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**Research Article** 

# Investigation of P Fimbriae Presence in *Escherichia coli* Strains Isolated from Urine Samples in Human, and Their Antibacterial Resistance

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# Abstract

**Background:** In Turkey, *Escherichia coli* causing urinary tract infections (UTI) do not have current data on antibacterial resistance, extended spectrum β-lactamase (ESBL) enzymes types and virulence factors.

**Objectives:** In this study, the isolation of *E. coli* from patients with UTI, determination of antimicrobial resistance of isolates, and investigation of the presence of ESBL enzymes and pyelonephritis-associated P fimbriae types, were aimed.

**Methods:** Isolation *E. coli* from urine samples and identification were performed with conventional methods. Antibiotic resistance profiles were performed with the disc diffusion test (DDT). The presence of ESBL enzyme was investigated by the double disk diffusion test (DDDT). The presence of ESBL genes and P fimbriae genes were also investigated with PCR. The Chi squared test was used to compare the relationship between antibiotic susceptibility and resistance status in ESBL producing or non-producing and P fimbriae carrying and non-carrying isolates.

**Results:** It was determined that 50.3% and 41.9% of isolates were resistant to ciprofloxacin, trimethoprim sulfamethoxazole. It was demonstrated that 49.7% of isolates were phenotypically ESBL positive, while 59.4% were genetically positive. The percentages of genes detected with PCR were 32.3%, 31.6%, 21.1%, 2.3% for *bla*TEM, *bla*CTX, *bla*OXA, *bla*SHV, respectively. A total of 51 *pap* genes were detected in 27.0% (36/133) of the total isolates.

**Conclusions:** Extended spectrum  $\beta$ -lactamase producing *E. coli* infections associated with UTI increased in last years in Turkey. Further studies should be designed to investigate plasmids of ESBL positive *E. coli* stains isolated from UTI and whether plasmids are responsible for ESBL production.

Keywords: Antibiotic Resistance, Escherichia coli, Virulence Factors

# 1. Background

Exointestinal pathogenic *Escherichia coli* are one of the most important causes of urinary tract infections (UTI). Since the bacteria have a large number of virulence factors and due to the ability to develop rapid resistance to antibiotics, *E. coli* causing UTI is called uropathogenic *E. coli* (UPEC). It has been reported that UTI are placed at the top of nosocomial and community-acquired infections. *Escherichia coli* in community-acquired UTI are reported to be the most common pathogen in Turkey (1, 2).

The most important virulence factor of *E. coli* that is commonly isolated in UTIs is pyelonephritis-associated P fimbriae (*pap*). Adhesion was the most important task of P fimbriae. P fimbriae; consists of two subunits, a large subunit, *pap*A and the gene *pap*G. The pap gene cluster consists of 11 genes encoding the foremost component of the pilus rod (papA), which determines 11 different serogroups, and a terminally located adhesion, papG(3, 4).

Antibacterial resistance is a problem faced by the medical world and is getting more and more important every day. *Escherichia coli* is known to be one of the important beta lactamase producers. Bacteria that produce the betalactamase enzyme have a serious resistance mechanism that neutralizes the beta-lactam antibiotics by hydrolysis. At least 350 different beta-lactamase enzymes have been identified. One of the most important enzyme groups are the plasmid encoded extended spectrum beta lactamases (ESBL). Extended spectrum beta lactamases are a group of enzymes that hydrolyze antibiotics, including those containing new cephalosporin. They are found in a significant percentage of *E. coli* isolates (5). Especially in the last decade, multi-drug resistant *E. coli* isolates producing ES-BLs are an important problem in Turkey as well as in the

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world (5, 6). Health Protection Agency of Turkey reported that the rate of ESBL-producing *E. coli* isolates were 21% in 2011 (6).

# 2. Objectives

This study was aimed at the isolation of *E. coli* from the urine of patients with UTI using conventional methods, the determination of antibiotic resistance by the disc diffusion test (DDT) and investigation of ESBL enzymes using the double disk diffusion test (DDDT) of isolates; presence of the ESBL genes (*bla*TEM, *bla*CTX, *bla*OXA, *bla*SHV) and P fimbriae genes (*papA*, *papG*) were also investigated using the polymerase chain reaction (PCR).

# 3. Methods

# 3.1. Ethics Statement

Ethics Committee of Adnan Menderes University Faculty of Medicine approved the study protocol (413/2014).

