

# Activity of Dalbavancin against *Staphylococcus* spp., *Streptococcus* spp., and *Enterococcus* spp. Isolated by Clinical Laboratories in Canadian Hospitals: CANWARD 2007-2017

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## Introduction

Dalbavancin is a semisynthetic lipoglycopeptide structurally related to teicoplanin (1, 2). Like other glycopeptides, dalbavancin inhibits growth of Gram-positive bacterial pathogens. Dalbavancin inhibits cell wall synthesis in Gram-positive bacteria by forming complexes with C-terminal d-alanyl-d-alanine residues of growing peptidoglycan chains. In addition, dalbavancin appears to have the unique ability to form dimers and to anchor its lipophilic side chain in bacterial membranes which increases its target affinity and increases its potency. Consequently, dalbavancin possesses more potent *in vitro* bactericidal activity than vancomycin or teicoplanin against many resistant Gram-positive organisms. Dalbavancin is indicated for acute bacterial skin and skin structure infections (ABSSSI) caused by designated susceptible strains of Gram-positive microorganisms. Dalbavancin can be used as a single dose regimen of 1500mg administered via intravenous infusion, using a total infusion time of 30 minutes.

The purpose of this study was to compare the activity of dalbavancin, vancomycin and comparators against Gram-positive pathogens causing infections in patients in Canadian hospitals.

## Materials and Methods

**Participating Sites:** From January 2007 to October 2017, tertiary-care medical centres in major population centres in 8 of the 10 provinces in Canada were recruited (2, 3). These sites were geographically distributed in a population based fashion.

**Bacterial Isolates:** Tertiary-care medical centres submitted pathogens from patients attending hospital clinics, emergency rooms, medical and surgical wards, and intensive care units. From January 2007 through October 2017, inclusive, each study site was asked to submit clinical isolates (consecutive, one per patient, per infection site) from inpatients and outpatients with respiratory, urine, wound, and bloodstream infections. The medical centres submitted clinically significant isolates from patients with a presumed infectious disease. Surveillance swabs, eye, ear, nose and throat swabs were excluded. We also excluded anaerobic organisms. Isolate identification was performed by the submitting site and confirmed at the reference site as required, based on morphological characteristics and antimicrobial susceptibility patterns. Isolates were shipped on Amies semi-solid transport media to the coordinating laboratory (Health Sciences Centre, Winnipeg, Canada), subcultured onto appropriate media, and stocked in skim milk at -80° C until minimum inhibitory concentration (MIC) testing was carried out.

**Antimicrobial Susceptibilities:** The *in vitro* activity of selected antimicrobials was determined by broth microdilution in accordance with (CLSI) guidelines (4). Antimicrobial MIC interpretive standards were defined according to CLSI breakpoints (5). The MICs of the antimicrobial agents were determined using 96-well custom designed microtitre plates. These plates contained doubling antimicrobial dilutions in 100µL/well of cation adjusted Mueller-Hinton broth and inoculated to achieve a final concentration of approximately 5 x 10<sup>5</sup> CFU/mL then incubated in ambient air for 24 hours prior to reading. Colony counts were performed periodically to confirm inocula. Quality control was performed using ATCC QC organisms including *S. pneumoniae* 49619, *S. aureus* 29213, *E. faecalis* 29212.

CLSI breakpoints were used for dalbavancin: *Staphylococcus aureus* (including methicillin-resistant isolates) susceptible (S) ≤0.25µg/mL; *Streptococcus pyogenes* and *Streptococcus agalactiae* (S) ≤0.25µg/mL; and *Enterococcus faecalis*, (S) ≤0.25µg/mL.

