J Anim Sci Technol 2020;62(2):159-173 https://doi.org/10.5187/jast.2020.62.2.159



# Replacement value of cottonseed meal for soybean meal in broiler chicken diets with or without microbial enzymes

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## Abstract

A 4×2 factorial feeding trial was designed to investigate the effect of replacing soybean meal (SBM) with cottonseed meal (CSM) in wheat/sorghum/SBM-based diets fed with or without microbial enzymes in diets on the performance, visceral organ development and digestibility of nutrients of broiler chickens. Four graded levels of CSM - none (0%), low (4%, 8%, and 12%), medium (5%, 10%, and 15%), and high (6%, 12%, and 18%) of complete diets in starter, grower and finisher, respectively were fed with or without 100 mg/kg of xylanase and  $\beta$ -glucanase blend. Eight isocaloric and isonitrogenous diets were formulated using least-cost method to meet the nutrient specifications of Ross 308 male broilers. Each treatment was randomly assigned to 6 replicates (10 birds per replicate). There were CSM-enzyme interactions (p < 0.05) on feed intake (FI) and weight gain (WG) in the starter phase. Enzyme supplementation improved (p < 0.05) feed conversion ratio (FCR) in the grower and finisher phases, and increased WG in growing and finishing birds. CSM inclusion reduced (p < 0.05) the weight of gizzard and proventriculus in starter chicks, while these organs were bigger (p < 0.05) in the grower phase. The test ingredient decreased (p < 0.05) small intestinal weight in starter and grower birds. The CSM increased the absolute weight of thighs (p < 0.05) while breast meat was increased (p< 0.01) by enzyme addition. Starch digestibility was improved (p < 0.01) by enzyme inclusion and decreased (p < 0.01) by CSM. Enzyme supplementation improved (p < 0.05) the ileal digestibility of gross energy and protein. The results demonstrate that CSM can substitute up to 90% SBM in broiler chicken diets without compromising performance, and the nutritive value of CSM-containing diets can effectively be improved by enzyme supplementation.

Keywords: Broilers performance, Cottonseed meal, Microbial enzymes, Nutrient digestibility, Soybean meal



Received: Jul 21, 2019 Revised: Nov 13, 2019 Accepted: Nov 13, 2019

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#### **Competing interests**

No potential conflict of interest relevant to this article was reported.

#### **Funding sources**

This work was supported by Feedworks Pty Ltd, Australia and the University of New England.

#### Acknowledgements

The authors gratefully acknowledge the Feedworks Pty Ltd, Australia for their financial support, and the University of New England.

#### Availability of data and material

Upon reasonable request, the datasets of this study can be available from the corresponding author.

#### Authors' contributions

Conceptualization: Iji PA, Abdallh ME. Data curation: Abdallh ME, Musigwa S, Ahiwe EU, Chang'a EP, Al-Qahtani M, Bhuiyan M. Formal analysis: Abdallh ME. Methodology: Abdallh ME. Software: Abdallh ME. Validation: Abdallh ME. Validation: Abdallh ME. Writing - original draft: Abdallh ME, Musigwa S Writing - review & editing: Abdallh ME, Musigwa S, Ahiwe EU, Chang'a EP, Al-Qahtani M, Bhuiyan M, Iji PA.

## Ethics approval and consent to participate

This article does not require IRB/IACUC approval but was approved by the University of New England Animal Ethics Committee (AEC15-084).

# INTRODUCTION

The price of conventional protein sources such as sovbean meal (SBM), which is the predominant plant protein source used in the poultry industry, keeps increasing. This may result in loss in volume of poultry production due to the cost of production or formulation of poor diets. Therefore, cheaper alternative protein ingredients are desperately needed in poultry feed formulation that can maintain feed quality and bird performances [1,2]. For this purpose, cottonseed meal (CSM) has extensively been investigated and it has been shown to be a potential source of protein and a substitute for costly SBM in poultry diets [1-3]. This oilseed residue is highly rich in protein, with values between 41% and 44% crude protein (CP), depending on the degree of dehulling before oil extraction [4]. On the other hand, using this ingredient in poultry feeds presents a number of challenges and limitations, including high fibre content, presence of gossypol, and unbalanced amino acids, particularly a low content of lysine [1,4]. The amount of gossypol in CSM has been reported to vary from 0.03% to 0.2% [2,5]. The high fibre content and gossypol presence in CSM have deleterious effects on feed intake (FI) and growth performance [6]. A number of reports have demonstrated different ways to optimize the use of CSM in poultry diets, including plant breeding, proper processing [2], supplementation with lysine [7] and other feed additives. Furthermore, special combination of dietary ingredients and feed additives can improve bird performance [8].

Exogenous enzymes are feed additives which can improve CSM utilization by poultry. One of the factors that can impede nutrient digestibility and reduce dietary protein utilization is the presence of non-starch polysaccharides (NSP). According to Perez-Maldonado and Australian Egg Corporation Limited [9] the total content of all NSP fractions of commonly used Australian CSM is reported to be as follows (g/kg dry matter): arabinose (37–44), xylose (31–33), mannose (3–4), rhamnose (3–4), fucose (1–1.5), galactose (10–11), and glucose (40–46). It is well established that supplementing NSP-degrading enzymes, such as  $\beta$  -glucanase, xylanase and pectinase in fibrous diets can potentially lower the intestinal content viscosity and improve digestibility [10,11]. Numerous reports have investigated the effect of using xylanase with or without  $\beta$  -glucanase in wheatbased diets on nutrient utilization and broiler performances [10–12]. However, the information on the impact of incorporating a  $\beta$  -glucanase and xylanase cocktail in diets containing CSM as a major source of protein for broiler is still limited. Therefore, the aim of this study was to investigate the effect of graded replacement of SBM by CSM with or without a  $\beta$  -glucanase and xylanase enzyme blend on broiler chicken growth performance and nutrient digestibility.

The overall objective of this study was to evaluate CSM as an alternative and cost-effective protein ingredient to SBM without compromising broiler performance. To achieve this, two specific objectives were developed: (a) To investigate the effect of CSM levels in broiler diets on nutrient digestibility and bird performance; and (b) To evaluate the response of birds to CSM-containing diets supplemented with a composite microbial enzyme product with xylanase and  $\beta$ -glucanase activities. Therefore, based on the above objectives, the following hypotheses were formulated: (a) The level of CSM inclusion in diets can influence the nutritive value of the diets; and (b) The nutritive value of CSM-containing diets can be improved by microbial enzyme supplementation.

