



RESEARCH ARTICLE

PREVALENCE OF ANTIMICROBIAL RESISTANT *ESCHERICHIA COLI* STRAINS ISOLATED FROM
POULTRY, MEGHALAYA, INDIA

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ABSTRACT

The frequent use of antimicrobial has rendered the risk of increase in resistance against *E. coli*. The objective of this study was to observe the resistance pattern against most commonly used drugs. A total of 92 *Escherichia coli* were isolated from tissues of diseased poultry birds. The antibiotic resistance-susceptibility profiles were conducted by standardized disc diffusion following CLSI method. Test for resistance to 14 antimicrobial agents were conducted. The most frequently encountered form of resistance in all the samples were to sulphafurazole (98.91%) trimethoprim-sulfamethoxazole (96.73%), colistin (88.04%), tetracycline (86.95%), co-trimoxazole (76.08%), enrofloxacin (75%) and ampicillin (70.65%). 67 (72.82%) of the isolates showed resistance to more than three antimicrobial classes. The high percentage of multi-drug resistance prevailing among the samples was mainly against sulphonamides, tetracycline and fluoroquinolones. Therefore, the increase in resistance against colistin, sulphonamides and fluoroquinolones which were considered highly effective against *E. coli* was highly notable.

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INTRODUCTION

Antimicrobial resistance (AMR) is one of the most burning concern and thus considered as a global problem in human as well as veterinary medicine (Boerlin *et al.*, 2005). The widespread risks developed can be attributed to the frequent use and misuse of the antimicrobials. The presence of antibiotic-resistant *Escherichia coli* in domestic poultry, beef and swine has been reported (Sandvang *et al.*, 1997; Madsen *et al.*, 2000; Guerra *et al.*, 2003). *Escherichia coli* play a crucial role either as commensal bacteria or as potential foodborne pathogens. It is ubiquitous in nature and can enter the food chain through contamination of faecal matter or food animal slaughter. Poultry is one of the most important farm animals as it serves as an integral food item for the meat eaters all over the world. Therefore, antimicrobials are often used as growth promoters for production purpose, particularly in poultry industry. Colibacillosis, a common disease in poultry is caused by *E. coli*. It not only renders huge economic losses but also can infect human with diseases such as haemorrhagic colitis

and haemolytic uremic syndrome (Chansiripornchai, 2009; Ferens and Hovde, 2011). The selection pressure rendered upon using antimicrobials as feed additives or therapy has resulted in the emergence of multiple antimicrobial resistant bacteria thus making treatment difficult (Witte, 1998; van den Bogaard *et al.*, 2000). The emergence and dissemination of antibiotic resistant genes among bacterial strains leading to the ineffectiveness of broad spectrum antibiotic treatment is an increasing problem (Maynard *et al.*, 2003). In recent times, many outbreaks of *E. coli* infection have been seen all around the world. ESBL (Extended Spectrum Beta Lactamases) producing bacteria which are resistant to the third and fourth generations of cephalosporins, penicillins and monobactams possess challenging infection control issues (Rupp and Fey, 2003). World Health Organization (WHO) has listed several antimicrobials belonging to ESC (Extended spectrum cephalosporins) used in the treatment of human and livestock as important since they are the sole therapeutic option available for curbing life-threatening diseases (WHO, 2007). Antimicrobial resistance is typically more frequent and higher among pathogens than commensal bacteria (DANMAP, 2004). Therefore, the main aim of this study was to observe the prevalence trends of the various classes of antimicrobials in the

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diseased poultry samples collected from the farms of Meghalaya, India, and also to investigate the antimicrobial resistant patterns.

MATERIALS AND METHODS

Collection of samples

The samples of poultry were randomly collected from organised and unorganised farms in and around Meghalaya, India. A total of 106 post-mortem tissue samples (heart, liver, spleen, intestine and kidney) were collected aseptically using disposable gloves and sterile vials. The samples were kept in ice-box and transported. They were then processed within 6h from the time of collection. Proper care was taken during collection and transportation for the viability of the samples.

