

Antibiogram and Bacterial Profile Analysis of Various Clinical Samples at Tertiary Healthcare Centre in North-West region of Punjab

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ABSTRACT

Background: Antimicrobial resistance is a significant barrier in achieving the sustainable development and is one of the leading global public health menaces to humanity. Present study determined the antibiogram profiles of bacteria isolated from different body site infections among patients admitted to tertiary care Hospital, Punjab, Northwest India.

Methods: A cross-sectional study was carried out using 11201 clinical samples at the tertiary care hospital Patiala from June 2023 to June 2024. Following the criteria of the Clinical and Laboratory Standards Institute *S. aureus* isolates and further tested for antimicrobial activity using standard procedures

Results: From the 11201 clinical specimens processed, a total of 1638 urine, 403 blood and 2126 other body fluids (pus and sputum) were culture positive for bacterial pathogens. The prevalence of infection was highest among middle aged males in blood and other fluids where middle aged female's predominance was found in urine infection. In urine samples, *Escherichia coli* (63.67%) from Gram-negative and *Enterococcus* (6.53%) and *S. Aureus* (4.45%) from gram positive were the predominant bacterial. Blood sample showed the prevalence of *K. pneumonia* (28.28%) among gram negative and *Enterococcus* (27.79%) among gram positive. *K. pneumonia* (20.93%) and *S. Aureus* (29.25%) were predominant in other fluid infections. antimicrobial resistance pattern in all clinical samples and results suggest that most of the bacteria isolated from studied clinical samples are less susceptible ranging from 0% to 35% mostly towards penicillin, norfloxacin, Cefuroxime and ampicillin. Intermediate susceptibility ranged from 40%-85% towards amikacin, Gentamicin and Meropenem. High susceptibility varied according to bacteria and sample type and it ranged between 90%-100%. In urine sample high susceptibility was observed for *P. aeruginosa* (100%) and *A. Baumannii* (100%). In Blood samples *S. aureus* showed the highest susceptibility towards amikacin (100%). Many antibiotics showed 100% susceptibility in other body fluids infections such as *P. aeruginosa* towards Norfloxacin, *S. aureus* towards amoxicillin, *enetrococcus* towards ampicillin, gentamicin and amikacin.

Conclusion: Overall result of the present investigation showed that penicillins are least susceptible and amikacin is highly susceptible antibiotic against the studied infections. By

focusing on locally prevalent antibiotic resistance patterns instead of regionally and internationally available data on antibiogram, we may ensure faster patient recovery at lower costs and avoid the unwarranted development of drug resistance.

Keywords: Antibiogram, Antibiotic resistance, Bacterial isolates, clinical specimen, Punja

INTRODUCTION

Antimicrobial resistance is a significant barrier in achieving the sustainable development and is one of the leading global public health menaces to humanity. Over the last two decades, antibiotic resistance has skyrocketed, and the primary cause of this can be irrational use, and the accessibility of these medications over-the-counter. [1] Microbial infections are responsible for at least 25% of the 60 million fatalities that occur worldwide each year.

Clinical infections such as bloodstream, UTIs and body fluid infections continue to be major causes of morbidity and mortality among hospitalized patients. [2] Bloodstream infections caused by bacteria or other germs are referred to as bloodstream infections (BSIs), septicemia, or bacteremia. Many pathogens, including as bacteria, viruses, and fungi, can cause bloodstream infections (BSIs). 7. *Staphylococcus*, *Escherichia*, and *Klebsiella* are the three most prevalent bacterial genera that cause bloodstream infections. [3] Urinary tract infection (UTI) is another most frequent hospital-acquired infection and the second most frequent reason for hospital visits. Gram positive organisms including *Staphylococcus saprophyticus*, *Staphylococcus aureus*, and *Enterococci*, as well as Gram negative enteric bacilli like *Escherichia coli* (*E. coli*), *Klebsiella spp.*, and *Proteus spp.*, are the most common causes of UTIs. [4] Lastly, bacterial infections of bodily fluids can result in invasive conditions like bacterial meningitis, bacteremia, sepsis, bacterial peritonitis, and other problems. [5]

