

**Antibiotic Susceptibility profile and prevalence Pattern of of  
Gram negative pathogens in Tertiary Care Hospital**

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25       **Abstract**

26       ***Background and objective***

27       Large amounts of antibiotics consumed by the human population has resulted in the  
28       culmination of pathogenic bacteria resistant to multiple drugs. The resistance profile of  
29       pathogens differ from one geographical location to another and keeps on changing  
30       continuously.

31       ***Methods***

32       A retrospective observational analysis of antibiogram data was performed to characterize  
33       the susceptibility pattern of different pathogen isolates from various clinical sources. A  
34       total of 213 clinical isolates identified from the period June 2015 to June 2016 were  
35       included in the study.

36       ***Results***

37       Of the 213 Gram negative isolates, 36.6% were from urine, 23.9% from respiratory  
38       specimens, 11.74% from blood, 10.33% from pus whereas 17.37% were from other  
39       sources. *E. coli* (42.25%) was most predominant pathogen isolated followed by *K.*  
40       *pneumoniae* . (25.35%) and *Pseudomonas spp.* (15.96%) while other Gram negative  
41       pathogens contributed 16.4%. Antibiogram analysis has shown CSE-1034 as the most  
42       susceptible drug exhibiting 91.1%, 77.8%, 82.4% and 82.3% susceptibility against *E. coli*,  
43       *K. pneumoniae*, *A. baumannii* and *P. aeruginosa*. Among carbapenems, both meropenem  
44       and imipenem-Cilastin were most effective against *E. coli*. Meropenem was least effective  
45       against *K. pneumoniae* (50%) and imipenem against                    *P. aeruginosa* (32.35%).  
46       Like imipenem, Piperacillin-Tazobactam was highest effective against *E. coli* (20%) and  
47       lowest against *P. aeruginosa* (26.47%).

48

49       ***Conclusion***

50 Susceptibility profile indicates CSE-1034 (a novel antibiotic resistance breaker) as the  
51 most susceptible drug among all the classes of antibiotics against the Gram-negative  
52 pathogens. A high resistance to piperacillin-tazobactam and penems, advocates use of  
53 CSE-1034 as empiric drug of choice in treatment of bacterial infectious diseases where  
54 the pathogen isolates are suspected resistant towards  $\beta$ -lactam and  $\beta$ -lactamase inhibitor  
55 combinations.

56 **Keywords:** Antibiotic, Clinical isolates, CSE-1034, Prevalence, Susceptibility,  
57 Resistance.

58

## 59 **1. Introduction**

60 The emergence of resistance among pathogenic bacteria towards potent antimicrobial  
61 agents has become a critical problem in modern medicine [1]. WHO has warned that the  
62 level of resistance to drugs used to treat common infectious diseases is arriving at a crisis  
63 point and if not controlled, entire population could be wiped out by these superbugs [2].  
64 The developing resistance towards currently available drugs increases the economic  
65 burden on the community by increasing the rates of hospitalization, length of hospital  
66 stays and cost of treatment [3] [4][5]. The rising antimicrobial resistance among the most  
67 common opportunistic Gram negative pathogens, are also associated with increased  
68 mortality and morbidity rates [2].

69  $\beta$ -lactam antibiotics used to be the most common treatment for bacterial infections but  
70 the constant exposure of bacteria to  $\beta$ -lactams drugs has created a selective pressure  
71 leading to ESBL and MBL producing strains. [6]. In past few years, a significant increase  
72 in the prevalence of ESBL, MBL and Carbapenemase producing strains has been  
73 observed throughout the globe [7]. These beta-lactamase producing Gram-negative  
74 pathogens are reported resistant to other classes of antibiotics also [12,13].

75 Taking into account such a situation, there is a need to optimize the antibiotic therapy  
76 against multidrug-resistant pathogens which may vary from one geographical locale to  
77 another. Surveillance data and hospital antibiogram profiles help clinicians in the  
78 prescription of appropriate antimicrobial therapy. Therefore, we aimed to study the  
79 susceptibility profile of clinical isolates collected from Noble Hospital, Pune towards  
80 commonly used 2<sup>nd</sup> line antibiotics including Ceftriaxone/Sulbactam/EDTA,  $\beta$ -lactam and  
81  $\beta$ -lactamase inhibitor combination (Piperacillin-tazobactam) and Carbapenems  
82 (meropenem and imipenem-cilastatin) drugs.

