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Antimicrobial Drug Resistance in *Escherichia coli* Including an O157:H7 Isolate from Feces of Healthy Goats in Grenada

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Authors' contributions

This work was carried out in collaboration between all authors. Authors VAA, RS and HH designed the study. Authors RS and SG managed the collection of all the samples. Authors VAA, EA, OAO and ZP managed the analyses and literature searches. Authors VAA and HH wrote the protocol and wrote the first and final drafts of the manuscript. All authors read and approved the final manuscript.

Article Information

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Original Research Article

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ABSTRACT

Aim: To determine: the occurrence of *Escherichia coli* including the O157:H7 serotype in feces of healthy goats in 10 randomly selected farms in Grenada and the antimicrobial drug resistance in *E. coli* isolated from feces of the tested goats.

Study Design: During the period of May to July, 2014, fecal samples were obtained from randomly selected healthy goats in 10 farms in Grenada and analyzed in the bacteriology laboratory, in the Pathobiology Department, School of Veterinary Medicine, St. George's University, Grenada.

Methodology: Fecal samples were obtained from 70 randomly selected healthy goats in 10 farms in Grenada and cultured for *E. coli* and tested for O157:H7 serotype by the presence of non-sorbitol fermenting colonies and a positive reaction to O157-agglutination latex kits.

Results: All the 70 tested goats were culture positive for *Escherichia coli*. A total of 140 *E. coli* isolates were recovered and analyzed for the presence of non-sorbitol fermenting colonies and O157-agglutination. Of the 140 yielded *E. coli*, 11 (8%) isolates were non-sorbitol fermenters but only one (<1%) out of the non-sorbitol fermenters gave a positive reaction to the two *E. coli* O157:H7 latex kits. Antimicrobial susceptibility tests against 12 drugs showed susceptibility of the single *E. coli* O157:H7 isolate recovered to all the tested antibiotics. Among the non-O157:H7 isolates, the susceptibility rates to amoxicillin-clavulanic acid, ciprofloxacin, enrofloxacin, gentamicin, cefepime and ceftazidime ranged from 99% to 100%. The resistance rate to ampicillin, gentamicin, nalidixic acid, streptomycin, trimethoprim-sulfamethoxazole, and tetracycline observed was very low except to streptomycin (19%). Resistance to two or more antibiotics was observed only in 5% of the 140 *E. coli* isolates.

Conclusion: Our study revealed that presently, healthy goats in Grenada are not major reservoirs for the *E. coli* O157:H7 serotype and for multiple resistant *E. coli* strains.

Keywords: Escherichia coli; O157; goats; drug resistance; Grenada.

1. INTRODUCTION

Grenada which is approximately 344 km² in size is a volcanic island in the south Caribbean [1]. Kumthekar et al. [2] indicated that small ruminants are important production animals in Grenada due to the controllable size and feed requirements as well as their ability to utilize sloped landscape and limited pasture area. According to these authors, Grenada has an estimate small ruminant population of 2,500 goats, and the pasture land suitable for grazing livestock totals approximately 11,655 hectares of the Caribbean tri-island nation. Meat and milk from sheep and goats are commonly exploited for food and sold for as a source of income for many individuals in Grenada. In spite of the importance of goats, little is known about the prevalence and characteristics of bacteria that are associated with goats in Grenada.

E. coli is a common inhabitant of the large and lower small intestines of variety of mammals [3-5] including goats. It is excreted in feces and can be easily spread via soil, food and water [3,4]. A large number of *E. coli* strains are nonpathogenic [6], however, the pathogenic strains may cause severe intestinal or extra intestinal disease in humans [7] and are also capable of causing zoonotic infections [8]. Globally, more than 100 million cases of gastrointestinal illnesses and approximately one million deaths per year have been associated with pathogenic *E. coli* [9].

Microbial foodborne illness remains a global concern. *E. coli* O157:H7 has been recognized as the most important enterohaemorrhagic *E. coli*

(EHEC) serotype associated with human disease [8]. It has also emerged as the most important foodborne, zoonotic pathogen in humans [8,10]. Its importance as a public health problem was recognized in 1982, following an outbreak in the United States of America [8]. This serotype can cause hemolytic uremic syndrome (HUS), hemorrhagic colitis, abdominal cramps and diarrhea in humans [11,12]. WHO [8] estimated that more than nine percent of patients with EHEC infection may develop HUS, with a casefatality rate of three to five percent. The intestinal tracts of cattle, sheep and goats can serve as reservoir of E. coli O157:H7 [13]. These animals can contaminate the environment through shedding of the organism in their feces [14,15].

