05586 Emergence of carbapenemase-producing *Klebsiella pneumoniae* clinical isolates with a pan-drug resistant (PDR) phenotype in Argentina

03. Bacterial susceptibility & resistance

3d. Resistance mechanisms incl. in vitro and in vivo studies, mobile elements (excl. TB) Likely attendance Onsite

Fernando Pasteran¹, Juan Manuel De Mendieta¹, Paola Ceriana¹, Laura A. Fernandez², Natalia Pujato³, Mariano Echegorry¹, Melina Rapoport¹, Ezequiel Albornoz¹, Celeste Lucero¹, Alejandra Menocal¹, Denise De Belder⁴, Sonia Gomez¹, Diego Faccone¹, Diego Corso¹

¹Antimicrobianos INEI ANLIS Malbrán - Buenos Aires (Argentina), ²Laboratorio Dr Rapela -Buenos Aires (Argentina), ³Instituto de Transplante de Alta Complejidad - Buenos Aires (Argentina), ⁴UOCNGB ANLIS Malbrán - Buenos Aires (Argentina)

Background

Carbapenemase producers represented the 33% of *K. pneumoniae* (Kpn) infections in Argentina 2021, being $bla_{\rm NDM}$ and $bla_{\rm KPC}$ responsible of 88% of those infections. NDM producers are usually susceptible to aztreonam plus ceftazidime/avibactam (AZA). We describe here the emergence of three closely related Kpn clinical isolates with a PDR phenotype, including high-level resistance to AZA.

Methods

For antimicrobial susceptibility testing we used: 1) agar dilution for AZA, ceftazidime/avibactam (CZA), imipenem/relebactam (IMR) (DBOs 4mg/L) and rifampin; 2) cefiderocol microdilution with iron depleted CAMHB; 3) colistin disks elution and 4) susceptibility to other antimicrobials, through automated systems. Synergistic activity of selected antimicrobials was assessed crossing gradient diffusion strips at a 90° angle at the intersection of each MIC. Synergy tests were carried-out on MHA plates, alone or supplemented with the indicated drug/DBOs. Relevant genes were characterized by two multiplex PCR (*bla*KPC/*bla*NDM/*bla*OXA-48-like/*bla*VIM/*bla*IMP and *bla*CTXM/*bla*PER/*bla*CIT). NGS was performed by Illumina MiSeq. MLST and alleles were confirmed by assemblies in silico and AMRFinder/ARIBA and/or by Sanger sequencing.

Results

Kpn1 to Kpn3 were recovered in a single Institution from urine samples of three respective patients undergoing kidney transplantation (male, 38-54 y.o). Strains were only susceptible to cefiderocol (Fig-1), which is not available in Argentina. Patients remained alive after compassionately treatment with a triple combination of aztreonam plus CZA plus clavulanic

acid (Fig-2) but the transplanted kidneys were rejected. Strains were confirmed as ST258, capsular type K107/O1/O2V2/wzi154. SNP analysis revealed a close evolutionary relationship between isolates: 59 SNP (Kpn1 vs Kpn2), 47 SNP (Kpn2 vs Kpn3) and 90 SNP (Kpn1 vs Kpn3). The 3 strains were confirmed as producers of bla_{NDM-5} , $bla_{CTXM-15}$, $bla_{SHV-5-like}$ (K234R, A237G), additionally, Kpn2 co-produced bla_{KPC-2} and bla_{OXA-1} . Main resistance mechanisms are depicted in Fig-3.

Conclusions

We described the emergence of Kpn isolates resistant to all antibiotics available in the country. PDR was multicausal, involving β -lactamases, modifying enzymes, efflux, porin deficiency and target mutations, including essential PBPs. The addition of two β -lactamase inhibitors was required to reduce aztreonam MICs to levels close to the PK/PD breakpoint for continuous infusion. The emergence of PDR strains should be considered of high epidemiological risk, maximizing efforts for rapid detection and containment.

