

# Risk factors for resistance in urinary tract infections in women in general practice: A cross-sectional survey

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#### Title: Risk factors for resistance in urinary tract infections in women in general practice: A cross-sectional

#### survey

## Running title: Risk factors for resistance in urinary tract infections

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## Highlights

- Individual factors associated with drug-resistant UTIs in general practice are scarce
- Only two factors were associated with drug-resistant UTIs:
  - o Penicillin use in the previous three months
  - o Exposure to a person returning from an area with a high risk of drug resistance
- A protective role of raw meat consumption has been found and should be explored

## Summary

## Objectives

In 2012 and 2013, a cross-sectional survey was conducted in women visiting a general practitioner for a urinary tract infection (UTI) to i) describe the patterns of antibiotic resistance of Enterobacteriaceae involved in community-acquired UTIs and ii) identify the factors associated with UTIs due to a multi-drug-resistant Enterobacteriaceae (MDREB).

## Methods

Urine analyses were performed systematically for all adult women presenting with signs of UTI. Characteristics of women with UTI due to MDREB were compared to those with UTI due to non-MDREB. Weighted logistic regressions were performed to adjust for the sampling design of the survey.

## Results

Significant factors associated with MDREB included the use of penicillin by the patient in the last three months (OR = 3.1; [1.2-8.0]); having provided accommodation in the previous 12 months to a resident from a country at high risk for drug resistance (OR = 4.0; [1.2-15.1]); and the consumption of raw meat within the previous three months (OR = 0.3; [0.1-0.9]).

## Conclusions

In the community, antibiotic use and exposure to a person returning from an area with a high risk of drug resistance are associated with UTIs due to MDREB. The potentially protective role of raw meat consumption warrants further study.

Key words: urinary tract infection, epidemiology, antibiotic resistance, risk factors, general population, general practitioner

## Introduction

## ACCEPTED MANUSCRIPT

Urinary tract infections (UTIs) are among the most common bacterial infections. In the United States, one-third of all women have had at least one physician-diagnosed presumed UTI by the age of 26 years (1). UTIs are typically caused by Enterobacteriaceae, such as *Escherichia coli* (*E. coli*), which are involved in 70-80% of cases (2).

The epidemiology of antimicrobial resistance varies among countries, particularly due to the impact of antibiotic prescriptions (3-5). In France, in 2009, the resistance rates of *E. coli* reached 55% for amoxicillin, 18% for fluoroquinolone, 8% for aminoglycoside and 10% for third-generation cephalosporin in healthcare-related infections (6). In 2011, the incidence of patients with at least one diagnostic sample testing positive for extended-spectrum  $\beta$ -lactamase Enterobacteriaceae (ESBLE) was 0.46 per 1000 days of hospitalization in French acute care hospitals (7). Data are scant for community-onset UTIs, as urine analysis is not mandatory in the treatment of uncomplicated cystitis. Thus, in previous epidemiological studies, urine cultures were not obtained for all women presenting with symptoms of UTI, but rather for women with complicated UTIs or UTIs seen in healthcare settings.

The empirical therapy of UTIs should be guided by knowledge of antimicrobial resistance and by identifying risk factors for resistance (8). Known risk factors include previous hospitalization, previous antibiotic use, urinary catheterization and recurrent UTIs (9, 10). Other risk factors have been suspected, although they are less clearly established, such as international travel, living with pets, eating fish or raw meat, and swimming (11-13). Almost all of these risk factors have been identified in hospitalized patients. However, most UTIs occur in the community and are frequently treated in the absence of urine analysis. Data on the epidemiology of drug resistance in this setting are scant. Therefore, to identify factors associated with antimicrobial resistance in UTIs caused by Enterobacteriaceae in the French community, we conducted a cross-sectional survey in women visiting a general practitioner (GP) for a suspected UTI. Urine analysis was performed for all included individuals.

