

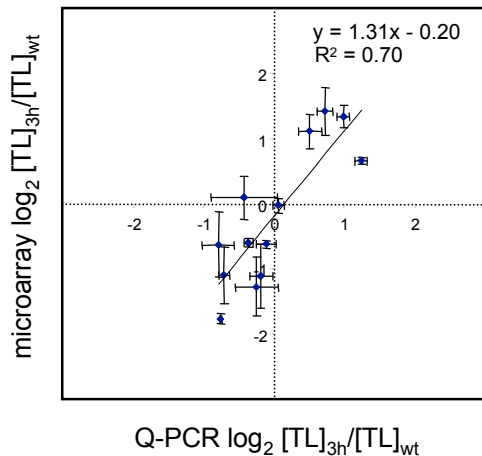
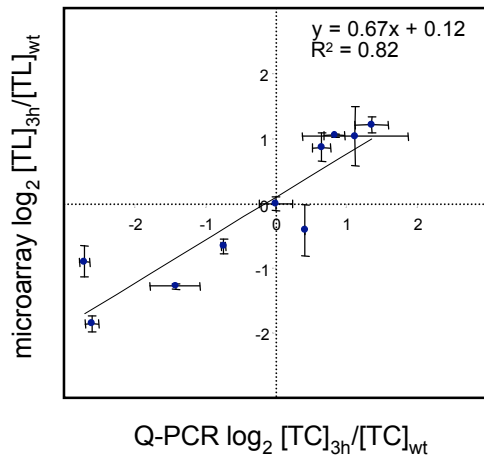
A**B**

Figure S2. Quantitative reverse-transcriptase PCR analysis of polysome loading for representative mRNAs detected in the microarray experiment. Microarray data (y-axis) on polysome loading state [TL] or total mRNA levels [TC] are correlated with corresponding quantitative RT-PCR data (x-axis).