

Impact of Glycopeptide Resistance in *Staphylococcus aureus* on the Dalbavancin *In Vivo* Pharmacodynamic Target

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Dalbavancin is a novel lipoglycopeptide with activity against *Staphylococcus aureus*, including glycopeptide-resistant isolates. The *in vivo* investigation reported here tested the effects of this antibiotic against seven *S. aureus* isolates with higher MICs, including several vancomycin-intermediate strains. Results of 1-log kill and 2-log kill were achieved against seven and six of the isolates, respectively. The mean free-drug area under the concentration-time curve (fAUC)/MIC values for net stasis, 1-log kill, and 2-log kill were 27.1, 53.3, and 111.1, respectively.

The increasing rates of resistance among hospital- and community-acquired bacterial pathogens such as *Staphylococcus aureus*, coagulase-negative staphylococci, and enterococci have prompted attempts to discover new antimicrobials with activities against multidrug-resistant Gram-positive pathogens (1–6). Dalbavancin is a new lipoglycopeptide antibiotic with activity against multidrug-resistant Gram-positive organisms (4, 7–9). In addition to enhanced antimicrobial potency, the compound possesses a unique pharmacokinetic (PK) profile that includes an extremely long elimination half-life of more than 1 week (10–12). Clinical development of the compound has thus far demonstrated success for the treatment of skin and soft tissue infections and catheter-related bloodstream infections (13–18). Once-weekly administration of the doses used in these trials has been shown to produce free-drug trough concentrations exceeding the MIC_{90s} of Gram-positive pathogens from large surveillance databases (17, 19–23).

The current studies were designed to define the pharmacodynamic (PD) target for dalbavancin against *S. aureus* strains with dalbavancin MICs at or above the current FDA breakpoint (≥ 0.12 $\mu\text{g/ml}$), some of which were vancomycin-intermediate *S. aureus* (VISA) strains (24–29). The results from these studies provide a pharmacodynamic rationale in support of the current clinical dosing regimens. Furthermore, the data provide a starting point for the development of revised susceptibility breakpoints for this new compound.

Seven strains of *Staphylococcus aureus* (including four vancomycin-intermediate *S. aureus* [VISA] strains) were studied (Table 1). The dalbavancin and vancomycin MIC values were determined in triplicate using CLSI reference broth microdilution methods, in the presence of polysorbate 80 (30). The dalbavancin MIC range for the *S. aureus* isolates was 0.12 to 0.50 $\mu\text{g/ml}$. Ani-

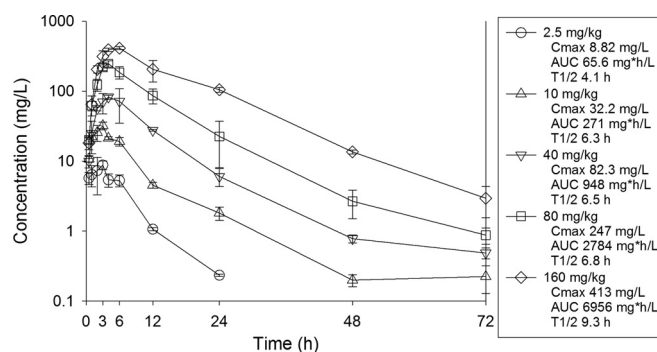


FIG 1 Plasma pharmacokinetics of dalbavancin in mice following intraperitoneal administration. Each symbol represents the mean and standard deviation from three mice. The drug concentration values presented represent total (protein-bound and unbound) drug. The AUC values represent 0 to infinity. C_{max}, maximal drug concentration; T_{1/2}, half-life.

imals were maintained in accordance with the criteria of the Association for Assessment and Accreditation of Laboratory Animal Care. All animal studies were approved by the Animal Research Committee of the William S. Middleton Memorial Veterans Hospital. The neutropenic murine thigh infection model was used for all studies. Mice were inoculated with 10^7 CFU/ml of each strain. Single-dose plasma pharmacokinetic studies were performed with thigh-infected mice given intraperitoneal doses (0.2 ml/dose) of dalbavancin (2.5, 10, 40, 80, or 160 mg/kg). Dalbavancin plasma concentrations were measured with a liquid chromatography-tandem mass spectrometry (LC-MS/MS) assay (Fig. 1); the lower limit of quantification for the assay was 0.05 $\mu\text{g/ml}$. Sample analysis precision (coefficient of variation [CV]) ranged from 5% to 6.4%, and accuracy (bias) ranged from -3.5% to -10.0% . Peak

