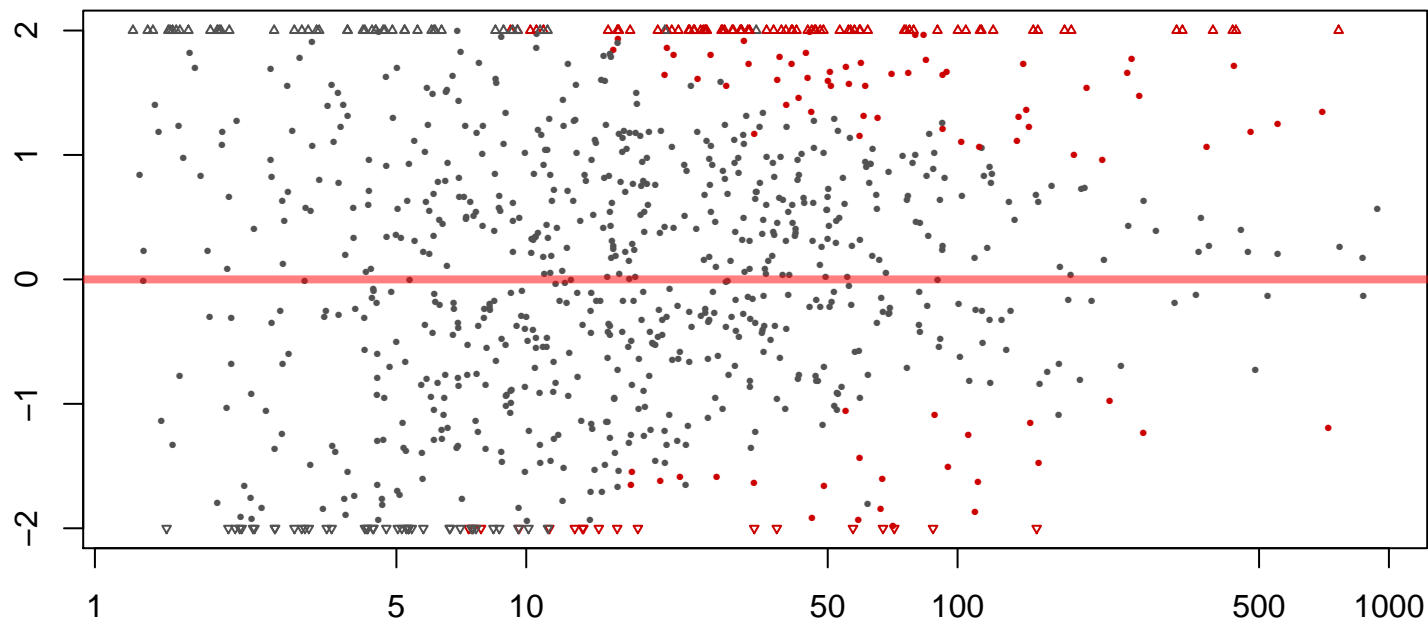


# LFC over mean norm counts on: example data

~genotype + condition + genotype:condition log2 fold change (MLE): condition\_B\_vs\_A effect

## No shrinkage



## Shrinkage

