

Effects of hydrolyzed gallotannin on intestinal physical barrier, immune function, and microbiota structure of yellow-feather broilers

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ABSTRACT The present study was performed to explore the effects of dietary supplementation of hydrolyzed gallotannin (**HGT**) on intestinal physical barrier, immune function and microbiota structure in yellow-feather broilers. A total of 288 male yellow-feather broilers were randomly allocated to 4 diet treatments: the basal diet (**CON**) and 3 diets supplemented with 150, 300, and 450 mg/kg HGT for 63 d, respectively, with 6 replicates per treatment and 12 birds per replicate. The findings demonstrated that 300 or 450 mg/kg HGT addition enhanced the expression of duodenal occludin (**OCN**) and tight junction protein1 (**TJP-1**) genes of birds at 21 d of age, and the expression of duodenal and ileal **OCN** gene in 63-day-old broilers was upregulated due to 450 mg/kg HGT treatment ($P < 0.05$). The dietary supplementation of 150 mg/kg HGT strengthened the expression of duodenal **IL-6** and **IL-4** genes and ileal

IL-4 gene of 21-day-old broilers, whereas the expression of jejunal **IL1B** and **IL-6** genes in birds at 63 d of age weakened because of 300 or 450 mg/kg HGT addition ($P < 0.05$). As for microbial community, the HGT addition altered the cecal microbiota structure of birds at 21 d of age based on analysis of similarities (**ANOSIM**) test and 450 mg/kg HGT treatment increased the relative abundance of *norank Eubacterium coprostanoligenes group* at 21 d of age and *unclassified Lachnospiraceae* at 63 d of age ($P < 0.05$). In short, diet supplemented with 300 to 450 mg/kg HGT may be the optimal for yellow-feather broilers to enhance intestinal barrier function. Altogether, our study clarified the regulatory role of HGT in broiler intestinal health in earnest, but the underlying mechanism is still unclear. Hence, more research is needed to carry out until the application of HGT as a new functional additive in broiler production.

Key words: tannin, chicken, mechanical barrier, immunity, bacterial community

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INTRODUCTION

Novel functional additives are in urgent need to substitute antibiotics, which had been used in husbandry to promote animal intestinal health, improve nutrient absorption and prevent pathogenic bacteria infection before its prohibition of use as a feed additive in European and China (Baumler and Sperandio, 2016). Hydrolyzed gallotannin (**HGT**), a class of polyphenolic compounds, can bind protein and produce 1 molecule of glucose and 10 molecules of gallic acid when hydrolyzed (Aelenei et al., 2009). Several previous studies indicated that dietary tannins supplementation could exert a positive influence on the health of broilers, especially

immune function and antioxidant capacity. A study has shown that there was a significant increase in $CD4^+CD8^+$ and $\gamma\delta^+$ cell populations in white-feather broilers when supplemented with 500 mg/kg tannins (Ramah et al., 2020). Another recent study reported a causal role of HGT administration in strengthening intestinal barrier of mice suffered from diquat-induced oxidative stress (Wang et al., 2019). In our previous study, we found that dietary supplementation with 150 mg/kg of HGT could improve the growth performance of broilers during the early growth phase (1–21 d of age) and HGT treatment protected against oxidative damage of yellow-feather broilers by increasing the content of liver total antioxidant capacity (**T-AOC**) and glutathione/glutathione disulfide ratio (**GSH/GSSG**) (Tong et al., 2022a,b), however, the effect of HGT in intestinal barrier function and microbiota structure was still obscure, and we hypothesized that the sounder intestine might contribute to the growth performance-promoting role of HGT on broilers.

The intestine acts as a defense system to prevent pathogens and toxins from entering the body, and the

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intestinal barrier system is composed of 4 parts: the physical, chemical, immunological, and microbiological barriers (Anderson et al., 2012). The physical barrier, mainly composed of mucosa epithelial cells and intercellular junctions, is the first line of defense (Odenwald and Turner, 2017). Tight junctions are critical in establishing the interepithelial barrier and maintaining epithelial polarity (Otani and Furuse, 2020). The chemical barrier mainly refers to the mucus layer, which is produced by goblet cells and some epithelial cells, and also serves as the first line of innate defense (Johansson and Hansson, 2016). In addition, digestive enzymes, gastric acid, and bile acids that can inhibit the growth of potential pathogens also play an undoubted role in intestinal chemical barrier (Huang et al., 2021). The immunological barrier consists of intestinal immune tissues (like Peyer's patch), immune cells, and immune factor which maintains immunologic homeostasis by orchestrating immune activation and tolerance (Lecocq et al., 2013). Intestinal microbiota, as the microbiological barrier, is essential in inducing cell renewal, fortifying the mucus layer and stimulating mucosa immunity, thereby affecting the host physiology (Hayes et al., 2018).

Consequently, this study was conducted to investigate the effects of dietary supplementation with HGT on intestinal barrier system of yellow-feathered broilers, in order to promote the use of HGT as a new functional additive to maintain intestinal health of yellow-feather broilers.

MATERIALS AND METHODS

Animals were cared for and handled in accordance with the animal care protocols approved by the Fujian Agriculture and Forestry University Animal Care and Use Ethics Committee (Fuzhou, Fujian, China, approval ID: PZCASFAFU23062).

Experimental Materials

HGT (extracted from gallnut, 50% purity and rice hull as the carrier) was obtained from Fujian Jinhualong Feed Co., Ltd. (Fujian, China). The 1-day-old male broilers were offered by WENS Foodstuff Group Co., Ltd. (Guangdong, China). Feedstuffs for broiler diet were purchased from Fujian Jinhualong Feed Co., Ltd. (Fujian, China).

