

Script_04.R

jairo

2025-09-25

```
# JAIRO ALBERTO LEAL GOMEZ
```

```
# 18/09/2025
```

```
# SEMANA 7
```

```
# SCRIPT 4
```

```
datos <- read.csv("crop.csv", header = T)
```

```
datos
```

##	density	block	fertilizer	yield
## 1	1	1	1	177.2287
## 2	2	2	1	177.5500
## 3	1	3	1	176.4085
## 4	2	4	1	177.7036
## 5	1	1	1	177.1255
## 6	2	2	1	176.7783
## 7	1	3	1	176.7463
## 8	2	4	1	177.0612
## 9	1	1	1	176.2749
## 10	2	2	1	177.9672
## 11	1	3	1	176.6013
## 12	2	4	1	177.0305
## 13	1	1	1	177.4795
## 14	2	2	1	176.8741
## 15	1	3	1	176.1144
## 16	2	4	1	176.0084
## 17	1	1	1	176.1083
## 18	2	2	1	178.3574
## 19	1	3	1	177.2624
## 20	2	4	1	176.9188
## 21	1	1	1	176.2390
## 22	2	2	1	176.5731
## 23	1	3	1	176.0393
## 24	2	4	1	176.8179
## 25	1	1	1	176.1606
## 26	2	2	1	177.2264
## 27	1	3	1	175.9385
## 28	2	4	1	177.1649
## 29	1	1	1	175.3608
## 30	2	2	1	177.2770
## 31	1	3	1	175.9454
## 32	2	4	1	175.8828
## 33	1	1	2	176.4793

## 34	2	2	2 176.0443
## 35	1	3	2 177.4125
## 36	2	4	2 177.3608
## 37	1	1	2 177.3855
## 38	2	2	2 176.9758
## 39	1	3	2 177.3798
## 40	2	4	2 177.9980
## 41	1	1	2 176.4349
## 42	2	2	2 176.9333
## 43	1	3	2 175.9835
## 44	2	4	2 177.0341
## 45	1	1	2 176.4368
## 46	2	2	2 176.0677
## 47	1	3	2 177.1210
## 48	2	4	2 177.1977
## 49	1	1	2 176.6037
## 50	2	2	2 177.2082
## 51	1	3	2 177.1488
## 52	2	4	2 176.8191
## 53	1	1	2 176.9991
## 54	2	2	2 178.1346
## 55	1	3	2 176.4292
## 56	2	4	2 176.6683
## 57	1	1	2 176.8959
## 58	2	2	2 177.7795
## 59	1	3	2 176.4145
## 60	2	4	2 176.8789
## 61	1	1	2 177.5807
## 62	2	2	2 176.9573
## 63	1	3	2 175.7475
## 64	2	4	2 177.3526
## 65	1	1	3 177.1042
## 66	2	2	3 178.0796
## 67	1	3	3 176.9034
## 68	2	4	3 177.5403
## 69	1	1	3 177.0327
## 70	2	2	3 178.2860
## 71	1	3	3 176.4054
## 72	2	4	3 176.4308
## 73	1	1	3 177.3963
## 74	2	2	3 176.9256
## 75	1	3	3 177.0550
## 76	2	4	3 177.3442
## 77	1	1	3 177.1284
## 78	2	2	3 177.1683
## 79	1	3	3 176.3539
## 80	2	4	3 179.0609
## 81	1	1	3 176.3005
## 82	2	2	3 177.5934
## 83	1	3	3 177.1152
## 84	2	4	3 177.7945
## 85	1	1	3 177.0040
## 86	2	2	3 178.0369
## 87	1	3	3 177.7014

```
## 88      2      4      3 177.6328
## 89      1      1      3 177.6523
## 90      2      2      3 177.1004
## 91      1      3      3 177.1880
## 92      2      4      3 177.4053
## 93      1      1      3 178.1416
## 94      2      2      3 177.7106
## 95      1      3      3 177.6873
## 96      2      4      3 177.1182
```

```
summary(datos)
```

```
##      density      block      fertilizer      yield
## Min.      :1.0    Min.    :1.00    Min.      :1    Min.      :175.4
## 1st Qu.:1.0    1st Qu.:1.75    1st Qu.:1    1st Qu.:176.5
## Median :1.5    Median :2.50    Median :2    Median :177.1
## Mean    :1.5    Mean    :2.50    Mean     :2    Mean     :177.0
## 3rd Qu.:2.0    3rd Qu.:3.25    3rd Qu.:3    3rd Qu.:177.4
## Max.     :2.0    Max.     :4.00    Max.      :3    Max.      :179.1
```

```
# Las variables estan en numeros integrales y hay que modificar a factores
# Ya que el experimento hara la comparación por factores
```

