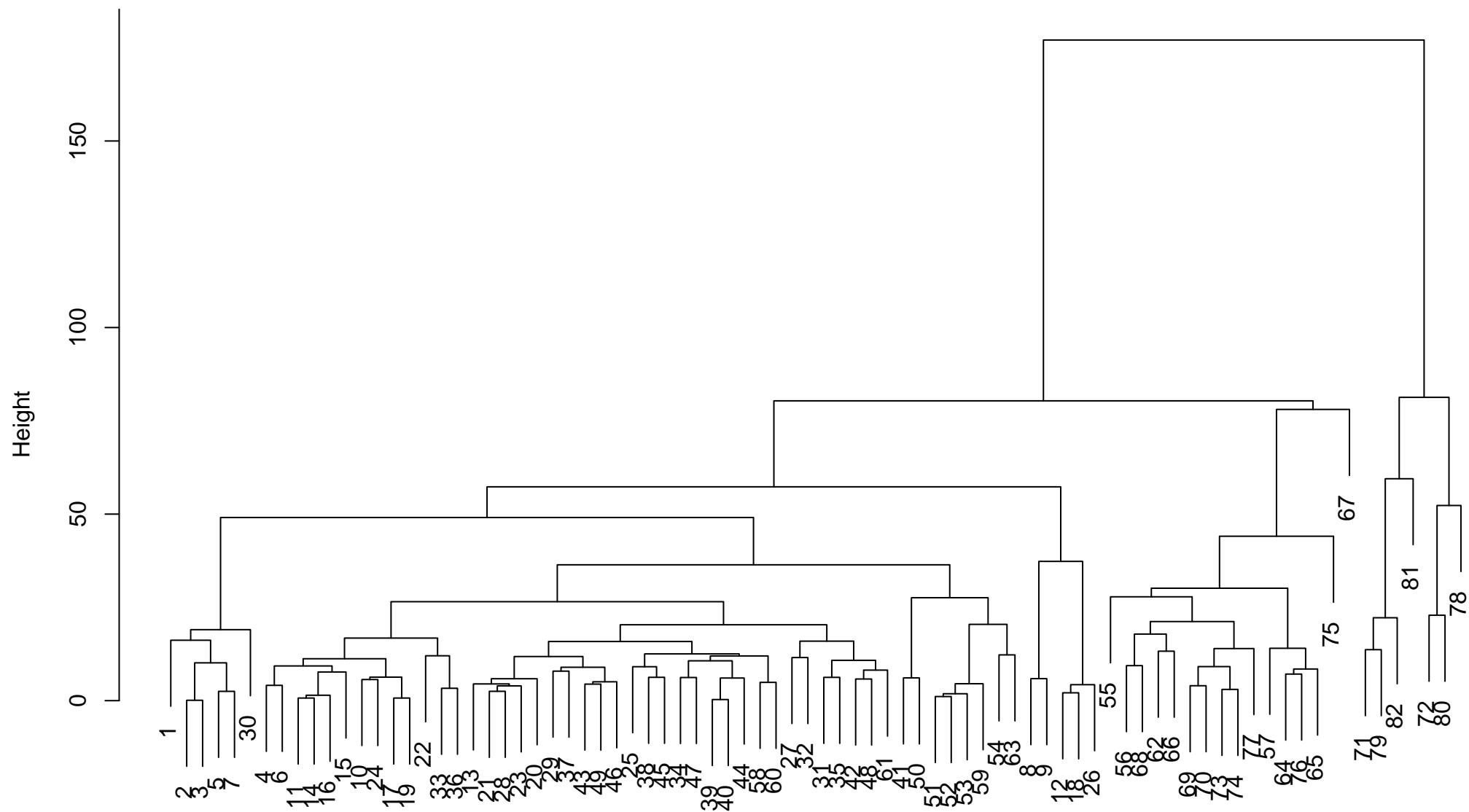
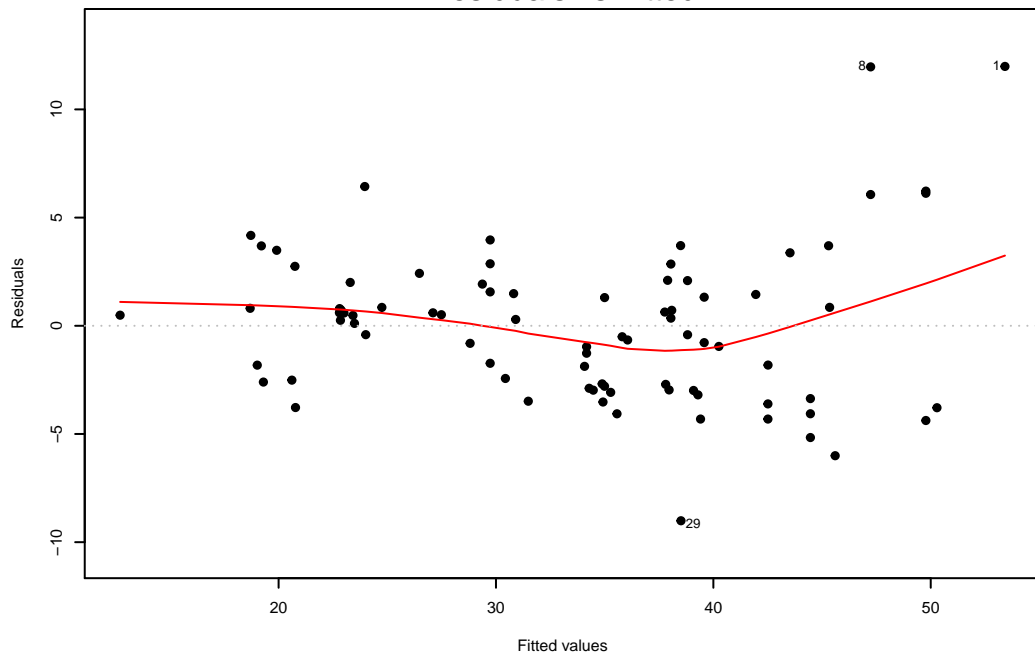


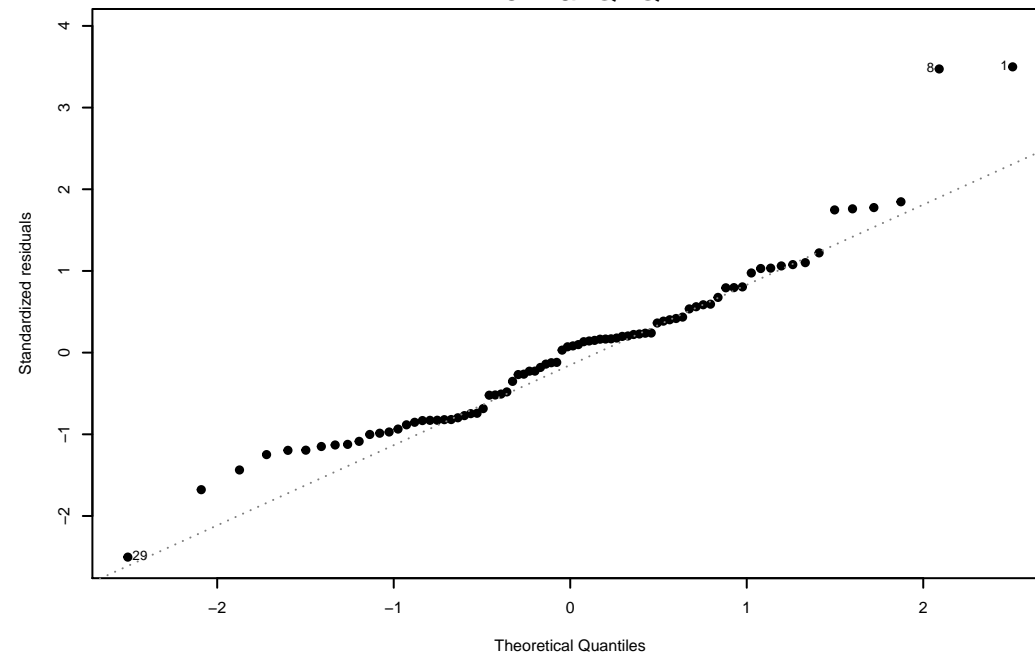
