Published online: 2024 July 27

**Research Article** 



# Epidemiological Characteristics and Drug Resistance Analysis of 187 Strains of Carbapenem-Resistant *Enterobacteriaceae* in a General Hospital

Maojie Zhang 🗓 <sup>1</sup>, Yanqiu Zhu <sup>2</sup>, Mei Yang <sup>2</sup>, Yamei Huang <sup>2</sup>, Yang Yu <sup>1</sup>, Shengwei Wu <sup>1,\*</sup>

<sup>1</sup> Department of Infection Management, First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine, Guizhou, China <sup>2</sup> Department of Laboratory, First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine, Guizhou, China

\*Corresponding author: Department of Infection Management, First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine, Guizhou, China. Email: 532159027@qq.com

Received 2024 June 22; Revised 2024 July 28; Accepted 2024 August 1

# Abstract

**Background:** Carbapenem antibiotics are currently the most broad-spectrum and potent  $\beta$ -lactam antibiotics. Without effective intervention measures, Carbapenem-Resistant *Enterobacteriaceae* (CRE) may become widespread in all healthcare institutions within ten years, potentially developing into an endemic issue.

**Objectives:** To understand the epidemiological characteristics and multilocus sequence typing (MLST) of CRE infections in hospitals, and to explore critical strategies for preventing and controlling CRE hospital infections.

**Methods:** The study focused on hospitalized patients with CRE strains isolated by the Laboratory Department of the First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine from 2020 to 2022. Patient data, bacterial isolation, culture, and drug sensitivity results were collected and analyzed to determine the epidemiological characteristics of CRE infections. Preserved Carbapenem-Resistant *Klebsiella pneumoniae* (CRKP) strains were reviewed, and the CRKP sequence type (ST) was obtained through MLST typing.

**Results:** Over the past three years, 187 CRE strains were isolated in the hospital, with infections predominantly occurring in males and individuals over 60 years old. The departments with the highest detection rates were the ICU (35.29%), geriatric department (11.23%), and rehabilitation department (8.02%). The top three specimen sources were sputum (43.85%), clean midstream urine (32.09%), and secretions (13.90%). carbapenem-resistant *Klebsiella pneumoniae* accounted for the majority of CRE isolates (80.21%), followed by carbapenem-resistant Escherichia coli (14.44%). Carbapenem-resistant *Enterobacteriaceae* strains exhibited a high resistance rate to most commonly used antibiotics but were sensitive to tigecycline. Multilocus sequence typing results from 11 CRKP strains showed that 90.91% were of the ST11 type, while 9.09% were of the ST15 type.

**Conclusions:** From 2020 to 2022, CRE infections at the hospital primarily affected elderly male patients over 60, with CRKP as the predominant pathogen and STI1 as the main sequence type. Carbapenem-resistant *Enterobacteriaceae* strains were highly resistant to most commonly used antimicrobial drugs. Hospitals should prioritize the prevention and control of CRE infections, emphasizing the management of antimicrobial drugs, patient identification, environmental hygiene, and infection control measures to prevent outbreaks caused by CRE.

*Keywords:* Carbapenem-Resistant *Enterobacteriaceae*, Drug Resistance, Multi-site Sequence Typing, Hospital Infection Prevention and Control

#### 1. Background

Carbapenem antibiotics are currently the most widely used and potent  $\beta$ -lactam antibiotics, with the broadest antibacterial spectrum, often referred to as the last line of defense against Gram-negative bacteria. However, due to their frequent use in clinical practice,

many *Enterobacteriaceae* have developed resistance to them. Carbapenem-resistant *Enterobacteriaceae* (CRE) refers to a group of *Enterobacteriaceae* that are resistant to any carbapenem antibiotic or capable of producing carbapenemases (1). Carbapenem-resistant *Enterobacteriaceae* is classified at the highest priority level on the WHO's list of antibiotic-resistant bacteria

Copyright © 2024, Zhang et al. This open-access article is available under the Creative Commons Attribution 4.0 (CC BY 4.0) International License (https://creativecommons.org/licenses/by/4.0/), which allows for unrestricted use, distribution, and reproduction in any medium, provided that the original work is properly cited.