# MATERIALS AND METHODS

#### **Dietary treatments**

Four wheat/sorghum/soybean-based diets were formulated using the least-cost method to meet the nutrient specifications for Ross 308 broiler chicks Aviagen [13]. The diets contained different levels of CSM as follows: control (0%) in all growing phases (starter, grower, and finisher), low (4%, 8%,

and 12%), medium (5%, 10%, and 15%) and high (6%, 12%, and 18%) respectively. Each diet was fed as such or supplemented with composite microbial enzymes (Axtra XB) at 100 mg/kg of diet. Axtra XB (Danisco Animal Nutrition, UK) is a blend of two microbial enzymes, xylanase and  $\beta$ -glucanase, providing 250 units of  $\beta$ -glucanase and 2,500 units of xylanase per kg of the formulated feeds. In addition, an indigestible marker, titanium dioxide (TiO<sub>2</sub>), was incorporated in the grower diets at a rate of 5 g/kg of diet, in order to assess nutrient digestibility. The starter chicks (from one to 10 days) were fed with crumbled feed, while the grower (from 11 to 24 days) and finisher (from 25 to 35 days) chicks were fed whole pellet diets. Diets were pelleted at 65 °C. Tables 1 and 2 show the ingredient composition of the diets used in the trial and the calculated nutrient composition of these diets respectively, while Table 3 shows the analyzed nutrient composition of the tested CSM, which were locally produced and purchased (Riverina, Milton, QLD, Australia).

#### Experimental design and bird husbandry

The trial was a  $4 \times 2$  factorial arrangement in which birds were randomly allocated into eight treatments. Each treatment was replicated six times, with 10 chickens per replicate, which made a total of 480 birds. One-day old male Ross 308 chicks were purchased from a commercial hatchery, (Baiada, Tamworth NSW, Australia), with average initial weight of 41.1 ± 0.8 g/chicks. The cages were placed in a climate controlled room. Light was provided for 24 h on the first day and then reduced to 18 h per day from the second day until the end of the experiment. The room temperature was also set as follows: 33 °C on day one, then gradually decreased to 24 °C from day 18 to the end of the feeding trial. Each cage was provided with a feeding trough and automated water system with two drinking nipples. The birds had *ad libitum* access to drinking water and feed. Bird observations were done on a regular basis to monitor health status, feed and water supplies, and environmental conditions. The experiment was ended at 35 days of age.

#### Sample collection and assessment

Feeds and birds were weighed at a cage basis on 0, 10, 24, and 35 days of age to determine the FI, WG, and feed conversion ratio (FCR). In addition, when there was a death, the dead bird, the remaining birds and feeds were weighed to correct FI and FCR. The visceral organs (small intestine, liver, gizzard and proventriculus, bursa and spleen) were collected from the birds killed on day 10 and 24 weighed and recorded. The relative weight of these organs was calculated as g/100g of body weight. To determine apparent nutrient digestibility in the ileum, diets and digesta samples were collected in plastic bags directly from the feed mill after pelleting and cooling, ground and kept in plastic containers for nutrient analysis. On the 24th day of the trial, three birds were randomly selected from each replicate, weighed and euthanized by cervical dislocation to collect digesta samples. The entire gastrointestinal tract (GIT) was exposed and the ileum, which is the small intestinal portion extending from the vitelline (Merkel's) diverticulum to a point about 40 mm above the ileo-caecal junction, was cut and its contents were then flushed into a plastic container, frozen and freeze dried. Lyophilized ileal digesta were then ground using a coffee grinder and kept in air-tight containers at 4°C for further analysis.

To determine the carcass yield, on day 35, two chickens were chosen from each cage, weighed, electrically stunned and euthanized using the cervical dislocation technique. The bird was scalded for 30 sec in hot water (approximately 63°C), and their feathers were plucked out in a batch defeatherer after which the carcass was eviscerated. The entire breast meat (pectoralis minor and pectoralis major), wings, thighs, and drumsticks (bones included) were separated from the carcass and their weight was recorded. The weight of these carcass parts was expressed as g/kg of bird live weight. The remaining carcass parts and feeds were disposed of according to the local regulations.

Table 1. Ingredient composition	of dietary tre	eatments (g/kç	1) <sup>1)</sup>									
		Star	ter			Gro	wer			Finis	sher	
ingreatent composition (g/Kg) -	Control	4% CSM	5% CSM	6% CSM	Control	8% CSM	10% CSM	12% CSM	Control	12% CSM	15% CSM	18% CSM
Sorghum	110.0	0.06	0.06	80.0	110.0	97.0	97.2	102.3	200.0	163.2	200.0	150.0
Wheat	489.7	538.1	523.6	534.7	489.7	551.1	551.1	551.1	484.8	506.5	524.1	577.6
SBM	309.6	235.9	240.0	228.2	309.6	186.0	166.5	138.0	227.0	120.4	30.2	0.0
Canola meal	20.0	20.0	20.0	20.0	10.0	3.0	1.5	1.3	10.0	8.9	5.0	3.3
CSM	0.0	40.0	50.0	60.0	0.0	80.0	100.0	120.0	0.0	120.0	150.0	180.0
Meat & bone meal	20.0	20.0	20.0	20.0	17.0	7.5	7.5	7.5	15.0	10.5	10.0	0.0
Canola oil	22.3	21.9	21.8	22.1	32.0	33.0	32.8	33.0	39.5	37.8	37.0	38.1
Limestone	11.2	11.3	11.4	11.5	10.6	11.9	11.9	12.0	9.8	10.7	11.0	12.2
Dical Phos	4.2	4.2	4.2	4.2	2.8	5.1	5.1	5.2	1.4	2.0	2.6	4.7
Axtra® PHY <sup>2)</sup>	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1
Salt	1.2	1.1	1.1	1.1	1.0	1.1	1.1	1.4	1.1	1.1	1.1	1.2
Na bicarb	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0
Marker (TIO <sub>2</sub> )	0.0	0.0	0.0	0.0	5.0	5.0	5.0	5.0	0.0	0.0	0.0	0.0
UNE VM premix <sup>3)</sup>	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0
Choline CI 70%	0.6	0.6	0.6	0.6	0.5	0.5	0.5	0.5	0.5	0.4	0.5	0.5
L-Lysine	2.4	4.5	4.7	5.0	3.1	5.6	5.8	6.8	3.4	5.4	8.3	9.5
DL-Methionine	3.1	3.7	3.8	3.9	2.7	3.8	4.0	4.3	2.4	3.6	4.5	4.9
L-Threonine	1.6	3.1	3.0	2.7	1.2	2.8	2.9	3.3	0.9	2.5	3.7	4.3
L-Arginine	0.0	1.5	1.7	1.9	0.0	2.5	3.0	3.9	0.0	2.9	5.7	6.9
L-Isoleucine	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0	1.1	1.4
L-Tryptophan	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.7
Total	1,000	1,000	1,000	1,000	1,000	1,000	1,000	1,000	1,000	1,000	1,000	1,000
<sup>1)</sup> Each diet in this table was divided into	two diets and s	supplemented wit	h or without mic	robial enzymes (,	Axtra XB; at 0.1c	J/kg).						
<sup>2)</sup> Axtra Phytase: unique bacterial phytas	ie enzyme deve	eloped by Danisc	o Animal Nutritic	on, UK.								
$^{3h}$ Vitamin-mineral concentrate supplied $r$	per kilogram of	diet, Retinol, 12,	000 IU; Choleca	lciferol, 5,000 IU;	Tocopheryl acet	ate, 75 mg; Mer	nadione, 3 mg; Tl	niamine, 3 mg; Ri	boflavin, 8 mg; l	Viacin, 55 mg; P	antothenate, 13	mg; Pyridoxine,
5 mg; Folate, 2 mg; Cyanocobalamin, 1	6 µg; Biotin, 20	00 µg, Cereal-ba	sed carrier, 149	mg; Cu (sulphatŧ	e), 16 mg; Fe (su	Ilphate), 40 mg;	l (iodide), 1.25 r	ng; Se (sodium s	elenite), 0.3 mg;	Mn (sulphate ai	nd oxide), 120 m	ig, Zn (sulphate

and oxide), 100 mg; Cereal-based carrier, 128 mg; Mineral oil, 3.75 mg.