Experimental Design and Bird Management

A total of 288 one-day-old male broilers (34.10 ± 0.08 g) were randomly assigned to 4 treatments: 1) corn-soybean-based diet (CON), 2) CON diet supplemented with 150 mg/kg HGT, 3) CON diet supplemented with 300 mg/kg HGT, 4) CON diet supplemented with 450 mg/kg HGT, with 6 replicates per group and 12 broilers per replicate. The doses of HGT were selected according to a previous report and the recommended dose of the manufacturer (Tonda et al., 2018). The basal

diets formulated according to the feeding standards of Chinese chickens (NY/T33-2004) are showed in [Supplementary Table S1](#). The test period was 63 d. Broilers were reared in cage ($1,562.5 \text{ cm}^2/\text{bird}$) with 23 h of light, ad libitum feeding and water intake. The temperature of the bird room was controlled at 35°C for the first week and then lowered by 2°C to 3°C per wk until 22°C . Additionally, standard immunization procedures were used throughout the experiment.

Sampling Procedure

On d 21 and d 63 of the experiment, 1 broiler from each replicate was randomly chosen to be stunned, exsanguinated, scalded, plucked and sampled. The mucosa of duodenum, jejunum, and ileum was scraped and frozen at -80°C for the detection of digestive enzymes activity and gene expression related to physical and immune barrier function. Cecal digesta were collected and stored at -80°C for 16S rRNA gene sequencing.

Determination of Lipase and α -Amylase Activities

The activities of lipase and α -amylase were determined according to the previous methods (Dahlqvist, 1962; Verduin et al., 1973). The mucosa collected from a middle section of duodenum was fully homogenized with cold PBS (pH = 7.4) with a weight-to-volume ratio of 1:50 and then centrifuged at $1,000 \times g$ for 10 min at 4°C . Total protein content in the intestinal mucosa was evaluated by using a BCA protein quantitative kit purchased from New cell § Molecular Biotech Co. Ltd. (Suzhou, China). The activities of lipase (Catalog number: A054-1-1) and α -amylase (Catalog number: C016-1-1) were determined by using assay kits from Nanjing Jiancheng Institute of Biological Engineering (Nanjing, China). The absorbance was detected by spectrophotometry (iMark, Bio Rad, Hercules, CA).

Gene Expression Analysis by qPCR

Total RNA was extracted from the duodenal, jejunal, and ileal mucosa by using the Eastep super Total RNA Extraction Kit (Directory number: LS1040) from Shanghai Promega Biological Products Co., Ltd. (Shanghai, China), and a Thermo Scientific NanoDrop 2000 Spectrophotometer (Wilmington) was used to determine the concentration and quality of RNA. Total RNA was reverse-transcribed via using the Eastep RT Master Mix Kit from Shanghai Promega Biological Products Co., Ltd. (Shanghai, China). The cDNA was used for quantitative real-time PCR on an Applied Biosystems. The primer sequences were synthesized by Sango Biotech Co., Ltd. (Shanghai, China) and displayed in [Supplementary Table S2](#). The relative mRNA expression levels of each target gene were analyzed according to the method described by Livak and Schmittgen (2001) and shown as $2^{-\Delta\Delta\text{CT}}$.

Bacterial DNA Extraction and 16S rRNA Gene Sequencing

The bacterial DNA of digesta was extracted by using Stool DNA Kit (D4015-01, Omega Bio-tek, Norcross, GA). The V3 to V4 regions of the bacterial 16S rRNA genes were amplified with the primers 338F (5'-ACTCC-TACGGGAGGCAGCAG-3') and 806R (5'-GGAC-TACHVGGGTWTCTAAT-3') by an ABI GeneAmp 9700 PCR Thermocycler (Waltham, MA). Purified amplicons were pooled in equimolar and paired-end sequenced on an Illumina MiSeq PE300 platform (San Diego, CA) according to the standard protocols by Majorbio BioPharm Technology Co. Ltd. (Shanghai, China).

Amplicon sequence variants (ASVs) with 99% similarity cutoff delineated were clustered using divisive Amplicon Denoising Algorithm 2 (DADA2). Rarefaction curves were made using Python 2.7 to assess whether the sequencing volume is sufficient (Gotelli and Colwell, 2001). The unique and common ASVs observed in different groups are showed in Venn diagram. The α -diversity was calculated using accumulated cyclone energy (ACE), Simpson, and Shannon diversity indices by mothur software (Schloss et al., 2009). Principal coordinate analysis (PCoA) based on unweighted UniFrac distance to analysis the similarities between samples and groups, and the intergroup difference analysis was performed by analysis of similarities (ANOSIM) test.

The raw microbial sequencing data have been deposited at the National Center for Biotechnology Information database (NCBI), under accession number PRJNA888548. The data that support the study findings and models are available from the authors upon reasonable request.

Data Analysis and Statistics

Statistical analysis was performed using SPSS 25.0 software (Chicago, IL). The intergroup differences were examined using 1-way ANOVA analysis with Duncan's post hoc test. The linear and quadratic comparisons were applied to determine the dose-effect of HGT on intestinal function of broilers. Specially, statistically significant differences in the relative abundance of the cecal microbiota were determined using Kruskal-Wallis test with Tukey's post hoc test. P value <0.05 was considered statistically significant.

RESULTS

Growth Performance

The effect of HGT supplementation on the growth performance of broilers is presented in [Supplementary Table S3](#) (Tong et al., 2022a,b). Compared with CON group, dietary supplementation with 150 mg/kg HGT significantly decreased the Feed:Gain of broilers at 1 to 21 d of age ($P < 0.05$), but no obvious effects of HGT treatment on ADFI and ADG were observed. Meanwhile, HGT supplementation had no effects on ADFI, ADG, and Feed:Gain of broilers at 1 to 63 d of age.

Digestive Enzyme Activity

The results of digestive enzyme activity are presented in [Table 1](#). At 21 d of age, the 150 mg/kg HGT supplementation notably increased the lipase activity in duodenum compared with the CON group ($P < 0.05$). However, α -amylase was not significantly different between groups. At 63 d of age, there were no effects of HGT treatment on α -amylase and lipase activity in duodenum of broilers.

