

Bacteriological profile and resistance of *Escherichia coli* to beta-lactam for patients consulted at Douala Laquintinie Hospital

ABSTRACT

Enterobacteriaceae are the most common causes of community-acquired and nosocomial infections. They are usually treated with beta-lactam antibiotics, i.e., penicillins, broad-spectrum cephalosporins, and carbapenems (Imipenem, meropenem, ertapenem). In order to evaluate the resistance profile of *Escherichia coli* (*E. coli*) to beta-lactam antibiotics, a 3-year retrospective study was carried out in the medical biology laboratory of the Laquintinie Hospital in Douala. The aim was to compare the behavior of *Escherichia coli* bacteria to penicillins, carbapenems, and cephalosporins. In order to achieve this objective, we used the results obtained from *E. coli* strains isolated from urine, cervico-vaginal fluid, puncture fluid, pleural fluid and pus samples. Identification was confirmed using the API 20 E *Enterobacteriaceae* system of biochemical testing and the VITEK 2 system. The VITEK 2 system was used to perform the antibiogram. Statistical analyses obtained from GraphPad Prim V 5.0 software allowed us to perform tests such as ANOVA, Kruskal-Wallis test, and Spearman correlations. Preliminary results show that *E. coli* bacteria are highly resistant to penicillins and significantly susceptible to cephalosporins of all generations if they are not ESBL (extended-spectrum beta-lactamases). Data on carbapenem behavior show less resistance and moderate susceptibility.

Keywords: Bacterial resistance, antibiotics, *Escherichia coli*, beta-lactams, Douala

1. Introduction

Escherichia coli, a gram-negative bacillus of the Enterobacteriaceae family and a commensal of the gastrointestinal tract (10⁸ per gram of feces), is the most frequently isolated uropathogenic bacterium in uncomplicated UTIs (70-95%) [1].

Other bacteria isolated include other enterobacteria (*Klebsiella spp.*, *Proteus spp.*). The bacterial epidemiology of complicated UTIs is more varied; *E. coli* remains the most common bacterium (50%), along with *Klebsiella spp.* and *Proteus spp.* but also *Pseudomonas aeruginosa*, enterococci and yeasts [2]. *Escherichia coli* is a bacterium commonly found in the digestive tract of humans and warm-blooded organisms. Most strains are harmless. Some, however, can cause severe food poisoning that can lead to serious illnesses, including urinary tract infections [3].

Urinary tract infections are a major public health problem. According to the World Health Organisation (WHO), *E. coli* urinary tract infections are the most common in hospitals and communities. The choice of an antibiotic, regardless of the species or resistance mechanism involved, must meet requirements to ensure clinical and microbiological success. In addition to the in-vitro sensitivity of the species concerned to the chosen antibiotic, the latter must have the best possible diffusion allowing it to reach the infectious site at sufficient tissue concentrations compared to the minimum inhibitory concentration

36 (MIC) of the isolated bacterium [4]. The beta-lactams are effective; they act by inhibiting the PLP
37 (penicillin-binding protein) enzymes essential for the final stage of peptidoglycan synthesis [5]. In
38 recent years, there has been an increase in antibiotic resistance to UTIs. The emergence of Broad
39 Spectrum Beta-Lactamase (ESBL) secreting Enterobacteriaceae is increasingly prevalent [6]. It
40 appears important to evaluate the behavior of antibiotics against *E. coli* in patients consulting the
41 Laquintinie Hospital of Douala for better therapeutic management. This approach allows us to
42 characterize the bacteriological resistance profile of *E. coli* to beta-lactam antibiotics in this hospital,
43 particularly penicillins, carbapenems, and cephalosporins.

44 **2. Materials and methods**

45 **Design of the study**

46 This is a retrospective descriptive study. The data for this study were obtained from bacterial
47 strains of *Escherichia coli* isolated and identified from patients attending the bacteriology laboratory of
48 Laquintinie Hospital in Douala during the period from 1 January to 31 December 2019.

49 **Sample Collection and Bacterial Isolation**

50 The different *E. coli* strains tested had been isolated from urine, cervico-vaginal fluid, puncture
51 fluid, pleural fluid and pus samples. their identification had been made through their morphological and
52 biochemical characteristics (Gram-negative bacilli, oxidase-negative) completed using the API 20 E
53 *Enterobacteriaceae* system of biochemical testing (BioMérieux, France) and The VITEK 2 automated
54 system (BioMérieux, France). A total of 2332 *E. coli* strains had been isolated and identified during
55 2019.

