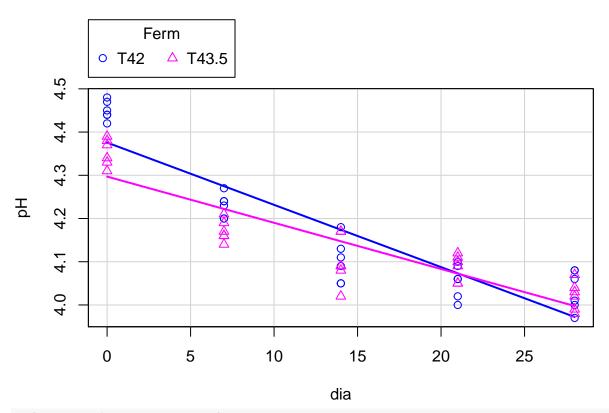
T1-E01-Iogurt.R

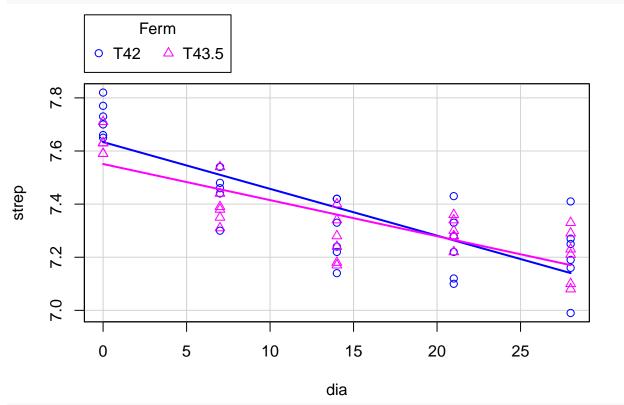
jordi

Thu Oct 11 17:40:43 2018

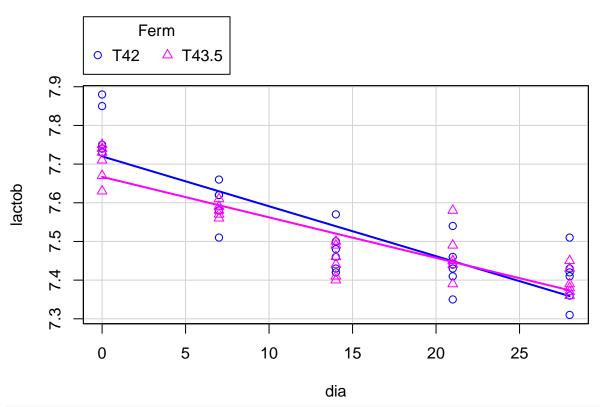
```
setwd("~/Documents/CURS 2018-2019/PIE2")
dd<-read.csv2("./Dades/Iogurt.csv")</pre>
head(dd)
    Ferm dia pH strep lactob
## 1 T42 21 4.10 7.43
                        7.46
## 2 T42 0 4.44 7.65 7.75
## 3 T42 21 4.02 7.10 7.35
## 4 T42 7 4.24 7.54 7.62
## 5 T42
         7 4.27 7.54 7.66
## 6 T42 28 4.01 7.25 7.41
library(car)
## Loading required package: carData
library(tables)
## Loading required package: Hmisc
## Loading required package: lattice
## Loading required package: survival
## Warning: package 'survival' was built under R version 3.4.4
## Loading required package: Formula
## Loading required package: ggplot2
##
## Attaching package: 'Hmisc'
## The following objects are masked from 'package:base':
##
      format.pval, units
#_____
# (a) Descriptiva
# Gràfiques
sp(pH~dia|Ferm,dd,smooth=F)
```



sp(strep~dia|Ferm,smooth=F,dd)



sp(lactob~dia|Ferm,smooth=F,dd)



Taules
dd\$Fdia<-as.factor(dd\$dia)
tabular((pH+strep+lactob)*Ferm*((n=1)+mean+sd)~Fdia,dd)</pre>

