| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| hicAB  | +      | 1507310      | 1507948    |
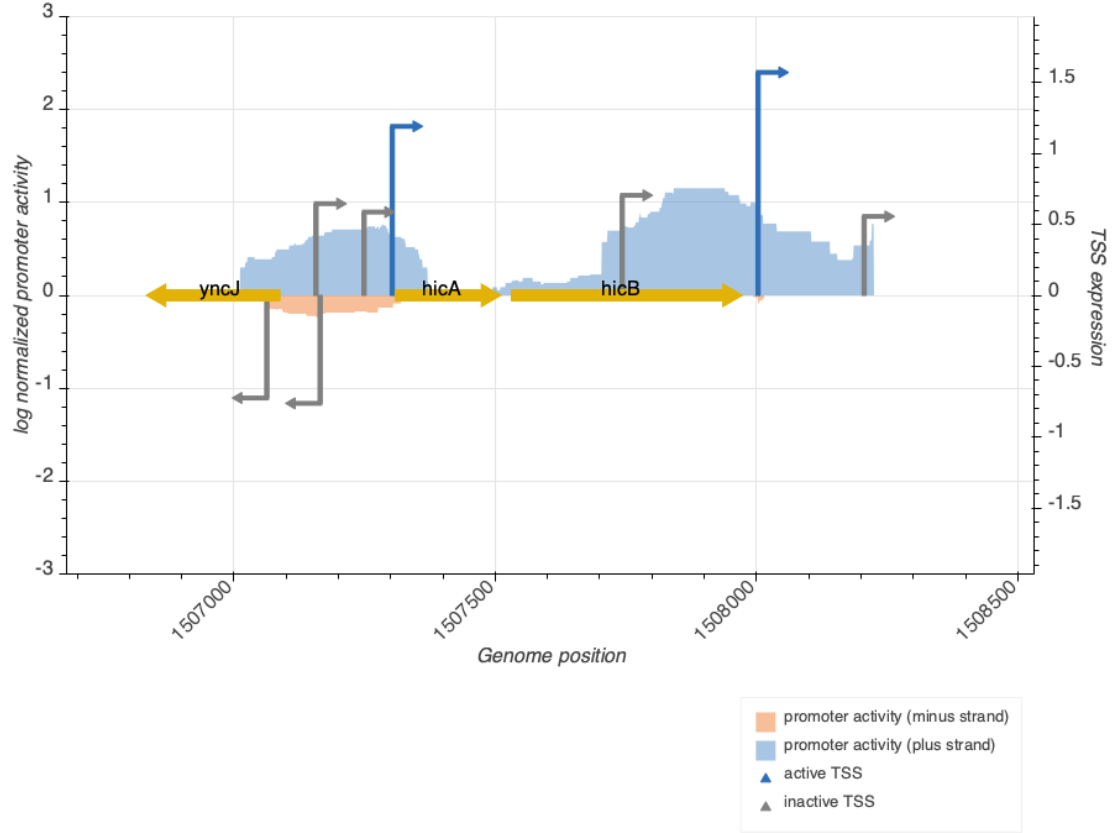


Figure 1: Figure 1: Promoter activity surrounding query operon.

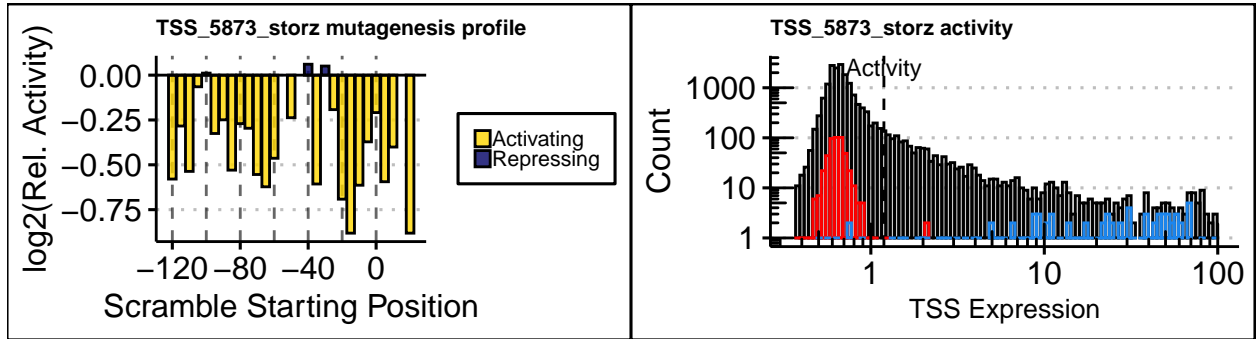| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_5870_storz | 1507153 | + | 0.6451065 | inactive |
| TSS_5873_storz | 1507299 | + | 1.1895994 | active |
| TSS_5872_regulondb | 1507245 | + | 0.5861024 | inactive |
| TSS_5869_storz_regulondb | 1507059 | - | 0.7223869 | inactive |
| TSS_5871_wanner | 1507161 | - | 0.7602557 | 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.