



Supplemental Figure 3. Correlation of protein-coding abundances.

mRNA profiles were highly correlated between the same samples prepared by InDA-C rRNA depletion and polyA mRNA enrichment. The y-axis represents data generated in this study from InDA-C libraries, and the x-axis data from poly-A enriched libraries sequenced by Rosengarten et al., 2015. Time points, in hours (h), descend from 0 to 24 on the y-axis, and ascend from left to right on the x-axis. Spearman's correlation values are shown as a heatmap relating mRNA transcriptome profiles at each time point (average of 2 biological replicates). Green shading shows low correlation whereas pink shows high correlation, as indicated in the legend.