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Review Article

An Update on Technical, Interpretative and Clinical Relevance of Antimicrobial Synergy Testing Methodologies

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Abstract

Testing for antimicrobial interactions has gained popularity in the last decade due to the increasing prevalence of drug-resistant organisms and limited options for the treatment of these infections. *In vitro* combination testing provides information, on which two or more antimicrobials can be combined for a good clinical outcome. Amongst the various *in vitro* methods of drug interactions, time-kill assay (TKA), checkerboard (CB) assay and *E*-test-based methods are most commonly used. Comparative performance of these methods reveals the TKA as the most promising method to detect synergistic combinations followed by CB assay and *E*-test. Various combinations of antimicrobials have been tested to demonstrate synergistic activity. Promising results were obtained for the combinations of meropenem plus colistin and rifampicin plus colistin against *Acinetobacter baumannii*, colistin plus carbapenem and carbapenem plus fluoroquinolones against *Pseudomonas aeruginosa* and colistin/polymyxin B plus rifampicin/meropenem against *Klebsiella pneumoniae*. Antagonism was detected in only few instances. The presence of synergy or antagonism with a combination seems to correlate with minimum inhibitory concentration of the agent and molecular mechanism involved in the resistance. Further studies need to be conducted to assess the utility of *in vitro* testing to predict clinical outcome and direct therapy for drug-resistant organisms.

Keywords: Acinetobacter baumannii, antimicrobial resistance, checkerboard assay, combination testing, Klebsiella pneumoniae, Pseudomonas aeruginosa, time-kill assay

INTRODUCTION

In recent times, need for synergy testing has been driven by the following reasons: (i) necessity to extend the antimicrobial spectrum, (ii) possibility of reducing the dosage and toxicity and (iii) possibility of reducing the development of resistance. ^[1] In addition, the emergence of multidrug resistance (MDR), extensive drug resistance (XDR) and pan-drug resistance (PDR) strains, combined with the lacunae in the development of newer antimicrobial agents, has contributed to the necessity for the synergy testing between various combinations of antimicrobial agents.

The development of drug-resistant organisms is the prime cause for the increase in healthcare-associated infections, especially ventilator-associated pneumonia (VAP) and bacteraemia. Among the hospital-acquired infections (HAIs) due to Gram-negative organisms, MDR-Gram negative bacilli (GNB) infections accounted for 36.8% in a tertiary care centre in Taiwan during a 7-year period (2002–2009).^[2] Similar

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trend was seen in South America, where a tertiary care centre in Brazil recorded 3.7-fold increase in the infection rates due to MDR-GNB during 1999–2008.^[3] The development of MDR and carbapenem resistance was increasingly seen, especially for *Acinetobacter baumannii*.^[4] However, good infection control practices were able to decrease the overall HAI rates, and the trend remains unchanged for GNB-HAI contributed by carbapenem resistance organisms.^[2]

Alternative therapies or treatment strategies for such XDR and carbapenem-resistant (CR) GNBs are limited. Nevertheless, old drugs such as colistin, fosfomycin and tigecycline can be used in combination with other agents. In the past decades, the use of colistin has been restricted by the

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concerns of toxicity and problems in optimisation of dosage. Tigecycline use was hampered by its large distribution volume, leading to sub-inhibitory levels and selection of resistant strains with increase in the geometric mean of minimum inhibitory concentration (MIC).^[5] Further, the Food and Drug Administration approved the use of tigecycline only for complicated skin and soft tissue infections, intra-abdominal infections and community-acquired pneumonia. However, it was not approved for the use in VAP because of higher mortality rate. Fosfomycin was reported to have superior in vitro activity against CR-Enterobacteriaceae isolates but was restricted for the treatment of urinary tract infection (UTI). Very few reports exist on the use of fosfomycin for other systemic infections. Moreover, fosfomycin must be used in combination with other antimicrobial agents because of high rate of resistance mutation. To overcome the aforementioned concerns, it is essential to test different antimicrobial combinations including the agent to which the organism has developed resistance.

The rationale behind the choice of combination therapy is that the antimicrobials will have a synergistic effect when given together. This review summarises the various methods available to determine synergy between different antimicrobial agents and to provide scientific evidence for utility of such combinations in the clinical setting. In particular, special focus is given on *in vitro* efficacy of the combined antimicrobials against drug-resistant *A. baumannii, Klebsiella pneumoniae* and *Pseudomonas aeruginosa*.

Technical Performance of Methods for the Determination of Interactions between Antimicrobial Agents

Although many test methods are available to determine the interaction between antimicrobial agents, they were not well standardised. Interpretation criteria followed for test results are not defined and remain uncertain. The various testing methods for determining the synergistic activity of antimicrobials are discussed below.

In vitro assay

Time-kill assay

The time-kill assay (TKA) is considered as the standard reference method for the determination of synergy between antimicrobial agents. TKA determines the actual reduction in the viable count of the organism after exposure to the drug combination compared to the most active single agent at different time intervals. This is done by adding a standard inoculum in broths containing the individual antimicrobial agents and its combination. Sub-culturing is done from the broth containing antimicrobials at different time intervals and the bacterial count is done. Colony count is done at shorter time intervals e.g., every 2 h over a 24-h period for drugs having concentration-dependent activity. For drugs having time-dependent killing activity, colony count is done every 3–4 h till 24–48 h. The determination of the synergistic

action by TKA is defined as $\geq 2 \log_{10}$ CFU/ml reduction in the bacterial growth in the combination when compared to the most active single agent. However, antagonism is defined by an increase of $\geq 2 \log_{10}$ CFU/ml in the combination compared to the most active single agent. Less than 2 \log_{10} CFU/ml difference is interpreted as indifference. Bactericidal effects of the combinations are determined by a decrease of $\geq 3 \log_{10}$ CFU/ml from the initial inoculum.

Another method of interpretation of the TKA is area under the killing curve (AUKC),^[6] where instead of measuring the log₁₀ difference; the result was plotted on a graph with the log₁₀ CFU/ml value in the Y-axis and the time at the X-axis. The AUKC is calculated for single agent and combinations as well. Any statistically significant difference with P < 0.05is taken as a synergistic interaction.^[6] This method of interpretation was found to be robust with high precision and less intra-experimental variation but not widely used.

This method allows the testing of one concentration and one ratio of the antimicrobials at one time. The test has to be repeated to observe interactions at other concentrations and ratios. There is also a lack of consensus as to a standard inoculum of the organism to be used though the inoculum size varied from 1 to 5×10^5 . The reported concentration of antimicrobials tested in other studies varies from $0.125 \times \text{MIC}$ to $4 \times \text{MIC}$.^[7] When drug combinations are tested at the MIC or more than MIC concentrations, the test may be hard to interpret because inhibition of the organism by the single agent may preclude demonstration of synergy.

Some authors prefer testing of drug concentrations that are achievable in human serum when standard dosing regimens are administered.^[8] Though this strategy incorporates the pharmacokinetic (PK) property of the tested drugs, it does not implicate the concentration of drug at tissues or other sites of infection. Thus, results may not be extrapolated to particular organ system infection such as VAP where the serum concentration of the drug may not reflect the tissue concentration. The drug concentration in the *in vitro* test does not vary, while in vivo, there is a variation in the concentration and ratio of the drugs used. This depends on the PK and pharmacodynamic (PD) property of the drugs, dosing interval, strength and route of administration. The drawbacks of TKA include testing of limited antimicrobial concentrations, non-standardised inoculum size and antimicrobial concentration, static concentration of the drug, labour intensive and time-consuming.

Checkerboard assay

The checkerboard (CB) assay utilises a panel of antimicrobial combinations at different concentrations either in the macrobroth (2 ml volume) or microbroth (100 μ l volume) method. The range of tested concentrations varies from four to eight times the MIC to at least 1/8–1/16 of the MIC. It is important to include broad range of concentrations because MIC can vary depending on the method used and also within the method (a variation of one/two-fold dilution is allowed within a test system). For the interpretation of result, the fractional

inhibitory concentration (FIC) is calculated for each antibiotic at a given concentration combination by the following formula:

FIC of agent A = MIC of agent A in combination/MIC of agent A alone

The cumulative FIC is then calculated by summing up the FIC of both the agents. 'Synergy' is interpreted when the FIC index is ≤ 0.5 , 'indifference' or 'no interaction' corresponds to the FIC index >0.5–4.0 and 'antagonism' when the FIC index is >4.0.^[9]

However, in some studies, authors have defined 'partial synergy' for FIC index between >0.5 and <1 and an 'additive interaction' for FIC index of 1. Reporting of such results has to carefully considered because of the acceptance of inherent one tube dilution variation with this method and possibility of reproducibility error.^[9] This was addressed by Rand *et al.*, who reported 25% discordance with the CB method and suggested testing in at least five replicates and considering the reading only with ≥80% agreement between the replicates.^[10] Another contentious issue with CB assays is the use of different criteria to interpret the test.

E-test

E-test strips containing gradient of antimicrobial agents have been used to determine the synergistic combinations. The different methods are (i) *E*-test cross method, (ii) *E*-test fixed ratio method, (iii) *E*-test agar method and (iv) *E*-test MIC: MIC method.

E-test cross method

Mueller-Hinton agar (MHA) plate is inoculated with 0.5 McFarland matched inoculum, to which *E*-test strips are placed one over the other at 90° angle crossing at the MICs of the individual agent of the organism determined earlier [Figure 1]. After incubation for 18 h, the zone of inhibition is read and

the FIC index is calculated and interpreted as described for CB assay.^[11]

E-test fixed ratio method

In this method, MHA plates are inoculated with 0.5 McFarland matched inoculum. *E*-test strip of the first agent is placed and incubated at room temperature for 1 h to allow the antimicrobial to diffuse into the medium. After 1 h, it is removed and saved as MIC template. The *E*-test strip for the second agent is then placed directly over the imprint of the first strip [Figure 2]. The FIC index is again calculated and interpreted as described for CB assay.^[12,13]

E-test agar method

In this method, MHA plates are incorporated with $0.5 \times \text{or}$ $0.125 \times \text{MIC}$ of one agent and the *E*-test strip of the second agent is placed over the inoculated surface [Figure 3]. The MIC obtained is compared with the MIC in drug-free medium. The synergy is interpreted when there is more than three-fold reduction in MIC on the drug-incorporated medium.^[13]

E-test minimum inhibitory concentration: minimum inhibitory concentration method

In this method, one test strip is placed on the inoculated MHA plate and incubated at room temperature for 1 h to allow diffusion of the agent. After 1 h, the agar is marked adjacent to the previously determined MIC of the agent and removed. The second *E*-test strip is then placed over the imprint of the previous strip such that the mark on the agar corresponds to the MIC of the second agent [Figure 4]. The resulting ellipse of inhibition is read after 20 h of incubation at 37°C. The FIC index is calculated and interpreted as like that of CB assay.

