

Activity of Oritavancin Against *S. aureus*, *S. epidermidis*, *Enterococcus* spp. and *S. pneumoniae* Isolated from Canadian Hospital: Results of CANWARD 2007-2009

G.G. ZHANEL, K. NICHOL, P. LAGACÉ-WIENS, M. DeCORBY, R. VASHISHT, J.A. KARLOWSKY and D.J. HOBAN

University of Manitoba and Health Sciences Centre, Winnipeg, Manitoba, Canada

REVISED ABSTRACT

Background: Oritavancin (ORI) is a new lipopeptide antimicrobial agent. We assessed the activity of ORI and comparators against Gram-positive pathogens causing infections in Canadian hospitals.

Methods: From January 2007 through November 2009, 15 Canadian hospitals (10 centres - 2008) submitted pathogens from hospitalized patients. A total of 6,433 Gram-positive pathogens were collected and 4,773 tested to activity. Susceptibility testing used CLSI M7-A7 and M100-S15 broth microdilution which, for ORI, includes 0.002% turbidity test to date were:

Results: MIC₅₀ and MIC₉₀ values for ORI, vancomycin (VAN) and linezolid (LZO) isolates tested to date were:

Organism (n isolates)	MIC ₅₀ ORI		VAN		LZO	
	S	R	MIC ₅₀ /MIC ₉₀	MIC ₅₀ /MIC ₉₀	MIC ₅₀ /MIC ₉₀	MIC ₅₀ /MIC ₉₀
SPN-RI (902)	<0.002/0.004	<0.25/0.25	1/1			
- PenR (74)	<0.002/0.004	<0.25/0.25	0.5/1			
- PenI (120A)	<0.002/0.008	<0.25/0.5	0.5/1			
- PenI (84)	0.004/0.008	<0.25/0.5	0.5/1			
MSSA (1979)	0.12/0.25	1/1	2/2			
MSSA (631)	0.12/0.25	1/1	2/2			
- CA-MRSA (173)	0.12/0.25	1/1	2/2			
- HA-MRSA (433)	0.12/0.25	1/1	2/2			
MSE (202)	0.12/0.25	1/1	2/2			
MRSE (34)	0.03/0.05	2/2	1/1			
VS-E faecalis (308)	0.12/0.25	1/2	2/2			
VS-E faecium (124)	0.03/0.12	0.5/4	2/2			
Enterococcus spp. (371)	0.06/0.25	1/2	2/2			
VRE (28)	0.12/0.25	>32/32	2/2			
S. pneumoniae (200)	0.06/0.25	0.5/0.5	1/1			

SPN, *S. pneumoniae*; PenS, penicillin-susceptible; PenI, penicillin-intermediate; PenR, penicillin-resistant; HA, healthcare-associated; CA, community-associated; SA, *S. aureus*; MS, methicillin-susceptible; MR, methicillin-resistant; SE, *S. epidermidis*; VS, VAN-susceptible; VRE, VAM-resistant enterococci.

By MIC₅₀, oritavancin was more active than vancomycin and linezolid versus MSSA, MRSA (CA and HA), MSE, MRSE, SPN and Enterococcus spp. including VRE. Methicillin susceptibility did not affect ORI activity against staphylococci. ORI activity was not adversely impacted by penicillin susceptibility in SPN or by vancomycin susceptibility in enterococci. **Conclusions:** In the face of increasing incidence of antimicrobial resistance, ORI displays potential utility as its activity against gram-positive pathogens is unaltered by common resistance mechanisms. That CA- and HA-MRSA are equally susceptible to ORI may provide healthcare providers with another treatment option as the epidemiology of MRSA infections continues to evolve.

BACKGROUND

Infections caused by antibiotic resistant pathogens is a Canadian and global crisis. Antibiotic resistant pathogens including methicillin-resistant *Staphylococcus aureus* (MRSA), community-associated-CA and healthcare-associated-HA, vancomycin-resistant *Enterococcus* species (VRE) and penicillin-resistant *Streptococcus pneumoniae* (PRSP) are increasing in prevalence in Canada and around the world. Available therapeutic options for the treatment of these antibiotic resistant organisms are severely limited as these organisms frequently display a multidrug resistant (MDR) phenotype.

Oritavancin (ORI) is a new semisynthetic-lipopeptide antimicrobial agent which is very active in vitro against a range of gram positive pathogens, including both susceptible and multidrug-resistant staphylococci and enterococci.¹ It was originally discovered and developed by scientists at Eli Lilly as a potential replacement for vancomycin.

