

In Vitro Activity of Daptomycin in Combination with β -Lactams, Gentamicin, Rifampin, and Tigecycline against Daptomycin-Nonsusceptible Enterococci

Janet A. Hindler,^a Annie Wong-Beringer,^b Carmen L. Charlton,^{a,c} Shelley A. Miller,^a Theodoros Kelesidis,^d Marissa Carvalho,^a George Sakoulas,^e Poochit Nonejuie,^f Joseph Pogliano,^f Victor Nizet,^e Romney Humphries^a

Pathology and Laboratory Medicine, University of California, Los Angeles, Los Angeles, California, USA^a; School of Pharmacy, University of Southern California, Los Angeles, Los Angeles, California, USA^b; Laboratory Medicine and Pathology, University of Alberta, and the Provincial Laboratory for Public Health (ProvLab), Edmonton, AB, Canada^c; Division of Infectious Diseases, University of California, Los Angeles, Los Angeles, California, USA^d; Department of Pediatrics, University of California San Diego School of Medicine, La Jolla, California, USA^e; Department of Biological Sciences, University of California, San Diego, La Jolla, California, USA^f

Enterococci that are nonsusceptible (NS; MIC > 4 μ g/ml) to daptomycin are an emerging clinical concern. The synergistic combination of daptomycin plus beta-lactams has been shown to be effective against vancomycin-resistant *Enterococcus* (VRE) species *in vitro*. This study systematically evaluated by *in vitro* time-kill studies the effect of daptomycin in combination with ampicillin, cefazolin, ceftriaxone, ceftaroline, ertapenem, gentamicin, tigecycline, and rifampin, for a collection of 9 daptomycin-NS enterococci that exhibited a broad range of MICs and different resistance-conferring mutations. We found that ampicillin plus daptomycin yielded the most consistent synergy but did so only for isolates with mutations to the *liaFSR* system. Daptomycin binding was found to be enhanced by ampicillin in a representative isolate with such mutations but not for an isolate with mutation to the *yycFGHIJ* system. In contrast, ampicillin enhanced the killing of the LL-37 human antimicrobial peptide against daptomycin-NS *E. faecium* with either the *liaFSR* or *yycFGHIJ* mutation. Antagonism was noted only for rifampin and tigecycline and only for 2 or 3 isolates. These data add support to the growing body of evidence indicating that therapy combining daptomycin and ampicillin may be helpful in eradicating refractory VRE infections.

Daptomycin is a cyclic lipopeptide antimicrobial agent with bactericidal activity against Gram-positive bacteria, including *Enterococcus* spp. Daptomycin is not approved by the U.S. Food and Drug Administration (FDA) for the treatment of infections caused by vancomycin-resistant *Enterococcus* (VRE) strains (e.g., vancomycin-resistant *Enterococcus faecalis*) or by vancomycin-susceptible or -resistant *E. faecium* strains. However, due to a limited number of available therapeutic options, daptomycin is frequently used in clinical practice for treatment of serious infections caused by these bacteria. No daptomycin resistance breakpoint has been established for the enterococci by the Clinical and Laboratory Standards Institute or the U.S. FDA. Isolates with MICs above the susceptible breakpoint of 4 μ g/ml are therefore referred to as daptomycin-nonsusceptible (DNS) isolates (1). The prevalence of DNS *Enterococcus* strains in the United States remains low, ranging from 0.02% for *E. faecalis* to 0.18% for *E. faecium* (2). Nonetheless, we and others have isolated DNS *Enterococcus* strains from both patients treated with and patients naive to daptomycin therapy (3–8) and much higher rates of DNS *E. faecium* have been reported in Europe (9).

The recent description of daptomycin-susceptible *Enterococcus* strains for which daptomycin exhibits only bacteriostatic activity (10, 11) brings further into question the role of daptomycin for the treatment of enterococcal infections. These isolates have decreased susceptibility to daptomycin (DSD), with daptomycin MICs ranging from 3 to 4 μ g/ml, which is higher than the wild-type modal daptomycin MICs of 0.5 μ g/ml for *E. faecalis* and 2.0 μ g/ml for *E. faecium* (12). These isolates harbor point mutations in the *liaFSR* genes, which encode a three-component regulatory system involved in cell membrane stress response (10, 11). Because *liaFSR* mutation is thought to be one of the first events in the

stepwise accumulation of genomic mutations that lead to the DNS phenotype (11, 13, 14), it is possible that the use of daptomycin for the treatment of infections caused by DSD isolates increases the risk of the organism acquiring further chromosomal mutations and DNS MICs. This concern is more than theoretical, as a recent clinical treatment failure was documented for a patient with a bloodstream infection caused by a DSD *E. faecium* isolate (daptomycin MIC of 3 μ g/ml) that harbored T120A and W73C substitutions in LiaS and LiaR, respectively. The patient was treated with high-dose daptomycin (8 mg/kg of body weight/day) plus gentamicin (3 mg/kg/day) but had recurrent bacteremia with an *E. faecium* strain that eventually became DNS. Daptomycin MICs for this isolate progressed from 16 μ g/ml to 256 μ g/ml over the course of several months, while the patient was on daptomycin therapy (15). Supporting *in vitro* data from a simulated endocardial vegetation pharmacokinetic/pharmacodynamic model dem-

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Address correspondence to Romney Humphries, rhumphries@mednet.ucla.edu.

