

Frequency and antibiotic resistance pattern of the clinical isolates of extended spectrum beta Lactamase producing *Escherichia coli* from the blood specimens in hospitals and medical centers in Tabriz, Iran

Sara Naebi ^{*1}, Ahmadreza Shahniani ², Marziyeh Sadat-Amini ²

¹Department of Microbiology, Ahar Branch, Islamic Azad University, Ahar, Iran

²Department of Microbiology, Kazeroon branch, Islamic Azad University, Kazeroon, Iran

*Corresponding author: Sara Naebi, Department of Microbiology, Ahar Branch, Islamic Azad University, Ahar, Iran. Email: naebisara@gmail.com

DOI: 10.22034/HBB.2019.24

Received: September 3, 2019; Accepted: October 13, 2019

ABSTRACT

Production of Extended Spectrum Beta Lactamase (ESBL) enzymes is one of the main reasons for drug resistance in *Escherichia coli* (*E. coli*) isolates. The aim of this study was to determine the ESBL producing *E. coli* isolates in the blood culture of patients in hospitals and medical centers in Tabriz, Iran. In this descriptive cross-sectional study, 200 isolates of *E. coli* from blood specimens in hospitals and medical centers were investigated from December 2018 to May 2019. The production of ESBLs was confirmed using the combined disc method and antimicrobial susceptibility test. Out of 200 isolates, 100 isolates (50 %) were able to produce ESBL. The highest antibiotic resistance of isolates (100 %) was to ceftriaxone, cefotaxime, cefixime and cefuroxime. The highest antimicrobial susceptibility was also observed to amikacin (12 %), meropenem (6 %) and imipenem (2 %), respectively. The results of this study showed the high resistance of *E. coli* isolates especially ESBL producing strains to different antibiotics. Given the relatively high prevalence of the ESBL producing *E. coli* strains and their resistance to common antibiotics, effective control strategies such as the restricted use of broad spectrum cephalosporins should be implemented.

Keywords: *Escherichia coli*, antibiotic resistance pattern, extended spectrum beta Lactamases

INTRODUCTION

Escherichia coli (*E. coli*), as a Gram-negative bacillus from the Enterobacteriaceae family [1] is one of the common pathogens responsible for nosocomial infections [2-3]. Among the different types of nosocomial infections, bloodstream infections are very serious health problems worldwide [4]. One of the most important mechanisms used by gram-negative bacteria against beta-lactam antibiotics is the production of beta-lactamase enzymes [5]. These enzymes hydrolyze a wide range of beta-Lactam antibiotics including penicillins, cephalosporins and monobactams. Production of beta-Lactamase enzymes has emerged as an important mechanism for the resistance against beta-Lactam drugs, which include approximately 50% of used common antibiotics [6-8]. Since the identification of these enzymes in 1989 until now, more than 150 types of ESBLs have been identified, which have been mostly detected in the Enterobacteriaceae family [9]. Extended-spectrum beta-Lactamases (ESBLs) are mainly produced by *Klebsiella* and *Escherichia coli* and other members of Enterobacteriaceae, respectively [10]. The prevalence of ESBL-producing strains in clinical isolates varies in different cities and counties. In the United States, the production

Antibiotic resistance pattern

of these enzymes by the members of Enterobacteriaceae family has increased by 0-25 %. The frequency of ESBL-producing bacteria in other countries varied from 3-8 % in Sweden, 34 % in Portugal, 4.8 % in Korea, 8.5 % in Taiwan, 12 % in Hong Kong to 58 % in Turkey [11-12]. Nowadays, numerous reports indicated that the prevalence of these strains is a worldwide problem. In these studies, the multidrug resistance bacteria have been identified as one of the major medical problems, and the need for investigation and control measures have been emphasized [13]. The administration of broad-spectrum antibiotics is necessary for the treatment of infections caused by these strains. This is particularly complicated in patients with weakened immune systems and those admitted to intensive care units. Therefore, microbiology laboratories play an important role in identifying and reporting the ESBL- producing bacteria and facilitating effective treatment for the patients [14]. Due to the importance of ESBL-producing bacteria, their prevalence needs to be determined in order to take appropriate measures to treat the infections caused by them. Since the prevalence of these strains is increasing in hospitals and medical centers, isolation of ESBL-producing strains of *E. coli* from clinical specimens and determination of their antibiotic resistance

pattern were the main objectives of this study.