The intestines of animals are considered as a major reservoir and an ideal environment for the selection and transfer of antimicrobial resistance genes. Studies have shown that *E. coli* can serve as reservoirs of antibiotic resistance genes [16] which have been efficiently transferred not only to other *E. coli* strains but also to other enteric pathogens of humans and animals [17].

Several reports on culture proven *E. coli* O157:H7 diarrheal illnesses have been documented in many African countries including Nigeria, Uganda, Gabon, Kenya, Ivory Coast, South Africa and Swaziland [13]. In 1997, in Trinidad and Tobago, a survey of wildlife revealed the presence of only one (1 out of 271, <1%), four (4 out of 175, 2%), and two (2 out of 373, <1%) *E. coli* isolates from free-ranging mammals, captive wild animals, and zoo animals, respectively, that were non-sorbitol fermenters which were not agglutinated by O157

antiserum [18]. In Grenada, a study by Sylvester et al. [19] revealed the presence of 12 (12 out of 42, 29%) *E. coli* isolates from green iguanas (*Iguana iguana*) that were non-sorbitol fermenters which were negative for agglutination using the *E. coli* O157:H7 latex kits.

There have been no published surveys on the prevalence of *E. coli* including the *E. coli* O157:H7 serotype in goats in Grenada. The objectives of the present study were to determine: the occurrence of the *E. coli* including the O157:H7 serotype in feces in a population of healthy goats and the antimicrobial drug resistance of the *E. coli* isolates.

2. MATERIALS AND METHODS

Fecal samples were obtained from 70 randomly selected healthy adult goats from 10 farms in Grenada during a three months period, May 2014 to July 2014. Each sample was immediately placed in a sterile plastic container (Starplex Scientific Corp, Cleveland, TN, USA) and stored in a cooler with ice packs and transported to the Bacteriology Laboratory, School of Veterinary Medicine, St. George's University where all the laboratory analysis were performed. The approximate time between sample collection and culture was three hours.

For the isolation of E. coli, the fecal samples were placed in 10 ml of tryptic soy broth (Remel, Lenexa, KS, USA) and incubated at 37°C for 24 hours. After incubation, an aliquot was then streaked onto MacConkey (MAC) agar (Remel, Lenexa, KS, USA) and incubated at 37°C for 24 hours. To increase the possibility of identifying E. coli O157:H7 in a sample, two (2) pink to red color colonies with or without a zone of precipitated bile, morphologically representing E. coli were subcultured via streaking onto individual MAC agar and incubated at 37°C for 24 hours for isolation of pure colonies. Colonies from the second MAC agar plate were Gram stained and further tested using the API20E (Analytical Profile Index; BioM6rieux, Hazelwood, MO) bacterial identification strips for confirmation as E. coli. Non-lactose fermenting isolates identified as E. coli by API20E were also added in the study despite the fact that they were nonlactose fermenting variants.

The identification of *E. coli* O157:H7 was performed using the methods previously described by Sylvester et al. [19]. The pure colonies were first plated on sorbitol-MacConkey

agar (Remel, Lenexa, KS, USA) and incubated at 37°C for 24 hours. After incubation, the sorbitol-MacConkey agar plates were examined for the presence of non sorbitol fermenting colonies. All the colonies (both the sorbitol and the non sorbitol fermenting colonies) were then subjected to slide agglutination using two (2) *E. coli* 0157:H7 latex kits: Remel Wellcolex* *E. coli* 0157 Rapid Latex Test (Remel Europe Ltd, Kent, UK) and ProlexTM *E. coli* 0157 Latex Kit (Pro-lab Diagnostics, Toronto, Canada). Two (2) latex kits were used to reduce the possibility of obtaining a false positive result. Any isolate giving a positive reaction to the latex test kits was considered to be *E. coli* 0157:H7.

The antimicrobial susceptibility tests were carried out using the disc diffusion method as recommended by the Clinical and Laboratory Standards Institute (CLSI) using Mueller Hinton agar (BBL), and the inhibition zone sizes were interpreted as per CLSI guidelines [20]. The antibiotics discs used were ampicillin, amoxicillinclavulanic acid. cefepime. cefotaxime. ceftazidime. ciprofloxacin. enrofloxacin. gentamicin. nalidixic acid, streptomycin. tetracycline and trimethoprim-sulfamethoxazole (Becton, Dickinson and Co., Sparks, MD, USA).