#### Design and study population

The study population was derived from a national cross-sectional prospective survey, titled Drug resistant Urinary Tract Infection (Druti), that was conducted in France in 2012 and 2013 by general practitioners (GPs) of the Sentinelles network (14) to estimate the annual incidence of UTIs due to antibiotic-resistant Enterobacteriaceae in women visiting a GP for a suspected UTI (the results were pending at the time that this manuscript was written). In brief, a two-step sampling design was performed. For the first step, GPs of the Sentinelles network were randomly selected for participation and stratified into five groups according to five regions of France: north, east, west, southwest, and southeast. For the second step, the GPs prospectively included all female patients 18 years of age and older who had presented in the last seven days with at least one of the following UTI symptoms: dysuria, frequent urination, or urgency of urination. Those patients who agreed to participate and had not taken an antibiotic in the last seven days were included. A urine sample was collected from all participants, and urine culture performed on all samples at the same laboratory. The GPs were blinded to the urine culture results.

For each included patient, the GP administered a questionnaire that was completed during the consultation. It included questions regarding the patient's demographic characteristics (age, household members, and nationality), clinical status (chronic diseases and comorbidities, particularly pregnancy, urinary tract disorder, previous UTIs, and urinary catheterization), history of pharmacological treatments before consultation (especially antibiotic use by the patient or a member of the patient's household), previous hospitalization, contact with a professional or personal healthcare worker by the patient or a member of the patient's household, international travel, contact with an animal (with information on possible antibiotic treatments administered to this animal) and the consumption of raw meat. To recover any missing data, a trained investigator telephoned the GP and the patient to verify the information within two weeks of inclusion.

According to French recommendations, complicated UTI was defined as UTI occurring in a woman with urinary tract anomalies, pregnant, aged 65 years and older or treated for a chronic disease (diabetes, cancer, renal insufficiency). To define countries at high risk of drug resistance, a wide definition was chosen based on previous publications (2, 15, 16); if a country was located in Central/South America, Eastern/Southern Europe, Africa or Asia, it was considered to be high risk.

#### Sample size

The sample size of the Druti study was calculated for the objective of incidence. A total of 280 *E. coli*-positive samples was calculated to estimate the incidence rate of UTIs caused by fluoroquinolone-resistant *E. coli*, according to an

### expected proportion of fluoroquinolone resistance of 18% and a precision of 4.5% (6). Based on the assumption of

70% positive urine cultures for suspected UTIs and 77% positive cultures caused by *E. coli* (2), it was necessary to include 520 urine samples. Considering a median number of consultations for UTIs of 16 per year per GP and a proportion of eligible patients of 34%, it was necessary to recruit 96 GPs to obtain 520 included patients.

#### Bacteriological analyses

Urine cultures, bacterial identification and susceptibility testing were performed at the Department of Microbiology of Ambroise Paré University Hospital, Paris. Leukocyte and erythrocyte counts per mm<sup>3</sup> of urine were acquired using Kova<sup>®</sup> slides (Greiner bio one, Courtaboeuf, France). Microscopic examination after Gram staining was carried out for each sample, and the presence of bacteria was noted. Urine cultures were performed in a semi-quantitative manner using calibrated loops for plating. Isolated colonies were obtained for identification and susceptibility testing. All samples were plated on a chromogenic medium for urine analysis (uriselect4©; Biorad, Marnes-la-Coquette, France). When Gram-positive bacteria were observed at the microscopic examination, a selective medium for Gram-positive bacteria (CAN agar + 5% sheep blood, bioMérieux, Craponne, France) was added. Media were incubated at 35°C for 24 hours. In the event of discordance between the microscopic examination and the culture results, media were incubated for 48 hours. Urine samples were maintained at 4°C for one week.

