

| Operon | Strand | Operon start | Operon end |
|---------|--------|--------------|------------|
| hemCDXY | - | 3988789 | 3984709 |

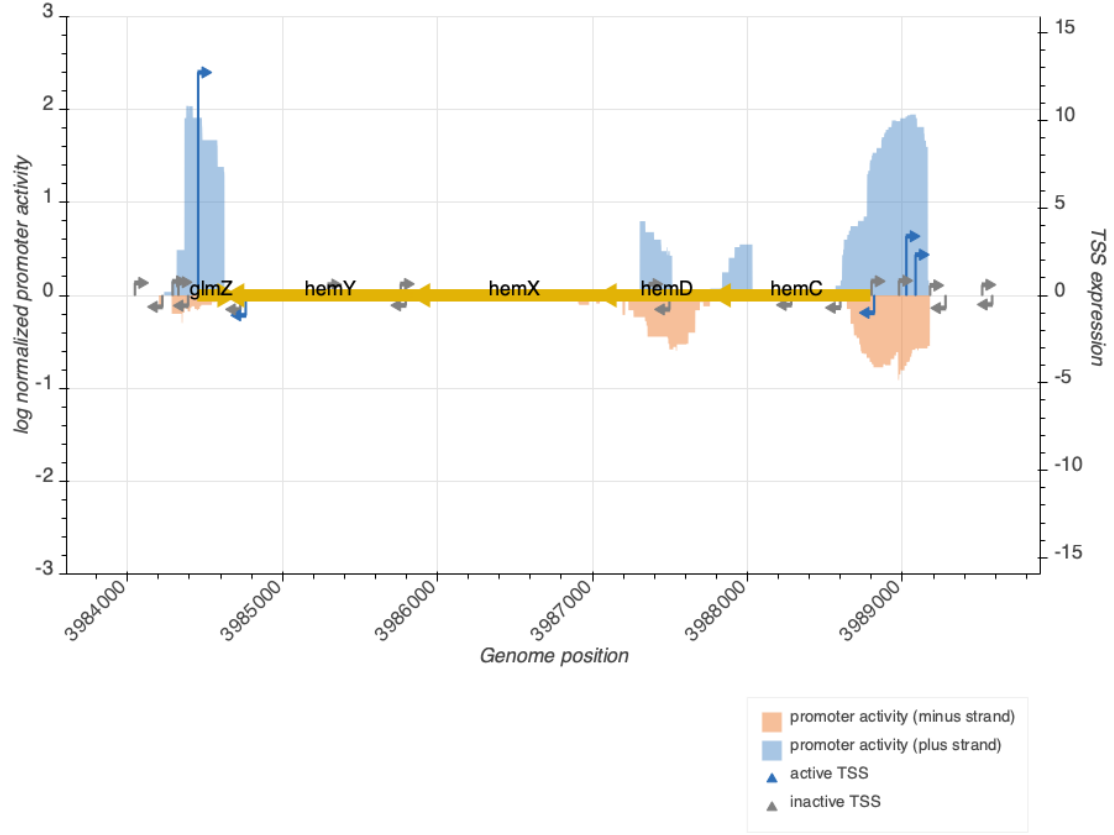


Figure 1: Figure 1: Promoter activity surrounding query operon.

| TSS name | TSS position | Strand | TSS activity | Category |
|---------------------------|--------------|--------|--------------|----------|
| TSS_15503_storz_regulondb | 3988797 | + | 0.8077013 | inactive |
| TSS_15509_regulondb | 3989175 | + | 0.5681475 | inactive |
| TSS_15507_storz_regulondb | 3989022 | + | 3.3753123 | active |
| TSS_15506_storz_regulondb | 3988975 | + | 0.8375499 | inactive |
| TSS_15510_storz | 3989274 | - | 0.7288652 | inactive |
| TSS_15504_regulondb | 3988814 | - | 0.9973037 | active |
| TSS_15508_regulondb | 3989081 | + | 2.3230045 | active |

