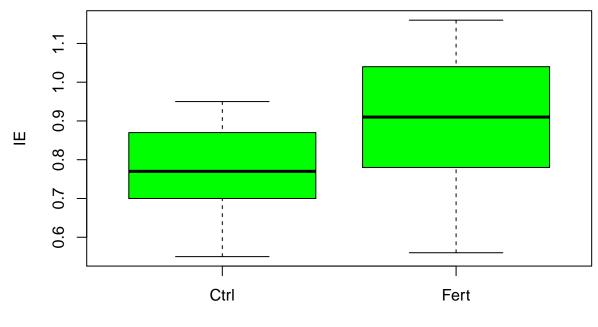
Script_Viveros.R

Usuario

2020-03-11

```
# Uriel Sarmiento Véliz -----
# Matricula: 1732196
# 11.03.2020
#Importar datos
setwd("C:/Tareas/108-Estadistica/Clases")
Vivero <- read.csv("viveros.csv", header =T)</pre>
summary(Vivero)
       planta
                        ΙE
                                  Tratamiento
## Min. : 1.00 Min.
                         :0.5500
                                  Ctrl:21
                 1st Qu.:0.7025
## 1st Qu.:11.25
                                  Fert:21
## Median :21.50 Median :0.7950
## Mean
         :21.50 Mean
                        :0.8371
## 3rd Qu.:31.75
                  3rd Qu.:0.9375
## Max.
         :42.00 Max.
                         :1.1600
boxplot(Vivero$IE ~ Vivero$Tratamiento, col="green",
       xlab = "Tratamientos", ylab = "IE")
```



Tratamientos

```
# Prueba de t -----
t.test(Vivero$IE ~ Vivero$Tratamiento) #aplicamos pureba de T
##
   Welch Two Sample t-test
##
##
## data: Vivero$IE by Vivero$Tratamiento
## t = -2.9813, df = 34.056, p-value = 0.00527
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.23382707 -0.04426816
## sample estimates:
## mean in group Ctrl mean in group Fert
                               0.9066667
##
            0.7676190
t.test(Vivero$IE ~ Vivero$Tratamiento, var.equal = T) #aplicamos pureba de T
##
##
   Two Sample t-test
##
## data: Vivero$IE by Vivero$Tratamiento
## t = -2.9813, df = 40, p-value = 0.004868
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.23331192 -0.04478332
## sample estimates:
```

mean in group Ctrl mean in group Fert ## 0.7676190 0.9066667