

Reduction of *Salmonella enterica* Serovar Enteritidis Colonization in 20-Day-Old Broiler Chickens by the Plant-Derived Compounds *trans*-Cinnamaldehyde and Eugenol

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The efficacies of *trans*-cinnamaldehyde (TC) and eugenol (EG) for reducing *Salmonella enterica* serovar Enteritidis colonization in broiler chickens were investigated. In three experiments for each compound, 1-day-old chicks ($n = 75$ /experiment) were randomly assigned to five treatment groups ($n = 15$ /treatment group): negative control (-ve *S. Enteritidis*, -ve TC, or EG), compound control (-ve *S. Enteritidis*, +ve 0.75% [vol/wt] TC or 1% [vol/wt] EG), positive control (+ve *S. Enteritidis*, -ve TC, or EG), low-dose treatment (+ve *S. Enteritidis*, +ve 0.5% TC, or 0.75% EG), and high-dose treatment (+ve *S. Enteritidis*, +ve 0.75% TC, or 1% EG). On day 0, birds were tested for the presence of any inherent *Salmonella* ($n = 5$ /experiment). On day 8, birds were inoculated with $\sim 8.0 \log_{10}$ CFU *S. Enteritidis*, and cecal colonization by *S. Enteritidis* was ascertained ($n = 10$ chicks/experiment) after 24 h (day 9). Six birds from each treatment group were euthanized on days 7 and 10 after inoculation, and cecal *S. Enteritidis* numbers were determined. TC at 0.5 or 0.75% and EG at 0.75 or 1% consistently reduced ($P < 0.05$) *S. Enteritidis* in the cecum ($\geq 3 \log_{10}$ CFU/g) after 10 days of infection in all experiments. Feed intake and body weight were not different for TC treatments ($P > 0.05$); however, EG supplementation led to significantly lower ($P < 0.05$) body weights. Follow-up *in vitro* experiments revealed that the subinhibitory concentrations (SICs, the concentrations that did not inhibit *Salmonella* growth) of TC and EG reduced the motility and invasive abilities of *S. Enteritidis* and downregulated expression of the motility genes *flhC* and *motA* and invasion genes *hilA*, *hilD*, and *invF*. The results suggest that supplementation with TC and EG through feed can reduce *S. Enteritidis* colonization in chickens.

Salmonella enterica serovar Enteritidis is one of the two most common bacterial agents that cause food-borne illness in the United States (4). Poultry and poultry products are epidemiologically attributed as the critical sources from which humans contract salmonellosis (26, 39). Chickens can harbor *S. Enteritidis* without showing any obvious clinical signs and disseminate the pathogen to the environment, raising significant public health concerns. Moreover, intestinal colonization of the pathogen in birds leads to carcass contamination during slaughter or contamination of eggs. Despite control measures adopted in reducing the pathogen by preharvest and postharvest approaches, *S. Enteritidis* is widespread in poultry, leading to the elevated incidence rates of human salmonellosis (4). Recently, the U.S. Centers for Disease Control and Prevention (CDC) reported that food-borne salmonellosis in the last decade has not decreased significantly, and efforts should be targeted for controlling *Salmonella* (5).

In chickens, *S. Enteritidis* predominantly colonizes in the cecum. Reducing *S. Enteritidis* in the intestinal tract of chickens could reduce contamination of poultry products (1). A number of approaches for reducing the colonization by the pathogen in poultry have been explored thus far, but with various degrees of success. These approaches include feeding chickens with competitive exclusion bacteria, bacteriophages, organic acids, oligosaccharides, antibiotics, and vaccines (9, 11, 17, 19, 22, 24, 27, 45, 46). The efficacy of natural antimicrobial compounds for killing pathogenic microorganisms has received recent attention due to the toxicity of synthetic chemicals and concern over emerging

antibiotic-resistant strains of bacteria (42). The antimicrobial activities of several plant-derived essential oils have been demonstrated, and a variety of active components in these oils has been identified. Among the various plant compounds, *trans*-cinnamaldehyde (TC), a major ingredient in cinnamon (*Cinnamomum zeylandicum*), and eugenol (EG), a component of clove oil (*Eugenia caryophyllis*), possess antibacterial properties against Gram-negative and Gram-positive bacteria (8). Both of these compounds are generally recognized as safe chemicals for use in foods (GRAS) by the U.S. FDA (TC, 21 CFR 182.60 [13]; EG, 21 CFR 582.60 [14]). Previously, we reported that TC was effective in killing *S. Enteritidis* in chicken drinking water (34). Additionally, we investigated the efficacy of various plant-derived compounds, including TC and EG, for reducing *S. Enteritidis* in chicken cecal contents *in vitro* (33). Among the molecules tested, TC and EG were most

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effective in significantly reducing the pathogen populations. Therefore, the present study was undertaken to determine the efficacies of TC and EG for reducing *S. Enteritidis* colonization in broiler chickens.

