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Co-occurrence of extended spectrum beta- lactamase genes and mcr-1 gene in strains of *Escherichia coli* isolated from clinical cases of bovine mastitis in Greece

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Background: Clinical bovine mastitis triggered by *Escherichia coli* is costly disease with clinical signs that vary from mild mastitis to very severe or even fatal forms. The disease has also a potential threat to public health since cow's milk infected with enterobacteria may represent vector of antibiotic resistance genes such as extended spectrum β -lactamase genes and *mcr-1*. Isolates of *E. coli*, therefore, harvested from animals, foodstuff, and human beings that expose co-resistance to b-lactam antibiotics and to colistin are globally reported. The prevalence of such microorganisms is poorly studied in Greece. This research investigates the prevalence of extended spectrum β -lactamase genes and *mcr-1* gene in *E.coli* isolates harvested from mastitic milk samples in dairy cows

Material/methods: Four hundred samples of mastitic milk from equal number of cows were enrolled during 2015-2016. Fifty μ l of each sample was plated on TBX agar. After incubation at 44° C for 24–48 h, plates were inspected for colonies growth. One β -glucuronidase positive colony per plate was purified on nutrient agar and further identified as *E. coli* by API 20E identification strips. Species identification of the isolates was confirmed by polymerase chain reaction that detects *usp(A)* gene, which is characteristic for *E. coli* strains. Antimicrobial susceptibility tests were done on Mueller-Hinton agar using Kirby Bauer disk diffusion method. The antimicrobial agents tested were: nalidixic acid (30

µg), ciprofloxacin (5 µg), doxycycline (30 µg) tetracycline (30 µg), gentamicin (10 µg), trimethoprim & sulfamethoxazole (1.75/23.75 µg), streptomycin (10 µg), ceftriaxone (30 µg), ceftazidime (30 µg), amoxicillin-clavulanic acid (10/20 µg), aztreonam (30 µg), chloramphenicol (30 µg), ampicillin (10 µg), and piperacillin (100 µg). Resistance data were interpreted according to CLSI guidelines (M100-21/2014). MICs of colistin were obtained on Mueller-Hinton agar by the agar dilution method in antibiotic concentrations ranging from 0.25 to 128 µg/ml and inoculum of 10⁴ colony-forming units per spot. PCR/DNA sequencing was used to characterize *bla* genes and *mcr-1* gene

Results: The overall prevalence of *E.coli* isolated from cows' mastitic milk was 22% (89/400). From the 89 *E. coli* isolates that were tested 18 exposed co-resistance to ampicillin, piperacillin, amoxicillin-clavulanic acid, ceftriaxone, ceftazidime, and aztreonam. All 18 isolates were further characterized by PCR as ESBL producers (16 CTX-M and 2 SHV). Interestingly MICs against colistin of 6 of the 18 ESBL producers ranged from 8 µg/ml to 32 µg/ml. Finally these 6 strains carried the *mcr-1* resistance gene.

Conclusions: This is the first report of the co-occurrence of extended spectrum β-lactamase genes and *mcr-1* gene in strains of *E. coli* isolated from clinical cases of bovine mastitis in Greece. Finally these findings are also important in relation to the use of β-lactam antibiotics for the treatment of bovine clinical mastitis.