Effects of encapsulated cinnamaldehyde and citral on the performance and cecal microbiota of broilers vaccinated or not vaccinated against coccidiosis

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ABSTRACT This study investigated the effects of encapsulated cinnamaldehyde (CIN) and citral (CIT) alone or in combination (CIN + CIT) on the growth performance and cecal microbiota of nonvaccinated broilers and broilers vaccinated against coccidiosis. Vaccinated (1,600) and nonvaccinated (1,600) 0-day-old male Cobb500 broilers were randomly allocated to 5 treatments: basal diet (control) and basal diet supplemented with bacitracin (BAC, 55 ppm), CIN (100 ppm), CIT (100 ppm), and CIN (100 ppm) + CIT (100 ppm). In general, body weight (BW) and feed conversion ratio were significantly improved in birds treated with BAC, CIN, CIT, and CIN + CIT (P < 0.05) but were all decreased in vaccinated birds compared with nonvaccinated birds (P < 0.05). Significant interactions (P < 0.05) between vaccination and treatments for average daily gain during the periods of starter (day 0–9) and BW on day 10 were noted. Broilers receiving vaccines (P < 0.01) or feed supplemented with BAC, CIN, CIT, or CIN + CIT (P < 0.01) showed reductions in mortality rate from day 0 to 28. The incidences of minor coccidiosis were higher (P < 0.05) in vaccinated birds than in nonvaccinated birds. Diet supplementation with BAC or tested encapsulated essential oils showed comparable effects on the coccidiosis incidences. Similar to BAC, CIN and its combination with CIT reduced both incidence and severity of necrotic enteritis (P < 0.05). No treatment effects were observed on the cecal microbiota at the phyla level. At the genus level, significant differences between vaccination and treatment groups were observed for 5 (Lactobacillus, Ruminococcus, Faecalibacterium, Enterococcus, and Clostridium) of 40 detected genera (P < 0.05). The genus Lactobacillus was more abundant in broilers fed with CIT, while Clostridium and Enterococcus were less abundant in broilers fed with CIN, CIT, or CIN + CIT in both the vaccinated and nonvaccinated groups. Results from this study suggested that CIN alone or in combination with CIT in feed could improve chicken growth performance to the level comparable with BAC and alter cecal microbiota composition.

Key words: broiler, cecal microbiota, coccidiosis, essential oil, growth performance

INTRODUCTION

Intestinal diseases such as coccidiosis and necrotic enteritis (NE) are responsible for immense economic losses (several billion USD annually) to the poultry industry worldwide (Dalloul and Lillehoj, 2006). The common Eimeria species involved in coccidiosis include Eimeria tenella, Eimeria maxima, and Eimeria acervulina. These species can multiply and damage the intestine epithelial layer of the chicken duodenum, mid-intestine, and ceca, reducing feed intake and nutrient digestibility (Long and Jeffers, 1986; Martin et al., 1997; Dahlia et al., 2006). The NE caused by Clostridium perfringens is an enteric disease characterized by severe necrosis of intestinal mucosa, which impairs broiler productivity (McDevitt et al., 2006). Necrotic enteritis B-like toxin produced by C. perfringens is considered the major toxin responsible for NE (Keyburn et al., 2008), although
many other toxins such as Beta 2, Tpel, and virulent factors also play a role (McDonald, 1980; Keyburn et al., 2008; Lepp et al., 2013; Prescott, 2016). Traditionally, coccidiosis and NE have been effectively controlled by the application of antimicrobials in feed (Reid, 1990). However, owing to the misuse and overuse of antimicrobials, resistance to antibiotics has become a public health issue, leading to new challenges to urgently develop effective alternatives to antibiotics to control enteric diseases in broilers (Casewell et al., 2003; Agyare et al., 2018). Recently, the Chicken Farmers of Canada revised its antimicrobial use to eliminate the preventive use of category II antibiotics in 2018 and that of category III antibiotics by the end of 2020 (Chicken Farmers of Canada, 2019).

To overcome the potential increase in mortality and morbidity of broilers due to the ban of in-feed antimicrobial use, probiotics, prebiotics, organic acid, essential oils (EOs), and vaccines are becoming studied as the alternatives to antibiotics (Griggs and Jacob, 2005; Diarra et al., 2007; Osman and Elhariri, 2013). In the 1970s, coccidiosis vaccines containing mixtures of living Eimeria species were introduced to control coccidiosis by stimulating intestinal immune T-cells of broilers (Lillehoj and Lillehoj, 2000; Wallach, 2010). The EOs are aromatic compounds derived from plants, and many are potent inhibitors of bacterial growth (Si et al., 2006; Bakkali et al., 2008). Natural EOs are extracted from parts of trees such as leaves, roots, and barks. Synthetic EOs may contain toxins although the purity could reach more than 95%. Although no studies have been conducted to compare the effects of natural and synthetic EOs on the performance of poultry, cinnamaldehyde (CIN) used in the present study was synthetic while citral (CIT) was a natural EO.

