Published online 2023 May 3.

Research Article

A Retrospective Cross-sectional Study on Prevalence and Susceptibility Pattern of Bacterial Pathogens Causing Urinary Tract Infections at a Teaching Hospital in Tehran, Iran

Maryam Noori¹, Shadi Aghamohammad², Fatemeh Ashrafian³, Mehrdad Gholami ¹/₀⁴, Hossein Dabiri^{1,*} and Hossein Goudarzi¹

¹Department of Microbiology, School of Medicine, Shahid Beheshti University of Medical Sciences, Tehran, Iran

²Department of Bacteriology, Pasteur Institute of Iran, Tehran, Iran

³Department of Clinical Research, Pasteur Institute of Iran, Tehran, Iran

⁴Department of Microbiology and Virology, Faculty of Medicine, Mazandaran University of Medical Sciences, Sari, Iran

Corresponding author: Department of Microbiology, School of Medicine, Shahid Beheshti University of Medical Sciences, Tehran, Iran. Email: hodabiri@gmail.com

Received 2022 August 26; Revised 2022 December 26; Accepted 2023 March 05.

Abstract

Background: Urinary tract infection (UTI) is the most prevalent infection among the community and hospitalized patients. **Objectives:** This study aimed to investigate the current antimicrobial susceptibility patterns among UTI agents in Tehran, Iran. **Methods:** This retrospective study analyzed 9836 urine samples collected from hospitalized patients within 2019 - 2020. The antibiotic susceptibility for commonly-used antibiotics was tested according to Clinical and Laboratory Standards Institute guidelines. **Results:** Based on the findings, *Escherichia coli* was the most prevalent etiological agent of UTIs (72.3%), followed by *Klebsiella* spp. (13.4%), *Pseudomonas aeruginosa* (4.8%), *Acinetobacter* spp. (2.8%), and other species (6.7%). Of isolated microorganisms, 943 cases (97%) belonged to gram-negative bacilli; however, 32 cases (3.05 %) were gram-positive cocci. The susceptibility rates of *E. coli* to amikacin, nitrofurantoin, gentamicin, imipenem, and cefoperazone were 88.4%, 87.5%, 68.3%, 65.9%, and 62.6%, respectively. The sensitivity rates of *Klebsiella* spp. isolates for amikacin, nitrofurantoin, and imipenem were 87.6%, 71.5%, and 68.9%, respectively. **Conclusions:** The results of the present study characterized the misuse of antibiotics in Iran. Iranian surveillance studies will assist clinicians in choosing the most appropriate empirical treatment and preventing infections caused by resistant organisms.

Keywords: Urinary Tract Infection, Antimicrobial Susceptibility, Gram-negative Bacteria, Gram-positive Bacteria, Iran

1. Background

One of the most common infectious diseases is urinary tract infection (UTI), accounting for more than 150 million cases globally per year (1). It is also considered one of the frequent reasons for referring to medical clinics and is the most common infection after upper respiratory tract infections and gastrointestinal infections (2, 3). Symptoms of recurrent UTI infections include renal hypertension and renal failure if untreated, which can lead to irreversible kidney damage (4). Acute UTI is associated with substantial morbidity and mortality and problems of recurrent infections in both outpatients and hospitalized patients (5).

Bacteria are the main etiologic agent for UTI cases; however, other microorganisms, such as fungi and viruses, can also be rare causes of UTI (6). *Escherichia coli* is the most prevalent uropathogen and is solely responsible for approximately 70 - 95% of UTIs, followed by other Enterobacteriaceae, such as Klebsiella and Proteus species (7).

The UTIs account for a significant portion of antibiotic consumption in hospitals and out of them, particularly in developing countries where there is no accurate antibiotic prescription control. The UTIs have a large socioeconomic impact and result in antibiotic-resistant strains in the hospital and the community (8). Urinary pathogens have been known to include numerous strains resistant to many of the commonly used antibiotics (9). Multidrug-resistant organisms (MDROs) causing UTIs are widely known as a major threat to the public health system (10).

The MDROs have been increasingly reported in community-acquired and hospital-acquired infections; however, the prevalence is different by region (11). Some bacteria have been recognized as very important in developing MDROs, including extended-spectrum ß-lactamase (ESBL) producing *E. coli, Klebsiella* spp., methicillinresistant *Staphylococcus aureus* (MRSA), vancomycin-

Copyright © 2023, Author(s). This is an open-access article distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 International License (http://creativecommons.org/licenses/by-nc/4.0/) which permits copy and redistribute the material just in noncommercial usages, provided the original work is properly cited.

resistant (VRE) *Enterococcus* spp., and multidrug-resistant (MDR) non-fermenting bacteria, such as *Acinetobacter* spp. and *Pseudomonas* spp. (12-15). The MDR microorganisms would increase morbidity and mortality rates if they were not covered by suitable antibiotics (16). The UTI manifestation is a widespread reason for referring or returning patients to hospitals in Iran, and unfortunately, the numbers have been rising, according to several studies (17-19).

2. Objectives

The present study aimed to distinguish the bacterial etiological agents from cases of UTI and their drug resistance pattern to commonly used antimicrobial agents among hospitalized patients. It is hoped that the gained information will help provide proper antibiotic therapy for cases of UTI in Tehran, Iran.