MATERIALS AND METHODS

In this descriptive cross-sectional study, 200 isolates of *E. coli* were provided from blood cultures of patients who were suspected to bacteremia in hospitals and health centers of Tabriz during a 6 months period (December 2018 to May 2019). A questionnaire was designed and coded for each patient. The sampling was performed after specifying the purpose of the study for each patient under study and the written consent was obtained from them. The isolates were confirmed using at least 14 biochemical tests including IMViC, amino acid decarboxylation, oxidase, urea and malonate utilization tests. Antibiotic resistance pattern of the isolates was determined using the Disc Diffusion Agar (DAD) method according to the guidelines of Clinical & Laboratory Standards Institute (CLSI). The used antibiotic discs were included: ceftazidime (30 µg), ceftriaxone (30 µg), cefotaxime (30 µg), cefepime (30 µg), aztreonam (15 µg), imipenem (10 µg), meropenem (10 µg), ampicillin (10 µg), gentamicin (10 µg), amikacin (30 µg), cotrimoxazole (25 µg), cefalexin (30 µg), cefixime (5 µg), ciprofloxacin (5 µg) and cefuroxime (30 µg). In the disc diffusion test, the diameter of

Antibiotic resistance pattern

inhibitory zones around each disc was measured and the results were reported according to CLSI standards. Combined disc method was used for phenotypical evaluation of ESBL-producing strains. In this method, the discs of ceftazidime plus cefotaxime (30+10 µg), ceftazidime plus clavulanic acid (30+10 µg) and cefotaxime plus clavulanic acid (30+10 µg) were placed on Mueller-Hinton agar medium (Oxoid, England) with a distance of 25 mm from each other (center to center). The plates were incubated at 35 °C for 18-24 h. If the inhibitory zone around clavulanic acid-containing discs was at least 5 mm higher than clavulanic acid-free discs, it would be considered as positive for ESBLs [15]. Standard strains of *Klebsiella pneumoniae* (ATCC 700603) and *E. coli* (ATCC 25922) were used as positive and negative controls, respectively.

RESULTS

In the present study, *E. coli* were isolated from 200 blood specimens of patients including 126 hospitalized (63 %) and 74 ambulatory patients (37 %). Out of total isolates, 119 (59.5 %) and 81 (40.5 %) isolates were belonged to female and male patients, respectively. In this study, antibiotic resistance pattern of the isolates was identified using the DAD method and 16 types of antibiotics. The highest resistance of

isolates (100 %) was observed to ceftriaxone, ceftriaxone, cefotaxime, cefixime, cefuroxime. Also, the lowest antibiotic resistance was detected to amikacin (12 %),

meropenem (6 %) and imipenem (2 %), respectively (Figure 1).

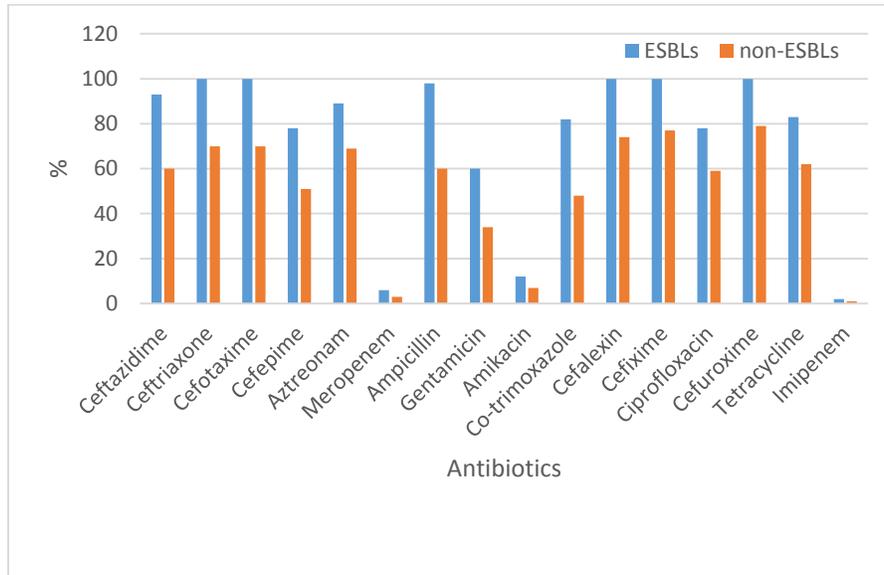


Figure 1. The Antibiotic resistance frequency of ESBL and non-ESBL-producing isolates of *Escherichia coli* against common antibiotics using disc diffusion agar method.