Dendrogram of `cluster::agnes(x = data[, c(covariates, response)])`



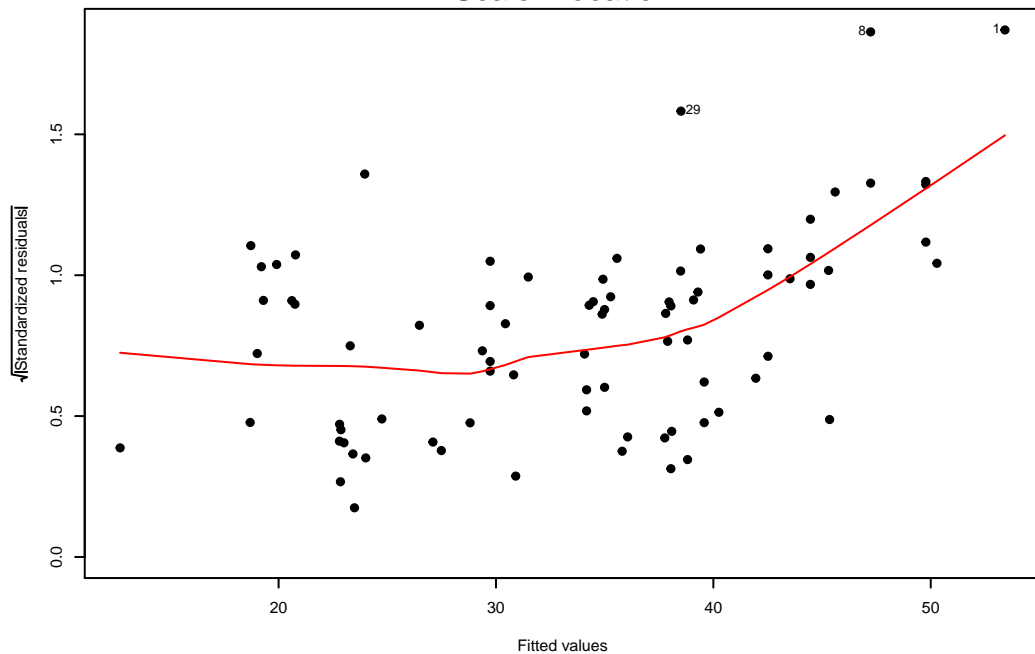
Residuals vs Fitted



