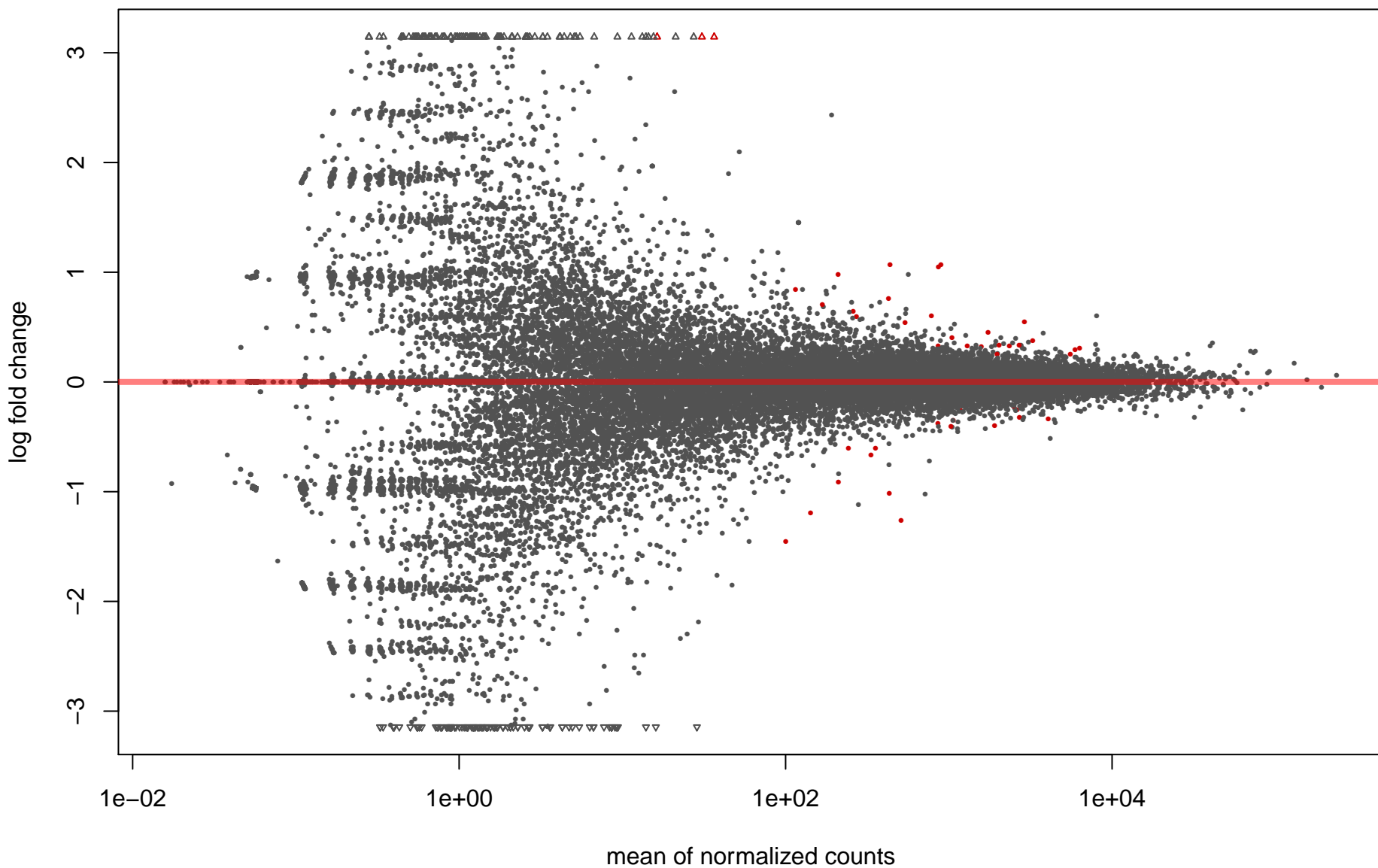
