



FICHA TÉCNICA DE PROYECTOS

Nombre del proyecto:	Estudio de Prevalencia y Resistencia a los antibacterianos de <i>Campylobacter</i> spp., <i>Escherichia coli</i> BLEE y <i>Salmonella</i> spp. en muestras gastrointestinales de pollos faenados en camales industriales de la Provincia de Pichincha-Ecuador.
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Organismo Financiador	SENPLADES
Monto de Financiamiento:	100.000 USD (ejecutados 71.065,35 USD)
Fecha de Inicio:	01 de Julio de 2014
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Instituciones participantes:	• UNIETAR. Facultad de Medicina Veterinaria y Zootecnia. Universidad Central del Ecuador.
Códigos:	DGI: 43 Código Único del Proyecto (CUP): 9175.0000.0000.377771
Objetivos generales:	Determinar la prevalencia y la resistencia a los antibacterianos de <i>Campylobacter</i> spp., <i>Escherichia coli</i> BLEE y <i>Salmonella</i> spp. en pollos faenados en camales industriales en la Provincia de Pichincha.
Resumen Ejecutivo:	<p><i>Salmonella</i></p> <p>Caeca content from 388 at random selected broiler batches were collected in 6 slaughterhouses during 1 year and analyzed by the ISO 6579/Amd1 protocol for the isolation for <i>Salmonella</i>. Isolates were serotyped and genotypic variation was acceded by pulsed field gel electrophoresis. MIC values for sulfamethoxazole, gentamicin, ciprofloxacin, ampicillin, cefotaxime, ceftazidime, tetracycline, streptomycin, trimethoprim, chloramphenicol, colistin, florfenicol, kanamycin and nalidixic acid were obtained. Presence of bla_{CTX-M}, bla_{TEM}, bla_{SHV} and bla_{CMY}; and mcr-1 plasmid genes was investigated in resistant strains to cefotaxime and colistin respectively. Prevalence at batch level was 16.0%. The most common serotype was S. Infantis (83.9%) followed by S. Enteritidis (14.5%) and S. Corvallis (1.6%).</p> <p>The pulsed field gel electrophoresis analysis showed that S. Corvallis, S. Enteritidis and S. Infantis isolates belonged to 1, 2 and 12 genotypes respectively. S. Infantis isolates showed high resistance rates to 12 antibiotics ranging from 57.7% (kanamycin) up to 98.1% (nalidixic acid and sulfamethoxazole). All S. Enteritidis isolates showed resistance to colistin.</p> <p>High multiresistant patterns were found for all the serotypes. The bla_{CTX-M} gene was present in 33 S. Infantis isolates while mcr-1 was negative in 10 colistin resistant isolates.</p> <p><i>Escherichia coli</i></p>

	<p>Sampling was performed from June 2013 to July 2014 in 6 slaughterhouses that slaughter broilers from 115 farms totaling 384 flocks. Each sample of collected caeca was streaked onto TBX agar supplemented with cefotaxime (3 mg/l). In total, 176 isolates were analyzed for AR patterns by the disk diffusion method and for bla_{CTX-M}, bla_{TEM}, bla_{CMY}, bla_{SHV}, bla_{KPC}, and mcr-1 by PCR and sequencing. ESBL and AmpC <i>E. coli</i> were found in 362 flocks (94.3%) from 112 farms (97.4%). We found that 98.3% of the cefotaxime-resistant isolates were multi-resistant to antibiotics. Low resistance was observed for ertapenem and nitrofurantoin. The most prevalent ESBL genes were the ones belonging to the bla_{CTX-M} group (90.9%), specifically the bla_{CTX-M-65}, bla_{CTX-M-55} and bla_{CTX-M-3} alleles. Most of the AmpC strains presented the bla_{CMY-2} gene. Three isolates showed the mcr-1 gene.</p> <p>Campylobacter</p> <p>Caecal content from 379 randomly selected broiler batches originating from 115 farms were collected from 6 slaughterhouses located in the province of Pichincha during 1 year. Microbiological isolation was performed by direct plating on mCCDA agar. Identification of <i>Campylobacter</i> species was done by PCR. Minimum inhibitory concentration (MIC) values for gentamicin, ciprofloxacin, nalidixic acid, tetracycline, streptomycin, and erythromycin were obtained.</p> <p>Genetic variation was assessed by RFLP-<i>flaA</i> typing and Multilocus Sequence Typing (MLST) of selected isolates. Prevalence at batch level was 64.1%. Of the positive batches 68.7% were positive for <i>C. coli</i>, 18.9% for <i>C. jejuni</i>, and 12.4% for <i>C. coli</i> and <i>C. jejuni</i>. Resistance rates above 67% were shown for tetracycline, ciprofloxacin, and nalidixic acid. The resistance pattern tetracycline, ciprofloxin, and nalidixic acid was the dominant one in both <i>Campylobacter</i> species. RFLP-<i>flaA</i> typing analysis showed that <i>C. coli</i> and <i>C. jejuni</i> strains belonged to 38 and 26 profiles respectively. On the other hand MLST typing revealed that <i>C. coli</i> except one strain belonged to CC-828, while <i>C. jejuni</i> except 2 strains belonged to 12 assigned clonal complexes (CCs). Furthermore 4 new sequence types (STs) for both species were described, whereby 2 new STs for <i>C. coli</i> were based on new allele sequences.</p>
Productos:	<p>Artículos científicos</p> <ul style="list-style-type: none"> • Vinuela-Burgos C, Cevallos M, Ron-Garrido L, Bertrand S, De Zutter L (2016). Prevalence and Diversity of <i>Salmonella</i> Serotypes in Ecuadorian Broilers at Slaughter Age. PLoS ONE 11(7): e0159567. DOI:10.1371/journal.pone.0159567 • Vinuela-Burgos C. <i>Salmonella</i> and <i>Campylobacter</i> in broilers at slaughter age: a possible source for carcasses contamination in Ecuador. (2017) Ghent University Press. DOI: 10.13140/RG.2.2.20687.48803 • Christian Vinuela-Burgos, Magali Wautier, Delphine Martiny, Marco Cisneros, Inge Van Damme, Lieven De Zutter; Prevalence, antimicrobial resistance and genetic diversity of <i>Campylobacter coli</i> and <i>Campylobacter jejuni</i> in Ecuadorian broilers at slaughter age. Poultry Science 2017 pew487. DOI: 10.3382/ps/pew487 • Christian Vinuela-Burgos, Lieven De Zutter; 2P4 - Important foodborne bacteria and their antimicrobial resistances in Ecuadorian poultry. Special Issue: Abstracts of the 10th European Congress on Tropical Medicine and International Health, 16-20 October 2017, Antwerp, Belgium October 2017 Volume 22, Issue Supplement S1, Page 144. DOI: 10.1111/tmi.12979

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 - Medina J. (2015). Cuantificación de *Campylobacter* spp. en un matadero semi-industrial de Aves y posterior identificación de especies (*C. jejuni* y *C. coli*) mediante técnicas de diagnóstico molecular.
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