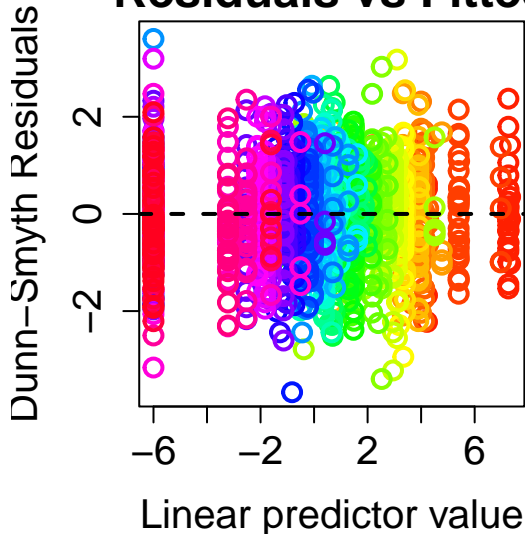
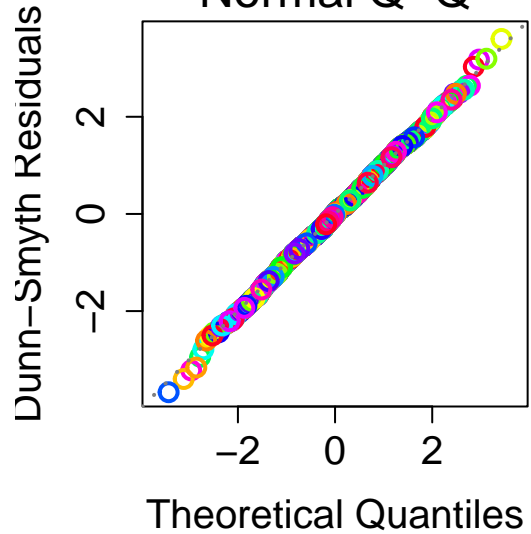


manyglm(Y ~ soi_phase ...)

Residuals vs Fitted



Normal Q-Q



Scale-Location