PURPOSE

The purpose of this study was to compare the activity of oritavancin and several comparators against gram-positive pathogens causing infections in patients affiliated with Canadian hospitals.

MATERIALS & METHODS

Bacterial Isolates

From Jan 2007-Dec 2009, inclusive, sentinel hospitals in major population centres in 8 of the 10 provinces in Canada were recruited. These sites were geographically distributed in a population based fashion. Each study site was asked to collect and submit pathogens (consecutive isolates, one organism per specimen source per patient) from patients using the following criteria: (a) 200 isolates from patients with respiratory tract infections (community or nosocomial); (b) 50 isolates from patients with skin/skin structure infections (wound/IV site infections); (c) 100 isolates from patients with urinary tract infections; (d) 20 blood stream infection isolates/month. All submitted isolates were deemed clinically significant by each site. All organisms were identified at each site using local site criteria and at the reference site, where indicated. At study sites, isolates were subcultured on appropriate solid media and incubated overnight. Isolates were shipped on Amies semi-solid transport media to the coordinating laboratory (Health Sciences Centre, Winnipeg, Canada), where isolates were subcultured on appropriate media and stocked in skim milk at -80°C.

Antimicrobial Susceptibilities

A custom microtitre panel was designed with a variety of antimicrobials. Antimicrobials were obtained as laboratory-grade powders from their respective manufacturers. Stock solutions were prepared and dilutions made as described by the Clinical and Laboratory Standards Institute (2009 CLSI).² Following two subcultures from frozen stock, the MICs were determined by the CLSI-approved broth microdilution method.² Briefly, 96-well custom-designed microtitre plates containing doubling antibiotic dilutions in 100µL/well of cation adjusted Mueller-Hinton broth (eg, for Enterobacteriaceae) with or without lysed horse blood (2.5-5% v/v) (eg, for *S. pneumoniae*) were inoculated to achieve a final concentration of approximately 5 x 10⁶ CFU/mL and incubated in ambient air (35°C) for 20-24 hours prior to reading. Polysorbate-80 (V025H) was added to both the stock solution of oritavancin and cation-adjusted broth used for panels. Quality control was performed periodically using a variety of ATCC QC organisms including: *S. pneumoniae* 49619, *S. aureus* 29212, *E. faecalis* 29212, *E. coli* 25922 and *P. aeruginosa* 27853. For all antimicrobials tested, MIC interpretive standards were defined according to CLSI breakpoints.² The following interpretive breakpoints (FDA) were used for teicoplanin susceptible (S), intermediate (I) and resistant (R) (µg/mL): *Staphylococcus aureus* (MSSA and MRSA) ≤ 0.5 (S); Enterococcus spp., ≤ 0.25 (S); Enterobacteriaceae, ≤ 2 (S), 4 (I), and ≥ 8 (R).

ACKNOWLEDGMENTS

This study was supported in part by the Medicines Company.

The authors would like to thank the investigators and laboratory site staff at each medical centre that participated in the Canadian Ward Surveillance Study (CANWARD) study:

Dr. D. Roscoe - Vancouver Hospital, Vancouver	Dr. M. Desjardins - The Ottawa Hospital, Ottawa
Dr. R. Remie - University of Alberta Hospital, Edmonton	Dr. M. Lavendyère - Hôpital Maisonneuve-Rosemont, Montreal
Dr. J. Blondeau - Royal University Hospital, Saskatoon	Dr. V. Loo - Montreal General Hospital, Montreal
Dr. S. Hoban-G. Zhanel - Health Sciences Centre, Winnipeg	Dr. V. Loo - Royal Victoria Hospital, Montreal
Dr. Z. Hussain - London Health Sciences Centre, London	Dr. M. Goyette - CHRTR Pavilion Site, Marie, Trois-Rivières
Dr. S. Poulakian - Mount Sinai Hospital, Toronto	Dr. M. Kuhn - South East Regional Health Authority, Moncton
Dr. L. Makris - St. Michael's Hospital, Toronto	Dr. R. Davidson - Queen Elizabeth II HSC, Halifax
Dr. F. Chan - Children's Hospital of Eastern Ontario, Ottawa	

RESULTS

Table 1. Activity of oritavancin and comparators against gram-positive cocci from CANWARD 2007-2009

