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1 **Field trial evaluating changes in prevalence and patterns of**
2 **antimicrobial resistance among *Escherichia coli* and *Enterococcus* spp.**
3 **isolated from growing broilers medicated with enrofloxacin,**
4 **apramycin and amoxicillin**

5
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27

28 Abstract

29 The present study investigates, under field conditions, the influence of antimicrobial
30 administration on prevalence and patterns of antimicrobial resistance among
31 *Escherichia coli* and *Enterococcus* spp. isolated from growing broilers. For this
32 purpose, a group of 16 000 commercial broiler chickens was treated with enrofloxacin
33 from day 1 to day 3, gentamicin from day 19 to day 21, and ampicillin from day 26 to
34 day 28. A control group of 16 000 broilers was placed in the same controlled
35 environment poultry house. Fecal (from both groups) and feed samples were collected
36 at regular intervals. Few *E. coli* isolates were obtained from either farm environment or
37 poultry feed samples, while enterococci were found to be ubiquitous among these
38 samples. The frequency of resistance against most antimicrobials tested was
39 significantly higher ($P < 0.05$) in *E. coli* isolated from broilers receiving intermittent
40 antimicrobial pressure than that from non-medicated broilers, whereas in enterococci
41 these differences were only observed among structurally related antimicrobial drugs and
42 over a short period of time. By the time the broilers reached market age (33 days),
43 several multi-resistant *E. coli* and enterococci were detected in the feces of the
44 medicated group. Results suggest that antimicrobial resistance in *E. coli* was mainly
45 medication-dependent, whereas among enterococci, changes observed over time were
46 apparently influenced by factors apart from antimicrobial exposure, namely the
47 resistance organisms previously present in farm environment and those present in
48 feedstuffs.

49

50 Keywords: antimicrobials; resistance; *Escherichia coli*; enterococci; poultry.

51

52

53 1. Introduction

54 Historically, the assessment of the biological consequences associated with
55 antimicrobial use in food animals was restricted to zoonotic enteropathogens. During
56 the last decade, there has been an increasing awareness about the effects of
57 antimicrobial drugs on resistance in animal commensal flora, such as *Escherichia coli*
58 and enterococci (van den Bogaard et al., 2002; McDermott et al., 2005; Diarra et al.,
59 2007; Smith et al., 2007). Antimicrobial resistance determinants, generated and/or
60 amplified at the farm level, interfere with therapeutic efficiency (increasing morbidity
61 and mortality associated with disease outbreaks), and may be later transferred to other
62 animals or humans either through direct contact, contamination of meat or, more
63 indirectly, through environmental pathways (Dancer, 2004). Broiler production is
64 unique among animal husbandry, as their fast growing cycle (30 to 42 days) and their
65 high stocking density (14 to 26 birds m⁻²) result in no tolerance for disease episodes.
66 Thus, preventive mass medication of the growing broilers, administering drugs at
67 therapeutic levels for short periods of time is carried out. Shortly after European Union
68 banned the use of growth promoters in animal production, there were some increases in
69 morbidity and mortality among farm animals that were controlled by a proportional
70 increase in therapeutic and preventive administration of antimicrobials (McEwen and
71 Fedorka-Cray, 2002). This has created serious concerns because the majority of these
72 antimicrobials are used in human medicine, or at least are structurally related.

73 To better understand the selective pressure exerted by antimicrobials under field
74 conditions, this study examined the effects on development and persistence of
75 antimicrobial resistance among commensal *E. coli* and *Enterococcus* spp. isolated from
76 broiler chickens medicated during their rearing cycle with enrofloxacin, gentamicin and
77 amoxicillin.

78

79 **2. Materials and Methods**80 **2.1. Animals, housing and treatments**

81 A controlled environment poultry house with 1 600 m² area was selected for this
82 study. Inside, 32 000 broiler chicks originating from the same breeder were divided in
83 two groups on a random basis: (i) Medicated group (M) (n = 16 000) medicated with
84 enrofloxacin (Baytril 10%[®] - Bayer, Carnaxide, Portugal), gentamicin (Gentamicina
85 Oral[®] - Vetlima, Lisboa, Portugal) and amoxicillin (Paracilina SP[®] - Intervet, Mem
86 Martins, Portugal); (ii) Control group (C) (n = 16 000). Isolated by a barrier, both
87 groups shared exactly the same rearing, environmental and sanitary conditions.
88 Medicated and control group were fed the same commercial corn-soy meal broiler diet,
89 containing anticoccidial products: narasin at 100 mg kg⁻¹ in starter and grower diets and
90 salinomycin at 60 mg kg⁻¹ in finisher.

91 The birds in the medicated group were given, for a period of three days,
92 therapeutic doses of enrofloxacin (23.8 mg kg⁻¹ of live weight), gentamicin (26.0 mg kg⁻¹
93 ¹) and amoxicillin (25.1 mg kg⁻¹) in their drinking water when they were 1, 19 and 26
94 days old, respectively. The antimicrobial concentrations in drinking water for these
95 treatments were as follows: enrofloxacin (50 mg l⁻¹), gentamicin (115 mg l⁻¹) and
96 amoxicillin (140 mg l⁻¹).

97 Both compartments had independent drinking systems and exterior entrances.
98 Strict biosecurity measures were observed, implying the use of disposable boots and
99 coveralls before each room entry. The broiler house was geographically remote from
100 others and operated under the principle of all-in/all-out; prior to flock placement, the
101 broiler house was thoroughly cleaned and disinfected and was left empty for 14 days

102 before the flock placement. The previous history of antimicrobial usage was known for
103 this commercial farm.

