



# Occurrence of multi-carbapenemases producers among carbapenemase-producing Enterobacterales and in vitro activity of combinations including cefiderocol, ceftazidime-avibactam, meropenem-vaborbactam, and aztreonam in the COVID-19 era

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## Abstract

**Purpose** To evaluate the prevalence of multi-carbapenemase-producing Enterobacterales (EB) and the activity of cefiderocol (CFDC), meropenem-vaborbactam (MEV), ceftazidime-avibactam (CZA), and combinations of CZA plus aztreonam (ATM), MEV plus ATM and CFDC plus CZA against them.

**Methods** A collection of carbapenemase-producing EB clinical isolates ( $n = 1242$ ) was investigated by lateral flow immunoassay NG-Test CARBA-5 and molecular testing. Cefiderocol MICs were determined using broth microdilution Sensititre™ panel. MICs of CZA and MEV were determined by the gradient diffusion method. Antimicrobial synergy testing was performed using gradient diffusion strip crossing.

**Results** KPC were the most frequent carbapenemases (83.2%), followed by VIM (9.2 %), OXA-48-like (4.3 %) and NDM enzymes (4.1%). Multi-carbapenemase producers were found in 10 (0.8%) isolates. Three combinations of two different carbapenemases were observed: KPC+VIM ( $n = 4$ ), NDM+OXA-48-like ( $n = 4$ ), and VIM+OXA-48-like ( $n = 2$ ). CFDC showed potent activity against eight out of ten dual-carbapenemases producers, while resistance or reduced susceptibility was shown towards CZA and MEV. CFDC in combination with CZA showed no synergistic effects and only two additive effects on seven (87.5%) of the CFDC-susceptible strains. Conversely, CZA plus ATM and MEV plus ATM combinations were synergistic against all ATM-resistant strains regardless of dual-carbapenemases phenotype.

**Conclusions** The occurrence of multi-carbapenemase producers is not uncommon in Northern Italy area. MEV in combination with ATM might be considered as a potential therapeutic option, alternative to CZA plus ATM. CFDC susceptibility testing and synergy evaluation of ATM-based combinations should be performed in the lab routine to evaluate the most in vitro active antimicrobial regimen.

**Keywords** Carbapenemase · Metallo- $\beta$ -lactamase · NG-Test CARBA-5 · Cefiderocol · Ceftazidime-avibactam · Meropenem-vaborbactam · COVID-19 infections

## Introduction

Carbapenem-resistant Enterobacterales (CRE) rank among the top three multi-drug-resistant pathogens on WHO's priority list [1]. The subset of CRE that produces carbapenemases, carbapenemase-producing Enterobacterales (CPE), is of high clinical and public health concern, because it has been associated with increasing mortality and high ability to spread in healthcare settings. Three major classes of carbapenemases are largely associated with the global spread of CRE: KPC (Ambler class A), metallo- $\beta$ -lactamases (M $\beta$ Ls)

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(Ambler class B, e.g., NDM, VIM, and IMP), and OXA-48-like (Ambler class D) carbapenemases [2].

Management and treatment of patients with infections due to CPE is a daily challenge in clinical practice since CPE strains co-harbor antimicrobial resistance determinants towards more classes of antimicrobials. Recently, several active drugs against CPE have been approved for clinical use, including  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations, such as ceftazidime-avibactam (CZA) and meropenem-vaborbactam (MEV), and the novel siderophore cephalosporin named cefiderocol (CFDC) [3, 4].

Knowledge of carbapenemase type is important to guide antibiotic therapy since not all  $\beta$ -lactamase inhibitor combinations have activity against all classes of enzymes. Indeed, both avibactam and vaborbactam are potent inhibitors of class A carbapenemases but are ineffective against M $\beta$ Ls-producing strains, while only avibactam shows inhibitory activity against OXA-48-like producers [5, 6]. CFDC showed broad activity against meropenem-non-susceptible Gram-negative bacilli, including carbapenem-resistant Enterobacterales (EB), *Pseudomonas aeruginosa* and *Acinetobacter baumannii*. Its broad activity is explained by its distinctive mechanism of penetrations via the iron transport system of Gram-negative bacteria that overcomes resistance mechanisms including efflux pump upregulation and porin channel loss. Moreover, the side-chain properties render high stability against hydrolysis by  $\beta$ -lactamases, including serine- $\beta$ -lactamases and M $\beta$ Ls [7].

