| Operon | Strand | Operon start | Operon end |  |
|--------|--------|--------------|------------|--|
| epmA   | +      | 4380666      | 4381643    |  |

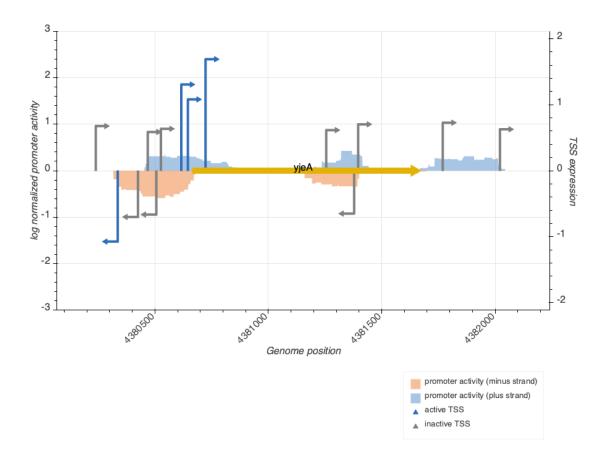


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                         | TSS position | Strand | TSS activity | Category |
|----------------------------------|--------------|--------|--------------|----------|
| TSS_17118_storz                  | 4380464      | +      | 0.5861024    | inactive |
| TSS_17119_wanner                 | 4380502      | -      | 0.6648691    | inactive |
| TSS_17115_storz                  | 4380235      | +      | 0.6755228    | inactive |
| $TSS\_17121\_storz$              | 4380612      | +      | 1.3048303    | active   |
| TSS_17122_storz_wanner_regulondb | 4380641      | +      | 1.0791152    | active   |
| $TSS\_17120\_storz$              | 4380522      | +      | 0.6338880    | inactive |
| TSS_17117_regulondb              | 4380421      | -      | 0.7010062    | inactive |
| $TSS_17116_{regulondb}$          | 4380332      | -      | 1.0737537    | active   |

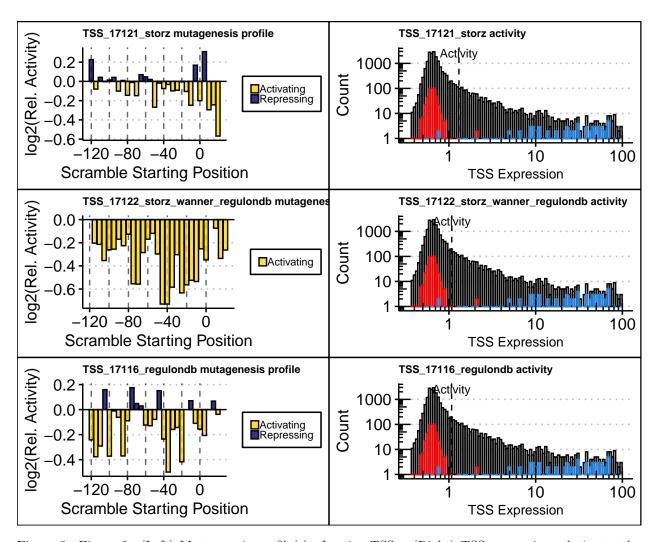


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.