



## *In vitro* evaluation of meropenem-vaborbactam against clinical CRE isolates at a tertiary care center with low KPC-mediated carbapenem resistance

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### ARTICLE INFO

#### Article history:

Received 6 August 2018

Received in revised form 26 September 2018

Accepted 30 September 2018

Available online 4 October 2018

#### Keywords:

Meropenem-vaborbactam

CRE

KPC

Antimicrobial stewardship

### ABSTRACT

The *in vitro* activity of meropenem-vaborbactam was examined against clinical carbapenem-resistant *Enterobacteriaceae* isolates collected over 3 years at our medical center. Only 3 KPC-producers were identified. Susceptibility to meropenem-vaborbactam was noted in 15/16 (94%) isolates (MIC<sub>90</sub> 2 mg/L) that were nonsusceptible to meropenem. Meropenem-vaborbactam may have utility at centers where non-KPC-producers are more frequent.

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Carbapenem-resistant *Enterobacteriaceae* (CRE) are a significant global threat with substantially poorer outcomes and limited treatment options (Bassetti et al., 2016; Doi et al., 2017; Duin and Doi, 2017; Logan and Weinstein, 2017). Meropenem-vaborbactam was FDA-approved in 2017 for use in adult patients with complicated urinary tract infections (Drugs@FDA). Vaborbactam is a cyclic boronic acid derivative with potent activity against Ambler class A and class C  $\beta$ -lactamases (Lomovskaya et al., 2017). Its major utility has been demonstrable potency *in vitro* and efficacy in clinical trials against Ambler class A carbapenemase-producing *Enterobacteriaceae*, specifically isolates with KPC-mediated resistance (Castanheira et al., 2016, 2017; Hackel et al., 2018; Kaye et al., 2017; Lomovskaya et al., 2017; Pfaller et al., 2018). While these results were particularly encouraging, there is still considerable variation in KPC prevalence between medical centers even in similar geographical locations. At the University of Wisconsin Hospitals and Clinics, we have had approximately 1 KPC-producing clinical isolate per year (0.3 per 100 000 patient days and 0.007% of all *Enterobacteriaceae* isolates processed annually at our clinical laboratory). In contrast, non-KPC-producing CRE isolates are much more common, numbering 15–20 per year. Given the very low incidence of KPC-producing isolates,

we sought to answer whether meropenem-vaborbactam could have potential benefit in our patient population by evaluating its *in vitro* potency against CRE clinical isolates from our institution.

Retrospective review of the clinical microbiology database at the University of Wisconsin Hospitals and Clinics from 1/1/2015 to 12/30/2017 was performed. Over this period, it was standard practice to test and report ertapenem and meropenem susceptibility results on all *Enterobacteriaceae* isolates. A clinical isolate had to be resistant by CLSI interpretive criteria to at least 1 carbapenem to be included in analysis. Forty-six CRE isolates were identified from unique clinical encounters; 44 were able to be recovered from frozen stocks and included in the analysis. These isolates included *E. cloacae* complex ( $n = 19$ ), *E. coli* ( $n = 10$ ), *K. pneumoniae* ( $n = 8$ ), *S. marcescens* ( $n = 5$ ), *Klebsiella* (formerly *Enterobacter*) *aerogenes* ( $n = 1$ ), and *K. oxytoca* ( $n = 1$ ). Isolates were most commonly recovered from urine ( $n = 21$ ), followed by intraabdominal source ( $n = 9$ ), lung ( $n = 6$ ), blood ( $n = 3$ ), skin and soft tissue ( $n = 3$ ), bone ( $n = 1$ ), and sinus ( $n = 1$ ). Antimicrobial susceptibility testing performed by CLSI broth microdilution techniques (MicroScan, Beckman Coulter, Indianapolis, IN) at the time of isolation (CLSI, 2018a, 2018b) from the clinical specimen revealed 2 distinct groups of isolates: 1) resistant to ertapenem but susceptible to meropenem ( $n = 28$ , 64%) and 2) resistant to ertapenem and nonsusceptible (intermediate  $n = 2$ , resistant  $n = 14$ ) to meropenem

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**Table 1***In vitro* susceptibility of clinical CRE isolates to meropenem-vaborbactam<sup>a</sup> and select comparator antibiotics (University of Wisconsin Hospitals and Clinics, 2015–2017).

