



The *in vitro* activity of polymyxin B and tigecycline alone and combination with other antibiotics against carbapenem-resistant *Enterobacter cloacae* complex isolates, including high-risk clones

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Background: The emergence of carbapenem-resistant Enterobacteriaceae (CRE) has become a significant problem for global public health. Currently, treatments program is minimal. This study aimed to evaluate the molecular mechanisms of carbapenem-resistant *Enterobacter cloacae* complex isolates (CREC) infections.

Methods: Resistance genes were detected using PCR with specific primers. Multilocus sequence typing (MLST) was also performed. Furthermore, we evaluated the effects of polymyxin B (PMB) and tigecycline (TGC) antibiotics (Abs) alone and in combination with meropenem (MEM), amikacin (AMK), and levofloxacin (LEV) against CREC isolates. The results were then compared with *in vitro* synergy testing results obtained from time-kill assays (TKAs), and the microdilution checkerboard method.

Results: The synergistic efficiency of PMB + TGC was also evaluated. Abs use clinically achievable concentrations to determine the antibacterial effects of the Ab. Similar sequence type (ST) classifications had a comparably resistant phenotype; PMB-based combination therapy is better than TGC-based combination therapy.

Conclusions: we found that the combination of PMB + AMK is promising for the treatment of AMK-sensitive CREC. The high-risk ST93 carrying the *bla*_{KPC-2} gene should be monitored.

Keywords: Synergism; carbapenemases; *Enterobacter cloacae* complex (ECC); polymyxin B (PMB); tigecycline (TGC); ST93

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Introduction

The World Health Organization (WHO) has listed carbapenem-resistant enterobacteria (CRE) as one type of bacteria that is in urgent need of new antibiotics (Abs) development. Most *in vitro* CRE data concerns *Klebsiella*

pneumoniae with *Enterobacter cloacae* complex (ECC) data being poorly understood (1). With the emergence of the carbapenem-resistant *Enterobacter cloacae* complex (CREC), combination antimicrobial therapies are more often being considered. Polymyxin B (PMB) Abs have been used since

the late 1950s, but can cause ototoxicity and neurotoxicity; less toxic aminoglycosides in the 1970s gradually replaced PMB Abs. However, due to the lack of new Ab development, PMBs continued to be used in clinical practice, even with the pharmacokinetic/pharmacodynamic limitations and increased risk of severe renal insufficiency. It was found that PMBs might promote membrane transport of other Abs, thereby enhancing bactericidal or bacteriostatic activities (2,3). The broad-spectrum antimicrobial agent, tigecycline (TGC), was also active against most Gram-negative isolates *in vitro*, including ESBL producers. TGC is indicated for the treatment of complicated skin and soft tissue infections (4). PMB or TGC has been used in Ab combination therapy studies (2,5,6). However, past *in vitro* studies has ignored the importance of using clinically achievable concentrations to determine antibacterial effects. In our study, we used the checkerboard method and time-kill analysis (TKAs) to evaluate the effect of PMB or TGC combined with meropenem (MEM), amikacin (AMK), and levofloxacin (LEV) against seven CREC isolates. We also studied the effects of TGC combined with PMB. The purpose of this study was to provide more guidance for the medical treatment of CREC infection.

Methods

Bacterial isolates

Seven non-repeated clinical CREC strains were collected from the First Affiliated Hospital of Jiamusi University between September 2016 and December 2018, and all isolates were stored frozen until further use. Species identifications and minimum inhibitory concentrations (MICs) of the Abs evaluated were analyzed using the Vitek2 compact system (BioMerieux, France). MICs of MEM, PMB, TGC, AMK, and LEV were further determined by broth microdilution according to the guidelines of the 2016 Clinical and Laboratory Standards Institute (CLSI 2016).

Antibiotics

Antimicrobial agents included PMB (Sigma Chemical, St. Louis, MO, USA), AMK (Tianjin Chemical Co., Ltd., Tianjin, China), LEV (CISEN Chemical Industries, Ltd., Shandong, China), MEM (Haibin Chemical Industry Co. Ltd., Shenzhen, China), and TGC (Pfizer Co. Ltd., Hangzhou, China). A PMB concentration of 2 µg/mL is considered a sensitive breakpoint (7).