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TABLE 1 DNS *Enterococcus* isolates included in this study^a

Strain	DAP Etest MIC (μg/ml)	DAP MIC (μg/ml)	DAP MBC (μg/ml)	MIC (μg/ml)						Susceptibility to Gent (500 μg/ml)	Predicted amino acid change(s) associated with daptomycin resistance in indicated proteins						
				VAN	AMP	CPT	TIG	RIF	LiaF		LiaS	LiaR	ClsA	ClsB			
<i>E. faecalis</i>																	
Efc01	>256	64	>180	1	≤2	4	≤0.25	4	S	171Idel							
Efc04	24	32	128	>32	>64	1	≤0.25	>4	S	171Idel							
<i>E. faecium</i>																	
Efm12	256	64	128	>32	>64	>8	≤0.25	4	R	N251E		G53R, R215K		G174V			
Efm13	>256	64	180	>32	>64	>8	≤0.25	>4	S	T120A		W73C N23T					
Efm15	>256	16	180	>32	>64	>8	≤0.25	>4	R						V38L G174V S298T		
Efm16	>256	64	128	>32	>64	>8	≤0.25	>4	R	N251E					G174V		
Efm19	48	16	128	>32	>64	>8	≤0.25	>4	R						V38L G174V		
Efm23	>256	64	128	>32	>64	>8	≤0.25	>4	R	N251E					V38L G174V S298T		
Efm25	>256	64	>180	>32	>64	>8	≤0.25	>4	S	T120A		W73C N23T					

^a MICs and MBCs were determined by BMD unless otherwise indicated. VAN, vancomycin; AMP, ampicillin; CPT, ceftaroline; TIG, tigecycline; RIF, rifampin; Gent, gentamicin; S, susceptible; R, resistant.

onstrated that exposure of enterococci to daptomycin concentrations equivalent to FDA-cleared doses (i.e., 4 to 6 mg/kg/day) results in the development of the DNS phenotype (16).

In order to mitigate the development of DNS, a minimum dose of 10 mg/kg/day, particularly in cases with high bacterial burden, such as cases of endocarditis, has been suggested for the enterococci (16). Alternatively, combination therapy, such as with a β-lactam plus daptomycin, has been suggested for successful treatment of DSD *Enterococcus* infections (13, 16). The combinations of daptomycin plus ampicillin and daptomycin plus ceftaroline have been shown to enhance the activity of daptomycin against the enterococci, by improving binding to the target cytoplasmic membrane, even in ampicillin-resistant isolates (17–19). However, such synergy is not observed for all DNS *Enterococcus* isolates and may occur only for isolates of *E. faecium* with a DSD phenotype associated with mutation to the LiaFSR pathway. A second pathway to the DSD phenotype has also been previously described and is associated with mutations to the YycFGHIJ system, a second regulatory system involved in cell wall homeostasis in Gram-positive bacteria (13). Two DSD *E. faecium* isolates associated with mutations to *yycFGHIJ* did not display *in vitro* synergy between daptomycin and ampicillin in a recent study (13).

The intent of the present study was to further explore the *in vitro* killing kinetics of daptomycin against a collection of 9 DNS *Enterococcus* isolates with a variety of daptomycin MICs and previously identified genetic mutations conferring DSD. The daptomycin concentrations evaluated ranged from 0.5× the daptomycin MIC to 180 μg/ml, the mean total serum concentration achievable with maximal daptomycin dosing (12 mg/kg/day). In addition, as daptomycin may be combined with broad-spectrum β-lactams in critically ill hospitalized patients with DNS enterococcal infections, the effect of the combination of daptomycin with five β-lactams and three other antimicrobial agents plus a host defense peptide, LL-37, was evaluated for these isolates and was correlated with resistance mechanisms.

MATERIALS AND METHODS

Bacterial isolates. Seven clinical isolates of *E. faecium* and two of *E. faecalis* were included in this study, all with daptomycin MICs of >4 μg/ml

(Table 1). Typing of the isolates was performed as described previously, by repetitive sequence-based PCR (rep-PCR) analysis (bioMérieux, Durham, NJ), to confirm that the isolates were not clonal (3–5). Daptomycin MICs were determined by Etest (bioMérieux, Durham, NJ) on Mueller-Hinton agar according to the manufacturer's instructions, and by broth microdilution (BMD), in cation-adjusted Mueller-Hinton broth (CA-MHB; BBL, Sparks, MD) supplemented with 50 mg/liter CaCl₂, on panels prepared in-house (1). Ampicillin, cefazolin, ceftriaxone, ceftaroline, eraptenem, rifampin, and tigecycline MICs were also determined by BMD, following Clinical and Laboratory Standards Institute standards (1). As expected, all isolates had high ceftriaxone, cefazolin, and eraptenem MICs (≥32 μg/ml; data not shown). High-level gentamicin resistance was determined by BMD, by the ability to grow in 500 μg/ml gentamicin in brain heart infusion (BHI) medium (BBL) (1). The minimum bactericidal concentration (MBC) of daptomycin was determined for each isolate, as described elsewhere (20), in CA-MHB supplemented with 50 mg/liter CaCl₂, as the concentration of daptomycin that resulted in a ≥3 log₁₀ reduction in CFU compared to the inoculum, after 24 h of incubation. MIC and MBC testing was performed in triplicate for each isolate, on separate testing days, and modal MIC and MBC were reported. Use of clinical isolates for this study was approved by the institutional review board of the University of California, Los Angeles (UCLA).

Time-kill assays and synergy testing. The bactericidal activity of daptomycin was performed by time-kill assays with an initial inoculum of 6 × 10⁶ CFU/ml in 10 ml of CA-MHB supplemented with 50 mg/liter CaCl₂. Daptomycin concentrations tested against each isolate were 0.5×, 1×, and 2× the daptomycin MIC (Table 1). In addition, each isolate was tested in the presence of 180 μg/ml daptomycin (DAP₁₈₀), as 183.7 μg/ml is the mean maximum concentration of drug in serum (C_{max}) reported in the CUBICIN package insert for a cohort of subjects administered 12 mg/kg/day daptomycin. Bacterial colony counts were performed at 0, 6, and 24 h, in duplicate, by removal of two 100-μl aliquots of the culture, serial dilution in sterile saline solution, and plating of 25 μl on sheep blood agar plates (BD, Sparks, MD). Preliminary experiments were performed to ensure that this method did not result in antimicrobial carry-over (not shown [21]). The limit of detection for the time-kill experiments was 100 CFU/ml, assuming maximal plating efficiency. Bactericidal activity was defined as a ≥3 log₁₀ reduction in CFU/ml at 24 h in comparison to the CFU/ml at 0 h.