MATERIALS AND METHODS

All experiments were approved by the Institutional Animal Care and Use Committee (IACUC) at the University of Connecticut.

Experimental birds. Day-old commercial, straight-run broiler chicks (Ross × Ross) were procured from Burr Farm Inc., Hampton, CT. They were allocated to floor pens with access *ad libitum* to nonmedicated feed (Blue Seal Feeds Inc., Londonderry, NH), *Salmonella*-free water, and arrangements for age-appropriate temperatures and bedding (23) at the Isolation Facility, University of Connecticut.

Experimental design. Three separate experiments were conducted for each compound. In each experiment, 75-day-old chicks were randomly allocated to five treatment groups ($n = 15$). In the experiments with TC (99% purity; Sigma-Aldrich, St. Louis, MO), the treatment groups included a negative control (no *S. Enteritidis* challenge and no supplemental TC), a compound control (no *S. Enteritidis* challenge but 0.75% [vol/wt] supplemental TC), a positive control (*S. Enteritidis* challenge but no supplemental TC), a low-dose treatment (*S. Enteritidis* challenge and 0.5% supplemental TC), and a high-dose treatment (*S. Enteritidis* challenge and 0.75% supplemental TC). Likewise, in the experiments with EG (99% purity; Sigma-Aldrich), the treatments included a negative control (no *S. Enteritidis* challenge and no supplemental EG), a compound control (no *S. Enteritidis* challenge but with 1% [vol/wt] supplemental EG), a positive control (*S. Enteritidis* challenge and no supplemental EG), a low-dose treatment (*S. Enteritidis* challenge and 0.75% supplemental EG), and a high-dose treatment (*S. Enteritidis* challenge with 1% supplemental EG). On day 0, five birds per experimental group were randomly selected and sacrificed to confirm that the birds were initially devoid of any *Salmonella*. TC or EG was supplemented in the feed for 20 days, starting on day 0. The appropriate volume of each plant compound was measured using a graduated cylinder, added into feed, and mixed thoroughly to obtain the desired concentrations in the feed (0.5 and 0.75% for TC and 0.75 and 1% for EG). On day 8, birds in the positive control, low-dose, and high-dose treatment groups were challenged with *S. Enteritidis* ($8 \log_{10}$ CFU/bird) by crop gavage. After 24 h (day 9), two birds from each treatment group were sacrificed to determine pathogen colonization in the cecum ($n = 10$ /experiment). After 7 and 10 days of challenge, six birds per treatment group were sacrificed by CO₂ asphyxiation, and the cecum with its contents from each bird was collected in 5 ml of sterile phosphate-buffered saline (PBS) for bacteriological analysis.

Bacterial strains and dosing. A four-strain mixture of *S. Enteritidis* isolated from chickens (obtained from the Connecticut Veterinary Diagnostic Medical Laboratory, University of Connecticut) was used to colonize the birds, as described previously (32). The isolates were *S. Enteritidis* 12 (chicken liver, phage type 14b), *S. Enteritidis* 21 (chicken intestine, phage type 8), *S. Enteritidis* 28 (chicken ovary, phage type 13a), and *S. Enteritidis* 31 (chicken gut, phage type 13a). Each strain was preinduced for resistance to 50 µg/ml of nalidixic acid (NA; Sigma-Aldrich, St. Louis, MO) for selective enumeration. One hundred microliters of each nalidixic acid-resistant strain was grown separately in 10 ml tryptic soy broth (TSB; Difco) overnight, transferred into separate conical flasks containing 100 ml TSB with 50 µg/ml NA, and incubated overnight at 37°C with shaking (100 rpm). The cultures were combined and sedimented by centrifugation ($3,600 \times g$, 15 min, 4°C), and the pellet was resuspended in 100 ml of PBS (pH 7.0) and used as the inoculum ($\sim 10^8$ CFU/ml). The bacterial counts in the individual cultures and the four-strain mixture were confirmed by plating 0.1-ml portions of appropriate dilutions on xylose-lysine-desoxycholate agar (XLD; Difco) plates containing NA (XLD-NA) and incubating the plates at 37°C for 24 h.