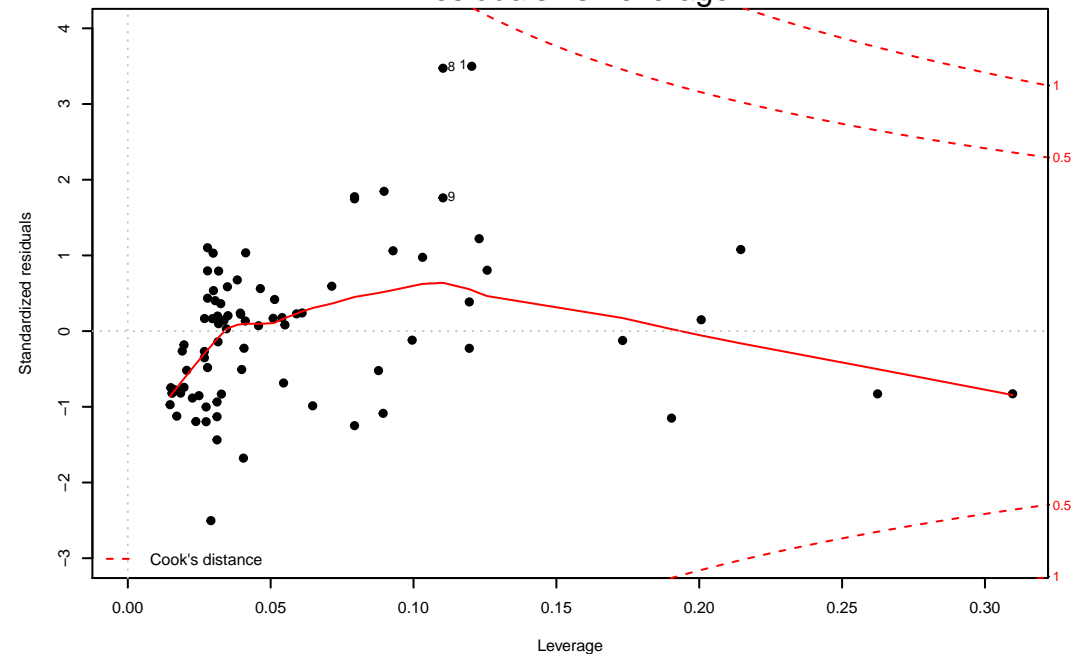
Normal Q-Q



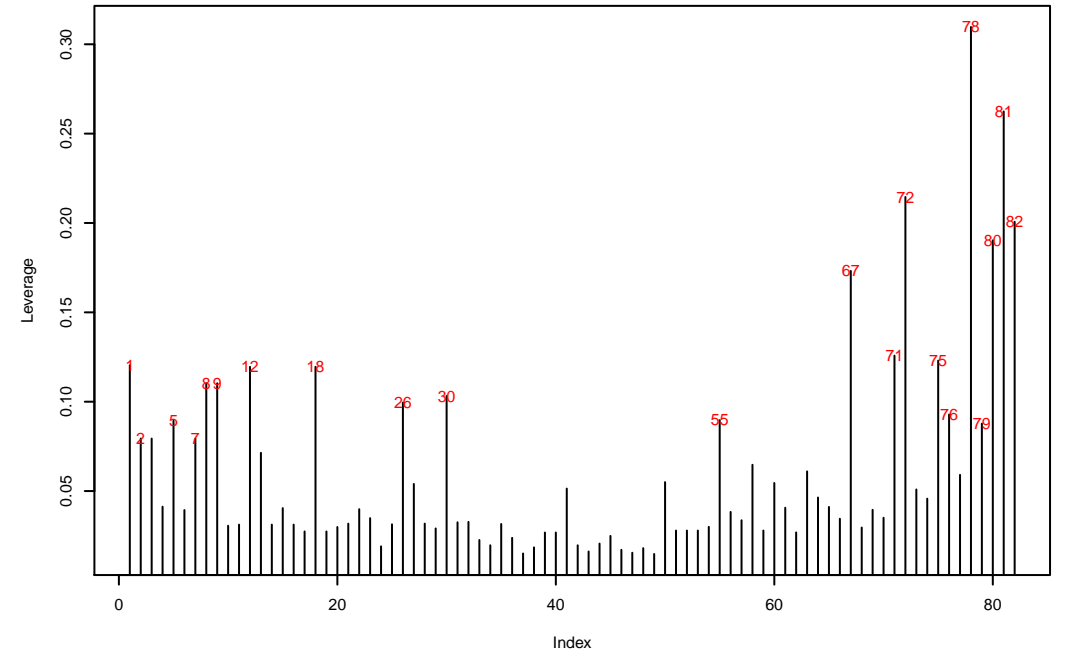
