

Cufflinks Expression Estimates

Fig. S1: Comparison of the log of the mean gene expression values determined by GENE-counter and Cufflinks. The average of the log transformed normalized expression values derived for genes from Cufflinks (x-axis) and GENE-counter (y-axis) were plotted against each (only $\Delta hrcC$ shown; mock was similar (r = 0.86)). Only genes tested in both packages with an estimated expression > 0 by either package ($\Delta hrcC$ = 19168 and mock = 19047) were included. In cases where a gene had more than one expressed isoform, the isoform with the highest expression level was used. All logtransformed values were treated as described in the main text. Pearson's correlation coefficients and the scatter plot were calculated and created as described in the main text.