Cecal *S. Enteritidis* determination. *S. Enteritidis* population numbers in ceca were determined as described previously (32). The ceca with

their contents from each bird were weighed and homogenized. Each homogenate was serially diluted (1:10) in PBS, and appropriate dilutions were plated on XLD-NA plates. The plates were incubated for 48 h at 37°C before counting colonies. Representative colonies from XLD-NA plates were confirmed as *Salmonella* by using the *Salmonella* rapid detection kit (Microgen Bioproducts Ltd., Camberley, United Kingdom). When colonies were not detected by direct plating, samples were tested for surviving cells by enrichment for 48 h at 37°C in 100 ml selenite-cysteine broth (SCB; Oxoid) (17, 18), followed by streaking on XLD-NA plates. Representative colonies from the plates were confirmed as *Salmonella* as previously mentioned.

Determination of cecal endogenous bacteria and cecal pH. Appropriate dilutions of the samples from ceca were plated on duplicate thio-glycolate agar plates (TGA; Difco) (15) and incubated at 40°C under 5% CO₂ for 24 h. The pH of cecal contents was also recorded for all treatment groups by using a pH meter and direct immersion of the electrode into the samples (33).

Body weight and feed consumption. The average feed consumption and body weights of birds were also determined for each experiment. Birds were weighed individually at the start and end of each experiment. The average feed consumption per bird was calculated by dividing the total amount of feed consumed per treatment group by the number of birds in the respective treatment group.

Determination of SICs of TC and EG. The effects of subinhibitory concentrations (SICs) of TC and EG against *S. Enteritidis* strains were determined as described previously (35). Duplicate 50-ml tubes containing 20 ml Luria-Bertani (LB; Difco) broth or cell culture medium (Dulbecco's modified Eagle medium [DMEM; Gibco, Invitrogen]) supplemented with 10% fetal calf serum (FCS; Gibco, Invitrogen) were separately inoculated with 4×10^6 to 5×10^6 CFU/ml of each *S. Enteritidis* strain. TC or EG was added with an increment of 1 mg/µl each from 0 to 10 mg/µl to corresponding tubes, and tubes were incubated at 37°C for 24 h. Samples were drawn from each tube, diluted in sterile PBS (pH 7.2), and plated on TSA plates at 0, 2, 4, 6, 8, 10, 12, and 24 h, and the plates were incubated at 37°C for 48 h. The experiment was repeated three times. The highest concentration of TC or EG that did not inhibit the bacterial growth after 24 h of incubation was taken as the SIC for that compound.

Motility assay. The effects of TC and EG on *S. Enteritidis* motility were determined according to the modified procedure described in reference 40. Ten microliters ($\sim 10^6$ CFU/ml) of a mid-log-phase *S. Enteritidis* culture grown in the presence or absence of the SIC of TC (0.01% [vol/vol]) or EG (0.04% [vol/vol]) was inoculated onto the centers of duplicate plates containing LB plus 0.3% agar. The plates were incubated at 37°C for 8 h. The zone of motility (distance the bacteria traversed [in cm] after incubation) was measured. In addition, tubes containing 10 ml of LB broth added with appropriate volumes of TC or EG and bacterial cultures (10^6 CFU/ml) were also incubated at 37°C for 8 h. The tubes were diluted and plated to determine the number of surviving bacteria.

Cell culture. (i) Intestinal epithelial cell line. Budgerigar abdominal tumor cells (BATCs), a permanent avian intestinal epithelial cell line, a kind gift of Margie Lee, College of Veterinary Medicine, University of Georgia, Athens, was used for the study. The cell line is a published model for studying *Salmonella* invasion and pathogenesis in avian species (16, 21, 25). This experiment was carried out to determine if TC and EG reduced *S. Enteritidis* invasion of avian intestinal epithelial cells. The cells were cultured in DMEM with FCS (16, 21, 25), and after three successful propagations were seeded into wells of 24-well tissue culture plates containing 1 ml DMEM with FCS at 1×10^5 cells/well. The cells were incubated at 37°C with 5% CO₂, to a confluence of >95% within 48 h. The viability of cells was confirmed using a trypan blue vital dye exclusion assay (41). Briefly, 50 µl of the diluted BATC suspension prior to each experiment was mixed with 10 µl of trypan blue dye, and 10 µl of the mixture was loaded in the counting chambers of a hemocytometer. After 1 min, the number of nonstained cells was counted under the low-power objective of the microscope.