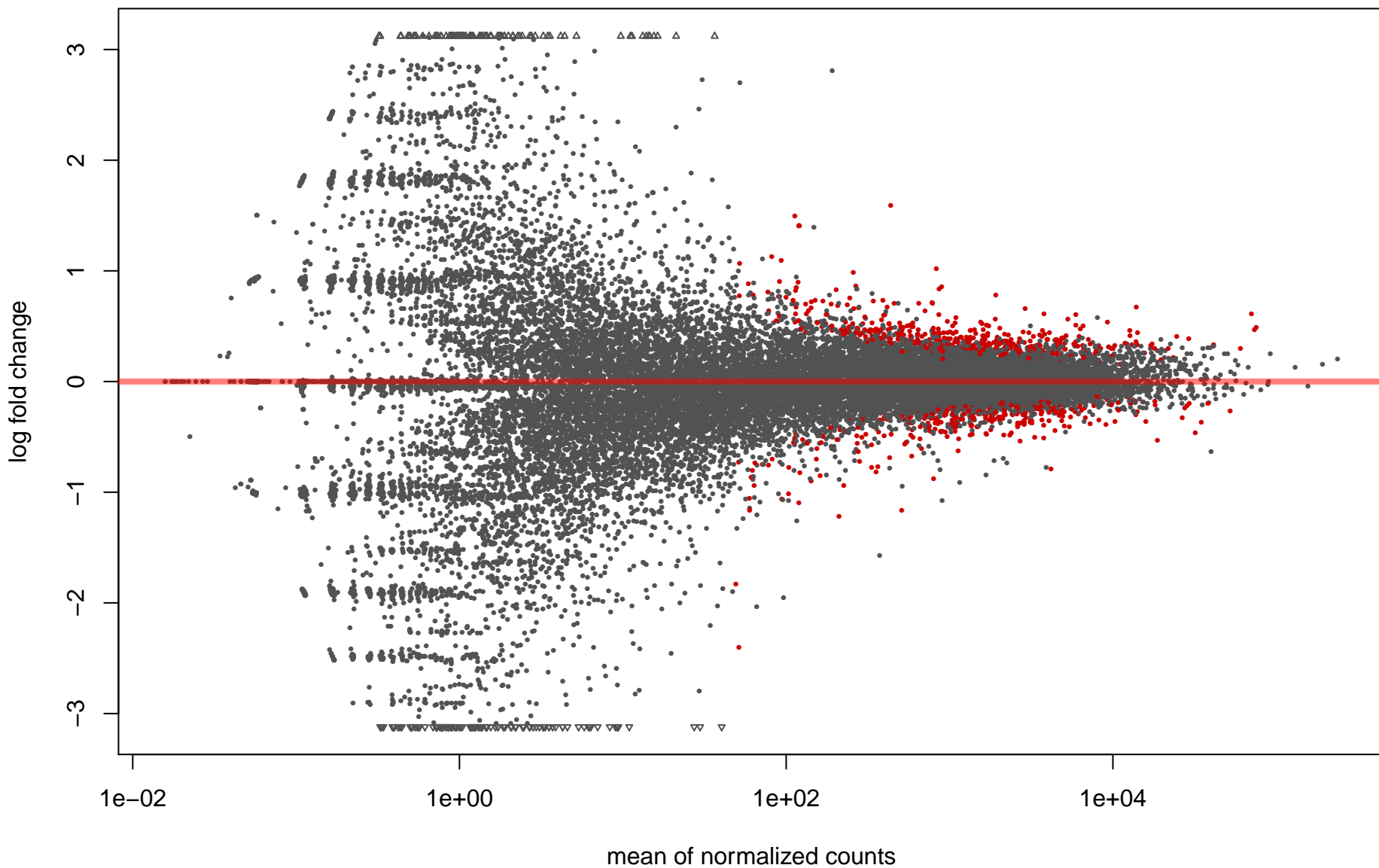


