

Operon	Strand	Operon start	Operon end
yifL-dapF-yigA-xerC-yigB	+	3992545	3995922

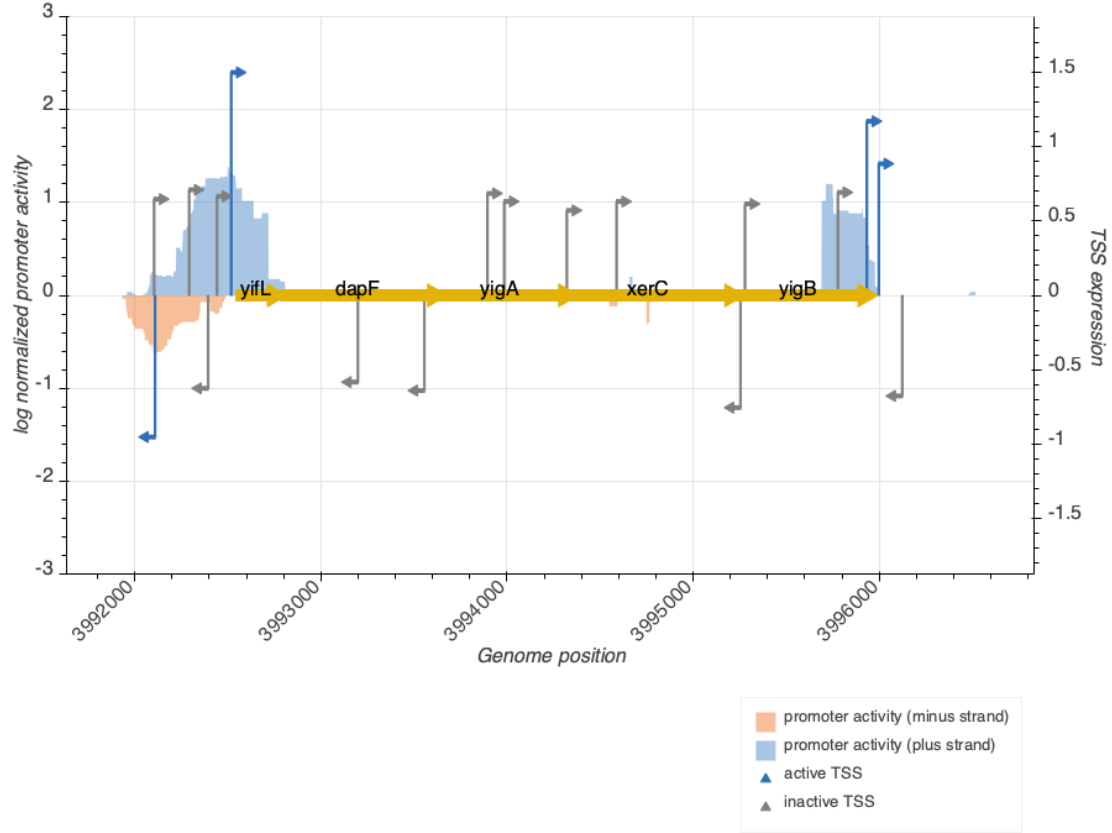


Figure 1: Figure 1: Promoter activity surrounding query operon.

TSS name	TSS position	Strand	TSS activity	Category
TSS_15521_storz	3992100	+	0.6464243	inactive
TSS_15525_storz	3992438	+	0.6655643	inactive
TSS_15524_storz	3992390	-	0.6249061	inactive
TSS_15523_storz	3992288	+	0.7086318	inactive
TSS_15526_regulondb	3992514	+	1.4972367	active
TSS_15522_wanner	3992105	-	0.9514336	active

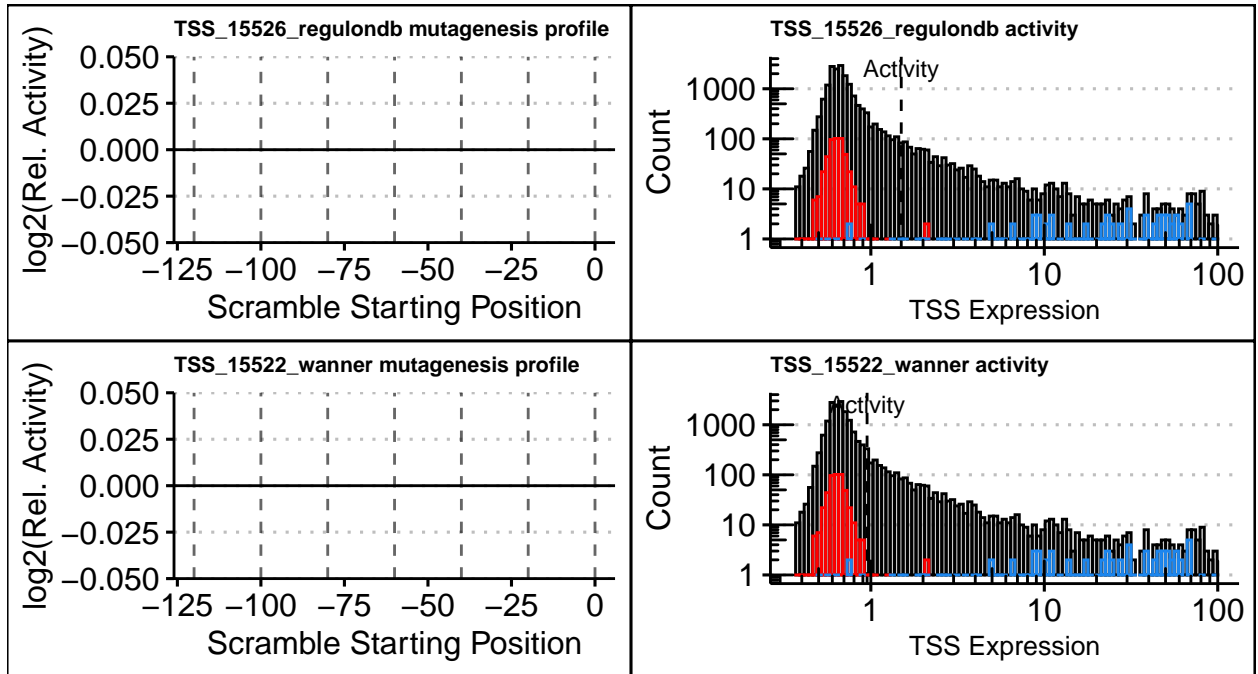


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the *nth-ydgR* intergenic locus for quantification.