CSM, cottonseed meal; SBM, soybean meal.

Table 2. Nutrient composition of	f dietary treat	tments (g/kg)										
Variation of the second s		Sta	rter			Gro	wer			Finis	sher	
Nutrient composition (g/kg)	Control	4% CSM	5% CSM	6% CSM	Control	8% CSM	10% CSM	12% CSM	Control	12% CSM	15% CSM	18% CSM
ME (MJ/kg)	12.6	12.6	12.6	12.6	13.0	13.0	13.0	13.0	13.4	13.4	13.4	13.4
CP	230.0	230.0	230.0	230.0	215.0	215.0	215.0	215.0	195.0	195.0	195.0	195.0
Crude fat	42.5	42.6	41.8	42.0	51.7	51.7	51.7	51.9	59.9	58.5	57.0	56.8
Crude fibre	26.3	26.7	26.9	27.1	25.3	26.0	26.2	26.4	24.5	26.4	26.0	26.2
Arginine	14.3	13.9	13.8	13.7	13.2	12.3	12.3	12.3	11.6	10.9	10.9	10.9
Lysine	12.8	12.8	12.9	12.8	12.5	11.5	11.5	11.5	11.4	10.2	10.2	10.2
Methionine	6.2	6.5	6.5	6.5	5.6	6.2	6.1	6.3	5.1	5.6	6.0	6.1
Met + cystine	9.5	9.5	9.5	9.5	8.7	8.7	8.7	8.7	8.0	8.0	8.0	8.0
Tryptophan	2.8	2.6	2.4	2.3	2.6	1.8	1.9	1.7	2.3	1.6	1.6	1.6
Isoleucine	9.6	0.0	8.9	8.8	9.1	7.8	7.8	7.8	8.2	7.2	7.0	7.0
Threonine	8.6	8.8	0.0	8.6	7.7	7.7	7.7	7.7	6.8	6.8	6.8	6.8
Valine	10.8	10.4	10.2	10.0	10.1	9.4	9.2	8.9	9.2	8.7	7.8	7.8
Calcium	9.6	9.6	9.6	9.6	8.7	8.7	8.7	8.7	7.8	7.8	7.8	7.8
Available P	4.8	4.8	4.8	4.8	4.4	4.4	4.4	4.4	3.9	3.9	3.9	3.9
Sodium	1.9	1.9	1.9	1.9	1.8	1.8	1.8	1.9	1.8	1.8	1.8	1.8
Chlorine	1.9	2.2	2.3	2.4	1.9	2.3	2.4	2.8	2.0	2.3	2.9	2.3
Choline	17.0	17.0	17.0	17.0	16.0	16.0	16.0	16.0	15.0	15.0	15.0	15.0
CSM, cottonseed meal; CP, crude prot∈	ein.											

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Nutrient	Value
ME (MJ/kg)	9.7
Crude protein (%)	45.1
Crude fat (%)	1.7
Crude fibre (%)	8.6
Ash (%)	6.8
Indispensable amino acids (g/kg)	
Arginine	46.7
Lysine	21
Methionine	6.9
Threonine	14.6
Tryptophan	5
Valine	21.5
Histidine	13
Leucine	26.7
Isoleucine	14.4
Phenylalanine	24.8
Minerals (g/kg)	
Calcium	1.8
Total phosphorus	9.1
Sodium	0.45
Potassium	1.3

#### Nutrients analyses and digestibility calculations

The concentration of  $TiO_2$  in the diets and ileal digesta was measured according to the colorimetric procedure of Short et al. [14], using about 0.1 g of digesta and 0.2 g of the diet samples. Ground diet samples were oven-dried at 105 °C for 24 h to determine DM content of experimental diets using a forced air convection oven (Qualtex Universal Series 2000, Watson Victor Ltd, Perth, Australia). The DM content of the ileal digesta samples was obtained directly after freeze-drying (Martin Christ freeze dryer, Germany) at  $-50\,^\circ$ C for at least 72 h. The nitrogen content of ileal digesta and diet samples was measured using the Dumas combustion technique, following the method described by Sweeney [15], through direct electronic reading on a LECO® FP-2000 automatic nitrogen analyzer (Leco Corporation, St. Joseph, MI, USA). The obtained nitrogen results were then converted to CP content, using a factor of 6.25. Around 0.5 g of finely ground (0.5 mm) diet and ileal digesta samples were weighed in metallic crucibles to determine gross energy (GE), using an IKA®-WERKE bomb calorimeter (C7000, GMBH & Co., Staufen, Germany), and the GE value of samples measured as MJ/kg, was read from the calorimeter digital system. The method developed by McCleary et al. [16] was followed to assess the total starch content of diets and ileal digesta. The Megazyme total starch assay kitAA/AMG (Megazyme International Ireland Limited, Bray, Co. Wicklow, Ireland) was used for total starch analysis. Absorbance was read at 510 nm on a spectrophotometer (Varian Cary 50 Bio UV - Visible Spectrophotometer Mulgrave, Victoria, Australia) against milli-Q water.

The index method proposed by Moughan and Marlies Leenaars [17] and reviewed by Kong and Adeola [18] with titanium dioxide as an index compound was used to determine nutrient digestibility. The following equation was used to calculate apparent ileal digestibility of nutrients:

$$AIDN = 1 - \frac{Ti_{Diet} \times N_{Digesta}}{TiO_{2 Digesta} \times N_{Diet}}$$

where, AIDN is the coefficient of apparent ileal digestibility of nutrients,  $N_{Digesta}$  is the nutrient concentration in digesta (g/kg), TiO<sub>2 Digesta</sub> is the titanium concentration in digesta (g/kg),  $N_{Diet}$  is nutrient concentration in diets (g/kg) and TiO<sub>2 Diet</sub> is the titanium concentration in feeds (g/kg).

#### **Statistical analysis**

Broiler chicken's growth, visceral organs development, meat yield and nutrient digestibility data were statistically analyzed according to  $4 \times 2$  factorial arrangement. The obtained data were subjected to analysis of variance (ANOVA) in the procedures of general linear model of Minitab statistical software, version 17 [19]. Fisher's least significant difference (LSD) method for multiple comparisons was used to separate significant differences between mean values at  $p \le 0.05$  level of probability.

#### **Animal ethics**

The experimental design and methodology were approved by the Animal Ethics Committee of the University of New England, Australia under approval number AEC15-084. All husbandry and management procedures used were in accordance with The Code of Practice for the Use of Animals for Scientific Purposes was issued by the Australian Bureau of Animal Health.

