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Histomorphometric traits, microbiota, nutrient digestibility, growth performance, carcass traits and meat quality parameters of chickens fed diets supplemented with different levels of *Bacillus protease*

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ABSTRACT

The effects of dietary *Bacillus protease* on broiler performance were examined. Three hundred 'Cobb 500' day-old broilers were randomly assigned to five dietary groups with five replicates of 12 birds each. Treatments include: PROT0; without protease addition (0 g/kg; control), PROT10, PROT15, PROT20 and PROT25 diets supplemented with *Bacillus protease* at the level of 1, 1.5, 2 and 2.5 g/kg, respectively. Birds fed PROT25 had the highest ($p < 0.05$) villi height and villi height to crypt depth ratio, as well as crypt depth at the duodenum, jejunum, and ileum of intestine. At starter and finisher phases, BWG was the highest ($p < 0.05$), whereas FI and FCR had their lowest values in PROT25 group ($p < 0.05$). PROT25 broilers had the best ($p < 0.05$) energy efficiency ratio. Crude protein, dry matter, and crude fibre digestibility were improved ($p < 0.05$) with increased inclusion levels of *Bacillus protease*. *Lactobacillus* and *Bifidobacteria* count increased ($p < 0.05$) as the inclusion levels of *Bacillus protease* increased in both ileum and caecum, while *E. coli* population decreased. Chickens fed with the PROT25 diets had the highest ($p < 0.05$) values for carcass, thigh, breast and drumstick weight. Cooking loss decreased ($p < 0.05$) as the inclusion levels of *Bacillus protease* increased, while water holding capacity increased ($p < 0.05$). Highest *Bacillus protease* (2.5 g/kg) inclusion improved digestibility, gut integrity, and health status with a better FCR, body weight gain, and retail cut yields.

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KEYWORDS

Crypt depth; *E. coli*; gut integrity; villus height; water holding capacity

1. Introduction

Diet formulation attracts great attention in poultry production, since it constitutes about 60–70% of the production cost (Oyeagu et al. 2019a; Wu et al. 2020). One of the biggest challenges to commercial poultry production is quality feeds at sustainable and stable prices. Kamel et al. (2015) reported 80–85% digestibility of protein compared to 90% for starch in corn-soy diets. They argued that a certain amount of protein passed via gastrointestinal tract (GIT) undigested (Lemme et al. 2004; Schoenfeld and Aragon 2018). Feed ingredients of protein origin (soybean meal and sunflower meal) contain several anti-nutritional factors. Non-starch polysaccharides (NSP) are one of the anti-nutrient which reduce the nutrient bioavailability in poultry. These anti-nutritional factors suppress the production of insulin, which can impair glucose and amino acids uptake and utilization in the peripheral intestinal tissues such as striated muscle by non-ruminant animals, resulting in poor feed efficiency and growth (Masey O'Neill et al. 2014; Qaid and Abdelrahman 2016). An improved performance of birds was a result of exogenous enzyme effect (Shakouri et al. 2008; Oyeagu et al. 2019a). Furthermore, protein digestion in non-ruminant animals is promoted by the inherent proteases in two major stages: protein breakdown (gastric digestion) in an acidic setting and pancreatic digestion in the gut. The inherent proteases are produced and released in the GIT.

They are relatively sufficient to optimize feed protein use (Ndzigaruye et al. 2019), but a significant amount of undigested protein (18–20%) escapes via the GIT (Lemme et al. 2004; Angel et al. 2011). Exogenous protease can enhance animal feed (Oyeagu et al. 2019b). According to Ahmed et al. (2020) the supplementation of exogenous serine-protease (Ronozyme® ProAct) at levels of 200–300 mg/kg can enhance animal protein utilization and resulted in positive effects on growth performance, nutrients utilization, cut yields and encourage beneficial bacteria in the intestine of broilers. Epithelium of the intestine operates as a natural obstruction against pathogenic bacteria and toxic substances. These bio-activities take place in the intestinal lumen. Some of the stressors (anti-nutritional factors and pathogens) cause disturbances in the normal micro-flora epithelium. They change the permeability of these natural obstructions and increase pathogens' attacks, altering the metabolism process. They also impair the capacity to digest and absorb nutrients, resulting in chronic inflammatory activities at the intestinal mucosa (Markowiak and Śliżewska 2017; Pickard et al. 2017). The addition of protease induced some changes in microbial richness and diversity, and tended to change some specific microbial taxa (e.g. increase the abundance of *Bacteroidetes*). These microbes (*Bacteroidetes*) are very important because they produce the favourable metabolites, including SCFAs, which have been correlated

