica arizonica
## [11] arizonica arizonica arizonica arizonica arizonica
## [16] arizonica arizonica arizonica arizonica arizonica
## [21] arizonica arizonica arizonica arizonica arizonica
## [26] arizonica arizonica arizonica arizonica arizonica
## [31] durangensis durangensis durangensis durangensis durangensis
## [36] durangensis durangensis durangensis durangensis durangensis
## [41] durangensis durangensis durangensis durangensis durangensis
## [46] durangensis durangensis durangensis durangensis durangensis
## [51] durangensis durangensis durangensis durangensis durangensis
## [56] durangensis durangensis durangensis durangensis durangensis
## Levels: arizonica durangensis
```

```
# separar factor
```

```
ariz <- subset(rascon, SP == "arizonica")
ariz
```

```
##      arbol  DAP EDAD      SP
## 1      1 27.4  59 arizonica
## 2      2 19.5  29 arizonica
## 3      3 20.0  24 arizonica
## 4      4 22.0  40 arizonica
## 5      5 34.0  50 arizonica
## 6      6 33.1  44 arizonica
## 7      7 32.0  44 arizonica
## 8      8 10.0  17 arizonica
## 9      9 14.0  15 arizonica
## 10     10 11.0  16 arizonica
## 11     11 21.6  23 arizonica
## 12     12 11.2  14 arizonica
## 13     13  8.0  18 arizonica
## 14     14 22.1  22 arizonica
```

```
## 15    15 10.0   20 arizonica
## 16    16 16.8   20 arizonica
## 17    17 17.9   22 arizonica
## 18    18 26.1   40 arizonica
## 19    19  7.7   15 arizonica
## 20    20 11.3   25 arizonica
## 21    21 13.5   26 arizonica
## 22    22 10.7   28 arizonica
## 23    23 17.0   34 arizonica
## 24    24 20.0   33 arizonica
## 25    25 24.7   46 arizonica
## 26    26 21.9   42 arizonica
## 27    27 25.2   39 arizonica
## 28    28 28.9   43 arizonica
## 29    29 29.2   44 arizonica
## 30    30 12.6   29 arizonica
```

```
ariz.lm <-lm(ariz$EDAD ~ ariz$DAP)
ariz.lm
```

```
##
## Call:
## lm(formula = ariz$EDAD ~ ariz$DAP)
##
## Coefficients:
## (Intercept)      ariz$DAP
##          5.333          1.313
```

```
dura <-subset(rascon, SP == "durangensis")
dura
```

```
##   arbol  DAP EDAD      SP
## 31   31 16.8   82 durangensis
## 32   32 25.9   32 durangensis
## 33   33 43.7  125 durangensis
## 34   34 41.2  103 durangensis
## 35   35 24.2   45 durangensis
## 36   36 37.1  108 durangensis
## 37   37 35.5   58 durangensis
## 38   38 45.6   75 durangensis
## 39   39 38.0   89 durangensis
## 40   40 21.9   40 durangensis
## 41   41 18.5   36 durangensis
## 42   42 22.3   59 durangensis
## 43   43 16.1   56 durangensis
## 44   44 17.6   34 durangensis
## 45   45 24.8   57 durangensis
## 46   46 16.1   41 durangensis
## 47   47 15.2   38 durangensis
## 48   48 25.8   63 durangensis
## 49   49 23.2   42 durangensis
## 50   50 32.0  106 durangensis
## 51   51 33.2   82 durangensis
## 52   52 27.1   60 durangensis
## 53   53 16.0   37 durangensis
## 54   54 29.7   69 durangensis
```

```
## 55    55 39.2  128 durangensis
## 56    56 35.5  113 durangensis
## 57    57 24.8   54 durangensis
## 58    58 21.1   55 durangensis
## 59    59 25.7   32 durangensis
## 60    60 21.3   42 durangensis

# Regresion dos factores
cov.edad <-lm(rascon$EDAD ~ rascon$DAP + rascon$SP)
cov.edad

##
## Call:
## lm(formula = rascon$EDAD ~ rascon$DAP + rascon$SP)
##
## Coefficients:
##           (Intercept)           rascon$DAP  rascon$SPdurangensis
##                -7.657                1.986                19.063

summary(cov.edad)