Synergy testing was performed for daptomycin, at the concentrations listed above, and for eight other antimicrobials at C_{max} concentrations based on pharmacokinetic studies in adults. The following concentrations

were tested for the respective drugs based on routine doses given intravenously: ampicillin at 90 µg/ml to 1 to 2 g every 6 h (22); cefazolin at 185 µg/ml to 1 g every 8 h (23); ceftriaxone at 200 µg/ml to 1 g every 24 h (24), ceftaroline at 21 µg/ml to 600 mg every 12 h (25); ertapenem at 115 µg/ml to 1 g every 24 h (26); gentamicin at 25 µg/ml to 6 mg/kg daily (27); rifampin at 10 µg/ml to 600 mg every 24 h (28); and tigecycline at a loading dose of 0.8 µg/ml to 100 mg followed by 50 mg every 12 h (29). Synergy was defined as a decrease of $\geq 2 \log_{10}$ CFU/ml in bacterial counts at 6 or 24 h for the combination, compared to the counts for the most active agent alone at the respective time point, provided that the counts for the combination were $\geq 2 \log_{10}$ CFU/ml below the starting inoculum. Bactericidal activity of the combination was defined as a $\geq 3 \log_{10}$ CFU/ml reduction in bacterial counts at 24 h compared to the starting inoculum (30). Antagonism was defined as an increase of $\geq 2 \log_{10}$ CFU/ml in bacterial counts at 6 or 24 h for the combination, compared to the counts for the most active agent alone.

Mutational analysis. Mutations in genes previously associated with DNS were evaluated by Sanger sequencing of PCR products, as described by Werth and colleagues (16). The following genes were evaluated: *liaFSR*, encoding a three-component regulatory system that is part of the cell envelope response to stress, and *cls*, which encodes cardiolipin synthetase. Sequences were compared against the genomes of *E. faecalis* V583 and *E. faecium* DO, two daptomycin-susceptible enterococcal isolates whose genomes are sequenced and publicly available.

BDP-daptomycin assays. Tested strains were grown overnight (14 to 16 h) to stationary phase in Luria broth (LB), diluted 1:100 in fresh antibiotic-free LB or LB containing ampicillin at 50 mg/liter, grown at 37°C with shaking at 200 rpm to an optical density at 600 nm (OD_{600}) of 0.6 (approximately 5 to 6 h), and stained for 20 min with boron-dipyrromethene (BODIPY)-labeled daptomycin (BDP-daptomycin) at 32 mg/liter (supplied courtesy of Cubist Pharmaceuticals, Lexington, MA) and with 50 mg/liter $CaCl_2$ as previously described (18). The concentration of labeled daptomycin was established by pilot studies as optimal for fluorescence microscopy (data not shown). Excess unincorporated label was removed by washing the cells three times in antimicrobial-free LB. The cells were counterstained with 2 mg/liter DAPI (4',6-diamidino-2-phenylindole) in the final LB wash to visualize the nucleoid and then imaged using a Delta Vision Deconvolution microscope (Applied Precision, Inc., Issaquah, WA) as previously described (18).

Human cathelicidin LL-37 killing assays. Human cathelicidin LL-37 (net charge, +6 at pH 7.5) was purchased from AnaSpec, Inc. (Fremont, CA), and killing assays were performed at $1 \times$ MIC (2 µM) as previously described (18). Bacteria were grown overnight (14 to 16 h) in LB in the absence or presence of ampicillin at 50 µg/ml, pelleted, washed in phosphate-buffered saline (PBS), and resuspended to an OD_{600} of 0.5 in PBS (approximately 10^8 CFU/ml). Bacteria were diluted to 10^3 CFU/ml in RPMI medium–5% LB containing $1 \times$ MIC of LL-37 and incubated at 37°C. Aliquots (10 µl) were plated on sheep blood agar after 2 h of incubation, and colonies were enumerated after 24 h to determine the percentages of surviving bacteria (\pm standard deviations [SD]). Results represent experiments performed in quadruplicate.

RESULTS

Effect of daptomycin and of other antimicrobials alone against 9 DNS *Enterococcus* isolates. The accompanying mutations and daptomycin MICs and MBCs for the 9 isolates are shown in Table 1. Modal MICs ranged from 16 to 64 µg/ml by BMD and 24 to >256 µg/ml by Etest (Table 1), and MICs for each isolate were reproducible within a \log_2 dilution range on different testing days. Modal daptomycin MBCs ranged from 128 to >180 µg/ml (Table 1 and Fig. 1). Growth curves for the 9 isolates, in the absence of antimicrobials, were not appreciably different, with an average increase in CFU/ml of $2.45 \log_{10}$ (± 0.11) after 24 h of incubation (not shown). Daptomycin was bactericidal (i.e., ≥ 3 log reduction

in CFU/ml versus the starting inoculum) at $2 \times$ the daptomycin MIC ($DAP_{2 \times MIC}$) for 5 of 9 (55.6%) isolates (Fig. 1A) and was bactericidal at 180 µg/ml daptomycin (DAP_{180}) for 7 of 9 (77.8%) isolates (Fig. 1A). Time-kill curves for the two isolates in this study for which DAP_{180} was not bactericidal are presented in Fig. 1B. Isolate *E. faecalis* c01 (Efc01) had a $2.51 (\pm 1.01) \log_{10}$ decrease in CFU/ml at 24 h, compared to the inoculum, in the presence of DAP_{180} . In contrast, isolate Efm25, an *E. faecium* isolate, demonstrated $2.50 (\pm 1.72) \log_{10}$ killing at 6 h in the presence of DAP_{180} , but by 24 h, the CFU/ml had increased $1.38 \log_{10}$ (± 0.09) from the 6-h reading, yielding an overall 1.2 log decrease in CFU/ml at 24 h compared to 0 h (Fig. 1B). Both of these isolates had daptomycin MICs of 64 µg/ml by BMD and >256 µg/ml by Etest. For the other 7 isolates, there was no growth from the 6-h or 24-h subcultures following incubation in the presence of DAP_{180} (i.e., <100 CFU/ml was present). No clear correlation between *liaFSR* or *cls* mutation and response to daptomycin in the time-kill studies was identified, and these differences may be attributed to other mutations in the genomes of these isolates.

