

Determination of Susceptibility Breakpoints of Eiores in Extended Spectrum Beta-lactamases and Metallo Beta-lactamases Producing *Acinetobacter* Species

Madhav Prabhu¹, Mohit Arora¹, Zubin Ruttonji², Satish Patil³, Vineeta Dhyani⁴, Mohd Amin Mir⁵

ABSTRACT

Introduction: Infections caused by *Acinetobacter* species have now been becoming troublesome due to their ability to develop resistance. Current study was aimed and to find microbiological cut-off of Eiores against *Acinetobacter* species.

Material and Methods: In this study, 457 clinical isolates of *Acinetobacter* species recovered from patients were involved. Breakpoint of Eiores was established using the methods of pharmacokinetic and pharmacodynamic correlation.

Results: Among 457 isolates, 26.9% were ESBL (Extended Spectrum Beta-Lactamases) positive, 51.2% were MBL (Metallo Beta-Lactamases) positive and remaining 21.8% isolates were ceftriaxone susceptible. Out of identified ESBL and MBL producers 20 strains (10 ESBL and 10 MBL) were picked randomly for gene characterization and it was observed that 3 were TEM-1 positive, 1 was SHV-1 positive, 1 was AMP-C positive, 2 were OXA-1 positive, 3 was CTX-M positive, 3 were VIM-1 positive, 4 were IMP-1 positive, 3 were NDM-1 positive. In ceftriaxone, resistant *Acinetobacter* group, the MIC₅₀ and MIC₉₀ for ceftriaxone were 16 and 128 µg/ml and for Eiores MIC₅₀ and MIC₉₀ were 4 and 8 µg/ml, respectively. The MIC cut off for Eiores against ceftriaxone resistant *Acinetobacter* species was ≤8 µg/ml. Similarly, the AST cut off for Eiores against ceftriaxone susceptible *Acinetobacter* species was ≥21 mm. The AST and MIC cut off for Eiores against ceftriaxone resistant was same as of ceftriaxone susceptible.

Conclusion: Our data apparently indicates that Eiores could be a better option to curb ESBL and MBL producing pathogens which synergistically provides clinically relevant concentration at higher susceptible MIC values.

Keywords: Adjuvant Therapy, Extended Spectrum Beta-Lactamases, *E. coli*, Pharmacodynamic, Pharmacokinetic

INTRODUCTION

Acinetobacter, an opportunistic pathogen, known as dominant cause of nosocomial infections such as meningitis, bacteraemia, wound infections, bloodstream infection, nosocomial-hospital acquired pneumonia or ventilator-associated pneumonia in ICU patients.¹⁻⁵

Acinetobacter infections are commonly treated with cephalosporins, aminoglycosides, carbapenems, and tetracyclines.⁴ However, infections caused by *Acinetobacter* species have now been becoming troublesome due to their ability to develop resistance against these drugs by acquiring various mechanisms predominantly extended spectrum beta lactamases (ESBLs) and metallo beta lactamses (MBLs), representing a major challenge in health care setting.⁶⁻⁹ In India prevalence of resistance due to ESBLs and MBLs (70.9%) varies greatly.¹⁰⁻¹² Polymyxins and tigecycline have also been reported to be resistant to *A. baumannii*.¹³⁻¹⁴

Almost 35% *Acinetobacter* species have been reported to be resistant to carbapenem drugs.^{1,15-16} According to a surveillance study conducted at 40 centers in 12 countries revealed a substantial increase in resistance rates in *Acinetobacter* species for meropenem (43.4%) and imipenem (42.5%).¹⁷ The prevalence of imipenem resistance in *Acinetobacter baumannii* isolated from a burns unit of United State America was found to be 87%.¹⁸ Similarly, according to a surveillance study conducted in several regions of Greece between 1996 and 2007. *A. baumannii* showed resistance to imipenem upto 85% (ICUs), 60% (medical wards), and 59% (surgical wards) [Greek System for Surveillance of Antimicrobial Resistance (GSSAR): <http://www.mednet.gr/whonet/>].

In light of the above background, the soaring rate of the antibiotic resistance and its impact on treatment failure, Eiores, a combination of ceftriaxone, sulbactam and non-antibiotic adjuvant disodium EDTA (ethylene diamine tetraacetic acid) collectively called as antibiotic adjuvant entity (AAE) is a ray of new hope of clinicians.

