



# Study of Biofilm, Virulence Genes, and Risk Factors in Urinary Catheter-Associated *Escherichia coli* Infections

Eman Hamdy Mohamed<sup>1</sup>, Maysaa El Sayed Zaki<sup>2,\*</sup>, Mostafa Abdelsalam Ramadan<sup>3</sup>, Mohamed Mohsen Motawea<sup>4</sup>, Dina F Badr<sup>5</sup>, Ahmed Gomaa Ahmed El Sayed<sup>5</sup>, Mai Esam Ahmed Hamed<sup>1</sup>

<sup>1</sup> Clinical Pathology Department, Beni-Suef Faculty of Medicine, Beni-Suef, Egypt

<sup>2</sup> Clinical Pathology Department, Mansoura Faculty of Medicine, Mansoura, Egypt

<sup>3</sup> Dialysis Unit, Internal Medicine Department, Mansoura Faculty of Medicine, Mansoura, Egypt

<sup>4</sup> Internal Department of Medicine, Mansoura Faculty of Medicine, Mansoura, Egypt

<sup>5</sup> Medical Microbiology and Immunology Department, Mansoura Faculty of Medicine, Mansoura, Egypt

\* Corresponding author: Clinical Pathology Department, Mansoura Faculty of Medicine, Mansoura, Egypt. Email: may\_s@mans.edu.eg

Received 2023 December 31; Revised 2024 April 22; Accepted 2024 April 29.

## Abstract

**Background:** Catheter-associated urinary tract infections (CAUTIs) are prevalent among intensive care unit (ICU) patients and pose significant health risks. The causative agent, *Escherichia coli*, is commonly associated with these infections.

**Objectives:** The primary objective of our study was to evaluate various risk factors associated with CAUTIs. Additionally, we investigated the capacity of *E. coli* to form biofilms and the presence of virulence genes in these infections.

**Methods:** We conducted a prospective study involving 204 patients admitted to ICUs with indwelling urinary catheters who subsequently developed CAUTIs. The isolated pathogen in all cases was *E. coli*. We subjected the isolated *E. coli* strains to microbiological analyses, including biofilm formation assays and polymerase chain reaction (PCR) for detecting virulence genes (*pap-C*, *fim-H*, *sfa*, *afa*, *hlyA*, and *cnfI*).

**Results:** Our findings revealed that 111 out of 204 isolated *E. coli* strains (54.4%) could form biofilms in vitro. Among the detected virulence genes, *fim-H* was the most frequently observed (32.4%), followed by *pap-C* (14.2%), *sfa* (12.3%), and *afaI* (11.3%). The least frequently detected genes were *cnfI* (3.9%) and *hlyA* (2.5%). Notably, there was a significant association between the presence of *fim-H*, *pap-C*, *afaI*, and *hlyA* genes and the ability of *E. coli* to form biofilms ( $P = 0.001$ ,  $P = 0.001$ ,  $P = 0.04$ ,  $P = 0.046$ , respectively). Regarding the timing of CAUTIs, most cases occurred 7 - 10 days after catheter insertion (48.53%), followed by 3 - 6 days after insertion (45.59%). The least frequent occurrence was after 10 days from catheter insertion (5.86%). Early presence of CAUTIs (within 3 - 6 days) was significantly associated with diabetes mellitus (DM) and post-operative complications (both with  $P = 0.046$ ). Furthermore, we observed a significant increase in the presence of the *cnfI* gene in cases of early CAUTIs (3 - 6 days after catheter insertion,  $P = 0.001$ ).

**Conclusions:** Our study highlights the importance of understanding the risk factors, virulence genes, and biofilm formation capacity in *E. coli*-related CAUTIs. These findings may guide preventive strategies and improve patient outcomes.

**Keywords:** *Escherichia coli*, Urinary Tract Infections, Biofilm, Virulence Genes

## 1. Background

Catheter-associated urinary tract infections (CAUTIs) are common in healthcare-associated infections (1). Around 40% of healthcare-associated infections are urinary tract infections, with about 80% of cases attributed to the use of urinary catheters (2). The use of urinary catheters ranges from 12% to 16% among adult hospital patients, and the incidence of CAUTIs ranges

from 3% to 7% in these patients (3). The total number of urinary tract infections worldwide is 222 million (3). Diagnosing CAUTIs relies on both clinical and laboratory diagnoses of bacteriuria. Signs and symptoms of these infections include fever, suprapubic tenderness, and flank pain (4, 5). Asymptomatic bacteriuria alone, without predisposing risk factors for infections, does not indicate antibiotic therapy (6). The removal of the catheter, along with antibiotics and other drugs, must

be indicated by the clinician (7, 8). Infection control guidelines for preventing CAUTIs depend on knowledge of the data concerning risk factor analysis of CAUTIs (9).

The most common pathogen associated with urinary tract infections is *Escherichia coli*, responsible for about 90% of community-acquired infections and around 50% of healthcare-associated infections (3). Various virulence factors in uropathogenic *E. coli* facilitate the invasion, growth, and persistence in the urinary tract epithelium (10). The host defense mechanisms include cytokine secretion such as interleukin 8, uroepithelial defensin, cytopeptide expression like Tamm-Horsfall protein, and urinary flow (11). The formation of biofilm by *E. coli* represents a virulence mechanism where microbial cells are irreversibly bound to the epithelial surface with the formation of polysaccharide material (12). The biofilm prevents the access of antimicrobial agents and antibodies to the bacterial cells and provides nutrients to the bacterial cells (13).