PCR studies

The resistance genes were detected using the polymerase chain reaction (PCR), including MBL (*bla*_{NDM-1}, *bla*_{IMP-4}, *bla*_{VIM}) and others β-lactamases (*bla*_{TEM}, *bla*_{SHV}, *bla*_{CTX-15}, *bla*_{CTX-9}, *bla*_{KPC}, *bla*_{OXA-48}, *bla*_{OXA-1}, *bla*_{OXA-23}, *bla*_{OXA-24}, *bla*_{OXA-51}, and *bla*_{OXA-58}). AmpC enzyme genes (*bla*_{ACC}, *bla*_{DHA}, *bla*_{CMY}), quinolone resistance determining regions (QRDRs), *gyrA* and *parC*, and plasmid-mediated quinolone resistance (PMQR) determinants [*qnrA*, *qnrB*, *qnrS*, *aac*(6′)-Ib, and *qepA*] (8,9) were also detected. The colistin resistance gene *mcr-1* primer sequences and conditions were previously reported by Liu *et al.* (10).

Multilocus sequence typing (MLST)

MLST for seven housekeeping genes, including *dnaA*, *fusA*, *gyrB*, *leuS*, *pyrG*, *rplB*, and *rpoB*, were performed according to a previously described method (<https://pubmlst.org/ecloacae/>).

Synergy evaluations using the checkerboard assay

The activities of PMB + TGC, and PMB or TGC combined with AMK, LEV, and MEM were measured against seven bacterial strains derived from FICI values using the microdilution checkerboard method. Five Ab concentrations were prepared, 1/8× MIC, 1/4× MIC, 1/8× MIC, 1× MIC, and 2× MIC. Along with the horizontal and vertical wells, 50 µL of each Ab was added into each well, and then a 5×10⁵ CFU/mL bacterial suspension was inoculated into each well. The negative control wells contained only MH broth (MHB). The results were interpreted as follows: a FICI ained only MH bro, 0.5< FICI ≤4, indifferent, and FICI >4, antagonistic (11). *E. Cloacae* (National Center for Medical Culture Collections, CMCC-45301) were used as quality control strains.

The minimum bactericidal concentration (MBC) method of detection

An inoculum that contained 5×10⁵ CFU/mL bacterial suspensions in the exponential growth phase was used for these experiments. Microtiter plates were inoculated with the bacterial suspension and diluted antimicrobials. After incubating the plates at 37 °C for 24 h, suspensions corresponding to 1×, 2×, 4×, and 8× the MICs of the compound were spread onto MH agar plates and incubated

overnight at 37 °C. Plates that had a colony counts of less than 10 were considered to be the MBC (12).

Time-kill assays (TKA)

TKA were conducted for each strain according to previously described methods. The Ab concentrations used for PMB were 0.25 and 1 µg/mL. The Ab concentrations used for TGC were 0.25 and 1 µg/mL. The concentrations for MEM with PMB or TGC were 4 and 16 µg/mL. For AMK with PMB or TGC, the concentrations were 8 and 16 µg/mL. The concentrations of LEV with PMB or TGC were 6.4 µg/mL. All concentrations were determined according to clinically achievable concentrations (7,13). A 5×10^5 CFU/mL inoculum of the tested organism with into 10 mL of the corresponding broth (containing the Abs alone or in combination). The samples were obtained aseptically at 0, 2, 4, 8, and 24 hours. Regrowth was defined as an increase in colony counts from the previous time point. Synergistic effects were determined by a decrease of $\geq 2 \log_{10}$ CFU/mL when compared with the combined Abs, with the most active single antimicrobial alone at that time point, while an increase of $>2 \log_{10}$ was considered antagonistic.

Results

Antibiotic susceptibility and level of resistance

Susceptibilities for seven strains are listed in *Table 1*. In summary, all isolates were susceptible to PMB (MICs were 2–4 µg/L), MEM (MICs were 8–256 µg/L); AMK (MICs were 0.5–20,480 µg/L), TGC (MICs were 0.25–1 µg/L), and LEV (MICs were 0.0625–32 µg/L). Four strains that were resistant to PMB (MIC =4 µg/L), but a *mcr-1* gene was not found.