##
## Call:
## lm(formula = rascon$EDAD ~ rascon$DAP + rascon$SP)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -30.844  -8.515  -1.731   7.473  38.741
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -7.6573     5.2903  -1.447   0.153
## rascon$DAP       1.9861     0.2342   8.480 1.10e-11 ***
## rascon$SPdurangensis 19.0629     4.2942   4.439 4.19e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 15.03 on 57 degrees of freedom
## Multiple R-squared:  0.7269, Adjusted R-squared:  0.7174
## F-statistic: 75.87 on 2 and 57 DF,  p-value: < 2.2e-16

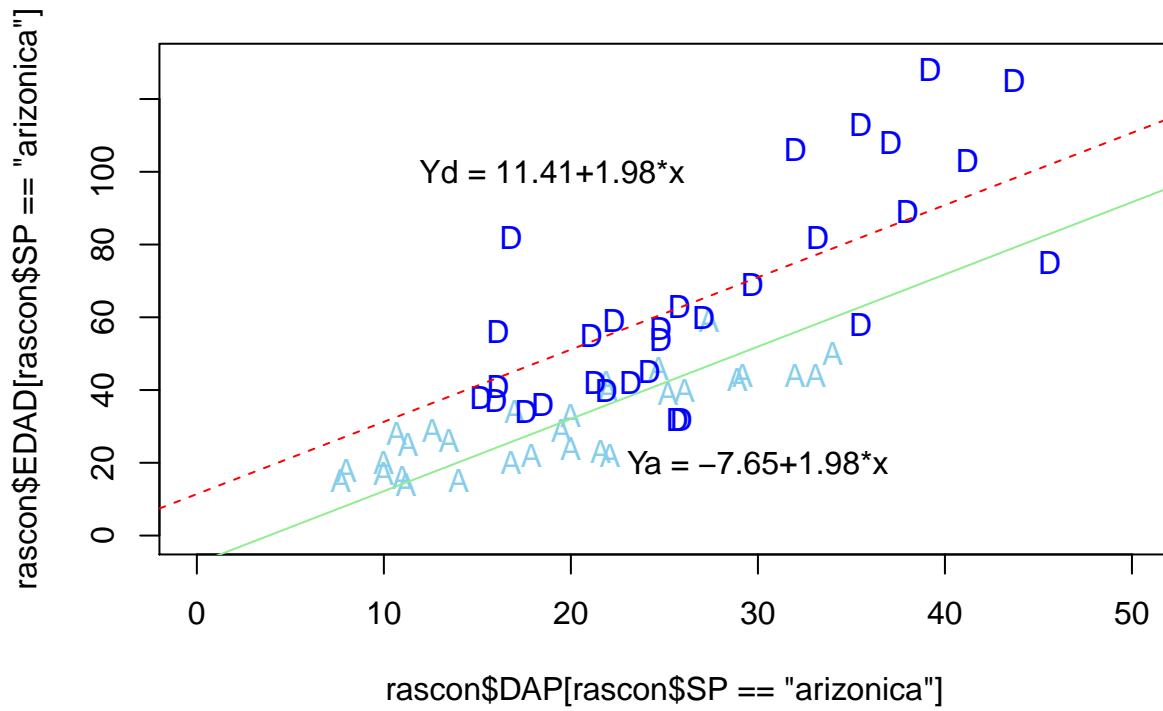
#existe diferencia significativas
#p-value que ambas lineas son difenrentes significativas

#aqui se compara en diferentes especies
plot(rascon$DAP[rascon$SP == "arizonica"], rascon$EDAD[rascon$SP == "arizonica"],
     col="sky blue", pch="A", xlim=c(0,50),ylim=c(0,130))
abline(cov.edad$coefficients[1], cov.edad$coefficients[2], col="light green")
text(30,20, "Ya = -7.65+1.98*x")

points(rascon$DAP[rascon$SP == "durangensis"], rascon$EDAD[rascon$SP == "durangensis"],
       col="blue", pch="D")

abline(cov.edad$coefficients[1] + cov.edad$coefficients[3],
       cov.edad$coefficients[2], col="red", lty="dashed")
```

```
text(19,100, "Yd = 11.41+1.98*x")
```



```
#h0.no EXISTE COVARIACION ENTRE LA LINEA DE REGRESION
#h1= EXISTE covaraicion entre la linea de regresion
```