# Análise Doutorado Débora

## Biorigin

### Carregando pacotes

library(tidyverse)  
library(ExpDes.pt)  
library(lme4)  
library(agricolae)  
library(readxl)  
library(janitor)  
library(MASS)  
library(car)

### Entrada de dados

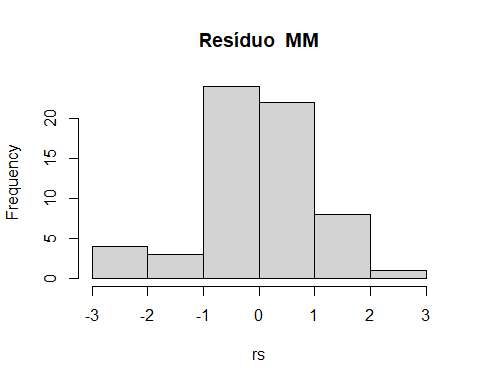
## Lendo os dados  
dados <- read\_xlsx("data/dados\_01.xlsx",  
 sheet = "Análises Fezes",  
 na=".") %>%   
 clean\_names() %>%   
 mutate(  
 dose = case\_when(  
 tr == 1 ~ 0,  
 tr == 2 | tr == 5 ~ 0.2,  
 tr == 3 | tr == 6 ~ 0.4,  
 tr == 4 | tr == 7 ~ 0.8  
 ),  
 ingrediente = ifelse(tr==1,"PCL",  
 ifelse(tr > 1 & tr<=4 ,"PCL","PCLs")),  
   
 # transformação dos dados  
 # mo = log(mo),  
 # fibra\_in = log(fibra\_in),  
 # ee\_in = log(ee\_in),  
 # amido\_in = log(amido\_in),  
 # ph=ph^(-2),  
 # am=am^2  
 )  
  
#extraindo o controle  
da <- dados %>% filter(dose==0) %>%   
 mutate(  
 ingrediente="PCLs"  
 )  
  
#duplicando o controle  
dados <- rbind(dados,da)

### Vislumbre

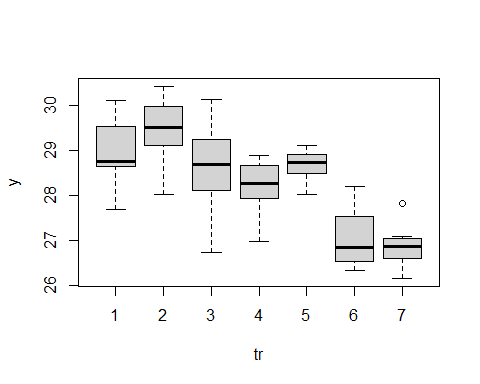
# glimpse(dados)

## Normalidade de Homocedasticidade

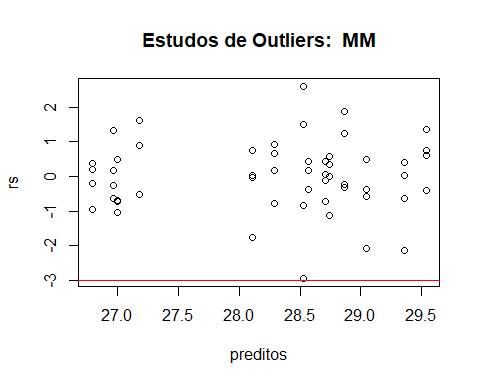
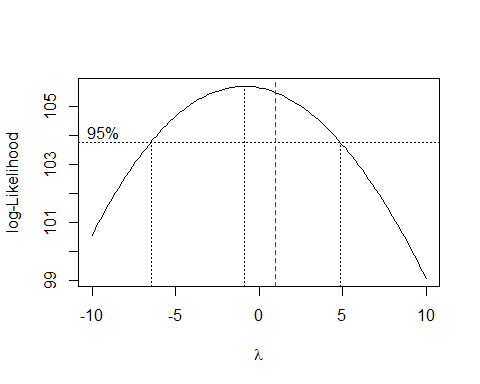
variaveis <- dados %>%   
 dplyr::select(mm:tripta\_63,-score)  
  
for(i in seq\_along(variaveis)){  
 nome <- names(variaveis[i])  
 print("====================================")  
 print(str\_to\_upper(nome))  
 print("====================================")  
  
 daux <- variaveis[i]  
 names(daux) <- "y" ### <-  
   
 y <- daux$y ### <-  
 daux$bl <- dados$bl %>% as\_factor()  
 daux$tr <- dados$tr %>% as\_factor()  
   
 daux <- daux %>% drop\_na()  
 y <- daux$y ### <-  
 bl <- daux$bl %>% as\_factor()  
 tr <- daux$tr %>% as\_factor()  
   
 mod <- aov(y ~ tr + bl)  
 rs <- mod %>% rstudent()  
 hist(rs, main = paste("Resíduo ", str\_to\_upper(nome)) )  
  
 print(shapiro.test(rs))  
  
 plot(y ~ tr)  
 cat("Médias por tratamentos\n")  
 medias <- data.frame(y, tr) %>%   
 group\_by(tr) %>%   
 summarise(media = mean(y,na.rm=TRUE))  
 print(medias)  
  
 filtro <- y != 0  
 mod\_box <- aov(y[filtro] ~ tr[filtro] + bl[filtro])  
 boxcox(mod\_box,seq(-10,10,.5))  
 abline(v=1,col="red",lty=2)  
   
 preditos <- predict(mod)  
 plot(preditos,rs,  
 main=paste("Estudos de Outliers: ", str\_to\_upper(nome)))  
   
 dfa <- data.frame(Trat = tr, Bloco = bl, Y = y, Pred = preditos, RS =rs) %>%   
 filter(RS >= 3 | RS <= -3) %>%   
 dplyr::select(Trat, Bloco, Y)  
   
 cat("Outliers")  
 print(dfa)  
 abline(h=c(3,-3),col="red")  
 cat("\n")  
}  
#> [1] "===================================="  
#> [1] "MM"  
#> [1] "===================================="



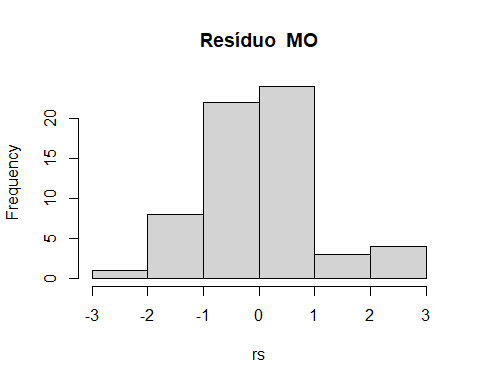
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98071, p-value = 0.4366



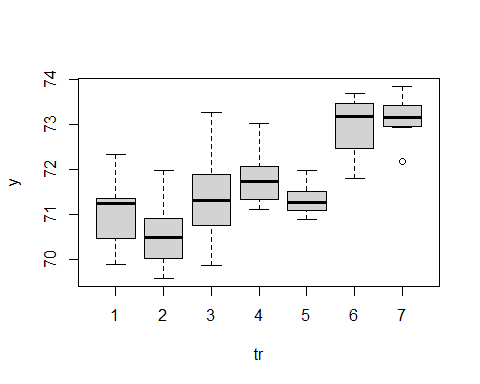
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 29.0  
#> 2 2 29.5  
#> 3 3 28.6  
#> 4 4 28.2  
#> 5 5 28.7  
#> 6 6 27.1  
#> 7 7 26.9



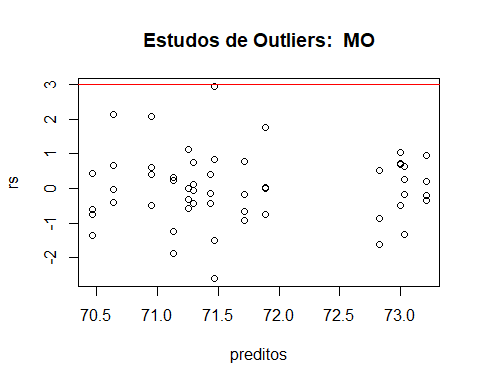
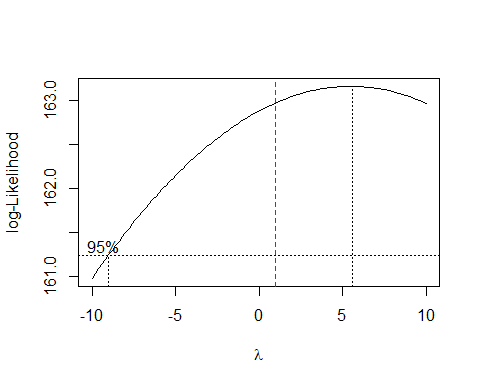
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "MO"  
#> [1] "===================================="



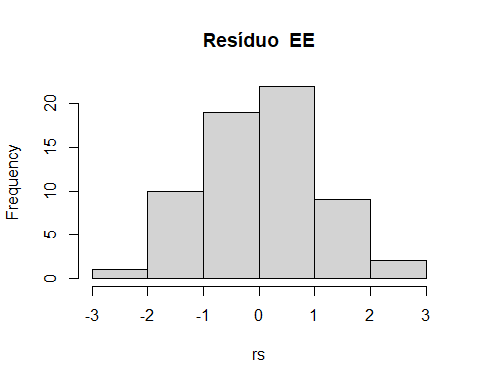
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98071, p-value = 0.4366



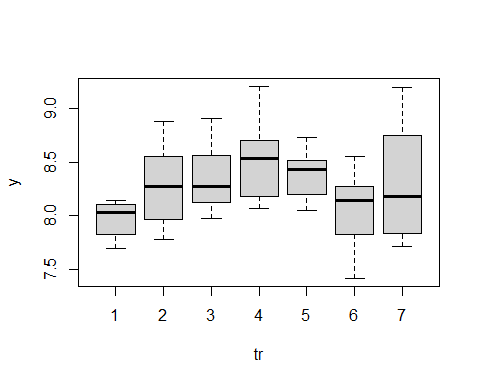
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 71.0  
#> 2 2 70.5  
#> 3 3 71.4  
#> 4 4 71.8  
#> 5 5 71.3  
#> 6 6 72.9  
#> 7 7 73.1



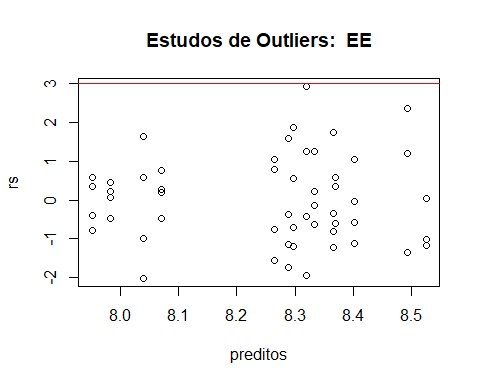
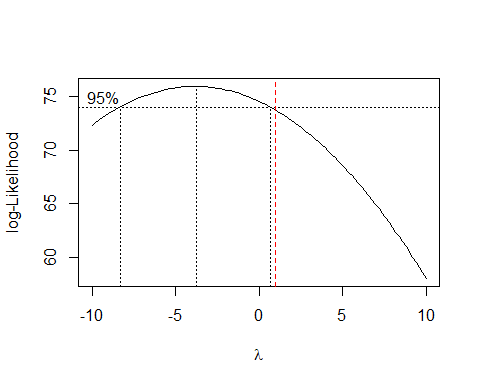
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "EE"  
#> [1] "===================================="



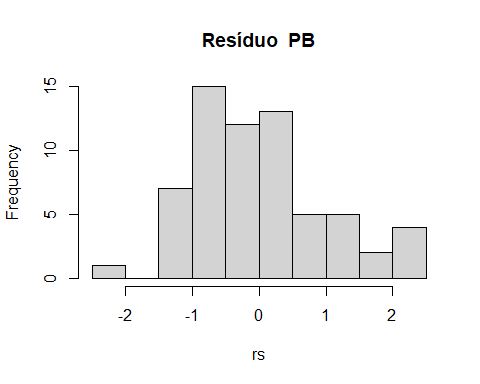
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98394, p-value = 0.5829



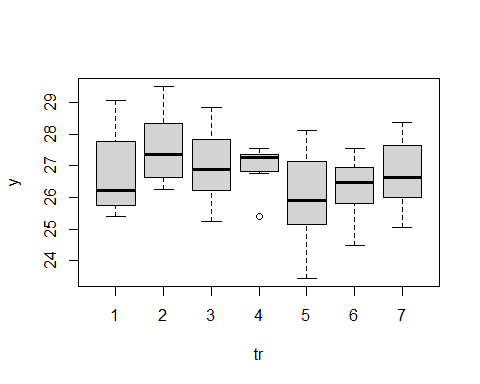
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 7.97  
#> 2 2 8.28  
#> 3 3 8.35  
#> 4 4 8.51  
#> 5 5 8.39  
#> 6 6 8.06  
#> 7 7 8.30



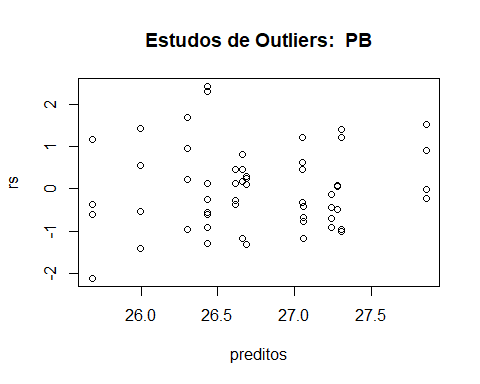
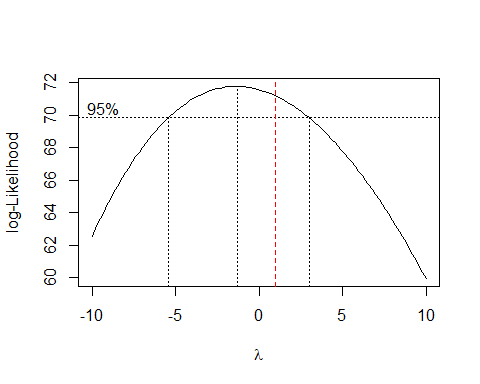
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PB"  
#> [1] "===================================="



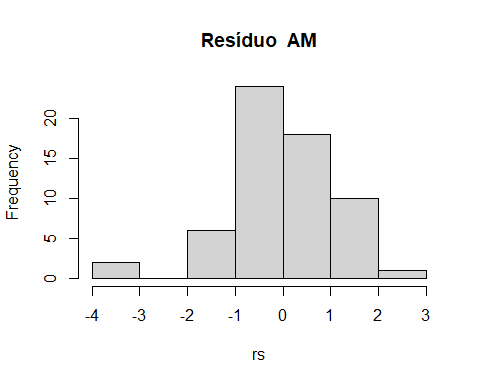
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95028, p-value = 0.01181



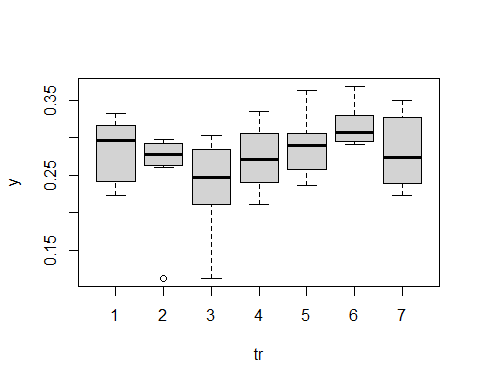
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 26.7  
#> 2 2 27.5  
#> 3 3 27.0  
#> 4 4 27.0  
#> 5 5 26.0  
#> 6 6 26.3  
#> 7 7 26.7



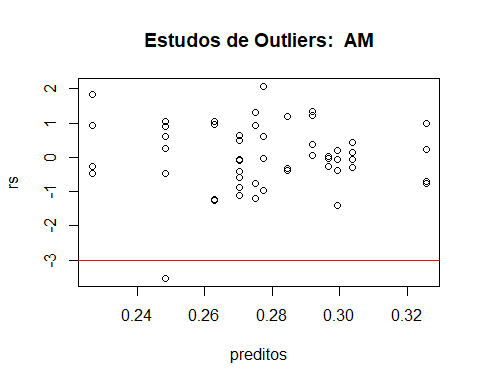
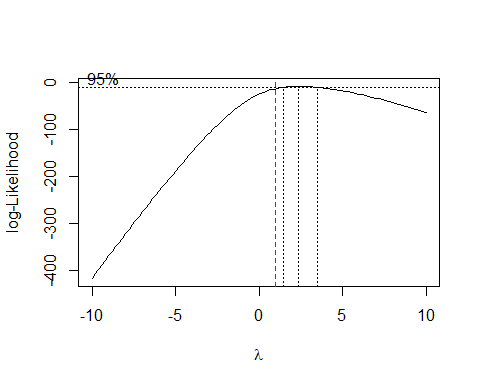
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AM"  
#> [1] "===================================="



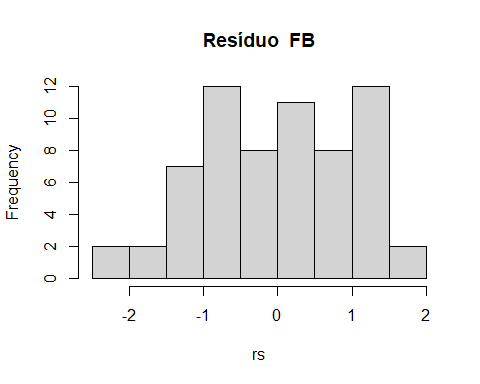
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.93004, p-value = 0.001804



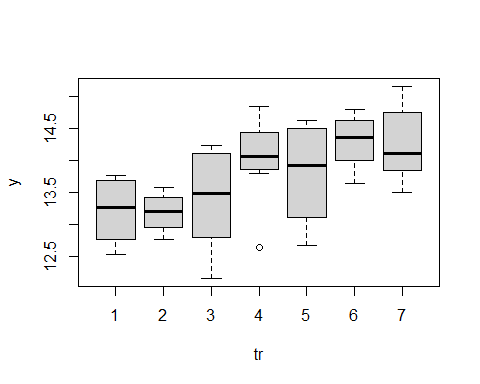
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 0.284  
#> 2 2 0.259  
#> 3 3 0.238  
#> 4 4 0.272  
#> 5 5 0.288  
#> 6 6 0.315  
#> 7 7 0.281



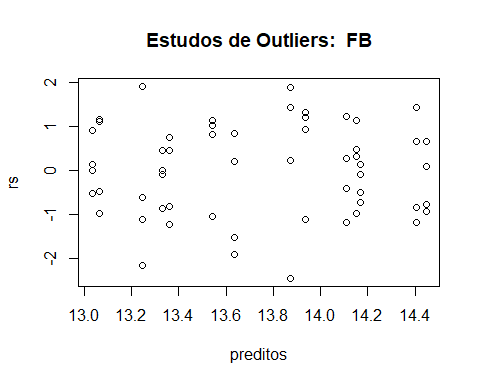
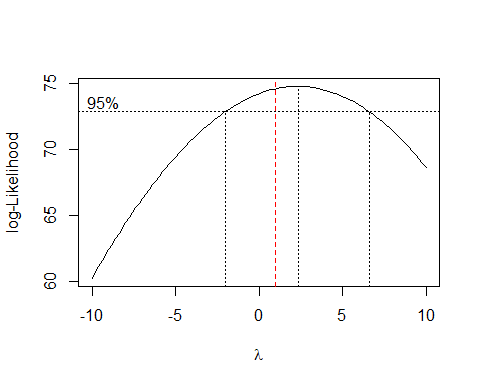
#> Outliers Trat Bloco Y  
#> 11 2 1 0.1123946  
#> 22 3 2 0.1123452  
#>   
#> [1] "===================================="  
#> [1] "FB"  
#> [1] "===================================="



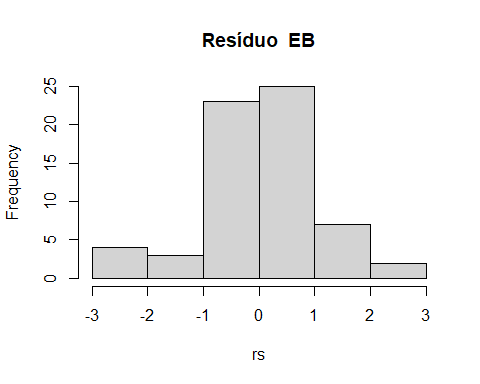
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96898, p-value = 0.107



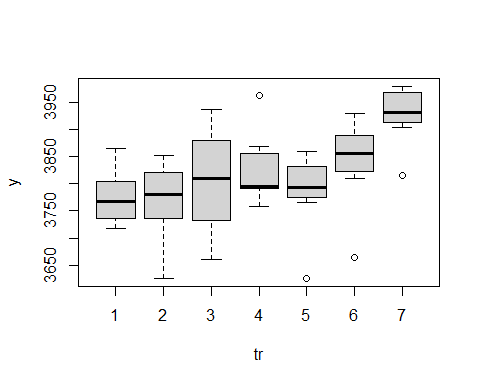
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 13.2  
#> 2 2 13.2  
#> 3 3 13.4  
#> 4 4 14.0  
#> 5 5 13.8  
#> 6 6 14.3  
#> 7 7 14.3



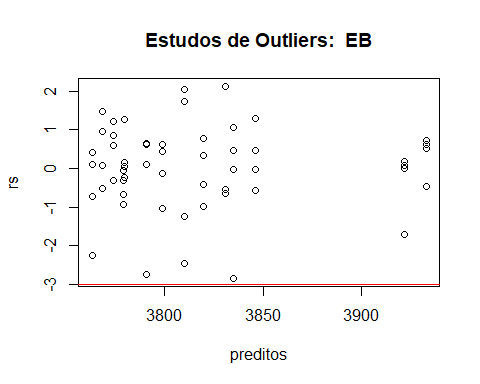
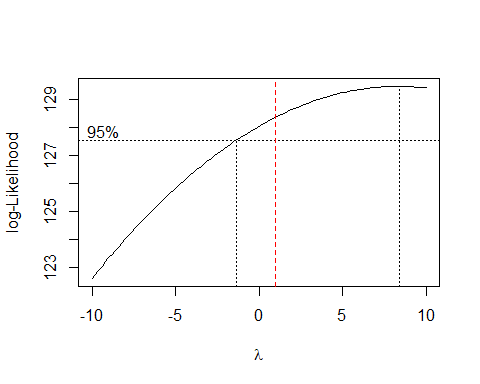
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "EB"  
#> [1] "===================================="



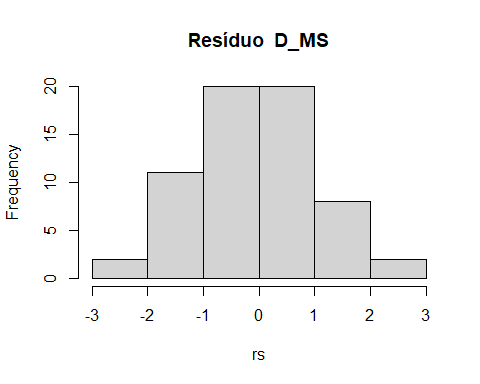
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96403, p-value = 0.05902



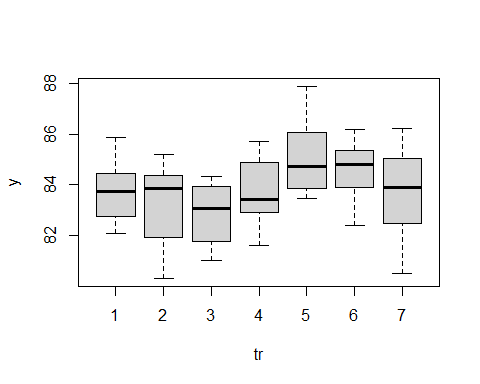
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 3774.  
#> 2 2 3769.  
#> 3 3 3805.  
#> 4 4 3825.  
#> 5 5 3785.  
#> 6 6 3841.  
#> 7 7 3927.