Epidemiological links and drug resistance gene of CREC

Seven isolates were clustered into five sequence types (STs), including ST520, ST528, ST1119, ST1120, and ST93. Among them, ST1119 and ST1120 are newly discovered STs (*Table 1*). The clustering of CREC isolates was also apparent in the analysis of the antimicrobial resistance gene contents, as shown in *Figure 1*. Similar ST types had a similar drug resistance phenotype. The isolates were classified according to the content of resistance determinants using a binary distance matrix and the UPGMA clustering method. In brief, except for the

CREC6 isolate, all strains carried the metal β -lactamase gene. Interestingly, the CREC3 strain co-produced the *bla*_{NDM-1} and *bla*_{IMP-4} genes. The *bla*_{OXA-1} carbapenemase class D family gene was detected in four isolates. Among the AmpC-producing enzymes, *bla*_{ACC} and *bla*_{DHA} were found. Focusing on PMQR harbored by these isolates, we showed that the *qnr* and *aac(6')-Ib-cr* genes were detected. Two *qnr* family genes were also found (*qnrB* and *qnrS*), no *qnrA* or *qepA* genes were detected, and QRDR DNA sequences of the *gyrA* and *parC* genes were identified. Mutations were seen in three sites of the CREC4 strain and the two sites of the CREC6 strain. The MICs of the two isolates were 16 and 32 µg/mL, respectively. In addition, mutations in the one site of the CREC3 and CREC7 strains were found, and the MIC of the two isolates were 8 µg/mL. Substitutions in the QRDR were detected in LEV resistant strains, as shown in *Figure 1*.

Checkerboard synergy analysis

The results of the checkerboard combination studies are shown in *Table 1*. The combination of PMB and MEM resulted in FICI values of 0.5, 0.25, and 0.5 against CREC1, CREC2, and CREC3, respectively. The combination of PMB and AMK resulted in FICI values of 0.25 and 0.375 against CREC5 and CREC7, respectively. PMB and LEV displayed FICI values of 0.5 against CREC6 and CREC7, and PMB + TGC resulted in FICI values of 0.25 against CREC6, which indicated synergistic interactions. The results of the checkerboard combination studies based on TGC showed no synergistic interactions. It is worth noting that antagonism was not found with any combination. No MBC values exceeded 4× MIC, as shown in *Table 1*.

Time-kill assays results

All monotherapies were evaluated against the seven isolates using the TKAs. Specifically, MEM used alone at 4, and 16 µg/mL concentrations were bacteriostatic, resulting in regrowth by 8 hours for all isolates. PMB used alone at 0.25, and 1 µg/mL was rapidly bactericidal for two isolates by 2 hours (the CREC1 and CREC5 strains). Other strains exhibited bacteriostatic actions, and regrowth was observed by 8 hours except for the CREC1 strain. TGC used alone at 0.25, and 1 µg/mL was bacteriostatic for all isolates by 2 hours, but regrowth was observed by 8 hours. AMK used alone at 16 µg/mL was bactericidal by 8 hours for the

Table 1 Clinical information (MLST, MICs, MBCs and FICIs of polymyxin B; tigecycline; meropenem; amikacin; levofloxacin against CREC isolates)

Patient	Sex	Age (year)	Clinical sample	MLST	Clinical wards	PMB (MIC/MBC)	MEM (MIC/MBC)	AMK (MIC/MBC)	TGC (MIC/MBC)	LEV (MIC/MBC)	FICI						
											PMB-MEM	PMB-AMK	PMB-TGC	PMB-LEV	TGC-LEV	TGC-AMK	TGC-MEM
1	M	31	Wound swab	1,119	Emergency department	2/4	16/16	2/8	1/1	0.5/1	0.5*	0.625	0.625	0.625	1	0.625	0.625
2	F	59	BAL	520	Emergency ICU	4/4	16/16	20,480/40,960	1/1	0.0625/0.0625	0.25*	No	0.625	0.625	0.625	No	0.625
3	M	83	Urine	1,120	Infectious department	2/4	32/128	0.5/0.5	1/1	8/16	0.5*	0.625	1	0.75	1	0.75	0.625
4	F	82	Pus	93	Emergency ICU	4/4	64/64	8/16	0.5/0.5	16/32	0.75	0.625	0.625	0.75	0.625	1	0.625
5	F	46	BAL	528	Emergency department	2/2	8/8	16/16	0.5/0.5	0.25/0.5	0.75	0.25*	0.75	1	1	0.625	1
6	M	16	BAL	93	Emergency department	4/4	256/256	8/8	0.25/0.5	32/32	0.625	0.625	0.25*	0.5*	1	0.625	0.625
7	F	65	Pus	1,120	Neurosurgery department	4/4	32/32	64/64	1/1	32/32	0.625	0.375*	0.75	0.5*	1	1	0.75

