

Risk Factors Associated with ESBL and CPE Acquisition among Pediatrics: A Systematic Review

ARTICLE INFO

Article Type

Systematic Review

Authors

Ghasemian A.M.1 PhD, Norouzi F.1 PhD. Shokouhi Mostafavi S.Kh.2 PhD, Hasanvand F.3 MSc, Nojoomi F.* PhD

How to cite this article

F, Nojoomi F. Risk Factors Associated with ESBL and CPE Acquisition among Pediatrics: A Systematic Review, Infection Epidemiology and Microbiology. 2018 ;4(1):35-40.

Ghasemian A.M, Norouzi F, Shokouhi Mostafavi S Kh, Hasanvand

*Microbiology Department, Medicine Faculty, AJA University of Medical Sciences, Tehran, Iran

¹Microbiology Department, Fasa University of Medical Sciences, Fasa,

²Microbiology Department, Medicine Faculty, AJA University of Medical Sciences, Tehran, Iran

³Bacteriology Department, Medical Sciences Faculty, Tarbiat Modares University, Tehran, Iran

Correspondence

Address: AJA University of Medical Sciences, Gisha Bridge, North Kargar, Tehran, Iran

Phone: +98 (21) 88432157

bacteriology94@gmail.com

Article History

Received: December 5, 2017 Accepted: February 3, 2018 ePublished: March 20, 2018

ABSTRACT

Aims Infections by extended-spectrum beta-lactamase (ESBL) and carbapenemase-producing Enterobacteriaceae (CPE) are increasing problems in pediatrics and are usually associated with higher hospital costs and mortality rates. The aims of this study were the statistical investigation of the worldwide prevalence and risk factors of ESBL and CPE family members among pediatric population.

Instruments & Methods From October 1, 1995 to July 27, 2017, some keywords including "ESBLs", "carbapenemase", "pediatrics", "children", and "risk factor" were searched in the searching databases such as Google Scholar, Embase, Scopus, PubMed, and Web of Science among original research articles. The univariate and multivariate analysis of the collected data was performed by Graph Pad Prism 6.1 software.

Findings The mean percentage of ESBL production was 20.23±22.31 and the mean percentage of CPE was 1.81±2.77. E. coli (n=991) and K. pneumonia (n=627) were the predominant ESBLproducers. Nephrology (n=5005) and NICU (n=1805) were predominant hospital wards. ESBL-PE had significantly higher prevalence in the infants unit (OR=0.9832, 95% CI=12.271-19.519; p<0.001). Moreover, ICU ward was a significant and independent risk factor for CPE acquisition (OR=0.849, 95% CI=2.211-5.415; p=0.0035). ESBL-PE and CPE were significantly isolated from blood samples (OR=0.9276, 95% CI=1.508-2.433, p<0.0001) and fecal specimens (OR=0.968, 95% CI=2.829-5.133, p<0.0001), respectively.

Conclusion Most of risk factors between ESBL-PE and CPE are similar including previous hospitalization and prolonged use of antibiotics, cephalosporins, and previous colonization. Other possible potential risk factors that should be considered include presence of catheters and travel history. Detection of risk factors provides useful information for formulation of infection control policy.

Keywords Enterobacteriaceae; Extended-spectrum beta-lactamase; Carbapenemases; Risk Factors; **Pediatrics**

CITATION LINKS

[1] Nosocomial outbreak of Klebsiella infection ... [2] The epidemiology of candida ... [3] β-Lactams: Chemical structure, mode of ... [4] An overview of carbapenem ... [5] Prevalence of clinically isolated ... [6] Federal funding for the study of ... [7] NHSN annual update: Antimicrobialresistant ... [8] Ecology drives a global network ... [9] Effect of overgrowth or decrease ... [10] Antibiotic resistance patterns and ... [11] Carbapenemases: The versatile ... [12] Carbapenemase genes and ... [13] Biochemical sequence analyses ... [14] Co-resistance to different ... [15] Outbreak of multiply ... [16] Detection of gastrointestinal pathogens ... [17] Influx of extended-spectrum ... [18] Faecal carriage of extended-spectrum ... [19] Prevalence and spread of ... [20] Epidemiology and risk ... [21] Risk factors for acquisition ... [22] Fecal colonization with ... [23] Colonisation with ... [24] Molecular epidemiology ... [25] Diagnosis and management ... [26] The mother as most ... [27] Prevalence of ESBL-producing ... [28] Risk factors for ... [29] Extended-spectrum ... [30] Risk factors for the ... [31] Risk factors in community ... [32] Risk factors for community ... [33] Fecal carriage of ... [34] The prevalence of plasmid ... [35] Antimicrobials increase ... [36] International travel is a risk factor ... [37] Risk factors associated with ... [38] Multidrug-resistant ESBL ... [39] Extended-spectrum beta-lactamase-producing ... [40] Bloodstream infections by ... [41] Risk factors for extended ... [42] Molecular epidemiology of an SHV-5 extended-spectrum ... [43] Infection due to extended-spectrum ... [44] Multi-drug-resistant Enterobacteriaceae in a Portuguese ... [45] ESBL-producing Escherichia ... [46] Fecal carriage of ESBL-producing ... [47] High rate of fecal carriage of ... [48] Risk factors for carbapenem-resistant ... [49] Molecular characterization and risk factors for carbapenem-resistant ... [50] High gastrointestinal colonization rate with ...