(2). Without effective intervention measures, CRE could become widespread in healthcare institutions within the next ten years, potentially developing into an endemic issue (3). Bloodstream infections caused by CRE are associated with alarmingly high mortality rates, ranging from 30% to 80% (4). The severe drug resistance posed by CRE may result in limited or no effective treatment options. Additionally, significant geographical differences exist in the epidemiological distribution and multilocus sequence typing (MLST) of CRE.

#### 2. Objectives

This study conducted an epidemiological analysis and MLST typing of carbapenem-resistant *Klebsiella pneumoniae* (CRKP) strains isolated from clinical samples in hospitals from 2020 to 2022. The aim was to understand the transmission and epidemiological characteristics of CRE, ultimately providing evidence for the prevention and control of CRE infections and improving clinical anti-infection treatment strategies.

## 3. Methods

## 3.1. Basic Information Collection of Carbapenem-Resistant Enterobacteriaceae Infected Individuals

Based on drug sensitivity results, 187 patients with CRE strains isolated from the First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine between 2020 and 2022 were screened (excluding patients with duplicate CRE strains) and included in the study. Relevant information on the infected individuals was collected. Inclusion criteria for CRE-infected patients excluded those with incomplete medical records and test results. Each patient with the same hospitalization number and CRE bacteria was enrolled only once. This study was approved by the hospital ethics committee.

#### 3.2. Main Instruments and Reagents

The VITEK 2 Compact fully automated microbial identification analyzer (BioMerieux, France), VITEK MS fully automated microbial mass spectrometry detection system (BioMerieux, France), nutrient agar plates (Dijing, Guangzhou), Simmons citrate agar (HB0115, Qingdao Haibo), inositol (Macklin), meropenem (Macklin), and bacterial DNA extraction kit (Genefist) were used. All reagents were used within their validity period.

3.3. Isolation, Cultivation, Identification, and Drug Sensitivity Experiments of Carbapenem-Resistant Enterobacteriaceae Strains

Microbiological samples submitted for clinical testing were inoculated onto nutrient agar plates and incubated at 36°C for 18 - 24 hours. Bacteria were identified using the VITEK MS fully automated microbial mass spectrometry detection system, followed by drug sensitivity testing with the VITEK 2 Compact fully automated microbial identification analyzer. The results were interpreted according to the standards published by the Clinical and Laboratory Standards Institute (CLSI) in the United States.

# 3.4. Review Identification and Bacterial DNA Extraction of Carbapenem-Resistant Klebsiella pneumoniae Strains

Resuscitate CRKP strains on a Simmons citrate inositol agar plate containing meropenem at a final concentration of 2  $\mu$ g/mL (5). Select the colonies growing on the plate and extract bacterial DNA according to the instructions provided with the reagent kit.

# 3.5. Multilocus Sequence Typing

Referring to the MLST website of the Pasteur Institute (https://bigsdb.pasteur.fr/), Beijing Tianyi Huiyuan Biotechnology Co., Ltd. was commissioned to synthesize the amplification products and sequencing primers for seven housekeeping genes: GapA, infB, mdh, pgi, phoE, rpoB, and tonB. The PCR amplification products were subjected to bidirectional sequencing, and the sequencing results were submitted to the MLST database (https://bigsdb.pasteur.fr/klebsiella/primers-used/). The sequences were aligned, the allele profile was obtained, and the sequence type (ST) of the tested strain was determined.

# 3.6. Data Analysis and Processing

The data were organized using WHONET 5.6 software, with quantitative variables presented as mean  $\pm$  standard deviation (SD) and categorical variables expressed as frequency (percentage). Statistical analysis was performed using SPSS version 23.0 for Windows (IBM, Armonk, New York).