Isolation and identification procedure

The tissue samples were cut into small pieces and weighed around 1g approximately. It was inoculated in MacConkey broth (HiMedia) and then incubated at 37°C for 18 to 24h. A loopful of the broth culture was taken and streaked onto MacConkey Lactose Agar plate (MLA, HiMedia) and incubated for 18 to 24h at 37°C. A suspected pink colony was selected and streaked on Eosin Methylene Blue Agar (EMB, HiMedia). Colonies showing green metallic sheen were then subjected to biochemical tests for further confirmation. The tests include IMViC (indole, methyl red, Voges Proskauer and citrate) as well as oxidase and urease tests. The confirmed isolates were then stored in BHI (Brain Heart Infusion) broth containing 15% glycerol at -80°C.

Antimicrobial Sensitivity testing

Antimicrobial sensitivity test was performed on all the 92 *E. coli* isolates as per recommendation of Clinical Laboratory Standards Institute (CLSI, 2014) (Figure 1). Mueller-Hinton Agar (HiMedia, Mumbai, India) was used for the purpose. The following antimicrobial discs belonging to nine different classes were commercially procured (HiMedia, India) and screened: tetracycline (30µg), neomycin (30µg), ciprofloxacin (5 µg), colistin (30µg), enrofloxacin (5 µg), ampicillin (10 µg), chloramphenicol (10 µg), streptomycin (25 µg), ceftriaxone (30µg), ceftazidime (30µg), trimethoprim (30µg), gentamycin (120 µg), co-trimoxazole (25 µg), sulphafurazole (300 µg).

The results were interpreted according to CLSI (2014) and EUCAST (2014). The standard strain of *Escherichia coli* (ATCC 29522) was used as a quality control. An isolate was regarded as 'multi-resistant' if it exhibited resistance to drugs of ≥ 3 antimicrobial types (Tenover, 2006).

RESULTS

Out of the 106 post-mortem tissue samples collected from diseased poultry birds, 92 (86.79%) were biochemically tested and confirmed as *E. coli* isolates. The 92 *E. coli* isolates were subjected to antimicrobial susceptibility tests to 14 antimicrobials belonging to 9 different classes. One strain from each positive sample was used for further analysis. The highest resistance was registered in sulfafurazole. 98.91% isolates showed resistant to it followed by trimethoprim-sulfamethoxazole (96.73%), colistin (88.04%), tetracycline (86.95%), co-trimoxazole (76.08%), enrofloxacin (75%) and ampicillin (70.65%) (Table 1). Combined resistance to sulfafurazole, trimethoprim-sulfamethoxazole, colistin, tetracycline, co-trimoxazole, and ampicillin was found in 54.34% isolates. Ceftazidime and ceftriaxone, the third generation antibiotics belonging to the class Cephalosporins showed resistance up to 10.86% and 6.52% respectively. None of the isolates were found resistant to gentamicin, eventually making it the most sensitive with 91.30% as shown in Figure 1. 79.34% of the isolates were sensitive to ceftriaxone followed by chloramphenicol (77.17%).

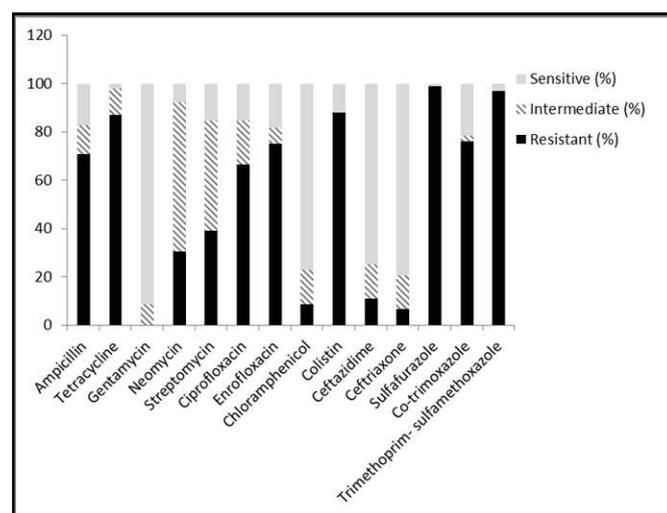


Figure 1. Graphical representation of the sensitivity pattern of the *E. coli* isolates

Table 1. Numbers (%) of *Escherichia coli* isolates from diseased poultry sample depicting antimicrobial resistance profile

