| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| sspAB  | -      | 3375442      | 3374301    |
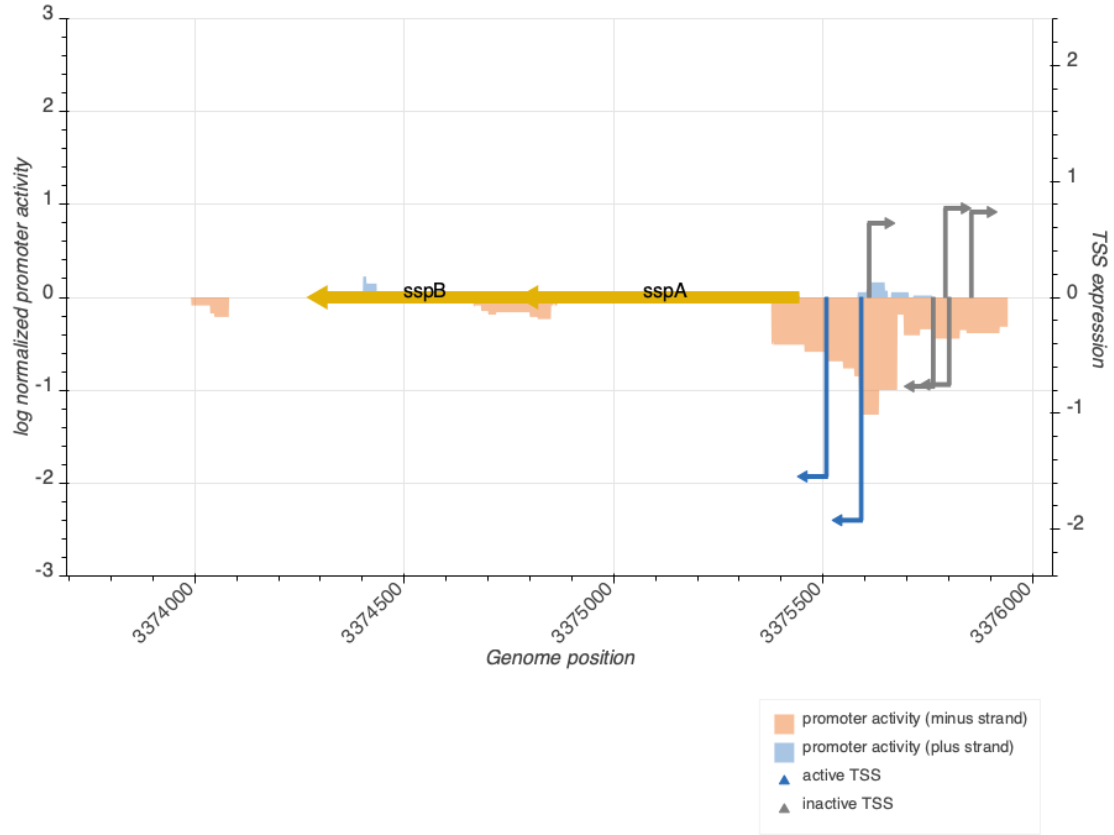


Figure 1: Figure 1: Promoter activity surrounding query operon.

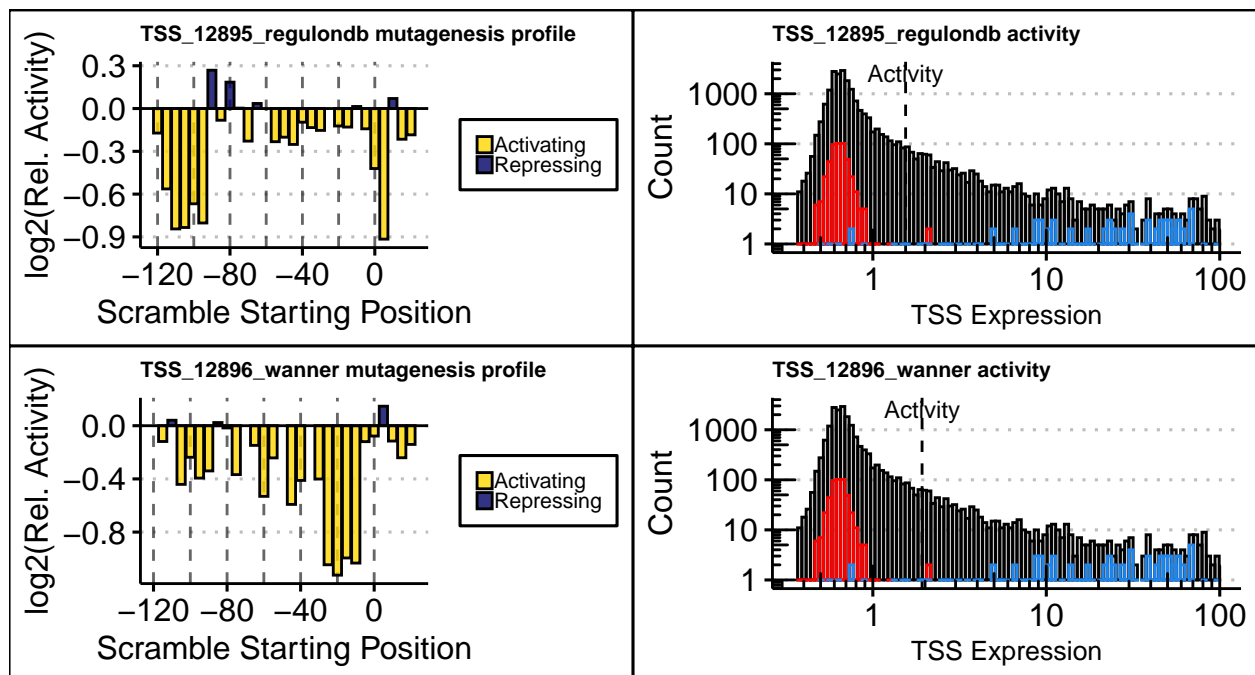| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_12899_storz     | 3375788 | + | 0.7680428 | inactive |
| TSS_12897_storz     | 3375606 | + | 0.6391195 | inactive |
| TSS_12901_storz     | 3375850 | + | 0.7358152 | inactive |
| TSS_12900_storz     | 3375797 | - | 0.7526635 | inactive |
| TSS_12898_storz     | 3375759 | - | 0.7679535 | inactive |
| TSS_12895_regulondb | 3375504 | - | 1.5451355 | active   |
| TSS_12896_wanner    | 3375587 | - | 1.9223959 | 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.