

# Errata

## Title page

Fluoroquinolone resistance in the environment and the human gut

## List of papers, page iv

- Paper I      **Acquired genetic mechanisms of a multiresistant bacterium isolated from a treatment plant receiving wastewater from antibiotic production**  
Anna Johnning, Edward R. B. Moore, Liselott Svensson-Stadler, Yogesh S. Shouche, D. G. Joakim Larsson & Erik Kristiansson  
*Applied and Environmental Microbiology* 79.23 (2013): 7256-7263.
- Paper II      **Isolation of novel broad host fluoroquinolone resistance plasmids from an antibiotic-polluted lake**  
Carl-Fredrik Flach, Anna Johnning, Ida Nilsson, Kornelia Smalla, Erik Kristiansson & D. G. Joakim Larsson  
*Manuscript*
- Paper III     **Resistance mutations in *gyrA* and *parC* are common in bacterial communities of both pristine and fluoroquinolone-polluted environments**  
Anna Johnning, Erik Kristiansson, Jerker Fick, Birgitta Weijdegård & D.G. Joakim Larsson  
*Submitted*
- Paper IV     **International travel affects the abundance of chromosomal quinolone resistance mutations in the human gut microbiome**  
Anna Johnning, Erik Kristiansson, Martin Angelin, Nachiket Marathe, Yogesh S. Shouche, Anders Johansson & D.G. Joakim Larsson  
*Submitted*

## Methodological Considerations, page 12

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