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with an alleviation of gut inflammation (Jeferson et al. 2020). Exogenous proteases may complement pancreatic enzymes and increase the rate of intestinal protein degradation (Mahmood et al. 2017). It has been demonstrated that exogenous protease can change gut morphology, pancreatic enzyme production and secretion, the microbial populations along the gastrointestinal tract, and the short chain fatty acid profile in the digesta (Cowieson 2010). Consequently, shorter villi speed up cell turnover and decrease digestive and absorptive activities (Kai 2021). According to Rodrigues and Choct (2018), dietary phytase and protease reduced the viscosity of the digesta, the relative weight and length of the small intestine, and the number of goblet cells in the duodenum. The microbial community and its activities in the intestinal tract of broilers are influenced by diet composition. It has been demonstrated that anti-nutritional effects on broiler chicken are partly mediated through intestinal microbial activity (Adeokun and Olojede 2019; de Carvalho et al. 2021). However, the restriction on antibiotic growth promoters increased interest concerning the study of micro-flora in broiler GIT. Due to the ban, the persistent proliferation of diseases such as *Clostridium perfringens* (necrotic enteritis) and *E. coli* are expected (Yadav and Jha 2019; Jha et al. 2020). The exogenous enzymes help to remove/reduce anti-nutritional factors' challenging effect (Oyeagu et al. 2015; 2016; 2019a). The use of protease as a dietary enzyme may offer an opportunity to overcome some

of the potential drawbacks imposed by vegetable–protein-based diets that may negatively affect the carcass yield of chicken (Mushtag et al. 2009; Alagawany et al. 2018). Factors that influence meat quality can be controlled throughout the chicken production chain, mainly through diets or during slaughter and processing. The carcass yield is closely linked with the nutritional status of broilers since animals supplied with adequate nutrients will promote the development of muscle tissue (Dalólio et al. 2015; Oyeagu et al. 2019b). The inclusion of protease in the diet has improved protein digestibility and enhanced nutrient availability and absorption for feed conversion to meat yield (Dessimoni et al. 2019; Gervais et al. 2019). Protease inclusion in the corn-soybean diet can alter the nutritional status and improve the retail cut yields and meat quality of broiler chickens (Buyse et al. 2002; Erdaw et al. 2019). According to Medhi et al. (2003), yield of chickens fed diets supplemented with multi-enzyme (xylanase, protease, and amylase) did not differ from those that received the control diet. On the other hand, the inclusion of a blend of protease and xylanase to a fibrous diet improved carcass and retail cut yields (Khan et al. 2006). There is limited information on *Bacillus* protease effect on histomorphometric characteristics, gut microflora, and meat quality of broilers. Therefore, the present study aims to determine *Bacillus* protease's impact on intestinal micro-flora, histomorphology, meat quality, growth performance and carcass traits of broilers.

Table 1. Ingredient (%) and chemical composition (g/kg DM unless otherwise stated) of experimental diets for broiler chicks at the starter phase (0–3 weeks).

Ingredient	PROTO	PROT10	PROT15	PROT20	PROT25
Yellow maize	62.98	62.98	62.98	62.98	62.98
Soybean meal	27.38	27.38	27.38	27.38	27.38
Sunflower meal	4.00	4.00	4.00	4.00	4.00
Fish meal	2.50	2.50	2.50	2.50	2.50
Canola oil	0.11	0.11	0.11	0.11	0.11
Limestone	1.32	1.32	1.32	1.32	1.32
MonoCaP	0.44	0.44	0.44	0.44	0.44
Salt	0.28	0.28	0.28	0.28	0.28
Methionine	0.23	0.23	0.23	0.23	0.23
Threonine	0.05	0.05	0.05	0.05	0.05
Lysine	0.32	0.32	0.32	0.32	0.32
Choline Cl	0.10	0.10	0.10	0.10	0.10
¹ VMP	0.20	0.20	0.20	0.20	0.20
² Maxiban	0.05	0.05	0.05	0.05	0.05
³ Surmax	0.04	0.04	0.04	0.04	0.04
Protease	0.00	0.10	0.15	0.20	0.25
Total	100.00	100.00	100.00	100.00	100.00
Chemical composition					
Moisture	11.11	11.19	11.00	11.18	11.11
ME (kcal/kg)	2988.96	2988.86	2988.06	2988.99	2988.69
Crude protein	23.83	23.90	23.79	23.85	23.88
Crude fat	4.01	4.03	4.00	4.01	4.02
NDF	15.32	15.27	15.39	15.28	15.30
ADF	3.99	3.95	3.97	4.00	3.98
Calcium	0.93	0.94	0.92	0.93	0.94
Phosphorus	0.66	0.68	0.65	0.69	0.67

Notes: MonoCaP = Monocalcium Phosphate. Choline Cl = Choline Chloride. VMP = Vitamine mineral premix. ME = Metabolizable energy. NDF = Neutral detergent fibre. ADF = Acid detergent fibre. PROTO (only basal diet; BD), PROT10 (BD + 1 g protease), PROT15 (BD + 1.5 g protease), PROT20 (BD + 2 g protease) and PROT25 (BD + 2.5 g protease). ¹2.5 kg of vitamin premix contained: 2700 mg retinal, 400 mg caldiol, 18 g tocopheryl acetate, 2000mg menadione, 1800 mg thiamine, 6600 mg riboflavin, 10 g niacin, 30 g calcium pantothenate, 3 g pyridoxine, 1 g folic acid, 15 mg cobalamin, 250 g choline chloride, 100 mg biotin. ²2.5 kg of trace mineral premix contained: 100 g Mn, 50 g Fe, 100 g Zn, 10 g Cu, 1 g I, 200 mg Se. ³1000 g of Maxiban contained: 80 g/kg narasin, 80 g/kg nicarbazin. ³1000 g of Surmax contained: 100 g/kg avilamycin.

2. Materials and methods

2.1. Animal welfare statement

The authors confirm that the ethical policies that guide research of this nature were adhered to. The appropriate ethical review committee approval was received from the University of Fort Hare, South Africa.