56 **Antimicrobial Susceptibility Testing**

57 The VITEK 2 automated system (BioMérieux, France) was used with antimicrobial
58 susceptibility testing (AST), respectively, of gram-negative bacilli. The antimicrobial agents tested were
59 amikacin, amoxicillin-clavulanic acid, ampicillin, cefalotin, cefepime, cefotaxime, ceftazidime,
60 cefpodoxime, ceftazidime, cefuroxime, ciprofloxacin, gentamicin, meropenem, nitrofurantoin,
61 norfloxacin, ofloxacin, piperacillin, piperacillin-tazobactam, tobramycin, and trimethoprim-
62 sulfamethoxazole.

63 **Statistical analysis**

64 Statistical analysis was carried out using different software packages, initially using Excel from
65 office 2010 as the spreadsheet. The data was then exported to analysis software such as Graph Pad
66 Prism V.5.0 to perform Kruskal Wallis tests and spearman correlations. A value of $p < 0.05$ were
67 considered statistically significant.

68 **3. Results**

69 **Different sampling sites**

70 During the study period, 2332 bacterial strains of *E. coli* were isolated, identified, and
71 classified into three clinical categories: two susceptible categories (standard dose susceptible (**S**) and

72 high exposure susceptible (**I**) and one resistant category (**R**) according to the recommendations of the
 73 Antibiotic Committee of the French Society of Microbiology (CASFM) V1.0 Mai 2022. The strains
 74 isolated were mainly from urine (59.61%), pus (19.47%), pleural fluid (9.01%), puncture fluid (4.03%),
 75 and finally, cervical-vaginal swab (7.89%), as shown in Table 1. Resistance was found in 48.8% of
 76 cases, with 2.3% in intermediate cases and 48.7% in sensitive cases. Of the resistant germs isolated,
 77 more than half were found in the urine and more than a quarter in the pus. Among the susceptible
 78 germs, 61.16% were found in urine and 11.07% in pus. Of the highly exposed susceptible germs,
 79 53.70% were found in urine and 29.63% in pus. In general, urine and pus were the specimens in
 80 which a high frequency of resistance was observed.

81 **Table 1.** Antimicrobial susceptibility profile of the Romanian fresh-cheese-origin *E. coli* strains tested
 82 with the VITEK®

sample	Susceptibility Test Result of 2332 Strains (n/%)			Total
	R	I	S	
Urine	665 (58.3)	29 (53.7)	696 (61.2)	1390
Cervical-vaginal swab	49 (4.3)	3 (5.6)	132(11.6)	184
Puncturefluid	34 (3)	3 (5.6)	57 (5)	94
Pleural fluid	80 (7)	3 (5.6)	127(11.2)	210
Pus	312 (27.4)	16 (29.6)	126 (11.1)	454
Total	1140	54	1138	2332
P Value	0.0003	0.005	<0.001	

83 **Description:** **R:** resistant category; **S:** standard dose susceptible; **I:** high exposure susceptible.

84 **The behavior of germs towards the different antibiotics tested in different samples.**

85 The evaluation of the distribution of the germ's behavior towards the different antibiotics tested shows
 86 a high resistance of the germ isolated in the urine to penicillins (17.1%), carbapenems (2.6%), and
 87 cephalosporins (28.1%). The *Escherichia coli* germs found in pus also showed high resistance to
 88 penicillins (26.3%), carbapenems (7.6%), and cephalosporins (28.1%). The behavior of *Escherichia*
 89 *coli* in specimen media other than urine and pus showed less than 10% resistance to the three
 90 different styles of antibiotics, as did cervical-vaginal specimens (CVS), which showed 1.2% resistance
 91 to penicillins, 0.6% to carbapenems and 1.7% to cephalosporins. Concerning the sensitivities, we
 92 observe a high sensitivity for *E. coli* strains isolated in urine, notably for cephalosporins (29.8%) and
 93 carbapenems (13.2%).