TABLE 1 Primers used in the study

Gene	Forward primer	Reverse primer
<i>hilA</i>	5'-TTACTGTGCGCTGGCAGAAT-3'	5'-TCGCCTTAATCGCATGTTCTT-3'
<i>hilD</i>	5'-GGCGGTACCCACAGAAAAG-3'	5'-TCGTACAGGAGAACGCCGTT-3'
<i>invF</i>	5'-ACGCCATAGTCTTCTCCAGC-3'	5'-TCAGTCAACCAGCGGCAAC-3'
<i>flhC</i>	5'-TTGGCGCTCGTCTACAAATG-3'	5'-GACCACGGGTGAGCTGTGTT-3'
<i>motA</i>	5'-GATTTGCTGGCGTTGCTCTAT-3'	5'-CCCCTGCTGACGTGATTTG-3'
16S	5'-GTATGCGCCATTGTAGCACG-3'	5'-TCATCATGGCCCTTACGACC-3'

(ii) ***Salmonella* invasion assay.** *S. Enteritidis* strains were grown separately in DMEM containing FCS with the SIC of TC (0.01% [vol/vol]) or EG (0.04% EG [vol/vol]) or without compound (control) at 37°C until the mid-log phase. The cultures were sedimented by centrifugation (3,600 × g, 10 min, 4°C), and the pellet was resuspended in DMEM and used as the inoculum. The bacterial counts in the individual cultures were confirmed by plating 0.1-ml portions of appropriate dilutions on TSA plates. A multiplicity of infection (MOI) of ~50 was used for the study. The tissue culture plates were then centrifuged at 1,000 × g for 3 min at 23°C and incubated for 45 min at 37°C with 5% CO₂. Thereafter, the medium was removed from the wells and replaced with DMEM supplemented with 100 µg/ml of gentamicin (Gibco, Invitrogen). The samples were incubated for 1 h to kill all the extracellular bacteria. The wells were then washed with PBS three times, and the intestinal cells were lysed using 0.1% Triton X-100 (Invitrogen), followed by incubation at 37°C with 5% CO₂ for 15 min to release the internalized bacteria. The cell lysates were serially diluted, plated on TSA plates, and incubated at 37°C for 24 h before counting colonies. Duplicate samples of each treatment were included, and the experiment was repeated three times.

Effects of SICs of TC and EG on expression of *S. Enteritidis* motility and invasion genes. (i) **RNA isolation and cDNA synthesis.** Each strain of *S. Enteritidis* was grown in LB with and without the SIC of TC (0.01% [vol/vol]) or EG (0.04% [vol/vol]) to mid-log phase at 37°C. Three milliliters of bacterial culture was centrifuged at 12,000 × g for 2 min at 4°C, and the resultant pellet was incubated with 1 ml of RNeasy protect reagent (Qiagen, Valencia, CA) for 5 min at room temperature. Total RNA was extracted from the control and treated *S. Enteritidis* cells by using the RNeasy minikit (Qiagen) according to the manufacturer's instructions. Quantitation of RNA was done by measuring the absorbance at 260 and 280 nm (Nanodrop; Bio-Rad). cDNA was synthesized using the SuperScript II reverse transcriptase kit (Invitrogen, Carlsbad, CA).

(ii) **RT-qPCR.** The cDNA was used as the template for the amplification of *Salmonella* motility genes *flhC* and *motA* and invasion genes *hilA*, *hilD*, and *invF*. The specific primers for the aforementioned genes and for 16S rRNA (endogenous control) (Table 1) were designed using Primer Express software (Applied Biosystems) based on the *Salmonella enterica* serovar Enteritidis strain P125109 genome (NCBI reference sequence NC_011294.1). The primers were custom synthesized by Integrated DNA Technologies (Foster City, CA). Real-time quantitative PCR (RT-qPCR) was performed with the ABI Prism 7900 sequence detection system (Applied Biosystems) using the SYBR green assay (Applied Biosystems) under custom thermal cycling conditions. The biological replicates were analyzed in duplicate and normalized against 16S rRNA gene expression. The comparative threshold cycle (C_T) method ($2^{-\Delta\Delta C_T}$) was used to assess the relative changes in mRNA expression levels between the control and TC-treated or EG-treated *S. Enteritidis* cells (7).

