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Analysis of Oritavancin Activity Against Gram-Positive Clinical Isolates **Responsible for Bacterial Endocarditis in United States and European** Hospitals (2008-2016) MA Pfaller, HS Sader, D Shortridge, RK Flamm, RE Mendes JMI Laboratories, North Liberty, Iowa, USA

Introduction

- Infective endocarditis (IE) is a life-threatening infectious disease
- Despite recent diagnostic and treatment advances, mortality by IE remains high, with more than 30% of patients affected dying within 1 year after diagnosis
- Presently, staphylococci and streptococci combined cause approximately 80% of IE cases
- Staphylococcus aureus remains the dominant pathogen, accounting for 25–30% of cases
- About 30% of all S. aureus causing IE are methicillinresistant (MRSA)
- Vancomycin is an agent of choice for MRSA IE despite poor penetration into vegetations and slow bactericidal activity
- Clinical failures are frequently observed
- Coagulase-negative staphylococci (CoNS) account for ~11% of cases
- Viridans group streptococci (VGS) cause ~30% of cases
- Enterococci cause ~10% of cases
- Oritavancin belongs to the lipoglycopeptide class of antimicrobial agents. It interrupts bacterial cell wall synthesis in addition to disrupting cell membrane integrity. Together, these mechanisms result in rapid bactericidal activity
- Oritavancin demonstrates potent in vitro activity against S. aureus (including MRSA), streptococci, and vancomycinsusceptible (VSE) and -resistant (VRE) enterococci, and its potency has remained stable over time
- This study analyzed the in vitro activity of oritavancin against gram-positive organisms causing endocarditis. Isolates were collected from United States (US) and European (EU) sites

Materials and Methods

Bacterial isolates

- A total of 424 organisms recovered from patients with a diagnosis of bacterial endocarditis at US and EU sites during the SENTRY Antimicrobial Surveillance Program (2008–2016) were included (Table 1)
- Isolates were determined to be clinically significant based on local guidelines and were submitted to a central monitoring laboratory (JMI Laboratories, North Liberty, Iowa, USA)
- Participating laboratories initially identified isolates and JMI confirmed bacterial identifications by standard algorithms supported by matrix-assisted laser desorption ionization-time of flight mass spectrometry (Bruker Daltonics, Bremen, Germany)

Antimicrobial susceptibility testing

- Isolates were tested for susceptibility by broth microdilution following guidelines in the CLSI M07–A10 document
- Testing used reference 96-well panels manufactured by JMI Laboratories
- Quality assurance was performed by concurrently testing CLSI-recommended quality control reference strains (S. aureus ATCC 29213, Enterococcus faecalis ATCC 29212, and Streptococcus pneumoniae ATCC 49619)

Breakpoint criteria for comparator agents were from CLSI (M100-S27) and EUCAST (Breakpoint tables v7.1)

• Oritavancin had similar MIC₉₀ values (0.06 μg/mL) against *S. aureus* and CoNS, inhibiting 98.8% of these isolates at $\leq 0.12 \,\mu$ g/mL

Results

- Oritavancin MIC₅₀ values were 8- to 32-fold lower than those for vancomycin, daptomycin, and ceftaroline against staphylococci
- Oritavancin showed MIC values against *E. faecium* (MIC_{50/90}, ≤0.008/0.03 µg/mL) that were 2-fold lower than against *E. faecalis* (MIC_{50/90}, 0.015/0.03 μg/mL; 97.5% susceptible against all or 100% susceptible against indicated vancomycin-susceptible isolates)

Table 1 Antimicrobial activity of oritavancin tested against the main organisms and organism groups of isolates

Organism /	No. of isolates at MIC (µg/mL; <i>cumulative %</i>)								
organism group (number of isolates)	≤0.008	0.015	0.03	0.06	0.12	0.25	0.5	MIC ₅₀	MIC ₉₀
S. aureus (212)	7 3.3	74 38.2	73 72.6	42 92.5	14 99.1	2 100.0		0.03	0.06
MRSA (67)	2 3.0	21 34.3	25 71.6	13 91.0	4 97.0	2 100.0		0.03	0.06
MSSA (145)	5 3.4	53 40.0	48 73.1	29 93.1	10 100.0			0.03	0.06
CoNS (47)	11 23.4	6 36.2	14 66.0	13 93.6	2 97.9	1 100.0		0.03	0.06
<i>E. faecalis</i> (81)	37 45.7	33 86.4	8 96.3	0 96.3	1 97.5	0 97.5	2 100.0	0.015	0.03
E. faecium (21)	13 61.9	3 76.2	4 95.2	0 95.2	1 100.0			≤0.008	0.03
Vancomycin-susceptible (12)	11 91.7	0 91.7	1 100.0					≤0.008	0.008
VanA-phenotype (9)	2 22.2	3 55.6	3 88.9	0 88.9	1 100.0			0.015	
β-hemolytic streptococci (24)	1 <i>4.2</i>	6 29.2	6 54.2	8 87.5	2 95.8	1 100.0		0.03	0.12
Viridans group streptococci (39)	19 <i>48.7</i>	5 61.5	5 74.4	3 82.1	4 92.3	3 100.0		0.015	0.12

