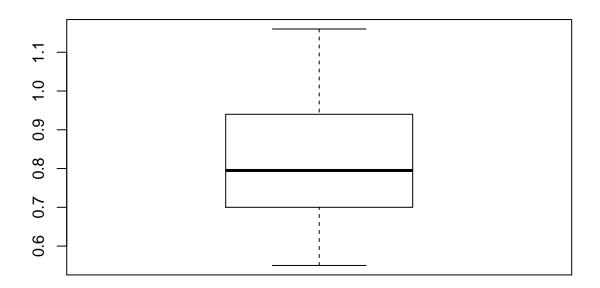
$clase_2.R$

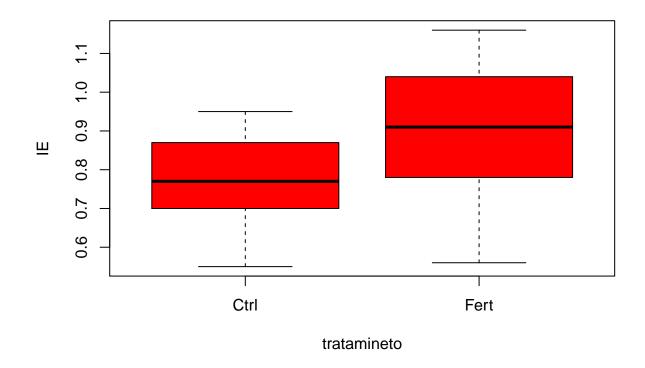
52618 2019-08-06

```
#cipriano guerrero cabrera
#06/08/2019
#clase 2
# importar datos viveros -----
vivero <- read.csv("C:/MCF 202-2019/MCF202/Datos/Clase2.csv", header = T)
summary(vivero)</pre>
```

```
##
       planta
                       ΙE
                                 Tratamiento
## Min. : 1.00 Min. :0.5500
                                Ctrl:21
## 1st Qu.:11.25 1st Qu.:0.7025
                                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
# prueba de t una muestra -----
par(mfrow=c(1,1))
boxplot(vivero$IE)
```



```
t.test(vivero$IE, mu = 0.85)
## One Sample t-test
##
## data: vivero$IE
## t = -0.5049, df = 41, p-value = 0.6163
## alternative hypothesis: true mean is not equal to 0.85
## 95 percent confidence interval:
## 0.7857153 0.8885704
## sample estimates:
## mean of x
## 0.8371429
#la media observada no es diferente estadisticamente ya que el valor de p
\#es mayor que el alfa establecido(0.05). ademas la media teoretica se
#encuentra dentro del rango del rango de los valores de intervalo sde confianza.
t.test(vivero$IE, mu = 0.90)
##
## One Sample t-test
##
## data: vivero$IE
## t = -2.4684, df = 41, p-value = 0.01783
## alternative hypothesis: true mean is not equal to 0.9
## 95 percent confidence interval:
## 0.7857153 0.8885704
## sample estimates:
## mean of x
## 0.8371429
#la media observada es diferenre estadisticamente ya que el valor de p (0.01)
#es menor que el valor de alfa establecido (0.05).por lo tanto se acepta H1.
# pruebas de t muestras independientes ------
boxplot(vivero$IE ~ vivero$Tratamiento, col="red", xlab="tratamineto",
       ylab="IE")
```



