

| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| serX | - | 1096875 | 1096788 |

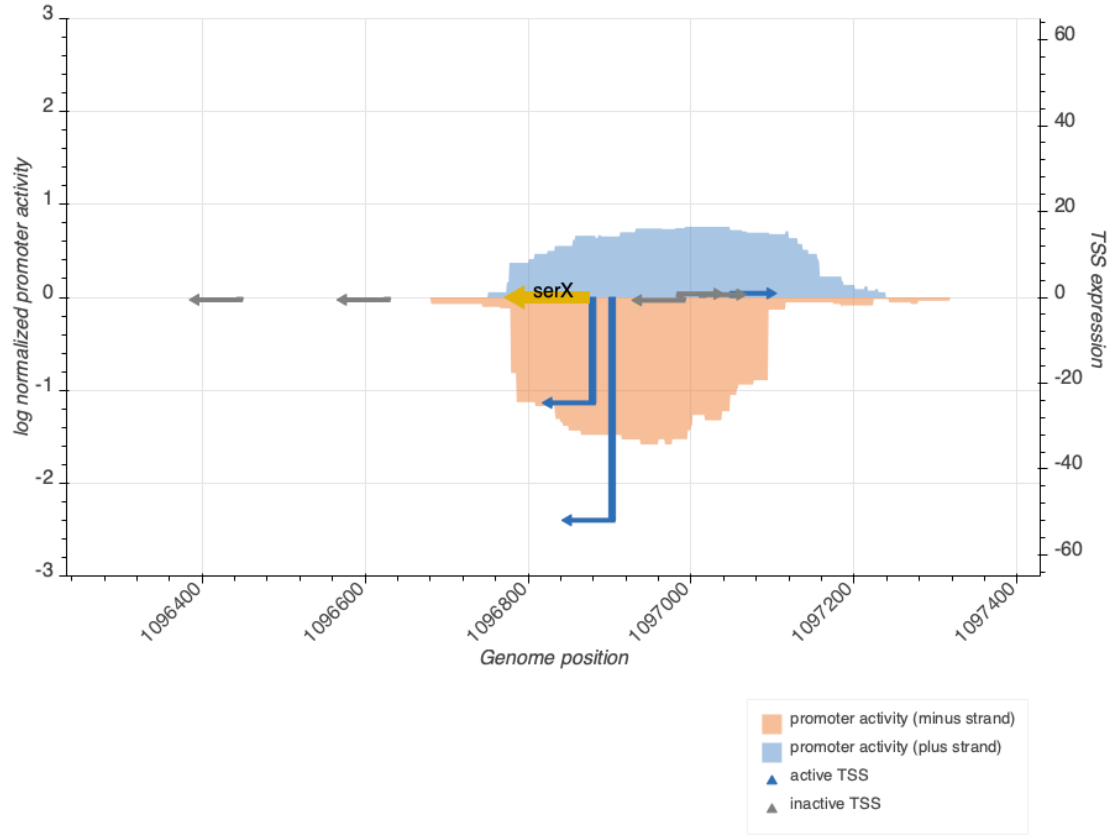


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|---------------------------|--------------|--------|--------------|----------|
| TSS_4222_regulondb | 1097010 | + | 0.6293075 | inactive |
| TSS_4220_wanner_regulondb | 1096982 | + | 0.8094255 | inactive |
| TSS_4219_regulondb | 1096899 | - | 51.9400171 | active |
| TSS_4223_regulondb | 1097047 | + | 0.9347025 | active |
| TSS_4221_storz | 1096985 | - | 0.6918560 | inactive |

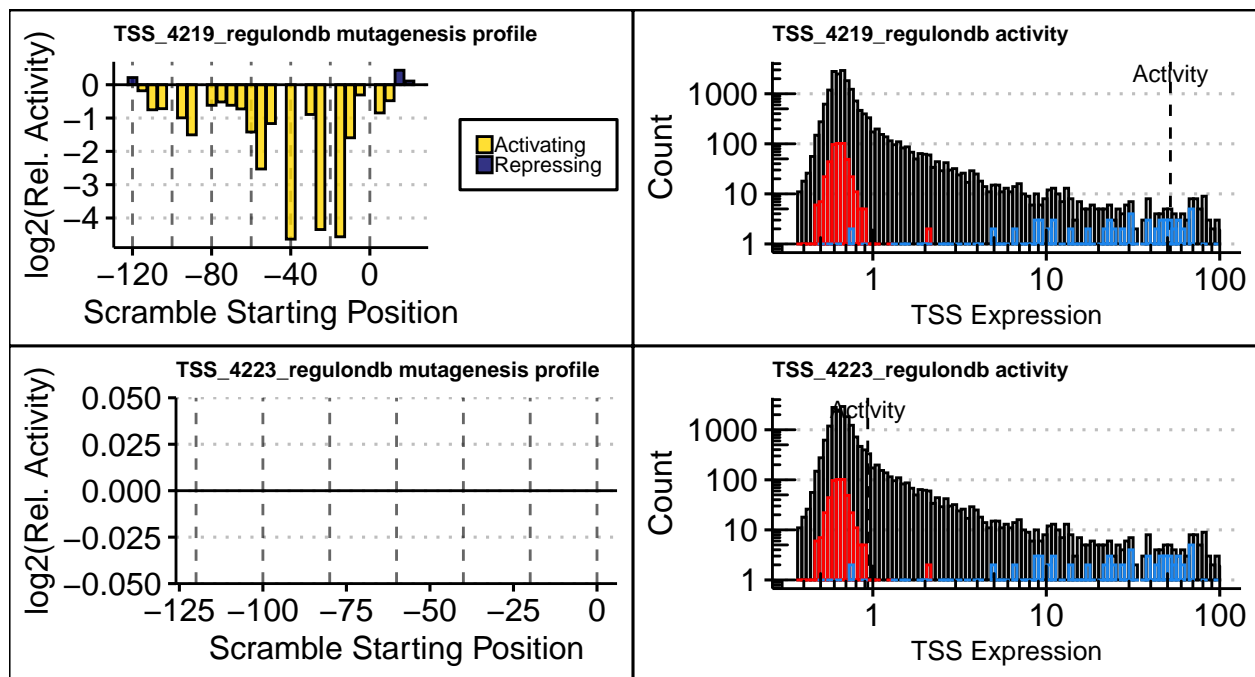


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the *nth-ydgR* intergenic locus for quantification.