Gene Expression of Tight Junction Proteins

As shown in [Figure 1](#), compared with the CON group, HGT supplementations of 300 or 450 mg/kg could increase the gene expression of occludin (*OCLN*) and tight junction protein1 (*TJP-1*) in duodenal mucosa on d 21 ($P < 0.05$) ([Figure 1A](#)). But no significant difference in gene expression of tight junction proteins in jejunal and ileal mucosa of 21-day-old broilers was observed between groups ([Figure 1C](#) and [E](#)). At 63 d of age, the 450 mg/kg HGT supplementation in the diet of broilers increased the expression of *OCLN* in duodenal and ileal mucosa compared with the CON group ($P < 0.05$) ([Figure 1B](#) and [F](#)). However, there were no special effects of HGT treatment on gene expression of tight junction proteins in jejunal mucosa ([Figure 1D](#)).

Gene Expression of Cytokines

Compared with the CON group, dietary supplementation of 150 mg/kg HGT increased the expression of *IL-6* and *IL-4* in duodenal mucosa of 21-day-old broilers ($P < 0.05$) ([Figure 2A](#)). Notably, dietary

Table 1. The effect of HGT on digestive enzyme activity in duodenum of broilers.¹

Items	HGT, mg/kg				SEM	ANOVA	P value	
	0	150	300	450			Linear	Quadratic
21 d of age								
α -Amylase, U/g prot	79.88	78.07	60.34	74.37	4.365	0.388	0.466	0.303
Lipase, U/g prot	0.554 ^b	1.955 ^a	0.629 ^b	0.677 ^b	0.198	0.023	0.299	0.102
63 d of age								
α -Amylase, U/g prot	48.98	56.67	55.90	56.76	3.616	0.870	0.522	0.664
Lipase, U/g prot	1.088	1.243	1.193	1.821	0.161	0.390	0.154	0.463

¹Values are means of 6 replicates per treatment with 1 bird each.

^{a-b}Means in the same row without the same superscript differ significantly ($P < 0.05$).

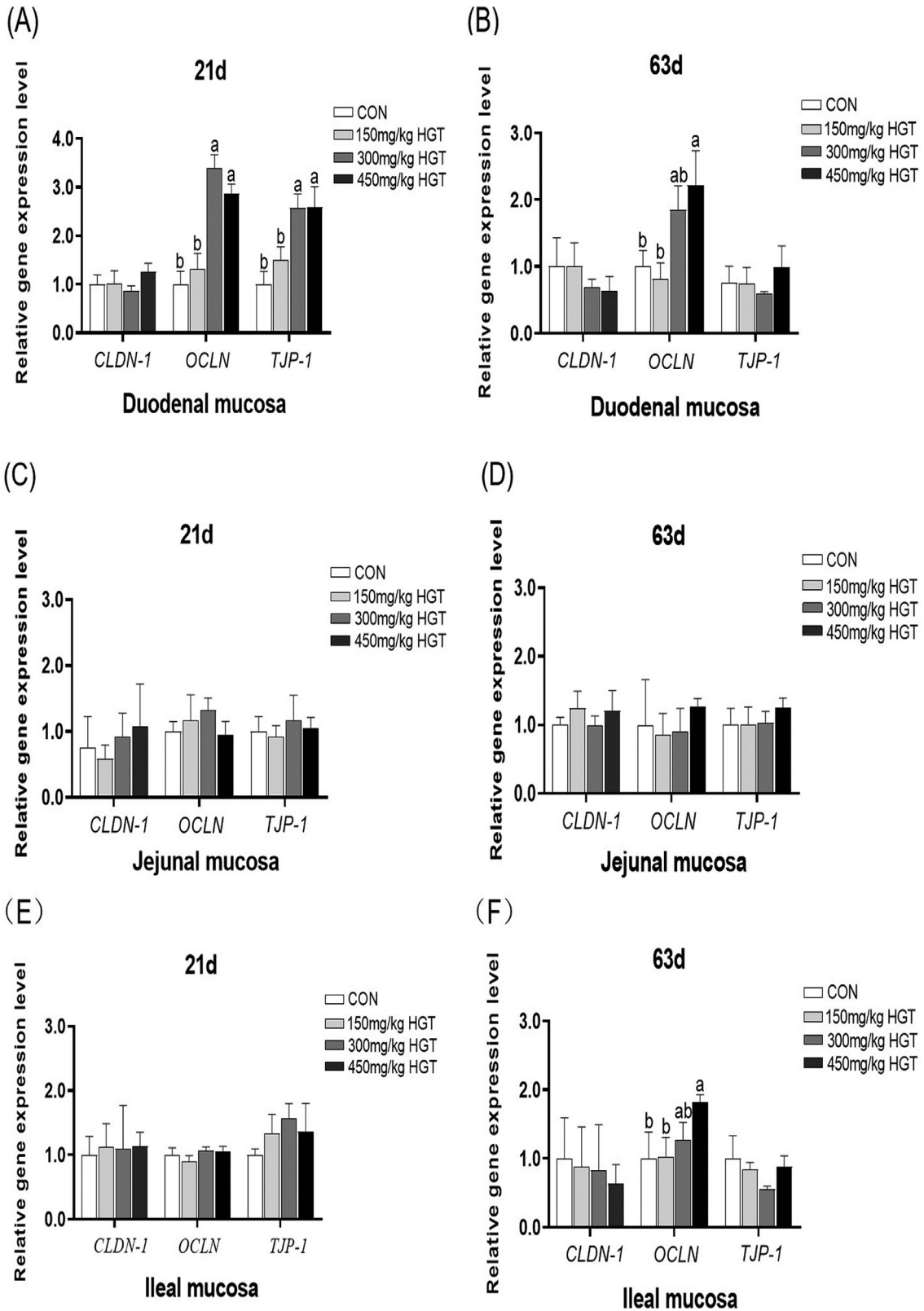


Figure 1. The effect of HGT on gene expression of tight junction proteins in intestinal mucosa of broilers. Values are means of 6 replicates per treatment with 1 bird each. *CLDN-1*, *OCLN*, and *TJP-1* are the genes that encode claudin1, occludin, and tight junction protein1, respectively. CON group: basal diet; 150 mg/kg HGT group: basal diet added with 150 mg/kg HGT; 300 mg/kg HGT group: basal diet added with 300 mg/kg HGT; 450 mg/kg HGT group: basal diet added with 450 mg/kg HGT. ^{a-b}Means in the same row without the same superscript differ significantly ($P < 0.05$). Error bar stands for SEM.

supplementation with 450 mg/kg HGT could obviously increase the expression of *IL-6* in jejunal mucosa on d 21 ($P < 0.05$) (Figure 2C). Interestingly, HGT supplementations of 150 mg/kg could increase

the expression of *IL-4* in ileal mucosa at 21 d of age ($P < 0.05$) (Figure 2E).

