THE EMERGENCE OF *bla*_{OXA-48} AND *bla*_{NDM} AMONG ESBL-PRODUCING *KLEBSIELLA PNEUMONIAE* IN CLINICAL ISOLATES OF A TERTIARY HOSPITAL IN IRAN

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The aim of this study was to investigate the prevalence of carbapenem-resistant Klebsiella pneumoniae (CRKP) and the most common types of carbapenemases, metallo-beta-lactamases (MBLs), and extended-spectrum beta-lactamases (ESBLs) among CRKP isolates in a tertiary hospital in Isfahan, Iran. Eighty non-repetitive clinical isolates of K. pneumoniae were obtained from different clinical specimens. Antibiotic resistance pattern of isolates was determined by disk diffusion method and production of carbapenemases and MBLs was confirmed using modified Hodge test and E-test, respectively. Molecular detection of the antibiotic resistance genes was performed using PCR. Fifty-one (63.8%) isolates have decreased susceptibility to carbapenems, of which 46 (90.2%) isolates were as carbapenemase producer and four (7.8%) isolates were positive for MBLs, phenotypically. The results of PCR showed that the prevalence of *bla*OXA-48, *bla*NDM, *bla*SHV, *bla*CTX-M, and *bla*TEM genes among CRKP isolates were 90.2%, 15.7%, 98%, 96.1%, and 90.2%, respectively. No isolates carrying the $bla_{\rm KPC}$, $bla_{\rm GES}$, $bla_{\rm IMI}$, $bla_{\rm VIM}$, and $bla_{\rm IMP}$ genes were detected. This study showed that the production of OXA-48 is one of the main mechanisms of resistance to carbapenems in CRKP isolates in Isfahan. In addition, the dissemination of NDM-producing CRKP isolates is a potential risk for the health care system of this area in the near future.

Keywords: Klebsiella pneumoniae, CRKP, OXA-48, NDM, ESBLs

Introduction

Klebsiella pneumoniae is an important opportunistic bacterial pathogen associated with both community- and hospital-acquired infections and potentially

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causing mortality, especially in hospitalized patients [1]. An important part of hospital infections (4%-8%) is due to this bacterium. K. pneumoniae isolates are increasingly resistant to several antibiotics. Hospital outbreaks caused by K. pneumoniae isolates are frequent and the transmission of resistant strains between the hospital and community has been described previously [2]. Treatment of infections caused by drug-resistant K. pneumoniae isolates, especially strains producing extended-spectrum beta-lactamases (ESBLs), as well as isolates with multidrug-resistant and extensively drug-resistant phenotypes is very complicated and costly, and usually not associated with satisfactory results [3, 4]. Carbapenems are a group of beta-lactam antibiotics that have been used as the "last-line" treatment for Enterobacteriaceae-associated infections, especially those producing ESBLs [5, 6]. Compared to cephalosporins, penicillins, or beta-lactam/ beta-lactamase inhibitors, carbapenems have a broader antimicrobial spectrum [7]. During the past decade, the emergence of resistance to carbapenems in glucose non-fermenting (e.g., Pseudomonas aeruginosa and Acinetobacter baumannii) and glucose-fermenting (e.g., Enterobacteriaceae) Gram-negative bacilli has been reported from all around the world [8, 9]. Recently, the spread of carbapenemresistant Enterobacteriaceae, especially carbapenem-resistant K. pneumoniae (CRKP), is the most important clinical problem in the field of antibiotic resistance that should be monitored seriously and prevented from its progress [10]. This study aims to investigate the prevalence of CRKP and the most common types of carbapenemases, metallo-beta-lactamases (MBLs), and ESBLs among CRKP in a tertiary hospital in Isfahan, Iran.