## 83 **2. Materials and Methods**

### 84 ***2.1 Sample collection***

85 Various clinical specimens used for pathogen isolation included urine, stool, blood, pus,  
86 endotracheal tube secretions (ETT), tracheal tube (TT) secretions, sputum, wound, , gall  
87 bladder specimens, abscess, drain, ear swab, vitreous eye, abdominal fluid, vitreous fluid,  
88 semen, peritoneal fluid and tissue specimens collected from 347 infected patients at Noble  
89 Hospital, Pune (India), during the period of July 2016 to February 2017. The collection  
90 and processing of the samples were done as per common standard operating procedures of  
91 hospital.

### 92 ***2.2 Isolation and Identification of microbes***

93 All the samples were collected aseptically in sterile containers and inoculated on the  
94 different selective and non-selective culture media as per the standard microbiological  
95 techniques. Details of the culture media used for the isolation of pathogens from various  
96 clinical samples are given in Table 1. Blood samples were collected in Bactec bottles and  
97 incubated in Bactec machine. These samples were further sub-cultured on the selective or  
98 non-selective media and incubated aerobically overnight at 37°C. Organisms were  
99 identified on the basis of colony morphology, gram staining, motility and biochemical

100 reactions. Biochemical reactions were performed by inoculating the bacterial colony in a  
101 nutrient broth at 37°C for 2– 3 hours [15].

### 102 **2.3 Antibiotic susceptibility testing**

103 Antimicrobial susceptibility study was performed by Kirby–Bauer disk diffusion method  
104 as recommended by the Clinical Laboratory Standards Institute (CLSI) guidelines [16]. In  
105 brief, inoculum of 0.5 McFarland standards turbidity was prepared in a Mueller-Hinton  
106 broth (MHB, Hi-Media, Mumbai, India) from isolated colony of pathogens selected from  
107 18–24 hour agar plates. A sterile cotton swab was dipped into the inoculum and streaked  
108 many times on the dried surface of a Mueller-Hinton agar (MHA) plate. After 5 minutes,  
109 antibiotic discs were applied and pressed down to check absolute contact with agar  
110 surface. The discs were apportioned in a minimum distance of 24 mm from the center.  
111 The plates were then incubated for 16-18 hrs aerobically at 37° C. The discs of  
112 meropenem (10 µg), imipenem-cilastatin (20 µg) and piperacillin-tazobactam (110 µg)  
113 were obtained from Microexpress Goa, India and CSE-1034 (45 µg) was obtained from  
114 third party.

### 115 **3. Results**

116 A total of 347 clinical specimens were obtained from the suspected patients out of  
117 which 213 (61.38%) clinical samples tested positive for Gram negative pathogens. Out of  
118 these 213 Gram negative isolates, the maximum isolates were obtained from urine  
119 specimens (36.62%) followed by respiratory specimens (13.62%), blood (11.74%), pus  
120 (10.33%) and wound (6.10%) while all other samples contributed a total of 6.58% (Table  
121 2).

122 On the basis of morphological and biochemical screening, eight bacterial species were  
123 obtained including *E. coli*, *K. pneumoniae*, *P. aeruginosa*., *A. baumannii*, along with other  
124 less prevalent Gram negative bacilli such as *Proteus spp.*, *Salmonella spp.*, *Serratia spp.*

125 and *Enterobacter spp.* which contributed 8.45% (% cumulatively) to the total clinical  
126 isolates. The detailed profile of various pathogens isolated from clinical specimens is  
127 shown in Table 3.

128 Table 2 represents the prevalence of different clinical isolates in different samples.  
129 Data revealed the maximum prevalence of *E. coli* in urine samples, pus and stool samples.  
130 *K. pneumoniae* was mostly isolated from blood and respiratory specimens whereas  
131 *P. aeruginosa* isolates were mostly retrieved from wound and respiratory specimens. *A.*  
132 *baumannii* was least prevalent in all the specimens.

133 Susceptibility profile of pathogens isolated from clinical specimens is presented in  
134 Table 4. Overall, 85.4% (182) of the total number of isolates were reported susceptible to  
135 CSE-1034, 59.6% (127) to Pip-taz, 66.2% (141) to Meropenem and 64.8% (138) to  
136 Imipenem. The susceptibility rates of CSE-1034 were *E. coli* (91.9%), *K. pneumoniae*  
137 (77.8%), *A. baumannii* (82.4%) and *P. aeruginosa* (82.3%). Among all the antibiotics  
138 tested, the least susceptibility was reported to Pip/Taz. *E. coli* exhibited the highest  
139 susceptibility (80%) to Pip-taz whereas the lowest was reported by *P. aeruginosa*  
140 (26.47%). Among Carbapenems, almost similar susceptibility of meropenem and  
141 imipenem-cilastin was reported against *E. coli* (77-82%) and *K. pneumoniae* (48-50%).  
142 The meropenem was marginally better than imipenem-cilastin against *A. baumannii*  
143 (58.8% vs 52.9%) whereas was reported significantly better than imipenem-cilastin  
144 against *P. aeruginosa* (58.8% vs 32.3%).

