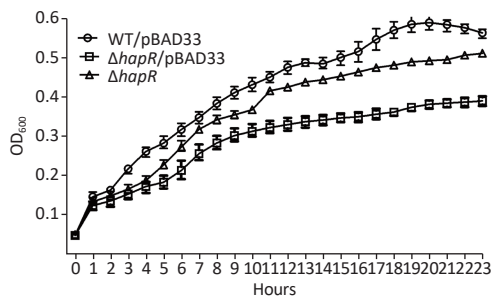


Supplementary Table S1. Oligonucleotide primers used in this study

| Target | Primers (forward/reverse, 5'-3') |
|-----------------------------|---|
| Protein expression | |
| <i>hapR</i> | GCGGGATCCATGGACGCATCAATCGAAAAAC/GCGAAGCTTCTAGTCTTATAGATACACAG |
| DNase I footprinting | |
| <i>mtlA</i> | GTAAAACGACGGCCAGTTTACGTATAGTGACG/CAGGAAACAGCTATGACGTTGGATGTTCCGTTTG |
| M13 | FAM-GTAAAACGACGGCCAGT/CAGGAAACAGCTATGAC-HEX |
| Luminescence assay | |
| <i>mtlA</i> | GCGGAGCTCGTGCATGACATTATCC/GCGGGATCCCGCGTCCCGGTTGGATG |



Supplementary Figure S1. Growth curves of the *Vibrio cholerae* strains in mannitol fermentation media. WT/pBAD24, *ΔhapR*/pBAD24, and C-*ΔhapR* were statically cultured in mannitol fermentation media at 37°C, and the OD₆₀₀ values for each strain were monitored with a 1-h interval. The experiments were performed at least twice, with three biological replicates for each strain per time.



Supplementary Figure S2. Structural organization of the *mtlADR* promoter. The transcription and translation starts are marked with bent arrows. The Shine-Dalgarno (SD) box and -10/-35 elements are enclosed in boxes. The MQR box-like sequence is enclosed in a yellow shadowed box. The Fur, CRP, and HapR binding sites (references in [5,9] and the current study) are underlined with solid, dotted, and break point lines, respectively.