**Table S3. Primer set used in real time qPCR Validation**

| Primer | Sequences (5′→3′) | Fragment Size (bp) |
|---|---|---|
| *mur*F-Fw | CGTCACAGGGTCTAATGGTAAA | 224 |
| *mur*F-Rw | CAATATCTGGTTGAGCGAGGTT | |
| *hld*-Fw | CTGAGTCCAAGGAAACTAACTCTAC | 235 |
| *hld*-Rw | ATCTTGTGCCATTGAAATCA | |
| *hla*-Fw | AGGTTCCATATTGATGAATCCTG | 321 |
| *hla*-Rw | GCTACTTCATTATCAGGTAGTTGC | |
| *dltA*-Fw | ATGTTTAGCATCAGGCGGTAC | 247 |
| *dltA*-Rw | ACTTGGGAAACGGCTCACTAA | |
| *mpr*F-Fw | GAACCACCGTTTTCAACTGAA | 244 |
| *mpr*F-Rw | GTAAATCTAACTCTGGCAACCATC | |
| *spa*-Fw | AGAACAACGCAATGGTTTCA | 215 |
| *spa*-Rw | GATCGTCTTTAAGGCTTTGGAT | |
| *agrA*-Fw | TTAACAACTAGCCATAAGGATGTG | 310 |
| *agrA*-Rw | CGTTACGAGTCACAGTGAACTTAC | |
| *icaA*-Fw | CATTGAACAAGAAGCCTGACA | 301 |
| *icaA*-Rw | ATATGATTATGTAATGTGCTTGGATG | |
| *sdrD*-Fw | AGTACACAGTGGGAACAGCATCA | 225 |
| *sdrD*-Rw | CATTACCTTGAGATGATACCATTTCT | |
| gyrB-Fw | CAACTATGAAACATTACAGCAGCGT | 256 |
| gyrB-Rw | TGTGGCATATCCTGAGTTATATTGAAT | |