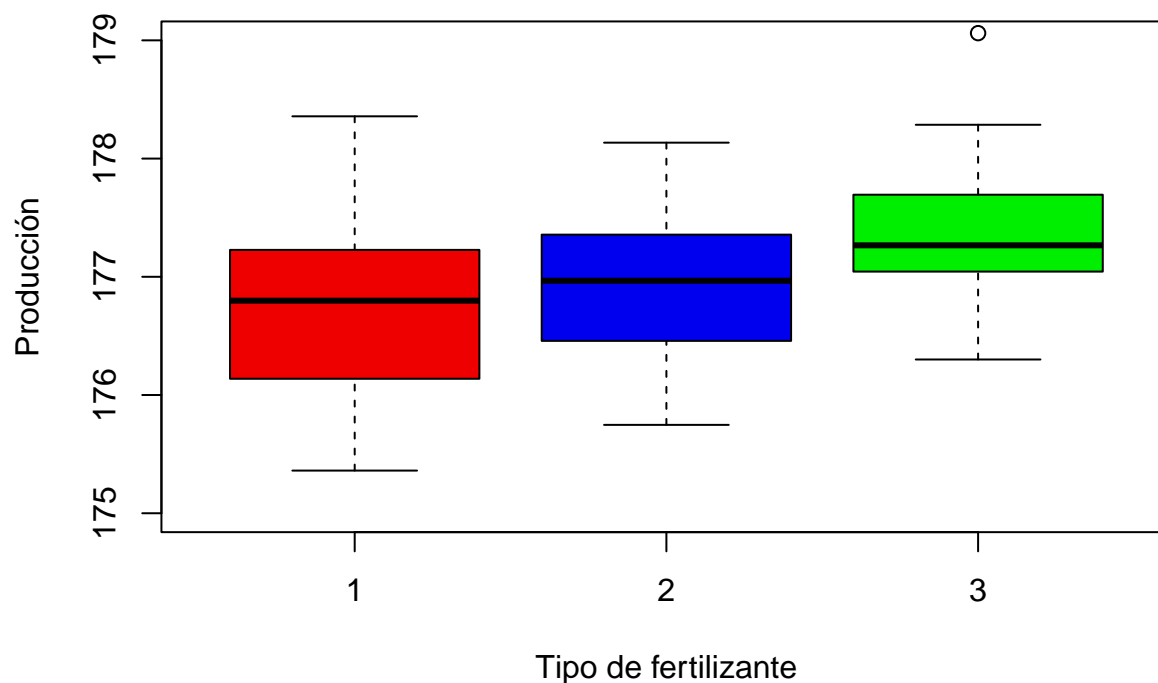
```
datos$density <- as.factor(datos$density)
datos$block <- as.factor(datos$block)
datos$fertilizer <- as.factor(datos$fertilizer)
```

```
summary(datos)
```

```
## density block fertilizer      yield
## 1:48    1:24    1:32      Min.      :175.4
## 2:48    2:24    2:32      1st Qu.:176.5
##        3:24    3:32      Median :177.1
##        4:24      Mean    :177.0
##              3rd Qu.:177.4
##              Max.     :179.1
```

```
boxplot(datos$yield ~ datos$fertilizer,
        main = "Producción por fertilizante",
        col= c("red2", "blue2", "green2"),
        ylab = "Producción",
        xlab = "Tipo de fertilizante",
        ylim = c(175, 179))
```

Producción por fertilizante



```
library(vioplot)
```

```
## Cargando paquete requerido: sm
```

```
## Package 'sm', version 2.2-6.0: type help(sm) for summary information
```

```
## Cargando paquete requerido: zoo
```

```
##
```

```
## Adjuntando el paquete: 'zoo'
```