3. Methods

3.1. Patents and Sampling

Overall, the current study included 9836 patients referred to a large teaching hospital (Loghman Hospital, Tehran, Iran) from March 2019 to February 2020. These patients were suspected of UTI according to clinical manifestations checked by a specialist. In inpatient or outpatient midstream clean catch, urine samples were collected in a sterile container from each patient. The samples were immediately transferred to a clinical microbiology laboratory, and direct examination and culture tests were performed.

3.2. Bacterial Isolation and Identification Procedures

Media were incubated aerobically at 37°C for 24 hours, and those cultures which became negative at the end of 24 hours of incubation were further incubated for 48 hours. A sample was considered positive for UTI if a single organism was cultured at 10⁵ cfu/mL (5). The organisms were identified based on phenotypic features, such as gram staining and colony morphology, and biochemical tests. For biochemical analysis, some catalase, oxidase, coagulase, Triple Sugar Iron agar, citrate utilization (Simmons's citrates medium), urease (Christensen's Urea Agar), indole, motility, H₂S production (Sulfide Indole Motility Medium), methyl red, Voges-Proskauer, Lysine agar, mannitol salt agar, Dnase agar, esculin hydrolysis (Bile-esculin agar), and sugar fermentation tests were considered according to suspected microorganisms.

3.3. Antimicrobial Susceptibility Test

According to the Kirby-Bauer method, antibiotic susceptibility testing was conducted against the most common causative UTI pathogens. The antibiotics used for gram-negative bacteria were trimethoprim/sulfamethoxazole (1.25/23.75 mcg), cefotaxime (30 mcg), cefoperazone (75 mcg), ceftazidime (30 mcg), imipenem (10 mcg), nitrofurantoin (300 mcg), nalidixic acid (30 mcg), amikacin (30 mcg), norfloxacin (10 mcg), ampicillin (10 mcg), and gentamicin (10 mcg), and those used for gram-positive bacteria were trimethoprim/sulfamethoxazole (1.25/23.75 mcg), cefotaxime (30 mcg), cefoperazone (75 mcg), ceftazidime (30 mcg), imipenem (10 mcg), norfloxacin (10 mcg), ampicillin (10 mcg), gentamicin (10 mcg), chloramphenicol (30 mcg), clindamycin (2 mcg), erythromycin (15 mcg), vancomycin (30 mcg), penicillin (10 units), and oxacillin (1 mcg). The results were interpreted according to the recommendation of Clinical and Laboratory Standards Institute criteria as sensitive (S), intermediate (I), and resistant (R) (16).

Reference strains of *E. coli*, ATCC 25922, *Klebsiella pneumoniae*, ATCC700603, and *P. aeruginosa* ATCC 27853 were used as controls for the gram-negative bacteria and included in all daily runs. *Staphylococcus aureus* ATCC 29213 and *Enterococcus faecalis* ATCC 29212 were used as grampositive controls (16). An MDRO isolate was determined as resistant to at least three of the following antimicrobial categories:

Folate pathway antagonists (trimethoprimsulfamethoxazole), aminoglycosides (amikacin or gentamicin), fluoroquinolones (norfloxacin or nalidixic acid), and nitrofurantoin

3.4. Statistical Analysis

Finally, the chi-square test was used with the help of the SPSS software (version 25.0) for the statistical analyses of this study. A P-value less than 0.05 was considered to be significant.

4. Results

Overall, 972 (9.88%) of 9836 patients with suspected UTI were referred for urine culture. The mean age of the subjects was 35 ± 19.4 years. In addition, 608 (62.6%) female and 364 (37.4%) male subjects were positive in bacterial culture (Table 1). According to the obtained results, females were more susceptible to developing UTIs (P < 0.01). There was no significant correlation between ages with culture results regardless of the microorganisms' type (P > 0.05) since all mixed infections were excluded from the

able 1. Distribution of Isolated Bacteria by Gender						
Bacteria	No. (%)	Ge	P-Value			
buccerna	(70)	Male, No. (%)	Female, No. (%)	-value		
Escherichia coli	703 (72.3)	235 (33.4)	468 (66.6)	0.0001		
Klebsiella spp.	130 (13.4)	56 (43.1)	74 (56.9)	0.2368		
Pseudomonas aeruginosa	47(4.8)	18 (38.3)	29 (61.7)	0.2129		
Acinetobacter spp.	27 (2.8)	19 (70.4)	8 (29.6)	0.0992		
Proteus spp.	17 (1.7)	9 (52.9)	8 (47.1)	1.0000		
Enterobacter spp.	13 (1.3)	9 (69.2)	4 (30.8)	0.3179		
Coagulase-negative Staphylococcus aureus	10 (1.0)	7 (70)	3 (30.9)	0.6828		
Streptococcus spp.	9 (0.9)	3 (33.3)	6 (66.7)	0.6828		
S. aureus	8 (0.8)	3 (37.5)	5 (62.6)	0.6792		
Enterococcus spp.	5 (0.5)	3(60)	2(40)	1.0000		
Citrobacter spp.	3 (0.3)	1(33.3)	2 (66.7)	1.0000		
Total	972	364 (37.4)	608 (62.6)	0.0001		

