

Assessing the possible association between veterinary antimicrobial consumption and resistance in indicator E. coli isolated from farm animals in Belgium

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Introduction

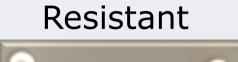
Antimicrobial use is significant factor accounting for the selection and spread of antimicrobial resistance in commensal and pathogenic bacteria (Burow et al., 2014; Horigan et al., 2016). Bacteria are frequently found resistant to many antimicrobials to the point that both animal and public health are now seriously challenged (Megha et al., 2014). Exploring the trend possibly associating antimicrobial consumption and resistance is a highly desirable exercise, that was tentatively completed in the present study focused on indicator *Escherichia coli* from farm animals in Belgium.

Materials and methods

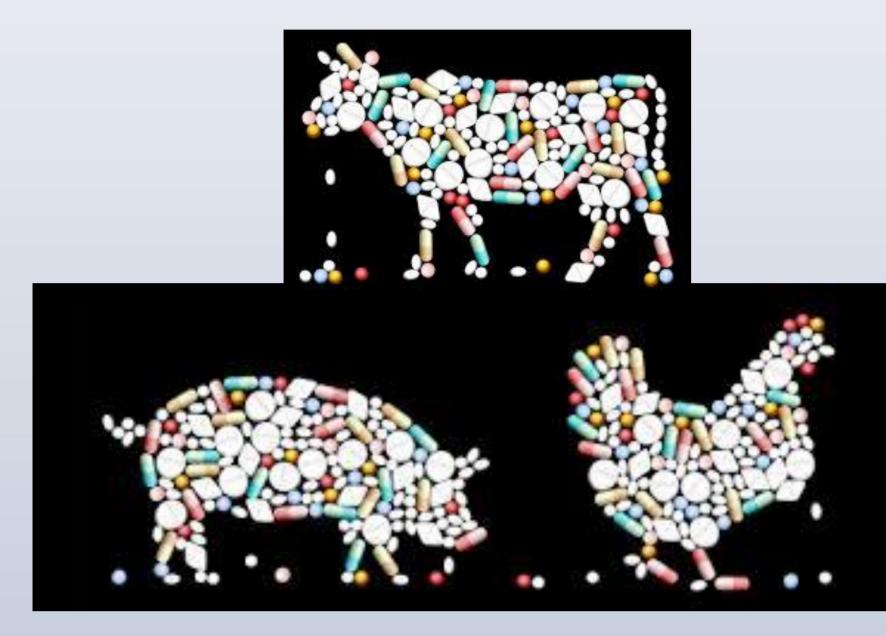
Antimicrobial consumption evaluation

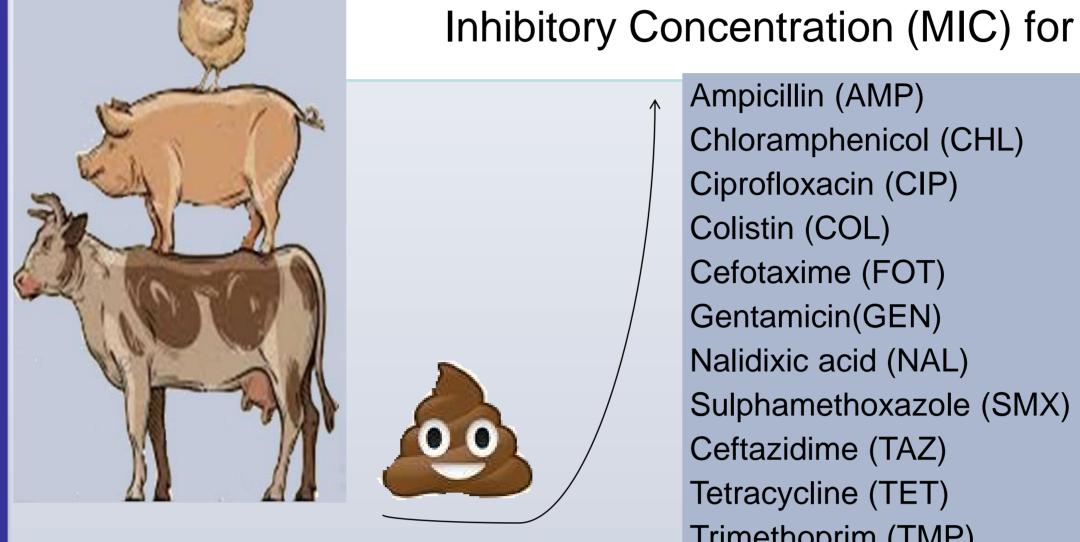
Total of all veterinary antimicrobials sold in Belgium (2011 to 2015) adjusted with the biomass

Antimicrobial resistance evaluation



0





R m

≥170 fecal isolates per year (2011-2015) for veal calves, beef cattle, broiler chickens, pigs

Ampicillin (AMP) Chloramphenicol (CHL) Ciprofloxacin (CIP) Colistin (COL) Cefotaxime (FOT) Gentamicin(GEN) Nalidixic acid (NAL) Sulphamethoxazole (SMX) Ceftazidime (TAZ) Tetracycline (TET) Trimethoprim (TMP)

