Whole-Genome Sequences of Two *Campylobacter coli* Isolates from the Antimicrobial Resistance Monitoring Program in Colombia

Johan F. Bernaletal., Pilar Donado-Godoy, María Fernanda Valencia, Maribel León, Yolanda Gómez, Fernando Rodríguez, Richa Agarwala, David Landsman, Leonardo Mariño-Ramírez

Corporación Colombiana de Investigación Agropecuaria (CORPOICA), Centro de Investigación Tibaíta, Cundinamarca, Colombia; Instituto Colombiano Agropecuario (ICA), Laboratorio Nacional de Diagnostico Veterinario, Bogotá, Colombia; National Center for Biotechnology Information, National Library of Medicine, National Institutes of Health, Bethesda, Maryland, USA

*Campylobacter coli,* along with *Campylobacter jejuni,* is a major agent of gastroenteritis and acute enterocolitis in humans. We report the whole-genome sequences of two multidrug-resistance *C. coli* strains, isolated from the Colombian poultry chain. The isolates contain a variety of antimicrobial resistance genes for aminoglycosides, lincosamides, fluoroquinolones, and tetracycline.

**TABLE 1** *Campylobacter coli* genome annotation statistics

| Strain | NCBI BioSample | No. of genes | No. of CDSes | No. of pseudogenes | No. of CRISPR arrays | No. of rRNAs | No. of tRNAs | No. of ncRNAs | GenBank accession no. |
|--------|----------------|--------------|--------------|-------------------|---------------------|--------------|---------------|---------------|----------------------|
| M1483  | SAMN04353893    | 1,782        | 1,739        | 55                | 0                   | 3            | 37            | 3             | LQXL00000000          |
| M1486  | SAMN04353891    | 1,916        | 1,873        | 58                | 1                   | 3            | 37            | 3             | LQXK00000000          |

a CDSs, coding sequences.

b ncRNAs, noncoding RNAs.
quences, pseudogenes, CRISPR arrays, rRNAs, tRNAs, and non-coding RNAs are summarized in Table 1.

A search for resistance-associated genes present in the isolates was performed using ResFinder version 2.1 (15) and enriched using RAST version 2.0 (16), both with default parameters. We found antimicrobial resistance genes for aminoglycosides (Aph 3′-III), lincosamides (InuC), fluoroquinolones (gyrA and gyrB), and tetracyclines (EF-G and TetO). Additionally, we found efflux pump genes (CmeA, CmeB, TolC, MATE, MFS, MacA, MacB, RND, AcrB, and OM) and CmeABC operon genes, both associated with increased multidrug resistance.

**Nucleotide sequence accession numbers.** This whole-genome shotgun project has been deposited in DDBJ/EMBL/GenBank under the accession numbers listed in Table 1. The versions described in this paper are the second versions.

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