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Emergence of IMP and VIM Metallo-ß-Lactamase Producing P. aeruginosa Strains in Mexico: Report from the SENTRY Antimicrobial Surveillance Program



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ABSTRACT

Background: Acquired carbapenem (CARB) resistance (R) due to metallo-ß-lactamase (MBL) production is increasing rapidly in Asia, Europe and South America, but has been rarely reported in North America and particularly in Mexico. We report the first discovery of IMP and VIM among P. aeruginosa (PSA) clinical strains isolated in Mexican medical

Methods: As part of the SENTRY Program, all isolates are tested for susceptibility (S) by CLSI broth microdilution methods to >30 antimicrobials. Strains R to imipenem (IMP), meropenem and ceftazidime are screened for MBL production by a disk approximation (DA) test. Strains with a positive MBL screen test (enzyme inhibited by EDTA and 2-MPA) were evaluated by PCR for blaimp, blavim, blaspm and blagim, followed by gene sequencing. Results: 28 PSA strains were collected from a Mexican center (115) in 2004, 4 (14.3%) were R to IMP and 3 of those organisms (75%) showed positive MBL DA test results. All strains had positive PCR results for *bla_{IMP}* and gene sequence analysis demonstrated greatest homology with IMP-5. The strains showed the same ribotype/PFGE pattern and were isolated in April, July and October, 2004 from different patients with long-term hospitalization and exposure to multiple antimicrobial treatments. A fourth strain with a positive MßL screen test was isolated in January 2005. This strain had a positive PCR result for blavim, and negative results for blaimp and blaspm. All 4 PSA strains were R to all antimicrobials tested except the polymyxins.

Conclusions: This is the first report of MBL-producing strains (IMP and VIM) in Mexico, and MßL production appears to be the main mechanism of CARB-R among PSA in the index medical center. These results raised the concern of the dissemination of this potent MBL mechanism in this institution, as well as, in the region or nation.

INTRODUCTION

Metallo-beta-lactamase (MBL) production is an emerging mechanism of carbapenem resistance among enteric and non-fermentative Gram-negative bacilli. MBLs of the IMPfamily were first reported from Japan and are now prevalent in Asia, Europe and Latin America. Additional families of these enzymes namely VIM (Italy), SPM (Brazil), GIM (Germany) and more recently SIM (Korea) have emerged over time. Certain MßL types, such as SPM have become widely endemic in the geographic regions where they were first identified while others have spread across and between continents (e.g. VIM).

Inspite of increasing incidence of MBLs in various parts of the world, reports of MBLproducing pathogens from North America are a recent event and still relatively rare. There are four reports of VIM-producing *Pseudomonas aeruginosa* (VIM-7 [Texas, two reports] and VIM-2 [Texas and Illinois]) from the USA and one report of an outbreak of IMP-7 producing *P. aeruginosa* from Canada. However, there are no reports of strains producing acquired MBL from Mexico.

In this report, we characterized MBL-producing strains isolated in a Mexican medical center through the SENTRY Antimicrobial Surveillance Program.

MATERIALS AND METHODS

Bacterial isolates. In 2004, a medical center in Mexico submitted 28 non-duplicate P. aeruginosa isolates to the SENTRY Antimicrobial Surveillance Program. These strains were isolated from bloodstream infections, skin and skin structure infections and pneumonia in hospitalized patients according to protocols used by all SENTRY Program participant centers. Species level identification was confirmed by standard biochemical tests and Vitek cards where necessary.

Susceptibility testing. The isolates were susceptibility tested (minimum inhibitory concentration [MIC]) against >30 antimicrobials by broth microdilution procedure as described by the Clinical and Laboratory Standards Institute (CLSI, formerly the NCCLS) using dry-form panels manufactured by Trek Diagnostics (Cleveland, OH, USA). Interpretations of susceptibility test results were as per CLSI criteria, where available. Escherichia coli ATCC 25922 and P. aeruginosa ATCC 27853 were routinely tested in parallel with clinical isolates for quality assurance.

Screening for metallo-ß-lactamases. P. aeruginosa isolates resistant to imipenem, meropenem and ceftazidime (MIC >16 µg/ml), were tested for MBL production. Disk approximation screen tests for MBL detection were performed using imipenem, meropenem and ceftazidime as substrates and EDTA as well as 2-mercaptopropionic acid (MPA) as inhibitors. MßL Etest strips with imipenem ± EDTA (AB BIODISK, Solna, Sweden) were used to confirm positive disk approximation test results when needed. Isolates with a positive phenotypic MßL test were screened for *bla*_{IMP}, *bla*_{VIM} and *bla*_{SPM} using PCR primers described elsewhere. Primers derived from conserved sequences of the class 1 integron were used to study genetic context of the MBL genes.