CIT and cinnamon oil are 2 EOs that have been used in medication over the last century (Burt, 2004). CIT compounds displaying a pale-yellow color are extracted from the bark of cinnamon trees or from other species of the genus Cinnamomum (Tisserand and Balacs, 1995). Cinnamon oil has antimicrobial effects but has additionally been used for food flavoring in sweets and chewing gum (Nabavi et al., 2015). It has been demonstrated that cinnamon powder in feed could improve meat quality and growth quality of broiler chickens (Sang-Oh et al., 2013) and alleviate intestinal injury (Wang et al., 2015). CIT (3,7-dimethyl-2,6-octadienal) is extracted from different plants including lemongrass (around 76% by gas chromatography) (Silva et al., 2008), lemon myrtle (>90%) (Tisserand and Balacs, 1995), and Lindera citriodora (about 65%) (Ohtsuru et al., 1967). The antimicrobial activities of CIT have been demonstrated as effective against several bacterial pathogens including C. perfringens (Onawummi, 1989; Si et al., 2006, 2009; Yang et al., 2016). Previous studies have demonstrated that protection is required for effective delivery of EOs to the animal gut (Zhang et al., 2014, 2015; Ma et al., 2016; Yang et al., 2015a; Omonijo et al., 2017). The use of CIT and cinnamon in feed as antibiotic alternatives could be limited because of physical and chemical instabilities during storage and in the gastrointestinal tract of poultry (Kimura et al., 1981; Tian et al., 2016). A recent study indicated that the incorporation of a soy protein-polysaccharide Maillard reaction product stabilized CIT and offered protection to CIT during the storage, upon low pH in the stimulated gastrointestinal tract fluid and heat treatment (Yang et al., 2015b). The protection could be due to the incorporation of soy protein-polysaccharide Maillard reaction product that may have shield peptide bonds against proteolysis and thus retard the release of CIT from the droplets (Yang et al., 2016).

In this study, the effectiveness of encapsulated CIT and cinnamaldehyde alone or in combination in feed on growth performance, gut health, and cecal microbiota was evaluated in broilers vaccinated or not against coccidiosis.

**MATERIALS AND METHODS**

**Essential Oils**

CIT (a mixture of cis and trans isomers, 95% purity) and CIN (≥95% purity) were purchased from Sigma-Aldrich Chemical Co. (St. Louis, MO). The CIT and CIN were encapsulated separately using the materials and methods described previously with minor modifications (Yang et al., 2015b; Ma et al., 2016).

**Experimental Design**

A total of 3,200 0-day-old male Cobb 500 broiler chickens were housed in 40 floor pens (80 birds per pen). The pens were assigned to 2 groups, half of the birds (1,600) received a commercial coccidiosis vaccine via spraying at 0-day of age and the other half (1,600) did not receive the vaccine. Pens in each of these 2 groups were randomly allocated to 5 dietary treatments (4 pens per treatment) in a complete randomized design. The dietary treatments were 1) basal diet (enrich with animal by-products) serving as the control; 2) basal diet with 55 mg/kg BAC (positive control); 3) basal diet with 100 mg/kg encapsulated CIT; 4) basal diet with 100 mg/kg encapsulated CIN; and 5) basal diet with a combination of 100 mg/kg encapsulated CIT and 100 mg/kg CIT (CIN + CIT). Broilers were fed a starter diet from the age of 0 to 9 D, grower diet from age 10 to 19 D, and a finisher diet from age 20 to 28 D. The EOs were fed from day 10 to 28 (grower, finisher), and BAC was fed from day 0 to 28 (starter, grower, and finisher). The starter, grower, and finisher diets (Table 1) were formulated and pelleted with wheat and corn as the principal cereals and soybean meal, fish meal, and meat meal as protein sources according to the nutritional recommendation by Cobb 500 (2012). No coccidiostats were provided in the diets to prevent coccidiosis.
Table 1. Feed ingredients and nutrient composition for starter (days 0–9), grower (days 10–19), and finisher (days 20–28) of chicken (% as feed basis otherwise indicated).

