**S1 Table:** **Primers and probes used in this study and RACE sequences**

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| **Northern blotting probes** |
| Oligo dT | 5’-/5biosg/T20-3’ |
| 5s rRNA | 5’-/5biosg/ACACTTGTCAGCCAGTCAAC-3’ |
| sR073 | 5’-/5biosg/GAAATCATCATCTACCAGCGCAGC-3’ |
| Ysr28 | 5’-/5biosg/GGTGATTATTATCCCACCGTGCG-3’ |
| Ysr72 | 5’-/5biosg/CACGTGTTTTAGCGACAGACTCAC-3’ |
| Ysr99 | 5’-/5biosg/GCATAACGAAATAGCCCTTGCTCC-3’ |
| Ysr100 | 5’-/5biosg/GATGCATATGTAGGGGTACAGCAG-3’ |
| Ysr104 | 5’-/5biosg/GAATGAAGGGTCTACTTCCCCTAC-3’ |
| Ysr114 | 5’-/5biosg/CTGCTTTGTTGCTGGATATCCAC-3’ |
| Ysr165 | 5’-/5biosg/CTAATGCCACTCGCTTAACTGTGC-3’ |
| Ysr170 | 5’-/5biosg/AGTGATGACACCCAAACCCAAG-3’ |
| Ysr172 | 5’-/5biosg/GCTGCCAGCGTTATTCAGTATTCG-3’ |
| Ysr260 | 5’-/5biosg/CTTCTGGAGTGACAGCCCAAATAG-3’ |
| Ysr273 | 5’-/5biosg/CCTTACAAAGCATTTCACGGTGC-3’ |
| Ysr277 | 5’-/5biosg/GTTTGACTCCGTCAGGTGATACG-3’ |
| Ysr283 | 5’-/5biosg/GAAATAACAAAGACGCCGTGCGC-3’ |
| Ysr283 | 5’-/5biosg/GAAATAACAAAGACGCCGTGAGC-3’ |
|  |
| **Cloning of ysr170 and Hfq** |
| 170senF | 5’-TTTCTCACAGTATTGGAAGCATGCC-3’ |
| 170senR | 5’-ATAAAAGCCGGGTGTGAGAAGACC-3’ |
| HfqF | 5’-GATCGGATCCGAATGGCTAAGGGGCAATCTTTGCAA-3’ |
| HfqR | 5’-GATCAAGCTTTTATTCAGCGTCATCACTGTCCTG-3’ |
|  |
| **RACE primers** |
| 72 RACE-F | 5’-GTGAGTCTGTCGCTAAAACACGTG-3’ |
| 72 RACE-R | 5’-TCGCCAAACCCAGTGTGATACG-3’ |
| 114 RACE-F | 5’-GCCGCATGCCAAAATGAAAATAAC-3’ |
| 114 RACE-R | 5’-GACCATCACAAACTCAATTTTTTTCAC-3’ |
| 165 RACE-F | 5’-GCACAGTTAAGCGAGTGGCATTAG-3’ |
| 165 RACE-R | 5’-AGCCCATTTAGCTGGCTGAACG-3’ |
| 170 RACE-F | 5’-CTTGGGTTTGGGTGTCATCACTC-3’ |
| 170 RACE-R | 5’-CACACGGCCGGCTTCTTG-3’ |
| 283 RACE-F  | 5’-GCTCACGGCGTCTTTGTTATTTC-3’ |
| 283 RACE-R  | 5’-GCCTTTATGAGGGCTCGAAC-3’ |
|  |
| **Real-time PCR primers**  |
| 114PCR-F | 5’-TGAAAATAATTACGCTTGTAAAAACA-3’ |
| 114PCR-R | 5’-TTATCTGCTTTGTTGCTGGA-3’ |
| 170PCR-F | 5’-AACCTTTGGCAAAAATACGG-3’ |
| 170PCR-R | 5’-TTAAGAGTGATGACACCCAAACC-3’ |
| 172PCR-F | 5’-GCCTGAGCGATCACCTTTAAT-3’ |
| 172PCR-R | 5’-ACAAGCGGCGAATACTGAAT-3’ |
| 283PCR-F | 5’-CAAACCCCGATAATTCGATCT-3’ |
| 283PCR-R | 5’-GCAGCGGGAGAAGAGAGAAA-3’ |
|  |
| **RACE sequences** |
| **Ysr72** (2348534-2348614) 81ntAGTTAATTATTCTAACTTCATGAGGGTTTTAGGCGTATCACACTGGGTTTGGCGAGTGAGTCTGTCGCTAAAACACGTGTA***Homology:*** 100% to *Y. pseudotuberculosis* |
| **Ysr114** (1543770-1543972) 203ntGTCATGACCCGAGGTGAAAATAATTACGCTTGTAAAAACAATTGCATTTGTGAAAAAAATTGAGTTTGTGATGGTCGCCGCATGCCAAAATGAAAATAACGGCTTGATTGTTTTTTATTGGAATGTGGATATCCAGCAACAAAGCAGATAAAAGCGGGAAAACAGATAAAAAAGCGGGTGGCCCATGTAGCCACCCGTATAGA***Homology:*** 100% to *Y. pseudotuberculosis* |
| **Ysr165** (1846560-1846676) 117ntGTACCAAGAACCCGCGGAGATAGTGTTAGTTCTGATCGTTTAGCCAGAAAGAGCGTTCAGCCAGCTAAATGGGCTGCACAGTTAAGCGAGTGGCATTAGCATTAACGCGGGTTTCTC***Homology:*** ~99-100% to *Y. pseudotuberculosis* |
| **Ysr170** (2266341-2265980) 362ntTTTCTCACAGTATTGGAAGCATGCCATGCATGAGTCCAAAACCCCGCTAATATAGTTAGCGGATTACTTTATCTCAAGAAGATAGGAGTCATATTATGACTAAAAATACTGCGACTAAAGTAAAAAGCATCAAACTCGTAACCTTTGGCAAAAATACGGCTCTGGCGGGCGCAGTGCCTAGAACATTATCAGGCCAAGAAGCCGGCCGTGTGCTTGGGTTTGGGTGTCATCACTCTTAATTTAAGCATTTAATATACTGGACGTGACCTAGAAGTAGATATTGGAATAATTTGTTCCAATAATCTTCACATAACTTAGTGACTCAAGCCGGGAGTGGCGGTCTTCTCACACCCGGCTTTTAT***Homology:*** 100% to *Y. pseudotuberculosis* |
| **Ysr283** (4122174-4122369) 196ntGGTACCAGACTGCTGACAAACCCCGATAATTCGATCTTAAATGGTGAGGATAGGCAGGGAAGGCGCGGTTCGAGCCCTCATAAAGGCGCTCACGGCGTCTTTGTTATTTCTCTCTTCTCCCGCTGCAGGCACGTTGTCATCAACCTCGGTACCTTCCTTCTTCGCCTCCCTTTGATGGGTGGCATCATCTCCTCAG***Homology:*** ~99% to *Y. pseudotuberculosis* |