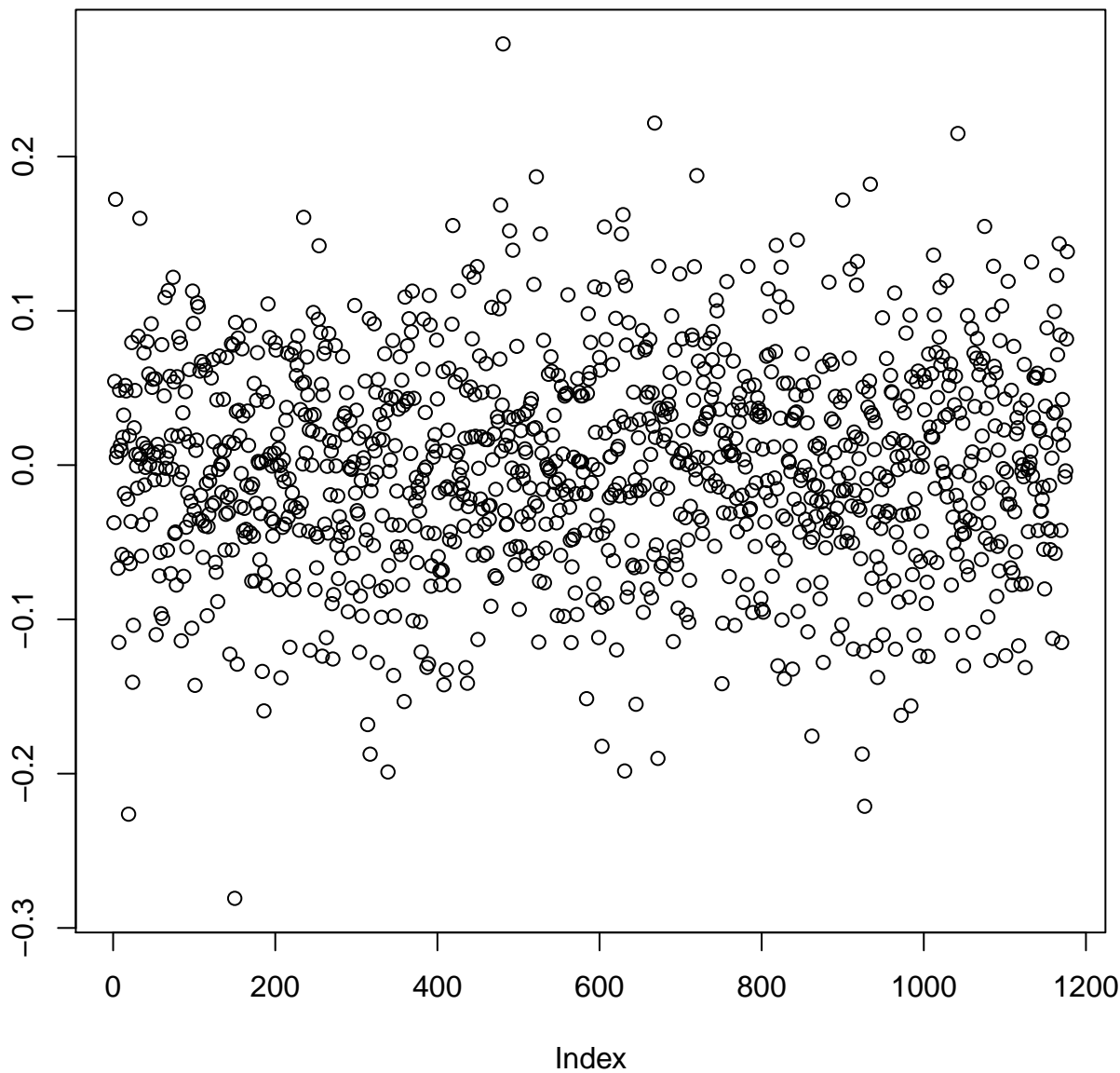
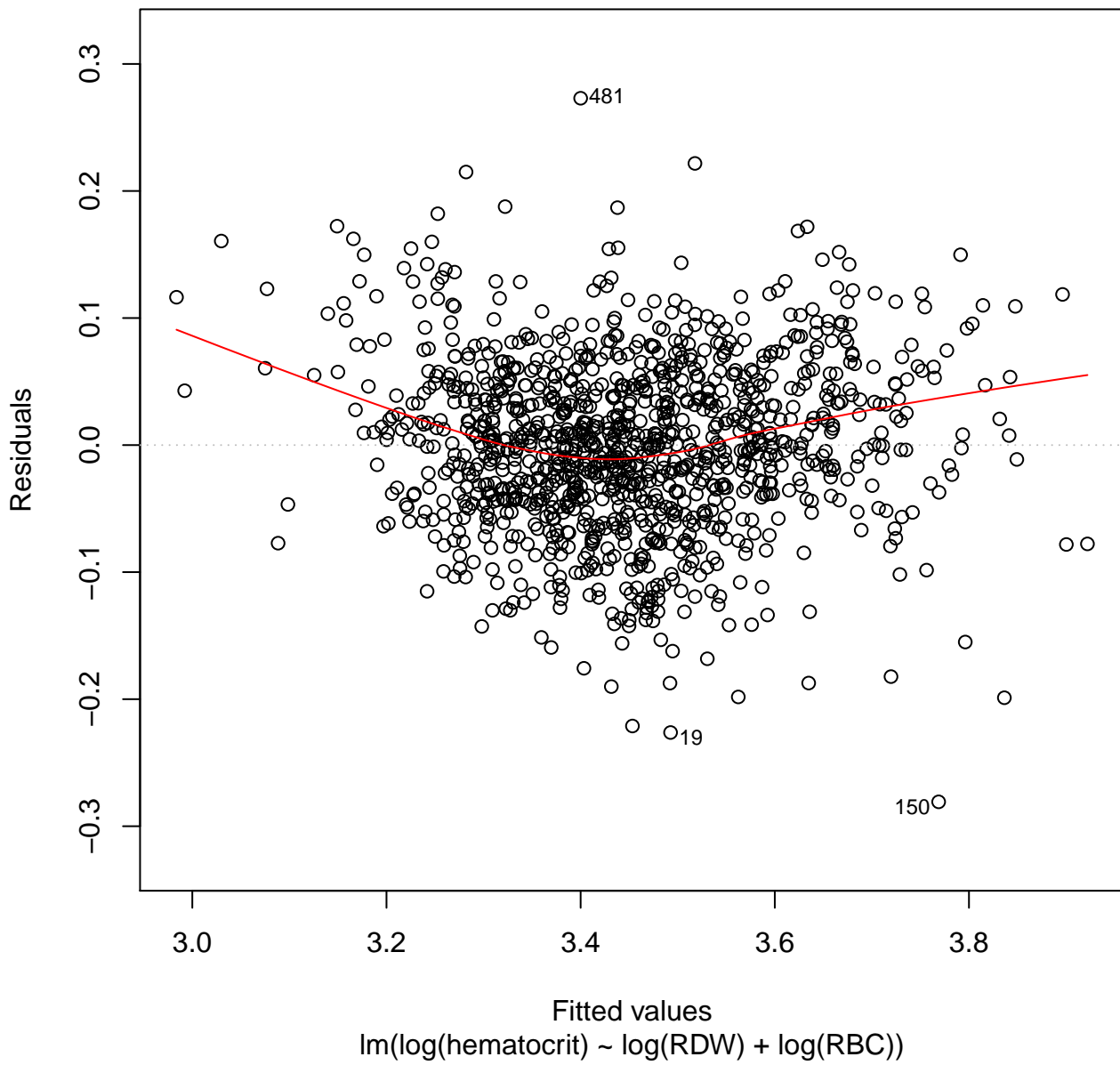


