| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| ybjM | + | 889312 | 889689 |

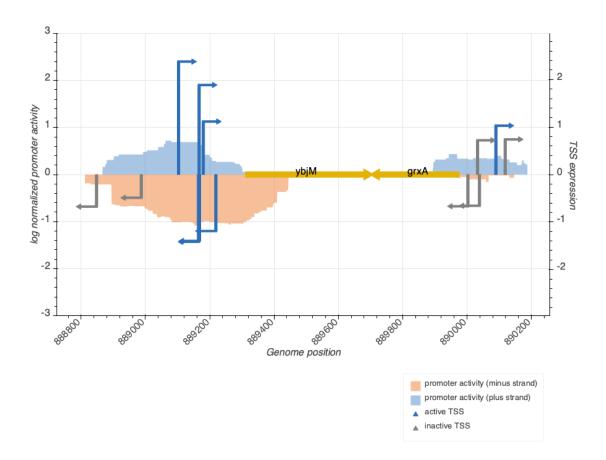


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|-----------------------|--------------|--------|--------------|----------|
| TSS_3397_storz | 889099 | + | 2.3785841 | active |
| TSS_3402 _wanner | 889215 | - | 1.1898100 | active |
| TSS_3398_regulondb | 889161 | - | 1.4161375 | active |
| TSS_3395_storz | 888844 | - | 0.6759383 | inactive |
| TSS_3400 _regulondb | 889164 | - | 1.3981530 | active |
| TSS_3396_storz | 888983 | - | 0.4885926 | inactive |
| TSS_3399_regulondb | 889163 | + | 1.8853930 | active |
| TSS_3401_wanner | 889176 | + | 1.1164884 | active |

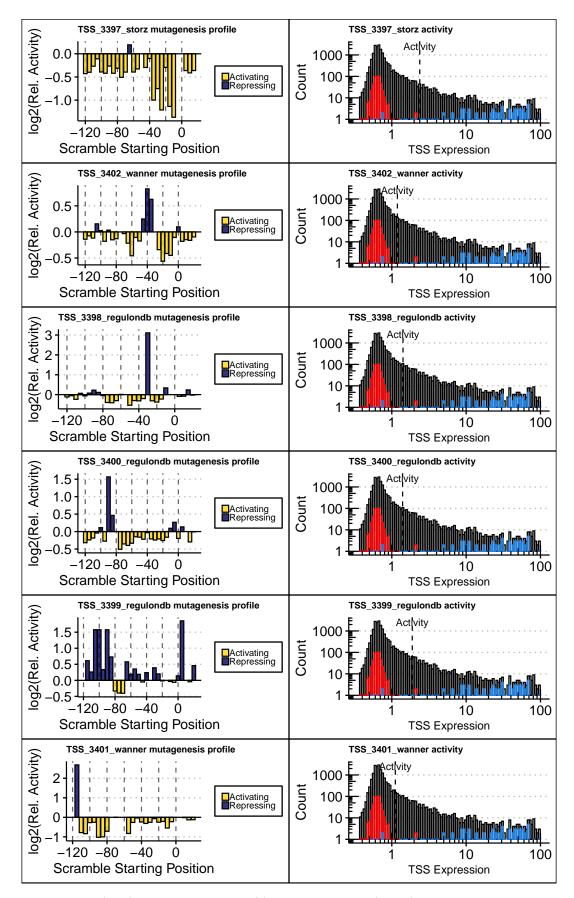


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. T\$Ss were integrated into the nth-ydgR intergenic locus for quantification.