

Supplemental Data Table S1. Publicly available genome sequences of VSSA and VISA strains

Strain	Phenotype	MLST	GenBank accession
<i>Staphylococcus aureus</i> N315	VSSA	ST5	NC_002745
<i>S. aureus</i> MW2	VSSA	ST1	NC_003923
<i>S. aureus</i> MSSA476	VSSA	ST1	NC_002953
<i>S. aureus</i> CN1	VSSA	ST72	NC_022226
<i>S. aureus</i> FPR3757	VSSA	ST8	NC_007793
<i>S. aureus</i> TW20	VSSA	ST239	NC_017331
<i>S. aureus</i> COL	VSSA	ST250	NC_002951
<i>S. aureus</i> Newman	VSSA	ST8	NC_009641
<i>S. aureus</i> NCTC 8,325	VSSA	ST8	NC_007795
<i>S. aureus</i> RF122	VSSA	ST151	NC_007622
<i>S. aureus</i> JH1	VSSA	ST105	NC_009632
<i>S. aureus</i> MRSA252	VSSA	ST36	NC_002952
<i>S. aureus</i> Mu50	VISA	ST5	NC_002758
<i>S. aureus</i> Mu3	VISA	ST5	NC_009782
<i>S. aureus</i> JH9	VISA	ST105	NC_009487
<i>S. aureus</i> JKD6008	VISA	ST239	NC_017341

Abbreviations: VSSA, vancomycin-susceptible *S. aureus*; VISA, vancomycin-intermediate resistant *S. aureus*; MLST, multilocus sequence type.