104 Once a day, the animals were observed by a veterinarian for checking general
105 health status and for any clinical symptoms of disease, to verify feed and water supply,
106 and to check the proper functioning of heating and ventilation systems.

107

108 **2.2. Sample collection**

109 Fresh broiler feces were collected randomly along two main diagonals of the
110 broiler compartments, from both medicated and control groups, at study days 4, 9, 14,
111 11, 22, 25, 29 and 33.

112 To investigate transmission of *E. coli* and enterococci between consecutive
113 broiler flocks, one day before chicks arrival (day -1), 12 cotton swabs of rearing
114 equipment and inner surfaces of the poultry house were obtained. Additionally, one
115 composite sample of pine shavings (used as bedding) was collected from the transport
116 vehicle and one sample of drinking water (1 000 ml) was obtained from the main water
117 reservoir. Cloacal swabs from 100 broiler chicks were collected at arrival (day zero),
118 before the chickens were placed inside the two shed compartments. Feed samples were
119 also obtained from the storage silo at the time of fecal sampling of broiler groups.

120

121 **2.3. Isolation and confirmation of enterococci and *E. coli***

122 Twenty five grams of each sample (poultry feces and feed) were suspended in
123 225 ml of Buffered Peptone Water (Oxoid, Basingstoke, United Kingdom) (BPW) and
124 directly plated onto Tergitol BCIG Agar (Biokar Diagnostics, Beauvais, France)
125 (Tergitol) and Kanamycin Aesculin Azide Agar Base (Oxoid) (KAA). To facilitate
126 isolation of *E. coli* and enterococci from feed and poultry house environment samples,

127 enrichment was performed in BPW (1:10 dilution) and incubated at 37 °C for 16 hours
128 before plating.

129 Tergitol and KAA agar plates were incubated at 37 °C for 24 hours and then,
130 after careful examination with a binocular dissecting microscope (magnification, × 20),
131 a maximum of 26 *E. coli* and 12 *Enterococcus* spp. characteristic colonies were selected
132 for susceptibility tests to twelve and ten antimicrobial agents, respectively. *E. coli*
133 confirmation and presumptive identification of the genus *Enterococcus* spp. were
134 performed as previously described (Martins da Costa et al., 2007).

135 One hundred milliliters of drinking water were, at each time, filtered through a
136 0.45 µm-pore-size membrane filters (Pall Corporation). The filters were then placed on
137 TBX agar (Biokar Diagnostics) and Slanetz & Bartley Medium (Oxoid) (SB) agar
138 plates and then incubated at 44 °C for 24 hours and 37 °C for 48 hours for
139 detection/enumeration of *E. coli* and *Enterococcus* spp., respectively.

140

141 **2.4. Antimicrobial susceptibility testing**

142 Antimicrobial susceptibility was determined using disk diffusion assay
143 following the guidelines of Clinical and Laboratory Standards Institute (CLSI 2007),
144 using the following antimicrobial drugs (Oxoid) for enterococci: ampicillin (AMP, 10
145 µg), chloramphenicol (CHL, 30 µg), ciprofloxacin (CIP, 5 µg), erythromycin (ERY, 30
146 µg), gentamicin (GEN, 120 µg), quinupristin-dalfopristin (Q-D, 15 µg), nitrofurantoin
147 (NIT, 300 µg), rifampicin (RIF, 5 µg), tetracycline (TET, 30 µg) and vancomycin
148 (VAN, 30 µg); and for *E. coli*: amoxicillin/clavulanic acid (AMC, 30 µg), ampicillin
149 (AMP, 10 µg), apramycin (APR, 15 µg), cephalothin (CEF, 30 µg), chloramphenicol
150 (CHL, 30 µg), enrofloxacin (ENR, 5 µg), gentamicin (GEN, 10 µg), kanamycin (KAN,

151 20 µg), nitrofurantoin (NIT, 300 µg), streptomycin (STR, 10 µg), tetracycline (TET, 30
152 µg) and trimethoprim/sulfamethoxazole (SXT, 25 µg).

153 Reference strains *E. coli* ATCC 25922 and *Enterococcus faecalis* ATCC 29212
154 were included as control. Interpretation of the diameter of the inhibition zone was made
155 according to the recommendations of CLSI and the manufacturer in the case of
156 apramycin (Lilly Farma, Algés, Portugal). Organisms considered intermediate by this
157 method were recorded as sensitive.

158

159 **2.5. Data analysis**

160 Statistical analysis was carried out using the SPSS version 15.0 for windows.
161 Differences in the prevalence of antimicrobial resistance between both groups were
162 assessed by the Chi-Square test. Alternatively, we used the Fisher's exact test. The null
163 hypothesis that antimicrobial resistance occurred equally in medicated and non-
164 medicated broilers was tested by the Mann-Whitney test. *P* values less than an alpha of
165 0.05 (Probability of Type I Error) were considered significant throughout this study.

166

167 **3. Results**

168 No infectious disease outbreak was observed in either group, avoiding non-
169 planned antibiotic treatments.

170 One feed sample (starter diet) and one environmental sample (swab collected
171 from nipple drinkers) was positive for *E. coli* isolation. All strains were resistant to
172 tetracycline, co-trimoxazol, streptomycin and kanamycin. Only one strain, isolated from
173 the rearing equipment, was resistant to ampicillin and enrofloxacin, whereas no
174 resistance was found against gentamicin.

