| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| yjbLM  | +      | 4258344      | 4259329    |

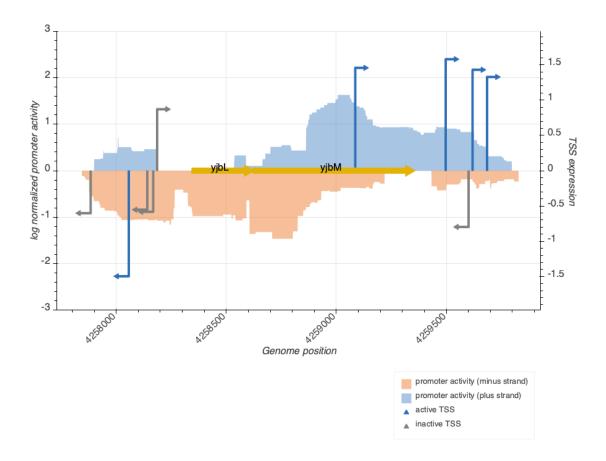


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                | TSS position | Strand | TSS activity | Category |
|-------------------------|--------------|--------|--------------|----------|
| TSS_16674_regulondb     | 4257880      | -      | 0.6023692    | inactive |
| $TSS\_16675$ _wanner    | 4258053      | -      | 1.4963635    | active   |
| TSS_16677_regulondb     | 4258164      | -      | 0.5824313    | inactive |
| $TSS_16678_{regulondb}$ | 4258182      | +      | 0.8706269    | inactive |
| $TSS\_16676\_regulondb$ | 4258137      | -      | 0.5533823    | inactive |

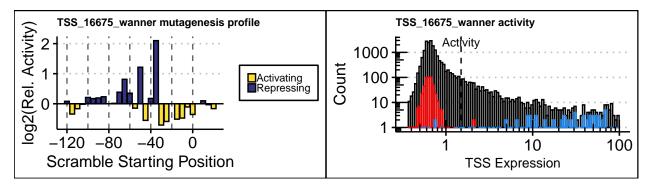


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.