Ampicillin, cefazolin, ceftriaxone, ceftaroline, ertapenem, gentamicin, tigecycline, and rifampin did not display bactericidal activity at the concentrations used in this study (see Fig. S1 in the supplemental material), with one exception. Ertapenem was bactericidal against isolate Efc01 alone (see Fig. S1). Efc01 was susceptible to ampicillin ($MIC \leq 2$ µg/ml), which predicts susceptibility to imipenem but not necessarily susceptibility to ertapenem (1).

***liaFSR* mutations associated with synergy between daptomycin and ampicillin.** The changes in \log_{10} CFU/ml after 24-h antimicrobial exposures, for experiments with ampicillin (90 µg/ml) in combination with 4 concentrations of daptomycin, are shown in Table 2. Ampicillin alone was not bactericidal for any of the DNS enterococci tested. At $0.5 \times$ the daptomycin MIC ($DAP_{0.5 \times MIC}$), the addition of ampicillin yielded synergy in 7/9 isolates (77.8%), including both *E. faecalis* isolates, despite the fact that only one isolate, Efc01, was susceptible to ampicillin (Table 1). These 7 isolates all harbored mutation to the *liaFSR* system, whereas the 2 isolates for which the ampicillin-daptomycin combination was not synergistic with $DAP_{0.5 \times MIC}$ were the only isolates included in this study without *liaFSR* mutation. The combination of ampicillin and $DAP_{0.5 \times MIC}$ yielded bactericidal activity against only 3 of the isolates: Efc01, Efc04, (both *E. faecalis*), and Efm23 (an *E. faecium* isolate; Table 2). Ampicillin plus $DAP_{1 \times MIC}$ was both synergistic and bactericidal for 6/9 isolates (66.6%), all of which harbored mutations in *liaFSR*. Ampicillin, which was synergistic but not bactericidal, in combination with $DAP_{0.5 \times MIC}$ for isolate Efm13, no longer demonstrated synergy for this isolate when tested at this isolate's daptomycin MIC ($DAP_{1 \times MIC}$). Daptomycin at twice the MIC ($DAP_{2 \times MIC}$) was bactericidal for isolates Efm4, Efm12, Efm16, Efm19, and Efm23 (Fig. 1). For the 4 remaining isolates for which an effect of adding a second antimicrobial could be discerned, ampicillin yielded synergy for 3 (75.0%; Table 2). Daptomycin at 180 µg/ml (DAP_{180}) was bactericidal against all but two isolates: Efc01 and Efm25 (Fig. 1A). Ampicillin acted synergistically with DAP_{180} for both isolates (Fig. 2A and Table 2).

***liaFSR* mutations associated with synergy between daptomycin and other β -lactams.** The changes in \log_{10} CFU/ml after 24-h antimicrobial exposures, for experiments with cefazolin, ceftriaxone, ceftaroline, and ertapenem in combination with 4 concentra-

