Frequency of Carbapenemase-Encoding Genes among Imipenem-Resistant Gram-Negative Bacilli Isolated from Latin America: Is there a Role for Imipenem/Relebactam? Results from SMART 2017-2018

INTRODUCTION

Gram-negative bacterial (GNB) infections have become increasingly difficult to treat due to raising antimicrobial resistance rates in certain geographic regions such as Latin America^{1,2}. Imipenem/relebactam (IMI/REL) is a novel beta-lactam/beta-lactamase inhibitor combination that was recently developed for clinical use to overcome the emergence and carbapenemase-producing spread Of Enterobacterales³. IMI/REL was recently approved in the USA but it is not available in Latin America yet. The Study for Monitoring Antimicrobial Resistance Trends (SMART) evaluates the in vitro activity of various GNB against selected antimicrobial agents since 2002 and includes IMI/REL since 2015.

The objective of this analysis was to gain insight of the possible therapeutic role of IMI/REL in this region, evaluating the activity of IMI/REL and comparator agents against the four most frequent GNB isolates and the frequency of carbapenemase-encoding genes (CEG) among imipenem-resistant GNB and PSA isolated from Latin America through the SMART Program from 2017-2018.

METHODS

13,843 GNB isolates including Enterobacterales (ENT) and *P. aeruginosa* (PSA) were collected from 36 medical sites of 10 Latin American countries during 2017-2018. Each site collected up to the indicated number of isolates in 2017/2018 from lower respiratory tract infections (RTI) (100/100), complicated urinary infections (cUTI) (75/50), complicated intra-abdominal infections (cIAI) (75/50) and blood (50 only in 2018).

Samples were consecutive and non-duplicate. MICs were determined using CLSI broth microdilution and also interpreted by CLSI criteria, except for IMI/REL, where United States Food and Drug Administration (US FDA) breakpoints were applied.

A subset of IMI-resistant isolates were selected for characterization of carbapenemase encoding genes via multiplex PCR and DNA sequencing. β-lactamase genes codifying ESBL, carbapenemases, and plasmidmediated AmpC were investigated in ENT and PSA.



*76 samples of unknown body sites were counted in the total sum





Figure 2. Most frequent CEG detected among carbapenemase producing, IMI-resistant Enterobacterales (Panel A), and MBL among *P. aeruginosa* (Panel B) isolates in Latin American countries

Table 1. Antimicrobial susceptibility to IMI/REL and comparators by enterobacterales and *P. aeruginosa*^a

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to IMI/REL

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ogen (n) Nicrobial Agonts	MIC _{50/90}	S(%)	l(%)	R(%)
erichia coli (4877)				
acin	<4/8	98.5%	0.8%	0.7%
zidime	<1/><1/>	72.3%	5.4%	22.3%
in	<1/<1	99.1%	0.0%	0.9%
enem	<0 12/0 12	97.9%	0.5%	1.6%
nem	<0.5/<0.5	98.5%	0.4%	1.0%
nem/relebactam ^b	0 12/0 25	99.6%	0.1%	0.3%
nenem	0 12/0 12	99.0%	0.2%	0.8%
acillin Tazobactam	≤2/16	90.1%	5.8%	4.1%
siella pneumoniae (2718)				
acin	≤4/16	93.0%	4.0%	3.0%
zidime	8/>32	48.3%	4.8%	46.9%
in	≤1/≤1	94.5%	0.0%	5.5%
enem	0.06/>4	77.4%	0.9%	21.7%
nem	≤0.5/>8	80.9%	1.2%	17.9%
nem/relebactam ^b	0 25/0 5	96.5%	0.8%	2 7%
benem	0 12/>8	79.8%	2.0%	18.2%
acillin Tazobactam	16/>64	58.7%	12.7%	28.6%
domonas aeruginosa (2108)				
acin	≤4/>32	84.9%	3.6%	11.5%
zidime	4/>32	71.6%	5.2%	23.2%
in	≤1/≤1	99.8%	0.0%	0.2%
nem	2/32	63.2%	5.1%	31.7%
nem/relebactam ^b	0.5/8	82.8%	5.5%	11.7%
benem	1/>16	65.2%	6.0%	28.8%
acillin Tazobactam	8/>64	67.9%	13.9%	18.2%
obacter cloacae (578)				
acin	≤4/8	94.5%	2.4%	3.1%
zidime	1/>32	62.1%	1.7%	36.2%
in	≤1/≤1	93.6%	0.0%	6.4%
enem	0.06/2	82.0%	6.4%	11.6%
nem	≤0.5/1	91.0%	2.6%	6.4%
nem/relebactam ^b	0.25/0.5	96.7%	0.3%	3.0%
benem	0.12/0.25	93.1%	1.7%	5.2%
acillin Tazobactam	≤4/>64	72.3%	9.2%	18.5%