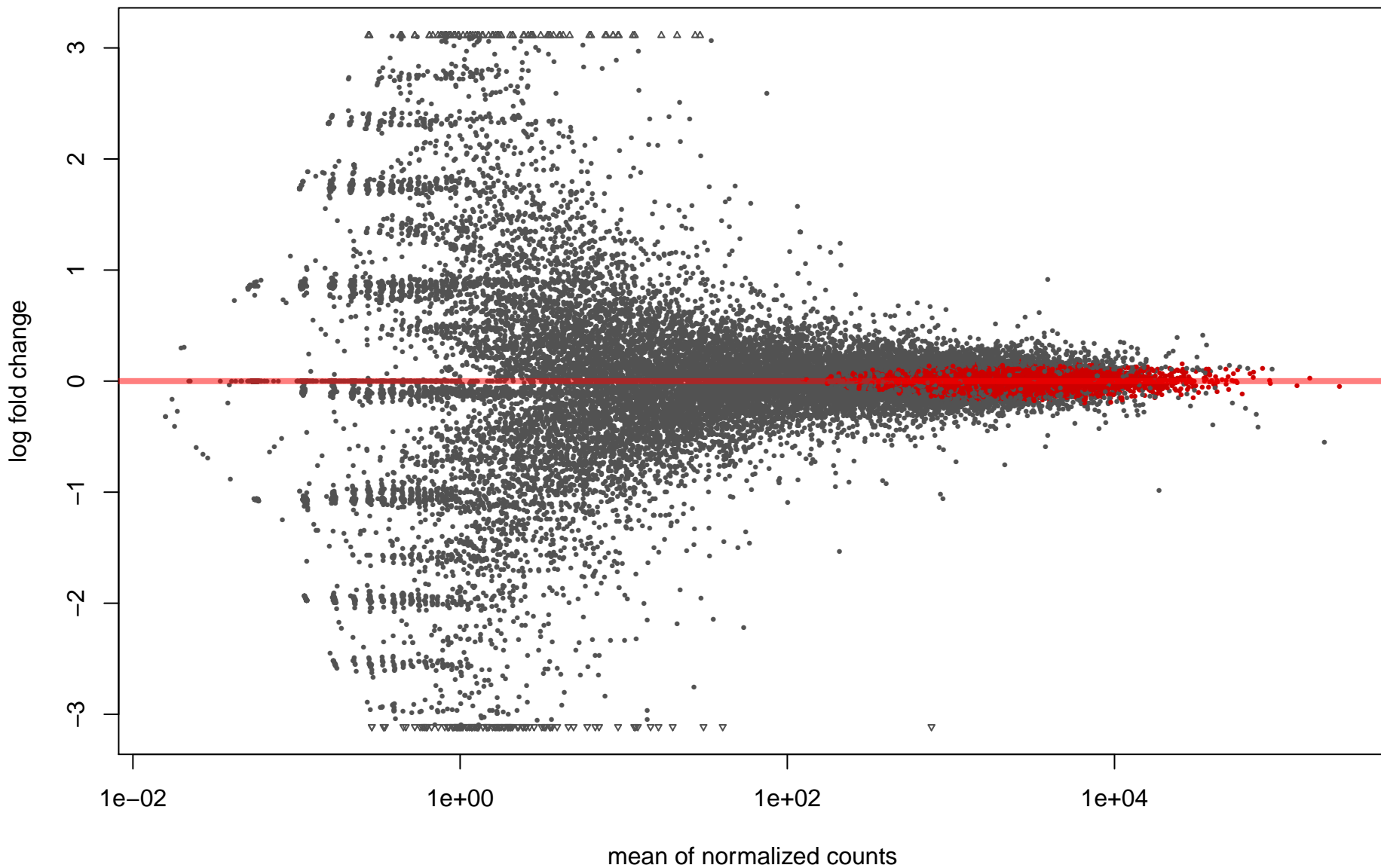
res48h



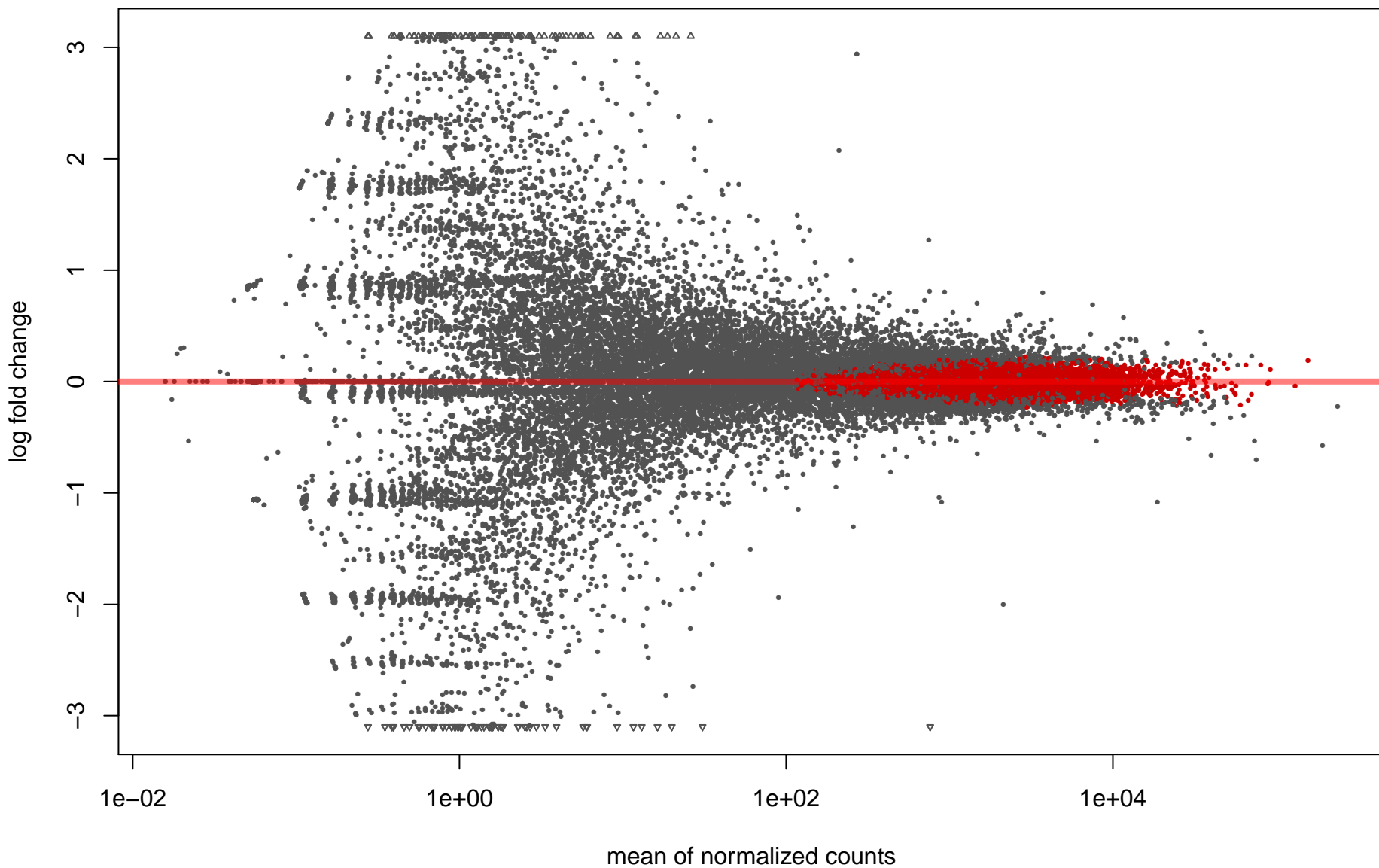
res96h