Introduction

Infections due to the multidrug-resistant Gramnegative bacteria are rising in the pediatric population and they pose serious problems in the limited therapeutic options and long-time hospitalization.

Showing resistance to penicillin and cephalosporins as well as to non-beta-lactam antibiotics, Extended-spectrum beta-lactamaseproducing Enterobacteriaceaee (ESBL-PE) have been described since the 1990s as nosocomial pathogens^[1-3]. Furthermore, the recent emergence and spread of carbapenemase-bearing strains (CPE) among children have limited the therapeutic choices as the carbapenems are the last resorts of eradication of infections due to ESBL[4, 5]. These types of infections are associated with the high rate of mortality as the pediatrics is a vulnerable population.

Two members of the Enterobacteriaceaee family, *Klebsiella pneumonia* and Enterobacter spp., are placed in the ESKAPE pathogens, standing for *Enterococcus faecium, Staphylococcus aureus, Klebsiella pneumoniae, Acinetobacter baumannii, Pseudomonas aeruginosa,* and Enterobacter species group^[6] and this family members are among the most common causes of device-related infections^[7].

Because of plasmid-borne or extrachromosomal expression of ESBLs and carbapenemases, the transmission of encoding genes between human micobiota as well as from livestock bacteria occurs rapidly and easily^[8, 9]. It has been reported that gastrointestinal infections are the primary causes of death in children; thus, the detection of drugresistant Enterobacteriaceae and uncovering most significant risk factors is critical in order to foster the proper treatment regimens^[10].

The ESBLs are encoded by CTX-M, TEM and SHV genes, which have been reported worldwide in both healthcare and community settings and are associated with the high rate of mortality among pediatrics. Furthermore, carbapenemases are encoded by various classes including OXA-48, KPC and GES, and metallo-β-lactamases such as IMP, VIM, and NDM1 genes[11, 12]. The spread of these genes will make the treatment options very limited, especially for vulnerable population of pediatrics, because both **ESBL** carbapenemase-producing strains carry resistance elements to other classes of antibiotics[13, 14].

Risk factors of fecal colonization and infection with ESBL-PE in adults include hospital stay more than 5 days, chronic care facilities, catheterization, antibiotic exposure, comorbid conditions, sputum suction, meat use associated with the agricultural antibiotic apply, underlying diseases, and international travel, especially for those persons with systemic febrile illnesses,

immunosuppression, diarrheal illnesses, dermatologic disorders[15-25]. Risk factors of ESBL infections in pediatrics have been determined to some extent; they include healthcare exposure, exposure of antibiotics, previous use or particularly cephalosporins, maternal transmission, and comorbid conditions such as urologic or neurologic conditions^[26-29]. In addition, there have been several risk factors associated with the acquisition of ESBL and carbapenemaseproducing Enterobacteriaceae in the community including age over 60 years, antibiotic treatment in the past 3 months, previous hospitalization in the past 3 months, within-household transmission, diabetes and some other underlying disorders, male gender, previous infection, nursing home stay and previous use of cephalosporins, penicillin, and quinolones[30-33]; thus, it is an urgent issue to increase knowledge, validate, and hinder the transmission of these multidrug-resistant species among children.

The aims of the current study were the statistical investigation of the worldwide prevalence and risk factors of ESBL and carbapemenase-producing Enterobacteriaceaee family members among pediatric population.

Information and Methods

We conducted a systematic review over the prevalence and risk factors associated with the acquisition and spread of ESBLs and carbapemase-producing species of Enterobacteriaceae and statistically analyzed them.

For this purpose, from October 1, 1995 to July 27, 2017, some keywords including "ESBLs", "carbapenemase", "pediatrics", "children", and "risk factor" were searched in the searching databases such as Google Scholar, Embase, Scopus, PubMed, and Web of Science among original research articles. The national publications were also considered to be used.

Inclusion and exclusion criteria: All publications regarding the prevalence of ESBLs and carbapenemases and risk factors among children (both patients and healthy population) conducted in all countries were considered. Pediatric patients from both healthcare and community were included. Pediatrics with ages between <1 year to 16 years were included. Furthermore, both colonization and infection by ESBL-producing and carbapenemase-producing Enterobacteriaceae members were included. All the infection sources caused by the Enterobacteriaceae family in children (UTI, fecal infection, respiratory tract sampling, etc.) and hospital wards (Pediatrics, NICU, ICU, Emergency, etc.) were included. Patients with asymptomatic status, who were colonized or infected with **ESBL** carbapenemases-non producers, were excluded.