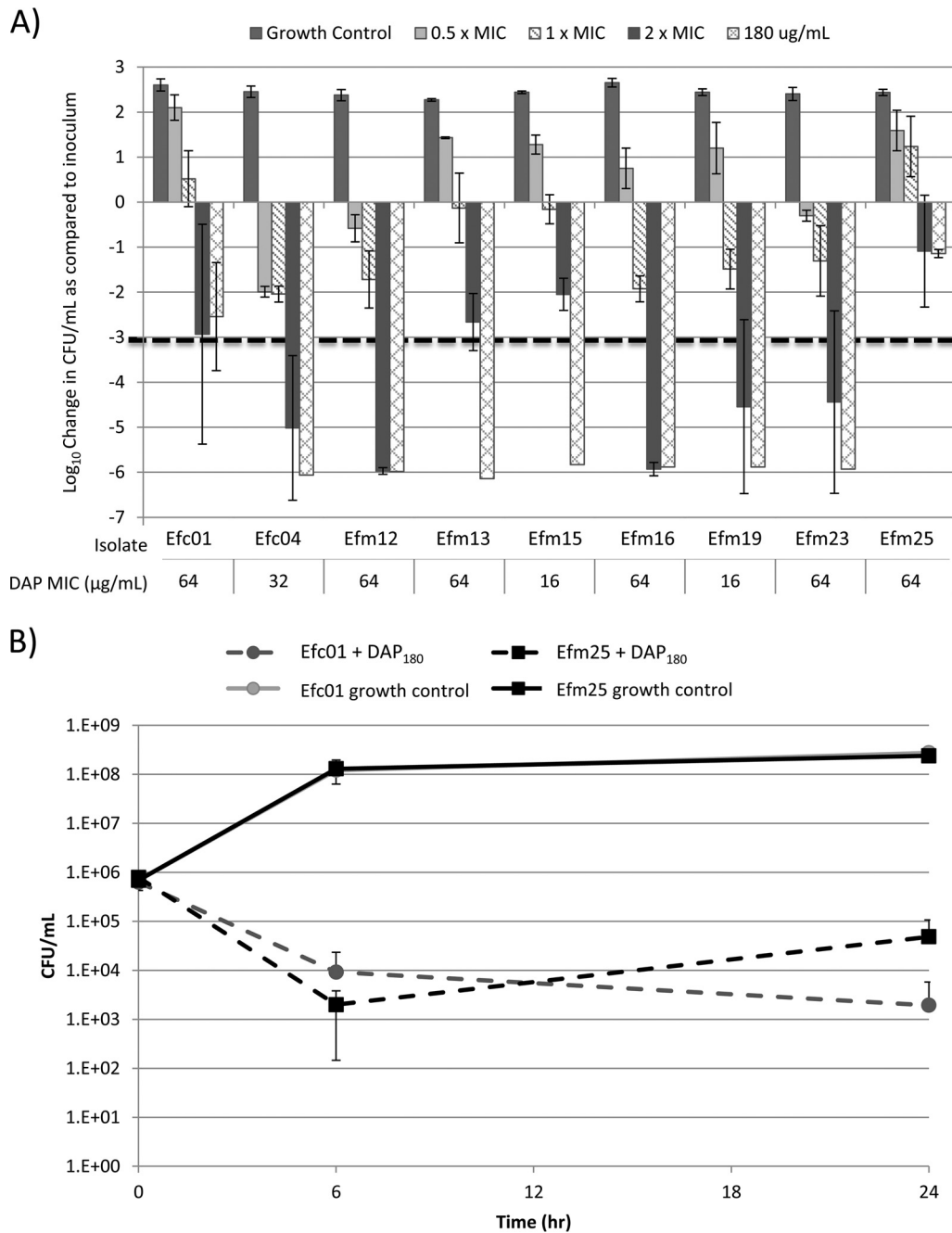


FIG 1 (A) Change in \log_{10} CFU/ml after 24 h of antibiotic exposure to various concentrations of daptomycin in the kill curve. (B) Kill curves for isolates Efc01 and Efm25, the only two isolates that displayed growth after 24 h of incubation in 180 $\mu\text{g/ml}$ daptomycin, in combination with test antimicrobials.

tions of daptomycin, are shown in [Table 3](#). When combined with $\text{DAP}_{0.5\text{xMIC}}$, ceftazidime demonstrated synergy for only 2 isolates (22.2%; [Table 3](#)) and was bactericidal for 1 (11.1%). At $\text{DAP}_{1\text{xMIC}}$, the addition of ceftazidime yielded synergy for 5 isolates (55.5%), and for all 5, the combination was bactericidal ([Table 3](#)). At twice the daptomycin MIC, ceftazidime yielded synergy for 2 of the 4 isolates (50.0%) for which $\text{DAP}_{2\text{xMIC}}$ alone was not bactericidal ([Table 3](#)). Ceftazidime in combination with DAP_{180} was synergistic for both isolates that were not killed by 180 $\mu\text{g/ml}$ daptomycin (Efc01 and Efm25).

Ceftriaxone (200 $\mu\text{g/ml}$) demonstrated synergy for 4/9 (44.4%) isolates when combined with $\text{DAP}_{0.5\text{xMIC}}$ and for 7/9 (77.7%) when combined with $\text{DAP}_{1\text{xMIC}}$ ([Table 3](#)). The latter combination was bactericidal for 6 (66.6%) of the isolates, the same 6 isolates for which synergy was found for ampicillin with $\text{DAP}_{1\text{xMIC}}$ ([Tables 2](#) and [3](#)). In addition to these 6 isolates, isolate Efm13 yielded synergy with the ceftriaxone-plus- $\text{DAP}_{1\text{xMIC}}$ combination, but this combination was not bactericidal ([Table 3](#)). The combination of ceftriaxone plus $\text{DAP}_{2\text{xMIC}}$ was synergistic and bactericidal for 2/4 (50.0%) isolates not killed by $\text{DAP}_{2\text{xMIC}}$, and

TABLE 2 Summary of 24-h time-kill testing for daptomycin at 4 concentrations in combination with 90 $\mu\text{g/ml}$ ampicillin

Isolate (daptomycin MIC)	Result for daptomycin at indicated tested concn or synergy result ^a			
	0.5 \times MIC	1 \times MIC	2 \times MIC	180 $\mu\text{g/ml}$
Efc01 (64 $\mu\text{g/ml}$)	BC	BC	BC	—
Efc04 (32 $\mu\text{g/ml}$)	BC	BC	NA	NA
Efm12 (64 $\mu\text{g/ml}$)	—	BC	NA	NA
Efm13 (64 $\mu\text{g/ml}$)	—	—	BC	NA
Efm15 (16 $\mu\text{g/ml}$)	—	—	—	NA
Efm16 (64 $\mu\text{g/ml}$)	—	BC	NA	NA
Efm19 (16 $\mu\text{g/ml}$)	—	—	NA	NA
Efm23 (64 $\mu\text{g/ml}$)	BC	BC	NA	NA
Efm25 (64 $\mu\text{g/ml}$)	—	BC	BC	—
Total no. (%) of isolates with synergy ^b	7 (78)	6 (67)	3 (75)	2 (100)

^a Boldface data indicate synergy. —, daptomycin plus ampicillin yielded a colony count at 24 h that was $\geq 2 \log_{10}$ CFU/ml lower than that seen with the most active antimicrobial alone. BC, bactericidal combination, where daptomycin plus ampicillin yielded a $\geq 3 \log_{10}$ CFU/ml reduction in colony counts compared to the starting inoculum. NA, daptomycin at this concentration was bactericidal and synergy could not be assessed.

^b Data indicate the total numbers and percentages of isolates for which neither antimicrobial was bactericidal alone.

the combination of ceftriaxone plus DAP₁₈₀ was synergistic for the two isolates not killed by 180 $\mu\text{g/ml}$ daptomycin alone (Table 3 and Fig. 2).

