

| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| tig    | +      | 454357       | 455655     |

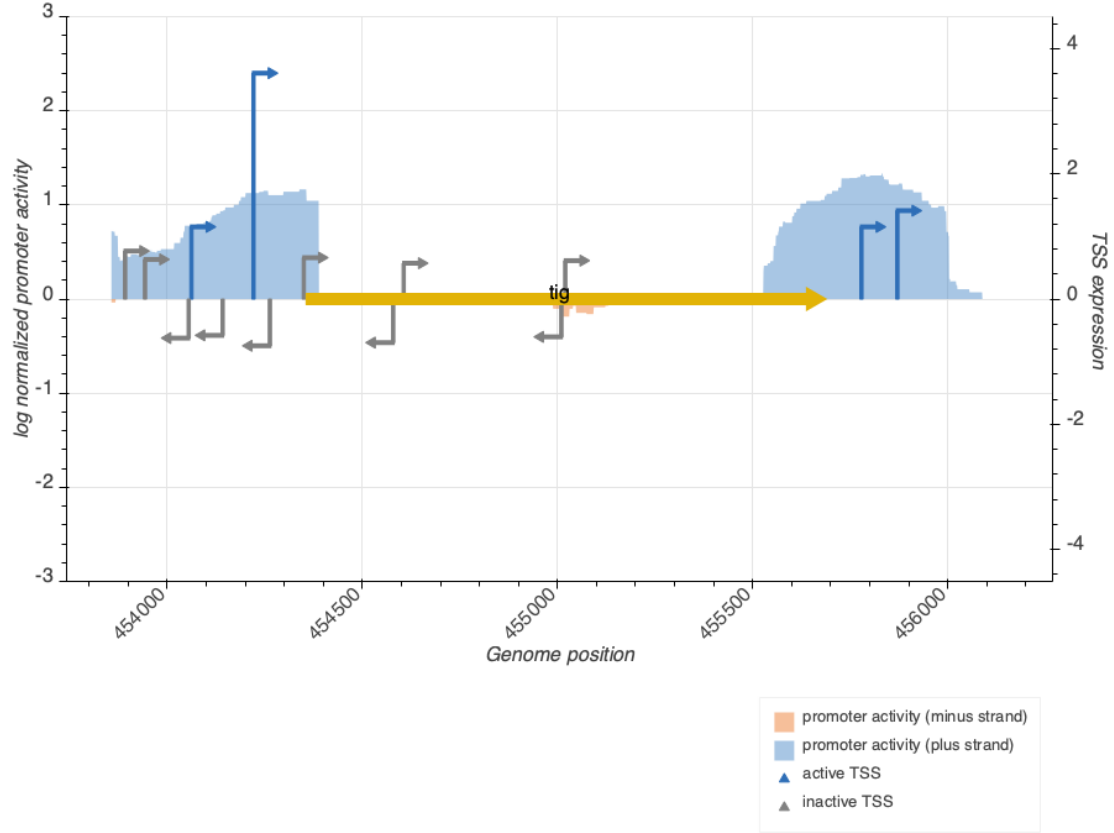


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                 | TSS position | Strand | TSS activity | Category |
|--------------------------|--------------|--------|--------------|----------|
| TSS_1798_storz           | 454051       | -      | 0.6235306    | inactive |
| TSS_1803_regulondb       | 454346       | +      | 0.6613027    | inactive |
| TSS_1797_storz           | 453939       | +      | 0.6331588    | inactive |
| TSS_1799_storz           | 454058       | +      | 1.1534318    | active   |
| TSS_1801_storz_regulondb | 454217       | +      | 3.6091553    | active   |
| TSS_1800_storz           | 454138       | -      | 0.5800841    | inactive |
| TSS_1802_storz           | 454259       | -      | 0.7466460    | inactive |
| TSS_1796_storz_regulondb | 453888       | +      | 0.7669279    | inactive |

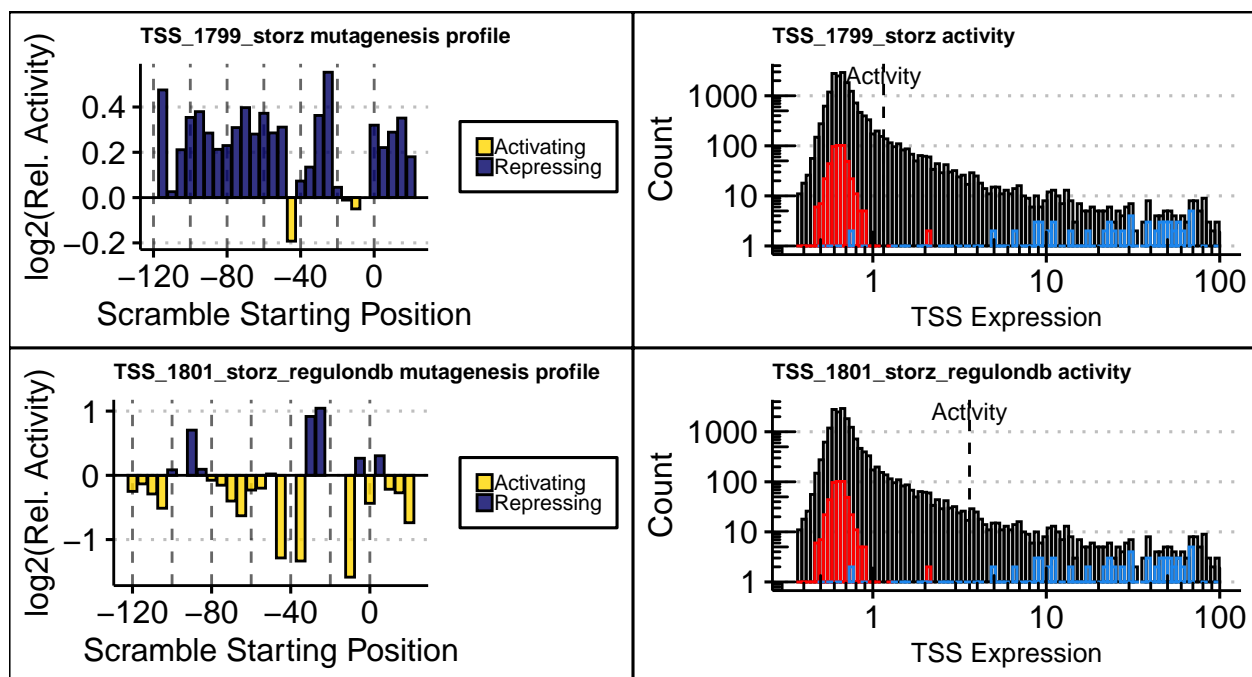


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the *nth-ydgR* intergenic locus for quantification.