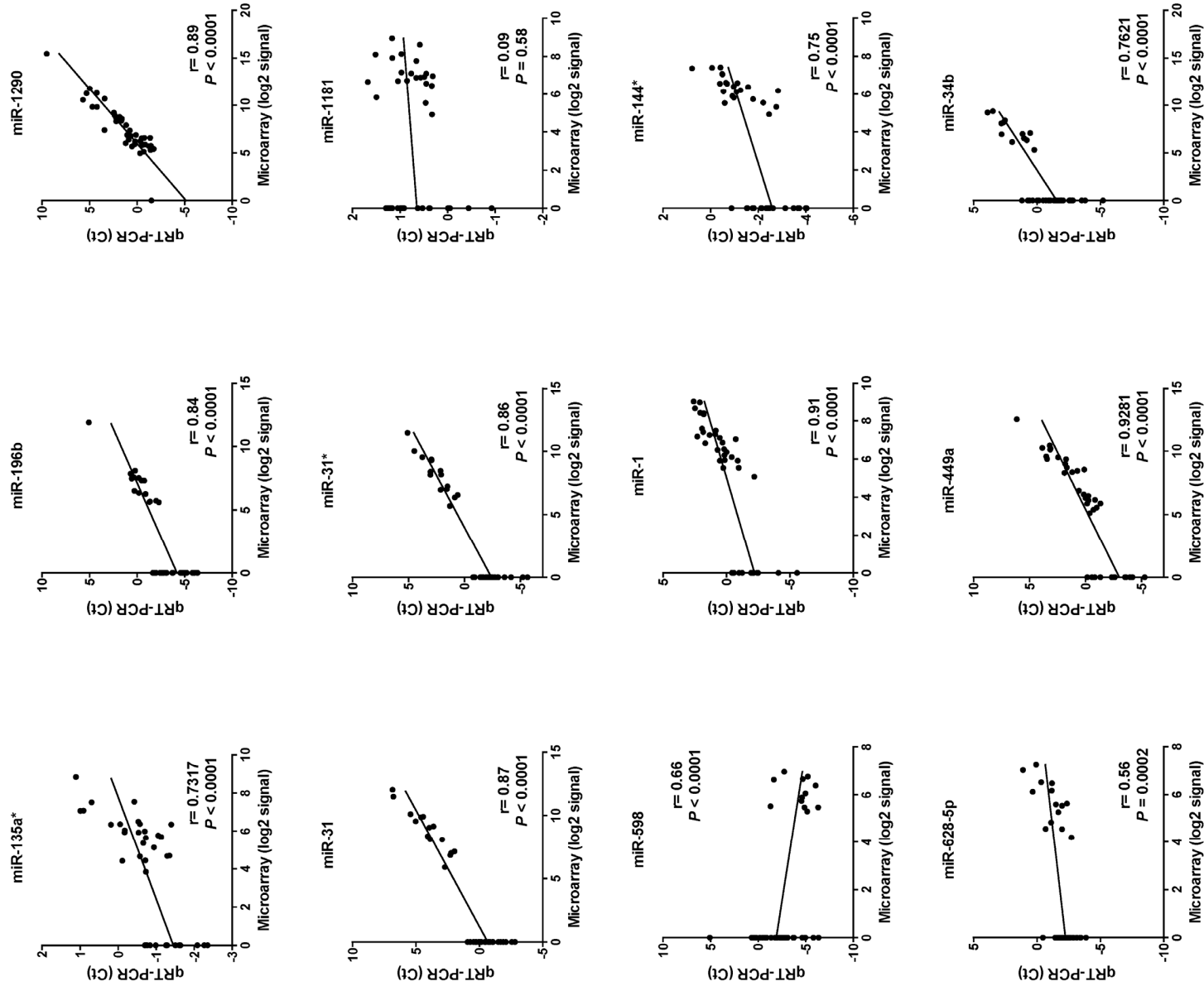


Supplemental Figure S1.



Supplemental Figure S1. Correlation of microarray data with quantitative RT-PCR data for 12-miRNAs expression in the test cohort. Expression levels of miRNAs assessed by qRT-PCR are shown by relative threshold cycle values against *RNU66*, a control gene. Expression levels of miRNAs assessed by microarray are shown by log2 signals. P -values by Spearman r test are shown.