

# EVOLUTION OF IMIPENEM RESISTANCE IN ESCHERICHIA COLI : A 9-YEAR RETROSPECTIVE SURVEILLANCE REPORT IN A HOSPITAL POPULATION OF SOUTHERN CHINA

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**Abstract:** *Escherichia coli* (*E. coli*) is one of the most common causes of hospital-acquired infection. Our aim was to assess the evolution of Imipenem Resistance in *Escherichia coli* at the First Affiliated Hospital of Jinan University (Guangzhou, China) from 2010 to 2018. 3412 non-duplicate strains of *E. coli* were isolated from inpatients and outpatients during the researched period. The majority of isolates appeared in the urine sample (57.68%) were derived from Urology(33.76%). With an average imipenem resistance rate of 0.61%, 27 strains of *E. coli* were detected. From 2010 to 2018, the resistance rates of the 3412 strains of *E. coli* to imipenem were 0, 0.00%, 0.35%, 0.30%, 0.26%, 0.44%, 1.08%, 0.70%, 2.33%, respectively. The overall imipenem resistant rate of *E. coli* was gradually increased in recent years. Moreover, the imipenem-resistant *E. coli* was multi-drug resistant to the antibiotics used clinically.. Therefore, restraining the use of antibiotics is extremely urgent to avoid severe infection by resistant pathogens.

**Key words:** *Escherichia coli*; Imipenem; Antibiotic resistance.

## INTRODUCTION

*Escherichia coli* (*E. coli*), is a Gram-negative bacillus belonging to the normal intestinal flora of humans and other mammals. Depending on its conditional pathogenicity, Various intestinal and extra-intestinal infections are caused by *E. coli*. It has also occupied the first in the detection rate of pathogenic bacteria in clinical specimens, mainly isolated from urine samples [1, 2].

Carbapenems, including imipenem, possess broad spectrum antibacterial activity and have a unique structure that is defined by a carbapenem coupled to a  $\beta$ -lactam ring which confers protection against most  $\beta$  lactamases such as metallo- $\beta$ -lactamase (MBL) as well as extended spectrum  $\beta$ -lactamases. Consequently, carbapenems are considered as one of the most reliable drugs for treating bacterial infections [3]. However, with the widespread use of carbapenems, carbapenem-resistant

*Enterobacteriaceae* (CRE) has gradually emerged in clinical practice all over the world. In 2017, the European Centre for Disease Prevention and Control (ECDC) published the bacterial resistance monitoring of 30 European countries, it showed that the drug-resistant rate of CRE gradually increased from 6.0% (2012) to 8.1% (2015) [4]. Meanwhile, the China Antimicrobial Resistance Surveillance System(CARSS) reported the similar increasing resistance rate of *E. coli* to imipenem, which was elevated from 0.3% (2005) to 1.9% (2017) [5].

The mechanisms of resistance of *Enterobacteriaceae* to carbapenems are complex including the productions of carbapenemase, the changes of penicillin-binding protein targets, the overexpression of Amp C or ESBLs combined with mutations and deletions of outer porin channel proteins, the activation of external Enhancement of the exhaust system [6-9], the above mechanisms can

mediate resistance to carbapenems through individual or synergistic effects [10]. Ambler classification divides this enzyme into type A (KPC, SME, NMC-A, IMI), type B (NDM, VIM, IMP) and type D (OXA), in which type A and type D are serine enzymes, and their active sites have serine structure, while type B is metalloenzyme, which needs metal ion  $Zn^{2+}$  to play the activity and can be inhibited by EDTA [11].

CRE-induced drug resistance is often characterized by multiple drug resistance, which makes clinical treatment of patients with CRE infection very difficult [12, 13]. The aim of this study was to evaluate the drug-resistance changes of *E. coli* to imipenem at the First Affiliated Hospital of Jinan University (Guangzhou, China) from 2010 to 2018.

## MATERIALS AND METHODS

### *Hospital setting and strains collection*

The *E. coli* isolates were collected at the First Affiliated Hospital of Jinan University (Guangzhou, China) from 2010 to 2018. Only the first pathogens isolated from the patients were selected, the strains with unclear clinical information were excluded. The samples included urine, blood, sputum, ascites and wound secretions. Bacterial identification to the species level on all tested strains was performed by standard procedures reported previously [14, 15].