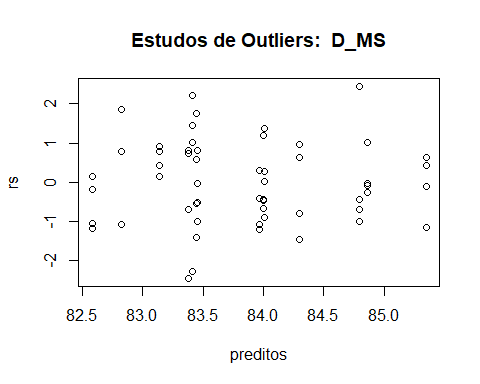
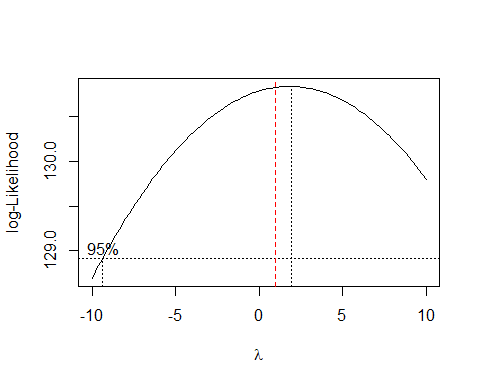
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_MS"  
#> [1] "===================================="



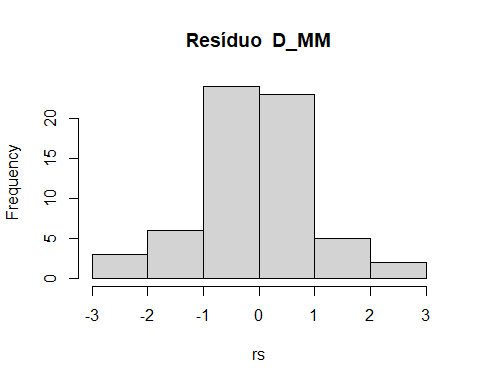
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98786, p-value = 0.7921



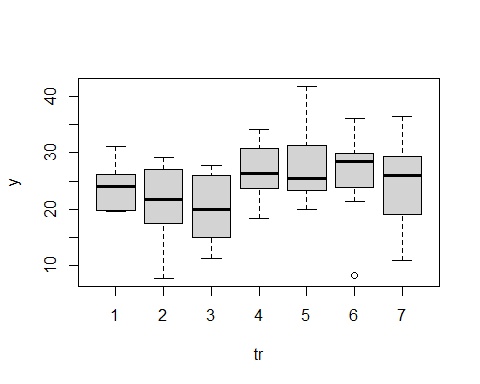
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 83.7  
#> 2 2 83.1  
#> 3 3 82.9  
#> 4 4 83.7  
#> 5 5 85.1  
#> 6 6 84.6  
#> 7 7 83.7



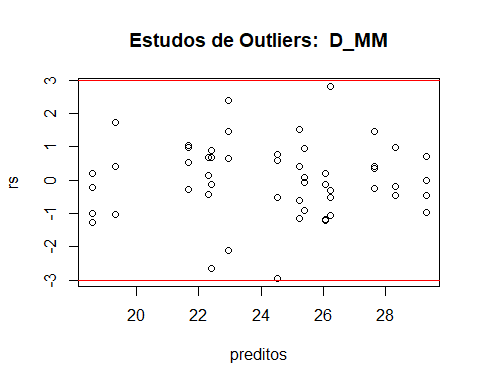
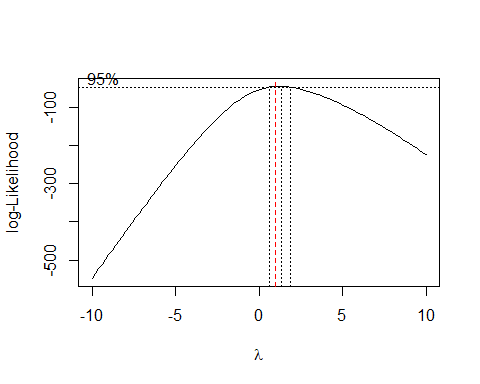
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_MM"  
#> [1] "===================================="



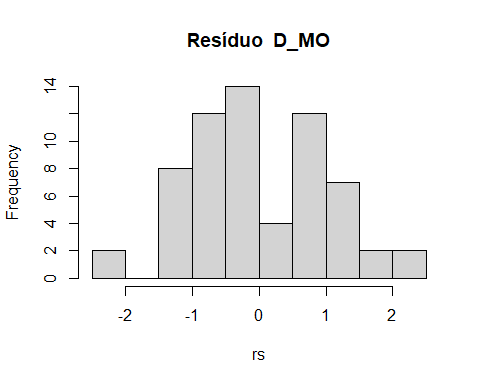
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97765, p-value = 0.3068



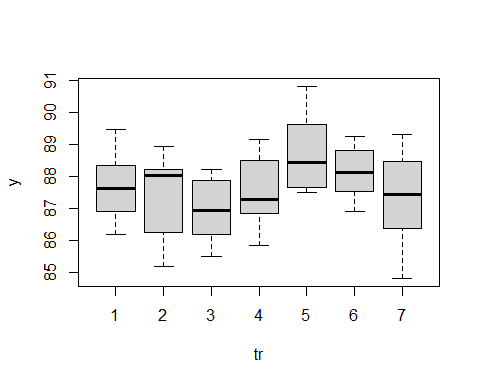
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 23.8  
#> 2 2 21.1  
#> 3 3 20.1  
#> 4 4 26.8  
#> 5 5 27.8  
#> 6 6 26.1  
#> 7 7 24.5



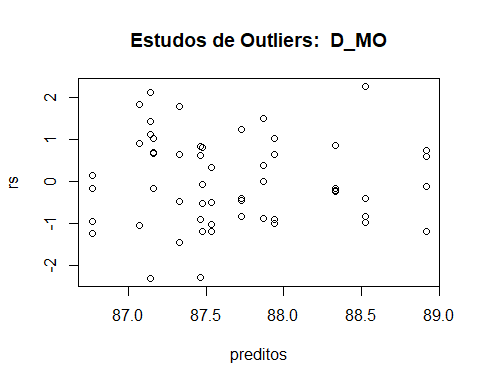
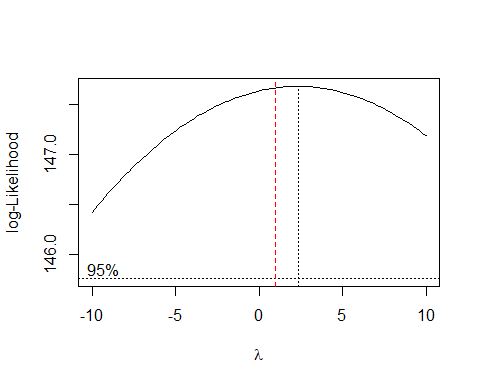
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_MO"  
#> [1] "===================================="



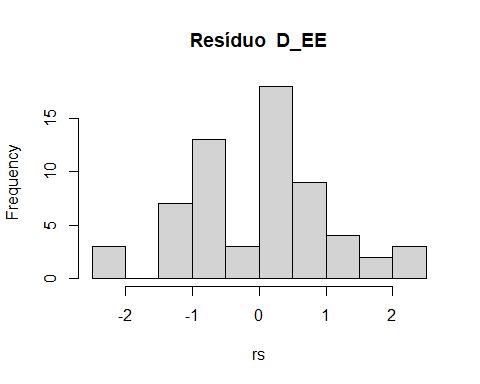
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.9779, p-value = 0.3154



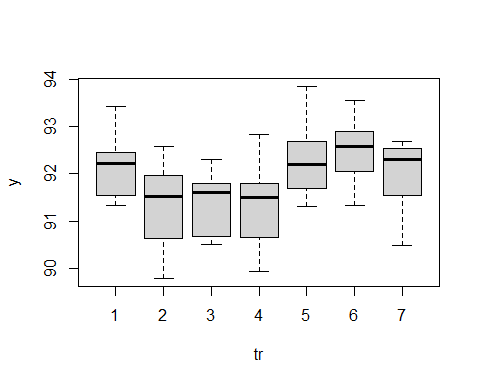
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 87.7  
#> 2 2 87.3  
#> 3 3 87.0  
#> 4 4 87.5  
#> 5 5 88.7  
#> 6 6 88.1  
#> 7 7 87.3



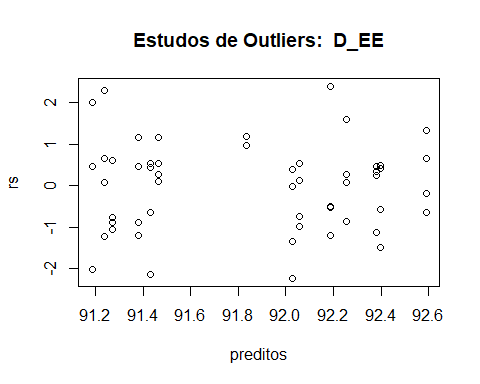
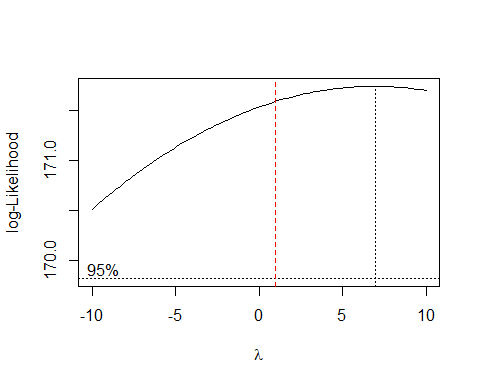
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_EE"  
#> [1] "===================================="



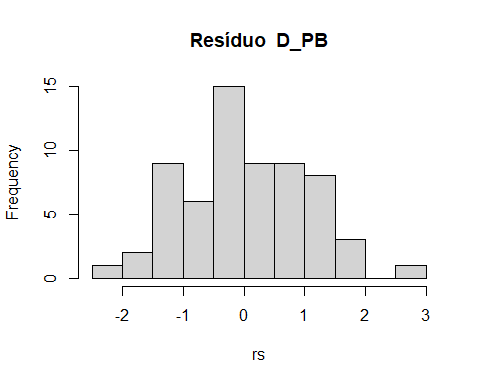
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97913, p-value = 0.371



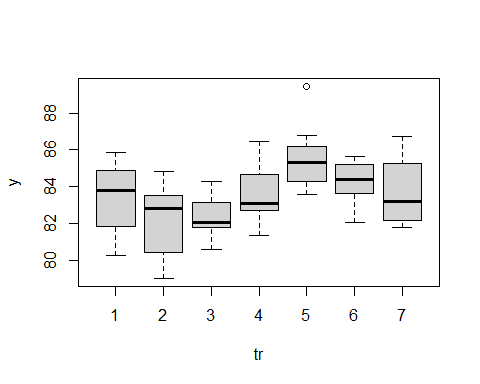
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 92.2  
#> 2 2 91.3  
#> 3 3 91.4  
#> 4 4 91.3  
#> 5 5 92.3  
#> 6 6 92.5  
#> 7 7 91.9



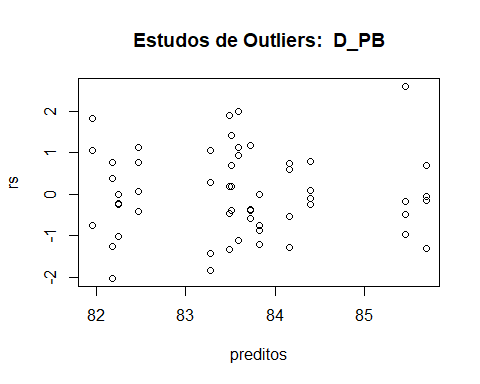
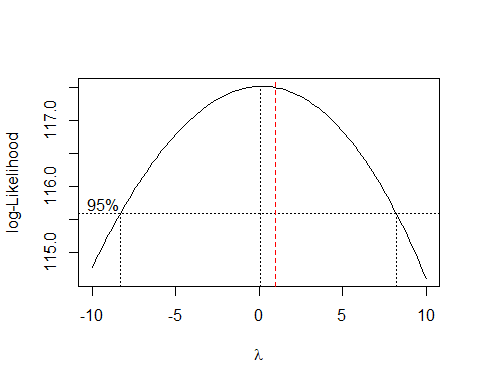
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_PB"  
#> [1] "===================================="



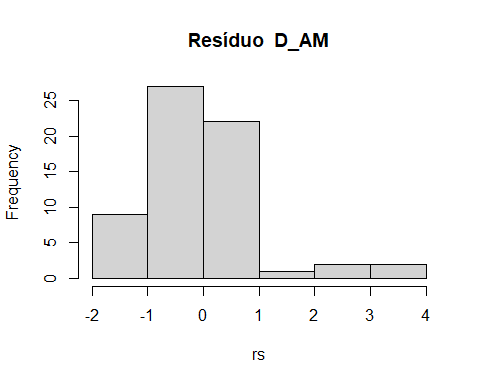
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98741, p-value = 0.7689



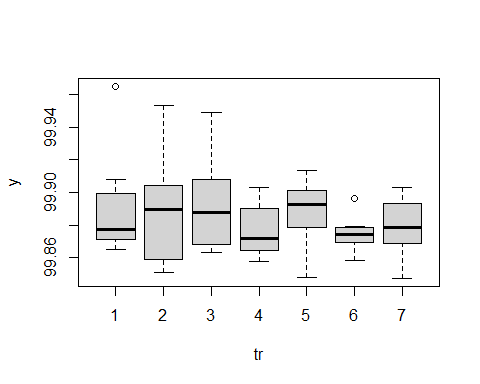
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 83.4  
#> 2 2 82.1  
#> 3 3 82.4  
#> 4 4 83.6  
#> 5 5 85.6  
#> 6 6 84.3  
#> 7 7 83.7



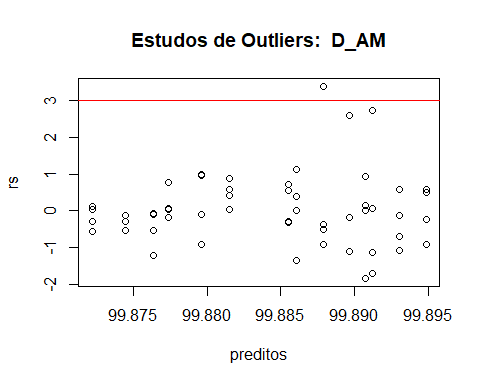
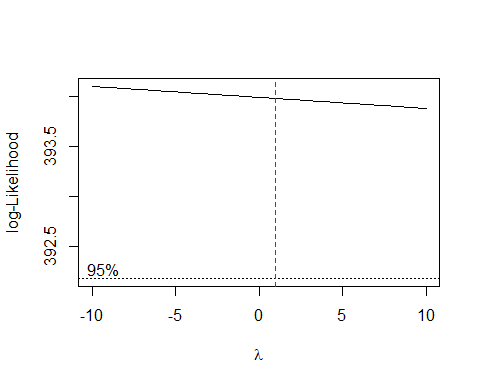
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_AM"  
#> [1] "===================================="



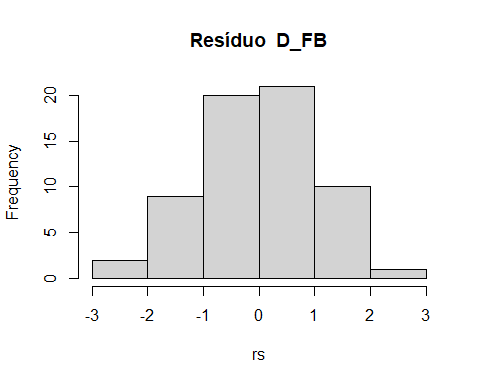
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.89009, p-value = 3.933e-05



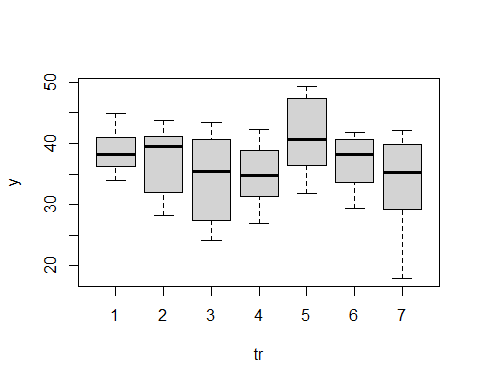
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 99.9  
#> 2 2 99.9  
#> 3 3 99.9  
#> 4 4 99.9  
#> 5 5 99.9  
#> 6 6 99.9  
#> 7 7 99.9



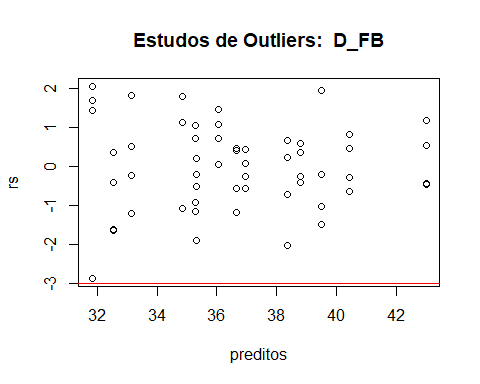
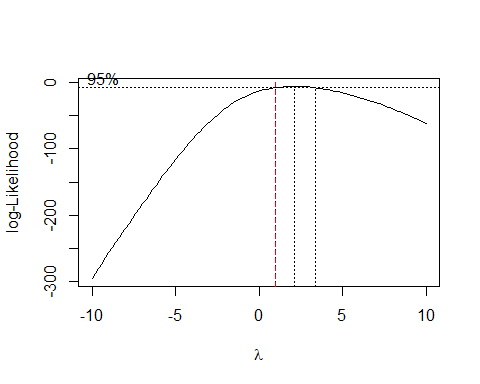
#> Outliers Trat Bloco Y  
#> 6 1 2 99.96496  
#> 61 1 2 99.96496  
#>   
#> [1] "===================================="  
#> [1] "D\_FB"  
#> [1] "===================================="



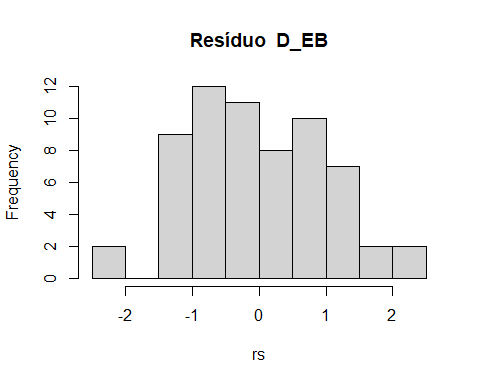
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98762, p-value = 0.78