94 **Table 2.**The behavior of the different antibiotics at the sampling sites

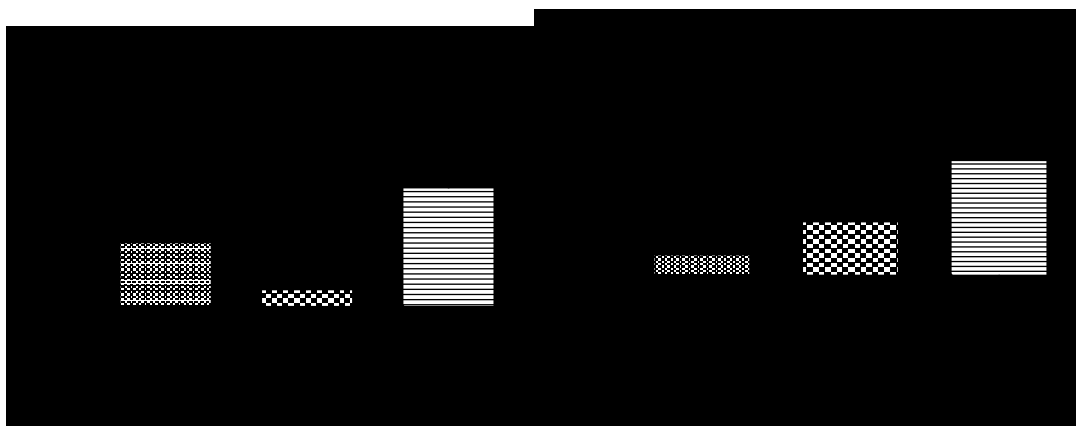
sample	Susceptibility Test Result of 2332 Strains (n/%)									P value
	R			I			S			
	Rp	Rc	Rce	Ip	Ic	Ice	Sp	Sc	Scce	
Urine	17.1	2.6	28.1	0.4	0.2	1.5	7.1	13,2	29.8	0.006

Cervico-vaginal swab	1.2	0.6	1.7	0.0	0.0	0.2	0.9	2,2	6.3
PunctureLiquid	1.4	0.1	0.9	0.0	0.1	0.1	0.9	1,8	1.4
Pleural fluid	0.9	0.1	4.7	0.0	0.2	0.0	0.3	2.4	6.4
Pus	5.7	4.0	12.7	0.1	0.1	0.9	0.2	3.7	5.1

95 **Description:** **Rp:** Resistant to penicillins; **Rc:** Resistant to carbapenems; **Rce:** Resistant to
 96 cephalosporins; **Ip:** Susceptible at a high exposure to penicillins; **Ic:** Susceptible at a high exposure to
 97 carbapenems; **Ice:** Susceptible at a high exposure to cephalosporins.

98 **Distribution of antibiotic behavior on *Escherichia coli***

99 Figure 1 below shows the behavior of the various antibiotics against *E. coli*. It can be seen
 100 that, in general, cephalosporins were more active on the different strains of *Escherichia coli* in terms of
 101 resistance (more than 18%) with a significant P value (P Value=0.016) and terms of sensitivity (more
 102 than 20%) with significant P values (0.0017). Carbapenems (more than 7%) were more sensitive than
 103 penicillins (less than 5%) and consequently less resistant than penicillins (more than 10%).



104 **Figure 1: Distribution of the behavior of different antibiotics (Betalactamins) towards**
 105 *Escherichia coli*

107 **Description:** A: Sensible; B: Resistances; Significant < 0,005 (Kruskal-Wallis and Friedman statistic
 108 via ANOVA Analysis); **Rp:** Resistant to penicillin's; **Rc:** Resistant to carbapenems; **Rce:** Resistant to
 109 cephalosporins; **Ip:** Susceptible at a high exposure to penicillin's; **Ic:** Susceptible at a high exposure to
 110 carbapenems; **Ice:** Susceptible at a high exposure to cephalosporins.

111 **Bacteriological profile of *Escherichia coli* and penicillin resistance**

112 Amongst the Beta-lactam antibiotics tested for antibiotic susceptibility, there was a resistance
 113 of over 8.58% compared to a sensitivity of 1.8% for ampicillin and less than 1% for oxacillin and
 114 ticarcillin.