CRE phenotype	N (%)	Meropenem-vaborbactam MIC <sub>50</sub> (mg/L)	Meropenem-vaborbactam MIC <sub>90</sub> (mg/L)	Number (%) susceptible to meropenem-vaborbactam <sup>b</sup>	Number (%) susceptible to ciprofloxacin <sup>c</sup>	Number (%) susceptible to piperacillin/tazobactam <sup>c</sup>	Number (%) susceptible to tobramycin <sup>c</sup>
Ertapenem resistant, meropenem susceptible <sup>d</sup>	28 (64)	0.125	0.5	28 (100)	18 (64)	5 (18)	23 (82)
Ertapenem resistant, meropenem nonsusceptible <sup>d</sup>	16 (36)	0.125	2	15 (94)	8 (50)	7 (44)	10 (63)

<sup>a</sup> Meropenem-vaborbactam MIC testing is performed with fixed vaborbactam concentration of 8 mg/L per manufacturer instructions.<sup>b</sup> 2018 US FDA MIC interpretative criteria for meropenem-vaborbactam as follows: susceptible  $\leq 4/8$  mg/L, intermediate 8/8 mg/L, and resistant  $\geq 16/8$  mg/L.<sup>c</sup> Approved CLSI breakpoints at the time of testing were used for determining susceptibility to ciprofloxacin, piperacillin/tazobactam, and tobramycin.<sup>d</sup> Ertapenem resistance was determined based on CLSI breakpoints (MIC  $\leq 0.5$  mg/L). Meropenem nonsusceptible includes intermediate ( $n = 2$ ) and resistant ( $n = 14$ ) organisms based on CLSI interpretive criteria. MIC<sub>50</sub> and MIC<sub>90</sub> were not able to be calculated for meropenem alone as the data included truncated MICs for those isolates that were  $\leq 1$  mg/L.

(total  $n = 16$ , 36%). Each isolate was sent to the Wisconsin State Laboratory of Hygiene for molecular detection of carbapenemase genes for KPC, NDM-1, and OXA-48-like *via* real-time PCR. Only 3 were positive, and all were KPC-producing *K. pneumoniae* isolates.

Each isolate recovered from stock underwent additional antimicrobial susceptibility testing against meropenem-vaborbactam by antimicrobial gradient diffusion (Liofilchem Inc., Waltham, MA) and disk diffusion (Mast Group Ltd., Bootle, UK) per manufacturer instructions with a fixed vaborbactam concentration of 8 mg/L (CLSI, 2015, 2018a). Quality control testing was performed with ATCC reference strains using CLSI performance guidelines, and all values were in range (CLSI, 2018b). Breakpoints established by the FDA and included in the package insert were used for interpretation (The Medicines Company, 2017). As expected, the combination of meropenem-vaborbactam against isolates that were resistant to ertapenem but susceptible to meropenem was highly efficacious with all 28 isolates demonstrating susceptibility to the combination with an MIC<sub>50</sub> of 0.125 mg/L and MIC<sub>90</sub> of 0.5 mg/L (Table 1). In the group resistant to ertapenem and nonsusceptible to meropenem, meropenem-vaborbactam continued to demonstrate efficacy with an MIC<sub>50</sub> of 0.125 mg/L and MIC<sub>90</sub> of 2 mg/L (Table 1). Susceptibility to meropenem-vaborbactam was noted in 15 of 16 (94%) isolates in this group (Table 2). Notably, 11 of 16 isolates had

a  $\geq 4$ -fold decrease and 9 of 16 had a  $\geq 16$ -fold decrease in MIC for meropenem-vaborbactam compared to meropenem alone. On average, the meropenem-vaborbactam MIC decreased by almost 128-fold compared with meropenem alone. There were no discrepancies between susceptibility interpretation using gradient diffusion and disk diffusion.

This study demonstrates a number of important findings. First, we found that a majority (64%) of the CRE isolates at our center are resistant to ertapenem but are susceptible to other carbapenems such as meropenem. This has been reported as due to the combination of ESBL or AmpC  $\beta$ -lactamase production with contribution of active efflux pumps and/or porin loss (Woodford et al., 2007), and other carbapenems usually maintain activity against these isolates. Secondly, for CRE isolates that were resistant to ertapenem and nonsusceptible to meropenem, non-KPC-producers are much more frequently encountered at our center than KPC-producing isolates (81% vs. 19%). Nonetheless, we found that vaborbactam significantly decreased the meropenem MIC in most isolates compared to meropenem alone for this group. Previously published large *in vitro* studies have demonstrated a relative decrease in potency for meropenem-vaborbactam against non-KPC-producing CRE (Castanheira et al., 2017; Lomovskaya et al., 2017; Pfaller et al., 2018). For example, Pfaller et al. (2018)

**Table 2***In vitro* susceptibility results for meropenem-vaborbactam<sup>a</sup> by gradient diffusion and disk diffusion for CRE clinical isolates that were resistant to ertapenem and nonsusceptible to meropenem (University of Wisconsin Hospitals and Clinics, 2015–2017).

