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Comparative pharmacodynamics of four different carbapenems in combination with polymyxin B against carbapenem-resistant Acinetobacter baumannii

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Abstract

The objective of this study was to determine the comparative pharmacodynamics of four different carbapenems in combination with polymyxin B (PMB) against carbapenem-resistant Acinetobacter baumannii isolates using time–kill experiments at two different inocula. Two A. baumannii strains (03-149-1 and N16870) with carbapenem minimum inhibitory concentrations (MICs) ranging from 8 to 64 mg/L were investigated in 48-h time–kill experiments using starting inocula of 10^6 CFU/mL and 10^8 CFU/mL. Concentration arrays of ertapenem, doripenem, meropenem and imipenem at $0.25 \times$, $0.5 \times$, $1 \times$, $1.5 \times$ and $2 \times$ published maximum serum concentration (C_{max}) values (C_{max} concentrations of 12, 21, 48 and 60 mg/L, respectively) were investigated in the presence of 1.5 mg/L PMB. Use of carbapenems without PMB resulted in drastic re-growth. All carbapenem combinations were able to achieve a $\frac{3 \log_{10} CFU/mL}{2 \log_{10} CFU/ML}$ reduction by 4 h against both strains at 10^6 CFU/mL, whereas maximum reductions against strain 03-149-1 at 10^8 CFU/mL were 1.0, 3.2, 2.2 and 3.3 \log_{10} CFU/mL for ertapenem, doripenem, meropenem and imipenem, respectively. None of the combinations were capable of reducing $10⁸$ CFU/mL of N16870 by $2 \log_{10}$ CFU/mL. Ertapenem combinations consistently displayed the least activity, whereas doripenem, meropenem and imipenem combinations had similar activities that were poorly predicted by carbapenem MICs. As doripenem, meropenem, or imipenem displayed similar pharmacodyanmics in combination, the decision of which carbapenem to use in

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combination with PMB may be based on toxicodynamic profiles if drastic discordance in MICs is not present.

Keywords

Acinetobacter baumannii; Carbapenem resistance; Polymyxins

1. Introduction

Acinetobacter baumannii is an invasive, opportunistic, Gram-negative pathogen responsible for an alarming rate of morbidity and mortality in the intensive care population [1]. Historically, carbapenems have been the most reliable treatment option for serious nosocomial A. baumannii infections. However, identification of the ideal carbapenem for combatting multidrug-resistant A. baumannii has been confounded by conflicting reports of minimum inhibitory concentration (MIC) testing and in vitro kill assessments. Whilst ertapenem reportedly possesses the least intrinsic activity [2], opposing studies have suggested that either doripenem [2], meropenem [3] or imipenem [4] is the most active against multidrug-resistant A. baumannii. Unfortunately, acquisition of carbapenem resistance mechanisms has further obscured the ideal treatment of A. baumannii.

To counter the increasing prevalence of carbapenem-resistant A. baumannii, clinicians are now forced to utilise a polymyxin [colistin or polymyxin B (PMB)] as a drug of last resort [5]. However, the emergence of colistin heteroresistance and the increasing frequency of polymyxin resistance have precipitated the search for polymyxin combinations that elicit greater bacterial killing than is possible with a polymyxin alone [6]. Enhanced activity against A. baumannii has been observed when a polymyxin is paired with a carbapenem in vitro [7], and successful use of carbapenem/polymyxin combinations has been reported clinically [8]. Although a meta-analysis of in vitro carbapenem/polymyxin killing identified meropenem and doripenem as the most likely candidates for enhancing polymyxin activity, the study results were based on rates of synergy and did not examine the rate and extent of killing for each combination [9]. It is also unknown whether the density of the A . baumannii inoculum influences selection of the optimal carbapenem. In the present study, we sought to characterise the comparative pharmacodynamics of each carbapenem in combination with PMB against carbapenem-resistant A. baumannii utilising time–kill experiments conducted at two different starting inocula.

