



Tigecycline in combination with other antibiotics against clinical isolates of carbapenem-resistant *Klebsiella pneumoniae in vitro*

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Background: To investigate the activity of 5 antibiotic monotherapies, including colistin (COL), meropenem (MEM), amikacin (AMK), levofloxacin (LEV), and tigecycline (TGC), when combined with 4 other antibiotics against clinical isolates of carbapenem-resistant *Klebsiella pneumoniae* (CRKP) *in vitro*.

Methods: The minimum inhibitory concentrations (MICs) of 5 antibiotics against 40 CRKP isolates were determined by micro-broth dilution method. There were synergistic effects between TGC combinations in the 10 CRKP isolates detected with checkerboard microdilution method. Time-kill assay was used to assess the monotherapies and the TGC combinations against 4 distinct sequence typing (STs) CRKP isolates. Polymerase chain reaction (PCR) tests were used to detect the carbapenemase genes, extended-spectrum beta lactamase (*ESBL*) genes, colistin resistance gene, and quinolone resistance genes, while multilocus sequence typing (MLST) was performed for 10 CRKP isolates.

Results: The MICs of TGC, COL, MEM, AMK, and LEV were 0.5–2, 2–32, 4–256, 1–16,384, and 0.5–64 µg/mL, respectively. The combinations exerted a significant synergism or additive effect via the checkerboard technique for most tested CRKP isolates, but a portion of the CRKP isolates had an indifferent effect except for the TGC-AMK combination. In addition, time-kill assays revealed that TGC enhanced the bactericidal activity of the 4 other antibiotics. Among 10 CRKP isolates, *blaKPC-2* (90%), *blaSHV* (100%), and *blaacc(6)-Ib* (100%) were the most common carbapenemase genes, *ESBL* genes, and quinolone resistance genes, respectively. ST76 (70%) was the most predominant clone, followed by ST11 (10%), ST375 (10%), and ST530 (10%).

Conclusions: In contrast to the currently recommended TGC therapy, our *in vitro* data suggest that TGC combinations may be a valid therapeutic option against CRKP, even in the presence of 1 antibiotic resistant isolate in TGC combination therapy. TGC-AMK combination is a cost-effective option for treating CRKP in the eastern region of Heilongjiang Province. In addition, TGC combinations might circumvent the overuse of carbapenems during the era of multi-drug resistance in *Klebsiella pneumoniae* (KP).

Keywords: Carbapenem-resistant *Klebsiella pneumoniae* (CRKP); tigecycline; combination therapy; synergistic effect; resistance genes

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Introduction

Klebsiella pneumoniae (KP) are *Enterobacteriaceae* responsible for various human infections such as bacteremia, meningitis, and pneumonia (1). Standard treatment involves the use of carbapenems whose wide application has led to the emergence of carbapenem-resistant *Enterobacteriaceae* (CRE) (2). The resistance occurring in China is mainly due to the emergence of carbapenems hydrolases producing *Klebsiella pneumoniae* carbapenems (KPC), for example. The infection caused by carbapenem-resistant *Klebsiella pneumoniae* (CRKP) is complicated. CRKP usually carries multiple drug resistance genes which limit the treatment effect. At present, only a few drugs, such as colistin (COL), tigecycline (TGC), some aminoglycosides, and ceftazidime-avibactam, still have favorable *in vitro* activity against CRKP. The potential nephrotoxicity of COL and aminoglycosides restricts their broad use, especially among elderly patients with renal insufficiency. Ceftazidime-avibactam only inhibits the activity of class A enzymes. TGC, the 9-t-butylglycylamido derivative of minocycline, is regarded as the last resort for the management of difficult-to-treat carbapenem-resistant isolates infections (3). This limitation in treatment options further amplifies the need for new antibiotics (4,5). The dearth of novel antibiotics introduced to the market has been attributed to regulatory hurdles, high research costs, and low investment returns (6). A possible solution is drug repurposing. Since an approved drug has been cleared in terms of safety, pharmacological profile, and manufacturing process, it can be rapidly made available for a new disease indication (7). Based on this, we screened TGC combinations with other antibiotics, such as COL, meropenem (MEM), amikacin (AMK), and levofloxacin (LEV), to study antibacterial activity against clinical isolates of CRKP *in vitro*. Our research may provide laboratory data on antibiotic the susceptibility for drug-resistant patients and compensate for the lack of domestic antibiotic-related studies in the east of Heilongjiang.