Statistical analysis. The number of *S. Enteritidis* colonies in the ceum was logarithmically transformed (\log_{10} CFU/g) before analysis to achieve homogeneity of variance (10). These data were analyzed using the PROC-MIXED procedure of the SAS statistical analysis software (version 9.2; SAS Institute Inc., Cary, NC). Differences among the means were detected at a P level of ≤ 0.05 using Fisher's least significance difference (LSD) test. For motility, cell culture, and RT-qPCR assays, the results are provided as mean values and standard errors of the means (SEM). Differ-

ences between two independent treatments were analyzed using two-tailed t tests, and a P value of < 0.05 was considered statistically significant.

RESULTS

No *S. Enteritidis* was recovered from the cecal samples of negative or compound control birds (both TC and EG controls) during the entire duration of the study. After 24 h of inoculation, 1×10^7 to 3×10^7 CFU of *S. Enteritidis*/g of cecal contents was recovered from all the chicks necropsied from the inoculated treatment groups.

The results for the effects of TC and EG on *S. Enteritidis* counts in cecal samples are depicted in Fig. 1. The *S. Enteritidis* counts recovered from the cecal samples of the control birds ranged from 6 to 7 \log_{10} CFU/g after 10 days postinfection (p.i.), with the exception of experiment 2 with TC, where the recovery was ~4.5 \log_{10} CFU/g (Fig. 1). It was observed that both doses of TC resulted in similar reductions of *S. Enteritidis* populations in the cecal samples ($P < 0.05$). The lower dose (0.5% TC) reduced *S. Enteritidis* by 3.1, 3.7, and 3.6 \log_{10} CFU/g, whereas the higher dose (0.75% TC) resulted in 4, 3.3, and 4.3 \log_{10} CFU/g reductions in experiments 1, 2, and 3, respectively (Fig. 1). Similar results were observed with EG at both concentrations on *S. Enteritidis* in the ceum (Fig. 1). EG at 0.75 and 1% resulted in 3.3, 4.6, 3.0 and 2.8, 5.1, 2.6 \log_{10} CFU/g reductions of *S. Enteritidis*, respectively, in experiments 1, 2, and 3 (Fig. 1).

The cecal endogenous bacterial counts and pH did not differ ($P > 0.05$) between various treatment groups (Table 2). The supplementation with TC at 0.5 or 0.75% did not significantly alter ($P > 0.05$) the body weights of birds compared to the positive,

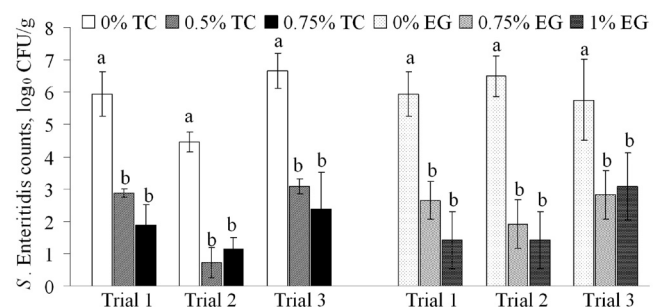


FIG 1 Effects of TC and EG on *S. Enteritidis* in ceum and cecal contents on the 10th day postinfection. Error bars represent SEM ($n = 6$ /treatment group/experiment). In each experiment, chicks (except for negative and TC controls) were challenged on day 8 posthatch with a mixture of four *S. Enteritidis* isolates at 8 \log_{10} CFU/bird. TC was added to the feed prophylactically from day 0 until the completion of the study (20 days). Columns with different letters (a or b) denote significant differences between the treatments ($P < 0.05$). Negative and compound controls were not included in the statistical analysis, since *S. Enteritidis* was not recovered from those treatment group birds.

TABLE 2 Effects of TC and EG on cecal pH, cecal endogenous bacteria, body weight, and cumulative feed consumption of 20-day-old chickens^a

Treatment	pH ^b	Cecal bacteria (log ₁₀ CFU/g) ^b	Avg body wt (g) ^c	Cumulative feed intake (g) ^c
Negative control	6.2 ± 0.1	8.7 ± 0.2	790.5 ± 19.3 ^A	960.0 ± 5.8 ^A
TC control	6.2 ± 0.1	8.6 ± 0.1	808.8 ± 26.3 ^A	952.9 ± 6.0 ^A
EG control	6.2 ± 0.1	8.8 ± 0.2	582.0 ± 39.7 ^B	760.3 ± 5.8 ^E
Positive control	6.1 ± 0.1	9.4 ± 0.2	786.4 ± 20.3 ^A	896.9 ± 9.1 ^B
0.5% (vol/wt) TC	6.3 ± 0.1	8.5 ± 0.3	799.9 ± 36.8 ^A	885.5 ± 6.2 ^B
0.75% (vol/wt) TC	6.0 ± 0.1	8.9 ± 0.2	792.1 ± 40.8 ^A	883.3 ± 6.0 ^B
0.75% (vol/wt) EG	6.3 ± 0.1	9.3 ± 0.1	536.4 ± 26.4 ^B	821.8 ± 5.9 ^C
1% (vol/wt) EG	6.3 ± 0.1	9.2 ± 0.1	530.6 ± 17.2 ^B	785.5 ± 6.2 ^D

