| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| lptFG | + | 4484241 | 4486423 |

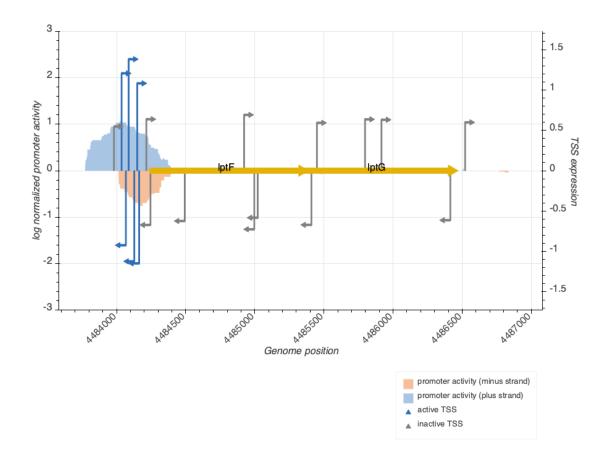


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|-------------------------|--------------|--------|--------------|----------|
| TSS_17540_regulondb | 4484145 | + | 1.0817341 | active |
| TSS_17536_storz | 4484032 | + | 1.2057975 | active |
| TSS_17538_storz | 4484083 | + | 1.3809351 | active |
| TSS_17541_storz | 4484159 | - | 1.1481961 | active |
| $TSS_17539_{regulondb}$ | 4484124 | - | 1.1234121 | active |
| TSS_17535_storz | 4483977 | + | 0.5468420 | inactive |
| TSS_17542 _regulondb | 4484211 | + | 0.6373126 | inactive |
| $TSS_17537_regulondb$ | 4484063 | - | 0.9237764 | active |

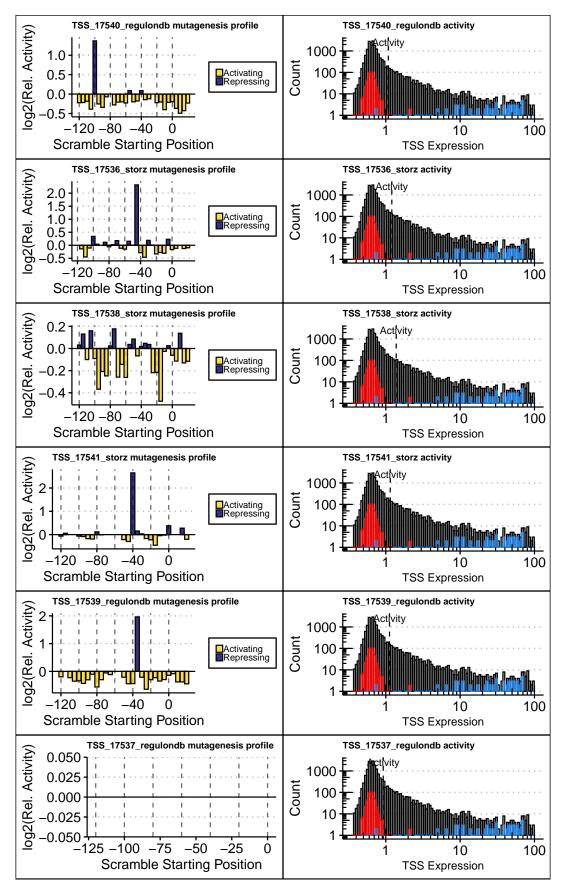


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. T\$Ss were integrated into the nth-ydgR intergenic locus for quantification.