| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| dtpD | - | 741779 | 740298 |

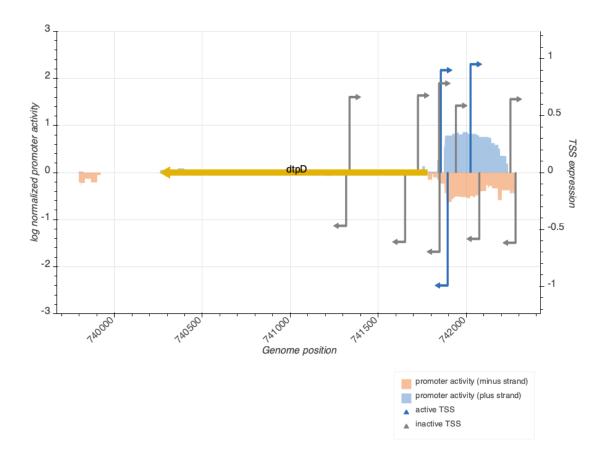


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|--------------------------|--------------|--------|--------------|----------|
| TSS_2838_regulondb | 741842 | + | 0.7828982 | inactive |
| TSS_2844_storz | 742071 | - | 0.5830922 | inactive |
| TSS_2839_regulondb | 741845 | - | 0.6961009 | inactive |
| TSS_2842 _wanner | 741938 | + | 0.5874936 | inactive |
| TSS_2840_storz | 741852 | + | 0.8990107 | active |
| TSS_2846_storz | 742277 | - | 0.6170700 | inactive |
| TSS_2845_storz | 742248 | + | 0.6442553 | inactive |
| TSS_2841_wanner | 741891 | - | 0.9922070 | active |
| TSS_2843_storz_regulondb | 742021 | + | 0.9514336 | active |

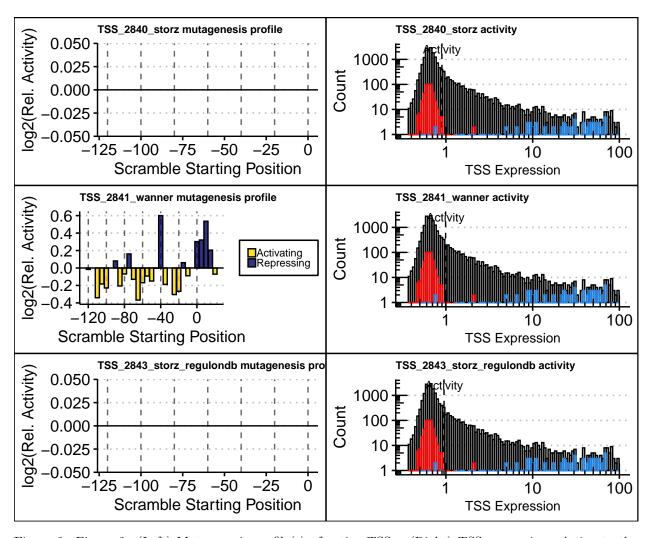


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.