Of the total 200 isolates, 100 isolates (50 %) were identified as ESBL-producing strains which had a higher percentage of resistance to different antibiotics than others. Seventy percent (70 %) of ESBL-producing strains were isolated from females while 30 % of them were detected in males. The mean age of patients with ESBL-producing *E. coli* was

50 ± 24.4 years. The resistance of isolates was significantly more than 50 % for about 13 types of antibiotics (p<0.05).

DISCUSSION

Bacteremia is one of the leading causes of death among patients. Drug-resistant Gram-

Naebi et al.

negative bacilli, particularly the members of the Enterobacteriaceae, are the most common organisms responsible for bacteremia in hospitals, and the infections caused by them are associated with high mortality [16-17]. Production of beta-Lactamases is the main resistance mechanism of gram-negative bacteria against beta-Lactams. As beta-Lactam antibiotics are used clinically, beta-Lactamases evolved along with these drugs and played a major role in the therapeutic failures in antibiotic therapy [18]. Since the fifteen years ago, there have been numerous outbreaks of infections caused by beta-Lactamase-producing organisms throughout the world. This phenomenon is a major threat to the use of cephalosporins. It also has been well established that treatment for such cephalosporin-resistant infections will not be led to satisfactory results. The mortality rate caused by ESBL-producing strains is significantly higher than the others. Another issue is whether cephalosporins treatment is appropriate for ESBL-positive organisms whose MIC is in the sensitive range [19]. The emergence and distribution of ESBL-producing bacteria occur due to the widespread use of broad-spectrum beta-lactams, and the prevalence of these strains are increasing in the hospitals in recent years. In this study, the prevalence rate of ESBL-producing *E. coli* strains in clinical samples

Antibiotic resistance pattern

was 50 %. In similar studies, the prevalence rate of these strains was reported as 17 % in Turkey [20], 3.3–7.7 % in Colombia [21], 29.1 % in India [22], 4.7 % in the USA [23] and 16.2 % in France [24]. On the other hand, in the study of Zhou [25] in Shanghai, it was shown that 47.4 % of *E. coli* isolates from the patients were positive for ESBL. In another study conducted by Wu. [26] at Taiwan hospitals, ESBL-producing *E. coli* was 18.18 %. However, this prevalence was reported as 28.1 % in Beirut, Lebanon [27]. In the study of Fazeli [28] on the clinical specimens of a hospital in Isfahan, 150 (53.9 %) of 278 *E. coli* isolates were positive for the production of ESBL. Mehrgan [29] also reported a prevalence rate of 16 % for beta-Lactamase-producing *E. coli* in a tertiary care hospital in Tehran. Comparison of the above results with the findings of present study shows that the prevalence of ESBL-producing isolates varies between different countries as well as from one hospital to the another one in a country, depending on the infection control system and the treatment level of patients [30]. In the study of Mardaneh [31], 58 % of *E. coli* isolates from blood cultures, were positive for beta-lactamase production, and the most effective antibiotics against these strains were colistin, polymyxin B, imipenem, meropenem, amikacin and piperacillin-tazobactam. Jafari Sales and

Naebi et al.