| Ingredients         | Inclusion in basal diet |
|---------------------|-------------------------|
|                     | Starter | Grower | Finisher |
| Wheat               | 30.00   | 35.00  | 35.00    |
| Soybean meal        | 27.79   | 21.89  | 16.76    |
| Vegetable oil       | 3.74    | 3.54   | 4.41     |
| Corn                | 23.39   | 24.81  | 29.15    |
| Corn gluten meal    | 5.00    | 5.00   | 5.00     |
| Limestone           | 0.96    | 0.96   | 0.92     |
| Biof 1              | 0.31    | 0.11   | 0.00     |
| Mineral and vitamin Mix 2 | 0.25 | 0.25   | 0.25     |
| L-Lysine HCl        | 0.12    | 0.13   | 0.25     |
| Sodium Bicarbonate  | 0.10    | 0.10   | 0.10     |
| DL-Methionine       | 0.23    | 0.21   | 0.19     |
| Avizyme 15024       | 0.05    | 0.05   | 0.05     |
| Sodium Bicarbonate  | 0.14    | 0.34   | 0.31     |
| Phytase 1           | 0.01    | 0.01   | 0.01     |
| Choline             | 0.12    | 0.10   | 0.10     |
| Fish meal           | 4.50    | 4.50   | 4.50     |
| Meat meal           | 3.00    | 3.00   | 3.00     |
| Calculated Nutrients|         |        |          |
| Crude Protein       | 25.30   | 23.30  | 21.30    |
| Methionine          | 0.62    | 0.58   | 0.53     |
| Methionine & Cysteine| 1.08   | 1.01   | 0.94     |
| Lysine              | 1.37    | 1.23   | 1.19     |
| Metabolisable Energy, kcal/kg | 3,022 | 3,063  | 3,152    |
| Crude Fat           | 6.19    | 6.06   | 7.00     |
| Crude Fiber         | 2.31    | 2.24   | 2.11     |
| Calcium             | 0.90    | 0.85   | 0.80     |
| Total Phosphorus    | 0.66    | 0.60   | 0.55     |
| Available Phosphorus| 0.40   | 0.35   | 0.32     |
| Sodium              | 0.20    | 0.17   | 0.16     |

1Feed-grade monocalcium phosphate.
2Supplied per kilogram of diet: vitamin A, 9,000 IU; cholecalciferol, 5,000 IU; vitamin E, 30 IU; vitamin K, 0.5 mg; cobalamin, 0.007 mg; thiamine, 0.4 mg; riboflavin, 6 mg; folic acid, 1 mg; biotin, 0.15 mg; niacin, 135 mg; pyridoxine, 4 mg; Fe, 125 mg; Mn, 60 mg; Cu, 5 mg; Se, 0.10 mg; I, 0.35 mg; Zn, 50 mg.
3Multi-Enzyme System for Wheat-Based Poultry Feed (Halchemix Canada Inc., Toronto, ON, Canada) containing 600 U/g of xylanase, 8,000 U/g of protease, and 800 U/g of amylase.
4Ronozyme P5000 (DSM Nutritional Products Canada Inc., Ayr, ON, Canada).

Birds were weighed at the start of the trial (day 0); body weight (BW) and feed intake were measured on day 10, 20, and 28 from each pen, and average daily feed intake (ADFI) and feed conversion ratio (FCR) were calculated. Birds were inspected at least twice per day, and mortalities or culls were removed and necropsied by the “Services Vétérinaires Ambulatoires Triple-V Inc.” (Acton Vale, QC, Canada). The mortality rate was calculated based on the average mortality in each pen on day 0 to 28.

General and Gut Health

Fecal samples were collected on days 6, 9, 13, 16, 20, 23, and 27 from each pen (3/pen; 60/vaccination group/sampling time for a total of 420 samples) for oocyst counts (log10/g). At day 21–22, 4 birds/pen (80 per vaccinated or nonvaccinated group for a total of 160 birds) were sacrificed for necropsy by the “Services Vétérinaires Ambulatoires Triple-V Inc.” The intestines of all sacrificed birds (4 birds per pen: 16/treatment) were examined for evidence coccidiosis and NE. The intestinal health was scored for NE and coccidiosis lesions according to the study by Collier et al., (2003). Intestines were longitudinally opened to score mucosa on a scale of 0 to 3 for NE lesions for each of the upper gut and lower gut (including ceca). Coccidiosis lesions were scored on a scale of 0 to 4 for each of E. maxima which induces bleeding in the middle of the small intestines, mucosa, E. tenella causing severe inflammation of ceca and E. acervulina causing white plaques in the duodenum. The body weights of killed birds were determined.

Genomic DNA Isolation and 16S Ribosomal RNA Gene Sequencing

Cecal contents of the aforementioned 160 sacrificed birds (4 birds/treatment for a total of 40 pooled contents) were used for genomic DNA extraction. Genomic DNA was extracted from frozen cecal content using the QIAamp DNA Stool Mini Kit (QIAGEN, Toronto, Canada) according to the manufacturer’s instructions. The purity and concentrations of the extracted DNA were determined using an Invitrogen Qubit 2.0 Fluorometer (Life Technologies Inc., Carlsbad, CA). Sequencing libraries of the 16S ribosomal RNA gene (rRNA) were prepared according to the Illumina 16S Metagenomic Sequencing Library Preparation Guide Rev. B and sequenced on a MiSeq instrument (Illumina). Briefly, a 444-bp fragment spanning the V3-V4 hypervariable region (Escherichia coli 16S rRNA position 340 - 784) was amplified with primers Bakt_341F (5'- CCTACGGGNGGCWGGCAG3'-3') and Bakt_805R (5'-GACTACHVGGGTATCTAATAC-3') (Klindworth et al., 2013) containing 5' Illumina overhang adapter sequences (5'-TGTCGGGCGGCAGCGATGTGATAGATACAG-3' and 5'-GTCTCGTGGGCTCGGA GATGTGTAGTATAAGACAG-3', respectively) using 2x KAPA HiFi HotStart ReadyMix (VWR, CA89125-042) and purified with AMPure XP beads (Beckman).
Coulter A63880). Unique 8-bp dual indexes were added by PCR using the Nextera XT Index Kit (Illumina Inc., FC-131-1002), and PCR products were cleaned up with AMPure XP beads. Samples were pooled together at equimolar concentrations and sequenced using a 600-cycle v3 reagent kit (Illumina, MS-102-3003).