					Fdia		
	Ferm		0	7	14	21	28
pН	T42	n	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	4.45000	4.23000	4.10167	4.05500	4.03333
		sd	0.02191	0.02683	0.04997	0.03886	0.04633
	T43.5	n	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	4.35333	4.17167	4.10333	4.08667	4.02167
		sd	0.03141	0.02483	0.05785	0.03011	0.03312
strep	T42	\mathbf{n}	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	7.72167	7.46000	7.29500	7.24667	7.21167
		sd	0.06555	0.08854	0.11415	0.12644	0.13891
	T43.5	\mathbf{n}	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	7.63000	7.40167	7.26833	7.29667	7.20667
		sd	0.04382	0.08035	0.09042	0.04967	0.10013
lactob	T42	\mathbf{n}	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	7.78000	7.59500	7.47667	7.43833	7.40667
		sd	0.06693	0.05128	0.05465	0.06242	0.06772
	T43.5	\mathbf{n}	6.00000	6.00000	6.00000	6.00000	6.00000
		mean	7.70500	7.58333	7.45000	7.46667	7.39667
		sd	0.04637	0.01751	0.04099	0.06408	0.03559

o bé per separat tabular(pH*Ferm*((n=1)+mean+sd)~Fdia,dd) ...

#----

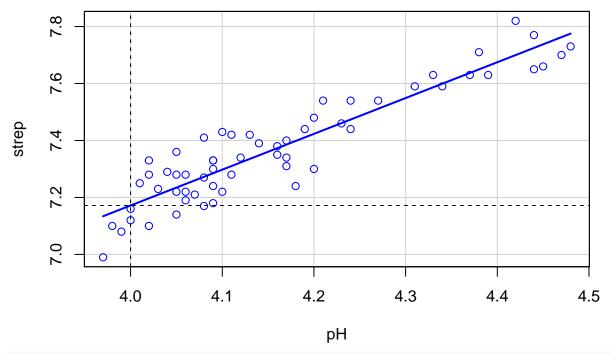
(b) Comparacions de 2

```
# pH dia 0
t.test(pH~Ferm,dd[dd$dia==0,])
##
##
   Welch Two Sample t-test
##
## data: pH by Ferm
## t = 6.1828, df = 8.9338, p-value = 0.0001673
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.06125851 0.13207482
## sample estimates:
##
    mean in group T42 mean in group T43.5
##
              4.450000
                                  4.353333
t.test(pH~Ferm,var.equal=T,dd[dd$dia==0,])
##
##
   Two Sample t-test
##
## data: pH by Ferm
## t = 6.1828, df = 10, p-value = 0.0001038
\#\# alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## 0.06183034 0.13150299
## sample estimates:
    mean in group T42 mean in group T43.5
##
              4.450000
                                  4.353333
var.test(pH~Ferm,dd[dd$dia==0,])
##
## F test to compare two variances
##
## data: pH by Ferm
## F = 0.48649, num df = 5, denom df = 5, p-value = 0.4479
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.06807452 3.47661819
## sample estimates:
## ratio of variances
            0.4864865
# strep dia 21
t.test(strep~Ferm,dd[dd$dia==21,])
##
##
   Welch Two Sample t-test
##
## data: strep by Ferm
## t = -0.90159, df = 6.5071, p-value = 0.3994
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.18317752 0.08317752
## sample estimates:
```