## Results

**Table 1. Activity of dalbavancin and comparators against Gram-positive cocci from CANWARD 2007-2017**

Organism (no. tested) / Antimicrobial Agent	% of Isolates per Category			MIC <sub>50</sub>	MIC <sub>90</sub>	Range Min	Range Max
	S	I	R				
<b>Methicillin-susceptible <i>S. aureus</i> (MSSA, n=5280)</b>							
Cefazolin	NA <sup>a</sup>			≤ 0.5	1	≤ 0.5	32
Clarithromycin	75.3	0.6	24.1	0.25	> 32	≤ 0.03	> 32
Clindamycin	93.0	0.5	6.5	≤ 0.25	≤ 0.25	≤ 0.03	> 8
<b>Dalbavancin</b>	100.0			0.06	0.06	≤ 0.03	0.25
Daptomycin	100.0			0.25	0.25	≤ 0.03	1
Levofloxacin	90.1	0.3	9.6	0.25	1	≤ 0.06	> 32
Linezolid	100.0			2	4	≤ 0.12	4
Meropenem	NA <sup>a</sup>			0.12	0.25	≤ 0.03	16
Pip-Tazo	NA <sup>a</sup>			≤ 1	≤ 1	≤ 1	64
Tigecycline	99.8			0.2	0.25	≤ 0.03	2
TMP/SMX	99.5			≤ 0.12	≤ 0.12	≤ 0.12	> 8
Vancomycin	100.0			1	1	≤ 0.12	2
<b>Methicillin-resistant <i>S. aureus</i> (MRSA, n=1482)</b>							
Cefazolin	NA <sup>a</sup>			64	> 128	≤ 0.5	> 128
Clarithromycin	14.0	0.1	85.9	> 32	> 32	≤ 0.03	> 32
Clindamycin	51.1	0.1	48.8	0.25	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	99.9			0.06	0.06	≤ 0.03	1
Daptomycin	99.9			0.1	0.25	0.5	0.06
Levofloxacin	14.1			85.9	> 32	0.12	> 32
Linezolid	100.0			2	4	≤ 0.12	4
Meropenem	NA <sup>a</sup>			4	32	0.12	> 64
Pip-Tazo	NA <sup>a</sup>			32	128	≤ 1	512
Tigecycline	99.7			0.3	0.5	0.06	1
TMP/SMX	93.3			≤ 0.12	≤ 0.12	≤ 0.12	> 8
Vancomycin	99.8	0.2		1	1	≤ 0.25	4
<b>Community-associated methicillin resistant <i>S. aureus</i> (CA-MRSA, n=484)</b>							
Cefazolin	NA <sup>a</sup>			16	64	≤ 0.5	> 128
Clarithromycin	24.8	0.2	75.0	> 32	> 32	≤ 0.12	> 32
Clindamycin	86.0			14.0	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	99.8			0.2	0.06	≤ 0.03	1
Daptomycin	100.0			0.25	0.5	0.12	1
Levofloxacin	40.0			60.0	8	0.12	32
Linezolid	100.0			2	2	0.5	4
Meropenem	NA <sup>a</sup>			2	4	0.12	32
Pip-Tazo	NA <sup>a</sup>			16	64	≤ 1	256
Tigecycline	100.0			0.25	0.25	0.06	0.5
TMP/SMX	99.8			≤ 0.12	≤ 0.12	≤ 0.12	4
Vancomycin	99.8	0.2		1	1	≤ 0.5	4
<b>Healthcare-associated methicillin resistant <i>S. aureus</i> (HA-MRSA, n=949)</b>							
Cefazolin	NA <sup>a</sup>			128	> 128	≤ 0.5	> 128
Clarithromycin	5.9			94.1	> 32	≤ 0.03	> 32
Clindamycin	31.5	0.1	68.4	> 8	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	100.0			0.06	0.06	≤ 0.03	0.25
Daptomycin	99.9			0.1	0.25	0.5	0.06
Levofloxacin	2.9			97.1	> 32	0.12	> 32
Linezolid	100.0			2	4	≤ 0.12	4
Meropenem	NA <sup>a</sup>			16	> 32	0.12	> 64
Pip-Tazo	NA <sup>a</sup>			64	128	≤ 1	512
Tigecycline	99.6			0.4	0.25	0.5	0.06
TMP/SMX	89.7			10.3	≤ 0.12	8	≤ 0.12
Vancomycin	99.8	0.2		1	1	≤ 0.25	4
<b>Methicillin-susceptible <i>S. epidermidis</i> (MSSE, n=637)</b>							
Cefazolin	NA <sup>a</sup>			1	4	≤ 0.5	8
Clarithromycin	35.5	2.0	62.5	> 16	> 32	≤ 0.03	> 32
Clindamycin	65.3	1.1	33.6	≤ 0.12	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	NA <sup>a</sup>			≤ 0.03	0.06	≤ 0.03	1
Daptomycin	100.0			0.12	0.25	≤ 0.03	1
Levofloxacin	51.9	1.1	47.0	0.25	> 32	0.12	> 32
Linezolid	100.0			0.5	2	≤ 0.12	2
Meropenem	NA <sup>a</sup>			1	8	≤ 0.03	> 32
Pip-Tazo	NA <sup>a</sup>			≤ 1	4	≤ 1	64
Tigecycline	99.5			0.5	0.12	0.5	0.03
TMP/SMX	67.2			32.8	≤ 0.12	8	≤ 0.12
Vancomycin	100.0			1	2	≤ 0.12	4
<b>Methicillin-resistant <i>S. epidermidis</i> (MRSE, n=90)</b>							
Cefazolin	NA <sup>a</sup>			128	> 128	32	> 128
Clarithromycin	16.7			83.3	> 32	≤ 0.03	> 32
Clindamycin	18.9			81.1	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	NA <sup>a</sup>			≤ 0.03	0.06	≤ 0.03	0.06
Daptomycin	100.0			0.12	0.25	≤ 0.03	0.5
Levofloxacin	2.2			95.6	> 32	> 32	> 32
Linezolid	100.0			1	1	≤ 0.12	2
Meropenem	NA <sup>a</sup>			32	32	4	64
Pip-Tazo	NA <sup>a</sup>			32	64	8	128
Tigecycline	100.0			0.12	0.25	0.06	0.5
TMP/SMX	20.0			80.0	4	8	> 8
Vancomycin	100.0			1	2	≤ 0.12	2