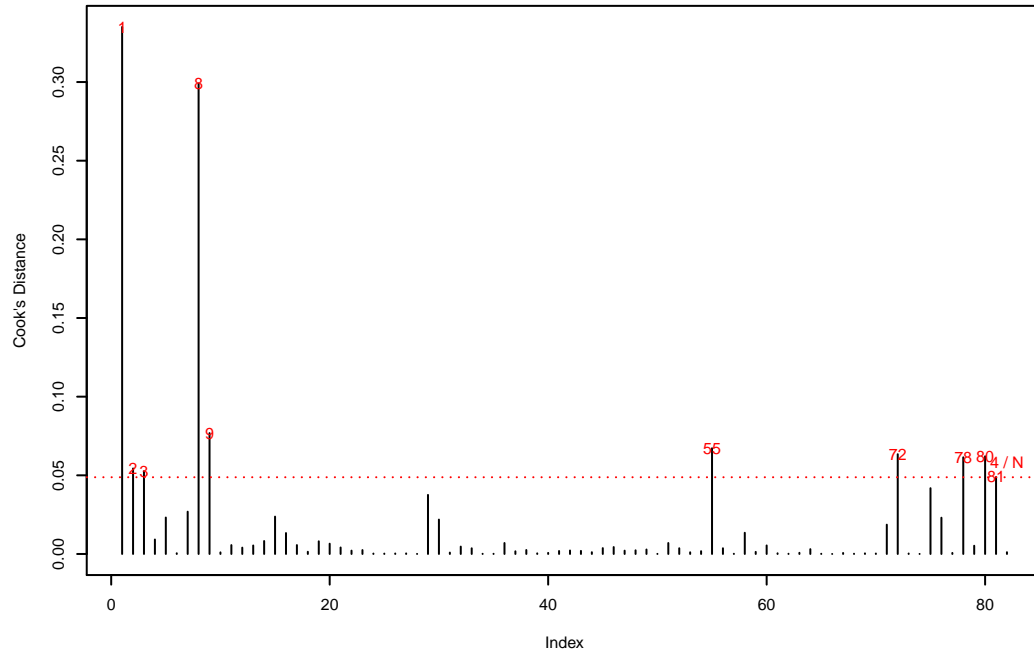
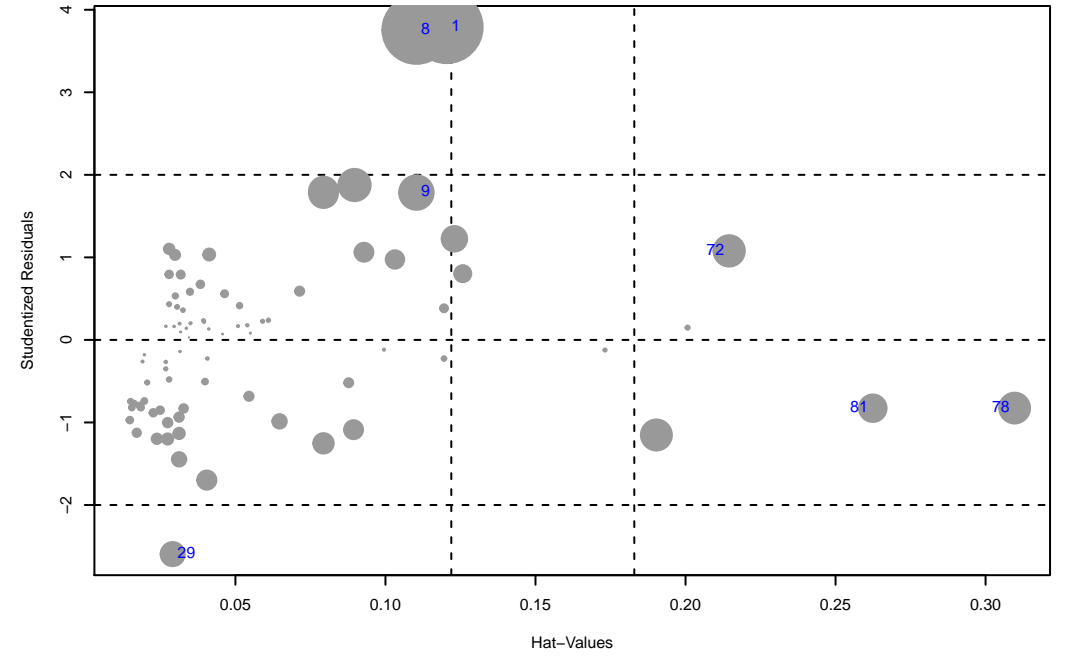
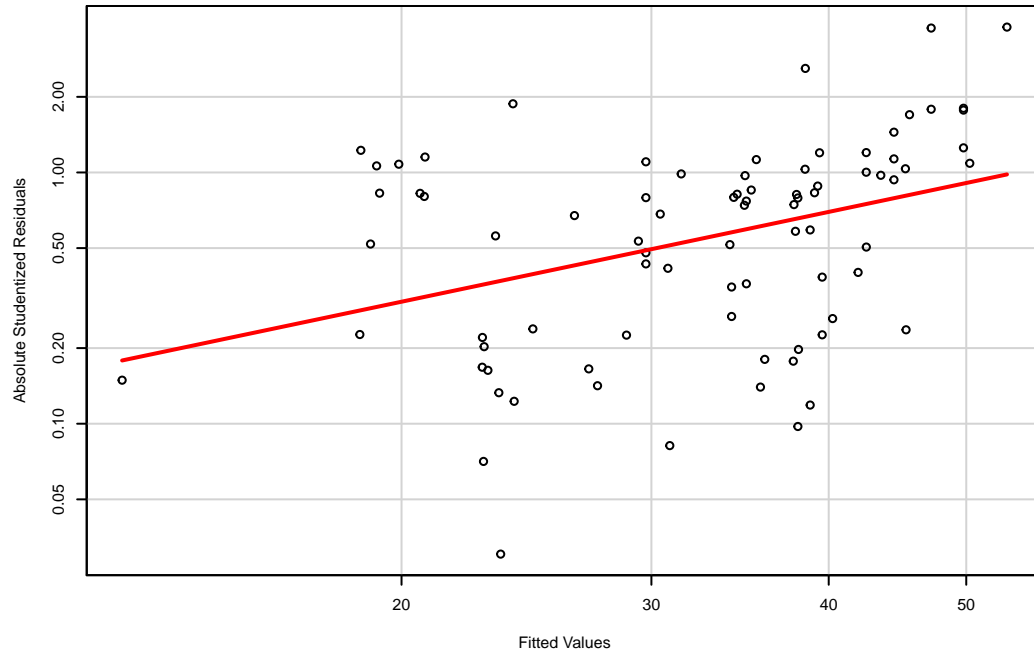
Scale-Location

