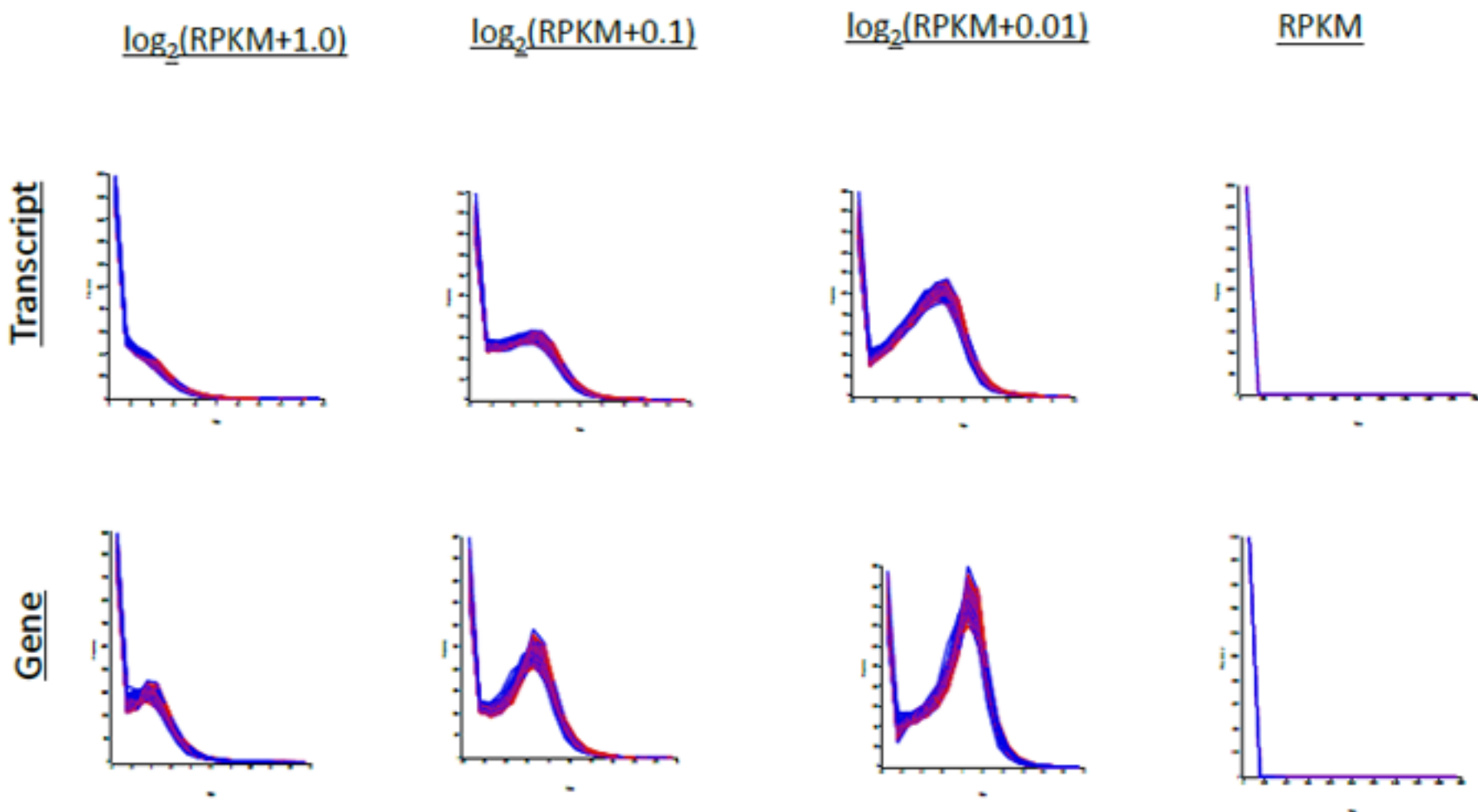
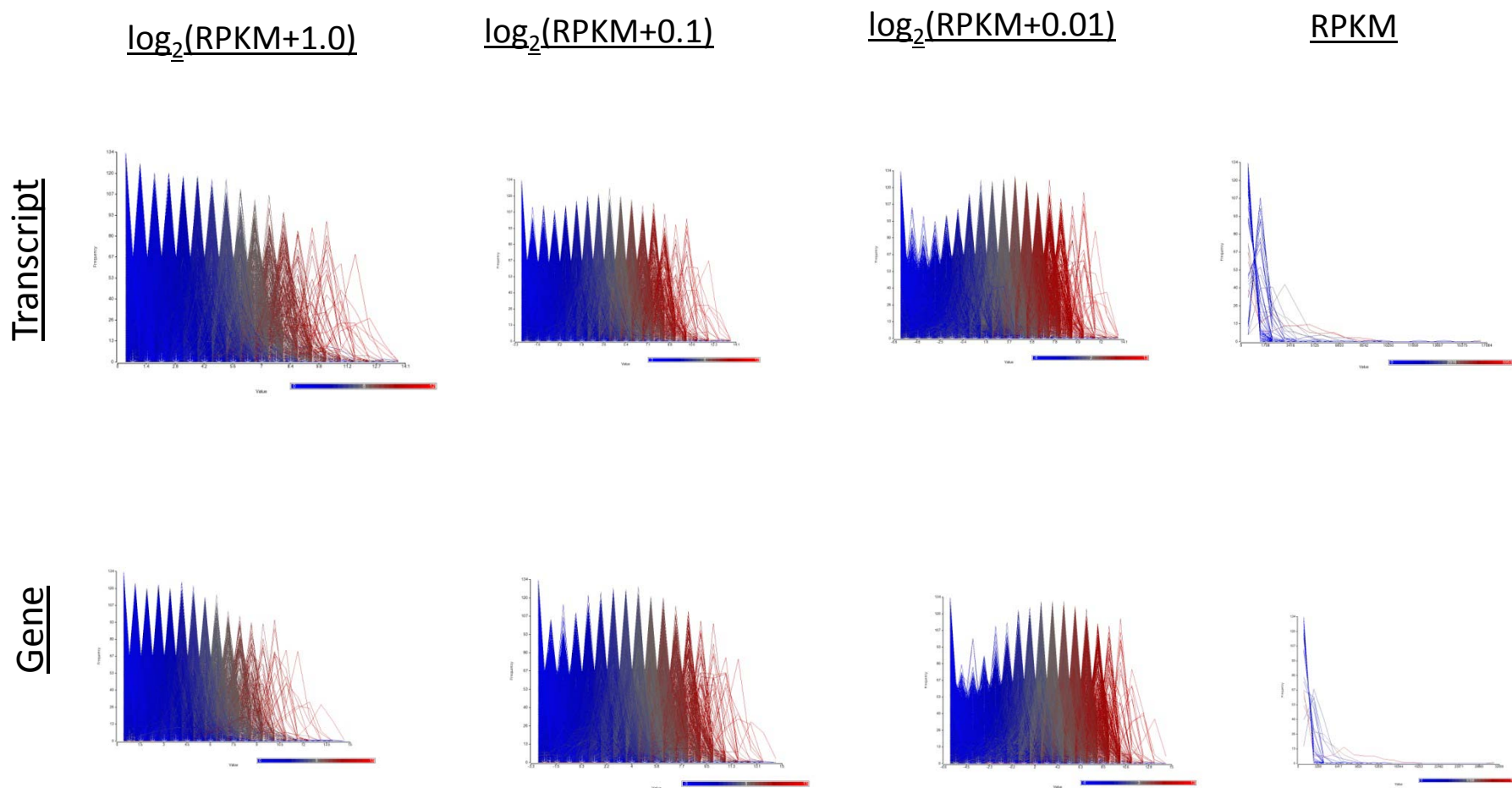


