

## ANTIBACTERIAL SUSCEPTIBILITY PROFILE OF *ESCHERICHIA COLI* IN A PRIVATE HOSPITAL, INDIA

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### ABSTRACT

*E.coli* is a freely available gram negative bacteria among the natural resources like gut of animals which at the same time emerges to be one of the most causative organisms for stomach and urine related infections in humans. The presence of Extended-spectrum beta-lactamase ESBL type of drug resistance is constantly evolving and are under dynamic flux posing a global threat to public health programs. Also, there is a significant necessity for regular antimicrobial sensitivity surveillance not only for the presence and spread of ESBL genes but also for both urban and rural populations for more informed and structured treatment. At an institutional level, practices that can minimize the spread of such organisms include clinical and bacteriological surveillance of patients admitted to intensive care units and antibiotic cycling, as well as policies of restriction, especially on the empirical use of broad-spectrum antimicrobial agents. The present study focuses on the trends of antibacterial susceptibility and resistance among clinically isolated *E.coli* for rational prescribing. Susceptibility and resistance data of *E.coli* were collected from a tertiary care hospital's Microbiology Department over a period of three years. The data collected included patient's source (ICU or ward), specimen of the isolate, antibacterial susceptibility and resistance profile. *E.coli* was identified from the sample on the basis of colony and microscopic morphology. Commercially available antibiotic disks were used for antimicrobial susceptibility testing as per Kirby-Bauer disk diffusion method and Clinical Laboratories Standard Institute CLSI guidelines. The pattern of antibiotics used within prescribing pattern for *E.coli* were found out to be 33.3%, 30.3% and 27.5% in the study period. The identification of ESBL, AmpC, Carbapenamase were carried out as they were identified to be as resistant strains and more complicated in terms of identification and treatment. Susceptibility to third generation Cephalosporins, Gentamycin, Imipenem, Meropenem, Amikacin showed a narrow increase in resistance level against *E. coli*.

**KEYWORDS:** *Escherichia Coli, ESBL, AmpC, Carbapenamase, Cephalosporins and Meropenem*

### INTRODUCTION

*Escherichia coli* (*E.coli*) is the most prevalent facultative anaerobic bacteria in the gastrointestinal tract of humans and animals. It is usually a harmless microbe, also causing a number of significant illnesses<sup>1</sup>. The discovery of *E.coli* as an 'emerging pathogen' was made in the same year when the association of sporadic cases of Hemolytic Uremic Syndrome (HUS) with Cytotoxin - producing fecal *E. coli* was been observed<sup>2</sup>. World Health Organization (WHO) grades *E.coli* as one of the major agents of concern

associated both with hospital and community acquired infections<sup>3</sup>. *E.coli* has come into existence in several countries as a cause of prevalent bloody and non-bloody diarrhea, HUS and Thrombotic Thrombocytopenic Purpura. This can lead to fatal complications that occur approximately at a rate of 5-10% of all cases<sup>4,5</sup>. The emergence of Extended-spectrum  $\beta$ -lactamases (ESBL) type of resistance offered by *E.coli* is supported by many reports<sup>6</sup>. This is a major threat to the already hostile community of physician in the hospitals who are looking for alternative and novel antibiotics to tackle ESBL type of infections<sup>7</sup>. ESBLs are a rapidly evolving group of  $\beta$ -lactamases which share

the ability to hydrolyze third-generation Cephalosporins and Aztreonam but are inhibited by Clavulanic acid. They represent the first example in which  $\beta$ -lactamase - mediated resistance to  $\beta$ -lactam antibiotics resulted from fundamental changes in the substrate spectra of the enzymes<sup>8</sup>. Identifying ESBL-producing organisms is a major challenge for the clinical microbiology laboratory. Multiple factors contribute to this, including production of multiple different  $\beta$ -lactamase types by a single bacterial isolate<sup>9</sup>. At an institutional level, practices that can minimize the spread of such organisms include clinical and bacteriological surveillance of patients admitted to intensive care units and antibiotic cycling; as well as policies of restriction, especially on the empirical use of broad-spectrum antimicrobial agents such as the third and fourth-generation Cephalosporins and Quinolones<sup>10-12</sup>. The present study focuses on the trends of antibacterial susceptibility and resistance among clinically isolated *E.coli*. The data gathered is anticipated for rational prescribing.