2.2. Research trial location

The research trial was conducted at the poultry section of the North-West University research farm (Molelwane) in the North-West province of South Africa. The research area normally experience summer from August to March and has an ambient temperature within 22–35°C. The average yearly rainfall ranges between 200 and 450 mm.

2.3. Additive traits

Exogenous enzyme under investigation (RONOZYME® ProAct (RPA), DSM Nutritional Products Johannesburg South Africa) is a granulated heat-stable formulated product from *Bacillus strain E A.1* with an enzyme activity of 75,000 PROT/g. One PROT is one protease unit and is defined as the amount of enzyme that releases one mmol of p-nitroaniline from one mM substrate (Suc-Ala-Ala-Pro-Phe-pNA) per Adler minute at pH 9.0 and 37°C. The *Bacillus* protease was selected as a feed enzyme due to its inherent bioactive traits. According to the manufacturer, at peptic and acidic conditions (pH), the enzyme retains more than 90% residual activity after 2hrs at 40°C (DSM).

2.4. Experimental diets

Iso-nitrogenous and iso-caloric experimental maize-soybean meal diets were used for the research trial. Starter (0–21 d) and finisher (22–42 d) feeds were formulated (Tables 1 and 2) to satisfy the birds' dietary nutrient needs (NRC 1994). At each feeding phase (i.e. starter and finisher), five dietary treatments were formulated by adding the *Bacillus protease* at five different levels. The five experimental diets for starter and finisher generated were Protease0, Protease10, Protease15, Protease20 and Protease25 represented as PROT0 (only basal diet; BD), PROT10 (BD + 1 g protease/kg diet), PROT15 (BD + 1.5 g protease/kg diet), PROT20 (BD + 2 g protease/kg diet) and PROT25 (BD + 2.5 g protease/kg diet) respectively. Composition and proximate analysis of the five dietary groups (starter and finisher) are presented in Tables 1 and 2, respectively.

2.5. Research chickens and management

Three hundred-day-old chicken (Cobb 500[®]) was used for the research trial and was randomly allocated to the five dietary groups (PROT0, PROT10, PROT15, PROT20, or PROT25). Each dietary group was replicated into five cells, while 12 birds were allotted to each cell. The cells had wood shaving as litter. Chickens were offered diets and clean water *ad libitum* throughout the research trial. The research duration was 6-weeks.

Table 2. Ingredient (%) and chemical composition (g/kg DM unless otherwise stated) of experimental diets for broilers at the finisher phase (4–6 weeks).

Ingredient	PROT0	PROT10	PROT15	PROT20	PROT25
Yellow maize	72.36	72.36	72.36	72.36	72.36
Soybean meal	24.55	24.55	24.55	24.55	24.55
Canola oil	0.24	0.24	0.24	0.24	0.24
Limestone	1.25	1.25	1.25	1.25	1.25
MonoCaP	0.17	0.17	0.17	0.17	0.17
Salt	0.39	0.39	0.39	0.39	0.39
Methionine	0.21	0.21	0.21	0.21	0.21
Tryptophan	0.05	0.05	0.05	0.05	0.05
Threonine	0.05	0.05	0.05	0.05	0.05
Lysine	0.34	0.34	0.34	0.34	0.34
Choline Cl	0.10	0.10	0.10	0.10	0.10
¹ VMP	0.20	0.20	0.20	0.20	0.20
² Maxiban	0.05	0.05	0.05	0.05	0.05
³ Surmax	0.04	0.04	0.04	0.04	0.04
Protease	0.00	0.10	0.15	0.20	0.25
Total	100.00	100.00	100.00	100.00	100.00
Chemical composition					
Moisture	11.01	11.03	11.00	11.02	11.00
ME (kcal/kg)	3110.94	3111.76	3111.00	3111.85	3111.93
Crude protein	19.75	19.91	19.70	19.83	19.85
Crude fat	4.13	4.12	4.12	4.13	4.12
NDF	18.23	18.32	18.28	18.21	18.22
ADF	4.85	4.97	4.82	4.95	4.80
Calcium	0.94	0.92	0.94	0.93	0.93
Phosphorus	0.61	0.62	0.61	0.61	0.61

Notes: MonoCaP = Monocalcium Phosphate. Choline Cl = Choline Chloride. VMP = Vitamine mineral premix. ME = Metabolizable energy. NDF = Neutral detergent fibre. ADF = Acid detergent fibre. PROT0 (only basal diet; BD), PROT10 (BD + 1 g protease), PROT15 (BD + 1.5 g protease), PROT20 (BD + 2 g protease) and PROT25 (BD + 2.5 g protease). ¹2.5 kg of vitamin premix contained: 2700 mg retinal, 400 mg caldiol, 18 g tocopheryl acetate, 2000mg menadione, 1800 mg thiamine, 6600 mg riboflavin, 10 g niacin, 30 g calcium pantothenate, 3 g pyridoxine, 1 g folic acid, 15 mg cobalamin, 250 g choline chloride, 100 mg biotin. ²2.5 kg of trace mineral premix contained: 100 g Mn, 50 g Fe, 100 g Zn, 10 g Cu, 1 g I, 200 mg Se. ³1000 g of Maxiban contained: 80 g/kg narasin, 80 g/kg nicarbazin. ³1000 g of Surmax contained: 100 g/kg avilamycin.