Ceftriaxone, a third generation cephalosporin, has been used for the disposition of bounteous bacterial infections some of which are very severe such as endocarditis, meningitis, pneumonia, bone and joint infections, intra-abdominal infections, skin infections and urinary tract infection.¹⁹⁻²³ Sulbactam, well known β-lactamase inhibitor, has been reported to have significant *in vitro* activity towards *Acinetobacter* species.²⁴ Its efficacy with combination of ampicillin was shown to have synergistic antibacterial activity.²⁴ Wood et al.²⁵ observed no differences in clinical outcomes between sulbactam-treated and imipenem-treated patients.

Eiores exhibits antibacterial activity by inhibiting bacterial cell wall synthesis, irreversible inhibition of beta-lactamase enzymes. Moreover, EDTA synergizes the activity of ceftriaxone and sulbactam when administered along with these antibiotics.²⁶⁻²⁸ Several recent published articles proved its in

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vitro and in vivo activity towards *Acinetobacter* species.²⁹⁻³³ The aim of the current study is to provide the data to clinicians for their understanding and implementation of Elores (an antibiotic adjuvant entity (AAE) of ceftriaxone+sulbactam+EDTA) breakpoints for *Acinetobacter* species.

MATERIAL AND METHODS

In this study, a total of 457 clinical isolates of *Acinetobacter* family including *A. baumannii*, *A. iwoffii*, *A. junii*, *A. nosocomialis* and *A. pittii* were used. This study was conducted between year 2013 to 2015 in Jawaharlal Nehru Medical College-Belagavi, Karnataka. Re-identification of these isolates was carried out as described earlier.³⁴ These isolates were characterized into ceftriaxone susceptible and ceftriaxone resistant based on the CLSI breakpoints. Quality control (QC) was assured by concurrent testing of *E. coli* ATCC 25922 (ESBL-ve), *E. coli* ATCC 35218 (TEM+ve), *K. pneumoniae* ATCC 700603 (SHV+ve), *P. aeruginosa* ATCC 27853 (MBL-ve), *K. pneumoniae* ATCC BAA-2146 (NDM-1+ve), *K. pneumoniae* NCTC 13439 (VIM-1+ve) and *E. coli* NCTC 13476 (IMP+ve).

Reagents and media: SCD Broth Medium, Mueller-Hinton (MH) agar, Cation-adjusted MH broth (CAMHB), Barium chloride and Sulphuric acid were used for the study. All media and reagents were procured from Hi-Media, India.

Preparation of inoculum, McFarland standard and agar plates: The collected bacterial isolates of *Acinetobacter* species were spread onto SCDA plates and incubated at 35°C to 37°C for 18 to 24 hrs. Following incubation, 3 to 5 morphologically identical colonies of bacterial cells from SCDA plates were inoculated into 10 ml of MHB to achieve the turbidity of the 0.5 McFarland standard (18-24 hrs) and were used for further identification and analysis. The 0.5 McFarland Standard was prepared in MHB medium. Briefly, 0.5ml of 0.048 M BaCl₂ (1.172% w/v BaCl₂.H₂O) was added to 99.5 ml of 0.18 M H₂SO₄ (1% v/v) with constant stirring and the absorbance was monitored at wavelength of 625 nm. SCDA and Mueller Hinton Agar (MHA) plates were prepared by pouring freshly prepared, sterilized, and hot (~50°C) agar media (20-25 ml) into 90 mm flat-bottom sterilized petri dishes. These petri dishes were then allowed to solidify at room temperature. The SCDA and MHA plates were stored in sealed plastic bags for one week at 2 to 8 °C, when not in use. MHB was prepared according to instructions of manufacturer.

Drugs: The drugs used in the study were: ceftriaxone plus EDTA plus sulbactam; Elores (1.5 g) and ceftriaxone (1.0 g). Both the drugs were reconstituted in water for injection.

Screening for ESBL and MBL production: Ceftriaxone resistant isolates were further subjected for extended spectrum beta-lactamases (ESBL) and metallo beta-lactamases (MBL) characterization using the methods described earlier.³⁵⁻³⁶ Randomly selected twenty isolates which were found to be ESBL and MBL positive (ten of ESBL and ten of MBL) were further processed for gene characterization using the previous reported method (Table 1).