The sequencing data were analyzed by Quantitative Insights Into Microbial Ecology (QIIME, version 1.9.1; Caporaso et al., 2010). Paired-end reads were joined with fastq-join (Aronesty, 2011), and quality filtered and demultiplexed in QIIME using default settings. The reads were clustered at 97% sequence identity (similar as the species level) using ucldst (Edgar, 2010), and operational taxonomy units (OTUs) were picked against the Greengenes database (gg_otorus_13_8) using an open-reference approach (DeSantis et al., 2006). Taxonomic assignment of the sequences was performed using the ucldst consensus taxonomy assigner. Taxa that could not be assigned were presented as ‘unclassified’ using the highest taxonomic level that could be assigned to them. The sequences were aligned against the Greengenes core set with PyNast (Caporaso et al., 2010), and a phylogenetic tree was constructed with FastTree (Price et al., 2009). Alpha-diversity (within groups) metrics were then calculated by QIIME, and a beta diversity (between groups) distance matrix based on unweighted UniFrac metric (Lozupone and Knight, 2005) was calculated, which was used for principal co-ordinate analysis (PCoA).

**Statistical Analysis**

The experiment was arranged as 2 × 5 factorial design with 2 groups (vaccinated or nonvaccinated) and 5 feeding treatments (Control, BAC, CIN, CIT, CIN + CIT). Effects of vaccination and feeding treatments on growth performance, oocysts counts (log10/g), gut lesions, and incidences and the relative microbial abundances and diversity were analyzed as a randomized complete block design (RCBD) by the ANOVAs using the MIXED procedure followed by Tukey’s multiple comparison test of SAS 9.4 (SAS Institute Inc., Cary NC). Pens (as replicates) were included as the block. Only taxa with >0.05% mean relative abundance in at least one treatment group were included in the analysis. Block, vaccination, treatments, and the interaction between vaccination and treatments were considered as fixed effects. The results were expressed as the least square means and standard error of the mean (SEM). A P value < 0.05 was used to declare significance.

**RESULTS**

**Growth Performances**

In general, vaccination significantly decreased (P < 0.05) BW on days 20 and 28, while BAC, CIN, CIT, and CIT + CIN significantly increased (P < 0.05) BW compared with the control on day 28. At day 10, only BAC increased (P < 0.05) the BW compared with the control. The interactions (P < 0.05) between vaccination and treatments (control, BAC, CIN, CIT, and CIN + CIT) on affecting BW were noted on day 10. For ADFI, no differences were observed between vaccinated and nonvaccinated broilers in the starter, grower, finisher, and the whole phase (day 0–28). Similarly, the broilers fed CIN, CIT, and CIN + CIT showed ADFI that is similar to the control and BAC. In this study, birds vaccinated against coccidiosis showed a decreased (P < 0.05) ADG in the whole phase. During the grower, a significant effect on ADG was noted only with BAC and CIN + CIT (P < 0.05). Vaccination was found to reduce (P < 0.05) FCR only in the starter, grower, and whole phase. The CIN, CIT, and CIN + CIT showed the effects similar to the BAC treatment group, whereby there was a significant reduction (P < 0.05) of FCR in the grower, finisher, and whole phase. Like BAC, tested encapsulated oils (either alone or in combination) significantly reduced mortality rates compared to the control (P < 0.01, Table 2).

**General and Intestinal Health**

No treatment effects were noted on the oocyst counts in fecal materials collected from 6 to 27 D (Figure 1). As expected, birds that received vaccine showed a significant highest oocyst counts compared with birds that did not receive this vaccine (P < 0.01). In nonvaccinated birds, the highest oocyst counts were observed on day 27, while increased oocyst counts were observed in vaccinated birds from days 6 to 27. On day 27, the nonvaccinated group showed the highest oocyst counts in the control and CIN + CIT-fed birds (P < 0.01).

**Coccidiosis**: In general, subclinical (minor low lesion scores) coccidiosis which was more prevalent in vaccinated than nonvaccinated birds, were observed (Figures 2A and 2B).

In the nonvaccinated group, coccidiosis lesions due to *E. acervulina* were observed in the control, BAC-, CIN-, CIT-, and CIN + CIT-treated birds with an incidence being 37.5%, 6.3%, 6.3%, 18.8%, and 56.3%, respectively. Birds with a lesion core of 3 (numerous coalescent lesions in the duodenum) were found only in the control. Data suggested that encapsulated CIN at 100 ppm in feed could provide the results similar to BAC at 55 ppm in controlling coccidiosis due to *E. acervulina* in nonvaccinated broiler chicken.