Methods

Bacterial isolates

Forty CRKP isolates were collected from patients at the 1980-bed First Affiliated Hospital of Jiamusi University in Heilongjiang Province, northeast China, from October 2015 to January 2019. The isolates were identified as CRKP isolates by the Vitek 2 system (bioMérieux) and the AST-

GN card (bioMérieux, France). Carbapenemase resistance (either to ertapenem, imipenem, or meropenem) was determined according to Clinical and Laboratory Standard Institutes (CLSI-2016) criteria.

Antimicrobial susceptibility testing

The concentration ranges for the antibiotics tested were 0.25–128 µg/mL for TGC, 0.0625–64 µg/mL for COL, 0.5–256 µg/mL for MEM, 0.03125–16,384 µg/mL for AMK, and 0.25–128 µg/mL for LEV. The minimum inhibitory concentrations (MICs), defined as the lowest compound concentrations (µg/mL) required to stop bacterial growth, of the five antibiotics tested were determined using the microbroth dilution method per CLSI recommendations (8). Strain KP ATCC 700603 was *ESBL*-positive and was used as the reference strain. The MEM, AMK, and LEV results were interpreted based on CLSI criteria (8), whereas the TGC and COL results were interpreted based on the European Committee on Antimicrobial Susceptibility Testing (EUCAST) breakpoint recommendations (9). Finally, the MIC results were read on an enzyme-labeled instrument via optical density (OD) at 570 nm.

Checkerboard technique

TGC was tested for synergistic activity against 10 CRKP isolates via the checkerboard technique with the TGC-COL, TGC-MEM, TGC-AMK, and TGC-LEV combinations. The concentration ranges were based on the MICs determined above. Fifty microliters of each antibiotic at 5 increasing (4-fold) concentrations (0.125× MIC to 2× MIC) were used, and each well was inoculated with 100 µL of a 7.5×10^5 CFU/mL suspension of the test CRKP isolates in a final volume of 200 µL in duplicate. Results were measured using an enzyme-labeled instrument through OD at 570 nm after incubating at 37 °C for 24 hours.

The effects of the different antimicrobial combinations were defined according to the fractional inhibitory concentration index (FICI) as follows: FICI ≤0.5, synergism; 0.5 < FICI ≤1, additive; 1 < FICI ≤2, indifferent; or FICI >2, antagonistic (10).

Antibacterial time-kill assay

The antibacterial time-kill assay was used to assess the results of the antibiotic monotherapies and TGC combinations

against 4 CRKP isolates (including 3 *KPC*-producing isolates and 1 *KPC*-free isolate) based on CLSI guidelines. Antibiotic concentrations were calculated using the mean value of the steady-state concentrations of the nonprotein-bound drug in humans as described previously (11). The monotherapies were carried out with TGC 0.1 and 1 mg/L; COL 0.25 and 1 mg/L; MEM 4 and 16 mg/L; AMK 8 and 16 mg/L; and LEV 7 and 10 mg/L (12–14). The TGC combinations used were 1 dose of TGC combined with 1 dose of COL, MEM, AMK, or LEV. A 5×10^5 CFU/mL inoculum of the tested organism was used to inoculate 10 mL of the corresponding broth (containing the antibiotics alone or in combination). The samples were obtained aseptically at predetermined timepoints (0, 2, 4, 8, 12, and 24 hours). Time-kill curves were then constructed as a function of time, and the results are represented as the difference in \log_{10} between the CFU/mL at 0 and 24 hours. A decrease of $\geq 3 \log_{10}$ compared with the initial CFU/mL indicated a bactericidal effect. Bacteriostatic activity was defined as a $< 3 \log_{10}$ CFU/mL decrease in colony counts. Regrowth was defined as an increase in colony counts from the previous timepoint (15). Synergistic effects were determined by a decrease of $\geq 2 \log_{10}$ CFU/mL when comparing the combined antibiotics with the most active drug at that timepoint, while an increase of $> 2 \log_{10}$ was considered antagonism. Additivity and indifference were interpreted as any other outcome that did not meet the criteria for either synergism or antagonism (16).