^a Values are means ± SEM ($n = 18$ per treatment group).

^b Means within the column did not differ significantly ($P > 0.05$).

^c Values with different superscript capital letters (A to E) differed significantly from each other within a column ($P < 0.05$).

negative, or compound controls (Table 2). However, all of the EG-supplemented treatment groups had reduced body weights compared to other treatment groups ($P < 0.05$). The cumulative feed consumption levels of the birds supplemented with 0.75 or 1% EG were also lower than other inoculated treatment groups ($P < 0.05$) (Table 2). In addition, among the nonchallenged treatment groups, EG lowered cumulative feed consumption compared to the negative and TC controls ($P < 0.05$).

The SICs of TC and EG were 0.01 and 0.04%, respectively (Fig. 2). The average initial *S. Enteritidis* population in the control and TC- and EG-treated samples was approximately $5.0 \log_{10}$ CFU/ml. After incubation at 37°C for 24 h, approximately $8.0 \log_{10}$ CFU/ml of bacteria were recovered from control and treated samples, thereby confirming that the aforementioned concentrations of TC and EG were not inhibitory for *S. Enteritidis* (Fig. 2).

The effects of the SICs of TC and EG on motility, invasion, and gene expression on the four strains of *S. Enteritidis* are shown in Table 3. Both plant compounds at their SICs significantly reduced the zone of motility in all strains without reductions in bacterial counts after 8 h of incubation compared to the controls ($P < 0.05$) (Table 3). The SICs of TC and EG were also found to significantly decrease ($P < 0.05$) the invasion by *S. Enteritidis* for BATCs (Ta-

ble 3). *trans*-Cinnamaldehyde at 0.01% reduced invasion by 60 to 80%, whereas EG at 0.04% decreased *Salmonella* invasion by 75 to 85% ($P < 0.05$). Further, the results of the gene expression studies revealed that TC and EG at their corresponding SICs significantly downregulated expression levels of *Salmonella* motility and invasion genes (Table 3).

DISCUSSION

Based on our previous observations that TC and EG were effective in significantly reducing *S. Enteritidis* populations in chicken cecal contents *in vitro* (33), we investigated the prophylactic efficacies of these molecules as a preharvest treatment for reducing *S. Enteritidis* populations in chickens. The tested doses of TC (0.5 and 0.75%) and EG (0.75 and 1%) were selected based on our *in vitro* data (33). The chicken cecum represents the major colonization site for *S. Enteritidis* (30, 32, 38, 47), thereby allowing further spread of the infection to a healthy uninfected flock. Therefore, the aim of our study was to reduce *S. Enteritidis* populations in the ceca of chickens.

The results indicated that supplementation with TC and EG consistently decreased *S. Enteritidis* counts in birds. It was observed that supplementation of both plant molecules in feed significantly reduced the pathogen populations in the cecal samples (Fig. 1). It was also observed that the two concentrations of TC and EG did not significantly differ ($P > 0.05$) in their efficacies in reducing *Salmonella*, and both plant compounds were effective after day 10 p.i.

Although supplementation of TC did not significantly alter ($P > 0.05$) the body weights of chickens, all the EG-supplemented treatment groups of birds had significantly lower body weights than the negative, positive, and TC controls ($P < 0.05$) (Table 2). The reductions in the body weights in the EG-supplemented treatment groups could have been due to the decreased feed consumption ($P < 0.05$), as the birds reduced their intake compared to the respective controls (Table 2). However, among the various EG-supplemented treatment groups, the body weights of birds were not different from each other ($P > 0.05$). Previously, based on the observation that EG supplemented at 850 ppm (0.085%) impaired the absorption of alanine in the rat jejunum (36), Lee and colleagues postulated that EG may impair the normal digestion process (37). While these findings have not been reported in chickens, we believe that the aroma or flavor induced by EG could have reduced the likeability of the feed for the chickens, thereby leading to decreased body weights in EG-supplemented birds.