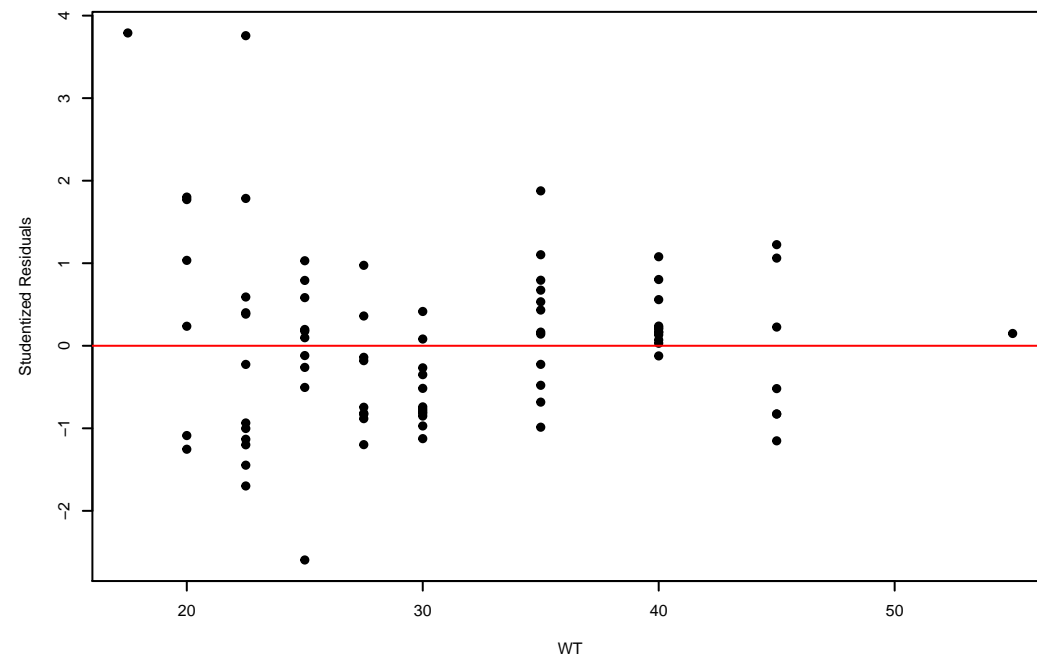
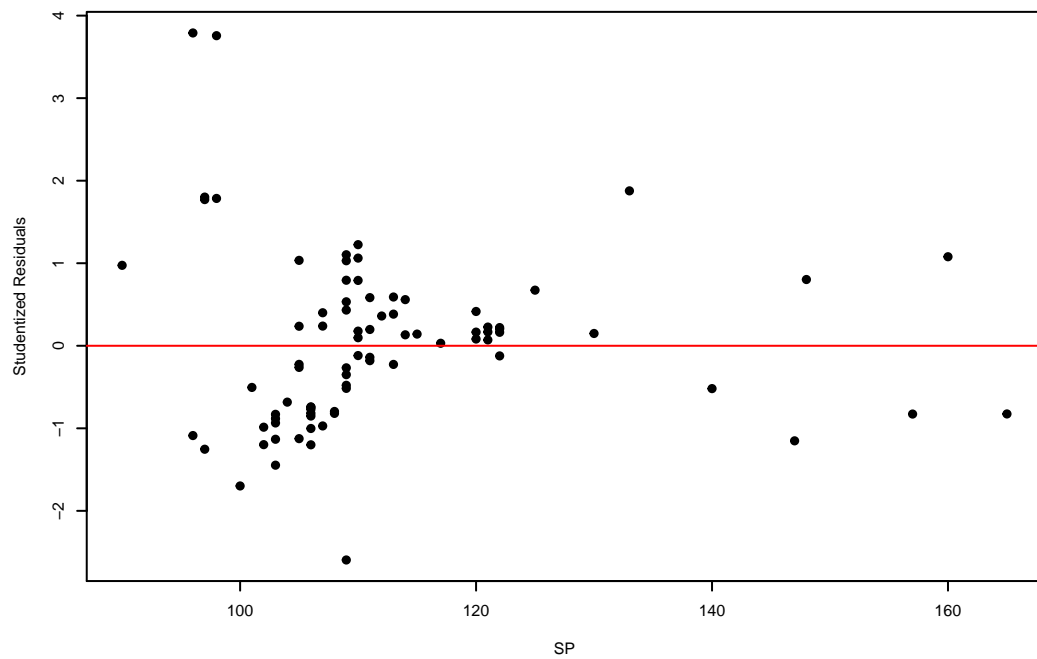
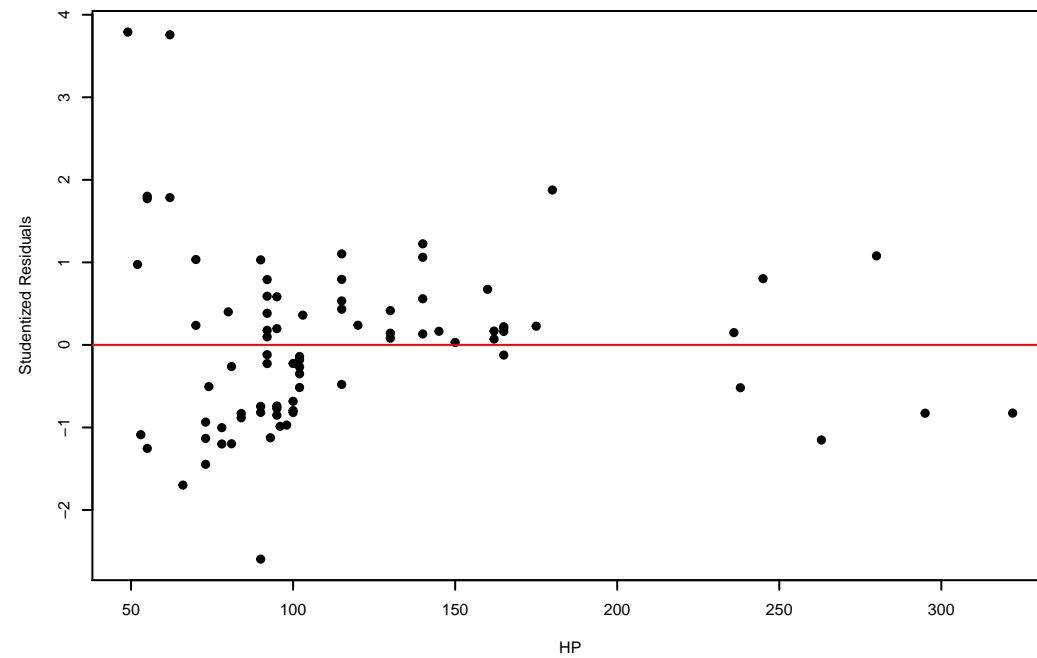
