

Operon Promoter Landscape

Operon	Strand	Operon start	Operon end
rpsP-rimM-trmD-rplS	-	2744207	2742205

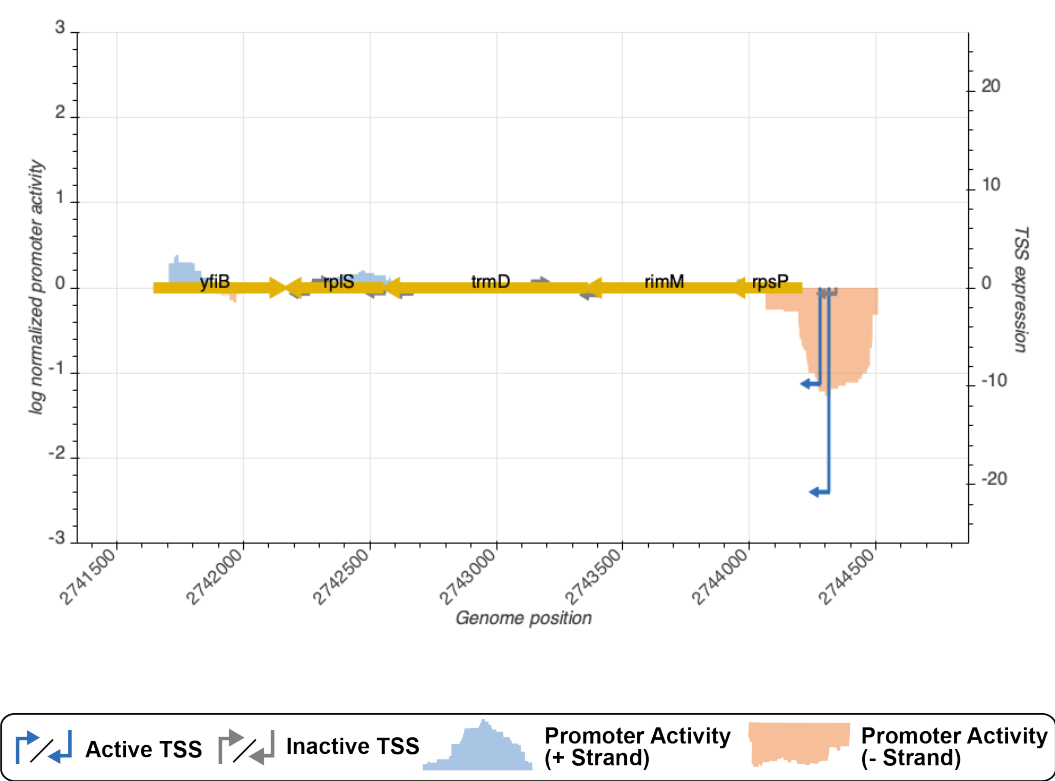


Figure 1: **Promoter activity in rich media (LB) surrounding query operon.** 17,767 previously reported TSSs were evaluated by measuring the promoter activity (right Y-axis) of the 150 bp surrounding the TSS (-120 to +30) to determine which were active or inactive. The genome-wide promoter activity (left Y-axis) was determined by measuring expression of over 300,000 genomic fragments spanning the *E. coli* genome and averaging promoter activity at all nucleotide positions in a strand-specific fashion. Genome coordinates corresponds to *E. coli* genome version U00096.2.

TSS Summary

TSS name	TSS position	Strand	TSS activity	Category
TSS_10376_regulondb	2744338	-	0.5914951	inactive
TSS_10374_wanner	2744275	-	9.7557143	active
TSS_10375_regulondb	2744310	-	20.7593214	active

TSS Scanning Mutagenesis