Materials and Methods

Bacterial isolates

During a 9-month period, April–December 2017, 80 non-repetitive *K. pneumoniae* isolates were obtained from various specimens of inpatients in Alzahra hospital, Isfahan, Iran. All isolates were identified in microbiology laboratory of medical school using the conventional methods, such as Gram staining and biochemical (oxidase, sugar fermentation, IMViC, Kliger's iron agar, nitrate reduction, motility, etc.) tests [11]. In addition, to confirm the species, a PCR detection based on the 16S–23S internal transcribed spacer sequence of *K. pneumoniae* was conducted [12]. This study was evaluated and approved by the Ethics Committee of Isfahan University of Medical Sciences (project no. 395891).

Antibiotic susceptibility testing

Susceptibility to antibiotics was determined using Kirby–Bauer's disk diffusion method according to the recommendations of Clinical Laboratory Standard Institute (CLSI) [13]. For this purpose, 18 antibiotics (MAST, UK and Liofilchem, Italy) including imipenem (10 μ g), meropenem (10 μ g), ertapenem (10 μ g), gentamicin (10 μ g), ceftaroline (30 μ g), piperacillin/tazobactam (100/10 μ g), cefazolin (30 μ g), cefuroxime (30 μ g), ceftazidime (30 μ g), cefepime (30 μ g), cefoxitin (30 μ g), ciprofloxacin (5 μ g), trimethoprim/sulfamethoxazole (1.25/23.75 μ g), tigecycline (15 μ g), aztreonam (30 μ g), amoxicillin/clavulanic acid (20/10 μ g), chloramphenicol (30 μ g), and tetracycline (30 μ g) were used. *Escherichia coli* ATCC 25922 was used for quality control of disk diffusion method [13].

Carbapenemase and MBL screening assays

To determine carbapenemase activity, the modified Hodge test (MHT) using ertapenem disk (MAST) was performed according to CLSI guidelines [13]. Moreover, to detect MBL activity, E-test method using strips containing meropenem/meropenem + EDTA (Liofilchem) was carried out based on manufacturer's guidelines.

PCR for detection of antibiotic resistance genes

DNA was extracted using the boiling method and used as template for PCR [14]. To detect antibiotic resistance genes including bla_{KPC} , bla_{GES} , bla_{IMI} , bla_{VIM} , bla_{IMP} , bla_{NDM} , $bla_{\text{OXA-48}}$, bla_{SHV} , bla_{TEM} , and $bla_{\text{CTX-M}}$, separate PCR reactions were performed. The list of all target genes and corresponding primers is presented in Table I [15–21]. PCR was performed using commercially available PCR Master Mix (AMPLIQON, Denmark) according to the manufacturer's instructions. Briefly, 1 µl template DNA (~100 ng/µl), 1 µl of each primer (10 pmoles/µl), and 9.5 µl DNase-free distilled water were added to 12.5 µl of Master Mix in a final volume of 25 µl. Thermocycling was carried out with an initial denaturation step at 95 °C for 5 min, followed by 35 cycles of denaturation at 95 °C for 30 s, annealing for 45 s at primer-specific temperatures (Table I), extension at 72 °C for 1 min, and a final extension step at 72 °C for 10 min. PCR products were resolved by standard electrophoresis on 1.5% agarose gel containing DNA safe stain.