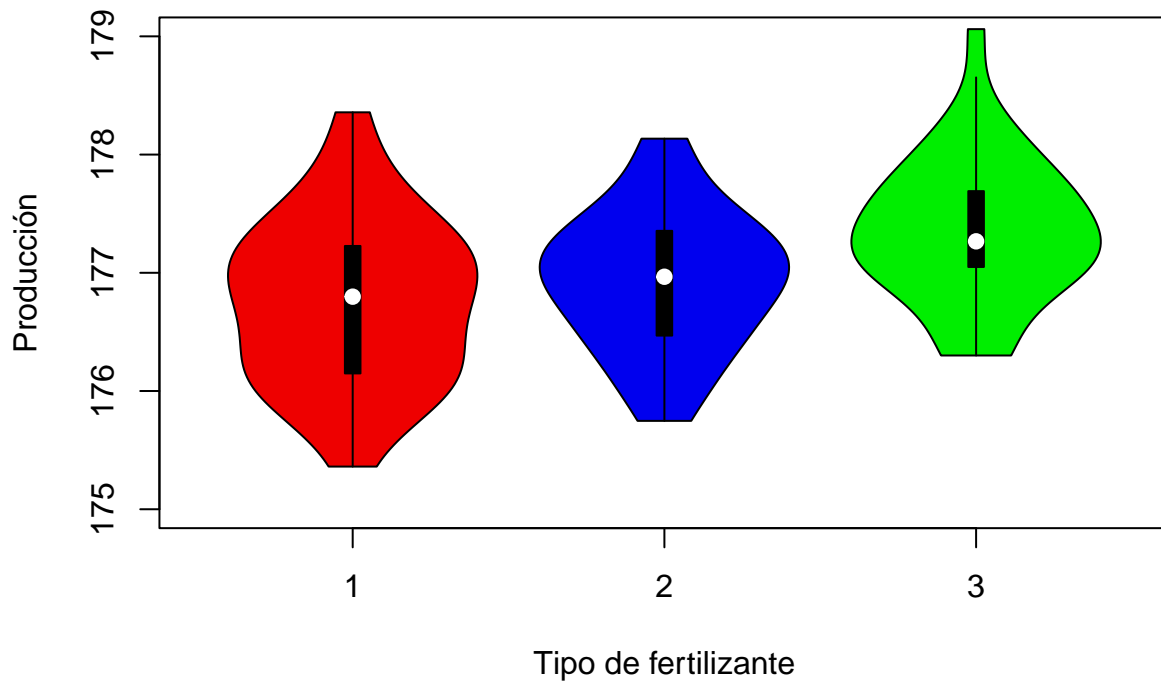
```
## The following objects are masked from 'package:base':
```

```
##
```

```
## as.Date, as.Date.numeric
```

```
vioplot(datos$yield ~ datos$fertilizer,  
  main = "Producción por fertilizante",  
  col= c("red2","blue2", "green2"),  
  ylab = "Producción",  
  xlab = "Tipo de fertilizante",  
  ylim = c(175, 179))
```

