| Operon          | Strand | Operon start | Operon end |
|-----------------|--------|--------------|------------|
| lapAB-pyrF-yciH | +      | 1338267      | 1341008    |

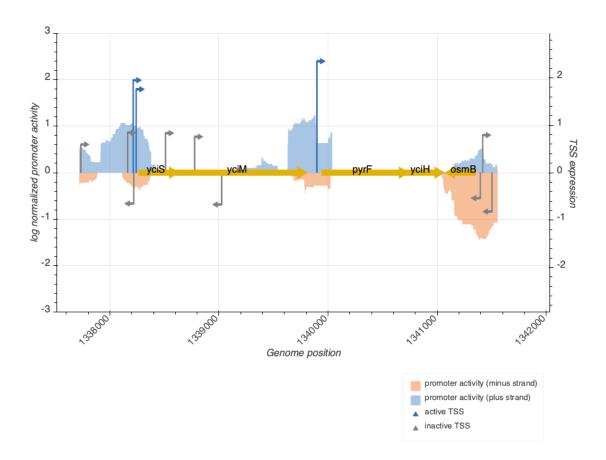


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                 | TSS position | Strand | TSS activity | Category |
|--------------------------|--------------|--------|--------------|----------|
| TSS_5300_storz_regulondb | 1338159      | +      | 0.8444035    | inactive |
| $TSS\_5302\_storz$       | 1338215      | -      | 0.6524152    | inactive |
| $TSS\_5301\_regulondb$   | 1338210      | +      | 1.9508267    | active   |
| $TSS\_5303\_regulondb$   | 1338237      | +      | 1.7590886    | active   |

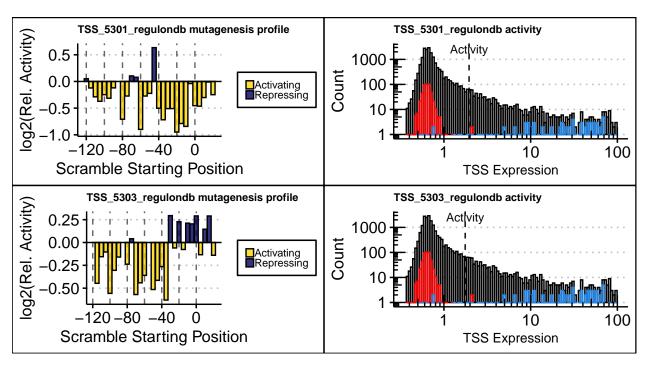


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.