Target gene	Primer sequence (5'-3')	Annealing temperature (°C)	Amplicon size (bp)	References
bla _{KPC}	F: GATACCACGTTCCGTCTGG	58	246	[15]
	R: GCAGGTTCCGGTTTTGTCTC			
bla_{GES}	F: GTTTTGCAATGTGCTCAACG	53	371	[16]
	R: TGCCATAGCAATAGGCGTAG			
bla_{IMI}	F: ATGTCATTAGGTGATATGGC	50	879	[17]
	R: GCATAATCATTTGCCGTACC			
$bla_{\rm VIM}$	F: TTTGGTCGCATATCGCAACG	66	500	[18]
	R: CCATTCAGCCAGATCGGCAT			
bla _{IMP}	F: GTTTATGTTCATACATCG	45	440	[18]
	R: GGTTTAACAAAAAAAACAACCAC			
bla _{NDM}	F: GGGCAGTCGCTTCCAACGGT	52	475	[19]
	R: GTAGTGCTCAGTGTCGGCAT			
bla _{OXA-48}	F: GCGTGGTTAAGGATGAACAC	60	438	[20]
	R: CATCAAGTTCAACCCAACCG			
$bla_{\rm SHV}$	F: ATGCGTTATATTCGCCTGTG	60	753	[18]
	R: TGCTTTGTTATTCGGGCCAA			
bla_{TEM}	F: AAACGCTGGTGAAAGTA	45	752	[18]
	R: AGCGATCTGTCTAT			
bla _{CTX-M}	F: TTTGCGATGTGCAGTACCAGTAA	51	544	[21]
	R: CGATATCGTTGGTGGTGCCATA			

Table I. List of primers, expected amplicon size, and annealing temperatures

Results

Bacterial isolates

Table II shows demographic information and clinical characteristics of patients infected with *K. pneumoniae*. The mean age of patients was 52.7 years.

Antibiotic susceptibility testing

Table III shows the results of antibiotic susceptibility testing in considered isolates. The highest resistance rate was observed to cefazolin (87.5%), followed by amoxicillin/clavulanic acid (78.8%), cefuroxime (77.5%), tetracycline (77.5%), and aztreonam (76.2%). Tigecycline and chloramphenicol were the most effective antibiotics with 13.7% and 18.7% resistance rate, respectively. About 62.5% of isolates were resistant to meropenem and ertapenem as well as overall, 51 isolates showed the decreased susceptibility to carbapenems.

Demographics	Number (%)	
Age range (years)	1–92	
Sex		
Male	45 (56.3)	
Female	35 (43.7)	
Hospital wards		
Intensive care unit	48 (60.0)	
Internal medicine	12 (15.0)	
Surgery	8 (10.0)	
Emergency	7 (8.8)	
Coronary care unit	3 (3.7)	
Hematology/oncology	2 (2.5)	
Specimens		
Urine	29 (36.3)	
Trachea	21 (26.3)	
Blood	8 (10.0)	
Cerebrospinal fluid	7 (8.8)	
Wound	6 (7.5)	
Sputum	3 (3.7)	
Abscess	2 (2.5)	
Catheter	2 (2.5)	
Bronchoalveolar lavage	1 (1.2)	
Drainage fluid	1 (1.2)	

 Table II. Demographic information and clinical characteristics of patients infected with K. pneumoniae (n = 80)

Table III. Antibiotic susceptibility of *K. pneumoniae* isolates (n = 80)

Antibiotic	Susceptible [n (%)]	Intermediate [n (%)]	Resistant [n (%)]
Imipenem	31 (38.8)	3 (3.7)	46 (57.5)
Meropenem	30 (37.5)	0 (0.0)	50 (62.5)
Ertapenem	30 (37.5)	0 (0.0)	50 (62.5)
Gentamicin	31 (38.8)	1 (1.2)	48 (60.0)
Ceftaroline	20 (25.0)	1 (1.2)	59 (73.8)
Piperacillin/tazobactam	24 (30.0)	4 (5.0)	52 (65.0)
Cefazolin	7 (8.8)	3 (3.7)	70 (87.5)
Cefuroxime	16 (20.0)	2 (2.5)	62 (77.5)
Ceftazidime	17 (21.3)	3 (3.7)	60 (75.0)
Cefepime	17 (21.3)	4 (5.0)	59 (73.7)
Cefoxitin	24 (30.0)	2 (2.5)	54 (67.5)
Ciprofloxacin	19 (23.8)	1 (1.2)	60 (75.0)
Trimethoprim/sulfamethoxazole	37 (46.3)	1 (1.2)	42 (52.5)
Tigecycline	57 (71.3)	12 (15.0)	11 (13.7)
Aztreonam	17 (21.3)	2 (2.5)	61 (76.2)
Amoxicillin/clavulanic acid	13 (16.2)	4 (5.0)	63 (78.8)
Chloramphenicol	36 (45.0)	29 (36.3)	15 (18.7)
Tetracycline	17 (21.3)	1 (1.2)	62 (77.5)