3. RESULTS

In this present study, all the 70 tested goats were culture positive for *E. coli*. A total of 140 *E. coli* isolates were recovered and analyzed for the presence of non-sorbitol fermenting colonies and agglutination. Of the 140 yielded *E. coli*, 11 (8%) isolates were non-sorbitol fermenters but only one (<1%) out of the non-sorbitol fermenters gave a positive reaction (O157-agglutinating) to the two *E. coli* O157:H7 latex kits.

Table 1 presents the details of the antimicrobial susceptibility of the 140 E. coli isolates recovered in this present study. The results revealed a low resistance rate (0 to 7%) for all the 140 E. coli isolates to five of the tested antibiotics except to streptomycin (19%). However, a number of the E. coli isolates revealed a moderate to high intermediate resistance rate (1 to 34%) to some of the tested antibiotics. The E. coli isolates recovered in this present study, were 99 to 100% susceptible to amoxicillin-clavulanic acid. ciprofloxacin, enrofloxacin, gentamicin, cefepime and ceftazidime and showed zones of inhibition measuring greater than 28 mm around each disc. Five percent (7 out of 140) E. coli isolates in this present study showed resistance to two or more

antibiotics. The single *E. coli* O157:H7 isolate recovered in this present study was susceptible to all the tested antibiotics.

4. DISCUSSION

The results of this study indicate that presently, healthy goats in Grenada harbor E. coli in their gastrointestinal tract and the occurrence is widespread among goats in the investigated farms. However, they are not major reservoirs for the E. coli O157:H7 serotype. Our study revealed the presence of only one (<1%) E. coli O157:H7 isolate. This is somewhat similar to the low rates reported in Ethiopia in 2010 where only two out of 60 (3.3%) fecal samples from goats showed the presence of E. coli O157:H7 serotype. The low occurrence rate of E. coli O157:H7 serotype observed in this present study also concurred with the low occurrence rate observed in different animals from other countries. E. coli O157:H7 serotype was recovered from 5.5% of sheep in Ethiopia [13], 3% of lambs in Spain [21], 0.2% of lambs in Italy [14], 4% of ewes and 4% of lambs in the Netherlands [22], and 1.4% of sheep, with monthly variations from 0% to 4.8% in the United Kingdom [15,23]. In contrast with the low occurrence rate of E. coli O157:H7 observed in this present study, higher occurrence rates ranging from 55% to 95% of the goats tested were reported in France [24]. In the United States of America, a prevalence rate of 43% was detected in sheep [25,26], and in Australia [27-29], occurrence rates of 40% was reported for goats, while rates ranging from of 56% to 68%

were reported for sheep. In the United Kingdom, a survey of 1000 sheep at slaughter revealed an occurrence rate of 2% for *E. coli* O157:H7 [30]. On the contrary, in other reports from different parts of the world, no *E. coli* O157:H7 was found in any of the samples examined [31-34]. The difference in the number of *E. coli* O157:H7 recovered in this study and those recovered in other studies carried out in different parts of the world were probably due to the differences in husbandry practices, agro-climatic variations, sampling, methods of detection, breeds and the age of animals [14,35].

In the present study, 10 (7%) of the E. coli isolates were non-sorbitol fermenters that gave a negative reaction (no O157-agglutinating) to the two E. coli O157:H7 latex kits. This present study was designed to target only the E. coli O157:H7 serotype which is typically non-sorbitol fermenters that give a positive reaction to the E. coli O157:H7 latex kits. Hence, the 10 (7%) E. coli isolates that were non-sorbitol fermenters that gave a negative reaction to the two E. coli O157:H7 latex kits were not identified in relation to their serotypes. It is pertinent to note that some pathogenic non-O157 serotypes of E. coli including O26, O103, O145, O172, O174, O113 and O111, which are non-sorbitol fermenters that do not give positive reaction to E. coli O157:H7 latex kits exist. Some of these non-O157 groups have been previously associated with infections in humans [14,21].