Various virulence genes in uropathogenic *E. coli* increase the severity of urinary tract infections. These genes include P fimbriae (*pap*), type 1-fimbriae (*fim-H*), fimbrial-adhesin (*afai*), S-fimbriae (*sfa*), hemolysin (*hly*), and cytotoxic necrotizing factor (*cnfI*) (14). These genes are responsible for colonizing the host epithelium, evading host mechanisms, and invading with injuries to host cells (15). There is a link between the presence of these virulence genes, biofilm formation, and antibiotic resistance in *E. coli*. These factors need careful assessment to avoid antibiotic therapy failure (16). Despite the known role of these virulence factors in the severity of urinary tract infections, few studies have explored their association with CAUTIs, particularly in the context of biofilm formation and antibiotic resistance. This represents a significant gap in the literature, as understanding these associations could inform more effective treatment strategies and help prevent antibiotic therapy failure.

## 2. Objectives

This study aims to address these gaps by evaluating the various risk factors associated with CAUTIs, with a particular focus on the capacity of *E. coli* to form biofilms and the presence of various virulence genes in these infections.

## 3. Methods

This retrospective study included patients with CAUTIs admitted to the intensive care units (ICU) of Mansoura University Hospital, Egypt, from January 2019 to March 2022. The inclusion criteria were patients aged

>18 years diagnosed with healthcare-associated CAUTIs and inserted urinary catheters, according to the criteria identified by the centers for disease control (CDC) (17), with bacterial infection due to *E. coli*. The exclusion criteria were patients with community-acquired urinary tract infections and those with urinary tract infections not associated with catheters. The sample size depended on the non-probability sampling method and the time frame of the study, during which patients with CAUTIs due to *E. coli* were diagnosed.

### 3.1. Urine Sample

Urine samples were obtained from the urinary catheter and rapidly transported to the laboratory. A microscopic examination was carried out for pus cells, followed by a microbiological culture (18). *Escherichia coli* identification was performed using Gram stain and complete biochemical identification with triple sugar iron agar, Simmon citrate, urease, lysine, phenylalanine, Methyl red, Voges Proskauer test, and subculture on MacConkey agar (19). A pure suspension of *E. coli* was prepared by subculturing the isolates in brain heart infusion broth overlaid with glycerol, and the suspension was kept frozen at -20°C for polymerase chain reaction (PCR) (20).

### 3.2. Antibiotics Sensitivity Test

An antibiotic sensitivity test was performed using the disc diffusion method. The antibiotics discs used were imipenem (10 µg), meropenem (10 µg), cefepime (30 µg), ceftriaxone (30 µg), gentamicin (10 µg), aztreonam (30 µg), cefotaxime (30 µg), ceftazidime (30 µg), ciprofloxacin (5 µg), trimethoprim/sulfamethoxazole (25 µg), and nitrofurantoin (300 µg) (Oxoid, United Kingdom). Pure colonies of *E. coli* were suspended in Muller-Hinton broth, and half McFarland concentrations were prepared. The suspension was then spread on Muller-Hinton agar. Antibiotic discs were placed on the agar, and the plates were incubated at 37°C for 24 hours. The diameter of the inhibition zones around the antibiotic discs was measured, and the results were interpreted as sensitive or resistant according to the guidelines of the Clinical Laboratory Standards Institute (CLSI) (21).

### 3.3. Biofilm Formation

*Escherichia coli* were grown in tryptic soy broth for 24 hours at 37°C. Then, a 1:20 dilution of the cultures was performed in tryptic soy broth, and 200 microliters were added to each well in a 96-well microplate, with incubation for 48 hours at 37°C. The wells were stained

with 2% crystal violet and incubated for 15 minutes. The wells were then washed with sterile distilled water, and 150 microliters of 95% ethanol was added to the wells. The microplate was read using a microplate reader at a wavelength of 570 nm (22).

#### 3.4. Extraction of DNA

The DNA of *E. coli* isolates was extracted using the boiling method as previously described (23).

#### 3.5. Amplification and Detection

The primers used are summarized in Table 1 and (24). The amplification mixture was prepared using a Qiagen ready-to-use mixture. The amplification procedure included the following steps using an Eppendorf Master Cycler: Initial denaturation at 94°C for 2 minutes, followed by thirty cycles of denaturation at 94°C for 60 seconds, annealing at 63°C for 30 seconds, and extension at 72°C for 90 seconds, with a final extension step at 72°C for 5 minutes. Sterile distilled water was used as a negative control. Detection was performed by electrophoresis on a 1.5% agarose gel stained with ethidium bromide for 20 minutes, followed by ultraviolet visualization of the bands.

#### 3.6. Statistical Analysis

The data from the study was analyzed using SPSS 22. Qualitative data was expressed as numbers and percentages, with comparisons performed using the chi-square test. Significance was indicated by a P-value of < 0.05. Quantitative data was expressed as mean and standard deviation (SD).