Archive 35f SID Ghasemian A.M.et al.

Furthermore, case reports, review articles, systematic review papers, and those available only in the abstract form were excluded from this study. **Statistical analysis:** The univariate and multivariate analysis of the collected data was performed by the Graph Pad Prism 6.1 software. The Mean±SD of factors, comparison of results, and possible relations were performed by the software. The unpaired t-test and one-way ANOVA tests were applied for the purpose of risk factor analysis and 95% confidence interval (95% CI) and p value<0.05 were considered the breakpoint for a significant result.

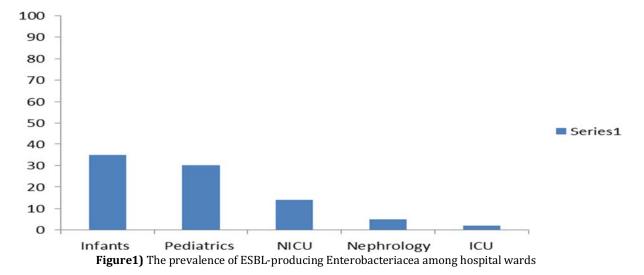
Findings

Risk factors for resistance development: A total of 8605 (5782 in the hospital and 2823 in the community) isolates were enrolled for ESBLPE and 1488 (all belonging to the hospital-associated

infections) isolates were investigated for CPE prevalence among 51 publications in children population, which met the criteria. The mean percentage of ESBL production was 20.23±22.31 and the mean percentage of CPE was 1.81±2.77. Moreover, among ESBL producers, E. coli (N=991) and K. pneumonia (N=627) were the predominant ESBL-producers, followed by Enterobacter cloacae (N=53), K. oxytoca (N=19), Proteus mirabilis (N=19), Citrobacter freundii (N=9), C. koseri (N=7), Seratia spp (N=7), morganella morganii (N=6), and Salmonella spp (N=3) (Table1). The children age ranges were as follow: 0 to 1 year=2894, 1 to 5 years=3177, 5 to 10 years=2510, and >10 years=924. The majority of isolates (N=5005) were collected from children in Nephrology ward, followed by 1805 isolates in NICU, 817 isolates in ICU, 687 in Pediatric ward, and 301 isolates in infants units. The prevalence of ESBLs among hospital wards was shown (Figure 1, Table 1).

Table1) The prevalence of ESBL and carbapenemase producing Enterobacteriacea in hospital wards and infection and colonization sites, ESBL PE: ESBL-producing Enterobacteriacea, *CPE: carbapenem-producing E (percentage among the total of 1488 isolates), ND: not detected

among the total of 1 foo isolates), Not not detected								
Variables	ESBL-PE No (%)	OR	95% CI	p value	*CPE No (%)	OR	95% CI	p value
Hospital ward								
Nephrology (N=5005)	158 (3.81)		1.124-3.141		4 (2.69)		1.111-3.141	
NICU (N=1805)	155 (8.56)		1.421-5.574		11 (7.39)		2.132-6.162	
ICU (N=817)	11 (1.34)	0.9832	0.982-2.111	< 0.0001	12 (8.06)	0.849	2.211-5.415	0.0035
Pediatrics (N=687)	134 (19.50)		11.187-15.596		ND		-	
Infants (N=301)	81 (26.9)		12.271-19.519		ND		-	
Sampling site								
Fecal (N=5939)	400 (6.73)		2.141-3.207		21 (3.53)		2.829-5.133	
Urinary tract (N=3789)	123 (3.24)		1.569-3.161		5 (1.58)		4.837	
Blood (N=599)	14 (2.33)	0.9276	1.508-2.433	< 0.0001	1 (1.6)	0.968	4.838	< 0.0001
Respiratory (N=63)	2 (3.17)		2.642-3.111		ND		-	
Other (N=15)	ND		ND		ND		-	



ESBL-PE had significantly higher prevalence in the infants unit (OR=0.9832, 95% CI=12.271-19.519; p<0.001), which can be considered as an independent risk factor for spread of ESBL-PE

(Figure 1, Table 1). Moreover, ICU ward was a significant and independent risk factor for CPE acquisition (OR=0.849, 95% CI=2.211-5.415; p=0.0035).

Furthermore, fecal samples (N=5939) were mostly investigated, followed by urine specimens (N=3789), blood samples (N=599), respiratory samples (N=63) and other (N=15). The prevalence of ESBL and carbapenemase-producing species among both infection and colonization sites were depicted (Table1).

ESBL-PE and CPE were significantly isolated from

blood samples (OR=0.9276, 95% CI=1.508-2.433, p<0.0001) and fecal specimens (OR=0.968, 95% CI=2.829-5.133, p<0.0001), respectively.