Antimicrobial susceptibility testing (AST): The AST was determined using the method described earlier in CLSI.³⁵ The discs of ceftriaxone (30 µg) and Elores (ceftriaxone 30 µg;

sulbactam 15µg) were used. Each disc of Elores contain 100 µg of EDTA and because EDTA is considered as non-antibiotic content hence not disclosed on the disc. Quality control (QC) was assured by concurrent testing of *E. coli* ATCC 25922 (ESBL-ve), *E. coli* ATCC 35218 (TEM+ve), *K. pneumoniae* ATCC 700603 (SHV+ve), *P. aeruginosa* ATCC 27853 (MBL-ve), *K. pneumoniae* NCTC 13439 (VIM-1+ve), *K. pneumoniae* ATCC BAA-2146 (NDM-1 +ve) and *E. coli* NCTC 13476 (IMP+ve).

Minimal inhibitory concentration (MIC): Minimum inhibitory concentration (MIC) was determined using broth micro dilution assay following the CLSI guidelines.³⁵ MIC can be defined as the lowest concentration of antimicrobial agent that prevented turbidity after 16 to 20 hours of incubation at 35 ± 2°C.

MIC₅₀ and MIC₉₀ determination: The concentrations that inhibited 50% (MIC₅₀) and 90% (MIC₉₀) of the strains were calculated for each of the antimicrobial agents as described earlier³⁷ using the formula of geometric.

MIC breakpoint determination: The MIC breakpoint for susceptible microorganisms was determined as per equation 1.³⁸ Breakpoint concentration = C_{max} * f * s / (e*t)

Where, C_{max} = maximum serum concentration; f is a protein binding factor, which is equal to 1 (or <70% protein binding), 0.5 (for 70-90% protein binding) and 0.2 (for >90% protein binding); s is a shift (or reproducibility) factor, which is usually equal to 1; e is factor by which the C_{max} should exceed the MIC and t is factor accounts for the serum elimination half-life, which is equal to 2 (for T_{1/2} <1 hr), 1 (for T_{1/2} = 1-3 hrs), and 0.5 (for T_{1/2} >3 hrs). Values were taken from literature reference. The MIC breakpoint for resistant microorganisms was determined based on PKPD index %T>MIC (equation 2).³⁹ %T>MIC = ln (Dose*/ (V_d*MIC)) *(T_{1/2}/0.693) *(100/DI) Where V_d (L) is the apparent volume of distribution in the central compartment, MIC is minimum inhibitory concentration (µg/mL) of Elores, T_{1/2} (hr) is half-life of antibiotic, and DI is the dosing interval (hrs).

AST breakpoint determination: The AST breakpoints were obtained by drawing a perpendicular towards X axis which when intersected at horizontal line from the MIC breakpoints and the point of intersection of regression line, that value at x-axis, was considered as AST breakpoint corresponding to that MIC. Based on this AST was categorized as the vulnerability of the bacterial isolates into susceptible, intermediate, and resistant against the target antibiotic.^{38,40}

STATISTICAL ANALYSIS

The scatter plots were drawn between MICs and corresponding AST values of the Elores and ceftriaxone against the bacterial isolates. Least square regression analysis was then performed to identify the best possible correlation between MIC and AST values of Elores and ceftriaxone.

RESULTS

Bacterial isolate characterization

Out of 457 isolates of *Acinetobacter* species (*A. baumannii*, *A. iwoffii*, *A. junii*, *A. nosocomialis*, *A. pittii*), 357 (78.1%) isolates were ceftriaxone resistant and outstanding 100 (21.8%)

isolates were ceftriaxone susceptible. Among ceftriaxone resistant isolates, 26.9% (123/457) were ESBL producers and 51.2% (234/457) of population were MBL producers. The distribution of ESBL and MBL are illustrated in Table 2. Out of identified ESBL and MBL producers 20 strains (10 ESBL and 10 MBL) were picked randomly for gene characterization and it was observed that 3 were TEM-1 positive (2 *A. baumannii*; 1 *A. iwoffii*), 1 was SHV-1 positive (1 *A. baumannii*), 1 was AMP-C positive (*A. baumannii*), 2 were OXA-1 positive (1 *A. baumannii*, 1 *A. nosocomialis*), 3 were CTX-M positive (1 *A. junii* and 2 *Acinetobacter pittii*), 4 were VIM-1 positive (3 *A. baumannii*, 1 *Acinetobacter pittii*), 3 were IMP-1 positive (2 *A. baumannii*, 1 *A. junii*), 3 were NDM-1 (2 *A. baumannii*, 1 *A. iwoffii*) positive.