**Figure S1: Sample Histograms for Various Normalizations**



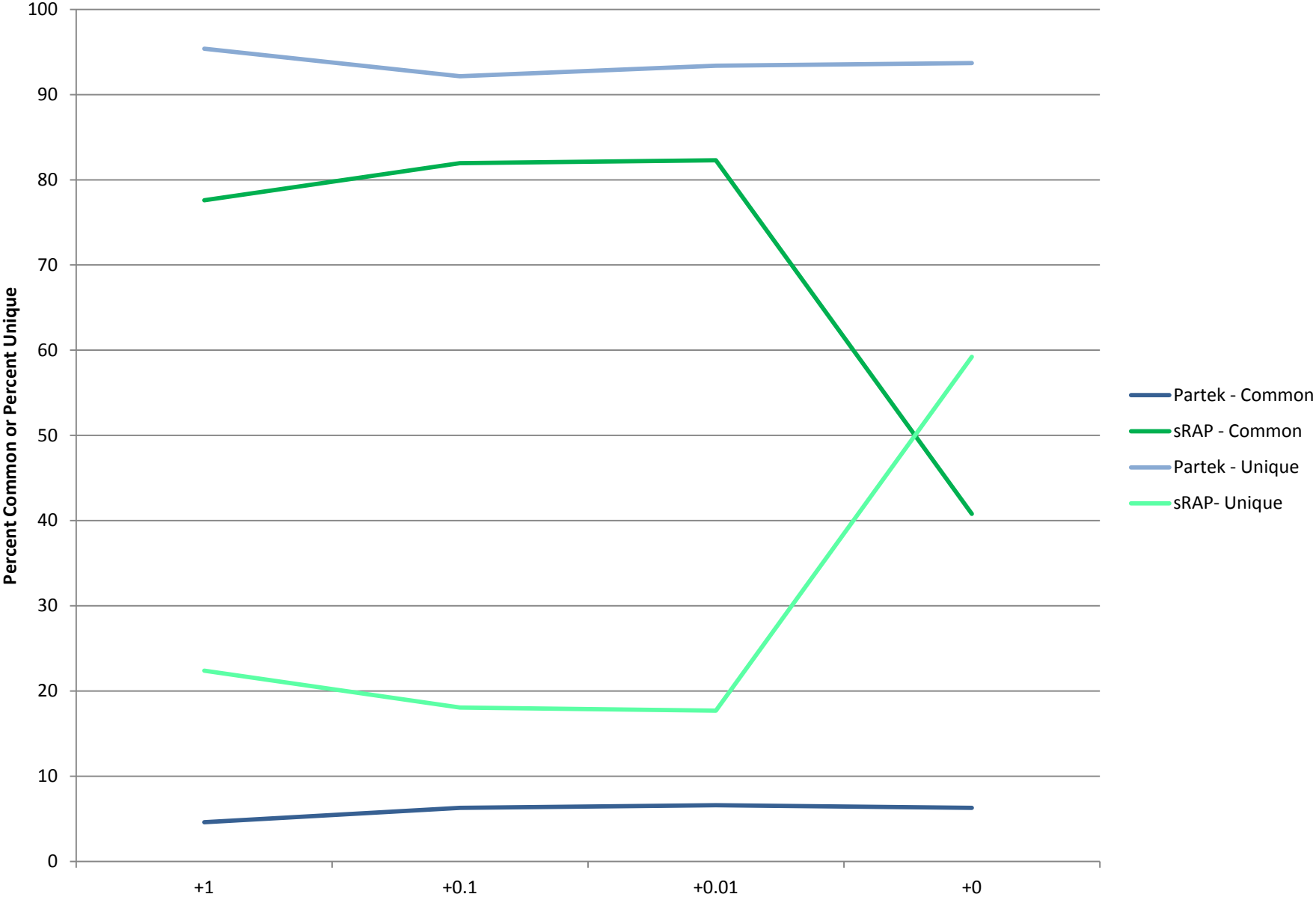
Tumor samples are colored in blue. Normal samples are colored in red.