After incubation, the number of bacteria and the identity of the infecting microorganism were determined, and its antimicrobial susceptibility was tested. Identification was carried out by matrix-associated laser desorption ionization time-of-flight mass spectrometry (MALDI TOF, Bruker, Ettlingen, Germany). Antimicrobial susceptibility testing was conducted using the disk diffusion method according to the recommendations of the French Society of Microbiology and the European Committee on Antimicrobial Susceptibility Testing (17, 18). ESBL production was detected by the API 20E and VITEK 2 systems (bioMérieux, Marcy l'Etoile, France), the double-disc synergy test, the MicroScan ESBL plus ESBL confirmation panel (Dade Behring, Sacramento, CA), and the Etest ESBL (AB Biodisk, Piscataway, NJ) (17, 19, 20). ESBL was identified by specific PCR and sequencing, as previously described (20, 21).

Isolates were considered to be resistant when tests revealed resistance or intermediate susceptibility to a particular antimicrobial agent; otherwise, they were considered to be susceptible. Multi-resistance was defined as acquired resistance to at least three of the following antimicrobial categories: penicillins, penicillins and  $\beta$ -lactamase inhibitors, antipseudomonas penicillins and  $\beta$ -lactamase inhibitors, monobactams, carbapenems, non-extended spectrum cephalosporins, extended-spectrum cephalosporins, anti-MRSA (methicillin-resistant *Staphylococcus aureus*) cephalosporins, cephamycins, aminoglycosides, tetracyclines, glycylcyclines, folate pathway inhibitors, fluoroquinolones, phenicols, phosphonic acids and polymyxins (antimicrobial categories were dropped if species had intrinsic resistance) (22). This definition of multi-resistance Enterobacteriaceae (MDREB) (acquired resistance to at

least three classes of antibiotic) is the definition proposed by the ECDC; it responds to a need for standardization at an international level, even if it does not take into account specific antibiotics used preferentially to treat UTIs. It does not overlap with the commonly accepted definition of multi-resistant bacteria, such as MRSA or ESBLE.

#### Statistical analyses

The study sampling plan and design variables (stratum and weighting variables) were taken into account during analyses to permit inference of the findings to the general population (23). Stratified and sampling steps were used to accurately estimate the associated variances. For the first step, all GPs participating in the study in each geographical area were considered to be a random sample of all GPs in that region; thus, the probability of inclusion at the first stage for each geographical stratum was identical for each GP and was equal to the participation rate of the GPs in the survey. For the second step, the GPs prospectively enrolled women with a suspected UTI. Of these eligible women, only those who agreed to participate, had not taken an antibiotic in the last seven days and were able to provide a urine sample were included. Thus, the probability of inclusion at this stage was calculated for each woman according to the number of women with a urine culture result divided by the number of eligible women consulted for a UTI. Finally, estimates were post-stratified using the number of consultations for all GPs in France, according to the national health insurance system (CNAM).

For descriptive analysis, we expressed the estimated proportions of the qualitative variables with 95% confidence intervals (Cls) and the estimated mean with standard error of the quantitative variables in the population. The main outcome variable was UTI due to MDREB, as explained by the associated factors using univariate or multivariate logistic regression models and taking into account the sampling design. Those analyses were also performed for UTIs due to Enterobacteriaceae resistance only to penicillins or folate pathway inhibitors (trimethoprim-sulfamethoxazole). Variable selection was conducted as follows. First, all variables collected were assessed by univariate analysis, and those achieving a *p*-value of < 0.20 (using the Wald test for logistic regression) were included in multivariate analysis. A backward stepwise variable selection procedure was then used to remove factors with a *p*-value of > 0.05. Adjusted odds ratios (ORs) and 95% Cls were calculated for the determinants that remained in the final model. Missing values were indicated and were excluded from the models. Data were collected with EPI-Data and analysed with R 2.10.1 statistical package, particularly the *survey* package (24).

#### Ethical considerations

The study obtained research authorization from the French independent administrative authority protecting privacy and personal data (CNIL), number 911 485 (CCTIRS number 11.474), and from the local human investigation committee of *Ile de France V*.