## RESULTS

#### Effect of CSM and microbial enzyme (Axtra XB) on gross responses

The FI, WG, and FCR in response to the four CSM levels in diets supplemented with or without Axtra XB enzyme are summarized in Table 4. In starter broilers (1–10 days), there was an interaction between the enzyme and CSM on FI (p < 0.028) and WG (p < 0.036). In the starter and grower phases (1–24 days) and entire cycle (1–35 days), FI did not vary (p > 0.05) between dietary treatments. The FCR was improved (p < 0.023) by CSM in the starter phase mostly in lower and medium CSM diets, and in the grower (p < 0.045) phase, particularly in chickens fed with the maximal level of CSM. Supplementation with Axtra XB enzyme increased WG in the grower (p < 0.037) and finisher (p < 0.017) phases. Enzyme addition significantly improved FCR in the grower (p < 0.008) and finisher (p < 0.017) phases. There was no interaction between enzyme and CSM for gross responses over 1–24 and 1–35 days.

#### Effect of CSM and Axtra XB enzyme on visceral organ weights

Incorporation of CSM in diets decreased the weight of the small intestine (p < 0.008) and the combined weight of gizzard plus proventriculus (p < 0.048) in starter chicks (Table 5). In grower birds, CSM contributed increased (p < 0.029) the weight of gizzard plus proventriculus with the highest weight recorded in the birds fed medium-CSM diet. The test ingredient decreased (p < 0.013) the weight of the small intestine in growing chickens except for those fed the medium level of CSM. There was no treatment effect (p > 0.05) on the weight of the other visceral organs at 10 or 24 days of age.

#### Effect of CSM and Axtra XB enzyme on meat yield

There was no interaction between CSM levels and Axtra XB enzyme on the weight of carcass parts. The enzyme supplement significantly increased (p < 0.007) the absolute weight of breast meat (Table 6). Diets containing CSM also resulted in an increase in thigh weight both in absolute (p < 0.024)