Among various risk factors, previous hospitalization and prolonged antibiotic use, especially cephalosporins, are independent risk factors for the acquisition of HA-ESBL-PE among pediatrics (Table 2).

Table2) The significant risk factors found among hospital acquired (HA) ESBL-PE, Community-acquired (CA) ESBL-PE, and CPE

Risk factor	p value: HA-ESBL-PE	p value: CA-ESBL-PE	p value: CPE (HA)
Previous hospitalization	0.05	-	0.001
Prolonged antibiotic use 1886	0.002	0.001	0.001
Nursery	0.007	-	-
Day care home 1886	0.013	0.003	-
Travel history 1886	0.003	0.001	0.01
prolonged out residence	-	0.041	-
Gestational age/birth weight	0.02	0.02	0.001
Previous cephalosporin	<0.001	<0.001	< 0.001
Previous stay	0.001	0.001	0.001
Previous UTI	0.001	0.001	0.001
Previous nitrofurantoin	<0.001	-	-
Previous quinolones	<0.001	-	0.001
Previous use of carbapenems	0.007	0.01	0.001
Neurologic comorbidity	0.01	-	-
central venous catheters	-	0.035	-
ESBL-E-positive mother	0.002	-	-
Underlying disease	0.006	0.006	0.004
Gastrointestinal comorbidity	0.006	-	-
tracheostomy or gastrostomy	0.018	-	-
Birth in hospital	-	-	0.027
Cesarean birth	0.036	-	-
Male gender	0.035	-	-
cephalosporin plus	0.02	_	_
aminoglycoside			
central vascular catheter	0.003	-	-
Previous colonization	0.002	0.003	0.001
Lower age	0.02	-	-
History of infection (last 3 month)	0.001	0.001	0.001
Suppression	0.001	-	-
Septicemia System in faction	0.001	-	-
Systemic infection	0.001 0.001	0.002	-
Renal abnormality Recurrent UTI	0.001	0.002	-
TMP/SMX-nitrofurantoin	0.004	0.004	-
I MIP / SMA-III U OI UF all tolli	0.003	-	-

Trends in ESBL and carbapenemase enzymes development: Pediatrics population is important regarding the acquisition of ESBL-PE and CPE for several reasons including vulnerability of them, unawareness of children about the issue, and, thus, rare observing the hygiene and having more contact among them and other populations.

For neonates, mother colonization with ESBL producing Enterobacteriaceae (p<0.001) and length of stay in hospital (0.037) were significant risk factors [26].

The presence of wound or drain and the use of cephalosporins, carbapenems, and PPIs (p=0.005) in the preceding 6 months were independent risk factors in the multivariable analysis.

Discussion

The aims of the current study were the statistical investigation of the worldwide prevalence and risk factors of ESBL and carbapemenase producing Enterobacteriaceaee family members among pediatric population. In Chinese pediatric hospitals, among pediatric patients, the DHA-1 type AmpC enzymes had the highest prevalent rate without any risk factor stated, which were mainly isolated from neonatology ward and respiratory medicine ward (41.2% and 20.5% for *K. pneumonia* and 24.4% and 27.9% for *E. coli*, respectively) [34]. Increasing age has also been stated as a risk factor for the development of carbapenem-resistant Enterobacteriaceae and

likewise antimicrobials can enhance these strains acquisition^[35].

Among 1- to 17-year-old pediatrics in a study, a recent hospital exposure was compared to the control group (90% vs. 63%; p=.008) and having a comorbid gastrointestinal (60% vs. 18.3%, p<.001) or neurologic (53% vs. 20% p=.01) impair were potential risk factors. However, a comorbid gastrointestinal illness was the only significant risk factor associated with non-ESBL infection (43.3% vs. 18.3%, p=.012); most of the children were diagnosed in NICU (37%) and PICU wards (23%) and 30% of pediatrics were outpatients^[29].

A study among children between 0 to 18 years old, on multivariate analysis showed that the significant risk factors for ESBL acquiring were Asian race (OR=2.56, 95% CI=1.34-4.89; p=0.005), international travel (OR=8.93, 95% CI=2.92-27.78; p<0.001), prior UTI (OR=8.06, 95% CI=3.47-18.87; p<0.001), and comorbid GI condition (OR=2.65, 95% CI=1.36-5.15; p=0.002) $^{[36]}$.

In a study on pediatrics, tracheostomy or gastrostomy (OR=3.62, 95% CI=1.24-10.53; p=0.018) and antibiotic therapy in previous 3 months (OR=4.07, 95% CI=1.29-12.81; p=0.016) were the significant risk factors of ESBL spread[37]. In a study in Lebanon, of 117 rectal swabs collected from 1- to 12-year-old pediatrics in the community, 58 (49.6%) of them were carriers of ESBL-PE. The male gender, caesarean delivery, hospital birth, and being formula-fed were the significant determined risk factors for colonization of ESBL-producing Enterobacteriaceae. Moreover, CTX-M9 and CTX-M15 were the most ESBL-encoding genes[38].