### *Susceptibility testing*

The MIC (Minimum Inhibitory Concentration) concentration was automatically tested by a VITEK 2 Compact automatic microorganisms analyzer. The Drug susceptibility cards (VITEK 2 AST-N335, Merrier diagnostics (Shanghai) co., LTD) were used to absorb the pure bacterial colony suspension of the positive samples with McKessler's unit turbidity required by the drug sensitivity test. The results were interpreted according to the 2018 criteria of Clinical and Laboratory Standards Institute (CLSI). The ATCC25922 *E. coli* from the Guangdong Center for Clinical Laboratory was used as control.

### *Statistical analysis*

Antimicrobial susceptibility results and organization were managed by WHONET (version 5.6). A chi-square test or Fisher's exact test was performed to exam the difference between the resistance rates of inpatients and outpatients. A  $p < 0.05$  was defined as statistically significant.

## RESULTS

### *Detection quantity and detection rate of E. coli*

A total of 3412 strains were isolated from January 2010 to December 2018, including 176 strains in 2010, 198 strains in 2011, 283 strains in 2012, 328 strains in 2013, 389 strains in 2014, 451 strains in 2015, 462 strains in 2016, 568 strains in 2017 and 557 strains in 2018. The average detection rate of *E.*

*Coli* nosocomial infection in the study years was 13.78%. The specific numbers were 13.84%, 12.32%, 14.40%, 15.12%, 14.47%, 14.67%, 12.35%, 11.44% and 17.01%, from 2010 to 2018, respectively (Figure 1).

### *Detection time distribution of E. coli*

The third quarter (July to September) showed the highest detection rate (27.67%) and the first quarter (January to March) showed the lowest (19.7%) of *E. coli* during the study period (Figure 2).

### *Departments distribution and source of specimens of E. coli*

The urology department had the highest detection rate of *E. coli* (33.76%), followed by obstetrics and gynecology (8.65%) and pediatrics (8.03%) (Table 1). *E. coli* was mainly detected from urine (57.68%), blood (12.57%) and sputum (8.70%) (Table 2).

### *Age distribution of E. coli*

The study cohort was grouped as infant (<1 year), youth (1-40 years), middle-aged (40-60 years) and agedness (>60 years) according to the World Health Organization (WHO) age classification standard. Most of the patients diagnosed with *E. coli* infection were agedness (44.61%), followed by the middle-aged group (24.97%), and the infant group had the lowest detection rate (5.80%) (Figure 3).

### *Imipenem-resistance rates of E. coli*

A total of 27 non-repeat *E. coli* strains were resistant to imipenem. The average resistance rate was 0.61%. The specific numbers were 0.00%, 0.00%, 0.35%, 0.30%, 0.26%, 0.44%, 1.08%, 0.70% and 2.33%, from 2010 to 2018, respectively (Figure 4).

By the year of 2011, the sensitivity rate of *E. coli* to imipenem was still 100.00%. The imipenem resistant began to appear in 2012. There was a significantly increasing of resistant strains since 2015. The highest number of resistant strains was detected in 2018 (Figure 5).

The total number of strains is the number of positive strains isolated and cultured in the hospital within one year, including all common and uncommon gram-positive bacteria and gram-negative bacteria. The number of *E. coli* strains is the number of *E. coli* strains isolated and cultured in the hospital within one year.

Composition ratio: The number of *E. coli* strains/ The total number of strains.

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Composition ratio: The number of *E. coli* strains/ The total number of strains