In vaccinated birds, *E. acervulina* was the most prevalent *Eimeria* ssp examined (regardless of treatments) with more than 80% of birds in each treatment group showing intestinal lesions due to this parasite. However, the average lesion scores due to *E. acervulina* were 1.3, 1.2, 1.0, 1.6, and 1.3 for the control, BAC-, CIN-, CIT- and CIN + CIT-treated groups, respectively. Low incidence and minor coccidiosis lesion scores by *E. maxima* (6.3 to 18.8% of birds with average lesion scores of 0.2 to 0.1) and *E. tenella* (6.3 to 12.5% of birds with average lesion scores of 0.1) were observed. No lesions due to *E.
Necrotic enteritis: Significant effects of the tested EOs were observed on reducing NE incidence and severity, which were comparable to the effects of bacitracin. The nonvaccinated control birds showed a 19% incidence of NE with an average lesion score being 0.2 of severity (the maximum severity was 3 found mainly in the control and CIT-fed birds). In general, the CIT-treated birds showed a similar level of NE incidence and severity as the control birds, while birds treated with CIN showed 0% NE incidence and severity. An incidence of 6.3% and an average severity score of 0.1 of NE were observed in the bacitracin-treated group. The CIN + CIT-treated birds resulted in a similar level of NE incidence and severity as the BAC-treated group.

Among the birds vaccinated against coccidiosis, the CIT-treated birds showed NE lesion scores. The CIT-treated birds resulted in 6% incidence and severity (the maximum severity was 3) found mainly in CIN-treated birds and no lesions due to E. tenella were observed in each of CIN- and CIT-treated birds.

Cecal Microbiota

Four bacterial phyla (Firmicutes, Proteobacteria, Tenericutes, and Actinobacteria) were detected by 16S rRNA gene sequencing analysis, with Firmicutes (90%) being the predominant phylum. Vaccination and treatments alone or their interaction did not affect the relative abundance of phyla (Table 3). At the genus level, a significant reduction in the relative abundance of Lactobacillus (P < 0.05) in the vaccinated group was observed. Ceca from broilers received feed supplemented with CIT, showed an increased relative abundance of Lactobacillus (P < 0.01) but a decreased Ruminococcus (P < 0.05) compared with the control and BAC-fed birds. In addition, feed supplemented with CIT, CIN, and CIN + CIT reduced relative abundances of Enterococcus (P < 0.05) and Clostridium (P < 0.01) in ceca of birds compared with control feed which is similar to BAC. In addition, Clostridium was reduced (P < 0.05) in the vaccinated group with interactions between vaccination and treatments being found. However, interactions between vaccination and treatments were observed for the relative abundances of Ruminococcus (P < 0.05), and Faecalibacterium (P < 0.01), but not for, Oscillospira and Dorea (Table 3).

The microbiota richness was estimated using observed OTUs, and the diversity was evaluated by Chao1, Shannon and Simpson indices. Vaccination did not affect microbiota richness and diversity. Feed supplemented with BAC, CIN, CIT alone showed lower (P < 0.05)
with 100 ppm encapsulated citral; CIN, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde; CIT, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde and 100 ppm citral. Data were affected by vaccination ($P < 0.05$) but it was unaffected by BAC, CIN, CIT, and CIN + CIT.

Simpson compared to the control and BAC supplemented feed (Table 4). The PCoA of the microbiota based on unweighted UniFrac phylogenetic distances followed with PERMANOVA showed that the majority of the samples from the nonvaccinated group (red circle) and vaccinated groups (green circle) clustered separately (Figure 3A); however, no significant differences between dietary treatments in the microbiota composition were observed (Figure 3B).

**DISCUSSION**

Since the restrictions on antibiotic use in feed-additives, the search for affordable alternatives has been a growing discipline (Manges et al., 2007). In this study, vaccine and encapsulated CIN and CIT were used alone and in combination as alternatives for preventing naturally incidences of coccidiosis and NE in broilers. Compared with previous studies (Aguilar et al., 2013; Khattak et al., 2014), the EOs in this research were fed at the beginning of the grower (day 10–28) instead of the starter. This is based on our previously reported experimental infection studies, in which EOs were applied from day 10 to day 28 and NE was reduced to the level similar to the treatment with antibiotics in the feed (Liu et al., 2016; Yang et al., 2016). In addition, we speculated that feeding EOs to broilers from the grower instead of starter could receive higher efficacy to promote the growth of broilers because the gut microbiota is not established and stable in the starter (Lu et al., 2003; Gong et al., 2008). As the cecum harbors more diverse and stable microbial communities than the ileum in broilers (Gong et al., 2002; 2007), we selected cecal digesta for analyzing microbiota in the present study.

