

Supplemental Data Table S1. Antimicrobial resistance patterns of MLST ST72-methicillin-resistant *Staphylococcus aureus* clinical isolates in Korea

Reference	Total (N)	Antimicrobial resistance (N)							
		ERY	CLI	CIP	GEN	Fusidic acid	TET	RIF	TMP_SMX
Kim, <i>et al.</i> 2008 [14]	25	14	14	0	2		1	0	0
Moon, <i>et al.</i> 2010 [17]	2	1	1	0	0		0	0	0
Lee, <i>et al.</i> 2011 [11]	29	18	18				0		
Kwon, <i>et al.</i> 2011 [18]	58	45	44			39			
Joo, <i>et al.</i> 2012 [9]	84	38	21			2			
Park, <i>et al.</i> 2015 [19]	79	28	18	5	12	0		0	0
Shin, <i>et al.</i> 2016 [10]	36	16	0	1	11		5		
Bae, <i>et al.</i> 2019 [20]	16	4	4	4	4	2	1		
Choi, <i>et al.</i> (Unpublished data) [22]	299	83	61	34	47	8	9	6	2
Overall resistance, N (%)	628	200/559 (35.78)	182/628 (28.98)	88/515 (17.09)	76/457 (16.63)	12/478 (2.51)	55/459 (11.98)	6/405 (1.48)	2/405 (0.49)

Abbreviations: MLST, multilocus sequence typing; ST, sequence type; CLI, clindamycin; ERY, erythromycin; CIP, ciprofloxacin; GEN, Gentamycin; TET, Tetracycline; RIF, Rifampicin; TMP_SMX, trimethoprim/sulfamethoxazole.