MIC₅₀ and MIC₉₀

In ceftriaxone, resistant *Acinetobacter* group, the MIC₅₀ and MIC₉₀ for ceftriaxone were 16 and 128 µg/ml and for Elores MIC₅₀ and MIC₉₀ were 4 and 8 µg/ml respectively, indicating 16 folds' reduction in MIC₉₀ of Elores compared to ceftriaxone. The ceftriaxone susceptible *Acinetobacter* isolates were susceptible to both ceftriaxone and Elores with MIC₅₀ and MIC₉₀ were ranging 2-4 µg/ml. The ceftriaxone susceptible *Acinetobacter* isolates, wild type strains and ESBL and MBL negative quality control strains were susceptible to both ceftriaxone and Elores with MIC₅₀ values were ranging 0.25 to 0.5 and MIC₅₀ and MIC₉₀ values were identical for both drugs for ceftriaxone susceptible isolates that is 4 and 2 µg/ml (Table 3, Table 4, and Table 5).

MIC and AST breakpoint determination

As per PK-PD study, the MIC breakpoint values for ceftriaxone and sulbactam were 13.5 and 7.7 µg/ml respectively. Since, the MIC breakpoint values fall between the MIC of 8 and 4 µg/ml, hence MIC breakpoint value of 8 µg/ml was selected to categorize susceptible (Table 6). Ceftriaxone shows time dependent killing of bacteria, hence percentage of time above the MIC (%T>MIC) was used to set a breakpoint between resistant and intermediate pathogens. As depicted in (Table 7), %T>MIC of ceftriaxone was 68.58 and 43.58 at the MIC of 32 and 64 µg/ml, respectively. In Elores, %T>MIC was noted to be below 50%, pressing the synergy of all components in Elores (Table 8). Hence the MIC cut off 16-32 µg/ml and ≥64 µg/ml were used to categories intermediate and resistant.

AST percentage for isolates are depicted in Figure 1. To perform regression analysis, a graph was plotted between MIC and AST values of Elores against *Acinetobacter* species (Figure 2). Least square correlation analysis was performed to obtain regression line and correlation coefficients (Figure 3). The AST cut off for Elores against *Acinetobacter* species was ≥21 mm (susceptible), 14-20 mm (intermediate) and ≤13 mm (resistant). Detailed AST breakpoint of AAE, are depicted in Figure 4. AST breakpoint of AAE, are depicted in Figure 4. The AST and MIC breakpoints of all the bacterial species are shown in (Table 9). The AST and MIC cut off for Elores against ceftriaxone resistant was same as of ceftriaxone susceptible.

DISCUSSION

Acinetobacter species are a common nosocomial infection and is one of the most challenging healthcare-associated burden. Infections due to this bug often lead to significant

mortality and morbidity and is a serious concern including in intensive care units.⁴¹ Based on substantial studies it has been found that resistance for *Acinetobacter* species is increasing worldwide, not only to third generation cephalosporins but also

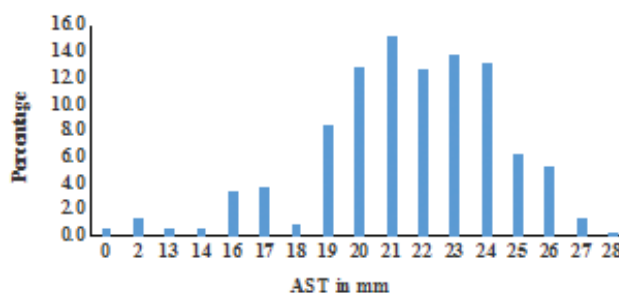


Figure-1: Elores AST percentage in ceftriaxone resistant *Acinetobacter* strains.