2.6. Growth performance

Growth traits assessed were body weight, feed intake, and feed conversion ratio. Feed consumed was determined on a daily basis by removing the feed refusal from the provided feed, divided by the number of birds in each cell/replicate. Live weight was measured on a weekly basis for all the birds. The feed and live weight grams were determined by using a 10,100 g (10.1 kg) capacity precision weighing balance (A and D Weighing GF-10 K industrial balance, Japan). Feed conversion ratio was calculated by dividing feed consumed with weight gain.

2.7. Slaughter procedure

The chickens were sent to the abattoir (Rooigrond poultry abattoir, North West province, South Africa) for slaughter at the end of the feeding trial (day 42). The chickens were starved for 12 h before slaughter and gas stunned. The chickens were gas stunned by exposing them to relatively low concentrations of carbon dioxide (< 40% by volume in air), and then, once they were unconscious, exposed to a higher concentration (approximately 80% to 90% by volume in air). At the abattoir, all the chickens were hung onto a movable metal rack that held them upside down by their feet. Chickens were then sacrificed by cutting the jugular vein with a sharp knife and they were left hanging until bleeding stopped.

2.8. Carcass parameters and meat quality assessments

Carcass and meat quality traits were determined using 5 birds per pen selected at random immediately after slaughter when the feathers were plucked and the gastro intestinal tract (GIT) was removed. After cutting the carcass into different parts, the breast muscle, neck, wings, shank, thighs, drumsticks and vertebrae (back) were weighed separately. Furthermore, dressing percentage and cut yield of different cuts were also determined according to the following formulae:

$$\text{Carcass yield (dressing percentage)\%}$$

$$= \text{Carcass weight/Slaughter weight} \times 100$$

$$\text{Cut yield(\%)} = \text{Cut weight/Carcass weight} \times 100$$

2.9. Meat pH and temperature measurements

Measurements on meat pH and temperatures were recorded immediately after slaughter and 24 h post slaughter on the breast muscle (central area of the breast) using a Corning Model 4 pH-temperature meter (Corning Glass Works, Medfield, MA) equipped with an Ingold spear-type electrode (Ingold Messtechnik AG, Urdorf, Switzerland).

2.10. Meat colour

The meat colour was assessed as L^* = Lightness, a^* = Redness, and b^* = Yellowness, 24 h after slaughter on the breast muscle using a Minolta colour-guide (BYK-Gardener GmbH, Geretsried, Germany).

2.11. Water holding capacity (WHC)

The water holding capacity was carried out using the pressure method as described by Delezie et al. (2007). About 100 g chicken meat sample from the pectoral major (PM) was cut and weighed as initial weight using a digital sensitive scale up to 0.01 g. The meat sample was placed in between 2 filter papers, and placed on a flat surface. Furthermore, approximately 60 kg weight was applied on the sample for 5 min. The meat sample was weighed again to generate the final (second) weight of the sample. Water holding capacity was calculated as the ratio of the amount of water retained over the initial sample weight.

2.12. Drip loss

The drip loss involved the use of approximately 30 g meat strips samples from the breast muscle parallel to the fibre direction. The samples were weighed for the initial weight using a digital scale sensitive to 0.01 g. The samples were stored inside a plastic container, sealed under atmospheric pressure, and placed at 2°C for 72 h after which they were removed from the container. The samples were blotted with paper towels to remove excess surface moisture and were then reweighed. The drip loss was then calculated by subtracting the blotted sample weight from the initial sample weight.

2.13. Cooking loss measurement

Raw meat cubes were cut from the breast muscle part of the chicken, weighed *in natura*, then placed in a plastic bag and cooked in a water bath at 75°C for 45 min (Rizz et al. 2007). The samples were cooled in running water for 15 min, dried and weighed (Sanka and Mbanga 2014). Cooking loss was calculated as percentage loss of weight after cooking relative to the weight of raw muscle samples before cooking (Gopinger et al. 2014) according to the following formula:

$$\text{Cooking loss(\%)} = \frac{\text{weight before cooking} - \text{weight after cooking}}{\text{weight before cooking}} \times 100$$

2.14. Tenderness

Samples of breast chicken muscles were wrapped in aluminium foil and baked to reach an internal temperature of 85°C, which was maintained for 30 min. Warner Bratzler shear device mounted on Universal Instron apparatus (cross head speed = 200 mm / minute, one shear in the centre of each core), was used for tenderness analysis. The average peak force value of each sample (in Newtons) was recorded using the Texture analyser (TA XT plus).

2.15. Gut histomorphological analysis

Gut histological analysis was implemented using two chickens from each cell at the end of the research trial. About 2 cm segment of different intestinal sections (duodenum, jejunum and ileum) were collected from each selected bird, rinsed

with a 0.9% saline and fixed at 10% buffered formalin solution. The samples were maintained in the formalin at room temperature until analyses. Samples were transferred to the laboratory for histological examination and this involved washing, trimming, dehydration with alcohol, clearing with xylene and impregnation with paraffin wax. For each segment of intestine, tissue sections of 2 µm were cut by a microtome, fixed on slides and stained using haematoxylin and eosin (H and E) for histological analysis. A 2.5× magnification objective lens of a Zeiss Axio light microscope, equipped with a digital camera, was used to examine slides. Images were analysed using Axiovision image-analysis software, version 4.7.2 (Carl Zeiss microscopy). Images was viewed to measure morphometric characteristics of the villi height (VH), villi width (VW), crypt depth (CD), VH/CD ratio, thickness of epithelium and thickness of muscularis.