The results suggested that vaccinations and EOs have interactions on increasing ADG in starter broilers and BW in grower broilers, respectively. This could be explained by the relationship between coccidiosis and NE, which reports that coccidiosis due to *Eimeria* could enhance mucus production and release plasma proteins by damaging gut epithelial cells, increasing nutrient availability for *C. perfringens* to grow (Williams et al., 2003). The results indicated that the combination of vaccines and EOs may improve the performance of broilers by controlling pathogen including *Eimeria* and *C. perfringens*. In the whole phase, the broilers supplemented with CIN, CIT, and CIT + CIN had similar effects as the BAC treatment, whereby BW was increased and FCR was decreased compared with the control. A previous study has shown that the higher nutrient digestibility associated with growth performance was due to increased secretions of endogenous digestive enzymes stimulated by EOs in the whole phase (Lee et al., 2003). Whether this is applied to the observation reported in the present study remains to be determined. Studies regarding the synergistic effects of EOs containing terpenes on growth performance have been conducted but no studies have examined synergism in an aldehyde and terpene blend, which were the compounds of CIN and CIT, respectively (Siani et al., 2013). In the present study, however, there were no significant synergistic effects of CIN and CIT on growth performance compared with CIN or CIT alone on growth performance. In addition, there were no observed differences among treatment groups regarding ADFI, suggesting that chickens may not sense the flavor of EOs because of their encapsulation. This finding is consistent with previously published studies reporting that feed intake was not significantly reduced by dietary inclusion of EOs (Brenes and Roura, 2010; Bozkurt et al., 2014). This study also indicated that CIN, CIT, and CIN + CIT could reduce mortality, similar to the BAC treatment. In contrast, the mortality in the control was 15%. It could be associated with the environment, management, and diet during the animal trial. For example, high dietary protein content in this study was 15%. It could be associated with the environment, management, and diet during the animal trial. For example, high dietary protein content in this study was 15%. It could be associated with the environment, management, and diet during the animal trial. For example, high dietary protein content in this study was 15%. It could be associated with the environment, management, and diet during the animal trial. For example, high dietary protein content in this study was 15%.
Figure 2. Effects of bacitracin and encapsulated cinnamaldehyde and citral in diets of broilers on (A) severity and (B) prevalence of coccidiosis lesions due to *E. acervulina*, *E. maxima*, *E. tenella*; necrotic enteritis lesions on (C) severity and (D) prevalence due to *C. perfringens* in both vaccinated (PV) or nonvaccinated (PNV) chickens at 21–22 D of age. CTRL, birds fed basal diet; BAC, birds fed basal diet with 55 ppm bacitracin; CIN, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde; CIT, birds fed basal diet with 100 ppm encapsulated citral; CIN + CIT, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde and 100 ppm citral. Data were not affected by the Vac X Trt ($P > 0.10$).
themselves against *Eimeria* infections (Williams, 2002). The lower FCR in the grower and whole phase indicated that vaccination could increase feed efficiency. The results were in accordance with a previous study indicating that the *Eimeria* challenge induced by vaccination could improve the growth performance of broilers by lowering the FCR (Lee et al., 2011). However, higher FCR in starter suggested that vaccination may reduce FCR in young birds (Yang et al., 2011).

The minor coccidiosis incidences and lesions due to *E. maxima* and *E. tenella* may be due to the high hygienic and biosecurity practice in this study (Diarra et al., 2007). However, coccidiosis due to *E. acervulina* and NE incidences increased after vaccination. The appearance of coccidiosis lesions could be explained that the immune response to the live coccidiosis vaccine could repeat reinfection through the ingestion of sporulated oocysts sprayed over the surface of feed (Reid, 1990). In addition, it has been reported that factors including *Eimeria* infection and dietary fish meal could be responsible for inducing NE lesions (Stanley et al., 2014; Wu et al., 2014). In this study, the increased NE incidences in vaccinated birds could be at least partially due to the appearance of *Eimeria* infection (coccidiosis lesions) and fish meal applied in the diet.

Many EOs including CIN and CIT have been studied in vitro for their potential to inhibit pathogens that cause diseases in chickens (Friedman et al., 2002, 2004;

### Table 3. Relative abundance of phyla and major genera (each representing >1.0% of total sequences on average) in ceca.

