

Fluoroquinolone-Resistant *Escherichia coli* and *Proteus mirabilis* in Poultry of Middle Moravia, Czech Republic

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Abstract

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The aim of the study was to detect antibiotic resistance of *Escherichia coli* and *Proteus mirabilis* isolates from farm-reared poultry.

During the period of June 2003 to June 2004, samples of cloacal swabs obtained from 5 poultry farms located in the central part of Moravia, Czech Republic were examined using aerobic cultivation and afterwards *E. coli* and *P. mirabilis* isolates were identified. Minimal inhibitory concentrations of antibiotics tested were determined by a microdilution method. Pulsed-field gel electrophoresis (PFGE) was performed using *Sma*I restriction endonuclease.

A total of 300 samples of cloacal swabs from healthy layer hens were cultivated and 239 *E. coli* and 127 *P. mirabilis* strains were isolated. In the case of *E. coli*, 7 isolates (3%) from 4 different farms were identified manifesting resistance to ofloxacin and ciprofloxacin. Out of 24 *P. mirabilis* isolates resistant to the tested fluoroquinolones, 20 strains were isolated from one of the farms and PFGE analysis of DNA proved that 19 isolates were probably identical and represented one clonal type.

The study confirmed the occurrence of multiresistant bacterial isolates with resistance to fluoroquinolones in poultry in the Czech Republic. Their clonal spread in farm-reared poultry can be suggested, too.

Poultry, faecal bacteria, ofloxacin, ciprofloxacin, resistance

Nowadays, increasing bacterial resistance to antibiotic agents including fluoroquinolones poses a serious problem (Jones et al. 1997; Neu 1992). This adverse trend is documented both in human and veterinary medicine and apparently its solution is to be seen in the synergism of preventive measures (Caprioli et al. 2000; Bogaard and Stobberingh 2000). The increasing bacterial resistance to fluoroquinolones both in human and animal populations is evidently connected with their overuse (Aguilar et al. 1992; Garau et al. 1999; Pena et al. 1995).

In veterinary medicine, fluoroquinolones are used in the treatment of both farm and pet animals. According to the data provided by the Institute for the State Control of Veterinary Biologicals and Medicaments, the overall usage of quinolones in Czech veterinary medicine in 2003 represented 1532.9 kg (Hera 2005). Many fluoroquinolones used in veterinary medicine belong to “antibiotic agents with indication limits”, a group of antibiotics that should be applied in serious animal infections only, based on clinical experience, diagnostic confirmation of the causal microorganism and resistance to “common” antimicrobial agents.

The aim of the study was to detect resistance to antimicrobial agents in *Escherichia coli* and *Proteus mirabilis* isolates from poultry bred in the central part of Moravia (Czech Republic), a region with extensive agricultural production. In fluoroquinolone-resistant

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isolates, the analysis of genomic DNA was performed and the degree of their similarity determined.

Materials and Methods

Collection of isolates

During the period of June 2003 to June 2004, samples of cloacal swabs from 5 poultry farms were examined. The farms were located in the central part of Moravia, Czech Republic. The parental breeds produced the final crossbreed of Brown Hissex. In each poultry farm, a total of 60 samples of cloacal swabs were taken from 60 healthy layer hens, i.e. 1 sample per hen. Each clinical sample was cultivated under aerobic conditions on conventional selective media (Endo agar, Xylose-Lysine-Desoxycholate agar). *E. coli* and *P. mirabilis* isolates were determined by standard biochemical procedures using Enterotest 24 (Pliva Lachema CZ).

Antibiotic susceptibility testing

Minimal inhibitory concentrations (MICs) of antibiotics tested for *E. coli* and *P. mirabilis* isolates were determined by a microdilution method according to the National Committee for Clinical Laboratory Standards guidelines (NCCLS 2000). Concentrations of tested isolates in Mueller-Hinton broth (Oxoid UK) were prepared to obtain 0.5 McFarland turbidity. MICs were read after 18 h of incubation at 37 °C. The MIC was interpreted as the lowest concentration of the antibiotic that visibly inhibited bacterial growth. As MIC breakpoints, the following values based on the National Institute of Public Health's National Reference Laboratory for Antibiotics recommendations (Urbášková 1998) were used: 1 mg/l for ciprofloxacin, 2 mg/l for ofloxacin and tetracycline, 4 mg/l for ampicillin, cefazolin, cefuroxime, cefotaxime, ceftazidime, chloramphenicol, gentamicin, colistin and meropenem, 8 mg/l for amikacin, oxolinic acid, trimethoprim and ampicillin/sulbactam, and 32 mg/l for trimethoprim-sulfamethoxazole and nitrofurantoin. MIC₉₀ and MIC₅₀ include 90% and 50% isolates with the given or lower MIC value of the respective antibiotics in the appropriate bacterial species.

Reference strains *E. coli* ATCC 25922, *E. coli* ATCC 35218 and *Pseudomonas aeruginosa* ATCC 27853 were used for protocol quality control.