Drug resistance is a significant issue in low-income nations due to the high rate of illnesses, the inappropriate use of antibiotics, the easy availability of medications, and the

lack of access to antimicrobial susceptibility testing. [6] The Indian populace is recognized as the world's largest consumer of antibiotics worsening AMR situation in India thus an action plan for its control is considered crucial. [7]

The region-wide report of culture and sensitivity data is provided by hospital antibiogram which indicates how an organism's susceptibility pattern is evolving. [8] Consequently, it will assist clinicians in selecting the right antibiotic, reducing the inappropriate use of antibiotics and the emergence of resistance. Thus, the purpose of this study is to ascertain the antibiotic susceptibility pattern and bacteriological profile of UTI bacterial isolates from a tertiary healthcare facility.

MATERIALS & METHODS

From June 2023 to June 2024, a prospective cross-sectional investigation was carried out at the tertiary care hospital, Punjab. The current investigation comprises 11201 isolates from different clinical specimens. Ethical clearance from the Research and Ethical committee was taken.

This study includes all the clinical samples that were properly collected, labeled, properly transported and processed for aerobic bacterial cultures. Isolates obtained from blood, urine, pus, and sputum, were included in this study. According to standard microbiological procedures received samples were processed.

Using the conventional Kirby Bauer disk diffusion technique, all isolates' antimicrobial susceptibility was assessed in accordance with Clinical Laboratory Standards Institute (CLSI) guidelines. Antibiotics included were amoxicillin (20mcg), ampicillin (50mcg),

gentamicin (30mcg), amikacin (15mcg), cefoxitin (30mcg), clindamycin (20mcg), doxycycline (50mcg), erythromycin (15mcg).

RESULT

The results show the distribution of several bacterial species that were isolated from clinical samples in the North West India during 2023. A total of 5053 urine samples, 2127 Blood samples and 4021 other body fluids such as pus, sputum samples were studied. Out of which 1638 (32.41%) urine isolates, 403 (18.94%) blood isolates and 2126 (52.87%) other body fluids isolates were reported in Table 1.

Among gram negative urine isolates *Esch.coli* (63.67%) is the most common bacterial isolates found followed by *K. pneumonia* (17.52%), *P. aeruginosa* (2.62%), *ABComplex* (1.28%), *Citrobacter* (1.09%), *Proteus* (0.97%), *Enterobacter* (0.79%), while *enterococcus* (6.53%) is most prevalent among gram positive urine isolates followed by *S. aureus* (4.45%) and *MRSA* (1.03%).

In case of Blood isolates Gram negative show different trend with *K. pneumonia* (28.28%) highest prevalent bacteria followed by *ABComplex* (8.68%), *Esch. coli* (7.94%), *P. aeruginosa* (5.45%), *Citrobacter* (2.72%), *Enterobacter spp* (1.98%) and *Proteus spp* (1.24%). Gram positive blood isolates showed similar trend as urine isolates with high prevalence of *Enterococcus* (27.79%) afterward *S. aureus* (13.39%) and *MRSA* (2.48%).

Other body fluids such as pus and sputum showed highest number of positive isolated among all samples with highest prevalence of *S.aureus* (29.25%) in gram positive bacteria and *K. pneumonia* (20.93%) in gram negative bacteria. After *K. pneumonia* among gram negative prevalence of *ABComplex* (15.75%) was observed followed by *Esch.coli* (11.52%), *P. aeruginosa* (3.80%), *Proteus* (0.65%), *Citrobacter* (0.61%), *Enterobacter* (0.42%) and among gram positive after *S.aureus* prevalence of *enterococcus* (15.85%) and *MRSA* (1.17%) was observed.