## Results

## ACCEPTED MANUSCRIPT

Of the 96 expected GPs, 87 participated in the study. During the study period, 1,569 women presented to their GP with symptoms of a UTI. Figure 1 shows the flow chart of the study. Urine samples were collected from 538 of the eligible women. The three main reasons for non-inclusion were unavailability for the transport of the urine sample (34%), lack of time of the GP for inclusion (18%), and previous antibiotic treatment in the last 7 days (14%). A total of 393 urine cultures were positive (74%): 366 urine samples (93%) had only one species of bacteria, and 26 samples (7%) had two or three species. Majority of bacteria were *Enterobacteriaceae* (87%), especially *E. coli* (79%). In summary, 363 urine cultures tested positive for Enterobacteriaceae, thus, a total of 363 women were included to permit analyses.

The median age of the 363 women with Enterobacteriaceae-positive urine cultures was 45 years old. Dysuria was reported in 95% of cases, frequent urination in 92%, urgency of urination in 80%, pelvic pain in 40%, fever in 6%, and vaginal itching in 5% of cases. There was no significant difference between the clinical characteristics (dysuria, frequent urination, urgency associated with urination, pelvic pain, fever, vaginal itching) of the women with a urine culture that tested positive for MDREB and those with a culture that was positive for non-MDREB.

The following bacterial species were identified: *E. coli* (329 samples), *Proteus mirabilis* (16), *Klebsiella pneumoniae* (6), *Citrobacter koseri* (5), *Klebsiella oxytoca* (4), *Enterobacter aerogenes* (1), *Enterobacter cloacae* (1) and *Raoultella planticola* (1). Among the isolates, 68 (19%) were MR. Estimated resistance rates for each antibiotic are listed in Table 1. No differences in the distribution of resistance between the five regions had been shown (for instance, the p value for MDREB distribution across regions was 0.36).

Univariate analysis revealed that the following factors were associated with MDREB UTIs: antibiotic treatment taken by a household member in the previous three months (OR = 2.4; 95% CIs [1.1-5.2]); contact with animals that had taken an antibiotic in the previous three months (OR = 2.9; [1.0-8.1]); having provided accommodation to a resident from a country at high risk of drug resistance in the previous 12 months (OR = 5.0; [1.3-19.0]); and the consumption of raw meat in the previous three months (OR = 0.3; [0.1-0.9]) (Table 2). Prior antibiotic use was higher among women with a urine culture that tested positive for MDREB, although this increase was not statistically significant.

The results of multivariate analysis are reported in Table 3. Significant factors identified in the final model included penicillin use by the patient in the previous three months (OR = 3.1; [1.2-8.0]); having provided accommodation for a resident from a country at high risk of drug resistance in the previous twelve months (OR = 4.0; [1.2-15.1]); and the consumption of raw meat in the previous three months (OR = 0.3; [0.1-0.9]).

Regarding monoresistance, either in univariate or multivariate analyses, only a complicated UTI was associated with resistance to penicillin, while a history of UTIs in the last 12 months was associated with trimethoprim-sulfamethoxazole resistance (Tables 4 and 5).

Among the 538 included women, only six had a UTI due to ESBLE (Table 1). Due to the small number, subgroup analyses were precluded. However, all women with a UTI due to ESBLE presented at least one of the factors identified above, including international travel in the previous twelve months (n = 3) and/or antibiotic treatment in the previous three months (n = 4).

#### Discussion

## ACCEPTED MANUSCRIPT

Among women visiting a GP in a general population setting for a UTI, the factors associated with MDREB infection are related to previous antibiotic use and exposure to a person returning from an area with a high risk of drug resistance. Furthermore, raw meat consumption could have a potentially protective role.