Molecular analysis of fluoroquinolone-resistant isolates

Pulsed-field gel electrophoresis (PFGE) was performed using *Sma*I restriction endonuclease. Genomic DNA isolation was carried out using protocols published by Pantůček et al. (1997).

Restriction cleavage of the blocks 1 × 1 × 5 mm was done in restriction solution [8 µl of restriction buffer for *Sma*I (Sigma-Aldrich), 65 µl of deionised water and 10 U of restriction enzyme *Sma*I]. The blocks were incubated at 25 °C overnight.

PFGE was prepared in 1.2% agarose gel (Amresco) in 1xTBE buffer. PFGE was carried out at BioRad CHEF-DR II for 28 h at a pulse time of 0.1 to 30 s, 5.0 V/cm, with angle 120°. The gel was stained in ethidium bromide solution (Sigma-Aldrich) (1 µg/ml).

The data about administration of antibiotics in observed poultry farms are not available.

Results

In total, 300 samples of cloacal swabs from poultry (parental breed of Brown Hissex) were cultivated and 239 *E. coli* and 127 *P. mirabilis* isolates were identified. Antibiotic resistance of the investigated species is given in Table 1. A higher rate of resistant strains in the case of *E. coli* was proved in tetracycline (48%) and ampicillin (29%). As for the *P. mirabilis* strains, higher resistance to trimethoprim (31%) and trimethoprim-sulfamethoxazole (28%) was documented, apart from their natural resistance to colistin, nitrofurantoin and tetracycline.

Among *E. coli* isolates, 7 (3%) resistant to ofloxacin and ciprofloxacin were isolated from 4 poultry farms, while 4 isolates were detected in farm IV. Out of the 24 *P. mirabilis* isolates resistant to the tested fluoroquinolones, 20 were isolated from farm IV (Table 2).

In *E. coli*, 5 other isolates were detected, resistant to oxolinic acid (MICs ≥ 16 mg/l), but with persisting sensitivity to ofloxacin and ciprofloxacin. However, MICs of these isolates reached higher values (1-2 mg/l in ofloxacin and 0.5 - 1 mg/l in ciprofloxacin).

Fluoroquinolone-resistant isolates of *E. coli* were also resistant to tetracycline, trimethoprim-sulfamethoxazole and trimethoprim; 4 isolates out of 7 were resistant even to gentamicin. *P. mirabilis* isolates resistant to fluoroquinolones were also resistant to trimethoprim-sulfamethoxazole, trimethoprim and, in 67%, to gentamicin.

In total, 20 fluoroquinolone-resistant isolates of *P. mirabilis* isolated from the same farm

Table 1. Resistance of *Escherichia coli* and *Proteus mirabilis* to antibiotics with MIC range, MIC₅₀ and MIC₉₀

Antimicrobial agents	<i>Escherichia coli</i>				<i>Proteus mirabilis</i>			
	MIC range [mg/l]	MIC ₅₀ [mg/l]	MIC ₉₀ [mg/l]	resistance [%]	MIC range [mg/l]	MIC ₅₀ [mg/l]	MIC ₉₀ [mg/l]	resistance [%]
amikacin	0.125-64	1	4	1	0.5-16	1	4	2
ampicillin	0.5-128	2	128	29	0.25-128	1	4	9
ampicillin-sulbactam	0.125-8	1	4	0	0.125-4	0.5	2	0
cefazolin	0.5-64	2	4	4	1-128	2	8	7
cefotaxime	0.06-0.5	0.03	0.125	0	0.03-1	0.03	0.125	0
ceftazidime	0.03-0.5	0.03	0.125	0	0.03-0.5	0.03	0.125	0
cefuroxime	0.25-32	0.25	2	1	0.25-4	0.25	1	0
ciprofloxacin	0.03-8	0.03	0.25	3	0.03-8	0.06	2	19
colistin	0.06-2	0.25	1	0	≥ 64	-	-	100*
trimethoprim-sulfamethoxazole	0.5-512	8	128	10	0.5-512	8	64	28
chloramphenicol	0.25-128	1	4	4	0.5-128	1	4	17
gentamicin	0.125-32	0.5	2	2	0.125-128	0.5	4	13
meropenem	0.03-2	0.125	0.5	0	0.06-0.5	0.06	0.25	0
nitrofurantoin	0.5-32	2	8	0	≥ 64	-	-	100*
ofloxacin	0.03-16	0.06	1	3	0.03-32	0.5	4	19
oxolinic acid	0.125-64	1	4	5	0.25-128	2	16	19
tetracycline	0.25-128	2	64	48	≥ 64	-	-	100*
trimethoprim	0.125-128	2	8	11	0.5-64	2	16	31

* natural resistance

Table 2. Occurrence of fluoroquinolone-resistant *Escherichia coli* and *Proteus mirabilis* isolates in observed poultry farms

Poultry farm	No. of fluoroquinolone-resistant isolates	
	<i>E. coli</i>	<i>P. mirabilis</i>
I	1	1
II	0	0
III	1	2
IV	4	20
V	1	1
Total	7	24

were analyzed by PFGE. The results are shown in Fig. 1 (see Plate XIII). Of these, 19 isolates were identical and represented one clonal type. A unique isolate shows 60% similarity of a DNA profile in *Sma*I spectrum with other probably identical isolates.