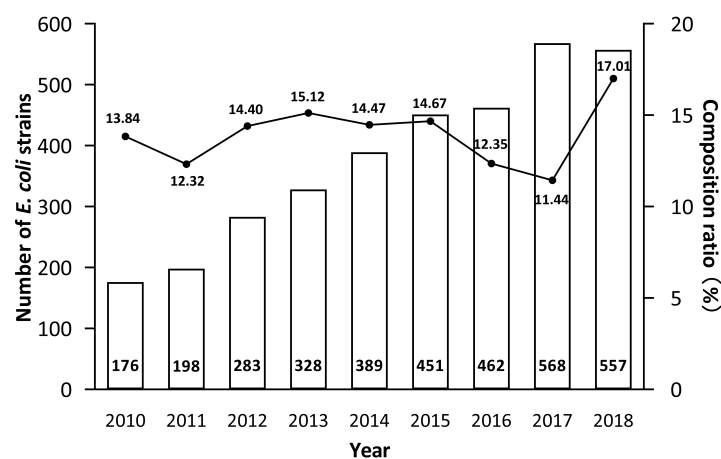


Figure 1 Statistics of detected *E. coli* strains from 2010 to 2018

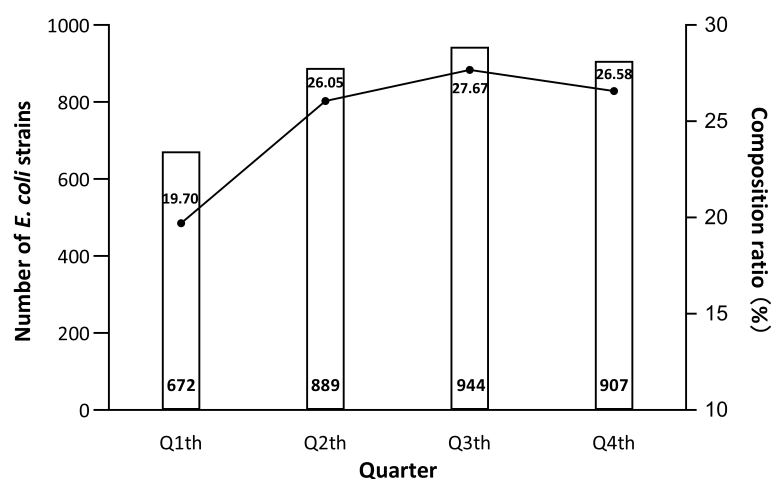


Figure 2 Time distribution of detection of *E. coli* strains from 2010 to 2018

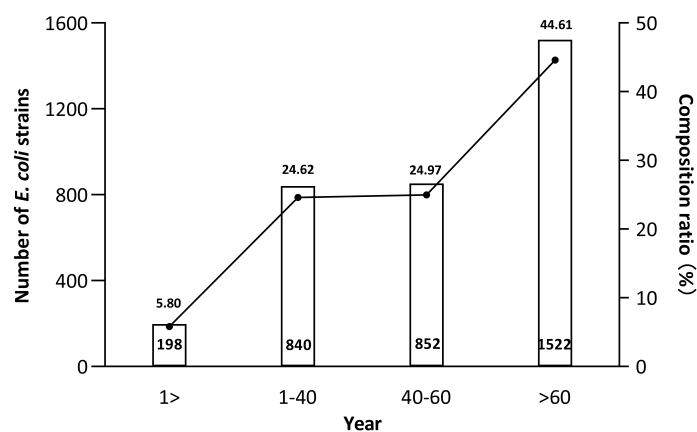
Table 1 Department distribution of detection of *E. coli* strains from 2010 to 2018

Department	Number of <i>E. coli</i> strains (n)	Composition ratio(%)
Urology	1152	33.76
Obstetricand Gynecology	295	8.65
Pediatrics	274	8.03
orthopedics	177	5.19
Hepatobiliary surgery	164	4.81
Respiratory	163	4.78
Digestion	153	4.48
Cardiovascular	133	3.90
Neurology	127	3.72
ICU	107	3.14
Others	667	19.55

Table 2 Specimen sources of *E. coli* strains from 2010 to 2018

Specimen sources	Number of <i>E. coli</i> strains (n)	Composition ratio(%)
Urine	1968	57.68
Blood	429	12.57

Sputum	297	8.70
Others	189	5.54
Secretions	112	3.28
Purulence	78	2.29
Amniotic fluid	45	1.32
Cerebrospinal fluid, pleural fluid, joint	44	1.29
Bile, gastric juices	250	7.33