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146

#### 147 **4. Discussion**

148 The predominant species isolated was *E. coli* (42.2%) followed by *K. pneumoniae*  
149 (25.3%). A good number of studies have reported *E. coli* and *K. pneumoniae* as most

150 common and opportunistic clinical pathogens [17, 18]. Similar results with high  
151 prevalence of *E coli* (54.9%) were reported by Sikka *et al.* [19]. Sachdeva [20] has also  
152 reported the prevalence of *E. coli* to a tune of 51.7%. A similar prevalence of *K.*  
153 *pneumoniae* has been reported by Makkar *et al.* [21] who demonstrated 22% of *K.*  
154 *pneumoniae* from clinical isolates. Sahu *et al.* [22] reported the prevalence of *K.*  
155 *pneumoniae* to a tune of 32% which sustains our data. *Pseudomonas spp.* (15.9%) also  
156 contributed significantly to the isolated pool of pathogens. As compared to other studies,  
157 less number of *A. baumannii* isolates were identified in this study. The similar prevalence  
158 pattern of *Proteus spp.*, *Salmonella spp.*, *Serratia spp.* and *Enterobacter spp* is also  
159 reported by many other studies. [17, 24, 25]

160 Similar to our observations, Ruppe *et al.* [14] have also reported 90% prevalence of *E.*  
161 *coli* in stool samples. Majority of *E. coli* (54%) isolates were recovered from urine during the  
162 study performed by Kumar *et al.* [17]. Ibrahim *et al.* [26] have also reported 40-50%  
163 prevalence of *E. coli* in pus samples. *K. pneumoniae* isolates were mostly isolated from  
164 blood and respiratory specimens whereas *P. aeruginosa* was mostly isolated from wound and  
165 respiratory specimens.

166 Among all the antibiotics tested, the least susceptibility was reported to Pip/Taz and  
167 highest was reported towards CSE-1034. A high rate of resistance observed to Pip/taz which  
168 is normally recommended second line of treatment in our hospital could be possibly the  
169 indiscriminate consumption of pip-taz. The AMR surveillance study conducted in India has  
170 shown resistance against pip-taz has risen to 65-70% [19]. Among carbapenems, the average  
171 susceptibility rates were 65% against all the pathogen isolates. The emergence of  
172 carbapenem-resistant strains, which ranges from 18-68% in different isolates is a matter of  
173 big concern as carbapenems are considered as the last resort drugs for MDR bacterial

174 infections. Singh *et al.* [11] have reported that MBLs to a tune of 15-22% among the Gram-  
175 negative isolates in their study.

176 The high rate of carbapenem resistant strains reported in this surveillance study is a  
177 matter of grave concern and needs to be addressed on priority at the global level. One of the  
178 approaches that the clinicians have adopted to reduce selective pressure on last resort drugs is  
179 by pumping in the use the antibiotic resistance breakers “ARBs” along with antibiotics to  
180 revive them for clinical purposes. CSE-1034 is one such combination of beta-lactam/beta-  
181 lactamase inhibitor (BL/BLI) combination with ARB “EDTA”. Interestingly, a significant  
182 number of isolates were sensitive to CSE-1034 i.e., *E. coli* (98.8%), *K. pneumoniae* (90.5%),  
183 *P. aeruginosa* (89.9%) and *Acinetobacter spp* (81%). Surprisingly, 131 isolates reported as  
184 Meropenem resistant were susceptible to CSE-1034 (Table 4). The higher susceptibility of  
185 Gram-negative pathogens to CSE-1034 has been reported by several other studies also. CSE-  
186 1034 is a novel combination of Ceftriaxone, Sulbactam and disodium edetate and the high  
187 susceptibility of CSE-1034 could be attributed to the synergistic effect of Ceftriaxone,  
188 disodium edetate and Sulbactam. The non-antibiotic adjuvant, EDTA mediates various  
189 antimicrobial effects by enhancing the penetration of antibiotic into cell membrane, decreases  
190 over-expression of efflux pumps, bio-film eradication, de-activates carbapenemases-MBL by  
191 chelating Zinc ions.