*, indicates a synergistic combination. F, female; M, male; MLST, multiple sequence locus typing; PMB, polymyxin B; TGC, tigecycline; MEM, meropenem; AMK, amikacin; LEV, levofloxacin; MIC, minimum inhibitory concentrations; MBC, minimum bactericidal concentration; No, since the single drug MIC is too high (20,480 µg/mL), no experiment was performed. CREC, carbapenem-resistant *Enterobacter cloacae*.

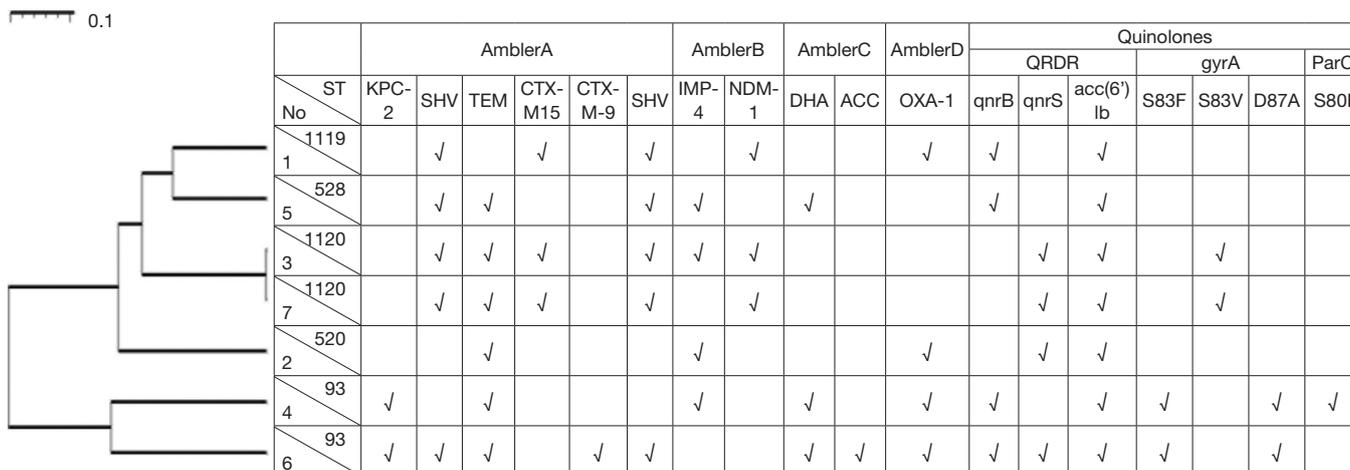


Figure 1 Genetic resistance determinants in *Enterobacter Cloacae* complex isolates from Jiamusi, Heilongjiang, China, January 12, 2014–December 31, 2015. The isolates were classified according to the content in resistance determinants by using a binary distance matrix and UPGMA clustering method. Scale bar shows the dissimilarity in resistance gene content. ST, sequence type.

