| Operon | Strand | Operon start | Operon end | |
|--------|--------|--------------|------------|--|
| pgi | + | 4231781 | 4233430 | |

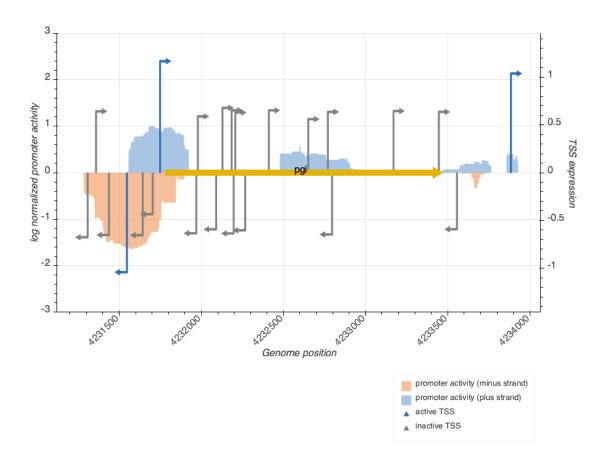


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|-------------------------|--------------|--------|--------------|----------|
| TSS_16574_regulondb | 4231304 | - | 0.6755228 | inactive |
| TSS_16576_storz | 4231434 | - | 0.6524152 | inactive |
| TSS_16579_storz | 4231700 | - | 0.4349434 | inactive |
| $TSS_16578_regulondb$ | 4231639 | - | 0.6544967 | inactive |
| $TSS_16580_{regulondb}$ | 4231745 | + | 1.1661844 | active |
| TSS_16575 _wanner | 4231355 | + | 0.6419117 | inactive |
| TSS_16577_wanner | 4231543 | - | 1.0417293 | active |

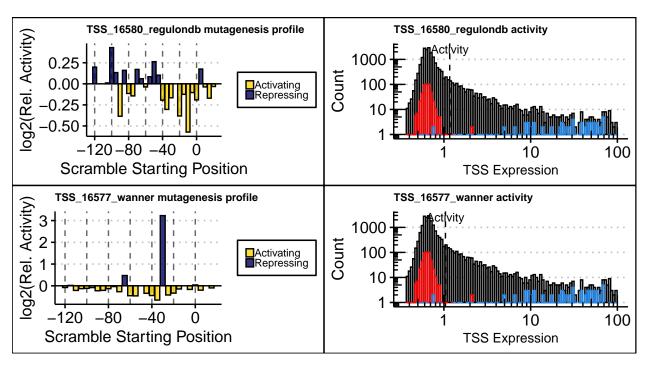


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.