192 About last line therapy agents for MDR infections, our study has shown Carbapenems  
193 as the most active agent only against *E.coli* (82%). Around 36-45% of *P. aeruginosa* and 45%  
194 of *Acinetobacter spp.* were Carbapenem resistant. Resistance to meropenem was found  
195 highest in *Klebsiella spp.* (54%). Chauhan *et al.* [19] have reported a Carbapenem resistance  
196 of 14.6% in *E. coli* and 29.6% in *Klebsiella spp.* in hospital isolates from various in and  
197 outpatient areas. Gupta *et al.* [21] have reported a Carbapenem resistance ranging from 17-  
198 22% in different strains of Enterobacteriaceae from North India.



199 Based on pathogen type, *E. coli* exhibited the highest susceptibility rate whereas the  
200 lowest was reported against *P. aeruginosa*. *E. coli* was found to be most susceptible clinical  
201 isolate among major pathogens which displayed 80%, 75.5% and 82.2% sensitivity against  
202 piperacillin-tazobactam, meropenem and imipenem-cilastatin respectively. Correspondent  
203 results were observed by many authors who reported 100%, 90% and 96.5% sensitivity of *E.*  
204 *coli* against meropenem, piperacillin-tazobactam and imipenem-cilastatin respectively  
205 [35,36]. *Klebsiella spp.* exhibited intermediate susceptibility i.e. 44.4%, 50% and 48.1%  
206 towards piperacillin-tazobactam, meropenem and imipenem-cilastatin. Similar results were  
207 noted by many authors who revealed 40-60% sensitivity of *Klebsiella spp.* against  
208 piperacillin-tazobactam, meropenem and imipenem-cilastatin [37,38]. As reported earlier  
209 also, *Acinetobacter spp.* experienced highest susceptibility (96.3%) towards CSE-1034CSE-  
210 1034 only while extreme resistance (96.3% each) against rest of the antibiotics which is due  
211 to sulbactam (a  $\beta$ -lactamase inhibitor) which owns intrinsic whole-cell activity against  
212 *Acinetobacter spp.* [39]. Surprisingly, contrary to expectations, *Pseudomonas spp.*  
213 documented 73.5%, 73.5% and 67.6% resistance against piperacillin-tazobactam, meropenem  
214 and imipenem-cilastatin respectively. Mohammadi and Feizabadi [37] reported >60%  
215 resistance of piperacillin+tazobactam against gram negative bacilli isolated from clinical  
216 samples which supports our data. Similarly, Hout *et al.* [40] revealed 70-100% resistance of  
217 meropenem towards *Acinetobacter spp.* and *Pseudomonas spp.*. Likewise, Shour and El-  
218 Sharif, [36] and Eldomany and Abdelaziz noticed significant resistance (>50%) of  
219 imipenem-cilastatin in *Acinetobacter spp.* and *Pseudomonas spp.* which is in accordance with  
220 our present data [41].

221 The emergence of antimicrobial resistance against BL-BLI and carbapenem drugs is due to  
222 numerous elements which assists the scattering of resistance among clinical pathogens which  
223 includes production of MBL enzymes, biofilm formation, over expression of efflux pumps

224 and accumulation of the drug [42,43]. None of these mechanisms are dressed by either  
225 piperacillin-tazobactam, meropenem or imipenem-cilastatin and probably this could be one  
226 reason for CSE-1034 super performance that it is supplemented with EDTA as ARB. The  
227 progressive and relentless resistance towards BL-BLI and carbapenem antibiotics is probably  
228 the result of overuse of antibiotics, improper processing and inappropriate prescribing [44]. In  
229 the light of above discussion it is evident that Antibiotic adjuvant therapy which has ARB can  
230 be used as the prime choice of therapeutics to overcome the resistance raised among gram  
231 negative pathogens towards  $\beta$ -lactam and  $\beta$ -lactamase inhibitor combinations and penems in  
232 the treatment of bacterial infectious diseases.

### 233 **5. Conclusion**

234 This retrospective study indicates the rise in resistance among most prevalent and  
235 opportunistic gram negative pathogens against  $\beta$ -lactam and  $\beta$ -lactamase inhibitor  
236 combinations and penems. Present data strongly advocates precedence of CSE-1034 over  $\beta$ -  
237 lactam and  $\beta$ -lactamase inhibitor combinations and penems as CSE-1034 has scored 85-100%  
238 susceptibility which excels the antimicrobial activity of rest of the drugs. Therefore, CSE-  
239 1034, a novel product with antibiotic resistance breaker can be used as an empirical and  
240 alternate choice of drug over potent therapeutics in encountering multidrug resistance among  
241 healthcare-associated pathogens.

