| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| ybdJ | - | 605422 | 605174 |

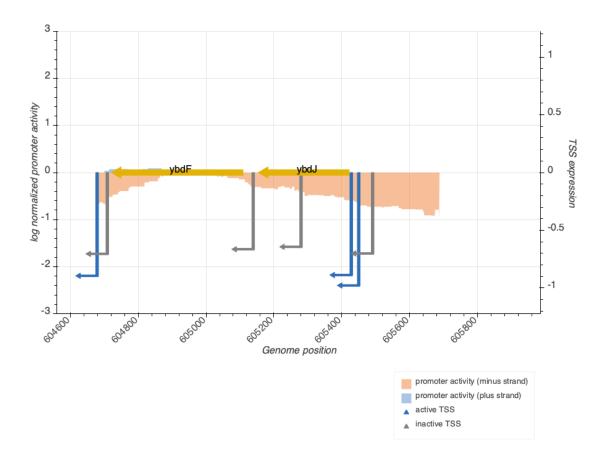


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|--------------------|--------------|--------|--------------|----------|
| TSS_2336_regulondb | 605487 | - | 0.7013527 | active |
| TSS_2335_regulondb | 605446 | - | 0.9788620 | |
| TSS_2334_wanner | 605424 | - | 0.8885093 | |

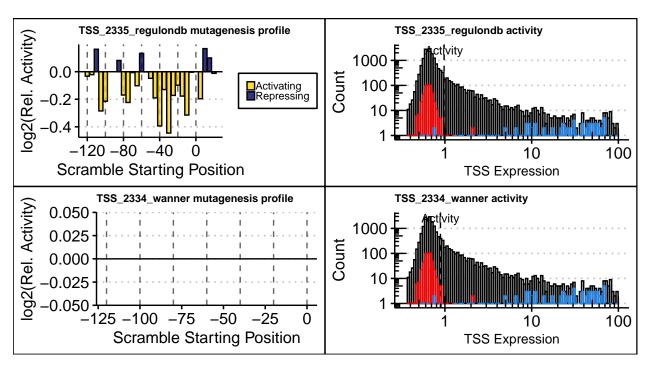


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.