The carriage of more than one type of carbapenemases, although not common in Europe [8], is a relevant challenge for antimicrobial treatment, especially when M $\beta$ Ls co-occur with KPC or OXA-48 like carbapenemases. Aztreonam (AZT) in combination with avibactam has shown activity against M $\beta$ L-producing EB, including strains co-harboring KPC or OXA-48 like carbapenemases [9–11]. However, evidence on the activity of new  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations and novel CFDC against multi-carbapenemases producers is limited. In this study, we evaluated the prevalence of multi-carbapenemases producers in a collection of clinical EB isolates collected from patients hospitalized during the COVID pandemic 2019–2021. Subsequently, the activity of CFDC, MEV, CZA and CZA+ATM, MEV+ATM, and CFDC+CZA combinations against multi-carbapenemases producers was evaluated.

## Material and methods

### Clinical isolates and carbapenemase detection

CPE clinical strains were retrospectively collected from the program of surveillance and control of healthcare-associated multi-drug resistant Gram-negative infections based at our

Microbiology Laboratory. The isolates were isolated from blood, urine, rectal swabs, lower respiratory tract specimens, and wound swabs of patients admitted at the reference Centre “Città della Salute e della Scienza di Torino” and other four hospitals in Piedmont, in North-western Italy, during COVID pandemic 2019–2021. Duplicate isolates of the same species from a single patient were excluded. Species identification was carried out by matrix-assisted laser desorption ionization-time of flight mass spectrometry (MALDI-TOF MS) (Bruker DALTONIK GmbH, Bremen, Germany). Carbapenemase production was investigated by lateral flow immunoassay NG-Test CARBA 5 (NG Biotech, Guipry, France) [12]. Multi-carbapenemase producers were confirmed by Xpert Carba-R molecular assay (Cepheid, Sunnyvale, CA, USA) and were also investigated for carriage of *bla*<sub>CTX-M-like</sub> genes (ESBL ELITE MGB Kits, ELITEchGroup Molecular Diagnostics, Turin, Italy).

### Antimicrobial susceptibility testing

Multi-carbapenemase-producing EB strains were subjected to antimicrobials susceptibility testing. Minimal inhibitory concentrations (MICs) of meropenem (MEM), amikacin, colistin, fosfomycin, cotrimoxazole, levofloxacin, and aztreonam were determined by a commercially available microdilution assay (Panel NMDR, Microscan WalkAway 96 Plus, Beckman Coulter, Switzerland) according to the manufacturer’s instructions. Cefiderocol susceptibility was determined using Sensititre™ panel (Thermo Scientific™) broth microdilution panels. MICs of CZA and MEV were determined by the gradient diffusion method (Liofilchem®, Roseto degli Abruzzi, Italy). All MIC values were interpreted according to the European Committee of Antimicrobial Susceptibility Testing (EUCAST) clinical Breakpoints (v.11 2021) (<https://eucast.org>).

### Evaluation of synergistic activity

Antimicrobial synergy testing was performed on multi-carbapenemase-producing clinical strains. The following combinations of antimicrobials were evaluated for synergistic activity: CZA+ATM, MEV+ATM, CFDC+CZA.

Synergy testing was carried out using gradient diffusion strip crossing, as previously described [13]. Briefly, the strip of drug A was placed perpendicularly at a 90° angle to strip off the drug B at their respective MICs onto cation-adjusted Mueller-Hinton agar plates using 0.5 MacFarland inoculum and incubated at 37 °C for 18–24 h. MICs were read at the point at which the elliptical inhibition area touched the strips. The mean fractional inhibitory concentration index (FICI) was calculated by dividing the mean MIC of each drug in combination with the MIC of each drug alone and adding the results. The FICI results were interpreted as

follows:  $\leq 0.5$  as synergy;  $> 0.5$  to  $\leq 4$  as no interaction; and  $> 4$  as antagonism.