Table 2. Frequency/Percentage of Gram-negative Strains Susceptible to Selected Antibiotics ^a

Bacteria	SXT	стх	CP	CAZ	IMP	FM	NA	AN	NOR	AM	GM
Escherichia coli	321 (45.9)	402 (57.9)	430 (62.6)	408 (58.9)	373 (65.9)	608 (87.5)	295 (44.5)	609 (88.4)	424 (61.1)	39 (48.1)	473 (68.3)
Klebsiella spp.	67 (51.5)	75 (57.7)	83 (64.8)	74 (56.9)	71 (68.9)	93 (71.5)	63 (51.2)	113 (87.6)	87 (66.9)	10 (62.5)	81(62.3)
Pseudomonas aeruginosa	10 (21.3)	26 (55.3)	40 (85.1)	37 (78.7)	31 (79.5)	7 (14.9)	9 (19.6)	36 (78.3)	39 (83)	4(50)	38 (80.9)
Acinetobacter spp.	5 (18.5)	4 (14.8)	5 (18.5)	4 (14.8)	3 (14.3)	3 (11.1)	4 (15.4)	6 (22.2)	6 (22.2)	1(25)	11 (40.7)
Proteus spp.	4 (23.5)	9 (52.9)	12 (70.6)	10 (58.8)	8 (61.5)	8 (47.1)	6 (35.3)	12 (70.6)	12 (70.6)	0(0)	14 (82.4)
Enterobacter spp.	9 (69.9)	8 (61.5)	9 (69.2)	8 (61.5)	6 (75)	4 (30.8)	8 (61.5)	11 (84.6)	9 (69.2)	1(33.3	12 (92.3)
Citrobacter spp.	1 (33.3)	2 (66.7)	2 (66.7)	2 (66.7)	2 (66.7)	3 (100)	1 (33.3)	3 (100)	2 (66.7)	ND	2 (66.7)

Abbreviations: SXT, trimethoprim/sulfamethoxazole; CTX, cefotaxime; CP, cefoperazone; CAZ, ceftazidime; IPM, imipenem; FM, nitrofurantoin; NA, nalidixic acid; AN, amikacin; NOR, norfloxacin; AM, ampicillin; GM, gentamicin; ND, not determined.

^a Values are expressed as No. (%).

study. Out of 972 bacterial strains, *E. coli* was the predominant bacterial isolate and accounted for 703 (72.3%), followed by *Klebsiella* spp. with 130 (13.4%) and *Pseudomonas aeruginosa* with 47 (4.8%) isolates (Table 1). Of isolated microorganisms, 96.81% and 3.18% belonged to gram-negative and gram-positive bacteria, respectively. The finding indicated that 83% of the isolates belonged to the six common pathogens, including *E. coli*, *Klebsiella* spp., *P. aeruginosa*, *Acinetobacter* spp., *Proteus* spp., and *Enterobacter* spp.

Trimethoprim/sulfamethoxazole showed less activity (n = 550, 56.9%) against commonly isolated pathogens regardless of the type of microorganisms. However, amikacin (n = 787, 84.9%), nitrofurantoin (n = 723, 77.1%), gentamicin (n = 629, 65.6%), imipenem (n = 492, 63.2%), and norfloxacin (n = 577, 61.7%) were observed to be more effective for common pathogens (Table 2). For *E. coli* as a predominant isolate, 88.4% and 87.5% of the isolates were susceptible to amikacin and nitrofurantoin, respectively. Additionally, 55.5% and 54.1% of the isolates were resistant to nalidixic acid and trimethoprim-sulfamethoxazole, re-

spectively.

Antibiotic susceptibility pattern for isolated *Klebsiella* spp. showed susceptibility to amikacin (87.6%) and nitro-furantoin (71.5%); however, 48.8%, 48.5%, and 43.1% were resistant to nalidixic acid, trimethoprim/sulfamethoxazole, and ceftazidime, respectively. Among *P. aeruginosa* isolates, as a third common isolated bacteria, cefoperazone (85.1%), norfloxacin (83%), gentamicin (80.9%), and imipenem (79.5%) had the highest rates of susceptibility; however, 85.1%, 80.4%, and 78.7% of the isolates were resistant to nitrofurantoin, nalidixic acid, and trimethoprim/sulfamethoxazole, respectively.

For Acinetobacter spp., only 40.7% of the isolates were susceptible to gentamicin, and a large number of the isolates were resistant to nitrofurantoin (88.9%), ceftazidime (85.2%), and cefotaxime (85.2%). Of 8 isolated *S. aureus*, 7 (87.5%) were MRSA (oxacillin/methicillin resistant). Only one isolate was susceptible to oxacillin, and one isolate (20%) was detected as VRE *Staphylococcus aureus*. Among *Enterococcus* spp., of 5 isolates, 3 (60%) were VRE (Table 3).

