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| **TABLE S1** Primers and oligos used in this study. |
| **Name** | **Sequencea** | **Function** |
| **Primers** |  |  |
|  NB094 | AAACTGCAGCAATAAATCCGCCTAAATTAAAATTAAGATCTATAATTG | *xrrA* deletion construct, 1kb downstream of *xrrA* locus |
|  NB095 | TATAACTCCCCAACATAGCTAAAATATACATCAATTTATAAAGGGG  | Overlap extension of *xrrA* deletion construct |
|  NB096 | GCTATGTTGGGGAGTTATAACTAACTCCCTTATTTCT | Overlap extension of *xrrA* deletion construct |
|  NB097 | AAAGTCGACCATTTCATATATTCACATTATAATATCAACATCATGC | *xrrA* deletion construct, 1kb upstream of *xrrA* locus |
|  NB088 | AAACTGCAGGAATATATCTCTAACCTTATTGTTAATTTTTGATATTTCG  | *xrrB* deletion construct, 1kb downstream of *xrrB* locus |
|  NB089 | TGTATTATTTAGGTTAATTAAAAATACCTAAAACAAATGCTGTTTTAGG  | Overlap extension of *xrrB* deletion construct |
|  NB090 | TTTTTAATTAACCTAAATAATACAATCAATATGATGTAGAGCGG  | Overlap extension of *xrrB* deletion construct |
|  NB091 | AAAGTCGACCAATGATTTGATGGAACGTCCAGA | *xrrB* deletion construct, 1kb upstream of *xrrB* locus |
|  IC23 | AAAGTCGACGAAGTGCGGAATAAGAAACTGAC  | *hfq1* deletion construct, 1kb downstream of *hfq1* locus |
|  IC24 | CGAGTGTCTTACATCTCTCTCCTTTTTGTTTAATATC  | Overlap extension of *hfq1* deletion construct |
|  IC25 | GGAGAGAGATGTAAGACACTCGTGCTGAGTGT  | Overlap extension of *hfq1* deletion construct |
|  IC26 | AAACTGCAGCTTCTAAAGAAAGTAGGCCGTG  | *hfq1* deletion construct, 1kb upstream of *hfq1* locus |
|  IC49 | AAAGTCGACTAATTCTGCTTCATTCATTCAAG  | *hfq2* deletion construct, 1kb downstream of *hfq2* locus |
|  IC50 | GGAGGATTCGACGAGTAGTAATGAATTATTGTACATAAC | Overlap extension of *hfq2* deletion construct |
|  IC51 | TCATTACTACTCGTCGAATCCTCCTCTTTTTCT | Overlap extension of *hfq2* deletion construct |
|  IC52 | AAACTGCAGATGTACCGTAATGAAAGAACTT  | *hfq2* deletion construct, 1kb upstream of *hfq2* locus |
|  IC01 | AAAGTCGACCGATGAAATTGAACATGCTGAAT | *hfq3* deletion construct, 1kb downstream of *hfq3* locus |
|  IC02 | TTTTAGATGGTCATGAATAGTTTCTTCCTCTC | Overlap extension of *hfq3* deletion construct |
|  IC03 | AACTATTCATGACCATCTAAAAGTCTAATTTG | Overlap extension of *hfq3* deletion construct |
|  IC04 | AAACTGCAGGGATGAAGCTACCTCCATGTG | *hfq3* deletion construct, 1kb upstream of *hfq3* locus |
|  IC91 | AAAGCATGCAGAAATAAGGGAGTTAGTTATAACTCCCTTGGG | *xrrA* expression construct for ligation to pUTE657 |
|  IC92 | AAAGTCGACATGTTGTTGAATGTTATAGTTCAAGAG | *xrrA* expression construct for ligation to pUTE657 |
|  IC108 | GTTCAAGAGTTACTGAAACTAGCC | *xrrA* DNA probe template |
| **TABLE S1** (Continued) |
| **Name** | **Sequence** | **Function** |
|  IC109 | CCATACCCATACATTTAATTTTAG | *xrrA* DNA probe template |
|  IC106 | GGGGAAACGTAGGGATTTAAAC | *xrrB* DNA probe template |
|  IC107 | CAACGCTTACAAATAATGTAAGTG | *xrrB* DNA probe template |
|  IC31 | ATTTGCCTGGCAACGTCCTAC | 5S rRNA DNA probe template |
|  IC32 | GAATACAGTCTGGTAATGATGGC | 5S rRNA DNA probe template |
|  IC200 | GCTCTTATGAAGTTAGCGGCGGAC | 16S rRNA DNA probe template |
|  IC201 | CCACTTTCCTCTTCTGCACTCAAG | 16S rRNA DNA probe template |
|  IC202 | CCTGGAGGAAGAGAAAGCAAATGC | 23S rRNA DNA probe template |
|  IC203 | CCCTGGACATGGGTAGATCACC | 23S rRNA DNA probe template |
|  IC191 | GAGTACGTAAGTGCTCGTTTAG | *rpsO* DNA probe template |
|  IC192 | CTTTAACACAAGAGCGTAAAAATG | *rpsO* DNA probe template |
|  IC46 | GCGCCAGAACCAAATTGAATCACG | *xrrA*-specific 5’ RACE primer |
|  IC53 | CAAATACATATCCTAATATGGAG | *xrrA*-specific 5’ RACE primer |
|  IC47 | GTTCAAGAGTTACTGAAACTAGCCA  | *xrrA*-specific 3’ RACE primer |
|  IC48 | GAAATTAGGTGAAGAAGCGATAGCC | *xrrA*-specific 3’ RACE primer |
|  IC68 | CAAGAGTTACTGAAACTAGCC | *xrrA*-specific 3’ RACE primer |
|  IC54 | GTGATGCATGTTTAATAATTC | *xrrB*-specific 5’ RACE primer |
|  IC55 | CATTAGACTACATACAGCTATCG | *xrrB*-specific 5’ RACE primer |
|  IC69 | GTCCAGTATCTATTCTATTCAC | *xrrB*-specific 3’ RACE primer |
|  IC70 | CAATTAGGCAATAAATACCATTCCC | *xrrB*-specific 3’ RACE primer |
|  IC105 | GTGACAAGAATCGCCGGCGTACGAG | 5’ RACE Adapter-specific primer |
|  IC104 | GAGCATGCGGCCGCTAAGAAC | 3’ RACE Adapter-specific primer |
|  TH195 | GAACGTTGCTCGAGGGTAAATG | pUTE657-specific primer |
|  TH196 | GGTACGTACGATCTTTCAGCC | pUTE657-specific primer |
|  IC218 | GTGAAAGATGCGAAAGCAGATAC | *inhA1*-specific qPCR primer |
|  IC219 | CGCTGGAGATGTTGGTACTT | *inhA1*-specific qPCR primer |
|  IC214 | AAGGTGGAACACATGAAGTAGG | *gyrB*-specific qPCR primer |
|  IC215 | CGAACGTCCTCACCAGTTAAA | *gyrB*-specific qPCR primer |
| **Oligos** |  |  |
|  5’ RACE  Adapter | UUCACUGUUCUUAGCGGCCGCAUGCUC | 5’ RACE Adapter RNA oligo |
|  3’ RACE  Adapter | (p)UUCACUGUUCUUAGCGGCCGCAUGCUC(dT) | 3’ RACE Adapter RNA oligo, contains 5’ mono-phosphate and 3’ inverted dT modifications |