## MATERIALS AND METHODS

Susceptibility and resistance data of *E.coli* were collected from a tertiary care hospital's Microbiology Department over a period of three years (calendar year 2012, 2013 and 2014) in a prospectively designed data collection form. The data collected includes patient's source (ICU or ward), specimen of the isolate, antibacterial susceptibility and resistance profile. *E.coli* was

identified from the sample on the basis of colony, microscopic morphology and biochemical reactions<sup>13</sup>. Antimicrobial susceptibility testing was done by Kirby-Bauer disk diffusion method as recommended by the Clinical Laboratory Standards Institute (CLSI) guidelines<sup>14</sup>. Commercially available antibiotic disks were used for antimicrobial susceptibility testing. The antibiotic disks used are Ampicillin (10  $\mu$ g), Piperacillin (100  $\mu$ g), Piperacillin / Tazobactam(100/10 $\mu$ g), Amoxicillin / Clavulanic acid (20/10 $\mu$ g), Cefoperazone/Sulbactam (75/10  $\mu$ g), Ceftazidime/Clavulanate (30/10  $\mu$ g), Cefoperazone (75  $\mu$ g), Cefoxitin (30  $\mu$ g), Ceftazidime (30 $\mu$ g), Cefotaxime (30  $\mu$ g), Ceftriaxone (30  $\mu$ g), Cefepime (30  $\mu$ g), Aztreonam (30  $\mu$ g), Imipenem(10  $\mu$ g), Amikacin (30  $\mu$ g), Gentamycin (10  $\mu$ g), Ciprofloxacin (30  $\mu$ g), Ofloxacin (5  $\mu$ g),Norfloxacin (10  $\mu$ g), and Nitrofurantoin (300  $\mu$ g). The quality control of antibiotic sensitivity was done using *E.coli* ATCC 25922 and *E.coli* ATCC 35218 (for  $\beta$ -lactam/ $\beta$ -lactamase inhibit or combination).

## RESULTS

Specimen wise distribution of *E. coli* is given in Table – 1. Prevalence of *E. coli* among the positive cultures is found to be 33.3%, 30.3% and 27.5% in 2012, 2013 and 2014 respectively. This shows that the pattern of antibiotics used was well versed within the prescribing pattern for *E.coli*.

**Table 1**  
*Specimen wise distribution of E.coli*

| Specimen                          |      | 2012  |                                  | 2013  |                                  | 2014  |                                  |
|-----------------------------------|------|---|----------------------------------|---|----------------------------------|---|----------------------------------|
|                                   |      | Total Number of samples with positive culture | Number of <i>E.coli</i> isolates | Total Number of samples with positive culture | Number of <i>E.coli</i> isolates | Total Number of samples with positive culture | Number of <i>E.coli</i> isolates |
| Urine                             | Ward | 1247  | 724                              | 1332  | 714                              | 336   | 169                              |
|                                   | ICU  | 190   | 48                               | 249   | 83                               | 105   | 28                               |
| Pus/EENT Swabs/Stool/BF           | Ward | 614   | 110                              | 574   | 89                               | 237   | 58                               |
|                                   | ICU  | 117   | 17                               | 110   | 12                               | 45  | 7                                |
| Respiratory                       | Ward | 224   | 14                               | 301   | 17                               | 106   | 13                               |
|                                   | ICU  | 300   | 19                               | 420   | 18                               | 205   | 18                               |
| Blood                             | Ward | 301   | 72                               | 238   | 34                               | 90  | 30                               |
|                                   | ICU  | 191   | 58                               | 175   | 46                               | 77  | 8                                |
| <b>Total no. of <i>E.coli</i></b> |      | <b>1062</b>                                   |                                  | <b>1013</b>                                   |                                  | <b>331</b>                                    |                                  |

The identification of EBSL, AmpC, Carbapenemase were carried out as they were identified to be resistant strains and are more complicated in terms of identification and treatment. The EBSL strain substantially increased from 52.5 % to 72.3 % in the three years of study.