Bacteria	SXT	стх	СР	CAZ	IPM	NOR	AM	GM	c	сс	E	v	р	ox
Staphylococcus aureus	5 (71.4)	3 (50)	5 (83.3)	2(100)	6 (100)	2(100)	1(100)	5 (83.3)	4 (66.7)	5 (83.3)	4 (57.1)	4 (80)	0(0)	7(12.5)
CONS	7(70)	6 (75)	6 (75)	2 (66.7)	7(100)	ND	1(1000)	7 (87.5)	7 (87.5)	8 (88.9)	5(50)	10 (100)	4 (57.1)	ND
Streptococcus spp.	2 (25)	5 (71.4)	4 (57.1)	ND	3 (42.9)	1 (100)	3 (100)	3 (33.3)	1(33.3)	4 (57.1)	5 (62.5)	8 (100)	4 (80)	ND
Enterococcus spp.	0(0)	1(20)	1(25)	ND	1(20)	ND	1(50)	1(25)	2(40)	0(0)	1(20)	2(40)	1(25)	ND

Abbreviations: SXT, trimethoprim/sulfamethoxazole; CTX, cefotaxime; CP, cefoperazone; CAZ, ceftazidime; IPM, imipenem; NOR, norfloxacin; AM, ampicillin; GM, gentamicin; C, chloramphenicol; CC, clindamycin; E, erythromycin; V, vancomycin; P, pericillin; OXOS, coagulase-negative Staphylococcus aureus; ND, not determined. Values are expressed as No. (%).

Of the common isolates collected in this study, 547 isolates (56.27%) were MDR. The Acinetobacter spp. isolates had the highest value (n = 24, 88.8%) for multidrug resistance, followed by 31 (65.9%) of P. aeruginosa, 80 (61.5%) of Klebsiella spp., and 399 (56.7%) of E. coli isolates, respectively (Table <mark>4</mark>).

able 4. Frequency of Bacterial Isolates Showing Multidrug-resistant Phenotyp					
Bacteria	Multidrug Resistance, No. (%)				
Escherichia coli	399 (56.7)				
Klebsiella spp.	80 (61.5)				
Pseudomonas spp.	31 (65.9)				
Acinetobacter spp.	24 (88.8)				
Total	534 (56)				

5. Discussion

Several geographical regions, including Iran, show decreased susceptibility rates to common urinary pathogens; therefore, the global trend to empirically treat community-acquired UTIs might not apply to these regions (20, 21). Antimicrobial resistance in uropathogens should be monitored to improve treatment recommendations. This study was conducted to determine the frequency and antimicrobial susceptibility patterns of community-acquired uropathogens in Iran. The current study demonstrated that E. coli is the most common cause of UTI in Tehran, Iran. This finding corresponds with the data obtained by other investigators (22-24). Klebsiella spp. was the second most common organism, followed by Pseudomonas spp. and Acinetobacter spp., which is similar to a report from Mohammadi-Mehr, and Feizabadi (25) and different from a report from Pouladfar et al. indicating Klebsiella spp. and Enterococcus spp. as prevalent strains next to E. coli (26).

The results of the present study showed that females are more likely to get UTI (P < 0.05) which is similar to nearly all the other reports (17, 19, 24). Fluoroquinolones or nitrofurantoin have been suggested for the empirical treatment of uncomplicated UTIs (27). However, the emergence of high levels of resistance of uropathogenic E. coli against trimethoprim/sulfamethoxazole has been reported in both developing countries (54 - 82%) and developed countries (14.6 - 37.1%) (28-30). The present study also discovered an elevated resistance rate to trimethoprim/sulfamethoxazole (54.1%), which is in accordance with other Iranian studies (19, 24, 31). According to these results, trimethoprim-sulfamethoxazole should no longer be used as the primary empirical treatment in Iran.

The results of the fluoroquinolone susceptibility test in the current study (norfloxacin) showed good action against E. coli (61%), which is in line with other Iranian studies carried out in 2006, indicating constant sensitivity of E. coli isolates to fluoroquinolones. Nitrofurantoin, as the second preferred antibiotic for the treatment of UTI, is effective for the prophylaxis and the treatment of MDR uropathogens in adults, children, and pregnant women. Additionally, it is a relatively safe drug with minimal effects on the resident bowel and vaginal flora (27, 32). Although nitrofurantoin demonstrated better activity against E. coli isolates (87.5% susceptible), it should not be used for serious upper UTIs or for those with systemic involvement (14).

In the present study, Klebsiella spp., as the second common cause of UTI, was resistant to commonly-used antibiotics, except amikacin (87.6%). Therefore, amikacin still remains the best choice for the empirical treatment of severe UTI caused by Klebsiella spp. The susceptibility to norfloxacin has remained constant during the past 3 years. The present study's sensitivity results (67%) are similar to a previous report from Shenagari et al., with 55% sensitivity (33). This finding might contribute to the limited usage of norfloxacin in Iranian patients.

Considering the current study's results, P. aeruginosa, with a 5% incidence, was the third most common cause of hospital-acquired UTIs. The currently studied Pseudomonas strains were susceptible to the second-line drugs, such as cefoperazone, norfloxacin, and gentamycin, with more than 80% of cases; however, most of these isolates were associated with high resistance to the firstline used antibiotics, namely nitrofurantoin, trimethoprim/sulfamethoxazole, and nalidixic acid. These findings are in agreement with another Iranian report in which

80% of isolates were sensitive to norfloxacin, and only 11% were sensitive to trimethoprim/sulfamethoxazole(34). Increased susceptibility was observed for nitrofurantoin, nalidixic acid, and trimethoprim/sulfamethoxazole, compared to the results of a previous study (35). This status might contribute to the reduced use of these antibiotics in Iran.