2. Materials and methods

Two polymyxin-susceptible A. baumannii strains (N16870 and 03-149-1) were utilised in this study. Time–kill experiments were conducted over 48 h at starting inocula of 10⁶ CFU/mL and $10⁸ CFU/mL$ in cation-adjusted Mueller–Hinton broth as detailed previously [10]. Solutions of ertapenem, doripenem, meropenem and imipenem (Sigma Chemical Co., St Louis, MO) were freshly prepared on the day of each experiment. An array of five antibiotic concentrations was prepared for each carbapenem. The highest unbound maximum serum concentration (C_{max}) resulting from the largest clinical dose reported in the package insert of each carbapenem was used to standardise the concentration of each agent

to clinically achievable levels $[11-14]$. The other four concentrations used in the arrays corresponded to 0.25 \times , 0.5 \times , 1.5 \times and 2 \times the chosen C_{max} value. The following carbapenem concentrations were investigated: ertapenem at 3, 6, 12, 18 and 24 mg/L; doripenem at 5.25, 10.5, 21, 31.5 and 42 mg/L; meropenem at 12, 24, 48, 72 and 96 mg/L; and imipenem at 15, 30, 60, 90 and 120 mg/L. Bactericidal activity was defined as a $\frac{3 \log_{10} CFU/mL}{2 \log_{10} CFU/mL}$ reduction within 24 h.

Carbapenem arrays were investigated in the presence of 1.5 mg/L PMB (Sigma Chemical Co.) to approximate the average free steady-state plasma concentration (C_{ss}) of PMB achieved by a 1.5 mg/kg every 12 h regimen proposed by Sandri et al [15]. PMB alone and the highest investigational carbapenem concentration alone served to control for the independent activity of each agent. Reaction vessels were incubated in a 37 °C water-bath with constant shaking and samples were collected at 0, 1, 2, 4, 8, 24, 26, 28, 32 and 48 h, were serially diluted with saline and were plated onto Mueller–Hinton agar. MIC testing was performed on 03-149-1 and N16870 in quadruplicate following Clinical and Laboratory Standards Institute (CLSI) guidelines M07-A10. Although both strains were susceptible to PMB (MIC of 0.5 mg/L for both isolates), the respective carbapenem MICs for 03-149-1 and N16870 were both 64 mg/L for ertapenem, 32 mg/L and 16 mg/L for doripenem, 64 mg/L and 16 mg/L for meropenem, and 32 mg/L and 8 mg/L for imipenem.

The log ratio change (Eq. 1) was calculated for the ertapenem, doripenem, meropenem and imipenem combinations that achieved the greatest A. baumannii killing [16]. As the log ratio change is only sensitive to A . baumannii counts at a discrete time and does not reflect the bacterial burden throughout the 48-h experiments, the log ratio area (Eq. 2) was also calculated for the most active carbapenem combinations to provide insight into antimicrobial activity achieved over the course of the entire experiment. The log ratio change and the log ratio area of each carbapenem combination were compared with one another for both A. baumannii strains at both starting inocula to determine differences in the rate and extent of killing for each combination.

Log ratio change=
$$
log_{10} \left(\frac{CFU_{48h}}{CFU_{0h}} \right)
$$
 (1)

Log ratio area=log₁₀
$$
\left(\frac{\text{AUCFU}_{\text{drug}}}{\text{AUCFU}_{\text{control}}} \right)
$$
 (2)

where AUCFU is the area under the CFU–time curve.

3. Results

The results of the time–kill experiments at the 10^6 CFU/mL inoculum are displayed in Fig. 1. For strain 03-149-1, all four carbapenems were able to achieve a $3 \log_{10} CFU/mL$ reduction by 6 h [Fig. 1(A1–A4)]. Ertapenem combinations displayed the least activity, with

bacterial counts that began to recover by 8 h independent of the ertapenem concentration used in the combination (mean standard deviation $0.4 \log_{10} CFU/mL$). The majority of doripenem combinations also resulted in regrowth by 8 h, with a maximum reduction of 3.8 log₁₀ CFU/mL by 8 h. Meropenem and imipenem combinations achieved sustained killing up to 8 h, with maximum reductions at 8 h of 4.5 log_{10} CFU/mL for both carbapenem combinations.

Despite relatively similar MICs to strain 03-149-1, carbapenem combinations achieved more drastic killing against 10^6 CFU/mL of N16870 [Fig. 1(B1–B4)]. Ertapenem combinations again displayed the least activity, with all five combinations ascending above 10^7 CFU/mL by 32 h. Both meropenem and imipenem alone were capable of sustained killing against N16870, albeit more slowly than in combination treatments, whilst doripenem alone regrew by 24 h. Doripenem, meropenem and imipenem combinations all displayed sustained killing that was dependent on the carbapenem concentration, with maximum reductions at the limit of detection (100 CFU/mL) for all three combinations. The lowest doripenem concentration of 5.25 mg/L was also the only concentration that resulted in complete regrowth by 48 h when in combination with PMB.