## 4. Results

#### 4.1. Demographic Characteristics and Clinical Data

The study included 204 patients with healthcare-associated CAUTIs. Among them, 107 (52.5%) were males and 97 (47.5%) were females, with a mean age of 49.53 ± 14.7 years. The common comorbidities were diabetes mellitus (DM) (33.3%), renal disorders (23.5%), and post-operative complications (11.3%). Various clinical conditions led to ICU admission, including common malignancy (24.0%), intracranial hemorrhage (20.6%), and renal failure (15.2%). Recurrent urinary tract infections were reported in 24 patients (11.4%). The most common clinical symptoms and signs were pain on clinical examination (55.4%), pain on micturition (46.6%), and fever (45.6%) (Table 2).

#### 4.2. Biofilm Formation

The capacity to form biofilm in vitro was detected in 111 (54.4%) of the isolated *E. coli* strains.

#### 4.3. Antibiotic Resistance

The most frequently detected resistances among isolated *E. coli* were to ciprofloxacin (64.7%), followed by resistance to the third-generation cephalosporin ceftazidime (46.6%) and cefotaxime (43.6%). Additionally, resistance to trimethoprim/sulfamethoxazole was observed in 42.2% of the isolates (Figure 1).

#### 4.4. Virulence Gene Detection

The most frequently detected genes were *fim-H* (32.4%), followed by *pap-C* (14.2%), *sfa* (12.3%), and *afaI* (11.3%). The least frequently detected genes were *cnfI* (3.9%) and *hly* (2.5%) (Figure 2). There was a significant association between the presence of *fim-H*, *pap-C*, *afaI*, and *hly* genes ( $P = 0.001$ ,  $P = 0.001$ ,  $P = 0.04$ ,  $P = 0.046$  respectively) and the capacity to form biofilm in isolated *E. coli* (Table 3).

Urinary tract infections were most commonly detected 7 - 10 days after catheter insertion (48.53%), followed by 3 - 6 days after insertion (45.59%). The least frequent occurrence of urinary tract infections was more than 10 days after catheter insertion (5.86%). There was a significant association between the early presence of CAUTIs (from 3 to 6 days) and the presence of DM ( $P = 0.046$ ) and postoperative complications ( $P = 0.046$ ). Regarding the presence of virulence genes, there was a significant increase in the presence of the *cnfI* gene ( $P = 0.001$ ) with early CAUTIs (3 - 6 days and 7 - 10 days) (Table 4).

## 5. Discussion

Catheter-associated urinary tract infections represent a major health problem worldwide. Many risk factors lead to CAUTIs despite the efforts of infection control guidelines (25). In the present study, the common comorbidities associated with CAUTIs were DM, renal disorders, and previous surgery history. There was a significant association between the early presence of CAUTIs (from 3 to 6 days) and the presence of DM ( $P = 0.046$ ) and postoperative complications ( $P = 0.046$ ). It is well known that DM predisposes individuals to various infections due to the compromised immune system (26). Different factors increase infections in diabetic patients, such as high glucose levels, which are considered a source of nutrient material for the

**Table 1.** The Sequences of the Primers Used for Gene Detection and the Product Base Pair (bp) Length

Gene	The Sequences of the Primers	Bp
PapC	F GACGGCACTGCTGCAGGGTGTGGCG	328
	R ATATCCTTTCTGCAGGGATGCAATA	
Fim-H	F TGTACTGCTGATGGGCTGGTC	564
	R GGGTAGTCCGGCAGAGTAACG	
Sfa-	F CTCGGGAGAAGTGGTGCATCTTAC	410
	R CGGAGGAGTAATTACAACTGGCA	
Afa	F GCTGGGCAGCAAAGTATAACTCTC	750
	R CATCAAGCTGTTTTCGTCGGCCG	
HlyA	F AACAAAGGATAAGCACTGTTCTGGCT	1177
	R ACCATATAAGGGTCATCCCGT	
CnfI	F AAGATGGAGTTTCTATGCAGGAG	498
	R TGGAGTTTCTATGCAGGAG	

**Table 2.** Demographic and Clinical Data of the Studied Patients<sup>a</sup>

Demographic and Clinical Data	Values
<b>Sex</b>	
Male	107 (52.5)
Female	97 (47.5)
<b>Age (y)</b>	(49.53 ± 14.79)
<b>Comorbidities</b>	
Diabetes mellitus	68 (33.3)
Kidney diseases	48 (23.5)
Post-operative complications	23 (11.3)
Underlying etiology	49 (24.0)
Malignancy	19 (9.3)
Diabetic complications	31 (15.2)
Renal failure intracranial hemorrhage	42 (20.6)
Surgical complications	22 (10.8)
Myocardial infarctions	6 (2.9)
Hepatic disorders	26 (12.7)
Miscellaneous	9 (4.4)
Recurrent urinary tract infections	24 (11.8)
<b>Clinical signs and symptoms</b>	
Fever	93 (45.6)
Urgency	48 (3.5)
Frequency	22 (10.8)
Pain in the micturition	95 (46.6)
Pain on examination	113 (55.4)

<sup>a</sup> Values are expressed as No. (%) or mean ± SD.

bacteria, decreased T cell-mediated immunity, and impaired emptying of the bladder due to autonomic nervous system neuropathy (27).