Acinetobacter spp. is known to be important in nosocomial UTIs (36). Acinetobacter spp. isolates demonstrated high resistance to most antibiotics, such as nitrofurantoin, imipenem, ceftazidime, cefotaxime, norfloxacin, ampicillin, trimethoprim/sulfamethoxazole, and nalidixic acid with an average of 83%. Despite most reports and in agreement with the results of studies by Rahimi and Rezaie Keikhaie et al., Acinetobacter spp. isolates showed partly good sensitivity to gentamicin (40%) (37, 38). In contrast to the results of the current study, Mortazavi et al. (39) reported very high resistance to gentamicin and amikacin simultaneously among 80 A. baumannii strains from Ahvaz, south-west Iran.

Previous studies reported the prevalence of S. aureus among UTI patients from 0.8%, 1%, and 6.92 to 11.65% (24, 40-42). The present study's results showed that 0.8% of patients were infected with S. aureus. The current study showed a high resistant rate (87%) to methicillin/oxacillin (MRSA) in comparison to previous national reports in 2012 (48%) and 2015 (28%) (43, 44). This difference might be related to the non-suitable usage of antibiotics or the low number (5) of studied organisms. A significant increase in MDR pathogenic strains to different antibiotics has been reported worldwide (45). Accordingly, 534 MDR (56%) isolates were detected. Of the 534 MDR isolates, 399 (57%) were E. coli. A lower percentage of MDR E. coli (63%) was found in Poland (22%) and Venezuela (25%) among isolates from community-acquired and hospital-acquired UTIs (7, 46). This diversity in MDR frequency reflects differences in antibiotic prescription and infection control policies in any region worldwide. In conclusion, a relatively high frequency of bacterial resistance was observed in the urine samples collected from Loghman Hospital in Tehran.

The data also indicated that most isolated microorganisms belonged to gram-negative bacilli (97%), and *E. coli* was the most frequent agent of UTIs (72.3%) in the current study. Considering bacterial diversity causing UTIs, aminoglycosides, such as amikacin, are recommended as the first choice, and nitrofurantoin as the second choice for the treatment of UTIs in Tehran. Nalidixic acid and trimethoprim/sulfamethoxazole, due to reduced efficiency against UTI causative agents, are no longer suggested for the empirical therapy of UTI.

Despite the precious findings on the resistance rate of uropathogens, there are some limitations in the present work. The major drawback pertained to the retrospective design of the study and the inability to have access to the patient's health records; therefore, the authors were unable to analyze and report the patients' demographic data and the correlation between the risk factors and underlying pathologies conditions with UTIs. In addition, the lack of molecular characterization of the resistance determinants in the studied isolates and no detection of ESBLs are other limitations of the current study. Further studies are essential to monitoring the rate of bacterial resistance among UTI patients in other hospitals in Iran.

5.1. Conclusions

The potential antimicrobial resistance is one crucial consideration for physicians when selecting an antibiotic for the treatment of infectious diseases, particularly for patients with UTIs. In most cases, antimicrobial chemotherapy is often empiric and should be determined by identifying the most common etiological agents and their antimicrobial susceptibility profiles.

Footnotes

Authors' Contribution: MN: Laboratory working and writing the manuscript; SA, FA, MG, and HD: Laboratory working; HG: Revision of the manuscript; HD: Designing the study and reviewing/editing the manuscript.

Conflict of Interests: The authors declare that they have no conflict of interest.

Funding/Support: None to declare.

References

- Zowawi HM, Harris PN, Roberts MJ, Tambyah PA, Schembri MA, Pezzani MD, et al. The emerging threat of multidrug-resistant Gram-negative bacteria in urology. *Nat Rev Urol.* 2015;**12**(10):570–84. [PubMed ID: 26334085]. https://doi.org/10.1038/nrurol.2015.199.
- Byron JK. Urinary Tract Infection. Vet Clin North Am Small Anim Pract. 2019;49(2):211–21. [PubMed ID: 30591189]. https://doi.org/10.1016/j.cvsm.2018.11.005.
- Behzadi P, Urban E, Matuz M, Benko R, Gajdacs M. The Role of Gram-Negative Bacteria in Urinary Tract Infections: Current Concepts and Therapeutic Options. Adv Exp Med Biol. 2021;1323:35–69. [PubMed ID: 32596751]. https://doi.org/10.1007/5584_2020_566.
- Murray BO, Flores C, Williams C, Flusberg DA, Marr EE, Kwiatkowska KM, et al. Recurrent Urinary Tract Infection: A Mystery in Search of Better Model Systems. Front Cell Infect Microbiol. 2021;11:691210. [PubMed ID: 34123879]. [PubMed Central ID: PMC8188986]. https://doi.org/10.3389/fcimb.2021.691210.
- Medina M, Castillo-Pino E. An introduction to the epidemiology and burden of urinary tract infections. *Ther Adv Urol.* 2019;**11**:1756287219832170. [PubMed ID: 31105774]. [PubMed Central ID: PMC6502976]. https://doi.org/10.1177/1756287219832172.