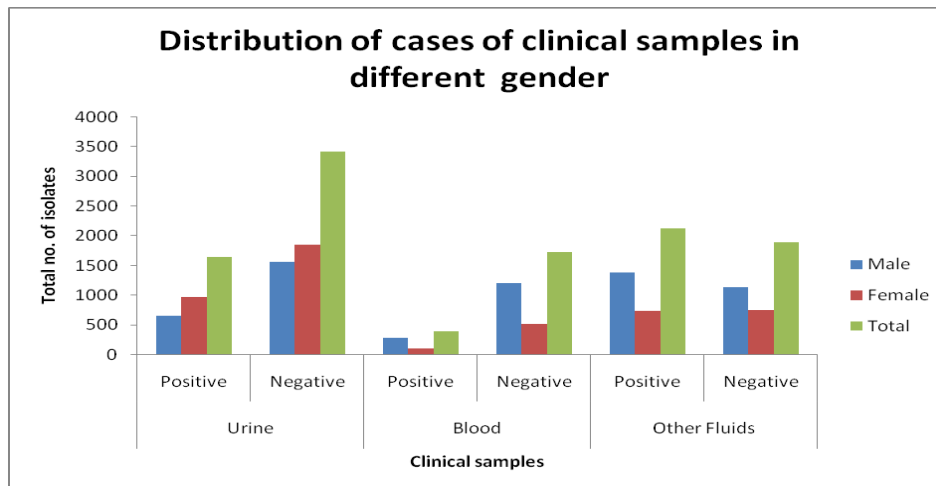
Table 1: Distribution of various isolates among clinical samples.

Organism	Gram staining	URINE SAMPLES	BLOOD SAMPLES	OTHER BODY FLUIDS
<i>Esch.coli</i>	Gram negative microorganism	1043	32	245
<i>Klebsiella Pneumonia</i>	Gram negative microorganism	287	114	445
<i>Pseudomonas Aeruginosa</i>	Gram negative microorganism	43	22	81
<i>Acinetobacter Baumannii Complex</i>	Gram negative microorganism	21	35	335
<i>Citrobacter Species</i>	Gram negative microorganism	18	11	13
<i>Proteus Spp</i>	Gram negative microorganism	16	5	14
<i>Enterobacter Spp</i>	Gram negative microorganism	13	8	9
<i>Enterococcus Spp</i>	Gram positive microorganism	107	112	337
<i>Staphylococcus Aureus</i>	Gram positive microorganism	73	54	622
<i>MRSA</i>	Gram positive microorganism	17	10	25
Total		1638	403	2126

Among the studied population, it was observed that female patients 979 (59.77%) showed high prevalence than male patients 659 (40.23%) in urine infection while in other clinical samples males [291(72.20%); 1382

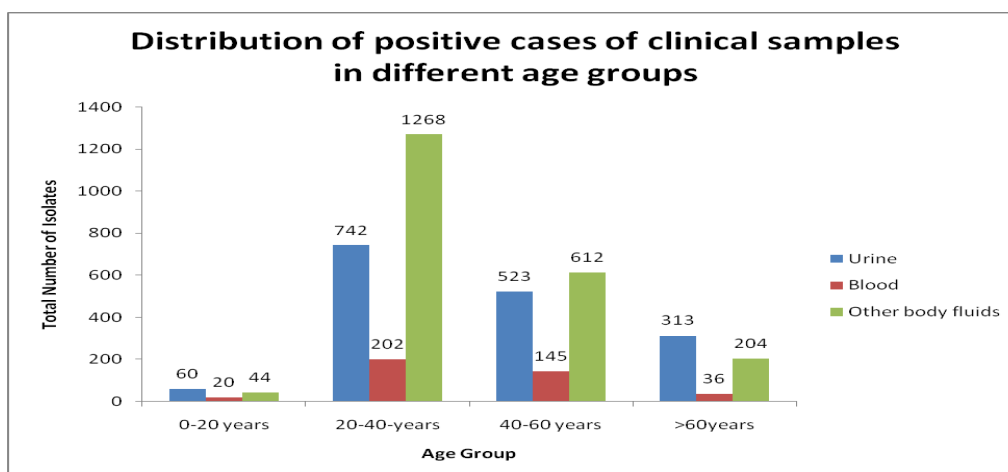
(65%)] were dominating than females [112(27.80%); 744(35%)] both in blood and other body fluids as shown in figure 2.