Antimicrobial Class	Antimicrobials	Disc content (µg)	Resistance breakpoint (mm)	<i>E. coli</i> isolates (n=92)		
				S (%)	I (%)	R (%)
Penicillin	Ampicillin	10	≤13	16(17.39)	11(11.95)	65(70.65)
Cephalosporins	Ceftazidime	30	≤14	69(75)	13(14.13)	10(10.86)
	Ceftriaxone	30	≤13	73(79.34)	13(14.31)	6(6.5)
Aminoglycosides	Gentamycin	120	≤12	84(91.3)	8(8.69)	0(0)
	Neomycin	30	≤12	7(7.6)	57(61.95)	28(30.43)
	Streptomycin	25	≤11	14(15.21)	42(45.65)	36(39.13)
Fluoroquinolones	Ciprofloxacin	5	≤15	14(15.21)	17(18.47)	61(66.3)
	Enrofloxacin	5	≤16	17(18.47)	6(6.52)	69(75)
Phenolics	Chloramphenicol	10	≤12	71(77.17)	13(14.13)	8(8.69)
Tetracycline	Tetracycline	30	≤14	2(2.17)	10(10.86)	80(86.95)
Polypeptides	Colistin	10	≤10	11(11.95)	0(0)	81(88.04)
Sulphonamides	Sulfafurazole	300	≤12	1(1.08)	0(0)	91(98.91)
	Co-trimoxazole	25	≤10	20(21.73)	2(2.17)	70(76.08)
Trimethoprim-sulfamethoxazole		30	≤10	3(3.26)	0(0)	89(96.73)

With regard to multidrug resistance profiles, out of the 12 antimicrobials tested, all isolates were resistant to at least one antimicrobial agent, 73.91% were resistant to more than five and 59.78% were resistant to more than seven. Two particular isolates were resistant to 11 antimicrobial agents and intermediately resistant to 3 antimicrobials. 67 (72.82%) of the isolates showed total resistance to more than three antimicrobial classes.

DISCUSSION

The higher resistance rate in tetracycline, ampicillin and trimethoprim-sulfamethoxazole has been in accordance with previous studies (Mooljunttee S. *et al.*, 2010; Jiang *et al.*, 2011; Momtaz *et al.*, 2012; Rasheed *et al.*, 2014). Also a zero percentage of resistance was monitored in gentamycin among the poultry isolates (Mooljunttee *et al.*, 2010). The finding of the present work somewhat corresponds with the study done by Hamisi *et al.* (2014) on tropical free-range chickens from Tanzania. According to their study, 83.1% isolates has shown resistance to at least one antimicrobial agent. The increase in frequency of antibiotic resistance obtained is also in accord with that stated by Van *et al.* (2007, 2008).

The presence of multidrug resistant *E. coli* in chicken has been previously reported from different countries (Kolar *et al.*, 2001; Schroeder *et al.*, 2003). Such alarming rates of multi resistance might be due to the frequent administration and over-usage of the antimicrobials (Moniri and Dastehgoli, 2007; Miranda *et al.*, 2008). Moreover, during slaughter, resistant strains from the gut can easily contaminate poultry carcasses, therefore, contaminating the poultry meats with resistant *E. coli* (Van den Bogaard *et al.*, 2001; Vaidya *et al.*, 2005; Akond *et al.*, 2009; Altekruise *et al.*, 2009). According to Overdeest *et al.* (2011), close relationship has been established between ESBL-producing *E. coli* strains of poultry and humans. Such links can be alarming for the poultry industry as well as humans. Third generation cephalosporin is considered a major drug for human disease treatments. The occurrence of cephalosporin resistant *E. coli* in retail meat is of great concern. The intestinal tract of many food producing animals has been known to possess cephalosporin resistant *E. coli* (Riano *et al.*, 2006; Cloeckaert *et al.*, 2007; Liu *et al.*, 2007). Moreover, 35% of *E. coli* strains isolated from live broilers were found to be resistant to third generation cephalosporin (Depoorter *et al.*, 2012). In many European countries, the genetic background for cephalosporin resistance in *E. coli* isolates, from broilers and retail chicken meat, is heterogeneous including genes of the CTX-M, TEM and SHV groups (Ewers *et al.*, 2012). Therefore, an attempt should be made to keep the prevalence of cephalosporin-resistant *E. coli* in retail meat low, thus, somewhat curbing the probability of zoonotic transmission.

In conclusion, the results of this study indicate that *E. coli* strains isolated from post-mortem tissue samples of diseased poultry showed a higher resistance rate for important antimicrobials including fluoroquinolones and colistin. The presence of high percentage multi-drug resistance (MDR) in the isolates is quite a concern. A proper monitoring is required while prescribing or giving feed additives.

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