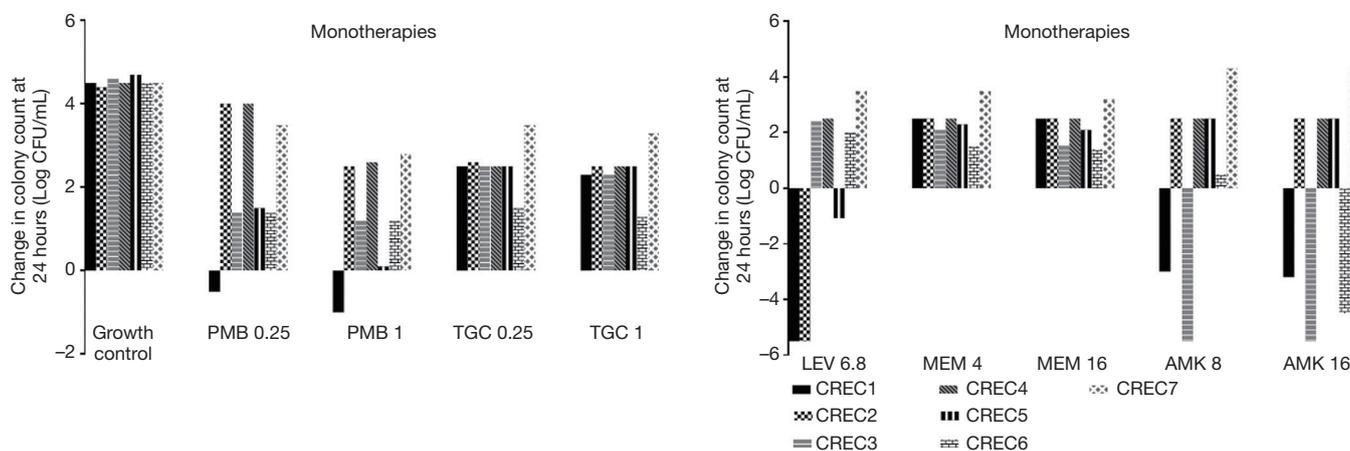


Figure 2 24-hour change in colony count for polymyxin B (PMB); tigecycline (TGC); meropenem (MEM); amikacin (AMK) and levofloxacin (LEV) against CREC isolates. Data are differences of geometric means at time points 0 and 24 h ,experiment repeated twice (n=2) PMB or TGC (0.25 and 1 µg/mL); MEM (4 and 16 µg/mL); and AMK (8 and 16 µg/mL) and LEV 6.8 µg/mL. CREC, carbapenem-resistant *Enterobacter cloacae*.

CREC1, CREC3, and CREC6 isolates; for these three strains, bactericidal activity was maintained for 24 hours. LEV used alone at 6.8 µg/mL was bactericidal by 24 hours for the CREC1 and CREC2 isolates; log change = logCFU₂₄ – logCFU₀, as shown in *Figure 2*.

Because the monotherapies were effectively bactericidal against the CREC1, CREC2, CREC3, and CREC6 isolates, we selected three clinical isolates (CREC4, CREC5, and CREC7) that the monotherapies were not bactericidal

against and evaluated the *in vitro* activity of PMB + TGC and PMB or TGC in combinations with MEM, AMK, and LEV.

In the TKA, synergy was not observed for all of the combinations. In general, PMB-based combination had a better antibacterial effect against the three isolates compared with the TGC-based combinations.

For CREC4 isolate, PMB+AMK combination had a better antibacterial effect with the TGC+AMK

combinations. TGC + MEM had a bacteriostatic and synergistic effect against CREC4 and increasing MEM concentrations had more pronounced synergistic effects. However, for the PMB + MEM combination, synergistic interactions were not observed. TGC + LEV appeared to be indifferent against the CREC4 isolates (Figure 3A,B,C,D).

For CREC5 isolate, PMB + AMK combination also had a better antibacterial effect with the TGC + AMK combinations. The TGC + MEM and PMB + MEM synergistic interactions were not observed. TGC + LEV and PMB + LEV appeared to be indifferent against the CREC5 isolates (Figure 3E,F,G,H).

For CREC7 isolate, the PMB + AMK and TGC + AMK combination had no bactericidal effects, for the PMB + MEM and TGC + MEM combination, synergistic interactions were not observed. High PMB + LEV concentrations have bacteriostatic and synergistic effects against the CREC7 isolate (Figure 3I,J,K,L).

We also combined PMB + TGC, which showed synergistic effects against all three strains; however, the bactericidal effects were not obvious, as shown in Table 2 and Figure 3.

Discussion

The presence of CRECs is due to the loss of porin, overproduction of a chromosomally encoded cephalosporinase, AmpC, activation of the efflux system, and the presence of important carbapenem hydrolases, such as metallo-beta-lactamase and *bla*_{KPC} (14). In China, the first reported CREC was a KPC-producing strain isolated from Shanghai in 2010 (15). Later, the CREC *bla*_{NDM}, *bla*_{IMP}, and *bla*_{VIM} genes were reported from different geographical regions (16). A previous CREC study showed that KPC-producing ECC exhibited lower MEM MICs of 2–8 µg/mL (7). In our study, KPC-producing CREC isolates exhibited higher MEM MICs (64–256 µg/mL) than IMP- and NDM-producing isolates (16–32 µg/mL). Fluoroquinolone (FQ) resistance is caused by mutations in DNA gyrase and topoisomerase IV chromosomal genes, and DNA gyrase is the primary target enzyme where substitutions occur to resist FQs. Our research results show that alterations in both GyrA and ParC often confer high-level resistance, consistent with conclusions by Hooper *et al.* (17).