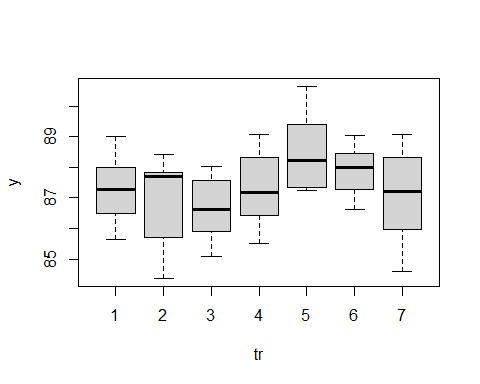
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 38.7  
#> 2 2 36.8  
#> 3 3 34.3  
#> 4 4 34.9  
#> 5 5 41.2  
#> 6 6 37.0  
#> 7 7 33.6



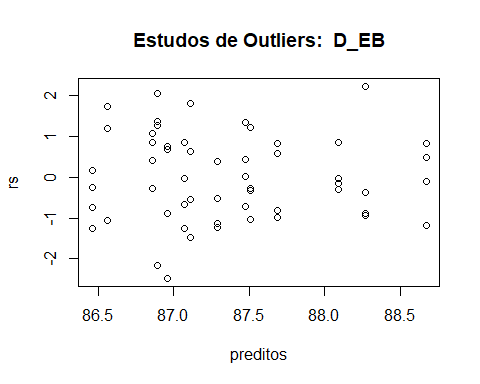
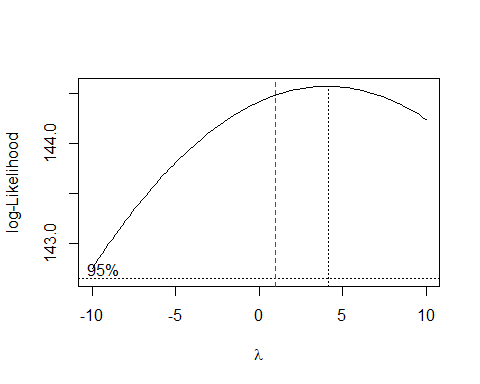
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "D\_EB"  
#> [1] "===================================="



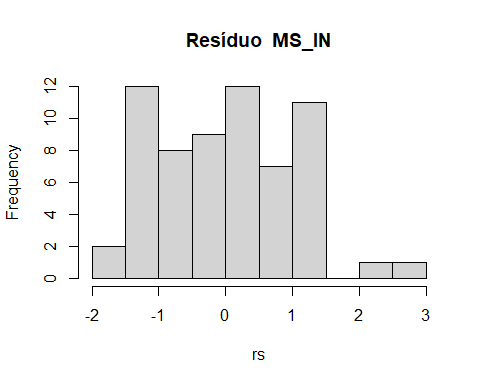
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98407, p-value = 0.5897



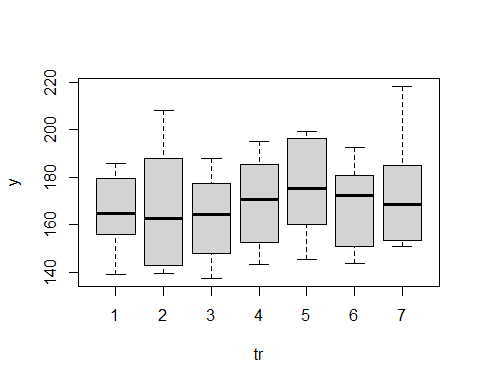
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 87.3  
#> 2 2 86.8  
#> 3 3 86.7  
#> 4 4 87.3  
#> 5 5 88.5  
#> 6 6 87.9  
#> 7 7 87.1



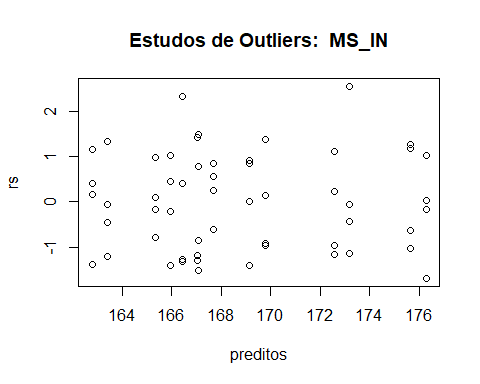
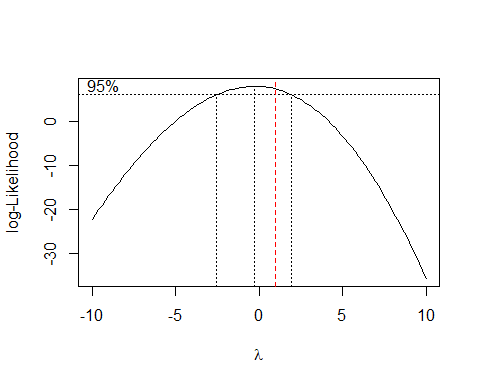
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "MS\_IN"  
#> [1] "===================================="



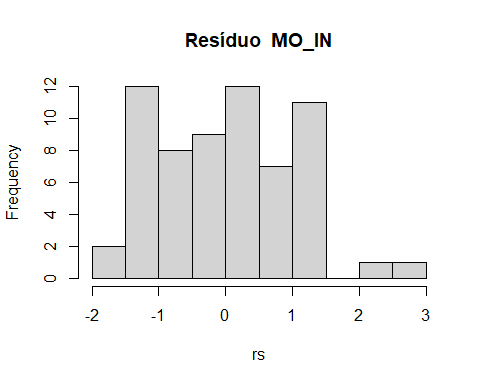
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95983, p-value = 0.03812



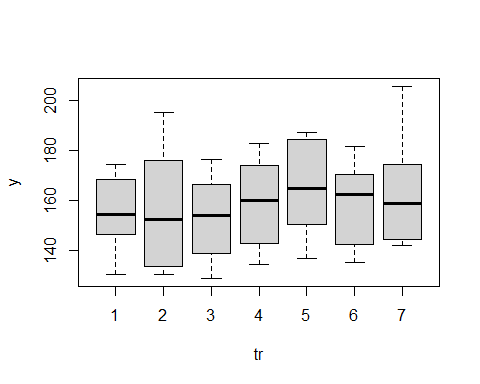
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 166.  
#> 2 2 167.  
#> 3 3 163.  
#> 4 4 169.  
#> 5 5 176.  
#> 6 6 167.  
#> 7 7 173.



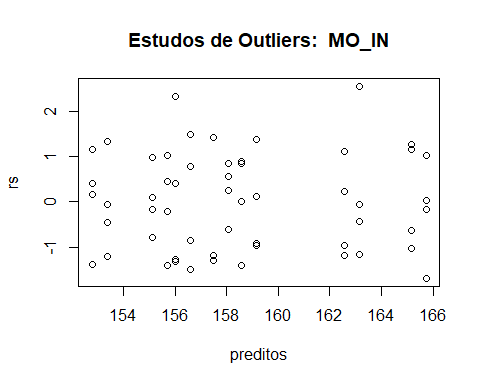
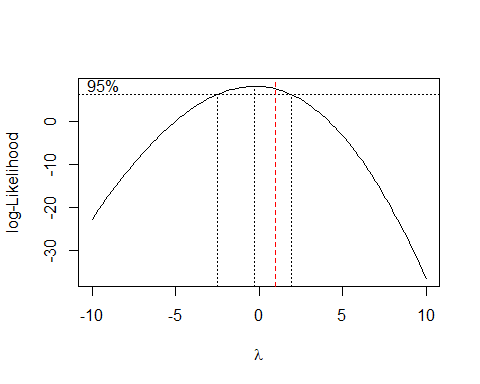
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "MO\_IN"  
#> [1] "===================================="



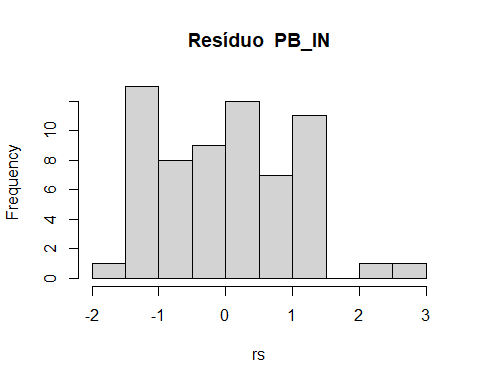
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95982, p-value = 0.0381



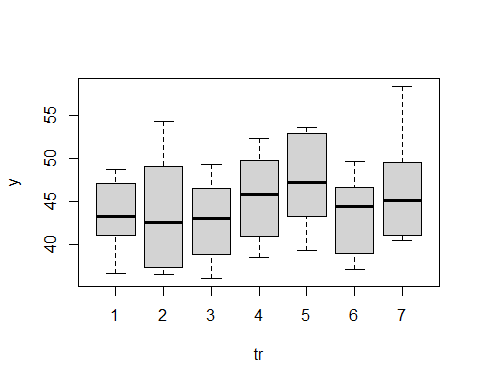
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 155.  
#> 2 2 156.  
#> 3 3 153.  
#> 4 4 159.  
#> 5 5 165.  
#> 6 6 158.  
#> 7 7 163.



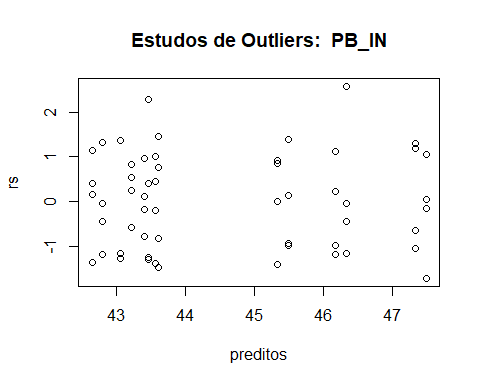
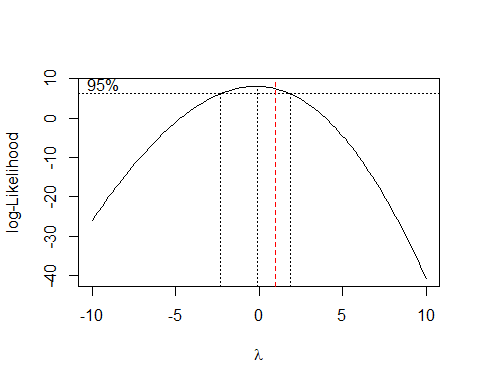
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PB\_IN"  
#> [1] "===================================="



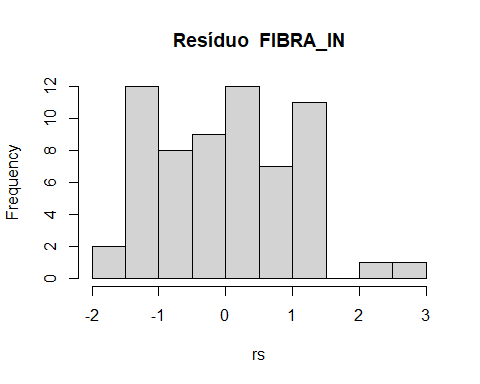
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96066, p-value = 0.04203



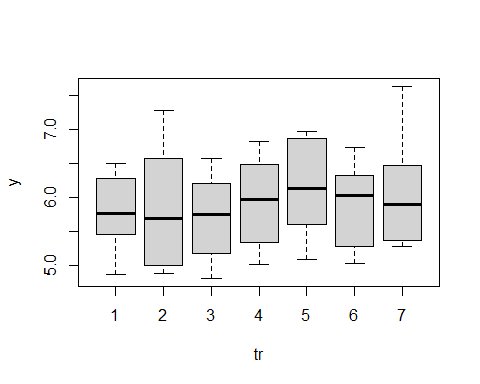
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 43.5  
#> 2 2 43.5  
#> 3 3 42.7  
#> 4 4 45.4  
#> 5 5 47.4  
#> 6 6 43.1  
#> 7 7 46.3



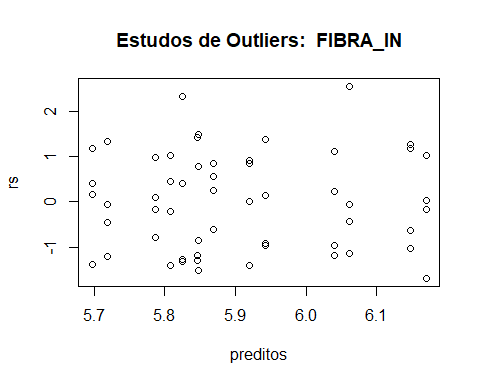
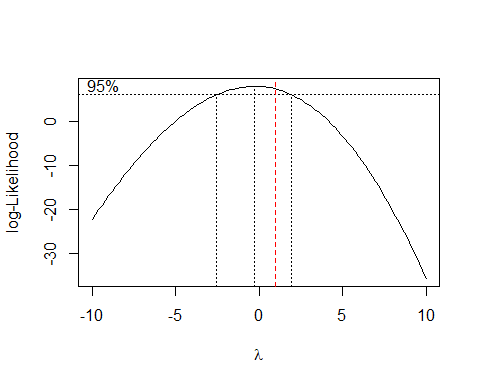
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "FIBRA\_IN"  
#> [1] "===================================="



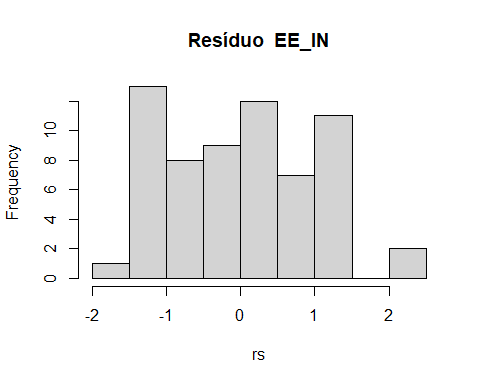
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95979, p-value = 0.03794



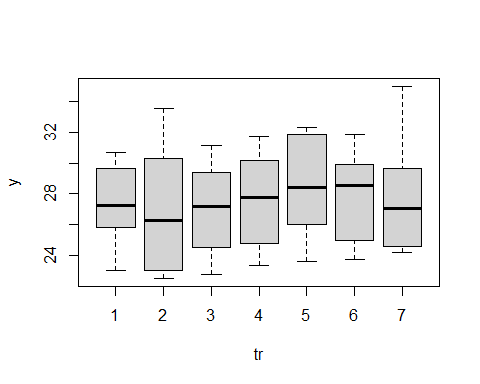
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 5.80  
#> 2 2 5.84  
#> 3 3 5.71  
#> 4 4 5.93  
#> 5 5 6.16  
#> 6 6 5.86  
#> 7 7 6.05



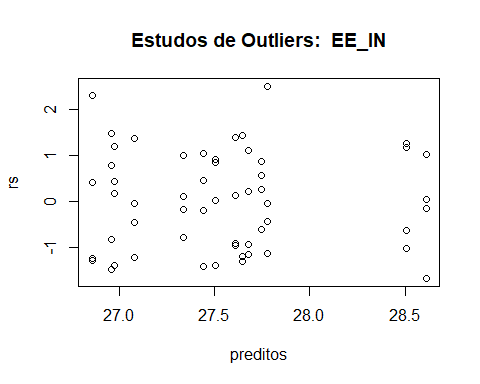
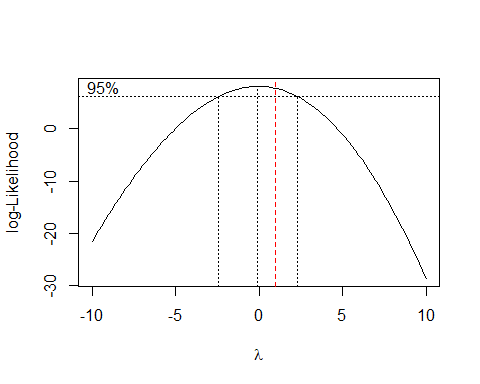
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "EE\_IN"  
#> [1] "===================================="



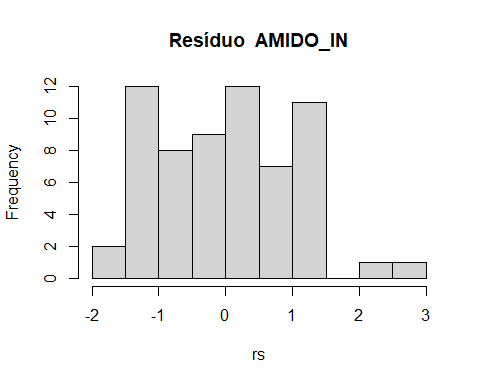
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95975, p-value = 0.03781



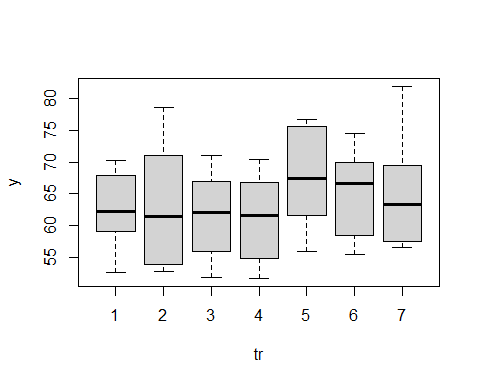
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 27.4  
#> 2 2 26.9  
#> 3 3 27.0  
#> 4 4 27.6  
#> 5 5 28.6  
#> 6 6 27.7  
#> 7 7 27.7



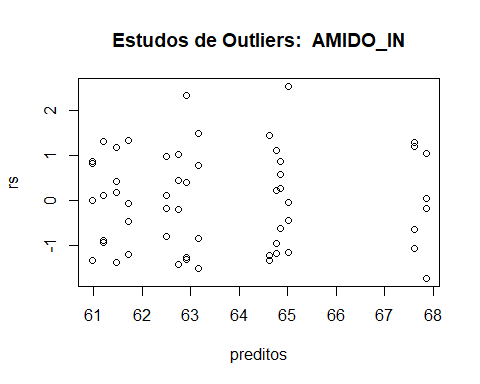
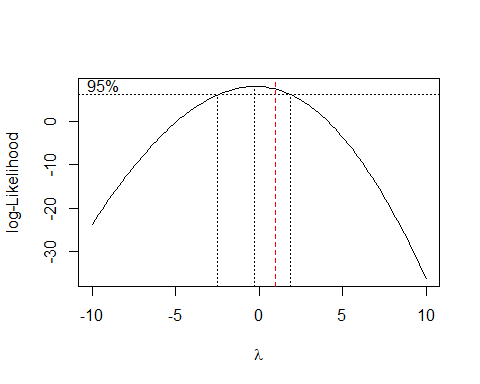
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AMIDO\_IN"  
#> [1] "===================================="



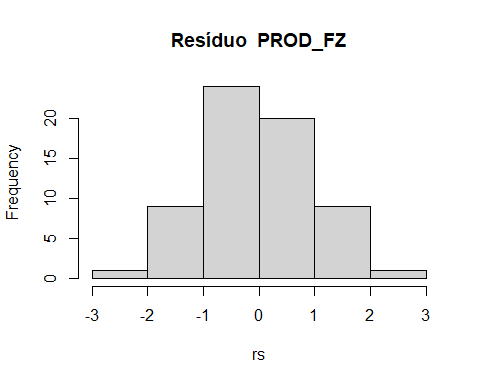
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.9612, p-value = 0.04478



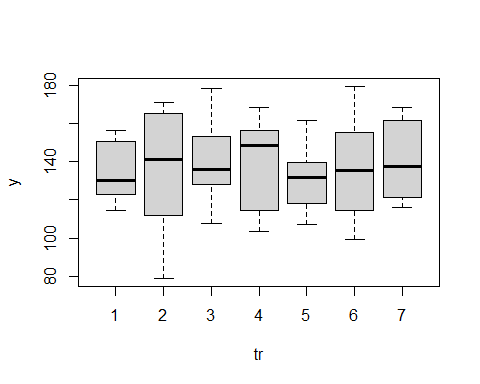
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 62.6  
#> 2 2 63.0  
#> 3 3 61.6  
#> 4 4 61.1  
#> 5 5 67.7  
#> 6 6 64.8  
#> 7 7 64.9



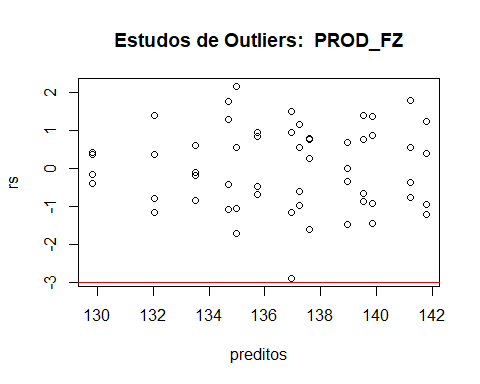
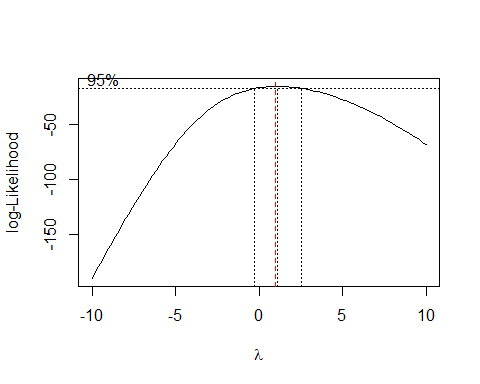
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PROD\_FZ"  
#> [1] "===================================="



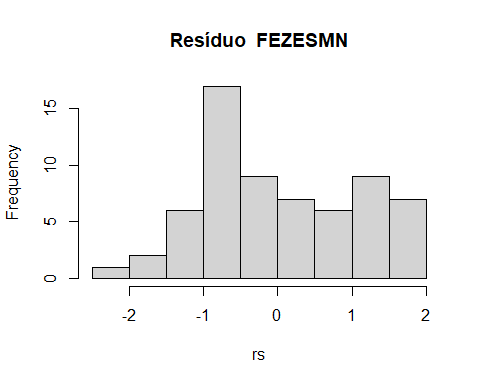
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97881, p-value = 0.3381



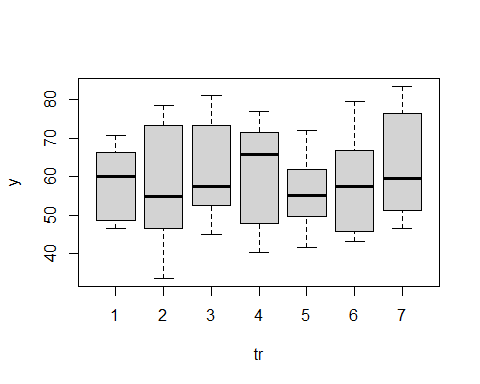
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 135.  
#> 2 2 136.  
#> 3 3 140.  
#> 4 4 139.  
#> 5 5 131.  
#> 6 6 136.  
#> 7 7 141.