In contrast to the bactericidal activity observed at 10^6 CFU/mL, the majority of carbapenem combinations were unable to substantially reduce the 10^8 CFU/mL A. baumannii burden by 24 h (Fig. 2). The reductions in 03-149-1 counts at 8 h achieved by the highest concentration combinations were 1.0, 3.2, 2.2 and 3.3 log_{10} CFU/mL for ertapenem, doripenem, meropenem and imipenem, respectively [Fig. 2(A1–A4)]. The highest concentration of doripenem (42 mg/L) in combination with PMB resulted in the most sustained killing, with a 3.2 \log_{10} CFU/mL reduction of 03-149-1 at 8 h that was maintained until 26 h.

Although killing of N16870 was more drastic compared with 03-149-1 at 10⁶ CFU/mL, at the 10^8 CFU/mL inoculum N16870 was more tolerant to the carbapenem and PMB combinations [Fig. 2(B1–B4)]. At 8 h, maximum reductions achieved by carbapenem combinations were 0.5, 1.6, 1.2 and 1.1 \log_{10} CFU/mL for ertapenem, doripenem, meropenem and imipenem, respectively. The only combination capable of sustained killing was 96 mg/L meropenem in combination with PMB, which resulted in a nadir of 6.5 log₁₀ CFU/mL at 32 h followed by subsequent regrowth. PMB alone was unable to achieve sustained killing against either strain at either inoculum.

A comparison of the maximum activities displayed by all four carbapenem combinations expressed as either the log ratio area or log ratio change at 48 h is summarised in Fig. 3. After 48 h of antimicrobial exposure, none of the carbapenem combinations achieved substantial reductions in bacterial counts against strain 03-149-1 at either inoculum [Fig. 3(A1)]; however, the log ratio areas of doripenem, meropenem and imipenem combinations were 1.1 lower than the ertapenem combination at the 10^6 CFU/mL inoculum [Fig. 3(A2)]. Against the N16870 strain, both the log ratio change and the log ratio areas of all four carbapenem combinations were comparable at the 10^8 CFU/mL inoculum, whereas the log ratio change of the most active ertapenem combination was ≥5.2 higher than the other carbapenem combinations at 10^6 CFU/mL [Fig. 3(B1)]. Similarly, ertapenem with PMB

achieved a log ratio area of −2.0 that was ≥2.7 higher than the other carbapenem combinations at 10^6 CFU/mL [Fig. 3(B2)].

4. Discussion

The ability of A. baumannii to acquire resistance mechanisms to commonly used antibiotics has made the pathogen particularly troubling in the nosocomial environment [1]. Here we sought to characterise the pharmacodynamics of four different carbapenems in combination with PMB at two different inocula to better understand the killing of antimicrobial combinations against carbapenem-resistant A . baumannii. Regardless of the inoculum, a PMB concentration of 1.5 mg/L was unable to achieve sustained killing for up to 48 h despite PMB MICs of 0.5 mg/L. When carbapenems were added to PMB, killing drastically improved, but was more pronounced at the 10^6 CFU/mL inoculum. The utility of colistin in combination with meropenem is currently being investigated in two clinical trials (NCT01732250 and NCT01597973) that will help illuminate the clinical utility of polymyxin combinations.

A previous hollow-fibre infection analysis of PMB alone against A. baumannii with susceptible PMB MICs also failed to eradicate the pathogen in vitro [17]. Despite rapid initial killing, polymyxin-resistant subpopulations amplified by over 5 log_{10} CFU/mL within 24 h of PMB exposure to allow for continued growth in the presence of PMB. In the present study, the rapid initial killing by PMB alone at the 10^6 CFU/mL inoculum was likely followed by a similar amplification of polymyxin-resistant subpopulations. When the A. baumannii was exposed to PMB and a carbapenem, PMB likely permeabilised the outer membrane of the bacterial cells and increased the accessibility of penicillin-binding proteins for the β-lactams [18].

Using the measures of log ratio area and log ratio change, doripenem, meropenem and imipenem combinations resulted in similar killing profiles against both A. baumannii strains. However, the use of only two A . baumannii isolates limits our ability to generalise the findings to other A . baumannii clinical strains. Similar to how ertapenem was previously identified as the least active agent against carbapenem-susceptible A . baumannii [2], ertapenem in combination with PMB displayed the least killing against both strains regardless of the inoculum. Interest in the utility of ertapenem against carbapenem-resistant Gram-negative organisms has risen following the reported use of ertapenem combination therapies targeted at carbapenem-resistant Klebsiella pneumoniae [19]. Although the results of the current investigation suggest that ertapenem may offer unfavourable pharmacodynamics in combination with PMB against carbapenem-resistant A. baumannii, dynamic in vitro models are needed to fully define the combinatorial pharmacodynamics of such a combination.