## Results

Overall, 1242 non-duplicate CPE clinical strains were collected. Of these, 1034 (83.2%) produced KPC enzymes that were the most frequent carbapenemases. VIM, OXA-48-like, and NDM enzymes were produced by 114 (9.2%), 53 (4.3%), and 51 (4.1%) CPE strains, respectively. The most common CPE specie was *Klebsiella pneumoniae* (87.6%) that was seven times more frequent than other EB species. In *K. pneumoniae*, the KPC enzyme (93%) was the most prevalent carbapenemase, followed by NDM (3.4%), VIM (2.5%), and OXA-48-like (1.7%). Conversely, a more homogeneous distribution of carbapenemases in the other Enterobacterales species was observed (Table 1).

Multi-carbapenemase producers were found in 10 (0.8%) CPE isolates. Three combinations of two different carbapenemases were observed: KPC+VIM ( $n = 4$ ), NDM+OXA-48-like ( $n = 4$ ), and VIM+OXA-48-like ( $n = 2$ ). Concerning antimicrobial susceptibility, eight of them were resistant to both MEM and at least three other antimicrobials among amikacin, colistin, fosfomycin, cotrimoxazole, levofloxacin, and aztreonam (Table 2). The expression of a M $\beta$ L by all the dual-carbapenemase-producing strains also led to resistance or reduced susceptibility to

the new combinations of  $\beta$ -lactam/ $\beta$ -lactamases inhibitor. In detail, MEV showed no activity towards all strains except the KPU04 (*K. pneumoniae* KPC+VIM co-producer), PRW05 (*P. rettgeri* VIM+OXA-48-like co-producer), and KPB09 (*K. pneumoniae* NDM+OXA-48-like co-producer), whereas CZA achieved high MICs, from 24 to  $> 256$  mg/L, for all the strains tested.

CFDC showed potent activity against all dual-carbapenemase producers except against strains KPU04 and ECR10, which tested resistant with MIC of 16 mg/L.

Synergy testing results were shown in Table 3. All KPC+VIM and VIM+OXA-48-like co-producing strains tested negative for *bla*<sub>CTX-M-like</sub> genes, whereas the remaining four strains were NDM+OXA-48-like and CTX-M-like co-producers. As previously observed [14], CFDC MICs determined by gradient diffusion testing were lower than those obtained with reference broth microdilution method, leading to consider strain ECR10 as false susceptible to CFDC in comparison to broth microdilution.

CFDC in combination with CZA showed no synergistic effects against dual-carbapenemase producer strains tested. However, additive effects against seven (70%) of the tested strains were observed. Conversely, CZA+ATM and MEV+ATM combinations were synergistic against all ATM-resistant strains regardless of dual-carbapenemases phenotype. Of note, ATM reduced MIC values of both CZA and MEV below resistance breakpoint (8 mg/L) in two of ten (20%) and seven of seven (100%) of tested resistant strains, respectively.

**Table 1** Prevalence of Enterobacterales species and carbapenemase enzymes detected during the study period (2019–2021)

Prevalence	KPC carbapenemase (%)	OXA-48-like carbapenemase (%)	Metallo- $\beta$ -lactamase VIM (%)	Metallo- $\beta$ -lactamase NDM (%)	KPC+VIM (%)	NDM+OXA-48-like (%)	VIM+OXA-48-like (%)
<i>Klebsiella pneumoniae</i> ( $n = 1088$ )	1008 (92.6)	16 (1.5)	23 (2.1)	34 (3.1)	4 (0.4)	3 (0.3)	
<i>Escherichia coli</i> ( $n = 62$ )	14 (22.9)	26 (42.6)	10 (16.4)	11 (17.7)		1 (1.6)	
<i>Enterobacter cloacae</i> ( $n = 47$ )	4 (8.5)		42 (89.3)	1 (2.1)			
<i>Citrobacter freundii</i> ( $n = 16$ )			16 (100)				
<i>Providencia rettgeri</i> ( $n = 6$ )		4 (66.7)					2 (33.3)
<i>Klebsiella oxytoca</i> ( $n = 6$ )	3 (50)		3 (50)				
<i>Providencia stuartii</i> ( $n = 5$ )		1 (20)	4 (80)				
<i>Klebsiella aerogenes</i> ( $n = 4$ )			4 (100)				
<i>Morganella morganii</i> ( $n = 3$ )			3 (100)				
<i>Citrobacter farmeri</i> ( $n = 2$ )			2 (100)				
<i>Proteus mirabilis</i> ( $n = 1$ )				1 (100)			
<i>Citrobacter brakii</i> ( $n = 1$ )			1 (100)				
<i>Serratia marcescens</i> ( $n = 1$ )	1 (100)						
Total ( $n = 1242$ )	1030 (82.9)	47 (3.8)	108 (8.7)	47 (3.8)	4 (0.3)	4 (0.3)	2 (0.2)