The addition of ceftaroline (21 $\mu\text{g/ml}$) to DAP_{0.5 \times MIC} yielded synergy for 5/9 (55.5%) of the isolates. Synergy was observed for isolate Efc04 only with this combination at 6 h (not shown). Similarly, ceftaroline plus DAP_{0.5 \times MIC} was bactericidal only against isolate Efc04. The ceftaroline MIC for isolate Efc04 was 1 $\mu\text{g/ml}$ by BMD, but at the 21 $\mu\text{g/ml}$ concentration used in the time-kill studies, ceftaroline was bacteriostatic for this isolate (see Fig. S1 in the supplemental material). When combined with DAP_{1 \times MIC}, ceftaroline demonstrated synergy with 4 (44.4%) isolates and bactericidal activity against 3 (33.3%) isolates (Table 3). Ceftaroline plus DAP_{2 \times MIC} was synergistic for 3/4 (75.0%) of the isolates not killed by DAP_{2 \times MIC} and bactericidal for 2 (50.0%) of these isolates (Table 3). The combination of ceftaroline plus DAP₁₈₀ was synergistic for the two isolates not killed by 180 $\mu\text{g/ml}$ daptomycin (Table 3 and Fig. 2) and bactericidal against only Efm25.

Ertapenem was synergistic with DAP_{0.5 \times MIC} for 4/8 isolates (50.0%, excluding Efc01, for which ertapenem alone was bactericidal) (see Fig. S1 in the supplemental material) and bactericidal in combination with DAP_{0.5 \times MIC} for 2 isolates (25.0%) (isolates Efc4 and Efm23) (Table 3). When tested with DAP_{1 \times MIC}, ertapenem yielded synergy with 3 isolates, and when tested with DAP, ertapenem was synergistic with DAP_{2 \times MIC} for isolate Efm25 alone (Table 3). When combined with DAP₁₈₀, ertapenem was synergistic for isolate Efm25.

Of note, none of the β -lactams when combined with any concentration of daptomycin demonstrated synergy against isolates Efm15 and Efm19, the two isolates evaluated without mutation to *liaFSR* genes.

Synergy between daptomycin and gentamicin, tigecycline, and rifampin. Four of the *Enterococcus* isolates included in this study, Efc01, Efc04, Efm13, and Efm25, did not display high-level

gentamicin resistance (Table 1). Gentamicin at 25 $\mu\text{g/ml}$ was consistently synergistic in combination with daptomycin for 3 (75.0%) of these isolates, Efc01, Efc04, and Efm25, at 24 h (Table 4). Synergy was not observed for isolate Efm13 when gentamicin was combined with DAP_{0.5 \times MIC} or DAP_{1 \times MIC}, but gentamicin was synergistic with DAP_{2 \times MIC} (Table 4). No synergy was observed at 6 h for any of these isolates. In addition, for isolate Efm16, there was synergy and bactericidal killing by the combination of DAP_{1 \times MIC} plus gentamicin (Table 3), despite the fact that this isolate expressed high-level gentamicin resistance (Table 1).

Tigecycline was synergistic in combination with DAP_{0.5 \times MIC} for 2 isolates and in combination with DAP_{1 \times MIC} for 3 isolates (Table 3). Tigecycline was synergistic with DAP_{2 \times MIC} for 2 isolates, Efm15 and Efm25, but was antagonistic for isolate Efc01. The addition of tigecycline to DAP₁₈₀ was also antagonistic for 2 isolates, Efm12 (5 \log_{10} less killing by the combination than by DAP₁₈₀ alone) and Efm13 (3.2 \log_{10} less killing by the combination than by DAP₁₈₀ alone) (Table 3 and Fig. 2B).

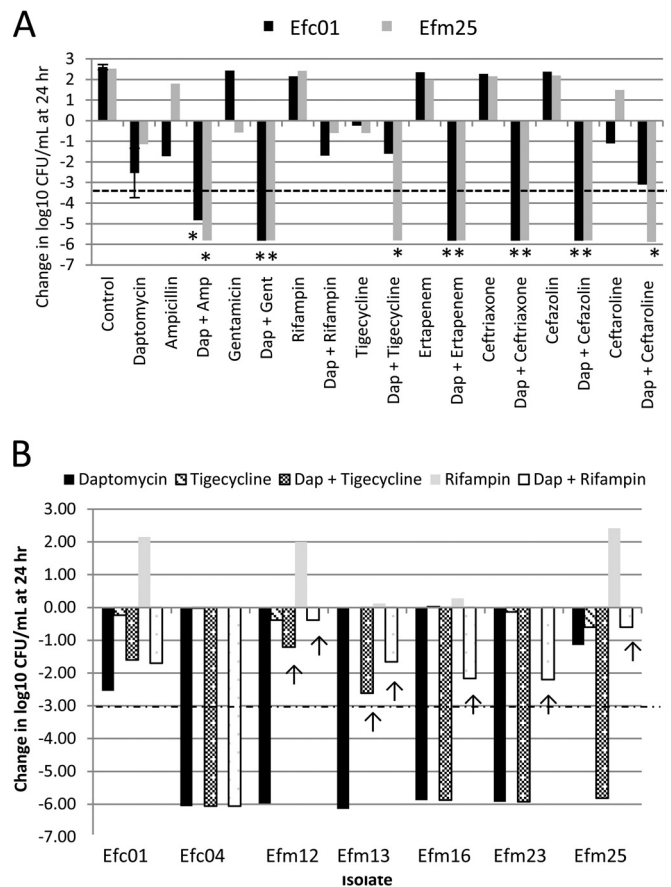


FIG 2 Change in \log_{10} CFU/ml after 24 h of antibiotic exposure to antimicrobials, alone or in combination, at 180 $\mu\text{g/ml}$ daptomycin. (A) Results for isolates Efc01 and Efm25, the only two isolates that displayed growth after 24 h of incubation in 180 $\mu\text{g/ml}$ daptomycin, in combination with test antimicrobials. Asterisks (*) represent synergistic interactions (e.g., $\geq 2 \log$ change in CFU/ml compared to daptomycin and/or the test antimicrobial alone). (B) Results of tigecycline and rifampin in combination with 180 $\mu\text{g/ml}$ daptomycin. Arrows (\uparrow) indicate antagonistic interactions (e.g., $\geq 2 \log$ higher CFU/ml compared to daptomycin at 180 $\mu\text{g/ml}$ alone). In both panels, a dashed line indicates a bactericidal effect (e.g., $\geq 99.9\%$ reduction in CFU/ml from time zero). Amp, ampicillin; Dap, daptomycin; Gent, gentamicin.