### 3.2. Patients

Uropathogenic *E. coli* isolates were collected from the Infectious Diseases and Clinical Microbiology Laboratory of Adnan Menderes University Hospital, during a twelve months period in 2015. In the study, 20 - 30 mL of morning sterile urine was taken from patients who had not received antibiotic treatment for at least 15 days as material. Materials were delivered to the lab within latest 20 minutes at the latest.

#### 3.3. Isolation of UPEC

Escherichia coli for the study were isolated from patients with urinary infections hospitalized in Adnan Menderes University Hospital. Collected samples were cultured in blood agar (Merck, Germany) and EMB Agar (Merck, Germany) and incubated for 24 hours at 37°C. Suspicious colonies were chosen and streaked on EMB Agar passaged. Gram staining, oxidase and other biochemical tests (indole, methyl red-voges proskauer, and citrate) were then performed (7). Isolation and identification were performed by conventional biochemical tests, as well as by PCR. We examined the mid-stream clean-catch urine samples of 264 adult patients with the diagnosis of acute uncomplicated UTI. Microorganisms were included in our work when they occurred as a pure culture and at a concentration of more than 105 CFU/ml. The isolates were kept in brain heart infusion broth (Merck, Germany) with 20% glycerol at -20°C.

3.4. Antimicrobial Susceptibility Test and Double Disk Diffusion Test

The Kirby Bauer disc diffusion method was used to detect antibiotic susceptibilities of isolates on Mueller-Hinton agar (Oxoid Ltd, UK) according to recommendations of the Clinical and Laboratory Standards Institute (CLSI)(8). The following Bioanalyse AST (Turkey) discs were used: Amoxicillin-clavulanate (20/10  $\mu$ g) was placed in the center of the plate with ceftriaxone (30  $\mu$ g), cefotaxime (30  $\mu$ g), cefpodoxime (10  $\mu$ g), aztreonam (30  $\mu$ g), amikacin (30  $\mu$ g), gentamicin (10  $\mu$ g), piperacillin-tazobactam (100/10  $\mu$ g), ampicillin/sulbactam (10/10  $\mu$ g), cefazolin (30  $\mu$ g), cefepime (30  $\mu$ g), cefuroxime (30  $\mu$ g), cefoperazone sulbactam (30  $\mu$ g), imipenem (10  $\mu$ g), ciprofloxacin (5  $\mu$ g), norfloxacin (10  $\mu$ g), colistin (10  $\mu$ g), nitrofurantoin (300  $\mu$ g), trimethoprim-sulfamethoxazole (1.25/23.75  $\mu$ g), fosfomycin (200  $\mu$ g). The CLSI recommends that, in DDDT a zone size of  $\geq 5$  mm difference between the cephalosporin disks should be considered as significant, and indicates ESBL production (8). Escherichia coli ATCC 25922 strains were used as quality control strains in antibiotic susceptibility tests.

# 3.5. DNA Extraction

We inoculated all isolates aerobically on tryptic soy agar (Merck, Germany) for 18 - 24 hours at 37°C. Genomic bacterial DNA was extracted from bacteria using a commercial DNA extraction kit (InstaGene Matrix, Bio-Rad, USA) according to the manufacturer's protocol. For this, isolated bacterial colony suspended in 1 ml sterile distilled water. After centrifugation at 10.000 rpm the supernatant was removed. The pellet was suspended in 200  $\mu$ L of Instagene matrix and vortexed, followed by heating at 56°C for 15 min. The samples were vortexed again and heated at 100°C for 8 min and then centrifuged. The prepared DNA was stored at -20°C and 2  $\mu$ L supernatant was used template DNA in each PCR.

### 3.6. Polymerase Chain Reaction

The primers used in this study are listed in Table 1. We performed PCR amplifications in 50  $\mu$ L reaction mixtures containing final concentrations of 10X Taq buffer solution 1X, 25 mM magnesium chloride (MgCl<sub>2</sub>) 2 mM, 10 mM dNTP 0.2 mM, 100 pmol primer 0.4 pmol (for each), 5U Taq DNA polymerase (Fermentas) 1.5 and 2  $\mu$ L of each template DNA. We amplified each gene separately. The DNA was amplified using the following protocol: held at 94°C for 3 minutes, followed by 35 cycles of denaturation (94°C for 30 seconds), annealing (44 - 63°C for 30 seconds) and extension (72°C for 1 minute), with a single final extension of 7 minutes at 72°C. Using a 1.5% agarose gel for 45 minutes at 100 V and stained

with Safe View (ABM, Canada), we visualized PCR products by electrophoresis. The primers used, together with sequences, amplicon lengths, target genes and sources are shown in Table 1. (9-15).