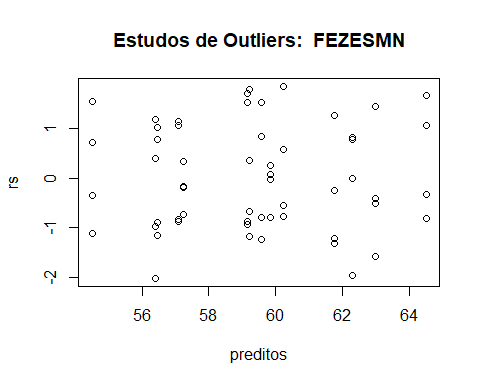
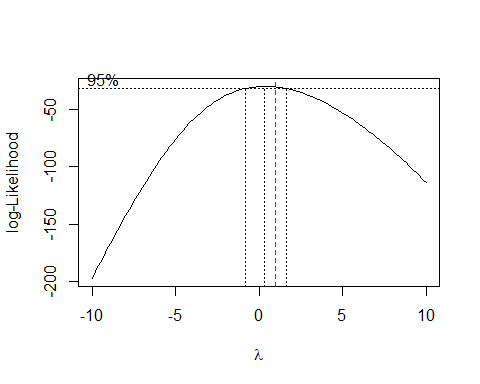
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "FEZESMN"  
#> [1] "===================================="



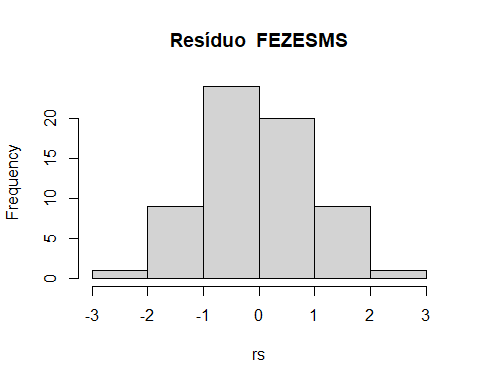
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95491, p-value = 0.02011



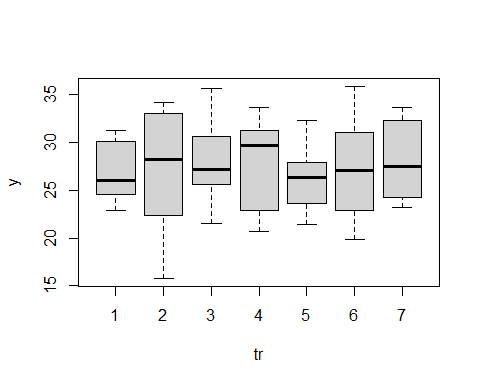
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 58.5  
#> 2 2 57.8  
#> 3 3 61.6  
#> 4 4 60.9  
#> 5 5 55.9  
#> 6 6 57.8  
#> 7 7 63.1



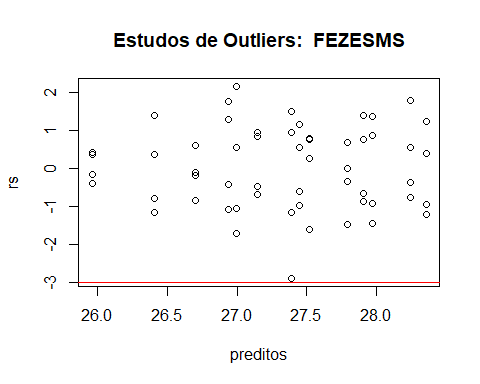
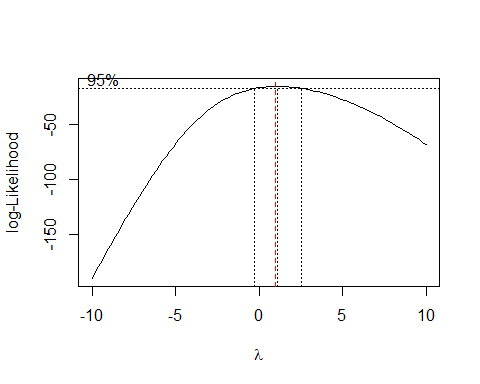
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "FEZESMS"  
#> [1] "===================================="



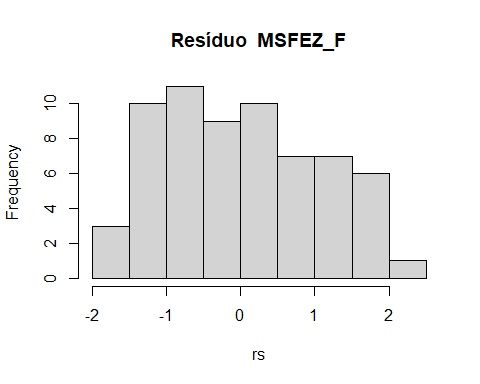
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97881, p-value = 0.3381



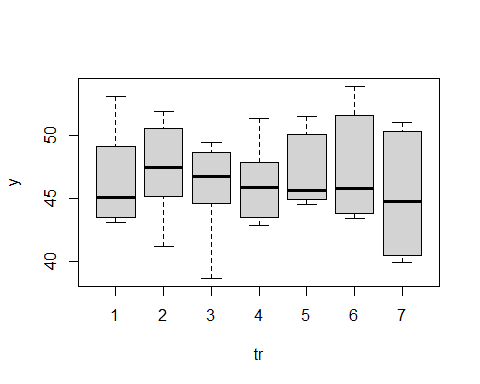
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 26.9  
#> 2 2 27.2  
#> 3 3 28.0  
#> 4 4 27.7  
#> 5 5 26.2  
#> 6 6 27.2  
#> 7 7 28.1



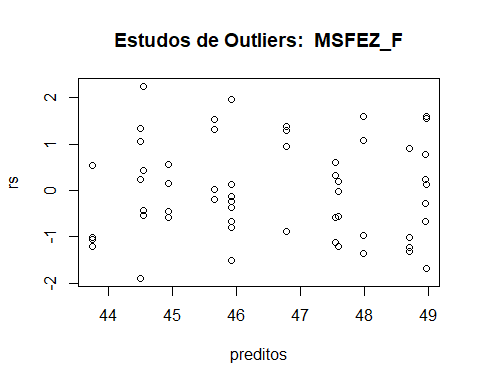
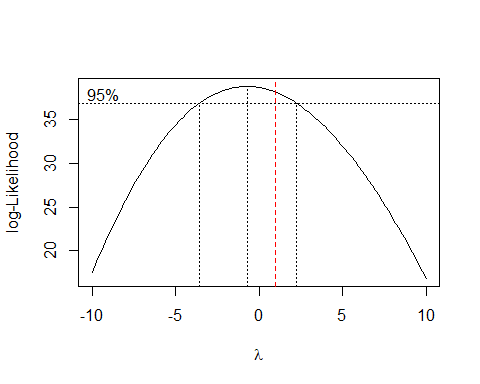
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "MSFEZ\_F"  
#> [1] "===================================="



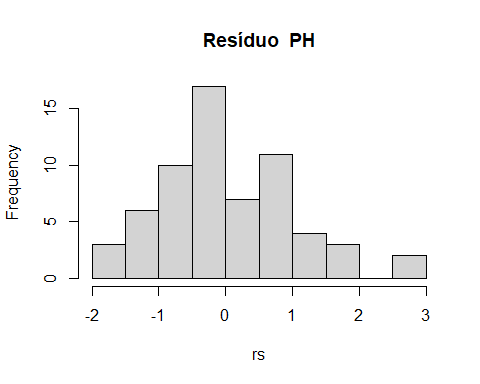
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96968, p-value = 0.1164



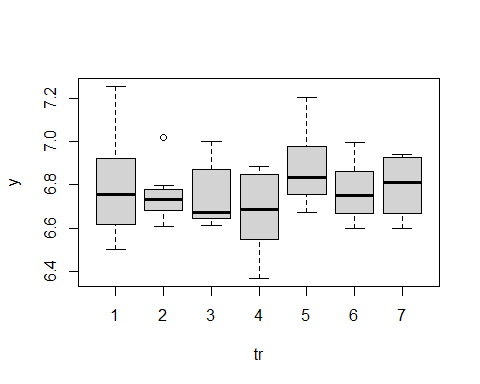
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 46.5  
#> 2 2 47.4  
#> 3 3 46.0  
#> 4 4 46.1  
#> 5 5 47.2  
#> 6 6 47.4  
#> 7 7 45.3



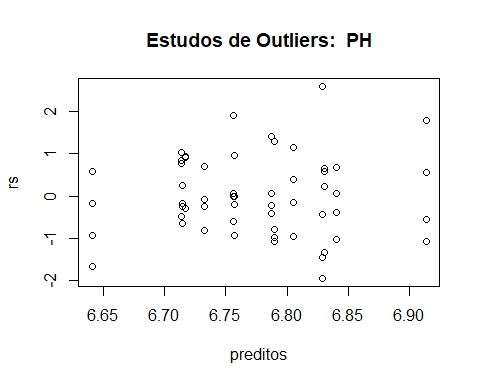
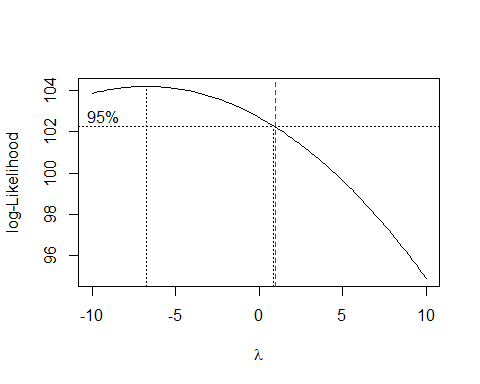
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PH"  
#> [1] "===================================="



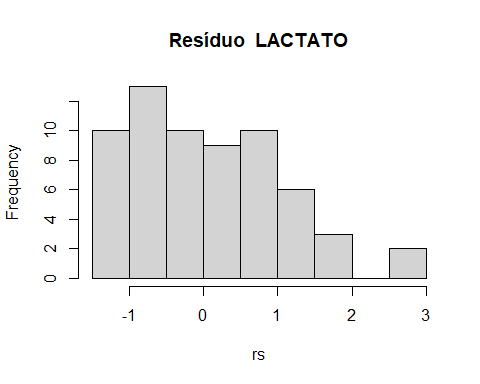
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97713, p-value = 0.2893



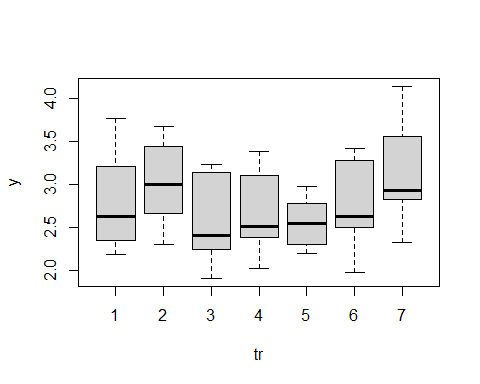
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 6.79  
#> 2 2 6.75  
#> 3 3 6.76  
#> 4 4 6.68  
#> 5 5 6.88  
#> 6 6 6.77  
#> 7 7 6.79



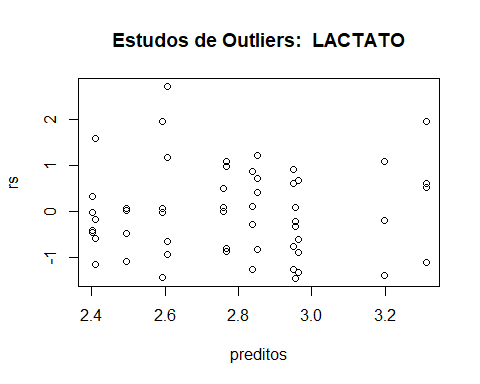
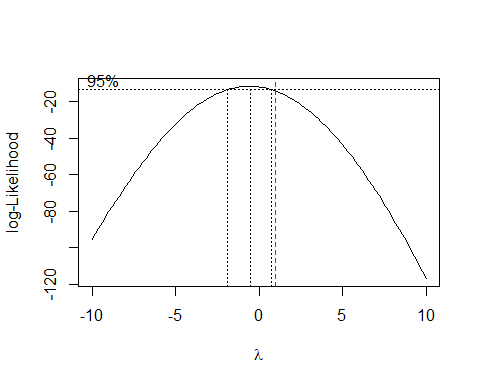
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "LACTATO"  
#> [1] "===================================="



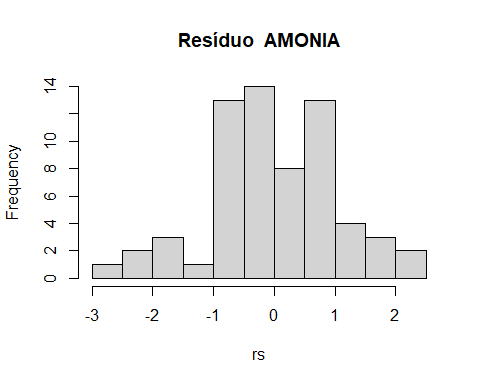
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95087, p-value = 0.01362



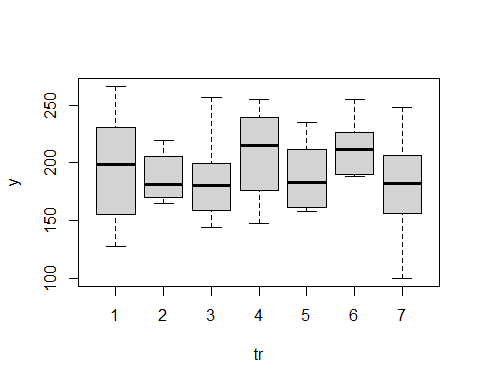
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 2.79  
#> 2 2 3.02  
#> 3 3 2.59  
#> 4 4 2.67  
#> 5 5 2.55  
#> 6 6 2.77  
#> 7 7 3.13



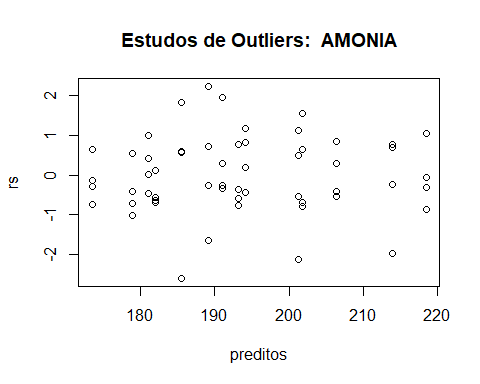
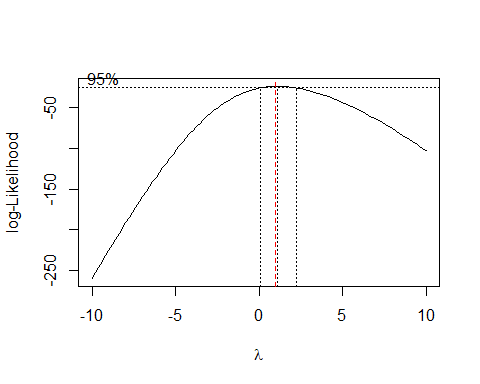
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AMONIA"  
#> [1] "===================================="



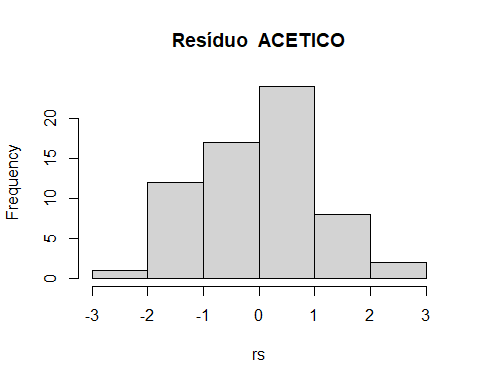
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.9752, p-value = 0.2242



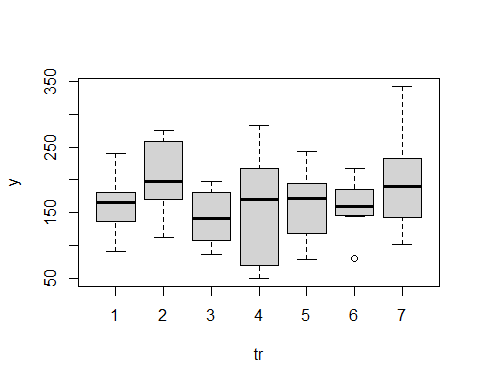
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 195.  
#> 2 2 187.  
#> 3 3 185.  
#> 4 4 208.  
#> 5 5 188.  
#> 6 6 212.  
#> 7 7 180.



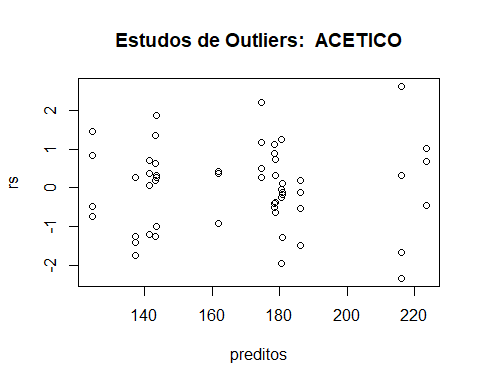
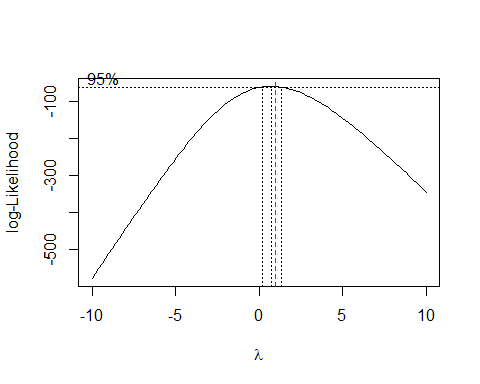
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "ACETICO"  
#> [1] "===================================="



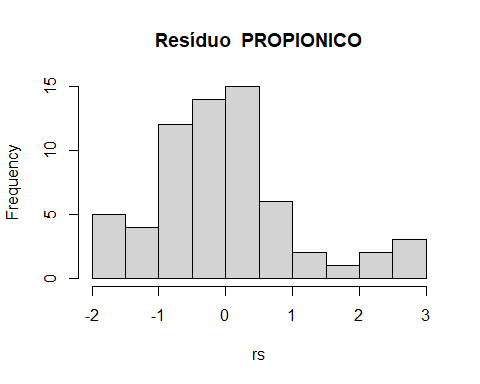
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98983, p-value = 0.8792



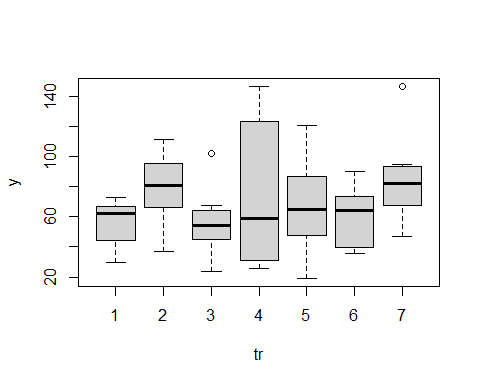
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 162.  
#> 2 2 205.  
#> 3 3 143.  
#> 4 4 156.  
#> 5 5 162.  
#> 6 6 160.  
#> 7 7 197.