Tathmatic         1-0 days         1-2 days         1-3 days         1-3 days         1-3 days           Tathmatic         1-0 days         1-0 days         1-2 days         1-3 days         1-3 days         1-3 days           CSMIerei         Exyme (mg/kg)         H (gh/m)         W (g)         FCR         1-3 days         1-3 days         1-3 days           CSMIerei         Exyme (mg/kg)         H (gh/m)         W (g)         FCR         1-3 days         1-3 days         1-3 days           100         345 °         323 °         1.08         1/7 (%)         1/7 (%)         1/7 (%)         1/2 (%)         3/4 (%)         1/3 (%)         1/3 (%)           100         367 °         323 °         1.08         1/7 (%)         1/7 (%)         1/7 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)         1/3 (%)	Table 4. Effect of C	SM level with or withou	ut enzyme blend c	on FI, WG, and Ft	CR of broilers						
CSM level         Expression(Med/C)         F (q)trict)         VG (q)         F (q)trict)         F (q)trirct)         F (q)trict)         F (q)trict)<	Trea	tments		1–10 days			1–24 days			1–35 days	
Control         0         34.5*         32.3*         1.08         1,784         1,25         3,3470         2.5895         1.35           100         367.6*         32.3*         1.07         1,754,7         1,4726         1.20         3,4843         2.5972         1.36           100         367.6*         32.9*         1.07         1,764,7         1,4726         1.20         3,4843         2.5972         21440         1.35           Medlum <sup>1</sup> 0         347,7*         341.6*         1.02         1,7764         1,4930         1.21         3,6164         1.36         1.36           Medlum <sup>1</sup> 0         347.7*         341.6*         1.05         1,7764         1,4930         1.21         3,5164         1.36           Medlum <sup>1</sup> 0         347.7*         341.6*         1.05         1,4935         1.97         1.36         1.36           Medlum <sup>1</sup> 0         357.7*         341.6*         1.05         1,4935         1.97         1.36         1.36           Medlum <sup>1</sup> 100         356.6         1.06*         1,776.6         1.4930         1.27         2.574.4         1.36           Medlum         356.6         33	CSM level	Enzyme (mg/kg)	FI (g/bird)	WG (g)	FCR	FI (g/bird)	WG (g)	FCR	FI (g/bird)	WG (g)	FCR
10         367 b         342 b         107         1,74 T         1,47 b         120         3,46 5         2,61 40         135           Low <sup>1</sup> 0         360.7         330.7*         105         1,75 A         1,45 A         1,2 C         3,46 B         2,57 Z         1,36           Medum <sup>1</sup> 0         340.7         330.7*         100         1,7 B         3,46 B         1,66 B         1,46 A         1,2 C         3,46 B         2,57 Z         1,36           Hgh <sup>1</sup> 0         340.7*         330.7*         100         346.9*         1,03         1,46 A         1,2 C         3,44 B         2,57 Z         1,36           Hgh <sup>1</sup> 0         347.9*         330.7*         1,01         1,76 A         1,2 C         3,44 B         2,57 Z         1,36           Hgh <sup>1</sup> 100         357.9*         1,00         1,76 A         1,48 A         1,2 C         3,49 B         2,57 Z         1,3 B           Ken         215         1,80         1,60 A         1,76 B         1,48 A         1,2 C         1,3 B         1,3 B           Ken         216         1,7 B         1,4 B         1,7 B         1,4 B         1,2 C         1,3 B	Control	0	345.3ª	323.7 <sup>b</sup>	1.08	1,769.9	1,438.4	1.25	3,547.0	2,589.5	1.37
		100	367.8ª	342.9ª	1.07	1,754.7	1,472.6	1.20	3,485.2	2,614.0	1.35
	Low <sup>1)</sup>	0	350.2 <sup>b</sup>	332.9 <sup>ab</sup>	1.05	1,786.8	1,450.4	1.22	3,484.6	2,557.2	1.36
		100	346.5 <sup>b</sup>	338.7 <sup>a</sup>	1.02	1,798.4	1,525.8	1.19	3,491.8	2,609.5	1.34
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	Medium <sup>2)</sup>	0	347.1 <sup>b</sup>	333.1 <sup>ab</sup>	1.04	1,771.9	1,454.1	1.23	3,516.1	2,577.8	1.36
High <sup>1</sup> 0 $3577^{\circ}$ $3416^{\circ}$ $106$ $18005$ $14835$ $18916$ $25732$ $136$ $1360$ $25732$ $136$ $1360$ $25732$ $136$ $1361$ $131$ Rem $100$ $345.6^{\circ}$ $332.0^{\circ}$ $104$ $17686$ $14980$ $120$ $34433$ $26607$ $131$ Rem $215$ $180$ $0.01$ $971$ $911$ $0.01$ $1360$ $100$ Main effects $215$ $180$ $0.01$ $971$ $911$ $0.01$ $136$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$ $1360$		100	354.0 <sup>ab</sup>	342.4ª	1.03	1,750.4	1,493.0	1.21	3,539.0	2,656.4	1.33
100         345 %         332 0 %         104         178 %         149 %         120         349 %         26607         131           SEM         2.15         1.80         0.01         9.71         9.11         0.01         1390         1960         0.01           Main effects         2.15         1.80         0.01         9.71         9.11         0.01         1390         1960         0.01           Main effects         2.15         1.80         0.01         9.71         9.11         0.01         1390         1960         0.01           Main effects         2.56 33.3         1.08*         1.784 5         1.485.5         122*         3.516         2.6017         1.361           Control         335.8         1.04*         1.762.3         1.486.1         1.20*         3.474.1         1.361           Uow         351.8         337.8         1.04*         1.761.1         1.495.6         3.474.1         2.617.1         1.361           High         1.00         3.557.6         2.514.4         1.364         1.364           Fazyme (mg/kg)         0         3.557.6         2.617.1         2.667.7         1.364           High         1.00	High <sup>3)</sup>	0	357.7 <sup>ab</sup>	341.6ª	1.05	1,800.5	1,493.5	1.97	3,498.9	2,573.2	1.36
		100	345.8 <sup>b</sup>	332.0 <sup>ab</sup>	1.04	1,768.6	1,498.0	1.20	3,449.3	2,660.7	1.31
Main effects       Main effects       1,784.5       1,455.5       1,22"       3,516.1       2,601.7       1,361         CSM levels       335.6       333.3       1,08"       1,784.5       1,455.5       1,22"       3,516.1       2,601.7       1,361         Control       366.6       333.3       1,08"       1,784.5       1,488.1       1,20"       3,488.2       2,637.4       1,362         Low       351.8       1,04"       1,762.3       1,488.1       1,20"       3,488.2       2,637.4       1,362         Medium       351.6       337.8       1,04"       1,762.3       1,488.1       1,20"       3,488.2       2,617.1       1,361         Medium       351.6       337.8       1,04"       1,762.6       1,473.5       1,20"       3,474.1       2,617.0       1,349         Fuzyme (mg/g)       0       350.1       1,04"       1,760.4"       1,20"       3,474.1       2,617.0       1,364"         Fuzyme (mg/g)       0       355.6       1.04"       1,760.4"       1,22"       3,511.6       2,617.4"       1,364"         Fuzyme (mg/g)       0       335.5       1.04"       1,752.6"       1,20"       3,491.3       2,617.4"       1,364"	SEM		2.15	1.80	0.01	9.71	9.11	0.01	13.90	19.60	0.01
CSM levels Control 356.6 333.3 1.08° 1,784.5 1,455.5 1.22° 3.516.1 2.601.7 1.361 Low 348.2 2.583.4 1.361 Low 348.3 335.8 1.04° 1,762.3 1,488.1 1.20° 3,488.2 2.583.4 1.352 Medium 350.6 337.8 1.04° 1,772.3 1,488.1 1.20° 3,488.2 2.583.4 1.349 High 1.22° 3,51.6 2,617.1 1.349 High 1.22° 3,51.6 2,617.1 1.349 Firstme (mg/g) 351.8 336.8 1.04° 1,761.1 1,495.8 1.20° 3,474.1 2,617.0 1.341 Firstme (mg/g) 353.5 339.0 1.04° 1,761.1 1,495.8 1.20° 3,474.1 2,617.0 1.341 Firstme (mg/g) 333.5 330.0 1.04° 1,782.3 1,456.4 1.20° 3,474.1 2,617.0 1.349 Firstme (mg/g) 333.5 339.0 1.04° 1,782.3 1,456.4 1.20° 3,491.3 2,635.1° 1.364° Significance 100 333.5 339.0 1.04 1,788.0 1,525.8° 1.20° 3,491.3 2,635.1° 1.346° Significance 2.574.4° 1.22° 3,511.6 2,574.4° 1.346° Significance 2.574.4° 1.22° 3,511.6 2,574.4° 1.346° Significance 2.574.4° 1.22° 3,511.6 2,574.4° 1.346° Significance 2.574.4° 1.20° 3,511.6 2,574.4° 1.346° Significance 2.574.8° 2,574.4° 1.355.8° 0.036 0.037 0.036 0.077 0.056 0.077 CSM × Enzyme 2.028 0.023 0.036 0.001 0.573 0.130 0.659 0.076	Main effects										
	<b>CSM</b> levels										
	Control		356.6	333.3	1.08 <sup>a</sup>	1,784.5	1,455.5	1.22 <sup>a</sup>	3,516.1	2,601.7	1.361
Medium         350.6         337.8         1.04 <sup>b</sup> 1,792.6         1,473.5         1.22 <sup>b</sup> 3,527.6         2,617.1         1.349           High         351.8         336.8         1.04 <sup>ab</sup> 1,761.1         1,495.8         1.20 <sup>b</sup> 3,474.1         2,617.0         1.361           Fuzyme (mg/kg)         0         351.8         336.8         1.04 <sup>ab</sup> 1,761.1         1,495.8         1.20 <sup>b</sup> 3,474.1         2,617.0         1.361           Fuzyme (mg/kg)         0         350.1         332.8         1.04 <sup>ab</sup> 1,768.0         1,761.4         1.20 <sup>b</sup> 3,474.1         2,617.0         1.364 <sup>a</sup> Inzyme (mg/kg)         0         353.5         339.0         1.04 <sup>ab</sup> 1,768.0         1,526.8 <sup>a</sup> 1.20 <sup>b</sup> 3,491.3         2,635.1 <sup>a</sup> 1.364 <sup>a</sup> Significance         100         353.5         339.0         1.04 <sup>b</sup> 1,768.0         0,613         0,491.3         2,635.1 <sup>a</sup> 1.364 <sup>b</sup> Significance         100         3537         0.802         0.023         0.613         0.402         0,402         0.787         0.365         0.374 <sup>b</sup> 1.34 <sup>b</sup> Significance         1.04 <sup>b</sup>	Low		348.3	335.8	1.04 <sup>b</sup>	1,762.3	1,488.1	1.20 <sup>ab</sup>	3,488.2	2,583.4	1.352
High $351.8$ $336.8$ $1.04^{46}$ $1.761.1$ $1.495.8$ $1.20^{5}$ $3.474.1$ $2.617.0$ $1.361$ Enzyme (mg/kg)0 $350.1$ $332.8$ $1.04^{46}$ $1.761.1$ $1.495.8$ $1.20^{5}$ $3.474.1$ $2.617.0$ $1.361^{4}$ 0 $350.1$ $332.8$ $1.06$ $1.782.3$ $1.450.4^{5}$ $1.22$ $3.511.6$ $2.574.4^{5}$ $1.364^{4}$ 100 $353.5$ $339.0$ $1.04$ $1.768.0$ $1.525.8^{4}$ $1.20$ $3.491.3$ $2.635.1^{4}$ $1.34^{4}$ Significance $0.022$ $0.023$ $0.613$ $0.402$ $0.045$ $0.787$ $0.806$ $0.527$ Significance $0.402$ $0.045$ $0.045$ $0.787$ $0.806$ $0.527$ Significance $0.402$ $0.045$ $0.787$ $0.806$ $0.527$ Significance $0.627$ $0.036$ $0.036$ $0.037$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$ $0.036$	Medium		350.6	337.8	1.04 <sup>b</sup>	1,792.6	1,473.5	1.22 <sup>ab</sup>	3,527.6	2,617.1	1.349
Enzyme (mg/kg)       0       350.1       332.8       1.06       1,782.3       1,450.4 <sup>b</sup> 1.22       3,511.6       2,574.4 <sup>b</sup> 1.364 <sup>a</sup> 100       353.5       339.0       1.04       1,782.3       1,450.4 <sup>b</sup> 1.20       3,491.3       2,635.1 <sup>a</sup> 1.34 <sup>a</sup> Significance       100       353.5       339.0       1.04       1,768.0       1,525.8 <sup>a</sup> 1.20       3,491.3       2,635.1 <sup>a</sup> 1.34 <sup>b</sup> Significance       0.053       0.802       0.023       0.013       0.402       0.787       0.806       0.527         CSM       0.537       0.802       0.215       0.486       0.037       0.086       0.627       0.366       0.017         Enzyme       0.028       0.028       0.036       0.613       0.677       0.036       0.017	High		351.8	336.8	1.04 <sup>ab</sup>	1,761.1	1,495.8	1.20 <sup>b</sup>	3,474.1	2,617.0	1.361
0         350.1         32.8         1.06         1.782.3         1,450.4 <sup>b</sup> 1.22         3,511.6         2,574.4 <sup>b</sup> 1.364 <sup>a</sup> 100         353.5         339.0         1.04         1,768.0         1,525.8 <sup>a</sup> 120         3,491.3         2,635.1 <sup>a</sup> 1.334 <sup>b</sup> Significance           0.04         1,525.8 <sup>a</sup> 120         3,491.3         2,635.1 <sup>a</sup> 1.334 <sup>b</sup> Significance           0.637         0.802         0.023         0.613         0.402         0.787         0.806         0.527           CSM          0.400         0.072         0.215         0.486         0.037         0.036         0.627         0.036         0.017           CSM × Enzyme          0.028         0.036         0.801         0.890         0.573         0.036         0.017	Enzyme (mg/kg)										
100         353.5         339.0         1.04         1,768.0         1,525.8 <sup>a</sup> 1.20         3,491.3         2,635.1 <sup>a</sup> 1.334 <sup>b</sup> Significance                1.34           Significance               2,635.1 <sup>a</sup> 1.34 <sup>b</sup> Significance                 1.34 <sup>b</sup> Significance		0	350.1	332.8	1.06	1,782.3	1,450.4 <sup>b</sup>	1.22	3,511.6	2,574.4 <sup>b</sup>	1.364ª
Significance CSM 0.537 0.802 0.023 0.613 0.402 0.045 0.787 0.806 0.527 Enzyme 0.400 0.072 0.215 0.486 0.037 0.008 0.627 0.036 0.017 CSM × Enzyme 0.028 0.028 0.036 0.801 0.890 0.573 0.130 0.859 0.855		100	353.5	339.0	1.04	1,768.0	$1,525.8^{a}$	1.20	3,491.3	2,635.1 <sup>ª</sup>	1.334 <sup>b</sup>
CSM         0.537         0.802         0.023         0.613         0.402         0.045         0.787         0.806         0.527           Enzyme         0.400         0.072         0.215         0.486         0.037         0.008         0.627         0.036         0.017           CSM × Enzyme         0.028         0.036         0.801         0.890         0.573         0.130         0.859         0.855	Significance										
Enzyme         0.400         0.072         0.215         0.486         0.037         0.008         0.627         0.036         0.017           CSM × Enzyme         0.028         0.036         0.801         0.890         0.573         0.130         0.855	CSM		0.537	0.802	0.023	0.613	0.402	0.045	0.787	0.806	0.527
CSM × Enzyme 0.028 0.028 0.036 0.801 0.890 0.573 0.130 0.859 0.855	Enzyme		0.400	0.072	0.215	0.486	0.037	0.008	0.627	0.036	0.017
	CSM × Enzyme		0.028	0.028	0.036	0.801	0.890	0.573	0.130	0.859	0.855