^aSusceptibility values ≥90% shaded in blue

^bUnited States Food and Drug Administration (US FDA) breakpoints for imipenem/cilastatin/relebactam were applied

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RESULTS SUMMARY

E. coli was the most frequent species isolated from all body infections, except for RTI infections, where P. aeruginosa was the most frequent species. *K. pneumoniae* ranked as the second most frequent pathogen in all body sites infections (Figure 1).

• Among the 422 imipenem-resistant ENT further selected for detection of carbapenemase encoding genes, 394 (93.4%) isolates were detected as carbapenemase producers (Figure 2).

• bla_{KPC-2} and bla_{KPC-3} were the most common CEG found in ENT in all countries except in Guatemala and Mexico. *bla*OXA-48 and *bla*OXA-163 were only detected in Guatemala and Argentina, respectively. bla_{NDM-1} was identified in ENT isolated from all countries except Argentina, Chile and Puerto Rico (Figure 2A)

• A total of 415 imipenem-resistant P. aeruginosa were selected for characterization of carbapenemase content. The presence of metallo-betalactamase (MBL) encoding gene was confirmed in only 92 (22.2%) of these isolates (Figure 2).

bla_{VIM-2} was the most common MBL encoding gene identified, except for Ecuador, Guatemala, and Puerto Rico. *bla*_{IMP} variants were observed in Brazil (*bla*_{IMP-1}, *bla*_{IMP-74}), Colombia (a new *bla*_{IMP-variant}), Mexico (*bla*_{IMP-18} and bla_{IMP-75}, bla_{IMP-83}), and Panama (bla_{IMP-18}). As already expected, bla_{SPM-1} was only encountered in Brazil (Figure 2B).

CONCLUSION

CEG were frequently detected among imipenem-resistant ENT. In contrast, CEG was less frequently detected in imipenem-resistant P. aeruginosa.

 bla_{KPC-2} and bla_{VIM-2} were the most frequent CEG detected in ENT and P. aeruginosa isolates collected from Latin American medical centers. However, important local variations were observed in ENT and P. aeruginosa isolates with some CEG being detected in specific countries. For example, *bla*_{SPM-1} was identified only in Brazilian *P. aeruginosa* isolates, while bla_{OXA-48} and its derivatives only found in ENT isolated from Argentina and Guatemala.

IMI/REL showed relevant in vitro activity against the most frequent ENT and P. aeruginosa isolates and will represent an important addition to the therapeutic options to treat GNB infections in Latin America.

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A Comparison of Treatment Outcomes Between Analysis Populations in the RESTORE-IMI 1 Phase 3 Trial of Imipenem/Cilastatin/Relebactam Versus Colistin Plus Imipenem/Cilastatin in Patients With Imipenem-Nonsusceptible Bacterial Infections Keith S. Kaye, Helen W. Boucher, Michelle L. Brown, Angela Aggrey, Ireen Khan, Hee-Koung Joeng, Robert W. Tipping, Jiejun Du, Katherine Young, Joan R. Butterton, Amanda Paschke Antimicrobial Agents and Chemotherapy Feb 2020, AAC.02203-19; DOI: 10.1128/AAC.02203-19



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