

Operon Promoter Landscape

Operon	Strand	Operon start	Operon end
codBA	+	354146	356678

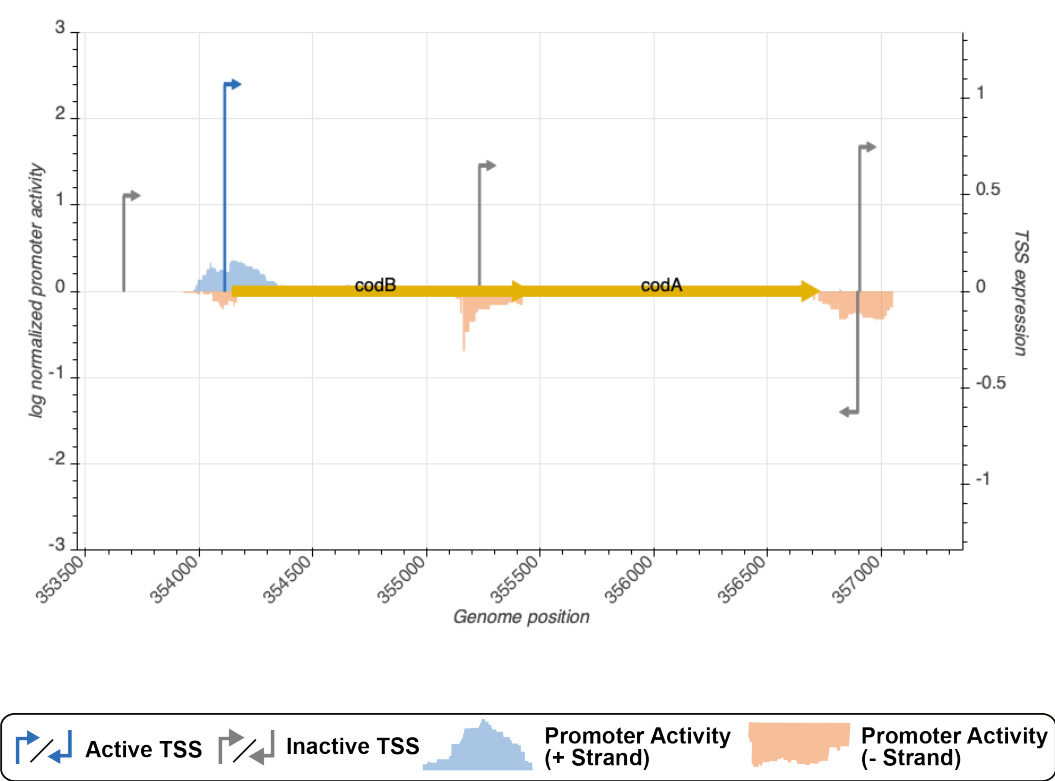


Figure 1: **Promoter activity in rich media (LB) surrounding query operon.** 17,767 previously reported TSSs were evaluated by measuring the promoter activity (right Y-axis) of the 150 bp surrounding the TSS (-120 to +30) to determine which were active or inactive. The genome-wide promoter activity (left 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_1435_storz	353664	+	0.4962869	inactive
TSS_1436_storz_regulondb	354108	+	1.0737537	active

TSS Scanning Mutagenesis