Compared to the other commonly used methods such as TKA and CB assay, *E*-test methods are technically simpler to



Figure 1: *E*-test cross method. In this example, minimum inhibitory concentration of A is 6 μ g/ml and minimum inhibitory concentration of B is 8 μ g/ml. After combination of A and B, minimum inhibitory concentration of A is 0.094 μ g/ml and minimum inhibitory concentration of B is 0.75 μ g/ml. Σ FIC = 0.1 (synergy). FIC: Fractional inhibitory concentration

perform and reproducible. The limitations of *E*-test methods are the inability to determine interaction of more than two antimicrobial combinations and the limited gradient of antimicrobial on the paper strip. For organisms where the MIC is more than the highest concentration on the strip, difficulties may be encountered with calculation of the FIC index and may result in the false interpretations. In addition, detection of antagonistic combinations will be limited for such isolates. With the *E*-test cross method, mild degree of antagonism may not be detected because of overlapping of strips.^[11]

In vitro pharmacokinetic model

The various *in vitro* tests for the determination of antimicrobial interactions involve testing of drugs at a static concentration without any change in concentration with time. However, the *in vivo* drug concentrations and ratio keep changing with time. To better simulate these changing conditions, PK models were designed. In a single-compartment model, a glass apparatus with inlet and outlet is maintained at 37°C. Fresh media with antimicrobial agents are loaded and the media from the apparatus are withdrawn using a peristaltic pump at a constant rate mimicking the elimination kinetics of the drug and the half-life at the standard dosing regimen of the drug tested. The compartment is charged with a standard inoculum and



Figure 2: *E*-test fixed ratio method. In this example, minimum inhibitory concentration of A is 16 μ g/ml and minimum inhibitory concentration of B is 32 μ g/ml. Minimum inhibitory concentration of combination A and B is 1 μ g/ml. Σ FIC = 0.09 (synergy). FIC: Fractional inhibitory concentration

the change in organism load is compared between single agent and the combination.^[14]

In a two-compartment model, a similar central compartment as above was used with a constant volume with changing antimicrobial concentration. The compartment is connected to three or four dialyser unit which acts as the peripheral compartment. Each peripheral compartment containing 150 ml of the organism culture is exposed to a changing antimicrobial concentration similar to the central compartment. The whole system was then kept at 37°C. This method enables simultaneous testing of up to four isolates.^[15,16]

The two models mimic the *in vivo* PK property of the individual agents at the standard dose and regimen. The change in CFU/ml was compared using a standard inoculum for the single-drug administration and the combination regimen at regular time intervals. Synergism is interpreted by decrease of $\geq 2 \log_{10} \text{ CFU/ml}$ compared to the best monotherapy regimen or AUKC analysis can be used to detect synergistic interactions.^[15,16]

Hollow fibre infection model

The hollow fibre bioreactor is an important advancement in the in vitro combination testing. Currently available in vitro testing methods have a drawback of not examining time and concentration of the drug at various exposure concentrations. Hollow fibre model has an advantage of considering PK and PD parameters; thereby it mimics the *in vivo* conditions with dynamic concentration of drug over time. The bioreactor module contains thousands of filters with 200 μ in diameter. The peripheral chamber containing the bacteria is separated from the central compartment via semi-permeable membranes, which allows the flow of nutrients and other molecules in and out while retaining the bacteria. These fibres are designed in such a way that the fibre acts as barriers for the flow of contents. Drug concentration is adjusted through infusions at different intervals and by supplying fresh medium to promote dilution of the drug. By adjusting the volume of central reservoir, a state of dynamic concentration of the drug is created, without



Figure 3: *E*-test agar method. In this example, minimum inhibitory concentration of A is 12 μ g/ml and in combination 0.38 μ g/ml. Fractional inhibitory concentration of A = 0.03. Minimum inhibitory concentration of B is 96 μ g/ml and in combination is 6 μ g/ml. Fractional inhibitory concentration of B = 0.06. Σ FIC = 0.09 (synergy). FIC: Fractional inhibitory concentration

diluting the bacterial load in the peripheral compartment. Sampling is done from the peripheral compartment at different intervals to quantify the drug concentration and the bacterial count [Figure 5]. This phenomenon provides the reliable PK and PD profiles, which could be considered for clinical decision-making. Two-compartment hollow fibre infection models provide advantages over one-compartment model with respect to the variable concentration of drug exposure over time. Such *in vitro* PK/PD models are cost-effective and resource intensive. Moreover, it permits the investigation over considerable duration, which is not feasible to perform in animal models. However, this method is technically demanding and requires complex instruments and difficult to standardise.^[17]

Critical inhibitory concentration

Determination of critical inhibitory concentration (CIC) was shown to help predict in vivo synergistic effect.^[18] For the determination of CIC, a pour plate of media inoculated with the organism is prepared. Ten-millimetre holes are made and filled with combinations of the antimicrobials at different concentration ratios and at graded concentrations [Figure 6]. After incubation for 20 h at 37°C, the distance from the edge of the well till the edge of the zone of inhibition is measured (d). The square of d (d^2) was then plotted against the concentration of antibiotic at time zero (log, m₀). A straight line was obtained intercepting the log_a m_a axis, and antilog of this point of interception gives the CIC value of the combination. A lower CIC indicates a higher killing effect. Using CIC, Chan *et al.* demonstrated the synergistic activity for the combination of amikacin and piperacillin at the ratio of 70:30 for P. aeruginosa and was confirmed by TKA and in vivo mouse model.^[18]

Double disc synergy

Double disc synergy test was conventionally used for the detection of extended-spectrum beta-lactamase (ESBL) production and can also be used for the detection of synergy between antimicrobial combinations. In this method, discs containing the antimicrobials are placed 20 mm (or sum of radii of the zone of inhibition of each drug separately) apart over a lawn culture of the organism and incubated at 37°C. Synergy was indicated by an increase in the zone diameter of ≥ 2 mm compared to the single agent or bridging of the zone of inhibition [Figure 7].^[19,20] An increase of <2 mm in the zone of inhibition is classified as weak synergy, and antagonism is indicated by truncation of the zone of inhibition at the junction of the two antimicrobials. For P. aeruginosa, this method was shown to give more synergism for a combination of antimicrobials than CB assay.^[19] In addition, double disc synergy test was observed to show more synergy than other methods such as agar-based and broth-based dilution method.^[21] Despite the simplicity and easy interpretation of results, this method has not been widely used because of its qualitative nature and subjective interpretation.



Figure 4: *E*-test minimum inhibitory concentration: minimum inhibitory concentration method. In this example, minimum inhibitory concentration of A is 12 μ g/ml and minimum inhibitory concentration of B is 6 μ g/ml. Minimum inhibitory concentration of combination A and B is 0.5 μ g/ml. Σ FIC = 0.12 (synergy). FIC: Fractional inhibitory concentration



Figure 5: Hollow fibre infection model for *in vitro* antimicrobial combination testing



Figure 6: Scheme for critical inhibitory concentration determination. 'd' is the distance between the edge of the well and the edge of zone of inhibition in mm. 'd²' plotted against concentration of drug gives value of critical inhibitory concentration

Paper strip diffusion

In this method, filter paper strips soaked in different antimicrobial solutions at or above MIC were placed at right angles on the MHA plate inoculated with the test organism. Antibiotics in the filter paper strips are allowed to diffuse in the medium and are removed after several hours and the plates are incubated for 18–24 h at 37°C. Alternately, the antibiotic soaked strips can be overlaid onto un-inoculated plate media for 24 h for diffusion and the organism inoculated using a membrane transfer technique [Figure 8].^[22] The pattern of growth of the organism was interpreted as follows: indifferent (additive) effect is considered as two oval area of inhibition joining at right angles, synergism is indicated by broadening of the inhibition around the angle and antagonism is indicated by indentation or narrowing around the angle.^[23] This method provides qualitative result and has not been widely evaluated.

Multiple-combination bactericidal test

Multiple-combination bactericidal test is done in 96-well microtiter plates. Different combination of antimicrobials with a standard inoculum is added into each well and incubated for 48 h. All the non-turbid wells following incubation is



Figure 7: Double disk synergy test. (a) Synergy (bridging of zone of inhibition); (b) synergy (appearance of zone of inhibition in between agent A and B); (c) antagonism (flattening of zone of inhibition); (d) indifference/additive (no effect on zone of inhibition)

sub-cultured onto antimicrobial-free medium and checked for 99.9% killing. Antagonism is defined as growth of the organism on addition of a second antibiotic to a single agent which was bactericidal when tested alone. Though this method detects the extent of bacterial killing, the outcome is not clearly defined. Enhancement of bactericidal activity of a previous non-bactericidal drug in combination can only be made out in terms of a synergistic combination. Its use is limited to the detection of antagonistic combinations rather than a synergistic combinations for agents used for the treatment of respiratory infection in cystic fibrosis patients.^[24,25]

Overlay inoculum susceptibility disc method

In this method, solid media incorporated with half the MIC of one agent were used as an agar base over which molten antibiotic-free agar with a standard inoculum of the organism is poured to obtain an overlay inoculum layer. Similar control plates are prepared without antibiotic containing base. Antimicrobial discs are placed over the plate and incubated [Figure 9]. An increase in the inhibition zone diameter (IZD) by 19% corresponds to synergy, <19% synergy corresponds to additive effect and no variation in IZD is an indicative of indifference.^[26] Nworu and Esimone demonstrated agreement of this technique with CB with both techniques, showing synergistic interaction between ampicillin and ciprofloxacin for *Staphylococcus aureus* and *Escherichia coli*.^[27] However, this method has not been widely evaluated.

Serum bactericidal titre

For the better prediction of the PK property of the antimicrobials tested, synergy can be tested using the serum bactericidal titre (SBT) method. This method takes into account not just the drug elimination kinetics but also the protein binding and the effect of metabolic congeners of the antimicrobial agents.^[22] Here, serum from patient or volunteer is collected to get the



Figure 8: Paper strip diffusion test, (a) synergy (broadening of zone of inhibition at the angle); (b) synergy (appearance of zone of inhibition at the angle); (c) antagonism (indentation and narrowing of zone of inhibition at the angle); (d) indifference/additive (no effect in the zone of inhibition)



Figure 9: Overlay inoculum susceptibility disk method for determination of synergy. Increase in inhibition zone diameter >19% indicates synergy

peak and the trough level of the antimicrobial in single doses and in combination. The serum is serially diluted and standard inoculum of the organism inoculated. The highest dilution of the patient serum which results in 99.9% killing is designated as the SBT. The minimum bactericidal concentrations of the antimicrobials are determined in Mueller-Hinton broth and the free drug concentrations (drug-f) in serum are determined. The drug interaction is determined using the formula given below:

Drug-A-f/(MBC-A)(SBT) + Drug-B-f/(MBC-B)(SBT)

A value of ≤ 0.25 indicates synergy, 0.25–4 indicate additive effect and ≥ 4 indicate antagonism.

Robinson *et al.* compared the SBT with *in vitro* TKA and CB in patients receiving multiple antimicrobial combinations for endocarditis, osteomyelitis or severe septicaemia.^[28] Compared to CB assay, SBT detected synergy in 3/10 tests while CB detected synergy in 2/10 tests. One antagonistic combination detected by SBT was determined as synergistic by CB, while two of the synergistic combinations by SBT were determined as antagonistic by CB. TKA detected more number of synergy than either SBT or CB (6/10 at $0.5 \times$ MBC and 5/10 at $1 \times$ MBC). There was no concordance among the three methods when strict definitions are used. However, for four additive combinations tested by SBT, results of synergy or indifference were achieved in the TKA and CB.

Technical difficulties encountered with SBT include difficulty in measuring the drug-f concentration and the need to compare SBT following removal of the antimicrobials from the sample to exclude bactericidal effect due to complement or other inhibitors in the sample.

In vivo models

In vivo studies are essential for the translation of *in vitro* combination testing data to clinical trials for implementation in the clinical setting. The *in vitro* methods does not consider the following factors: pharmacokinetics of the antimicrobials in combination, difference in the route of delivery, humoral and cellular immunity of the host, site of involvement, inoculum of the organism at the infected site, virulence factors of the organism and continuous changing concentration

of the antimicrobials as single agent and relative to one another (changing ratio of drug concentration).^[29] Animal model studies may confirm or contradict *in vitro* findings based on the PD properties of the antimicrobial agents as well as the host immune response. In addition, *in vivo* models are necessary to determine the optimum dosing strategy.^[30]

To better simulate the *in vivo* conditions, various experimental models of infection have been used. Synergy between different drug combinations is determined by statistically significant survival rate or organism load reduction in the combination therapy compared to the most active single-drug regimen. However, using these criteria, additive effect cannot be differentiated from a synergistic activity. Fantin and Carbon suggested to define *in vivo* synergy as 'a significant bactericidal effect of the drug combination in comparison with the sum of the bactericidal effect of each agent alone in comparison with the effect in untreated animal'.^[30]

For the mouse pneumonia model, the organisms are inoculated intranasal and kept in hyperoxic condition.^[31] For a systemic infection model, organisms are inoculated intraperitoneally in neutropenic mice.^[18] Due to ethical and technical considerations, invertebrate models of infections have become an attractive option to study pathogenesis. In vivo models involving larva of Galleria mellonella (wax moth) has been used for the study of antimicrobial efficacy as infection in this model is amenable to treatment.^[32] Hornsey and Wareham demonstrated combination of colistin and vancomycin to be highly effective (>90%) in protecting the larva against infection with both a susceptible and a bla_{OXA-23} producing-resistant strain of A. baumannii which showed synergism in vitro by CB assay.[33] On the other hand, combination of colistin and teicoplanin was more effective in controlling infection by the susceptible strain than the resistant strain. Monotherapy with vancomycin also showed in vivo activity. This has been postulated to be due to ability of vancomycin to enhance the immune response in the larva. O'Hara et al. reported a significant improvement in the survival of larva using combinations of doripenem and vancomycin and triple combination of colistin, doripenem and vancomycin in colistin-resistant A. baumannii infection. In the same experiment, TKA did not demonstrate synergy with doripenem and vancomycin combination.[34] Hornsey et al. also demonstrated the synergistic activity of telavancin and colistin against A. baumannii.^[35] In spite of its good turnaround time (96 h), simplicity of procedures and clearly defined endpoints, results obtained in the invertebrate model need to be confirmed in vertebrate model as this model may not reflect the exact mammalian in vivo milieu.