175 No cultivable *E. coli* or enterococci were obtained from either drinking water or
176 pine shavings used as bedding, while enterococci were widely distributed in the broiler
177 chicken environment and also in the feed samples. These strains were predominantly
178 tetracycline-resistant. No resistance was found against gentamicin or ciprofloxacin, and
179 only two isolates, obtained from the farm environment, were resistant to ampicillin.

180

181 **3.1. *E. coli***

182 Twenty-six *E. coli* isolates were obtained from cloacal swabs taken from day-old
183 chick and none of these isolates displayed resistance to cephalothin, kanamycin,
184 apramycin, chloramphenicol and enrofloxacin. However, resistance to these
185 antimicrobial drugs were intermittently found in control broilers during the rearing
186 period. Within this group, no resistance against amoxicillin/clavulanic acid was found
187 and resistance rates to ampicillin, tetracycline, trimethoprim/sulfamethoxazole and
188 streptomycin revealed a slight decrease during the rearing period (Table 1).

189 For the medicated group, the prevalence of resistance was considerably
190 increased upon exposure to the three antimicrobials. Enrofloxacin use selected for *E.*
191 *coli* with resistance to structurally unrelated antimicrobials, such as the beta-lactams
192 (ampicillin and cephalothin), aminoglycosides (streptomycin and kanamycin),
193 trimethoprim/sulfamethoxazole and chloramphenicol. However, 11 days after the end of
194 enrofloxacin use, only enrofloxacin and tetracycline resistance frequencies were
195 significantly different between both groups. After medication with gentamicin,
196 enrofloxacin resistance frequency had increased again to 100 % and then dropped,
197 progressively, until the end of the rearing period. The more pronounced decrease in the
198 overall resistance five days after antimicrobial treatment was found for ampicillin, as a

199 result of the emergence of more sensitive strains having resistance phenotypes
200 equivalent to those found in non-medicated broilers.

201 Four days after the placement of day-old chicks, were found 14 new resistance
202 patterns in *E. coli* isolated from the control group (Table 2). This shift in resistance
203 profiles remained until the end of the study with the gradual isolation of new patterns
204 and, simultaneously, with the “return” of early resistance patterns obtained from cloacal
205 swabs collected from chicks and broiler feces.

206 Profiles of multi-resistance were more frequent and diverse in the medicated
207 group; among 208 *E. coli* strains isolated in this group, 47, 20, 37, and 9 displayed
208 simultaneous resistance to six (13 patterns), seven (seven patterns), eight (five patterns)
209 and nine (three patterns) antimicrobials, respectively, while in the control group only
210 two *E. coli* strains (two patterns) showed resistance to more than six antimicrobials. The
211 most frequent resistance pattern in control broilers was TET^R (n = 38), whereas in
212 broilers receiving intermittent antimicrobial pressure it was
213 AMP^RCEF^RTET^RSXT^RSTR^RKAN^RCHL^RENR^R (n = 25).

214

215 **3.2. Enterococci**

216 Comparing resistance rates in enterococci isolated from feces collected from
217 both groups, significant differences were only found for ciprofloxacin, gentamicin,
218 ampicillin and erythromycin ($P < 0.05$) (Table 3). However, after the first medication,
219 and throughout the study, no enterococcal isolates were susceptible to the full range of
220 antimicrobial drugs tested even though sensitive strains were isolated from the cloacal
221 swabs of the day-old chicks. As was the case with *E. coli*, multi-resistant enterococci
222 had only persisted in medicated broilers.

223 Correspondence between resistance patterns found in growing broilers and those
224 found in cloacal swabs from day-old chicks, feed diets and farm environment prior to
225 birds placement was, respectively, 26.0, 34.4 and 57.3 % for the medicated birds and
226 44.8, 72.9 and 82.3 % for the control group (Table 4).

227 None of the seven enterococci which displayed resistance to gentamicin was
228 isolated neither from control group nor from medicated broilers before the use of
229 gentamicin. While among ampicillin-resistant strains, there were three isolated from
230 control group and only five, out of 26 found in medicated broilers, were obtained after
231 the medication with amoxicillin. Additionally, it was found a greater incidence of
232 resistance to ampicillin after gentamicin medication rather than after amoxicillin use.
233 There were no strains exhibiting vancomycin resistance.

234

235 **3.3. *E. coli* and enterococci**

236 The overall resistance to all drugs tested reached the highest level after
237 medication with enrofloxacin for *E. coli* and gentamicin for enterococci. One day after
238 enrofloxacin, gentamicin and ampicillin had been used, 100, 58 and 100 % of *E. coli*
239 were resistant to these drugs, respectively, whereas 75, 42 and 42 % of the enterococci
240 were resistant to ciprofloxacin, gentamicin and ampicillin.

241 After the first medication, and throughout the study, no enterococci and only one
242 *E. coli* revealed susceptibility to all antimicrobial drugs tested, whereas simultaneous
243 resistances against fluoroquinolones, gentamicin and ampicillin were detected in two
244 and 43 isolates of *Enterococcus* spp. and *E. coli*, respectively.

245 Regarding enterococci, only ampicillin resistance was co-selected through
246 gentamicin medication between days 19 and 21, while in *E. coli* significant differences
247 in resistance incidence between groups had often occurred, without a direct selective

248 pressure imposed by the use of the same antimicrobial drugs or even by structurally
249 related ones.

250 At the end of the study, there was a reduction in the differences in resistance
251 profiles in the enterococcal isolates.

252

253 **4. Discussion**

254 Under field conditions, it is impossible to control the numerous factors that
255 interact in the development and persistence of resistance. Thus, this study was mainly
256 focused in visualizing the selective effects of the sequential medication of a broiler
257 flock with enrofloxacin, gentamicin and amoxicillin on the susceptibility profile of a
258 representative gram-positive (enterococci) and gram-negative enteric bacteria (*E. coli*).