The rate of third-generation cephalosporin susceptibility reported in our study (98%) was in line with the rates observed in other studies that had been published since the end of the present study, i.e., from 96 to 99% (25, 26). For fluoroquinolone, we also observed a susceptibility rate of 98%, while the latest published studies in France reported rates of 89% to 97% (25, 26). In the study with the lowest susceptibility rate estimates, data had been transmitted by a laboratory network based on urine collected mostly for complicated UTIs, thus resulting in higher resistance rates than observed in the general population. ESBLE represented 1.6% of our sample, which is within the range of the most recently published studies, i.e., 0 to 3% of UTIs (25, 26). Thus, our resistance rates were comparable to the rates observed in other recent French studies. In other European countries, the resistance rates of fluoroquinolone ranged from 0.5 to 7.6%, also in line with our data (27). In parallel, the total outpatient quinolone use in France in 2009 was estimated to be 2.0 defined daily dose of quinolones per 1000 habitants per day, while it ranged from 0.48 to 4.13 in other European countries (28).

The strength of the association between drug-resistant infection and antibiotic use by the patient in the present study was almost equal to that found for the risk of ampicillin-resistant *E. coli* infection in a study performed on a UK community (OR = 3.91) (29). However, other studies reported higher odds ratios of 18.6 for previous fluoroquinolone use and fluoroquinolone resistance in febrile *E. coli* UTIs (30), and 23.7 for previous antibiotic use and ESBLE infection (31). In these previous studies, patients were recruited from healthcare facilities, where a higher prevalence of antibiotic exposure and drug resistance are usually observed, possibly accounting for the differences in the magnitude of the observed risk estimates (29-32).

Previous hospitalization and urinary catheterization were not found to be associated with MDREB UTIs in this study. In a previous study of an outpatient population treated in primary healthcare centres and emergency departments for febrile *E. coli* UTI, these factors have been associated with fluoroquinolone resistance (OR = 2.28 for previous hospitalization and OR = 6.02 for urinary catheterization) (30). Again, the lower rates reported in our study may be due to our evaluation of the general population versus the hospital-based populations that have been assessed in previous studies. Unexpectedly, raw meat consumption was shown to be inversely associated with MDREB UTIs. This result contradicts other studies reporting that raw chicken consumption increases the risk of antibiotic-resistant infection (33). However, a recent study has found this same "protective" association between raw meat consumption and CTX-M-producing *E. coli* (34). The authors have suggested that raw meat consumption is linked to nationality and that its consumption may have been more prevalent in the native population than in the foreign population studied, which may have a higher rate of antibiotic resistance. However, in the present study, raw meat consumption was not found to be linked to nationality or occupational grade (data not shown). Thus, further studies are needed to determine whether this association between raw meat consumption and antimicrobial-resistant UTIs is mainly an artefact or is due to a direct protective impact on commensal flora, to a bias related to nationality (raw meat consumption is rather unusual in many countries), or to other socio-professional characteristics.

Several studies have identified an association of geographical factors, including nationality and international travel, with antibiotic-resistant UTIs (9-11, 34). In this study, we did not detect these associations. However, contact with a resident from a country with a high risk of drug resistance was strongly associated with MDREB UTIs, suggesting the cross-transmission of resistant bacteria.

This study had a few limitations. There were lower rates of resistance than expected, particularly for fluoroquinolone, resulting in a lack of statistical power. This possibly explains why we did not confirm other previously identified risk factors for resistant UTIs, such as previous UTI, nationality, travel, previous hospitalization, or urinary catheterization (9-11, 34). For antibiotics with a high rate of resistance (penicillin or trimethoprim-sulfamethoxazole), association with previously identified risk factors was not evidenced in our community study. As carriage of antibiotic resistance of *E. coli* in faecal flora is important, nearly 20% for ampicillin or trimethoprim-sulfamethoxazole (13, 35), it might be hypothesized that individual factors play only an accessory role in resistance acquisition. This acquisition would be more passive than active, resulting from "living there" more than "doing this".

