S1 Table : Bacterial Strains used in this study

Strain	Description	Source
Escherichia coli		
DH5a	fhuA2 lac(del)U169 phoA qlnV44 Ф80' lacZ(del)M15	BRL Life
	gyrA96 recA1 relA1 endA1 thi-1 hsdR17	Technology
Staphylococcus aureus		
RN4220	NCTC 8325-4 <i>sau1<sup>-</sup>, hsdR<sup>-</sup>,</i> laboratory strain accepting	[48]
	foreign DNA; β-toxin producer, no β-hemolysis	
RN4220 p0182	RN4220 with complementation plasmid p0182	This study
6850	methicillin-sensitive; <i>spa</i> type t185, sequence type 50	[57]
	[38], Isolated from a patient with a skin abscess,	
	progressed to bacteremia, osteomyelitis, septic	
	arthritis, and multiple systemic abscesses	
LAC	USA300 CA-MRSA, staphylococcal chromosomal	[53]
	cassette <i>mec</i> (SCC <i>mec</i> ) type IV, <i>spa</i> -type 1, Sequence	
	type 8 [ST8].	
LAC Δαβδ	LAC with deletions of the $psm\alpha$ and $psm\beta$ operons, as	[54]
	well as a point mutation in the initiation codon of $\delta$ -	
	toxin	
JE2	USA300, $rsp^+$ . Derivative of LAC, which was cured of	[18]
	three plasmids.	
JE2 NE119 (ausA)	JE2 ausA::bursa, deficient in aureusimine production	[18]
JE2 NE964 (ausB)	JE2 ausB::bursa, deficient in aureusimine production	[18]
JE2 NE1908	JE2 <i>pmt</i> C:: <i>bursa</i> , mutant in the phenol-soluble	[18]
(pmtC)	modulin transporter component, pmtC	
JE2 NE1532	JE2 agrA::bursa, non-hemolytic insertional mutant in	[18]
(agrA)	the quorum sensing system agr	
NE964 p0182	NE964 carrying plasmid p0182,	This study
	aureusimine producer	