

Operon Promoter Landscape

Operon	Strand	Operon start	Operon end
prmB-aroC-mepA-yfcA-epmC-yfcL	-	2446462	2441913

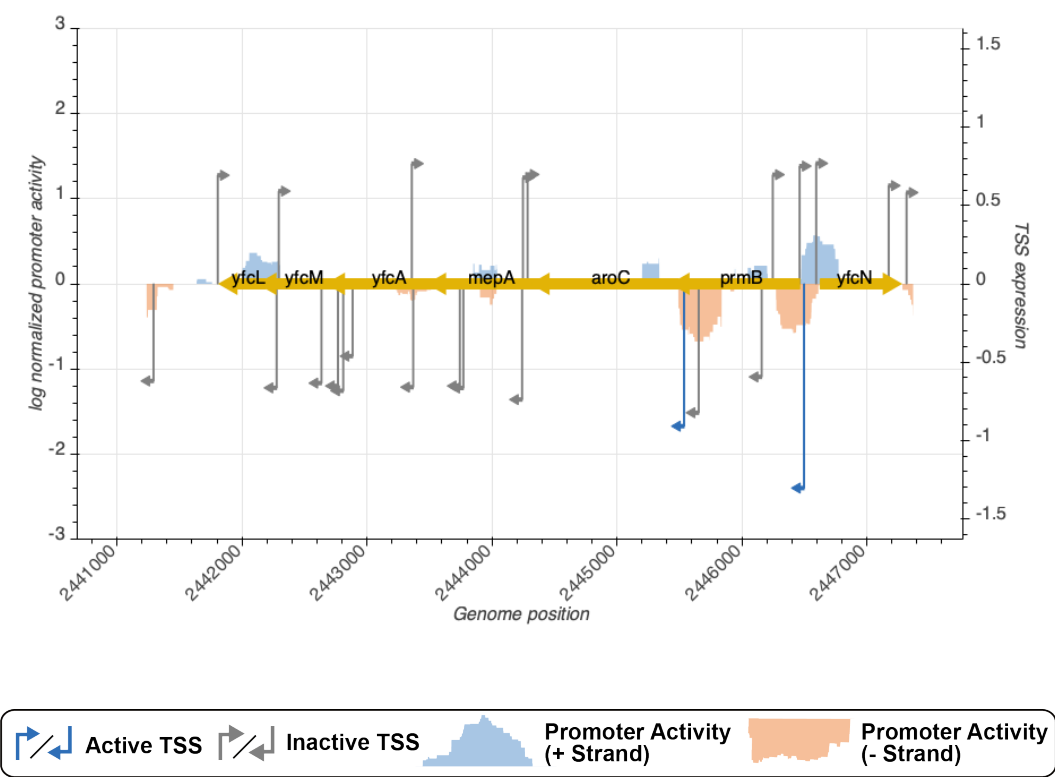


Figure 1: **Promoter activity in rich media (LB) surrounding query operon.** 17,767 previously reported TSSs were evaluated by measuring the promoter activity (right Y-axis) of the 150 bp surrounding the TSS (-120 to +30) to determine which were active or inactive. The genome-wide promoter activity (left 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_9252_wanner	2446493	-	1.3048303	active
TSS_9253_regulondb	2446590	+	0.7696107	inactive

TSS Scanning Mutagenesis

