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Brief report

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ABSTRACT

Objectives. Our objective was to evaluate the *in vit-ro* activity of ceftolozane-tazobactam against multidrug resistant (MDR) and extensively drug-resistant (XDR) non metallo- β -lactamase producing *Pseudomonas aerugino-sa* clinical isolates at Hospital Universitario Miguel Servet (Zaragoza, Spain) from February 2016 to October 2017.

Material and methods. We evaluated the *in vitro* activity of ceftolozane-tazobactam and other antipseudomonal antibiotics against 12 MDR and 117 XDR non metallo- β -lactamase producing *P. aeruginosa* isolates. Ceftolozane-tazobactam minimal inhibitory concentrations (MICs) were determined by MIC gradient diffusion test strip.

Results. Among the 129 MDR/XDR isolates included, 119 (92.2%) were susceptible to ceftolozane-tazobactam, and ten (7.8%) were resistant. MIC_{50} was 2 mg/L, and MIC_{90} 4 mg/L. Ceftolozane-tazobactam was the second most active antibiotic after colistin, overtaking amikacin.

Conclusions. Ceftolozane-tazobactam is a valuable treatment option for MDR and XDR *P. aeruginosa* infections in our setting.

Keywords: *Pseudomonas aeruginosa*, multidrug resistance, extensively drug resistance, ceftolozane-tazobactam, beta-lactams

Actividad de ceftolozano-tazobactam en aislados clínicos de *Pseudomonas aeruginosa* multirresistente y extremadamente resistente en un hospital español

RESUMEN

Objetivos. Nuestro objetivo fue evaluar la sensibilidad in vitro de ceftolozano-tazobactam en aislados clínicos de *P. aeruginosa* multirresistente (MDR) y extremadamente resistente (XDR) desde Febrero de 2016 a Octubre de 2017 en el Hospital Universitario Miguel Servet, Zaragoza (España).

Material y métodos. Evaluamos la actividad *in vitro* de ceftolozano-tazobactam y otros antibióticos anti-pseudomónicos en 12 aislados de *P. aeruginosa* MDR y en 117 aislados XDR, no productores de metalo- β -lactamasas. Se determinó la concentración mínima inhibitoria (CMI) de ceftolozano-tazobactam mediante tiras de difusión en gradiente.

Resultados. Entre los 129 aislados MDR/XDR incluidos, 119 (92,2%) fueron sensibles a ceftolozano-tazobactam, y diez (7,8%) presentaron resistencia. La CMI₅₀ fue de 2 mg/L, y la CMI₉₀ de 4 mg/L. Ceftolozano-tazobactam fue el segundo antibiótico más activo después de colistina, superando a amikacina.

Conclusiones. Ceftolozano-tazobactam es una opción de tratamiento válida para infecciones causadas por *P. aeruginosa* MDR y XDR en nuestro entorno.

Palabras clave: *Pseudomonas aeruginosa*, multirresistente, extremadamente resistente, ceftolozano-tazobactam, beta-lactámicos

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Antimicrobial activity of ceftolozane-tazobactam against multidrug-resistant and extensively drug-resistant Pseudomonas aeruginosa clinical isolates from a Spanish hospital

INTRODUCTION

Pseudomonas aeruginosa is a ubiquitous non-fermentative Gram-negative bacterium and one of the leading causes of nosocomial infections. A point-prevalence study lead by the European Centre for Disease Prevention and Control in 2011-2012 found that almost 9% of hospital-acquired infections were caused by P. aeruginosa, whereas a Spanish EPINE survey in 2016 identified it as the second cause of nosocomial infections (after Escherichia coli), being present in 10.5% of isolates [1]. P. aeruginosa is intrinsically resistant to many antibiotics due to the high inducible expression of multidrug efflux pumps, low levels of non-specific porins, necessary for penetration of hydrophilic antibiotics, and due to the inducible expression of β -lactamases, such as AmpC [2]. *P. aerugi*nosa is also highly capable of developing acquired resistance through chromosomal mutations that can lead to overexpression of AmpC or efflux pumps or to downregulation or loss of OprD porin, one of the mechanisms responsible of imipenem and meropenem resistance. It can also acquire resistance by horizontal gene transfer: plasmid-borne expression of extended-spectrum β -lactamases (ESBL), aminoglycoside-modifying enzymes and 16S rRNA methylases have been described in P. aeruginosa [2].

