

Antimicrobial Resistance Patterns of *Campylobacter* from Pig and Cattle Carcasses in Poland

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Abstract : *Campylobacter* is recognized as the main cause of bacterial gastrointestinal infections in Europe. A main source of the pathogen is poultry and poultry meat; however, other animals like pigs and cattle can also be reservoirs of the bacteria. Human *Campylobacter* infections are often self-limiting but in some cases, macrolide and fluoroquinolones have to be used. The aim of this study was to determine antimicrobial resistance patterns (AMR) of *Campylobacter* isolated from pig and cattle carcasses. Between July 2009 and December 2015, 735 swabs from pig (n = 457) and cattle (n = 278) carcasses were collected at Polish slaughterhouses. All samples were tested for the presence of *Campylobacter* by ISO 10272-1 and confirmed to species level using PCR. The antimicrobial susceptibility of *Campylobacter* isolates was determined by a microbroth dilution method with six antimicrobials: gentamicin (GEN), streptomycin (STR), erythromycin (ERY), nalidixic acid (NAL), ciprofloxacin (CIP) and tetracycline (TET). It was found that 167 of 735 samples (22.7%) were contaminated with *Campylobacter*. The vast majority of them were of pig origin (134; 80.2%), whereas for cattle carcasses *Campylobacter* was less prevalent (33; 19.8%). Among positive samples *C. coli* was predominant species (123; 73.7%) and it was isolated mainly from pig carcasses. The remaining isolates were identified as *C. jejuni* (44; 26.3%). Antimicrobial susceptibility indicated that 22 out of 167 *Campylobacter* (13.2%) were sensitive to all antimicrobials used. Fourteen of them were *C. jejuni* (63.6%; pig, n = 6; cattle, n = 8) and 8 was *C. coli* (36.4%; pig, n = 4; cattle, n = 4). Most of the *Campylobacter* isolates (145; 86.8%) were resistant to one or more antimicrobials (*C. coli*, n = 115; *C. jejuni*, n = 30). Comparing the AMR for *Campylobacter* species it was found that the most common pattern for *C. jejuni* was CIP-NAL-TET (9; 30.0%), whereas CIP-NAL-STR-TET was predominant among *C. coli* (47; 40.9%). Multiresistance, defined as resistance to three or more classes of antimicrobials, was found in 57 *C. coli* strains, mostly obtained from pig (52 isolates). On the other hand, only one *C. jejuni* strain, isolated from cattle, showed multiresistance with pattern CIP-NAL-STR-TET. Moreover, CIP-NAL-STR-TET was characteristic for most of multiresistant *C. coli* isolates (47; 82.5%). For the remaining *C. coli* the resistance patterns were CIP-ERY-NAL-TET (7 strains; 12.3%) and for one strain of each patterns: ERY-STR-TET, CIP-STR-TET, CIP-NAL-GEN-STR-TET. According to the present findings resistance to erythromycin was observed only in 11 *C. coli* (pig, n = 10; cattle, n = 1). In conclusion, the results of this study showed that pig carcasses may be a serious public health concern because of contamination with *C. coli* that might features multiresistance to antimicrobials.

Keywords : antimicrobial resistance, *Campylobacter*, carcasses, multi resistance

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