Organism	Meropenem MIC (mg/L) <sup>b</sup>	Meropenem-vaborbactam MIC (mg/L) <sup>c</sup>	Meropenem-vaborbactam inhibition zone diameter (nearest whole mm) <sup>c</sup>	Carbapenemase detected <sup>d</sup>
<i>Enterobacter cloacae</i> complex	2	2	21	
<i>Enterobacter cloacae</i> complex	2	0.5	28	
<i>Enterobacter cloacae</i> complex	4	2	20	
<i>Escherichia coli</i>	8	8	14	
<i>Escherichia coli</i>	>8	0.03	27	
<i>Escherichia coli</i>	4	0.06	30	
<i>Escherichia coli</i>	4	2	18	
<i>Klebsiella</i> (formerly <i>Enterobacter</i> ) <i>aerogenes</i>	4	4	17	
<i>Klebsiella oxytoca</i>	4	0.03	29	
<i>Klebsiella pneumoniae</i>	>8	0.125	20	KPC
<i>Klebsiella pneumoniae</i>	8	0.06	23	KPC
<i>Klebsiella pneumoniae</i>	>8	0.125	21	KPC
<i>Serratia marcescens</i>	>8	0.06	28	
<i>Serratia marcescens</i>	4	0.06	26	
<i>Serratia marcescens</i>	4	0.125	27	
<i>Serratia marcescens</i>	>8	0.06	29	

<sup>a</sup> Meropenem-vaborbactam MIC testing was performed with fixed vaborbactam concentration of 8 mg/L per manufacturer instructions.<sup>b</sup> 2018 CLSI M100 interpretative criteria for meropenem as follows: susceptible  $\leq 1$  mg/L, intermediate 2 mg/L, and resistant  $\geq 4$  mg/L.<sup>c</sup> 2018 US FDA MIC interpretative criteria for meropenem-vaborbactam as follows: susceptible  $\leq 4/8$  mg/L or  $\geq 17$  mm, intermediate 8/8 mg/L or 14–16 mm, and resistant  $\geq 16/8$  mg/L or  $\leq 13$  mm.<sup>d</sup> Isolates were tested for KPC, NDM-1, and OXA-48-like carbapenemases *via* real-time PCR.

recently published meropenem-vaborbactam activity against 330 CRE isolates from a worldwide collection during 2015. In this comprehensive study, meropenem-vaborbactam demonstrated an MIC<sub>50</sub> and MIC<sub>90</sub> against non-KPC-producers of 16 mg/L and 32 mg/L, respectively, due to the large number of MBL (52/121; 43%) and OXA-48 (40/121; 33%) producing isolates in this subset. Against the 16 isolates that were resistant to ertapenem and nonsusceptible to meropenem in our study, the MIC<sub>50</sub> was 0.125 mg/L and MIC<sub>90</sub> only 2 mg/L. Thus, we noted relatively preserved effectiveness despite the vast majority of our isolates in this group being non-KPC-producers. We believe this retained activity noted in our study can be explained by a relatively higher comparative incidence of other class A carbapenemase resistance mechanisms and lack of NDM and OXA-48-like isolates. The data presented in Table 2 demonstrate that most of the enhanced potency in our non-KPC-producing CRE was noted in 4 *S. marcescens* isolates, 2 *E. coli*, and a single *K. oxytoca* isolate. The *S. marcescens* isolates mimicked phenotypically a classic SME pattern, a class A carbapenemase. Based on the dramatic decrease in MIC in the presence of vaborbactam, it is possible that the *E. coli* and *K. oxytoca* isolates may have other less common class A carbapenemase genes present. Unfortunately, we were not able to assess for other serine Ambler class A carbapenemases (e.g., SME, IMI, GES, NMC, etc.) in these isolates. In general, these are considered relatively uncommon at many health care centers; however, our data suggest that these mechanisms may be as common as or more common than KPC-mediated resistance at some centers.

In conclusion, our study demonstrates that meropenem-vaborbactam activity is highly dependent on local epidemiology, may have retained *in vitro* potency against non-KPC-producers, and could be an important treatment option for patients with CRE at our institution despite a very low incidence of KPC-mediated resistance. Our *in vitro* evaluation serves as a template for other institutions to assess the activity of novel antimicrobial agents in the context of the local epidemiological distribution of resistance mechanisms circulating at their centers to best inform clinicians, guide antimicrobial stewardship efforts, and ensure their patients have access to optimized therapies.

## Acknowledgments

Meropenem-vaborbactam gradient test strips and disk diffusion test disks were provided by The Medicines Company (Parsippany, NJ).

## Funding

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

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