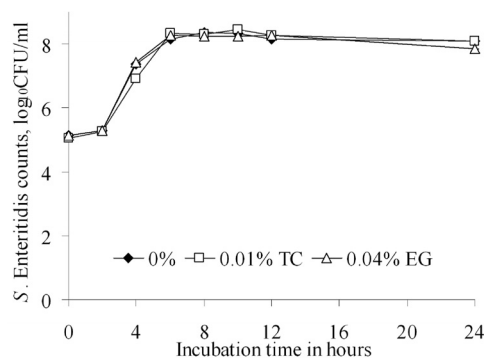


FIG 2 Effect of subinhibitory concentrations of TC and EG on *S. Enteritidis* growth ($n = 6$ /treatment group). The treatment groups did not differ significantly from the controls ($P > 0.05$). Tubes containing 20 ml LB broth were inoculated with $\sim 10^6$ CFU/ml of *S. Enteritidis*. TC or EG was added at increments of 1 mg/ μ l (from 0 to 10 mg/ μ l) and incubated at 37°C for 24 h. Samples drawn at various intervals of incubation were diluted and plated on TSA, before enumeration after incubation at 37°C for 24 h. The highest concentration of TC or EG that did not inhibit *S. Enteritidis* growth after 24 h was taken as the respective SIC.

TABLE 3 Effects of subinhibitory concentrations of TC and EG on motility, invasion, and virulence gene expression in *S. Enteritidis* strains used in the study^a

Strain and virulence gene	Motility ^{b,c} (cm)			% invasion ^c (CFU/ml)			Relative fold change in RT-qPCR assay ^d	
	0%	0.01% TC	0.04% EG	0%	0.01% TC	0.04% EG	0.01% TC	0.04% EG
<i>S. Enteritidis</i> 12	8.5 ± 0.0 (FL) ^c	7.15 ± 0.1 ^B	5.30 ± 0.0 ^A	100.00 ± 0.0 ^B (4.16 ± 0.2)	33.65 ± 17.8 [*] (1.34 ± 0.7)	17.31 ± 17.3 ^A (0.65 ± 0.7)	-1.34 ± 0.2 [*]	-1.90 ± 0.3 [*]
<i>hilA</i>							-1.35 ± 0.2 [*]	-1.98 ± 0.2 [*]
<i>hilD</i>							-0.87 ± 0.2 ^{NS}	-2.52 ± 1.0 ^{NS}
<i>invF</i>							-0.81 ± 0.0 ^{NS}	-1.25 ± 0.0 [*]
<i>flhC</i>							-0.94 ± 0.2 ^{NS}	-1.62 ± 0.1 [*]
<i>motA</i>								
<i>S. Enteritidis</i> 21	8.5 ± 0.0 (FL) ^c	6.20 ± 0.0 ^B	4.60 ± 0.1 ^A	100.00 ± 0.0 ^B (4.29 ± 0.2)	21.48 ± 21.5 [*] (0.89 ± 0.9)	14.30 ± 14.3 ^A (0.59 ± 0.6)	-1.72 ± 0.2 [*]	-1.06 ± 0.1 [*]
<i>hilA</i>							-1.94 ± 0.0 [*]	-1.32 ± 0.0 [*]
<i>hilD</i>							-2.14 ± 0.1 [*]	-1.42 ± 0.2 [*]
<i>invF</i>							-1.99 ± 0.3 [*]	-1.25 ± 0.0 [*]
<i>flhC</i>							-1.65 ± 0.1 [*]	-1.18 ± 0.0 [*]
<i>motA</i>								
<i>S. Enteritidis</i> 28	8.5 ± 0.0 (FL) ^c	4.20 ± 0.0 ^A	4.55 ± 0.1 ^B	100.00 ± 0.0 ^B (4.10 ± 0.2)	28.78 ± 18.7 [*] (1.12 ± 0.7)	27.81 ± 14.4 ^A (1.15 ± 0.6)	-1.28 ± 0.2 [*]	-2.10 ± 0.3 [*]
<i>hilA</i>							-1.62 ± 0.1 [*]	-2.14 ± 0.1 [*]
<i>hilD</i>							-1.52 ± 0.1 [*]	-2.32 ± 0.0 [*]
<i>invF</i>							-1.66 ± 0.1 [*]	-1.75 ± 0.1 [*]
<i>flhC</i>							-1.45 ± 0.1 [*]	-2.06 ± 0.0 [*]
<i>motA</i>								
<i>S. Enteritidis</i> 31	8.5 ± 0.0 (FL) ^B	3.70 ± 0.0 ^A	3.80 ± 0.0 ^A	100.00 ± 0.0 ^B (4.21 ± 0.2)	43.45 ± 11.1 ^A (1.79 ± 0.4)	27.68 ± 17.8 ^A (1.12 ± 0.7)	-2.42 ± 0.5 [*]	-4.46 ± 0.4 [*]
<i>hilA</i>							-2.36 ± 0.1 [*]	-4.57 ± 0.2 [*]
<i>hilD</i>							-2.50 ± 0.1 [*]	-5.52 ± 0.4 [*]
<i>invF</i>							-2.28 ± 0.1 [*]	-3.81 ± 0.0 [*]
<i>flhC</i>							-2.07 ± 0.2 [*]	-4.03 ± 0.2 [*]
<i>motA</i>								

