Correction to: Antimicrobial resistance and the presence of extended-spectrum beta-lactamase genes in *Escherichia coli* isolated from the environment of horse riding centers

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The original publication of this paper contains a mistake. The correct presentation of Table 5 and caption of fig. 2 is shown in this paper.

Fig. 2 UPGMA dendrograms of *E. coli* strains isolated from various environments (A-air, M-manure, S-swabs) of three horse riding centers. **a** – KJK Szary, **b** – OJK Pegaz, **c** – SKH Nielepice

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Table 5 Characteristics of most frequent^a haplotypes within strains isolated from the analyzed horse riding centers

Haplotype	No. of isolates	Detected in (environment)			Detected in (facility)		
		Air (n=24)	Manure (n=75)	Swabs (n=83)	SKH Nielepice (n=30)	OJK Pegaz (n=58)	KJK Szary (n=94)
Ec_1	27	3	15	9	5	9	13
Ec_2	17	6	4	7	1	2	14
Ec_3	14	1	6	7	0	11	3
Ec_4	10	2	4	4	3	5	2
Ec_5	8	0	5	3	3	4	1
Ec_6	8	0	5	3	2	4	2
Ec_7	7	0	6	1	6	0	1
Ec_8	7	0	3	4	0	7	0
Ec_9	6	0	2	4	0	4	2
Ec_10	5	0	2	3	1	0	4
Ec_11	4	3	0	1	1	3	0
Ec_12	4	0	0	4	2	2	0
Ec_13	3	1	1	1	0	0	3
Ec_14	3	0	2	1	1	0	2
Ec_15	3	0	2	1	0	0	3
Ec_16	3	2	0	1	1	2	0

^a Only haplotypes specific for three or more isolates are presented