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Biofilm formation and antibiotic resistance of urinary catheter associated bacteria from hospitalized patients, Bangladesh

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ABSTRACT

Aims: Biofilm formation of bacteria inside the surface of urinary catheters triggers severe urinary tract infections (UTIs). This study aims to determine the biofilm forming capacity of bacteria isolated from urinary catheters of patients diagnosed with UTIs as well as comparison of antibiotic sensitivity patterns between biofilm and non-biofilm forming isolates.

Methodology and results: A total of 40 urinary catheters were collected from 96 h catheterized patients. The isolated uropathogenic bacteria were identified and examined for biofilm formation using the microtiter plate method. Later, the isolates were subjected to antimicrobial susceptibility towards 12 antibiotics that commonly used for treating UTIs using the disk diffusion method. All the catheters were found colonized with two to five different bacterial species individually. Out of the 131 isolates from 40 catheters, *Pseudomonas aeruginosa* (38/131, 29%) was the predominant isolated bacteria followed by *Escherichia coli* (31/131, 24%), *Proteus vulgaris* (24/131, 18%), *Klebsiella pneumoniae* (21/131, 16%) and *Staphylococcus aureus* (17/131, 13%). Among these, the highest biofilm forming capacity was observed in *P. aeruginosa* (26%), followed by *P. vulgaris* (16%) and *K. pneumoniae* (13%). Regarding antibiotic resistance, biofilm forming bacteria showed resistance to multiple drugs except for carbapenems. Moreover, biofilm formers exhibited higher resistance than non-biofilm formers against antibiotics such as trimethoprim/sulfamethoxazole (100% vs 82%), amoxicillin-clavulanic acid (81% vs 55%), cefixime (85% vs 55%), ceftriaxone (81% vs 45%), cefalexin (93% vs 55%), amikacin (70% vs. 45%), ampicillin (89% vs 73%), ciprofloxacin (70% vs. 36%) and ceftriaxone (81 vs 45%), (*p*-value<0.05).

Conclusion, significance and impact of study: Most of the isolated uropathogenic bacteria from catheters were biofilm formers and multiple antibiotic resistant. Appropriate selection of antibiotics, meticulous hygiene practices in hospital settings and limiting the duration of catheterization can reduce biofilm formation and the emergence of antibiotic resistance.

Keywords: Urinary catheter, biofilm, antibiotic resistance, biofilm formers, non biofilm formers

INTRODUCTION

One of the leading causes of morbidity and mortality throughout the world is infections associated with hospital set up commonly known as hospital-associated infections (HAIs) which includes surgical site infections, bloodstream infections, urinary tract infections (UTIs), respiratory infections, gastroenteritis, pneumonia and meningitis and other soft tissue infections (Nicolle, 2012; Feleke et al., 2018; Haque et al., 2018). It was reported that about 40% of all HAIs are UTIs with a significant rate of morbidity and mortality (Hague et al., 2018). This percentage is also prominent in developing countries like Ethiopia (68.71%) (Ali et al., 2018), Nigeria (43%) (Ige et *al.*, 2011; Iliyasu *et al.*, 2018) as well as in Bangladesh (15.4%) (Afroz *et al.*, 2017; Alam *et al.*, 2019).

Several risk factors like age, pregnancy, diabetics, neurogenic bladder, menopause, frequent intercourse, change of the local bacterial flora, history of UTIs during pre-menopause or in childhood, family history have been made known to trigger UTIs (Storme *et al.*, 2019). On top of this, catheterization is also a leading source of UTIs and comprises around 80% of all hospital-associated UTIs (Jacobsen *et al.*, 2008; Sabir *et al.*, 2017). Catheters are standard medical devices that commonly used in hospitals to retain urine and eliminate urinary irregularities from patients. The predominant pathogens associated with catheter associated urinary

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tract infections (CAUTIs) include Escherichia coli, Enterococci, Serratia spp., Providencia stuartii, Pseudomonas aeruginosa, Klebsiella pneumoniae, Candida albicans, Enterobacter sp., Proteus mirabilis, and Morganella morganii coagulase-negative Staphylococci (Nicolle, 2012; Sabir et al., 2017; Maharjan et al., 2018). These pathogens can become a threat to treatment failure because of the emergence of multidrug resistance. The heavy use of wide-spectrum antibiotics to any infections has puts a high selective pressure that triggers the emergence of multidrug resistant bacteria (Soltani et al., 2016). This situation becomes jeopardizes when these resistant bacteria form biofilms and cause persistent infections (Almalki and Varghese, 2020). Biofilm, a layer of microorganism that composed of a complex assembly of bacterial protein, polysaccharide and DNA in an extracellular polymeric matrix and can be found on various surfaces, including, natural aquatic or potable water systems, living tissues and medical devices (O'Toole et al., 2000; Awoke et al., 2019). The surface of the urinary catheter also provides opportunities for microorganisms to form biofilms. Biofilm helps bacteria to survive, spread and resist antimicrobial therapy (Donlan, 2002). Several studies mentioned that biofilm-formers exhibited 10-1000 times more resistance to antimicrobials compare to non-formers (Potera, 2010; Sharma et al., 2019).

This study aimed to assess the distribution of catheterized bacterial pathogens, their biofilm-forming capacity as well as the comparison of antibiotic sensitivity patterns between biofilm and non-biofilm-forming isolates.