Table 1. Antimicrobial susceptibility profiles of 140 *E. coli* recovered from feces of goats inGrenada between May and July, 2014

Antimicrobial (Disc conc. ^a (μg))	Resistant # (%)**	Intermediate	Susceptible
Ampicillin (10)	3 (2)	15 (11)	122 (87)
Amoxicillin-clavulanic acid (20, 10)	0 (0)	2 (1)	138 (99)
Cefepime (30)	0 (0)	0 (0)	140 (100)
Cefotaxime (30)	0 (0)	6 (4)	134 (96)
Ceftazidime (30)	0 (0)	0 (0)	140 (100)
Ciprofloxacin (5)	0 (0)	1 (1)	139 (99)
Enrofloxacin (5)	0 (0)	1 (1)	139 (99)
Gentamicin (10)	1 (1)	0 (0)	139 (99)
Nalidixic acid (30)	6 (4)	0 (0)	134 (96)
Streptomycin (10)	26 (19)	48 (34)	66 (47)
Tetracycline (30)	10 (7)	10 (7)	120 (86)
Trimethoprim-sulfamethoxazole (1.25, 23.75)	3 (2)	0 (0)	137 (98)

#: number, % (percentage): values are rounded up and down to the nearest whole number

^aResistant, intermediate or susceptible according to CLSI guideline for all drugs

In comparison with other studies, a survey carried out on wildlife in Trinidad and Tobago, revealed the presence of non-sorbitol fermenting E. coli that were not agglutinated by O157 antiserum from: free-ranging mammals (<1%), captive wild animals (2%) and animals in a zoo (0.5%) [18]. Sylvester et al. [19] reported a 29% (12 out of 42) prevalence rate of non-O157 serotypes of E. coli in green iguanas in Grenada that did not ferment sorbitol and gave negative reaction to the E. coli O157:H7 latex kits. This basically means that both wild and domestic animals in Grenada and other parts of the world may be harboring non-O157 group in their gastrointestinal tracts. Further studies that targets the non-O157 group is required to determine whether the 10 (7%) of the E. coli isolates that were non-sorbitol fermenters that gave a negative reaction to the E. coli O157:H7, are the pathogenic non-O157 group. This information will enable us to determine whether goats in Grenada harbor non-O157 group of E. coli in their gastrointestinal tracts and may possibly be а contributing source of contamination.

In this study, the result obtained from the disc diffusion test revealed low resistance rate ranging from 1% to 19% for the 140 isolates to six out of the 12 antibiotics tested (Table 1). This is somewhat similar to the findings of Sylvester et al. [19] who tested E. coli isolates from green iguanas to similar types of antibiotics. Their result revealed resistance rates of 7% to ampicillin, 12% to streptomycin, and 2% each to nalidixic trimethoprim-sulfamethoxazole, acid. and tetracycline. We also observed a zero resistance ciprofloxacin ceftazidime, rate to and enrofloxacin. This is also similar to the findings of Sylvester et al. [19]. In contrast, the E. coli isolates in this present study showed zero resistance rate to amoxicillin-clavulanic acid and cefotaxime, and 1% resistance rate to gentamicin whereas the E. coli isolates from the study of Sylvester et al. [19] showed a resistance rate of 12% to amoxicillin-clavulanic acid and 2% to cefotaxime, and zero resistance rate to gentamicin. On the other hand, the low resistance rate to ampicillin (2%) observed in this present study was in contrast to the high resistance rate of 84% to ampicillin reported in eastern Ethiopia [36]. Interestingly, the one E. coli O157:H7 isolate recovered in this present study was 100% susceptible to all the tested antibiotics. A number of the E. coli isolates recovered in this present study showed moderate to high intermediate resistance rates (1 to 34%)

to some of the tested antibiotics and 5% (7 out of 140) of our *E. coli* isolates were resistant to two or more antibiotics. This is of public health concern since these multiple antibiotic resistant organisms can be transmitted from animals to humans.

5. CONCLUSION

Our study revealed that presently, healthy goats in Grenada are not major reservoirs for the *E. coli* O157:H7 serotype. The single *E. coli* O157:H7 isolate recovered in this present study was susceptible to all the tested antibiotics. Among the non-O157:H7 isolates, the resistance rate to drugs other than tetracycline and streptomycin was very low. Resistance to two or more antibiotics was observed only in 5% of the 140 *E. coli* isolates. This indicates that healthy goats in Grenada are presently not main reservoirs for multiple resistant *E. coli* strains.

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COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

- Central Intelligence Agency, World Fact Book; 2013. Accessed 11 October 2013. Avialable:<u>https://www.cia.gov/library/public</u> ations/the-world-factbook/fields/2147.html
- Kumthekar S, Manning EJB, Ghosh P, Tiwari K, Sharma RN, Hariharan H. Mycobacterium avium subspecies paratuberculosis confirmed following serological surveillance of small ruminants in Grenada, West Indies. Journal of Veterinary Diagnostic Investigation. 2013; 25(4):527-30.
- Sabarinath A, Tiwari KP, Deallie C, Belot G, Vanpee G, Matthew V, et al. Antimicrobial resistance and phylogenetic groups of commensal *Escherichia coli* isolates from healthy pigs in Grenada. Webmed Central Veterinary Medicine. 2011;2(5):1-10.
- 4. Markey B, Leonard F, Archambault M, Cullinane A, Maguire D. Clinical veterinary

microbiology. Edwards R and Hewat C, editors. 2nd ed. New York: Mosby; 2013.