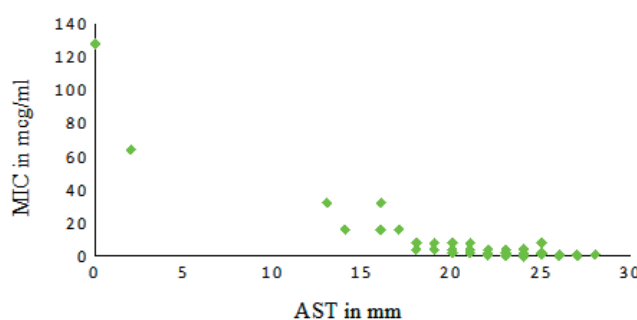


Figure-2: Scatter plot for Elores in ceftriaxone resistant *Acinetobacter* strains.

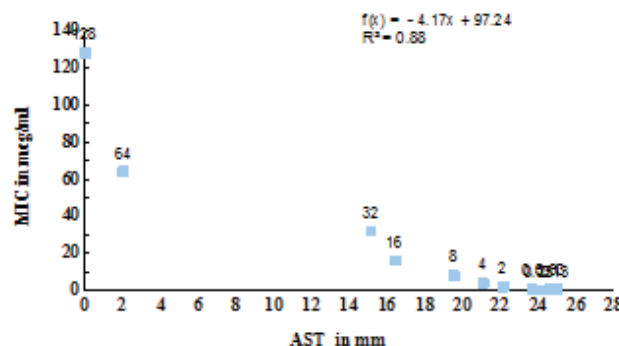


Figure-3: Regression plot for Elores in ceftriaxone resistant *Acinetobacter* strains.

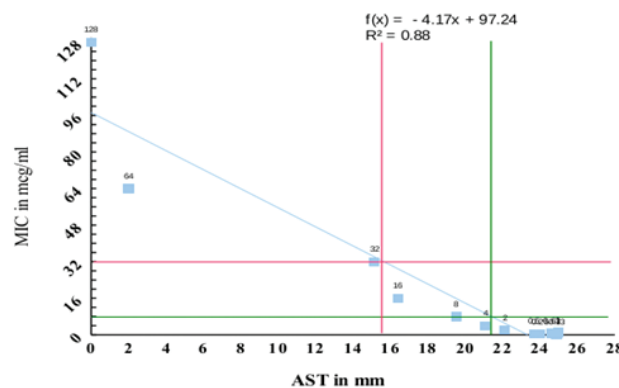


Figure-4: Breakpoints for Elores in ceftriaxone resistant *Acinetobacter* strains.

to multidrug categories of antibiotics restricting the treatment options.⁴² *Acinetobacter* species has been seen to be resistant to beta-lactams in both ESBL and MBL producers.⁴³ The present scenario of increasing resistance due to ESBL and MBL amongst *Acinetobacter* species lead us to restrict our study in details in terms of resistance profile with ceftriaxone and Elores. Even though, addition of beta-lactamase inhibitor to parent

beta-lactam can counter some of the ESBL infections, but for MBL infections, there is no effective alternative available. Therefore, the objective of current study was not only to evaluate the breakpoint sin ESBL producing Ceftriaxone resistant *Acinetobacter* spp but also in MBL producing resistnat strains isolated from clinical settings along with known positive controls.

Primer	Primer sequences (5'-3')	Amplicon (base pair)	References
TEM-1	F-5' CTGGGAAACGGAAGTGAATG-3' R-5' GGGGTATCCCGCAGATAAAT-3'	858	52
SHV-1	F-5'ATGAGTATTCAACATTCCG-3' R-5' CCAATGCTTATTCAGTGAGG-3'	308	52
AmpC	F-5CCC CGCTTA TAGAGC AAC AA-3 R-5TCA ATG GTC GAC TTC ACA CC-3	634	53
OXA-2	5'- GCCAAAGGCACGATAGTTGT-3' 5'- GCGTCCGAGTTGACTGCCGG-3'	701	54
CTX-M-1	F:5' GACGATGTCACCTGGCTGAGC 3' R: 5'AGCCGCCGACGCTAATACA 3'	499	54
NDM-1	F- 5'GGTTTGCGATCTGGTTTTC-3' R- 5'CGGAATGGCTCATCAGATC-3'	621	55
IMP-1	F- 5'GGAATAGAGTGGCTTAAYTCTC- 3' R -5'CCAAACYACTASGTTATCT- 3'	188	56
VIM-1	F- 5'TTATGGAGCAGCAACGATGT-3' R- 5'CAAAAGTCCCGTCCAACGA-3'	920	57