• Among the 424 isolates, 212 (50.0%) were S. aureus (31.6% MRSA), 47 (11.1%) were CoNS, 81 (19.1%) were Enterococcus faecalis, 21 (5.0%) were *Enterococcus faecium*, 24 (5.7%) were beta-hemolytic streptococci (BHS), and 39 (9.2%) were VGS (Table 1)

- Oritavancin inhibited 98.1% of all enterococci. including vancomycin-resistant isolates at ≤0.12 µg/mL
- Vancomycin, daptomycin, linezolid (MIC_{50/90}, $1/2 \mu g/mL$), and ampicillin (MIC_{50/90}, \leq 1/2 µg/mL) were similarly active against *E. faecalis*, while daptomycin and linezolid had coverage (100.0% susceptibility) against *E. faecium*
- Overall, BHS isolates were highly susceptible to all agents tested, except for erythromycin (70.8% susceptible), clindamycin (75.0% susceptible), and tetracycline (43.5% susceptible)
- Oritavancin was the most active agent (MIC₉₀, 0.12 μ g/mL; 100.0% susceptible) tested against VGS

Table 2. Comparative antimicrobial activity of oritavancin and comparator agents tested against the main organisms and organism groups of isolates causing IE in Europe and the United States

Organism / Organism / Organism / esistance group CLSI/ resister no tested) FUCASTa (use)		Organism / resistance group				CLSI/ FUCAST ^a							
Antimicrobial agent	MIC ₅₀	MIC ₉₀	Range	eucasiª %S	(number tested/ antimicrobial agent)	MIC ₅₀	MIC ₉₀	Range	8 %S				
MSSA (145)					E. faecium (21)								
Oritavancin	0.03	0.06	0.008 - 0.12	100.0/100.0	Oritavancin	≤0.008	0.03	≤0.008 – 0.12	_/_				
Ceftaroline	0.25	0.5	0.06 - 0.5	100.0/100.0	Ampicillin	>8	>8	1 – >8	4.8/4.8				
Clindamycin	≤0.25	≤0.25	≤0.25 - >2	96.6/96.6	Ceftaroline	>8	>8	2 -> 8	_/_				
Daptomycin	0.25	0.5	≤0.12 – 1	100.0/100.0	Daptomycin	2	2	0.12 – 4	100.0 /				
Erythromycin	≤0.25	>2	≤0.25 - >2	80.0/80.0	Ervthromvcin	>2	>2	0.5 ->2	5.3/-				
Levofloxacin	≤0.5	≤0.5	≤0.5 – >4	95.2/95.2		>4	>4	1 ->4	4 8/4 8				
Linezolid	1	2	0.5 – 2	100.0/100.0	Linezolid	1	2	05-2	100 0/100 0				
ТМХ	≤0.5	≤0.5	≤0.5	100.0/100.0	Toicoplanin	۱ دی	2 \16	0.0 − 2 <2 >16	57 1/57 1				
Vancomycin	1	1	0.25 – 2	100.0/100.0	Vanaamvoin	<u> </u>	>10	$\leq 2 = >10$	57.1/57.1				
Teicoplanin	≤2	≤2	≤2	100.0/100.0	vancomycin	I	>10	≤0.5 - >10	57.1/57.1				
MRSA (67)					Beta-hemolytic stre	eptococci ((24) ^c						
Oritavancin	0.03	0.06	0.008 - 0.25	97.0/97.0	Oritavancin	0.03	0.12	≤0.008 – 0.25	100.0/100.0				
Ceftaroline	1	2	0.25 – 2	80.6/80.6	Ceftaroline	≤0.015	≤0.015	≤0.015 – 0.03	100.0/100.0				
Clindamycin	≤0.25	>2	≤0.25 – >2	60.6/60.6	Clindamycin	≤0.25	>2	≤0.25 - >2	75.0/75.0				
Daptomycin	0.25	0.5	0.25 – 1	100.0/100.0	Daptomycin	0.12	0.25	≤0.06 – 0.5	100.0/100.0				
Erythromycin	>2	>2	≤0.25 ->2	20.9/20.9	Erythromycin	≤0.25	>2	≤0.25 – >2	70.8/70.8				
Levofloxacin	>4	>4	≤0.5 – >4	31.3/31.3	Levofloxacin	≤0.5	1	≤0.5 – 2	100.0/100.0				
Linezolid	1	2	0.25 – 2	100.0/100.0	Linezolid	1	1	0.5 – 1	100.0/100.0				
TMX	≤0.5	≤0.5	≤0.5-2	100.0/100.0	Penicillin	≤0.06	≤0.06	≤0.06	100.0/100.0				
Vancomycin	1	1	0.5 - 2	100.0/100.0	Teicoplanin	≤2	≤2	≤2	- /100.0				
Teicopianin	≤Z	≤Z	≤2 – 4	100.0/98.5	Tetracycline	8	>8	≤2 – >8	43.5/38.1				
CoNS (47) ^b					Vancomycin	0.25	0.5	0.25 – 0.5	100.0/100.0				
Oritavancin	0.03	0.06	≤0.008 - 0.25	_/_	Viridans group stre	ntococci (39)d						
Ceftaroline	0.25	0.5	≤0.06 – 2	- / -	Oritavancin	0.015	0.12		100 0/100 0				