259 Enterococci were widely distributed in the broiler chicken environment and also
260 in feed samples. The isolation of 11 different phenotypes of multi-resistant enterococci
261 from environmental swabs indicates that there was significant carry-over of resistant
262 enterococci from previous batches of broilers housed in this shed. Previous
263 investigations reported the long-lasting persistence of multi-resistant enterococci in
264 broiler house structure or on production equipment, as they are able to survive several
265 rounds of cleaning and disinfection (Chaslus-Dancla et al., 1987; Heuer et al., 2002).

266 Though of lesser magnitude, multi-resistant *E. coli* strains were also recovered
267 from the farm environment. The antimicrobial prescription protocol included:
268 enrofloxacin after day-old chicks placement, for prevention of early chick mortality and
269 to reduce possible colonization or horizontal spread of pathogens; gentamicin to
270 stabilize the gut flora promoting a greater uniformity of growth; and, finally,
271 amoxicillin to suppress *Clostridium perfringens* growth and to reduce microbial action
272 on dietary nutrients. In vivo, enrofloxacin is converted to a more potent intermediate

273 product, ciprofloxacin and both agents are completely cross-resistant (van den Bogaard
274 et al., 2001; Yan and Gilbert, 2004). In general, all antimicrobials administered were
275 extremely efficient in selecting for an increase in resistance and their use resulted in the
276 emergence of new resistance phenotypes. This shift in profile was more evident among
277 fecal *E. coli* and it continued beyond the period of antimicrobial exposure. This could
278 be explained by the well known capacity of *E. coli* to exchange resistance genes, as
279 earlier observed in commercial broilers (Bass et al., 1999; Smith et al., 2007), coupled
280 with changes in *E. coli* community structure, i.e. selective enrichment of previously non
281 detected phenotypes.

282 Since antimicrobials were used successively, it was difficult to elect the “more
283 selective” antimicrobial. However, following medication with enrofloxacin, there was a
284 marked change of the resistance patterns found in *E. coli* strains isolated from day-old
285 chicks by enrofloxacin-resistant strains. In the present study, only one enrofloxacin-
286 resistant *E. coli* was detected in the broiler house environment; however two previous
287 studies have described the propensity for some resistant strains to persist in the farm
288 environment and colonize new flocks (Diarra et al., 2007; Diarrassouba et al., 2007). As
289 fluoroquinolones had been used to treat two previous batches of chickens in the shed,
290 this could have contributed to the presence of enrofloxacin-resistant *E. coli* in all fecal
291 samples collected in both groups. However, it was clear from the results that exposure
292 to Gentamicin and amoxicillin were the driving forces in maintaining high frequency of
293 resistance to enrofloxacin during the second half of the rearing period of medicated
294 broilers. The strength of the co-selection effect was particularly evident with the *E coli*
295 isolates and it is probable that it contributed to the emergence and persistence of multi-
296 resistant strains, including resistance to structurally unrelated antibiotics.

297 It was found a greater incidence of resistance to ampicillin after gentamicin
298 medication, than following amoxicillin use. In addition, 37 of the 42 different
299 phenotypes which displayed resistance to ampicillin were isolated from medicated
300 broilers before the use of ampicillin. The majority of the resistance profiles included
301 either resistance to enrofloxacin or gentamicin to resistance to both antimicrobial drugs.
302 The extent of these co-selective effects is consistent with phenotypic or genotypic
303 evidence found in other studies (Berge et al., 2005; Harada et al., 2007).

304 Antimicrobial resistance in control birds not exposed to antimicrobials also
305 exhibited interesting changes over time, as early reported by others (Diarra et al., 2007;
306 Pleydell et al., 2007). Although *E. coli* strains sensitive to all antimicrobial, as well as
307 tetracycline-resistant ones, persisted only in these non-medicated broilers, multi-
308 resistant patterns were also found in the absence of selective antimicrobial pressure.

309 Similarly, as shown by the results obtained within the control group regarding to
310 enterococci, antimicrobials were not necessarily the cause of the increase in resistance
311 against ampicillin, tetracycline, erythromycin and nitrofurantoin. This shift in resistance
312 frequencies might be explained by colonization of the broilers' intestines with strains
313 found in the feed and in the broiler house environment, since their resistance patterns
314 were often equivalent. These data are in agreement with previously published studies,
315 which indicated that farm indigenous microbiota and feed-associated bacteria may have
316 a greater influence on the prevalence of antimicrobial resistance than antimicrobial
317 usage (McDermott et al., 2005; Smith et al., 2007).

318 While enterococci found in the feed were not expected to be as well adapted to
319 the broiler intestine as those present within the farm environment, their massive
320 numbers could compensate for this, considering the great amount of feed ingested
321 (almost 3.340 kg⁻¹ bird) during the broilers rearing period. In an earlier study, involving

322 22 broiler feed samples, it was observed that cultivable enterococci had ranged between
323 1.32 and 3.99 log CFU g⁻¹ (Martins da Costa et al., 2007).