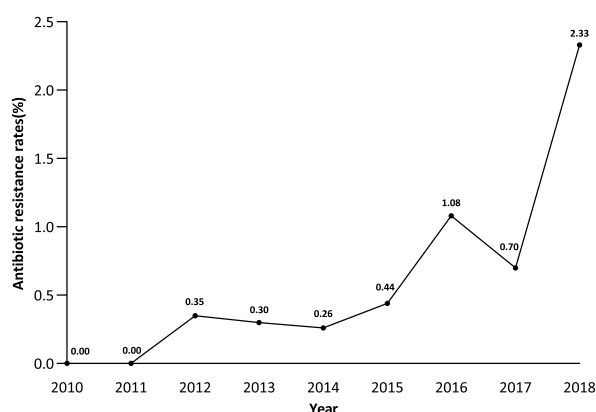
**Figure 3** The age distribution of patients with *E. coli* detected from 2010 to 2018

The total number of strains is the number of positive strains isolated and cultured in the hospital within one age group, including all common and uncommon gram-positive bacteria and gram-negative bacteria. The number of *E. coli* strains is the number of *E. coli* strains isolated and cultured in the hospital within one age group.

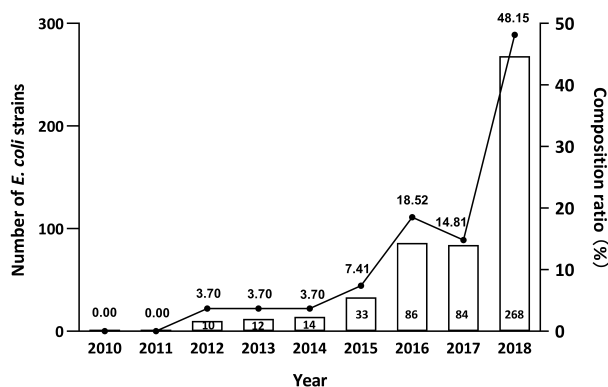
Composition ratio: The number of *E. coli* strains/ The total number of strain

The total number of strains is the number of positive strains isolated and cultured in the hospital within one year, including all common and uncommon gram-positive bacteria and gram-negative bacteria. The number of imipenem resistant *E. coli* strains is the number of imipenem resistant *E. coli* strains isolated and cultured in the hospital within a year.

Antibiotic resistance rates: The number of imipenem resistance *E. coli* strains/ The total number of strains



**Figure 4** The antibiotic resistance rate of *E. coli* to imipenem from 2010 to 2018



**Figure 5** Detection time distribution of imipenem resistant *E. coli* strains from 2010 to 2018

The total number of *E. coli* strains is the number of positive *E. coli* strains isolated and cultured in the hospital within one year. The number of imipenem resistant *E. coli* strains is the number of imipenem resistant *E. coli* strains isolated and cultured in the hospital within one year.

Composition ratio: The number of imipenem resistant *E. coli* strains / The total number of *E. coli* strains

**Department distribution and specimen source of imipenem resistant *E. coli***

From January 2010 to December 2018, the departments with the highest detection rate of imipenem-resistant *E. coli* were the urology department (25.93%), followed by the respiratory department (14.81%) and the gastroenterology department (11.11%), as shown in Table 3. Imipenem-resistant *E. coli* was mainly detected from urine (44.44%), sputum (22.22%) and blood (18.52%), as shown in Table 4.

**Table 3** Department distribution of imipenem resistant *E. coli* strains from 2010 to 2018

Department	Number of imipenem resistant <i>E. coli</i> strains (n)	Composition ratio(%)
Urology	7	35.93
Respiratory	4	14.81
Digestion	3	11.1
Pediatrics	2	7.41
Neurology	2	7.41
ICU	1	3.70
Cardiovascular	1	3.70
Hepatobiliary	1	3.70
Orthopedics	1	3.70
Others	5	18.52

**Table 4** Specimen sources of imipenem resistant *E. coli* strains from 2010 to 2018

Specimen sources	Number of imipenem	Composition ratio(%)
Urine	12	44.44
Sputum	6	22.22
Blood	5	18.52
Wound secretion	2	7.41
Ascites	1	3.70
Others	1	3.70