Catheter-associated urinary tract infection as a complication of surgery is known due to the insertion of catheters for long durations in various surgical

operations (28). Renal disorders, such as chronic renal failure, can be associated with increased urinary tract infections, especially in catheterized patients, due to a deficiency in the immune response and other comorbidities such as urinary obstruction and the presence of DM (27).

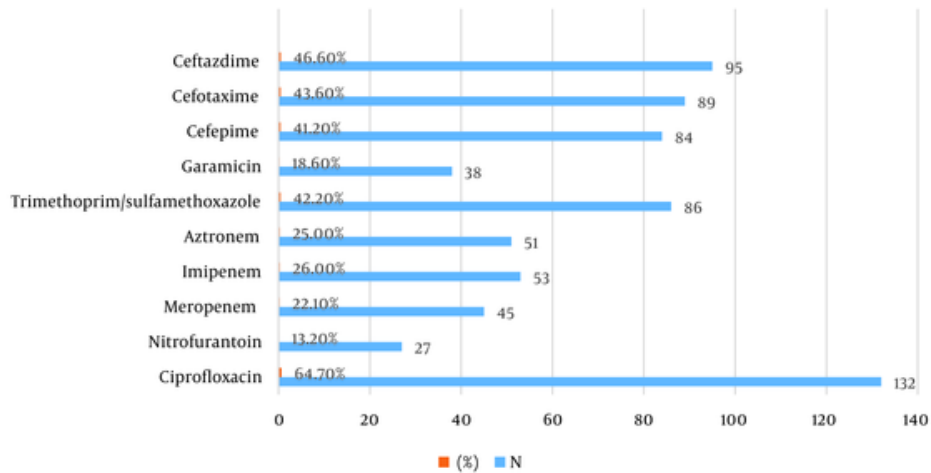


Figure 1. Antibiotic resistance among isolated *Escherichia coli*

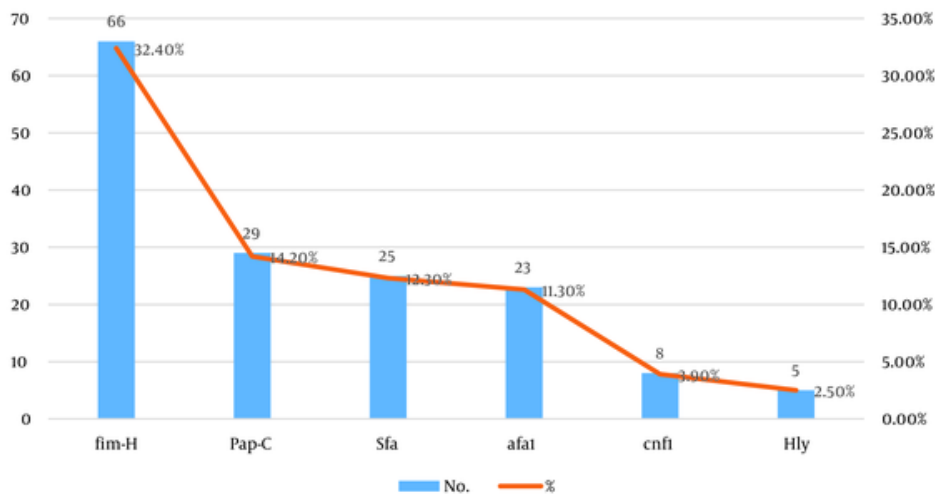


Figure 2. The frequency of virulence genes in isolated *Escherichia coli*

The early removal of the catheter is a cornerstone of the prevention of CAUTIs (29). However, the duration in which catheterization may lead to infections has not been fully studied. In the present study, urinary tract infections were commonly detected after 7 - 10 days of catheter insertion (48.53%), and then after 3 - 6 days of insertion (45.59%). A previous study reported a non-linear increase in the cumulative risk hazard as the increase in time of catheter insertion suggests that

everyday increases in catheterization lead to an increased risk of developing CAUTIs (25). The present study provides interesting data about antimicrobial resistance models, biofilm production, and virulence genes associated with *E. coli* causing CAUTIs.

The capacity to form biofilm in an in vitro study was detected among 111 (54.4%) of isolated *E. coli*. The processes that predispose biofilm-bound *E. coli* to progress to CAUTI remain unclear but are of diagnostic

**Table 3.** Association Between Biofilm and the Presence of Virulence Genes <sup>a</sup>

Virulence Genes	Positive Biofilm (n = 111)	Negative Biofilm (n = 93)	P-Value
<i>fim-H</i>	64 (57.7)	2 (2.2)	0.001
<i>Pap</i>	25 (22.5)	4 (4.3)	0.001
<i>Sfa</i>	13 (11.7)	12 (12.9)	0.48
<i>Cnf</i>	2 (1.8)	6 (6.4)	0.1
<i>Afa-I</i>	17 (15.3)	6 (6.4)	0.04
<i>Hyl</i>	5 (4.5)	0 (0)	0.046
<i>Cnf</i>	2 (1.8)	6 (6.4)	0.1

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

**Table 4.** Association Between Catheter Days of the Insertion and Demographic, Clinical Data, Biofilm Formation, and the Presence of Virulence Genes <sup>a</sup>

