Operon	Strand	Operon start	Operon end
yhjR	-	3694208	3694020

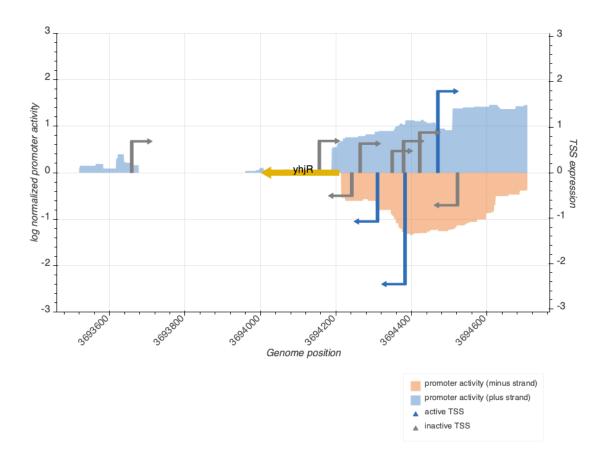


Figure 1: Figure 1: Promoter activity surrounding query operon.

TSS name	TSS position	Strand	TSS activity	Category
TSS_14304_storz	3694260	+	0.6385152	inactive
$TSS_14308_{regulondb}$	3694379	-	2.4521378	active
$TSS_14305_{regulondb}$	3694306	-	1.0737537	active
TSS_14309_regulondb	3694418	+	0.8826286	inactive
$TSS\_14311\_storz$	3694518	-	0.7156143	inactive
$TSS_14306$ _regulondb	3694345	+	0.4757168	inactive
$TSS_14307$ _wanner	3694375	+	0.6924437	inactive
TSS_14310_regulondb	3694466	+	1.7894869	active
TSS_14303_regulondb	3694238	-	0.5087278	inactive

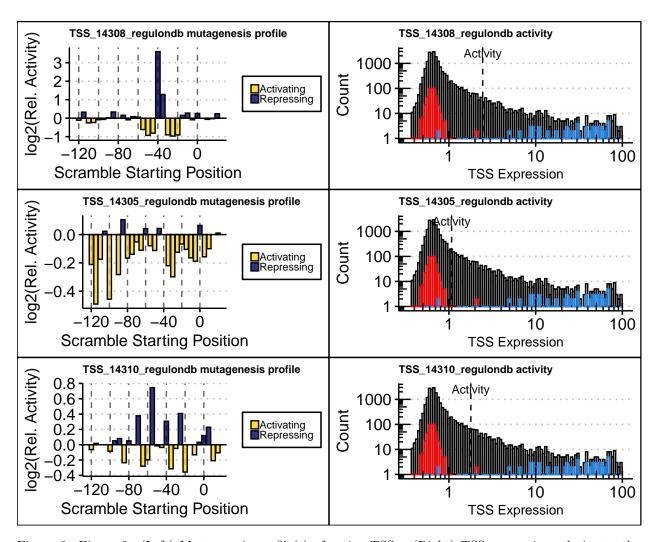


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.