**S6 Table. Oligonucleotides used in this study.**

|  |  |  |
| --- | --- | --- |
| Oligonucleotide | Oligonucleotide sequence (5’ →3’)*a* | Use |
| For mutant construction | | |
| CABH01\_F | GAGCTCAGGTTACCCGCATGAACGTGACCATCGATGCG | Deletion of *cabH* ORF |
| CABH01\_R | AATCAAACAGCGTTGTTCCAGACATAGATTATTCC |
| CABH02\_F | TGGAACAACGCTGTTTGATTTTGCTTGATAAGTGC |
| CABH02\_R | CGACCCTCGAGTACGCGTCACACTGTTGTGCCGTGACC |
| BRPN01\_F | CTCAGGTTACCCGCATGGTACAGCACTTAGGAGTC | Deletion of *brpN* ORF |
| BRPN01\_R | GGTGATAATCCACGGATGCAGCAAGGCAGTACAACC |
| BRPN02\_F | GGTTGTACTGCCTTGCTGCATCCGTGGATTATCACC |
| BRPN02\_R | CTCGAGTACGCGTCAGTATTGAACAGGCCAATG |
| For mutant complementation | | |
| CABH03\_F | GGCTAGCAGGAGGAATTCACATGTCTGGAACAACGGTTG | Amplification of *cabH* ORF |
| CABH03\_R | AAAACAGCCAAGCTTGCATGAGTGACGATACTCGCACTTATC |
| BRPN03\_F | GGCTAGCAGGAGGAATTCACATGTTCATTCAATCGATACACAAC | Amplification of *brpN* ORF |
| BRPN03\_R | AAAACAGCCAAGCTTGCATGGTATGGAATGATGGCGAAAG |
| For EMSA | | |
| CABHUP\_F | GTCTTACCAAGCTAGACCCGT | Amplification of *cabH* upstream region |
| CABHUP\_R | TGATGATATTGACTGCAAGCGT |
| BRPNUP\_F | GCATCAGTCTAAACACCGCAC | Amplification of *brpN* upstream region |
| BRPNUP\_R | TGGCGATGGCTCTAAAGTTGT |
| For qRT-PCR | | |
| BRPT\_qRT\_F | GAAGCTGTGTCGCGGGATTG | Quantification of *brpT* expression |
| BRPT\_qRT\_R | TGTGGCTCTTCCTTCTTCGCTC |
| CABH\_qRT\_F | ATAAGTTCGTCGGCTTCGGG | Quantification of *cabH* expression |
| CABH\_qRT\_R | CCGCCAAAGAGGGTGTCATT |
| BRPN\_qRT\_F | GCTCAACCGCAGATTATGGA | Quantification of *brpN* expression |
| BRPN\_qRT\_R | ATGTAGTGGAGCGCAAAGGA |
| RRSH\_qRT\_F | TAACGCTCGCACCCTCCGTA | Quantification of *rrsH* expression |
| RRSH\_qRT\_R | CATGCCGCGTGTGTGAAGAA |

*a* Regions of oligonucleotides not complementary to the corresponding genes are underlined.