#### 3.7. Statistical Analyses

Statistical analyses were performed by the commercial statistical software SPSS version 21.0 (SPSS Inc., USA). The Chi square test was used to compare the relationship between antibiotic susceptibility and resistance status in ESBL producing and non-producing strains, and in isolates carrying and non-carrying P fimbriae. The results were evaluated at a confidence interval of 95%. A P value < 0.05 was considered as significant.

# 4. Results

# 4.1. Isolation and Identification

From 264 urine specimens 133 (133/264: 50.4%) *E. coli* were isolated and identified using standard biochemical tests. The results were confirmed genotypically. Distribution of infected *E. coli* isolates according to sex demonstrated that 57.1% (76/133) of the patients were female and 42.9% (57/133) were male.

# 4.2. Antimicrobial Susceptibility Test and Double Disk Diffusion Test

It was determined that 133 *E. coli* isolates were 50.3% resistant to ciprofloxacin, 42.1% to trimethoprim sulfamethoxazole, 13.5% to cefazolin, 9.8% to amoxicillinclavulanic acid and 8.3% to norfloxacin resistant; resistance rates against the other 15 antibiotics used ranged from 5.3% to 0.8% (Table 2).

According to DDDT results 49.6% of isolates were found to be ESBL positive while 50.4% were ESBL negative. Of the 66 isolates identified as ESBL positive, 71.1% were ciprofloxacin, 63.6% were trimethoprimsulfamethoxazole, 10.6% were cefazolin, 9.1% were amoxicillin-clavulanic acid, gentamycin, norfloxacin and piperacillin-tazobactam resistant; resistance rates against the other 13 antibiotics ranged from 6.0% to 1.5% (Table 3).

Of the 67 isolates identified as ESBL negative, 29.8% were ciprofloxacin, 23.9% were trimethoprimsulfamethoxazole, 16.4% were cefazolin, 9.0% were amoxicillin-clavulanic acid and norfloxacin and 8.5% were resistant to nitrofurantoin; the resistance rates of the other 14 antibiotics varied between 6.0% and 0.0% (Table 4).

### 4.3. Examination of GSBL Enzyme Types

A total of 133 *E. coli* were examined for their ability to carry the ESBL genes by PZR. TEM, SHV, CTX-M and OXA primers were used for typing the ESBL enzyme. These isolates had four types of ESBL genes. 59.4% (79/133) of the isolates carried at least one of the examined ESBL genes. Extended spectrum beta lactamases genotypes were found among these isolates with the predominance of TEM-type (43/133: 32.3%), followed by CTX-M-type (42/133: 31.6%), OXA-type (28/133: 21.1%) and SHV-type (3/133: 2.3%) (Figure 1).

# 4.4. Examination of P Fimbriae Genes (papA and papG)

A total of 51 *pap* (*papA* + *papG*) genes were detected in 27.0% (36/133) of isolates; 12.0% (16/133) of them were found to be *papA* while 26.3% (35/133) were found to be *papG*. *papG* allele genotypes were found among these isolates with the predominance of allele II (33/133: 24.8%), followed by allele III (7/133: 5.3%) and allele I (1/133: 0.8%). Allele II and III were present in 26.3% (35/133) of the isolates.

#### 4.5. Statistical Analyses

As a result of the Chi-square test conducted to compare antibiotic susceptibility and resistance status, (1) it was detected that ciprofloxacin and trimethoprimsulfamethoxazole antibiotics were significantly different between the groups, with regards to ESBL-producing and nonproducing isolates, (2) it was detected that there was a significant difference between the ciprofloxacin resistance and *papG* presence in the groups, with regards to P fimbriae carrying and non-carrying isolates (P < 0.05).

# 5. Discussion

Escherichia coli isolates from human clinical specimens especially UTI cause significant public health problems because the majority of these strains carry multidrug resistance genes (16). However, mobile genetic elements for example plasmids can spread ESBL-related genes via horizontal gene transfer in other Gram-negative bacteria (17). The vast majority of UTIs are caused by a single species of bacteria, and E. coli have been reported as the most frequently isolated infectious agent from UTI in many studies conducted in Turkey and around the world. In this study, E. coli was isolated in a high proportion (50.4%) of UTIs, similar to the above studies; a large portion of these isolates (57.1%) were obtained from female patients. This is consistent with previous findings which report that UTI is more common in females (2, 6, 16, 17). This difference could be due to anatomic differences, hormonal effects or several clinical factors.

