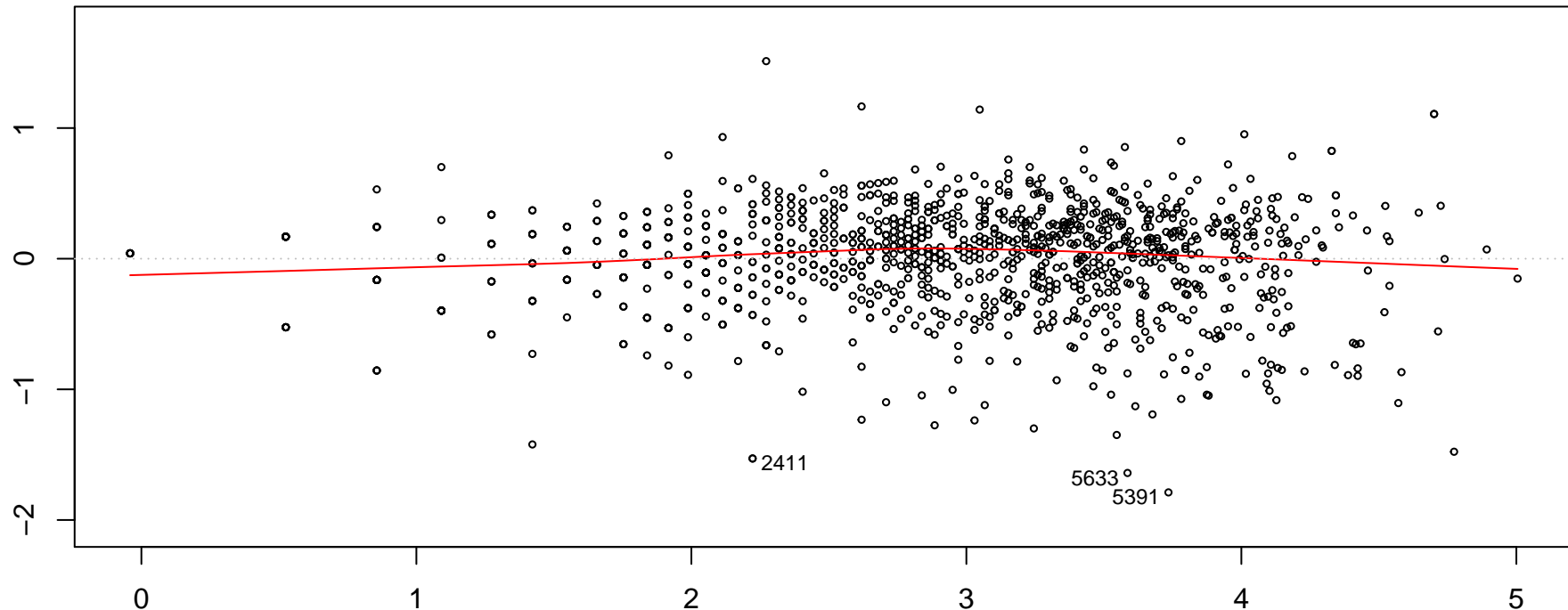


# Residuals plot, log-log LM

Residuals vs Fitted

Residuals



Fitted values  
lm(log(fams) ~ log(juvs))