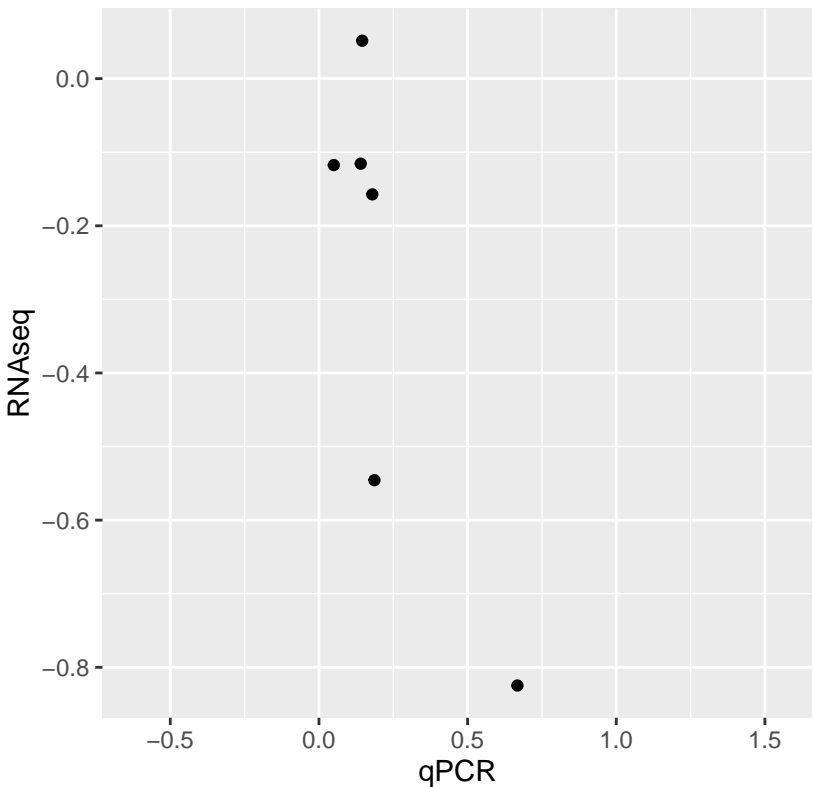


# Timepoint acute fold change, timepoint only

Lib size normalized



(Based on data from qpcr and gene sequencing)