2.16. Caecal and ileum microflora composition

Two chickens from each cell were used for microbial examination at the end of the research trial. The ileum and caeca parts of the gut were collected immediately under aseptic conditions into sterile glass bags and put on ice before they were transported to the laboratory for microbial examination. The contents of caecal and ileum were emptied in a new sterile bag and were immediately diluted 10-fold (ie 10% wt/vol) with sterile ice-cold anoxic PBS (0.1 M; pH 7.0) and subsequently homogenized for 3 min in a stomacher (Bagmixer 100 Minimix, Interscience, Arpents, France). After the homogenization of each of the caecal and ileum digesta, they were subjected to serial dilution from 10⁻¹–10⁻⁷. The dilutions were later plated on duplicate selective agar media for inventory of target bacterial groups. In particular, *E. Coli*, *Lactobacillus spp* and *Bifidobacterium spp* were enumerated using VRB agar (MERCK, 1.01406), Rogosa agar (MERCK, 1.10660), and Beerens agar respectively according to Tuohy et al. (2002). Plates were incubated at 39°C for 48–120 h anaerobically (Beerens, Rogosa agars) or 24–48 h anaerobically at 37°C (VRB agar). The colonies of the bacteria were recorded, and the average number of live bacteria was calculated based on the weight of original ileum and caecum contents. All the quantitative data generated were converted into logarithmic colony forming units (cfu/g) (Koc et al. 2010).

2.17. Nutrient utilization efficiency

2.17.1. Protein utilization efficiency

Daily protein intake (g); this was calculated as the CP (crude protein) percentage of the diet multiplied with the DFI (daily feed intake). Calculations were made for each bird, where, Total protein intake (TPI) was calculated and was used to calculate protein efficiency ratio (PER).

$$\text{Total protein intake(TPI)} = \text{FI} \times \text{CP} (\%)$$

where TPI = total protein intake, FI = feed intake, CP = crude protein.

$$\text{Protein efficiency ratio(PER)} = \frac{\text{weight gain}}{\text{total protein intake(TPI)}}$$

2.17.2. Energy utilization efficiency

From the two phases (starter and finisher) of feeding trial, Total metabolizable energy intake (TMEI) was calculated and was used to calculate energy efficiency ratio (EER). TMEI was calculated as ME of a diet in kcal multiplied by the feed intake during the whole experimental period.

$$\text{Total metabolizable energy intake (TMEI)} = \text{FI} \times \text{ME}$$

where TMEI = Total metabolizable energy intake, FI = feed intake, ME = metabolizable energy

$$\text{Energy efficiency ratio (EER)} = \text{weight gain} / \text{TMEI}$$

2.18. Nutrient digestibility trial

One week before the end of the research, two birds were removed from each cell to clean metabolic cages. There were three days of adaptation before the four-day sample (droppings) collection. The samples (droppings) were air dried at room temperature. Dried faecal samples were ground and analysed according to AOAC (2006) methods. The original sample was weighed, the weighed sample was placed in a 105°C oven for 12–16 h, and the sample was reweighed. The dry matter and moisture content of the feedstuff were calculated. Crude protein was subjected to Kjeldahl method where a dried sample was boiled in sulphuric acid, was then diluted with water and neutralized with sodium hydroxide, and finally, the sample was distilled and the distilled ammonium was titrated with a known concentrate of sulphuric acid. To determine crude protein content, the nitrogen value from the procedure was multiplied by 6.25. The crude fat or aether extract (EE) procedure estimates the quantity of lipids. The dried samples were ground and extracted with an organic solvent for 4 h and the remaining residue was dried and weighed. Aether extract was then calculated as the difference between the original sample and the aether extract residue. The sample was boiled in a sulphuric acid solution for 30 min and rinsed, and then boiled in a sodium hydroxide solution for 30 min and rinsed. Finally, the residue was dried, weighed, ashed, and reweighed. The crude fibre content was estimated as the difference between the pre-ash weight and the post-ash weight. Nutrient digestibility was calculated as:

$$\begin{aligned} \text{Nutrient digestibility (ND)} &= \text{DI} \times \text{nutrient in feed} - \text{F} \\ &\quad \times \text{nutrient in feed} / \text{DI} \\ &\quad \times \text{nutrient in feed} \times 100 \end{aligned}$$

2.19. Statistical design and analysis

The study was subjected to a completely randomized design (Steel and Torrie 1980) using SAS's General Linear Model Procedure (2010). The applied model is shown below:

$$A_{ij} = \mu + B_i + E_{ij}$$

where A_{ij} = Dependent variable; μ = Overall mean; B_i = Treatment effect; and E_{ij} = Residual error.

The Tukey test was used for mean significance at $p < 0.05$.