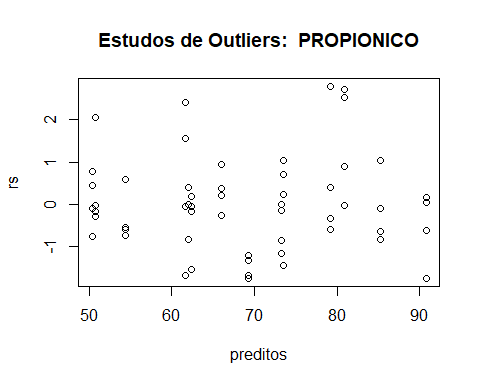
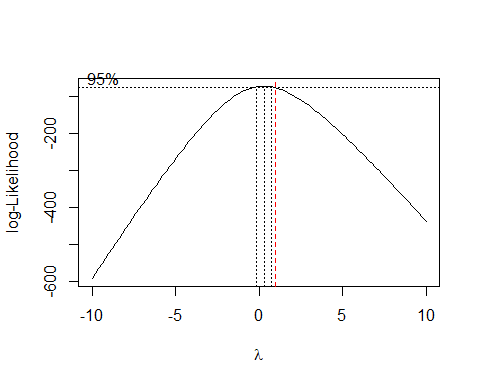
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PROPIONICO"  
#> [1] "===================================="



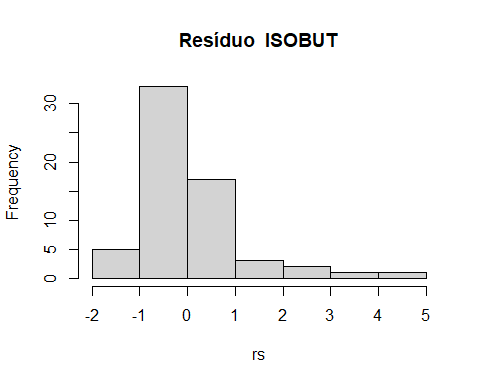
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.94001, p-value = 0.0038



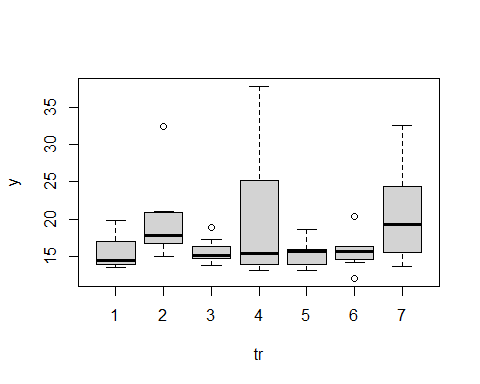
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 56.2  
#> 2 2 79.4  
#> 3 3 56.6  
#> 4 4 75.0  
#> 5 5 67.5  
#> 6 6 60.2  
#> 7 7 85.0



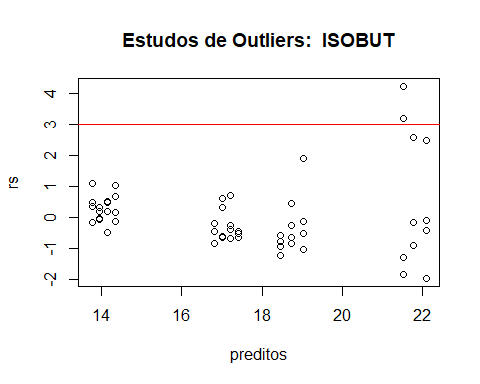
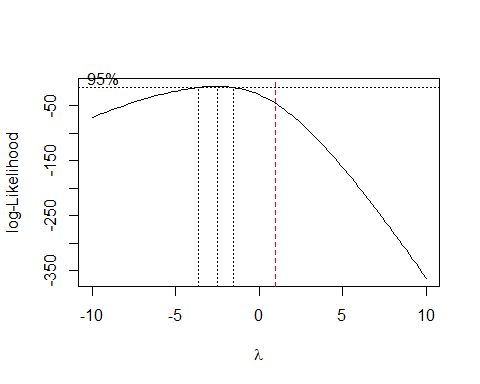
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "ISOBUT"  
#> [1] "===================================="



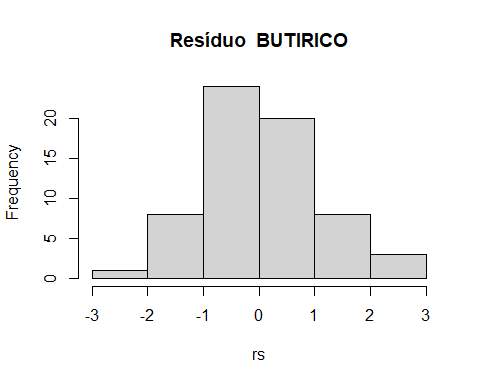
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.86126, p-value = 4.934e-06



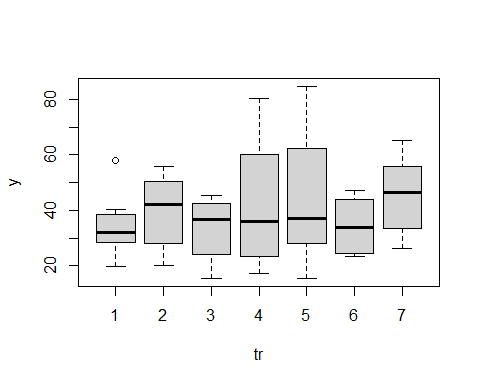
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 15.5  
#> 2 2 20.0  
#> 3 3 15.7  
#> 4 4 20.0  
#> 5 5 15.3  
#> 6 6 15.7  
#> 7 7 20.6



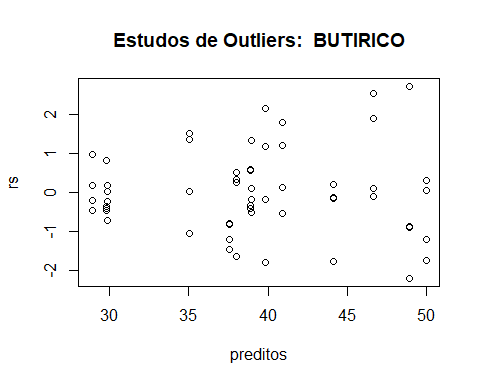
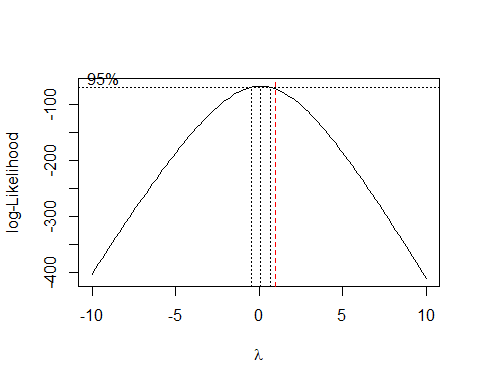
#> Outliers Trat Bloco Y  
#> 28 4 2 37.85974  
#> 29 4 2 34.54399  
#>   
#> [1] "===================================="  
#> [1] "BUTIRICO"  
#> [1] "===================================="



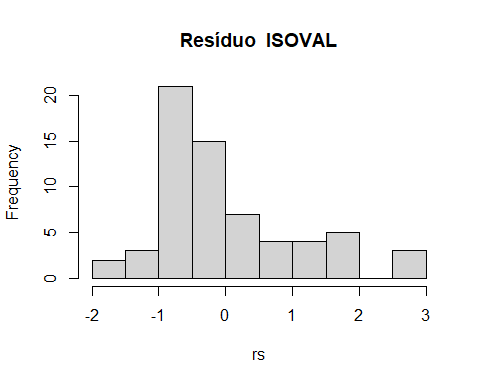
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96947, p-value = 0.1136



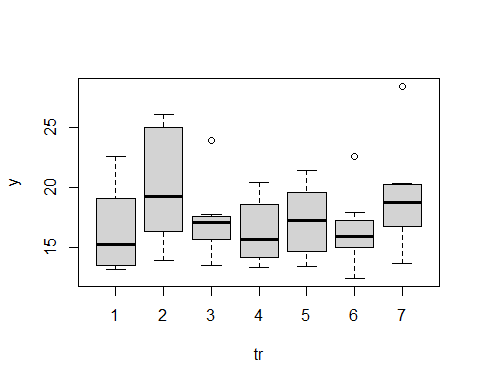
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 34.4  
#> 2 2 39.6  
#> 3 3 33.4  
#> 4 4 42.1  
#> 5 5 44.4  
#> 6 6 34.3  
#> 7 7 45.4



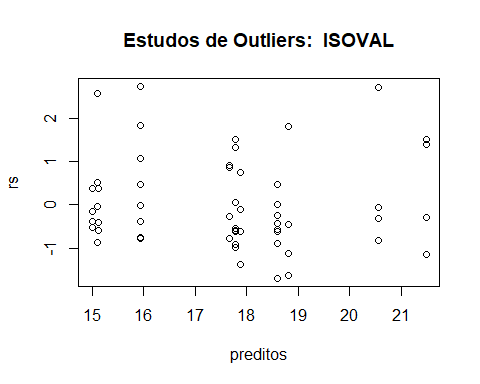
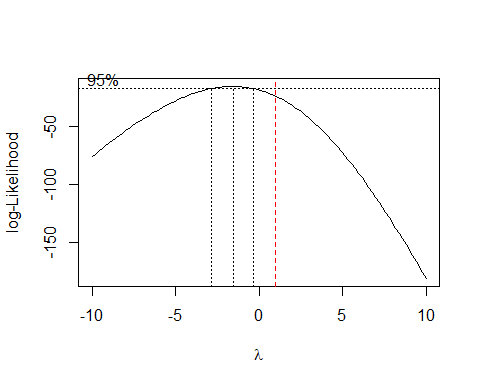
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "ISOVAL"  
#> [1] "===================================="



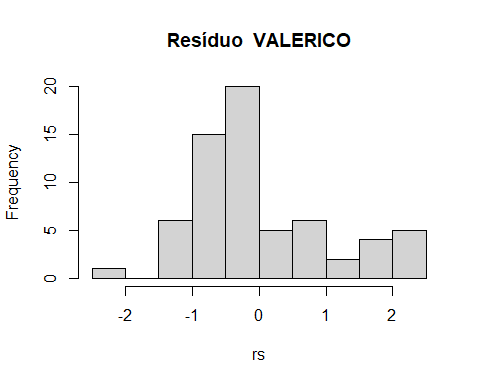
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.91185, p-value = 0.0002308



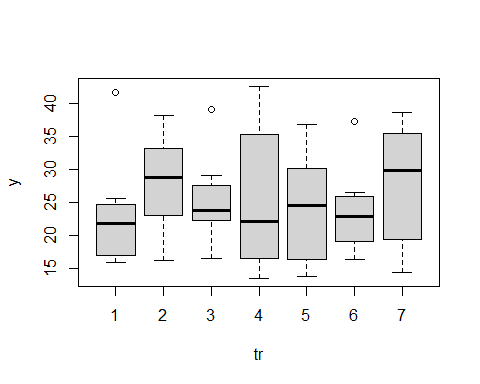
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 16.4  
#> 2 2 20.2  
#> 3 3 17.3  
#> 4 4 16.3  
#> 5 5 17.3  
#> 6 6 16.4  
#> 7 7 19.2



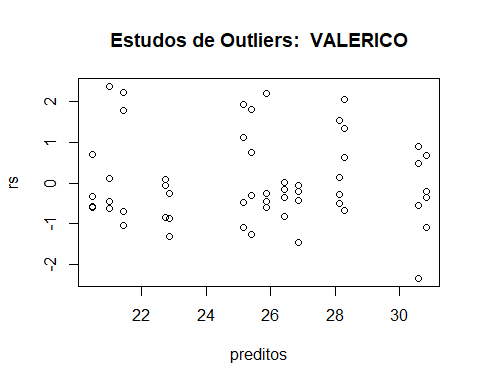
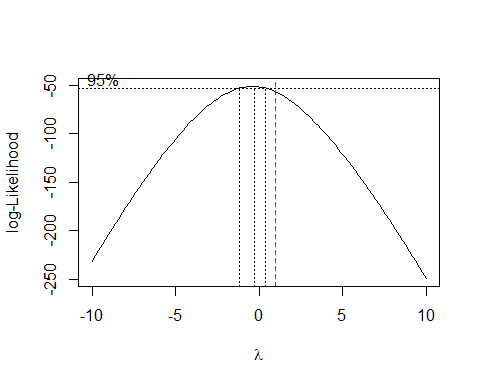
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "VALERICO"  
#> [1] "===================================="



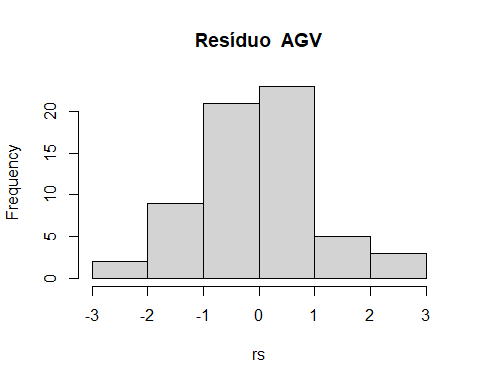
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.90926, p-value = 0.0001822



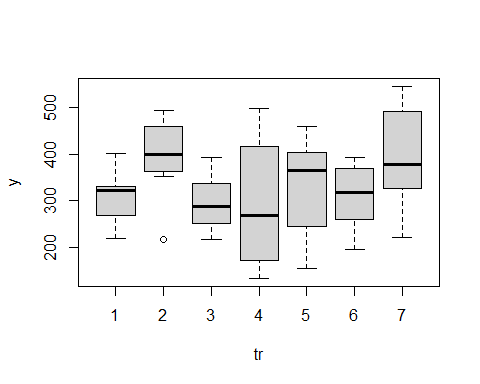
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 23.2  
#> 2 2 28.1  
#> 3 3 25.4  
#> 4 4 25.6  
#> 5 5 24.2  
#> 6 6 23.7  
#> 7 7 27.9



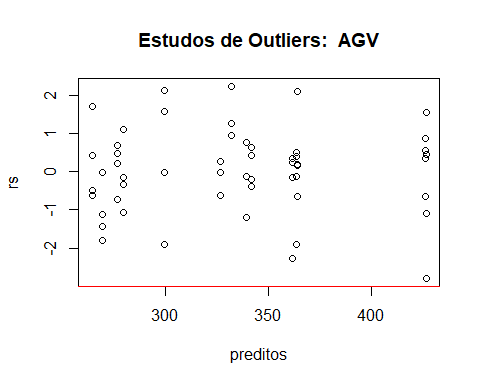
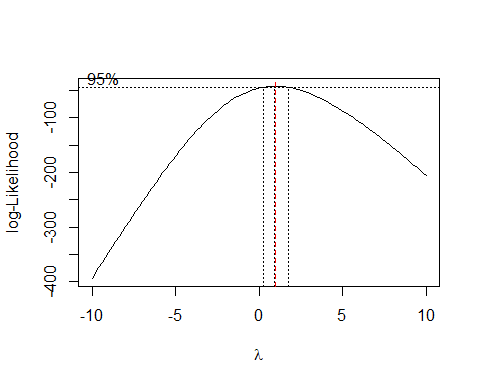
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV"  
#> [1] "===================================="



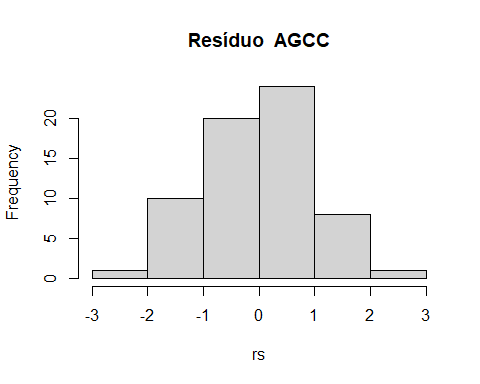
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98103, p-value = 0.4399



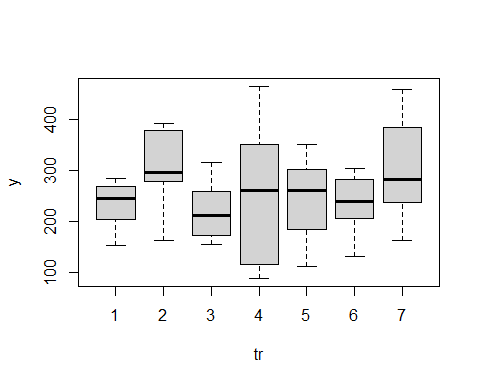
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 308.  
#> 2 2 395.  
#> 3 3 295.  
#> 4 4 296.  
#> 5 5 330.  
#> 6 6 310.  
#> 7 7 395.



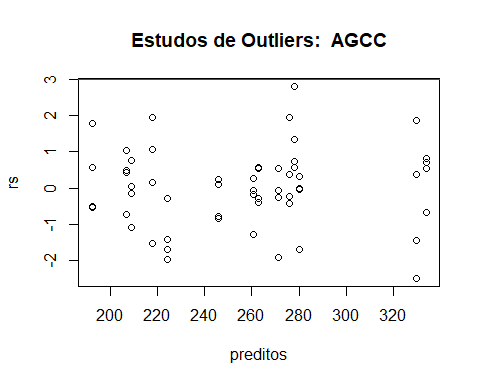
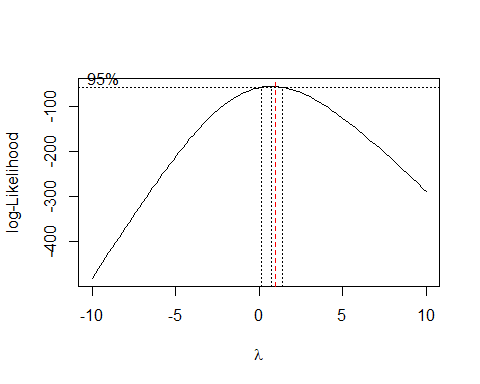
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGCC"  
#> [1] "===================================="



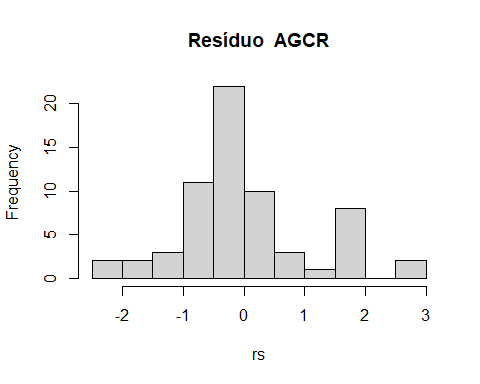
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98299, p-value = 0.523



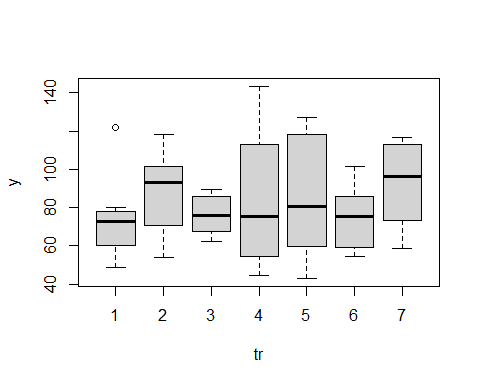
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 234.  
#> 2 2 307.  
#> 3 3 219.  
#> 4 4 251.  
#> 5 5 245.  
#> 6 6 236.  
#> 7 7 303.