Fig-1. Antimicrobial susceptibility profile

	Кр	n1	Кр	n 2	Kpn3				
	Date: 08	-28-2022	Date: 09	-01-2022	Date: 09	-15-2022			
	MIC	Cat.	MIC	Cat.	MIC	Cat.			
Aminopenillins, Piperacillin+tazobactam extended spectrum cephalosporins, cefoxitin	>64	R1	>64	R1	>64	R1			
Meropenem, imipenem, ertapenem	>32	R1	>32	R1	>32	R1			
Amikacin, gentamicin	>32	R1	>32	R1	>32	R1			
Ciprofloxacin, levofloxacin	>4	R1	>4	R1	>4	R1			
Minocycline	>8	R1	>8	R1	>8	R1			
Tigecycline	>2	R ²	>2	R ²	>2	R ²			
Fosfomycin	>64	R ²	>64	R ²	>64	R ²			
Trimethoprim sulfamethoxazole	>2	R1	>2	R1	>2	R1			
Rifampin	>4	R ³	>4	R ³	>4	R ³			
Nitrofurantoin	>64	R1	>64	R1	>64	R1			
Chloramphenicol	>16	R1	>16	R1	>16	R1			
CZA, IMR	>32	R1	>32	R1	>32	R1			
Colistin	>8	R ²	>8	R ²	>8	R ²			
Aztreonam	>256	R ^{1,2}	>256	R ^{1,2}	>256	R ^{1,2}			
AZA	32	R ⁴	>256	R ⁴	>256	R ⁴			
Cefiderocol ⁵	0,25	S ^{1,2}	0,25	S ^{1,2}	0,25	S ^{1,2}			

S: susceptible. R: resistant 1. CLSI breakpoint (M100, Ed 32-2022). 2 EUCAST breakpoint (v12, 2022) 3. Société Française de Microbiologie. 4. EUCAST breakpoint for aztreonam alone. 5. Cefiderocol is not available in Argentina

Fig.2 Evaluation of therapeutic options

	SELECTED ANTIMICRO	MICs values in mg/L.								
		-	I	Kpn1	Kpn2	Kpn3				
COMPOUND 1	COMPOUND 2	COMPOUND 3	COMPOUND 4							
AZTREONAM	AVIBACTAM 4ug/ml	-	-	32	>256	>256				
AZTREONAM	AVIBACTAM 4ug/ml	CLAVULANIC Ac. 4ug/ml	-	1	24	32				
AZTREONAM	AVIBACTAM 10ug/ml	CLAVULANIC Ac. 4ug/ml	-	1	24	32				
AZTREONAM	AVIBACTAM 4ug/ml	COLISTIN 1 ug/ml	-	24	>256	>256				
AZTREONAM	AVIBACTAM 4ug/ml	FOSFOMYCIN 10ug/ml	-	24	>256	>256				
AZTREONAM	AVIBACTAM 4ug/ml	RIFAMPIN 4ug/ml	-	32	>256	>256				
AZTREONAM	RELEBACTAM 4ug/ml	-	-	24	>256	>256				
AZTREONAM	RELEBACTAM 4ug/ml	CLAVULANIC Ac. 4ug/ml	-	1	12	12				
AZTREONAM	AVIBACTAM 4ug/m	RELEBACTAM 4ug/ml	-	24	>256	>256				
AZTREONAM	AVIBACTAM 4ug/m	RELEBACTAM 4ug/ml	CLAVULANIC Ac. 4ug/ml	0,047	12	12				
MEROPENEM	COLISTIN 1ug/ml	-	-	>32	>32	>32				
MEROPENEM	RIFAMPIN	-	-	>32	>32	-				
AZTREONAM	AVIBACTAM 4ug/m	CLAVULANIC Ac. 4ug/ml	COLISTIN	-	8	8				
AZTREONAM	AVIBACTAM 4ug/m	CLAVULANIC Ac. 4ug/ml	TIGECYCLINE	-	16	16				
		Most active combinat	ions are highlighted	in green						

Fig-3. Main resistant mechanisms identified by NSG

									Pr	eser	nce/a	absen	ce of	resis	tanc	e ge	nes a	and/	or ch	romo	soma	al m	utatio	ns								
		AMINOGLYCOSIDES β-LACTAMICS							TETPACYCLINES		PHENICOL	PHENICOL TRIMETHOPRIM SULFONA MIDES RIFAMICINAS				GUINOLONES		MACROUDES		FOSFOMYCIN	EFFLUX PUMPS		COLISTIN		(/ɛɬ/)		PBP2 (mrdA)					
	ST	aac(6')-1b	aac(3)-lid	aadA2	rmtB1	bla _{TEM-1}	blashv-s-uke	blaoxa-i	blacrxm-15	blakec-2	blanom-s	ompK36_D135DGD	rpsJ_V57L	tet(D)	catB3	dfrA12	sul1	arr-3	gyrA_S831	parC_S801	mph(A)	erm(B)	qac£∆1	fosA	oqxA	oqxB	pmrB_R256G	pmrB_T140P	V375A	A413V	D354A	T529N
Kpn1	258																															
Kpn2	258																															
Kpn3	258																															
											Р	resenc	eoft	neind	icate	d me	char	nism	is hig	nlight	ed in	red										

Keyword 1 Antibiotic stewardship (AMS) Keyword 2 Antimicrobial resistance (AMR) Keyword 3 pan-drug resistance

Conflicts of interest

Do you have any conflicts of interest to declare? No