- Tessema NN, Ali MM, Zenebe MH. Bacterial associated urinary tract infection, risk factors, and drug susceptibility profile among adult people living with HIV at Haswassa University Comprehensive Specialized Hospital, Hawassa, Southern Esthiopia. *Sci Rep.* 2020;**10**(1):10790. [PubMed ID: 32612139]. [PubMed Central ID: PMC7330024]. https://doi.org/10.1038/s41598-020-67840-7.
- Kot B, Gruzewska A, Szweda P, Wicha J, Parulska U. Antibiotic Resistance of Uropathogens Isolated from Patients Hospitalized in District Hospital in Central Poland in 2020. *Antibiotics (Basel)*. 2021;10(4). [PubMed ID: 33923389]. [PubMed Central ID: PMC8071495]. https://doi.org/10.3390/antibiotics10040447.
- Kostakioti M, Hultgren SJ, Hadjifrangiskou M. Molecular blueprint of uropathogenic Escherichia coli virulence provides clues toward the development of anti-virulence therapeutics. *Virulence*. 2012;3(7):592-4. [PubMed ID: 23154288]. [PubMed Central ID: PMC3545937]. https://doi.org/10.4161/viru.22364.
- Khan MI, Xu S, Ali MM, Ali R, Kazmi A, Akhtar N, et al. Assessment of multidrug resistance in bacterial isolates from urinary tract-infected patients. J Radiat Res Appl Sci. 2020;13(1):267-75. https://doi.org/10.1080/16878507.2020.1730579.
- Rodriguez-Villodres A, Martin-Gandul C, Penalva G, Guisado-Gil AB, Crespo-Rivas JC, Pachon-Ibanez ME, et al. Prevalence and Risk Factors for Multidrug-Resistant Organisms Colonization in Long-Term Care Facilities Around the World: A Review. *Antibiotics (Basel)*. 2021;**10**(6). [PubMed ID: 34200238]. [PubMed Central ID: PMC8228357]. https://doi.org/10.3390/antibiotics10060680.
- Almomani BA, Hayajneh WA, Ayoub AM, Ababneh MA, Al Momani MA. Clinical patterns, epidemiology and risk factors of community-acquired urinary tract infection caused by extendedspectrum beta-lactamase producers: a prospective hospital casecontrol study. *Infection*. 2018;46(4):495–501. [PubMed ID: 29748840]. https://doi.org/10.1007/s15010-018-1148-y.
- Gajdacs M, Burian K, Terhes G. Resistance Levels and Epidemiology of Non-Fermenting Gram-Negative Bacteria in Urinary Tract Infections of Inpatients and Outpatients (RENFUTI): A 10-Year Epidemiological Snapshot. Antibiotics (Basel). 2019;8(3):143. [PubMed ID: 31505817]. [PubMed Central ID: PMC6784256]. https://doi.org/10.3390/antibiotics8030143.
- Horcajada JP, Montero M, Oliver A, Sorli L, Luque S, Gomez-Zorrilla S, et al. Epidemiology and Treatment of Multidrug-Resistant and Extensively Drug-Resistant Pseudomonas aeruginosa Infections. *Clin Microbiol Rev.* 2019;**32**(4). [PubMed ID: 31462403]. [PubMed Central ID: PMC6730496]. https://doi.org/10.1128/CMR.00031-19.
- Kyriakidis I, Vasileiou E, Pana ZD, Tragiannidis A. Acinetobacter baumannii Antibiotic Resistance Mechanisms. *Pathogens*. 2021;10(3). [PubMed ID: 33808905]. [PubMed Central ID: PMC8003822]. https://doi.org/10.3390/pathogens10030373.
- Agyepong N, Govinden U, Owusu-Ofori A, Essack SY. Multidrugresistant gram-negative bacterial infections in a teaching hospital in Ghana. *Antimicrob Resist Infect Control.* 2018;7:37.
 [PubMed ID: 29541448]. [PubMed Central ID: PMC5845144]. https://doi.org/10.1186/s13756-018-0324-2.
- Tanwar J, Das S, Fatima Z, Hameed S. Multidrug resistance: an emerging crisis. *Interdiscip Perspect Infect Dis*. 2014;**2014**:541340.
 [PubMed ID: 25140175]. [PubMed Central ID: PMC4124702]. https://doi.org/10.1155/2014/541340.
- Nozarian Z, Abdollahi A. Microbial Etiology and Antimicrobial Susceptibility of Bactria Implicated in Urinary Tract Infection in Tehran, Iran. Iran J Pathol. 2015;10(1):54–60. [PubMed ID: 26516326]. [PubMed Central ID: PMC4539787].
- Mostafavi SN, Rostami S, Rezaee Nejad Y, Ataei B, Mobasherizadeh S, Cheraghi A, et al. Antimicrobial Resistance in Hospitalized Patients with Community Acquired Urinary Tract Infection in Isfahan, Iran. Arch Iran Med. 2021;24(3):187-92. [PubMed ID: 33878876]. https://doi.org/10.34172/aim.2021.29.
- 19. Mamishi S, Shalchi Z, Mahmoudi S, Hosseinpour Sadeghi R, Haghi

Ashtiani MT, Pourakbari B. Antimicrobial Resistance and Genotyping of Bacteria Isolated from Urinary Tract Infection in Children in an Iranian Referral Hospital. *Infect Drug Resist.* 2020;**13**:3317-23. [PubMed ID: 33061479]. [PubMed Central ID: PMC7535122]. https://doi.org/10.2147/IDR.S260359.