3. Results

3.1. Gut histology

Table 3 shows the gut mucosal morphological development of birds fed different inclusion levels of *Bacillus protease*. The histomorphometric characteristics measured in the present study were influenced ($p < 0.05$) by the different supplemental levels of *Bacillus protease*. The thickness of muscularis in jejunum and that of epithelium in ileum section of the intestine did not differ ($p > 0.05$). Chickens that consumed PROT25 recorded a higher ($p < 0.05$) villi height, and villi height to crypt depth ratio, with a lower ($p < 0.05$) crypt depth in the duodenum, jejunum and ileum sections. Chickens that consumed the experimental diets (PROT10, PROT15, PROT20 and PROT25) had greater ($p < 0.05$) villi height than those of the PROT0 group (control diet). On the other hand, birds in the treatment groups had a decreased ($p < 0.05$) crypt depth compared to the control group in the three sections of the gut. There was an increased ($p < 0.05$) villi height to crypt depth ratio for the three gut sections (duodenum, jejunum, and ileum) in protease supplemented chickens compared to the controls (PROT0). Figures 1 and 2 present the histomicrograph of broiler jejunum and ileum as affected by dietary supplementation with *Bacillus protease*. Figure 1 showed the histomicrograph of broiler Jejunum fed different levels of *Bacillus protease*, while Figure 2 represents the histomicrograph of ileum intestinal part of broilers.

3.2. Growth traits

Table 4 shows the effect of *Bacillus protease* dietary supplementation on growth traits. The data for the starter phase (1 to 21 d) showed a lower ($p < 0.05$) value of FCR for chickens that consumed PROT25 than that of the other dietary groups (PROT0, PROT10, PROT15, and PROT20). It was noticed that BWG was higher ($p < 0.05$) for chickens fed PROT25, while their feed intake was the lowest ($p < 0.05$). Although, chickens that consumed dietary PROT0, PROT10, PROT15, and PROT20 had greater feed intake, yet they recorded lower values for BWG ($p < 0.05$). During the finisher phase (22 to 42d), the BWG of PROT25 birds was the highest ($p < 0.05$) but similar to that of PROT20 group. PROT10 birds displayed the lowest ($p < 0.05$) BWG similar to that of PROT0 group. Chickens that consumed the PROT25 diet had the lowest ($p < 0.05$) FCR of 1.23 while a higher ($p < 0.05$) FCR of 1.91 and 1.79 was observed in birds fed with PROT0 and PROT10 diets, respectively. Chickens fed PROT25 consumed less ($p < 0.05$) feed compared to that of PROT0, PROT10, PROT15 and PROT20 group. As shown on the overall phase performance, birds fed PROT25 recorded the lowest FI with a better ($p < 0.05$) BWG of 3060 g, than those that received dietary PROT0 and PROT10. Birds fed the PROT25 diet recorded a better FCR than those received the PROT0 and PROT10 diets.

3.3. Utilization efficiency of energy and protein

The impact of dietary *Bacillus protease* inclusion on protein/energy utilization efficiency is presented in Table 5. As

Table 3. The effect of dietary *Bacillus* protease supplementation on the gut histomorphometric traits of broiler birds.

	Diets PROTO	PROT10	PROT15	PROT20	PROT25	SEM	<i>p</i> -value
Duodenum							
Villus height (µm)	1301.75 ^e	1407.15 ^d	1512.81 ^c	1610.74 ^b	1650.35 ^a	19.87	0.05
Crypt depth (µm)	280.46 ^a	251.99 ^b	231.79 ^{bc}	183.14 ^c	148.91 ^d	13.55	0.04
VH/CD ratio	4.64 ^d	5.58 ^c	6.53 ^c	8.81 ^b	11.08 ^a	3.03	0.00
TE (µm)	53.44 ^b	73.32 ^{ab}	72.59 ^{ab}	80.18 ^a	78.29 ^a	5.92	0.02
TM (µm)	196.44 ^b	212.48 ^{ab}	215.54 ^{ab}	217.43 ^{ab}	238.86 ^a	12.45	0.04
Jejunum							
Villus height (µm)	1197.63 ^{de}	1295.81 ^d	1418.72 ^c	1445.51 ^b	1481.62 ^a	18.99	0.05
Crypt depth (µm)	225.66 ^a	185.78 ^b	167.84 ^{bc}	166.71 ^{bc}	124.94 ^c	13.01	0.03
VH/CD ratio	5.31 ^c	6.97 ^c	8.45 ^b	8.67 ^b	11.86 ^a	4.07	0.01
TE (µm)	42.92 ^c	48.69 ^b	50.91 ^b	55.88 ^{ab}	61.66 ^a	5.81	0.05
TM (µm)	147.13	146.55	151.32	150.91	154.84	11.00	0.10
Ileum							
Villus height (µm)	1016.95 ^e	1127.47 ^d	1227.87 ^c	1339.08 ^b	1374.86 ^a	17.98	0.04
Crypt depth (µm)	202.54 ^a	160.38 ^{bc}	161.92 ^{bc}	175.35 ^b	146.54 ^c	11.99	0.05
VH/CD ratio	5.79 ^c	7.03 ^b	7.54 ^b	6.52 ^b	9.38 ^a	4.05	0.00
TE (µm)	48.91	47.79	45.57	50.99	51.46	6.42	0.08
TM (µm)	187.71 ^c	207.44 ^b	211.93 ^b	230.53 ^a	245.86 ^a	11.11	0.03

Notes: ^{a,b,c}Row means with different superscripts differ significantly. SEM = Standard error of the mean. PROTO = Basal Diet: BD (without PROT; protease). PROT10 = BD + 1 g PROT. PROT15 = BD + 1.5 g PROT. PROT20 = BD + 2 g PROT. PROT25 = BD + 2.5 g PROT. VH/CD ratio = Villi height: Crypt depth ratio. TE = Thickness of epithelium. TM = Thickness of muscularis. Two birds from each cell/replicate were used for the analysis.