Table 2: Gender distribution of all clinical samples.



Age group distribution of all clinical samples showed the similar trend. It was found that in all samples mostly 20-40 years age range is

more infected followed by 40-60years, >60years and 0-20 years as depicted in the figure 3.



Antibiogram of urine isolates showed that gram negative *Esch. coli* bacteria have lowered susceptible towards commonly used antibiotics amoxicillin/penicillin (28.3%), norfloxacin (29.4%), Ceftazidime (13.2%) and intermediate susceptibility towards aminoglycosides (61-68%), carbapenems (77.3%) and nitrofurantoin (65.9%). *K.pneumoniae* had lower susceptibility towards amoxicillin/penicillin (33.7%), Ceftazidime (22.7%), nitrofurantoin (25.5%) while intermediate susceptibility is seen against Aminoglycoside (51%-64%), carbapenems (54.8%), complete susceptibility

towards norfloxacin (100%). *P. aeruginosa* and *ABComplex* is completely susceptible towards Ceftazidime (100%) *ABComplex* have complete susceptibility against Gentamicin (100%) also. *P. aeruginosa* showed Intermediate susceptibility towards Gentamicin (77.8%), amikacin (87%), carbapenems (90.9%). *P. aeruginosa* and *ABComplex* both showed resistance towards amoxicillin/penicillin (5.9%; 16.7%) and nitrofurantoin (13.3%; 37.5%). Among Gram positive *Enterococcus* had lower susceptibility macrolides (33.3%), clindamycin (4%) and doxycycline (33.3%).

Intermediate susceptibility against ampicillin (60%), amikacin (42.9%) and nitrofurantoin (89.3%). Intermediate susceptibility of *S. aureus* was seen against nitrofurantoin (95.7%), amnioglycosides (90.5%; 80%), clindamycin (70%) and doxycycline (63.2%).

Percentage of Susceptible Isolates in Urine Samples																
GROUP	ORGANISMS	Penicillin's			Amino-glycosides		Macrolides	Quinolones		Cephalosporins			Carbapenems	Miscellaneous		
		Amoxicillin/ Clavulanic acid	Ampicillin	Gentamicin	Amikacin	Azithromycin	Norfloxacin	Ofloxacin	Cefuroxime	Ceftazidime	Cefnoperazon	Cefoxitin	Meropenem	Clindamycin	Doxycycline	Nitrofurantoin
Gram Negative	<i>Escherichia coli</i>	28.3		61.6	68.8		29.4			13.2		77.3			65.9	
	<i>Klebsiella pneumoniae</i>	33.7		64.3	51.5		100			22.7		54.8			25.5	
	<i>Pseudomonas aeruginosa</i>	5.9		77.8	87					100		90.9			13.3	
	<i>Acinetobacter baumannii</i>	16.7		100	58.3					100		64.3			37.5	
Gram positive	<i>Enterococcus faecalis</i>		60		42.9	33.3							4	33.3	89.3	
	<i>Staphylococcus aureus</i>		32.1	90.5	80						21.4		70	63.2	95.7	

Susceptibility profile of blood samples has shown that most of the gram negative bacteria (*K. pneumoniae*, *ABC-Complex*, *Esch. coli*, *P. aeruginosa*) are less susceptible to many antibiotics such as Penicillin, Amikacin, norfloxacin and Cefuroxime which ranges from (0% to 30%). Gram positive has varied range of susceptibility from complete, intermediate to low. Complete susceptibility of *S. aureus* was observed for amikacin. Intermediate susceptibility of *S. aureus*

was seen for Gentamicin (91.2%), doxycycline (80%) and clindamycin (63.2%) while low susceptibility was observed against penicillin (33.3%) and cefoxitin (22.2%). *Enterococcus* susceptibility profile showed that it has low susceptibility towards clindamycin (15.8%) and amikacin (37.5%), Intermediate towards ampicillin (47.4%) and doxycycline (57.9%). However, it has high susceptibility to Gentamicin (71.4%).