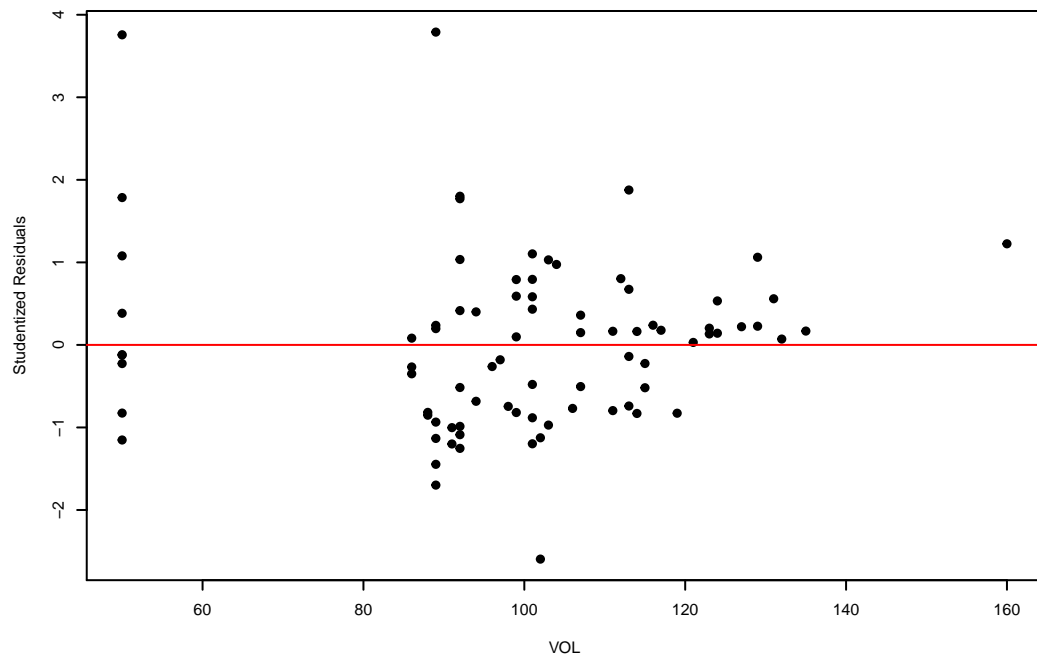


Residuals vs Leverage

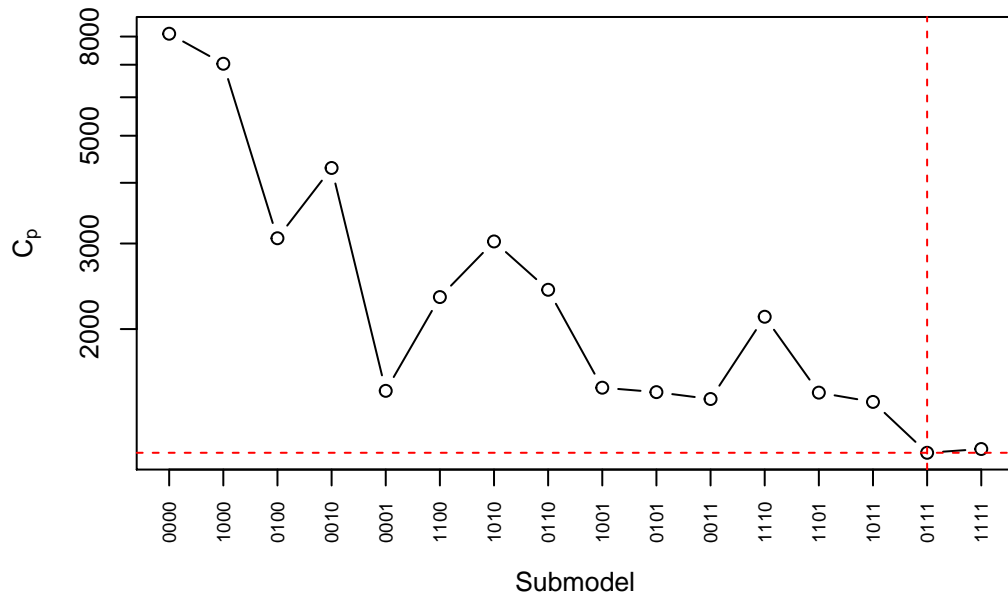


Spread–Level Plot for fit

