

Poster 11

High occurrence of multidrug-resistant *Escherichia coli* from Holstein-Friesian cattle in Northern Portugal: a One Health challenge

S. Quinteira^{1,2,3,*}, A. B. Dias⁴, L. Pinho⁵, N. V. Brito¹, C. Campos^{6,7}, A. R. Freitas^{1,8,9} and C. Miranda^{1,10}

¹ TOXRUN-Toxicology Research Unit, University Institute of Health Sciences, CESPU, CRL., Avenida Central de Gandra 1317, 4585-116 Gandra PRD, Portugal

² BIOPOLIS/CIBIO-InBIO, Centro de Investigação em Biodiversidade e Recursos Genéticos, Universidade do Porto, Rua Padre Armando Quintas, n° 7, 4485-661, Vairão, Portugal

³ Departamento de Biologia, Faculdade de Ciências, Universidade do Porto, Rua do Campo Alegre s/n, 4169-007 Porto, Portugal

⁴ University Institute of Health Sciences, CESPU, Avenida Central de Gandra 1317, 4585-116 Gandra PRD, Portugal

⁵ Department of Veterinary Clinics, Abel Salazar Biomedical Sciences Institute, University of Porto, Porto, Portugal

⁶ Instituto Português de Oncologia do Porto Francisco Gentil, 4200-072, Porto, Portugal

⁷ Escola Superior de Saúde, Instituto Politécnico do Porto, 4200-072, Porto, Portugal

⁸ UCIBIO/REQUIMTE, Applied Molecular Biosciences Unit, Department of Biological Sciences, Laboratory of Microbiology, Faculty of Pharmacy, University of Porto, Porto, Portugal

⁹ Associate Laboratory i4HB, Institute for Health and Bioeconomy, Faculty of Pharmacy, University of Porto, Portugal

¹⁰ Associated Laboratory for Green Chemistry (LAQV-REQUIMTE), University NOVA of Lisbon, Caparica, Portugal

* Correspondence: sandra.quinteira@ipsn.cespu.pt

Abstract

Background: *Escherichia coli* is currently a leading pathogen for deaths associated with antimicrobial resistance (AMR) [1]. AMR related to food-producing animals is a public health risk requiring a One Health approach [1,2]. Animals are major reservoirs of antibiotic resistant *E. coli* which can easily reach humans through the food chain, direct contact, or the environment [3]. **Objective:** To analyze the AMR profiles of *E. coli* in fecal samples from Holstein-Friesian cattle, as information on AMR from dairy farming in Portugal is scarce. **Methods:** Samples (n=112) collected from 7 different farms at Northern Portugal were pooled during February-March 2023 based on age group (8 calves and 8 cows per farm). Characteristic *E. coli* colonies were selected from MacConkey Agar supplemented with or without antibiotics (4 µg/ml cefotaxime; 3 µg/ml colistin), for confirmation by MALDI-TOF mass spectrometry and antimicrobial susceptibility testing (AST), according to EUCAST/CLSI guidelines. ESBL phenotype were searched using the Double-Disk Synergy Test. **Results:** Presumptive *E. coli* isolates (n=110) representing different farming production systems were obtained and 51 representatives were identified by MALDI-TOF and tested by AST. From both calves and cows, 95% and 63% of the *E. coli* isolates exhibited resistance to ≥ 1 antimicrobial classes and multidrug resistance (MDR, resistance to ≥ 3 antimicrobial classes), respectively. Isolates were mostly resistant to gentamycin (88%), ampicillin (72%), amoxicillin plus clavulanic acid (63%), tetracycline (63%) and cefotaxime (53%). Only two isolates showed susceptibility to all tested antibiotics. ESBL activity was observed in 21% of *E. coli* isolates. All analyzed pools were classified as MDR. **Conclusions:** This is one of the first studies on AMR rates of *E. coli* from dairy cattle in Northern Portugal. It reveals a high prevalence of MDR *E. coli*, which is a worrying finding and emphasize the need of a multisectoral One Health approach to minimize its impact.

Keywords: One Health; antimicrobial resistance; livestock; *Escherichia coli*

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References

1. Aslam B, Khurshid M, Arshad MI, Muzammil S, Rasool M, Yasmeen N, Shah T, Chaudhry TH, Rasool MH, Shahid A, Xueshan X and Baloch Z. Antibiotic Resistance: One Health one world outlook. *Front Cell Infect Microbiol* **2021**,11,771510.
2. Ma F, Xu S, Tang Z, Li Z, Zhang L. Use of antimicrobials in food animals and impact of transmission of antimicrobial resistance on humans. *Biosafety and Health* **2021**,3,32-38.
3. Samtiya M, Matthews KR, Dhewa T, Puniya AK. Antimicrobial resistance in the food chain: trends, mechanisms, pathways, and possible regulation strategies. *Foods* **2022**,11,2966.



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