Carbapenemase and MBL screening assays

The MHT was performed on 51 isolates with decreased susceptibility to carbapenems and following results were obtained: 46 (90.2%) isolates were positive, 4 (7.8%) isolates were negative, and 1 (2%) isolate was indeterminate. Similarly, phenotypic MBL production testing on desired isolates identified 4 (7.8%) isolates as positive, 4 (7.8%) isolates as negative, and 43 (84.4%) isolates with non-determinable results.

PCR for detection of antibiotic resistance genes

All 51 isolates with decreased susceptibility to carbapenems were tested in PCR for the presence of the selected antibiotic resistance genes. Forty-six (90.2%) isolates harbored bla_{OXA-48} and eight (15.7%) isolates were positive for bla_{NDM} . Moreover, seven (13.7%) isolates had both bla_{OXA-48} and bla_{NDM} , simultaneous-ly. ESBLs-encoding genes including bla_{SHV} , bla_{CTX-M} , and bla_{TEM} were identified in 98%, 96.1%, and 90.2% isolates, respectively. Similarly, these (ESBLs) genes were detected concurrently in 45 (88.2%) isolates. In addition, six (11.8%) isolates were positive for the presence of bla_{OXA-48} , bla_{NDM} , bla_{SHV} , bla_{CTX-M} , and bla_{TEM} genes, together. The results of PCR were negative for bla_{KPC} , bla_{GES} , bla_{IMI} , bla_{VIM} , and bla_{IMP} genes.

Discussion

One of the main problems in dealing with clinical strains of *K. pneumoniae* is the increasing incidence of resistance to antibiotics. These resistant strains are rapidly expanding and have caused many problems with the treatment of infections in various health care settings [22]. Carbapenems are a group of beta-lactam antibiotics that are used to treat serious infections in hospitals. The constant uses of these antibiotics and the selective pressure resulting from it have led to resistance to these antibiotics in Gram-negative bacteria, especially *K. pneumoniae* [5]. Usually, infections caused by CRKP threaten patients in hospitals, nursing homes, and other health care centers and do not occur in immunocompetent persons [23, 24]. In this study, more than half of patients were over 56.5 years of age. In addition, almost half of CRKP isolates (45%) were obtained from hospitalized patients in intensive care unit, which highlights the importance of aging and long-term hospitalization in these infections. PCR results showed that 90.2% of CRKP isolates were positive for bla_{OXA-48} . OXA-48-producing *Enterobacteriaceae* was first identified in Turkey in 2001, subsequently reported from several countries