The expression of *IL1B* and *IL-6* in jejunal mucosa from 63-day-old broilers fed with 300 and 450 mg/kg

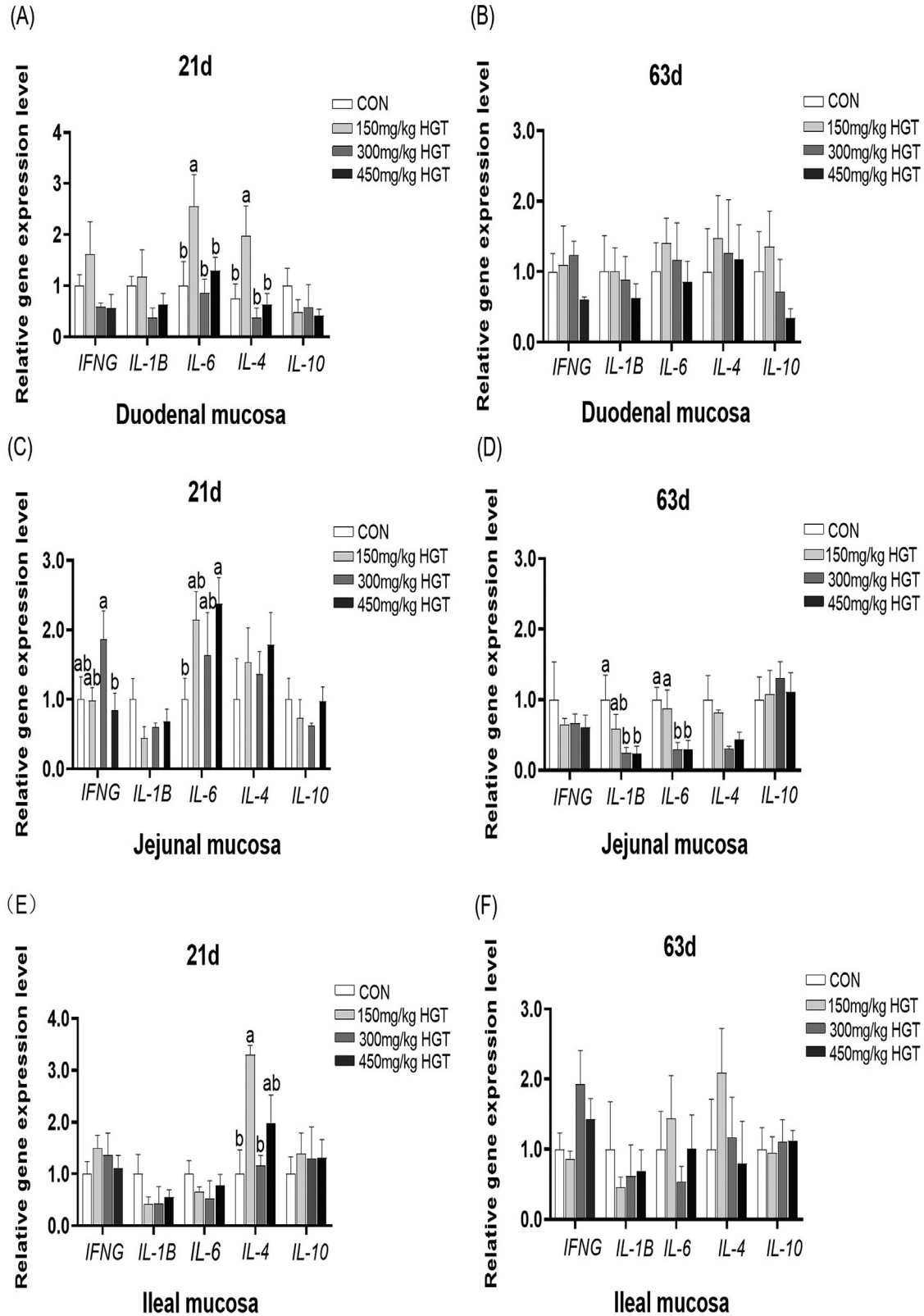


Figure 2. The effect of HGT on gene expression of cytokines in intestinal mucosa of broilers. Values are means of 6 replicates per treatment with 1 bird each. Abbreviation: IFNG, interferon gamma. CON group: basal diet; 150 mg/kg HGT group: basal diet added with 150 mg/kg HGT; 300 mg/kg HGT group: basal diet added with 300 mg/kg HGT; 450 mg/kg HGT group: basal diet added with 450 mg/kg HGT. ^{a-b}Means in the same row without the same superscript differ significantly ($P < 0.05$). Error bar stands for SEM.

HGT was lower than broilers fed with basal diet ($P < 0.05$) (Figure 2D). However, there were no obvious differences among diets in the HGT of the expression of *IFNG*, *IL1B*, *IL-6*, *I-L4*, and *IL-10* in duodenal and ileal mucosa on d 63 (Figure 2B and F).