Not only were similar killing profiles observed for doripenem, meropenem and imipenem, but a large inoculum effect was consistently observed among the three carbapenem combinations as well. Killing was the most drastic against N16870 in both the log ratio area and log ratio change analyses at a 10⁶ CFU/mL inoculum, yet the most substantial killing at a 10^8 CFU/mL inoculum was achieved against strain 03-149-2. The magnitude of the

inoculum effect in carbapenem-resistant A . *baumannii* is therefore difficult to predict, as the attenuation in antimicrobial activity varied markedly between both strains. Given the similar declines in bacterial killing as the inoculum increased from 10^6 CFU/mL to 10^8 CFU/mL, the bacterial burden of A. baumannii may not influence whether doripenem, meropenem or imipenem is the ideal agent for polymyxin combination therapy.

Although both A. baumannii strains possessed different carbapenem MICs that varied by multiple dilutions, the MIC of each carbapenem was not able to fully predict the activity of each combination. When used in monotherapy, the prevailing dogma for β-lactams asserts that obtaining a sufficient $\% T_{>MIC}$ (percentage of the dosing interval that the plasma level exceeds the MIC of the causative pathogen) is necessary for bactericidal activity [20]. However, the ability of PMB to perturb the integrity of both the outer and inner membranes of A. baumannii and increase carbapenem permeability is not accounted for by simply using the MIC of a carbapenem to predict the activity of combination treatment [18]. At a 10^6 CFU/mL inoculum, doripenem, meropenem and imipenem combinations utilising carbapenem concentrations below the MIC achieved bactericidal activity (0% $T_{\rm MIC}$), suggesting that a concentration dependence exists during combination therapy that is not entirely accounted for by the % $T_{>MIC}$ metric.

A significant limitation of the current study is the use of static time–kill experiments that do not simulate the dynamic pharmacokinetics of carbapenems and PMB. Although doripenem, meropenem and imipenem all share a similar half-life of ca. 1 h in healthy volunteers [11,13,14], ertapenem possesses a half-life of 4 h that may confer additional bacterial killing in comparison with the other carbapenems [12]. Further work with dynamic in vitro models or in vivo animal models is needed to fully define the combinatorial pharmacodynamics of carbapenems in combination with PMB.

5. Conclusions

During A. baumannii time–kill experiments, ertapenem was the least active carbapenem in combination with PMB, whilst doripenem, meropenem and imipenem combinations displayed slight differences in their activities that were influenced by the A. baumannii strain and bacterial burden. Individual carbapenem MICs were also poorly predictive of how each carbapenem performed when in combination with PMB. Given the similar activities of doripenem, meropenem and imipenem when paired with PMB, the decision of which carbapenem to use for combination treatment may be driven by the toxicity profiles of each carbapenem if MICs are similar. Further investigations that use dynamic models to completely simulate the pharmacokinetics of carbapenems in combination with PMB are needed to better translate the findings into the clinical setting.

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Fig. 1.

Time–kill experiments of (A) strain 03-149-1 and (B) strain N16870 at a starting inoculum of 10⁶ CFU/mL. Five concentrations of ertapenem (A1/B1), doripenem (A2/B2), meropenem (A3/B3) and imipenem (A4/B4) were investigated with 1.5 mg/L polymyxin B (PolyB). The highest concentrations of each carbapenem alone and PolyB alone were investigated separately to account for the independent activity of each agent.

Fig. 2.

Time–kill experiments of (A) strain 03-149-1 and (B) strain N16870 at a starting inoculum of 10^8 CFU/mL. Five concentrations of ertapenem (A1/B1), doripenem (A2/B2), meropenem (A3/B3) and imipenem (A4/B4) were investigated with 1.5 mg/L polymyxin B (PolyB). The highest concentrations of each carbapenem alone and PolyB alone were investigated separately to account for the independent activity of each agent.

Fig. 3.

(A) Log ratio change and (B) log ratio area after 48 h of exposure to the most active combinations of polymyxin B with either ertapenem, meropenem, imipenem or doripenem for strain 03-149-1 (A1/B1) and strain N16870 (A2/B2) at two different initial inocula.