TABLE 4 Summary of 24-h time-kill testing for daptomycin at 4 concentrations in combination with gentamicin, rifampin, and tigecycline

Isolate (daptomycin MIC)	Result for indicated concn of daptomycin or synergy or antagonism result ^a											
	Gentamicin (25 µg/ml)				Tigecycline (0.8 µg/ml)				Rifampin (10 µg/ml)			
	0.5× MIC	1× MIC	2× MIC	180 µg/ml	0.5× MIC	1× MIC	2× MIC	180 µg/ml	0.5× MIC	1× MIC	2× MIC	180 µg/ml
Efc01 (64 µg/ml)	—	—	BC	—								
Efc04 (32 µg/ml)	BC	BC	NA ^c	NA								
Efm12 (64 µg/ml)			NA	NA								
Efm13 (64 µg/ml)			BC	NA								
Efm15 (16 µg/ml)				NA	BC	BC	BC	NA				
Efm16 (64 µg/ml)		BC	NA	NA	—	—	NA	NA				
Efm19 (16 µg/ml)			NA	NA			NA	NA				
Efm23 (64 µg/ml)			NA	NA			NA	NA				
Efm25 (64 µg/ml)	BC	BC	BC	—		BC	BC	—				A
Total no. (%) of isolates with synergy ^b	3 (33)	4(44)	3(75)	2(100)	2 (22)	3 (33)	2 (22)	1 (50)	0 (0)	2 (22)	1 (14)	0(0)
Total no. (%) of isolates with antagonism ^b	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)	0 (0)	1 (11)	2 (22)	0(0)	0(0)	3 (33)	5 (55)

^a Boldface data indicate synergy. —, daptomycin plus antimicrobial yielded a colony count at 24 h that was $\geq 2 \log_{10}$ CFU/ml lower than that seen with the most active antimicrobial alone. BC, bactericidal combination, where daptomycin plus antimicrobial yielded a $\geq 3 \log_{10}$ CFU/ml reduction in colony counts compared to the starting inoculum. A, antagonism, where daptomycin plus antimicrobial yielded a $\geq 2 \log_{10}$ CFU/ml increase in bacterial counts at 24 h compared to most active agent in combination. NA, daptomycin at this concentration was bactericidal and synergy could not be assessed.

^b Data indicate the total numbers and percentages of isolates for which neither antimicrobial was bactericidal alone.

larger collection of isolates and further demonstrated that, for those isolates that harbor *liaFSR* mutations, synergy between ampicillin and daptomycin exists even for isolates with very high (i.e., $>256 \mu\text{g/ml}$) daptomycin MICs. In contrast, the isolates evaluated by Diaz and colleagues had daptomycin MICs of 3 to 16 µg/ml. It should be noted that, while we found synergistic activity

between ampicillin and daptomycin for all 7 isolates with mutations to the *LiaFSR* pathway at $\text{DAP}_{0.5\times\text{MIC}}$, this combination was bactericidal for only 3 of the isolates. Similarly, at $\text{DAP}_{1\times\text{MIC}}$, the addition of ampicillin was synergistic for 6/7 (85.7%) of the isolates with mutations to *liaFSR* and bactericidal against all 6 of these (Table 2). Together, these findings suggest that, while mu-

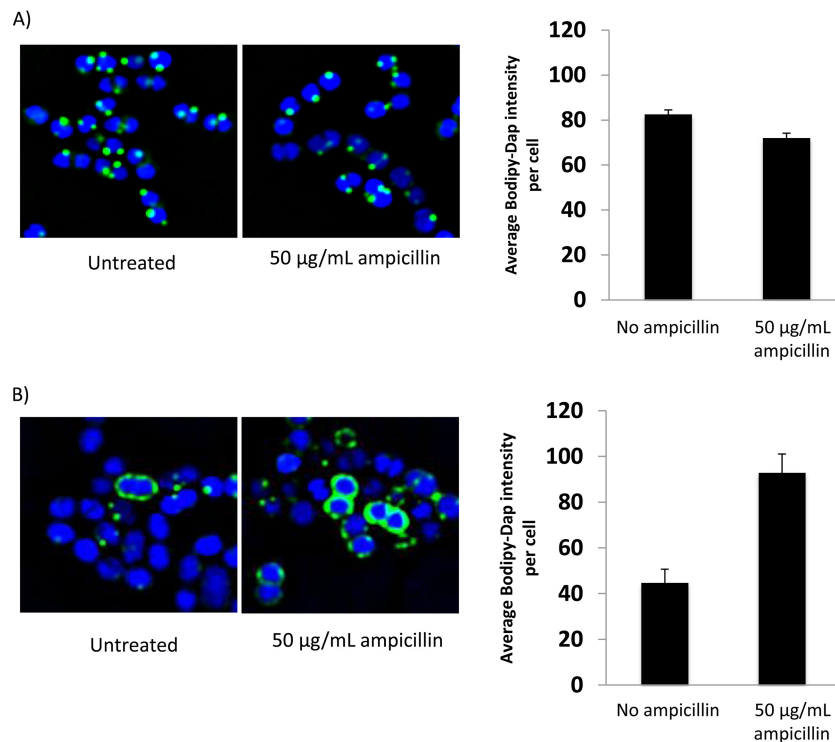


FIG 3 BDP-daptomycin (32 µg/ml) binding (20 min) studies for VRE Efm19 (no *liaFSR* mutation) (A) and Efm25 (with *liaFSR* mutation) (B) grown in the presence or absence of ampicillin at 50 mg/liter.