 $^{ab}$  Means with different superscripts within the columns are different ( $\rho < 0.05$ ).

<sup>3</sup>CSM high level: 6%, 12%, and 18% of complete diet introduced in starter, grower, and finisher phases of growth, respectively. <sup>2</sup>CSM medium level: 5%, 10%, and 15% of complete diet introduced in the starter, grower, and finisher period, respectively. <sup>1</sup>CSM low level: 4%, 8%, and 12% of complete diet introduced in the starter, grower, and finisher period respectively. CSM, cottonseed meal; FI, feed intake; WG, weight gain; FCR, feed conversion ratio; SEM, standard error of means.

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Table 5. Effect CSM I	evel with or without $\epsilon$	enzyme blend	l on visceral	organ weigh	t (g/100 g B\	V) at 10 and 3	24 days of aç	Je					
Treatr	nents			At 10	days					At 24	days		
<b>CSM</b> levels	Enzyme (mg/kg)	G + D	Pan	SI	Liver	Spleen	Bursa	G + P	Pan	ิง	Liver	Spleen	Bursa
Control	0	4.70	0.37	7.61	3.71	60.0	0.18	2.00	0.20	4.73	2.50	0.07	0.19
	100	4.71	0.36	7.19	3.37	0.09	0.19	2.21	0.21	3.81	2.33	0.08	0.17
Low <sup>1)</sup>	0	3.93	0.33	6.36	3.27	0.10	0.21	2.23	0.20	4.30	2.31	0.07	0.17
	100	4.52	0.37	6.93	3.24	0.08	0.15	2.03	0.18	4.30	2.38	0.07	0.16
Medium <sup>2)</sup>	0	4.32	0.34	6.33	3.35	0.09	0.17	2.60	0.22	3.91	2.33	0.07	0.16
	100	4.57	0.29	6.24	3.38	0.08	0.15	2.42	0.19	3.95	2.30	0.08	0.16
High <sup>3)</sup>	0	4.28	0.33	6.89	3.19	0.08	0.16	2.19	0.19	3.75	2.20	0.06	0.18
	100	4.06	0.33	6.52	3.81	0.08	0.17	2.25	0.18	3.44	2.22	0.07	0.16
SEM		0.08	0.01	0.12	0.53	0.00	0.01	0.05	0.00	0.09	0.04	00.0	0.01
Main effects													
<b>CSM</b> levels													
Control		4.70 <sup>a</sup>	0.37	7.40 <sup>a</sup>	3.54	0.09	0.18	2.11 <sup>b</sup>	0.20	4.27 <sup>a</sup>	2.4	0.08	0.18
Low		4.23 <sup>b</sup>	0.35	6.64 <sup>ab</sup>	3.23	0.08	0.18	2.13 <sup>ab</sup>	0.19	$4.30^{a}$	2.35	0.07	0.16
Medium		4.44 <sup>ab</sup>	0.32	6.28 <sup>b</sup>	3.30	0.08	0.16	2.50 <sup>a</sup>	0.20	3.93 <sup>ab</sup>	2.31	0.08	0.16
High		4.17 <sup>b</sup>	0.33	6.70 <sup>ab</sup>	3.28	0.08	0.17	$2.22^{ab}$	0.19	3.60 <sup>b</sup>	2.21	0.07	0.16
Enzyme (mg/kg)													
	0	4.31	0.34	6.80	3.38	0.09	0.18	2.25	0.20	4.17	2.32	0.07	0.18
	100	4.47	0.34	6.72	3.30	0.08	0.17	2.23	0.19	3.87	2.31	0.08	0.17
Significance													
CSM		0.048	0.326	0.008	0.157	0.441	0.588	0.029	0.335	0.013	0.447	0.352	060.0
Enzyme		0.312	0.827	0.725	0.402	0.080	0.183	0.797	0.111	0.078	0.848	0.134	0.144
CSM × Enzyme		0.294	0.481	0.362	0.369	0.773	0.165	0.420	0.270	0.166	0.869	0.387	0.313
Values are means of 6 re <sup>ab</sup> Means with different su	plicates (10 birds per repli perscripts within the colurr	cate). ins are different	(p < 0.05).										