The AmpC and Carbapenemase strains showed a significant drift in 2013 but tend to increase in 2014 as shown in Table – 2. This suggests choosing the right antibiotics is required to minimize the increase of these strains.

**Table 2**  
*Distribution of E.coli and resistant strains*

|       | 2012                               |             |             |                      | 2013                   |             |             |                      | 2014                   |             |             |                      |
|-------|------------------------------------|-------------|-------------|----------------------|------------------------|-------------|-------------|----------------------|------------------------|-------------|-------------|----------------------|
|       | Total<br><i>E.coli</i><br>isolates | EBSL<br>(%) | AMPC<br>(%) | CARBAPENAMASE<br>(%) | Total<br><i>E.coli</i> | EBSL<br>(%) | AMPC<br>(%) | CARBAPENAMASE<br>(%) | Total<br><i>E.coli</i> | EBSL<br>(%) | AMPC<br>(%) | CARBAPENAMASE<br>(%) |
| Ward  | 920                                | 61          | 12          | 8                    | 386                    | 65          | 1           | 3                    | 51                     | 78          | 4           | 1.8                  |
| ICU   | 142                                | 44          | 12          | 5                    | 110                    | 63          | 1           | 8                    | 51                     | 66.6        | 5.8         | 2                    |
| Total |                                    | 52.5        | 12          | 6.5                  | 507                    | 64          | 1           | 5.5                  | 107                    | 72.3        | 4.9         | 1.9                  |

The susceptible profiles (in terms of % of isolates) of *E.coli* over three year period is given in Table – 3. Susceptibility to third generation Cephalosporins (Figure-1), Gentamycin (Figure-3), Imipenem, Meropenem (Figure-5), Amikacin (Figure-4) showed a narrow increase in resistance level against *E. coli*. The Fourth generation Cephalosporin,

Cefepime(Figure-2), showed a slight decrease in resistance when compared to 2012 and 2014. Even though, antibiotics like Colistin, Piperacillin/Tazobactam were lately introduced, it remains to be more susceptible to the gram negative bacteria