### 242 **Ethical Approval:**

243

244 As per international standard or university standard written ethical approval has been collected and  
245 preserved by the author(s).

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374 **Table 1: Selective culture medium used for isolation of different pathogens.**

<b>Pathogen</b>	<b>Selective media</b>
<i>Klebsiella spp.</i>	Hicrome Klebsiella selective agar base medium
<i>E. coli</i>	Eosine Methylene Blue (EMB) agar medium
<i>Acinetobacter spp.</i>	Leeds acinetobacter agar base medium
<i>Pseudomonas spp.</i>	Citrimide agar medium
<i>Proteus spp.</i>	EMB agar medium
<i>Salmonella spp.</i>	Wilson and Blair bismuth sulphite medium
<i>Serratia spp.</i>	Caprylate-thallos agar medium
<i>Enterobacter spp.</i>	EMB agar medium

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384 **Table 2: A profile of clinical samples used as a source of the pathogenic isolates.**

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Sr. No.	Name of clinical samples	Total no. of samples collected	Number of samples showing growth of pathogens (%)	Number of samples not showing growth of pathogens
1	Urine	94	78 (36.62%)	16
2	Respiratory specimens	84	51	
3	Blood	39	25 (11.74%)	14
4	Pus	34	22 (10.33%)	12
5	Wound	22	13 (6.10%)	9
6	Stool	19	10 (4.69%)	9
7	Other samples	55	14 (6.58%)	41
	<b>Total</b>	<b>347</b>	<b>213 (61.38%)</b>	<b>134</b>

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**Table 3: Prevalence of different clinical isolates in different samples.**

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Samples	Clinical Isolates					
	No. of isolates	<i>E. coli</i> (%)	<i>Klebsiella spp.</i> (%)	<i>Acinetobacter baumannii</i> (%)	<i>Pseudomonas aeruginosa</i> (%)	Other pathogens (%)
Urine	78	59 (75.64)	11 (14.10)	0	7 (8.97)	1 (1.28)
Respiratory specimens	51	6 (3.45)	17 (31.03)	10 (44.83)	12 (23.52)	6 (11.76)
Blood	25	2 (8)	10 (40)	4 (36)	5 (20)	4 (16)
Pus	22	10 (45.45)	6 (27.27)	2 (9.09)	2 (9.1)	2 (9.1)
Wound	13	2 (15.38)	3 (23.08)	1 (7.69)	5 (38.46)	2 (15.38)
Stool	10	8 (80)	2 (20)	0	0	0

Other samples	14	3 (21.43)	5 (35.71)	0	3 (21.43)	3 (21.43)
<b>Total</b>	<b>213</b>	<b>90</b>	<b>54</b>	<b>17</b>	<b>34</b>	<b>18</b>
<b>Total (%)</b>		<b>42.25 %</b>	<b>25.35 %</b>	<b>7.98 %</b>	<b>15.96 %</b>	<b>8.45%</b>

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**Table 4: Susceptibility pattern of clinical isolates.**

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<b>Susceptibility (%)</b>									
<b>Clinical isolates</b>	<b>No. of isolates</b>	<b>AAE</b>		<b>BL-BLI</b>		<b>Carbapenem</b>			
		<b>CSE-1034</b>		<b>Piperacillin-tazobactam</b>		<b>Meropenem</b>		<b>Imipenem-cilastatin</b>	
		<b>S</b>	<b>R</b>	<b>S</b>	<b>R</b>	<b>S</b>	<b>R</b>	<b>S</b>	<b>R</b>
<i>E. coli</i>	<b>90</b>	91.1 (82)	8.9 (8)	(80) 72	(20) 18	77.7 70	22.3 22	82.22 74	17.78 16
<i>Klebsiella spp.</i>	<b>54</b>	77.8 (42)	22.2 (12)	(44.44) 24	55.56 30	50 27	50 27	48.15 26	51.85 28
<i>Acinetobacter baumannii</i>	<b>17</b>	82.4 (14)	17.6 (3)	35.3 6	64.7 11	58.8 10	41.2 7	52.9 9	47.1 8
<i>Pseudomonas aeruginosa</i>	<b>34</b>	82.3 (28)	17.7 (6)	26.47 9	73.53 25	58.8 20	41.2 14	32.35 11	67.65 23
Other pathogens	<b>18</b>	(88.9) 16	(9.91) 2	88.9 16	9.91 2	88.9 16	9.91 2	100 18	0
<b>Total</b>	<b>213</b>	(85.4)	(14.5%)	(59.6)	(40.4)	(66.2)	(33.8)	(64.8)	(35.2)

		182	31	127	86	141	72	138	75
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