

1    **Short-Form Paper**

2    **Chicken Meat as Reservoir of Colistin-Resistant *Escherichia coli* Carrying**  
3    ***mcr-1* Genes in South America**

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14    **Running title:** *mcr-1* and *bla*<sub>CTX-M</sub> genes in chicken meat

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### Abstract

The detection and rapid spread of colistin-resistant Enterobacteriaceae carrying the *mcr-1* gene has created an urgent need to strengthen surveillance. In this study, eight clonally unrelated colistin-resistant *Escherichia coli* isolates carrying *mcr-1*, and *bla*<sub>CTX-M</sub> or *bla*<sub>CMY-2</sub> genes, were isolated from commercial chicken meat in Brazil. Most *E. coli* carried IncX4 plasmids, previously identified in human and animal isolates. These results highlight a new reservoir of *mcr-1*-harboring *E. coli* in South America.

## Text

The detection and rapid spread of colistin-resistant *Escherichia coli* isolates carrying the *mcr-I* gene has created an urgent need to strengthen surveillance. Nowadays, *mcr-I*-harboring Enterobacteriaceae isolates have been identified in food-producing animals, foods, aquatic environments, and humans (1-12). Although, the *mcr-I* gene has spread rapidly in Asia, Europe, Africa, North America and South America, studies reporting its presence in Enterobacteriaceae from foods have been unfrequent. In this regard, *mcr-I*-positive *E. coli* have been described in meats or vegetables in Europe (4-7, 9-11), Asia (8) and North America (12), so far. In this study, we report for the first time the identification of colistin-resistant *E. coli* carrying the *mcr-I* gene in commercial chicken meat, in Latin America.

As part of a local investigation conducted for monitoring the presence of colistin-resistant bacteria, carrying *mcr-I*, in chicken meat sold in markets in São Paulo, southeastern Brazil, forty-one samples including breasts ( $n=20$ ), thighs ( $n=20$ ), and liver ( $n=1$ ) were collected from 12 markets, between August and October of 2016. Samples (25 g) were dispensed in sterile plastic bags (Whirl-Pak, Nasco, WI, USA) containing 225 ml of MacConkey broth and incubated at 37°C for 24 h. After incubation, an aliquot of 1 ml of MacConkey broth was serially diluted on buffered peptone water and inoculated onto MacConkey agar plates containing colistin (2 µg/ml) (Sigma-Aldrich, St. Louis, MO), being incubated at 37°C for 24 h (13). Next, antimicrobial susceptibility profiles and MIC values of polymyxin B and colistin were determined by disc diffusion (14) and microdilution method (15), respectively, and *mcr-I*, ESBL and pAmpC genes were screened by PCR and sequencing (1, 16).

Eight colistin-resistant *E. coli* isolated from chicken meat samples (19.5%) collected from markets located in the North, South, and West region of São Paulo city, were positive for *mcr-I*

69 and *bla*<sub>CTX-M</sub> or *bla*<sub>CMY-2</sub> genes (Table 1). These isolates were found to be genetically unrelated by  
70 pulsed field gel electrophoresis (PFGE) (17), as well as to other *mcr-I*-positive *E. coli* isolates,  
71 previously identified in food-producing animals (2) and human (3), in Brazil. However, plasmid  
72 characterization by PCR-based replicon typing (PBRT) (18) revealed the presence of IncX4-type  
73 plasmids in five *mcr-I*-positive *E. coli* isolates (Table 1), which has been globally reported (3).

74 Genomic DNA from five representative colistin-resistant *E. coli* isolates (CF1.2, CF 101,  
75 CF 121, CF132, CF351) was extracted to construct a Nextera XT DNA library, which was  
76 sequenced using the MiSeq v3 platform (Illumina, San Diego, CA), using paired-end reads  
77 (300bp). *De novo* assembly was performed using A5-Miseq pipeline and this assembly was  
78 optimized using Geneious v.R9 (Biomatters Ltd, New Zealand). Serotypes, MLST, plasmid  
79 replicons, antimicrobial resistance genes, and *E. coli* virulence genes were identified using  
80 multiple databases as SerotypeFinder 1.1, MLST 1.8, PlasmidFinder 1.3, ResFinder 2.1 and  
81 VirulenceFinder 1.5 respectively, available from the Center for Genomic Epidemiology  
82 (<http://genomicepidemiology.org/>).

83 Most *E. coli* isolates exhibited a MDR phenotype and, indeed, clinically important genes  
84 conferring resistance to aminoglycosides, quinolones, sulphonamide and tetracyclines were  
85 identified by whole genome sequencing (WGS) (Table 1). On the other hand, MLST analysis  
86 from sequence reads identified the sequence types ST132, ST48, ST4419, ST522 and ST10  
87 (Table 1). In particular, the ST10 has been widely identified from animal, food, human and  
88 environmental samples, and associated to the production of CTX-M-type ESBLs and, more  
89 recently, the MCR-1 enzyme, denoting a great versatility of this lineage for adaptation to  
90 different hosts (3, 19-23). Furthermore, presence of IncX4 plasmids carrying the *mcr-I* gene in *E.*  
91 *coli* isolates CF1.2, CF131 and CF132 was confirmed, as previously reported in human and

92 animal clinical samples collected in this region (3, 19). In this regard, contigs of 5545-kbp,  
93 containing the *mcr-1* gene were found to bear an IncX4 replicon signature, as determined by the  
94 PlasmidFinder database (<https://cge.cbs.dtu.dk/services/PlasmidFinder/>). Moreover these contigs  
95 were found to host a hit showing 100% identity to another IncX4 plasmid harbouring *mcr-1*  
96 (GenBank accession no. CP015977). Partial IncX4 plasmid sequences were deposited in the  
97 GenBank database under the accession numbers: KY550358 (pCF1-2), KY550357 (pCF131) and  
98 KY550359 (pCF132). CTX-M-type-encoding genes were not located in the same plasmid that  
99 carried the *mcr-1* gene. In fact, IncX4 plasmids have only been associated with the mobilization  
100 of *mcr-1* (24).