Organism (n), Antibiotic	% of Isolates per Category			MIC ₅₀	MIC ₉₀	Range	Range Max
	S	I	R				
MSSA (1979)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	32	32
Clindamycin	74.9	0.3	24.7	< 0.25	> 16	> 0.25	> 32
Daptomycin	89.6	0.6	7.7	< 0.25	> 0.25	< 0.12	> 8
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
MSSA (631)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	89.6	0.1	10.3	< 0.12	> 0.008	1	1
Daptomycin	98.8	0.2	0.25	< 0.25	> 0.008	1	1
Linezolid	99.3	0.7	< 0.12	< 0.12	> 0.12	> 8	> 8
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	2	2
MSSA (173)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	11.7	0.1	88.2	> 16	> 128	> 128	> 128
Daptomycin	40.0	0.1	59.9	> 8	> 8	> 8	> 8
Linezolid	100.0	0.0	0.0	< 0.12	> 0.06	0.5	0.5
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
MSSA (433)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	44.0	0.1	55.9	> 8	> 8	> 8	> 8
Daptomycin	100.0	0.0	0.0	< 0.12	> 0.06	0.5	0.5
Linezolid	14.1	0.0	85.9	> 32	> 32	> 32	> 32
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
MSE (202)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
MRSE (34)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
VS-E faecalis (308)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
VS-E faecium (124)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Enterococcus spp. (371)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
VRE (28)							
Clarithromycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Clindamycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Daptomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
Linezolid	100.0	0.0	0.0	< 1	< 0.05	1	1
Vancomycin	100.0	0.0	0.0	< 1	< 0.05	1	1
S. pneumoniae (200)							
Clarithromycin	99.4	0.4	0.2	< 0.08	0.12	< 0.08	4
Clindamycin	81.3	4.3	14.9	< 0.03	> 0.03	< 0.03	> 32
Daptomycin	93.5	0.8	5.9	< 0.12	< 0.12	< 0.12	> 8
Daptomycin	93.2	2.8	4.0	< 0.25	> 1	< 0.08	> 16
Linezolid	99.1	0.3	0.8	< 1	< 1	< 0.08	32
Linezolid	100.0	0.0	0.0	< 1	< 0.12	> 2	> 2
Vancomycin	99.4	0.5	1.1	< 0.05	< 0.08	< 0.08	2
Oritavancin	100.0	0.0	0.0	< 0.004	0.004	< 0.002	0.12
Penicillin	99.1	14.5	4.4	< 0.01	< 0.02	< 0.01	> 8
Teicoplanin	98.8	0.6	1.5	< 0.003	< 0.003	0.5	0.5
TMP/SMX	100.0	0.0	0.0	< 0.1	< 0.12	> 8	> 8
Vancomycin	100.0	0.0	0.0	< 0.25	> 0.25	> 2	> 2
CA-MRSA (173)							
Clarithromycin	99.9	3.0	3.0	0.5	1	< 0.25	4
Clindamycin	4.3	1.8	85.9	> 0.25	> 16	> 0.03	> 0.03
Daptomycin	75.8	12.1	11.1	< 0.12	> 0.12	> 0.12	> 8
Daptomycin	75.8	12.1	11.1	< 0.12	> 0.12	> 0.12	> 8
Linezolid	100.0	0.0	0.0	< 1	< 0.12	> 2	> 2
Linezolid	100.0	0.0	0.0	< 1	< 0.12	> 2	> 2
Vancomycin	45.9	48.5	6.1	0.5	1	< 0.08	2
Oritavancin	100.0	0.0	0.0	< 0.004	0.004	< 0.002	0.12
Penicillin	100.0	0.0	0.0	< 0.02	> 0.02	> 0.02	> 8
Teicoplanin	100.0	0.0	0.0	< 0.01	< 0.01	< 0.01	0.12
TMP/SMX	30.3	6.1	63.6	4	8	< 0.12	> 8
Vancomycin	100.0	0.0	0.0	< 0.25	> 0.25	> 2	> 2

¹ Based on clinical trial data.

S, susceptible; I, intermediate; R, resistant.
No BP, no breakpoints available.
CA-MRSA, community-associated methicillin-resistant *S. aureus*.
HA-MRSA, healthcare-associated methicillin-resistant *S. aureus*.
Pen-Taz, piperacillin-tazobactam combination.
TMP/SMX, trimethoprim-sulfamethoxazole.

Table 2. MIC distribution of oritavancin against gram-positive cocci from CANWARD 2007-2009

Organism (no. tested)	Number (Cumulative percentage) at each MIC					
	<=0.004	0.008	0.015	0.03	0.12	0.25
Methicillin-susceptible <i>S. aureus</i> (1979)	107(5.4)	427(21.7)	339(14.1			