# 4. Results

# 4.1. General Situation

lassification	No. (%)
sge (y)	
≤ 35	4 (2.14)
35 ~ 60	37 (19.79)
$60 \sim 80$	53 (28.34)
≥80	93 (49.73)
ender	
Male	143 (76.47)
Female	44 (23.53)
pecimen source	
Sputum	82 (43.85)
Clean midstream urine	60 (32.09)
Secretion	26 (13.90)
Blood	12 (6.42)
Serosal cavity effusion	3 (1.60)
Catheter tip	1(0.53)
Genital secretions	2 (1.07)
Cerebrospinal fluid	1(0.53)
epartment	
Intensive care unit	66 (35.29)
Geriatrics	21 (11.23)
Neurosurgery	10 (5.35)
Urology	10 (5.35)
Rehabilitation	15 (8.02)
Anorectal	11 (5.88)
Hepatobiliary Surgery	10 (5.35)
Neurology	13 (6.95)
Cardiology	6 (3.21)
Emergency	9 (4.81)
Orthopedics	2 (1.07)
Dermatology	2 (1.07)
Respiratory medicine	5 (2.67)
Otolaryngology	1(0.53)
Gastroenterology	1(0.53)
Nephrology	2 (1.07)
Traditional Miao Medicine and Pharmacy	1(0.53)
Oncology	1(0.53)
Obstetrics	1(0.53)

From 2020 to 2022, the hospital isolated 187 strains of CRE bacteria, including 78 strains in 2020, 58 in 2021, and 51 in 2022. The majority of infections occurred in males, comprising 76.47% (143/187) of the cases. Among

the infected individuals, patients over 60 years old accounted for 78.07% (146/187), with those over 80 years old making up 63.70% (93/146) of the total infections in this age group.

Bacterial Classification	2020	2021	2022	Subtotal
Escherichia coli	11 (14.10)	7 (12.07)	9 (17.65)	27 (14.44)
Klebsiella pneumoniae	64 (82.05)	48 (82.76)	38 (74.51)	150 (80.21)
Strange Proteobacteria	0 (0.00)	2 (3.45)	0(0.00)	2 (1.07)
Enterobacter cloacae	3 (3.85)	1(1.72)	3 (5.88)	7 (3.74)
K. oxytoca	0 (0.00)	0 (0.00)	1 (1.96)	1(0.53)
Subtotal	78 (100.00)	58 (100.00)	51 (100.00)	187 (100.00)

<sup>a</sup> Values are expressed as No. (%).

# 4.2. Distribution of Carbapenem-Resistant Enterobacteriaceae Departments and Specimen Sources

A total of 19 departments in the hospital detected CRE, with the ICU wards, geriatric departments, and rehabilitation departments showing higher detection rates, accounting for 35.29% (66/187), 11.23% (21/187), and 8.02% (15/187), respectively. In terms of specimen sources, sputum, clean midstream urine, and secretions were the primary sources, accounting for 43.85% (82/187), 32.09% (60/187), and 13.90% (26/187), respectively. Please refer to Table 1 for details.

## 4.3. Composition of Carbapenem-Resistant Enterobacteriaceae Bacteria

From 2020 to 2022, the primary CRE bacteria isolated in the hospital were carbapenem-resistant *K. pneumoniae* and *Escherichia coli*, accounting for 80.21% (150/187) and 14.44% (27/187), respectively, with an additional 3.74% (7/187) comprising other *E. coli* strains. Please refer to Table 2 for details.

## 4.4. Drug Resistance Rate Carbapenem-Resistant Enterobacteriaceae Bacterial Resistance Rate

The analysis shows that CRE exhibits high resistance to commonly used antibiotics, including β-lactams, quinolones, nitrofurans, aminoglycosides, and К. tetracyclines. However, carbapenem-resistant pneumoniae has a low resistance rate to tigecycline and cefotaxime/avibactam, at 0% and 4.08%, respectively. Carbapenem-resistant *E. coli* shows a resistance rate of 0% to tigecycline and 5.41% to amikacin. Additionally, carbapenem-resistant E. coli exhibits 0% resistance to both tigecycline and amikacin. The resistance rates of CRE bacteria to the corresponding antibiotics are presented in Table 3.

# 4.5. Multilocus Sequence Typing Results

Out of 23 preserved strains of CRKP bacteria in the laboratory, 11 strains survived. CRKP bacterial DNA was extracted, and seven housekeeping genes were amplified using the PCR method. After aligning the sequences, the ST types were identified. The results revealed that two sequence types were detected among the 11 strains, with ST11 being the predominant type, accounting for 90.91% (10/11), and ST15 accounting for 90.9% (1/11).