Molecular detection of resistance genes and homology analysis

Polymerase chain reaction (PCR) tests were used to detect the carbapenemase genes (*blaKPC*, *blaNDM*, *blaVIM-1*, *blaVIM-2*, *blaIMP-4*, *blaIMP-8*, *blaOXA-23*, *blaOXA-24*, *blaOXA-48*, *blaOXA-51*, and *blaOXA-58*), extended-spectrum beta lactamase (*ESBL*) genes (*blaCTX*, *blaTEM*, *blaACC*, and *blaSHV*), colistin resistance gene (*blaMCR-1*) and quinolone resistance genes (*blaqnrA*, *blaqnrB*, *blaqnrS*, *blaqepA*, and *blaacc(6)-Ib*) in 10 CRKP isolates, as described in a previous study (17). Bioedit software was used to analyze the test data, and the results were compared using online blast software. In addition, multilocus sequence typing (MLST) was performed using 7 housekeeping genes of *K. pneumoniae* which were amplified using primers Available online online databases (http://bigsd.b.pasteur.fr/klebsiella/primers_used.html). The products of PCR were sequenced. Sequence types (STs) were determined using online database tools.

Results

Bacterial isolates

From October 2015 to January 2019 a total of 40 nonduplicated CRKPs were isolated from various clinical specimens. The CRKP isolates were isolated, one from each patient, with an age range of 16–86 years (median 61.5 years). Of the 40 patients, 25 (62.5%) were male. The mortality of the patients with CRKP infections was 20%. Three patients were excluded: 1 outpatient had no hospital records, and 2 nosocomial patients fell out of contact. The specimens with positive culture for CRKP included sputum (80%, $n=32$), blood (15%, $n=6$), wound secretion, and other specimens (2.5%, $n=1$, each). The CRKP isolates emerged from neurosurgery (32.5%, $n=13$); ICU (50.0%, $n=20$); emergency room (7.5%, $n=3$); and hematology, orthopedics, general surgery, and cardiac surgery departments (2.5%, $n=1$, each).

Antimicrobial susceptibility

In terms of the susceptibility patterns determined by the micro-broth dilution method, the MICs range of CRKP isolates against TGC, COL, MEM, AMK, and LEV were 0.5–2, 2–32, 4–256, 1–16,384, and 0.5–64 $\mu\text{g/mL}$, respectively. TGC was sensitive against all CRKP isolates. The resistance to COL, MEM, AMK, and LEV was 92.5%, 100%, 2.5%, and 15.0%, respectively. The 50% MIC (MIC_{50}) of TGC, COL, MEM, AMK, and LEV were 1, 8, 16, 4, and 1 $\mu\text{g/mL}$, respectively. The 90% MIC (MIC_{90}) of TGC, COL, MEM, AMK, and LEV were 2, 16, 128, 8, and 8 $\mu\text{g/mL}$, respectively (Table 1).

Synergistic activity and statistical analysis

Synergism and additive effects were detected in TGC combinations using the checkerboard technique, in which the FICI of the TGC-COL, TGC-MEM, TGC-AMK, and TGC-LEV combinations were 0.675 ± 0.188 , 0.613 ± 0.358 , 0.575 ± 0.237 , and 0.863 ± 0.314 , respectively (Table 2). TGC-AMK showed the best synergistic and additive effects with a FICI of 70% and 30% in TGC combinations, respectively. The synergistic effects of TGC-MEM and TGC-COL were 50% and 10%, respectively. The additive effects of TGC-COL, TGC-LEV, and TGC-MEM were 80%, 70%, and 40%, respectively. The indifferent effects of TGC-LEV, TGC-COL, and TGC-MEM were 30%, 10%, and 10%, respectively (Table 3). The TGC-COL, TGC-MEM,