**Table 3**  
*Susceptible profile (in %) of E.coli*

| Year / Antibacterial               | 2012                 |                     |         | 2013                 |                      |         | 2014           |                    |         |
|------------------------------------|----------------------|---------------------|---------|----------------------|----------------------|---------|----------------|--------------------|---------|
|                                    | Ward<br>(N =<br>920) | ICU<br>(N =<br>142) | Average | Ward<br>(N =<br>386) | ICU<br>(N =<br>=110) | Average | Ward<br>(N=51) | ICU<br>(N =<br>51) | Average |
| FLUOROQUINOLONES                   | 37                   | 19                  | 28      | 21                   | 4                    | 12.5    | -----          | -----              | -----   |
| 3 <sup>RD</sup> GEN CEPHALOSPORINS | 36                   | 15                  | 25.5    | 21                   | 4                    | 12.5    | 16             | 25                 | 20.5    |
| CEFIPIME                           | 36                   | 19                  | 22.5    | 28                   | 20                   | 24      | 16             | 25                 | 20.5    |
| GENTAMYCIN                         | 48                   | 40                  | 44      | 44                   | 48                   | 46      | 40             | 57                 | 48.5    |
| AMIKACIN                           | 93                   | 63                  | 78      | 29                   | 29                   | 29      | 96             | 98                 | 97      |
| CEFOPERAZONE/SULBACTAM             | 83                   | 75                  | 79      | 85                   | 82                   | 83.5    | 87             | 88                 | 87.5    |
| IMIPENAM                           | 95                   | 92                  | 93.5    | -----                | -----                | -----   | 98             | 98                 | 98      |
| MEROPENEM                          | 95                   | 92                  | 93.5    | 55                   | 48                   | 51.5    | 98             | 98                 | 98      |
| AMOXYCILLIN                        | -----                | -----               | -----   | 8                    | 9                    | 8.5     | 10             | 12                 | 16      |
| AMOXYCILLIN/CLAVULANATE            | -----                | -----               | -----   | 32                   | 46                   | 39      | 40             | 55                 | 47.5    |
| PIPERACILLIN/TAZOBACTAM            | -----                | -----               | -----   | -----                | -----                | -----   | 71             | 75                 | 73      |
| COLISTIN                           | -----                | -----               | -----   | -----                | -----                | -----   | 100            | 100                | 100     |
| CO-TRIMAXAZOLE                     | -----                | -----               | -----   | 1                    | 4                    | 2.5     | 0.7            | 2                  | 1.35    |

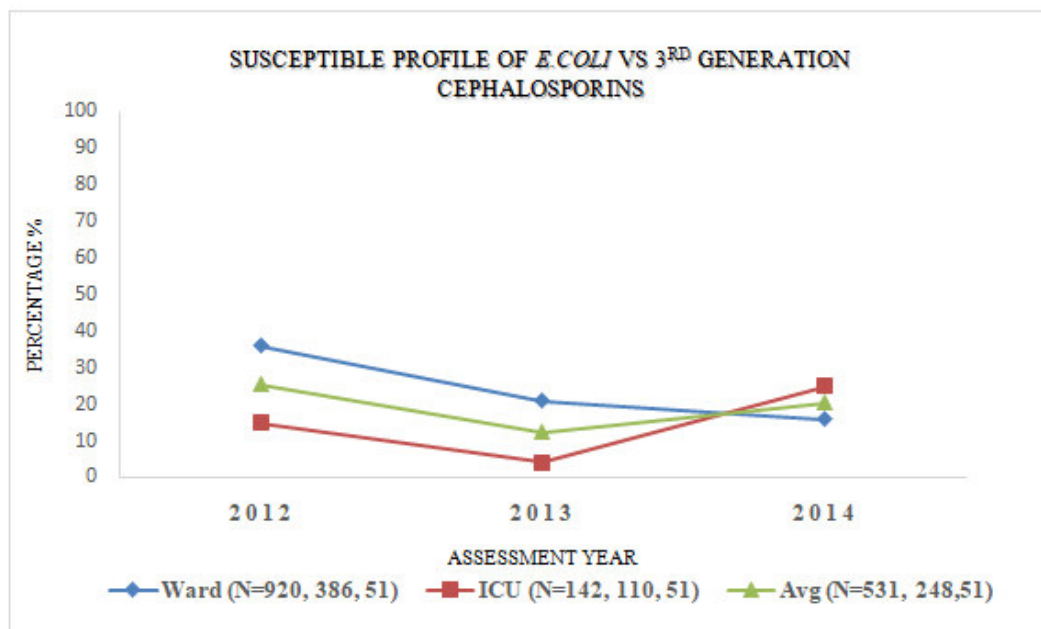
## DISCUSSION

The *E.coli* is a freely available gram negative bacteria among the natural resources. It is also found in the cattle and human gut flora. At the same time, it emerges to be one of the most causative organism for stomach and urine related infections in both children and adults. The rise of EBSL has become a growing threat in the clinics to identify and treat the resistant strains. A three year study was