115 **Table 3.** The behavior of *Escherichia coli* towardspenicillin's

Antimicrobial	Susceptibility Test Result of 2332 Strains (n%)		
	R	I	S
AMOX	2 (0.7)	0 (0.0)	0 (0.0)
AMP	200 (74.6)	0(0.0)	24 (8.9)

OXA	16 (5.9)	1 (0.37)	0 (0.0)
TIC	22 (8.2)	0(0.0)	3 (1.1)
P value	0.0276		

116 **Description:** **R:** resistant category; **S:** standard dose susceptible; **I:** high exposure susceptible;
 117 **AMOX:** Amoxicillin; **AMP:** Ampicillin; **OXA:** Oxacillin; **TIC:** Ticarcillin

118 **Bacteriological profile of *Escherichia coli* and carbapenem and monobactam resistance**

119 Several carbapenem and monobactam antibiotics were tested on *Escherichia coli* during this
 120 period. Almost 1% resistance was observed with imipenem and less than 1% with Meropenem.
 121 Meropenem showed rather high sensitivities compared to the other, up to more than 11%. A sensitivity
 122 of more than 2% was also observed with imipenem. The sensitivity of Azthreonam was the lowest
 123 observed, with less than 1%.

124 **Table 4.**Behavior towards carbapenems and Azthreonam

Antimicrobial	Susceptibility Test Result of 2332 Strains (n/%)		
	R	I	S
ATM	0 (0.0)	8 (1.9)	12 (2.9)
IMI	24 (5.9)	0 (0.0)	68 (16.8)
MEM	20 (4.9)	0 (0.0)	272 (67.3)
P value	0.0106		

125 **Description:** *Significant < 0.005 (Friedman test statistic); **R:** resistant category; **S:** standard dose
 126 susceptible; **I:** high exposure susceptible; **ATM:** Azthreonam; **IMI:** Imipenem; **MEM:** Meropenem.

127 **Bacteriological profile of *Escherichia coli* and cephalosporin resistance**

128 Several cephalosporin antibiotics were tested, with other resistance observed, for other
 129 cephalosporin antibiotics less than 1%. Compared to beta-lactam sensitivity in general.

130 **Table 5.**Behavior towards cephalosporins

Antimicrobial	Susceptibility Test Result of 2332 Strains (n/%)		
	R	I	S
CAZ	66(8.8)	12(1.6)	52(6.9)
CFM	64(8.5)	6(0.8)	40(5.3)
CMX	32(4.7)	0(0.0)	12(1.6)
CPD	76(10.1)	2(0.2)	78(10.4)
CRO	78(10.4)	2(0.2)	74(9.8)
CTX	30(4)	0(0.0)	64(8.5)
FIX	34(4.5)	0(0.0)	28(3.7)
P value	0.0012		

131 **Description :** *Significant< 0.005 (Kruskal-Wallis statistic) ;**R :** resistantcategory ;**S :** standard dose
 132 susceptible ;**I :** high exposure susceptible ; **CAZ :**Ceftazidime; **CFM:** Ceftriazone; **CXM:** Cefuroxime;
 133 **CPD:** Cefpodoxime; **CRO:** Ceftriazone; **CTX:** Cefotaxime; **FIX:** Cefixime.

134 **Correlational analysis of the behavioral dynamics of different beta-lactam antibiotics against E.**
 135 **coli strains.**

136 After observing the behavior of the different classes of beta-lactam antibiotics, it was important
 137 to establish a correlation to test a potential increase in sensitivity or a potential decrease in resistance
 138 effects of the latter.

139 **RESISTANCE**

140 A strong (0.9) significant correlation (P value=0.0045) on the possibility of E. coli resistance
 141 was obtained with a combination of penicillins and cephalosporins. Negative and non-significant
 142 correlations were obtained with the other antibiotic combinations.

143 **Table 6.**Correlation of resistance between different antibiotics

	Penicillins	Carbapenems	Cephalosporins
correlation coefficients			
Penicillin		-0.36	0.90
Carbapenem	-0.36		-0.13
Cephalosporin	0.90	-0.13	
P value			
Penicillin		0.35	0.0045
Carbapenem	0.35		0.75
Cephalosporin	0.0045	0.75	

144 **SENSITIVITY**

145 The comparison of the action of the different antibiotics about their capacity to increase
 146 susceptibility when combined shows no correlation between the different combinatory modes.
 147 Furthermore, all these combinations were non-significant.