Discussion

The relation between the application of antibiotics and the dissemination of bacterial resistance from animals to humans has been described by Hummel et al. (1996). Also, the study by van den Bogaard et al. (2001) confirmed that transmission of resistant clones and resistance plasmids of *E. coli* from poultry to humans commonly occurs.

E. coli is a major pathogen of worldwide importance in commercially produced poultry (Gross 1994). Although this bacterial species is commonly found in the intestinal tract of

hens, some serotypes can cause diseases. Antibiotic therapy is therefore an important tool for reducing mortality associated with *E. coli* infections (Dho-Moulin 1993; González et al. 1990). On the other hand, the use of antibiotics can select multiresistant bacteria. A study performed in Spain showed the resistance of avian *E. coli* isolates to ofloxacin and ciprofloxacin in 14% and 13%, respectively (Blanco et al. 1997). In the present study, the corresponding value reached 3%. In *P. mirabilis* isolates, the frequency of fluoroquinolone-resistant isolates was much higher. However, it must be stressed that 83% of the isolates were obtained from one farm, suggesting that these probably originated from one resistant clone. Thus the average resistance level cannot be determined.

Isolates of both species resistant to fluoroquinolones were multi-resistant, manifesting resistance to trimethoprim-sulfamethoxazole, trimethoprim and, in the case of *E. coli* isolates, even to tetracycline. Higher occurrence of gentamicin-resistant isolates was also notified. Similar results were reported by Bazile-Pham-Khac et al. (1996) who described simultaneous resistance to tetracycline, trimethoprim and tobramycin in fluoroquinolone-resistant *E. coli* isolates. It is to be noted that in *E. coli* five other isolates were identified, manifesting resistance to oxolinic acid. Ofloxacin and ciprofloxacin MIC ranges were 1 - 2 mg/l and 0.5 - 1 mg/l, respectively, and, according to the breakpoints applied, these isolates were sensitive to fluoroquinolones. These are probably isolates with single mutations in *gyrA*, i.e. with decreased sensitivity (McDonald et al 2001). This finding should be the first sign of the negative phenomenon of increasing resistance to fluoroquinolones. Therefore, sensitivity testing of gram-negative rods isolated from poultry should involve not only fluoroquinolones, such as ofloxacin or ciprofloxacin, but oxolinic acid as well.

Based on the analysis of the genomic DNA in *P. mirabilis* isolates resistant to fluoroquinolones and isolated from one of the farms, it is possible to assume that it is most probably one identical isolate spread in the farm.

In conclusion, the results of our study support the occurrence of multiresistant bacterial isolates with resistance to fluoroquinolones in poultry in the Czech Republic and underline the importance of antibiotic policy implementation in veterinary medicine, including monitoring bacterial isolates with dangerous phenotypes of resistance.

Výskyt fluorochinolon-rezistentních kmenů *Escherichia coli* a *Proteus mirabilis* v chovech drůbeže na střední Moravě

Celosvětově stoupající rezistence bakterií k antimikrobním léčivům představuje vážný problém při terapii infekčních onemocnění i v epidemiologické praxi. Problém bakteriální rezistence je aktuální i v oblasti veterinární medicíny.

Cílem studie bylo zjistit výskyt rezistence k fluorochinolonům (ofloxacinu a ciprofloxacinu) u kmenů *Escherichia coli* a *Proteus mirabilis* izolovaných v 5 chovech drůbeže na střední Moravě. Z každého chovu bylo odebráno 60 kloakálních výtěrů od 60 zdravých slepic. Celkem bylo izolováno 239 kmenů *E. coli* a 127 kmenů *P. mirabilis*. Četnost fluorochinolon-rezistentních kmenů *E. coli* dosáhla hodnoty 3%, v případě *P. mirabilis* byla rezistence prokázána u 24 kmenů, přičemž 20 pocházelo ze stejného chovu. Kmeny obou species se rezistencí k fluorochinolonům byly multirezistentní se současnou rezistencí k trimethoprim-sulfamethoxazolu, trimethoprimu, u kmenů *E. coli* rovněž k tetracyklinu, dále byla pozorována vyšší četnost gentamicin-rezistentních kmenů. Na základě výsledků analýzy genomové DNA 20 fluorochinolon-rezistentních kmenů *P. mirabilis* izolovaných z jednoho chovu, lze s velkou pravděpodobností předpokládat šíření rezistentního klonu jednoho kmene. Z našich výsledků je patrné, že v drůbežích chovech na střední Moravě se vyskytují multirezistentní kmeny *E. coli* a *P. mirabilis* s rezistencí k fluorochinolonům a je tedy nutné vzít v úvahu možnost jejich šíření do lidské populace.

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