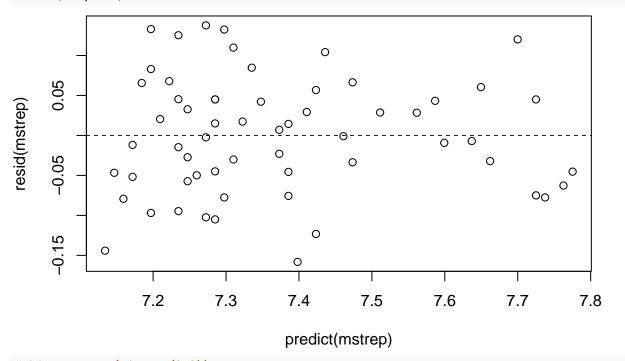
```
mean in group T42 mean in group T43.5
##
             7.246667
                                7.296667
t.test(strep~Ferm, var.equal=T,dd[dd$dia==21,])
##
##
   Two Sample t-test
##
## data: strep by Ferm
## t = -0.90159, df = 10, p-value = 0.3885
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.17356744 0.07356744
## sample estimates:
    mean in group T42 mean in group T43.5
##
##
             7.246667
                                7.296667
var.test(strep~Ferm,dd[dd$dia==21,])
## F test to compare two variances
##
## data: strep by Ferm
## F = 6.4811, num df = 5, denom df = 5, p-value = 0.06108
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
## 0.9069038 46.3162801
## sample estimates:
## ratio of variances
##
            6.481081
# (c) Predicció a partir del pH
sp(strep~pH|Ferm,dd,smooth=F,boxplot=F)
```

```
Ferm
            o T42
                      △ T43.5
                                                                           0
                                                                              0
                                                                             000
     7.6
                                                  0
                            0 00 0
     7.4
                                             Φ
                                          0
     7.2
     7.0
                                            4.2
                4.0
                                                         4.3
                              4.1
                                                                       4.4
                                                                                     4.5
                                               рΗ
sp(strep~pH,dd,smooth=F,boxplot=F)
summary(mstrep<-lm(strep~pH,dd))</pre>
##
## Call:
## lm(formula = strep ~ pH, data = dd)
##
## Residuals:
##
                  1Q
                       Median
                                     30
```

```
## -0.15814 -0.05035 -0.00171 0.04508 0.13758
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                          0.28205
                                    7.599 2.89e-10 ***
## (Intercept) 2.14327
                1.25715
                          0.06775 18.556 < 2e-16 ***
## pH
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.07359 on 58 degrees of freedom
## Multiple R-squared: 0.8558, Adjusted R-squared: 0.8533
## F-statistic: 344.3 on 1 and 58 DF, p-value: < 2.2e-16
(lstrep<-mstrep$coef[1]+mstrep$coef[2]*4)
## (Intercept)
     7.171852
sp(strep~pH,dd,smooth=F,boxplot=F)
abline(v=4,h=1strep,1t=2)
```

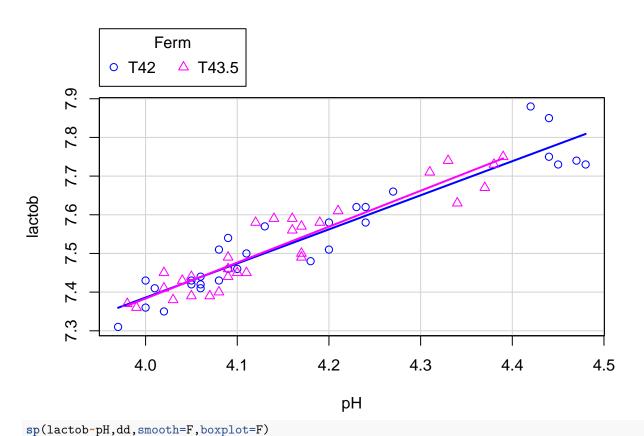


plot(predict(mstrep),resid(mstrep))
abline(h=0,lt=2)