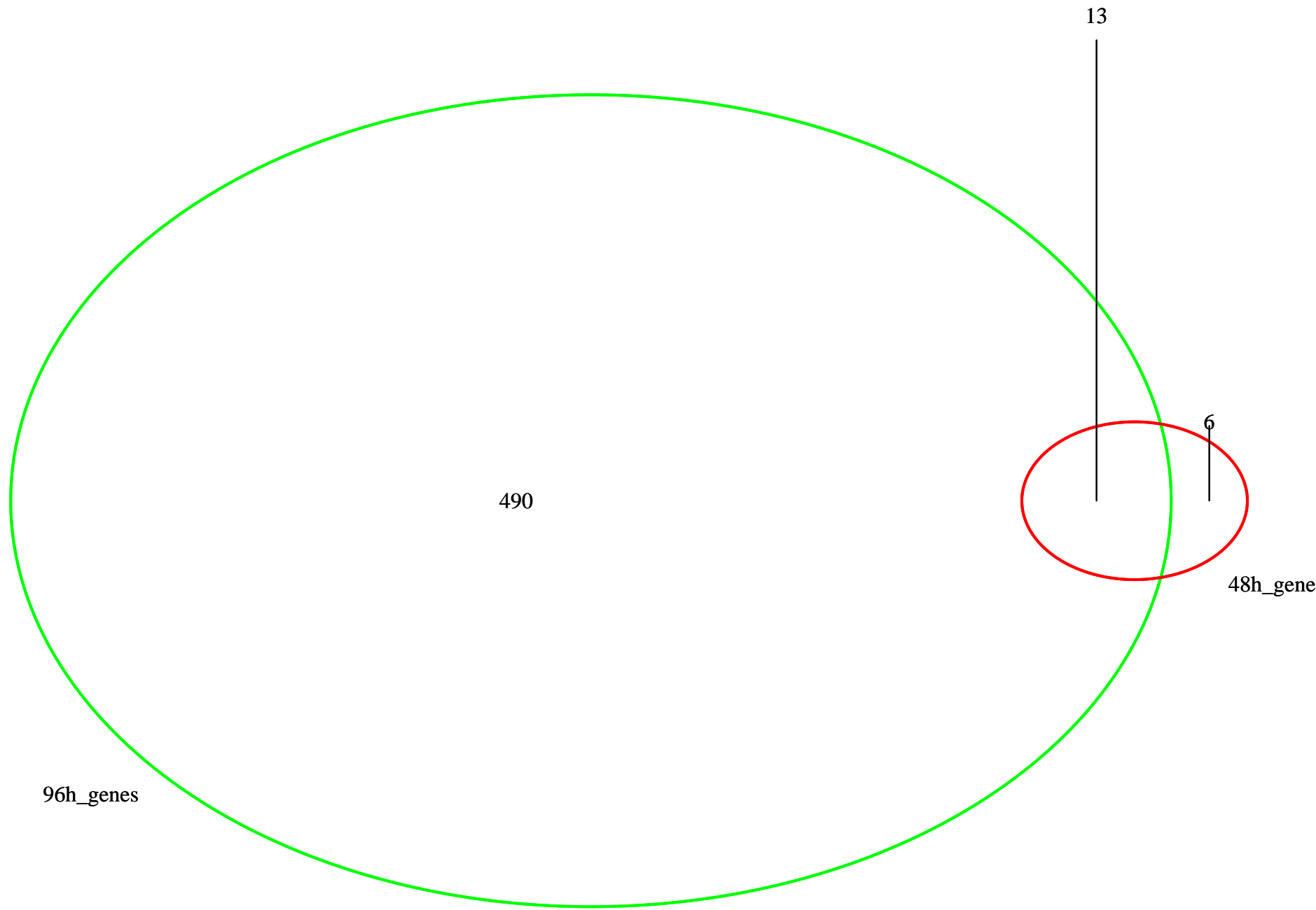


res48hControl_lowLFC

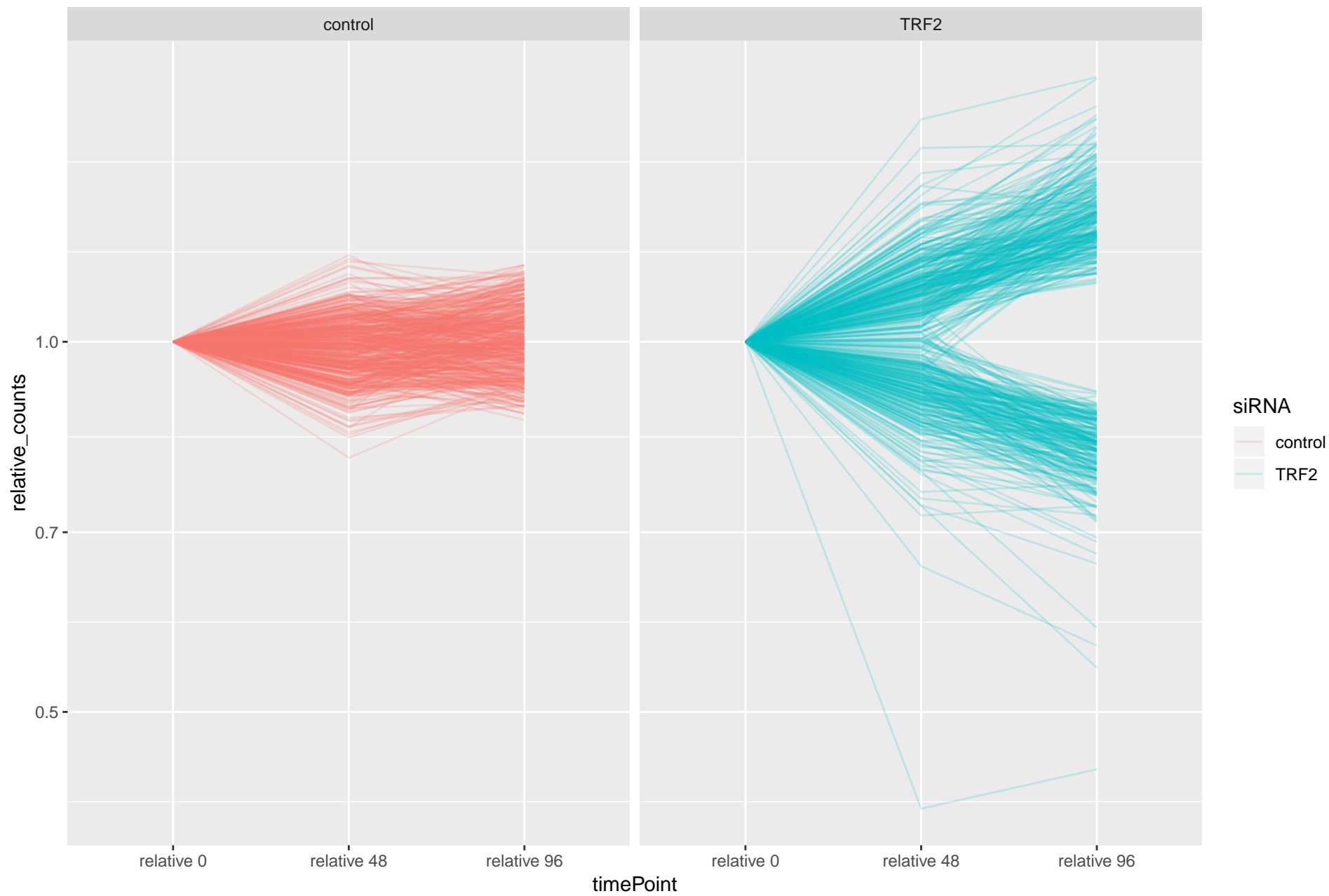


res96hControl_lowLFC