MATERIALS AND METHODS

Collection of samples

A total of 40 urinary catheters were collected randomly from 96 h catheterized postoperative female patients diagnosed with urinary tract infections and preserved in sterile polybag at 4 °C for 6 h. Samples were collected from two hospitals of Chittagong, Chittagong Medical College Hospital (CMCH) and University of Science and Technology Hospital, Chittagong (USTC), Bangladesh.

Catheters were cut into pieces of 1 cm in length, rinsed with sterile phosphate-buffered saline (PBS, pH 7.4). Then, 1 mL of the PBS was transferred into 5 mL of Brain Heart Infusion broth (BHIB) (Himedia M210, India) and incubated at 37 °C for 24 h.

Isolation and identification of bacteria

One loopful suspension from BHIB for each catheter sample was streaked on different selective media like Cetrimide agar (Oxoid[™] CM0579, UK), MacConkey agar (Oxoid[™] CM0007, UK), Casein Enzyme Hydrolysate (CLED) agar (Oxoid[™] CM0301, UK) and Mannitol Salt Agar (Himedia[™] M118, India) for selective isolation of *P. aeruginosa, E. coli, K. pneumoniae, P. vulgaris* and *S. aureus.* Then, the plates were incubated at 37 °C for 24 h. The individual isolates obtained from the selective media plates were presumptively identified using conventional microbiological methods including both culture-based methods and biochemical tests, respectively according to Bergey's Manual, 9th Edn. (Holt *et al.*, 1994).

Quantitative detection of biofilm formation

Stepanović et al. (2007) method was used for quantitative analysis of biofilm formation. To conduct the test, each bacterial isolate was inoculated into 10 mL of trypticase soy broth (Himedia LQ508) in addition with 1% glucose and incubated at 37 °C for 24 h. After that each culture was diluted for 100 times using fresh trypticase soy broth medium at room temperature. Each diluted culture was transferred to individual wells of sterile 96-well flat-bottom polystyrene tissue culture-treated plates (Sigma Aldrich, Costar, USA) in the amount of 125 µL and incubated at 37 °C for 24 h. The negative control wells contained same amount of fresh trypticase soy broth medium. After incubation, the contents of each well were gently tapped out. Each well was washed four times with 0.2 mL of phosphate buffer saline (PBS) (pH 7.2) to remove the free-floating bacteria from the wells before fixation. Finally, the wells were treated with sodium acetate (2%) for cell fixation. After fixation, the plate with sodium acetate was washed with 0.2 mL PBS thrice before staining. For staining, 125 µL of 0.1% crystal violet was added for five min. The excess crystal violate solution was removed and the plate was rinsed 3-4 times with PBS. For elution, 125 µL ethanol-acetone mixture (80:20) was used and left at room temperature for 30 min. The optical density (OD₆₃₀) of the stained adherent biofilm was determined by using a micro ELISA auto reader (Model 680, Biorad, UK) and analyzed for calculating the biofilmforming capacity of the bacterial isolates. The experiment was performed three times in triplicates and the results were interpreted according to the criteria as described by Stepanović et al. (2007).

In brief, OD₆₃₀ values were calculated as OD₆₃₀ = Ø OD – ODc, where Ø OD indicates optical density of each well containing the bacterial isolate and ODc indicates the cut-off value which was calculated using the following formula: ODc = Ø ODn + 3 × SDn. Here, Ø ODn indicates the mean of negative control and SDn indicates the standard deviation of the negative control. The isolates were categorized based on the OD₆₃₀ values:

 $OD_{630} \le ODc$: No biofilm $ODc < OD_{630} \le 2x ODc$: Weak biofilm $2x ODc \le OD_{630} \le 4x ODc$: Moderate biofilm $4x ODc < OD_{630}$: Strong biofilm

Antibiotic susceptibility patterns

Antibiotic susceptibility pattern of all isolated bacteria was screened using the disc diffusion method, towards Ampicillin, AMP (10 μ g), Amoxicillin/Clavulinic acid, AMC (30 μ g), Amikacin, AK (30 μ g), Gentamicin, CN (30 μ g), Erythromicin, E (15 μ g), Cephalexin, CL (30 μ g),

Cefixime, CFM (5 μ g), Ceftriaxone, CRO (30 μ g), Ciprofloxacin, CIP (5 μ g), Trimethoprim/sulfamethoxazole, SXT (1.25/23.75 μ g), Imipenem, IPM (10 μ g) and Meropenem, MEM (10 μ g). The inhibition zone diameter against tested antibiotics was measured in mm and interpreted based on the Clinical and Laboratory Standards Institute (CLSI) guidelines (CLSI, 2017). *Pseudomonas* spp. ATCC 27853 was used as control strains in this study.

Statistical analysis

Data were analyzed using Statistical Package for Social Sciences (SPSS) software version 22 (SPSS Inc, Chicago, IL, USA). The chi-squared test, odds ratio (risk) with 95% CI, was used to calculate possible associated factors between the resistance of biofilm formers and non-biofilm formers. All statistical test values of p<0.05 were considered statistically significant.