Peirano *et al.* reported that ST93 had a global distribution and was associated with different carbapenemases (*bla*_{IMP-8}, *bla*_{IMP-14}, *bla*_{VIM-1}, *bla*_{NDM-1}, *bla*_{KPC-2}, and *bla*_{OXA-48}) (18). Of the seven CREC isolates studied, two carried the *bla*_{KPC-2}

carbapenemase and belonged to ST93. Interestingly, although the two isolates have similar resistant phenotypes, the effects of antibiotic combination therapy vary widely. In addition, results from another study confirmed that antibacterial activity is strain-specific, consistent with our findings. In a recent Brazilian study, a high mortality rate associated with KPC-producing *Enterobacter cloacae* was found in 24 patients, where the mortality rate reached 41.8% (19). Therefore, for the ST93 KPC-2-producing high-risk clones, infection control measures should be performed, and stricter Ab policies should be required to control the spread of ST93.

The cost to treat CRE is higher than with other chronic diseases. A study evaluating the health and economic burden of CRE in the United States showed that, depending on the infection type, the cost to the family, hospital, and society for a single CRE infection had created a substantial economic burden (20). And, while the number of studies is increasing, exploration of effective combinations against CRE are few. Combination therapies include the follow advantages: (I) synergistic effects have been demonstrated in *in vitro* studies that showed better clinical outcomes, and helped to guide clinical decisions, (II) drug resistance could rapidly develop if a sole antimicrobial, such as PMB and TGC, is given, (III) combination drug therapy could reduce the dose of each Ab, reducing side effects and decreasing patient costs.

Combining antimicrobials with different mechanisms of action could provide better pharmacokinetic effects. In contrast to most previous studies, which employed colistin in combination with other Abs against CREC, we chose PMB as the basis for the antimicrobial combinations because of its lower nephrotoxicity and higher tissue concentrations compared with colistin (2). Randomized trials have indicated that using TGC alone could confer an increased mortality risk (21), which further illustrates that TGC is best suited for combination therapy.

The lack of a gold standard has brought significant challenges in being able to detect synergistic effects on the clinical microbiology laboratory using *in vitro* drug susceptibility. TKA is time-consuming and technically challenging, but articles about TKA are often considered more relevant for clinical situations (22). The checkerboard method measures only the inhibitory activity. E-test synergy methods are more rapid and easier to perform, but there are more affected by other factors. In our research, the checkerboard and TKA methods are different. The results of the TGC-based checkerboard method were indifference,

