| Operon                    | Strand | Operon start | Operon end |
|---------------------------|--------|--------------|------------|
| lptD-surA-pdxA-rsmA-apaGH | -      | 57109        | 50380      |

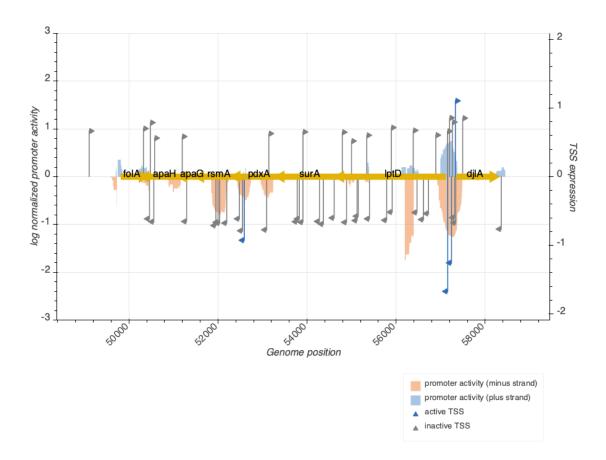


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                | TSS position | Strand | TSS activity | Category |
|-------------------------|--------------|--------|--------------|----------|
| TSS_249_storz           | 57311        | -      | 0.5962989    | inactive |
| $TSS\_248\_storz$       | 57259        | +      | 0.7957535    | inactive |
| $TSS_250$ _wanner       | 57327        | +      | 1.1054847    | active   |
| TSS_245_storz_regulondb | 57156        | -      | 1.6714832    | active   |
| TSS_251_regulondb       | 57336        | -      | 0.6733660    | inactive |
| $TSS\_252\_storz$       | 57492        | +      | 0.8535776    | inactive |
| TSS_247_storz_regulondb | 57241        | -      | 1.2556048    | active   |
| $TSS\_244\_storz$       | 57150        | +      | 0.6622984    | inactive |
| TSS_246_storz           | 57197        | +      | 0.8578889    | inactive |

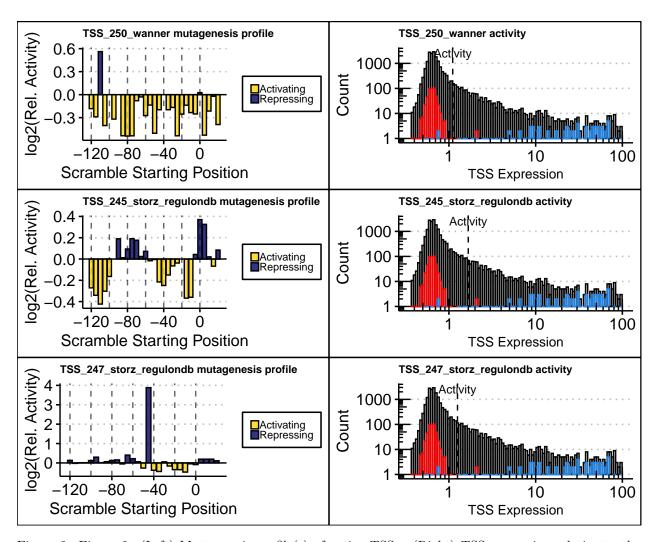


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.