RESULTS

Catheter colonization and distribution of uropathogens

All the urinary catheter samples collected were found to be culture-positive with two to five mixed bacterial species. The number of catheters with mixed colonization is summarized in Figure 1, which most of the catheters (n=18) show co-colonized with 3 species and the least number of catheters (n=3) shows co-colonized with 5 species. From 40 urinary catheters, a total of 131 bacterial isolates were isolated using selective media. Out of the 131 identified isolates, the most predominant bacteria were *P. aeruginosa* (38/131, 29%) followed by *E. coli* (31/131, 24%), *P. vulgaris* (24/131, 18%), *K. pneumoniae* (21/131, 16%) and *S. aureus* (17/131, 13%).

Assessment of biofilm formation ability of isolates

Next, the biofilm formation capability of these isolates was examined using the microtiter plate method. Out of 131 isolates, 71% (n=93) of them showed biofilm forming capacity. Among these, 87% (81/93) exhibited weak to moderate biofilm formation and 12.9% (12/93) exhibited strong biofilm formation (Table 1). If we linked the catheter colonization and biofilm forming ability, all the isolates found in catheter co-colonized with four to five different bacterial species were capable of forming biofilms. Among the 18 catheters that were co-colonized with three bacterial species, ten of them were colonized with biofilm forming bacteria. Among all the biofilm forming bacteria, P. aeruginosa (89.5%) was the most prevalent biofilm-former, followed by P. vulgaris (87.5%) and K. pneumoniae (85.7%) (Table 1). The optical density (OD₆₃₀) of all 93 biofilm formers is presented in Supplementary Table S1.

Antibiotic susceptibility test of biofilm forming isolates

In this study, 12 antibiotics were used to evaluate antibiotic sensitivity of catheter associated biofilm forming bacteria. All the biofilm-formers showed resistance to multiple antibiotics except for both carbapenems (meropenem and imipenem). The antibiotic pattern of each type of biofilm producer was recorded in Table 2. All P. vulgaris isolates showed pan resistance to 10 treated antibiotics. Other bacterial isolates such as P. aeruginosa have also shown pan resistance to ampicillin, cephalexin, cefixime and trimethoprim/sulfamethoxazole. Apart from these two, all K pneumoniae isolates were also resistant to ampicillin, amoxicillin/clavulanic acid, erythromycin and trimethoprim/sulfamethoxazole (Table 2). Gram-positive isolates S. aureus showed less resistance than all Gramnegative isolates. However, all the isolates of S. aureus showed resistance to cefixime and pan trimethoprim/sulfamethoxazole.

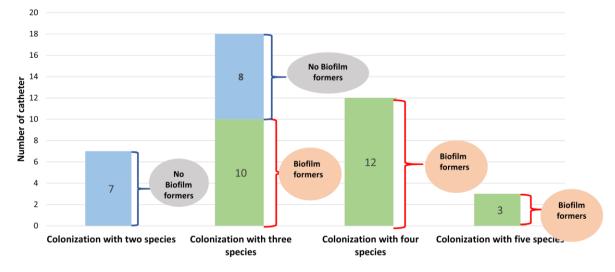


Figure 1: Number of catheters colonization by multiple species and biofilm formers.

Bacteria				Biofilm f	ormation*				
		Foi	mers (n=93) (71%	Non-formers (n=38) (29%)					
	Strong	Moderate	Weak	n	%	n	%	Total	
P. aeruginosa	6	9	19	34	89.5	4	10.5	38	
E. coli	2	1	7	10	32.3	21	67.7	31	
P. vulgaris	3	8	10	21	87.5	3	12.5	24	
K. pneumoniae	1	4	13	18	85.7	3	14.3	21	
S. aureus	0	2	8	10	58.8	7	41.2	17	
Total (%)	12	24	57	93					
	(12.9%)	(25.8%)	(61.3%)						

Table 1: Biofilm formation capacity of isolated to pathogen (n=131) using microtiter plate method.

*The isolates were categorized based on the OD₆₃₀ values (Stepanović et al., 2007).

Table 2: Antibiotic resistance patterns of the identified biofilm forming bacteria from urinary cathe

Isolates	Α	MP	A	ИС	A	K	C	N		E	(Ľ	С	FM	С	RO	С	IP	S	XT	IP	М	M	ΞM
	S	R	S	R	S	R	S	R	S	R	S	R	S	R	S	R	S	R	S	R	S	R	S	R
P. aeruginosa (n=34)	0	34	3	31	21	13	14	20	3	31	0	34	0	34	3	31	11	23	0	34	34	0	34	0
<i>E. coli</i> (n=10)	4	6	7	3	0	10	0	10	1	9	0	10	7	3	5	5	0	10	0	10	10	0	10	0
<i>P. vulgaris</i> (n=21)	0	21	0	21	0	21	0	21	0	21	0	21	0	21	0	21	0	21	0	21	21	0	21	0
K. pneumoniae (n=18)	0	18	0	18	7	11	11	7	0	18	1	17	7	11	4	14	11	7	0	18	18	0	18	0
S. aureus (n=10)	6	4	8	2	0	10	3	7	3	7	6	4	0	10	6	4	6	4	0	10	10	0	10	0

"S" denotes sensitive and "R" denotes resistant. The results are interpreted based on the inhibition zone diameter against tested antibiotics according to the Clinical and Laboratory Standards Institute (CLSI) guidelines (CLSI, 2017). The tested antibiotics are Ampicillin, AMP (10 μg), Amoxicillin/Clavulinic acid, AMC (30 μg), Amikacin, AK (30 μg), Gentamicin, CN (30 μg), Erythromicin, E (15 μg), Cephalexin, CL (30 μg), Cefixime, CFM (5 μg), Ceftriaxone, CRO (30 μg), Ciprofloxacin, CIP (5 μg), Trimethoprim/sulfamethoxazole, SXT (1.25/23.75 μg), Imipenem, IPM (10 μg) and Meropenem, MEM (10 μg).