## Figure S2: Gene Histograms for Various Normalizations



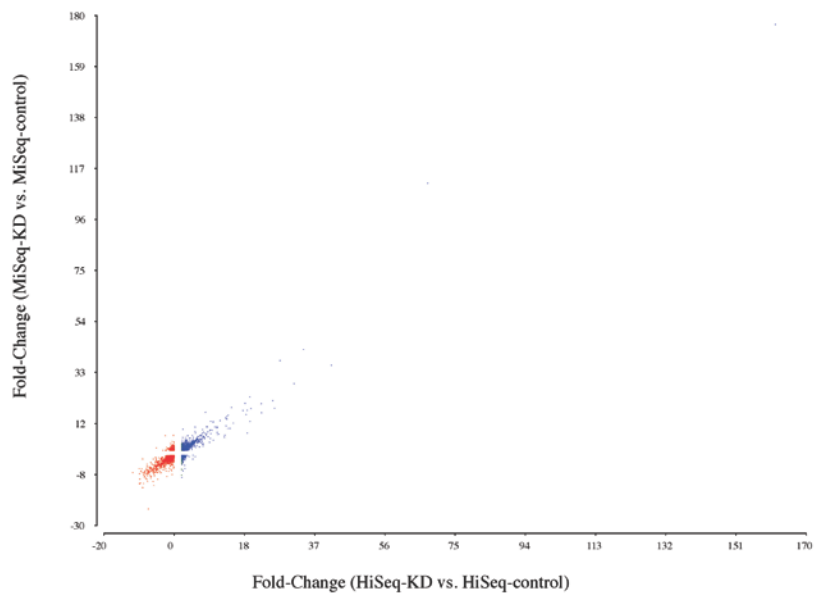
Gene distributions are colored based upon the expression values from an arbitrarily chosen sample (ERR164473)

**Figure S3: Most sRAP DEGs Overlap Partek DEGs in Patient Dataset**

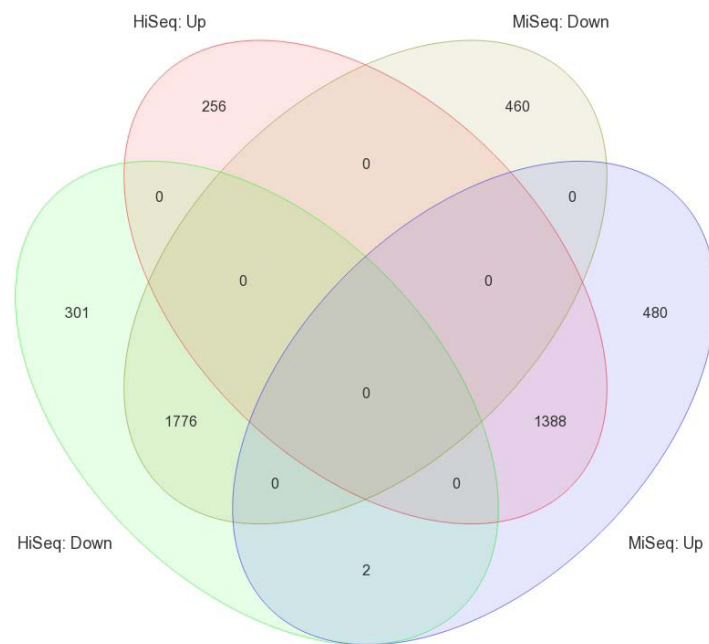


# Figure S4: Concordance between HiSeq and MiSeq Data

**A.**



**B.**



**Figure S5: Percentage Overlap with Microarray Data**