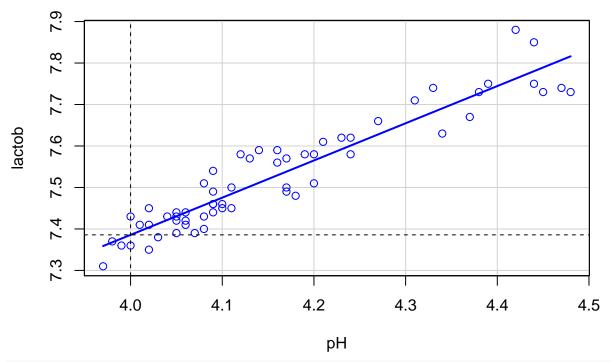


```
#oldpar <- par(mfrow=c(2,2))
#plot(mstrep,ask=F)
#par(oldpar)

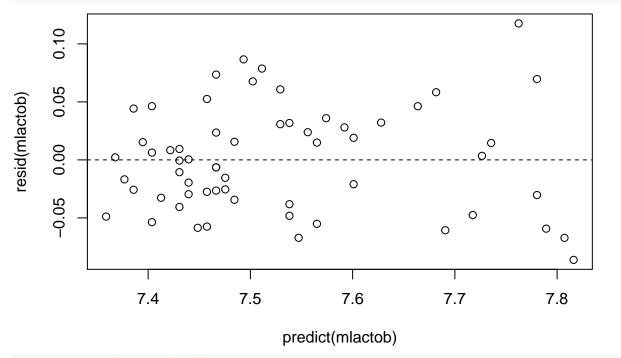
#lactob
sp(lactob~pH|Ferm,dd,smooth=F,boxplot=F)</pre>
```



```
summary(mlactob<-lm(lactob~pH,dd))</pre>
##
## Call:
## lm(formula = lactob ~ pH, data = dd)
## Residuals:
##
                    10
                          Median
                                        30
## -0.086181 -0.033098 -0.000082 0.031023 0.117622
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.79895
                           0.17497
                                     21.71
                                             <2e-16 ***
                0.89670
                           0.04203
                                     21.34
                                             <2e-16 ***
## pH
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.04565 on 58 degrees of freedom
## Multiple R-squared: 0.887, Adjusted R-squared: 0.885
## F-statistic: 455.2 on 1 and 58 DF, p-value: < 2.2e-16
(llactob<-mlactob$coef[1]+mlactob$coef[2]*4)</pre>
## (Intercept)
      7.385763
sp(lactob~pH,dd,smooth=F,boxplot=F)
abline(v=4,h=llactob,lt=2)
```



plot(predict(mlactob),resid(mlactob))
abline(h=0,lt=2)



```
#oldpar <- par(mfrow=c(2,2))
#plot(mlactob,ask=F)
#par(oldpar)

# R2 i lv
summary(mstrep)</pre>
```

##

```
## Call:
## lm(formula = strep ~ pH, data = dd)
##
## Residuals:
                 1Q
                     Median
                                   3Q
## -0.15814 -0.05035 -0.00171 0.04508 0.13758
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                          0.28205
                                   7.599 2.89e-10 ***
## (Intercept) 2.14327
## pH
               1.25715
                          0.06775 18.556 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.07359 on 58 degrees of freedom
## Multiple R-squared: 0.8558, Adjusted R-squared: 0.8533
## F-statistic: 344.3 on 1 and 58 DF, p-value: < 2.2e-16
summary(mlactob)
##
## Call:
## lm(formula = lactob ~ pH, data = dd)
##
## Residuals:
##
        Min
                   1Q
                         Median
                                       3Q
## -0.086181 -0.033098 -0.000082 0.031023 0.117622
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.79895
                          0.17497
                                    21.71
                                            <2e-16 ***
## pH
               0.89670
                          0.04203
                                    21.34
                                            <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.04565 on 58 degrees of freedom
## Multiple R-squared: 0.887, Adjusted R-squared: 0.885
## F-statistic: 455.2 on 1 and 58 DF, p-value: < 2.2e-16
c(strep=summary(mstrep)$r.squared,lactob=summary(mlactob)$r.squared)
##
      strep
               lactob
## 0.8558319 0.8869845
c(strep=logLik(mstrep),lactob=logLik(mlactob))
##
      strep
               lactob
## 72.43656 101.08524
```