# 5. Discussion

Enterobacteriaceae are significant pathogens in both community-acquired and hospital-acquired infections, and carbapenem antibiotics are the most effective treatment for severe Enterobacteriaceae infections. However, the emergence of carbapenem-resistant Enterobacteriaceae poses substantial challenges to clinical treatment, with the growing concern of having no viable therapeutic options. Studies have shown that the incidence and infection rates of CRE in China are steadily rising and surpassing those in other countries. According to data from the 2020 National Antibiotic Resistance Monitoring Network (CHINET), the resistance rate of K. pneumoniae to carbapenems increased from 4.8% in 2014 to 10.5% in 2019 (6). CRE represents the most urgent and threatening drug-resistant bacteria in hospital settings, requiring stringent monitoring and control

This study revealed that the CRE infection rate at a traditional Chinese medicine hospital primarily affected elderly patients, with those aged  $\geq$  60 years accounting for 78.07% of the total infection rate. Among them, elderly patients aged  $\geq$  80 years comprised 63.70%, a finding consistent with research both domestically and internationally (7, 8). As age increases, immune function declines, and elderly patients often have multiple underlying diseases. Frequent antibiotic use in this population can disrupt gut microbiota, making them more susceptible to CRE infections (9).

Antibiotics	Klebsiella pneumoniae			Escherichia coli			Enterobacter cloacae		
	No. of Specimens (Pieces)	Drug Resistance No. (Copies)	Drug Resistance Rate (%)	No. of Specimens (Pieces)	Drug Resistance No. (Copies)	Drug Resistance Rate (%)	No. of Specimens (Pieces)	Drug Resistance No. (Copies)	Drug Resistance Rate (%)
Amikacin	150	137	91.33	27	1	3.70	7	0	0.00
Amoxicillin/clavulanic acid	119	118	99.16	23	22	95.65	-	-	-
Ampicillin/Sulbactam	85	85	100.00		-	-		-	-
Amtrazumab	88	87	98.86	-	-	-	-	-	-
Ertapenem	122	118	96.72	23	21	91.30	3	3	100.00
Furantoin	85	84	98.82	-		-	-		-
Compound Xinnuomin	150	94	62.67	27	13	48.15	7	4	57.14
Meropenem	50	50	100.00	6	6	100.00	-	-	-
Piperacillin/Tazobactam	150	147	98.00	27	25	92.60	7	7	100.00
Gentamicin	85	79	92.94	-	-	-	-	-	-
Tigecycline	36	0	0.00	24	0	0.00	7	0	0.00
Cefepime	150	147	98.00	27	27	100.00	7	4	57.14
Cefuroxime	119	118	99.16	23	23	100.00	7	7	100.00
Cefuroxime axetil	119	118	99.16	23	23	100.00	7	7	100.00
Cefoperazone/Sulbactam	128	125	97.66	27	23	85.19	7	7	100.00
Ceftriaxone	150	149	99.33	27	27	100.00	-	-	-
Ceftazidime	150	146	97.33	27	26	96.30	5	5	100.00
Ceftazidime/Avibactam	98	4	4.08	-	-	-	-	-	-
Cefotetan	85	84	98.82	-	-	-	-	-	-
Cefoxitin	119	118	99.16	23	22	95.65	-	-	-
Cefozolin	95	95	100.00	-	-	-	-	-	-
Tobramycin	88	82	93.18	-	-	-	-	-	-
Imipenem	150	145	96.67	27	24	88.89	7	4	57.14
Ciprofloxacin	88	85	96.59	-	-	-	-	-	-
Levofloxacin	150	145	96.67	27	24	88.89	5	3	60.00

<sup>a</sup> "-" indicates that the antimicrobial drug has not been used for drug sensitivity testing.

The department with the highest CRE isolation rate was the ICU, which aligns with trends observed in most hospitals (10), followed by geriatric and rehabilitation departments. ICU patients often have multiple comorbidities and severe conditions, requiring invasive medical procedures (e.g., tracheostomy, catheter placement) and the use of broad-spectrum antibiotics. Additionally, their weakened immune systems make them highly vulnerable to CRE colonization and transmission (11).