In a study conducted in NICU among infected Brazilian children, of 383 neonates screened for the presence of ESBL-KP, the prevalence of infection and colonization of ESBL-KP was 13 (3.4%) and 206 (53.8%), respectively. The associated risk factors of colonization and infection were combination therapy with cephalosporin and aminoglycoside (HR=4.60; 95% CI=1.48±14.31), previous colonization (HR=5.19; 95% CI=1.58-17.08), and presence of central vascular catheter (HR=13.89; 95% CI=2.71±71.3), respectively^[39].

In a Korean study, of 157 blood isolates at the Seoul Children's Hospital, 17.9% and 52.9% of the *E. coli* and 52.9% of the *K. pneumoniae* isolates were ESBL producers and the fatality rate since ESBL producers was significantly higher (26.7% vs 5.7%) than those non-ESBL producers^[40].

Another study conducted among 225 children in Hong Kong demonstrated a high prevalence of ESBL-producing *E. coli* and *K. pneumonia* fecal carriage among both 0– to 5-year-old children at hospital admission and household members and showed both intra- and inter-household

transmission. CTX-M14 and 15 were detected higher than other types[33]. Among pediatrics with urinary tract infection (UTI), the presence of an underlying disease, infections, hospitalization, and use of antibiotics within the prior 3 months were potential risk factors of acquisition of ESBLbearing K. pneumonia and E. coli (p<0.001). In addition, E. coli was mostly isolated from UTI, but K. pneumonia was isolated more frequently from ESBL-positive children than ESBL-negative UTI[31]. The associated risk factors among a 4 cases outbreak in an NICU caused by K. pneumonia and K. oxytoca isolates, which produced ESBLs, were low gestational age and exposure to thirdgeneration cephalosporins[41]. The low birth weight and low gestational age and combination therapy using cephalosporins and aminoglycosides have been demonstrated in several other studies as significant risk factors for ESBL colonization and infection in children[39, 42, 43].

In a recent study among pediatrics admitted to an NICU ward, gestational age, NICU stay time, and birth weight were associated with infection or colonization with ESBLPE^[44]. In France, of 1886 rectal samples from children, 7.60% of *E. coli* isolates were ESBL producers and risk factors for the acquisition of the strains included being cared for at home, recent antibiotic use, and travel history^[45].

In another study in Guinea Bissau, of 408 <5 years children, 133 (32.6%) were ESBL-PE, and bed sharing and crowded population was shown as a potential risk factor for ESBL-PE spread^[46].

Another study in Spain among 8- to 16-year-old healthy children showed that 10.7% of 318 *E. coli* isolates from fecal samples were ESBL producers and the SHV-12, CTX-M-1, CTX-M-14, and TEM-52 were the predominant enzyme encoding genes^[47].

A study assessed the risk factors among carbapenem-resistant Enterobacteriaceae concluded that previous hospitalizations within a 3 month period, use of carbapenem, first generation cephalosporin antibiotics and **β-Lactamase** inhibitor, and use of a central catheter of peripheral insertion are significant risk factors^[48]. Among the risk factors of acquisition of CPE isolates, age <1 year, presence of underlying chronic diseases, nasogastric tube placement, surgical intervention, ampicillin usage, and carbapenem use were significant among children population^[49].

In a study conducted in Ethiopia, a high rate of ESBL-PE was determined among neonates and children (74% and 69%, respectively), who were higher than that of adults mostly due to K. pneumonia (76%) and E. coli (46%). In addition 1.9% of them produced carbapenemases^[50].

Most of the risk factors between ESBL producer and carbapenemase producer species of

Enterobacteriaceae are similar, including previous hospitalization, prolonged use of antibiotics, especially cephalosporins, previous colonization, and last 3 month stay in hospital. Other possible potential risk factors that should be considered include the presence of catheters and travel history. It seems that use of fluoroquinolones and nitrofurantoin should also be cautious.

Identifying the associated risk factor of ESBL-PE and CPE carriage in infant populations would help to define the population requiring more control measures to prevent the in-hospital dissemination of ESBL-PE and CPE. Although more studies are needed for the empirical therapy in children, cephalosporins and carbapenems should be used cautiously; likewise, response to the treatment must be followed closely. Limitations of this study were lack of other age groups for the assessment of the risk factors.

Conclusion

Tracking the risk factors associated with ESBL-PE and CPE acquisition among pediatrics has important implications on empiric antibiotic consumption as well as infection control management. The significant emergence of ESBL-PE and CPE pathogens among children populations indicate a serious health threat, which requires the implementation of control measures and surveillance at hospital and outside.

Acknowledgments: This study was performed by the authors.

Ethical Permissions: No ethical permission was stated by the authors.

Conflict of Interests: None declared by authors.