The increasing spread of multidrug-resistant (MDR) and extensively-drug resistant (XDR) P. aeruginosa is a growing concern worldwide [3]. Recent epidemiological studies reported MDR prevalence in P. aeruginosa of around 15% and XDR-prevalence ranging between 2.6% and 9.6% from in- and outpatients in North America [4, 5]. In Spain, the prevalence of MDR strains is already above 30% [3, 6].

Ceftolozane-tazobactam is a combination of a novel cephalosporin with the β -lactamase inhibitor tazobactam, approved for the treatment of complicated intra-abdominal infections, complicated urinary tract infections, and currently under investigation for the treatment of ventilator-associated pneumonia [7]. Unlike the previous-generation cephalosporins, ceftolozane-tazobactam has demonstrated increased stability to AmpC β -lactamases; it is also unaffected by the loss of expression of OprD porin and by the overexpression of multidrug efflux pumps [7]. The activity of ceftolozane is compromised in the presence of carbapenemases, such as metallo- β -lactamases (MBL) [7]. Ceftolozane-tazobactam constitutes a valuable treatment option for MDR gram-negative pathogens and is currently the cephalosporin with the highest activity against P. aeruginosa [3, 8].

The objective of the present study was to evaluate the in vitro activity of ceftolozane-tazobactam against MDR and XDR non MBL producing P. aeruginosa clinical isolates, in order to assess its suitability as treatment in infections caused by MDR and XDR P. aeruginosa in our setting.

MATERIALS AND METHODS

Samples were obtained from in- and outpatients at Hos-

pital Universitario Miguel Servet, Zaragoza (Spain), between February 2016 and October 2017. Bacterial identification was performed by MALDI-TOF mass spectrometry (Bruker Daltonics) and antimicrobial susceptibility testing by MicroScan WalkAway (Beckman Coulter). The following antibiotics were routinely tested: ceftazidime, cefepime, piperacillin-tazobactam, aztreonam, imipenem, meropenem, ciprofloxacin, levofloxacin, gentamicin, tobramycin, amikacin, and colistin. Current EUCAST breakpoints [9] were applied to define resistance/ susceptibility.

Ceftolozane-tazobactam minimal inhibitory concentrations (MICs) were determined by MIC gradient diffusion test strip (Liofilchem). MDR isolates were defined as those non-sus-

Table 1	Characteristics of the <i>P. aeruginosa</i> isolates.	129 MDR/XDR
Characteristics		n (%)
Type of sample		
Respiratory		59 (45.8)
Urine		51 (39.5)
Surgical wound		9 (7.0)
Blood		2 (1.6%)
Catheter		1 (0.8%)
Others		7 (5.4)
Origin of patient		
In-patient		99 (76.8%)
Out-patient		30 (23.2)
MDR		12 (9.3)
XDR		117 (90.7)
Isolates susceptible to:		
Amikacin		101 (78.2)
Gentamicin		31 (24.0)
Tobramycin		54 (41.9)
Ceftazidime		13 (10.1)
Cefepime		14 (10.8)
Piperacillin-tazobactam		8 (6.2)
Ceftolozane-tazobactam		119 (92.2)
Imipenem		16 (12.4)
Meropenem		7 (5.4)
Aztreonam		0 (0)
Ciprofloxacin		10 (7.0)
Levofloxacin		5 (3.9)
Colistin		128 (99.2)
Isolates only susceptible to colistin and amikacin		49 (38)
Isolates only susceptible to colistin		13 (10.1)



ceptible to \geq 1 agent in at least 3 antibiotic classes; XDR isolates were defined as those only susceptible to \geq 1 agent in maximum 2 antibiotic classes [10]. MBL producing isolates were excluded from the analysis since ceftolozane-tazobactam is not active against them. MBL production was screened by phenotypic methods: combined disk test with imipenem 10 µg (Oxoid) and imipenem 10 µg with 750 µg of EDTA (Sigma-Aldrich) in-house added and by Neo-Rapid CARB Kit (Rosco); and confirmed by real-time PCR (Check Direct CPE Kit Check-Points Health B.V.).

RESULTS

During the period studied, a total of 1,421 *P. aerugino-sa* non-duplicate isolates were obtained; of these, 141 (9.9%) were defined as MDR and 316 (22.2%) as XDR. A total of 68 (4.7%) isolates were MBL producing.