In the geriatric department, most patients are elderly with severe underlying conditions, often receiving prolonged antibiotic treatments and sometimes transferring to ICU wards. The rehabilitation department, a newly established unit specializing in traditional Chinese medicine and functional recovery, also saw a high rate of CRE detection. Many patients in this department are bedridden with pressure ulcers and require invasive procedures such as urethral catheterization, increasing the likelihood of CRE isolation. The relatively high CRE detection rate in non-ICU wards highlights the need to strengthen infection prevention and control measures beyond ICU settings.

Regarding specimen sources, the majority of CRE strains were isolated from sputum and clean midstream urine samples, accounting for 43.85% and 32.09% of cases, respectively. This may be due to the fact that CRE infections predominantly affect open systems, such as the respiratory and urinary tracts, which are more prone to infection when the body's resistance is compromised (12).

The prevalence of CRE isolates from sputum and clean midstream urine specimens in this study may be due to the ease of collecting these specimens and the large volume of samples submitted for testing. This contrasts slightly with other hospitals (13), where CRE isolates are mainly sourced from sputum, followed by purulent secretions and drainage fluids. The isolation of CRE strains from sterile sites holds greater clinical significance. However, the hospital in this study had fewer strains isolated from sterile sites (such as blood, serous fluid, or venous catheters), which warrants closer attention. Further analysis of CRE composition revealed that the predominant CRE bacteria in the hospital were *K. pneumoniae, E. coli*, and *Enterobacter cloacae*, which are resistant to carbapenem antibiotics. This aligns with both domestic and international trends (14, 15).

The drug sensitivity results indicated that CRE is resistant to most antibiotics, with susceptibility to only a few, such as tigecycline. This highlights the severe resistance situation of CRE strains, posing substantial limitations on clinical treatment options. Clinical treatment should be guided by drug sensitivity results, with rational selection and combination of antibiotics for the management of CRE infections (16, 17).

The MLST plays a pivotal role in CRE research by analyzing multiple conserved gene fragments to classify strains and identify their evolutionary relationships and transmission patterns. Multilocus sequence typing relies on nucleotide sequencing of housekeeping genes to accurately detect genetic variations in bacteria. This method enables the study of genetic evolution across regions and aids in epidemiological investigations. By comparing laboratory data from different areas, MLST helps determine whether pathogens are related, if they originate from specific clones, and if outbreaks have occurred in hospitals. It can also track CRE transmission routes, identify the source of infection, and, when combined with whole-genome sequencing, conduct indepth studies on drug resistance mechanisms. The global MLST database supports the monitoring of CRE strains at both local and international levels, assisting public health departments in formulating infection control policies to prevent the spread of CRE.

Carbapenem-resistant *Klebsiella pneumoniae* is a priority pathogen identified by the WHO as a severe threat to human health, primarily linked to hospital-acquired infections (18, 19). In this study, CRKP accounted for 80.21% of the CRE isolates, with MLST revealing two sequence types, predominantly STI1 (90.91%). This finding is consistent with the predominant genotype and clone type of CRKP in China (20, 21). In China, STI1 CRKP is the leading strain (22), known as the most widespread multidrug-resistant lineage in Asia (23), and was first identified as a hypervirulent strain in China (24).

The detection of the ST15 type is less frequent compared to the ST11 type, but it remains one of the

more commonly found strains in China (25). Research has shown (26) that the ST15 CRKP is a high-risk clonal strain that has emerged recently and often leads to hospital outbreaks. ST15, which carries plasmids containing both virulence and resistance genes, has been identified. The most widely prevalent CRKP multilocus sequence types are ST258 and ST11 (22). In the United States and Europe, ST258 CRKP is the dominant type, often associated with localized infections and high mortality rates (27-29). In contrast, ST11-type CRKP is most common in Asia, especially in China (27, 30). The limited diversity of ST types observed in this study could be due to the small number of strains analyzed through MLST typing.