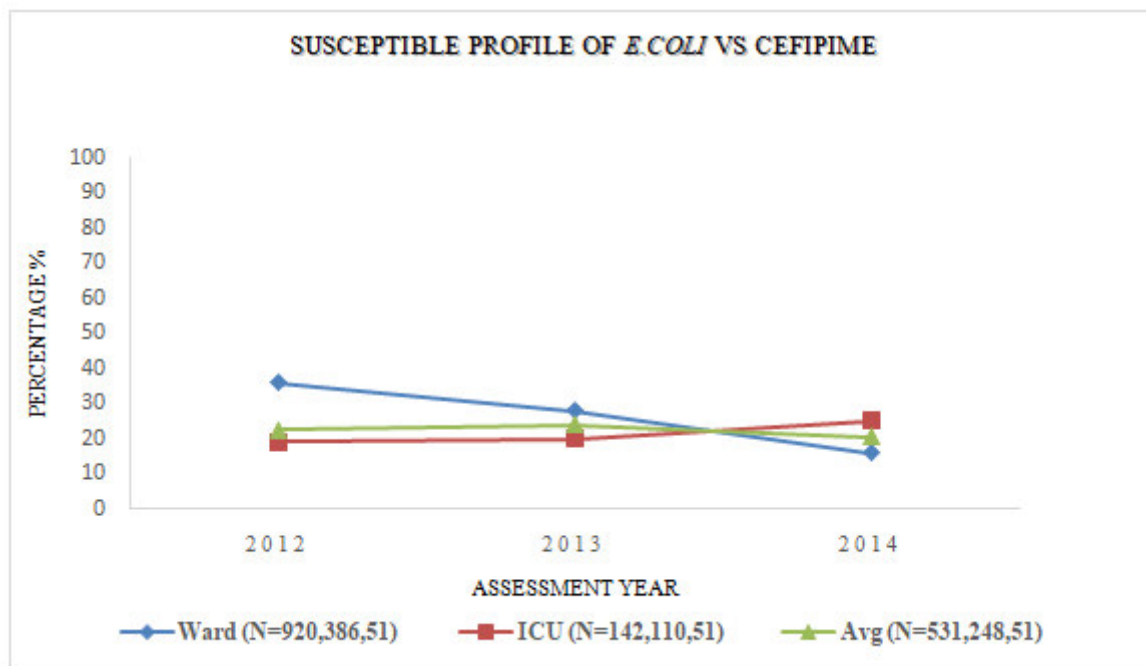
carried out to identify the antibacterial strains prominent in the ICU and wards. The study showed a major increase in the EBSL strains in comparison to AmpC and Carbapenemase strains. This can worsen the disease condition in the patients infected and can also be fatal. The increase in prevalence needs to be addressed at the earliest through creating awareness and sensitizing the prescribers regarding the threat of ESBL strains. The study conducted by Kibret *et.al* showed an increased number of positive cultures in urine sample which

was similar to our study<sup>15</sup>. The gram negative bacteria are prone to cause urinary tract infection and thus are found widely in urine samples. In another study taken up by Ugwu *et.al* found that in stool samples EBSL strains were mostly present among the *E.coli* strains. In the present study, the EBSL strains were mostly present along with AmpC and Carbapenemase which was not

