Fig. S9. Fitness scores of toxins (Tx) and antitoxin (Ant) genes of predicted TA systems in the strain B728a. Y-axis on the left shows number of reads that lie within the gene and normalized so that the median across all genes is 1, and also normalized for GC content. Y-axis on the right shows if the gene contained enough reads to be used in experiments. Genes that were not used were important for in vitro growth