Mobin [32] reported that 100 (51.02 %) of 196 *E. coli* isolates from medical centers in Marand city (Iran) were positive for ESBL production. Eighty-two percent (82 %) of the isolates were ampicillin-resistant while 7 % of them were sensitive to imipenem. In the study of Haghighi [33], 33.5 % of isolates were able to produce ESBLs. The highest resistance of isolates (71.35 %) was against amoxicillin and all strains (100 %) were sensitive to imipenem. In the study of Mohajeri [34] studied the frequency of ESBL-producing *E. coli* isolates from urinary tract infections in Kermanshah, Iran. They found that the highest susceptibility and resistance of isolates was to imipenem (100 %) ampicillin (77 %), respectively. Rahbar [35] also reported the highest rate of resistance in *E. coli* isolates to ampicillin (18 strains, 100 %). In the study of Mshana in Tanzania [36], the resistance rate of isolates to ampicillin was 92.7 %, whereas in the present study it was 98 %. In this study, 98 % of isolates were susceptible to imipenem, whereas in other studies this rate was varied from 91.7 % to 100 % [37-38]. In the present study, 100 % of ESBL-positive strains were resistant to cefalexin, ceftriaxone, cefotaxime, cefixime and cefuroxime. This result indicates the increased resistance to these drugs in the country. According to the above results, adequate precautions should be

Antibiotic resistance pattern

taken to prevent the increasing resistance to these antibiotics. The phenomenon of antibiotic resistance annually imposes huge costs on the healthcare systems of the countries. In addition to the misuse of antibiotics, aggressive therapeutic measures, increasing the number of immunocompromised patients and the lack of practical advices in infection control also play an important role in the increasing of the antibiotic resistance.

CONCLUSION

According to the results of this study, the high prevalence of resistance to ESBL-producing *E. coli* strains is an important health hazard for hospitalized and ambulatory patients. Also, producing ESBLs by these strains is a major threat to the use of extended spectrum cephalosporins. Therefore, in order to treat infections potentially caused by ESBL-producing strains, the appropriate antibiotic should be selected based on the results of antibiogram test. To reduce the prevalence of beta-lactamase-producing organisms, further studies in this area, the strict controlling measures as well as the reduced use of antibiotic are necessary.

REFERENCES

- [1]. Jafari-Sales A; Rasi-Bonab F. Detection of the antibiotic resistance pattern in *Escherichia coli* isolated from urinary tract infections in Tabriz City. *J Mol Microbiol*, 2017, 1(1): 101.
- [2]. Hidron AI, Edwards JR, Patel J, Horan TC, Sievert DM, Pollock DA. Antimicrobial-resistant pathogens associated with healthcare-associated infections: annual summary of data reported to the National Healthcare Safety Network at the Centers for Disease Control and Prevention. *ICHE*, 2008; 29(11): 996-1011.
- [3]. Jafari-Sales A, Shadi-Dizaji A. Molecular analysis of CTX-M genes among ESBL producing in *Pseudomonas aeruginosa* isolated from clinical samples by Mul-tiplex-PCR. *HOZAN J Environment Sci*; 2018; 2(5): 17-29.
- [4]. Taneja J, Mishra B, Thakur A. Nosocomial blood-stream infections from extended-spectrum-beta-lactamase-producing *Escherichia coli* and *Klebsiella pneumoniae* from GB Pant Hospital, New Delhi. *JIDC*, 2010, 4.08: 517-20.
- [5]. Mobasher Kare Jeddi A, Nahaei M, Mobayyen H, Pornour M, Sadeghi J. Molecular study of extended-spectrum beta-lactamase (SHVtype) in *Escherichia coli* and *Klebsiella pneumoniae* isolated from Medical Centers of Tabriz. *Iran J Med Microbiol*, 2009; 2 (4): 9-17.
- [6]. Rawat D, Nair D. Extended-spectrum β Lactamases in Gram negative bacteria. *J Glob Infect Dis*, 2010; 2: 263-74.
- [7]. Jafari-Sales A, jafari B, Bagherizadeh Y, Khalifehpour M, Abdoli-senejan M, Helali-Pargali R. Antibiotic resistance pattern and bla-TEM gene expression in *Acinetobacter baumannii* isolated from clinical specimens of Tabriz hospitals. *Zanko J Med Sci*, 2019; 20(65): 20-29.
- [8]. Jafari-Sales A. Study of antibiotic resistance and prevalence of bla-TEM gene in *Klebsiella pneumoniae* Strains isolated from children with UTI in Tabriz hospitals. *Focus on Sciences*, 2018, 4: 1.
- [9]. Rupp ME, Fey PD. Extended spectrum beta-lactamase (ESBL)-producing *Enterobacteriaceae*: considerations for diagnosis, prevention and drug treatment. *Drugs*, 2003; 63(4): 353-65.
- [10]. Thomas KS, Moland ES, Coudron PE. Occurrence and detection of extended spectrum beta- Lactamases in members of the family *Enterobacteriaceae* at a Veterans Medical Center: seek and you may find. *J Clin Microbiol*, 1997; 35: 2593-97.