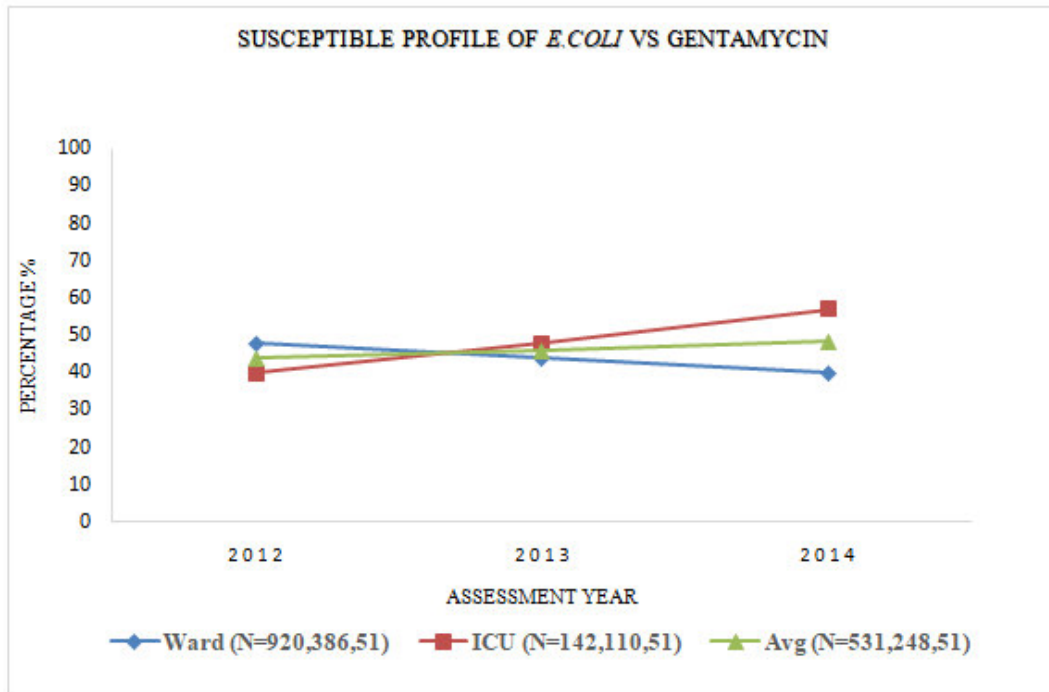
identified in the Ugwu *et.al* study<sup>16</sup>. The susceptibility of different antibiotics was tested in the following years and the susceptibility profile is graphically presented below. The new generation antibiotics showed high resistance in comparison to previously prescribed antibiotics. The prescribing pattern needs to be altered to come across the resistant strains.



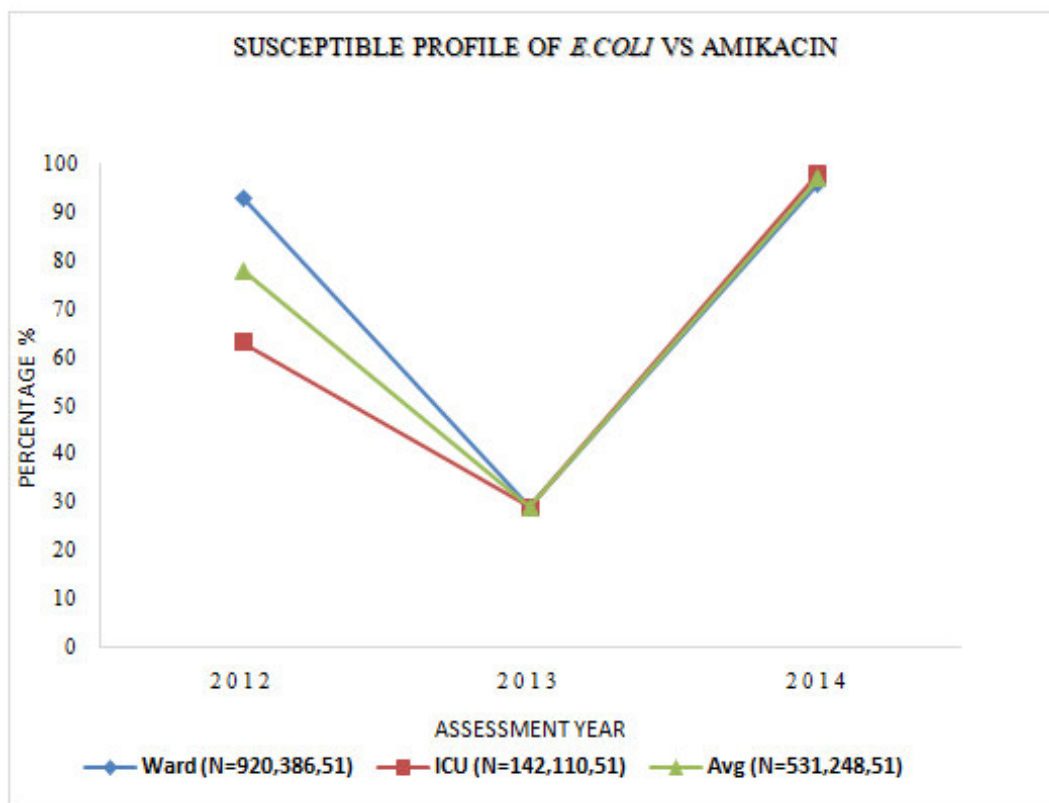
**Figure 1**  
*Susceptible profile of E.coli vs 3<sup>rd</sup> generation Cephalosporins*