324 Among enterococci, the overall resistance to all drugs tested reached the highest
325 level after medication with gentamicin. This was unexpected, as enterococci are
326 intrinsically resistant to aminoglycosides at a low level (8-128 µg ml⁻¹) (Klare et al.,
327 2003). However, gentamicin is a relatively stable drug (Yan and Gilbert, 2004). Thus,
328 considering that gentamicin given orally is poorly absorbed from the gastrointestinal
329 tract and that intestine content represents less than one tenth of the live broiler weight,
330 fecal gentamicin concentrations high enough to favor the selective enrichment of
331 gentamicin-resistant enterococci may have occurred. Furthermore, salinomycin and
332 narasin or any other stress factor within the intestinal environment, could damage the
333 enterococcal cell wall, increasing gentamicin penetration into enterococci and thus
334 contributing to a significant elimination of susceptible strains.

335 In contrast to the emergence of gentamicin-resistant enterococci, the appearance
336 of ampicillin-resistant strains had no chronological relationship with the use of
337 amoxicillin. This was most likely due to co-selection by the antimicrobials used earlier,
338 as 43 % and 29 % of these patterns displayed resistance against ciprofloxacin and
339 gentamicin, respectively.

340 Cessation of antimicrobial use did not appear to result in an immediate reduction
341 in antimicrobial resistance. The short growing cycle of modern broilers might not
342 provide sufficient time for resistance determinants to be lost once antimicrobial
343 treatment ceases. Furthermore, we must be aware of the potential risk of spreading these
344 multi-resistant strains to poultry farmers or even beyond farm boundaries, exposing
345 humans via food chain and/or environmental pathways.

346 The understanding of the association between antimicrobial use in food
347 producing animals and the development and persistence of resistant commensal bacteria
348 may help us to manage critical decisions concerning use of antimicrobials in these
349 animals.

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Table 1: Evolution and comparison of the percentage of antibiotic resistance in 26 isolates of *Escherichia coli* obtained from feces samples collected from medicated (M) and non-medicated (C) broilers on days 4, 9, 14, 18, 22, 25, 29 and 33. Medicated birds were treated with Enrofloxacin from day 1 to day 3, Gentamicin from day 19 to day 21, and Ampicillin from day 26 to day 28.

| Antibiotics ^a | Days | | | | | | | | | | | | | | | | | | | | | | | |
|--------------------------|------|-----|--------|----|-----|--------|----|----|--------|----|----|--------|----|-----|--------|----|-----|--------|----|-----|--------|----|----|--------|
| | 4 | | | 9 | | | 14 | | | 18 | | | 22 | | | 25 | | | 29 | | | 33 | | |
| | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P |
| AMP | 58 | 96 | 0.001 | 35 | 92 | <0.001 | 46 | 65 | 0.163 | 46 | 58 | 0.405 | 46 | 61 | 0.266 | 38 | 85 | 0.001 | 38 | 100 | <0.001 | 31 | 81 | <0.001 |
| AMC | 0 | 4 | 1.000 | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 11 | 0.235 | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - |
| CEF | 4 | 61 | <0.001 | 0 | 42 | <0.001 | 11 | 35 | 0.048 | 11 | 8 | 1.000 | 19 | 23 | 0.734 | 15 | 27 | 0.308 | 11 | 65 | <0.001 | 11 | 42 | 0.012 |
| TET | 96 | 100 | 1.000 | 69 | 96 | 0.024 | 77 | 81 | 0.734 | 54 | 85 | 0.016 | 69 | 96 | 0.024 | 69 | 100 | 0.004 | 73 | 100 | 0.010 | 65 | 81 | 0.211 |
| SXT | 38 | 85 | 0.001 | 46 | 81 | 0.010 | 31 | 61 | 0.026 | 31 | 46 | 0.254 | 27 | 77 | <0.001 | 35 | 58 | 0.095 | 46 | 61 | 0.266 | 27 | 42 | 0.244 |
| STR | 46 | 88 | 0.001 | 42 | 88 | <0.001 | 42 | 50 | 0.578 | 35 | 46 | 0.397 | 35 | 58 | 0.095 | 23 | 69 | 0.001 | 15 | 92 | <0.001 | 15 | 35 | 0.109 |
| KAN | 11 | 50 | 0.003 | 19 | 54 | 0.010 | 15 | 27 | 0.308 | 0 | 15 | 0.110 | 8 | 8 | 1.000 | 0 | 23 | 0.023 | 8 | 73 | <0.001 | 11 | 19 | 0.703 |
| GEN | 4 | 11 | 0.610 | 0 | 11 | 0.235 | 4 | 8 | 1.000 | 0 | 0 | - | 0 | 58 | <0.001 | 0 | 50 | <0.001 | 4 | 35 | 0.005 | 0 | 23 | 0.023 |
| APR | 0 | 0 | - | 0 | 8 | 0.490 | 0 | 4 | 1.000 | 0 | 0 | - | 0 | 58 | <0.001 | 0 | 15 | 0.110 | 4 | 15 | 0.350 | 4 | 11 | 0.610 |
| CHL | 15 | 73 | <0.001 | 8 | 27 | 0.140 | 4 | 27 | 0.050 | 8 | 27 | 0.140 | 0 | 15 | 0.110 | 0 | 19 | 0.051 | 4 | 31 | 0.024 | 0 | 27 | 0.010 |
| NIT | 0 | 11 | 0.235 | 0 | 8 | 0.490 | 0 | 11 | 0.235 | 0 | 8 | 0.490 | 0 | 0 | - | 0 | 19 | 0.051 | 4 | 19 | 0.191 | 0 | 4 | 1.000 |
| ENR | 50 | 100 | <0.001 | 19 | 100 | <0.001 | 15 | 88 | <0.001 | 11 | 65 | <0.001 | 8 | 100 | <0.001 | 19 | 81 | <0.001 | 15 | 65 | <0.001 | 8 | 54 | <0.001 |

^aAbbreviations: AMP, ampicillin; AMC, amoxicillin/clavulanic acid; CEF, cephalothin; TET, tetracycline; SXT, trimethoprim/sulfamethoxazole; STR, streptomycin; KAN, kanamycin; GEN, gentamicin; APR, apramycin; CHL, chloramphenicol; NIT, nitrofurantoin; ENR, enrofloxacin.

