Operon	Strand	Operon start	Operon end
chaA	-	1271072	1269972

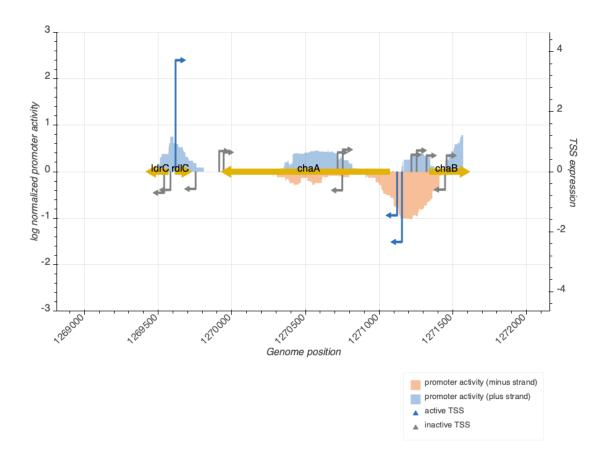


Figure 1: Figure 1: Promoter activity surrounding query operon.

TSS name	TSS position	Strand	TSS activity	Category
TSS_4938_storz_regulondb TSS_4940_regulondb TSS_4937_regulondb	1271318 1271454 1271251	+ + + + +	$0.5368317 \\ 0.5371504 \\ 0.7052432$	inactive inactive
TSS_4936_storz_regulondb TSS_4939_storz	1271231 1271216 1271444	+	$0.5647513 \\ 0.5978077$	inactive inactive
TSS_4935_storz TSS_4934_wanner	$1271151 \\ 1271118$	-	$2.3404971 \\ 1.4543874$	active active

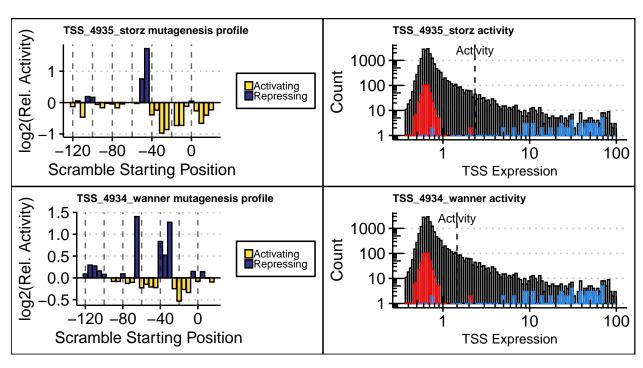


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.