indicated in all the phases (starter phase, finisher phase, and overall performance) of productions, the protein efficiency ratio (PER) of birds fed PROT25 had a greater value ($p < 0.05$) than that of birds of the other treatments (PROTO, PROT10, PROT15, and PROT20). The other treatment groups had lower ($p < 0.05$) intake of energy and a higher energy efficiency ratio compared to the control (PROTO) group at all phases or stages of production (starter phase, finisher phase, and overall performance). On the contrary, birds fed PROT25 consumed less ($p < 0.05$) energy and had the best energy efficiency ratio (EER) throughout the feeding trial. Daily protein intake increased ($p < 0.05$) in chickens that consumed the control, PROT10, PROT15, and PROT20 diets, while PROT25 fed birds consumed the lowest amount of protein. During the starter phase, protein intake for birds fed PROTO, PROT10, PROT15 and PROT20 was significantly ($p < 0.05$)

higher, while birds fed PROT25 consumed less ($p < 0.05$) protein. The protein intake of chickens at the finisher and overall phases showed that birds fed PROT25 consumed less ($p < 0.05$) protein than those in other groups (PROTO, PROT10, PROT15 and PROT20).

3.4. Nutrient digestibility

The apparent nutrient digestibility values in broiler chickens fed dietary *Bacillus* protease supplementation are presented in Table 6. Crude fat digestibility was the same ($p > 0.05$). Crude protein digestibility was higher ($p < 0.05$) for chickens that consumed PROT20 and PROT25. The level of digestibility recorded for crude fibre and dry matter was also higher ($p < 0.05$) for chickens provided PROT20 and PROT25 diets, but they are statistically the same ($p > 0.05$) with those that received dietary

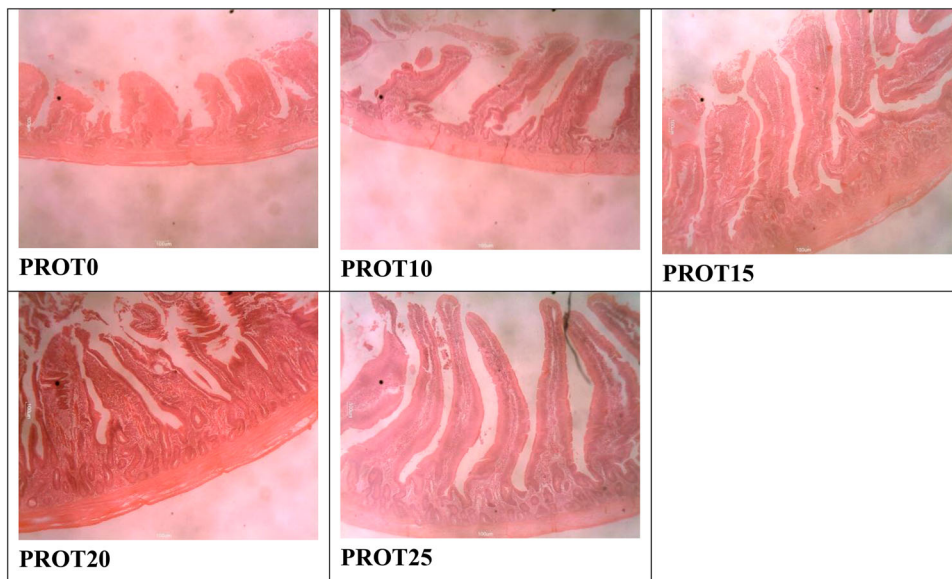


Figure 1. Histomicrograph of the Jejunum of broilers fed PROTO, PROT10, PROT15, PROT20 and PROT25 which shows the villi height, crypt depth, thickness of the epithelium, and muscularis. Two birds from each cell/replicate were used for the analysis.

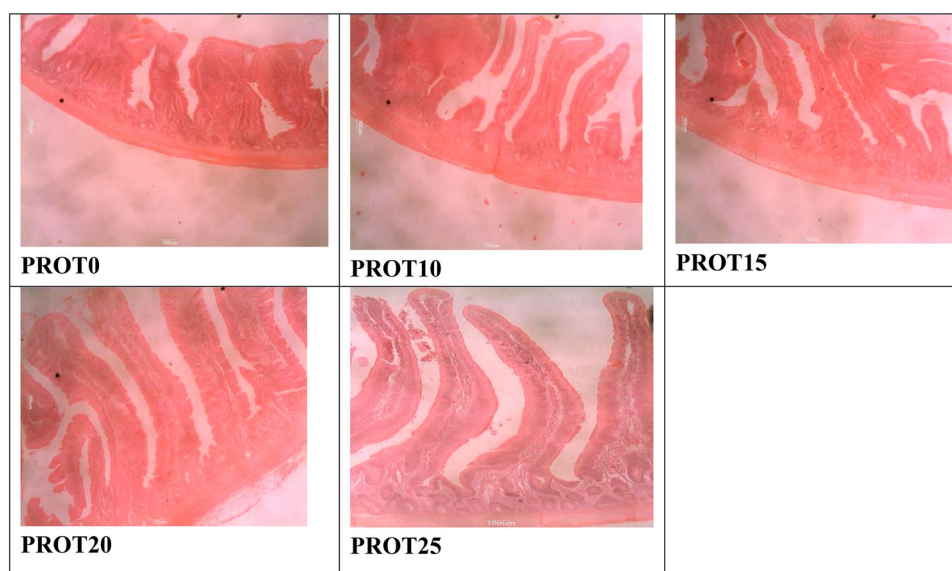


Figure 2. Histomicrograph of the ileum of broilers fed PROT0, PROT10, PROT15, PROT20 and PROT25 which shows the villi height, crypt depth, thickness of the epithelium, and muscularis. Two birds from each cell/replicate were used for the analysis.