TABLE 1 Study strains and dalbavancin *in vitro* susceptibility

<i>S. aureus</i> isolate	MIC (mg/liter)	
	Dalbavancin	Vancomycin
LSI1653	0.25	2
LSI1848	0.12	2
LSI1854	0.5	2
LSI1856	0.25	4 (VISA)
LSI1857	0.25	4 (VISA)
LSI1861	0.25	4 (VISA)
LSI1862	0.5	4 (VISA)

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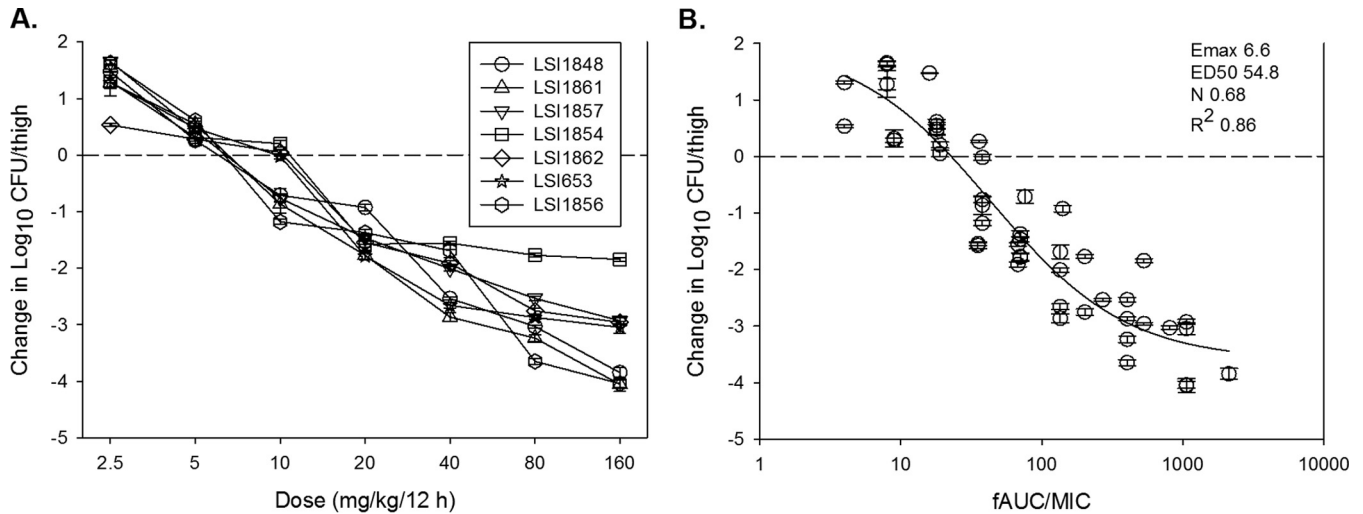


FIG 2 *In vivo* dose-dependent effects of dalbavancin against seven select *S. aureus* isolates in a neutropenic mouse thigh model. (A) Dalbavancin exposure expressed at dose level (mg/kg/12 h). (B) Exposure expressed as $fAUC/MIC$. Each symbol represents the mean and standard deviation from four thighs. Dalbavancin exposure is expressed as the 24-h $fAUC/MIC$. The burden of organisms was measured at the start and end of therapy. The horizontal dashed line at 0 represents the burden of organisms in the thighs of mice at the start of therapy. Data points below the line represent killing, and points above the line represent growth. R^2 represents the coefficient of determination. The 50% effective dose (ED_{50}) represents the AUC/MIC associated with 50% of the maximal effect (E_{max}), and N is the slope of the relationship or the Hill coefficient. The line drawn through the data points is the best-fit line based on the sigmoidal E_{max} formula.

levels were observed by 2 to 6 h. Dalbavancin exhibited relatively linear pharmacokinetics, based on the dose-area under the concentration-time curve (AUC) relationship. The half-life was long and varied from 4.1 to 9.31 h. A protein binding value of 98.4%, based on prior studies in this model (31), was used.