Although the design of our study had the above-mentioned limitations, it allowed for more precise evaluations of the weights of different factors for antibiotic-resistant UTIs in the general population. This study is one of the first to systematically collect urine samples from all women in a population presenting with symptoms of a UTI in general practice. Microbial data on UTIs are typically based on urine samples associated with healthcare-related UTI cases. Our findings expand upon the available data based on healthcare-related populations (34, 36). The sampling design and post-stratification corrected biases stem from subject dropout and geographical repartition. Lastly, another strength of this study was in the standardized collection of data, which limited information bias.

## **Competing interests**

The authors declare that they have no competing interests.

## **Authors' contributions**

LR, SM, SV, TB, BC and TH conceived and designed the experiments. LR, SM and BH performed the experiments. LR, AB and YLS analysed the data. LR, SM, SV, SK, AB, YLS, BH, TB, BC and TH wrote the paper.

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## Figure 1 - Flow chart

## Tables

## **ACCEPTED MANUSCRIPT**

## Table 1: Resistance rates\* for Enterobacteriaceae in urinary tract infections, France, 2012/2013.

	Percentages [95% confidence intervals] **		
Amoxicillin	41 [36-46]		
Amoxicillin-clavulanic acid	9 [6-12]		
C3G (cefotaxim, ceftazidime)	2 [1-4]		
Nalidixic acid	5 [3-8]		
Ofloxacin	5 [3-7]		
Ciprofloxacin	2 [1-4]		
Trimethoprim-sulfamethoxazole	18 [13-24]		
Nitrofurantoin	6 [3-10]		
Fosfomycin	0		
ESBLE	1 [1-4]		
At least one antibiotic	49 [43-54]		
Antibiotic multi-resistant	19 [15-26]		

Enterobacteriaceae were classified as being resistant when testing revealed resistance or intermediate susceptibility to a \* \*\*Proportions were estimated according to the sampling design; \*\*\* Acquired resistance to at least three classes of antibiotics (22);

ESBLE: extended-spectrum β-lactamase Enterobacteriaceae; C3G: third-generation cephalosporin.

# Table 2: Factors associated with antibiotic-multi-resistant Enterobacteriaceae (MDREB) urinary tract infections (UTIs): univariate analysis performed according to the sampling design

	MDREB N=68, (sample size, %**)	Non- MDREB N=295, (sample size, %**)	OR [95% CI]	p value*	Missing value
UTI in the last 12 months	58 (86)	242 (82)	1.3 [0.5-3.6]	0.58	1
Complicated UTI	3 (2)	27 (8)	0.3 [0.1-1.0]	0.06	2
Patients with ≥ four UTIs/year	5 (7)	23 (7)	1.1 [0.4-2.7]	0.91	1
Antibiotic use by the patient in the previous three months Penicillin in the previous three months	20 (29) 9 (18)	55 (20) 20 (7)	1.6 [0.6-4.1] 2.8 [1.0-8.0]	0.35 0.06	1 1
Quinolone in the previous three months Other antibiotics in the previous three months	5 (5) 7 (7)	13 (5) 23 (6)	0.9 [0.3-3.3] 1.2 [0.5-3.0]	0.89 0.72	1 1
Antibiotic use by a household member in the previous three months	17 (27)	39 (13)	2.4 [1.1-5.2]	0.03	1
Hospitalization in the previous 12 months	5 (6)	36 (14)	0.4 [0.2-1.1]	0.07	2
Healthcare contact in the previous month	16 (17)	66 (21)	0.8 [0.4-1.6]	0.52	1
Household healthcare contact in the previous month	16 (20)	46 (14)	1.6 [0.7-3.8]	0.31	1
International travel in the previous 12 months	19 (23)	72 (27)	0.8 [0.4-1.5]	0.50	1
International travel in the previous twelve months in countries at high risk of drug resistance	13 (15)	59 (24)	0.6 [0.3-1.1]	0.08	4
Providing accommodation for a resident from a country at high risk of drug resistance in the previous twelve months	5 (9)	7 (2)	5.0 [1.3-19.0]	0.03	1
Daily contact with animal Contact with animals having taken antibiotics in the previous three months	38 (56) 7 (13)	154 (49) 17 (5)	1.3 [0.7-2.4] <b>2.9 [1.0-8.1]</b>	0.40 <b>0.05</b>	1 3
Living alone	13 (17)	50 (18)	1.0 [0.4-2.4]	0.94	1
Foreign nationality	1 (1)	13 (4)	0.2 [0.03-2.0]	0.19	1
Age ≥ 65 years old	15 (18)	50 (15)	1.3 [0.6-3.1]	0.51	0
Consumption of raw meat in the previous three months	5 (6)	46 (19)	0.3 [0.1-0.9]	0.03	2