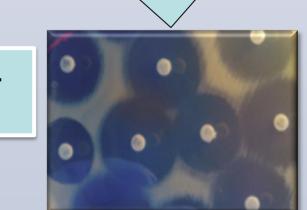
π=0.4

p-value=0.3272

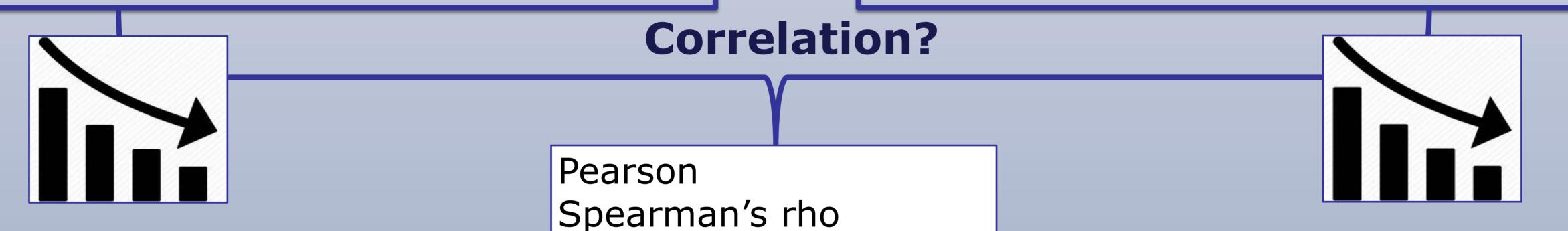
Epidemiological Cutoffs defined by the European Committee on Antimicrobial Susceptibility Testing

