| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| gdhA | + | 1840395 | 1841738 |
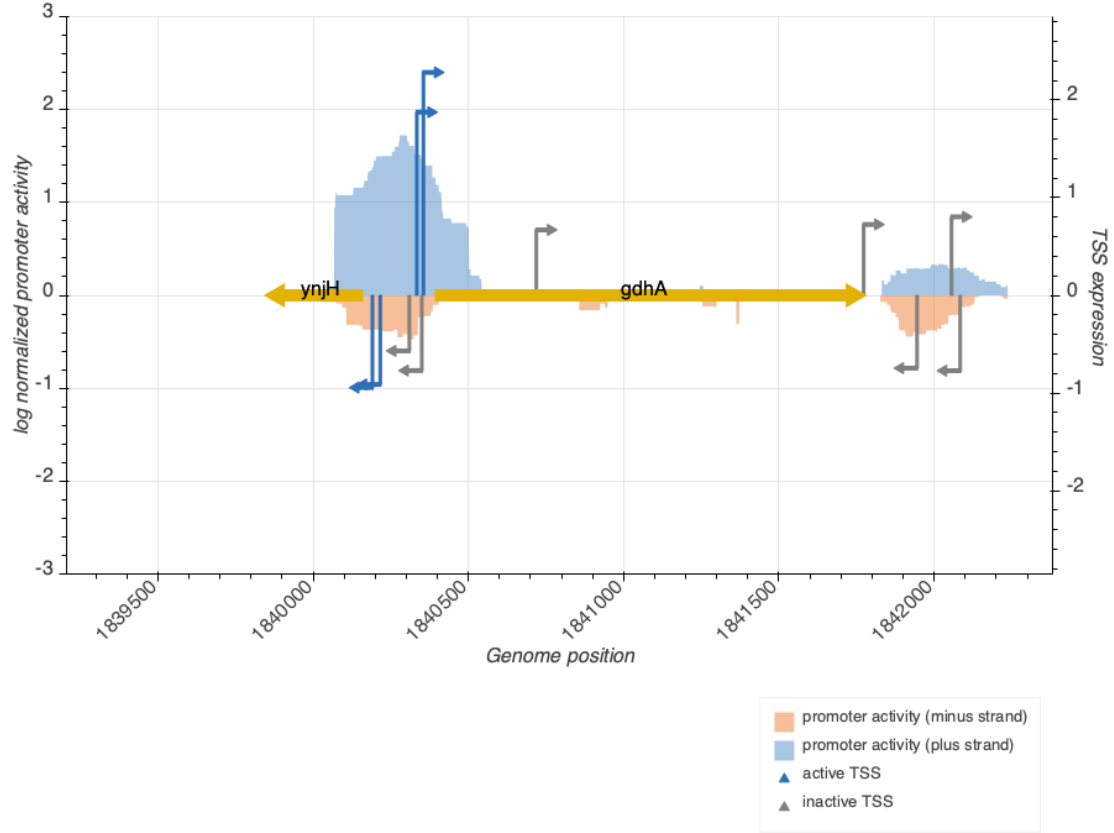


Figure 1: Figure 1: Promoter activity surrounding query operon.

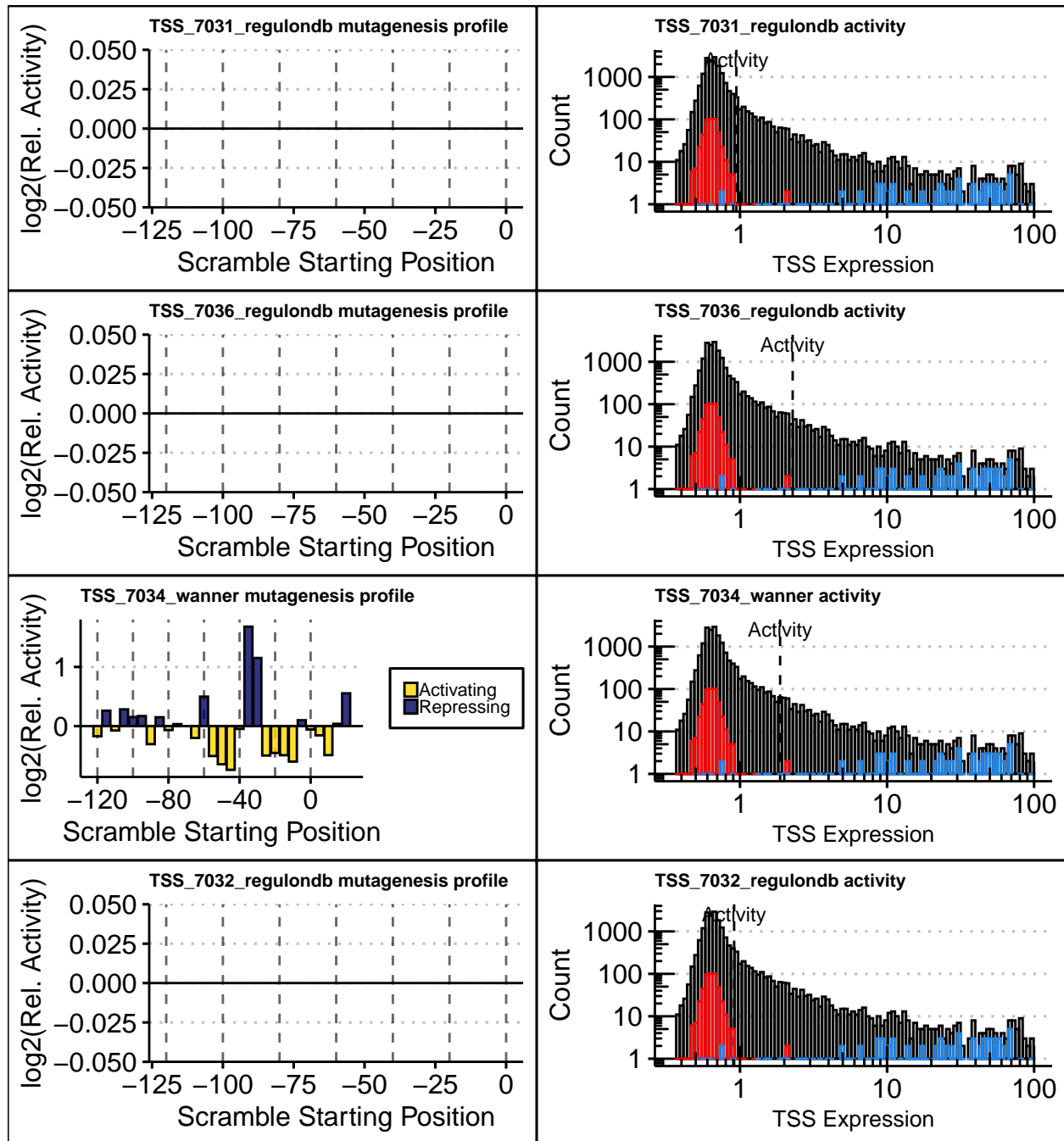| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_7031_regulondb | 1840187 | - | 0.9446381 | active |
| TSS_7036_regulondb | 1840351 | + | 2.2820156 | active |
| TSS_7035_storz | 1840346 | - | 0.7703292 | inactive |
| TSS_7034_wanner | 1840330 | + | 1.8747182 | active |
| TSS_7032_regulondb | 1840212 | - | 0.9124742 | active |
| TSS_7033_storz | 1840306 | - | 0.5686696 | 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.