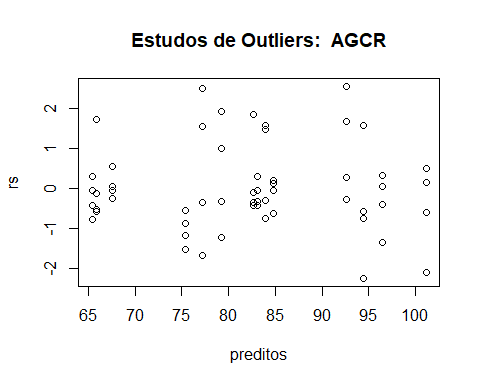
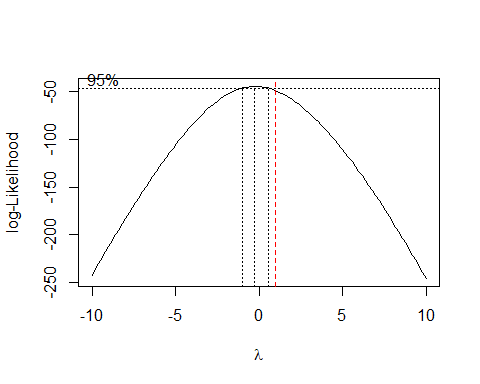
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGCR"  
#> [1] "===================================="



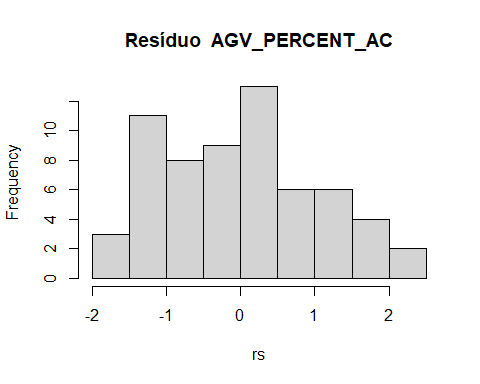
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.93463, p-value = 0.00215



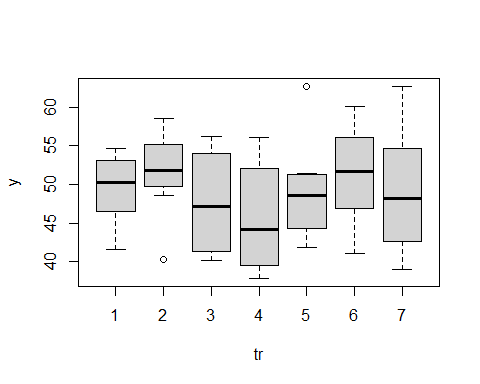
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 74.0  
#> 2 2 87.9  
#> 3 3 76.1  
#> 4 4 84.0  
#> 5 5 85.8  
#> 6 6 74.5  
#> 7 7 92.5



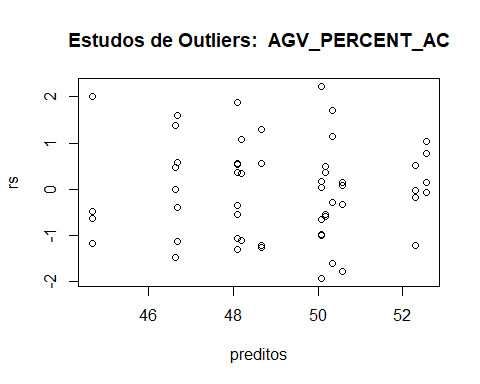
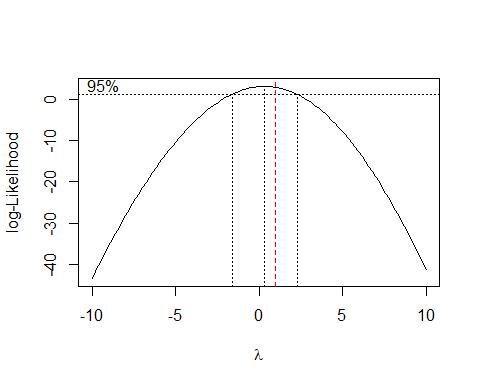
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_AC"  
#> [1] "===================================="



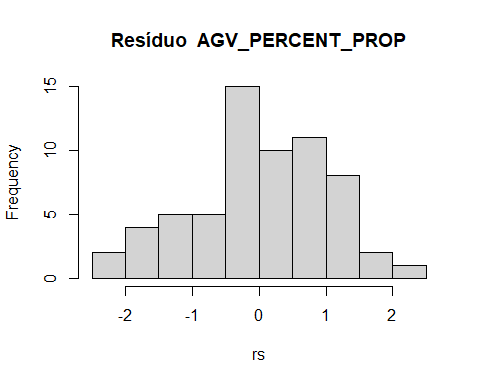
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97655, p-value = 0.2807



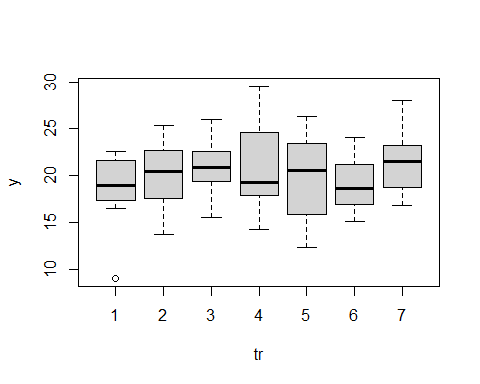
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 49.3  
#> 2 2 51.6  
#> 3 3 47.7  
#> 4 4 45.7  
#> 5 5 49.1  
#> 6 6 51.3  
#> 7 7 49.1



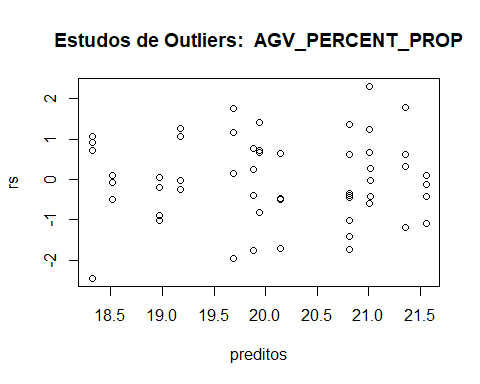
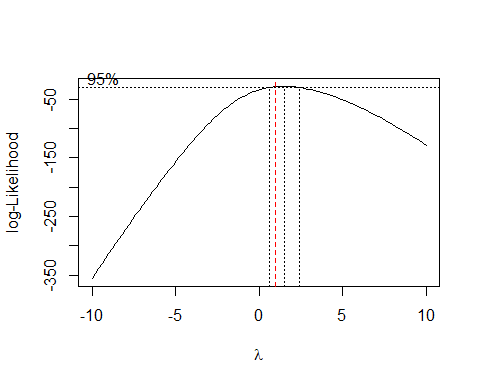
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_PROP"  
#> [1] "===================================="



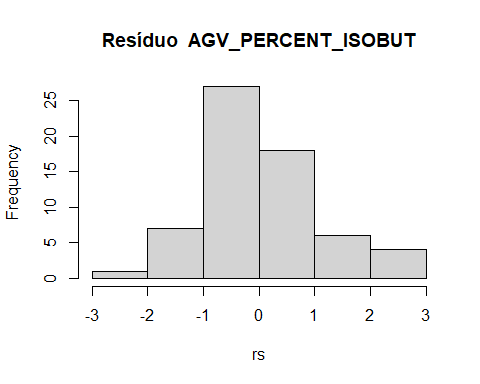
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98344, p-value = 0.5567



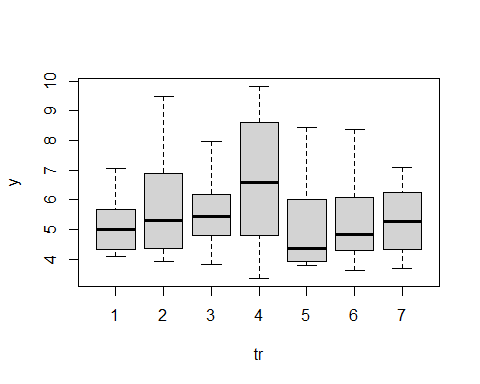
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 18.4  
#> 2 2 20.0  
#> 3 3 20.9  
#> 4 4 20.9  
#> 5 5 19.8  
#> 6 6 19.1  
#> 7 7 21.5



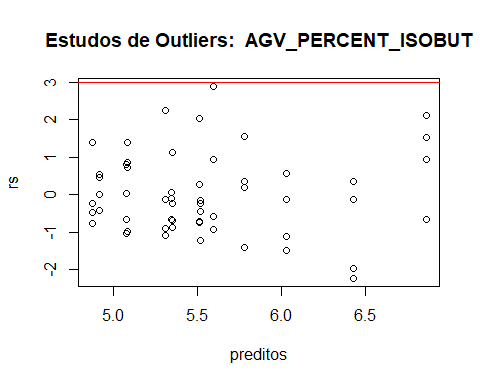
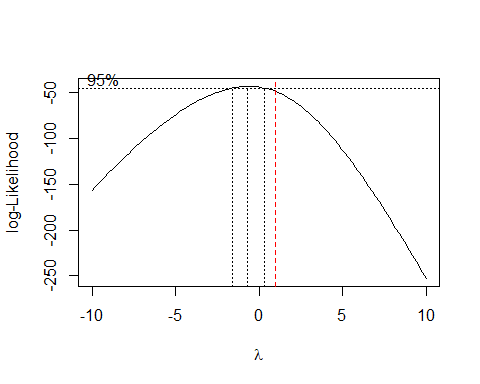
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_ISOBUT"  
#> [1] "===================================="



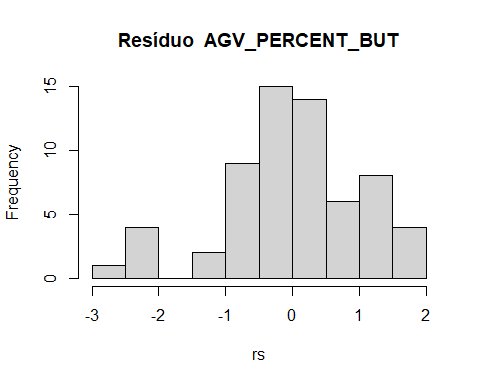
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97477, p-value = 0.2213



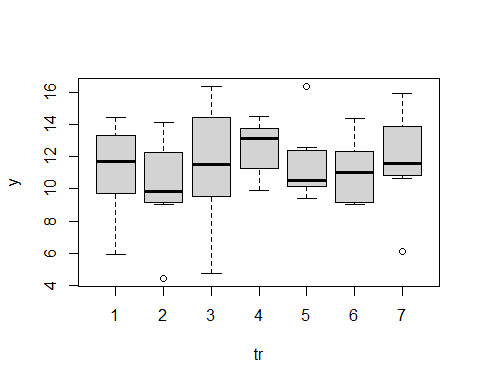
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 5.13  
#> 2 2 5.81  
#> 3 3 5.60  
#> 4 4 6.65  
#> 5 5 5.09  
#> 6 6 5.29  
#> 7 7 5.30



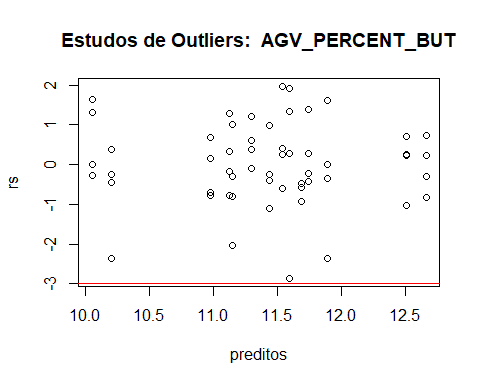
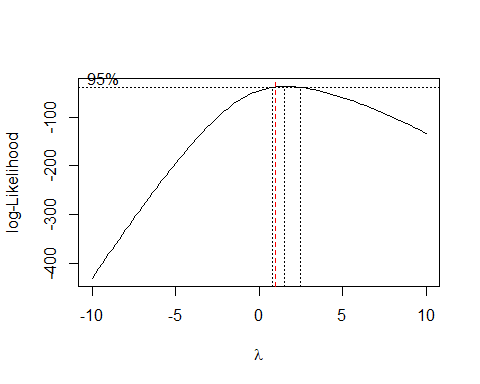
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_BUT"  
#> [1] "===================================="



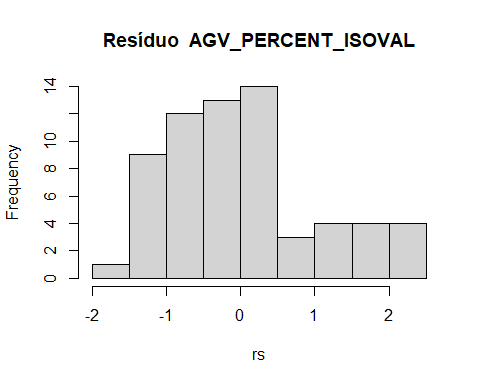
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96631, p-value = 0.0819



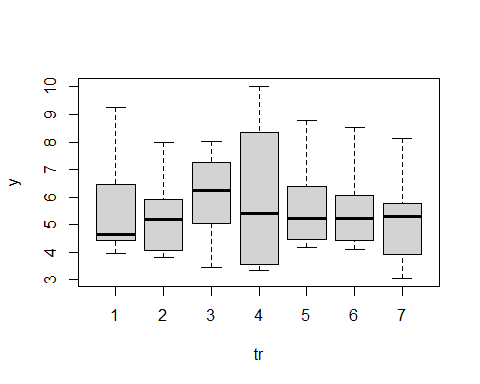
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 11.2  
#> 2 2 10.1  
#> 3 3 11.5  
#> 4 4 12.6  
#> 5 5 11.6  
#> 6 6 11.1  
#> 7 7 11.8



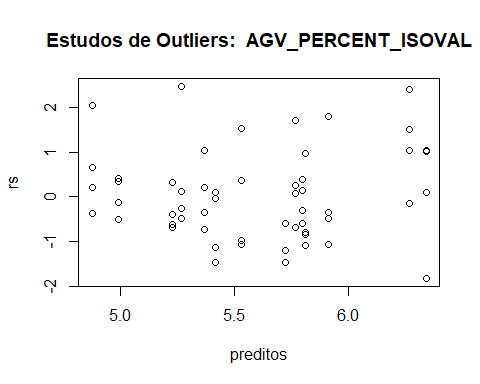
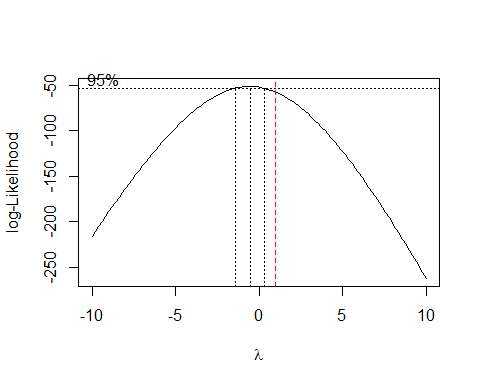
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_ISOVAL"  
#> [1] "===================================="



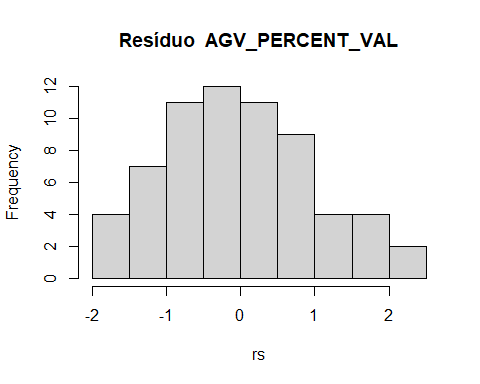
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.94875, p-value = 0.009936



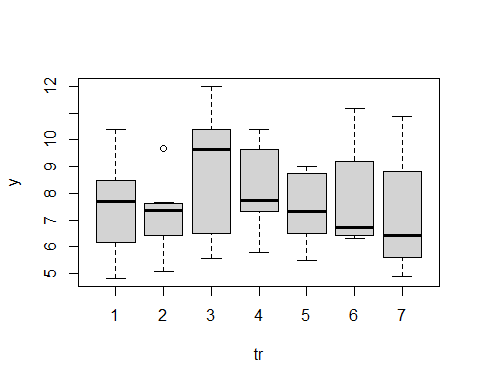
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 5.54  
#> 2 2 5.26  
#> 3 3 6.07  
#> 4 4 6.00  
#> 5 5 5.64  
#> 6 6 5.50  
#> 7 7 5.15



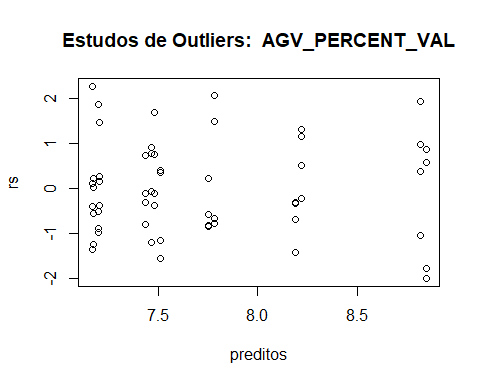
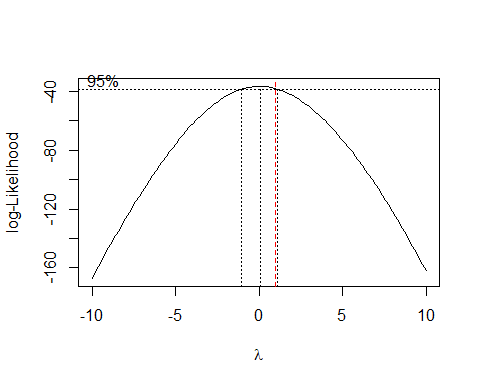
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "AGV\_PERCENT\_VAL"  
#> [1] "===================================="



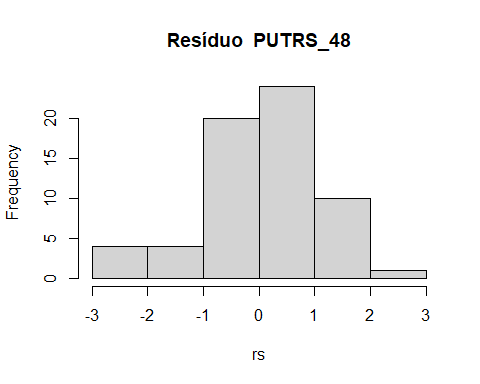
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98134, p-value = 0.4431



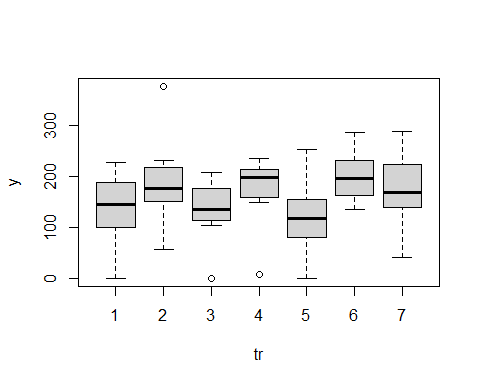
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 7.50  
#> 2 2 7.19  
#> 3 3 8.83  
#> 4 4 8.20  
#> 5 5 7.45  
#> 6 6 7.77  
#> 7 7 7.18



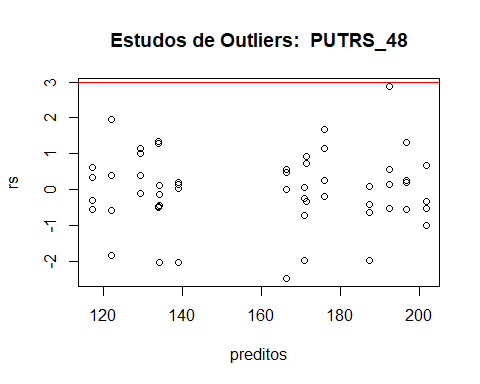
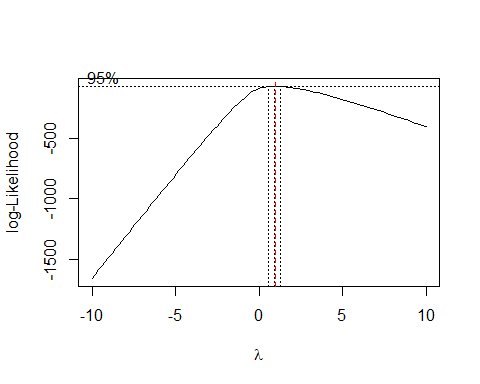
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "PUTRS\_48"  
#> [1] "===================================="



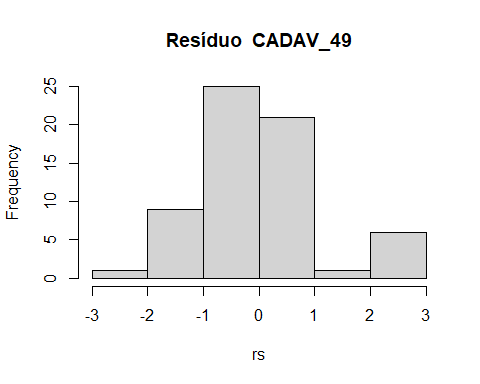
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96034, p-value = 0.04047



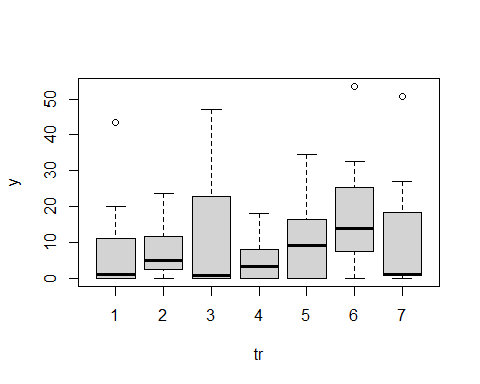
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 136.  
#> 2 2 190.  
#> 3 3 132.  
#> 4 4 168.  
#> 5 5 120.  
#> 6 6 199.  
#> 7 7 173.