Primer	Sequence (5' - 3')	Amplicon Length, bp	Target Gene	Source
usp-F	CCGATACGCTGCCAATCAG	004		0
usp-R	ACGCAGACCGTAGGCCAGAT	884	usp	g
SHV-F	CGCCTGTGTATTATCTCCCT	202	blaSHV	10
SHV-R	CGAGTAGTCCACCAGATCCT	233		
CTX-M-F	CGCTGTTGTTAGGAAGTGTG	569	blaCTX	10
CTX-M-R	GGCTGGGTGAAGTAAGTGAC	303	blactifi	10
TEM-F	ATAAAATTCTTGAAGACGAAA	1080	blaTEM	11
TEM-R	GACAGTTACCAATGCTTAATCA			
OXA-F	ACCAGATTCAACTTTCAA	598	blaOXA	12
OXA-R	TCTTGGCTTTTATGCTTG			
рарАН-F	ATGGCAGTGGTGTCTTTTGGTG	72.0	papA	13
papAH-F	CGTCCCACCATACGTGCTCTTC	,20		
papG allel I-F	TCGTGCTCAGGTCCGGAATTT	461	ngnC allel I	14
papG allel I-F	TGGCATCCCCCAACATTATCG	401	pupo anci i	14
papG allel II-F	GGGATGAGCGGGCCTTTGAT	190	papG allel II	15
papG allel II-R	CGGCGGCCCAAGTAACTCG			
papG allel III-F	GGCCTGCAATGGATTTACCTGG	258	papG allel III	15
papG allel III-R	CCACCAATTGACCATGCCAGAC			
pαpG allel II/III-F	CTGTAATTACGGAAGTGATTTCTG	1070	papG allel II-III	12
papG allel II/III-R	ACTATCCGGCTCCGGATAAACCAT	1070		13



Figure 1. 1 - 3 - 5 - 7: Gel electrophoresis images of GSBL (SHV, CTX, OXA, TEM) positive isolates. 2 - 4 - 6 - 8: Negative control (mastermix without DNA) M: 100 bp DNA ladder (Vivantis)

It is important to have knowledge about the resistance to antimicrobials at the country or even region level during UTI treatment. As a result, there is always a need for studies to determine the susceptibility profiles of agents. Numerous studies have been carried out on the antibiotic resistance of UPECs in Turkey and in the world and continue to be performed. The most commonly used drugs in the management of UTI are trimethoprimsulfamethoxazole and quinolones. In the study, *E. coli*  isolates were found to be most resistant to ciprofloxacin, trimethoprim sulfamethoxazole, cefazolin, amoxicillinclavulanic acid and norfloxacin, which are the most frequently, used antibiotics in empirical therapy. These findings have also been reported in other studies (18, 19). It is recommended to use other antibiotics such as imipenem, aztreonam, colistin in the treatment. As a result, these antibiotics, which are now less resistant, can be used for the treatment of UTI's. This may be due to increased use of



Figure 2. The gel electrophoresis images of the *pap* genes. 1: *pap*A positive isolate (720 bp) 3: *pap*G allele I positive isolate (461 bp) 5: *pap*G allele II positive isolate (190 bp) 7: *pap*G allele II positive isolate (258 bp) 9: *pap*G allele II - III positive (1070 bp) 2 - 4 - 6 - 8: negative control (mastermix without DNA) M: 100 bp DNA ladder (Vivantis<sup>®</sup>)

Table 2. Antibiotic Susceptibility Test Results <sup>a</sup>					
Antibiotic	Sensitive	Intermediate	Resistant		
Ciprofloxacin	63 (47.4)	3 (2.3)	67 (50.3)		
Trimethoprim- sulfametaxazole	76 (57.1)	1(0.8)	56 (42.1)		
Cefazolin	115 (86.5)	-(0.0)	18 (13.5)		
Amoxicillin-clavunic acid	120 (90.2)	-(0.0)	13 (9.8)		
Norfloxacin	122 (91.7)	-(0.0)	11 (8.3)		
Gentamicin	126 (94.7)	-(0.0)	7 (5.3)		
Cefuroxime	126 (94.7)	-(0.0)	7 (5.3)		
Nitrofurantoin	126 (94.7)	-(0.0)	7(5.3)		
Piperacillin-tazobactam	119 (89.4)	7(5.3)	7 (5.3)		
Cefotaxime	126 (94.7)	-(0.0)	7(5.3)		
Fosfomycin	127 (95.5)	-(0.0)	6(4.5)		
Cefepime	127 (95.5)	-(0.0)	6(4.5)		
Cefoperazone-sulbactam	122 (91.7)	5 (3.8)	6(4.5)		
Ampicillin-sulbactam	124 (93.3)	4 (3.00)	5 (3.7)		
Amikacin	127 (95.4)	3 (2.3)	3 (2.3)		
Cefpodoxime	130 (97.7)	-(0.0)	3 (2.3)		
Ceftriaxone	129 (97.0)	1(0.8)	3 (2.3)		
Aztreonam	132 (99.2)	-(0.0)	1(0.8)		
Colistin	132 (99.2)	-(0.0)	1(0.8)		
Imipenem	132 (99.2)	-(0.0)	1(0.8)		

