

Name	Sequence (5' – 3')	Restriction site
essCRN6390fw	AAATAGATCTAGGACTGAGGCAAG	<i>Bgl</i> II
essCRN6390rev	CCTATTGAATTCATTGCTATTTAAACC	<i>Eco</i> RI
essCfwdSacl	AAAGAGCTCTAGGACTGAGGCAAAGACAATGC	<i>Sac</i> I
essCr2EMRSA15	CAAATCTCATAGAGCTCTCGTTTTATTCAAATAA	<i>Sac</i> I
essCr2ST398	CATAATTGAGCTCCCTATTGAATTAATTTATTTT	<i>Sac</i> I
essCrevMRSA252	CTTTATGAGCTCTATCCCTCCATTAG	<i>Sac</i> I

Table S1. Oligonucleotides used in this study. The *essC* gene from RN6390 was amplified using oligonucleotides *essCRN6390fw* and *essCRN6390rev* and cloned into plasmid pRAB11 as a *Bgl*II-*Eco*RI fragment. The other three *essC* genes were amplified using the same forward primer (*essCfwdSacl*) and a strain-specific reverse primer and cloned into pRAB11 as *Sac*I fragments. All inserts were confirmed for directionality and fidelity by DNA sequencing.