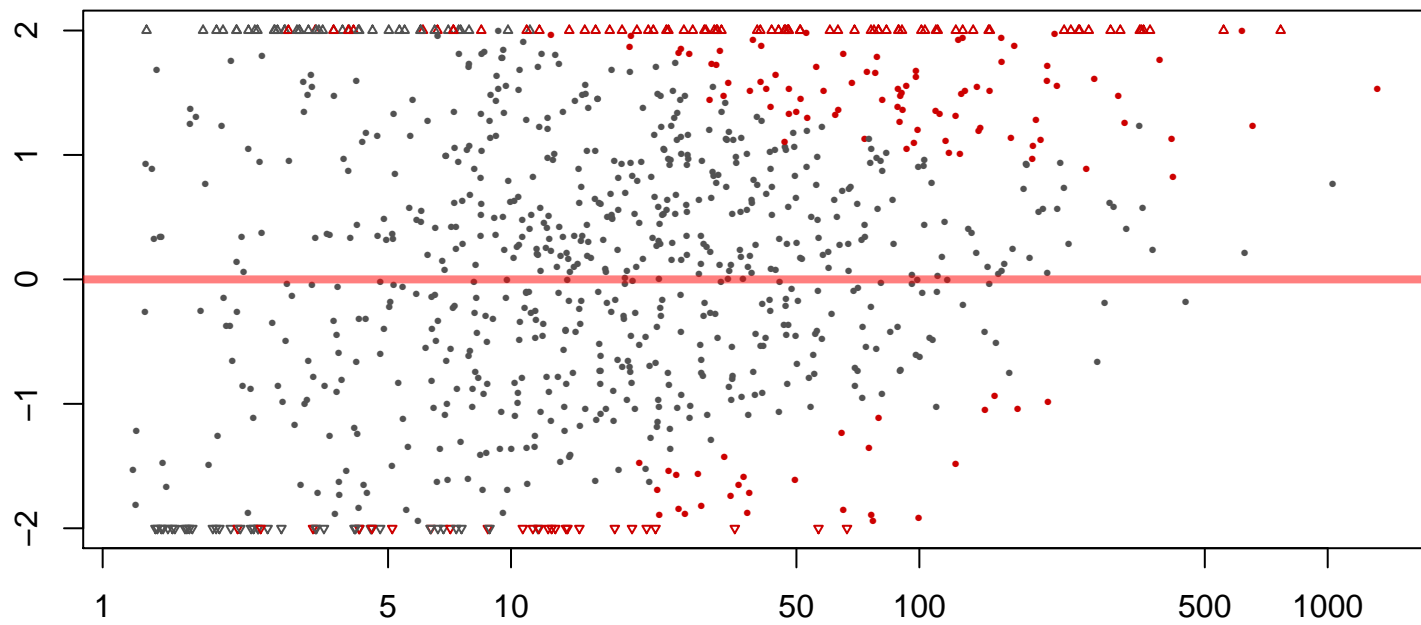


LFC over mean norm counts on: example data

~genotype + genotype:condition log2 fold change (MLE): genotypeI.conditionB effect

No shrinkage



Shrinkage