- Hadifar S, Moghoofei M, Nematollahi S, Ramazanzadeh R, Sedighi M, Salehi-Abargouei A, et al. Epidemiology of Multidrug Resistant Uropathogenic Escherichia coli in Iran: a Systematic Review and Meta-Analysis. Jpn J Infect Dis. 2017;70(1):19–25. [PubMed ID: 27000462]. https://doi.org/10.7883/yoken.JJID.2015.652.
- Naziri Z, Derakhshandeh A, Soltani Borchaloee A, Poormaleknia M, Azimzadeh N. Treatment Failure in Urinary Tract Infections: A Warning Witness for Virulent Multi-Drug Resistant ESBL- Producing Escherichia coli. *Infect Drug Resist.* 2020;**13**:1839– 50. [PubMed ID: 32606833]. [PubMed Central ID: PMC7306463]. https://doi.org/10.2147/IDR.S256131.
- Fallah F, Parhiz S, Azimi L. Distribution and antibiotic resistance pattern of bacteria isolated from patients with community-acquired urinary tract infections in Iran: a cross-sectional study. Int J Health Stud. 2019;4(2). https://doi.org/10.22100/ijhs.v4i2.631.
- Alizade H. Escherichia coli in Iran: An Overview of Antibiotic Resistance: A Review Article. *Iran J Public Health*. 2018;47(1):1–12. [PubMed ID: 29318111]. [PubMed Central ID: PMC5756583].
- 24. Ghanbari F, Khademi F, Saberianpour S, Shahin M, Ghanbari N, Naderi K, et al. An Epidemiological Study on the Prevalence and Antibiotic Resistance Patterns of Bacteria Isolated from Urinary Tract Infections in Central Iran. Avicenna J Clin Microbiol Infect. 2017;4(3):42214. https://doi.org/10.5812/ajcmi.42214.
- Mohammadi-Mehr M, Feizabadi M. Antimicrobial resistance pattern of Gram-negative bacilli isolated from patients at ICUs of Army hospitals in Iran. *Iran J Microbiol*. 2011;3(1):26–30. [PubMed ID: 22347579]. [PubMed Central ID: PMC3279800].
- Pouladfar G, Basiratnia M, Anvarinejad M, Abbasi P, Amirmoezi F, Zare S. The antibiotic susceptibility patterns of uropathogens among children with urinary tract infection in Shiraz. *Medicine (Baltimore)*. 2017;**96**(37). e7834. [PubMed ID: 28906365]. [PubMed Central ID: PMC5604634]. https://doi.org/10.1097/MD.000000000007834.
- Porreca A, D'Agostino D, Romagnoli D, Del Giudice F, Maggi M, Palmer K, et al. The Clinical Efficacy of Nitrofurantoin for Treating Uncomplicated Urinary Tract Infection in Adults: A Systematic Review of Randomized Control Trials. Urol Int. 2021;105(7-8):531–40. [PubMed ID: 33535221]. https://doi.org/10.1159/000512582.
- Kot B. Antibiotic Resistance Among Uropathogenic Escherichia coli. Pol J Microbiol. 2019;68(4):403-15. [PubMed ID: 31880885]. [PubMed Central ID: PMC7260639]. https://doi.org/10.33073/pjm-2019-048.
- Belete MA, Saravanan M. A Systematic Review on Drug Resistant Urinary Tract Infection Among Pregnant Women in Developing Countries in Africa and Asia; 2005-2016. *Infect Drug Resist.* 2020;13:1465-77. [PubMed ID: 32547115]. [PubMed Central ID: PMC7245001]. https://doi.org/10.2147/IDR.S250654.
- Sugianli AK, Ginting F, Parwati I, de Jong MD, van Leth F, Schultsz C. Antimicrobial resistance among uropathogens in the Asia-Pacific region: a systematic review. JAC Antimicrob Resist. 2021;3(1):dlab003. [PubMed ID: 34223081]. [PubMed Central ID: PMC8210283]. https://doi.org/10.1093/jacamr/dlab003.
- 31. Yekani M, Baghi HB, Sefidan FY, Azargun R, Memar MY, Ghotaslou R. The rates of quinolone, trimethoprim/sulfamethoxazole and aminoglycoside resistance among Enterobacteriaceae isolated from urinary tract infections in Azerbaijan, Iran. *GMS Hyg Infect Control.* 2018;13:Doc07. [PubMed ID: 30202721]. [PubMed Central ID: PMC6124734]. https://doi.org/10.3205/dgkh000313.
- Gardiner BJ, Stewardson AJ, Abbott IJ, Peleg AY. Nitrofurantoin and fosfomycin for resistant urinary tract infections: old drugs for emerging problems. *Aust Prescr.* 2019;**42**(1):14– 9. [PubMed ID: 30765904]. [PubMed Central ID: PMC6370609].

https://doi.org/10.18773/austprescr.2019.002.