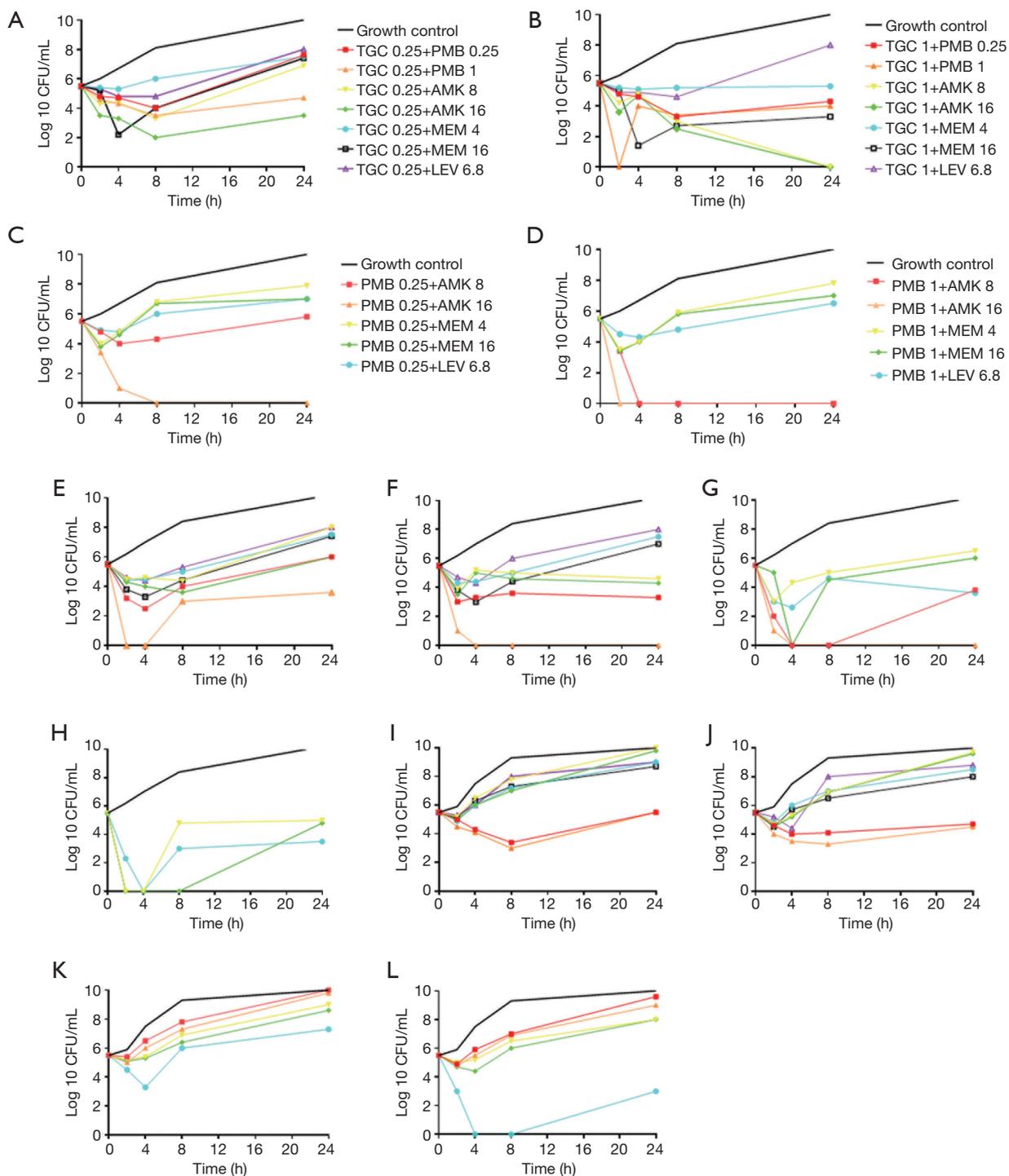


Figure 3 Time kill curves of polymyxin B and tigecycline combination with meropenem; amikacin and levofloxacin against CREC isolates. Data points are geometric means of replicate experiments (n=2). Antibiotic concentrations were PMB or TGC (0.25 and 1 µg/mL); MEM (4 and 16 µg/mL); and AMK (8 and 16 µg/mL) and LEV 6.8 µg/mL. (A,B,C,D) Represents CREC4; (E,F,G,H) represents CREC5; (I,J,K,L) represents CREC7. The legend of (A) is consistent with (E,I); the legend of (B) is consistent with (F,J); the legend of (C) is consistent with (G,K); The legend of (D) is consistent with (H,L). CREC, carbapenem-resistant *Enterobacter cloacae*; PMB, polymyxin B; TGC, tigecycline; MEM, meropenem; AMK, amikacin; LEV, levofloxacin.

Table 2 Three isolates selected for time-kill assay and their combination results at 24 h

Isolates, antimicrobials + concentrations (µg)	Effects of antibiotic combination, log change = (logCFU ₂₄ – logCFU ₀)		
	CREC4	CREC5	CREC7
T _{0.25} + P _{0.25}	2.1	0.5	-0.2*
T _{0.25} + P ₁	-0.8*	-1.9*	-0.3*
T _{0.25} + A ₈	1.4	2.5	4.5
T _{0.25} + A ₁₆	-2*	0.5	4.3
T _{0.25} + M ₄	2	2	3.5
T _{0.25} + M ₁₆	1.9	1.9	3.2
T _{0.25} + L _{6.8}	2.5	1	3.5
T ₁ + P _{0.25}	-1.2*	-2.2*	-0.8*
T ₁ + P ₁	-1.5*	-5.5*	-1*
T ₁ + A ₈	-5.5*	-0.9*	4.2
T ₁ + A ₁₆	-5.5*	-1.2*	4.1
T ₁ + M ₄	-0.2*	2	3
T ₁ + M ₁₆	-2.2*	1.5	2.5
T ₁ + L _{6.8}	2.5	1	3.3
P _{0.25} + A ₈	0.3*	-1.7*	4.5
P _{0.25} + A ₁₆	-5.5*	-5.5*	4.3
P _{0.25} + M ₄	2.4	1	3.5
P _{0.25} + M ₁₆	1.5	0.5	3.1
P _{0.25} + L _{6.8}	1.5	-1.9	1.8
P ₁ + A ₈	no	-5.5*	4.1
P ₁ + A ₁₆	no	no	3.5
P ₁ + M ₄	2.3	-0.5	2.5
P ₁ + M ₁₆	1.5	-0.7	2.5
P ₁ + L _{6.8}	1	-2	-2.5*