Demographic, Clinical Data, Biofilm Formation, and the Presence of Virulence Genes	Catheter Days 3 - 6 (n = 93)	Catheter Days 7 - 10 (n = 99)	Catheter Days > 10 (n = 12)	P-Value
Sex				0.13
Male	54 (58.1)	45 (45.5)	8 (66.6)	
Female	39 (41.9)	54 (54.5)	4 (33.3)	
<b>Diabetes mellitus</b>	68 (73.1)	58 (58.6)	10 (83.3)	0.046
<b>Kidney diseases</b>	72 (77.4)	77 (77.8)	7 (58.3)	0.31
<b>Post-operative complications</b>	9 (9.7)	10 (10.1)	4 (33.3)	0.045
<b>Fever</b>	54 (58.1)	53 (53.5)	4 (33.3)	0.26
<b>Dysuria</b>	68 (73.1)	75 (75.8)	7 (58.3)	0.43
<b>Urgency</b>	73 (78.5)	73 (73.7)	10 (83.3)	0.63
<b>Frequency</b>	86 (92.5)	87 (87.9)	9 (75)	0.16
<b>Pain on micturition</b>	41 (44.1)	47 (47.5)	7 (53.3)	0.63
<b>Biofilm</b>	54 (58.1)	49 (49.5)	8 (66.7)	0.33
<i>fim-H</i>	33 (35.5)	27 (27.3)	6 (50)	0.19
<i>Pap-C</i>	10 (10.8)	15 (15.2)	4 (33.3)	0.10
<i>Sfa</i>	12 (12.9)	11 (11.1)	2 (16.7)	0.83
<i>AfaI</i>	10 (10.8)	13 (13.1)	0	0.34
<i>CnfI</i>	2 (2.2)	3 (3.03)	3 (25)	0.001
<i>Hyl</i>	3 (3.2)	2 (2.02)	0	0.73

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

and therapeutic interest (30). These processes are presumably complex and include biofilm efflux, adhesion to host tissue, evasion of the immune system, and nutrient acquisition (31). The capacity of organisms to form biofilm in an in vitro study indicates their capacity to form biofilm on the surface of the catheter and the tissues of the renal system, a finding that alerts clinicians that this biofilm is part of the pathogenesis of CAUTIs (32). The most frequently detected resistances to antibiotics among isolated *E. coli* were toward ciprofloxacin, followed by resistance to the third-generation cephalosporins ceftazidime and cefotaxime, and resistance to trimethoprim/sulfamethoxazole.

The common antibiotics used for the treatment of UTIs include quinolones, cephalosporins, and trimethoprim-sulfamethoxazole. The unwise use of these antibiotics may explain the high frequency of resistance among isolated *E. coli* in the present study. The high resistance of the isolated *E. coli* from CAUTIs to ciprofloxacin may be attributed to the presence of plasmids that encode resistance to this antibiotic, leading to the endemicity of these isolates in hospitals associated with hospital outbreaks of UTIs. The most frequently detected genes were *fim-H* (32.4%), followed by *Pap-C* (14.2%), *SFA* (12.3%), and *afaI* (11.3%). The least frequently detected genes were *cnfI* (3.9%) and *hyl* (2.5%). Higher rates of virulence genes were reported



previously in *E. coli* isolated from UTI for *Pap-C*, *hyl*, and *cnfI* (50.4%, 50.4%, 50.4% respectively) and lower frequency for *afaI* (8.13%) (33). In another study, the prevalence rates for virulence genes *sfa*, *pap*, *cnfI*, *afa*, and *hylI* were 26%, 25%, 18.6%, 6%, and 5% respectively (34).

The high prevalence of the type 1 fimbria gene was in line with a previous study indicating a high incidence of type 1 fimbriae among uropathogenic *E. coli* (35), especially in CAUTIs. The difference in the prevalence of virulence genes among different studies can be attributed to the difference in virulence gene distribution among different strains of *E. coli*. There is also a known ability of *E. coli* to exchange its genetic material and transfer genetic virulence factors, besides the effects of various antibiotics intake that leads to the elimination of certain strains with different virulence genes. Moreover, the presence of some virulence genes such as *pap-c* may be associated with certain types of UTIs, such as pyelonephritis, more than with its presence in cystitis (35). There was a significant association between the presence of *fim-H*, *pap*, *afa-I*, and *hyl* genes ( $P = 0.001$ ,  $P = 0.001$ ,  $P = 0.04$ ,  $P = 0.046$  respectively) and the capacity to form biofilm in isolated *E. coli*.

In another similar study by Zamani and Salehzadeh, it was found that biofilm-producing UPEC were significantly associated with the *fim* gene (36). The association of these adhesion genes and biofilm formation may indicate that these factors play a role in biofilm formation. Regarding the presence of virulence genes, there was a significant increase in the presence of the *cnfI* gene and the early CAUTI insertion (3 - 6 days and 7 - 10 days). This finding may indicate that these adhesion genes play a role in the pathogenesis of CAUTIs shortly after catheter insertion.

The study's focus on biofilm formation and virulence genes in CAUTIs represents a novel approach. Prior research has primarily examined bacterial colonization and antibiotic resistance, but this study delves deeper into the mechanisms driving infection persistence. This study is important for understanding biofilm dynamics, which is crucial for preventing and managing CAUTIs. Identifying specific virulence genes in CAUTIs can inform targeted therapies and also shed light on the genetic diversity of uropathogens. Finally, novel insights into biofilm and virulence genes can lead to more effective treatment strategies. We can implicate these findings in:

1. Catheter care guidelines: Healthcare policies should emphasize proper catheter insertion, maintenance, and removal.

2. antibiotic stewardship: Understanding virulence genes can guide antibiotic selection.