 Table 3. Antibiotic Susceptibility and Resistance Rates in ESBL Positive Isolates<sup>a</sup>

1 5			
Antibiotic	Sensitive Intermediate Resistant		
Ciprofloxacin	19 (28.9)	-(0.0)	47 (71.1)
Trimethoprim- sulfametaxazole	22 (33.4)	2(0.0)	42 (63.6)
Cefazolin	59 (89.4)	-(0.0)	7 (10.6)
Amoxicillin-clavunic acid	60 (90.9)	-(0.0)	6 (9.1)
Gentamicin	60 (90.9)	-(0.0)	6 (9.1)
Norfloxacin	60 (90.9)	-(0.0)	6 (9.1)
Piperacillin-tazobactam	54 (81.2)	6 (9.1)	6 (9.1)
Cefuroxime	62 (94.0)	-(0.0)	4(6.0)
Cefepime	62 (94.0)	-(0.0)	4(6.0)
Cefoperazone-sulbactam	56 (84.9)	6 (9.1)	4(6.0)
Cefpodoxime	63 (95.5)	-(0.0)	3(4.5)
Ceftriaxone	63 (95.5)	-(0.0)	3(4.5)
Fosfomycin	63 (95.5)	-(0.0)	3(4.5)
Cefotaxime	63 (95.5)	-(0.0)	3(4.5)
Amikacin	62 (94.0)	2 (3.0)	2 (3.0)
Aztreonam	64 (97.0)	-(0.0)	2 (3.0)
Imipenem	64 (97.0)	-(0.0)	2 (3.0)
Colistin	65 (98.5)	-(0.0)	1 (1.5)
Ampicillin-sulbactam	65 (98.5)	-(0.0)	1(1.5)
Nitrofurantoin	65 (98.5)	-(0.0)	1(1.5)

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

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these antibiotics, and resistance transfer (17).

A different problem is the increased occurrence of ESBL producing organisms causing UTI, and the emergence of severe health problems due to antibiotic resistance. Extended spectrum beta lactamases can be transferred to other strains via plasmid and transposon, further increasing the significance of this ESBL-resistant isolates (3, 15, 19). Senbayrak et al. reported a significant

Antibiotic	Sensitive	Intermediate	Resistant
Ciprofloxacin	44 (65.7)	3 (4.5)	20 (29.8)
Trimethoprim- sulfametaxazole	51 (76.1)	-(0.0)	16 (23.9)
Cefazolin	56 (83.6)	-(0.0)	11 (16.4)
Amoxicillin-clavunic acid	61 (91.0)	-(0.0)	6(9.0)
Norfloxacin	61 (91.0)	-(0.0)	6(9.0)
Nitrofurantoin	61 (91.5)	-(0.0)	6 (8.5)
Cefotaxime	63 (94.0)	-(0.0)	4(6.0)
Fosfomycin	64 (95.5)	-(0.0)	3(4.5)
Ampicillin-sulbactam	61 (91.5)	4(6.4)	1(1.5)
Gentamicin	66 (98.5)	-(0.0)	1(1.5)
Piperacillin-tazobactam	66 (98.5)	-(0.0)	1(1.5)
Cefepime	66 (98.5)	-(0.0)	1(1.5)
Ceftriaxone	66 (98.5)	-(0.0)	1(1.5)
Amikacin	66 (98.5)	-(0.0)	1(1.5)
Cefuroxime	67 (100.0)	-(0.0)	-(0.0)
Cefoperazone-sulbactam	67 (100.0)	-(0.0)	-(0.0)
Cefpodoxime	67 (100.0)	-(0.0)	-(0.0)
Aztreonam	67(100.0)	-(0.0)	-(0.0)
Imipenem	67 (100.0)	-(0.0)	-(0.0)