Authors Contribution: Ghasemian A (First author), Original researcher (20%); Norouzi F (Second author), Introduction author (20%); Shokouhi Mostafavi SK (Third author), Introduction author (20%); Hasanvand F (Fourth author), Introduction author (20%); Nojoomi F (Fifth author), Introduction author/ Methodologist/ Original researcher or Assistant/ Statistical analyst/ Discussion author (20%).

Funding/Support: This study was supported by AJA University of Medical Sciences, Tehran, Iran.

References

- 1- Meyer KS, Urban C, Eagan JA, Berger BJ, Rahal JJ. Nosocomial outbreak of Klebsiella infection resistant to late-generation cephalosporins. Ann Intern Med. 1993;119(5):353-8.
- 2- Salehi M, Ghasemian A, Shokouhi Mostafavi SK, Nojoomi F, Ashiani D, Rajabi Vardanjani H. The epidemiology of candida species isolated from urinary tract infections. Arch Clin Infect Dis. 2016;11(4):e37743. 3- Fernandes R, Amador P, Prudêncio C. β -Lactams:

Chemical structure, mode of action and mechanisms of

- resistance. Rev Med Microbiol. 2013;24(1):7-17.
- 4- Spagnolo AM, Orlando P, Panatto D, Perdelli F, Cristina ML. An overview of carbapenem-resistant Klebsiella pneumoniae: Epidemiology and control measures. Rev Med Microbiol. 2014;25(1):7-14.
- 5- Ghasemian A, Salimian Rizi K, Rajabi Vardanjani H, Nojoomi F. Prevalence of clinically isolated metallo-betalactamase-producing pseudomonas aeruginosa, coding genes, and possible risk factors in Iran. Iran J Pathol. 2018;13(1):1-9.
- 6- Rice LB. Federal funding for the study of antimicrobial resistance in nosocomial pathogens: No ESKAPE. J Infect Dis. 2008;197(8):1079-81.
- 7- Hidron AI, Edwards JR, Patel J, Horan TC, Sievert DM, Pollock DA, et al. NHSN annual update: 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, 2006-2007. Infect Control Hosp Epidemiol. 2008;29(11):996-1011.
- 8- Smillie CS, Smith MB, Friedman J, Cordero OX, David LA, Alm EJ. Ecology drives a global network of gene exchange connecting the human microbiome. Nature. 2011;480:241-4.
- 9- Nojoomi F, Ghasemian A. Effect of overgrowth or decrease in gut microbiota on health and disease. Arch Pediatr Infect Dis. 2016;4(2):e34558.
- 10- De Francesco AS, Tanih NF, Samie A, Guerrant RL, Bessong PO. Antibiotic resistance patterns and beta-lactamase identification in Escherichia coli isolated from young children in rural Limpopo province, South Africa: The MAL-ED cohort. S Afr Med J. 2017;107(3):205-14.
- 11- Queenan AM, Bush K. Carbapenemases: The versatile beta-lactamases. Clin Microbiol Rev. 2007;20(3):440-58.
- 12- Diene SM, Rolain JM. Carbapenemase genes and genetic platforms in Gram-negative bacilli: Enterobacteriaceae, pseudomonas and acinetobacter species. Clin Microbiol Infect. 2014;20(9):831-8.
- 13- Poirel L, Le Thomas I, Naas T, Karim A, Nordmann P. Biochemical sequence analyses of GES-1, a novel class A extended-spectrum beta-lactamase, and the class 1 integron In52 from Klebsiella pneumoniae. Antimicrob Agents Chemother. 2000;44(3):622-32.
- 14- Tacão M, Moura A, Correia A, Henriques I. Coresistance to different classes of antibiotics among ESBL-producers from aquatic systems. Water Res. 2014;48:100-7.
- 15- Lucet JC, Chevret S, Decré D, Vanjak D, Macrez A, Bédos JP, et al. Outbreak of multiply resistant Enterobacteriaceae in an intensive care unit: Epidemiology and risk factors for acquisition. Clin Infect Dis. 1996;22(3):430-6.
- 16- Alberer M, Schlenker N, Bauer M, Helfrich K, Mengele C, Löscher T, et al. Detection of gastrointestinal pathogens from stool samples on hemoccult cards by multiplex PCR. Can J Infect Dis Med Microbiol. 2017;2017:3472537.
- 17- Ben-Ami R, Schwaber MJ, Navon-Venezia S, Schwartz D, Giladi M, Chmelnitsky I, et al. Influx of extended-spectrum beta-lactamase-producing Enterobacteriaceae into the hospital. Clin Infect Dis. 2006;42(7):925-34.
- 18- Rodríguez-Baño J, López-Cerero L, Navarro MD, de Alba PD, Pascual A. Faecal carriage of extended-spectrum beta-lactamase-producing Escherichia coli: Prevalence, risk factors and molecular epidemiology. J

