| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| ybeQ   | -      | 675770       | 674793     |

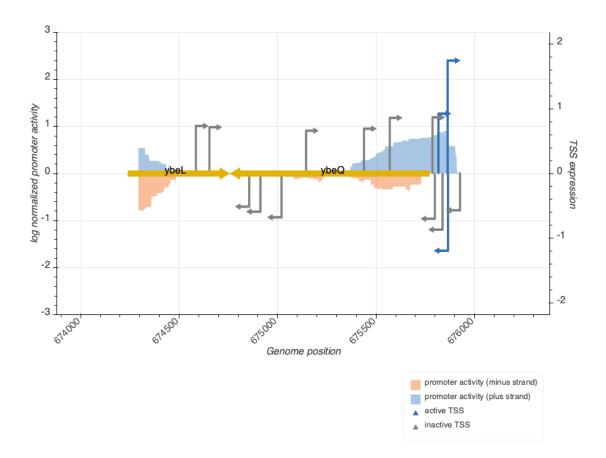


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name   | TSS position                                   | Strand                | TSS activity  | Category   |
|--|--|-----------------------|---|--|
| TSS_2609_regulondb<br>TSS_2606_regulondb<br>TSS_2604_regulondb<br>TSS_2605_regulondb<br>TSS_2608_regulondb | 675862<br>675813<br>675783<br>675795<br>675860 | -<br>+<br>+<br>-<br>+ | 1.1953860<br>0.9269696<br>0.8703654<br>0.7002913<br>1.7492766 | active<br>active<br>inactive<br>inactive<br>active |
| TSS_2610_regulondb<br>TSS_2607_regulondb   | $675923 \\ 675834$                             | -<br>-                | $0.5690664 \\ 0.8695032$                                      | inactive inactive                                  |

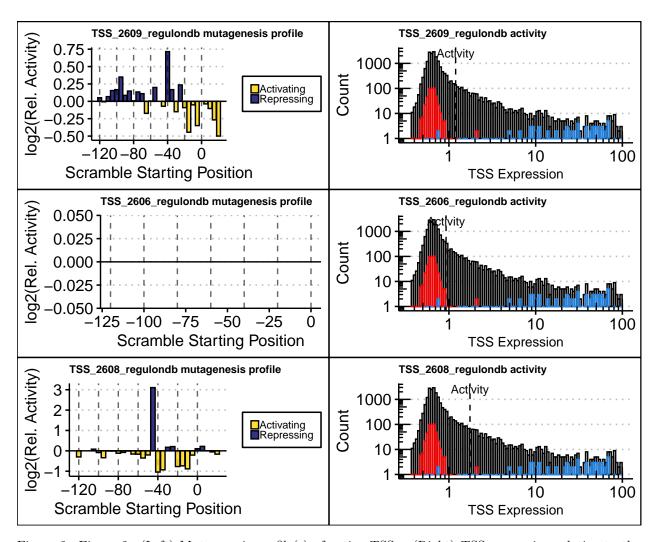


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.