101 Colistin has been widely used in animal feed as a growth promoter in Brazilian livestock,  
102 mainly in pigs and poultry. From 2008, the Ministry of Agriculture, Livestock, and Supply  
103 (MAPA) has established appropriate levels for colistin use in broilers (2-10 g/tonne of feed),  
104 poultry (4-10 g/tonne of feed), pigs (20-40 g/tonne of feed) and cattle (5-40 g/tonne of feed).  
105 However, after the presence of colistin-resistant *E. coli* carrying the *mcr-1* gene was confirmed in  
106 human and animals (including livestock), following international recommendations of the World  
107 Health Organization (WHO), the use of colistin in animal feed was banned by the MAPA  
108 (Regulatory Instruction no. 45) in November 2016, available at <http://www.agricultura.gov.br/>.

109 In summary, these results highlight that commercial chicken meat can be an important  
110 reservoir of *mcr-1*-carrying *E. coli*, which is a cause for public health concern, since this could  
111 contribute to the acceleration of the spread of the *mcr-1* gene. In fact, in the agribusiness, Brazil  
112 is the third largest chicken meat producer country and the largest exporter of this product with a  
113 high domestic consumption (25). Finally, the occurrence of *E. coli* carrying the *mcr-1* gene in  
114 chicken meat could be favoured by the versatility of *E. coli*, regarding host adaptability, ubiquity,

115 and persistence along the food chain; whereas IncX4 plasmids might be key vectors responsible  
116 for the dissemination of this gene. So, surveillance of colistin-resistant *E. coli* carrying *mcr-I*  
117 gene in the food chain needs to be established as a priority, to prevent their spread.

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#### 119 **CONFLICT OF INTEREST STATEMENT**

120 The authors have no conflict of interest to declare.

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237

238 **TABLE 1** Characteristics of colistin-resistant *E. coli* isolates, carrying the *mcr-1* gene, isolated from commercial chicken meat  
 239 samples.  
 240

<i>E. coli</i> strain	Date (m/y)	Sample meat	Region <sup>a</sup>	Resistance profile <sup>b</sup>	MIC (μg/ml) <sup>c</sup>			Additional resistance genes	Plasmid Inc group <sup>f</sup>	Virulence genotype	Phylogroup	PFGE profile <sup>g</sup> (MLST, ST)
CF 1.2	08/2016	Breast	W	CTF, CRO, CTX	8	4	CTX-M-2	<i>aadA1, aadB, sul1</i>	IncFIB, IncFIC, <u>IncX4</u>	<i>iss, iroN, lpfA, mchB, mchC, mchF, ireA,</i>	D	A (48)
CF 101	08/2016	Breast	W	CTF, CRO, CTX, FOX, SXT, TET, GEN	8	4	CTX-M-2	<i>aadA1, aac(3)-VIa, sul1, sul2, tetB</i>	IncHI2A, IncQ1	<i>gad, ireA, iss</i>	A	B (10)
CF 111	08/2016	Breast	S	AMC, CRO, CTX, FOX	4	2	- <sup>d</sup>	ND <sup>e</sup>	<u>IncX4</u>	ND <sup>e</sup>	A	C
CF 121	09/2016	Breast	S	AMC, CTF, CRO, CTX, FOX	2	2	CMY-2	<i>aadA2</i>	IncFII, IncN, IncR	-	A	D (522)
CF 131	09/2016	Breast	S	AMC, CTF, CRO, CTX, GEN	4	4	CTX-M-8	ND <sup>e</sup>	<u>IncX4</u>	ND <sup>e</sup>	B1	E
CF 132	09/2016	Breast	S	AMC, CTF, CRO, CTX, TET, GEN	4	4	CTX-M-8	<i>aadA1, aac(3)-VIa, sul1, sul2</i>	IncI1, IncX1, <u>IncX4</u>	<i>gad</i>	A	F (4419)
CF 341	10/2016	Breast	N	CTF, CRO, CTX, TET, GEN	4	2	CTX-M-2	ND <sup>e</sup>	<u>IncX4</u>	ND <sup>e</sup>	B2	G
CF 351	10/2016	Breast	N	CTF, CRO, CTX, TET, GEN	8	4	CTX-M-2	<i>aadA1, aadA5, aac(3)-VIa, aph(3')-Ic, sul1, tetA</i>	IncI1	-	B2	H (132)

241 <sup>a</sup> W, West; S, South; N, North.

242 <sup>b</sup> AMC, amoxicillin-clavulanic acid; CTF, ceftiofur; CRO, ceftriaxone; CTX, cefotaxime; FOX, cefoxitin; SXT, trimethoprim-sulfamethoxazole; TET, tetracycline; GEN, gentamicin.

243 <sup>c</sup> PMB, polymyxin B.

244 <sup>d</sup> ESBL phenotype was not confirmed by PCR.

245 <sup>e</sup> ND, not determined.

246 <sup>f</sup> The replicon type of plasmids carrying the *mcr-1* gene is underlined.

247 <sup>g</sup> PFGE patterns were analysed using the Dice similarity with coefficient optimisation set at 1% and tolerance at 2% (BioNumerics software; Applied Maths, Kortrijk, Belgium).