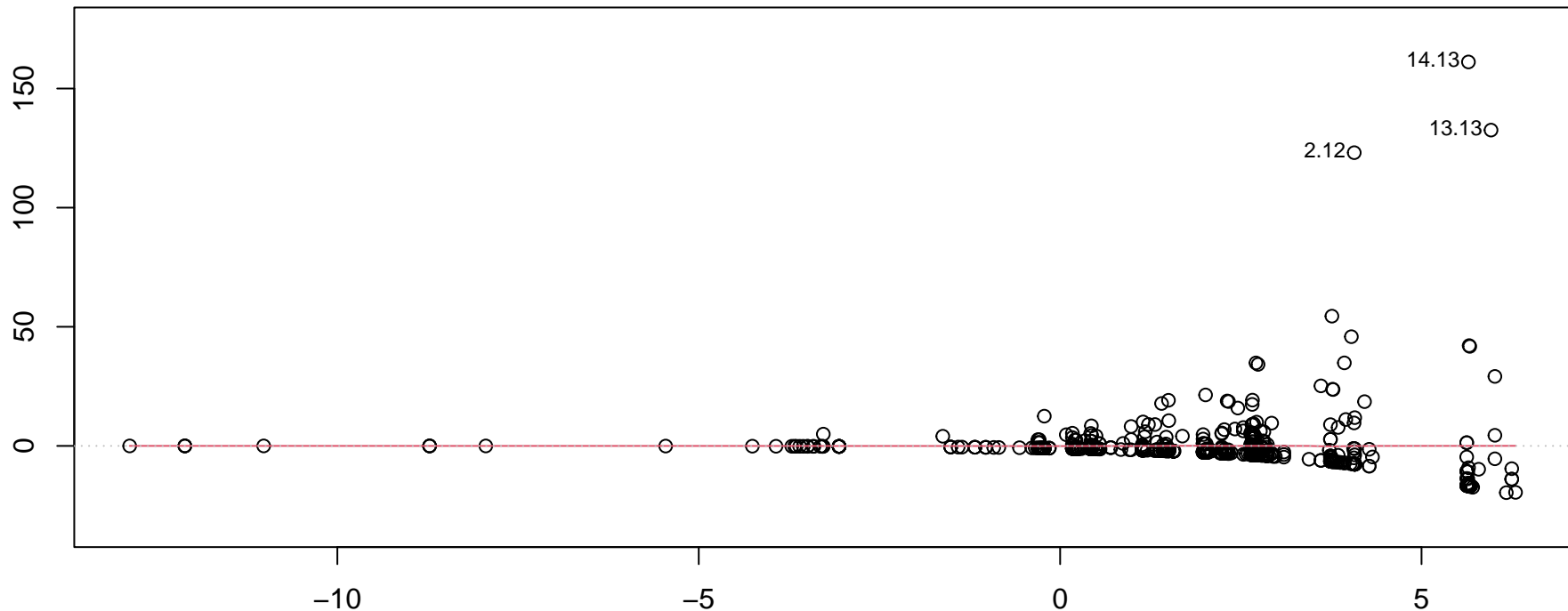


Pearson Residuals

Residuals vs Fitted



Predicted values  
glm(Count ~ 0 + Species + NO3 + NO3:Species)