Table 1 Antimicrobial resistance rates and MIC distribution of 40 CRKP isolates

| Antibiotic | MIC ($\mu\text{g/mL}$) | | | |
|------------|--------------------------------|----------|-------------------|-------------------|
| | Cutoff value of resistance (%) | Range | MIC ₅₀ | MIC ₉₀ |
| TGC | >2 (0%) | 0.5–2 | 1 | 2 |
| COL | >2 (92.5%) | 2–32 | 8 | 16 |
| MEM | ≥ 4 (100%) | 4–256 | 16 | 128 |
| AMK | ≥ 16 (2.5%) | 1–16,384 | 4 | 8 |
| LEV | ≥ 8 (15.0%) | 0.5–64 | 1 | 8 |

MIC, minimum inhibitory concentrations; MIC₅₀, MIC at which 50% of the isolates tested are inhibited; MIC₉₀, MIC at which 90% of the isolates tested are inhibited; TGC, tigecycline; COL, colistin; MEM, meropenem; AMK, amikacin; LEV, levofloxacin.

TGC-AMK, and TGC-LEV combinations decreased the MICs of TGC by 2.7-fold, 4.2-fold, 3.6-fold, and 3.1-fold, respectively.

Bacterial time-kill effect

Considerable regrowth occurred in the antibiotic monotherapies in the time-kill assay (Figure 1). TGC had bacteriostatic activity and AMK had bactericidal effect on 4 CRKP isolates. On 3 CRKP isolates, 0.25 mg/L COL had bacteriostatic activity, and had a bactericidal effect for 1 CRKP isolate (ST375). On 2 CRKP isolates (ST76 and ST530), 1 mg/L COL had bacteriostatic activity, while on the other 2 CRKP isolates, it had a bactericidal effect. MEM had bacteriostatic activity on 2 CRKP isolates (ST11 and ST530) and had bactericidal effect on the other 2 CRKP isolates. Seven mg/L LEV had bacteriostatic activity on 4 CRKP isolates; 10 mg/L LEV had bacteriostatic activity on 3 CRKP isolates and had a bactericidal effect for 1 CRKP isolate (ST11).

Time-kill assay showed a synergistic effect with TGC combination to 4 CRKP isolates (Figure 1). TGC-COL combination was synergic against 4 CRKP isolates. TGC-MEM combination, TGC-AMK combination, and TGC-LEV combination were synergic against 3 CRKP isolates (ST76, ST375, and ST530), 2 CRKP isolates (ST11 and ST76) and 2 CRKP isolates (ST76 and ST530), respectively.

Molecular detection of resistance genes and homology analysis

Among the 10 CRKP isolates, 9 (90%) were *KPC-2* producers. One CRKP isolate (10%) produced *NDM-5* carbapenemase gene. *ESBL* genes were found in 10 (100%),

9 (90%), 7 (70%), and 1 (10%) CRKP isolates carrying *blaSHV*, *blaTEM*, *blaCTX-M-15*, and *blaCTX-M-177* genes, respectively. No CRKP isolate was carrying the *blaMCR-1* gene. Quinolone resistance genes were found in 10 (100%), 8 (80%), and 2 (20%) CRKP isolates carrying *blaacc(6)-Ib*, *blaqnrB*, and *blaqnrS* genes, respectively. Four distinct STs were observed among all CRKP isolates and ST76 (n=7) was the most predominant clone, followed by ST11 (n=1), ST375 (n=1), and ST530 (n=1) (Table 2).

Discussion

Despite the increasing occurrence and the severity of infections due to CRKP, limited data exist on the efficacies of the available treatment schemes. We carried out this study to investigate both the *in vitro* activity of TGC, COL, MEM, AMK, and LEV, alone or in TGC combinations against CRKP, and to provide available treatment schemes applicable to clinical settings.

As for the antimicrobial susceptibility testing in this study, TGC was the most sensitive against all CRKP isolates and was the most effective regimen when used alone. These findings are consistent with the results of another recent survey that included 18 European countries, in which a susceptibility rate of TGC to CRE of 88.6% was found (18). With the investigation of TGC susceptibility in numerous selected pathogens, TGC was found to be one of the most active antimicrobial agents against gram-positive isolates and also to be effective against gram-negative isolates *in vitro*, including drug-resistant pathogens (19). Other drugs in our study with higher sensitivities were AMK and LEV. AMK had a higher sensitivity rate, possibly because it has only been used for a short time in this region or it has been re-enabled somehow. When AMK is used to treat CRKP, it