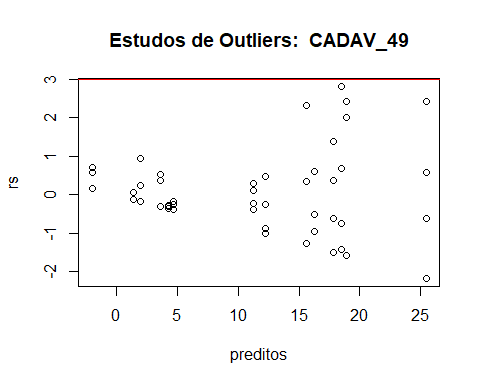
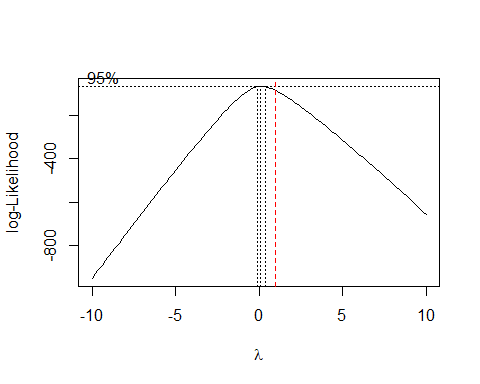
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "CADAV\_49"  
#> [1] "===================================="



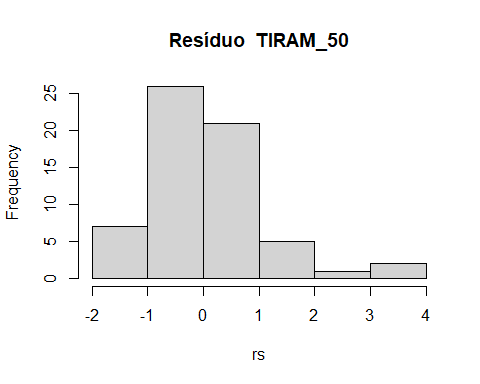
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.9299, p-value = 0.001458



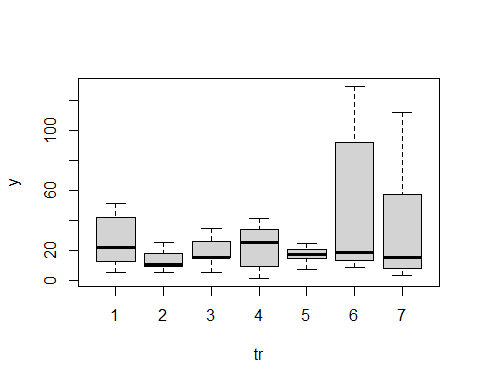
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 8.53  
#> 2 2 8.11  
#> 3 3 11.8   
#> 4 4 5.13  
#> 5 5 10.7   
#> 6 6 18.3   
#> 7 7 11.4



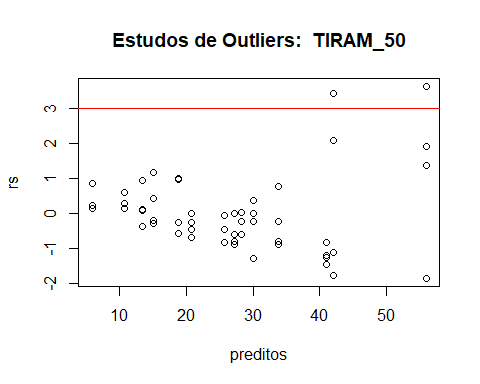
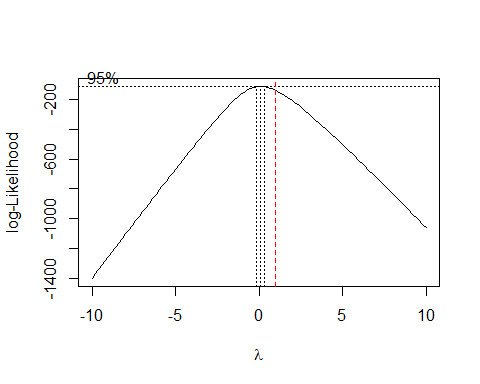
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "TIRAM\_50"  
#> [1] "===================================="



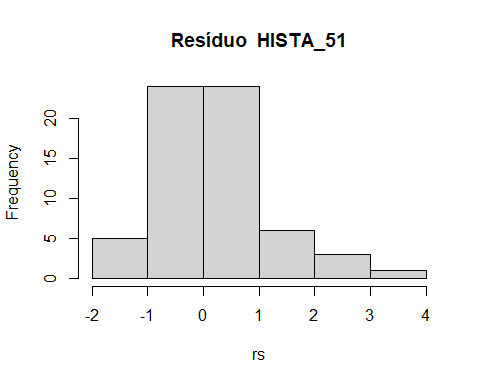
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.91921, p-value = 0.0005726



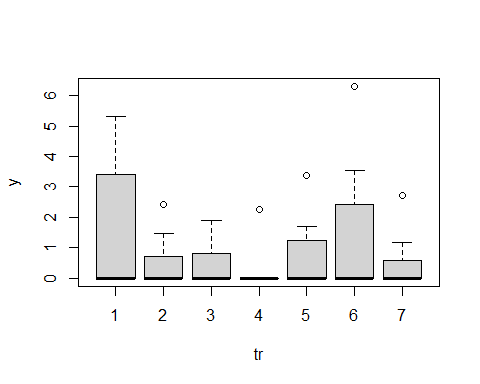
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 26.4  
#> 2 2 13.4  
#> 3 3 19.8  
#> 4 4 22.6  
#> 5 5 17.2  
#> 6 6 48.5  
#> 7 7 34.6



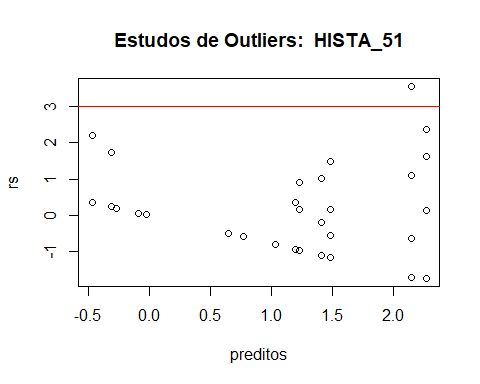
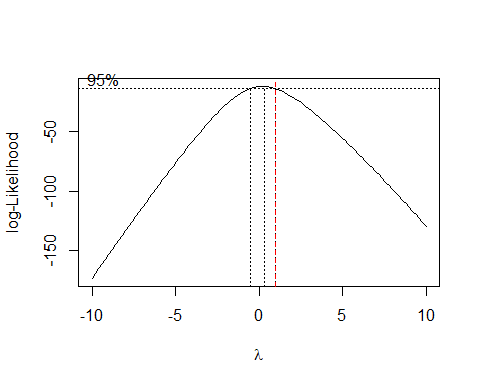
#> Outliers Trat Bloco Y  
#> 41 6 1 129.25  
#> 48 7 1 111.88  
#>   
#> [1] "===================================="  
#> [1] "HISTA\_51"  
#> [1] "===================================="



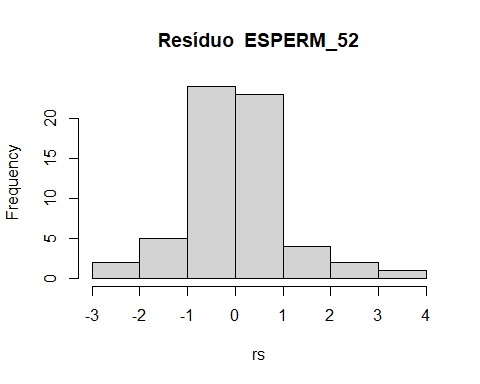
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.90948, p-value = 0.0002088



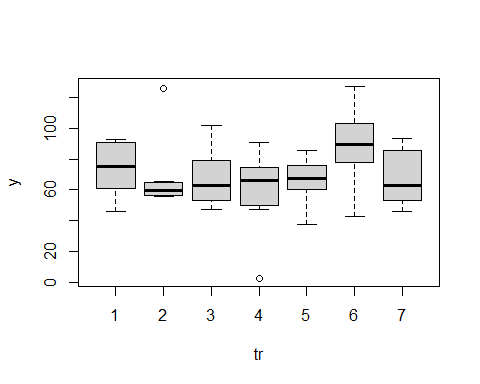
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 1.52   
#> 2 2 0.484  
#> 3 3 0.445  
#> 4 4 0.282  
#> 5 5 0.731  
#> 6 6 1.40   
#> 7 7 0.556



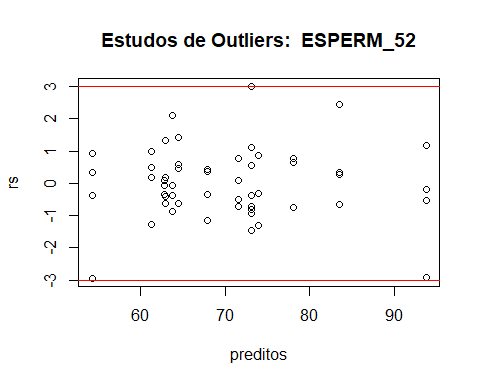
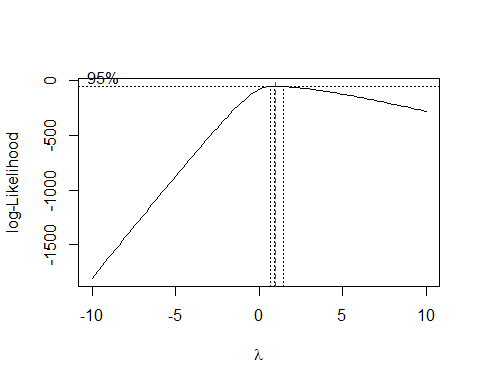
#> Outliers Trat Bloco Y  
#> 47 6 2 6.3  
#>   
#> [1] "===================================="  
#> [1] "ESPERM\_52"  
#> [1] "===================================="



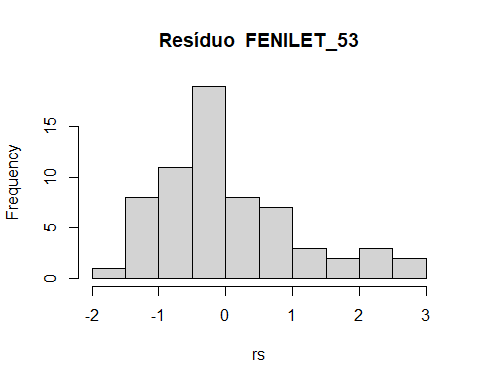
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.9632, p-value = 0.06369



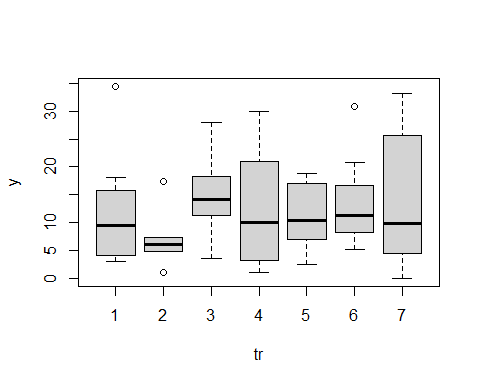
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 72.2  
#> 2 2 68.0  
#> 3 3 68.1  
#> 4 4 59.3  
#> 5 5 66.4  
#> 6 6 88.7  
#> 7 7 68.0



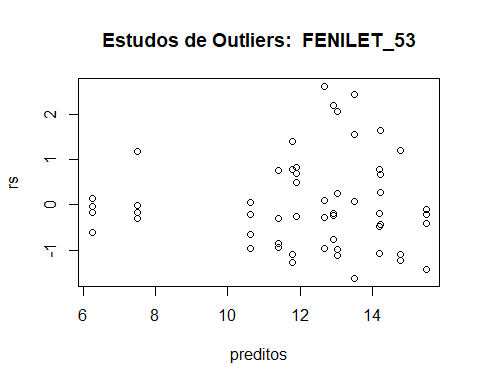
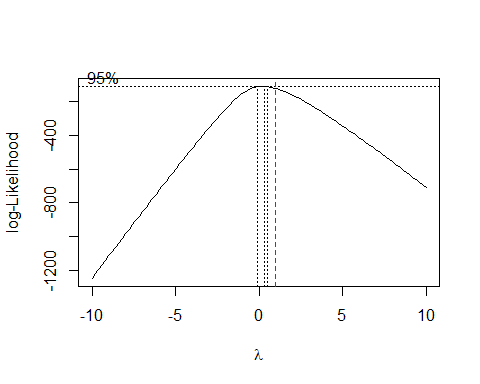
#> Outliers Trat Bloco Y  
#> 15 2 2 125.51  
#>   
#> [1] "===================================="  
#> [1] "FENILET\_53"  
#> [1] "===================================="



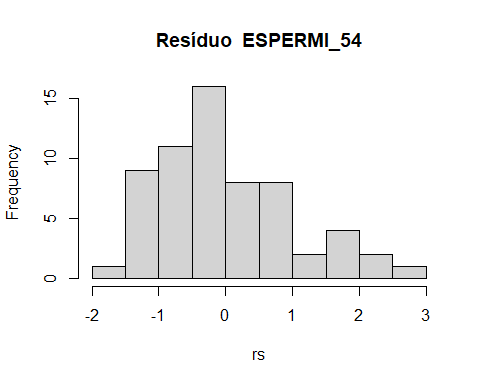
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.92578, p-value = 0.000874



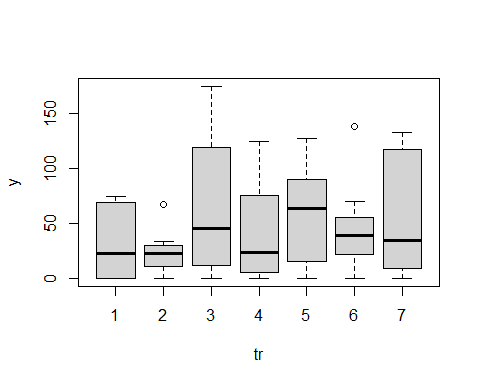
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 12.0   
#> 2 2 6.88  
#> 3 3 14.9   
#> 4 4 12.4   
#> 5 5 11.3   
#> 6 6 13.6   
#> 7 7 14.1



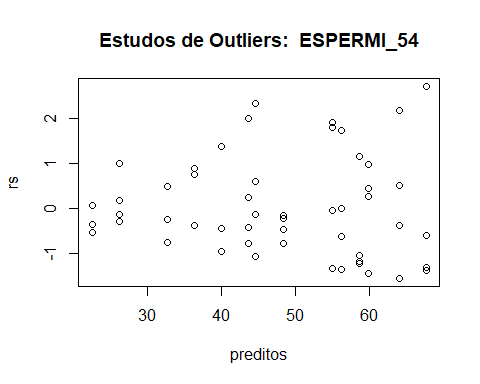
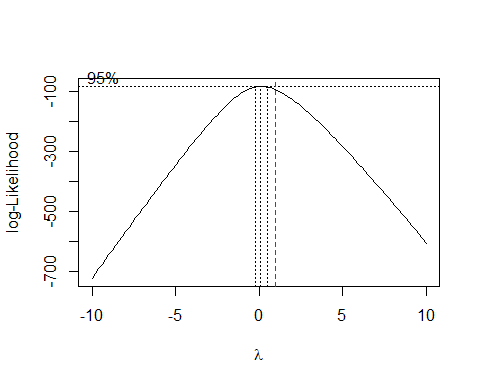
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "ESPERMI\_54"  
#> [1] "===================================="



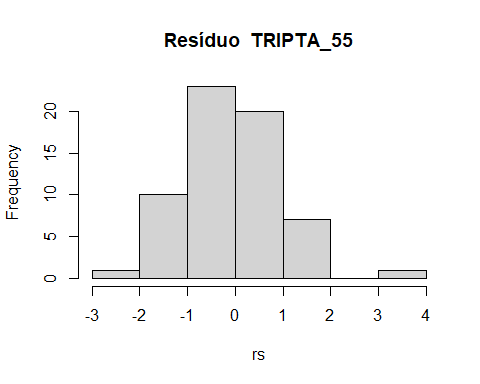
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.94504, p-value = 0.007757



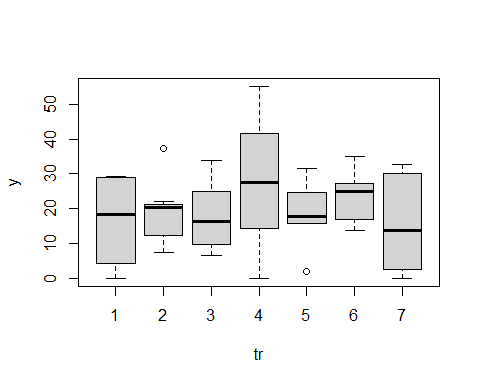
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 34.3  
#> 2 2 24.3  
#> 3 3 65.9  
#> 4 4 41.8  
#> 5 5 58.0  
#> 6 6 46.5  
#> 7 7 56.9



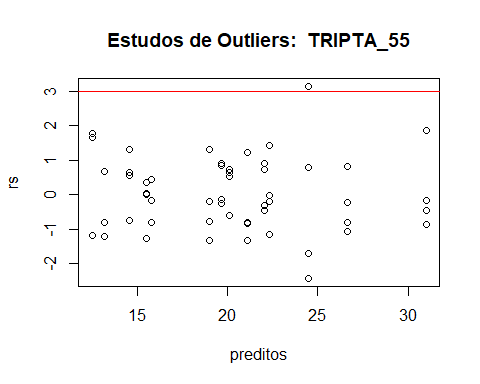
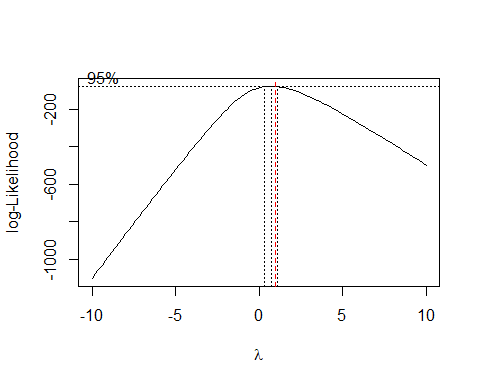
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "TRIPTA\_55"  
#> [1] "===================================="



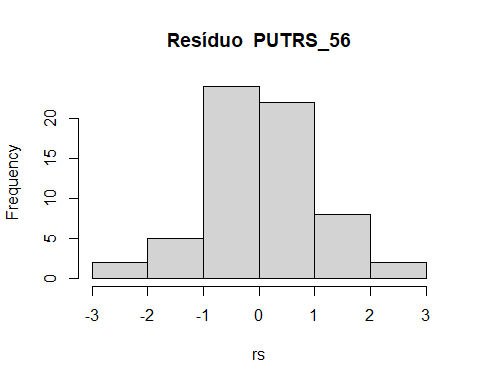
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.97657, p-value = 0.2812



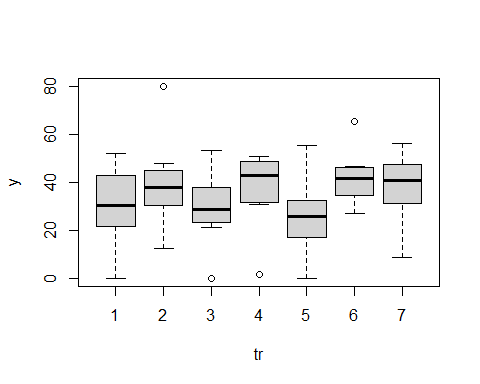
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 16.9  
#> 2 2 19.1  
#> 3 3 17.8  
#> 4 4 27.8  
#> 5 5 18.8  
#> 6 6 23.4  
#> 7 7 15.7



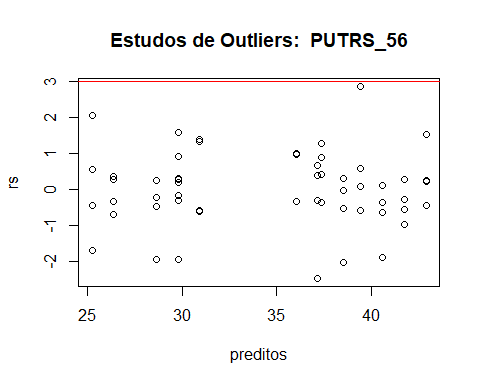
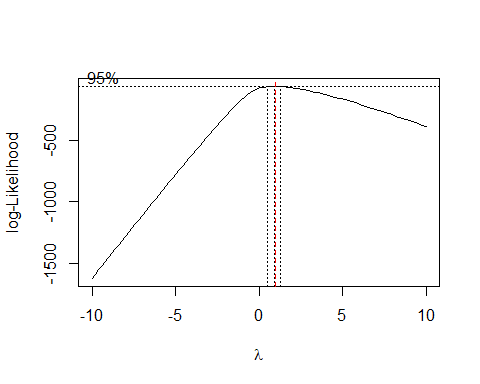
#> Outliers Trat Bloco Y  
#> 26 4 1 55.17  
#>   
#> [1] "===================================="  
#> [1] "PUTRS\_56"  
#> [1] "===================================="



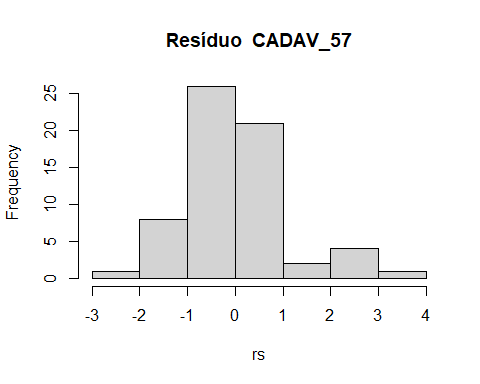
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.96672, p-value = 0.08595



