| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| proBA  | +      | 259612       | 261980     |

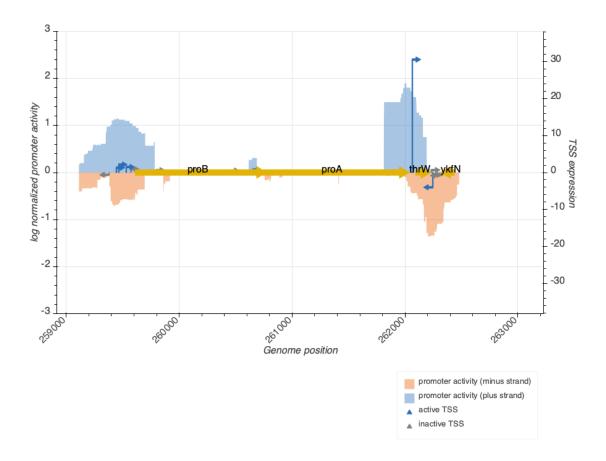


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name               | TSS position | Strand | TSS activity | Category |
|------------------------|--------------|--------|--------------|----------|
| TSS_1120_regulondb     | 259569       | +      | 0.8698869    | inactive |
| $TSS_1116$ _wanner     | 259374       | -      | 0.6139024    | inactive |
| $TSS\_1118\_storz$     | 259461       | +      | 2.1606908    | active   |
| TSS_1117_regulondb     | 259440       | +      | 1.3069773    | active   |
| $TSS\_1119\_regulondb$ | 259527       | +      | 1.5020866    | active   |

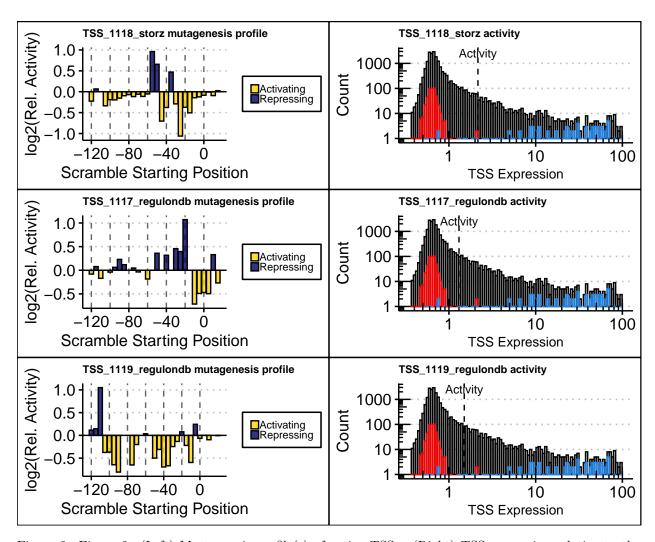


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.