BOOK OF ABSTRACTS

STH MEETING OF YOUNG RESEARCHERS OF UNIVERSITY OF PORTO IJUP'12

TJUP'12

. .

.

•

5th MEETING OF YOUNG RESEARCHERS OF UNIVERSITY OF PORTO



CREDITS

Livro de Resumos IJUP'12

5º Encontro de Investigação Jovem da U.Porto

© Universidade do Porto AA ID+i T. 22 040 81 46 secidi@reit.up.pt

Design

Tiago Campeã Rui Mendonça

Impressão e acabamentos Invulgar – artes gráficas

Tiragem 1300 exemplarès

Depósito Legal

340336/12

ISBN

978-989-8265-82-1

Tracking antibiotic resistance along the Silk Road

<u>M. Mendes¹</u>, C. Novais^{3,5}, P. Antunes^{3,4}, J. Campos³, A. Freitas³, A. Abdukadir⁷, L. Peixe³, N. Monteiro^{2,5} and S. Quinteira^{1,2,6}

¹ Department of Biology, Faculty of Sciences, University of Porto, Portugal.
² CIBIO - Research Center in Biodiversity and Genetic Resources. University of Porto, Portugal.
³ REQUIMTE. Faculty of Pharmacy, University of Porto, Portugal.
⁴ Faculty of Food Sciences and Nutrition, University of Porto, Portugal

⁵ CEBIMED. Health Sciences Faculty. Fernando Pessoa University, Porto, Portugal

⁶ CITS - Research Center in Health Technologies, Polytechnic Health Institute of the North (IPSN/CESPU), Famalicão, Portugal.

⁷ Xinjiang Institute of Ecology and Geography, Chinese Academy of Sciences, Urumqi, China.

Information on the bacterial communities that inhabit wild-animal species, some of which facing extinction in the near future, as well as information on the spreading of antibiotic resistance into relatively pristine ecosystems, remains scarce. Accordingly, the goal of this study was to evaluate if wild or feral animals, inhabiting remote areas of the globe (from the Tibetan plateau to the Gobi desert), carry antibiotic resistant bacteria. During the SilkRoad2010 expedition (CIBIO and Chinese Academy of Sciences), faecal samples were collected from nine iconic mammal species, most of which endangered or in the verge of extinction. Faecal samples (0.1ml) were plated in different selective culture media (e.g. Slanetz-Bartley agar for Enterococcus spp., MacConkey agar for Enterobacteriaceae and non fermenter Gram negative bacteria, XLD and SS agar for Salmonella), with/without antibiotic supplements, before and after a pre-enrichment step. Different bacteria morphotypes were selected for further characterization. Susceptibility to several antibiotics was tested by disk diffusion method (CLSI). Two hundred and seventy bacterial isolates (97 Gram positive and 173 Gram negative) were collected from 22 faecal samples, from different animal species: Mongolian wild ass, Dhole, Mongolian gazelle, Przewalski horse, Gray marmot and Bactrian camel. Different antibiotic resistance rates were detected amongst Enterococcus spp: tetracycline-49,5% (48/97), erythromycin-36% (35/97), High Level of Resistance (HLR) to gentamicin -15,5% (15/97), ampicillin - 6% (6/97), and Quinupristin/dalfopristin -23,7% (23/97). Decreased susceptibility (n=10) and resistance (n=6) to vancomycin were observed. Among Gram negative lactose fermentors, resistance was observed for ampicillin - 39% (31/79), ampicillin+clavulanic acid -15% (12/79) and cefotaxime -16,5% (13/79). Variable susceptibility to aminoglycosides was observed: gentamicin-19% (21/108); tobramycin-16% (20/123). Two isolates (obtained from Bactrian camel and Gray marmot) showed resistance to imipenem. Although preliminary, results emerging from this study show that even though inhabiting remote areas with extremely low human pressure, the critically endangered Silk Road fauna surprisingly harbours bacteria carrying antibiotic resistance. As so, drug resistance, far from limited to hospitals, may be spreading into the most remote areas of the globe.

Acknowledgments: This research was supported by Universidade do Porto/Santander Totta "Projectos Pluridisciplinares 2010".