| Phyla/Genera     | Vaccination | Control | BAC | CIN | CIT | CIT + CIN | SEM | Effects |
|------------------|-------------|---------|-----|-----|-----|-----------|-----|---------|
| **Actinobacteria** | PV          | 0.02    | 0.01| 0.01| 0.03| 0.04      | 0.019| ns      |
|                  | PNV         | 0.02    | 0.03| 0.03| 0.03| 0.02      |      | ns      |
| **Firmicutes**   | PV          | 97.44   | 94.16| 96.71| 96.76| 94.88     | 0.485| ns      |
|                  | PNV         | 96.88   | 96.26| 92.62| 93.71| 95.13     |      | ns      |
| **Proteobacteria** | PV         | 1.44    | 2.89| 2.23| 2.28| 4.53      | 4.460| ns      |
|                  | PNV         | 2.14    | 2.84| 6.51| 5.94| 3.82      |      | ns      |
| **Tenericutes**  | PV          | 0.83    | 2.86| 0.81| 0.80| 0.39      | 1.145| ns      |
|                  | PNV         | 0.91    | 0.73| 0.79| 0.46| 0.91      |      | ns      |
| **unassigned**   | PV          | 0.26    | 0.08| 0.24| 0.13| 0.12      | 0.120| ns      |
|                  | PNV         | 0.06    | 0.14| 0.05| 0.06| 1.63      |      | ns      |
| **Lactobacillus**| PV          | 10.70   | 8.90| 3.76| 7.76| 17.99     | 3.09 | ns      |
|                  | PNV         | 5.82    | 5.30| 3.75| 11.11| 1.74     |      | ns      |
| **Ruminococcus** | PV          | 13.05   | 11.60| 11.10| 7.42| 12.33     | 2.230| ns      |
|                  | PNV         | 11.39   | 12.25| 11.33| 10.33| 7.53     |      | ns      |
| **Oscillospira** | PV          | 8.87    | 11.53| 11.02| 9.02| 12.80     | 3.114| ns      |
|                  | PNV         | 10.18   | 10.71| 9.08| 9.96| 8.06     |      | ns      |
| **Faecalibacterium** | PV       | 2.78    | 7.55| 4.61| 6.09| 6.46      | 2.940| ns      |
|                  | PNV         | 8.62    | 5.36| 8.11| 5.25| 3.18      |      | ns      |
| **Dorea**        | PV          | 1.26    | 1.27| 1.14| 0.69| 1.40      | 0.498| ns      |
|                  | PNV         | 1.35    | 1.42| 1.29| 1.56| 0.87      |      | ns      |
| **Enterococcus** | PV          | 1.21    | 0.04| 0.05| 0.13| 0.08      | 0.107| ns      |
|                  | PNV         | 1.04    | 0.08| 0.07| 0.05| 0.02      |      | ns      |
| **Clostridium**  | PV          | 1.00    | 0.17| 0.20| 0.15| 0.19      | 0.214| ns      |
|                  | PNV         | 1.29    | 0.37| 0.23| 0.34| 0.16      |      | ns      |

1PV, birds vaccinated with a live coccidiosis vaccine; PNV, birds not vaccinated with vaccine.
2BAC, 55 ppm bacitracin; CIN, 100 ppm encapsulated cinnamaldehyde; CIT, 100 ppm encapsulated citral; CIN + CIT, a combination of 100 ppm encapsulated cinnamaldehyde and citral.
3Vac, main effect of vaccination; Trt, main effects of treatments; Vac × Trt, interaction between vaccination and treatments.
4Asterisks indicate significant statistically differences (1 asterisk means a significance level of 0.05 and 2 asterisks 0.01).

### Table 4. Summary of alpha-diversity measurements of microbiota in ceca of vaccinated and nonvaccinated broilers treated with bacitracin, cinnamon, citral alone, or in combination.

| Vaccination | Control | BAC | CIN | CIT | CIT + CIN | SEM | Vac | Trt | Vac × Trt |
|-------------|---------|-----|-----|-----|-----------|-----|-----|-----|-----------|
| **Observed OTUs** | PV      | 393.38| 396.95| 409.50| 389.60| 383.88| 18.88| ns  | ns  | ns  |
|              | PNV     | 398.67| 384.78| 399.75| 391.20| 400.10|      |     |     |     |
| **Chao1**   | PV      | 431.91| 436.71| 444.70| 427.54| 419.26| 15.72| ns  | ns  | ns  |
|              | PNV     | 437.41| 421.36| 436.50| 428.08| 438.48|      |     |     |     |
| **Shannon** | PV      | 6.47  | 6.34  | 6.67  | 5.85  | 6.60  | 0.36 | ns  | ns  | ns  |
|              | PNV     | 6.54  | 6.42  | 6.42  | 6.22  | 6.61  |      |     |     |     |
| **Simpson** | PV      | 0.97  | 0.97  | 0.98  | 0.94  | 0.98  | 0.016| ns  | *  | ns  |
|              | PNV     | 0.98  | 0.97  | 0.97  | 0.96  | 0.98  |      |     |     |     |

1PV, birds vaccinated with a live coccidiosis vaccine; PNV, birds not vaccinated with vaccine.
2BAC, 55 ppm bacitracin; CIN, 100 ppm encapsulated cinnamaldehyde; CIT, 100 ppm encapsulated citral; CIN + CIT, a combination of 100 ppm encapsulated cinnamaldehyde and citral.
3Vac, main effect of vaccination; Trt, main effects of treatments; Vac × Trt, interaction between vaccination and treatments. Asterisks indicate significant statistically differences (1 asterisk means a significance level of 0.05).
The 3D principal coordinate analysis (PCoA) graph shows the variation among distance matrixes (unweighted UniFrac) of cecal microbiota in (A) vaccinated or nonvaccinated status and (B) treatments with bacitracin, encapsulated cinnamaldehyde and citral, alone or in the combination. Percentages shown are percentages of variation explained by the PC1 (14.64%), PC2 (10.03%), and PC3 (8.79%). PV, birds were vaccinated against coccidiosis; PNV, birds were not vaccinated against coccidiosis; Control, birds fed basal diet; BAC, birds fed basal diet with 55 ppm bacitracin; CIN, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde; CIT, birds fed basal diet with 100 ppm encapsulated citral; CIN + CIT, birds fed basal diet with 100 ppm encapsulated cinnamaldehyde and 100 ppm citral.