Table 2: Number of antimicrobial resistance patterns in *Escherichia coli* isolates from day-old chick meconium and broiler feces of medicated and non-medicated control broilers.

| Antimicrobial ^a resistance pattern | Day-old chicks | Control group | | | | | | | | Medicated group | | | | | | | |
|---|----------------|---------------|---|----|----|----|----|----|----|-----------------|---|----|----|----|----|----|----|
| | | Day | | | | | | | | Day | | | | | | | |
| | | 4 | 9 | 14 | 18 | 22 | 25 | 29 | 33 | 4 | 9 | 14 | 18 | 22 | 25 | 29 | 33 |
| AMP TET SXT NIT | 7 | | | | | | | | | | | | | | | | |
| AMP TET STR GEN | 6 | | | | | | | 1 | | | | | | | | | |
| AMP TET SXT STR | 5 | | | 1 | | | | | | | | | | | | | |
| none | 4 | 1 | 6 | 3 | 7 | 4 | 4 | 3 | 4 | | | | 1 | | | | |
| TET | 3 | 6 | 1 | 4 | 1 | 4 | 9 | 3 | 10 | | | 1 | 1 | | | 1 | |
| AMP TET SXT | 1 | | | 2 | | | 2 | 1 | 2 | | | | | | | | |
| TET SXT STR | | 3 | 5 | | 3 | 1 | 1 | | | | | | | | | | |
| AMP TET SXT STR ENR | | 3 | 1 | | 1 | 1 | 4 | | 1 | 2 | 4 | 1 | 2 | | 1 | | |
| AMP TET STR KAN | | 2 | 4 | 4 | | 2 | | 1 | 2 | | | | 1 | | 1 | 6 | |
| TET ENR | | 1 | 1 | | | | | | | | | 1 | 3 | | | | |
| AMP TET | | 1 | 2 | 2 | | 1 | | | | | | | | | 1 | | |
| AMP TET ENR | | 1 | | 1 | 1 | | | 1 | | | 1 | | 2 | | 2 | 1 | |
| AMP TET STR ENR | | 1 | | | | | | | | | 1 | 1 | | | | | |
| AMP TET STR CHL ENR | | 1 | | | 1 | | | | | | | 2 | 4 | | 1 | | |
| AMP TET STR KAN ENR | | 1 | | | | | | | | | | | | | | | |
| AMP TET SXT ENR | | 1 | 1 | | | | | 1 | | 1 | | 3 | | | | | |
| AMP TET SXT CHL | | 1 | 2 | | | | | | | | | | | | | | |
| AMP TET SXT GEN CHL ENR | | 1 | | | | | | | | 2 | | 1 | | 2 | 2 | 3 | |
| AMP TET SXT STR CHL ENR | | 1 | | | | | | | | 2 | 1 | | 2 | | | | |
| AMP CEF TET ENR | | 1 | | | | | | | | | | | 1 | | | 3 | |
| TET SXT | | | 2 | 2 | 3 | 4 | 1 | 7 | | | | | 2 | | | 1 | |
| TET SXT STR KAN | | | 1 | | | | | | | | | | | | | | |
| TET SXT STR ENR | | | | 3 | | 1 | 1 | | | | | | | | | | |
| AMP CEF | | | | 2 | | 4 | 2 | 1 | 1 | | | | | | | 5 | |
| AMP TET SXT STR CHL | | | | 1 | 1 | | | | | | | | 1 | | | | |
| AMP CEF TET STR | | | | 1 | 3 | 1 | | | | | | | | | | | |
| AMP | | | | | 5 | | 2 | 1 | | | | | | | | | |
| AMP TET STR | | | | | | 3 | | | | | | | | | | | |
| AMP CEF TET | | | | | | | | 2 | 1 | | | | | | | | |
| SXT | | | | | | | | 1 | 3 | | | 2 | 2 | | | | |
| TET STR KAN | | | | | | | | 1 | | | | | | | | | |
| TET SXT APR CHL ENR | | | | | | | | 1 | | | | | | 1 | | 1 | |
| AMP SXT STR NIT ENR | | | | | | | | 1 | | | | | | | | | |
| TET SXT APR ENR | | | | | | | | | 1 | | | 1 | | 6 | | 1 | |
| AMP STR KAN | | | | | | | | | 1 | | | | | | | | |
| AMP CEF TET SXT STR KAN CHL ENR | | | | | | | | | | 9 | 6 | 2 | | | | 7 | |
| AMP CEF TET SXT STR KAN NIT ENR | | | | | | | | | | 2 | 1 | | 1 | | 1 | 3 | |
| TET STR ENR | | | | | | | | | | 1 | | | 1 | | | | |