Comparison of different methods of detection of synergy

Table 1 summarises the relative merits and demerits of the different methods for determination of synergy.

Table 2 gives the comparison of commonly used methods of determination of synergy for *A. baumannii*, *P. aeruginosa* and *K. pneumoniae*. Synergy is detected most often by TKA followed by CB. *E*-test detected least of synergistic interactions.

Method	Advantage	Disadvantage
In vitro methods		
E-test	Easy to perform	Different methods used
		Detects less synergy
		Restricted to two drugs combination only
Checkerboard	Relatively easy	Different methods for interpretation
	Multiple concentrations tested	Intra-assay variation
		Static concentration of drugs
Time-kill assay	Reference assay	Limited concentrations tested
	Measures bactericidal effect of combination (both	Time-consuming and labour intensive
	rate and extent of killing)	Static concentration of drugs
MCBT	Detects bactericidal effect	Only useful for determination of antagonistic combinations
Double disc synergy	Simple and easy	Qualitative measure
Paper strip diffusion	Simple and easy	Qualitative measure
		Not widely evaluated
OISDM	Relatively easy	Not widely evaluated
CIC	Detects synergism at a particular ratio of the	Technical expertise required for performance and
	combination	interpretation of assay
In vivo PK	Takes PK of drugs in consideration	Technically complex
		May not necessarily reflect in vivo conditions
SBT	Takes PK of drug in consideration	Technically complex
		Need for volunteers or patients
Hollow fibre infection	Takes PK of drug in consideration	Technically complex
model		
In vivo methods		
Mouse models	Takes PK and PD properties of the drugs into account	PK and PD in humans may be different from that of mouse
Larva of Galleria mellonella	Simpler model than mouse	Invertebrate model may not mimic conditions in mammals
		Needs confirmation in vertebrate model

Table 1: Relative merits and demerits of methods of determination of antimicrobial interaction

MCBT: Multiple combination bactericidal test, OISDM: Overlay inoculum susceptibility disc method, CIC: Critical inhibitory concentration, SBT: Serum bactericidal titre, PK: Pharmacokinetic, PD: Pharmacodynamic

Concordance rates between TKA, CB and *E*-test observed were as follows: 33.3%-100% between CB and TKA; 60%-80.6%between *E*-test and TKA; 83%-84.4% between CB and *E*-test; 52%-75% with all three methods. In spite of more conservative interpretation of synergy, Wareham and Wareham have argued that *E*-test methods may be clinically relevant by giving rapid results of combinations with marked synergy only.^[36] Clinical relevance of combinations with only weak synergistic interactions missed by the *E*-test method needs to be studied further by *in vivo* milieu to give evidence for the recommendation of *E*-test for rapid reporting of synergistic combinations.

In spite of the availability of different methods to determine interactions between different antimicrobial agents, lack of standardisation has hampered reliable comparison and compilation of results of different studies. It is also difficult to assess the difference in results due to strain difference, and thus, the reproducibility or clinical efficacy of the combination might vary.

In vitro Synergy of Antimicrobial Combinations in Multidrug-resistant -Gram-negative Bacilli

Emergence of MDR, carbapenem-resistant organism and PDR GNB has triggered the search for synergistic combinations of

antimicrobials in the last decade. *A. baumannii, P. aeruginosa* and *K. pneumoniae* are the most commonly studied organisms because of their major role as nosocomial pathogen with frequent drug resistance.

Acinetobacter baumannii

Among commonly studied drug combinations, colistin with either meropenem or rifampicin shows high synergy rates of 96.3% and 94.2% by TKA. Imipenem plus sulbactam/colistin shows moderate rate of synergy (66.6% and 59%, respectively, by TKA). There is a paucity of data to allow adequate comparison of differences among the different carbapenems. In general, all the carbapenems gave a wide range of synergy levels at different combinations which may be accounted by strain difference. Antagonism was noted with combinations sulbactam plus colistin/meropenem; colistin plus meropenem and polymyxin B plus meropenem in few studies. The significance of these observations needs to be further validated by *in vivo* model testing. Table 3 summarises *in vitro* studies done on sulbactam-based combinations and Table 4 summarises polymyxin-based *in vitro* studies done for *A. baumannii*.

Pseudomonas aeruginosa

Available data are very limited to give meaningful interpretation of the combinations tested. However, combinations of colistin plus carbapenem and carbapenem plus higher fluoroquinolone

Klebsiella pneumoniae,	Pseudomo	nas aeruginosa ai	nd Acinetoba	cter spp.		
Reference	Number of isolates	Drug combination	Percentage	ntibiotic combinations	Concordance	
			CB (%)	TKA (%)	E-test* (method) (%)	
			ATCC strai	ns		
White <i>et al.</i> , 1996 ^[11]	4	Czd/Cefe + Cipro/ Tobra	4/16 (25)	3/16-4/16 (18.7-25)	2/16 (12.5) (cross)	<i>E</i> -test and TKA - 63%-75% CB and TKA - 44%-88% All three - 75%
			Acinetobacter	r spp.		
Bajaksouzian et al., 1997 ^[51]	15	Ak + Levo/Oflo/ Cipro	0/45 (0)	35/45 (77.7)	-	-
Bonapace <i>et al.</i> , 2000 ^[52]	10	Pipe, Cefe, Tobra, Trova	12/40 (30)	15/30 (50)	0% (cross)	TKA and <i>E</i> -test - 72% TKA and CB - 51%
Pankey and Ashcraft, 2009[53]	8	PB + Mero	-	8/8 (100)	5/8 (62.5%) (MIC: MIC)	62.5%
Gordon et al., 2010 ^[54]	6	Col + Van	4/6 (66.6)	4/5 (80)	6/6 (100%) (E-test-agar)	-
Sheng et al., 2011 ^[40]	17	Imi + Ak/Cipro/Col/ Tige/Amp-sul	32/85 (37.6)	62/85 (72.9)	-	-
Sopirala et al., 2010 ^[13]	8	Tige, Col, Imi, Ak	4/40 (10)	14/20 (70)	16/32 (50) (E-test-agar)	CB and <i>E</i> -test - 84.4%
Tan <i>et al.</i> , 2011 ^[55]	16	PB, Rif, Tige	8/48 (17)	19/48 (40)	1/48 (2%) (cross)	All three - 52% TKA and <i>E</i> -test - 60% E-test and CB - 83%
Santimaleeworagun <i>et al.</i> , 2011 ^[56]	8	Sul + Fos	6/8 (75)	6/8 (75)	-	100%
Vidaillac et al., 2012[57]	4	Col + Van/SXT	12/12 (100)	12/12 (100)	-	100%
Principe <i>et al.</i> , 2013 ^[38]	22	Dori + Tige/Col/Ak/ Amp-sul/Rif	13/22 (54.2)	8/22 (36.4)	-	-
Galani <i>et al.</i> , 2014 ^[58]	10 COL-S 4 COL-R	Col + Dapto		16/30 isolate- concentration combination (53.3)	-	-
García-Salguero <i>et al.</i> , 2015 ^[59]	10	Ak + Imi/Mero/Fos/ Col/Tige/Plaz + Imi/ Mero/Col/Fos/Tige	33/100 (33)	9/64 (14)	-	-
Park et al., 2016[60]	69	Col, Dori, Tige	-	75/207 (36.2)	-	-
Hong et al., 2016 ^[61]	41	Col + Mero/Rif/Imi/ Doxy/Tige Tige + Mero/Imi	-	-	99/287 (34.4) (MIC: MIC)	-
Bae <i>et al.</i> , 2016 ^[62]	9	Col + Tige/Azi/Ak/ SXT/Amp-sul/Cefe/ Azt/Mero/Teico/ Van/Rif	38/99 (38.3)	-	-	-
Nepka et al., 2016 ^[63]	6	Col + SXT	-	6/6 (100)	-	-
Büyük <i>et al.</i> , 2017 ^[64]	15	Col + Mero/Rif/ Cipro/Moxi/Ami Tige + Mero/Rif/ Cipro/Moxi/Ak/Col	90/165 (54.5)	-	-	-
			P. aerugino	sa		
Visalli et.al., 1998[65]	12	Levo + Cefe/Czd/ Genta/Mero	2/48 (4.1)	34/48 (70.8)	-	33.3%
Di Bonaventura et al., 2004[66]	20	Grepa/Levo + Ctr/Ctx	35/80 (43.8)	-	23/80 (31.3)	71.2%
Pankey and Ashcraft, 2005 ^[67]	31	Cipro + Gati	-	13/31 (42)	6/31 (19) (MIC: MIC)	65%
Vidaillac et al., 2012 ^[57]	4	Col + Van/SXT	No synergism	2/12 (16.6)	-	83.3%
			K. pneumon	iae		
Pankey and Ashcraft.,	14	PB + Mero	-	9/14 (64)	6/14 (43) (MIC: MIC)	79%
2011 ^[68]		PB + Rif	-	14/14 (100)	3/14 (21)	21%

Table 2: Comparison of different methods for determination of synergy with different antibiotic combinations for *Klebsiella pneumoniae*, *Pseudomonas aeruginosa* and *Acinetobacter* spp.

Contd...

Reference	Number of isolates	Drug combination	Percentage s	ynergy of isolate/a tested	ntibiotic combinations	Concordance
			CB (%)	TKA (%)	E-test* (method) (%)	
Vidaillac et al., 2012[57]	4	Col + Vanco/SXT	3/12 (25)	3/12 (25)	-	100%
Pankey et al., 2013 ^[69]	31	PB + Mero	-	23/31 (74)	17/31 (54.8) (MIC: MIC)	80.6%
Gaibani <i>et al.</i> , 2014 ^[70]	8	Col + Rif Col + Tig Csol + Mero Rif + Tig Rif + Mero Col + Tec	3/8-37.5% (Col + Mero) 6/8-75% (Col + Tig) 8/8-100% (Col + Rif) 0/8-0% (Col + Tec)	8/8-100% (Col + Rif)	8/8-100 (Col+Rif) 2/8-25% (Col+Tig) 1/8-12.5% (Col+Mero) 0/8-0% (Rif+Tig) 0/8-0% (Rif+Mero) 0/8-0% (Col+Tec)	-

**E*-test method showing highest rate of synergy when more than one *E*-test method evaluated. CB: Checker board, TKA: Time-kill assay, Col: Colistin, Cefe: Cefepime, Cipro: Ciprofloxacin, Tobra: Tobramycin, Ak: Amikacin, Levo: Levofloxacin, Oflo: Ofloxacin, Pipe: Piperacillin, Trova: Trovafloxacin, PB: Polymyxin, Mero: Meropenem, Van: Vancomycin, Imi: Imipenem, Tige: Tigecycline, Amp-sul: Ampicillin/sulbactam, Rif: Rifampicin, Sul: Sulbactam, Fos: Fosfomycin, SXT: Trimethoprim/sulphamethoxazole, Dori: Doripenem, Dapto: Daptomycin. Plaz: Plazomicin, Doxy: doxycycline, Azi: Azithromycin, Czd: Ceftazidime, Azt: Aztreonam, Teico: Teicoplanin, Van: Vancomycin, Moxi: Moxifloxacin, Czd: Ceftazidime, Genta: Gentamicin, Ctr: Ceftriaxone, Ctx: Cefotaxime, Gati: Gatifloxacin, Tec: Teicoplanin, MIC: Minimum inhibitory concentration, COL-R: Colistin resistant, *P. aeruginosa: Pseudomonas aeruginosa, K. pneumoniae: Klebsiella pneumoniae*, COL-S: Colistin-susceptible

such as gatifloxacin and levofloxacin seem promising. Except for the combination of colistin with vancomycin and sulphonamides, antagonism was not seen in any of the combinations. Table 5 summarises *in vitro* studies done on antimicrobial combinations in *P. aeruginosa*.

Klebsiella pneumoniae

Combinations of polymyxin B/colistin plus rifampicin/ meropenem give promising results. It may be noted that antagonism was detected in combinations of colistin plus ertapenem/imipenem and was found to be correlating with the high MIC of colistin. This needs to be studied further with characterisation of the isolates to understand the underlying mechanism. Table 6 summarises *in vitro* studies done on antimicrobial combinations in *K. pneumoniae*.