**Susceptibility of imipenem-resistant *E. coli* to other antimicrobial agents**

The resistant rate of the 27 strains to other antimicrobial drugs was also evaluated. 10 antibiotics, including cephalosporins (ceftazidime and cefepime),

quinolones (ciprofloxacin and ofloxacin), aminoglycosides (amikacin and tobramycin), sulfa (sulfamethoxazole), monocyclic- $\beta$ -lactamases (aztreonam) and combinations of  $\beta$ -lactamase inhibitors (piperacillin / tazobactam and cefoperazone / sulbactam) were selected to conduct

the evaluation. More than 50% of the imipenem-resistant *E. coli* showed multi-drug resistance to the tested antibiotics.  $\beta$ -lactamases showed the highest resistant rate of 90%, followed by tobramycin (48.15%), and amikacin showed the lowest (25.93%) (Table 5).

**Table 5 Susceptibility of 27 imipenem resistant *E. coli* strains to 10 antimicrobial agents**

Antibacterial drugs	Resistance		Intermediary		Sensitive	
	strains	resistance rate (%)	strains	Intermediary rate (%)	strains	Sensitivity rate (%)
Ceftazidime	25	92.59	0	0.00	2	7.41
Cefepime	25	92.59	0	0.00	2	7.41
Ciprofloxacin	19	70.37	2	7.41	6	22.22
Levofloxacin	19	70.37	2	7.41	6	22.22
Amikacin	7	25.93	0	0.00	20	74.07
Tobramycin	13	48.15	3	11.11	11	40.74
Trimethoprim/ sulfamethoxazole	17	62.96	0	0.00	10	37.04
Aztreonam	24	88.89	0	0.00	3	11.11
Piperacillin/ Tazobactam	24	88.89	0	0.00	3	11.11
Cefoperazone/ sulbactam	25	92.59	0	0.00	2	7.41

## DISCUSSION

*E. coli* is the most common pathogen causing nosocomial infections and is widely found in hospital settings. When the patient's immune function is low, *E. coli*, which is a normal flora in the human intestine, will be ectopically colonized to other tissues or organs, causing urinary tract infections, respiratory infections, blood infections, and abdominal infections [16]. This article showed that from 2010 to 2018, the average detection rate of *E. coli* in the First Affiliated Hospital of Jinan University was 13.78%, and the average resistance rate to imipenem was 0.61%, which was lower than the 2017 CHINET drug resistance monitoring reported rate 1.90% [17]. This might be related to the different use of antibacterial drugs in different regions and different levels of hospitals. The detection rate of the number of *E. coli* isolates from 2010 to 2017 was stabilized and had no obviously increasing trend, but the detection rate rose sharply to 17.01% in 2018, which was much higher than the average detection rate of 13.78%. Meanwhile, the number of imipenem-resistant *E. coli* isolated in 2018 was 13 isolates, accounting for 48.15% of the resistant strains isolated during the nine-year period. This might be a result of the surge in outpatients, inadequate nursing staff, and insufficient management of clinical medications in 2018.

The nine years of continual monitoring results showed that among the 3412 detected strains of *E. coli*, urine specimens (57.68%) were the most common sample, followed by blood specimens

(12.57%) and sputum specimens (8.70%). *E. coli* was the main pathogen that causing urinary tract infections in our hospital, which was consistent with other domestic reports [18, 19]. It has been reported that *E. coli* is easy to cause urinary tract infection because the fimbriae of *E. coli* can adhere firmly to the epithelial cells of the urethra, thus weakening the peristalsis of the ureter, leading to ureteral dilatation, so that the urine can't wash away the bacteria in time [20]. On the perspective of department distribution, *E. coli* has the highest detection rate in the urology department (33.76%). It has more invasive procedures such as urinary catheters and cystoscopy with patients in this ward, which causes urinary tract mucosal damage and then more likely to cause *E. coli* infection. According to the age distribution, *E. coli* infection mainly occurred in the elderly group (44.61%) over 60 years old. On the one hand, with the increase of age, the patient's organs aged and atrophied, degenerative changes in the urinary tract, obstacles to urination reflexes, increased residual urine volume, and decreased body immunity, making urinary tract infections more prone. On the other hand, *E. coli* is the common pathogen of urinary infection [21].