The *in vivo* virulence of the *S. aureus* isolates was similar in the untreated control mice, based on the increase in thigh burden over the treatment period, i.e., $2.30 \pm 0.14 \log_{10}$ CFU/thigh. Two hours after infection, dalbavancin was administered via the intraperitoneal route, with one of seven 2-fold-escalating doses of dalbavancin (2.5, 5, 10, 20, 40, 80, and 160 mg/kg) being administered every 12 h for a 6-day treatment period. Untreated control groups were sampled at the start of therapy and at the end of the study. The thighs were removed from the animals and immediately processed for CFU determination. The results of these studies were analyzed by using a sigmoidal dose-effect model (32). The magnitude of the PK/PD index associated with each endpoint dose was calculated with the following equation: $\log_{10} D = \log_{10} [E/(E_{max} - E)]/(N +$

$\log_{10} ED_{50})$, where E is the control growth for the static dose (D), E is the control growth $- 1$ log unit for D for 1-log kill, and E is the control growth $- 2$ log units for D for 2-log kill.

Results of 1-log kill and 2-log kill were achieved against seven and six of the isolates, respectively (Fig. 2A and Table 2). The dalbavancin *in vivo* exposure-response data were also considered relative to the PK/PD-linked driver AUC/MIC , using concentrations of free drug. Drug accumulation was calculated and included in AUC estimates. Using a sigmoidal E_{max} model, the data fit was strong for the seven-strain data set ($R^2 = 0.86$), as shown in Fig. 2B. The numerical AUC/MIC values associated with each of the three treatment endpoints are also shown in Table 2. Net stasis was observed with a dalbavancin free-drug $AUC (fAUC)/MIC$ value near 25. $fAUC/MIC$ values near 50 and 100 were associated with 1-log and 2-log reductions, respectively, in organism burdens in the neutropenic mice.

These PK/PD targets are lower than those observed previously with wild-type *S. aureus* strains in the same model (31). This is partly due to lower pharmacokinetic values measured in the pres-

TABLE 2 *In vivo* efficacy of dalbavancin against select *S. aureus* isolates, using $fAUC/MIC$ as the predictive pharmacodynamic index

Strain	Stasis		1-log kill		2-log kill	
	24-h dose (mg/kg)	24-h $fAUC/MIC$	24-h dose (mg/kg)	24-h $fAUC/MIC$	24-h dose (mg/kg)	24-h $fAUC/MIC$
LSI1848	15.17	56.49	31.45	112.81	62.75	214.21
LSI1861	13.55	25.00	24.63	45.35	44.38	77.35
LSI1857	14.34	26.59	26.72	48.74	60.05	102.72
LSI1854	15.00	13.95	35.80	31.73		
LSI1862	12.64	11.60	32.92	29.39	85.46	76.66
LSI653	14.93	27.77	27.20	49.52	54.09	93.07
LSI1856	15.21	28.32	30.88	55.49	60.05	102.73
Mean	14.41	27.10	29.94	53.29	61.13	111.12
Median	14.93	26.59	30.88	48.74	60.05	97.90
SD ^a	0.98	14.62	3.93	27.93	13.62	51.81

^a SD, standard deviation.

ent study, perhaps due to differences in the drug assay method. Of note, the present kinetic study included a robust sampling scheme and a more sensitive and accurate drug assay method, compared to the prior animal model investigation; we used a specific LC-MS/MS assay, in contrast to the prior bioassay. The treatment studies were otherwise similar with respect to animal species, neutropenia, antibiotic (dalbavancin), drug preparation, route of administration, treatment duration, study endpoints, and data analysis.

The present studies were designed to discern the PK/PD impact of infection with less common *S. aureus* strains that had dalbavancin MICs at or above the current dalbavancin FDA breakpoint (≥ 0.12 $\mu\text{g/ml}$). Dalbavancin demonstrated potent *in vivo* activity against *S. aureus* strains with higher MICs, including those exhibiting a VISA phenotype. While it will be important to corroborate these preclinical findings with data from patients, consideration of the AUC/MIC targets from these studies in the context of human pharmacokinetics suggests a safe treatment margin against these higher-MIC isolates. If the steady-state kinetics of dalbavancin in patients are considered relative to the stasis, 1-log kill, and 2-log kill AUC/MIC targets in this study, then the MIC breakpoints would be revised to 4, 2, and 1 $\mu\text{g/ml}$, respectively.

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