

Operon Promoter Landscape

Operon	Strand	Operon start	Operon end
hinT-ycfL-lpoB-thiK-nagZ-ycfP	+	1161108	1164908

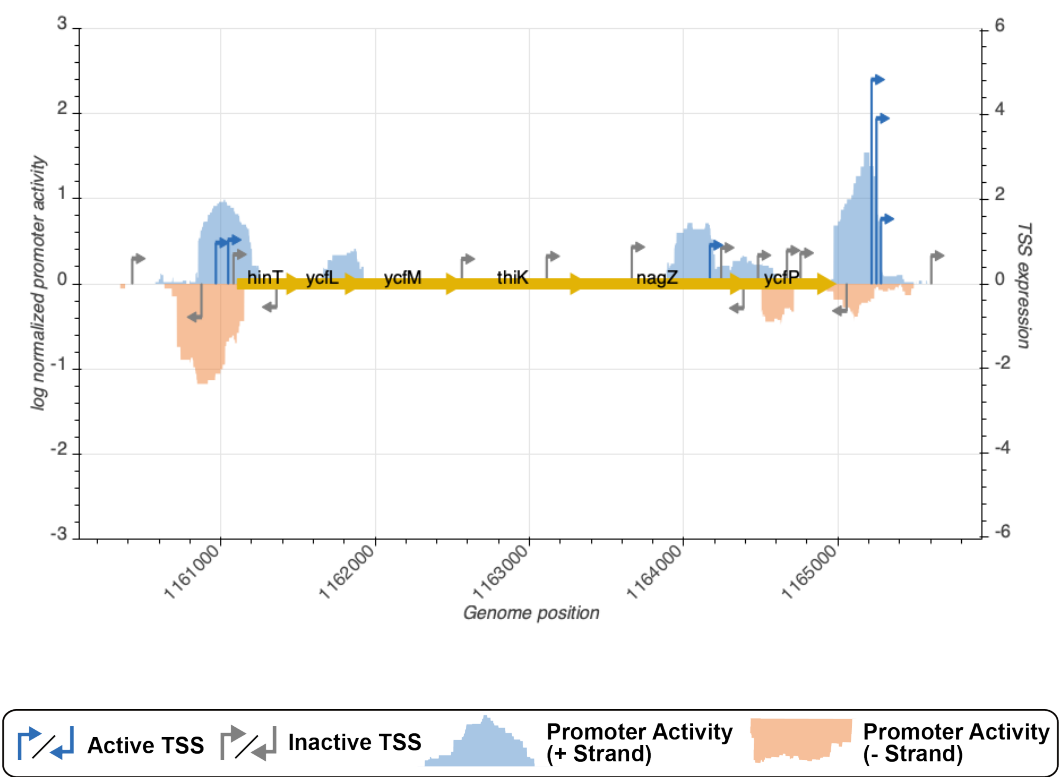


Figure 1: **Promoter activity in rich media (LB) surrounding query operon.** 17,767 previously reported TSSs were evaluated by measuring the promoter activity (right Y-axis) of the 150 bp surrounding the TSS (-120 to +30) to determine which were active or inactive. The genome-wide promoter activity (left 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_4501_regulondb	1161078	+	0.6968530	inactive
TSS_4498_regulondb	1160870	-	0.7872205	inactive
TSS_4500_regulondb	1161044	+	1.0407428	active
TSS_4499_storz	1160964	+	0.9718203	active

TSS Scanning Mutagenesis