Table 7 summarises the most commonly studied combinations of antimicrobials for *A. baumannii*, *P. aeruginosa* and *K. pneumoniae*. Combined rates were calculated from total number of synergy observed in different studies against the total number of isolates studied.

CORRELATION OF SYNERGISM WITH OTHER FACTORS Synergy in relation to minimum inhibitory concentration value

There seems to be some degree of relationship between MIC of the combination of antimicrobials tested against a particular organism. Some studies show more synergy in isolates with higher MIC, while other studies have reported contrary findings [Table 8].

For *A. baumannii*, combinations of sulbactam plus meropenem/ doripenem and doripenem plus colistin/tigecycline/amikacin/ rifampicin exhibited more synergy with isolates with higher MIC for either sulbactam or doripenem.^[37,38] However, the actual MIC for the agents seems to have an effect on the level of synergy. Lee *et al.* demonstrated synergism for isolates with moderately high MIC of about 16 µg/ml for subactam and 64 µg/ml for meropenem. In contrast, no synergism was noted for isolates with very high MIC of about 128 µg/ml and 256 µg/ml for subactam and meropenem, respectively.^[39] Combination of colistin and rifampicin also showed synergy for isolates with higher MIC for rifampicin,^[8] whereas a combination of ampicillin-subactam plus colistin/imipenem and imipenem plus amikacin/colistin/tigecycline showed more synergy with isolates which are colistin susceptible or with lower MIC for imipenem.^[40,41]

For *K. pneumoniae*, the combination of doripenem plus colistin showed higher synergy with isolates having high colistin MIC, whereas Clancy *et al.* reported higher synergy with isolates having low doripenem MIC.^[42,43] Combination of colistin and imipenem however showed more synergy in isolates with low colistin MIC, with antagonism detected at high MIC.^[44]

The variations observed between the tests may be due to the difference in the strain, methodology and geographical area. In particular, the mechanisms of resistance in these isolates were not fully characterised. The question of presence of synergism or antagonism as a function of MIC value for each agent needs to be investigated further to use MIC as a predictor for success of combination therapy. Henceforth, studies must be carried out to decipher the MIC value of individual agents, which is likely to yield synergism or antagonism for a particular combination.

Synergism as a correlate of molecular mechanism of resistance

Another aspect of synergy testing in resistant isolates is its correlation with a particular resistance mechanism involved. Table 9 gives the correlation of synergy with antimicrobial combinations. Very few studies have further investigated

Table 3: Sulbactam-based in vitro combination study for Acinetobacter baumannii								
Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism		
СВ	Ozseven et al. 2012 ^[71]	Turkey	34 CRAB	Amp/Sul + Imi	88.2	Nil		
				Cefe/Sul + Imi	70.6			
				Amp/Sul + Mero	94.1			
				Cefe/Sul + Mero	8.8			
	Kiffer <i>et al.</i> , 2005 ^[37]	Brazil	48	Sul + Mero	29.2	6.2		
	Ji et al., 2013 ^[72]	China	40 IMI-S	Sul + Imi/Mero/Cefe	20-27.5 in 40 S	Nil		
			40 IMI-R		7.5-25 in 40 R			
	Pongpech <i>et al.</i> , 2010 ^[73]	Thailand	30 MDR	Sul + Mero	70	6.7% in Sul +		
				Sul + Col	53.3	Col		
				Sul + Mero + Col	96.7			
	Sheng <i>et al</i> . 2011 ^[40]	Tiwan	12 CRAB	Amp/Sul + Imi	16	Nil		
	Lee <i>et al.</i> 2007 ^[49]	Taiwan	4 MDR CRAB	Sul + Imi	0	Nil		
				Sul + Mero	0			
	Ni <i>et al.</i> , 2013 ^[74]	China	70 MDR	Sul + Tige	64.4	Nil		
	Principe <i>et al.</i> , 2013 ^[38]	Italy	22 MDR	Amp/Sul + Dori	9	Nil		
	Pei et al., 2012 ^[75]	China	53 CRAB	Cefe/Sul + Mino	73.5	Nil		
	Tong et al., 2006 ^[76]	China	23 CRAB	Sul + Cefe	33.3	Nil		
	Santimaleeworagun et al., 2011 ^[56]	Thailand	8 CRAB	Sul + Fos	75	Nil		
	Turk Dagi et al., 2014 ^[77]	Turkey	40 CRAB	Sul + Imi/Mero/Cefe	45.8	Nil		
	Laishram et al., 2016 ^[78]	India	50 CRAB	Sul + Mero/Col	34	Nil		
	Marie et al., 2015 ^[79]	Riyadh	54 MDR	Sul + Mero/Col	49	Nil		
E-test	Kempf et al., 2012 ^[80]	France	1 COL-R	Sul + Col	100	Nil		
	Cikman et al., 2013 ^[41]	Turkey	33 IMI-R	Sul + Col	45.5	27.3		
	Kiratisin et al., 2010[81]	Thailand	40	Cefe/Sul + Dori/Imi/Mero	17.5-32.5	Nil		
	Marie et al., 2015 ^[79]	Riyadh	54 MDR	Sul + Mero/Col	42.5	Nil		
TKA	Ko et al., 2004 ^[82]	Tiwan	1 MDR	Sul + Mero	100	Nil		
	Sheng et al., 2011 ^[40]	Tiwan	12 CRAB	Amp/Sul + Imi	42	Nil		
	Song et al., 2007 ^[83]	Korea	8 IMI-R	Sul + Imi	87.5	Nil		
	Choi et al., 2004 ^[84]	Korea	2 IMI-R	Sul + Imi	100	Nil		
			2 IMI-S					
	Tripodi et al., 2007[8]	Italy	9 MDR CRAB	Amp/Sul + Rif	100	Nil		
	Principe <i>et al.</i> , 2013 ^[38]	Italy	22 MDR	Amp/Sul + Dori	0	Nil		
	Tatman-Otkun et al., 2004[85]	Turkey	8 MDR	Amp/Sul + Tobra	50	Nil		
	Santimaleeworagun et al., 2011 ^[56]	Thailand	8 CRAB	Sul + Fos	75	Nil		
	Laishram et al., 2016 ^[78]	India	50 CRAB	Sul + Mero/Col	40	Nil		

CB: Checker board, TKA: Time-kill assay, CRAB: Carbapenem-resistant *Acinetobacter baumannii*, IMI-S: Imipenem susceptible, IMI-R: Imipenem resistant, Amp-sul: Ampicillin/sulbactam, Imi: Imipenem, Cefe: Cefepime, Mero: Meropenem, Sul: Sulbactam, Dor: Doripenem, Fos: Fosfomycin, Col: Colistin, Rif: Rifampicin, COL-R: Colistin resistant, Mino: Minocyline

the resistance mechanism for the study isolates, for which combination testing has been done. In case of *K. pneumoniae*, studies have reported the role of porin channels in determining synergism of the combinations being tested. Similarly, *K. pneumoniae*-producing *bla*_{NDM} carbapenemase alone showed significantly more synergy than isolates producing *bla*_{OXA-48}-like carbapenemases.^[78] Such correlations with the specific resistance mechanism involved might help predict synergism for a particular combination of antimicrobials for treatment. Thus, determining molecular mechanisms would help direct combination therapy to improve therapeutic success.^[45]

In vitro Synergy as a Predictor of Clinical Response

The likelihood of the findings of the *in vitro* synergy studies to be translated into clinical efficacy still remains debatable. The

classical example of *in vitro* synergy between aminoglycoside and beta-lactam agents has not stood the test of time. Studies have reported no clinical benefit of combination of beta-lactam plus aminoglycoside combination for Gram-negative infection either in the neutropenic or in the non-neutropenic host.^[46,47] Combination therapy may result in adverse effects of nephrotoxicity.

Despite issues of toxicity, combination therapy is the only strategy available for treating infections due to PDR organisms. Very few studies have documented the clinical outcome with combination therapy supported by *in vitro* synergy. Biancofiore *et al.* reported successful treatment of multifocal infection of MDR *A. baumannii* in a 16-year-old female with a combination of colistin, rifampicin and meropenem after synergism between the combinations was proved by CB assay.^[48] Lee *et al.*

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Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism
СВ	Timurkaynak et al., 2006 ^[86]	Turkey	5 MDR	Col + Rif	80	Nil
				Col + Mero	60	
				Col + Azi	60	
	Biancofiore et al., 2007 ^[48]	Italy	1 CRAB	Col + Rif	100	Nil
				Col + Mero	0 (par)	
	Guelfi et al., 2008 ^[87]	Brazil	10	PB + Mero	0 (par)	Nil
	Arroyo <i>et al.</i> , 2009 ^[88]	Spain	35	Col + Tige	0	Nil
	Pongpech <i>et al.</i> , 2010 ^[73]	Thailand	30 MDR	Col + Imi	100	6.7 in Col +
				Col + Mero	73.3	Mero
	Sopirala <i>et al.</i> , 2010 ^[13]	USA	8 PDR	Col + Tige	0	Nil
				Col + Imi	25	
	Gordon <i>et al.</i> , 2010 ^[54]	UK	6	Col + Van	66.6	Nil
	Sheng et al., 2011 ^[40]	Tiwan	12 CRAB	Col + Imi	42	Nil
	Santimaleeworagun et al., 2011[56]	Thailand	8 CRAB	Col + Imi	0	Nil
				Col + Fos	12.5	
	Tan <i>et al.</i> , 2011 ^[55]	Singapore	16	PB + Rif	18.75	Nil
				PB + Tige	12.5	
	Ozseven <i>et al.</i> , 2012 ^[71]	Turkey	34 CRAB	PB + Imi	38.2	Nil
				PB + Mero	2.9	
	Ni et al., 2013 ^[74]	China	70 MDR	Col + Tige	24.3	Nil
	Principe <i>et al.</i> , 2013 ^[38]	Italy	22	Col + Dori	36	Nil
	O'Hara et al., 2013 ^[34]	USA	3 COL-R	Col + Dori	66.6	Nil
				Col + Van	100	
	Clock et al., 2013 ^[89]	USA	XDR	PB + Dori	2	Nil
				PB + Dori + Rif	10	
				PB + Dori + Tige	2	
	Vidaillac <i>et al.</i> , 2012 ^[57]	France	4	Col + Van	100	Nil
				Col + Tmp	100	
				Col + Cotri	100	
E-test	Wareham and Bean, 2006 ^[36]	UK	5 CRAB	Col + Imi	20	Nil
	Tan <i>et al.</i> , 2007 ^[90]	Singapore	13 CRAB	Col + Mino	0	Nil
	Pankey and Ashcraft, 2009 ^[53]	USA	8 CRAB	PB + Mero	62.5	Nil
	Shields et al., 2011 ^[91]	USA				
	Sopirala <i>et al.</i> , 2010 ^[13]	USA	8 PDR	Col + Tige	0	Nil
				Col + Imi	100	
	Gordon <i>et al.</i> , 2010 ^[54]	UK	6	Col + Van	100	Nil
	Tan <i>et al.</i> , 2011 ^[55]	Singapore	16	PB + Rif	6.2	6.2 in PB +
				PB + Tige	0	Tige
	Nastro et al., 2014 ^[50]	Argentina	4 COL-R	Col + Rif	100	Nil
	Miyasaki et al., 2012 ^[92]	USA	20 MDR	Col + Doxy	10	Nil
				Col + Imi	5	
				Col + Rif	5	
TKA	Yoon <i>et al.</i> , 2004 ^[93]	USA	8 IMI R	PB + Imi	87.5	Nil
				PB + Rif	87.5	
				PB + Imi + Rif	100	
	Tripodi et al., 2007 ^[8]	Italy	9 CRAB	Col + Rif	77	Nil
				Col + Imi	0	
	Song <i>et al.</i> , 2007 ^[83]	Korea	8 CRAB	Col + Rif	100	Nil
	Tan <i>et al.</i> , 2007 ^[90]	Singapore	13 CRAB	Col + Mino	92	Nil
	Pankuch et al., 2008 ^[94]	Germany	51	Col + Mero	96	Nil
	Pankey and Ashcraft, 2009 ^[53]	USA	8 CRAB	PB + Mero	100	Nil
	Rodriguez et al., 2010 ^[95]	Argentina	14 MDR	Col + Imi	100	Nil
				Col + Rif	100	

Contd...