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in the Middle East, North Africa, and Europe [25, 26]. The first report of OXA-48producing K. pneumoniae strains in Iran was presented by Azimi et al. [27]. In two separate studies conducted in Tehran hospitals (capital of Iran), the prevalence of bla_{OXA-48} among CRKP isolates was reported as 96.4% [27] and 4.1% [28]. Another study in southwestern Iran showed that 6.9% of CRKP isolates were positive for bla_{OXA-48} [29]. The results of this study regarding the prevalence of bla_{OXA-48} -harboring CRKP is in parallel with the study by Solgi et al. [30] between 2014 and 2016 in our region. In their study, all (100%) CRKP isolates harbored bla_{OXA-48} . The results of the phenotypic test for detection of MBLs showed that only four (7.8%) CRKP isolates produced MBLs, while in the PCR assay, eight (15.7%) CRKP isolates were identified as *bla*_{NDM}-positive. This finding suggests that the phenotypic test alone is not sufficient for the definitive detection of MBLs, and a molecular test such as PCR should be used along with it. ESBLs-encoding genes including bla_{SHV} , bla_{CTX-M} , and bla_{TEM} with 98%, 96.1%, and 90.2% were the most prevalent bla genes among CRKP isolates, respectively. In a meta-analysis article that we previously published in Iran, the overall relative frequency (RF) of bla_{CTX-M} gene among ESBLs-producing K. pneumoniae clinical isolates was 56.7% [31]. Our results show that the RF of bla_{CTX-M} among CRKP isolates in Isfahan is higher than its RF in Iran. Solgi et al. [30] in their study from Isfahan have reported that 95.8%, 94.8%, and 96.9% of CRKP isolates were positive for *bla*_{CTX-M}, *bla*_{TEM}, and *bla*_{SHV}, respectively. In this case, our results were fully consistent with their results. Similar to other studies that were carried out previously in Isfahan, we have not detected class A carbapenemases (KPC, GES, and IMI) among CRKP isolates [30, 32, 33]. In conclusion, this study showed that the production of class D carbapenemases (OXA-48) is one of the main mechanisms of resistance to carbapenems in CRKP isolates in Isfahan. Moreover, the dissemination of NDM-producing CRKP isolates is a potential hazard for the health care system of this area in the future that should be controlled.

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Conflict of Interest

The authors declare no conflict of interest.

References

- Kohlenberg, A., Schwab, F., Ruden, H.: Wide dissemination of extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli* and *Klebsiella* spp. in acute care and rehabilitation hospitals. Epidemiol Infect 140, 528–534 (2012).
- Monnet, D. L., Biddle, J. W., Edwards, J. R., Culver, D. H., Tolson, J. S., Martone, W. J., Tenover, F. C., Gaynes, R. P.: Evidence of interhospital transmission of extended-spectrum beta-lactam-resistant *Klebsiella pneumoniae* in the United States, 1986 to 1993. The National Nosocomial Infections Surveillance System. Infect Control Hosp Epidemiol 18, 492–498 (1997).
- Ben-David, D., Kordevani, R., Keller, N., Tal, I., Marzel, A., Gal-Mor, O., Maor, Y., Rahav, G.: Outcome of carbapenem resistant *Klebsiella pneumoniae* bloodstream infections. Clin Microbiol Infect 18, 54–60 (2012).
- Wang, Q., Li, B., Tsang, A. K., Yi, Y., Woo, P. C., Liu, C. H.: Genotypic analysis of *Klebsiella pneumoniae* isolates in a Beijing Hospital reveals high genetic diversity and clonal population structure of drug-resistant isolates. PLoS One 8, e57091 (2013).
- 5. Patel, G., Bonomo, R. A.: Status report on carbapenemases: Challenges and prospects. Expert Rev Anti Infect Ther **9**, 555–570 (2011).
- Morrill, H. J., Pogue, J. M., Kaye, K. S., LaPlante, K. L.: Treatment options for carbapenem-resistant *Enterobacteriaceae* infections. Open Forum Infect Dis 2, ofv050 (2015).
- Bedenic, B., Plecko, V., Sardelic, S., Uzunovic, S., Godic Torkar, K.: Carbapenemases in gram-negative bacteria: Laboratory detection and clinical significance. Biomed Res Int 2014, 841951 (2014).
- Poirel, L., Potron, A., Nordmann, P.: OXA-48-like carbapenemases: The phantom menace. J Antimicrob Chemother 67, 1597–1606 (2012).
- Rezaei, A., Fazeli, H., Moghadampour, M., Halaji, M., Faghri, J.: Determination of antibiotic resistance pattern and prevalence of OXA-type carbapenemases among *Acinetobacter baumannii* clinical isolates from inpatients in Isfahan, Central Iran. Infez Med 26, 61–66 (2018).
- Bialvaei, A. Z., Kafil, H. S., Asgharzadeh, M., Yousef Memar, M., Yousefi, M.: Current methods for the identification of carbapenemases. J Chemother 28, 1–19 (2016).
- Podschun, R., Ullmann, U.: *Klebsiella* spp. as nosocomial pathogens: Epidemiology, taxonomy, typing methods, and pathogenicity factors. Clin Microbiol Rev 11, 589–603 (1998).
- Liu Y, Liu C, Zheng W, Zhang X, Yu J, Gao Q, Hou Y, Huang X: PCR detection of *Klebsiella pneumoniae* in infant formula based on 16S-23S internal transcribed spacer. Int J Food Microbiol **125**, 230–235 (2008).
- Clinical and Laboratory Standards Institute (CLSI): Performance Standards for Antimicrobial Susceptibility Testing (27th ed.). CLSI supplement M100. CLSI, Wayne, 2017.
- 14. Amiri, A., Firoozeh, F., Moniri, R., Zibaei, M.: Prevalence of CTX-M-Type and PER extended-spectrum beta-lactamases among *Klebsiella* spp. isolated from clinical specimens in the teaching hospital of Kashan, Iran. Iran Red Crescent Med J 18, e22260 (2016).
- Hindiyeh, M., Smollen, G., Grossman, Z., Ram, D., Davidson, Y., Mileguir, F., Vax, M., Ben David, D., Tal, I., Rahav, G., Shamiss, A., Mendelson, E., Keller, N.: Rapid detection of *bla*_{KPC} carbapenemase genes by real-time PCR. J Clin Microbiol **46**, 2879–2883 (2008).