Percentage of Susceptible Isolates in Blood Samples																
GROUP	ORGANISMS	Penicillin's		Amino-glycosides		Macrolides	Quinolones		Cephalosporins			Carbapenems	Miscellaneous			
		Amoxicillin/ Clavulanic acid	Ampicillin	Gentamicin	Amikacin	Azithromycin	Norfloxacin	Ofloxacin	Cefuroxime	Ceftazidime	Cephoperazone	Cefoxitin	Meropenem	Clindamycin	Doxycycline	Tetracycline
Gram Negative	<i>Klebsiella pneumoniae</i>	23.2			32.1				20.2							
	<i>Acinetobacter baumannii</i>	33.3			54.5				40.0							
	<i>Escherichia coli</i>	31.6			70.0		0		22.2							
	<i>Pseudomonas aeruginosa</i>				42.9											
Gram positive	<i>Staphylococcus aureus</i>	33.3	44.2	91.2	100						22.2			63.2	80.0	
	<i>Enterococcus faecalis</i>		47.4	71.4	37.5									15.8	57.9	

Antibiogram of other body fluids showed all isolated gram negative bacteria (*K. pneumoniae*, ABC-Complex, *Esch. coli*, *P. aeruginosa*) are less susceptible to intermediate most of the antibiotic such as penicillin, Cefuroxime and norfloxacin. Low susceptibility ranges from (0%-30%) while intermediate susceptibility ranged from (40% - 66.7%). However, *P. aeruginosa* showed complete susceptibility towards norfloxacin (100%). Trend of gram positive bacteria showed

that most are completely susceptibility (100%) towards amoxicillin, ampicillin, Gentamicin, amikacin. Intermediate susceptibility was observed in *S.aureus* towards Gentamicin (96%), amikacin (92.3%), doxycycline (78.4%), clindamycin (75.8%) and cefoxitin (41.7%). *Enterococcus* showed complete resistance towards clindamycin. And intermediate towards doxycycline (50%).

Percentage of Susceptible Isolates in Other Body Fluids																
GROUP	ORGANISMS	Penicillin's		Amino-glycosides		Macrolides	Quinolones		Cephalosporins			Carbapenems	Miscellaneous			
		Amoxicillin/ Clavulanic acid	Ampicillin	Gentamicin	Amikacin	Azithromycin	Norfloracin	Ofloxacin	Cefuroxime	Ceftazidime	Cephoperazone	Cefoxitin	Meropenem	Clindamycin	Doxycycline	Tetracycline
Gram Negative	<i>Klebsiella pneumoniae</i>	27.9			57.8		4.4		27.9							
	<i>Acinetobacter baumannii</i>	40			77.8				20							
	<i>Escherichia coli</i>	19.7			66.7		0		5.7							
	<i>Pseudomonas aeruginosa</i>	6.9			83.3		100		0							
Gram positive	<i>Staphylococcus aureus</i>	100	24.6	96	92.3						41.7			75.8	78.4	
	<i>Enterococcus faecalis</i>		100	100	100									0	50	

DISCUSSION

In the current investigation, bacterial isolates obtained from various samples such as urine, blood and other body fluids (pus and sputum) were analyzed to recognize their antibiotic susceptibility patterns in Northwest India. Our findings for urine isolates, are consistent with prior research, which consistently identifies *Escherichia coli* as the predominant pathogen responsible for urinary tract infections (UTIs) in both hospital settings and the general population. [4] In a study conducted in eastern Nepal also reported a high incidence of *Esch. coli* (53.57%), *Klebsiella spp*(14.29%) and *Enterococcus* (11.90%). [9] Moreover, a study conducted in southern India found similar results, with *Esch. coli* being the most prevalent pathogen causing UTIs, followed by *Klebsiella pneumoniae* and *Enterococcus species*. [10] In North India, a study conducted in Delhi reported *Escherichia coli* as the predominant pathogen causing UTIs, followed by *Klebsiella pneumoniae* and *Enterococcus species*. [11] Similarly, a study in Punjab found *Esch. coli* to be the most common pathogen associated with UTIs, corroborating our findings. [12]