^a Values are means ± SEM.^b FL, full lawn.^c Values with different superscript capital letters (A to C) differ significantly within the row for the indicated experiment ($P < 0.05$).^d The fold change relative to the control. *, TC or EG downregulated expression of the gene compared to the control ($P < 0.05$). NS, not significant ($P > 0.05$).

It was also observed that both TC and EG at the tested concentrations did not cause significant reductions in the total endogenous bacterial populations in the chicken cecum (Table 2). The exact mechanism behind the selective inhibitory effects of TC and EG on *Salmonella* with no apparent effects on the normal gut flora is not known. However, many previous investigators have reported that plant molecules such as cinnamaldehyde, eugenol, carvacrol, and thymol can cause significant reductions in *S. Typhimurium* DT104, *Escherichia coli* O157:H7 (28, 43), pathogenic *E. coli*, and *Clostridium perfringens* (28) without any adverse effects on endogenous bacterial populations, including lactobacilli and bifidobacteria (28, 43).

In order to determine if TC and EG produced any effects on major *S. Enteritidis* colonization factors in chickens, we investigated the effects of SICs of the plant compounds on *Salmonella* colonization factors, namely, motility and host cell invasion. The SIC of a molecule is the highest concentration below the MIC that does not inhibit bacterial growth (2, 35). However, since the SIC can modify bacterial physico-chemical functions, including that of genes, it is used for studying the effect of antimicrobials on bacterial gene expression and virulence (reviewed in reference 20). Motility plays a critical role in host-bacterium interactions, colonization, and virulence (29). Flagellum-mediated and non-flagellum-mediated motility have been reported to aid in the invasion and colonization by *S. Enteritidis* (3, 31). The gene *motA* is associated with the regulation of flagellar assembly (44), and *flhC*, a transcriptional activator for motility genes, is a global regulator of flagellin production (12). In addition, it was previously demonstrated that *S. Enteritidis* attaches and invades various host cells prior to establishing an infection process (16, 21, 47). It is also well-established that genes such as *hil* and *inv* play major roles in *Salmonella* invasion (6, 48, 49).

Our experiments revealed that both motility and invasion of avian intestinal epithelial cells were substantially inhibited by TC and EG ($P < 0.05$) (Table 3). Decreases in these attributes without any concurrent reduction in the bacterial counts led us to hypothesize that these molecules may modulate *Salmonella* genes responsible for motility and invasion. Therefore, we investigated the effects of TC and EG on the expression of motility and invasion genes, namely, *motA*, *flhC*, *hilA*, *hilD*, and *invF*. The results from RT-qPCR experiments revealed that the plant molecules downregulated the virulence genes compared to controls ($P < 0.05$) (Table 3).

To summarize, supplementation with 0.5 and 0.75% TC or 0.75 and 1% EG was effective in reducing *S. Enteritidis* populations in 20-day-old commercial broiler chickens. Although both molecules were effective in reducing *S. Enteritidis* populations in the cecum, TC is more suitable than EG for prophylactic dosing, considering the negative impact of EG on body weight and feed consumption for birds. The results of our *in vitro* studies indicated that TC and EG at their SICs significantly reduced *S. Enteritidis* motility and invasion of avian intestinal epithelial cells, and this could be due to the downregulation of the motility genes *motA* and *flhC* and invasion genes *hilA*, *hilD*, and *invF*. We conclude that TC could be used as an antimicrobial feed additive to reduce *S. Enteritidis* colonization in chickens along with standard hygienic practices used on farms.

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