[11]. Ferreira CM, Ferreira WA, Almeida NC. Extended-spectrum beta-lactamase-producing bacteria isolated from hematologic patients in Manaus, State of A mazonas, *Braz J Microbiol*, 2011; 42: 1076-84.

[12]. Dizaji AS, Fathi R, Sales AJ. Molecular study of extended-spectrum beta-lactamase (TEM-1) gene in *Escherichia Coli* isolates collected from Ostad Alinasab Hospital in Tabriz Iran. *Marmara Med J*, 2016; 29(1): 35.

[13]. Palucha A, Mikiewicz B, Hryniewicz W. Concurrent outbreaks of ESBL producing organism of the family *Enterobacteriaceae* in a Warsaw Hospital. *J Antimicrobiol Chemother*, 1999; 44(4): 489-99.

[14]. Magdalena T, Fritz H, Herbert H. Survey and molecular genetics of SHV beta Lactamases in Enterobacteriaceae in Switzerland: two novel enzymes, SHV-11 and SHV-12. *Antimicrob Agents Chemother*, 1997; 41(5): 943-49.

[15]. Livermore DM, Woodford N. Guidance to diagnostic laboratories: laboratory detection and reporting of bacteria with extended spectrum β -Lactamases. Issued by standards unit, evaluations and standards laboratory. centre for infections. HPA. June 2004.

[16]. Sales AJ, Fathi R, Mobaiyen H. Molecular Study of the Prevalence of CTX-M1, CTX-M2, CTXM3 in *Pseudomonas aeruginosa* Isolated from Clinical Samples in Tabriz Town, Iran. *Electronic J Biol*, 2014; 13(3).

[17]. Jafari Sales A, Mobaiyen H, Farshbafi Nezhad Zoghi J, Nezamdoost Shadbad N, Purabdollah Kaleybar V. Antimicrobial resistance pattern of extended-spectrum β -Lactamases (ESBLs) producing *Escherichia coli* isolated from clinical samples in Tabriz city, Iran. *Adv Environ Biol*, 2014; 8(16): 179-82.

[18]. Sanders CC. Chromosomal cephalosporinases responsible for multiple resistances to newer β -lactam antibiotics. *Clin. Microbiol Rev*, 1987; 41: 573- 93.

[19]. Paterson DL, Ko WC, Gottberg VA, Casellas JM, Mulazimoglu L, Klugman KP. Outcome of cephalosporin treatment for serious infections due to apparently susceptible organisms producing extended spectrum Beta-Lactamases: Implications for the clinical microbiology laboratory. *J Clin Microbiol*, 2001; 39(6): 2206-12.

[20]. Tasli H, Bahar IH. Molecular characterization of TEM and SHV-derived extended spectrum beta-Lactamases in hospital-based

Enterobacteriaceae in Turkey. *Jpn J Infect Dis*, 2005; 58(3): 162-67.

[21]. Villegas MV, Correa A, Perez F, Miranda MC, Zuluaga T, Quinn JP. Prevalence and characterization of extended-spectrum beta-Lactamases in *Klebsiella pneumoniae* and *Escherichia coli* isolates from Colombian hospitals. *Diagn Microbiol Infect Dis*, 2004; 49(3): 217-22.

[22]. Duttaroy B, Mehta S. Extended spectrum beta Lactamases (ESBL) in clinical isolates of *Klebsiella pneumoniae* and *Escherichia coli*. *Indian J Pathol Microbiol*, 2005; 48: 45-48.

[23]. Saurina G, Quale JM, Manikal VM, Oydna E, Landman D. Antimicrobial resistance in *Enterobacteriaceae* in Brooklyn, NY: epidemiology and relation to antibiotic usage patterns. *J Antimicrob Chemother* 2000; 45: 895-98.