- Queenan, A. M., Bush, K.: Carbapenemases: The versatile beta-lactamases. Clin Microbiol Rev 20, 440–458 (2007).
- 17. Du, J., Li, P., Liu, H., Lu, D., Liang, H., Dou, Y.: Phenotypic and molecular characterization of multidrug resistant *Klebsiella pneumoniae* isolated from a university teaching hospital, China. PLoS One **9**, e95181 (2014).
- Hujer, K. M., Hujer, A. M., Hulten, E. A., Bajaksouzian, S., Adams, J. M., Donskey, C. J., Ecker, D. J., Massire, C., Eshoo, M. W., Sampath, R., Thomson, J. M., Rather, P. N., Craft, D. W., Fishbain, J. T., Ewell, A. J., Jacobs, M. R., Paterson, D. L., Bonomo, R. A.: Analysis of antibiotic resistance genes in multidrug-resistant *Acinetobacter* sp. isolates from military and civilian patients treated at the Walter Reed Army Medical Center. Antimicrob Agents Chemother **50**, 4114–4123 (2006).
- Voulgari, E., Gartzonika, C., Vrioni, G., Politi, L., Priavali, E., Levidiotou-Stefanou, S., Tsakris, A.: The Balkan region: NDM-1-producing *Klebsiella pneumoniae* ST11 clonal strain causing outbreaks in Greece. J Antimicrob Chemother 69, 2091–2097 (2014).
- Preechachuawong, P., Santimaleeworagun, W., Jitwasinkul, T., Samret, W.: Detection of New Delhi metallo-beta-lactamase-1-producing *Klebsiella pneumoniae* at a general hospital in Thailand. Southeast Asian J Trop Med Public Health 46, 1031–1036 (2015).
- Edelstein, M., Pimkin, M., Palagin, I., Edelstein, I., Stratchounski, L.: Prevalence and molecular epidemiology of CTX-M extended-spectrum beta-lactamase-producing *Escherichia coli* and *Klebsiella pneumoniae* in Russian hospitals. Antimicrob Agents Chemother 47, 3724–3732 (2003).
- Magiorakos, A. P., Srinivasan, A., Carey, R. B., Carmeli, Y., Falagas, M. E., Giske, C. G., Harbarth, S., Hindler, J. F., Kahlmeter, G., Olsson-Liljequist, B., Paterson, D. L., Rice, L. B., Stelling, J., Struelens, M. J., Vatopoulos, A., Weber, J. T., Monnet, D. L.: Multidrugresistant, extensively drug-resistant and pandrug-resistant bacteria: An international expert proposal for interim standard definitions for acquired resistance. Clin Microbiol Infect 18, 268–281 (2012).
- Akova, M., Daikos, G. L., Tzouvelekis, L., Carmeli, Y.: Interventional strategies and current clinical experience with carbapenemase-producing Gram-negative bacteria. Clin Microbiol Infect 18, 439–448 (2012).
- Shahraki-Zahedani, S., Moghadampour, M., Bokaeian, M., Ansari-Moghaddam, A.: Prevalence of CTX-M-8 and CTX-M-15 type extended-spectrum beta-lactamases between *Klebsiella pneumoniae* spp. isolated from Zahedan, Southeast Iran. J Chemother 28, 343–345 (2016).
- 25. Djahmi, N., Dunyach-Remy, C., Pantel, A., Dekhil, M., Sotto, A., Lavigne, J. P.: Epidemiology of carbapenemase-producing *Enterobacteriaceae* and *Acinetobacter baumannii* in Mediterranean countries. Biomed Res Int **2014**, 305784 (2014).
- Adler, A., Shklyar, M., Schwaber, M. J., Navon-Venezia, S., Dhaher, Y., Edgar, R., Solter, E., Benenson, S., Masarwa, S., Carmeli, Y.: Introduction of OXA-48-producing *Enterobacteriaceae* to Israeli hospitals by medical tourism. J Antimicrob Chemother 66, 2763–2766 (2011).
- Azimi, L., Nordmann, P., Lari, A. R., Bonnin, R. A.: First report of OXA-48-producing Klebsiella pneumoniae strains in Iran. GMS Hygiene Infect Control 9, Doc07 (2014).
- Hashemi, A., Fallah, F., Erfanimanesh, S., Hamedani, P., Alimehr, S., Goudarzi, H.: Detection of beta-lactamases and outer membrane porins among *Klebsiella pneumoniae* strains isolated in Iran. Scientifica **2014**, 726179 (2014).

