

Supplemental Table S1. Novel PCR primers used in the study to characterize novel SCC*mec*-SCC elements.

Gene or region amplified	Primer pair no.	Primer name	Nucleotide sequence (5'-3')	Nucleotide coordinates
J1 SCC <i>mec</i> IIE to the end of the <i>ccrAB4</i> -carrying element	1	J1_IIE_F3 Contig 101 R2	CAAGTTATCGGATAATGAA GCCAAGCAATTATCATTATG	184-166 ^a 4636-4617 ^b
	2	Contig 101 F2 ccrA4R5	CCTCGTCTTGAATTTCTTTC CAATATCGTTGAACAATCAC	4598-4617 ^b 7205-7186 ^b
	3	ccrA4F5 M1_R2	CTTCATTCCTTTTCAGGAGTT AACATGCTTAACGTGAAGTG	7159-7178 ^b 9560-9541 ^b
	4	M1_F2 M1_R4	TAGGAGCACACTTACACAAG CAGCGTATCGTTTATACTG	9498-9517 ^b 13933-13914 ^b
J1 SCC <i>mec</i> IVc/IVE to within adjacent <i>ccrAB4</i> -carrying element	5	J1_IVc_F2 Contig 101 R2	GACAGTATTCCAACAGTCAA GCCAAGCAATTATCATTATG	84-65 ^c 4636-4617 ^b
J1 SCC <i>mec</i> IVb/IVF to within adjacent <i>ccrAB4</i> -carrying element	6	J1_IVb_F2 Contig 101 R2	ACTTAAGTAGTAGCTAACAG GCCAAGCAATTATCATTATG	172-153 ^d 4636-4617 ^b
J1 SCC <i>mec</i> IVg to within adjacent <i>ccrAB4</i> -carrying element	7	J1_IVg_F1 Contig 101 R2	CGTGATAGCGACACAATAC GCCAAGCAATTATCATTATG	133-115 ^e 4636-4617 ^b

^aNucleotide coordinates based on the nucleotide sequence of SCC*mec* IIE (GenBank accession number AJ810120.1).

^bNucleotide coordinates based on the nucleotide sequence of SCC_{MI} of AR13.1/3330.2 (HE858191).

^cNucleotide coordinates based on the nucleotide sequence of SCC*mec* IVc (AB096217.1).

^dNucleotide coordinates based on the nucleotide sequence of from SCC*mec* IVb (AB063173.1).

^eNucleotide coordinates based on the nucleotide sequence of SCC*mec* IVg (DQ106887.1).

Supplemental Table S2. The main characteristics of the SCC*mec* types and subtypes investigated during the present study^a

SCC <i>mec</i> type	<i>ccr</i> type	<i>mec</i> class	Major characteristics of J regions ^b			Reference
			J1	J2	J3	
I	1	B	<i>pls</i>	SCC <i>mec</i> I specific J2 region	<i>dcs</i>	(1, 4, 6)
I – <i>pls</i>	1	B	Lacks <i>pls</i>	ND	<i>dcs</i>	(6)
Ia			<i>pls</i>	SCC <i>mec</i> I specific J2 region	pUB110 (<i>aadD</i>) & <i>dcs</i>	(1, 4)
II			<i>kdp</i>	Tn554 (<i>erm(A)</i> & <i>spc</i>)	pUB110 & <i>dcs</i>	(6)
IIA	2	A.4 ^c	SCC <i>mec</i> IVb specific J1 region	Tn554	pUB110 & <i>dcs</i>	(6)
IIB	2	A	SCC <i>mec</i> IVb specific J1 region	Lacks Tn554	pUB110 & <i>dcs</i>	(6)
IIC	2	A.3 ^d	SCC <i>mec</i> IVb specific J1 region	Tn554, lacks ORFs between Tn554 & <i>mec</i> complex in SCC <i>mec</i> II	pUB110 & <i>dcs</i>	(6)
IID	2	A.4 ^c	SCC <i>mec</i> IVb specific J1 region	Tn554	<i>dcs</i>	(6)
IIE ^e	2	A.3 ^d	SCC <i>mec</i> IVb specific J1 region	Tn554; lacks ORFs between Tn554 & <i>mec</i> complex	<i>dcs</i>	(6)
III & SCCHg	3	A	SCC <i>mec</i> IIIHg-specific J1 region	ψTn554	pT181 (<i>tet(K)</i>) & SCCHg with	(1, 5)

SCC <i>mec</i> type	<i>ccr</i> type	<i>mec</i> class	Major characteristics of J regions ^b			Reference
			J1	J2	J3	
III & SCC <i>Hg</i> – p1258/Tn554	3	A	ND	ND	p1258 (<i>mer</i>), Tn554 & <i>ccrC</i> pT181 & SCC <i>Hg</i> with p1258, Tn554 & <i>ccrC</i> but lacking ORFs between p1258 & Tn554	(6)
IVA	2	B	SCC <i>mec</i> IVc-specific J1 region	No J2 region	pUB110 & <i>dcs</i>	(1)
IVa	2	B	SCC <i>mec</i> IVa-specific J1 region	No J2 region	<i>dcs</i>	(1, 3)
IVc	2	B	SCC <i>mec</i> IVc-specific J1 region	No J2 region	<i>dcs</i>	(1)
IVE	2	B	SCC <i>mec</i> IVc-specific J1 region	ND	Lacks <i>dcs</i>	(6)
IVF	2	B	SCC <i>mec</i> IVb-specific J1 region	ND	Lacks <i>dcs</i>	(6)
IVg	2	B	SCC <i>mec</i> IVg-specific J1 region	No J2 region	<i>dcs</i>	(1, 2)

^a This table was adapted from the Guidelines on the Classification of SCC*mec* elements (1). For the main characteristics of additional SCC*mec* types and subtypes not investigated in the present study see reference (1) or http://www.sccmec.org/Pages/SCC_HomeEN.html.

^b The antimicrobial resistance genes carried on plasmids and transposons integrated within the SCC*mec* elements are written in parenthesis after the first time the relevant plasmid or transposon name is listed.

^c class A.4 *mec* complex differs from class A *mec* due to the presence of IS1182 within *mecI* (6).

^d class A.3 *mec* complex differs from class A *mec* due to the presence of IS1182 within *mecI* and deletion of the 3' end of *mec* regulatory gene *mecI* (6).

ND, not determined.

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