\*Wald test; OR: odds ratio; MDR: antibiotic multi-resistant; \*\* proportions were estimated with the sampling design, which explain why it is not equal to numerator / denominator; "occurring in women with urinary tract anomalies, pregnancy, those over 65 years old, or those with a chronic disease (diabetes, recent or recurring cancer, or renal insufficiency)

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Table 3: Factors associated with antibiotic-multi-resistant Enterobacteriaceae urinary tract infections: multivariate analysis performed according to the sampling design

	OR [95% Cls]	p value <sup>*</sup>
Penicillin use by the patient in the previous three months	3.1 [1.2-8.0]	0.02
Providing accommodation for a resident from a country at high risk of drug resistance in the previous 12 months	4.0 [1.2-15.1]	0.03
Consumption of raw meat in the previous three months	0.3 [0.1-0.9]	0.04

\*Wald test; OR: odds ratio; and CIs: confidence intervals. Two observations were deleted due to missing values.

# Table 4: Factors associated with penicillin-resistant (PN) *Enterobacteriaceae* (EB) urinary tract infections (UTIs): univariate analysis with the sampling design

		PN EB N=139, sample size (%**)	Not PN EB N=224, sample size (% **)	OR [95% CI]	p value*	Missing value
UTI in the I	ast twelve months	117 (84)	183 (81)	1.5 [0.7-3.0]	0.28	1
Complicate	ed UTI	10 (4)	20 (9)	0.4 [0.2-0.9]	0.04	2
Patients wi	th ≥ four UTIs/year	13 (10)	15 (4)	2.5 [0.9-6.4]	0.06	1
Antibiotic use by the patient in the previous 3 three months	37 (28)	38 (18)	1.8 [0.8-3.8]	0.15	1	
Pe	nicillin in the previous three months	14 (12)	15 (8)	1.6 [0.4-6.1]	0.47	1
Qu mo	inolone in the previous three nths	9 (7)	9 (4)	1.6 [0.5-5.3]	0.44	1
Otł mo	ner antibiotics in the previous three onths	14 (7)	16 (6)	1.2 [0.5-2.9]	0.65	1
Antibiotic u previous th	se by a household member in the ree months	27 (20)	29 (13)	1.8 [0.9-3.4]	0.10	1
Hospitaliza	tion in the previous 12 months	12 (9)	29 (14)	0.6 [0.2-1.4]	0.21	2
Healthcare	contact in the previous month	28 (16)	54 (23)	0.7 [0.4-1.2]	0.20	1
Household month	healthcare contact in the previous	24 (16)	38 (14)	1.1 [0.6-2.0]	0.63	1
Internationa	al travel in the previous 12 months	36 (28)	55 (25)	1.2 [0.7-2.0]	0.56	1
Inte twe of c	ernational travel in the previous elve months in countries at high risk drug resistance	25 (21)	47 (23)	0.9 [0.5-1.6]	0.68	4
Providing a country at h previous tw	accommodation for a resident from a high risk of drug resistance in the velve months	6 (5)	6 (2)	2.2 [0.6-8.3]	0.25	1
Daily conta Co ant	ict with animal ntact with animals having taken ibiotics in the previous three months	71 (51) 10 (8)	121 (50) 14 (6)	1.0 [0.6-1.9] 1.4 [0.5-3.8]	0.89 0.52	1 3
Living alon	е	29 (19)	34 (17)	1.1 [0.4-2.8]	0.88	1
Foreign nat	tionality	5 (4)	9 (3)	1.6 [0.4-7.1]	0.53	1
Age ≥ 65 y	ears old	25 (14)	40 (16)	0.8 [0.4-1.5]	0.51	0
Consumption months	on of raw meat in the previous three	18 (19)	33 (15)	1.3 [0.6-2.5]	0.51	2