3. Biofilm disruption: Develop interventions to prevent or disrupt biofilm formation.

4. Gene-Based diagnostics: Incorporate virulence gene detection into diagnostic protocols.

5. Targeted therapies: Develop drugs that specifically target biofilm-associated virulence factors.

This study's findings provide a fresh perspective on CAUTIs, emphasizing biofilm dynamics and virulence genes. By integrating these insights into healthcare policies and practice guidelines, we can enhance patient outcomes and reduce the burden of CAUTIs.

### 5.1. Conclusions

The present study highlights several risk factors leading to healthcare-associated urinary tract infections, including DM, renal disorders, and surgical intervention. Catheter-associated urinary tract infections usually occur within the first 10 days of catheter insertion. The presence of biofilm formation is a common finding with *E. coli* associated with CAUTIs. The virulence factors are common among isolated *E. coli* and are significantly associated with biofilm formation.

These findings can guide several aspects of healthcare policies, such as catheter care guidelines, which should emphasize proper catheter insertion, maintenance, and removal. Additionally, in antibiotic stewardship, understanding virulence genes can guide antibiotic selection and aid in developing interventions to prevent or disrupt biofilm formation. Finally, there is potential for developing drugs that specifically target biofilm-associated virulence factors. Further investigations are needed to identify novel targets for disrupting biofilms and enhancing antibiotic efficacy.

### Footnotes

**Authors' Contribution:** EHM, shared in the laboratory study, the draft preparation of the article data analysis of the study and revision of the draft of the article; MEZ, shared in the laboratory study and draft preparation of the article; MAR, shared in the clinical data collection, and draft preparation of the article; MMM, shared in the clinical data collection, and draft preparation of the article; DEFB, shared in the laboratory study, the draft preparation of the article data analysis of the study, and revision of the draft of the article; AGAE, shared in the laboratory study, the draft preparation of the article data analysis of the study and revision of the draft of the article; MEA, shared the laboratory study, the draft

preparation of the article, and the data analysis of the study; MEAH, shared in the laboratory study and draft preparation of the article. All authors have read and approved the final manuscript.

**Conflict of Interests Statement:** Authors declared no conflict of interests.

**Data Availability:** The datasets generated and analyzed during the current study are available in the Figshare repository at <https://doi.org/10.6084/m9.figshare.24783033.v1>.

**Ethical Approval:** All methods were performed by the ethical standards as laid down in the Declaration of Helsinki and its later amendments or comparable ethical standards. The ethical approval of the study was obtained from the ethical committee of Mansoura Faculty of Medicine (R. 23.12.2399).