In recent years, imipenem-resistant *E. coli* at home and abroad has been widely reported [22, 23]. In 2012, the first imipenem-resistant *E. coli* strain appeared in our hospital. Carbapenemase was the main mechanism of resistance of Enterobacter bacteria to carbapenems. According to the reports, clinically isolated carbapenem-resistant *E. coli* in

China commonly produce KPC enzymes and NDM-1 (New Delhi metallo- $\beta$ -lactamase-1) enzymes [24, 25]. Because the carbapenemase gene was mostly located on a conjugable plasmid, the resistance to carbapenem can also be transmitted between different strains and different species through plasmid transfer, which may also easily cause resistant strains widespread [26]. In this study, 13 imipenem-resistant *E. coli* were detected in 2018, accounting for 48.15% of the resistant strains isolated during the nine-year period, which was significantly higher than the number detected in 2017 and was most likely a plasmid transfer caused by the carbapenemase gene. At the same time, a statistical analysis of the susceptibility of 27 imipenem-resistant *Escherichia coli* strains revealed that Imipenem-resistant *E. coli* was multi-drug resistant to most commonly used antibacterials, but it had a high sensitivity rate to aminoglycosides amikacin. Aminoglycosides are one of the few drugs with in vitro resistance to carbapenem-resistant Enterobacteriaceae, which is consistent with Smith K [27]. Therefore, the clinicians should timely grasp the drug resistance characteristics of *E. coli*, try to use drugs reasonably according to the drug sensitivity results, and avoid the continuous resistance of imipenem-resistant *E. coli* to aminoglycosides increase. At the same time, more attention should be paid to the isolation of patients with multi-drug resistant bacterial infections to prevent their spread in hospitals.

## CONCLUSION

During the nine years from 2010 to 2018, urine specimens were the main source of specimens for clinical isolation of *E. coli*, while Urology was the main department for sending specimens for *E. coli* infection. The results were consistent with that of imipenem-resistant *E. coli*. The overall drug resistance rate of *E. coli* to imipenem was increasing gradually, and there were multiple drug resistance to common antibiotics.

## ETHICS APPROVAL

Given that this study was performed without accessing patient information, approval of the ethics committee was not required.

## ACKNOWLEDGMENTS

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## REFERENCES

1. Kaper JB, Nataro JP, and Mobley HLT. Pathogenic *Escherichia coli*. *Nat Rev Microbiol*. 2004; 2:123-140.
2. Alabsi MS, Ghazal A, Sabry SA, and Alasaly MM. Association of some virulence genes with antibiotic resistance among uropathogenic *Escherichia coli* isolated from urinary tract infection patients in Alexandria, Egypt: A hospital-based study. *Journal of Global Antimicrobial Resistance*. 2014;2:83-86.
3. Codjoe, F. S., Donkor, E. S. Carbapenem Resistance: A Review. *Med Sci*. 2017;(Basel). 6.
4. Magiorakos, A. P., Burns, K., Roiguez Bano, J., Borg, M., Daikos, G., Dumpis, U., Voss, A., and Weber, J. T. Infection prevention and control measures and tools for the prevention of entry of carbapenem-resistant Enterobacteriaceae into healthcare settings: guidance from the European Centre for Disease Prevention and Control. *Antimicrobial Resistance and Infection Control*. 2017;6:113-117.
5. Shaowei Zheng, Ping Li, Zhengliang Zhang, and Honghong Fei. Surveillance of carbapenem antibiotic resistance of common gram-negative bacteria in CHINET in China from 2005 to 2017. *Journal of Clinical Emergency*. 2019;20:45-49.
6. Karaikos I, Giamarellou H. Multidrug-resistant and extensively drug-resistant Gram-negative pathogens: current and emerging therapeutic approaches. *Expert Opin Pharmacother*. 2014;15: 1351-1370.
7. Yamachika, S., Sugihara, C., Kamai, Y., and Yamashita, M. Correlation between penicillin-binding protein 2 mutations and carbapenem resistance in *Escherichia coli*. *Journal of medical microbiology*. 2013;62:429.
8. Wozniak, A., Villagra, N. A., Undabarrena, A., Gallardo, N., Keller, N., Moraga, M., Román, J. C., Mora, G. C., and García, P. Porin alterations present in non-carbapenemase-producing Enterobacteriaceae with high and intermediate levels of carbapenem resistance in Chile. *Journal of medical microbiology*. 2012;61:1270-1279.
9. Skalweit MJ, Li M. Bulgecin A as a  $\beta$ -lactam enhancer for carbapenem-resistant *Pseudomonas aeruginosa* and carbapenem-resistant *Acinetobacter baumannii* clinical isolates containing various resistance mechanisms. *Drug design, development and therapy*. 2016;10:3013-3020.
10. Grundmann, H., Glasner, C., Albiger, B., Aanensen, D. M., Tomlinson, C. T., Andrasević, A. T., Cantón, R., Carmeli, Y., Friedrich, A. W., Giske, C. G., Glupczynski, Y., Gniadkowski, M., Livermore, D. M., Nordmann, P., Poirel, L., Rossolini, G. M., Seifert, H., Vatopoulos, A., Walsh, T., Woodford, N., Monnet, D. L., Koraqi, A., Lacey, D., Apfalter, P., Hartl, R., Glupczynski, Y., Huang, T., Strateva, T., Marteva-Proevska, Y., Andrasevic, A. T., Butic,