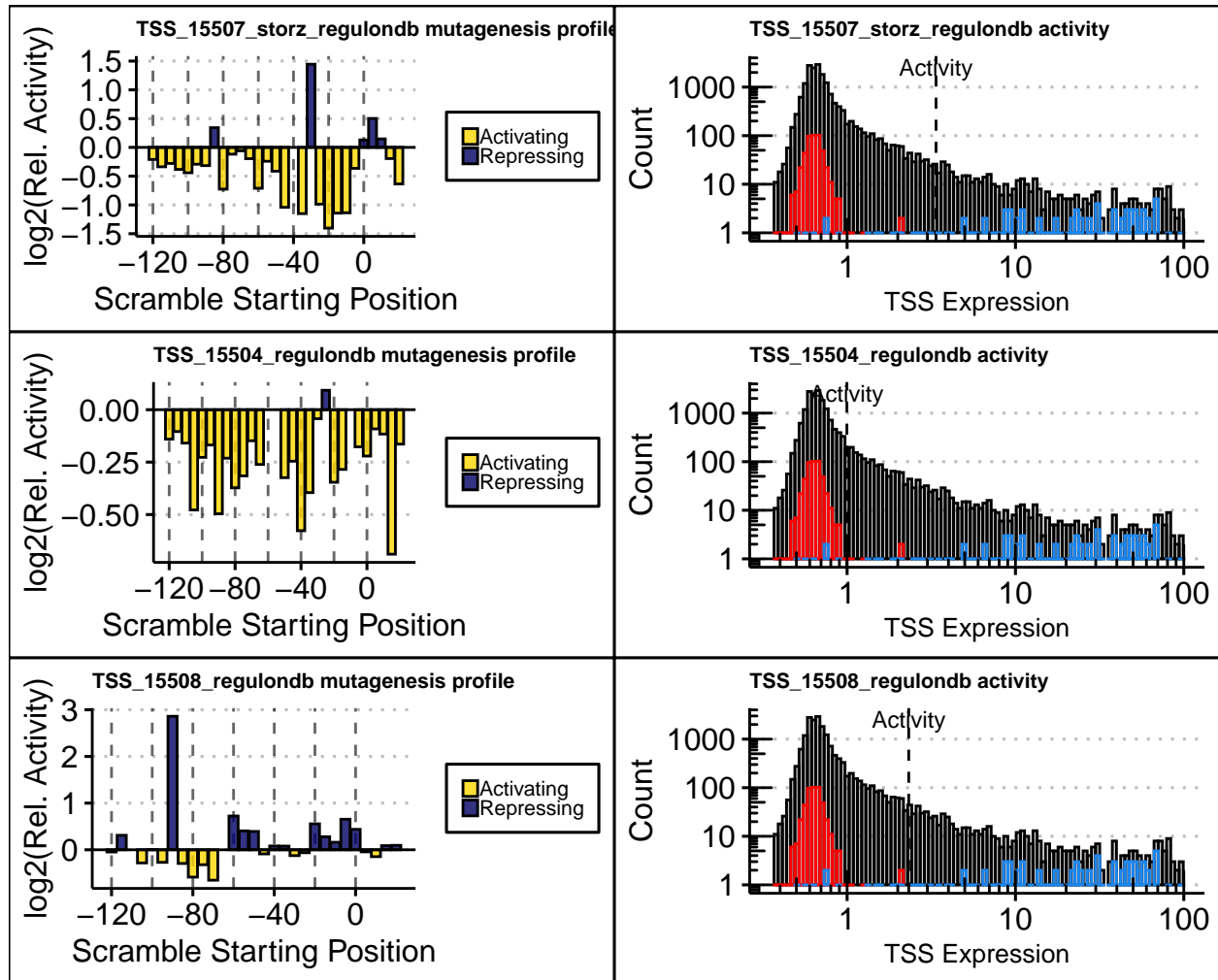


Figure 2: Figure 2: (Left) Mutagenesis profile(s) of active TSSs. (Right) TSS expression relative to the distribution of expression levels for all tested TSSs. TSSs were integrated into the *nth-ydgR* intergenic locus for quantification.