Cecal Microbiota

After denoise and quality control, 470,225 and 418,586 16S rRNA gene sequences were obtained from 16 cecal bacterial DNA samples of yellow-feather broilers at 21 d

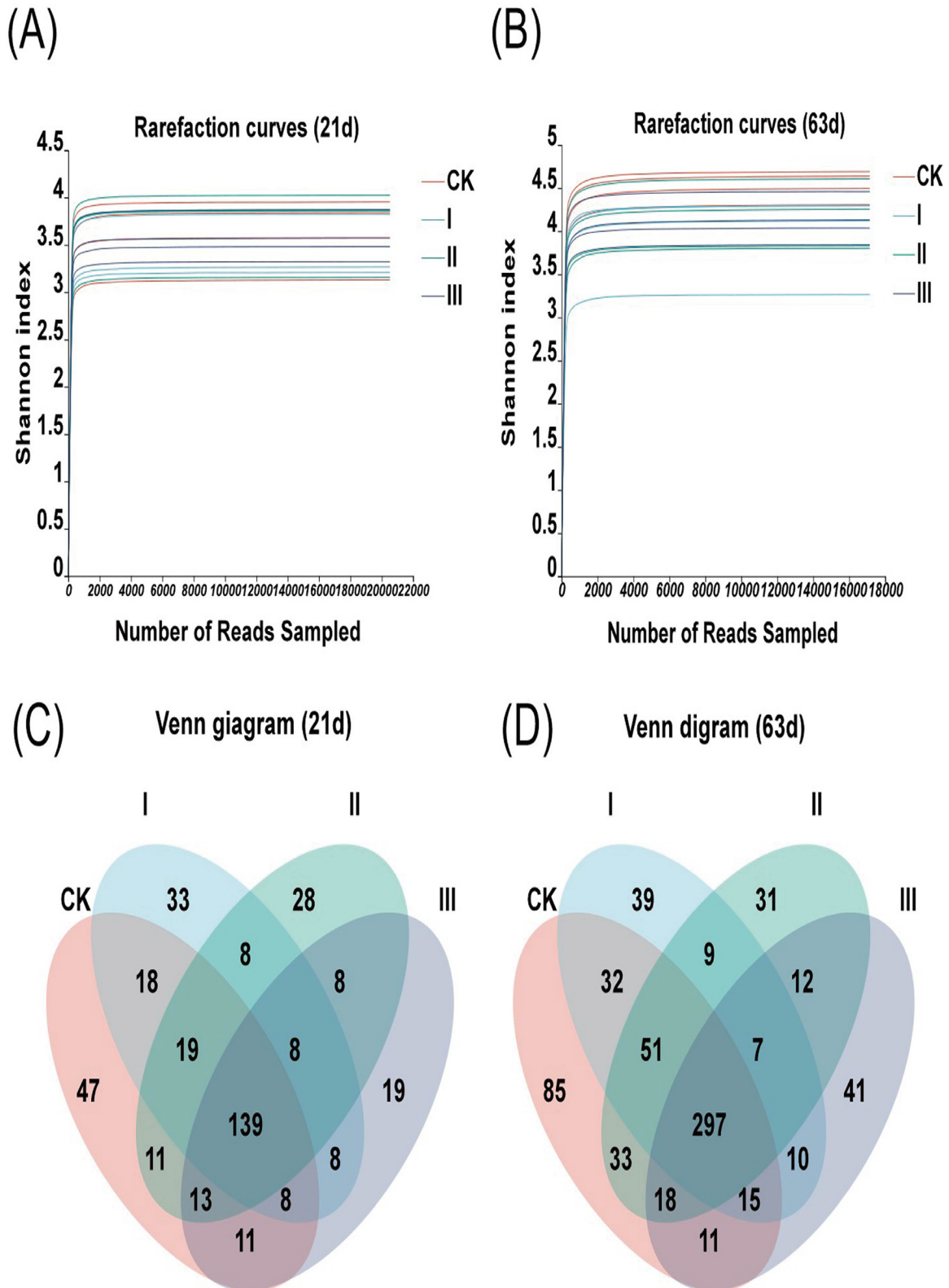


Figure 3. Rarefaction curves of sequencing reads and Venn diagram of ASVs in broilers. Values are means of 4 replicates per treatment with 1 bird each. Group CK: basal diet; Group I: basal diet added with 150 mg/kg HGT; Group II: basal diet added with 300 mg/kg HGT; Group III: basal diet added with 450 mg/kg HGT.

and 63 d of age, respectively. Rarefaction curve analysis showed that the sequencing data were sufficient to reflect the microbial diversity (Figure 3A and B).

Besides, 378 and 691 ASVs are obtained based on 99% sequence similarity from all samples on d 21 and d 63,

respectively (Figure 3C and D). On d 21, there were 139 shared ASVs and 47, 33, 28, and 19 ASVs unique in the Group CK (0 mg/kg HGT), Group I (150 mg/kg HGT), Group II (300 mg/kg HGT), and Group III (450 mg/kg HGT), respectively (Figure 3C). On d 63, there were 297

Table 2. Effects of HGT on alpha diversity in cecum microorganism of yellow-feather broilers.¹

Items	HGT, mg/kg				SEM	P value
	0	150	300	450		
21 d of age						
ACE index	150.7	136.8	136.8	120.8	4.774	0.226
Simpson index	0.065	0.062	0.053	0.054	0.006	0.571
Shannon index	3.63	3.54	3.73	3.56	0.078	0.766
63 d of age						
ACE index	337.8	263.8	269.3	227.3	15.40	0.106
Simpson index	0.026	0.056	0.057	0.045	0.007	0.178
Shannon index	4.53	4.00	4.12	4.12	0.093	0.150

¹Values are means of 4 replicates per treatment with 1 bird each. Abbreviation: ACE, accumulated cyclone energy.

ASVs shared among all groups, whereas 85, 39, 31, and 41 ASVs were identified only present in the Group CK, I, II and III, respectively (Figure 3D).

As shown in Table 2, the dietary addition of HGT did not affect the ACE index, Simpson index, and Shannon index of cecum microorganisms in birds at 21 and 63 d of age. As revealed in Figure 4A, PCoA based on unweighted UniFrac distance showed that the distance of samples between groups is far at 21 d of age ($P = 0.002$), indicating that the microbiota structure between groups were distinguishing. However, the cecal microbiome profiles of 63-day-old broilers were not significantly different between groups (Figure 4B).