| | | | | | | | | | | |
|-------------------------------------|--|--|--|--|--|--|---|---|---|---|
| TET STR CHL ENR | | | | | | | 1 | | | |
| AMP TET SXT STR KAN CHL ENR | | | | | | | 1 | | | |
| AMP CEF TET STR CHL ENR | | | | | | | 1 | | | |
| AMP CEF TET STR GEN ENR | | | | | | | 1 | | | 1 |
| AMP CEF TET SXT STR CHL ENR | | | | | | | 1 | | | |
| AMP CEF TET SXT STR KAN CHL NIT ENR | | | | | | | 1 | | | |
| AMP AMC CEF TET SXT STR CHL ENR | | | | | | | 1 | | | |
| AMP TET SXT STR KAN ENR | | | | | | | | | | |
| AMP TET SXT STR GEN APR ENR | | | | | | | 5 | 2 | 1 | |
| AMP CEF TET SXT STR KAN ENR | | | | | | | 2 | | | |
| TET NIT ENR | | | | | | | 2 | 2 | | |
| TET STR NIT ENR | | | | | | | 2 | 2 | 1 | |
| AMP CEF TET SXT ENR | | | | | | | | 1 | | |
| AMP CEF TET SXT STR KAN GEN NIT ENR | | | | | | | | 1 | | |
| TET SXT ENR | | | | | | | | 1 | | |
| TET SXT GEN APR CHL ENR | | | | | | | | 1 | | |
| AMP TET STR GEN APR ENR | | | | | | | | 1 | | |
| AMP CEF TET SXT STR GEN ENR | | | | | | | | 1 | | |
| AMP AMC SXT STR ENR | | | | | | | | 1 | | 1 |
| AMP AMC TET SXT STR KAN ENR | | | | | | | | 1 | | |
| AMP AMC CEF TET SXT STR KAN APR ENR | | | | | | | | 1 | | |
| AMP TET SXT STR GEN APR | | | | | | | | | | |
| TET STR GEN | | | | | | | | | 2 | |
| AMP TET SXT KAN ENR | | | | | | | | | 1 | 1 |
| AMP TET SXT STR GEN ENR | | | | | | | | | 1 | |
| AMP TET SXT STR KAN GEN CHL ENR | | | | | | | | | 1 | |
| AMP CEF TET SXT GEN CHL ENR | | | | | | | | | 1 | 1 |
| AMP CEF TET SXT KAN ENR | | | | | | | | | 1 | |
| AMP CEF TET SXT STR ENR | | | | | | | | | 1 | |
| AMP CEF TET STR GEN APR | | | | | | | | | 1 | |
| AMP TET STR GEN APR | | | | | | | | | | 2 |
| AMP TET SXT STR KAN GEN ENR | | | | | | | | | | 1 |
| AMP CEF TET SXT STR GEN APR ENR | | | | | | | | | | 1 |
| AMP CEF TET SXT STR | | | | | | | | | | 1 |

^a For antimicrobial abbreviations, see Table 1, footnote a.

Table 3: Evolution and comparison of the percentage of antibiotic resistance in 12 isolates of *Enterococcus* spp. obtained from feces samples collected from medicated (M) and non-medicated (C) broilers on days 4, 9, 14, 18, 22, 25, 29 and 33. Medicated birds were treated with Enrofloxacin from day 1 to day 3, Gentamicin from day 19 to day 21, and Ampicillin from day 26 to day 28.

| Antimicrobials ^a | Days | | | | | | | | | | | | | | | | | | | | | | | |
|-----------------------------|------|----|-------|----|----|-------|----|----|-------|-----|----|-------|-----|-----|-------|----|----|-------|----|-----|-------|----|-----|-------|
| | 4 | | | 9 | | | 14 | | | 18 | | | 22 | | | 25 | | | 29 | | | 33 | | |
| | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | C | M | P | | | |
| AMP | 0 | 0 | - | 17 | 0 | 0.478 | 8 | 17 | 1.000 | 0 | 17 | 0.478 | 0 | 58 | 0.005 | 0 | 25 | 0.217 | 0 | 42 | 0.037 | 0 | 17 | 0.478 |
| VAN | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - |
| Q-D | 0 | 25 | 0.217 | 8 | 8 | 1.000 | 0 | 0 | - | 17 | 8 | 1.000 | 0 | 0 | - | 0 | 17 | 0.478 | 0 | 0 | - | 17 | 0 | 0.478 |
| TET | 75 | 67 | 1.000 | 75 | 92 | 0.590 | 75 | 75 | 1.000 | 100 | 92 | 1.000 | 100 | 75 | 0.217 | 92 | 92 | 1.000 | 92 | 100 | 1.000 | 83 | 100 | 0.478 |
| RIF | 42 | 42 | 1.000 | 0 | 25 | 0.217 | 25 | 8 | 0.590 | 8 | 25 | 0.590 | 0 | 8 | 1.000 | 8 | 25 | 0.590 | 17 | 8 | 1.000 | 0 | 0 | - |
| ERY | 25 | 58 | 0.098 | 50 | 67 | 0.408 | 58 | 92 | 0.155 | 42 | 75 | 0.098 | 50 | 100 | 0.014 | 58 | 92 | 0.155 | 50 | 50 | 1.000 | 67 | 75 | 1.000 |
| GEN | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 42 | 0.037 | 0 | 8 | 1.000 | 0 | 0 | - | 0 | 8 | 1.000 |
| CHL | 0 | 8 | 1.000 | 8 | 0 | 1.000 | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 0 | - | 0 | 8 | 1.000 | 0 | 8 | 1.000 |
| NIT | 17 | 50 | 0.193 | 25 | 42 | 0.667 | 42 | 42 | 1.000 | 42 | 50 | 0.682 | 50 | 75 | 0.400 | 50 | 58 | 0.682 | 50 | 42 | 0.682 | 50 | 50 | 1.000 |
| CIP | 8 | 75 | 0.001 | 8 | 67 | 0.009 | 0 | 33 | 0.093 | 8 | 50 | 0.069 | 8 | 25 | 0.590 | 8 | 33 | 0.317 | 8 | 25 | 0.590 | 0 | 0 | - |

^aAbbreviations: AMP, ampicillin; VAN, vancomycin; Q-D, quinupristin-dalfopristin; TET, tetracycline; RIF, rifampicin; ERY, erythromycin; GEN, gentamicin; CHL, chloramphenicol; NIT, nitrofurantoin; CIP, ciprofloxacin.