**Table 2** Antimicrobial susceptibility of multi-carbapenemase-producing Enterobacterales isolates

Strain	Specie carbap-enemases	Sample	Meropenem MIC (mg/L)	Amikacin MIC (mg/L)	Colistin MIC (mg/L)	Fosfomicin MIC (mg/L)	Cotrimoxazole MIC (mg/L)	Levofloxacin MIC (mg/l)	Aztreonam MIC (mg/L)	Ceftazidime-avibactam MIC (mg/L)	Meropenem-vaborbactam MIC (mg/L)	Cefiderocol MIC (mg/L)
KPB01	<i>K. pneumoniae</i> KPC+VIM	Biopsy	> <b>32</b>	≤ 8	≤ 2	≤ 16	> 4	> 1	> 4	> <b>256</b>	<b>32</b>	0.25
KPU02	<i>K. pneumoniae</i> KPC+VIM	Urine	<b>32</b>	≤ 8	≤ 2	≤ 16	> 4	> 1	> 4	> <b>256</b>	<b>16</b>	0.5
KPB03	<i>K. pneumoniae</i> KPC+VIM	Blood	> <b>32</b>	> <b>16</b>	≤ 2	<b>64</b>	≤ 2	> 1	> 4	> <b>256</b>	<b>16</b>	0.25
KPU04	<i>K. pneumoniae</i> KPC+VIM	Urine	8	≤ 8	> 4	> <b>64</b>	> 4	≤ 0.5	> 4	> <b>256</b>	0.75	<b>16</b>
PRW05	<i>P. rettgeri</i> VIM+OXA-48-like	Wound exudate	<b>16</b>	≤ 8	> 4	≤ 16	> 4	> 1	2	> <b>256</b>	8	0.25
PRU06	<i>P. rettgeri</i> VIM+OXA-48-like	Urine	4	> <b>16</b>	> 4	> <b>64</b>	> 4	> 1	2	<b>64</b>	<b>12</b>	0.25
KPU07	<i>K. pneumoniae</i> NDM+OXA-48-like	Urine	<b>32</b>	> <b>16</b>	≤ 2	<b>32</b>	≤ 2	> 1	> 4	> <b>256</b>	<b>16</b>	0.5
KPR08	<i>K. pneumoniae</i> NDM+OXA-48-like	Rectal swab	> <b>32</b>	> <b>16</b>	> 4	≤ 16	> 4	> 1	> 4	> <b>256</b>	<b>24</b>	0.5
KPB09	<i>K. pneumoniae</i> NDM+OXA-48-like	Blood	> <b>32</b>	> <b>16</b>	> 4	> <b>64</b>	> 4	> 1	> 4	> <b>256</b>	6	2
ECR10	<i>E. coli</i> NDM+OXA-48-like	Rectal swab	<b>32</b>	≤ 8	≤ 2	≤ 16	> 4	> 1	> 4	> <b>256</b>	<b>16</b>	<b>16</b>

Bold characters indicate MICs in the range of resistance according to current EUCAST breakpoints (v.11 2021)