- I., Pieridou-Bagatzouni, D., Maikanti-Charalampous, P., Hrabak, J., Zemlickova, H., Hammerum, A., Jakobsen, L., Ivanova, M., Pavelkovich, A., Jalava, J., Österblad, M., Dortet, L., Vaux, S., Kaase, M., Gatermann, S. G., Vatopoulos, A., Tryfinopoulou, K., Tóth, Á., Jánvári, L., Boo, T. W., McGrath, E., Carmeli, Y., Adler, A., Pantosti, A., Monaco, M., Raka, L., Kurti, A., Balode, A., Saule, M., Miculeviciene, J., Mierauskaite, A., Perrin-Weniger, M., Reichert, P., Nestorova, N., Debattista, S., Mijovic, G., Lopacic, M., Samuelson, Ø., Haldorsen, B., Zabicka, D., Literacka, E., Caniça, M., Manageiro, V., Kaftandzieva, A., Trajkovska-Dokic, E., Damian, M., Lixandru, B., Jelesic, Z., Trudic, A., Niks, M., Schreterova, E., Pirs, M., Cerar, T., Oteo, J., Aracil, B., Giske, C., Sjöström, K., Gür, D., Cakar, A., Woodford, N., Hopkins, K., Wiuff, C., and Brown, D. J. Occurrence of carbapenemase-producing *Klebsiella pneumoniae* and *Escherichia coli* in the European survey of carbapenemase-producing Enterobacteriaceae (EuSCAPE): a prospective, multinational study. *The Lancet Infectious Diseases* 2017;17:153-163.
11. Kazi, M., Drego, L., Nikam, C., Ajbani, K., Soman, R., Shetty, A., and Rodrigues, C. 2015. Molecular characterization of carbapenem-resistant Enterobacteriaceae at a tertiary care laboratory in Mumbai. *European journal of clinical microbiology & infectious diseases* : official publication of the European Society of Clinical Microbiology 2015;34: 467-472.
12. Yin Danping, Zhou Guoqing. Study on the clinical distribution and drug resistance of carbapenem resistant Enterobacteriaceae. 2018. *Chinese Journal of nosocomial infection* 2018;28: 2752-2755.
13. Zilberberg MD, Shorr AF. Prevalence of multidrug-resistant *Pseudomonas aeruginosa* and carbapenem-resistant Enterobacteriaceae among specimens from hospitalized patients with pneumonia and bloodstream infections in the United States from 2000 to 2009. *J Hosp Med* 2013; 8: 559-563
14. Xu, Z., Li, L., Shirtliff, M. E., Peters, B. M., Peng, Y., Alam, M. J., Yamasaki, S., and Shi, L. 2010. First report of class 2 integron in clinical *Enterococcus faecalis* and class 1 integron in *Enterococcus faecium* in South China. *Diagnostic Microbiology and Infectious Disease* 2010; 68: 315-317.
15. Xie J, Peters BM, Li B, Li L, Yu G, Xu Z, and Shirtliff ME. Clinical features and antimicrobial resistance profiles of important Enterobacteriaceae pathogens in Guangzhou representative of Southern China, 2001-2015. *Microbial Pathogenesis* 2017; 107: 206-211.
16. van Duin D, Kaye KS, Neuner EA, and Bonomo RA. Carbapenem-resistant Enterobacteriaceae: a review of treatment and outcomes. *Diagnostic Microbiology & Infectious Disease* 2013; 75: 115-120.