Producción por fertilizante



```
tapply(datos$yield, datos$fertilizer, mean)
```

```
##          1          2          3  
## 176.7570 176.9332 177.3562
```

```
tapply(datos$yield, datos$fertilizer, sd)
```

```
##          1          2          3  
## 0.6849233 0.5740668 0.5991214
```

```
tapply(datos$yield, datos$fertilizer, var)
```

```
##          1          2          3  
## 0.4691199 0.3295526 0.3589464
```

```
# Prueba de normalidad
```

```
shapiro.test(datos$yield)
```

```
##  
## Shapiro-Wilk normality test  
##  
## data:  datos$yield  
## W = 0.989, p-value = 0.6135
```

```
# Prueba de normalidad
```

```
shapiro.test(subset(datos$yield, datos$fertilizer == "1"))
```

```
##
## Shapiro-Wilk normality test
##
## data: subset(datos$yield, datos$fertilizer == "1")
## W = 0.97914, p-value = 0.7743
shapiro.test(subset(datos$yield, datos$fertilizer == "2"))

##
## Shapiro-Wilk normality test
##
## data: subset(datos$yield, datos$fertilizer == "2")
## W = 0.98329, p-value = 0.8875
shapiro.test(subset(datos$yield, datos$fertilizer == "3"))

##
## Shapiro-Wilk normality test
##
## data: subset(datos$yield, datos$fertilizer == "3")
## W = 0.95878, p-value = 0.2542
# No hay diferencias significativas, los datos son normales

# Prueba homogeneidad de varianzas

# VAR. TEST ES PARA DOS FACTORES, DOS GRUPOS CON VARIANZAS
# var.test(datos$yield ~ datos$fertilizer)

bartlett.test(datos$yield ~ datos$fertilizer)

##
## Bartlett test of homogeneity of variances
##
## data: datos$yield by datos$fertilizer
## Bartlett's K-squared = 1.0622, df = 2, p-value = 0.5879
# No hay diferencias entre las varianzas

datos.aov <- aov(datos$yield ~ datos$fertilizer)
summary(datos.aov)

##              Df Sum Sq Mean Sq F value Pr(>F)
## datos$fertilizer  2   6.07   3.0340    7.863  7e-04 ***
## Residuals       93  35.89   0.3859
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# + para comparar dos variables
# * interacción

# LSD DETERMINAR EL VALOR Diferencia Mínima Significativa
qt(0.975, 93)

## [1] 1.985802
```

```

sqrt((2*0.3859)/32) * qt(0.975, 93)

## [1] 0.3083992
tapply(datos$yield, datos$fertilizer, mean)

##          1          2          3
## 176.7570 176.9332 177.3562
# F1 vs F2

176.9332 - 176.7570 # NO SUPERA EL VALOR DE LSD 0.3083992

## [1] 0.1762
# EL valor entre las medias, debe ser mayor que el LSD para que sean diferentes

# F2 vs F3

176.9332 - 177.3562 # SUPERA EL VALOR DE LSD 0.3083992

## [1] -0.423
# F1 vs f3

176.7570 - 177.3562 # SUPERA EL VALOR DE LSD 0.3083992

## [1] -0.5992
# LSD ES MAS ROBUSTA PARA LA COMPARACIÓN DE LAS DIFERENCIAS

sqrt((2*0.3859)/32) * qtkey(0.95, nmeans = 3, df = 93)

## [1] 0.5231185
0.3083992 # 300 kilos de dif con LSD mas sensible

## [1] 0.3083992
0.5231185 # 500 kilos de dif con Tukey menos sensible

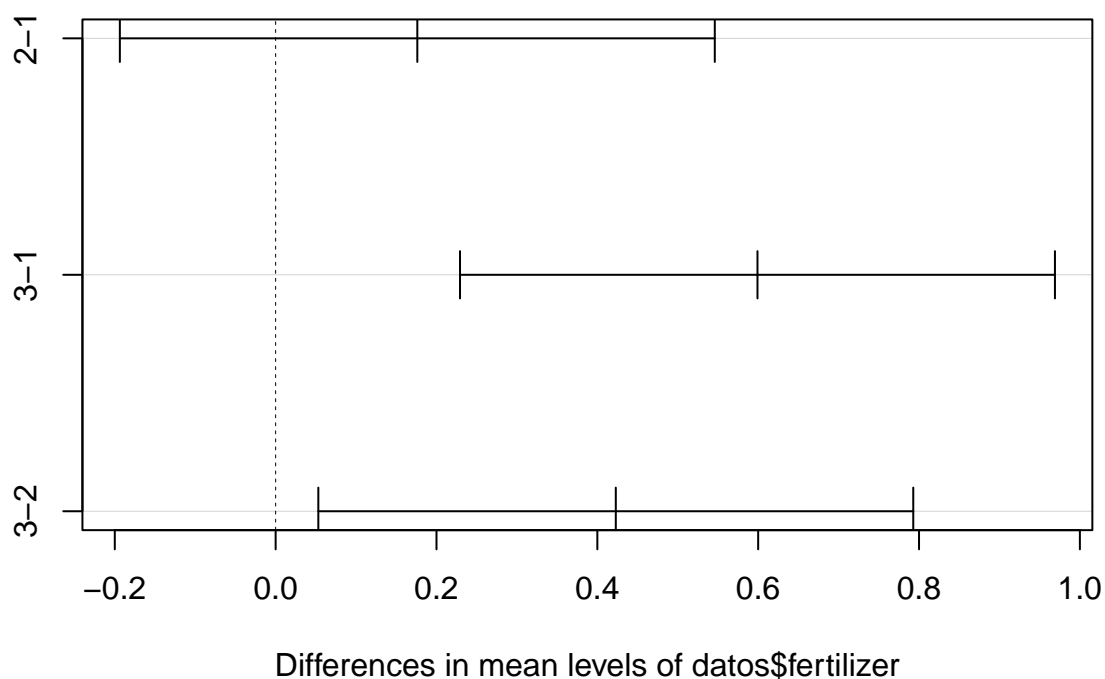
## [1] 0.5231185
TukeyHSD(datos.aov)

##      Tukey multiple comparisons of means
##      95% family-wise confidence level
##
## Fit: aov(formula = datos$yield ~ datos$fertilizer)
##
## $`datos$fertilizer`
##           diff           lwr           upr           p adj
## 2-1 0.1761687 -0.19371896 0.5460564 0.4954705
## 3-1 0.5991256 0.22923789 0.9690133 0.0006125
## 3-2 0.4229568 0.05306916 0.7928445 0.0208735

plot(TukeyHSD(datos.aov))

```

95% family-wise confidence level



```
library(ggplot2)

ggplot(datos, aes(x = fertilizer, y = yield, fill= fertilizer))+
  geom_violin()+
  geom_jitter(col = "indianred")+
  geom_boxplot(width=0.1, col = "white", alpha=0.5)+
  theme_light()+
  labs(x = "Fertilizante",
       y = "Producción")
```