## 5.1. Conclusions

The continuous emergence of CRE presents significant challenges to clinical treatment and poses a severe threat to patient outcomes. Preventing the occurrence or outbreak of CRE infections in hospitals is crucial. Managing hospital-acquired CRE infections is a complex and systematic endeavor; any oversight or insufficient isolation measures can lead to the spread or even outbreaks of these infections. Effective strategies for preventing and controlling CRE include antibacterial drug management, patient identification and management. Implementing these measures in a comprehensive and detailed manner can help reduce the incidence of CRE infections and further decrease mortality rates (31).

Analyzing the epidemiological characteristics and MLST subtypes of hospital CRE-infected patients can effectively aid clinical personnel in understanding the transmission patterns of CRE. This knowledge encourages the strict adherence to antibiotic usage guidelines, including the rational use of carbapenems, and ensures better patient safety.

# Footnotes

Authors' Contribution: S. W. W.: Major contributions to the idea and study design; Y. Y.: Data analysis; M. J. Z.: Writing and revising the paper; Y. Q. Z., M. Y., and Y. M. H.: Data collection and follow-up. **Conflict of Interests Statement:** There is no conflict of interest to he declared. **Data Availability:** The dataset presented in the study is available on request from the corresponding author submission during or after publication.

**Ethical Approval:** This research ethics was approved by the Science and Technology Fund of the Guizhou Provincial Health Commission with project number gzwkj2022-194.

**Funding/Support:** The paper was supported by the following funding projects: (1) science and Technology Fund of Guizhou Provincial Health Commission, Project Number: gzwkj2022-194. (2) the Doctoral Initiation Fund of the First Affiliated Hospital of Guizhou University of Traditional Chinese Medicine, Project Number: GYZYFY-BS-2020 (03).

#### References

- Centers for Disease Control Prevention. Guidance for control of infections with carbapenem-resistant or carbapenemase-producing Enterobacteriaceae in acute care facilities. *MMWR Morb Mortal Wkly Rep.* 2009;**58**(10):256-60. [PubMed ID: 19300408].
- 2. World Health Organization. WHO Publishes list of bacteria for which new antibiotics are urgently needed. 2024. Available from: https://www.who.int/news/item/27-02-2017-who-publishes-list-ofbacteria-for-which-new-antibiotics-are-urgently-needed.
- 3. Lee BY, Bartsch SM, Wong KF, McKinnell JA, Slayton RB, Miller LG, et al. The potential trajectory of carbapenem-Resistant Enterobacteriaceae, an emerging threat to health-care facilities, and the impact of the centers for disease control and prevention toolkit. *Am J Epidemiol.* 2016;**183**(5):471-9. [PubMed ID: 26861238]. [PubMed Central ID: PMC4772438]. https://doi.org/10.1093/aje/kwv299.
- Hu Q, Chen J, Sun S, Deng S. Mortality-related risk factors and novel antimicrobial regimens for carbapenem-resistant enterobacteriaceae infections: A systematic review. *Infect Drug Resist.* 2022;15:6907-26. [PubMed ID: 36465807]. [PubMed Central ID: PMC9717588]. https://doi.org/10.2147/IDR.S390635.
- Wei L, Wu L, Wen H, Feng Y, Zhu S, Liu Y, et al. Spread of carbapenemresistant Klebsiella pneumoniae in an intensive care unit: A wholegenome sequence-based prospective observational study. *Microbiol Spectr.* 2021;9(1). e0005821. [PubMed ID: 34259540]. [PubMed Central ID: PMC8552774]. https://doi.org/10.1128/Spectrum.00058-21.
- 6. China Antimicrobial Resistance Surveillance System. [Antimicrobial resistance of clinically isolated bacteria from elderly patients: Surveillance report from China antimicrobial resistance surveillance system in 2014 2019]. *Chinese J Infect Control*. 2021;(2):112-23. ZH.
- Saeed NK, Alkhawaja S, Azam N, Alaradi K, Al-Biltagi M. Epidemiology of carbapenem-resistant Enterobacteriaceae in a tertiary care center in the Kingdom of Bahrain. *J Lab Physicians*. 2019;**11**(2):111-7. [PubMed ID: 31160848]. [PubMed Central ID: PMC6543944]. https://doi.org/10.4103/[LP.]LP\_101\_18.
- 8. Yang H, Tao J, Wei T, Wang J, Mao J, Wang H. [Analysis of carbapenem resistant Enterobacteriaceae bacterial resistance in hospitalized patients of a combined traditional Chinese and Western medicine hospital in Shanghai from 2015 to 2019]. *Laboratory Med.* 2021;**36**(1). ZH.
- Li Y, Sun QL, Shen Y, Zhang Y, Yang JW, Shu LB, et al. Rapid increase in prevalence of carbapenem-resistant Enterobacteriaceae (CRE) and emergence of colistin resistance gene mcr-1 in CRE in a hospital in Henan, China. J Clin Microbiol. 2018;56(4). [PubMed ID: 29386265]. [PubMed Central ID: PMC5869811]. https://doi.org/10.1128/JCM.01932-17.
- 10. Liu J, Yu J, Chen F, Yu J, Simner P, Tamma P, et al. Emergence and establishment of KPC-2-producing ST11 Klebsiella pneumoniae in a