- Hosseinzadeh, Z., Sedigh Ebrahim-Saraie, H., Sarvari, J., Mardaneh, J., Dehghani, B., Rokni-Hosseini, S. M. H., Motamedifar, M.: Emerge of *bla*_{NDM-1} and *bla*_{OXA-48}-like harboring carbapenem-resistant *Klebsiella pneumoniae* isolates from hospitalized patients in southwestern Iran. J Chin Med Assoc **81**, 536–540 (2017).
- Solgi, H., Badmasti, F., Giske, C. G., Aghamohammad, S., Shahcheraghi, F.: Molecular epidemiology of NDM-1- and OXA-48-producing *Klebsiella pneumoniae* in an Iranian hospital: Clonal dissemination of ST11 and ST893. J Antimicrob Chemother **73**, 1517–1524 (2018).
- Eskandari-Nasab, E., Moghadampour, M., Tahmasebi, A.: Prevalence of *bla*_{CTX-M} gene among extended-spectrum β-lactamases producing *Klebsiella pneumoniae* clinical isolates in Iran: A meta-analysis. Iran J Med Sci **43**, 347–354 (2018).
- Fazeli, H., Norouzi-Barough, M., Ahadi, A. M., Shokri, D., Solgi, H.: Detection of New Delhi metallo-beta-lactamase-1 (NDM-1) in carbapenem-resistant *Klebsiella pneumoniae* isolated from a university hospital in Iran. Hippokratia 19, 205–209 (2015).
- Khorvash, F., Yazdani, M. R., Soudi, A. A., Shabani, S., Tavahen, N.: Prevalence of acquired carbapenemase genes in *Klebsiella pneumoniae* by multiplex PCR in Isfahan. Adv Biomed Res 6, 41 (2017).