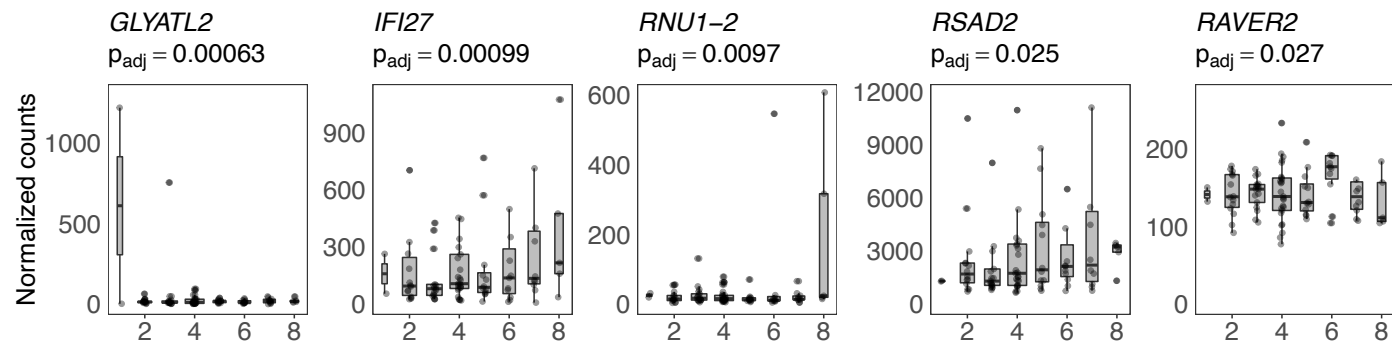
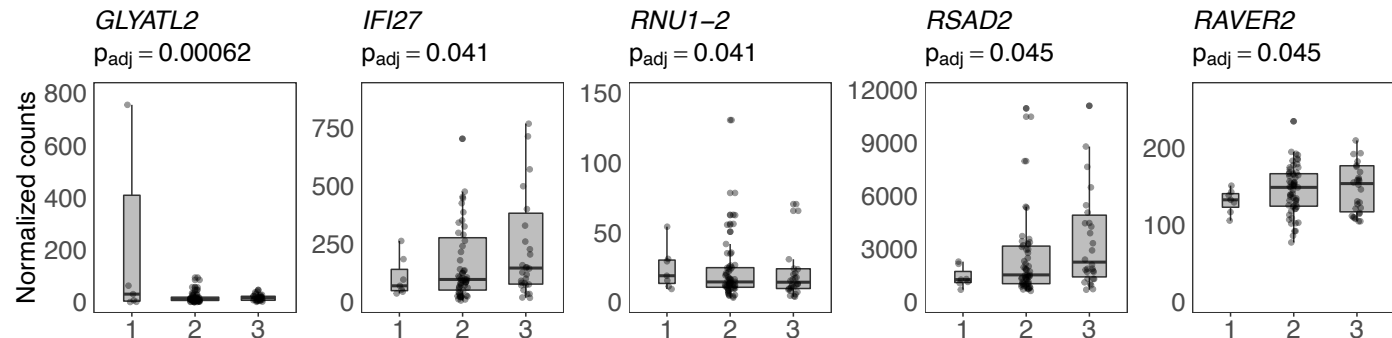


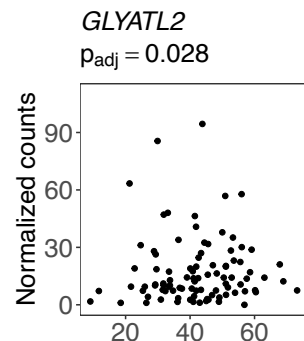
STIR volume



STIR composite



STIR intensity



Supplementary Figure 5: Protein-coding genes significantly differentially expressed across analyses. Expression levels of the genes in each sample normalized by sequencing depth and RNA composition (“Normalized counts”) vs STIR variable values. P-values are the adjusted significance values from the differential expression analyses (Table S2-S4).