Table-1: Oligonucleotides used in the study for each tested genes

Characterization	Types	No of isolate/total	Percentage
Ceftriaxone resistant	ESBL	123/457	26.9
	MBL	234/457	51.2
Ceftriaxone susceptible	Non-ESBL+MBL	100/457	21.8

Table-2: Prevalence of ESBL, MBL, and non-ESBL and MBL among *Acinetobacter* clinical isolates

	No of pathogens	Ceftriaxone		Elores	
		MIC ₉₀	MIC ₅₀	MIC ₉₀	MIC ₅₀
<i>Acinetobacter</i> (Ceftriaxone resistant strains)	357	128	16	8	4
<i>Acinetobacter</i> (Ceftriaxone susceptible strains)	100	4	2	4	2

Table 3: MIC₅₀ and MIC₉₀ values of ceftriaxone and Elores against *Acinetobacter* isolates.

Organisms	Number of occurrence at MIC (µg/ml) of Elores												
	0.0625	0.125	0.25	0.5	1	2	4	8	16	32	64	128	256
<i>E. coli</i> (ATCC 25922) (ESBL-)	6	-	-	-	-	-	-	-	-	-	-	-	-
<i>E. coli</i> (ATCC35218) (TEM+)	-	1	5	-	-	-	-	-	-	-	-	-	-
<i>K. pneumoniae</i> (ATCC700603) (SHV+)	-	1	5	-	-	-	-	-	-	-	-	-	-
<i>K. pneumoniae</i> ATCC BAA-2146 (NDM-1)	-	-	-	-	-	-	-	6	-	-	-	-	-
<i>K. pneumoniae</i> NCTC 13439 (VIM-1)	-	-	-	-	-	-	-	6	-	-	-	-	-
<i>E. coli</i> NCTC 13476 (IMP)	-	-	-	-	-	-	-	6	-	-	-	-	-
<i>P. aeruginosa</i> (ATCC27853) (MBL-)	-	-	-	-	6	-	-	-	-	-	-	-	-
Number of occurrence at MIC (µg/ml) of Ceftriaxone													
Organisms	0.0625	0.125	0.25	0.5	1	2	4	8	16	32	64	128	256
<i>E. coli</i> (ATCC 25922) (ESBL-)	5	1	-	-	-	-	-	-	-	-	-	-	-
<i>E. coli</i> (ATCC35218) (TEM+)	-	-	-	-	1	5	-	-	-	-	-	-	-
<i>K. pneumoniae</i> (ATCC700603) (SHV+)	-	-	-	-	-	1	5	-	-	-	-	-	-
<i>K. pneumoniae</i> ATCC BAA-2146 (NDM-1)	-	-	-	-	-	-	-	-	-	-	5	1	-
<i>K. pneumoniae</i> NCTC 13439 (VIM-1)	-	-	-	-	-	-	-	-	-	-	3	3	-
<i>E. coli</i> NCTC 13476 (IMP)	-	-	-	-	-	-	-	-	-	-	4	2	-
<i>P. aeruginosa</i> (ATCC27853) (MBL-)	-	-	-	-	-	-	-	-	3	3	-	-	-

Table-4: MIC values for quality control strains.

In the current investigation, 457 isolates of *Acinetobacter* species (*A. baumannii*, *A. iwoffii*, *A. junii*, *A. nosocomialis*, *A. pittii*) were used and susceptibility behaviour of Elores and ceftriaxone on these clinical isolates were evaluated in terms of MIC₅₀ and MIC₉₀, as both are important parameters of presenting susceptibility. These data indicate that >90% of the organisms were inhibited at MIC 8 µg/ml with Elores which is indicative of its susceptibility towards *Acinetobacter* resistant isolates. This is also in agreement with the “90-60 rule” which means 90-95% of the time therapy would respond,⁴⁴⁻⁴⁵ hence showing a good outcome for Elores. Earlier studies also demonstrate a higher susceptibility of Elores against resistant *Acinetobacter* isolates.^{30,31-32}

MIC breakpoints have been used to categorize susceptibility and resistant behaviour of any drugs.⁴⁶ It relies on pharmacokinetic-pharmacodynamic (PK/PD) variables such as protein binding, half-life, volume of distribution. The PK/PD data for AAE was calculated using earlier published paper.⁴⁷ The PK/PD data for AAE was calculated using earlier published paper.^{47,48} The regression analysis, a graph was plotted between MIC and AST values, showed equal to or near to 0.9 regression coefficients (R²) suggesting strong relationship between MIC values and AST distribution.