In a total of 129 MDR/XDR *P. aeruginosa* non-duplicate isolates (one isolate per patient) and non-MBL producing ceftolozane-tazobactam could be tested and were included in the study. Of the 129 isolates, 12 (9.3%) were defined as MDR and 117 (90.7%) as XDR. One hundred-and-eight (83.7%) patients were male, mean age (SD) was 63.9 (18.0) years; most were in-patients (76.8%). The majority of isolates were from respiratory samples (45.8%) and urine samples (39.5%). The rates of susceptibility to the antibiotics included in the panel and other characteristics are summarized in table 1. Most isolates were classified as XDR, with a median of two active antibiotics out of 12 tested (range: 0-8). All except one isolate were susceptible to colistin. Other antibiotics more frequently active against *P. aeruginosa* isolates were amikacin (78.2%)

and tobramycin (41.9%). A few isolates were susceptible to imipenem (12.4%), meropenem (5.4%), ceftazidime (10.1%), cefepime (10.8%) and piperacillin-tazobactam (6.2%). Only 34 (26.3%) isolates were susceptible to all the β -lactams tested. Thirteen (10.1%) isolates were only susceptible to colistin, and 49 (38%) were only susceptible to colistin and amikacin.

When susceptibility to ceftolozane-tazobactam was analyzed, 119 isolates (92.2%) were susceptible and 10 (7.8%) isolates showed MIC above the susceptibility breakpoint (4 mg/L); MIC₅₀ was 2 mg/L, and MIC₉₀ was 4 mg/L (figure 1). Among the 10 resistant isolates, five were susceptible to amikacin, and all were susceptible to colistin. Nine isolates were non-susceptible to the other β -lactams tested, and only one isolate was susceptible to imipenem and meropenem. Seven of the ten resistant isolates came from respiratory specimens.

DISCUSSION

Although β -lactams are the mainstream therapy for *P. aeruginosa* infections, resistances to these drugs are increasingly common [1,3,4,5]. Thus, oprD porin mutation-mediated suppression causes resistance to carbapenems; overexpression of β -lactamase AmpC inactivates penicillins, cephalosporins and monobactams, whereas efflux pump up-regulation causes resistance to various β -lactams, fluoroquinolones and aminoglycosides. Plasmid-encoded ESBLs and MBLs are emergent and worrisome mechanisms of resistance in *P. aeruginosa* [1].

The combination of the fifth-generation cephalosporin ceftolozane, which is less affected by loss of OprD, overexpression of efflux pumps or AmpC than other available β -lactams, joined to tazobactam, which inhibits a wide variety of beta-lactamases (but not AmpC or MBLs), covers most of the pathogens causing β -lactam resistance [7]. Indeed, in the present study, only 7.8% of MRD/XDR non-MBL isolates were resistant to ceftolozane-tazobactam, whereas almost 75% of them were non-susceptible to the rest of the tested β -lactams. Two previous studies reported similar rates (from 4.8% to 7%) of non-MBL mediated resistance to ceftolozane-tazobactam in MDR and XDR *P. aeruginosa* [11,12].

In our study, we did not further explore the mechanisms of such resistance. When Moya et al. explored the mechanisms of pan- β -lactam resistance in *P. aeruginosa* (including ceftolozane), they found that all pan-resistant isolates lacked OprD, overexpressed AmpC and efflux pumps, and 5 out of 6 isolates had modifications in penicillin-binding protein (PBP) profiles [13]. An epidemiological study on a collection of 150 XDR isolates from Spanish hospitals reported an overall rate of non-susceptibility to ceftolozane-tazobactam of 31%, mediated by horizontally acquired carbapenamases, although potentially relevant PBP3 mutations were also detected [14]. Other studies identified AmpC overexpression and chromosomal mutations as potential mechanism of resistance to ceftolozane/ tazobactam [15-17]. In our study, the ceftolozane-tazobactam MIC for non-MBL producing isolates ranged between 8 and 16 mg/L, which is only 2-4 fold higher that the breakpoint. It is reasonable to suggest that a massive overexpression of AmpC and/or the mutations, such as target modification, could lead to the observed non-MBL-mediated resistance.

Nonetheless, the activity of ceftolozane-tazobactam against the tested MDR and XDR isolates was high (92.2%) and was only surpassed by colistin. We conclude that ceftolozane-tazobactam is a valuable treatment option for complicated infections caused by MDR and XDR non-MBL producing *P. aeruginosa* isolates in our setting.

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CONFLICT OF INTEREST

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