general hospital in Shanghai, China. Eur J Clin Microbiol Infect Dis. 2017;**37**(2):293-9. https://doi.org/10.1007/s10096-017-3131-4.

- Zhang T, Liu Z, Xu T; et al. [Analysis of antibiotic resistance and disinfectant resistance genes in carbapenem resistant Enterobacteriaceae bacteria from a tertiary hospital]. *Chin J Hospital Infect.* 2023;33(3). ZH.
- 12. Hu Z, Pan K, Pan X, Zhu J, Zhou J, Tang J. [Bacterial resistance monitoring in a tertiary hospital in Anhui Province in 2017]. *Chin J Antibiotics*. 2019. ZH.
- 13. Lan M, Zhao Z, Kang Y, Wang Y, Li J, Liu C, et al. [Molecular epidemiological characteristics and drug resistance of carbapenem resistant Enterobacterales]. *Chin J Infect Control*. 2022;(11):1053-9. ZH.
- Ahn JY, Ahn SM, Kim JH, Jeong SJ, Ku NS, Choi JY, et al. Clinical characteristics and associated factors for mortality in patients with carbapenem-resistant enterobacteriaceae bloodstream infection. *Microorganisms*. 2023;11(5). [PubMed ID: 37317095]. [PubMed Central ID: PMC10220897]. https://doi.org/10.3390/microorganisms11051121.
- Miao X, Meng X. [Research advances in active screening as well as prevention and control strategies on CRE from patients and hospital environment]. *Chin J Infect Control*. 2022;21(12):1257-63. ZH.
- Carrara E, Bragantini D, Tacconelli E. Combination versus monotherapy for the treatment of infections due to carbapenemresistant Enterobacteriaceae. *Curr Opin Infect Dis.* 2018;31(6):594-9. [PubMed ID: 30299357]. https://doi.org/10.1097/QCO.000000000000495.
- Trecarichi EM, Tumbarello M. Therapeutic options for carbapenemresistant Enterobacteriaceae infections. *Virulence*. 2017;8(4):470-84. [PubMed ID: 28276996]. [PubMed Central ID: PMC5477725]. https://doi.org/10.1080/21505594.2017.1292196.
- 18. Tacconelli E. Global priority list of antibiotic-resistant bacteria to guide research, discovery, and development. *infect control africa net*. 2024;(7-1).
- Navon-Venezia S, Kondratyeva K, Carattoli A. Klebsiella pneumoniae: A major worldwide source and shuttle for antibiotic resistance. *FEMS*  Microbiol Rev. 2017;41(3):252-75. [PubMed ID: 28521338]. https://doi.org/10.1093/femsre/fux013.
- Zhang Y, Wang Q, Yin Y, Chen H, Jin L, Gu B, et al. Epidemiology of carbapenem-resistant Enterobacteriaceae infections: Report from the china CRE network. *Antimicrob Agents Chemother*. 2018;62(2). [PubMed ID: 29203488]. [PubMed Central ID: PMC5786810]. https://doi.org/10.1128/AAC.01882-17.
- 21. Hu X, Yuan G, Wu Y, Liu W, Yang B, Han J, et al. [Molecular epidemiological characteristics and antimicrobial resistance of carbapenem-resistant Klebsiella pneumoniae in three general hospital in southwest China]. *Chin J Infect Control*. 2022;**21**(2):121-7. ZH.
- Liao W, Liu Y, Zhang W. Virulence evolution, molecular mechanisms of resistance and prevalence of ST11 carbapenem-resistant Klebsiella pneumoniae in China: A review over the last 10 years. J Glob Antimicrob Resist. 2020;23:174-80. [PubMed ID: 32971292]. https://doi.org/10.1016/j.jgar.2020.09.004.
- Xu L, Sun X, Ma X. Systematic review and meta-analysis of mortality of patients infected with carbapenem-resistant Klebsiella pneumoniae. Ann Clin Microbiol Antimicrob. 2017;16(1):18. [PubMed ID: 28356109]. [PubMed Central ID: PMC5371217]. https://doi.org/10.1186/s12941-017-0191-3.
- Yang X, Sun Q, Li J, Jiang Y, Li Y, Lin J, et al. Molecular epidemiology of carbapenem-resistant hypervirulent Klebsiella pneumoniae in China. *Emerg Microbes Infect*. 2022;11(1):841-9. [PubMed ID: 35236251].
  [PubMed Central ID: PMC8942559]. https://doi.org/10.1080/22221751.2022.2049458.
- 25. ChunMei Z, ShengLei H, JinNan C, Yan M, YuZhang S, JiaJin S, et al. [Drug resistance genes and molecular epidemiological