Comparison of antibiotic sensitivity among biofilm-formers and non-biofilm-formers

When biofilm-forming uropathogenic bacteria were compared with nonbiofilm-formers, the biofilm-formers showed significantly higher antibiotic resistance, trimethoprim/sulfamethoxazole (100% vs 82%), amoxicillinclavulanic acid (81 vs 55%), cefixime (85% vs 55%), ceftriaxone (81% vs 45%), cephalexin (93% vs 55%), amikacin (70% vs 45%), ampicillin (89% vs 73%) and ciprofloxacin (70% vs 36%), with a *p*-value of less than 0.05. However, in the remaining antibiotics tested including erythromycin and gentamicin, there was no significant difference (p>0.05) in their action among biofilm and non-biofilm-formers (Table 3).

DISCUSSION

Catheter associated urinary tract infection is becoming one of the major hospital-associated infections (HAIs) globally. Longer duration of the urinary catheter in patients allows habituating bacteria in the device to form biofilms which eventually find their way to the urinary tract system and leads to infections (Donlan, 2002; Dougnon *et al.*, 2016; Awoke *et al.*, 2019). This

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Table 3: Comparison of antibiotic resistance pattern of biofilm-forming and non-biofilm forming bacteria.

Antibiotic used	Resistance by	Resistance by	Odd ratio	95 % CI	P value	X ²
	biofilm-formers	non-biofilm formers				
	(n) (%)	(n) (%)				
SXT	93 (100)	31 (82)	0.296	0.130-0.674	0.003*	8.887
E	86 (93)	35 (91)	0.950	0.232-3.883	0.943	0.005
CL	86 (93)	21 (55)	0.101	0.037-0.274	0.000*	24.961
AMP	83 (89)	28 (73)	0.337	0.127-0.895	0.025*	5.051
CFM	79 (85)	21 (55)	0.219	0.09-0.515	0.000*	13.158
CRO	75 (81)	17 (45)	0.194	0.086-0.441	0.000*	16.637
AMC	75 (81)	21 (55)	1.226	1.054-1.426	0.000*	18.099
AK	65 (70)	17 (45)	0.349	0.160-0.759	0.007*	7.291
CN	65 (70)	24(63)	0.738	0.334-1.634	0.454	0.562
CIP	65 (70)	14 (36)	0.251	0.114-0.556	0.000*	12.310
IPM**	0 (0)	0 (0)				
MEM**	0 (0)	0 (0)				

*Biofilm-forming bacteria are more antimicrobial resistant compared with non-biofilm-forming ones (p≤0.05).

** Chi square tests not performed for "0" value.

Ampicillin, AMP (10 μg), Amoxicillin/Clavulinic acid, AMC (30 μg), Amikacin, AK (30 μg), Gentamicin, CN (30 μg), Erythromicin, E (15 μg), Cephalexin, CL (30 μg), Cefixime, CFM (5 μg), Ceftriaxone, CRO (30 μg), Ciprofloxacin, CIP (5 μg), Trimethoprim/sulfamethoxazole, SXT (1.25/23.75 μg), Imipenem, IPM (10 μg) and Meropenem, MEM (10 μg).

study provides data to quantify the biofilm forming ability of the isolated bacteria from urinary catheters and checked the status of the antibiotic resistance pattern of these biofilm formers.

All the catheters collected in this study were found culture positive with multiple bacterial species. The cocolonization by three different bacterial species in a single catheter was the highest. This type of multiple colonization in a single catheter was also recorded in previous studies (Macleod and Stickler, 2007; Verma *et al.*, 2016). We also observed the mixed bacterial community up to five species which also exhibited in another few studies (Kolter and Losick, 1998; Macleod and Stickler, 2007).

Among all colonized catheters, Gram-negative bacteria *P. aeruginosa* (29%), *E. coli* (24%), *K. pneumoniae* (16%) and Gram-positive *S. aureus* (16%) were predominant in this study. The high prevalence of *K. pneumoniae*, *E. coli*, *P. mirabilis*, *S. aureus* and *P. aeruginosa* were also reported from UTIs and CAUTIs by several studies from Ethiopia (Feleke *et al.*, 2018; Awoke *et al.*, 2019), Canada (Karlowsky *et al.*, 2011), Pakistan (Sabir *et al.*, 2017) and Saudia Arabia (Almalki and Varghese, 2020). We also observed the biofilm-forming capability of isolated bacteria quantitatively and 71% (93 out of 131) isolates exhibited biofilm-forming capacity *in vitro*.