- Amadi VA, Matthew-Belmar V, Tiwari K, Brathwaite E, Ravindra S, Hariharan H. Antimicrobial susceptibility profiles of *Escherichia coli* recovered from feces of young healthy domestic pigs in Grenada, West Indies. British Microbiology Research Journal. 2015;5(3):300-6.
- Clermont O, Bonacorsi S, and Bingen E. Rapid and simple determination of the *Escherichia coli* phylogenetic group. Applied and Environmental Microbiology. 2000;66(10):4555-8.
- Santos AC, Zidko AC, Pignatari AC, Silva RM. Assessing the diversity of the virulence potential of *Escherichia coli* isolated from bacteremia in Sao Paulo, Brazil. Brazilian Journal of Medical and Biological Research. 2013;46(11):968-73.
- WHO. Enterohaemorrhagic Escherichia coli (EHEC). Fact sheet N°125; 2011. Accessed 13 February 2015. Available:<u>http://www.who.int/mediacentre/f</u> actsheets/fs125/en/
- Wirth T, Falush D, Lan R, Colles F, Mensa P, Wieler LH, et al. Sex and virulence in *Escherichia coli*: An evolutionary perspective. Molecular Microbiology. 2006; 60(5):1136-51.
- Perna NT, Plunkett G, 3rd, Burland V, Mau B, Glasner JD, Rose DJ, et al. Genome sequence of enterohaemorrhagic *Escherichia coli* O157:H7. Nature. 2001; 409(6819):529-33.
- 11. Fratamico P, Smith J. *Escherichia coli* infections. In: Riemann H and Cliver D, editors. Foodborne Infections and Intoxications. New York: Elsevier Science. 2006;3-15.
- Zschock M, Hamann HP, Kloppert B, Wolter W. Shiga-toxin-producing *Escherichia coli* in faeces of healthy dairy cows, sheep and goats: Prevalence and virulence properties. Letters in Applied Microbiology. 2000;31(3):203-8.
- Mersha G, Asrat D, Zewde BM, Kyule M. Occurrence of *Escherichia coli* O157:H7 in faeces, skin and carcasses from sheep and goats in Ethiopia. Letters in Applied Microbiology. 2010;50(1):71-6.
- Battisti A, Lovari S, Franco A, Di Egidio A, Tozzoli R, Caprioli A, et al. Prevalence of *Escherichia coli* O157 in lambs at slaughter in Rome, central Italy. Epidemiology and Infection. 2006;134(2): 415-9.

- Chapman PA, Cerdan Malo AT, Ellin M, Ashton R, Harkin. *Escherichia coli* O157 in cattle and sheep at slaughter, on beef and lamb carcasses and in raw beef and lamb products in South Yorkshire, UK. International Journal of Food Microbiology. 2001;64(1-2):139-50.
- Lanz R, Kuhnert P, Boerlin P. Antimicrobial 16. resistance and resistance gene determinants in clinical Escherichia coli from different animal species in Switzerland. Veterinary Microbiology. 2003;91(1):73-84.
- Oguttu JW, Veary CM, Picard JA. Antimicrobial drug resistance of *Escherichia coli* isolated from poultry abattoir workers at risk and broilers on antimicrobials. Journal of the South African Veterinary Association. 2008;79(4):161-6.
- 18. Adesiyun AA. Absence of *Escherichia coli* O157 in a survey of wildlife from Trinidad and Tobago. Journal of Wildlife Diseases. 1999;35(1):115-20.
- Sylvester WRB, Amadi V, Hegamin-Younger C, Pinckney R, Macpherson CNL, McKibben JS, et al. Occurrence of antibiotic resistant *Escherichia coli* in green iguanas (*Iguana iguana*) in Grenada, West Indies. International Journal of Veterinary Medicine: Research & Reports. 2014;1-8. Article ID 260412. DOI:10.5171/2014.260412.
- Jorgenson JH, Turnidge JD. Susceptibility test methods: dilution and disk diffusion methods. In: Murray PR, editor. Manual of clinical microbiology. Washington, DC: ASM Press. 2003;1108-27.
- Blanco J, Blanco M, Blanco JE, Mora A, Gonzalez EA, Bernardez MI, et al. Verotoxin-producing *Escherichia coli* in Spain: Prevalence, serotypes, and virulence genes of O157:H7 and non-O157 VTEC in ruminants, raw beef products, and humans. Experimental Biology and Medicine (Maywood). 2003;228(4):345-51.
- 22. Heuvelink AE, van den Biggelaar FL, de Boer E, Herbes RG, Melchers WJ, Huis in 't Veld JH, et al. Isolation and characterization of verocytotoxin-producing *Escherichia coli* O157 strains from Dutch cattle and sheep. Journal of Clinical Microbiology. 1998;36(4):878-82.
- 23. Paiba GA, Gibbens JC, Pascoe SJ, Wilesmith JW, Kidd SA, Byrne C, et al. Faecal carriage of verocytotoxin-producing *Escherichia coli* O157 in cattle and sheep