Firmicutes was predominant and Proteobacteria and Actinobacteria were subdominant in cecal microbiota community of 21-day-old broilers. There were no significant differences in the relative abundance of these 3 phyla levels among groups (Table 3 and Figure 5A). At the genus level, the cecal microbiota community were dominated by *unclassified Lachnospiraceae*, *Faecalibacterium*, *Ruminococcus torques* group, *Blautia*, and *Subdoligranulum*, but no remarkable intergroup differences were observed. Only the relative abundances of *norank Eubacterium coprostanoligenes* group in Group II and

Table 3. Relative richness of cecal dominant phyla and genera in 21-day-old broilers.¹

Items (%)	HGT, mg/kg				SEM	P value
	0	150	300	450		
Phylum level						
Firmicutes	98.45	97.46	97.69	98.70	0.368	0.431
Proteobacteria	1.13	1.13	1.08	0.66	0.260	0.830
Actinobacteria	0.42	1.40	1.23	0.64	0.221	0.131
Genus level						
<i>Unclassified</i>	30.50	53.02	33.75	43.03	3.275	0.065
<i>Lachnospiraceae</i>						
<i>Faecalibacterium</i>	26.47	1.312	11.21	2.090	3.800	0.069
<i>Ruminococcus torques</i> group	7.82	7.97	9.90	14.35	1.323	0.368
<i>Blautia</i>	5.99	12.03	8.66	6.34	0.925	0.057
<i>Subdoligranulum</i>	4.35	3.36	2.09	5.23	1.241	0.687
<i>Norank Eubacterium coprostanoligenes</i> group	1.45 ^b	2.52 ^{ab}	4.22 ^a	4.42 ^a	0.410	0.010
<i>Romboutsia</i>	1.00	0.27	4.34	5.08	1.015	0.056
<i>Erysipelatoclostridium</i>	2.13	1.70	2.86	3.03	0.284	0.235
<i>Sellimonas</i>	2.17	1.54	3.04	1.92	0.299	0.562
<i>Unclassified</i>	2.42	0.42	1.28	3.26	0.643	0.524
<i>Peptostreptococcaceae</i>						

¹Values are means of 4 replicates per treatment with 1 bird each.

^{a-b}Means in the same row without the same superscript differ significantly ($P < 0.05$).

Group III were significantly higher than Group CK ($P < 0.05$) (Table 3 and Figure 5B).

As presented in Table 4 and Figure 6, at the phylum level, the cecal microbiota community in 63-day-old broilers was dominated by Firmicutes, Bacteroidota, and Campylobacterota and their proportions accounted for more than 99%. The relative abundance of Campylobacterota decreased from 1.95 to 0.39% because of 450 mg/kg HGT treatment ($P < 0.05$). At the genus level, *Faecalibacterium*, *unclassified Lachnospiraceae*, *Romboutsia*, *norank Clostridia vadinBB60* group, *norank Clostridia UCG-014*, and *Blautia* were dominant genera. The relative abundance of *unclassified Lachnospiraceae* in Group III was significantly higher than Group CK ($P < 0.05$).

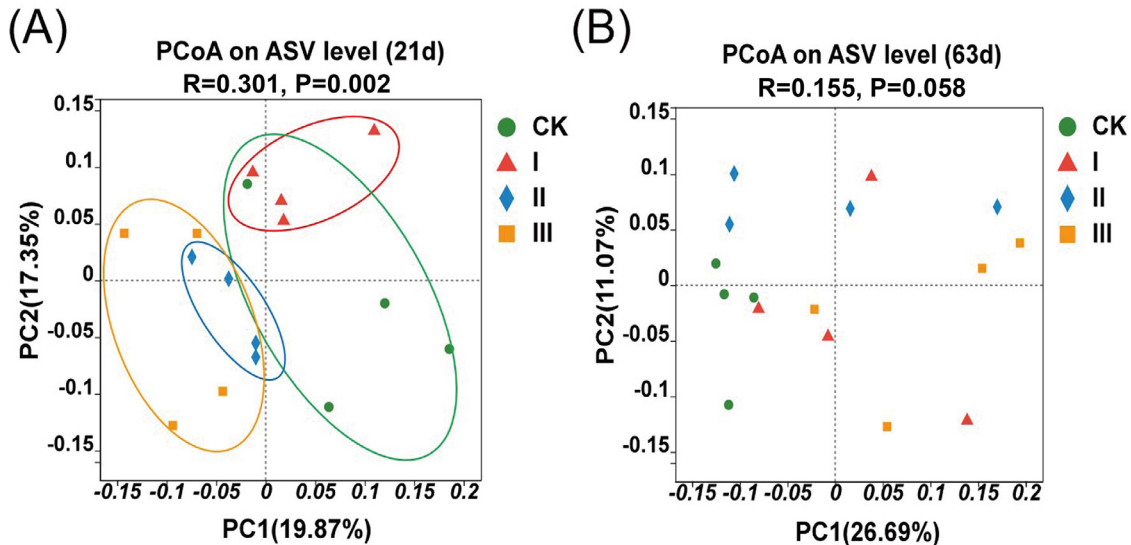


Figure 4. PCoA analysis based on unweighted UniFrac distance and ANOSIM analysis. Group CK: basal diet; Group I: basal diet added with 150 mg/kg HGT; Group II: basal diet added with 300 mg/kg HGT; Group III: basal diet added with 450 mg/kg HGT. Abbreviations: ANOSIM, analysis of similarities; ASV, amplicon sequence variants; PCoA, principal coordinates analysis.

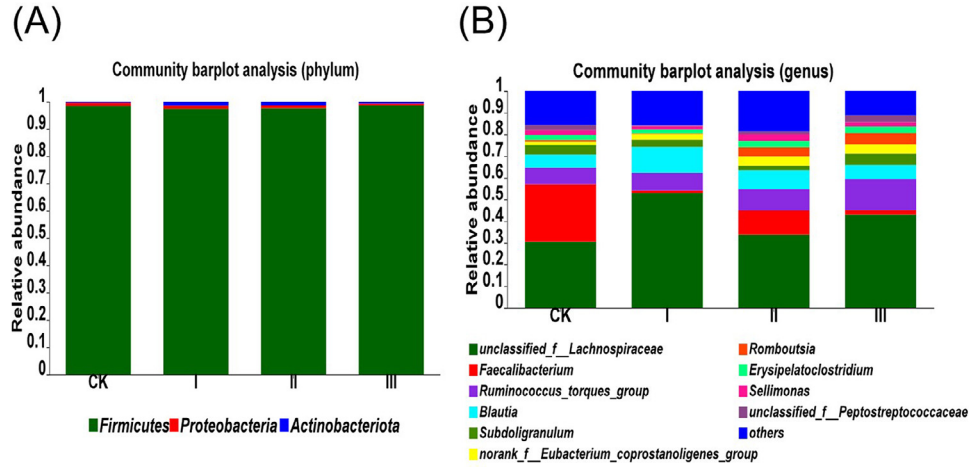


Figure 5. Relative abundance at phylum level (A) and genus level (B) in cecum microbiota of 21-day-old broilers. Values are means of 4 replicates per treatment with 1 bird each. Group CK: basal diet; Group I: basal diet added with 150 mg/kg HGT; Group II: basal diet added with 300 mg/kg HGT; Group III: basal diet added with 450 mg/kg HGT.