148 **Table 7.**Correlation on sensitivity between different antibiotics

	Penicillins	Carbapenems	Cephalosporins
correlation coefficients			
Cephalosporins		-0.14	-0.50
Carbapenems	-0.14		-0.07
Penicillins	-0.50	-0.07	
P value			
Cephalosporins		0.73	0.19
Carbapenems	0.73		0.86
Penicillins	0.19	0.86	

149 **DISCUSSION**

150 Based on the characterization of the bacteriological profile and resistance of *Escherichia coli*
151 to beta-lactam in patients consulting the medical biology laboratory of the Laquintinie Hospital in
152 Douala, this study enabled us to obtain varied results. The analyses were carried out in various
153 samples and showed that among the 2332 non-repetitive bacterial strains isolated, the majority of the
154 latter came from urine (59.61%). This can be explained by the fact that *E. coli* are normally harmless
155 intestinal bacteria for humans when they remain confined to the digestive tract. [7]. The
156 preponderance of *Escherichia coli* in urine has been found by several authors, as presented by
157 Cheung and colleagues in 2020[8] and Wilson and collaborateurs[9]. Given the ubiquity of *E. coli*, it
158 was also found in various other samples, such as pus (19.47%), data that is consistent with the work
159 of Serraino and colleagues in 2018, who found *E. coli* at a frequency of 26.5% in pyogenic liver
160 abscesses (PAA) [10]. In this study, *E. coli* was also isolated from pleural fluid (9.01%) as
161 demonstrated by Cartelle and colleagues, by isolating a clinical strain of *E. coli* from pleural fluid with
162 high levels of resistance to cefotaxime, ceftazidime, and aztreonam has a novel CTX-M gene
163 (bla(CTX-M-32)) whose amino acid sequence differs from that of CTX-M-1 by a single Asp240-Gly
164 substitution [11]. The strains found in the puncture fluid (4.03%) corroborated with the ebongue and
165 collaborators data obtained at the Douala General Hospital in 2015, CERVICAL-VAGINAL SWAB
166 (7.89%)[12]. The resistance observed was more in urine, with a rate for penicillins of 17.1%; this could
167 be due to the location of the bacteria in the digestive tract. This observation is consistent with data
168 obtained by Vazouras and colleagues in 2020, showing that *E. coli* was highly resistant to ampicillin
169 [13]. For carbapenems, resistances have been observed; these resistances to carbapenems are
170 particularly notable in Gram-negative bacilli species; this could be since these species naturally have
171 relatively low transmembrane diffusion coefficients to beta-lactams as presented by Nordmann Patrice
172 in 2010[14]. Carbapenems are mainly prescribed for the treatment of enterobacterial infections of
173 nosocomial and, more rarely, community origin. Some publications show the extension of this
174 resistance in enterobacteria: it is no longer restricted to certain regions of the world and now affects
175 species that may be typically community-acquired, as is *E. coli* [15]. The cephalosporin susceptibility
176 observed against *E. coli* is believed to be due to the production of CTX-M ESBLs, CTX-M ESBLs, and
177 CMY-type AmpC enzymes and an absence of CTX-M-15 ESBL production as presented by Dhanji
178 and colleagues in 2010 [16].

179 **CONCLUSION**

180 This retrospective study of 2332 samples were initiated to evaluate the resistance profile of *E. coli* to
181 Beta-lactam antibiotics, including penicillins, carbapenems, and cephalosporins. In general,
182 cephalosporins were more active on the different strains of *Escherichia coli* in terms of resistance and
183 sensitivity. Carbapenems were more sensitive than penicillins and, therefore, less resistant than
184 penicillins. However, *E. coli* had higher resistance to penicillins, and carbapenems and less than
185 cephalosporins, with low susceptibility, compared to cephalosporins and carbapenems. Some
186 cephalosporins were intermediate, showing a strong ability to either shift in resistance or sensitivity.

187 **Declarations**

188 **Ethics approval and consent to participate**

189 The protocol of this study was approved by the National Ethics Committee for Research in Human
190 Health (N° 2020/18/082/CE/CNERSH/SP). As this was a retrospective study, the confidentiality of the
191 information obtained on the study subjects was respected.

192 **Availability of data and materials**

193 The data will be available upon reasonable request to the corresponding author.

194 **Competing interests**

195 Authors' have none to declare.

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197 This study did not receive any funding in any form.

198 **Author's contributions**

199 All authors contributed to the design and execution of the study, participated in article drafting and
200 critical revision, and read and approved the final version of the manuscript.

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