The biofilm-forming capacity of these bacteria is similar or stronger than the isolates from previous studies. Awoke *et al.* (2019) reported that 80% of the total study isolates recovered from urinary catheterized inpatients were biofilm-formers where Sabir *et al.* (2017) also reported 73.4% (785/1070) of study isolates as biofilm-formers isolated from CAUTIs. Similar to our study, Murugan *et al.* (2016) also reported strong biofilm production by *S. aureus, E. faecalis* and *P. aeruginosa.*

Moreover, we reported biofilm-forming capacity increased with multiple co-colonization as the bacteria that cocolonized with three or four mixed-species showed higher biofilm-forming capacity compared to two mixed-species. Cooperation among the mixed species was found to be increased in biofilm formation as all the co-colonizing species can help each other. Poor biofilm formers get benefited from this mutualistic relationship by utilizing the nutrients produced by one species and promote their growth inside the biofilm matrix (Rao *et al.*, 2020; Yuan *et al.*, 2020). Besides, it had been reported that co-culturing of *Lactococcus lactis* and *P. fluorescens* increased bacterial adhesion by up to 20,000- and 100-fold, respectively (Kives *et al.*, 2005).

Given to the complex microbial interactions, a mixedspecies biofilm often achieves substantially more biomass than a monospecies biofilm without the need for extra nutrients such as in case of *Stenotrophomonas rhizophila*, *Xanthomonas retroflexus*, *Microbacterium oxydans* and *Paenibacillus amylolyticus* (Ren *et al.*, 2015). Similarly, previous studies also revealed the presence of mixed bacterial species in biofilm crystals observed from catheters (Kolter and Losick, 1998). Mixed community of *P. mirabilis*, *P. aeruginosa*, *K. pneumoniae* and few other Gram-negative bacteria in crystalline biofilm on urinary catheter and their interaction in the biofilm also observed by other studies (Macleod and Stickler, 2007; Balasubramanian *et al.*, 2012).

Biofilm-producing bacteria is more resistant to antibiotics than the bacteria that do not form biofilms (Hashemzadeh *et al.*, 2020; Luo *et al.*, 2021). All the biofilm-forming isolates in our study were resistant to multiple antibiotics except for both carbapenems (meropenem and imipenem). These findings correlated with various other studies where multidrug-resistant pathogens were also reported (Nicolle, 2012; Dougnon *et*

al., 2016; Awoke *et al.*, 2019). According to Prakash *et al.* (2013), most of the uropathogens were susceptible to carbapenem-group antibiotics imipenem and meropenem. This susceptibility was reported to be 92.26% for imipenem and 84.52% for meropenem (Prakash *et al.* 2013).

Among Gram-negative biofilm-forming bacteria, the most resistant pathogens in our study were *P. vulgaris* and *P. aeruginosa*, which showed maximum resistance to multiple antibiotics. This resistance pattern indicated that current antibiotics used to treat acute urinary tract infections seem ineffective in the eradication of isolated uropathogens. One of the possible reasons for high antibiotic resistance in biofilm formers can be the effect of selective antibiotic pressure on isolates. As all the isolates were collected from hospitalized postoperative patients with UTIs and the practice of frequent antibiotic pressure on isolates leading to the acquisition of antibiotic resistance by nosocomial isolates (Kolář *et al.*, 2001).

Moreover, biofilm formers can also acquire drug resistance by delayed penetration through inactivation of antimicrobial molecules or limiting the diffusion of antimicrobial molecules through the polymer matrix of biofilm, decreasing antimicrobial uptake by slower the growth rate of biofilm associated cells, enzymatic transformation of biocidal agents, genetic adaptation and producing resistance phenotype and exchange of extrachromosomal DNA (Ehlers and Bouwer, 1999; Hausner and Wuertz, 1999; Roberts et al., 1999; Donlan, 2002). It has been reported that when the planktonic bacterial cells form biofilm, they became resistant to antibiotics and to treat such infections, the concentration of antibiotics required to increase three- to four-times higher to achieve the bactericidal level depending on the species and drug combination (Gurung et al., 2013).

In our study, the highest resistance was observed among biofilm-formers with trimethoprim/ sulfamethoxazole (100%), followed by erythromycin and cephalexin (93%), ampicillin (89%) and cefixime (85%). A study from Bangladesh reported the highest resistance against trimethoprim/ sulfamethoxazole, amoxicillin/ clavulanic acid, ceftazidime, ceftriaxone and cefixime by E. coli and P. aeruginosa (Hossain et al., 2014). However, this study did not specify the biofilm-forming activities of the isolates. Karigoudar et al. (2019) also reported high resistance by biofilm-former against trimethoprim/ sulfamethoxazole. Sabir et al. (2017) also showed the highest resistance of ampicillin followed by ciprofloxacin amond biofilm-formers. As trimethoprim/ sulfamethoxazole is one of the most frequently used antibiotics for UTI, unjudicial and overuse of the drugs may raise the resistance (Alam et al., 2019).

The study data concluded that apart from both carbapenems, all the drugs exhibited less sensitivity against biofilm-formers. However, ciprofloxacin (36%), ceftriaxone (45%) and amikacin (45%) were effective against non-biofilm-formers. Excessive and inappropriate way of using antibiotics in hospital settings are the major factors which lead to the development of such antibiotic

resistant bacteria. Furthermore, the biofilm-forming capability of these bacteria on the surface of medical devices (such as indwelling vascular catheters, cardiac pacemakers, prosthetic heart valves) that are installed for patient care increased the threat as biofilms protect the uropathogens from environmental stresses and it also leads to decreased susceptibility of the colonizing bacteria to the antibiotics. This will be more hazardous if infection prevention practices are not followed during patient care. Routine cleaning of the urethral meatus surface, using a small size catheter with good urinary drainage flow, using sterile and closed unobstructed urinary drainage, and cleaning the catheter and collecting system junction with chlorhexidine gluconate, a povidoneiodine solution, or a 70% isopropyl alcohol solution are effective infection prevention practices to reduce urinary catheter-related infections (Assadi, 2018).