**Table 3** Synergy testing results of dual-carbapenemase-producing Enterobacterales isolates tested

Strain	Specie Carbapenemases and CTX-M content	Minimum inhibitory concentrations (µg/mL)										FICI			
		CFDC*	CZA	CFDC	ATM	MEV	CZA+CFDC	CZA+ATM	MEV+ATM	CZA+CFDC	CZA+ATM	MEV+ATM	CZA+CFDC	CZA+ATM	MEV+ATM
		CFDC sin	CZA sin	CFDC sin	ATM sin	MEV sin	CZA+CFDC sin	CZA sin	ATM sin	MEV sin	ATM sin	MEV sin	ATM sin	ATM sin	ATM sin
KPB01	<i>K. pneumoniae</i> KPC+VIM producer, CTX-M negative	0.25	> 256	0.125	> 256	32	128	0.064	12	48	4	64	1.01	0.23	0.37
KPU02	<i>K. pneumoniae</i> KPC+VIM producer, CTX-M negative	0.5	> 256	0.047	> 256	16	96	0.016	32	32	2	64	0.71	0.25	0.37
KPB03	<i>K. pneumoniae</i> KPC+VIM producer, CTX-M negative	0.25	> 256	0.016	> 256	16	96	0.006	8	32	1.5	64	0.75	0.16	0.34
KPU04	<i>K. pneumoniae</i> KPC+VIM producer, CTX-M negative	16	> 256	3	> 256	0.75	128	2	16	16	0.047	16	1.25	0.12	0.12
PRW05	<i>P. rettgeri</i> VIM+OXA-48-like producer, CTX-M negative	0.25	24	0.25	2	8	8	0.094	6	0.75	6	1.5	0.71	0.62	1.25
PRU06	<i>P. rettgeri</i> VIM+OXA-48-like producer, CTX-M negative	0.25	> 256	0.125	2	12	64	0.032	32	1	6	1	0.51	0.62	1
KPU07	<i>K. pneumoniae</i> NDM+OXA-48-like producer, CTX-M positive	0.5	> 256	0.064	> 256	16	128	0.016	32	32	1	16	0.75	0.25	0.12
KPR08	<i>K. pneumoniae</i> NDM+OXA-48-like producer, CTX-M positive	0.5	> 256	0.125	> 256	24	64	0.064	24	32	2	32	0.76	0.22	0.21
KPB09	<i>K. pneumoniae</i> NDM+OXA-48-like producer, CTX-M positive	2	> 256	0.047	64	6	128	0.016	16	3	0.75	8	0.84	0.11	0.25
ECR10	<i>E. coli</i> NDM+OXA-48-like producer, CTX-M positive	16	> 256	1	> 256	16	128	0.5	24	16	2	64	1	0.16	0.37

\*Cefiderocol MIC values determined by broth microdilution panel

Ceftazidime-avibactam, cefiderocol, aztreonam, and meropenem-vaborbactam MICs were determined by the gradient diffusion method

Bold characters indicate MICs in the range of resistance according to current EUCAST breakpoints (v.11 2021). FICI results were interpreted as follows: ≤ 0.5 as synergy; > 0.5 to ≤ 1 as additive effect; > 1 to 4 as indifferent; and > 4 as antagonism. Values in italics indicate a synergistic effect

Abbreviations: FICI, fractional inhibitory concentration index; CZA, ceftazidime-avibactam; CFDC, cefiderocol; ATM, aztreonam; sin: synergism

## Discussion

The emergence of multi-carbapenemases producing EB has clinical, laboratory testing, and public health implications. The new combinations of  $\beta$ -lactam/ $\beta$ -lactamase inhibitor have increased treatment options for organisms producing serine- $\beta$ -lactamases (Ambler class A and D), but the growth in the proportion of isolates that co-harbor M $\beta$ Ls jeopardizes their usefulness [15]. In this study, we analyzed the prevalence of multi-carbapenemases producers in a large collection of CPE strains collected in an area of Northern Italy during a 2-year period overlapping the COVID-19 pandemic. We also evaluated in vitro activity of the new siderophore cephalosporin CFDC and the synergy of CZA+ATM, CFDC+CZA, and MEV+ATM combinations against phenotypes co-expressing M $\beta$ Ls and serine-carbapenemases.

The highlights of this study are the following findings: (1) prevalence of dual-carbapenemases-producing EB was notable; (2) resistance to CFDC occurred among M $\beta$ L- and serine-carbapenemase-co-producing EB; (3) MEV+ATM showed synergistic activity similar to CZA+ATM against M $\beta$ L- and serine-carbapenemase-co-producers.