characteristics of CRKP strains isolated from ICU patients with bloodstream infection]. *Chin J Hospital Infect*. 2022;**32**(5):659-63. ZH.

- Zhao H, He Z, Li Y, Sun B. [Epidemiology of carbapenem-resistant Klebsiella pneumoniae ST15 of producing KPC-2, SHV-106 and CTX-M-15 in Anhui, China]. *BMC Microbiol*. 2023;**22**(1):262. ZH. [PubMed ID: 36319965]. [PubMed Central ID: PMC9624029]. https://doi.org/10.1186/s12866-022-02672-1.
- 27. Liao W, Liu Y, Zhang W. Virulence evolution, molecular mechanisms of resistance and prevalence of ST11 carbapenem-resistant Klebsiella pneumoniae in China: A review over the last 10 years. *J Glob Antimicrob Resist.* 2020;**23**:174-80. [PubMed ID: 32971292]. https://doi.org/10.1016/j.jgar.2020.09.004.
- Dautzenberg MJ, Haverkate MR, Bonten MJ, Bootsma MC. Epidemic potential of Escherichia coli ST131 and Klebsiella pneumoniae ST258: A systematic review and meta-analysis. *BMJ Open.* 2016;6(3). e009971. [PubMed ID: 26988349]. [PubMed Central ID: PMC4800154]. https://doi.org/10.1136/bmjopen-2015-009971.
- 29. DeLeo FR, Chen L, Porcella SF, Martens CA, Kobayashi SD, Porter AR, et al. Molecular dissection of the evolution of carbapenem-resistant multilocus sequence type 258 Klebsiella pneumoniae. *Proceedings National Academy Sci.* 2014;**111**(13):4988-93. https://doi.org/10.1073/pnas.1321364111.
- Munoz-Price LS, Poirel L, Bonomo RA, Schwaber MJ, Daikos GL, Cormican M, et al. Clinical epidemiology of the global expansion of Klebsiella pneumoniae carbapenemases. *Lancet Infect Dis.* 2013;13(9):785-96. https://doi.org/10.1016/s1473-3099(13)70190-7.
- 31. Wei T, Zou C, Qin J, Tao J, Yan L, Wang J, et al. Emergence of hypervirulent ST11-K64 Klebsiella pneumoniae poses a serious clinical threat in older patients. *Front Public Health*. 2022;**10**. https://doi.org/10.3389/fpubh.2022.765624.
- Qiao F, Chen Y, Zong Z. [Interpretation of the prevention and control standards for carbapenem resistant Enterobacteriaceae west China medicine]. West China Medicine. 2024;39(3):362-6. ZH. https://doi.org/10.7507/1002-0179.202402116.