There are some limitations in our study, such as the catheters were collected randomly without linking patients' profiles and hospital stays. We also acknowledged that the small sample size and absence of clinical records describing the types and severity of the urinary tract infections and profile of patients providing urinary catheters. However, our study provides data on antibiotic resistance patterns among catheter-associated biofilm-forming bacteria in two hospital settings of Bangladesh which prioritized the management guidelines for antibiotic use against such infections.

CONCLUSION

The diverse resistance pattern of uropathogenic bacteria highlights the importance of studying the pattern of urinary infection in every setting and the rational use of antibiotics in the management of such infections. In addition, rigorous hygiene practices and limited duration of catheterization should be implemented so that biofilm formation, as well as further emergence of antibiotic resistance can be reduced.

CONFLICT OF INTEREST

Disclose no conflict of interest.

ETHICAL APPROVAL

The study had been performed to fulfill the academic purpose and the approval to conduct the proposed study in the Department of Microbiology was taken. In this study, no patient data and specimens were being used, verbal consent was only taken from patients at the time of catheter collection.

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SUPPLEMENTARY INFORMATION

Table S1: The OD₆₃₀ values of 131 isolates and interpretation of biofilm-forming capacity.

No.	Isolate	OD ₆₃₀	Interpretation	56	E. coli-18	0.11	Weak Biofilm
1	P. aeruginosa-1	0.01	No Biofilm	57	<i>E. coli</i> -19	0.06	No Biofilm
2	P. aeruginosa-2	0.11	Weak Biofilm	58	E. coli-20	0.04	No Biofilm
3	P. aeruginosa-3	0.22	Moderate Biofilm	59	E. coli-21	0.02	No Biofilm
4	P. aeruginosa-4	0.24	Moderate Biofilm	60	E. coli-22	(0.04)	No Biofilm
5	P. aeruginosa-5	0.40	Strong Biofilm	61	E. coli-23	0.05	No Biofilm
6	P. aeruginosa-6	0.08	No Biofilm	62	E. coli-24	0.03	No Biofilm
7	P. aeruginosa-7	0.07	No Biofilm	63	E. coli-25	0.03	No Biofilm
8	P. aeruginosa-8	0.44	Strong Biofilm	64	E. coli-26	0.06	No Biofilm
9	P. aeruginosa-9	0.13	Weak Biofilm	65	E. coli-27	0.03	No Biofilm
10	P. aeruginosa-10	0.18	Weak Biofilm	66	E. coli-28	0.02	No Biofilm
11	P. aeruginosa-11	0.23	Moderate Biofilm	67	E. coli-29	0.05	No Biofilm
12	P. aeruginosa-12	0.07	No Biofilm	68	E. coli-30	0.05	No Biofilm
13	P. aeruginosa-13	0.39	Strong Biofilm	69	E. coli-31	0.04	No Biofilm
14	P. aeruginosa-14	0.39	Strong Biofilm	70	P. vulgaris-1	0.07	No Biofilm
15	P. aeruginosa-15	0.29	Moderate Biofilm	71	P. vulgaris-2	0.08	No Biofilm
16	P. aeruginosa-16	0.384	Strong Biofilm	72	P. vulgaris-3	0.16	Weak Biofilm
17	P. aeruginosa-17	0.12	Weak Biofilm	73	P. vulgaris-4	0.04	No Biofilm
