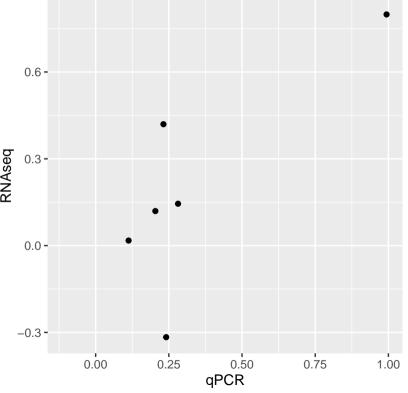
Timepointw12 fold change, timepoint only Model: Lib size normalized



(Based on data from qpcr and gene sequencing)