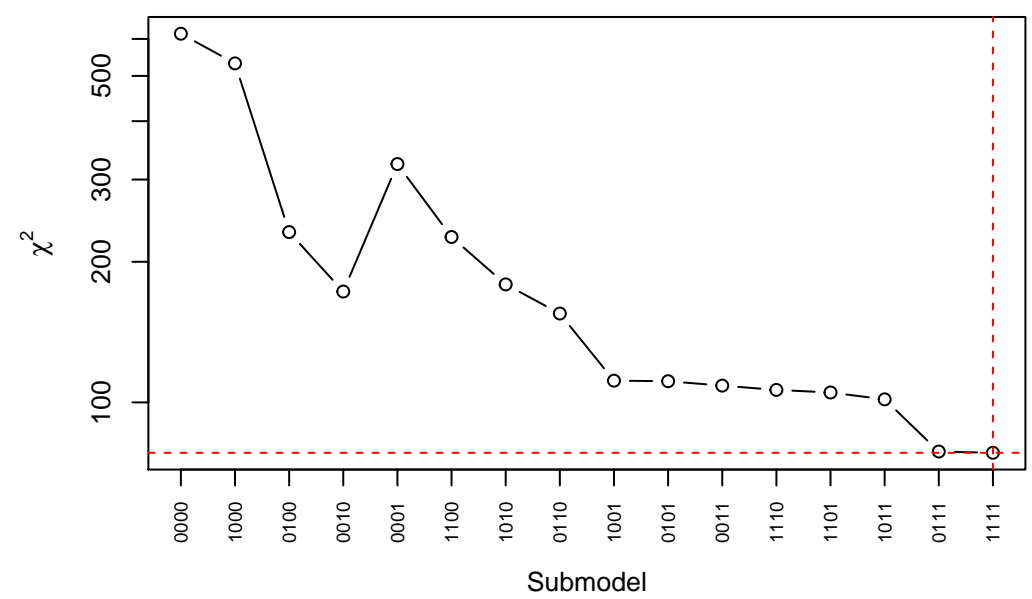




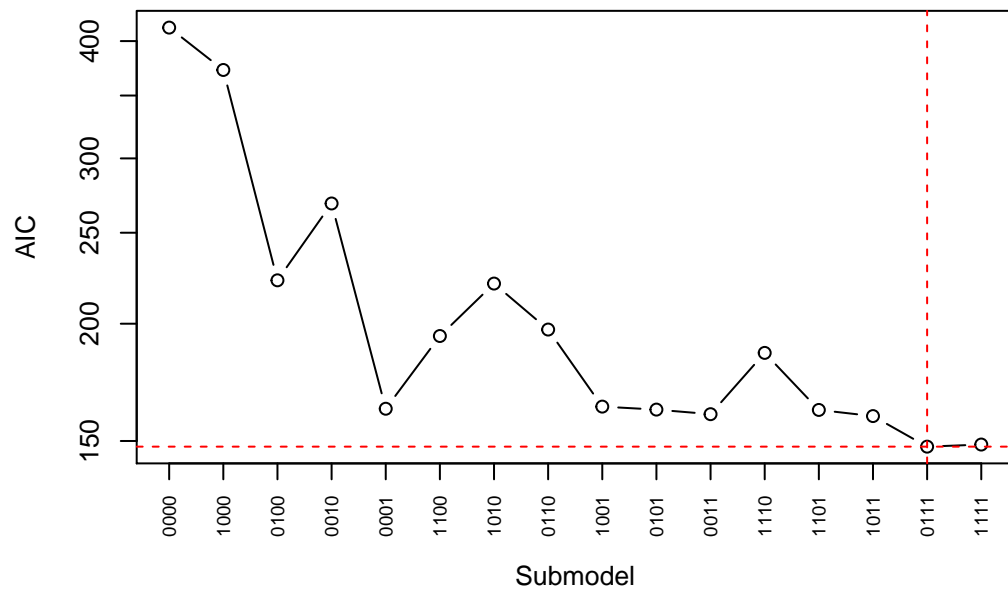
**(Linear) Model Selection (Cp)**



**(Linear) Model Selection (Chi-Squared)**



**(Linear) Model Selection (AIC)**



**(Linear) Model Selection (BIC)**