18	P. aeruginosa-18	0.38	Moderate Biofilm	74	P. vulgaris-5	0.13	Weak Biofilm
19	P. aeruginosa-19	0.39	Strong Biofilm	75	P. vulgaris-6	0.46	Strong Biofilm
20	P. aeruginosa-20	0.18	Weak Biofilm	76	P. vulgaris-7	0.19	Weak Biofilm
21	P. aeruginosa-21	0.24	Moderate Biofilm	77	P. vulgaris-8	0.12	Weak Biofilm
22	P. aeruginosa-22	0.15	Weak Biofilm	78	P. vulgaris-9	0.19	Moderate Biofilm
23	P. aeruginosa-23	0.21	Moderate Biofilm	79	P. vulgaris-10	0.19	Moderate Biofilm
24	P. aeruginosa-24	0.18	Weak Biofilm	80	P. vulgaris-11	0.20	Moderate Biofilm
25	P. aeruginosa-25	0.16	Weak Biofilm	81	P. vulgaris-12	0.13	Weak Biofilm
26	P. aeruginosa-26	0.10	Weak Biofilm	82	P. vulgaris-13	0.38	Strong Biofilm
27	P. aeruginosa-27	0.16	Weak Biofilm	83	P. vulgaris-14	0.15	Weak Biofilm
28	P. aeruginosa-28	0.16	Weak Biofilm	84	P. vulgaris-15	0.10	Moderate Biofilm
29	P. aeruginosa-29	0.18	Weak Biofilm	85	P. vulgaris-16	0.21	Moderate Biofilm
30	P. aeruginosa-30	0.15	Weak Biofilm	86	P. vulgaris-17	0.12	Weak Biofilm
31	P. aeruginosa-31	0.13	Weak Biofilm	87	P. vulgaris-18	0.12	Moderate Biofilm
32	P. aeruginosa-32	0.29	Moderate Biofilm	88	P. vulgaris-19	0.194	Moderate Biofilm
33	-	0.29	Weak Biofilm	89	P. vulgaris-20	0.13	Weak Biofilm
33 34	P. aeruginosa-33	0.18	Weak Biofilm	90	P. vulgaris-20	0.46	Strong Biofilm
35	P. aeruginosa-34	0.17	Weak Biofilm	91	P. vulgaris-22	0.13	Weak Biofilm
36	P. aeruginosa-35	0.17	Weak Biofilm	92	P. vulgaris-23	0.13	Moderate Biofilm
30 37	P. aeruginosa-36	0.19		93	-	0.194	Weak Biofilm
	P. aeruginosa-37		Weak Biofilm	93 94	P. vulgaris-24 K. proumoniao 1	0.12	Moderate Biofilm
38	P. aeruginosa-38	0.24	Moderate Biofilm	94 95	K. pneumoniae-1	0.22	Weak Biofilm
39 40	E. coli-1	0.05 0.05	No Biofilm No Biofilm	95 96	K. pneumoniae-2 K. pneumoniae-3	0.12	No Biofilm
40 41	E. coli-2			90 97	-	0.07	Weak Biofilm
41	E. coli-3	0.21	Moderate Biofilm	98	K. pneumoniae-4		Weak Biofilm
	E. coli-4	0.45	Strong Biofilm		K. pneumoniae-5	0.10	
43	E. coli-5	0.03	No Biofilm	99 100	K. pneumoniae-6	0.03	No Biofilm
44	E. coli-6	0.05	No Biofilm	100	K. pneumoniae-7	0.21	Moderate Biofilm Weak Biofilm
45	E. coli-7	0.38	Strong Biofilm	101	K. pneumoniae-8	0.14	
46	E. coli-8	0.06	No Biofilm	102	K. pneumoniae-9	0.20	Moderate Biofilm
47	E. coli-9	0.03	No Biofilm	103	K. pneumoniae-10	0.12	Weak Biofilm
48	E. coli-10	0.05	No Biofilm	104	K. pneumoniae-11	0.39	Strong Biofilm
49	E. coli-11	0.15	Weak Biofilm	105	K. pneumoniae-12	0.11	Weak Biofilm
50	E. coli-12	0.06	No Biofilm	106	K. pneumoniae-13	0.20	Moderate Biofilm
51	E. coli-13	0.17	Weak Biofilm	107	K. pneumoniae-14	0.11	Weak Biofilm
52	E. coli-14	0.17	Weak Biofilm	108	K. pneumoniae-15	0.10	Weak Biofilm
53	E. coli-15	0.18	Weak Biofilm	109	K. pneumoniae-16	0.07	No Biofilm
54	E. coli-16	0.18	Weak Biofilm	110	K. pneumoniae-17	0.12	Weak Biofilm
55	E. coli-17	0.13	Weak Biofilm	111	K. pneumoniae-18	0.10	Weak Biofilm