Table 2 Results of clinical information, bla genotype, MLST, MICs, and FICIs of tigecycline, colistin, meropenem, amikacin, and levofloxacin against 10 CRKP isolates

| Isolate | Source | Section | Carbapenemase genotype | ESBL genotype | Quinolone-resistance genotype | MLST | MIC ($\mu\text{g/mL}$) | | | | | | FICI | | | | | | |
|---------|--------|----------------------|------------------------|---------------------|-------------------------------|-------|--------------------------|-----|-----|-----|-----|-----------|-----------|-----------|-----------|-----------|-----------|-------|-------|
| | | | | | | | TGC | COL | MEM | AMK | LEV | TGC + COL | TGC + MEM | TGC + AMK | TGC + COL | TGC + MEM | TGC + AMK | LEV | |
| 700603 | - | - | - | - | - | - | 2 | 4 | 1/4 | 1/2 | 1/2 | 1/2 | - | - | - | - | - | - | - |
| CRKP1 | Sputum | Intensive care unit | KPC-2, NDM-5 | SHV, TEM, CTX-M-177 | qnrB, qnrS, acc(6')-lb | ST76 | 1/2 | 8 | 64 | 2 | 1 | 1 | 0.625 | 0.375 | 1 | 0.625 | 0.375 | 1 | 0.75 |
| CRKP2 | Blood | Intensive care unit | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST76 | 1 | 4 | 16 | 4 | 8 | 8 | 0.75 | 0.75 | 0.5 | 0.625 | 0.75 | 0.5 | 0.625 |
| CRKP3 | Blood | Neurosurgery | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST76 | 1 | 4 | 16 | 2 | 1 | 1 | 0.625 | 0.625 | 0.5 | 1.25 | 0.625 | 0.5 | 1.25 |
| CRKP4 | Blood | Intensive care unit | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST76 | 1/2 | 4 | 64 | 2 | 4 | 4 | 1.125 | 0.25 | 0.75 | 1.125 | 0.25 | 0.75 | 1.125 |
| CRKP5 | Sputum | Intensive care unit | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST76 | 1 | 2 | 64 | 2 | 1 | 1 | 0.625 | 1.5 | 0.375 | 1.5 | 0.625 | 1.5 | 0.375 |
| CRKP6 | Sputum | Neurosurgery | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST76 | 1 | 4 | 16 | 2 | 1 | 1 | 0.625 | 0.625 | 0.375 | 0.75 | 0.625 | 0.375 | 0.75 |
| CRKP7 | Blood | Blood specialty | KPC-2 | SHV | qnrS, acc(6')-lb | ST11 | 1/2 | 4 | 128 | 1 | 64 | 64 | 0.75 | 0.375 | 0.5 | 0.625 | 0.375 | 0.5 | 0.625 |
| CRKP8 | Sputum | Neurosurgery | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, qnrS, acc(6')-lb | ST76 | 1 | 8 | 16 | 2 | 1/2 | 1/2 | 0.625 | 0.5 | 0.375 | 0.625 | 0.5 | 0.375 | 0.625 |
| CRKP9 | Sputum | Intensive care unit | KPC-2 | SHV, TEM, CTX-M-15 | qnrB, acc(6')-lb | ST375 | 2 | 2 | 8 | 1 | 4 | 4 | 0.625 | 0.75 | 0.625 | 0.625 | 0.75 | 0.625 | 0.625 |
| CRKP10 | Sputum | Emergency department | - | SHV, TEM | acc(6')-lb | ST530 | 2 | 2 | 4 | 1 | 1 | 1 | 0.375 | 0.375 | 0.375 | 0.375 | 0.375 | 0.375 | 0.375 |

MLST, multilocus sequence typing; MIC, minimum inhibitory concentrations; FICI, fractional inhibitory concentration index; TGC, tigecycline; COL, colistin; MEM, meropenem; AMK, amikacin; LEV, levofloxacin.