DISCUSSION

Recently, polyphenol compounds have gained substantial attention because of their antioxidant properties (Surai, 2014). The growth-promoting and health-improving benefits due to phenolic compounds supplementation in broiler diets have been reported in several studies (Starcevic et al., 2015; Mahfuz et al., 2021). Specially, HGT, an endogenous plant phenol, is mainly derived from Chinese gallnuts (Haslam, 1996). Our previous studies indicated that dietary supplementation with 150 mg/kg of HGT could improve the feed conversion ratio (FCR) of broilers during the early growth phase (1–21 d of age), and elevate the content of liver T-AOC and GSH/GSSG (Tong et al., 2022a,b). We speculated that a stronger intestine might facilitate the FCR-promoting role of HGT in broilers, and HGT-generated reinforced antioxidant capacity can also contribute to a healthier intestine. As we expected, in our

Table 4. Relative richness of cecal dominant phyla and genera in 63-day-old broilers.¹

Items (%)	HGT, mg/kg				SEM	P value
	0	150	300	450		
Phylum level						
Firmicutes	92.40	88.49	90.78	90.34	0.993	0.409
Bacteroidota	5.02	9.43	6.57	8.91	1.029	0.309
Campylobacterota	1.95	1.45	2.48	0.39	0.322	0.030
Genus level						
<i>Faecalibacterium</i>	17.93	28.43	28.39	22.80	2.865	0.128
<i>Unclassified</i>	14.28 ^b	10.31 ^b	12.92 ^b	19.87 ^a	1.080	0.024
<i>Lachnospiraceae</i>						
<i>Romboutsia</i>	5.35	6.24	6.36	9.50	0.816	0.382
<i>Norank Clostridia vadinBB60 group</i>	7.96	6.97	7.36	1.82	1.064	0.117
<i>Norank Clostridia UCG-014</i>	6.37	4.74	4.14	2.11	0.676	0.080
<i>Blautia</i>	3.38	4.10	2.84	5.59	0.511	0.497
<i>Ruminococcus torques group</i>	2.94	3.78	4.14	4.58	0.306	0.343
<i>Parabacteroides</i>	2.56	1.76	3.28	3.32	0.526	0.589

¹Values are means of 4 replicates per treatment with 1 bird each.

^{a–b}Means in the same row without the same superscript differ significantly ($P < 0.05$).

study, the higher lipase activity in duodenum was observed in the 21-day-old broilers fed with 150 mg/kg HGT. It is widely accepted that the higher digestive enzyme activity indicates the more complete utilization of nutrient (Chen et al., 2022). Similarly, it was found that the tannins at low concentration (200 and 300 mg/d) had a positive effect on the activities of lipase, amylase and trypsin to strengthen nutrient digestion and uptake (Majumdar and Moudgal, 1994). However, Quesada et al. (1995) reported that extracts of pears, lentils and cocoa beans, rich in condensed and hydrolyzed tannins, could inhibit the activity of α -amylase, and the higher the dose, the stronger the inhibition effect. It is common knowledge that the phenolic hydroxyl in tannins acts as the hydrogen donor and the carbonyl peptide in enzyme serves as the hydrogen acceptor. Carbonyl peptide of the enzyme was excessively exposed to the solution of hydrolyzed tannin, resulting in its precipitation and the loss of enzyme activity (Zhu et al., 1997). However, there is a great discrepancy in the affinities between tannins and enzymes. Hence, the reason for the above different results on enzyme activity might be due to the effective content of tannins and the discrepant affinities between tannins and enzymes (Nyamambi et al., 2000).

The interepithelial physical barrier primarily consists of tight junction, adhere junction, and desmosome and is essential to maintain intestinal integrity and sustain the polarity of epithelial cells (Farquhar and Palade, 1963). Herein, tight junction is a multiprotein complex including the transmembrane barrier proteins (claudins, occludin, etc.) and peripheral scaffolding proteins (ZO-1, ZO-2, ZO-3, etc.) (Van Itallie and Anderson, 2014). Claudins were first identified in junctional fraction of chicken liver, and the gene overexpression of Claudins (*CLDNs*) in fibroblasts was reported sufficient to reconstitute tight junction-like networks of strands (Günzel and Yu, 2013). Additionally, Umeda et al. (2004) observed a marked retardation of tight junction formation when *TJP-1* expression of epithelial cell was hampered. Our results showed that dietary