*, indicates a synergistic combination. PMB or TGC (0.25 and 1 µg/mL); MEM (4 and 16 µg/mL); and AMK (8 and 16 µg/mL) and LEV 6.8 µg/mL. No: since the low concentration can play a bactericidal effect, there is no synergistic killing experiment at this concentration. CREC, carbapenem-resistant *Enterobacter cloacae*; PMB, polymyxin B; TGC, tigecycline; MEM, meropenem; AMK, amikacin; LEV, levofloxacin.

but with the TKS method, the TGC-based checkerboard method showed synergy with AMK and MEM. At present, based on the limited data available, it is not possible to decide which *in vitro* test method is the most relevant for

clinical outcomes, so further research is urgently needed. It is worth noting that with the TKA, the superiority of PMB + MEM in combination with CREC was not found, but the synergistic effect against the CREC4 isolate was found with TGC + MEM. However, in a recent clinical study, TGC + MEM produced an antagonistic effect in the treatment of a KPC-2-producing *Klebsiella pneumoniae* infection, and eventually, the patient was cured with an increased dose of TGC (23). This result is inconsistent with our study. In our study, we found that TGC + MEM had a synergistic effect on the CREC4 isolate producing KPC-2, and no antagonism was found. The likely reason is that the combined effect differs among different strains. In addition, further clinical studies should be performed for the combination of TGC + MEM.

Limited studies have explored the activity of PMB or TGC + AMK against CREC. Cai *et al.* reported that PMB + AMK was synergistic against PMB-sensitive *Enterobacter cloacae* in 2/4 cases; however, no synergistic effect was evident against the PMB-resistant strains (2). Therefore, we evaluated KPC-producing PMB-resistant CREC isolates and found that PMB + AMK had bactericidal and synergistic activity against AMK-susceptible CREC isolates that produce KPC and/or IMP-4. Compared with PMB, TGC requires high concentrations to act on AMK-sensitive strains (CREC4). Therefore, the combination of PMB + AMK is promising for the treatment of AMK-sensitive CREC.

Cai and Betts *et al.* found that PMB + TGC was promising in the treatment of CREC (2,24). In contrast, PMB and TGC (at 0.25 or 1 µg/mL) were used in combination with other Abs, and we did not find sustained bactericidal activity against all three CREC isolates within 24 hours except for PMB-sensitive CREC5. Therefore, TGC + PMB should be used with caution, against the PMB-resistant CREC. The CREC7 strain was resistant to both PMB, AMK, and LEV, so no bactericidal effect was observed for all of our experimental combinations. Only high PMB concentrations combined with LEV showed synergistic and bacteriostatic effects.

The antimicrobial concentrations used in the tests are important when considering the antimicrobials combinations assays used. In this study, we used clinically achievable concentrations to determine antibacterial effects, making the obtained results more useful to clinical medicine. Our study has some limitations. First, the number of isolates was small, and those in Jiamusi, China, only represented the hospitals. Second, further clinical studies

determining the relevance of these data are warranted.

Conclusions

In this study, we showed that the ST93 of the KPC-2 producing high-risk clones are dangerous, and research should be intensified to combat these bacteria with stricter Ab policy requirements to control the spread of CREC. In addition, we found that the combination of PMB + AMK is promising for the treatment of AMK-sensitive CREC. Future studies should investigate these drug combinations in animal models, and studies of the pharmacokinetic parameters of these drug combinations should be performed.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The Ethics Committee of Jiamusi University Clinical Medical College approved the study protocol for research. The committee's reference number is 20180326. Individual informed consent was waived by the ethics committee listed above because this study used currently existing samples collected during routine medical care and did not cause any additional risks to the patients.

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