17. Fupin Hu, Demei Zhu, Fu Wang, Xiaofei Jiang, Qing Yang, Yingchun Xu, Xiaojiang Zhang, Ziyong Sun, Zhongju Chen, Chuanqing Wang, Aimin Wang, Yuxing Ni, Jingyong Sun, Yunsong Yu, Jie Lin, Bin Shan, Yan Du, Yuanhong Xu, Jilu Shen, Hong Zhang, Jing Kong, Chao Zhou, Danhong Su, Zhaoxia Zhang, Ping Ji, Yunjian Hu, Xiaoman Ai, Wenxiang Huang, Bei Jia, Lianhua Wei, and Ling Wu. 2017 China bacterial resistance monitoring. *China Journal of infection and chemotherapy* 2018; 18: 241-251.
18. Luo Ling, Liu Xiaoqiang, Ou Yangying, et al. Drug resistance analysis of 1411 strains of *Escherichia coli*. *Chinese Journal of health inspection* 2012; 22: 1706-1707.
19. Ke Zhao, Pengcheng Xia, Zhijun Zhang, and Shuping Zhao. Clinical distribution and drug resistance analysis of 4828 strains of *Escherichia coli*. *International Journal of laboratory medicine* 2018; 39: 1765-1768.
20. Zhang, R., Liu, L., Zhou, H., Chan, E. W., Li, J., Fang, Y., Li, Y., Liao, K., and Chen, S. Nationwide Surveillance of Clinical Carbapenem-resistant Enterobacteriaceae (CRE) Strains in China. *EBioMedicine* 2017; 19: 98-106.
21. Toner, L., Papa, N., Aliyu, S. H., Dev, H., Lawrentschuk, N., and Al-Hayek, S. Extended-spectrum beta-lactamase-producing Enterobacteriaceae in hospital urinary tract infections: incidence and antibiotic susceptibility profile over 9 years. *World J Urol* 2016; 34: 1031-1037.
22. Wang, Q., Zhang, Y., Yao, X., Xian, H., Liu, Y., Li, H., Chen, H., Wang, X., Wang, R., Zhao, C., Cao, B., and Wang, H. Risk factors and clinical outcomes for carbapenem-resistant Enterobacteriaceae nosocomial infections. *Eur J Clin Microbiol Infect Dis* 2016; 35: 1679-1689.
23. Pannaraj, P. S., Bard, J. D., Cerini, C., and Weissman, S. J. Pediatric carbapenem-resistant Enterobacteriaceae in Los Angeles, California, a high-prevalence region in the United States. *The Pediatric infectious disease journal* 2015; 34: 11-16.
24. Cai, J. C., Zhang, R., Hu, Y. Y., Zhou, H. W., and Chen, G. X. Emergence of *Escherichia coli* sequence type 131 isolates producing KPC-2 carbapenemase in China. *Antimicrob Agents Chemother* 2014; 58: 1146-1152.
25. Yang, Q., Fang, L., Fu, Y., Du X, Shen, Y., and Yu, Y. Dissemination of NDM-1-Producing Enterobacteriaceae Mediated by the



- IncX3-Type Plasmid. PLoS One. 2015;10:e129454.
26. Qu, H., Wang, X., Ni, Y., Liu, J., Tan, R., Huang, J., Li, L., and Sun, J. NDM-1-producing Enterobacteriaceae in a teaching hospital in Shanghai, China: IncX3-type plasmids may contribute to the dissemination of blaNDM-1. International Journal of Infectious Diseases 2015; 34: 8-13.
27. Smith K.P, Kirby J E. Evaluation of apramycin activity against carbapenem-resistant and susceptible strains of Enterobacteriaceae. Diagnostic Microbiology and Infectious Disease 2016; 86: 439-441.