Clindamycin	≤0.25	>2	≤0.25 - >2	78.7/78.7	Coffaralina	<0.015	0.12	$\leq 0.000 - 0.23$	/				
Daptomycin	0.25	0.5	≤0.12 – 0.5	100.0/100.0	Clindomycin	≤0.015 <0.25	<0.12	$\leq 0.013 - 2$	-/-				
Erythromycin	>2	>2	≤0.25 ->2	46.8/46.8	Doptomycin	≤0.25 0.25	<u>≤</u> 0.25	$\leq 0.23 - 32$	92.3/92.3				
Levofloxacin	2	>4	≤0.5 - >4	46.8/46.8	Enythromycin	0.25	0.0	$\leq 0.00 - 1$	100.07 -				
	0.5	1	≤0.12 - 8	97.9/97.9	Lovoflovooin	۲ ۲	2	=0.25 - 2	41.0/-				
	2	>2	≤0.25 - >2	25.5/25.5			<u>ک</u>	0.25 - 2	100.0 / -				
Vanaamvain	<u></u> <u></u>	8	$\leq 2 - 16$	97.9/85.1		0.10	1 0	0.25 - 1	71.9/76.0				
vancomycin	1	Z	0.25 – 2	100.0/100.0	Teicoplanin	0.1∠ ≤2	∠ ≤2	≤0.06 - >8 ≤2	- / 100 0				
E. faecalis (81)					Tetracycline	<2	>8	<2 - >8	79.5/-				
Oritavancin	0.015	0.03	≤0.008 – 0.5	97.5/-	Vancomycin	0.5	0.5	0.25 - 1	100 0/100 0				
Ampicillin	≤1	2	≤1 – 4	100.0/100.0	vanoonnyonn	0.0	0.0	0.20	100.0/100.0				
Ceftaroline	2	8	0.12 ->8	- / -	MRSA, methicillin-resistant S. aure	us; MSSA, methic	illin-susceptible S	. aureus; CoNS, coagula	se-negative staphylo				
Daptomycin	1	2	0.25 – 2	100.0 /	^a Criteria as published by CLSI (2017) and EUCAST (2017).Oritavancin breakpoint for vancomycin-susceptible <i>E. fae</i> applied to all isolates, including 2 vancomycin-resistant isolates. All vancomycin-susceptible <i>E. faecalis</i> were								
Erythromycin	2	>2	≤0.25 ->2	13.7 / —	susceptible to oritavancin.								
Levofloxacin	2	>4	≤0.5 – >4	70.4/72.8	^b Organisms include: Staphylococcus capitis (2), S. epidermidis (28), S. haemolyticus (3), S. hominis (3), S. lugdunensis (4), S. pasteuri (1), unspeciated coagulase-negative staphylococci (6)								
Linezolid	1	2	0.5 – 2	100.0/100.0	° Organisms include: Streptococcus agalactiae (11), S. dysgalactiae (1), S. equisimilis (2), S. pyogenes (10)								
Teicoplanin	≤2	≤2	≤2 – >16	97.5/97.5	^a Organisms include: Streptococcus anginosus (3), S. bovis group (1), S. gallolyticus (4), S. gordonii (2), S. mitis grou (6), S. mitis/oralis (4), S. mutans (1), S. oralis (5), S. parasanguinis (2), S. salivarius group (1), S. sanguinis (8)								
Vancomycin	1	2	≤0.5 – >16	96.3/96.3	S. sinensis (1), S. vestibularis (1)	,, (o), o .	,	,,					



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100.0 /100.0 75.0 /100.0 70.8 /100.0 /100.0)/100.0 0.00 /38.1 /100.0

/100.0 _

/92.3 0/-/ — 0 / – 0/-/76.9 0.00 / —

/100.0 staphylococci ible *E. faecalis*

mitis group

Conclusions

- Oritavancin showed potent in vitro activity against isolates recovered from patients with IE in US and EU sites
- Oritavancin coverage against this gram-positive collection was comparable to that of other antigram-positive agents, including daptomycin, linezolid, teicoplanin, and vancomycin
- Oritavancin, daptomycin, and linezolid demonstrated activity against VRE, including VanA-type organisms
- Results presented here provide invaluable information on the antimicrobial resistance profile of gram-positive isolates that cause infections in patients with IE and indicate that oritavancin exhibits potent in vitro activity against these organisms, including MRSA and VRE, isolated from IE patients hospitalized in EU and US medical centers

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