

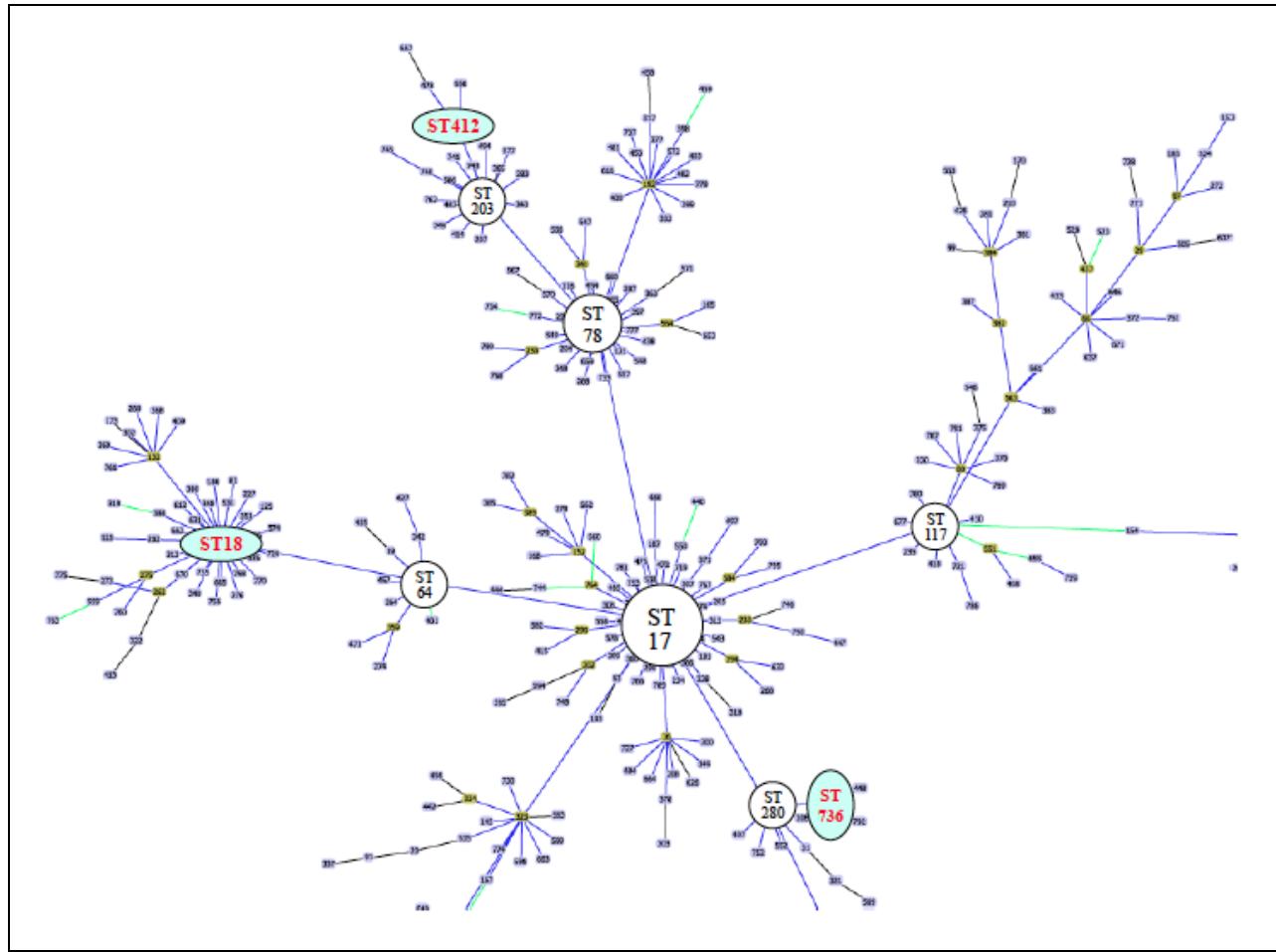
AAC02683-14**Supplemental Table S1.** Antimicrobial susceptibility profiles of DNSE and daptomycin-susceptible *E. faecium* (DSE) clinical isolates

Antimicrobial agent	DNSE isolates (n = 42, % ^a)	DSE isolates (n = 43, % ^a)	p value
Daptomycin ^b	0	43 (100)	<0.0001
Vancomycin	5 (11.9)	1 (2.3)	0.1096
Ampicillin	0	0	-
Linezolid	40 (95.2)	41 (95.3)	1.0000
Nitrofurantoin	23 (54.7)	24 (55.8)	1.0000
Rifampin	3 (7.1)	4 (9.3)	1.0000
Quinupristin/dalfopristin	40 (95.2)	42 (97.7)	0.6160
Tetracycline	1 (2.4)	6 (13.9)	0.1096
Levofloxacin	1 (2.4)	0	-
Erythromycin	0	0	-
Gentamicin synergy screen ^c	37 (88.1)	37 (86.1)	1.0000

^a The data shown in the parenthesis are % susceptible.

^b Refer to Table 1 for the MICs of DNSE isolates. The MIC distributions for the 43 DSE isolates are: 1 µg/mL (n = 1), 2 µg/mL (n = 8), 3 µg/mL (n = 6), 4 µg/mL (n = 28).

^c High level of aminoglycoside resistance if gentamicin MIC > 500 µg/mL).



Supplemental Figure 1. Genetic relatedness of daptomycin-nonsusceptible *E. faecium* clinical isolates with vancomycin-resistant enterococci of the cluster complex 17 (CC17). The three major clones (ST18, ST412 and ST736) detected in this study are shown in red.