\*Wald test; OR: odds ratio; \*\* proportions were estimated with the sampling design, which explain why it is not equal to numerator / denominator; \*\*\*occurring in women with urinary tract anomalies, pregnancy, those over 65 years old, or those with a chronic disease (diabetes, recent or recurring cancer, or renal insufficiency

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Table 5: Factors associated with trimethoprim-sulfamethoxazole-resistant (TMP) *Enterobacteriaceae* (EB) urinary tract infections (UTIs): univariate analysis with the sampling design

		TMP EB N=58, sample size (%**)	Not TMP EB N=305, sample size (%**)	OR [95% CI]	p value*	Missing value
UTI in t	he last twelve months	52 (93)	248 (80)	3.4 [1.3-8.8]	0.02	1
Complie	cated UTI	4 (3)	26 (8)	0.4 [0.1-1.3]	0.13	2
Patients	s with $\geq$ four UTIs/year	6 (9)	22 (6)	1.4 [0.5-3.5]	0.51	1
Antibiotic use by the patient in the previous three months Penicillin in the previous three months	17 (31)	58 (20)	1.8 [0.9-3.8]	0.11	1	
	Penicillin in the previous three months	6 (16)	23 (8)	2.2 [0.7-6.8]	0.17	1
	Quinolone in the previous three months	4 (5)	14 (5)	0.9 [0.3-3.1]	0.87	1
	Other antibiotics in the previous three months	6 (8)	24 (6)	1.3 [0.4-4.1]	0.62	1
Antibiot previou	ic use by a household member in the s three months	10 (17)	46 (16)	1.1 [0.5-2.6]	0.84	1
Hospita	lization in the previous 12 months	4 (6)	37 (13)	0.4 [0.1-1.3]	0.13	2
Healthc	are contact in the previous month	10 (19)	72 (20)	0.9 [0.4-2.1]	0.81	1
Househ month	old healthcare contact in the previous	10 (11)	52 (16)	0.6 [0.3-2.1]	0.24	1
Internat	ional travel in the previous 12 months	18 (37)	73 (24)	1.8 [0.8-4.0]	0.15	1
	International travel in the previous twelve months in countries at high risk of drug resistance	12 (27)	60 (21)	1.4 [0.5-3.6]	0.49	4
Providir country previou	ng accommodation for a resident from a at high risk of drug resistance in the s twelve months	3 (6)	9 (3)	2.7 [0.6-1.1]	0.19	1
Daily co	ontact with animal Contact with animals having taken antibiotics in the previous three months	30 (58) 4 (7)	162 (49) 20 (6)	1.4 [0.7-2.8] 1.2 [0.3-4.3]	0.30 0.76	1 3
Living a	lone	14 (23)	49 (17)	1.5 [0.8-3.1]	0.24	1
Foreign	nationality	1 (1)	13 (4)	0.3 [0.03-2.2]	0.22	1
Age ≥ 6	5 years old	13 (20)	52 (14)	1.5 [0.7-2.9]	0.30	0
Consun months	nption of raw meat in the previous three	9 (24)	42 (14)	1.8 [0.7-1.8]	0.22	2

\*Wald test; OR: odds ratio; \*\* proportions were estimated with the sampling design, which explain why it is not equal to numerator / denominator; \*\* occurring in women with urinary tract anomalies, pregnancy, those over 65 years old, or those with a chronic disease (diabetes, recent or recurring cancer, or renal insufficiency

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