Gene sequencing. PCR reactions were examined for amplification product by electrophoresis on 1% agarose gels. Amplicons cleaned using QIAquick PCR purification kit (QIAGEN GmbH, Germany) were sequenced using Sanger based dideoxy sequencing strategy involving the incorporation of fluorescent-dye-labeled terminators into the sequencing reaction products. Sequences obtained were compared to the available sequences via NCBI BLAST search.

Molecular Typing. Multiple isolates showing a similar antimicrobial susceptibility pattern and a positive disk approximation test for MBL were epidemiology typed using the RiboprinterTM Microbial Characterization system. Isolates with identical ribotypes were further characterized by pulsed-field gel electrophoresis (PFGE).

RESULTS

- Three of the 28 (10.7%) P. aeruginosa isolates submitted by the Mexican medical center in 2004 were resistant to most antimicrobials tested, including B-lactam-B-lactamase inhibitor combinations, cephalosporins, carbapenems, quinolones and aminoglycosides, but were susceptible to polymyxin B (Table 1).
- All three strains showed positive results for the MßL disk approximation test as well as MßL Etest (Table 2).
- PCR screens for MBLs showed amplification products with blamp primers from all three isolates. Sequences obtained from PCR amplicon of index strain (1686D) showed closest similarity with the blaimp-5 sequence.
- IMP-5 shares greatest amino acid identity with IMP-1 and IMP-4 (94.0%) and also with IMP-8 (89.8%). Comparison with these three IMP-types led to the identification of bla_{IMP} from 1686D as IMP-5 (Figure 1).
- A class I integron with an aminoglycoside modifying enzyme gene AadA6 was identified in 1686D, but the *bla*IMP-5 was not present in this integron.

- Case reports of patients infected with IMP-5-producing strains.
- Case 1: A 63 year-old Mexican male patient, with no previous hospitalizations and poorly controlled type II diabetes mellitus, was admitted for the treatment of a diabetic foot infection. During the initial evaluation renal failure and chronic obstructive pulmonary disease were diagnosed. Antibiotic treatment for a severe mixed bacterial foot infection, that included MRSA, was initiated with ciprofloxacin and clindamycin and was followed with piperacillin/tazobactam, linezolid, gentamicin, trimethoprim/sulfamethoxazole (TMP/SMX), rifampin and levofloxacin. After 60 days of hospitalization an IMP-5-producing *Pseudomonas* aeruginosa was isolated from the foot wound. P. aeruginosa infection was eradicated only after extensive débridement.
- Case 2: A 22 year-old Mexican male was admitted for the evaluation of acute abdomen. During an exploratory laparatomy a perforated appendix with diffuse peritonitis was found. The patient was started on cefotaxime plus clindamycin and after several days on mechanical ventilation he developed a nosocomial pneumonia. On July 12, 2004 an IMP-5-producing P. aeruginosa was isolated from sputum. Antimicrobial therapy was switched to meropenem, amikacin, and ciprofloxacin; sputum cultures subsequently became negative for P. aeruginosa.
- Case 3: A 54 year-old Mexican female with myelocytic leukemia under chemotherapy was admitted for the evaluation of fever. Initial empiric antimicrobial treatment included amoxicillin/clavulanic acid, TMP/SMX, and ciprofloxacin. The patient developed renal failure, C. difficile diarrhea and urinary tract infection. An IMP-5 producing *P. aeruginosa* was isolated from the blood on October 26, 2004. Meropenem was then started but after 3 days the patient's clinical condition deteriorated and she expired.

	MIC (μg/ml)				
Antimicrobial agent	1686D	2058C	10639A		
lmipenem ^a	>256	48	48		
Meropenem	>8	>8	>8		
Ceftazidime	>16	>16	>16		
Cefepime	>16	>16	>16		
Piperacillin/tazobactam	>128	128	128		
Aztreonam	>16	16	16		
Ciprofloxacin	>4	>4	>4		
Levofloxacin	>4	>4	>4		
Gentamicin	>8	>8	>8		
Amikacin	>32	>32	>32		
Tobramycin	>16	>16	>16		
Polymyxin B	0.5	0.5	0.5		

- All three isolates producing IMP-5 belonged to the same ribogroup and showed similar (<3 bands difference) PFGE patterns. Temporal separation of isolation dates of the three strains (April, July and October, 2004) indicates endemic nature of this cluster at the medical center (Table 1).
- A fourth strain (1112A) with a multidrug-resistance pattern (susceptible only to polymyxin B) was isolated in early 2005. It exhibited a positive MßL disk approximation test with imipenem, meropenem and ceftazidime as substrates and EDTA as enzyme inhibitor. MßL Etest showed marginally positive results (Table 2). PCR screens using blavim primers generated a PCR product, but presence of *bla*_{VIM} could not be confirmed.