**Funding/Support:** This article did not receive any funding.

**Informed Consent:** Written informed consent was obtained from the parent of each child.

## References

- Trautner BW. Management of catheter-associated urinary tract infection. *Curr Opin Infect Dis.* 2010;**23**(1):76-82. [PubMed ID: 19926986]. [PubMed Central ID: PMC2865895]. <https://doi.org/10.1097/QCO.0b013e328334dda8>.
- Wagenlehner F, Tandogdu Z, Bartoletti R, Cai T, Cek M, Kulchavenya E, et al. The global prevalence of infections in urology study: A long-term, worldwide surveillance study on urological infections. *Pathogens.* 2016;**5**(1). [PubMed ID: 26797640]. [PubMed Central ID: PMC4810131]. <https://doi.org/10.3390/pathogens5010010>.
- National Healthcare Safety Network. *Urinary tract infection (3-center disease control, nceqid, dhq). Catheter-associated urinary tract infection [cauti] and noncatheter-associated urinary tract infection [uti] and other urinary system infection [usi] events.* Atlanta, Georgia: National Healthcare Safety Network; 2019, [cited 2024]. Available from: <https://www.cdc.gov/nhsn/pdfs/pscmanual/7pscscauticurrent.pdf>.
- Marschall J, Zhang L, Foxman B, Warren DK, Henderson JP, C. D. C. Prevention Epicenters Program. Both host and pathogen factors predispose to Escherichia coli urinary-source bacteremia in hospitalized patients. *Clin Infect Dis.* 2012;**54**(12):1692-8. [PubMed ID: 22431806]. [PubMed Central ID: PMC3357479]. <https://doi.org/10.1093/cid/cis252>.
- Marschall J, Piccirillo ML, Foxman B, Zhang L, Warren DK, Henderson JP, et al. Patient characteristics but not virulence factors discriminate between asymptomatic and symptomatic E. coli bacteriuria in the hospital. *BMC Infect Dis.* 2013;**13**:213. [PubMed ID: 23663267]. [PubMed Central ID: PMC3658957]. <https://doi.org/10.1186/1471-2334-13-213>.
- Cope M, Cevallos ME, Cadle RM, Darouiche RO, Musher DM, Trautner BW. Inappropriate treatment of catheter-associated asymptomatic bacteriuria in a tertiary care hospital. *Clin Infect Dis.* 2009;**48**(9):1182-8. [PubMed ID: 19292664]. <https://doi.org/10.1086/597403>.
- Hooton TM, Bradley SF, Cardenas DD, Colgan R, Geerlings SE, Rice JC, et al. Diagnosis, prevention, and treatment of catheter-associated urinary tract infection in adults: 2009 International Clinical Practice Guidelines from the Infectious Diseases Society of America. *Clin Infect Dis.* 2010;**50**(5):625-63. [PubMed ID: 20175247]. <https://doi.org/10.1086/650482>.
- Nicolle LE, Gupta K, Bradley SF, Colgan R, DeMuri GP, Drekonja D, et al. Clinical practice guideline for the management of asymptomatic bacteriuria: 2019 update by the infectious diseases society of america. *Clin Infect Dis.* 2019;**68**(10):1611-5. [PubMed ID: 31506700]. <https://doi.org/10.1093/cid/ciz021>.
- Hariati H, Suza DE, Tarigan R. Risk factors analysis for catheter-associated urinary tract infection in medan, indonesia. *Open Access Maced J Med Sci.* 2019;**7**(19):3189-94. [PubMed ID: 31949514]. [PubMed Central ID: PMC6953942]. <https://doi.org/10.3889/oamjms.2019.798>.
- Terlizzi ME, Gribaudo G, Maffei ME. Uropathogenic Escherichia coli (UPEC) infections: Virulence factors, bladder responses, antibiotic, and non-antibiotic antimicrobial strategies. *Front Microbiol.* 2017;**8**:1566. [PubMed ID: 28861072]. [PubMed Central ID: PMC5559502]. <https://doi.org/10.3389/fmicb.2017.01566>.
- Spencer JD, Schwaderer AL, Becknell B, Watson J, Hains DS. The innate immune response during urinary tract infection and pyelonephritis. *Pediatric Nephrol.* 2014;**29**:1139-49.
- Donlan RM. Biofilms: Microbial life on surfaces. *Emerg Infect Dis.* 2002;**8**(9):881-90. [PubMed ID: 12194761]. [PubMed Central ID: PMC2732559]. <https://doi.org/10.3201/eid0809.020063>.
- Davies J, Davies D. Origins and evolution of antibiotic resistance. *Microbiol Mol Biol Rev.* 2010;**74**(3):417-33. [PubMed ID: 20805405]. [PubMed Central ID: PMC2937522]. <https://doi.org/10.1128/MMBR.00016-10>.
- Sarowska J, Futoma-Koloch B, Jama-Kmiecik A, Frej-Madrzak M, Ksiazczyk M, Bugla-Ploskonska G, et al. Virulence factors, prevalence and potential transmission of extraintestinal pathogenic Escherichia coli isolated from different sources: recent reports. *Gut Pathog.* 2019;**11**:10. [PubMed ID: 30828388]. [PubMed Central ID: PMC6383261]. <https://doi.org/10.1186/s13099-019-0290-0>.
- Luthje P, Brauner A. Virulence factors of uropathogenic E. coli and their interaction with the host. *Adv Microb Physiol.* 2014;**65**:337-72. [PubMed ID: 25476769]. <https://doi.org/10.1016/bs.ampbs.2014.08.006>.
- Katongole P, Nalubega F, Florence NC, Asimwe B, Andia I. Biofilm formation, antimicrobial susceptibility and virulence genes of Uropathogenic Escherichia coli isolated from clinical isolates in Uganda. *BMC Infect Dis.* 2020;**20**(1):453. [PubMed ID: 32600258]. [PubMed Central ID: PMC7325280]. <https://doi.org/10.1186/s12879-020-05186-1>.
- Healthcare Infection Control Practices Advisory Committee. *Guidelines for prevention of catheter - Associated Urinary Tract Infections 2009.* Atlanta, Georgia: Centers for Disease Control and Prevention; 2019, [cited 2024].
- Robinson JL, Finlay JC, Lang ME, Bortolussi R; Canadian Paediatric Society Infectious Diseases; Immunization Committee. Urinary tract infections in infants and children: Diagnosis and management. *Paediatr Child Health.* 2014;**19**(6):315-25. [PubMed ID: 25332662]. [PubMed Central ID: PMC4173959]. <https://doi.org/10.1093/pch/19.6.315>.
- Jackson EC. Urinary tract infections in children: knowledge updates and a salute to the future. *Pediatr Rev.* 2015;**36**(4):153-64. quiz 165-6. [PubMed ID: 25834219]. <https://doi.org/10.1542/pir.36-4-153>.
- Lupindu AM. *Isolation and characterization of Escherichia coli from animals, humans, and environment.* Houston, USA: InTech; 2017.
- Clinical and Laboratory Standards Institute. *Performance standards for antimicrobial susceptibility testing.* Wayne, Pennsylvania, United States: Clinical and Laboratory Standards Institute; 2018, [cited 2024]. Available from: [https://clsi.org/media/1930/m100ed28\\_sample.pdf](https://clsi.org/media/1930/m100ed28_sample.pdf).