Table 4:	Table 4: Contd								
Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism			
	Sopirala <i>et al.</i> , 2010 ^[13]	USA	8 PDR	Col + Imi	37	Nil			
	Pankuch et al., 2010 ^[96]	USA	25	Col + Dori	100	Nil			
	Liang et al., 2011[97]	China	4 CRAB	Col + Mero	100	Nil			
				Col + Mino	100				
				Col + Rif	100				
	Sheng et al., 2011 ^[40]	Tiwan	12 CRAB	Col + Imi	75	Nil			
	Tan et al., 2011 ^[55]	Singapore	16	PB + Rif	56.2	Nil			
				PB + Tige	43.75				
	Peck et al., 2012 ^[98]	Korea	6 CRAB	Col + Imi	50 at 0.5 × MIC	Nil			
				Col + Rif	100 at $1 \times MIC$				
				Col + Tige	33 at 0.5 × MIC				
					100 at $1 \times MIC$				
					67 at 0.5 × MIC				
					100 at $1 \times MIC$				
	Vidaillac <i>et al.</i> , 2012 ^[57]	France	4	Col + Van	100	Nil			
				Col + Tmp	100				
				Col + Cotri	100				
	Gordon et al., 2010 ^[54]	UK	6	Col + Van	83.3	Nil			

CB: Checker board, TKA: Time-kill assay, MDR: Multidrug resistant, CRAB: Carbapenem-resistant *Acinetobacter baumannii*, PDR: Pan-drug resistant, COL-R: Colistin resistant, IMI R: Imipenem resistant, Col: Colistin, Rif: Rifampicin, Mero: Meropenem, Azi: Azithromycin, PB: Polymyxin, Tige: Tigecycline, Imi: Imipenem, Van: Vancomycin, Fos: Fosfomycin, Dori: Doripenem, MIC: Minimum inhibitory concentration, Mino: Minocyline, TMP: Trimethoprim, Cotri: Trimethoprim/sulfamethoxazole

Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism
СВ	Santos et al., 2013 ^[99]	Brazil	4 (2 MDR)	Gen + Imi	25	Nil
				Gen + Pip/Tazo	50	
				Gen + Cefe	25	
				Gen + Czd	25	
				Gen + Cipro	0	
				Tobra + Imi	75	
				Tobra + Pip/Tazo	100	
				Tobra + Czd	75	
				Tobra + Cefe	50	
				Tobra + Cipro	25	
				Fos + Imi	100	
				Fos + Pip/Tazo	100	
				Fos + Cefe	25	
				Fos + Czd	75	
				Fos + Cipro	50	
				Fos + Tobra	50	
				Rif + Imi	75	
				Rif + Pip/Tazo	25	
				Rif + Cefe	25	
				Rif + Czd	50	
				Rif + Cipro	0	
				Rif + Tobra	75	
	Mitsugui et al., 2011[100]	Brazil	34	PB/Col + Czd/Cefe/Pip/Tazo	0	Nil
	Dundar and Otkun, 2010 ^[101]	Turkey	12 MDR	Czd + Tobra	67	Nil
				Pip/Tazo + Tobra	50	
				Cipro + Tobra	0	
				Imi + Tobra	0	
				Imi + Cipro	0	

Contd...

Table 5:	Contd					
Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism
			13 S	Czd + Tobra	31	Nil
				Pip/Tazo + Tobra	46	
				Cipro + Tobra	0	
				Imi + Tobra	15	
				Imi + Cipro	8	
	Aoki <i>et al.</i> , 2009 ^[31]	Japan	7 MDR	Col + Imi	28.5	Nil
				Col + Rif	100	
				Col + Cipro	14.2	
				Col + Ak/Tobra/Pipe/Czd		
	Piccoli <i>et al.</i> , 2005 ^[102]	Italy	102	Czd + Levo	69.6	6.8
				Czd + Ak	79.4	8.8
	Dawis <i>et al.</i> , $2003^{[103]}$	USA	10	Gati + Cefe	60	Nil
				Gati + Mero	70	
				Gati + Pipe	50	
				Gati + Genta	60	
	Visalli et al., 1998 ^[65]	USA	124	Levo + Cefe	7.2	Nil
				Levo + Czd	6.4	
				Levo + Genta	0.8	
				Levo + Mero	5.6	
	Visalli et al., 1997 ^[104]	USA	60	Trova + Czd	28.3	Nil
				Trova + Ak	8.3	
				Trova + Imi	23.3	
	Tessier and Quentin, 1997 ^[105]	France	40	Fos + Cipro	15	Nil
				Fos + Ak	7.5	
				Fos + Imi	0	
				Fos + Czd	0	
E-test	He <i>et al.</i> , 2012 ^[106]	USA	100 CR	Dori + Ak	20	Nil
				Dori + Col	3	
				Dori + Levo	9	
	Samonis <i>et al.</i> , 2012 ^[107]	Greece	15 MDR	Fos + Imi	46.7	Nil
				Fos + Mero	53.3	
				Fos + Dori	73.3	
				Fos + Col	13.3	
				Fos + Netil	13.3	
				Fos + Tige	13.3	
	Sueke <i>et al.</i> , 2010 ^[108]	UK	10	Mero + Cipro	10	Nil
	Balke et al., 2006 ^[109]	Germany	163 MDR	Czd + Tobra	28.8	Nil
				Mero + Tobra	19	
	Pankey and Ashcraft, 2005 ^[67]	USA	31 Cipro resistant	Gati + Cipro	19	Nil
TKA	Vidaillac <i>et al.</i> , 2012 ^[57]	France	4	Col + Van	0	100
				Col + Tmp	0	75
				Col + Cotri	25 (seen in COL-R)	75
	Pankuch et al., 2008 ^[94]	USA	51	Mero + Cipro	66.6	Nil
				Mero + Col	25.4	
	Pankey and Ashcraft, 2005 ^[67]	USA	31 Cipro resistant	Gati + Cipro	42	Nil
	Giamarellos-Bourboulis <i>et al.</i> , 2003 ^[110]	Greece	28 MDR	Col + Rif	11.8	Nil
	Ermertcan <i>et al.</i> , 2001 ^[111]	Turkey	18	Mero + Cipro	22 at 0.5 × MIC 61 at 1 × MIC	Nil
	Gradelski et al., 2001 ^[112]	USA	8	Gati + Cefe	37.5	Nil
	·			Gati + Czd	62.5	
				Gati + Cfper	62.5	

Table 5: Contd								
Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism		
				Gati + Imi	75			
				Gati + Ak	37.5			
	Visalli et al., 1998 ^[65]	USA	12	Levo + Cefe	83.3	Nil		
				Levo + Czd	75			
				Levo + Genta	33.3			
				Levo + Mero	91.6			
	Visalli et al., 1997 ^[104]	USA	3	Trova + Czd	100	Nil		
				Trova + Ak	100			
				Trova + Imi	100			

MDR: Multidrug resistant, CR: Carbapenem resistant, Cipro: Ciprofloxacin, Imi: Imipenem, pip/tazo: Piperacillin/tazobactam, Cefe: Cefepime, Czd: Ceftazidime, Tobra: Tobramycin, Fos: Fosfomycin, Rif: Rifampicin, PB: Polymyxin, Col: Colistin, Ak: Amikacin, Levo: Levofloxacin, Gati: Gatifloxacin, Genta: Gentamicin, Trova: Trovafloxacin, Mero: Meropenem, Dori: Doripenem, COL-R: Colistin resistant, MIC: Minimum inhibitory concentration, S: Susceptible, Netil: Netilmicin, TMP: Trimethoprim, Cotri: Trimethoprim/sulfamethoxazole

						_
Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism
СВ	Tascini et al., 2013[113]	Italy	13 Col R KPC	Col + Rif	100	Nil
				Col + Gen	38.5	
				Col + Mero	38.5	
				Col + Imi	38.5	
				Col + Tige	38.5	
				Tige + Gen	0	
				Tige + Mero	0	
				Tige + Imi	0	
	Clock et al., 2013 ^[89]	USA	48 XDR	Dori + Ak	10	4% with Dori
				Dori + Levo	0	+ Levo
				Dori + PB	4	
				Dori + Rif	23	
				Dori + Tige	0	
				Dori + PB + Rif	19	
				Dori + PB + Tige	8	
	Vidaillac et al., 2012[57]	France	4	Col + Van	25	Nil
				Col + Tmp	25	
				Col + Cotri	25 (in Col R)	
	Elemam <i>et al.</i> , 2010 ^[114]	USA	12 PB-R KPC	PB + Rif	100	Nil
				PB + Rif	100	
				PB + Imi	100	
				PB + Tige	100	
				PB + Gen	0	
	Dawis et al., 2003 ^[103]	USA	10 ESBL	Gati + Cefe	50	Nil
				Gati + Mero	20	
				Gati + Pipe	10	
				Gati + Gen	60	
	Gaibani et al., 2014 ^[70]	Italy	8 Col-R KPC-Kp	Col + Mero	37.5	Nil
	, ,	2	1	Col + Tige	75	
				Col + Rif	100	
				Col + Tec	0	
	Stein et al., 2015 ^[115]	Germany	20	Col + Mero	25	Nil
				Mero + Tige	10	
				Mero + Col + Tige	25	
E-test	Nastro et al., 2014 ^[50]	Argentina	27 Col R	Col + Rif	100	Nil
	Pankey et al 2013 ^[69]	USA	26 KPC	PB + Mero	57.7	Nil

Contd...

