| Operon | Strand | Operon start | Operon end |
|--------|--------|--------------|------------|
| fabA   | -      | 1015693      | 1015175    |
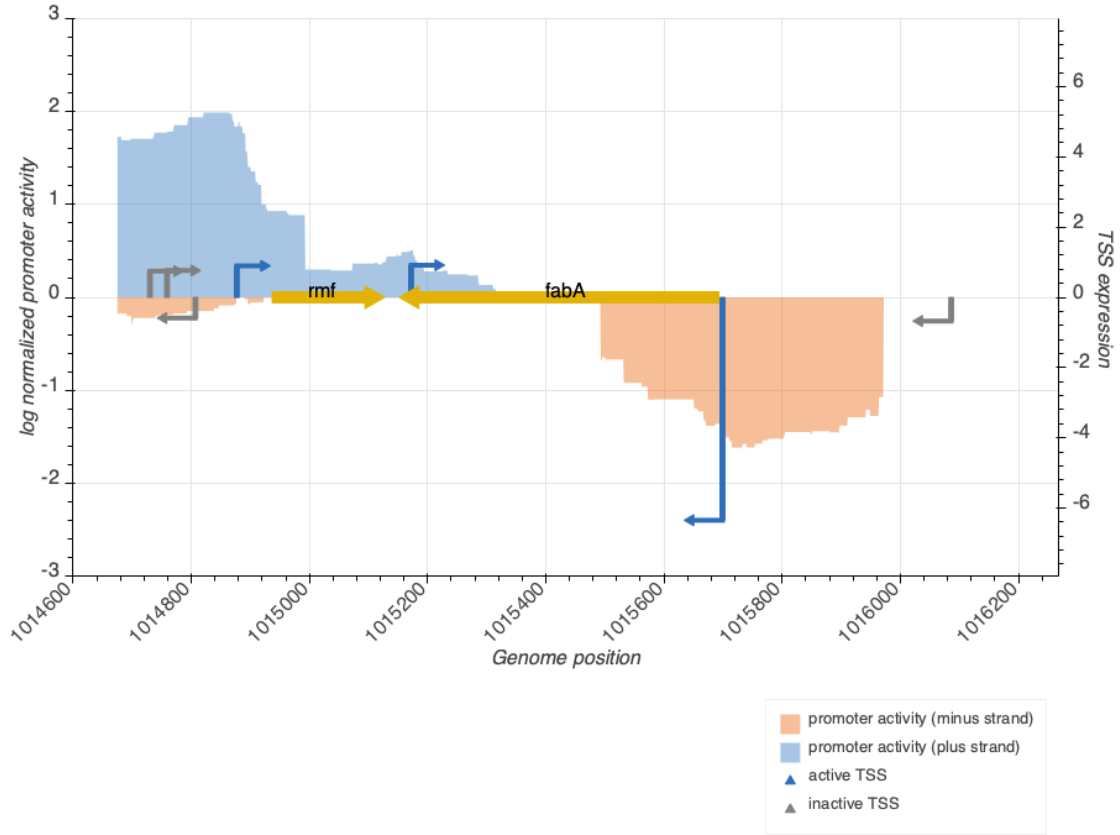


Figure 1: Figure 1: Promoter activity surrounding query operon.

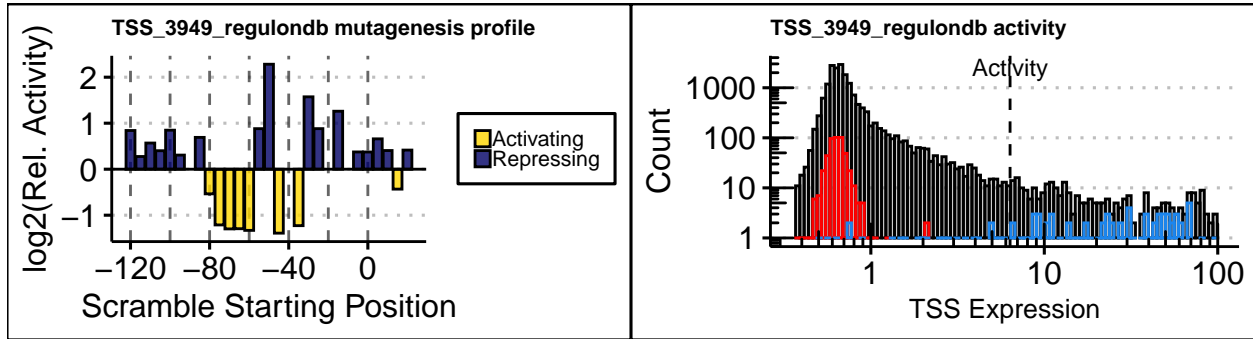| TSS name | TSS position | Strand | TSS activity | Category |
|----------|--------------|--------|--------------|----------|
| TSS_3950_regulondb | 1016082 | - | 0.6750625 | inactive |
| TSS_3949_regulondb | 1015695 | - | 6.3480034 | active |

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.