Si et al. 2009; Giteru et al., 2015). The reduction in coccidiosis incidences and lesions caused by E. acervulina after feeding CIN at 150 ppm has been demonstrated by a previous study (Orengo et al., 2012). However, in the present study, the coccidiosis incidence and lesions due to E. acervulina decreased in birds fed 100 ppm CIN. The results indicated that the protection of the EOs through encapsulation could result in lowering the concentration of CIN to decrease coccidiosis incidence and severity. Accordingly, the results in this study also indicated that both CIT and CIN supplementation possess the ability to reduce coccidiosis. As coccidiosis has been reported to promote NE (Williams et al., 2003), the decreased NE incidence in the present study could be due to the reduction of coccidiosis by CIN and CIN + CIT. Besides, CIN (aldehyde), and CIT (terpene) have been reported to act synergistically (Caldas et al., 2015). To fully understand the molecular mechanisms underlying the effects of CIN and CIT individually or in combination in promoting chicken gut health, more studies are required.

Oocyst counts in feces from broilers are the reflection of coccidial infection in the birds (Hodgson, 1970). The higher fecal oocyst counts in vaccinated birds was consistent with the results of coccidiosis lesions.

The cecal microbiota of broilers can reflect feed digestion and nutrient absorption (Rinttilä and Apajalahti, 2013), which is related to urine recycling and gut health (Karasawa, 1999). In the present study, 4 phyla of microbiota were detected, with Firmicutes (90%) being the predominant phylum, followed by Proteobacteria, Tenericutes, and Actinobacteria. The results were in
agreement with previous studies on broilers (Sakaridis et al., 2018; Biasato et al., 2019), although a higher relative abundance of phylum *Firmicutes* was observed. No *Bacteroidetes* were detected in the present study, which was also reported by others previously (Han et al., 2016; Pedroso et al., 2016; Lucke et al., 2018).

At the genus level, *Lactobacillus*, *Ruminococcus*, *Faecalibacterium*, *Enterococcus*, and *Clostridium* showed significant differences in relative abundance between the control and treatment groups in the present study. Previously *Lactobacillus* (*L. aviarus* in particular) has been reported to negatively correlate with the abundance of *C. perfringens* in broilers with NE disease (Feng et al., 2010). In addition, the reduction of *Lactobacillus johnsonii* by NE (Stanley et al., 2012) and independently by fish meal (Wu et al., 2014) has also been described. Stanley et al. (2012) observed a significant reduction of *Wissella confuse* in *C. perfringens* challenged birds of *Weissella confusa*, a heterofermentative lactic acid-producing bacterium that was previously classified as a member of *Lactobacillus* but later relocated to *Leuconostoc* (Collins et al., 1993; Bjorkroth et al., 2002). Antonissen et al. (2016) reported a shift in species composition of *Lactobacillus* and reduction in butyrate-producing strains that belong to *Ruminococcaceae* family in broilers treated with *Eimeria* and fish meal. A decrease of *Faecalibacterium* and *Oscillospira* in ceca were also detected in broilers with NE disease (Stanley et al., 2012; Lacey et al., 2018). In the present study, the CIT or vaccine treatment alone increased *Lactobacillus* abundance compared with control and bacitracin treatments, which could suggest a potential benefit to chicken gut health according to the reports above. In contrast, the CIT and vaccine treatments decreased *Ruminococcus* abundance compared with the control. *Ruminococcus* is responsible for the degradation of cellulolytic fibre in ruminants, but its role in broilers remains to be further clarified (Koike and Kobayashi, 2009; Mondot et al., 2016). *Faecalibacterium* is a group of bacteria able to produce butyrate that is a source of energy in broilers (Mondot et al., 2016; Bortoluzzi et al., 2017) and benefits animal gut health in general (Bedford and Gong, 2017). In the present study, neither vaccination nor EOs alone changed the abundance of *Faecalibacterium* but their combination did. The low relative abundance of *Enterococcus* may imply less potential for chicken infection, as the density of *Enterococcus* in feaces and digesta has been considered to be an indicator of fecal contamination that is associated with infections such as septicaemia in poultry (Gilmore, 2002; Boehm and Sassoubre, 2014). *Clostridium* contains some pathogenic, but largely nonpathogenic species (Num and Useh, 2017). Although it was proposed to be a factor for predicting the potential of infections (Udaondo et al., 2017), whether it can be well-established needs to be determined. The changes in microbiota composition beneficial to chicken gut health in response to EO treatment have also been suggested previously (Cooper et al., 2013; Rehman et al., 2018). To determine the cause-effect relationship, further studies are required.

In conclusion, encapsulated CIN alone or in combination with encapsulated CIT in feed altered cecal microbiota composition and improved the intestinal health and performance of broiler chickens similar to BAC. Further studies are required to determine if the cecal microbiota changes contribute to the improvement of intestinal health and performance.

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