Table 4: Number of antimicrobial resistance patterns in *Enterococcus* spp. isolated from the house environment (Farm, n=24), feed samples (n=61), day-old chicks meconium (n=12) and feces of non-medicated control broilers (n=96) and medicated broilers (n=96). Medicated birds were treated with Enrofloxacin from day 1 to day 3, Gentamicin from day 19 to day 21, and Ampicillin from day 26 to day 28.

| Resistance patterns | Farm | Feed | Day-old chicks | Control group | | | | | | | | Medicated group | | | | | | | | No. total |
|-------------------------|------|------|----------------|---------------|---|----|----|----|----|----|----|-----------------|---|----|----|----|----|----|----|-----------|
| | | | | day | | | | | | | | day | | | | | | | | |
| | | | | 4 | 9 | 14 | 18 | 22 | 25 | 29 | 33 | 4 | 9 | 14 | 18 | 22 | 25 | 29 | 33 | |
| TET NIT | 3 | 33 | | | | | 3 | 4 | 3 | 1 | 3 | | | 1 | | | | 1 | 52 | |
| TET | 7 | 8 | 2 | 4 | 4 | | 3 | 2 | | 2 | | | 2 | 1 | | | 4 | 1 | 43 | |
| TET ERY | 4 | 2 | 1 | 1 | 1 | 2 | 3 | 4 | 4 | | 2 | | 2 | 4 | 1 | | 1 | 5 | 39 | |
| TET ERY NIT | 2 | 5 | | 1 | 4 | | | 1 | 2 | 5 | 3 | | 1 | | 2 | | | 2 | 28 | |
| none | | 4 | 5 | | 2 | 2 | | | 1 | 1 | 1 | | | | | | | | 21 | |
| AMP TET ERY NIT | 2 | | | 2 | 1 | | | | | | | | 2 | 4 | 3 | 3 | 1 | 18 | | |
| ERY NIT CIP | 1 | | | 1 | 1 | | | | | | | 4 | 1 | 3 | 2 | 1 | | 14 | | |
| TET RIF | 1 | 1 | 1 | 3 | | 2 | | | 1 | 2 | | 1 | | | | | | 13 | | |
| TET ERY NIT CIP | | | | | | | | 1 | 1 | | | 2 | 3 | | 1 | | | 8 | | |
| TET RIF ERY | | 3 | 2 | | | | | | | | | | 1 | | | | | 8 | | |
| TET ERY CIP | | | | | | | | | | 1 | | | | 1 | 3 | 1 | 1 | 7 | | |
| Q-D TET ERY | 1 | | 1 | | 1 | | | | | | 2 | | | 1 | | | | 6 | | |
| TET RIF NIT | | 5 | | | | | | | | | | | | 1 | | | | 6 | | |
| TET RIF CIP | | | | | | | | | | | | 2 | 2 | | | | 1 | 5 | | |
| RIF | | | | 2 | | 1 | | | | | | | | | | | | 3 | | |
| ERY | 2 | | | | | | | | | | 1 | | | | | | | 3 | | |
| Q-D TET RIF ERY | | | | | | | 1 | | | | | | | | | 2 | | 3 | | |
| TET NIT CIP | | | | | | | 1 | | | | | | | | 1 | | | 2 | | |
| TET ERY GEN | | | | | | | | | | | | | 2 | | | | | 2 | | |
| Q-D TET RIF | | | | | | | | | | | | 2 | | | | | | 2 | | |
| AMP TET ERY CHL NIT | | | | | | | | | | | | | | | | 1 | 1 | 2 | | |
| AMP TET ERY GEN NIT | | | | | | | | | | | | | 2 | | | | | 2 | | |
| TET ERY CHL | | | | | 1 | | | | | | | | | | | | | 1 | | |
| Q-D TET | 1 | | | | | | | | | | | | | | | | | 1 | | |
| Q-D TET ERY NIT | | | | | | | 1 | | | | | | | | | | | 1 | | |
| RIF ERY | | | | | | | | | | | | | | 1 | | | | 1 | | |
| RIF ERY NIT CIP | | | | | | | | | | | | | | 1 | | | | 1 | | |
| TET GEN NIT | | | | | | | | | | | | | | | | | 1 | 1 | | |
| TET RIF ERY NIT CIP | | | | | | | | | | | | | 1 | | | | | 1 | | |
| TET RIF ERY GEN CIP | | | | | | | | | | | | | | 1 | | | | 1 | | |
| Q-D TET ERY CIP | | | | | | | | | | | | | 1 | | | | | 1 | | |
| Q-D TET ERY CHL CIP | | | | | | | | | | | | 1 | | | | | | 1 | | |
| AMP TET NIT CIP | | | | | | | | | | | | | | | | 1 | | 1 | | |
| AMP TET ERY GEN NIT CIP | | | | | | | | | | | | | | 1 | | | | 1 | | |
| AMP TET RIF ERY CIP | | | | | | | | | | | | | | 1 | | | | 1 | | |
| AMP Q-D TET ERY NIT CIP | | | | | | | | | | | | | | 1 | | | | 1 | | |

^a For antimicrobial abbreviations, see Table 3, footnote a.