[24]. Lavigne JP, Bouziges N, Chanal C, Mahamat A, Michaux-Charachon S, Sotto A. Molecular epidemiology of *Enterobacteriaceae* isolates producing extended-spectrum beta Lactamases in a French hospital. *J Clin Microbiol*, 2004; 42: 3805-8.

[25]. Zhou L. Pathogens and associated factors of infections in PICU. Shanghai second medical university afflicted shanghai children medical center: 2001.

[26]. Wu TL, Chia JH, Su LH, Kuo AJ, Chu C, Chiu CH. Dissemination of extended-spectrum beta-lactamase-producing *Enterobacteriaceae* in pediatric intensive care units. *J Clin Microbiol*, 2003; 41(10): 4836-38.

[27]. Daoud Z, Hakime N. Prevalence and susceptibility patterns of extended-spectrum betalactamase-producing *Escherichia coli* and *Klebsiella pneumoniae* in a general university hospital in Beirut, Lebanon. *Rev Esp Quimioter*, 2003; 16(2): 233-38.

[28]. Fazeli H, Hoseini M, Mohammadi Ghalaei P. Frequency and resistance pattern of extended spectrum beta lactamase producing *Escherichia coli* in clinical specimen of Alzahra hospital in Isfahan, Iran, 2007. *J Shahrekord Univ Med Sci*, 2009; 10 (4) :58-64.

[29]. Mehrgan H, Rahbar M. Prevalence of extended spectrum beta lactamase producing *Escherichia coli* in a tertiary care hospital in Tehran. *Iran J Antimicrob Agents*, 2008; 31: 147-51.

[30]. Shah AA, Hasan F, Ahmed S, Hameed A. Characteristics, epidemiology and clinical importance of emerging strains of Gram-negative bacilli producing extended-spectrum beta Lactamases. *Res Microbiol*, 2004; 155(6): 409-21.

[31]. Mardaneh J, Anvarinejad M, Abbasian A, Abbasi P, Razaatpour N, Dehyadegari M. Emergence of multi-drug resistant ESBL producing strains among *Enterobacteriaceae* members isolated from patients blood samples in south of Iran. *Iran South Med J*, 2015; 18(5): 970-81.

[32]. Jafari Sales A, Mobaiyen H. Frequency and resistance patterns in clinical isolates of *Escherichia coli* extended spectrum beta lactamase producing treatment centers in Marand city, Iran. *NCMBJ*, 2017; 7(26): 19-26.

[33]. Haghi F, Zeighami H, Keramati N, Hemmati F, Hajiahmadi F. Frequency of TEM extended spectrum beta lactamase producing *Escherichia coli* in clinical specimens by phenotypic and molecular methods in Zanjan. *ZUMS Journal*, 2013; 21(85): 55-63.

[34]. Mohajeri P, Izadi B, Rezai M, Falahi B, Khademi H, Ebrahimi R. Assessment of the frequency of extended spectrum beta Lactamases producing *Escherichia coli* isolated from urinary tract infections and its antibiotic resistance pattern in Kermanshah. *J Ardabil Univ Med Sci*, 2011; 11(1): 86-94.

[35]. Behrouzi A, Rahbar M, Vand Yousefi J. The prevalence of extended-spectrum β -lactamase (ESBLs) producing *Klebsiella pneumonia* and *Escherichia coli* isolated in Milad hospital of Tehran in 2010. *Iran J Med Microbiol*, 2010; 4 (2): 36-41.

[36]. Mshana SE, Kamugisha E, Mirambo M, Chakraborty T, Lyamuya EF. Prevalence of multiresistant gram negative organisms in a tertiary hospital in Mwanza, Tanzania. *BMC Res Notes*, 2009; 2(49): 325-28.

[37]. Shiju MP, Yashavanth R, Narendra N. Detection of extended spectrum β -lactamase production and multidrug resistance in clinical isolates of *E.coli* and *K.pneumoniae* in mangalore. *JCDR*, 2010; 4(3): 2442-45.

[38]. Yazdi M, Nazemi A, Mirsaed MN, Khataminezhad MR, SHarifi SHA, Babaei Kochaksaraei M. Prevalence SHV/CTX-M/TEM genes in *Escherichia coli* producing extended spectrum β -lactamase isolated from urinary infection in Tehran city. *L Sci J*, 2010; 4(1):120-24.