112	K. pneumoniae-19	0.14	Weak Biofilm	123	S. aureus-9	0.23	Moderate Biofilm
113	K. pneumoniae-20	0.11	Weak Biofilm	124	S. aureus-10	0.10	Weak Biofilm
114	K. pneumoniae-21	0.10	Weak Biofilm	126	S. aureus-11	0.01	No Biofilm
115	S. aureus-1	0.12	Weak Biofilm	127	S. aureus-12	0.25	Moderate Biofilm
116	S. aureus-2	0.10	Weak Biofilm	128	S. aureus-13	0.11	Weak Biofilm
117	S. aureus-3	0.10	Weak Biofilm	129	S. aureus-14	0.11	Weak Biofilm
118	S. aureus-4	0.08	No Biofilm	130	S. aureus-15	0.01	No Biofilm
119	S. aureus-5	0.06	No Biofilm	131	S. aureus-16	0.10	Weak Biofilm
120	S. aureus-6	0.05	No Biofilm	132	S. aureus-17	0.09	No Biofilm
121	S. aureus-7	0.08	No Biofilm	135	ODc	0.10	No Biofilm
122	S. aureus-8	0.11	Weak Biofilm				

 Table S2:
 Morphological, cultural, physiological and biochemical characteristics of isolates.

Vegetative cell Cell arrangement Gram staining Spore staining Acid fast staining Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test Tube coagulase test	isolates Rod Single Negative Non spore former Non-acid fast Motile Blue-green, circular, raised colony Not observed	isolates Rod Single Negative Non spore former Non-acid fast Motile Not observed Bright pinky red, circular, flat colonies	isolates Rod Single Negative Non spore former Non-acid fast Non motile Not observed Pink, circular, raised, mucoid	isolates Rod Single Negative Non spore former Non-acid fast Motile Not observed	isolates Coccus Cluster Positive Non spore former Non-acid fast Non motile Not observed
Cell arrangement Gram staining Spore staining Acid fast staining Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Single Negative Non spore former Non-acid fast Motile Blue-green, circular, raised colony Not observed	Single Negative Non spore former Non-acid fast Motile Not observed Bright pinky red, circular, flat colonies	Single Negative Non spore former Non-acid fast Non motile Not observed Pink, circular, raised, mucoid	Single Negative Non spore former Non-acid fast Motile Not observed	Cluster Positive Non spore former Non-acid fast Non motile Not observed
Gram staining Spore staining Acid fast staining Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Negative Non spore former Non-acid fast Motile Blue-green, circular, raised colony Not observed	Negative Non spore former Non-acid fast Motile Not observed Bright pinky red, circular, flat colonies	Negative Non spore former Non-acid fast Non motile Not observed Pink, circular, raised, mucoid	Negative Non spore former Non-acid fast Motile Not observed	Positive Non spore former Non-acid fast Non motile Not observed
Spore staining Acid fast staining Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Non spore former Non-acid fast Motile Blue-green, circular, raised colony Not observed	Non spore former Non-acid fast Motile Not observed Bright pinky red, circular, flat colonies	Non spore former Non-acid fast Non motile Not observed Pink, circular, raised, mucoid	Non spore former Non-acid fast Motile Not observed	Non spore former Non-acid fast Non motile Not observed
Acid fast staining Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	former Non-acid fast Motile Blue-green, circular, raised colony Not observed	former Non-acid fast Motile Not observed Bright pinky red, circular, flat colonies	former Non-acid fast Non motile Not observed Pink, circular, raised, mucoid	former Non-acid fast Motile Not observed	former Non-acid fast Non motile Not observed
Motility test Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Motile Blue-green, circular, raised colony Not observed	Motile Not observed Bright pinky red, circular, flat colonies	Non motile Not observed Pink, circular, raised, mucoid	Motile Not observed	Non motile Not observed
Growth on Cetrimide agar (Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Blue-green, circular, raised colony Not observed	Not observed Bright pinky red, circular, flat colonies	Not observed Pink, circular, raised, mucoid	Not observed	Not observed
(Oxoid [™] CM0579, UK) Growth on MacConkey agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	circular, raised colony Not observed	Bright pinky red, circular, flat colonies	Pink, circular, raised, mucoid		
agar (Oxoid [™] CM0007, UK) Growth on Casein Enzyme Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Not observed	red, circular, flat colonies	raised, mucoid	Not observed	Not observed
Hydrolysate (CLED) agar (Oxoid [™] CM0301, UK) Growth on Mannitol Salt Agar (Himedia [™] M118, India) Oxidase test Catalase test	Not observed	Not obcomicad	colonies		
Agar (Himedia [™] M118, India) Oxidase test Catalase test		Not observed	Not observed	Green-blue, circular, raised, slimy colony	Not observed
Oxidase test Catalase test	Not observed	Not observed	Not observed	Not observed	Circular, yellow, raised smooth colony
Catalase test	Positive	Negative	Negative	Negative	Negative
	Positive	Positive	Positive	Positive	Positive
	Not done	Negative	Negative	Negative	Positive
Citrate utilization	Positive	Negative	Positive	Positive	Positive
Gelatin hydrolysis test	Positive	Negative	Negative	Negative	Positive
Casein hydrolysis test	Positive	Negative	Positive	Negative	Negative
Starch hydrolysis test	Negative	Negative	Positive	Negative	Positive
Voges-Proskauer test	Negative	Negative	Positive	Negative	Positive
Methyl red test	Negative	Positive	Negative	Positive	Negative
Nitrate reduction test	Positive	Positive	Positive	Positive	Positive
	Alkaline slant, acid butt without gas	Acidic slant, acidic butt, Gas +ve	Acidic slant, acidic butt, Gas +ve	Alkaline slant, Acidic butt, Gas +ve	Negative
H ₂ S production	Negative	Negative	Negative	Positive	Negative
Indole test	Negative	Positive	Negative	Positive	Negative
Urease test	Negative	Negative	Positive	Positive	Positive
Arginine dehydrolase	Positive	Negative	Negative	Negative	Positive
Ornithine decarboxylase	Negative	Positive (28 isolates) Negative (3 isolates)	Negative	Positive	Negative
Lysine		Positive	Negative	Negative	

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		Fermentation te	st		
Arabinose	Negative	Positive	Positive	Negative	Negative
Cellobiose	Negative	Negative	Positive	Negative	Negative
DNase	Negative	Negative	Negative	Negative	Positive
Fructose	Positive	Positive	Negative	Not done	Positive
Galactose	Negative	Positive	Positive	Not done	Positive
Glucose	Positive	Positive	Positive	Positive	Positive
Lactose	Negative	Positive	Positive	Negative	Positive
Maltose	Negative	Positive	Positive	Negative	Positive
Mannitol	Positive	Positive	Positive	Negative	Positive
Mannose	Negative	Positive	Positive	Negative	Positive
Raffinose	Not done	Negative	Positive	Negative	Negative
Ribose	Positive	Positive	Not done	Negative	Positive
Salicin	Negative	Negative	Positive	Negative	Negative
Sucrose	Negative	Positive	Positive	Negative	Positive
Trehalose	Negative	Positive	Positive	Positive	Positive
Xylose	Negative	Negative	Positive	Positive	Negative