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

increase in ESBL prevalence from 12.5% to 44.7% in both male and female hospitalized patients, between 2004 -2012 (20). Toka Ozer et al. reported the rate of ESBL production in E. coli strains was 13.1% in 2018 (1). Antibiotic resistance rates have also been reported to increase for ESBL producing E. coli. These rates were higher than 70% for fluoroquinolones, amoxicillin-clavulanic acid and trimethoprim-sulfamethoxazole. The prevalence of ESBLproducing E. coli for inpatients in Switzerland is 6.6% (21), 10.4% in Saudi Arabia (22) and 52.9% in China (23). In our study, 49.7% of the isolates were phenotypically ESBL positive. Comparing antibiotic resistance profiles of ESBL producing and non-ESBL producing E. coli strains, there is great similarity in the order of resistant antibiotics, although there is a difference in resistance rates. Resistance rates were found to be higher in the presence of ESBL.

In ESBL-producing *E. coli*, resistance to different antibiotics is often observed because the coding genes are usually in the same plasmids (24). According to results of this study, 59.4% of the isolated bacteria carried at least one of the examined ESBL genes. The percentages of genes detected were 32.3%, 31.6%, 21.1%, 2.3% for *bla*TEM, *bla*CTX, *bla*OXA, *bla*SHV, respectively. Recent studies have unveiled the majority of *bla*CTX among ESBL positive phenotype all over the world. In these studies, the OXA gene was also not detected (25, 26). Mohajeri et al. in Iran demonstrated that 93.3%, 68.2%, 43.2%, 31.8%, and 22.7% of CTX-M, SHV, TEM, OXAI and OXAII genes were present in UPEC (27). In our study, the TEM gene was shown to be of a high frequency compared to CTX, OXA and SHV genes. However, further studies are required to identify other genes in ESBLs producing *E. coli* isolates.

P fimbriae are common virulence factors of UPEC, which play an important role in the pathogenesis of ascending UTIs and pyelonephritis in humans (2, 5). P fimbriae consist of heteropolymeric fibres of different protein subunits, encoded by the *papA*-K gene operon (28). *papA* is an important structural subunit of the pilus associated with pyelonephritis, and has critical prescription for pilus synthesis. papG is Gal(1-4)Gal-specific pilus tip adhesin molecule. papG has three alleles. Allele I is known as J96associated, allele II is known as pyelonephritis-associated, while allele III is known as cystitis-associated papG variant (5, 28). Of our isolates, 27% (36 of them) carried a total of 51 genes (papA or papG). In our isolates, 26.3% of papG was detected while 12% of papA was detected. A study from Ankara in Turkey, parallel to our study, revealed that almost 23% E. coli isolates were positive for pap genes (29). In Iran, Mohajeri (30), Farshad and Emamghorashi (31) and Rahdar et al. (32) reported the frequencies of pap genes as 20.5%, 30.2%, 57%, respectively. These differences may be due to the differences in geographical regions. The pap gene encodes for P-fimbriae that allows bacteria to adhere to epithelial surfaces and protect them against urine lavage, thus allowing them to ascend further to even cause serious disease (30).

It was reported that was extended spectrum  $\beta$ lactamase producing *E. coli* infections alarmingly increased in recent years in Turkey (1, 20). On the other hand, inappropriate and excessive use of antibiotics has resulted drug resistance. This limits treatment options. However, because of the rates of antibiotic resistance between regions are very variable, it is important to know the resistance profile of the region in the selection of antibiotics to be preferred in empirical treatment. When regional differences in antibiotic resistance rates are taken into consideration, antibacterial resistance should be regularly monitored in each geographical area in order to determine appropriate treatment strategies and reduce treatment costs.

# 5.1. Conclusions

Extended spectrum -lactamase producing E. coli infections associated with UTI increased in last years in Turkey. Further studies should be designed to investigate plasmids of ESBL positive *E. coli* stains isolated from UTI and whether plasmids are responsible for ESBL production

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#### Footnotes

Authors' Contribution: Conception and design of study: Emel Inegol Paykoc and Suheyla Turkyilmaz; acquisition of data (laboratory or clinical): Emel Inegol Paykoc and Suheyla Turkyilmaz; data analysis and/or interpretation: Emel Inegol Paykoc; drafting of manuscript and/or critical revision: Emel Inegol Paykoc and Suheyla Turkyilmaz; approval of final version of manuscript: Emel Inegol Paykoc and Suheyla Turkyilmaz.

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Jundishapur J Microbiol. 2018; 11(9):e66119.

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