at slaughter in Great Britain. Veterinary Record. 2002;150(19):593-8.

- 24. Bastian S, Carle I, Grimont F, Grimont P. Diversity of shiga toxin-producing *Escherichia coli* in herds of dairy cows and goats. Acta Clinica Belgica. 1999;54:49-50.
- 25. Kudva IT, Hatfield PG, and Hovde CJ. *Escherichia coli* O157:H7 in microbial flora of sheep. Journal of Clinical Microbiology. 1996;34(2):431-3.
- Kudva IT, Hatfield PG, Hovde CJ. Characterization of *Escherichia coli* O157:H7 and other Shiga toxin-producing *E. coli* serotypes isolated from sheep. Journal of Clinical Microbiology. 1997; 35(4):892-9.
- Sidjabat-Tambunan H, Bensink JC. Verotoxin-producing *Escherichia coli* from the faeces of sheep, calves and pigs. Australian Veterinary Journal. 1997;75:292 -3.
- Fagan PK, Hornitzky MA, Bettelheim KA, and Djordjevic SP. Detection of shiga-like toxin (*stx1* and *stx2*), intimin (*eaeA*), and enterohemorrhagic *Escherichia coli* (EHEC) hemolysin (EHEC *hlyA*) genes in animal feces by multiplex PCR. Applied and Environmental Microbiology. 1999; 65(2):868-72.
- Djordjevic SP, Hornitzky MA, Bailey G, Gill P, Vanselow B, Walker K, et al. Virulence properties and serotypes of Shiga toxinproducing *Escherichia coli* from healthy Australian slaughter-age sheep. Journal of Clinical Microbiology. 2001;39(5):2017-21.
- Chapman PA, Siddons CA, Gerdan Malo AT, Harkin MA. A 1-year study of *Escherichia coli* O157 in cattle, sheep, pigs and poultry. Epidemiology and Infection. 1997;119(2):245-50.

- Dontorou A, Papadopoulou C, Filioussis G, Apostolou I, Economou V, Kansouzidou A, et al. Isolation of a rare *Escherichia coli* O157:H7 strain from farm animals in Greece. Comparative Immunology, Microbiology & Infectious Diseases. 2004; 27(3):201-7.
- Johnsen G, Wasteson Y, Heir E, Berget OI, Herikstad H. *Escherichia coli* O157:H7 in faeces from cattle, sheep and pigs in the southwest part of Norway during 1998 and 1999. International Journal of Food Microbiology. 2001;65(3):193-200.
- Lenahan M, O'Brien S, Kinsella K, Sweeney T, Sheridan JJ. Prevalence and molecular characterization of *Escherichia coli* O157:H7 on Irish lamb carcasses, fleece and in faeces samples. Journal of Applied Microbiology. 2007;103(6):2401-9.
- 34. Synge B, Hopkins G. Studies of verotoxigenic *Escherichia coli* O157 in cattle in Scotland and association with human case. In: Karmali M and Goglio A, editors. Recent Advances in Verocytotoxin Producing *Escherichia coli* infections. New York: Elsevier Science. 1994;65-8.
- Reid CA, Small A, Avery SM, and Buncic S. Presence of food-borne pathogens on cattle hides. Food Control. 2002; 13(6-7):411-5.
- Mohammed O, Shimelis D, Admasu P, Feyera T. Prevalence and antimicrobial susceptibility pattern of *E. coli* isolates from raw meat samples obtained from abattoirs in Dire Dawa City, eastern Ethiopia. International Journal of Microbiological Research. 2014;5(1):35-9.

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