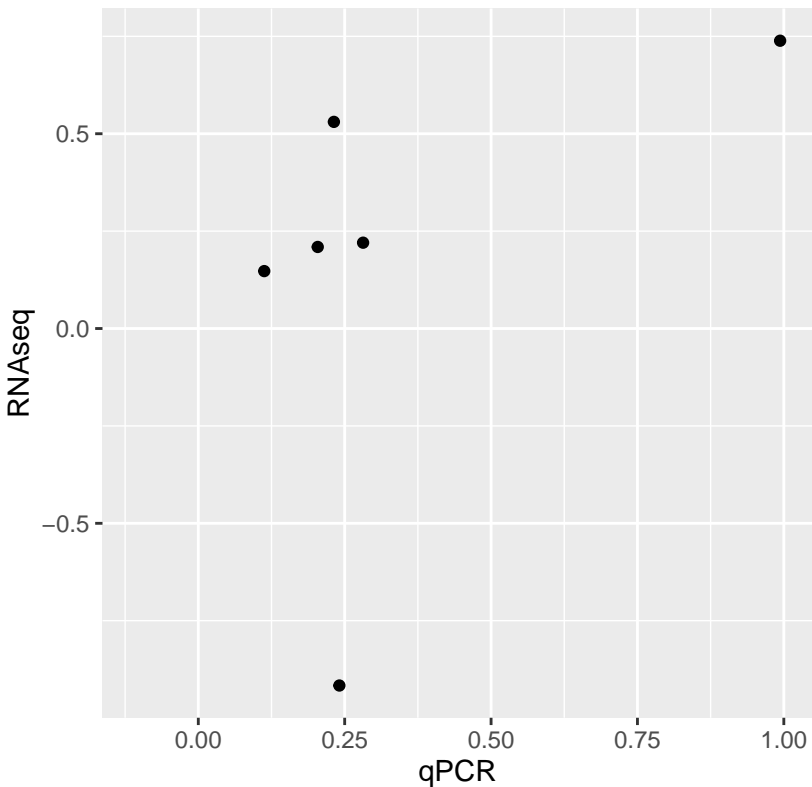


# Timepointw12 fold change, timepoint and sets

Model: Tissue offset lib size normalized



(Based on data from qpcr and gene sequencing)