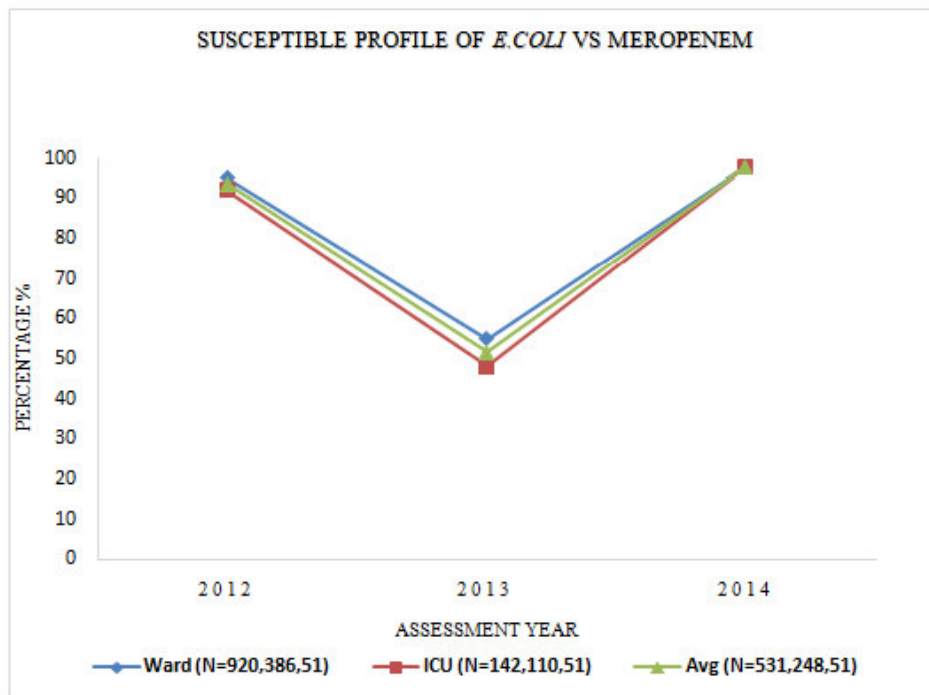
**Figure 2**  
*Susceptible profile of E.coli vs Cefipime*



**Figure 3**  
*Susceptible profile of E.coli vs Gentamycin*



**Figure 4**  
*Susceptible profile of E.coli vs Amikacin*



**Figure 5**  
*Susceptible profile of E.coli vs Meropenem*

## CONCLUSION

The presence of ESBL type of drug resistance is constantly evolving and is under dynamic flux. Hence, it poses a global threat to public health programs. Again, there is a significant necessity for regular antimicrobial sensitivity surveillance not

only for the presence and spread of ESBL genes but also for urban and rural populations for more informed and structured treatment.

## CONFLICT OF INTEREST

Conflict of interest declared none.

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