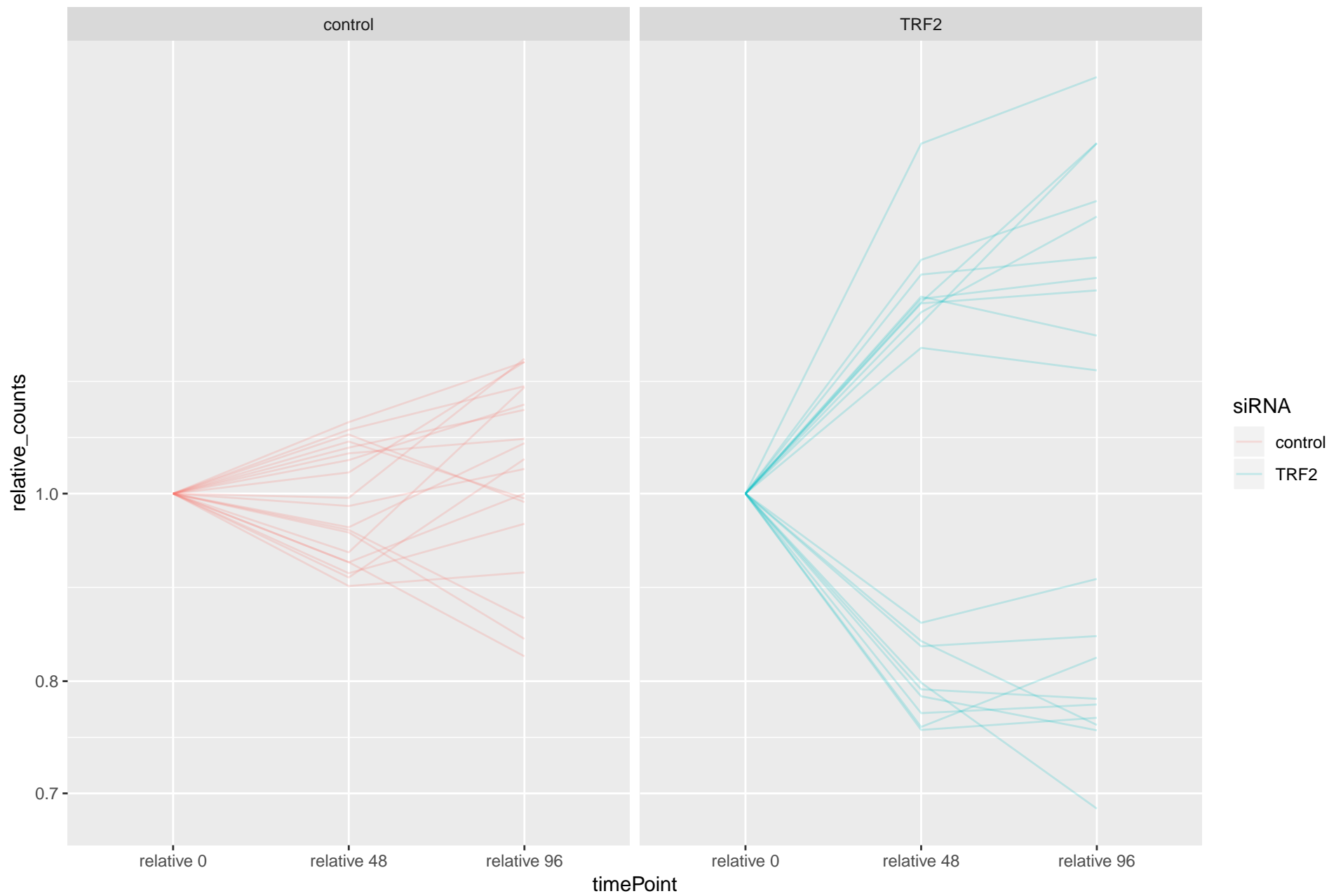




selected_genes96h

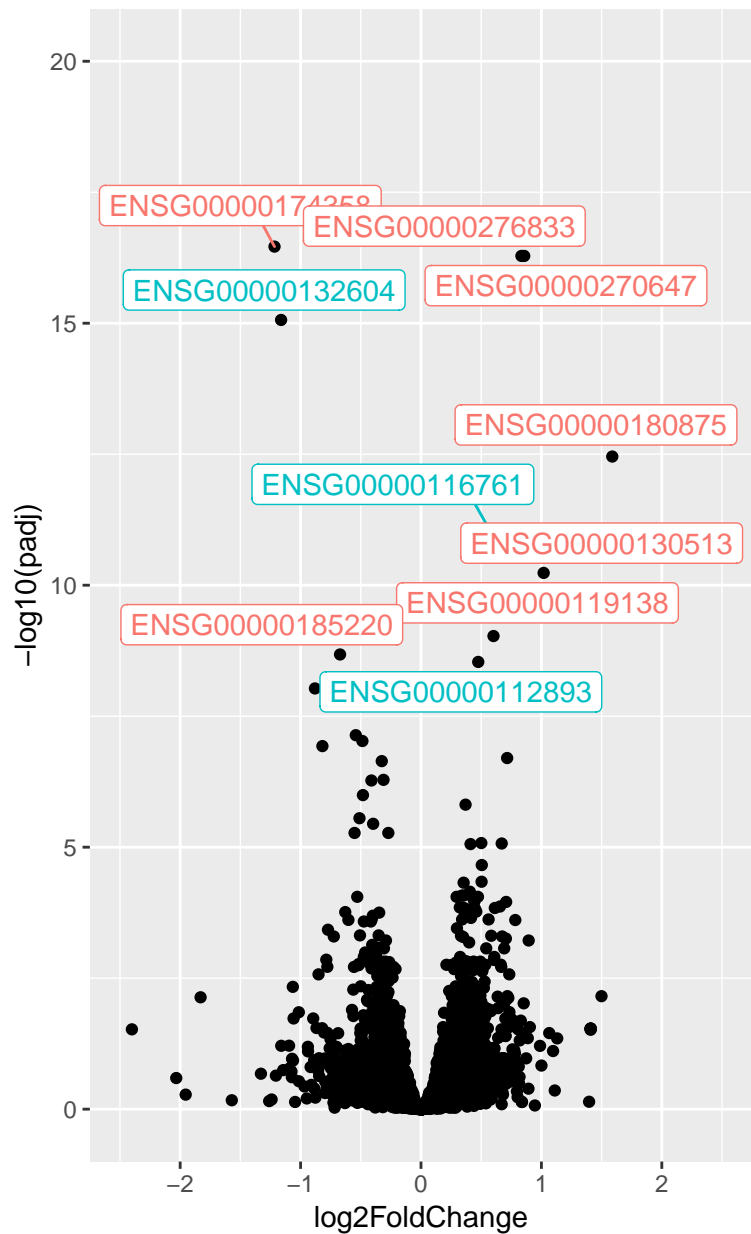


selected_genes48h



design = ~ condition

contrast = c("condition", "TRF2_96", "TRF2_0")



design = ~ condition

lfcThreshold = log2(1.25), altHypothesis = "lessAbs", contrast = c("condition", "TRF2_96", "TRF2_0")