- Antimicrob Chemother. 2008;62(5):1142-9.
- 19- Cantón R, Novais A, Valverde A, Machado E, Peixe L, Baquero F, et al. Prevalence and spread of extended-spectrum beta-lactamase-producing Enterobacteriaceae in Europe. Clin Microbiol Infect. 2008;14(Suppl 1):144-53
- 20- Zhao SY, Zhang J, Zhang YL, Wang YC, Xiao SZ, Gu FF, et al. Epidemiology and risk factors for faecal extended-spectrum β -lactamase-producing Enterobacteriaceae (ESBL-E) carriage derived from residents of seven nursing homes in western Shanghai, China. Epidemiol Infect. 2016;144(4):695-702.
- 21- Doernberg SB, Winston LG. Risk factors for acquisition of extended-spectrum β -lactamase-producing Escherichia coli in an urban county hospital. Am J Infect Control. 2012;40(2):123-7.
- 22- Karanika S, Karantanos T, Arvanitis M, Grigoras C, Mylonakis E. Fecal colonization with extended-spectrum beta-lactamase-producing Enterobacteriaceae and risk factors among healthy individuals: A systematic review and metaanalysis. Clin Infect Dis. 2016;63(3):310-8.
- 23- Kennedy K, Collignon P. Colonisation with Escherichia coli resistant to "critically important" antibiotics: A high risk for international travellers. Eur J Clin Microbiol Infect Dis. 2010;29(12):1501-6.
- 24- Zheng B, Dai Y, Liu Y, Shi W, Dai E, Han Y, et al. Molecular epidemiology and risk factors of carbapenemresistant Klebsiella pneumoniae infections in Eastern China. Front Microbiol. 2017;8:1061.
- 25- Solomkin JS, Mazuski JE, Bradley JS, Rodvold KA, Goldstein EJ, Baron EJ, et al. Diagnosis and management of complicated intra-abdominal infection in adults and children: Guidelines by the Surgical Infection Society and the Infectious Diseases Society of America. Clin Infect Dis. 2010;50(2):133-64.
- 26- Denkel LA, Schwab F, Kola A, Leistner R, Garten L, Von Weizsäcker K, et al. The mother as most important risk factor for colonization of very low birth weight (VLBW) infants with extended-spectrum β -lactamase-producing Enterobacteriaceae (ESBL-E). J Antimicrob Chemother. 2014;69(8):2230-7.
- 27- Flokas ME, Detsis M, Alevizakos M, Mylonakis E. Prevalence of ESBL-producing Enterobacteriaceae in paediatric urinary tract infections: A systematic review and meta-analysis. J Infect. 2016;73(6):547-57.
- 28- Zaoutis TE, Goyal M, Chu JH, Coffin SE, Bell LM, Nachamkin I, et al. Risk factors for and outcomes of bloodstream infection caused by extended-spectrum beta-lactamase-producing Escherichia coli and Klebsiella species in children. Pediatrics. 2005;115(4):942-9.
- 29- Logan LK, Meltzer LA, McAuley JB, Hayden MK, Beck T, Braykov NP, et al. Extended-spectrum β -lactamase-producing Enterobacteriaceae infections in children: A two-center case-case-control study of risk factors and outcomes in Chicago, Illinois. J Pediatric Infect Dis Soc. 2014;3(4):312-9.
- 30- Colodner R, Rock W, Chazan B, Keller N, Guy N, Sakran W, et al. Risk factors for the development of extended-spectrum beta-lactamase-producing bacteria in nonhospitalized patients. Eur J Clin Microbiol Infect Dis. 2004;23(3):163-7.
- 31- Topaloglu R, Er I, Dogan BG, Bilginer Y, Ozaltin F, Besbas N, et al. Risk factors in community-acquired urinary tract infections caused by ESBL-producing