<i>A. baumannii</i> (MTCC1425) (MIC µg/ml)
≤1
Table-5: MIC value of Elores in MTCC strain

PK-PD indices define the activity of an antibiotic is of great importance not only in treating infections but also to combat antibiotic resistance. For cephalosporin, %T>MIC PK/PD parameters are good predictors of antibiotic efficacy,⁴⁹ but scanty knowledge is available for AAEs.⁵⁰⁻⁵¹ In Elores, ceftriaxone shows time dependent killing of bacteria, hence percentage of time above the MIC (%T>MIC) was used to set a breakpoint between resistant and intermediate pathogens. The %T>MIC of ceftriaxone was 68.58 and 43.58 at the MIC of 32 and 64 µg/ml respectively. The MIC 32 µg/ml is the highest MIC value at which antibiotic concentration remains above MIC for more than 50% of the time, thus selected as a MIC breakpoint to categorize intermediate and resistance isolates. Elores has shifted the MIC distribution values, suggesting higher efficacy of the it at lower concentrations. Sulbactam is disposed along with ceftriaxone and EDTA to inhibit beta-lactamase enzyme.

CONCLUSION

Elores shows higher susceptibility towards ceftriaxone resistant *Acinetobacter* isolates while ceftriaxone failed to respond such strains because of degradation. The MIC cut off for Elores against ceftriaxone resistant *Acinetobacter* was ≤8 µg/ml (susceptible), 16-32 µg/ml (intermediate) and ≥64 µg/ml (resistant). From this study, it appears that Elores could be a better alternative than ceftriaxone alone for combating ESBL and MBL producing resistant pathogens which synergistically provides clinically relevant concentration at higher susceptible

MIC breakpoint for susceptible and intermediate = $C_{max} * f * s/e * t$							
Elores component	C _{max}	F	s	e	T	Breakpoint value	Cut off
Ceftriaxone	135	0.2	1	4	0.5	13.5	>8
Sulbactam	31.1	1	1	4	1	7.77	>4
Table-6: Calculation of MIC breakpoint of Elores 1.5 g for susceptible organisms.							

%T> MIC = $\ln(\text{Dose}/(\text{Vd} * \text{MIC})) * (t1/2/0.693) * (100/\text{DI})$							
MIC_ Elores (µg/ml)	MIC_of corresponding Ceftriaxone content (µg/ml)	Dose (mg)	V _d (L)	T1/2 (hrs)	DI (hrs)	%T> MIC	
4	2.67	1000	7	6	24	143.60	
8	5.34	1000	7	6	24	118.59	
16	10.68	1000	7	6	24	93.59	
32	21.34	1000	7	6	24	68.58	
64	42.69	1000	7	6	24	43.58	
Table-7: Calculation of MIC breakpoint for resistant isolates as per ceftriaxone component of Elores 1.5 g OD.							

%T> MIC = $\ln(\text{Dose}/(\text{Vd} * \text{MIC})) * (t1/2/0.693) * (100/\text{DI})$							
MIC_ Elores (µg/ml)	MIC_of corresponding Sulbactam content (µg/ml)	Dose (mg)	V _d (L)	T1/2 (hrs)	DI (hrs)	%T> MIC	
4	1.33	500	20	1.5	24	26.44	
8	2.67	500	20	1.5	24	20.19	
16	5.34	500	20	1.5	24	13.94	
32	10.68	500	20	1.5	24	7.77	
64	21.34	500	20	1.5	24	1.52	
Table-8: Calculation of MIC breakpoint for resistant strains as per sulbactam component of Elores 1.5 g OD.							

Bacterial Strain	AST breakpoints (mm)			MIC breakpoints (µg/ml)		
	Susceptible	Intermediate	Resistant	Susceptible	Intermediate	Resistant
<i>Acinetobacter</i> (ceftriaxone susceptible)	≥21	14-20	≤13	≤8	16-32	≥64
<i>Acinetobacter</i> (ceftriaxone resistant)	≥21	14-20	≤13	≤8	16-32	≥64
Table-9: MIC and AST breakpoints of Elores used to segregate different bacterial isolates in susceptible, intermediate, and resistant						

MIC values as observed through current study.

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