

Operon Promoter Landscape

Operon	Strand	Operon start	Operon end
galP	+	3086306	3087700

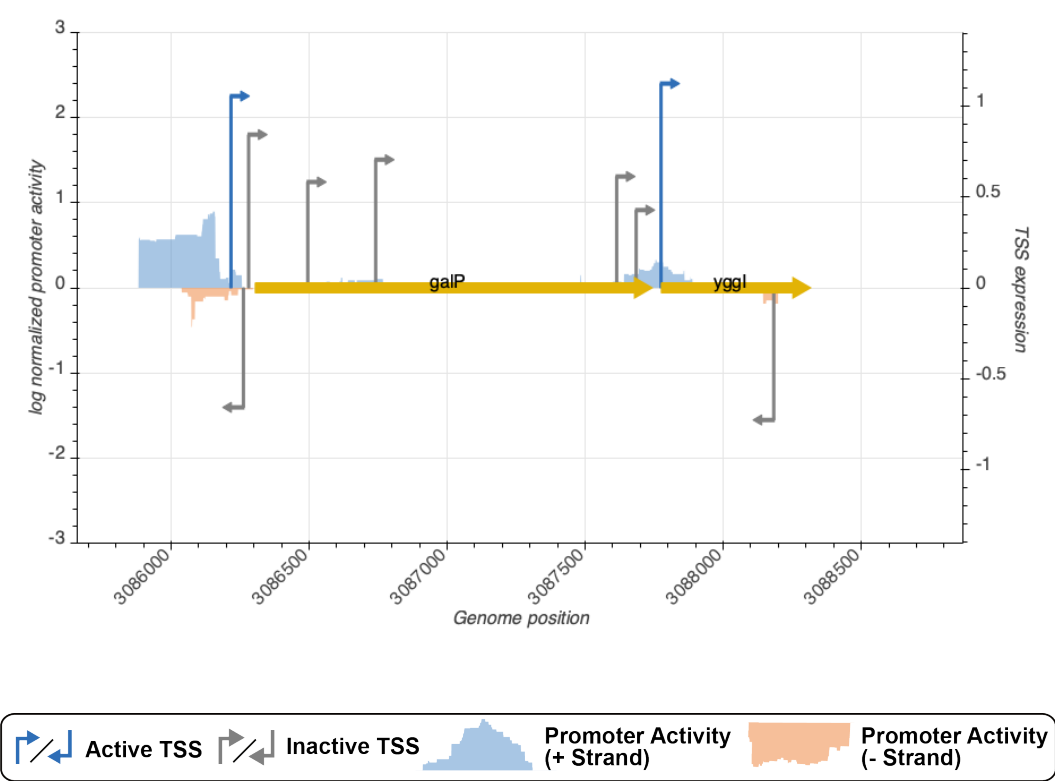


Figure 1: **Promoter activity in rich media (LB) surrounding query operon.** 17,767 previously reported TSSs were evaluated by measuring the promoter activity (right Y-axis) of the 150 bp surrounding the TSS (-120 to +30) to determine which were active or inactive. The genome-wide promoter activity (left 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_11666_wanner	3086213	+	1.0566665	active
TSS_11668_regulondb	3086277	+	0.8444035	inactive
TSS_11667_storz	3086258	-	0.6588053	inactive

TSS Scanning Mutagenesis