Method Reference Place of study Isolate Combination Percentage systems antagoins Gaibani et al., 2014 ¹⁰⁰ USA 14 KPC PB + Marin 21 Nil Gaibani et al., 2014 ¹⁰⁰ USA 14 KPC PB + Marin 21 Nil Gaibani et al., 2014 ¹⁰⁰ USA 6 Col - RF 100 Nil Col - Tage 0 Science 25 Nil Col - Maron 12 S.S Rif - Macon 0 Col - Maron 12 S.S Nil 100 Nil Pankey et al., 2013 ¹⁰¹ USA 26 KPC PB + Maron 73 Nil Hong et al., 2013 ¹⁰¹ USA 12 KPC Col + Era 67 Col + Era 67 Vabaillac et al., 2012 ¹⁰¹ France 4 Col - Dori + Era 67 50 Col + Era Fereingan et al., 2012 ¹⁰¹ USA 12 KPC Col + Dori + Gen 8 17 Gaibani et al., 2012 ¹⁰¹ USA 14 KPC PB + Maron 100 Nil	Table 6: (Contd					
Pankey and Asheraft, 2011 ^[9] USA 14 KPC PB + Mero 43 Nil Gaibani et al., 2014 ^[9] Indy 8 Col-R KPC-Kp Col + Faf 100 Nil Cal + Eaf 100 Nil Col + Faf 100 Nil Cal + Mero 0 Rif - Mero 0 Rif - Mero 0 Cal + Tee 0 Col + Tee 0 Nil Distribution TKA Lee and Burgese, 2013 ^[10] USA 26 KPC PB + Mero 0 Nil Pankey et al., 2013 ^[10] USA 26 KPC PB + Mero 73 Nil Idea et al., 2013 ^[10] USA 12 KPC Col + Pori 60 Col+Fran Cal + Dori 50 Col+Fran 67 Col+Fran 67 Vidaillae et al., 2012 ^[11] USA 12 KPC Col + Pori 50 60 Cal + Dory 50 0 Col + Pori 50 60 Col + Pori 50 60 Jermigan et al., 2011 ^[11] USA 12	Method	Reference	Place of study	Isolate	Combination	Percentage synergy	Percentage antagonism
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		Pankey and Ashcraft, 2011 ^[68]	USA	14 KPC	PB + Mero	43	Nil
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					PB + Rif	21	
KA Lee and Burgess, 2013 ^{10/41} USA 4 KPC Col + Tige (Col + Col + Control + Contro + Control + Control + Control + Control + Control + Control +		Gaibani et al., 2014[70]	Italy	8 Col-R KPC-Kp	Col + Rif	100	Nil
					Col + Tige	25	
TKA Lee and Burgess, 2013 ⁽¹⁾⁽⁴⁾ USA 4 KPC Col + Dri 00 Pankey et al., 2013 ⁽⁴⁾⁽⁴⁾ USA 26 KPC PH + Maro 73 Nil Pankey et al., 2013 ⁽⁴⁾⁽⁴⁾ USA 26 KPC Col + Era 42 25% with Col + Dori + 50 Col + Era 42 25% with Col + Dori + 50 Col + Era 50 Col + Era 50 50 Col + Era 50					Col + Mero	12.5	
TKA Lee and Burgess, 2013 ^[10] USA 4 KPC Col + Teo 0 TKA Lee and Burgess, 2013 ^[10] USA 26 KPC PB + Moro 73 Nil Pankey et al., 2013 ^[10] USA 26 KPC PB + Moro 73 Nil Hong et al., 2013 ^[10] USA 12 KPC Col + Era 42 25% with Col + Dori + Eria 67 Col + Dori + Eria 67 50 50 Gol + Dori + Eria 67 Col + Cori 50 (in Col-R) 50 50 Jernigan et al., 2012 ^[10] France 4 Col + Cori 50 (in Col-R) 50 Jernigan et al., 2012 ^[10] USA 12 KPC Col + Dori 50 60 50 Gen + Doxy 22 17 Gen + Doxy 22 17 50 50 16 50 17 Gen + Doxy 25 17 Gen + Doxy 25 17 50 16 16 100 Nil Purmaras et al., 2011 ^[10] USA 4 KPC Tige + Imi 100 Nil 17 100 100					Rif + Tige	0	
TKA Lee and Burgess, 2013 ⁽¹⁰⁾ USA 4 KPC Col + Dori 100 Nil Pankey et al., 2013 ⁽¹⁰⁾ USA 26 KPC PH + Mero 73 Nil Ilong et al., 2013 ⁽¹⁰⁾ USA 26 KPC PH + Mero 73 Nil Ilong et al., 2013 ⁽¹⁰⁾ USA 26 KPC PH + Mero 73 Nil Udaillac et al., 2012 ⁽¹⁰⁾ France 4 Col + Dori 50 Col + Dori Vidaillac et al., 2012 ⁽¹⁰⁾ France 4 Col + Van 25 S0 Col + Dori 50 0 Col + Dori 50 Dori + Dra 25 Fa Col + Dori<					Rif + Mero	0	
TKA Lee and Bargess, 2013 ⁽¹⁰⁾ USA 4 KPC Col + Dori 100 Nil Pankey et al., 2013 ⁽¹⁰⁾ USA 26 KPC PB + Merio 73 Nil Hong et al., 2013 ⁽¹⁰⁾ USA 12 KPC Col + Dori + Erta 42 25% with Viduillac et al., 2012 ⁽¹⁰⁾ France 4 Col + Dori + Erta 67 50 Jernigan et al., 2012 ⁽¹⁰⁾ France 4 Col + Torny 50 50 Jernigan et al., 2012 ⁽¹⁰⁾ USA 12 KPC Col + Torny 50 60 Col + Cotri 50(in Col-R) 0 0 0 0 0 Col + Cotri 50(in Col-R) 0 0 0 0 0 0 Jernigan et al., 2012 ⁽¹⁰⁾ USA 12 KPC Col + Dori + Erta 25 8 0					Col + Tec	0	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	TKA	Lee and Burgess, 2013 ^[116]	USA	4 KPC	Col + Dori	100	Nil
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					PB + Dori	100	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		Pankey et al., 2013 ^[69]	USA	26 KPC	PB + Mero	73	Nil
Col + Dori 50 Col + Erta Vidaillac et al., 2012 ^[10] France 4 Col + Vani 25 75 Col + Tmp 50 50 50 50 50 50 Jernigan et al., 2012 ^[10] USA 12 KPC Col + Dori 50 0 0 Col + Dori 50 0 Col + Dori 50 0 0 Col + Dori 50 0 0 Col + Dori 50 0 Col + Dori 50 0 0 0 0 0 0 Col + Dori 50 0 0 0 0 0 0 0 Le et al., 2011 ^[117] USA 4 KPC Ak + Imi 100 Nil Pournaras et al., 2011 ^[118] Greece 4 KPC Tige + Mero 0 0 Pankey and Ashcraft, 2011 ^[10] USA 14 KPC PB + Mero 64 Nil Yim et al., 2011 ^[119] Korea 35 ESBI+AmpC Tige + Ak 57.1 (8/14) 100 Yim et al., 2011 ^[119] Greece 17 KPC Fos + Gr		Hong <i>et al.</i> , 2013 ^[45]	USA	12 KPC	Col + Erta	42	25% with
Vidnillac et al., 2012 ^[157] France 4 Col + Van Col + Van 25 75 Hemigan et al., 2012 ^[127] USA 12 KPC Col + Carri 50 (in Col-R) 50 Identitian et al., 2012 ^[127] USA 12 KPC Col + Carri 50 0 Identitian et al., 2012 ^[127] USA 12 KPC Col + Carri 50 0 Identitian et al., 2011 ^[117] USA 4 KPC Ak + Hmi 100 Nil Identitian et al., 2011 ^[117] USA 4 KPC Ak + Hmi 100 Nil Pourmarns et al., 2011 ^[117] Greece 4 KPC Tige + Col 100 Nil Pankey and Asheraft, 2011 ^[168] USA 14 KPC PB + Mero 0 Nil Pim et al., 2011 ^[169] Korea 35 ESBL+AmpC Tige + Col 100 Nil Pim et al., 2011 ^[164] Greece 17 KPC Fos + Col 11.8 57.1 (8/14) Vidaillac et al., 2009 ^[159] USA 2 ESBL Cpt + Naro 0 0 Vidaillac et al., 2009 ^[159] USA 2 ESBL Cpt + Tizz 1000 Nil		<i>c ,</i>			Col + Dori	50	Col+Erta
Vidaillac et al., 2012 ^[27] France 4 Col + Van Col + Tamp 25 75 Jernigan et al., 2012 ^[27] USA 12 KPC Col + Corri 50 0 Col + Corri 50 0 0 0 0 0 Col + Corri 50 0 0 0 0 0 Col + Doxy 8 25 8 0					Col + Dori + Erta	67	
		Vidaillac <i>et al.</i> , 2012 ^[57]	France	4	Col + Van	25	75
$ \begin{array}{cccc} \operatorname{Col} + \operatorname{Corr} & 50 (\operatorname{in} \operatorname{Col} + \operatorname{R}) & 50 \\ \operatorname{Col} + \operatorname{Corr} & 50 (\operatorname{in} \operatorname{Col} + \operatorname{R}) & 50 \\ \operatorname{Col} + \operatorname{Corr} & 50 (\operatorname{in} \operatorname{Col} + \operatorname{R}) & 50 \\ \operatorname{Col} + \operatorname{Gen} & 25 & 8 \\ \operatorname{Col} + \operatorname{Gen} & 25 & 8 \\ \operatorname{Col} + \operatorname{Gen} & 8 & 17 \\ \operatorname{Dori} + \operatorname{Gen} & 8 & 17 \\ \operatorname{Dori} + \operatorname{Gen} & 8 & 17 \\ \operatorname{Dori} + \operatorname{Gen} & 8 & 17 \\ \operatorname{Ori} + \operatorname{Dori} & 1 & 00 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Doxy} & 42 & 17 \\ \operatorname{Re} + \operatorname{Mero} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Mero} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Rer} & 25 & 17 \\ \operatorname{Gen} + \operatorname{Doxy} & 42 & 17 \\ \operatorname{Re} + \operatorname{Rer} & 25 & 17 \\ \operatorname{Gen} + \operatorname{Doxy} & 42 & 17 \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Rer} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Rer} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Rer} & 25 & 17 \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 25 & 17 \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 & \operatorname{Nil} \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 57.1 (8/14) & \operatorname{Re} + \operatorname{Re} & 57.1 (8/14) \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 57.1 (8/14) & \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 \\ \operatorname{Re} + \operatorname{Re} + \operatorname{Re} + \operatorname{Re} + \operatorname{Re} & 100 & \operatorname{Re} + \operatorname{Re} +$					Col + Tmp	50	50
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $					Col + Cotri	50 (in Col-R)	50
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		Jernigan <i>et al</i> $2012^{[42]}$	USA	12 KPC	Col + Dori	50	0
$ \begin{array}{cccc} Col + Doxy & 8 & 25 \\ Dori + Gen & 8 & 17 \\ Dori + Doxy & 25 & 17 \\ Gen + Doxy & 42 & 17 \\ Dori + Doxy & 42 & 17 \\ Ak + Imi & 100 & Nil \\ Ak + Mero & 100 \\ Ak + Mero & 100 \\ Ak + Mero & 100 \\ Pankey and Asheraft, 2011[10] & Greece & 4 KPC & Tige + Col & 100 \\ Pankey and Asheraft, 2011[64] & USA & 14 KPC & Pi + Mero & 64 \\ Yim et al., 2011[119] & Korea & 35 ESBL+AmpC & Tige + Imi & 69.2 (18/26) \\ Yim et al., 2011[119] & Korea & 35 ESBL+AmpC & Tige + Imi & 69.2 (18/26) \\ Yim et al., 2011[14] & Greece & 17 KPC & Fos + Mero & 64.7 \\ Fos + Col & 11.8 \\ Cpt + Tax & 100 \\ Cpt + Cer^{i} & 0 \\ Cpt +$		501111Guil 67 47., 2012	0.011	12100	Col + Gen	25	8
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					Col + Doxy	8	25
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					Dori + Gen	8	17
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$					Dori + Doxy	25	17
Le et al., 2011 ^[117] USA 4 KPC Ak + Imi 100 Nil Le et al., 2011 ^[117] USA 4 KPC Ak + Imi 100 Nil Ak + Eta 25 Pournaras et al., 2011 ^[118] Greece 4 KPC Tige + Col 100 Nil Tige + Mero 0 Nil Tige + Mero 64 Nil PB + Rir 100 Yim et al., 2011 ^[119] Korea 35 ESBL+AmpC Tige + Imi 69.2 (18.26) Nil Tige + Ak 57.1 (8/14) Tige + Cipro 35 (7/20) Souli et al., 2011 ^[14] Greece 17 KPC Fos + Mero 64 Nil Fos + Gen 0 Vidaillac et al., 2009 ^[129] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Ak 100 Nil Cpt + Ak 100 Cpt + Ak 100 Cpt + Ak 100 Cpt + Ak 100 Cpt + Cele 0 Cpt + Cele 0 Cpt + Cele 0 Cpt + Tige 0 Souli et al., 2009 ^[129] USA 1 ESBL Gati + Mero 100 Nil Clancy et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[16] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[17] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2013 ^[17] USA 23 KPC Col + Dori 100 Nil PB + Mero 100 PB + Mero 7 PB 100 Nil					Gen + Doxy	42	17
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		$I = at al = 2011^{[117]}$	USA	AKDC	Ak + Imi	42	17 Nil
NR + Metho 100 Ak + Erta 25 Pournaras et al., 2011 ^[18] Greece 4 KPC Tige + Col 100 Nil Pankey and Asheraft, 2011 ^[83] USA 14 KPC PB + Mero 64 Nil PB + Mir 100 PB + Rif 100 Nil Yim et al., 2011 ^[119] Korea 35 ESBL+AmpC Tige + Imi 69.2 (18/26) Nil Souli et al., 2011 ^[141] Greece 17 KPC Fos + Mero 64.7 Nil Fos + Col 11.8 Fos + Col 11.8 Fos + Col 11.8 Vidaillac et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Azt 0 0 Cpt + Azt 0 0 Cpt + Cefe 0 Souli et al., 2009 ^[120] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[100] USA 1 ESBL Gati + Mero 100 Nil Gaibani et al., 2013 ^[41] USA 23 KPC Col + Rif		Le el ul., 2011	USA	4 KFC	Ak + IIII	100	INII
Pournaras et al., 2011 ^[134] Greece 4 KPC Tige + Col Tige + Mero 100 Nil Pankey and Ashcraft, 2011 ^[68] USA 14 KPC PB + Mero 64 Nil Pige + Mero 64 Nil PB + Rif 100 Nil Yim et al., 2011 ^[139] Korea 35 ESBL+AmpC Tige + Imi 69.2 (18/26) Nil Yim et al., 2011 ^[141] Greece 17 KPC Fos + Mero 64.7 Nil Fos + Gen 0 Fos + Gen 0 Nil Fos + Gen 0 Vidaillae et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Haro 0 Cpt + Azt 0 0 Cpt + Azt 0 Souli et al., 2009 ^[120] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[100] USA 1 ESBL Gati + Mero 100 Nil Gaibani et al., 2013 ^[41] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2017 ^[101] U					Ak + Nielo	100	
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		December of al. 2011[118]	Creation	ANDC	AK + LIta	23	N.:I
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		Pournaras et al., 2011 ^[110]	Greece	4 KPC	Tige + Col	100	INII
Pankey and Asheraft, 2011 ^[19] USA 14 KPC PB + Nir 100 PB + Rif 100 PB + Rif 100 PB + Rif 69.2 (18/26) Nil Tige + Ak 57.1 (8/14) Tige + Cipro 35 (7/20) Souli <i>et al.</i> , 2011 ^[14] Greece 17 KPC Fos + Mero 64.7 Nil Fos + Col 11.8 Fos + Col 11.8 Fos + Gen 0 Vidaillac <i>et al.</i> , 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Taz 100 Cpt + Azt 0 Cpt + Azt 0 Cpt + Cefe 0 Cpt + Cefe 0 Cpt + Cefe 0 Cpt + Cefe 0 Cpt + Tige 0 Souli <i>et al.</i> , 2009 ^[121] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis <i>et al.</i> , 2003 ^[103] USA 1 ESBL Gati + Mero 100 Nil Clancy <i>et al.</i> , 2017 ^[121] USA 23 KPC Col + Dori 26 Nil Gaibani <i>et al.</i> , 2017 ^[121] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 PB + Mero + Rif 100 Kulengowski <i>et al.</i> , 2017 ^[121] USA 4 KPC-Kp Mero + PB 100 Nil			110.4	14 800	Tige + Mero	0	N71
Yim et al., 2011 ^[119] Korea 35 ESBL+AmpC Tige + Ini 69.2 (18/26) Nil Tige + Ak 57.1 (8/14) Tige + Cipro 35 (7/20) Tige + Cipro 35 (7/20) Souli et al., 2011 ^[44] Greece 17 KPC Fos + Mero 64.7 Nil Fos + Col 11.8 Fos + Gen 0 11.8 Vidaillac et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Ak 0 O Cpt + Ak 00 Nil Souli et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 0 O Souli et al., 2009 ^[121] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[163] USA 1 ESBL Gati + Mero 100 Nil Gaibani et al., 2013 ^[43] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil PB + Nero 100 Nil		Pankey and Ashcraft, 2011 ^[00]	USA	14 KPC	PB + Mero	64	Nil
Yim et al., 2011 Korea 35 ESBL+AmpC lige + lmi 69.2 (18/26) Nil Tige + Ak 57.1 (8/14) Tige + Chro 35 (7/20) Souli et al., 2011 Greece 17 KPC Fos + Mero 64.7 Nil Fos + Gen 0 11.8 Fos + Gen 0 Nil Vidaillac et al., 2009 ⁽¹²⁰⁾ USA 2 ESBL Cpt + Ak 100 Nil Cpt + Haro 0 Cpt + Haro 0 0 Cpt + Azt 0 0 Souli et al., 2009 ⁽¹²⁰⁾ Greece 42 VIM-1 Col + Imi 33.3 23.8 Bawis et al., 2003 ⁽¹⁰³⁾ USA 1 ESBL Gati + Mero 100 Nil Gati et al., 2011 ⁽¹²¹⁾ Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ⁽¹⁰³⁾ USA 1 ESBL Gati + Mero 100 Nil Gati et al., 2014 ⁽¹⁰⁴⁾ Italy 8 Col-R KPC-Kp Col + Rif 100 Nil Diep et al., 2017 ⁽¹²²⁾ USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil			**		PB + Rif	100	N 711
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$		Yim <i>et al.</i> , $2011^{[119]}$	Korea	35 ESBL+AmpC	Tige + Imi	69.2 (18/26)	Nil
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$					Tige + Ak	57.1 (8/14)	
Souli et al., 2011 ^[44] Greece 17 KPC Fos + Mero 64.7 Nil Fos + Col 11.8 Fos + Col 11.8 Fos + Gen 0 Vidaillac et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Ak 0 Cpt + Akt 0 0 Cpt + Azt 0 Souli et al., 2009 ^[121] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[103] USA 1 ESBL Gati + Mero 100 Nil Gaibani et al., 2013 ^[43] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil Die et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil B + Mero 100 Nil PB + Mero 100 Nil					Tige + Cipro	35 (7/20)	
Vidaillac et al., 2009 ^[120] USA 2 ESBL Fos + Col 11.8 Fos + Gen 0 Nil Cpt + Ak 100 Nil Cpt + Taz 100 Cpt + Azt 0 Cpt + Levo 0 Cpt + Cefe 0 Cpt + Tige 0 Souli et al., 2009 ^[121] Greece 42 VIM-1 Dawis et al., 2003 ^[103] USA 1 ESBL Gati + Mero Gati + Gen 100 Nil Gati + Gen 100 Nil Clancy et al., 2013 ^[43] USA 2 SKPC Col + Dori 26 Nil Gaibani et al., 2014 ^[70] Italy 8 Col-R KPC-Kp Col + Rif 100 Nil Diep et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil PB + Mero 100 Nil Diep et al., 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil		Souli <i>et al.</i> , 2011 ^[44]	Greece	17 KPC	Fos + Mero	64.7	Nil
Fos + Gen 0 Vidaillac et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil Cpt + Taz 100 Cpt + Akz 0					Fos + Col	11.8	
Vidaillac et al., 2009 ^[120] USA 2 ESBL Cpt + Ak 100 Nil $Cpt + Taz$ 100 $Cpt + Mero$ 0 $Cpt + Azt$ 0 $Cpt + Levo$ 0 $Cpt + Cefe$ 0 $Cpt + Tige$ 0 Souli et al., 2009 ^[121] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[103] USA 1 ESBL Gati + Mero 100 Nil Gati + Gen 100 Nil Gati + Gen 100 Nil Clancy et al., 2013 ^[43] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil PB + Mero 100 Nil Videngowski et al., 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil					Fos + Gen	0	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		Vidaillac et al., 2009 ^[120]	USA	2 ESBL	Cpt + Ak	100	Nil
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					Cpt + Taz	100	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					Cpt + Mero	0	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$					Cpt + Azt	0	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$					Cpt + Levo	0	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$					Cpt + Cefe	0	
Souli et al., 2009 ^[121] Greece 42 VIM-1 Col + Imi 33.3 23.8 Dawis et al., 2003 ^[103] USA 1 ESBL Gati + Mero 100 Nil Gati + Gen 100 Gati + Gen 100 Nil Clancy et al., 2013 ^[43] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2014 ^[70] Italy 8 Col-R KPC-Kp Col + Rif 100 Nil Diep et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil PB + Mero + Rif 100 Nil Kulengowski et al., 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil					Cpt + Tige	0	
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $		Souli et al., 2009 ^[121]	Greece	42 VIM-1	Col + Imi	33.3	23.8
Clancy et al., 2013 ^[43] USA 23 KPC Col + Dori 26 Nil Gaibani et al., 2014 ^[70] Italy 8 Col-R KPC-Kp Col + Rif 100 Nil Diep et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Nil PB + Mero + Rif 100 Nil Kulengowski et al., 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil		Dawis et al., 2003 ^[103]	USA	1 ESBL	Gati + Mero	100	Nil
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$					Gati + Gen	100	
Gaibani et al., 2014 ^[70] Italy 8 Col-R KPC-Kp Col + Rif 100 Nil Diep et al., 2017 ^[122] USA 2 isolates PB + Rif 100 Nil PB + Mero 100 PB + Mero 100 Nil Kulengowski et al., 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil		Clancy et al., 2013 ^[43]	USA	23 KPC	Col + Dori	26	Nil
Diep <i>et al.</i> , $2017^{[122]}$ USA 2 isolates PB + Rif 100 Nil PB + Mero 100 Kulengowski <i>et al.</i> , $2017^{[123]}$ USA 4 KPC-Kp Mero + PB 100 Nil		Gaibani <i>et al.</i> , 2014 ^[70]	Italy	8 Col-R KPC-Kn	Col + Rif	100	Nil
PB + Mero + Rif 100 Kulengowski <i>et al.</i> , 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil		Diep <i>et al.</i> $2017^{[122]}$	USA	2 isolates	PB + Rif	100	Nil
Kulengowski et al., 2017 ^[123] USA4 KPC-KpMero + PB100Nil		r,,	~~~*		PB + Mero	100	
Kulengowski <i>et al.</i> , 2017 ^[123] USA 4 KPC-Kp Mero + PB 100 Nil					PB + Mero + Rif	100	
		Kulengowski et al., 2017 ^[123]	USA	4 KPC-Kp	Mero + PB	100	Nil

