| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| sohB | + | 1327356 | 1328405 |
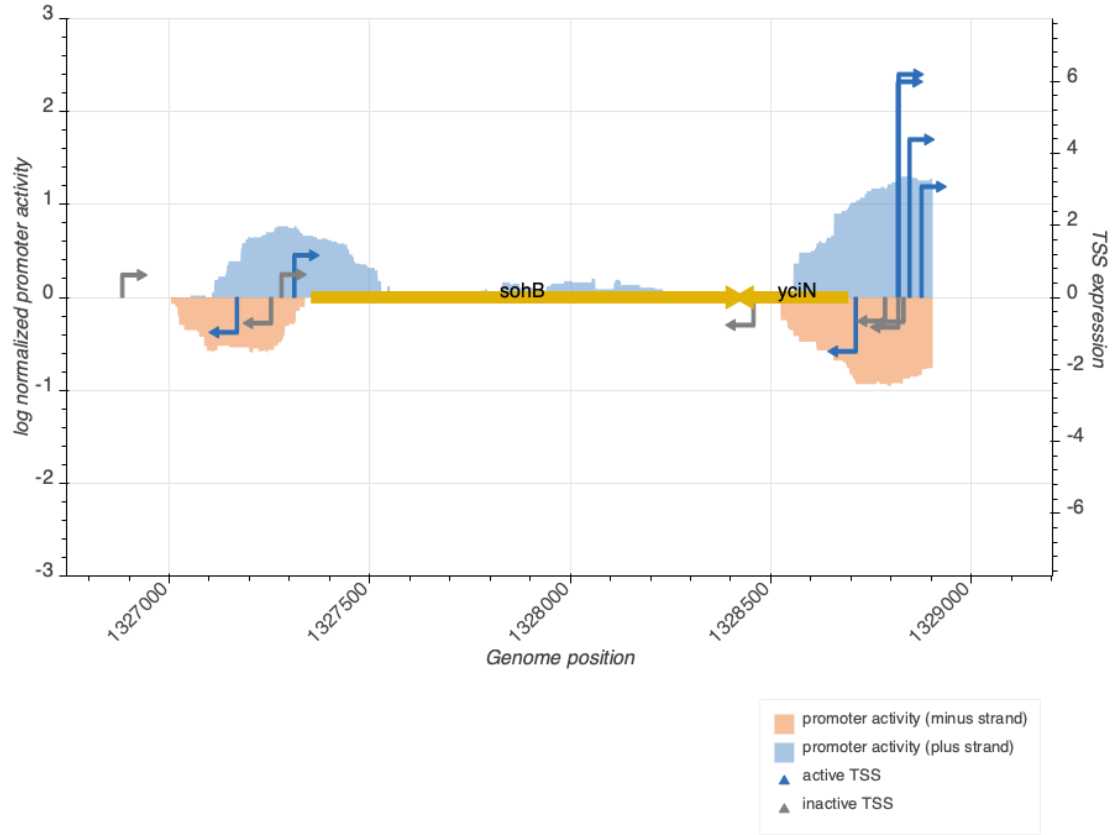


Figure 1: Figure 1: Promoter activity surrounding query operon.

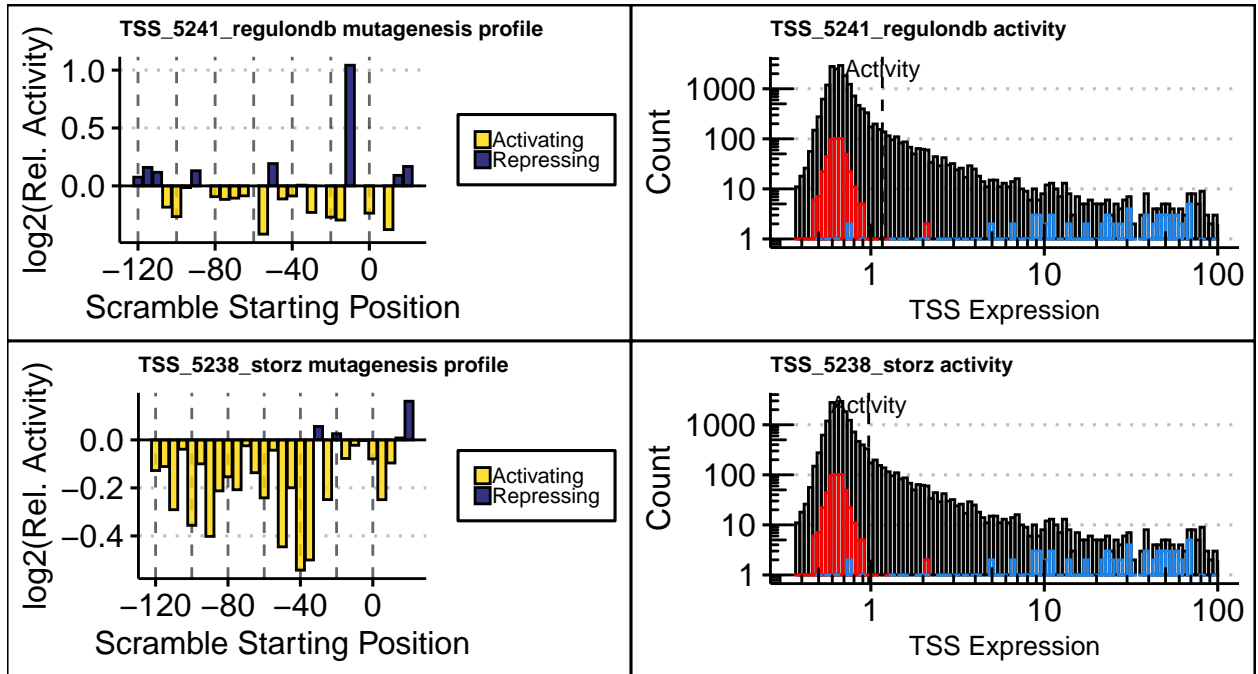| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_5241_regulondb | 1327308 | + | 1.1655919 | active |
| TSS_5240_wanner | 1327276 | + | 0.6320261 | inactive |
| TSS_5239_wanner | 1327250 | - | 0.7158358 | inactive |
| TSS_5237_storz | 1326879 | + | 0.6188789 | inactive |
| TSS_5238_storz | 1327165 | - | 0.9718203 | 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.