Despite KPC enzymes remain the most prevalent circulating carbapenemase type among CPE, increased frequencies of M $\beta$ Ls and OXA-48like enzymes in comparison to previous Italian reports have been observed [8, 16]. In our study, a notable prevalence of multi-carbapenemases-producing EB was found, and three dual-carbapenemase-producing phenotypes were observed: KPC+VIM, NDM+OXA-48-like, and VIM+OXA-48-like. Multi-carbapenemases detection may have important microbiological and therapeutic implications. Hence, co-carriage of serine-carbapenemases and M $\beta$ Ls explained the resistance towards CZA (10/10) and MEV (7/10). Eight strains (80%) showed resistance to ATM, related at least to the expression of serine- $\beta$ -lactamases (KPC or CTX-M). CFDC showed high efficacy since eight isolates (80%) tested susceptible with MICs ranging from 0.25 to 2 mg/L. The other two dual-carbapenemase-producing strains, that were KPC+VIM-producing *K. pneumoniae* and NDM+OXA-48-like producing *E. coli*, were resistant to CFDC with MICs of 16 mg/L. Although the number of strains tested was limited, CFDC plus CZA displayed no synergistic effect and additive effects on seven (87.5%) of the CFDC-susceptible dual-carbapenemase-producing strains.

Data have emerged supporting the activity of ATM in combination with avibactam against M $\beta$ L-producing EB, including strains co-harboring KPC or OXA-48 like carbapenemases [9–11]. In our synergy testing evaluation, we observed that CZA+ATM and MEV+ATM

were synergistic against all ATM-resistant dual-carbapenemases-producing strains tested, both KPC+VIM and NDM+OXA-48-like+CTX-M producing phenotypes including both CFDC-resistant strains. The synergy is explained by the inhibitory activity of avibactam or vaborbactam on serine- $\beta$ -lactamases (e.g., KPC and CTX-M enzymes) and refractoriness of ATM to M $\beta$ L hydrolysis [17]. Even if vaborbactam has no inhibitory activity on OXA-48-like enzymes, in vitro synergy between MEV and ATM against NDM+OXA-48-like+CTX-M producers could be explained by the abovementioned mechanism together with a probably poor carbapenemase activity of OXA-48-like enzymes.

Although both CZA+ATM and MEV+ATM combinations demonstrated synergy, important considerations regarding MIC values of CZA and MEV, tested alone or combined with ATM, should be done. CZA MICs were significantly higher compared to those of MEV for all dual-carbapenemase-producing strains tested (MICs ranged from 24 to > 256 mg/L and from 0.75 to 16 mg/L for CZA and MEV, respectively). The addition of ATM highly impacted on both MEV and CZA MIC values. However, MEV MIC values decreased below the resistance breakpoint more than CZA MICs.

Our study extends existing data on the synergy between CZA and ATM, and the limited data on the synergy between MEV and ATM against M $\beta$ L- and serine-carbapenemase-co-producing EB [18, 19]. Our findings suggest that CZA and MEV may be interchangeably combined with ATM for M $\beta$ Ls and serine- $\beta$ -lactamase co-producers. However, our data showed that although synergistic activities were observed for both antibiotics associations, clinically achievable concentrations were obtained more frequently for MEV in association with ATM than for CZA combined with ATM. Therefore, an association of MEV with ATM could represent an alternative to CZA plus ATM, especially in cases of selection of mutant strains resistant to avibactam. In fact, in vitro experiments showed that mutations in conserved amino acid residues of chromosomal and plasmid AmpC  $\beta$ -lactamase and other  $\beta$ -lactamases are associated with avibactam resistance in EB and *Pseudomonas aeruginosa* [20, 21].

Limitations of our study include the limited number of strains tested, the lack of a full characterization of  $\beta$ -lactamases genes content and cloning typing. Further studies including worldwide clinical isolates are necessary for the generalizability of our findings.

In conclusion, we showed that the occurrence of multi-carbapenemase producers is not uncommon in Northern Italy area. Carbapenemase detection methods able to differentiate the main carbapenemase types, such as immunochromatographic or molecular assays, are recommended to identify

multi-carbapenemase producers. Our findings suggest that MEV in combination with ATM might be considered as a potential therapeutic option, alternative to CZA plus ATM, for the treatment of infections caused by MβLs and serine-β-lactamase co-producing EB. CFDC susceptibility testing, and synergy evaluation of ATM-based combinations should be performed in lab routine to evaluate the most appropriate antimicrobial regimen according to clinical aspects.

**Data availability** The authors confirm that the data supporting the findings of this study are available within the article.

**Code availability** Not applicable

## Declarations

**Ethical approval** Not applicable.

**Consent to participate** Not applicable.

**Consent for publication** Not applicable.

**Competing interests** The authors declare no competing interests.

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