KPC: *Klebsiella pneumoniae* carbapenemase, Col: Colistin, Rif: Rifampicin, R: Resistant, Gen: Gentamicin, Mero: Meropenem, Imi: Imipenem, Tige: Tigecycline, XDR: Extensive drug resistance, Dori: Doripenem, Ak: Amikacin, Levo: Levofloxacin, PB: Polymyxin B, Van: Vancomycin, Cefe: Cefepime, Pipe: Piperacillin, ESBL: Extended spectrum beta-lactamase, Gati: Gatifloxacin, Tec: Teicoplanin, Doxy: Doxycycline, Cipro: Ciprofloxacin, Fos: Fosfomycin, Taz: Tazobactam, Azt: Aztreonam, Erta: Ertapenem, CB: Checker board, TKA: Time-kill assay, Cpt: Ceftaroline, TMP: Trimethoprim, Cotri: Trimethoprim/sulfamethoxazole

Table 7: Combined synergy by different methods for *Acinetobacter baumannii*, *Pseudomonas aeruginosa* and *Klebsiella* pneumoniae*

Drug combination	Percentage c	Remarks		
	ТКА	CB	<i>E</i> -test	
		A. baumannii		
Sul+Imi	66.6 (42-100)	64 (0-88.2)	17.5-32.5	Combination
Sul+Mero	100	57.7 (0-94)		of Col with
Sul+Dori	-	9		either Mero or
Sul+Col	-	53.3	45.5	Rif shows high
Sul+Rif	100	-	-	rate while
Col+Mero	96.3 (96-100)	69.4 (0-73.3)	-	combination of
Col+Imi	59 (0-100)	63.7 (0-100)	69.2 (20-100)	Imi with either
Col+Dori	100	40 (36-66.6)	-	Sul or Col shows
Col+Rif	94.2 (77-100)	83 (80-100)	100	moderate level of
Col+Tige	100	15 (0-24.3)	-	synergy
PB+Mero	100	2.2 (0-2.9)	62.5	
PB+Imi	87.5	38.2	-	
PB+Tige	43.7	12.5	0	
PB+Rif	66.6 (56.2-87.5)	18.7	6.2	
	、 /	P. aeruginosa		
Col + Rif	11.8	100	-	There are
Col + Imi	100	44.4 (28.5-100)	-	limited data
Col + Mero	25.4	12	-	on a particular
Col + Dori	76	-	3	combination.
Imi + Fos	-	12.7 (0-100)	46.7	combinations
Mero + Fos	-	-	53.3	include Col plus
Dori + Fos	-	-	73.3	carbapenem and
Imi + Cipro	_	-	carbapenem plus	
Imi + Trova	100	23.3	-	fluoroquinolone
Imi + Gati	75		-	combinations
Mero + Cipro	65.2 (61-66)		-	
Mero + Levo	91.6	5.6	-	
Mero + Gati	-	70	-	
Dori + Levo	_	-	9	
Tobra + Imi	_	68(0-15)	-	
Tobra + Mero	_	-	19	
Tobra + Pine/Taz	_	55 (46-100)	-	
Tobra + Czd	_	45.3 (31-75)	28.8	
Tobra + Cefe	_	50	-	
Tobra + Cipro	-	3.4 (0-25)	-	
		K. pneumoniae		
Col + Imi	33.3	38.5	_	Combination
Col + Mero	-	38.5	-	of Col/PB plus
Col + Dori	43.1 (26-100)	-	-	Rif/Mero gives
Col + Erta	42	-	-	promising result
Col + Tige 100		38.5	-	and needs to be
Col + Rif	-	100	100	<i>in vivo</i> models
PB + Dori 100		4	-	and clinical trials
PB + Mero	70 70 (64-73) - 524		52.5 (43-57.7)	Antagonism
PB + Rif	100 100 21			
Tige + Mero	0	0	<u>_ 1</u>	plus Imi/Erta
Tige + Imi	69.2	0	-	combinations
Tige + Dori	-	0	-	
Tige + Ak	57 1	-	-	
	C / . 1			

Contd...