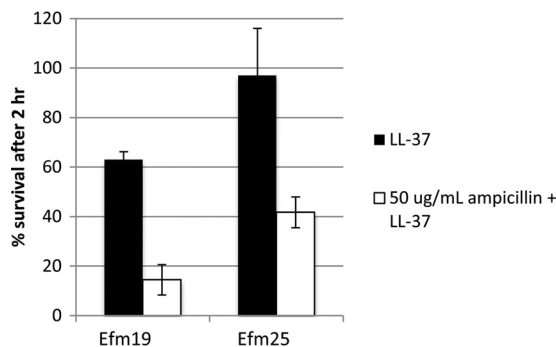


FIG 4 Effect of ampicillin on LL-37 killing of Efm19 (no *liaFSR* mutation) and Efm25 (with *liaFSR* mutation). Results are averages from 4 experiments, and error bars indicate standard deviations.

tation to *liaFSR* is more commonly associated with synergy between ampicillin and daptomycin, it is not predictive at all concentrations of daptomycin. To test the strains with high-level resistance, we chose the maximum total drug concentration achievable for a broad range of antibiotics in the study. However, it is notable that the free drug concentrations will likely differ depending on the degree of protein binding for each drug; it will be necessary to take this into consideration when extrapolating our results to the clinic. Given that a dose-dependent effect with daptomycin was observed to some extent for bactericidal activity and synergy with ampicillin against strains with *liaFSR* mutation, the maximally tolerated doses of daptomycin should be prescribed when daptomycin is used in combination with ampicillin. Regardless, no synergy was found between daptomycin and any β -lactam, at any concentration, for the two isolates with wild-type *liaFSR* alleles. As such, the absence of mutation to this region may rule out synergy. None of the other β -lactams evaluated here demonstrated consistent synergy with daptomycin.

The mechanism of synergy between ampicillin and daptomycin remains to be fully defined for enterococci. In a previous study, it was demonstrated that 50 $\mu\text{g/ml}$ ampicillin caused a net decrease in the relative positive surface charge of an ampicillin-resistant, daptomycin-susceptible (daptomycin MIC, 1 $\mu\text{g/ml}$) *E. faecium* isolate, associated with increased binding of daptomycin to the enterococcal cell membrane (17). Daptomycin exhibited bacteriostatic activity against this isolate, so the presence of a *liaFSR* mutation, although not examined by those authors, is assumed. These data suggest a charge-based mechanism for daptomycin-ampicillin synergy, and a similar effect has been noted for ceftaroline for both daptomycin-susceptible and DNS *E. faecium* isolates (18). In addition, we previously demonstrated for isolate 5938 (which is the same strain as Efm16) that treatment with 50 $\mu\text{g/ml}$ ampicillin caused an increase in cell wall thickness and increased LL-37 binding and activity (18). In the present study, treatment with 50 $\mu\text{g/ml}$ ampicillin significantly increased the binding of BODIPY-daptomycin to the cell membrane of Efm25 but not to that of Efm19, which does not harbor mutation to *liaFSR*. Interestingly, while ampicillin enhanced the binding of daptomycin only to Efm25, it sensitized both Efm19 and Efm25 to LL-37, the human cathelicidin host defense peptide (Fig. 4). Further studies are required to more fully define these different mechanisms, but the data suggest that the concept of the charge-based mechanism for synergy may be overly simplistic. However, it is

important that the relative levels of tolerance for LL-37 and, potentially, other cationic host defense peptides conferred by *liaFSR* mutations may be selected for by persistent endovascular infections, as has been shown in *mprF* for *Staphylococcus aureus* (40), perhaps even in the absence of daptomycin selective pressure.

The advantage conferred to the pathogen by its ability to resist killing by both the innate immune system and daptomycin appears, therefore, to result in an “Achilles’ heel” whereby β -lactams such as ampicillin may be employed as adjunctive agents, and only in these settings would such a practice be beneficial. This hypothesis is supported circumstantially by the study by Moise et al. (38) which shows that the addition of β -lactams to daptomycin for VRE bloodstream infections is beneficial in improving outcome only in cases where the daptomycin MIC is 3 to 4 $\mu\text{g/liter}$ (presumably in a *liaFSR* mutation-enriched subgroup, on the basis of prior data [10]) and not when the daptomycin MIC is ≤ 2 $\mu\text{g/liter}$ (38).

Synergy between daptomycin and gentamicin, for *E. faecium* isolates with high-level susceptibility to gentamicin, has been noted previously (41), but little to no clinical data exist for this combination (42). Unlike the case with the glycopeptides (43), it would appear that high-level gentamicin resistance does not necessarily abolish *in vitro* synergy, as was seen for isolate Efm16; however, this interaction requires further evaluation, and no data exist to suggest that this may be the case *in vivo*.

Several reports have demonstrated effective use of daptomycin plus tigecycline for the treatment of endocarditis caused by *E. faecium* (44–46). In our hands, this combination was frequently antagonistic, but again, *in vivo* data are required to confirm this finding. Similarly, we found antagonism for a significant number of isolates with daptomycin plus rifampin, but this has not been demonstrated by an *in vivo* model.

In summary, we systematically evaluated by *in vitro* time-kill studies the effect of daptomycin in combination with other antimicrobials for a collection of 9 DNS enterococcal isolates that exhibited a broad range of MICs and different resistance-confering mutations. We found that ampicillin plus daptomycin yielded the most consistent synergy but did so only for isolates with mutations to the *liaFSR* system. Daptomycin binding was found to be enhanced by the addition of ampicillin for such mutations but not for an isolate without mutations to this system. In contrast, ampicillin enhanced the killing of LL-37 against DNS *E. faecium*, regardless of the presence of *liaFSR* mutation. These data lend support to the growing body of evidence that combination therapy consisting of daptomycin plus ampicillin may be helpful in eradicating refractory VRE infections by counteracting the fitness advantages of reduced daptomycin susceptibility and resistance to killing by cathelicidin and other host defense peptides conferred by these mutations.

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