| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| ymfK | - | 1202156 | 1201482 |

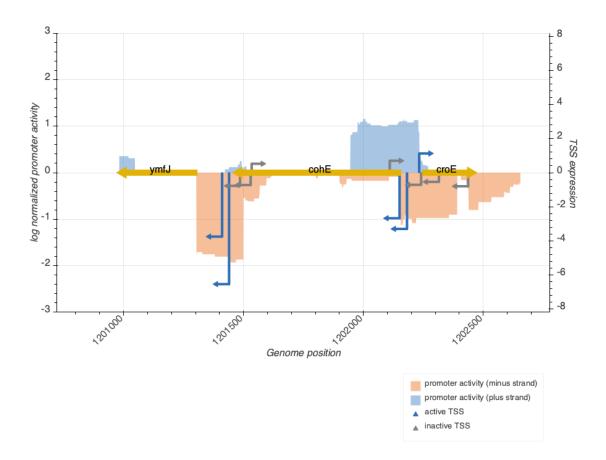


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|--|-------------------------------|-------------|--|--------------------------------|
| TSS_4661_regulondb TSS_4665_storz | 1202178 1202433 | - | 3.3086327 0.7987864 | active inactive |
| TSS_4663_regulondb TSS_4664_regulondb TSS_4662_regulondb | 1202237 1202311 1202229 | - - + | $\begin{array}{c} 0.7158358 \\ 0.5408609 \\ 1.1302096 \end{array}$ | inactive inactive active |

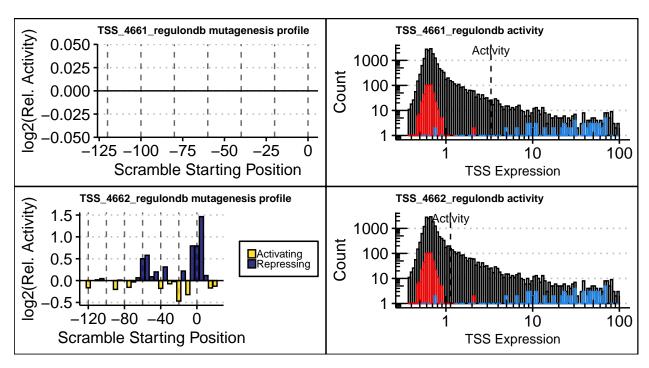


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.