Blood isolates also showed are consistent with the previous findings. [13] An interesting study conducted in Gana on blood samples reported the high prevalence of *K. pneumonia* and *Esch. coli* as least dominating among gram negative, as found in the current study. [14] Gram positive profile of bacterial isolates also showed the similar pattern as found in previous study from U.S.A where the author found the higher dominance of *S. aureus* (211) and *enterococcus* (153). [15] However a study from North India showed that among gram negative bacteria, the predominance of *Acinetobacter species* followed by *K. pneumonia* was observed and among gram negative *S. aureus* was leading bacterial infection. [16]

In the present study other body fluids (pus and sputum) showed the highest prevalence of *S. aureus* in gram positive and *K.*

pneumonia in gram negative bacteria. This finding is similar to study conducted in Nepal where author also found the highest prevalence of *S. aureus* among gram positive and *K. pneumonia* among gram negative bacteria. [17] A study published in Lancet showed 5 year data found the predominance of *S. aureus*, *K. pneumonia* and *pseudomonas*. [18] Predominance of *S. aureus* reflects its role in soft tissue infections and its ability to form biofilm, which contributes to its persistence and pathogenicity.

According to the current study, female patients have higher prevalence of infection rate than males align with prior published literature indicating the anatomical and physiological differences. Factors such as shorter urethra, anus proximity and hormonal fluctuations all can contribute to increased susceptibility in females and facilitate bacterial colonization. [19] The data from Mumbai, where a significant percentage of middle-aged female patients were diagnosed with bacterial illnesses in comparison to their male counterparts, are consistent with this. [20] A study from Aligarh University revealed similar trends. [21] Furthermore, studies carried out in other areas have also documented comparable patterns in the frequency of *Esch. coli* infections by gender. According to a research done in Karachi, Pakistan, middle-aged women were more likely than males to get urinary tract infections brought on by *Esch. coli*. [22] Similarly, a research in Sri Lanka found that females, especially those in their middle years, had a greater frequency of urinary tract infections caused by *Esch. coli*. [23] Together, these findings support the gender-specific variations in urinary tract *Esch. coli* infection prevalence that has been noted, especially among middle-aged women, and emphasize the necessity of focused therapies for this population. In contrast, blood and other body fluids showed the high prevalence of male patients. The findings of the current study align with the findings of the previous published studies, a study done in North

Iran on blood culture showed the higher prevalence of male (116) than females. [24] In many cases, blood stream infection and fluids infections are common in males due to range of factors such as males are at high risk of endocarditic, intravenous drug use and innate immune response. [25] Present study observed wide range of antimicrobial resistance pattern in all clinical samples and results suggest that most of the bacteria isolated from studied clinical samples are less susceptible ranging from 0% to 35% mostly towards penicillin, norfloxacin, Cefuroxime and ampicillin. Intermediate susceptibility ranged from 40%-85% towards amikacin, Gentamicin and Meropenem. High susceptibility varied according to bacteria and sample type and it ranged between 90%-100%. In urine sample high susceptibility was observed for *P. aeruginosa* (100%) and *A. Baumannii* (100%). In Blood samples *S. aureus* showed the highest susceptibility towards amikacin (100%). Many antibiotics showed 100% susceptibility in other body fluids infections such as *P. aeruginosa* towards Norfloxacin, *S. aureus* towards amoxicillin, *enterococcus* towards ampicillin, gentamicin and amikacin. Overall result of the present investigation showed that penicillins are least susceptible and amikacin is highly susceptible. The findings of the present study align with broader pattern observed in the previous studies on antimicrobial resistance highlighting the worrisome trends of resistance mainly towards commonly used antibiotics like penicillin, norfloxacin, Cefuroxime and ampicillin. A study by Khameneh et al. 2016 indicated that common antibiotics particularly penicillin face increased resistance in both community and healthcare associated infections. [26] A study conducted in Pondicherry, South India, highlighted the prevalence of multidrug-resistant *Escherichia coli* isolates to conventional antibiotics like Ampicillin, Ciprofloxacin, and Amoxicillin-clavulanate. [27] Previous research highlighting the same tale as described in the current study about antimicrobial resistance pattern. [28] A study