22. Stepanovic S, Vukovic D, Hola V, Di Bonaventura G, Djukic S, Cirkovic I, et al. Quantification of biofilm in microtiter plates: overview of testing conditions and practical recommendations for assessment of biofilm production by staphylococci. *APMIS*. 2007;**115**(8):891-9. [PubMed ID: 17696944]. [https://doi.org/10.1111/j.1600-0463.2007.apm\\_630.x](https://doi.org/10.1111/j.1600-0463.2007.apm_630.x).
23. Omar B, Atif H, Mogahid M. Comparison of three DNA extraction methods for polymerase chain reaction (PCR) analysis of bacterial genomic DNA. *Afr J Microbiol Res*. 2014;**8**(6):598-602. <https://doi.org/10.5897/ajmr2013.6459>.
24. Zhao R, Shi J, Shen Y, Li Y, Han Q, Zhang X, et al. Phylogenetic distribution of virulence genes among ESBL-producing uropathogenic *Escherichia coli* isolated from long-term hospitalized patients. *J Clin Diagn Res*. 2015;**9**(7):DC01-4. [PubMed ID: 26393125]. [PubMed Central ID: PMC4572956]. <https://doi.org/10.7860/JCDR/2015/13234.6157>.
25. Letica-Kriegel AS, Salmasian H, Vawdrey DK, Youngerman BE, Green RA, Furuya EY, et al. Identifying the risk factors for catheter-associated urinary tract infections: a large cross-sectional study of six hospitals. *BMJ Open*. 2019;**9**(2). e022137. [PubMed ID: 30796114]. [PubMed Central ID: PMC6398917]. <https://doi.org/10.1136/bmjopen-2018-022137>.
26. Ahmed AE, Abdelkarim S, Zenida M, Baiti MAH, Alhazmi AAY, Alfaifi BAH, et al. Prevalence and associated risk factors of urinary tract infection among diabetic patients: A cross-sectional study. *Healthcare (Basel)*. 2023;**11**(6). [PubMed ID: 36981518]. [PubMed Central ID: PMC10048613]. <https://doi.org/10.3390/healthcare11060861>.
27. Werneburg GT. Catheter-associated urinary tract infections: Current challenges and future prospects. *Res Rep Urol*. 2022;**14**:109-33. [PubMed ID: 35402319]. [PubMed Central ID: PMC8992741]. <https://doi.org/10.2147/rru.s273663>.
28. Salari N, Karami MM, Bokaei S, Chalesghar M, Shohaimi S, Akbari H, et al. The prevalence of urinary tract infections in type 2 diabetic patients: a systematic review and meta-analysis. *Eur J Med Res*. 2022;**27**(1):20. [PubMed ID: 35123565]. [PubMed Central ID: PMC8817604]. <https://doi.org/10.1186/s40001-022-00644-9>.
29. Kuo IC, Lee JJ, Hwang DY, Lim LM, Lin HY, Hwang SJ, et al. Pyuria, urinary tract infection and renal outcome in patients with chronic kidney disease stage 3-5. *Sci Rep*. 2020;**10**(1):19460. [PubMed ID: 33173137]. [PubMed Central ID: PMC7655801]. <https://doi.org/10.1038/s41598-020-76520-5>.
30. Ganz T. Iron and infection. *Int J Hematol*. 2018;**107**(1):7-15. [PubMed ID: 29147843]. <https://doi.org/10.1007/s12185-017-2366-2>.
31. Trautner BW, Darouiche RO. Role of biofilm in catheter-associated urinary tract infection. *Am J Infect Control*. 2004;**32**(3):177-83. [PubMed ID: 15153930]. [PubMed Central ID: PMC2963581]. <https://doi.org/10.1016/j.ajic.2003.08.005>.
32. Gunardi WD, Karuniawati A, Umbas R, Bardosono S, Lydia A, Soebandrio A, et al. Biofilm-producing bacteria and risk factors (gender and duration of catheterization) characterized as catheter-associated biofilm formation. *Int J Microbiol*. 2021;**2021**:8869275. [PubMed ID: 33688348]. [PubMed Central ID: PMC7920707]. <https://doi.org/10.1155/2021/8869275>.
33. Aazam K, Hassan M, Mahbobe M. Detection of uropathogenic *Escherichia coli* virulence factors in patients with urinary tract infections in Iran. *Afr J Microbiol Res*. 2012;**6**(39):6811-6. <https://doi.org/10.5897/ajmr12.1462>.
34. Oliveira FA, Paludo KS, Arend LN, Farah SM, Pedrosa FO, Souza EM, et al. Virulence characteristics and antimicrobial susceptibility of uropathogenic *Escherichia coli* strains. *Genet Mol Res*. 2011;**10**(4):4114-25. [PubMed ID: 22057993]. <https://doi.org/10.4238/2011.October.31.5>.
35. Tabatabaie A, Badamchi A, Ohadi E, Darbandi A, Javadinia S, Minaeian S, et al. Determination of virulence and quinolone resistance genes and biofilm production among uropathogenic *Escherichia coli* strains isolated from clinical specimens in Iran. *Rev in Med Microb*. 2021;**33**(3). <https://doi.org/10.1097/MRM.0000000000000252>.
36. Zamani H, Salehzadeh A. Biofilm formation in uropathogenic *Escherichia coli*: association with adhesion factor genes. *Turk J Med Sci*. 2018;**48**(1):162-7. [PubMed ID: 29479978]. <https://doi.org/10.3906/sag-1707-3>.