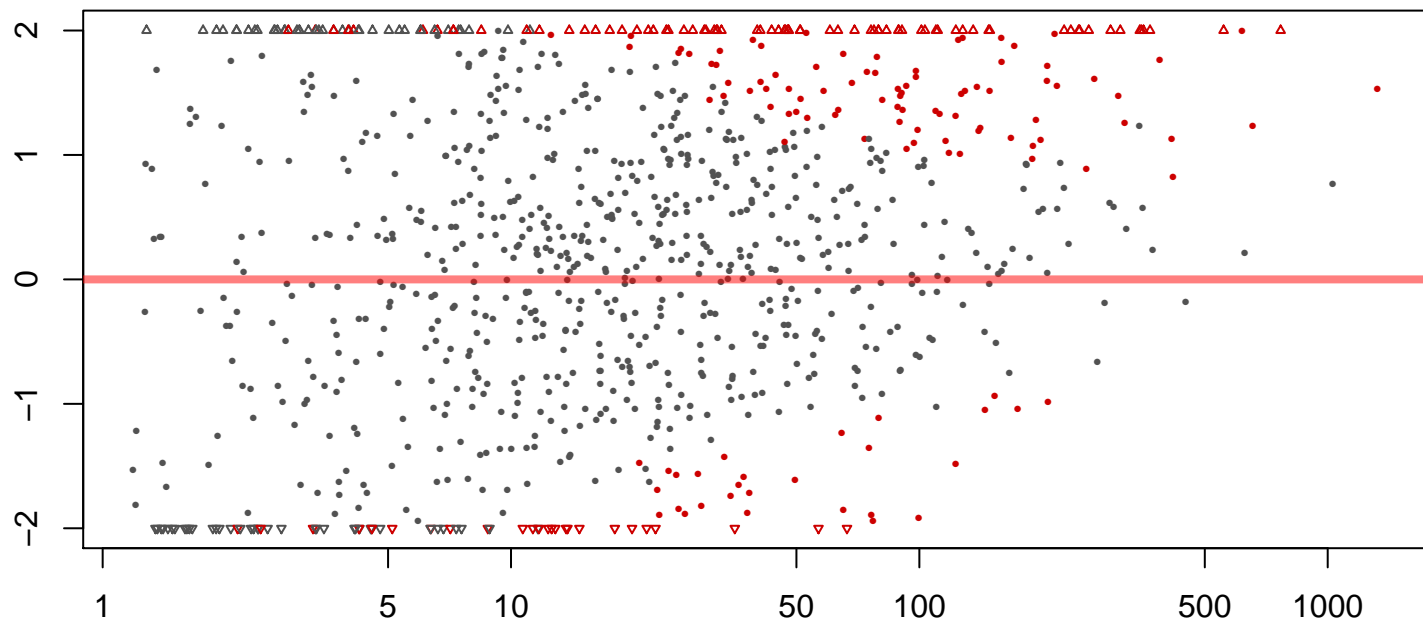# LFC over mean norm counts on: example data

~genotype + condition + genotype:condition log2 fold change (MLE): condition_B_vs_A effect

No shrinkage

Shrinkage