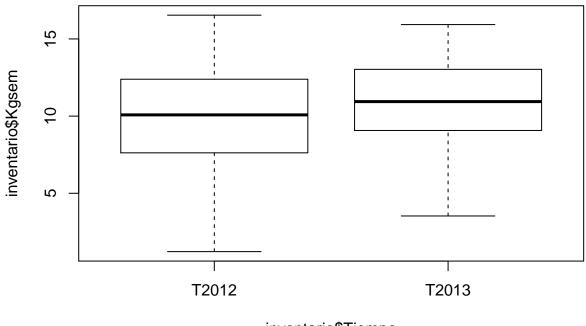
shapiro.test(vivero\$IE)

```
##
##
   Shapiro-Wilk normality test
##
## data: vivero$IE
## W = 0.96225, p-value = 0.1777
var.test(vivero$IE ~ vivero$Tratamiento)
## F test to compare two variances
## data: vivero$IE by vivero$Tratamiento
## F = 0.41068, num df = 20, denom df = 20, p-value = 0.05304
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.1666376 1.0121038
## sample estimates:
## ratio of variances
##
            0.4106757
#la varianza de ambos tratamientos son igual asi lo prueba el valor de p
#obtenido mediante una prueba de varianza (var.test).
t.test(vivero$IE ~ vivero$Tratamiento, var.ecual= T)
```

##

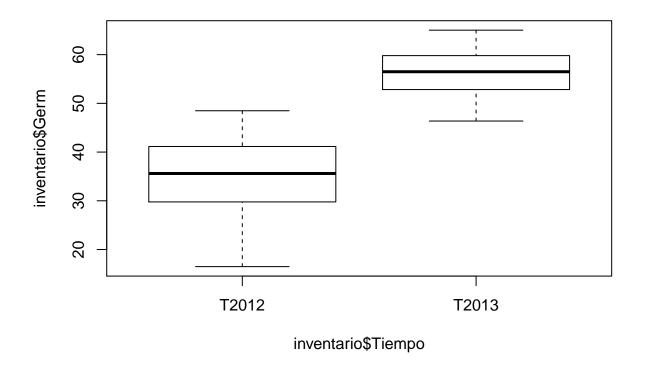
```
## Welch Two Sample t-test
##
## data: vivero$IE by vivero$Tratamiento
## t = -2.9813, df = 34.056, p-value = 0.00527
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.23382707 -0.04426816
## sample estimates:
## mean in group Ctrl mean in group Fert
           0.7676190
                             0.9066667
#existe una diferencia entre el IE de las plantulas fertilizadas
#el valor p comprueba nuestra hipotesis de que el fertilizante
#"power# mejora el IE
# inportar datos produccion -----
inventario <- read.csv("C:/MCF 202-2019/MCF202/Datos/produccion.csv", header = T)
summary(inventario)
##
     Tiempo
                                 BioRama
                                                  Germ
                  Kgsem
## T2012:50
                                    :44.54
              Min. : 1.220 Min.
                                             Min.
                                                    :16.49
## T2013:50
              1st Qu.: 8.492 1st Qu.:49.84 1st Qu.:35.61
##
              Median: 10.245 Median: 53.96 Median: 47.85
##
              Mean
                   :10.501 Mean :54.91 Mean
                                                    :45.83
              3rd Qu.:12.955 3rd Qu.:60.64
##
                                             3rd Qu.:56.30
##
              Max. :16.540 Max. :65.24 Max.
                                                    :65.02
##
         Н6
## Min. :-0.07
## 1st Qu.:14.16
## Median:16.56
## Mean
         :16.94
## 3rd Qu.:21.24
## Max.
          :29.71
```

boxplot(inventario\$Kgsem ~ inventario\$Tiempo)



inventario\$Tiempo

```
t.test(inventario$Kgsem ~ inventario$Tiempo, paired = T)
##
##
   Paired t-test
##
## data: inventario$Kgsem by inventario$Tiempo
## t = -1.2538, df = 49, p-value = 0.2159
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
   -2.0530953 0.4754953
## sample estimates:
## mean of the differences
##
                   -0.7888
tapply(inventario$Germ, inventario$Tiempo, mean)
     T2012
             T2013
## 35.5036 56.1628
boxplot(inventario$Germ ~ inventario$Tiempo)
```



t.test(inventario\$Germ ~ inventario\$Tiempo, paired = T)

```
##
## Paired t-test
##
## data: inventario$Germ by inventario$Tiempo
## t = -16.678, df = 49, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -23.14844 -18.16996
## sample estimates:
## mean of the differences
## -20.6592</pre>
```