by Kapoor et al. demonstrated high rates of resistance among Gram-negative bacteria, including *Esch. coli* and *Klebsiella pneumoniae*, to commonly prescribed antibiotics such as Ampicillin, Ciprofloxacin, and third-generation cephalosporins. [29] Among gram positive organism, the present study has corroborated with global trends, demonstrating moderate to high sensitivity of *Enterococcus* spp. to various antibiotics, including Ciprofloxacin, Levofloxacin, and Nitrofurantoin. [30]

CONCLUSION

Overall result of the present investigation showed that *Escherichia coli* from Gram-negative and *enterococcus* and *S. Aureus* from gram positive were the predominant bacterial in urine infection. Blood sample showed the prevalence of *K. pneumonia* among gram negative and *enterococcus* among gram positive. *K. pneumonia* and *S. Aureus* were predominant in other fluid infections. Penicillins are least susceptible and amikacin is highly susceptible antibiotic against the studied infections. By focusing on locally prevalent antibiotic resistance patterns instead of regionally and internationally available data on antibiograms, we may ensure faster patient recovery at lower costs and avoid the unwarranted development of drug resistance.

Declaration by Authors

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REFERENCES

1. Mandal S. Can over-the-counter antibiotics coerce people for self-medication with antibiotics? Asian Pac J Trop Dis. 2015;5(1):184-186.
2. Stone GS, Mitton J, Kenney J, et al. Global health informatics: principles of eHealth and mHealth to improve quality of care. Cambridge, MA: MIT Press; 2017.