- bacteria in children. Pediatr Nephrol. 2010;25(5):919-25
- 32- Søraas A, Sundsfjord A, Sandven I, Brunborg C, Jenum PA. Risk factors for community-acquired urinary tract infections caused by ESBL-producing Enterobacteriaceae a case control study in a low prevalence country. PloS One. 2013;8(7):e69581.
- 33- Lo WU, Ho PL, Chow KH, Lai EL, Yeung F, Chiu SS. Fecal carriage of CTXM type extended-spectrum beta-lactamase-producing organisms by children and their household contacts. J Infect. 2010;60(4):286-92.
- 34- Ding H, Yang Y, Lu Q, Wang Y, Chen Y, Deng L, et al. The prevalence of plasmid-mediated AmpC beta-lactamases among clinical isolates of Escherichia coli and Klebsiella pneumoniae from five children's hospitals in China. Eur J Clin Microbiol Infect Dis. 2008;27(10):915-21.
- 35- Kantele A, Lääveri T, Mero S, Vilkman K, Pakkanen SH, Ollgren J, et al. Antimicrobials increase travelers' risk of colonization by extended-spectrum betalactamase-producing Enterobacteriaceae. Clin Infect Dis. 2015;60(6):837-46.
- 36- Strysko JP, Mony V, Cleveland J, Siddiqui H, Homel P, Gagliardo C. International travel is a risk factor for extended-spectrum $$\beta$$ -lactamase-producing Enterobacteriaceae acquisition in children: A case-case-control study in an urban U. S. hospital. Travel Med Infect Dis. 2016;14(6):568-71.
- 37- Rivard-Yazigi L, Zahar JR, Le Guillou S, Chalouhi C, Lecuyer H, Bureau C, et al. Risk factors associated with extended-spectrum β -lactamase-producing Enterobacteriaceae carriage at admission in an infant cohort at a tertiary teaching hospital in France. Am J Infect Control. 2013;41(9):844-5.
- 38- Hijazi SM, Fawzi MA, Ali FM, Abd El Galil KH. Multidrug-resistant ESBL-producing Enterobacteriaceae and associated risk factors in community infants in Lebanon. J Infect Dev Ctries. 2016;10(9):947-55.
- 39- Pessoa-Silva CL, Meurer Moreira B, Câmara Almeida V, Flannery B, Almeida Lins MC, Mello Sampaio JL, et al. Extended-spectrum beta-lactamase-producing Klebsiella pneumoniae in a neonatal intensive care unit: Risk factors for infection and colonization. J Hosp Infect. 2003;53(3):198-206.
- 40- Kim YK, Pai H, Lee HJ, Park SE, Choi EH, Kim J, et al. Bloodstream infections by extended-spectrum beta-lactamase-producing Escherichia coli and Klebsiella pneumoniae in children: Epidemiology and clinical outcome. Antimicrob Agents Chemother. 2002;46(5):1481-91.
- 41- Linkin DR, Fishman NO, Patel JB, Merrill JD, Lautenbach E. Risk factors for extended-spectrum beta-lactamase-producing Enterobacteriaceae in a neonatal intensive care unit. Infect Control Hosp Epidemiol. 2004;25(9):781-3.
- 42- Venezia RA, Scarano FJ, Preston KE, Steele LM, Root TP, Limberger R, et al. Molecular epidemiology of an SHV-5 extended-spectrum beta-lactamase in Enterobacteriaceae isolated from infants in a neonatal intensive care unit. Clin Infect Dis. 1995;21(4):915-23.
- 43- Pessoa-Silva CL, Toscano CM, Moreira BM, Santos AL, Frota AC, Solari CA, et al. Infection due to extended-spectrum beta-lactamase-producing Salmonella enterica subsp. enterica serotype infantis in a neonatal unit. J Pediatr. 2002;141(3):381-7.

44- Pragosa H, Marçal M, Gonçalves E, Martins F, Lopo-Tuna M. Multi-drug-resistant Enterobacteriaceae in a Portuguese neonatal intensive care unit. J Hosp Infecti. 2017;96(2):130-1.

45- Birgy A, Levy C, Bidet P, Thollot F, Derkx V, Béchet S, et al. ESBL-producing Escherichia coli ST131 versus non-ST131: Evolution and risk factors of carriage among French children in the community between 2010 and 2015. J Antimicrob Chemother. 2016;71(10):2949-56.

46- Isendahl J, Turlej-Rogacka A, Manjuba C, Rodrigues A, Giske CG, Nauclér P. Fecal carriage of ESBL-producing E. coli and K. pneumoniae in children in Guinea-Bissau: A hospital-based cross-sectional study. PloS One. 2012;7(12):e51981.

47- Fernández-Reyes M, Vicente D, Gomariz M, Esnal O, Landa J, Oñate E, et al. High rate of fecal carriage of extended-spectrum- β -lactamase-producing Escherichia coli in healthy children in Gipuzkoa, Northern Spain. Antimicrob Agents Chemother. 2014;58(3):1822-4.

48- César García C J, Amaya S, Briceño C W, Rincón C, Pinzón J. Risk factors for carbapenem-resistant bacterial infection or colonization: A case control study. Acta Colombiana de Cuidado Intensivo. 2017;17(1):29-35. [Spanish]

49- Karaaslan A, Soysal A, Altinkanat Gelmez G, Kepenekli Kadayifci E, Söyletir G, Bakir M. Molecular characterization and risk factors for carbapenemresistant Gram-negative bacilli colonization in children: Emergence of NDM-producing Acinetobacter baumannii in a newborn intensive care unit in Turkey. J Hosp Infect. 2016;92(1):67-72.

50- Desta K, Woldeamanuel Y, Azazh A, Mohammod H, Desalegn D, Shimelis D, et al. High gastrointestinal colonization rate with extended-spectrum β -lactamase-producing Enterobacteriaceae in hospitalized patients: Emergence of carbapenemase-producing K. pneumoniae in Ethiopia. PloS One. 2016;11(8):e0161685.