- Shenagari M, Bakhtiari M, Mojtahedi A, Atrkar Roushan Z. High frequency of mutations in gyrA gene associated with quinolones resistance in uropathogenic Escherichia coli isolates from the north of Iran. *Iran J Basic Med Sci.* 2018;21(12):1226– 31. [PubMed ID: 30627365]. [PubMed Central ID: PMC6312683]. https://doi.org/10.22038/ijbms.2018.31285.7539.
- Kashef N, Djavid GE, Shahbazi S. Antimicrobial susceptibility patterns of community-acquired uropathogens in Tehran, Iran. J Infect Dev Ctries. 2010;4(4):202–6. [PubMed ID: 20440056]. https://doi.org/10.3855/jidc.540.
- Sadeghi Bojd S, Soleimani G, Teimouri A, Zarifi E, Rashidi S. Evaluation of antibiotic sensitivity of urinary tract pathogens among children in zahedan, south east of Iran. *Int J Pediatr.* 2017;5(10):5965–74. https://doi.org/10.22038/ijp.2017.22970.1924.
- 36. Jimenez-Guerra G, Heras-Canas V, Gutierrez-Soto M, Del Pilar Aznarte-Padial M, Exposito-Ruiz M, Navarro-Mari JM, et al. Urinary tract infection by Acinetobacter baumannii and Pseudomonas aeruginosa: evolution of antimicrobial resistance and therapeutic alternatives. J Med Microbiol. 2018;67(6):790–7. [PubMed ID: 29693543]. https://doi.org/10.1099/jmm.0.000742.
- Rahimi N, Honarmand Jahromy S, Zare Karizi S. Evaluation of Antibiotic Resistance Pattern of Meropenem and Piperacillin-Tazobactam in Multi Drug Resistant Acinetobacter baumannii Isolates by Flow Cytometry Method. *Iran J Med Microb.* 2019;13(3):194–209. https://doi.org/10.30699/ijmm.13.3.194.
- Rezaie Keikhaie K, Rezaie Keikhaie L, Bazi S, Javadian F. Antibiotic Resistance in Acinetobacter baumannii Isolated from Patients with Urinary Tract Infections in Zabol, Iran. Int J Infect. 2018;5(2). https://doi.org/10.5812/iji.60181.
- 39. Mortazavi SM, Farshadzadeh Z, Janabadi S, Musavi M, Shahi F, Moradi M, et al. Evaluating the frequency of carbapenem and aminoglycoside resistance genes among clinical isolates of Acinetobacter baumannii from Ahvaz, south-west Iran. New Microbes New In-

fect. 2020;**38**:100779. [PubMed ID: 33194209]. [PubMed Central ID: PMC7644744]. https://doi.org/10.1016/j.nmni.2020.100779.

- 40. Baghani Aval H, Ekrami Toroghi M, Haghighi F, Tabarraie Y. Common bacterial factors of urinary tract infections and determining their antibiotic resistance in hospitalized and out patients referred to the Vase'ee Hospital in Sabzevar in 2016. J Sabzevar Univ Med Sci. 2018.
- Pourmontaseri Z, Pourmontaseri M, Baziboroun M, Jorjani M, Bakhshizadeh S. Distribution and Antimicrobial Resistance Patterns of Urinary Tract Infection in Southern Iran. *Int J Nutr Sci.* 2018;3(2):113– 9.
- Rajabnia M, Bahadoram M, Fotoohi A, Mohammadi M. Antibiotic resistance patterns of urinary tract infections in the West of Iran. J Nephropharmacology. 2019;8(2):26. https://doi.org/10.15171/npj.2019.26.
- Dormanesh B, Siroosbakhat S, Khodaverdi Darian E, Afsharkhas L. Methicillin-Resistant Staphylococcus aureus Isolated From Various Types of Hospital Infections in Pediatrics: Panton-Valentine Leukocidin, Staphylococcal Chromosomal Cassette mec SCCmec Phenotypes and Antibiotic Resistance Properties. Jundishapur J Microbiol. 2015;8(11). e11341. [PubMed ID: 26862375]. [PubMed Central ID: PMC4741056]. https://doi.org/10.5812/jjm.11341.
- 44. Sabouni F, Mahmoudi S, Bahador A, Pourakbari B, Sadeghi RH, Ashtiani MT, et al. Virulence Factors of Staphylococcus aureus Isolates in an Iranian Referral Children's Hospital. Osong Public Health Res Perspect. 2014;5(2):96-100. [PubMed ID: 24955319]. [PubMed Central ID: PMC4064634]. https://doi.org/10.1016/j.phrp.2014.03.002.
- Esposito S, De Simone G. Update on the main MDR pathogens: prevalence and treatment options. *Infez Med.* 2017;25(4):301–10. [PubMed ID: 29286008].
- 46. Guzman M, Salazar E, Cordero V, Castro A, Villanueva A, Rodulfo H, et al. Multidrug resistance and risk factors associated with community-acquired urinary tract infections caused by Escherichia coli in Venezuela. *Biomedica*. 2019;**39**(s1):96–107. [PubMed ID: 31529852]. https://doi.org/10.7705/biomedica.v39i2.4030.