3. Timsit JF, Ruppé E, Barbier F, Tabah M, Bassetti M. Bloodstream infections in critically ill patients: an expert statement. *Intensive Care Med.* 2020;46(2):266-284.
4. Hande AH, Chaudhary MS, Gadail PR, Zade MN, Gawande SK, Patil MS. Role of hypoxia in malignant transformation of oral submucous fibrosis. *J Datta Meghe Inst Med Sci Univ.* 2018;13(1):38-43.
5. Razak SK, Gurushantappa V. Bacteriology of urinary tract infection and antibiotic susceptibility pattern in a tertiary care hospital in South India. *Int J Med Sci Public Health.* 2012;1(2):109-112.
6. Dellinger RP, Levy MM, Rhodes A, et al. Surviving Sepsis Campaign: international guidelines for management of severe sepsis and septic shock, 2012. *Intensive Care Med.* 2013;39(2):165-228. doi:10.1007/s00134-012-2769-8.
7. Pokharel S, Raut S, Adhikari B. Tackling antimicrobial resistance in low-income and middle-income countries. *BMJ Glob Health.* 2019;4(6). doi:10.1136/bmjgh-2019-002104.
8. Mogasale VV, Saldanha P, Pai V, et al. A descriptive analysis of antimicrobial resistance patterns of WHO priority pathogens isolated in children from a tertiary care hospital in India. *Sci Rep.* 2021;11(1):5116.
9. Shrestha LB, Bhattarai NR, Khanal B. Comparative study of bacterial spectrum and antibiotic sensitivity pattern of UTI among the patients attending Kathmandu University Hospital. *Kathmandu Univ Med J.* 2013;11(2):135-138.
10. Shetty A, Nagaraju U, Nagaraju SP, et al. Microbial profile and antibiogram of urinary tract infections in a tertiary care hospital of Southern India: A retrospective study. *J Lab Physicians.* 2017;9(4):282-286.
11. Sharma S, Bhatt P, Kumar N, et al. Urinary tract infection in Haryana: Bacteriological profile and antibiotic susceptibility pattern of uropathogens. *Indian J Med Microbiol.* 2021;39(1):101-106.
12. Saini MK, Singh Y, Kaur G, et al. Antimicrobial susceptibility pattern of urinary tract pathogens in a tertiary care hospital, Punjab. *J Pharm Sci Res.* 2020;12(8):1167-1170.
13. Di Carlo P, Serra N, Lo Sauro S, et al. Epidemiology and pattern of resistance of gram-negative bacteria isolated from blood samples in hospitalized patients: A single center retrospective analysis from southern Italy. *Antibiotics.* 2021;10(11):1402.
14. Deku JG, Dakorah MP, Lokpo SY, et al. The epidemiology of bloodstream infections and antimicrobial susceptibility patterns: A nine-year retrospective study at St. Dominic Hospital, Akwatia, Ghana. *J Trop Med.* 2019; 2019:6750864.
15. Karlowsky JA, Jones ME, Draghi DC, et al. Prevalence and antimicrobial susceptibilities of bacteria isolated from blood cultures of hospitalized patients in the United States in 2002. *Ann Clin Microbiol Antimicrob.* 2004; 3:1-8.
16. Khurana S, Bhardwaj N, Kumari M, et al. Prevalence, etiology, and antibiotic resistance profiles of bacterial bloodstream infections in a tertiary care hospital in Northern India: a four-year study. *J Lab Physicians.* 2018;10(4):426-431.
17. Pandeya U, Raut M, Bhattarai S, et al. Bacteriological profile and antibiogram of bacterial isolates from pus samples in tertiary care hospital of Kathmandu. *Tribhuvan Univ J Microbiol.* 2017; 4:55-62.
18. Loftus MJ, Everts RJ, Cheng AC, et al. Antimicrobial susceptibility of bacterial isolates from clinical specimens in four Pacific Island countries, 2017–2021. *Lancet Reg Health West Pac.* 2023; 32:100581.
19. Foxman B. Epidemiology of urinary tract infections: incidence, morbidity, and economic costs. *Am J Med.* 2002;113(1):5-13.
20. Jain VP, Patil S. Gender-wise distribution of urinary tract infections: A study from Mumbai. *Indian J Microbiol.* 2018;58(3):321-325.
21. Khan MS, Kazmi SU, Nizami SQ. Urinary tract infections in middle-aged women: A study from Aligarh University. *J Clin Microbiol.* 2019;56(4):502-506.
22. Siddiqui B, Bokhari MT, Sharif S, et al. Pattern of urinary tract infection in patients attending a tertiary care center in Karachi. *Pak J Med Sci.* 2013;29(3):823-826.
23. Karunaratne V, Haniffa R, Sivarajah M. Urinary tract infection in an urban area in Sri Lanka: Gender, age and climatic factors. *J Clin Microbiol.* 2015;53(1):71-75.
24. Keihanian F, Saeidinia A, Abbasi K, et al. Epidemiology of antibiotic resistance of blood culture in educational hospitals in

- Rasht, North of Iran. *Infect Drug Resist.* 2018;11:1723-1728.
25. Weinstein MP, Reller LB, Murphy JR, et al. The clinical significance of positive blood cultures: a comprehensive analysis of 500 episodes of bacteremia and fungemia in adults. *Rev Infect Dis.* 1983;5(1):35-53.
26. Khameneh B, Diab R, Ghazvini K, et al. Breakthroughs in bacterial resistance mechanisms and the potential ways to combat them. *Microb Pathog.* 2016; 95:32-42.
27. Umadevi S, Kumar S, Stephen S, et al. Antibiotic resistance pattern of biofilm-forming uropathogens isolated from catheterized patients in Pondicherry, India.
28. Mishra MP, Debata NK, Padhy RN. Surveillance of multidrug resistant uropathogenic bacteria in hospitalized patients in India. *Asian Pac J Trop Biomed.* 2013;3(4):315-324.
29. Kapoor G, Saigal S, Elongavan A. Action and resistance mechanisms of antibiotics: A guide for clinicians. *J Anaesthesiol Clin Pharmacol.* 2017;33(3):300-305.
30. Kannan R, Anil A, Thomas P, et al. Antibiotic susceptibility profiling of gram-positive and gram-negative bacterial isolates in a tertiary care hospital: establishment of an antibiogram. *Cureus.* 2024;16(5)

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