<sup>&</sup>lt;sup>2</sup>/CSM medium level: 5%, 10%, and 15% of complete diet introduced in the starter, grower, and finisher period, respectively. <sup>1</sup>/CSM low level: 4%, 8%, and 12% of complete diet introduced in the starter, grower, and finisher period, respectively.

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<sup>&</sup>lt;sup>3</sup>/CSM high level: 6%, 12%, and 18% of complete diet introduced in starter, grower, and finisher phases of growth, respectively. CSM, cottonseed meal; G + P, gizzard and proventriculus; Pan, pancreas; SI, small intestine; SEM, standard error of means.

Treat	ment		A	osolute weig	ht			Relative to	live weight	
CSM levels	Enzyme (mg/kg)	Dressing (%)	Breast	Thighs	Wings	Drum- sticks	Breast	Thighs	Wings	Drum- sticks
Control	0	76.0	537.9	258.2	198.3	236.5	202.2	97.2	74.8	88.9
	100	76.8	562.7	260.6	199.9	242.2	206.9	95.8	73.5	89.0
Low2	0	76.4	528.5	258.8	187.4	233.4	202.4	99.0	71.8	89.3
	100	76.9	587.1	278.7	208.9	257.0	210.4	99.8	74.8	91.9
Medium3	0	76.5	538.7	275.9	186.9	241.7	203.2	103.9	70.5	91.1
	100	77.2	552.1	290.9	189.1	250.8	203.2	103.3	71.7	90.1
High4	0	76.8	537.9	283.7	193.5	263.4	195.2	103.2	70.3	95.7
	100	77.0	562.7	283.8	198.8	250.1	203.7	102.3	71.7	90.3
SEM		0.18	5.58	3.41	2.20	2.87	0.15	0.09	0.05	0.06
Main effects										
CSM Levels										
Control		76.4	550.3	259.4b	199.3	239.3	204.6	96.5 <sup>b</sup>	74.1	90
Low		76.6	557.8	268.8ab	198.2	245.2	206.4	99.4 <sup>ab</sup>	73.3	90.6
Medium		76.8	545.4	283.4ab	188	246.2	203.2	103.6ª	71.1	90.6
High		76.9	550.3	283.7a	195.9	256.8	199.4	102.7ª	71.0	93.0
Enzyme (mg/kg)										
	0	76.4	535.8 <sup>b</sup>	269.1	191.7	243.7	200.8 <sup>b</sup>	100.8	71.8	91.3
	100	76.9	566.2ª	278.5	199.1	250.0	206.0 <sup>a</sup>	100.3	72.9	91.3
Significance										
CSM		0.756	0.875	0.024	0.234	0.165	0.404	0.015	0.068	0.122
Enzyme		0.126	0.007	0.149	0.084	0.252	0.083	0.745	0.271	0.440
CSM× Enzyme		0.944	0.481	0.630	0.289	0.136	0.722	0.975	0.476	0.125

Table 6. Effect of CSM level with or without enzyme blend on meat yield1 in absolute (g of carcass wt) and relative weight (g/100 of carcass wt) at 35 days of age

Values are means of 6 replicates (10 birds per replicate).

<sup>a,b</sup>Means with different superscripts within the columns are different (p < 0.05).

<sup>1)</sup>Meat yield refers to breast, thighs, wings, and drumsticks.

<sup>2)</sup>CSM low level: 4%, 8%, and 12% of complete diet introduced in the starter, grower, and finisher period, respectively.

<sup>3</sup>CSM medium level: 5%, 10%, and 15% of complete diet introduced in the starter, grower, and finisher period, respectively.

<sup>4)</sup>CSM high level: 6%, 12%, and 18% of complete diet introduced in starter, grower, and finisher phases of growth, respectively

SEM, standard error of means; CSM, cottonseed meal.

and relative (p < 0.015) terms. Apart from these carcass parts, there was no treatment effect (p > 0.05) observed in other body parts (meat) yield.

## Apparent ileal digestibility of nutrients

The enzyme increased apparent ileal digestibility of CP (p < 0.030), gross energy (p < 0.014) and starch (p < 0.008) in the ileum (Table 7). However, the apparent ileal starch digestibility was reduced (p < 0.010) by increasing CSM level in the diet, with the lowest starch digestibility noted in chickens fed the diets containing a high level of CSM. There was no effect (p > 0.05) of CSM on CP or gross energy digestibility.

## DISCUSSION

## **Gross responses**

Apart from the CSM-enzyme interaction observed on FI, feed consumption did not differ be-

CSM levels	Enzyme	Crude protein	Gross energy	Starch
Control	0	0.81	0.70	0.95
	100	0.81	0.72	0.98
Low <sup>1)</sup>	0	0.78	0.66	0.95
	100	0.82	0.71	0.97
Medium <sup>2)</sup>	0	0.80	0.70	0.95
	100	0.80	0.71	0.96
High <sup>3)</sup>	0	0.79	0.66	0.93
	100	0.82	0.69	0.96
SEM		0.00	0.01	0.00
Main effects				
CSM levels				
Control		0.81	0.70	0.97 <sup>a</sup>
Low		0.80	0.69	0.96 <sup>ab</sup>
Medium		0.80	0.70	0.95 <sup>ab</sup>
High		0.80	0.68	0.94 <sup>b</sup>
Enzyme (mg/kg))				
	0	0.80 <sup>b</sup>	0.68 <sup>b</sup>	0.94 <sup>b</sup>
	100	0.81ª	0.71ª	0.97 <sup>a</sup>
Significance				
CSM		0.729	0.231	0.010
Enzyme		0.030	0.014	0.008
CSM level × Enzyme		0.230	0.664	0.410

Table 7. Apparent Ileal digestibility of nutrients in response to different CSM levels with or without the enzyme blend

Values are means of 6 replicates (10 birds per replicate).

<sup>a,b</sup>Means with different superscripts within the columns are different (p < 0.05).