Table 3 Checkerboard synergy study results for the combinations of tigecycline with 4 different antibiotics (colistin, meropenem, amikacin, and levofloxacin) against 10 CRKP isolates

| Combination | Synergistic, n [%] | Additive | Indifferent | Antagonistic |
|-------------|--------------------|----------|-------------|--------------|
| TGC + COL | 1 [10] | 8 [80] | 1 [10] | 0 [0] |
| TGC + MEM | 5 [50] | 4 [40] | 1 [10] | 0 [0] |
| TGC + AMK | 7 [70] | 3 [30] | 0 [0] | 0 [0] |
| TGC + LEV | 0 [0] | 7 [70] | 3 [30] | 0 [0] |

TGC, tigecycline; COL, colistin; MEM, meropenem; AMK, amikacin; LEV, levofloxacin.

produces aminoglycoside-modifying enzymes and target *16S rRNA* gene mutations cannot be ignored (20). The sensitivity of LEV to CRKP was 85%, and some isolates may be resistant as they carry the quinolone-resistance genes like the *bla_{acc}(6')-Ib*, *bla_{qnrB}*, and *bla_{qnrS}* genes. However, there were other drugs with higher resistance including COL and MEM. The resistance rate to COL was over 92.5% and was negative for the *bla_{MCR-1}* gene. The *bla_{MCR-1}* gene had a low level of resistance to COL (8–16 µg/mL), but COL resistance may have the *PmrA/PmrB* and *PboP/PboQ* two-element regulation systems. All CRKP isolates were resistant to MEM, and most of them carried carbapenemase genes and *ESBL* genes that might have led to the resistance. CRKP isolates carrying multiple resistance genes render most antibiotic monotherapies ineffective. Meanwhile, it was noted that the higher dose regimen of either antibiotic monotherapy may not be tolerated due to serious side effects, which also becomes an important factor in discussing an optimal combination dosing regimen. Combinations may be the best choice for CRKP treatment through choosing an optimal combination to reduce the standard dose or use a lower dose in combinations.

In this study, we have used 4 antibiotic combinations and found synergism or additive effect for all of these combinations against most of CRKP isolates in the checkerboard microdilution. Overall, the combination of TGC-AMK was the most effective regimen, demonstrating synergism and additive effects against all 10 CRKP isolates. TGC and AMK both work together in bacterial ribosome by binding and interfering with bacterial protein synthesis to kill bacteria effectively (21,22). High-sensitive drugs may have better antibacterial effects in combinations, and combinations can have positive synergism and additive effects on CRKP isolates carrying multi-drug resistance genes. At the same time, our study found that the combination of TGC-COL and TGC-MEM were the best combinations against 4 CRKP isolates in the time-kill

assay. The combination of TGC-COL had a synergic effect against *KPC-2*-producing KP, which is consistent with the result that Toledo *et al.* reported; it did, however, have an antagonistic effect in combination with TGC-MEM, which is inconsistent with Toledo *et al.*'s reporting (23). At the same time, the clinical research conducted by Bi *et al.* showed that a TGC-MEM combination treatment failed in an infection caused by CRKP (24). Therefore, TGC-MEM combinations should be used with caution. The TGC-LEV combination only showed additive effect and even had an indifferent effect in the checkerboard microdilution while showing a synergistic effect against only one CRKP isolate in the time-kill assay. We also found that the 4 combinations had a synergic or additive effect against ST76 CRKP isolates in the checkerboard microdilution and showed a synergic effect in the time-kill assay.

The majority of CRKP isolates produced *KPC-2* in our study. First isolate of the *KPC-2*-producing KP was isolated in the First Affiliated Hospital of Zhejiang University School of Medicine in China in 2004 (25). Carbapenem resistance to KP has increased globally during the past decade and is typically caused by carbapenemase production in particular. *KPC-2*-producing CRKP isolates are of great clinical concern because of the frequent co-resistance to multiple antibiotic classes. Clinical studies have concluded that combination antibiotic therapy is associated with a better outcome than monotherapy for the treatment of severe infections with these isolates even if the isolated bacteria are susceptible to the individual drugs *in vitro* (26). This study showed that the 4 combinations had a synergistic or additive effect on most of *KPC-2*-producing CRKP isolates in the checkerboard microdilution, and the combination of TGC-COL, TGC-MEM, and TGC-AMK had a synergistic effect in the time-kill assay. The part of *KPC-2*-producing CRKP isolates were indifferent to all combinations except for the TGC-AMK combination in the checkerboard microdilution, indicating that the presence of the *KPC* gene

