| Operon            | Strand | Operon start | Operon end |
|-------------------|--------|--------------|------------|
| ivbL-ilvBN-uhpABC | -      | 3851011      | 3845328    |

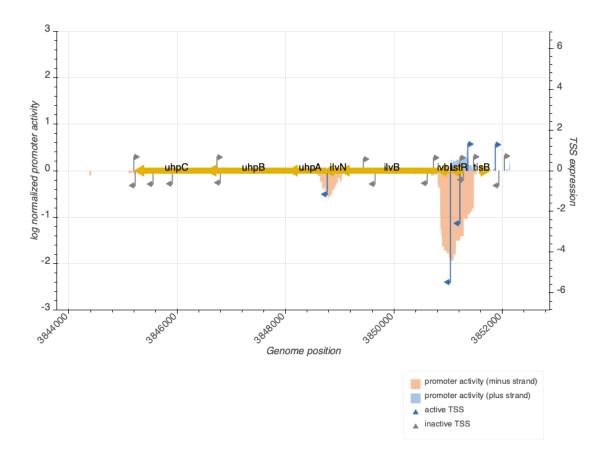


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name                         | TSS position | Strand | TSS activity | Category |
|----------------------------------|--------------|--------|--------------|----------|
| TSS_14926_storz                  | 3851216      | +      | 0.6357894    | inactive |
| $TSS\_14928\_regulondb$          | 3851360      | +      | 1.3048303    | active   |
| $TSS\_14927$ _wanner             | 3851279      | -      | 0.4450457    | inactive |
| $TSS_14929_{regulondb}$          | 3851465      | +      | 0.6820697    | inactive |
| TSS_14924_storz_wanner_regulondb | 3851039      | -      | 5.4677714    | active   |
| $TSS\_14925\_storz\_regulondb$   | 3851215      | -      | 2.5822146    | active   |

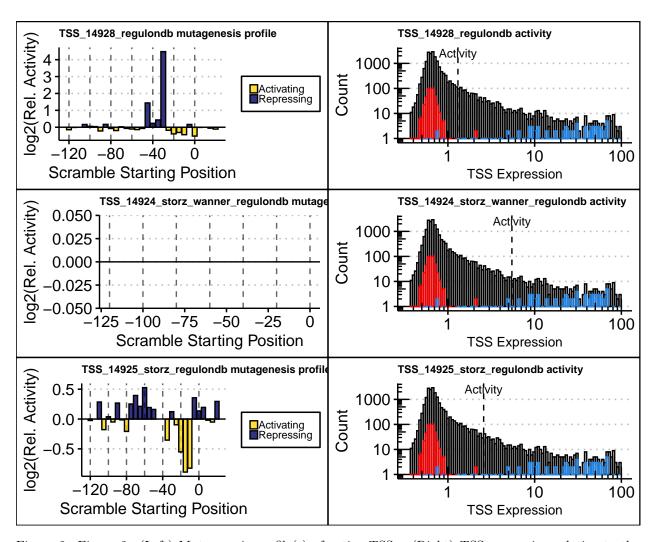


Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the nth-ydgR intergenic locus for quantification.