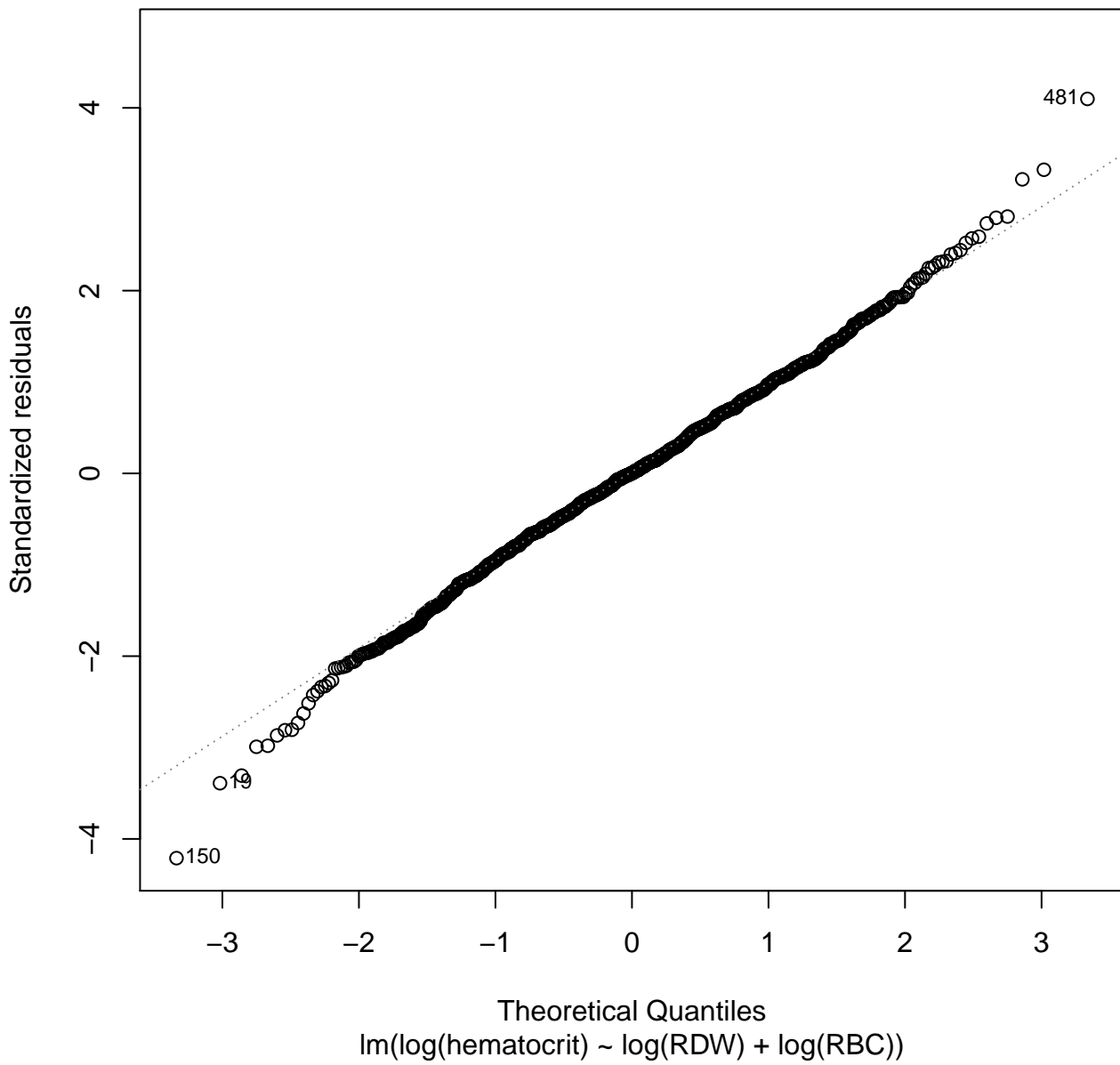
residuals(lm(log(hematocrit) ~ log(RDW) + log(RBC), data = dataset\_mortalidade))



Residuals vs Fitted

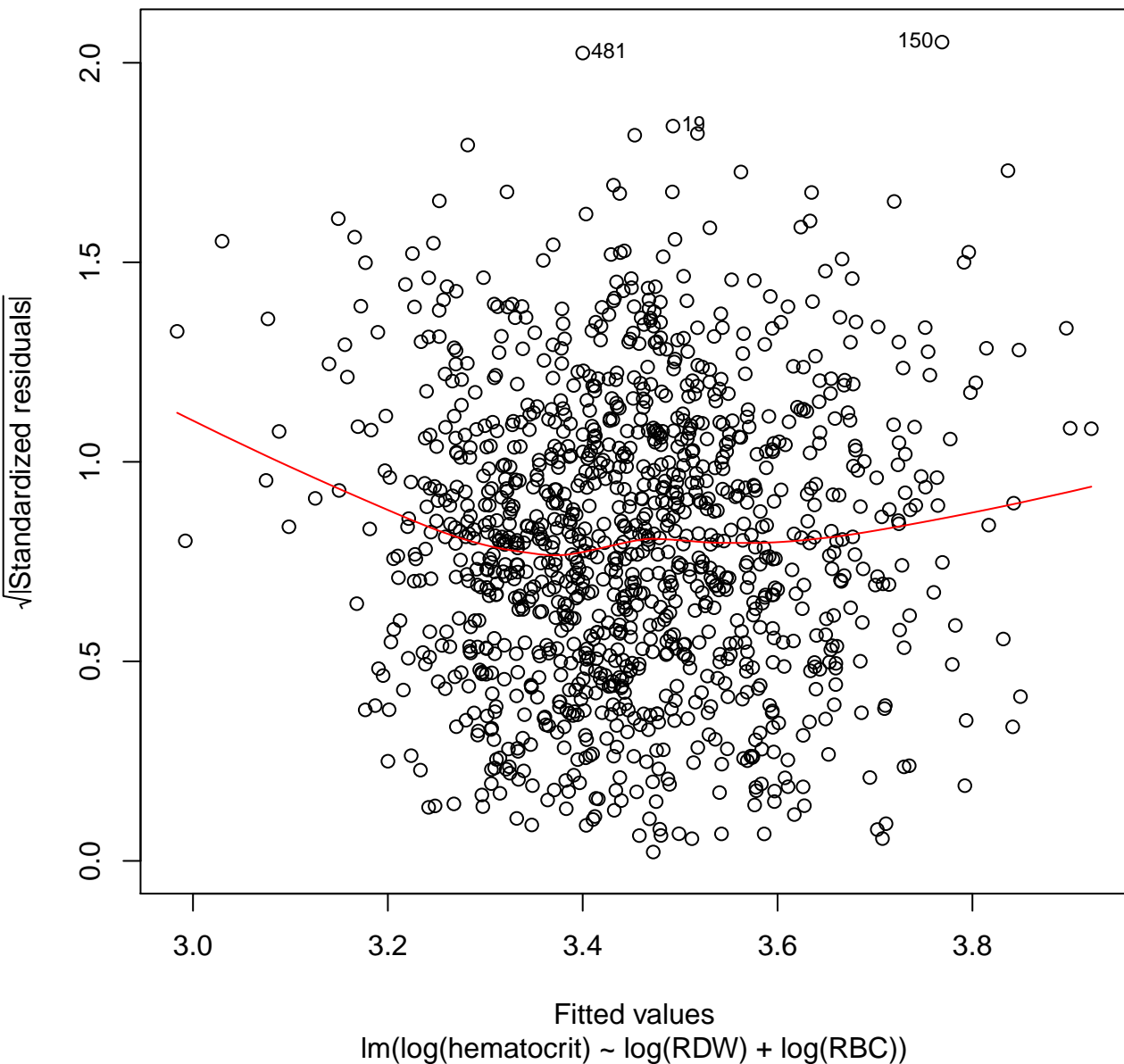


Normal Q-Q





# Scale-Location



Residuals vs Leverage