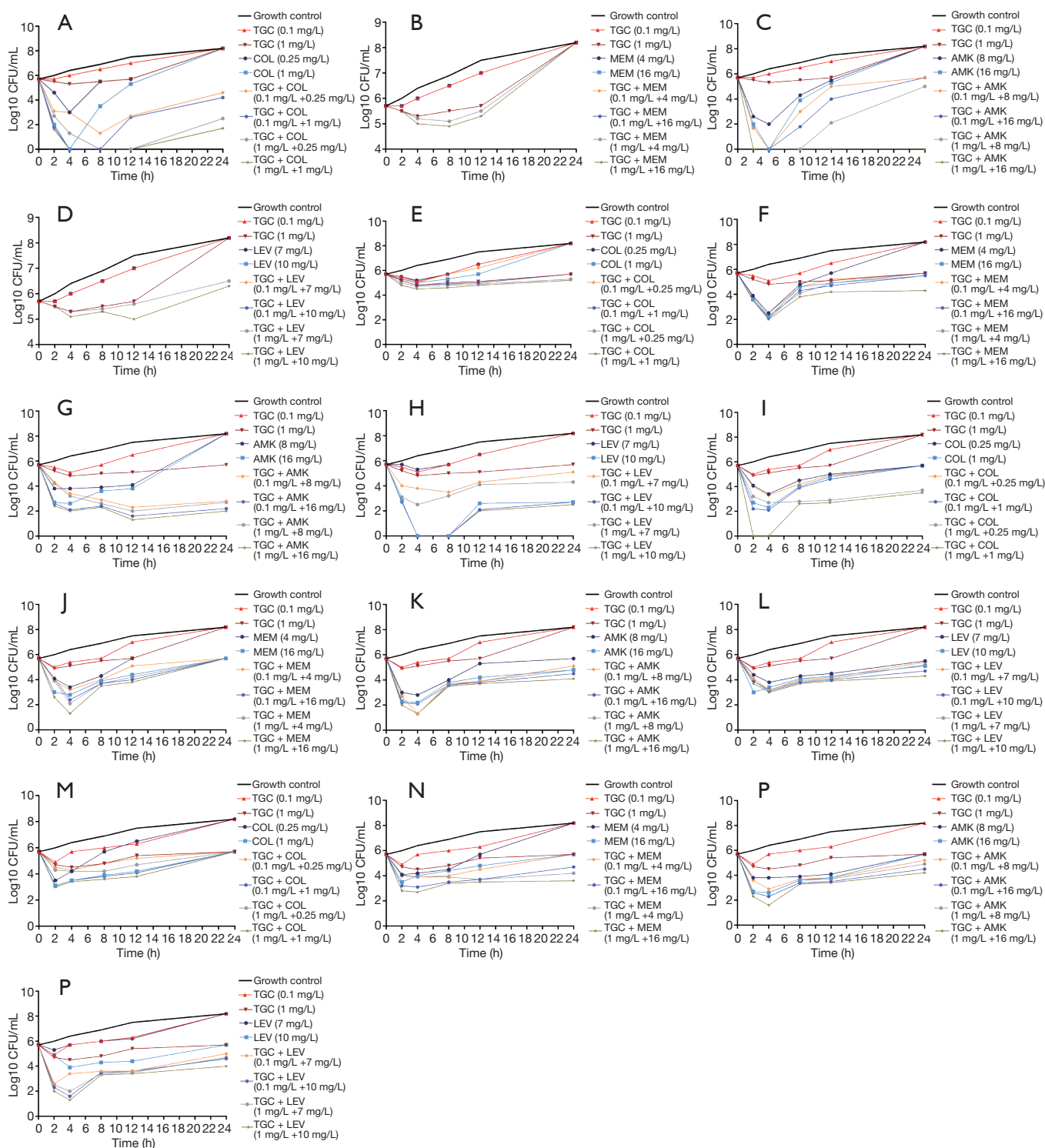


Figure 1 *In vitro* time-kill assays using serum concentrations of tigecycline (TGC), colistin (COL), meropenem (MEM), amikacin (AMK), and levofloxacin (LEV), either alone or in TGC combination against 4 CRKP isolates. (A-D) Monotherapies and combination therapies against the CRKP7 isolate, respectively; (E-H) monotherapies and combination therapies against the CRKP8 isolate, respectively; (I-L) monotherapies and combination therapies against the CRKP9 isolate, respectively; (M-P) monotherapies and combination therapies against the CRKP10 isolate, respectively.