If MIC>cut-off

If MIC≤cut-off

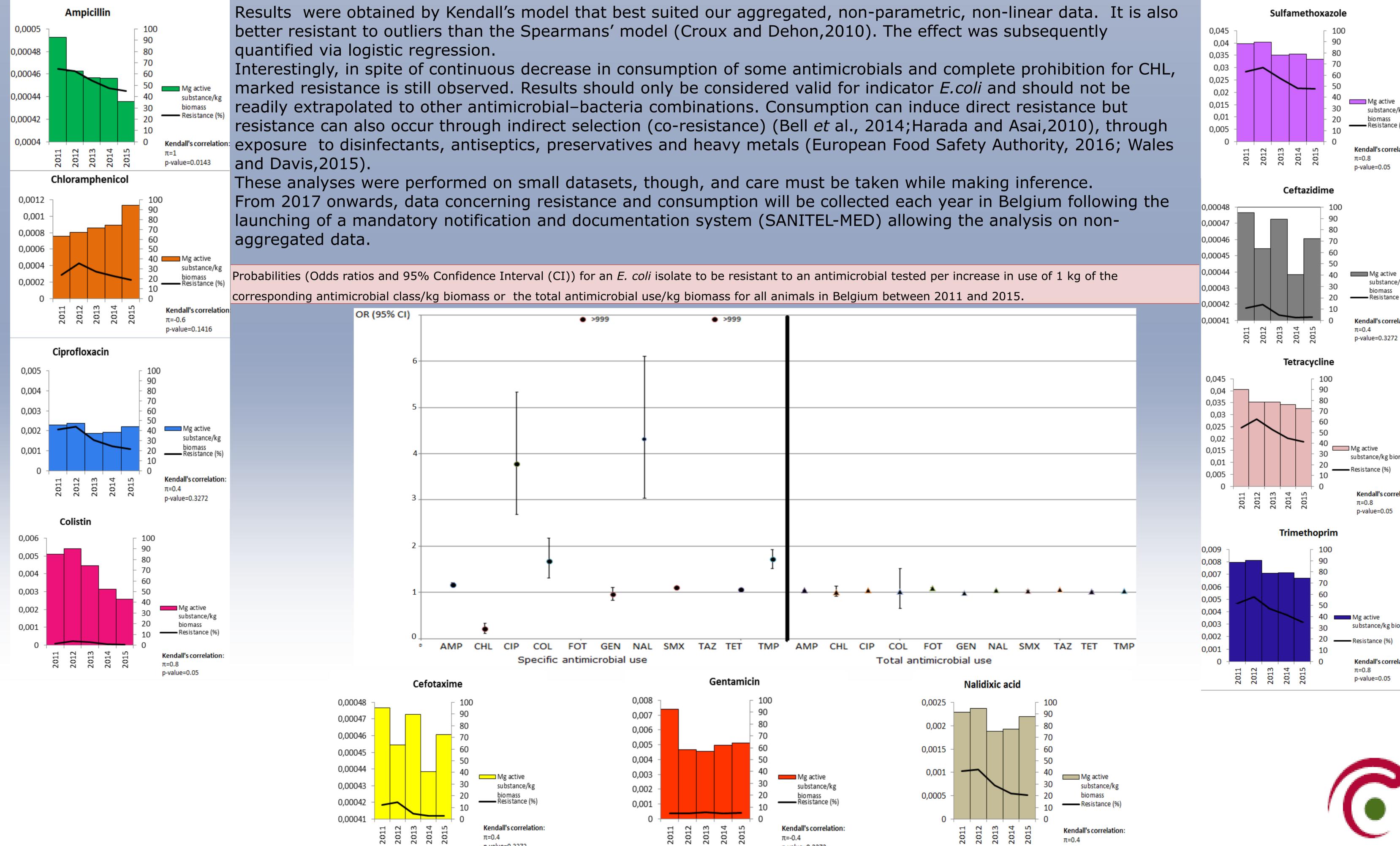


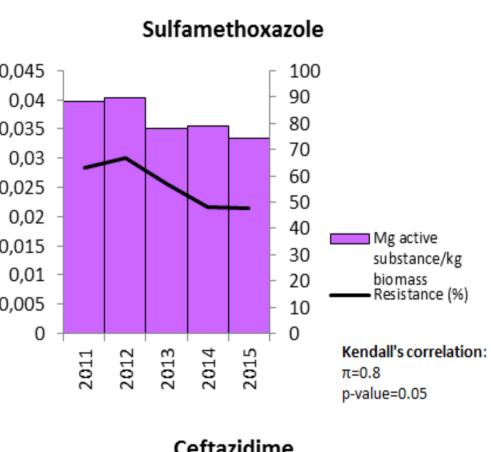
Susceptible

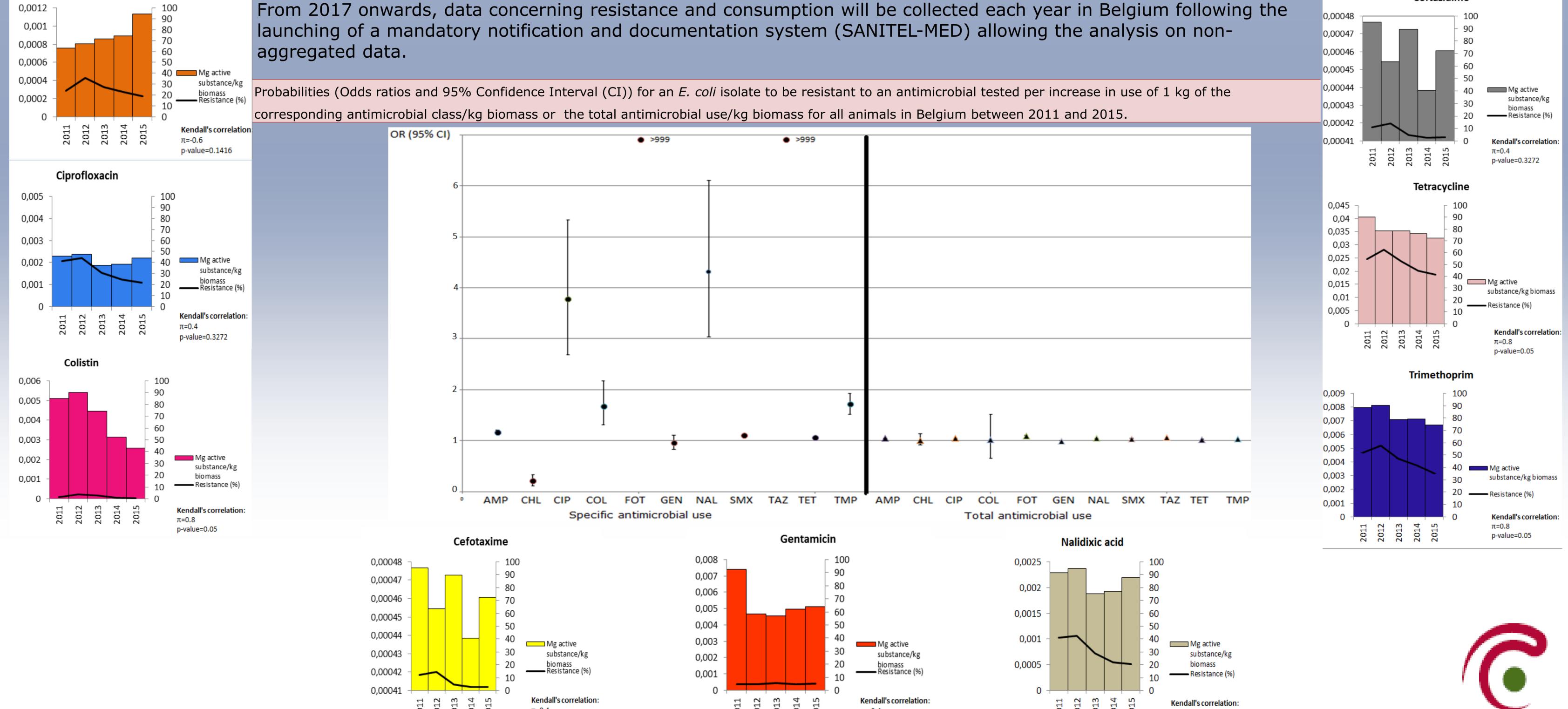


Kendall's tau Logistic regression

Results and perspectives







p-value=0.3272

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p-value=0.3272

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