**Table 2. MIC distribution of dalbavancin against Gram-positive cocci from CANWARD**

Organism	No. tested	MIC (Minimum Inhibitory Concentration µg/mL)										
		≤0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	>16
Methicillin-susceptible <i>S. aureus</i>	5280	1798	3383	94	5							
Methicillin-resistant <i>S. aureus</i>	1482	435	1013	30	3							
Community-associated MRSA	484	123	351	8	1							
Healthcare-associated MRSA	949	303	623	21	2							
Methicillin-susceptible <i>S. epidermidis</i>	637	416	206	14								
Methicillin-resistant <i>S. epidermidis</i>	90	70	20									
<i>S. pneumoniae</i>	1744	1718	21	5								
Penicillin-resistant <i>S. pneumoniae</i>	75	75										
<i>E. faecalis</i>	899	394	472	30	2	1						
<i>E. faecium</i>	338	56	126	85	11	8						
<i>Enterococcus</i> spp. *	570	224	309	35	2							
VRE	68	3	2	4	2	6	3	9	7	6	4	22

\* Not specified CANWARD 2007-2010

Organism (no. tested) / Antimicrobial Agent	% of Isolates per Category			MIC <sub>50</sub>	MIC <sub>90</sub>	Range Min	Range Max
	S	I	R				
<b><i>S. pneumoniae</i> (n=1744)</b>							
Ceftriaxone	99.4	0.4	0.2	≤ 0.12	≤ 0.12	≤ 0.12	4
Clarithromycin	79.5	4.1	16.4	≤ 0.03	4	≤ 0.03	> 32
Clindamycin	93.4	0.6	6.0	≤ 0.12	≤ 0.12	≤ 0.12	> 64
<b>Dalbavancin</b>	NA <sup>a</sup>			≤ 0.03	≤ 0.03	≤ 0.03	0.12
Doxycycline	87.1	1.4	11.5	≤ 0.25	1	≤ 0.25	> 16
Levofloxacin	99.1	0.1	0.8	1	1	≤ 0.06	32
Linezolid	100.0			0.5	1	≤ 0.12	2
Meropenem	95.9	2.7	1.3	≤ 0.06	≤ 0.06	≤ 0.06	2
Penicillin	81.3	14.4	4.3	≤ 0.03	0.25	≤ 0.03	> 8
Tigecycline	99.7			0.3	0.03	≤ 0.015	0.25
TMP/SMX	84.7	6.7	8.6	≤ 0.12	2	≤ 0.12	> 8
Vancomycin	100.0			≤ 0.25	≤ 0.25	≤ 0.25	1
<b>Penicillin-resistant <i>S. pneumoniae</i> (Pen-R, n=75)</b>							
Ceftriaxone	89.3	8.0	2.7	0.5	2	≤ 0.06	4
Clarithromycin	34.7	2.7	62.7	4	> 32	≤ 0.03	> 32
Clindamycin	54.7	2.7	42.7	≤ 0.12	> 64	≤ 0.12	> 64
<b>Dalbavancin</b>	NA <sup>a</sup>			≤ 0.03	≤ 0.03	≤ 0.03	≤ 0.03
Doxycycline	38.7			61.3	1	4	≤ 0.25
Levofloxacin	98.7			1.3	1	0.25	16
Linezolid	100.0			0.5	1	0.25	2
Meropenem	29.3	42.7	28.0	0.5	1	≤ 0.06	2
Penicillin	100.0			2	4	2	> 8
Tigecycline	98.7			1.3	0.03	0.06	≤ 0.015
TMP/SMX	18.7			6.7	74.7	4	> 8
Vancomycin	100.0			≤ 0.25	0.5	≤ 0.25	1
<b><i>E. faecalis</i> (n=899)</b>							
Ciprofloxacin	59.7	12.6	27.7	1	> 16	≤ 0.06	> 16
Clarithromycin	NA <sup>a</sup>			> 32	> 32	≤ 0.03	> 32
Clindamycin	NA <sup>a</sup>			> 8	> 8	≤ 0.12	> 8
<b>Dalbavancin</b>	99.9			0.06	0.06	≤ 0.03	0.5
Daptomycin	92.3	7.7		0.5	1	≤ 0.03	4
Ertapenem	NA <sup>a</sup>			8	16	0.25	> 32
Levofloxacin	65.6	1.0	33.3	2	> 32	0.25	> 32
Linezolid	90.2	9.8		2	2	0.5	4
Meropenem	NA <sup>a</sup>			4	8	≤ 0.06	> 32
Pip-Tazo	NA <sup>a</sup>						