Figure 1. Comparison of amino acid sequences of various IMP-types IMP-1: MSKLSVFFIFLFCSIATAAESLPDLKIEKLDEGVYVHTSFEEVNGWGVVPKH IMP-4: MSKLSVFFIFLFCSIATAAEPLPDLKIEKLDEGVYVHTSFEEVNGWGVVPKH IMP-5: MSKLFVFFMFLFCSITAAAESLPDLKIEKLDEGVYVHTSFEEVNGWGVVPKH IMP-8: MKKLFVLCVCFLCSITAAGAALPDLKIEKLEEGVYVHTSFEEVNGWGVVSKH GLVVLVNAEAYLIDTPFTAKDTEKLVTWFVERGYKIKGSISSHFHSDSTGGIEWLNSR GLVVLVDAEAYLIDTPFTAKDTEKLVTWFVERGYKIKGSISSHFHSDSTGGIEWLNSQ GLVVLVNTEAYLIDTPFTAKDTEKLVTWFVERGYKIKGSISSHFHSDSTGGIEWLNSQ GLVVLVNTDAYLIDTPFTATDTEKLVNWFVERGYKIKGTISSHFHSDSTGGIEWLNSQ SIPTYASELTNELLKKDGKVQATNSFSGVNYWLVKNKIEVFYPGPGHTPDNVVVWLPE SIPTYASELTNELLKKDGKVQAKNSFGGVNYWLVKNKIEVFYPGPGHTPDNLVVWLPE SIPTYASELTNELLKKDGKVQAKNSFSGASYWLVKKKIEVFYPGPGHTPDNVVVWLPE SIPTYASELTNELLKKDGKVQAKNSFSGVSYWLVKNKIEVFYPGPGHTQDNVVVWLPE RKILFGGCFIKPYGLGNLGDANIEAWPKSAKLLKSKYGKAKLVVPSHSEVGDASLLKL RKILFGGCFIKPYGLGNLGDANLEAWPKSAKLLISKYGKAKLVVPSHSEAGDASLLKL NRVLFGGCFVKPYGLGNLGDANVEAWPKSAKLLMSKYGKAKLVVPSHSEVGDASLLKR KKILFGGCFVKPDGLGNLGDANLEAWPKSAKILMSKYGKAKLVVSSHSEIGDASLLKR TLEOAVKGLNESKKPSKPSN TLEQAVKGLNESKKPSKLSN TLEQAVKGLNESKKPSKPSN TWEQAVKGLNESKKPSQPSN Green highlites: Different amino acid in IMP-5 compared to IMP-1 and/or -4 Pink highlites: Different amino acid in IMP-5 compared to IMP-8 Blue highlites: Different amino acid in IMP-5 compared to IMP-1, -4 and -8

CONCLUSIONS

- This study constitutes the first report of clonal dissemination of MßL-producing P. aeruginosa isolates in a Mexican medical
- This is also the first report of a IMP-5 producing *P. aeruginosa*. The only other report of IMP-5 was from an Acinetobacter baumannii strain isolated in Portugal.
- Continued surveillance and careful testing of multidrug-resistant P. aeruginosa and other common Gram-negative pathogens is imperative to monitor the spread and the epidemiology of this potent resistance mechanism.

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Case #	MßL Etest (µg/ml)		MßL disk approximation							
	Isolate #	Imipenem	IMI+EDTA	IMI+EDTA	MER+EDTA	IMI+MPA	MER+MPA	CAZ+MPA	Ribo/PFGE	MßL
1	1686D	>256	8	_	-	+	_	+	258.259.7/A	IMP-5
2	2058C	48	8	_	_	_	_	+	258.259.7/A1	IMP-5
3	10639A	48	8	_	_	_	_	+	258.259.7/A	IMP-5
	1112A	32	4	+	+	_	_	+	ND^a	_