

**Supplementary table: 1**

Primer Sequence (5'-3')			
Gene	Forward	Reverse	Reference
16S rRNA	ACTCCTACGGGAGGCAGCAG	ATTACCGCGGCTGCTGG	This study
<i>icaA</i>	ACACTTGCTGGCGCAGTCAA	TCTGGAACCAACATCCAACA	51
<i>icaD</i>	ATGGTCAAGCCCAGACAGAG	AGTATTTCATGTTAAAGCA	51
<i>sspB</i>	CCAGCAAATTGTTGTTGCTAG	AAGCCAAAGCCGATTCACACTC	21
<i>fnbA</i>	ATCAGCAGATGTAGCGGAAG	TTTAGTACCGCTCGTTGTCC	52
<i>fnbB</i>	AAGAACGCCGAAAAGTGTG	TCTCTGCAACTGCTGTAACG	52
<i>clfA</i>	ATTGGCGTGGCTTCAGTGCT	CGTTTCTTCCGTAGTTGCATTG	53
<i>altA</i>	TGTCGAAGTATTGCCGACTTCGC	TGGAATCCTGCACATCCAGGAAC	21
<i>sarA</i>	CCCAGAAATACAATCACTGTG	AGTGCCTAGTGCAAAACC	54
<i>hla</i>	CAACTGATAAAAAAGTAGGCTGGAAAGTGAT	CTGGTAAAAACCCCTGAAGATAATAGAG	51

51: B. Kouidhi, T. Zmantar, H. Hentati and A. Bakhrouf, *Microb. Pathog.*, 2010, **49**, 14.

52: N. M. Abraham and K. K. Jefferson, *Microb. Pathog.*, 2010, **49**, 388.

53: A. Tristan, L. Ying, M. Bes, J. Etienne, F. Vandenesch and G. Lina, *J. Clin. Microbiol.*, 2003, **41**, 4465.

54: J. H. Kim, C. H. Kim, J. Hacker, W. Ziebuhr, B. K. Lee and S. H. Cho, *J. Microbiol. Biotechnol.*, 2008, **18**, 28.