<sup>1)</sup>CSM low level: 4%, 8%, and 12% of complete diet introduced in the starter, grower, and finisher period, respectively.

<sup>2)</sup>CSM medium level: 5%, 10%, and 15% of complete diet introduced in the starter, grower, and finisher period, respectively. <sup>3)</sup>CSM high level: 6%, 12%, and 18% of complete diet introduced in starter, grower, and finisher phases of growth, respectively. SEM, standard error of the mean; CSM, cottonseed meal.

tween the dietary treatments and the control diet. This can be explained by the fact that, although CSM is high in fibre, the experimental diets were formulated in such a way that all nutrients were similar and balanced to meet the birds' nutrient requirements. This supports the results of a feeding trial conducted by Adeyemo and Longe [20] in which no specific FI pattern in meat chickens fed CSM-containing diets was found, although after the seventh week of their trial, differences between treatments means emerged. The enzyme interacted with CSM on WG response in starting birds and the values of this response were identical between birds fed CSM- and SBM-containing diets both in the growing and finishing phases. CSM can cause growth depression in chicks and this could, among other factors, be due to decreased amino acids digestibility, low lysine content and the presence of gossypol [21]. However, digestibility of amino acids, was improved in the present trial, which can counteract the gossypol effect and growth depression as stated by Henry et al. [22] and Nagalakshmi et al. [23]. These findings agree with the results obtained by Ojewola et al. [2] who reported that the WG of broilers fed CSM-containing diets was similar to that of birds fed SBM diet. However, this was not supported other work which recorded a significant differences in WG when CSM replaced SBM in broiler diets [20]. These authors demonstrated a similarity in gross response values among chickens fed diets in which soybean cake was replaced by cottonseed cake at 75% compared to those fed a control diet. They also stated that a complete replacement of SBM with CSM resulted in depressed growth performance. However, in the present trial, the replacement of SBM with CSM by even up to 90% (18% CSM diet) did not severely affect the gross responses. Enzyme supplementation in grower and finisher diets increased the WG and this may be due to its contribution to the improvement in nutrient digestibility, as previously reported that supplementation wheat-based diets with  $\beta$ -glucanase increased broiler growth [24].

It has been stated that starting chick diets containing high levels of CSM may have a depressing effect on growth coupled with increased feed-to-gain ratios due to the presence of gossypol [2]. Also it was stated that CSM-containing feeds consumed by birds were not converted as effectively as SBM diets due to poor utilisation of amino acids from CSM [20]. However, in the present trial, no statistical difference was observed in finisher chicks because of CSM incorporation in their diets. Most importantly, this proteinaceous ingredient contributed to a reduction in FCR in the starter and grower phases. As mentioned earlier, the enzyme supplement lowered the FCR in grower and finisher birds. These improvements could have resulted from enhanced energy and protein digestibility, initiated by the inclusion of the test enzyme. Feed conversion ratio has been shown to decrease in birds provided with diets containing a  $\beta$ -glucanase and xylanase cocktail [12].

#### Visceral organ weights

In the grower birds, CSM increased the weight of the gizzard and proventriculus. It is well established that the increased content of fibre in a diet can increase the length and size of the GIT [25,26]. Adding indigestible materials, such as wood shavings, can tremendously increase the weight of gizzard [27]. Other studies have linked the fibre content in dietary ingredients, such as CSM with hypertrophy of the GIT, including the gizzard [28,29]. Although, the fibre content of the formulated diets in this feeding trial was similar to the standard recommended by Aviagen [13], but CSM fibre tends to be physically bulky [30]. In the present trial, the activity of the test enzyme on wheatsorghum–CSM diets decreased the small intestinal weight, but the effect was not significant. The earlier work of Brenes et al. [31] also found that the weights of visceral organs (liver, pancreas, small intestine and proventriculus) were not affected by wheat or xylanase. Nayefi et al. [29] observed an increased weight of the gizzard and the rest of the GIT in birds fed CSM-containing diets.

#### Body parts (meat) yield

The yield of carcass parts and dressing percentage were expressed in relation to unit liveweight (g/ kg) and also absolute weight. The outstanding differences between these carcass parts included increased absolute weight of breast meat mediated through enzyme incorporation. The thighs obtained from birds fed CSM diets were also heavier than chickens fed the control diets. The increased weight of carcass parts can be associated with the improved brirds growth, due to increased nutrient digestibility, especially CP and apparent metabolizable energy (AME). The weights of other carcass parts were similar to those of the control group. There are no previous data on the effect of using this kind of enzyme product in diets containing CSM on carcass parts (meat) yield. The eviscerated yield of meat chickens fed CSM-based diets with or without transgenic modification did not differ from the control group fed SBM diets [6].

#### Apparent ileal digestibility of nutrients

Batonon-Alavo et al. [32] reported that CSM inclusion in diets resulted in a decrease in apparent ileal energy digestibility, but in this trial, the gross energy digestibility was not affected by CSM. In addition, different studies have reported that phytase and xylanase supplementation improved AME [33,34]. However, a feeding trial conducted by Munyaka et al. [11] revealed that there was no influence of a xylanase and  $\beta$ -glucanase blend on the AME in meat chickens fed with wheat-

and corn-based feeds. In the present feeding trial, starch digestibility decreased in birds fed high levels of CSM, with the lowest value observed in the group with the highest level of CSM. On the other hand, enzyme supplementation resulted in increased digestibility of starch. It has been found that incorporating a xylanase and  $\beta$ -glucanase blend in corn- and wheat-based diets resulted in increased starch digestibility, among other attributes [11]. This may be due to the role of  $\beta$ -glucanase and xylanase in reducing digesta viscosity caused by wheat arabinoxylans and pentosans and elimination of nutrient encapsulation of cell walls existing in many dietary ingredients, resulting in nutrient release [11,12,35]. The results from this feeding trial show that the graded substitution of SBM with CSM in broiler diets did not affect CP digestibility. Enzyme supplementation also resulted in increased apparent CP digestibility in the ileum by up to about 2%. However, Mushtaq et al. [4] found no impact of adding a  $\beta$ -glucanase and xylanase cocktail to CSM-containing diets. The findings of the current study were similar to those obtained by Garcia et al. [26] who indicated that incorporating xylanase in wheat-based feeds improved ileal digestibility of dietary nutrients, including CP.

# CONCLUSION

The results provided by this feeding trial show that the overall performance of broilers fed CSM-containing diets supplemented with the test enzyme product is not inferior to that of birds fed the more expensive diets containing SBM. For the gross responses, CSM-fed birds showed almost the same responses as the control groups, particularly in the grower and finisher periods. In addition, enzyme supplementation improved different production parameters, such as nutrient digestibility, some gross responses (WG and FCR in grower and finisher phases) and breast meat yield. Therefore, it is evident from this trial that CSM can be a good alternative to SBM, while enzyme incorporation can further improve the quality of such diets by enhancing nutrient digestibility, especially energy. Further tests should investigate the performance of broilers fed diets containing CSM supplemented with other exogenous enzyme combinations, including sodium selenite, lipase and protease.

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