Operon	Strand	Operon start	Operon end
rpmH-rnpA	+	3882359	3882875

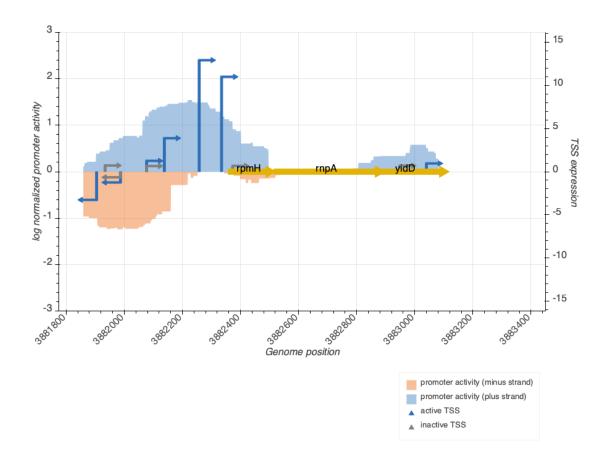


Figure 1: Figure 1: Promoter activity surrounding query operon.

TSS name	TSS position	Strand	TSS activity	Category
TSS_15044_storz	3882073	+	1.2504521	active
TSS_15043 _wanner	3881983	-	1.2504521	active
TSS_15042_storz	3881930	+	0.7155458	inactive
$TSS_15041_{regulondb}$	3881901	-	3.2690491	active
TSS_15044_storz	3882073	+	0.6524152	inactive
TSS_15043 _wanner	3881983	-	0.6524152	inactive
TSS_15045 _regulondb	3882134	+	3.8831791	active
TSS_15046_storz	3882254	+	12.9213519	active
TSS_15047_regulondb	3882331	+	10.9985871	active

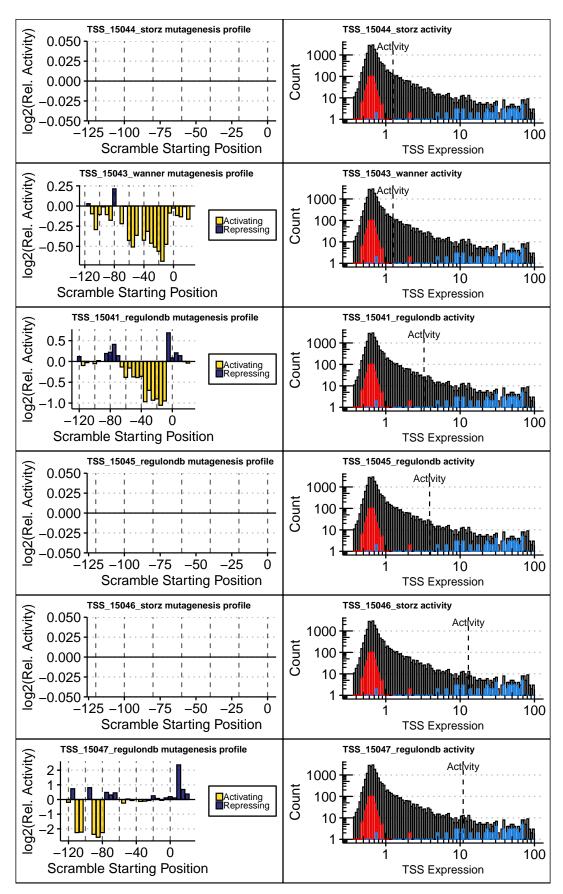


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. T\$Ss were integrated into the nth-ydgR intergenic locus for quantification.