| Operon | Strand | Operon start | Operon end |
|--------------------------|--------|--------------|------------|
| ribF-ileS-lspA-fkpB-ispH | + | 21407 | 27227 |

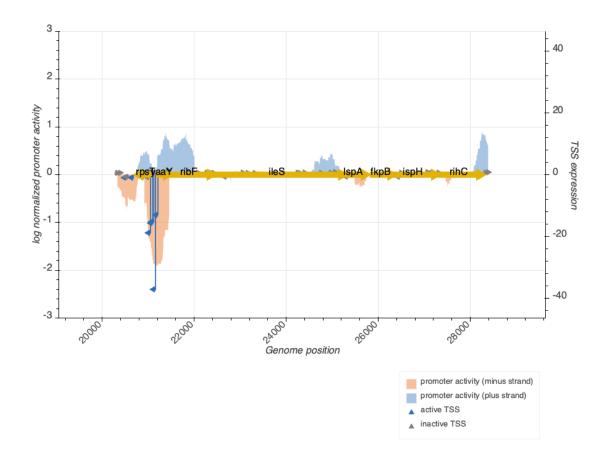


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|------------------------------|--------------|--------|--------------|----------|
| TSS_129_storz_wanner | 21207 | _ | 13.0686488 | active |
| TSS_122_storz | 21007 | + | 0.7030717 | inactive |
| TSS_128_regulondb | 21155 | - | 37.1633178 | active |
| $TSS_124_regulondb$ | 21045 | - | 18.8422646 | active |
| TSS_121_storz | 20984 | + | 0.8698869 | inactive |
| TSS_126_storz | 21113 | + | 0.7366829 | inactive |
| TSS_123_regulondb | 21020 | - | 0.7880935 | inactive |
| TSS_127_wanner | 21118 | - | 15.2200980 | active |
| $TSS_120_{\text{regulondb}}$ | 20910 | - | 0.6359097 | inactive |
| TSS_130_storz_regulondb | 21383 | + | 1.3422067 | active |
| $TSS_125_regulondb$ | 21094 | - | 15.6700183 | active |

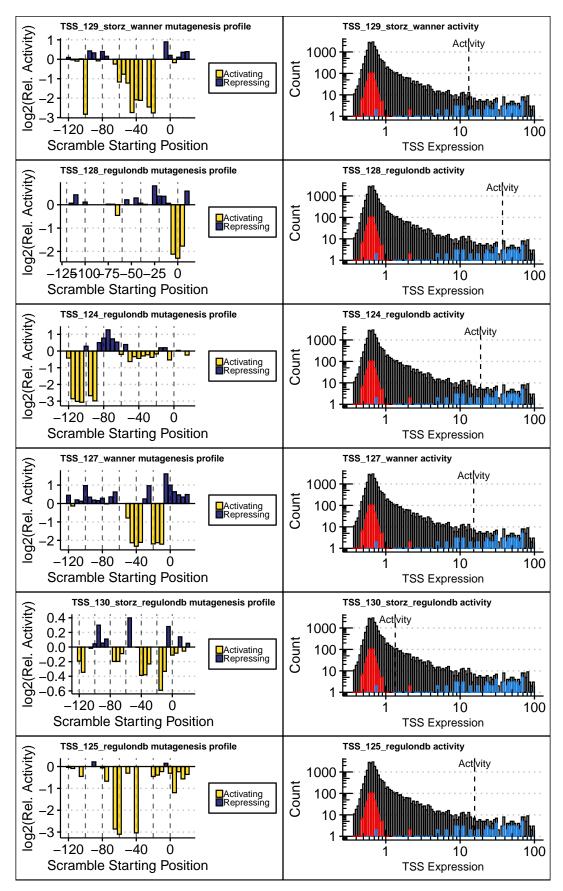


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. T\$Ss were integrated into the nth-ydgR intergenic locus for quantification.