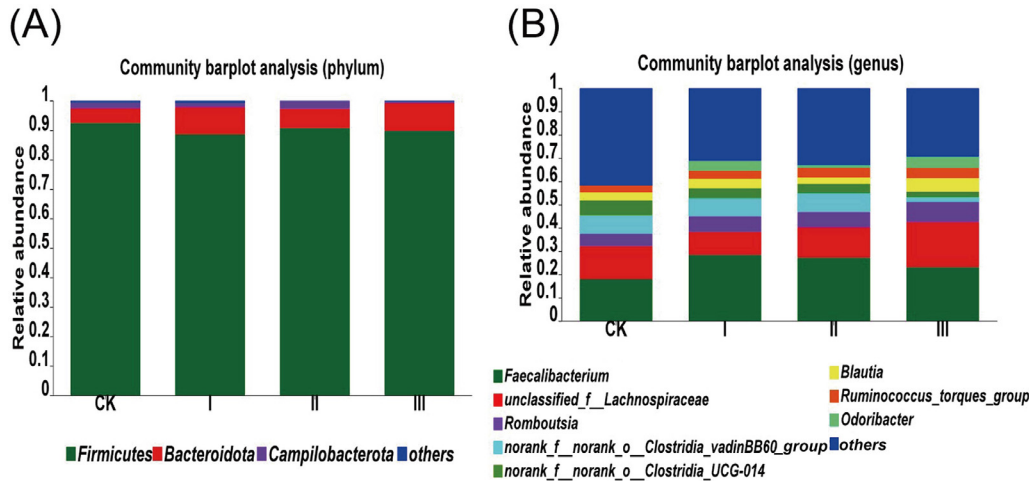


Figure 6. Relative abundance at phylum level (A) and genus level (B) in cecal microbiota of 63-day-old broilers. Values are means of 4 replicates per treatment with 1 bird each. Group CK: basal diet; Group I: basal diet added with 150 mg/kg HGT; Group II: basal diet added with 300 mg/kg HGT; Group III: basal diet added with 450 mg/kg HGT.

supplementation HGT increased the expression of *OCN* and *TJP-1* in duodenum and ileum mucosa of yellow-feather broilers, especially in broilers from 1 to 21 d of age. Similarly, Liu et al. (2018) found that addition of 2 g/kg HGT in diet strongly increased the expression of *TJP-1* in heat-stressed broilers to alleviate the negative effects of heat stress on intestinal function. Previous studies also found that 2.5 and 5 mg/kg pretreated tannins could significantly increase the expression of *TJP-1* in diquat-challenged mice, but the 10 mg/kg tannins addition significantly inhibited the expression of *CLDN*. So the effect of tannins might on intestinal tight junction might partly depends on the physical state of body and the dosage of tannins (Wang et al., 2019). Overall, despite a few inconsistent reports, it is reasonable to speculate that dietary HGT could upregulate the gene expression of tight junction protein, then improving physical barrier function of broilers.

Cytokines play an important role in the modulation of inflammatory response against intestinal mucosal damage (Shen and Turner, 2006). Th2 cytokines such as IL-4 play a crucial role in selective activation of macrophages, whereas IL-6 is a multifunctional cytokine that triggers intracellular signaling pathways to regulate immune response (Rose-John, 2012; Chaudhari et al., 2018). In our study, an increase in the expression of *IL-4* and *IL-6* on d 21 due to 150 mg/kg HGT supplementation and a decrease in the expression of *IL1B* and *IL-6* on d 63 on account of 300 or 450 mg/kg HGT addition were observed, indicating that HGT had a positive and significant effect on prevention of pathogen invasion during the early growth stage (1–21 d) and an energy-save and production-promotion effect through a reduction of immune response in the late growth stage (22–63 d) of broilers. Consistent with our results, Kawano et al. (2020) discovered that tannins enhanced IL-4 secretion from CD4⁺ T cells and CD8⁺ T cells in a dextran sodium salt-induced colitis model of mouse. Additionally, a previous study revealed that 25 mg/kg tannic acid reduced inflammatory response by downregulating the protein

expression of toll-like receptor 4 and restraining extracellular signal-regulated kinase 1/2 and p38 mitogen-activated protein kinase activation, then inhibiting the expression of *Tnfa* and *Il1b* in mice (Sivanantham et al., 2019). Nevertheless, Ramah et al. (2020) found that low-dose (500 mg/kg) tannins resulted in the expansion of CD4⁺CD8⁺ and $\gamma\delta^+$ cell populations, whereas high-dose (30 g/kg) tannins supplementation generated a smaller T and B cell subsets and inhibited the expression of *IL-4* in the spleen. The different results on cytokines expression and secretion might be due to the plant sources of tannins and animal species. Collectively, these observations suggested tannins could be a potential critical immunoregulator of intestinal health in broilers.

A growing body of scientific evidence has demonstrated that polyphenols derived from plants could alter the composition of gut microbial community (Lillehoj et al., 2018). Tannins were reported to induce morphological changes of bacteria in growth phase and to inhibit the microbial proliferation by limiting iron availability to metalloenzymes in microbial cells (Jones et al., 1994). Lee et al. (2010) reported that tannins had antimicrobial activity because there was a lower fecal population of coliform bacteria in weanling pigs fed tannins diets. It was widely accepted that tannins could disrupt membrane function of bacteria by inhibiting the electron transport system and oxidative phosphorylation (Goel et al., 2005). Furthermore, previous studies have described a reduction in genus *Bacteroides* but certain members of Ruminococcaceae and Lachnospiraceae were increased in broilers fed with tannins (Diaz Carrasco et al., 2018). In our study, dietary HGT supplementation notably affected the composition and structure of cecal microbial community in 21-day-old broilers, and the relative abundance of *norank Eubacterium coprostanoligenes* group at 21 d of age and *unclassified Lachnospiraceae* at 63 d of age were raised after 450 mg/kg HGT treatment. *Eubacterium coprostanoligenes* could decrease serum cholesterol concentration by modulating the conversion of absorbed cholesterol into coprostanol,

which plays a key role in cholesterol metabolism homeostasis (Martin et al., 1972; Li et al., 1998; Kriaa et al., 2019). In addition, some studies reported that the Lachnospiraceae family modified the immune statues of their hosts by producing short-chain fatty acids and converting primary bile acids into secondary bile acids (Sorbara et al., 2020), which are considered to have beneficial effects on intestinal health via regulation of gut microbiome community structure (Ridlon et al., 2014). Accordingly, it can be speculated that tannins might alter the relative abundance of beneficial intestinal bacteria but the underlying mechanisms are not to completely understand yet.

CONCLUSIONS

In conclusion, the supplementation of HGT in the diet of broilers could improve intestinal barrier function and regulate the dynamic balance of intestinal flora, thereby promoting the intestinal health of broilers. Overall, dietary supplementation of 300 to 450 mg/kg HGT is recommended for better intestinal function of yellow-feather broilers.

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DISCLOSURES

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in the present study.

SUPPLEMENTARY MATERIALS

Supplementary material associated with this article can be found in the online version at doi:10.1016/j.psj.2023.103010.

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