resulted in a weakened antibacterial effect in combinations, but on 1 CRKP isolate without the *KPC-2* gene, the combination showed a synergistic or additive effect in the checkerboard microdilution and had a synergistic effect in the time-kill assay. *In vitro* experiments can reduce the MIC value of drug monotherapy against CRKP isolates and have better antibacterial activity in combinations. Assessment conducted by using the mean value of the steady-state concentrations of the nonprotein-bound drug in humans against CRKP isolates showed that combinations had a synergistic effect. Therefore, the optimal combination for the *KPC-2*-producing CRKP isolates should be selected and used with caution.

This study also found that the CRKP9 isolate was a hypervirulent *Klebsiella pneumoniae* (hvKP) while the capsular serotyping was K2. CRKP9-associated virulence genes mainly included the capsular polysaccharide gene *rmpA*, siderophore-associated genes *iucBC*, *iutA*, *iroBD*, and aerobactin were present on the tig00000014 plasmid; fimbrial adhesin genes, *fimA-H* and *mrkD*, and siderophore-associated genes, *iutA* and *entAB*, were present on the chromosome. CRKP9 isolate was sensitive to the 5 tested antibiotic monotherapies in the microbroth dilution and not sensitive to MEM. In line with these findings, Liao *et al.*'s earlier study reported that ST375 belonged to the K2 serotype and was sensitive to most antibiotics (27). TGC combinations had additive effects against the CRKP9 isolate in the checkerboard microdilution, the combinations of TGC-COL and TGC-MEM had synergistic effects in the time-kill assay, and CRKP isolates carrying virulence-associated genes were more sensitive in both monotherapy or combination. However, the CRKP9 isolate, which was highly sensitive to serum complement-mediated killing, died within 3 hours. Whether the addition of serum complement in antibiotics can increase the antibacterial activity against hvKP isolates remains to be studied.

Among the few antibiotics that remain effective, COL, TGC, gentamicin (GEN), and fosfomycin (FM) are often used in combination therapy against CRKP isolates (28). However, the most recommended treatment regimens are confusing, and there is no consensus regarding which antimicrobial combinations should be used to treat these infections (13). In our previously study, COL combined with MEM and AMK showed synergistic and additive effects against CRKP isolates and decreased the MICs of COL by 5.8- and 5.3-fold, respectively. *In vitro* antibacterial activity of COL combination was better than the TGC combination because of the greater reduction in the MICs of the main

antibiotic in combination. However, the combination of TGC with MEM or AMK showed synergistic and additive effects, and even indifferent effects and a decrease of the MICs of TGC by 4.2- and 3.6-fold, respectively. TGC has been approved in recent years for clinical use in China, but resistance to TGC has emerged since its approval (29,30). Du *et al.* confirmed the transfer capacity of TGC resistance, which was found to be mediated by the mutated *tetA* through a transferable plasmid. This finding serves as a therapeutic warning, as the *tetA* gene is frequently carried by CRKP isolates (31). However, a COL combination can reduce the development of COL heteroresistance as proven in *in vitro* studies, but this has yet to be supported in clinical studies (32). Therefore, COL combination is superior to TGC combination in the treatment of infection caused by CRKP.

Based on our study's conclusions, we suggest to physicians that TGC combined with AMK is the last active group in the setting of low-dose regimens. Conversely, the combination of TGC with COL or MEM showed the most *in vitro* synergistic and bactericidal activities against the *ESBL*-producing, carbapenem-resistant, and even virulence-associated gene CRKP isolates; this regimen could be considered as a last resort approach for the infections caused by CRKP isolates. In addition, TGC combination might circumvent the overuse of carbapenems during the era of multi-drug resistance in KP.

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Footnote

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