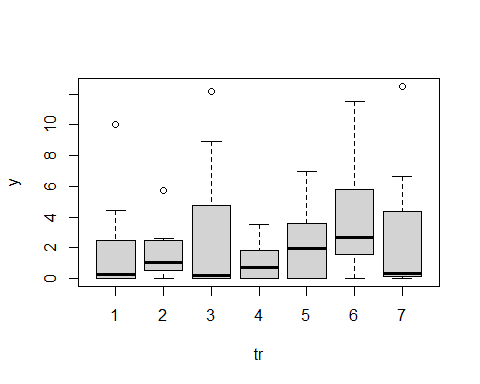
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 30.4  
#> 2 2 40.0  
#> 3 3 29.2  
#> 4 4 36.7  
#> 5 5 25.8  
#> 6 6 42.3  
#> 7 7 37.9



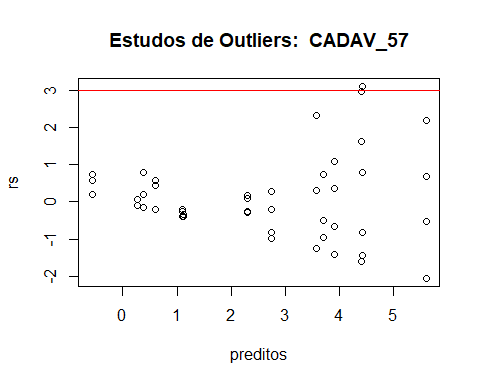
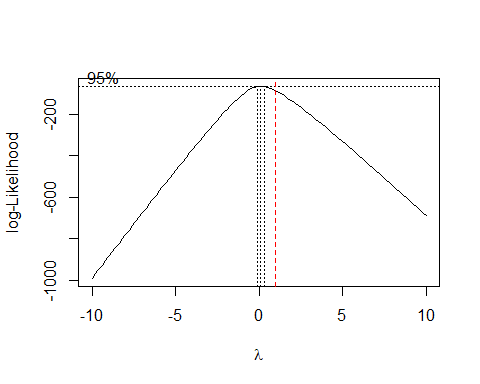
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "CADAV\_57"  
#> [1] "===================================="



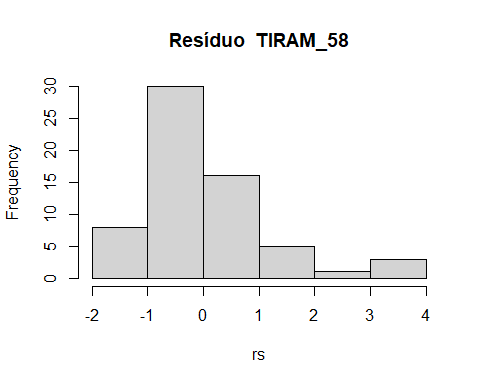
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.92458, p-value = 0.0008611



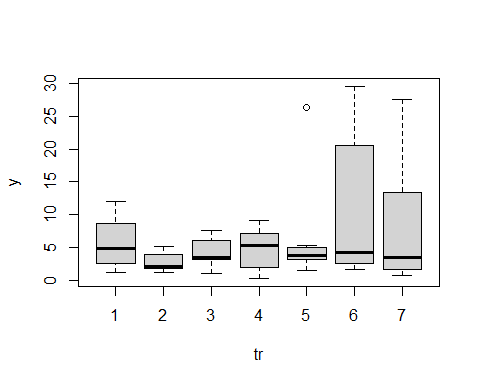
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 1.93  
#> 2 2 1.80  
#> 3 3 2.75  
#> 4 4 1.09  
#> 5 5 2.25  
#> 6 6 3.95  
#> 7 7 2.77



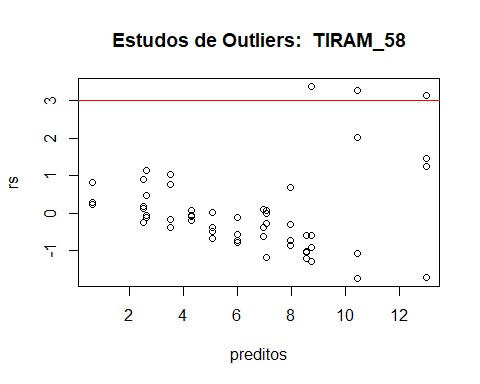
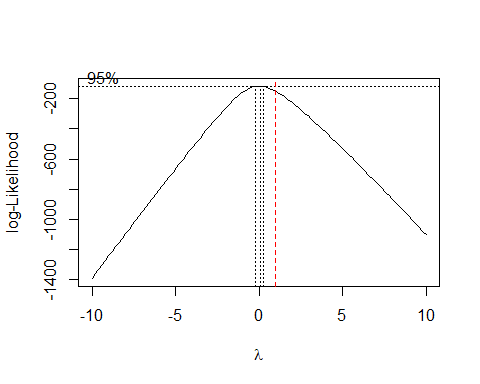
#> Outliers Trat Bloco Y  
#> 49 7 1 12.51207  
#>   
#> [1] "===================================="  
#> [1] "TIRAM\_58"  
#> [1] "===================================="



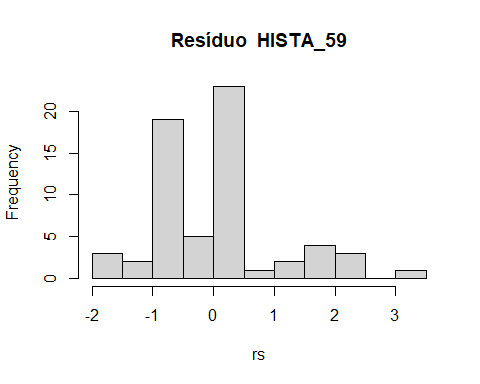
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.89192, p-value = 4.573e-05



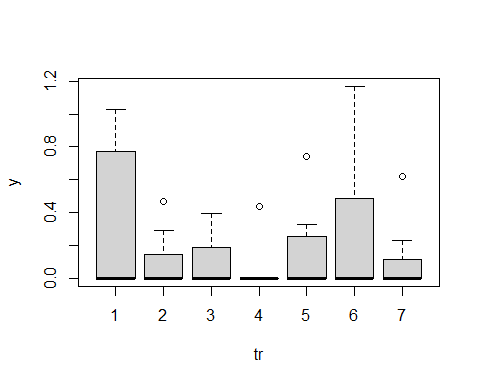
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 5.73  
#> 2 2 2.85  
#> 3 3 4.43  
#> 4 4 4.85  
#> 5 5 6.51  
#> 6 6 10.8   
#> 7 7 8.22



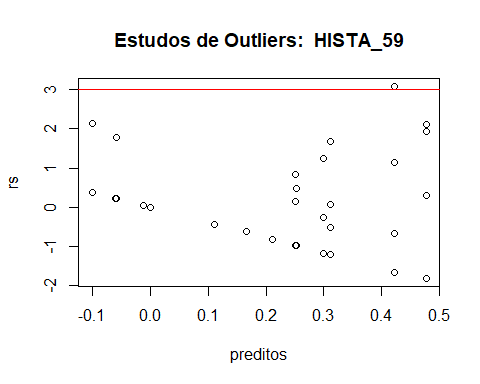
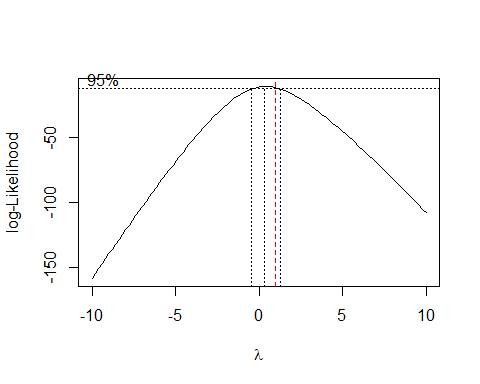
#> Outliers Trat Bloco Y  
#> 35 5 1 26.28803  
#> 42 6 1 29.51939  
#> 49 7 1 27.58869  
#>   
#> [1] "===================================="  
#> [1] "HISTA\_59"  
#> [1] "===================================="



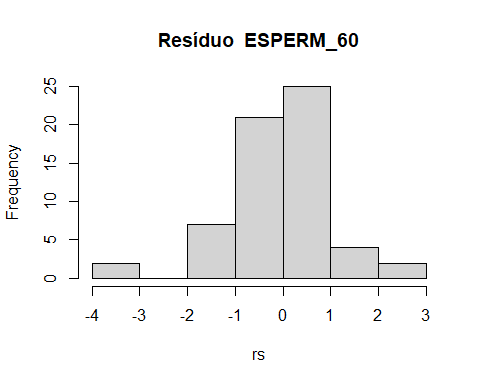
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.92634, p-value = 0.001024



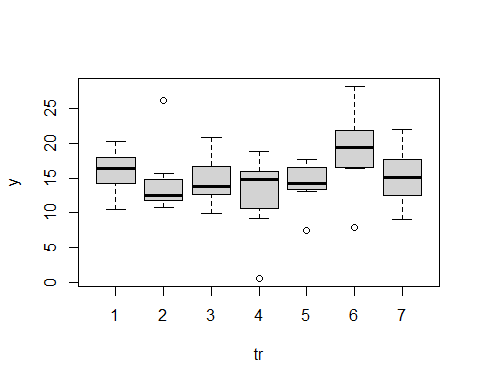
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 0.321   
#> 2 2 0.0953  
#> 3 3 0.0968  
#> 4 4 0.0550  
#> 5 5 0.156   
#> 6 6 0.267   
#> 7 7 0.121



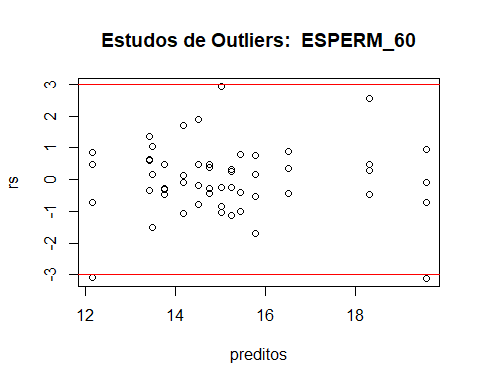
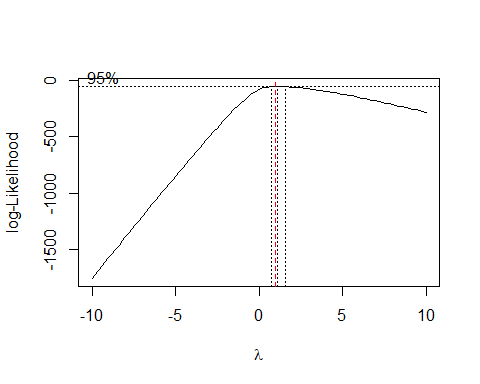
#> Outliers Trat Bloco Y  
#> 47 6 2 1.168114  
#>   
#> [1] "===================================="  
#> [1] "ESPERM\_60"  
#> [1] "===================================="



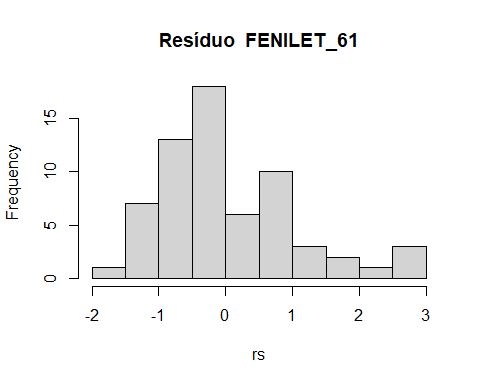
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.95222, p-value = 0.0184



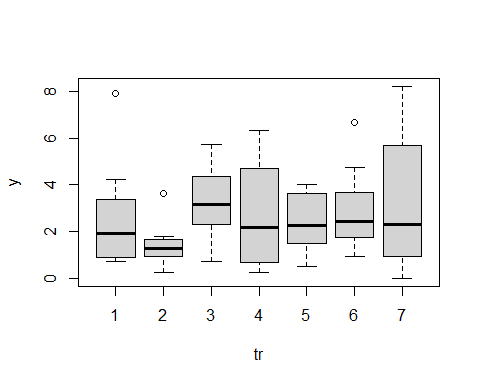
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 15.8  
#> 2 2 14.4  
#> 3 3 14.7  
#> 4 4 12.8  
#> 5 5 14.1  
#> 6 6 18.9  
#> 7 7 15.2



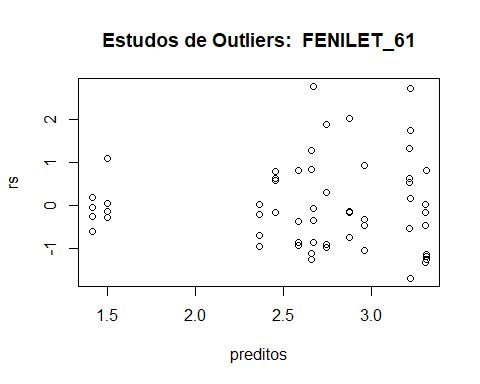
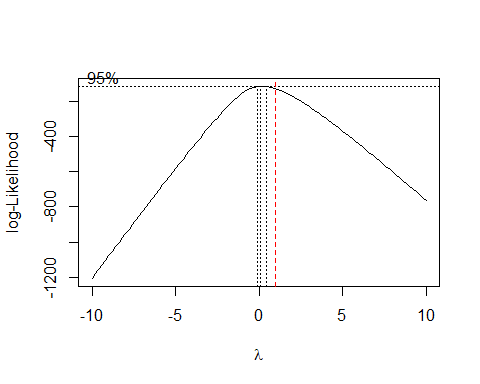
#> Outliers Trat Bloco Y  
#> 23 4 1 0.6213821  
#> 46 6 2 7.9147948  
#>   
#> [1] "===================================="  
#> [1] "FENILET\_61"  
#> [1] "===================================="



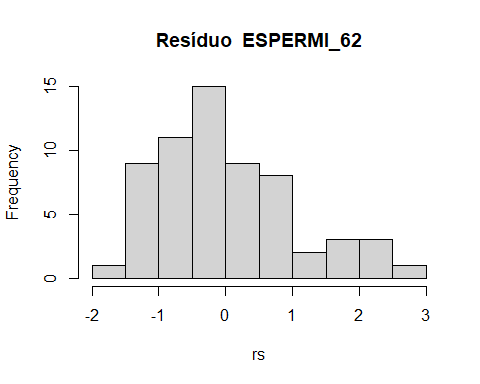
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.92406, p-value = 0.0007374



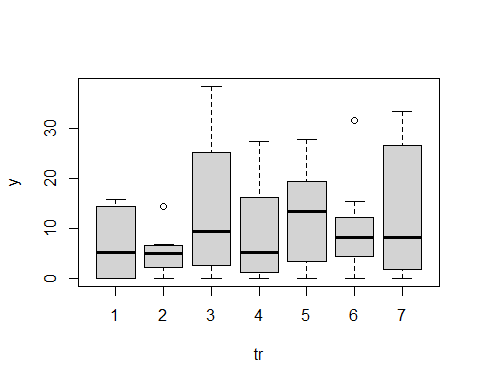
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 2.63  
#> 2 2 1.46  
#> 3 3 3.26  
#> 4 4 2.70  
#> 5 5 2.41  
#> 6 6 2.92  
#> 7 7 3.27



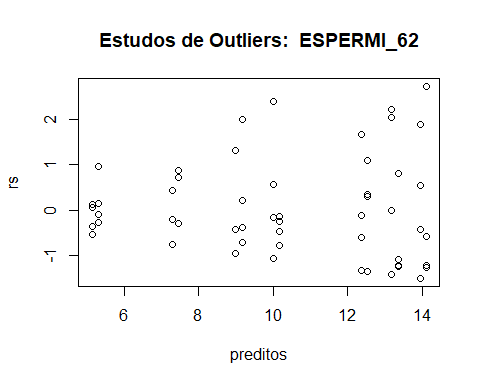
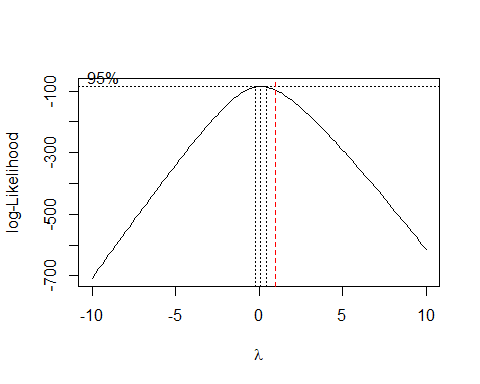
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "ESPERMI\_62"  
#> [1] "===================================="



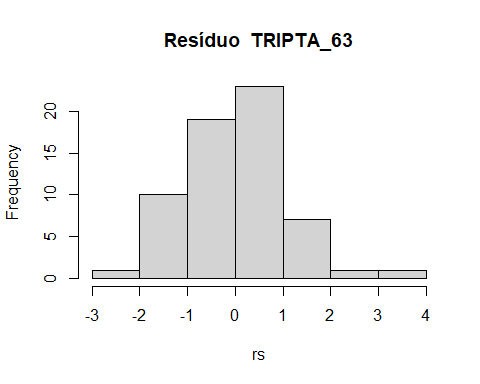
#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.93792, p-value = 0.003646



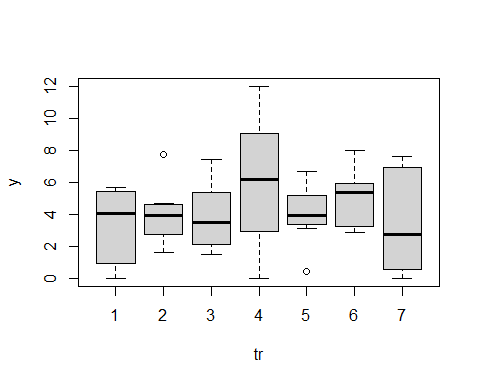
#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 7.36  
#> 2 2 5.22  
#> 3 3 14.0   
#> 4 4 9.08  
#> 5 5 12.4   
#> 6 6 10.1   
#> 7 7 13.3



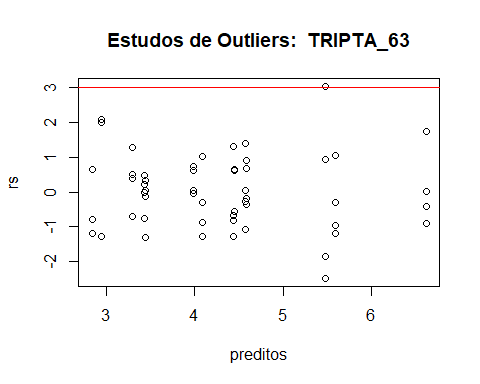
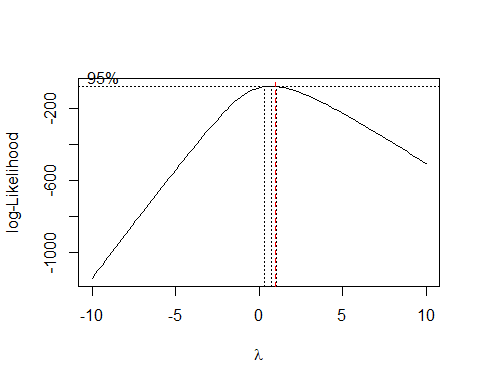
#> Outliers[1] Trat Bloco Y   
#> <0 linhas> (ou row.names de comprimento 0)  
#>   
#> [1] "===================================="  
#> [1] "TRIPTA\_63"  
#> [1] "===================================="



#>   
#> Shapiro-Wilk normality test  
#>   
#> data: rs  
#> W = 0.98308, p-value = 0.5494



#> Médias por tratamentos  
#> # A tibble: 7 x 2  
#> tr media  
#> <fct> <dbl>  
#> 1 1 3.49  
#> 2 2 4.00  
#> 3 3 3.86  
#> 4 4 6.05  
#> 5 5 4.01  
#> 6 6 5.02  
#> 7 7 3.52



#> Outliers Trat Bloco Y  
#> 26 4 1 12.01582

## Análise de variância

# for(i in seq\_along(variaveis)){  
# nome <- names(variaveis[i])  
# print("====================================")  
# print(str\_to\_upper(nome))  
# print("====================================")  
#   
# daux <- variaveis[i]  
# names(daux) <- "y" ### <-  
# y <- daux$y ### <-   
# ingred <- dados$ingrediente %>% as\_factor()  
# dose <- dados$dose  
# bl <- dados$bl %>% as\_factor()  
#   
# fat2.dbc(ingred,dose,bl,y,quali=c(TRUE,FALSE),  
# fac.names = c("Ingred.","Dose"))  
#   
# my\_plot<-tibble(dose,ingred,y) %>%  
# group\_by(dose,ingred) %>%   
# summarise(y=mean(y)) %>%   
# ggplot(aes(x=dose,y=y,color=ingred)) +  
# geom\_point(size=3) +  
# geom\_line() +  
# #facet\_wrap(~ingred,nrow = 2)+  
# theme\_bw() +  
# labs(y=str\_to\_upper(nome),x="DOSE",color="Ingrediente")+  
# theme(legend.position = "top")+  
# scale\_color\_manual(values = c("aquamarine4","red"))   
# print(my\_plot)  
# cat("\n")  
# }