Table 7: Contd							
Drug combination	Percentage c	Remarks					
	ТКА	CB	<i>E</i> -test				
Tige + Cipro	35	-	-				
Col + Gen	-	38.4	-				
Mero + Gen	-	38.4	-				
Imi + Gen	-	23	-				
Tig + Gen	-	0	-				
Mero + PB	100	-	-				
Mero + Rif + PB	100	-	-				

*Combined synergy is not provided for combinations with only one study. Sul: Sulbactam, Imi: Imipenem, Mero: Meropenem, Dori: Doripenem, Col: Colistin, Rif: Rifampicin, Tige: Tigecycline, PB: Polymyxin B, *P. aeruginosa: Pseudomonas aeruginosa, A. baumannii: Acinetobacter baumannii*, Fos: Fosfomycin, Cipro: Ciprofloxacin, Trova: Trovafloxacin, Gati: Gatifloxacin, Levo: Levofloxacin, Tobra: Tobramycin, Pipe: Piperacillin, Taz: Tazobactam, Czd: Ceftazidime, Cefe: Cefepime, Erta: Ertapenem, Ak: Amikacin, Gen: Gentamicin, *K. pneumonia: Klebsiella pneumonia*, TKA: Time-kill assay, CB: Checker board

Reference	Combination (method)	lsolate (MIC μg/ml)	п	Synergy (%)	Antagonism (%)
	A	. baumannii			
Kiffer et al.[37]	Sul + Mero (CB)	Sul R (≤4)	19	5.2	15.7
		Sul S (≥16)	13	76.9	0
Cikman et al.[41]	Amp/Sul + Col (CB)	Col R	12	8.3	75
		Col S	21	66.6	0
Principe et al.[38]	Dori + Tige/Col/Ak/Amp-Sul/Rif (CB)	Dori R	17	27	0
		Dori S	5	0	0
Sheng et al.[40]	Imi + Ak/Col/Tige/Amp-Sul (CB)	Imi ≥32	5	0	0
		Imi <32	7	20	0
	Imi + Col (TKA)	Imi ≥32	5	40	0
		Imi <32	7	100	0
Ji <i>et al</i> . ^[72]	Sul + Imi/Mero/Pan/Cef (CB)	Imi R	40	7.5-25	0
		Imi S	40	20-27.5	0
	К	. pneumonia			
Jernigan et al.[42]	Dori + Col (TKA)	Col≥8	9	66.6	0
		Col ≤4	3	0	0
Souli et al.[121]	Col + Imi (TKA)	Col≥16	18	11.1	55.5
		Col ≤8	24	50	0
Clancy et al.[43]	Dori + Col (TKA)	Dori≥16	19	32	0
-		Dori ≤8	4	100	0
Kulengowski et al.[123]	Mero + PB (TKA)	Mero ≥4	4	100	0
-	· ·	PB 0.06	2	100	0
		PB 0.125	2	100	0

*R: Resistant, S: Susceptible. MIC: Minimum inhibitory concentration, *A. baumannii: Acinetobacter baumannii*, Sul: Sulbactam, Mero: Meropenem, Col: Colistin, Ak: Amikacin, Amp: Ampicillin, Rif: Rifampicin, CB: Checker board, Imi: Imipenem, Tige: Tigecycline, Cef: Cefpirome, Dori: Doripenem, PB: Polymyxin B, *K. pneumonia: Klebsiella pneumonia*, TKA: Time-kill assay, Pan: panipenem

reported favourable outcome with carbapenem and sulbactam combination in four patients (two patients with VAP and two catheter-related bloodstream infection) caused by *A. baumannii*. *In vitro* synergy testing of all four isolates by CB assay showed partial synergy with FIC index ranging from 0.56 to 0.75 for combination of sulbactam and meropenem/imipenem.^[49] In one patient with post-neurosurgery bacteraemic meningitis due to CR *A. baumannii* (CRAB), combination of intravenous meropenem and sulbactam leads to reduction in the colony count in cerebrospinal fluid (CSF) from >50,000 CFU/ml to 10,000 CFU/ml in 4 days.^[39] Addition of intravenous and

intrathecal colistin resulted in clearance of the organism within 2 days both from CSF and blood. The SBT and CSF bactericidal titre increased from fourfold to 32-fold with the three-drug combination compared to two-drug regimen. *In vitro* TKA showed synergism with combination of colistin with meropenem, sulbactam or both. TKA with colistin alone and meropenem plus colistin showed re-growth at 24 h. Though the infection was cleared, the patient expired due to hypoxia secondary to respiratory distress. Nastro *et al.* reported successful treatment for cases with sepsis (n = 1), meningitis (n = 1) and UTI (n = 1) with a combination of

Table 9: Correlation of molec	ular mechanisms of resistance with	the result of in vit	ro antimicrobial interaction	on study
Reference	Molecular mechanism	п	Combination (method)	Percentage synergy
	A. bauman	nii		
Rodriguez et al., 2010 ^[95]	$bla_{0XA51} + bla_{0XA58}$	3	Imi + Rif (TKA)	0
	$bla_{0XA51} + bla_{0XA23}$	9	Imi + Rif (TKA)	0
Tripodi et al., 2007 ^[8]	bla _{OXAS8}	9	Imi/Amp-Sul + Rif	100
•	UAA38	he result of <i>in vitro</i> antimicrobial interationnCombination (method)ii3Imi + Rif (TKA)9Imi/Amp-Sul + Rif0Col + Rif (TKA)9Imi/Amp-Sul + Rif0Col + Rif (TKA)5Imi + PBRif + PBAzi + PB (E-test)8Col + SulCol + TimiCol + SulCol + FosSul + ImiCol + RifImiSul + Fos (CB)8Sul + ImiCol + Rif (If (<i>in vivo</i> mouse pneumonia model)1Col + RifImi + Rif1Col + RifImi + Rif5Ak + Imi/Col/Tige (<i>E</i> -test)10Sul + Col39Sul + Col8Col + Dori + Erta Col + Dori (TKA)4Col + Dori (TKA)8Col + Dori (TKA)8Col + Dori (TKA)8Col + Dori (TKA)8Col + Dori (TKA)90 (against Dori)Dori + Ak 	77	
Wareham and Bean, 2006 ^[36]	bla	5	Imi + PB	20
	07425		Rif + PB	20
			Azi + PB (E-test)	0
Santimaleeworagun et al., 2011 ^[56]	bla	8	Col + Sul	0
• · ·	UXA25		Col + Imi	0
			Col + Fos	12.5
			Sul + Imi	0
			Sul + Fos (CB)	75
Song <i>et al.</i> , 2007 ^[83]	bla	8	Sul + Imi	87.5
	07451		Col + Rif (TKA)	100
Song <i>et al.</i> , 2009 ^[124]	bla	1	Col + Rif	0
	· · · · IMP		Imi + Rif (<i>in vivo</i> mouse	100
	1.1	,		100
	bla _{OXA51}	1	Col + KII	100
	11	,	ImI + KII	100
	bla _{VIM}	1	Col + Rif	0
		-	Imi + Kii	100
Miyasaki <i>et al.</i> , $2012^{[92]}$	AME, Ser83leu substitution in GyrA,	5	Ak + Imi/Col/Tige	0
L	bla bla bla	10	(L-test)	40
	$bla_{\text{NDM}}, bla_{\text{OXA51}} + bla_{\text{OXA23}}$	10	Sul + Col	40
	$bla_{OXA51} + bla_{OXA23}$	39	Sul + Col	18
XX L 2012[45]	K. pneumon	iae		
Hong <i>et al.</i> , $2013^{[+3]}$	High <i>omp35/omp36</i> expression	8	Col + Dori + Erta	100
			Col + Dori (TKA)	63
	Low <i>omp35/omp36</i> expression	4	Col + Dori + Erta	0
			Col + Dori (TKA)	25
Clancy <i>et al.</i> , $2013^{[43]}$	Wild type <i>ompK36</i> or other mutations	8	Col + Dori (TKA)	75
	IS5 mutants	7	Col + Dori (TKA)	29
	Ins aa 134-135 GD mutants	8	Col + Dori (TKA)	25
Poirel <i>et al.</i> , 2016 ^[125]	$bla_{\rm KPC}$ (n=8)	20	Imp + Mem	40
	bla_{NDM} (n=4)		Imp + Dori	20
	$bla_{\text{OXA-48}}$ (n=6)		Imp + Etp	40
	$bla_{\rm NDM} + bla_{\rm OXA181}$ (2)		Mem + Dori	15
			Etp + Dori	15
			Etp + Mem	0
Laishram et al., 2016 ^[78]	bla_{OXA-48} (n=23)	50	Col + Mem	30
	bla_{NDM} (n=16)		Col + Mem	69
	$bla_{\text{NDM}} + bla_{\text{OXA-48}} (n=11)$		Col + Mem	46
	P. aerugino	sa		
He et al., 2012 ^[106]	Efflux	67 (against Dori)	Dori + Ak	23.8
		99 (against Dori/	Dori + Col	19.5
		Levo)	Dori + Levo	24.3
	Loss of porin	92	Dori + Ak	4.4
			Dori + Col	3.2
			Dori + Levo	7.3
	AmpC	41	Dori + Ak	9
			Dori + Col	8.6
			Dori + Levo	7.3

Contd...

Table 9: Contd				
Reference	Molecular mechanism	п	Combination (method)	Percentage synergy
	Metallo-β-lactamase	3	Dori + Ak	0
			Dori + Col	0
			Dori + Levo	0

Even though synergy levels vary with different mechanisms, antagonism was not noted in any of the combinations tested. TKA: Time-kill assay, Imi: Imipenem, Rif: Rifampicin, Amp: Ampicillin, Sul: Sulbactam, Col: Colistin, PB: Polymyxin B, *A. baumannii: Acinetobacter baumannii, K. pneumonia: Klebsiella pneumonia, P. aeruginosa: Pseudomonas aeruginosa*, Azi: Azithromycin, Fos: Fosfomycin, Tige: Tigecycline, Dori: Doripenem, Erta: Ertapenem, Levo: Levofloxacin, Ak: Amikacin, CB: Checker board, Imp: Imipenem, Mem: Meropenem, Etp: Ertapenem

Table 10: Correlation of in vitro synergy with clinical response							
Reference	Clinical condition	Organism	Combination administered	In vitro assay result	Clinical outcome		
Biancofiore et al., 2007 ^[48]	Multifocal (lung, skin, soft tissue) infection	A. baumannii	Col + Rif + Mero	Col + Rif CB FIC index 0.3 Mero + Rif CB FIC index 0.25 Col + Mero CB FIC index 1	Bacteria eradicated, patient survived		
Lee et al., 2007 ^[49]	Pneumonia	A. baumannii	Mero + Sul	CB FIC index 0.56	Bacteria eradicated, patient survived		
	VAP	A. baumannii	Imi + Sul	CB FIC index 0.56	Bacteria eradicated, patient survived		
	CRBSI	A. baumannii	Imi + Sul	CB FIC index 0.75	Bacteria eradicated, patient survived		
	CRBSI	A. baumannii	Imi + Sul	CB FIC index 0.56	Bacteria eradicated, patient survived		
Lee et al., 2008 ^[39]	Post-neurosurgery bacteraemic meningitis	A. baumannii	Col + Mero + Sul	Col + Mero synergy by TKA Col + Sul synergy by TKA Col + Mero + Sul synergy by TKA	Bacteria eradicated, patient expired due to respiratory distress		

A. baumannii: Acinetobacter baumannii, Col: Colistin, Rif: Rifampicin, Mero: Meropenem, CB: Checker board, FIC: Fractional inhibitory concentration, Sul: Sulbactam, TKA: Time-kill assay, VAP: Ventilator-associated pneumonia, CRBSI: Catheter-related bloodstream infection, Imi: Imipenem

colistin and rifampicin against carbapenemase producing GNB.^[50] The combination was found to be synergistic for all the isolates by the *E*-test-agar method. Table 10 summarises the studies that have correlated *in vitro* synergy with clinical response.

CONCLUSION

Combination therapy has gained attention due to increased efficacy and scope for decreasing the toxicity and development of resistance especially against drug-resistant strains. Therefore, it is imperative to investigate the antimicrobials that have to be used in combination for the clinical utility. At present, very few agents are available for treating infections due to PDR pathogens, and combination therapy is found to be the effective strategy to tackle this. Several methods exist for the assessment of synergistic activity of two or more antimicrobial agents. However, wide variation was observed in terms of their technical issues, complexity and interpretation of test results. This signifies the need for global-level standardisation of the various methods for the determination of synergy of antimicrobial combinations. At present, TKA is the reference method which yields considerable level of concordance rate among the various studies. To conclude, majority of the in vitro test methods could not predict the clinical success rates. Therefore, prospective clinical trials

with *in vitro* synergy testing data are needed to improve the clinical outcome.

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Conflicts of interest

There are no conflicts of interest.

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