| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| mmuPM | + | 274549 | 276871 |
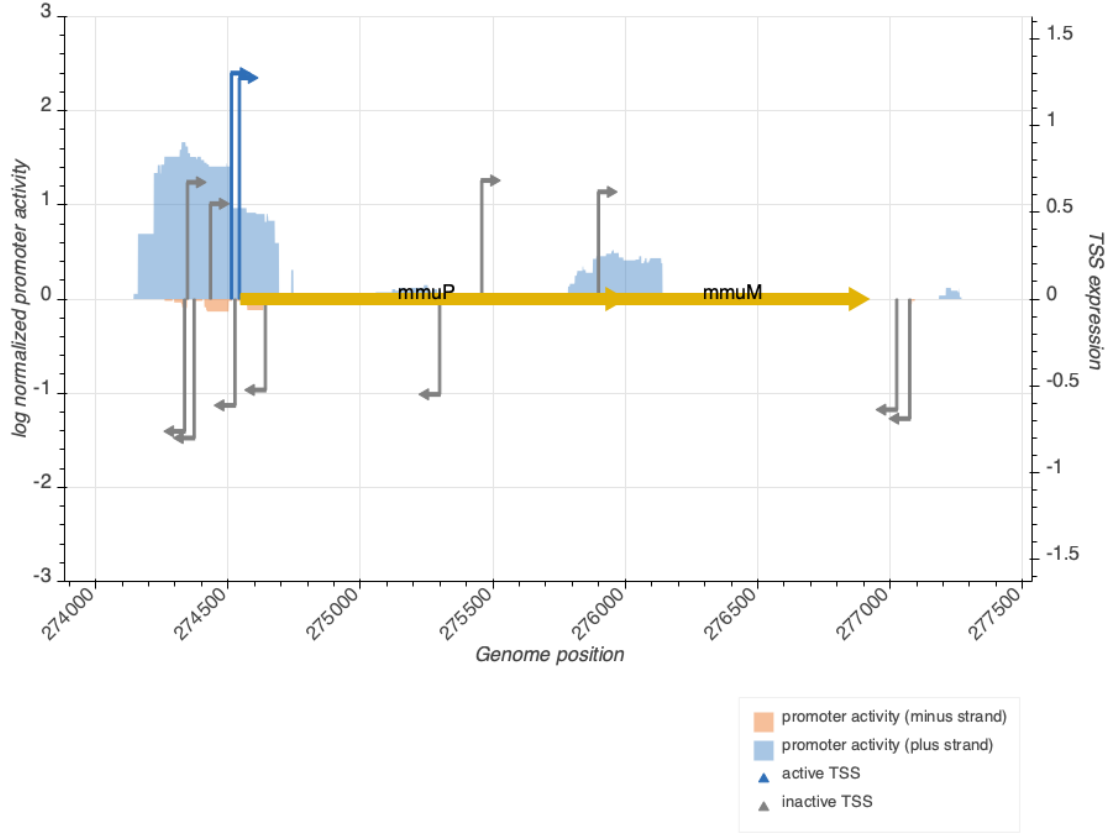


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_1173_storz | 274368 | - | 0.8024896 | inactive |
| TSS_1172_regulondb | 274343 | + | 0.6742580 | inactive |
| TSS_1177_regulondb | 274539 | + | 1.2773845 | active |
| TSS_1176_regulondb | 274522 | - | 0.6139024 | inactive |
| TSS_1175_regulondb | 274509 | + | 1.3043211 | active |
| TSS_1171_storz_regulondb | 274332 | - | 0.7638980 | inactive |
| TSS_1174_regulondb | 274430 | + | 0.5504818 | inactive |
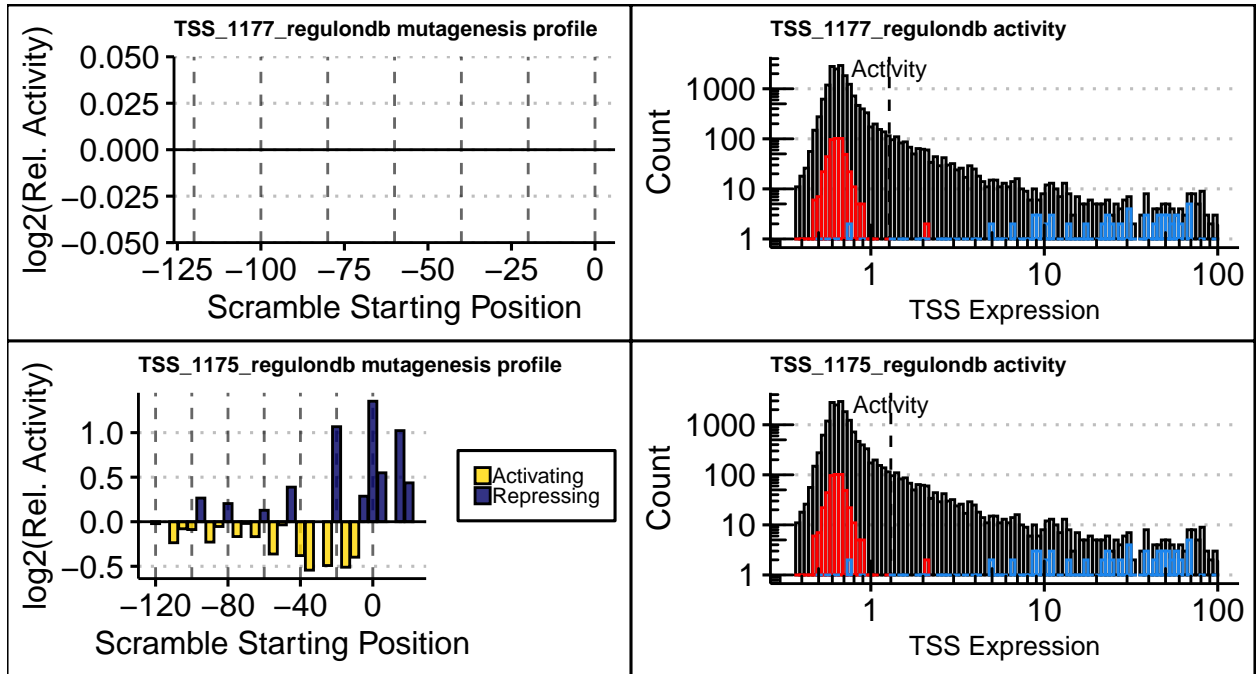
Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.