PROT15. The result showed that crude protein, dry matter, and crude fibre digestibility improved ($p < 0.05$) as the inclusion rate of *Bacillus* protease increased.

3.5. Intestinal micro examination

Figures 3 and 4 showed the ileum and caecum microbial population changes in birds fed dietary supplemented with *Bacillus* protease. From Figure 3, different *Bacillus* protease levels influenced ($p < 0.05$) the counts of *Lactobacillus*, *Bifidobacteria*, and *E.coli* found in the ileum section of the small intestine. *Lactobacillus* and *Bifidobacteria* count increased ($p < 0.05$) as the addition of *Bacillus* protease levels increased. The population of these bacteria (*Lactobacillus* and *Bifidobacteria*) was higher ($p < 0.05$) in the ileum section of the intestine for chickens that consumed the supplemented diets than the controls (PROT0) and PROT10 diet. Birds that consume PROT15, PROT20 and PROT25 recorded decreased ($p > 0.05$) counts for *E.coli* compared to the control (PROT0) and PROT10 group. Chickens that consume PROT25 recorded the lowest ($p < 0.05$) *E.coli* population in the ileum section of the gut. Meanwhile, Figure 4 represents the microbial population changes in the caecum section of the intestine. The *Lactobacillus* and *Bifidobacteria* populations in the caecum were the highest ($p < 0.05$) for chickens that consumed the PROT25 diet than those that consumed PROT0 (control diet). The lowest ($p < 0.05$) counts for *E. coli* at the caecum section of the small intestine were recorded for chickens fed PROT25 compared with those fed with PROT0 diet.

3.6. Carcass traits

Table 7 shows the carcass traits of chickens that were supplemented with *Bacillus* protease. The carcass weight, drumstick weight, thigh weight, and breast weight were significantly affected ($p < 0.05$) by the *Bacillus* protease inclusion. Conversely, neck, wing, and shank weights were

the same ($p > 0.05$). Carcass and thigh weights were higher ($p < 0.05$) for chickens provided with PROT25 than PROT0 and PROT10 diets. Breast weight was the highest ($p < 0.05$) for chickens fed PROT20 or PROT25 compared to other groups. The highest ($p < 0.05$) drumstick weight was also observed in birds fed PROT25 diets. Carcass yield (dressing percentage), drumstick yield, thigh yield, and breast yield percentage were influenced ($p < 0.05$) by *Bacillus* protease supplementation, while the neck, wing, and shank yields were the same ($p > 0.05$). Birds fed PROT20, and PROT25 recorded the highest ($p < 0.05$) breast yield percentage. The thigh yield percentage was higher ($p < 0.05$) for chickens that consumed PROT25 than those fed PROT0, PROT10, and PROT15. PROT25 birds had an improved ($p < 0.05$) percentage for carcass yield and drumstick yield than those receiving other dietary treatments.

3.7. Meat quality traits

The data for meat quality parameters of broilers fed dietary *Bacillus* protease is shown in Table 8. Only cooking loss (CL) and water holding capacity (WHC) was influenced ($p < 0.05$) by dietary *Bacillus* protease among the determined meat quality parameters. The values for CL decreased ($p < 0.05$) as the inclusion rate of *Bacillus* protease increased. Cooking loss decreased ($p < 0.05$) for birds fed PROT15, PROT20 and PROT25 compared to those provided PROT0 and PROT10. WHC was higher ($p < 0.05$) in birds fed all the dietary treatments except for those that consumed PROT0.

4. Discussion

4.1. Gut histology

Intestinal morphology involves the display and study of the structure of the small intestine under a microscope. Some of the microscopic structure includes villus height and crypt depth. The villus height and crypt depth are core pointers of

Table 4. The effect of *Bacillus* protease inclusion on feed intake, body weight gain, and feed conversion ratio of broiler chicken.

Treatment	PROTO	PROT10	PROT15	PROT20	PROT25	P-value	SEM
Items							
BWG (g)							
1–21 d	950.40 ^b	951.20 ^b	953.80 ^b	966.20 ^b	1075.00 ^a	0.04	19.05
22–42 d	1579.40 ^c	1580.60 ^c	1623.80 ^b	1725.00 ^{ab}	1950.00 ^a	0.03	19.08
1–42 d	2566.80 ^c	2579.80 ^c	2609.60 ^{bc}	2712.20 ^b	3060.00 ^a	0.05	20.33
FI (g/bird)							
1–21 d	1276.00 ^a	1221.00 ^a	1209.50 ^a	1265.00 ^a	1097.80 ^b	0.04	19.10
22–42 d	3014.00 ^a	2827.80 ^{ab}	2731.30 ^b	2768.90 ^{ab}	2396.50 ^c	0.02	21.64
1–42 d	4280.10 ^a	4039.40 ^{ab}	3930.80 ^b	4023.10 ^{ab}	3484.40 ^c	0.03	23.04
FCR							
1–21 d	1.34 ^a	1.30 ^a	1.31 ^a	1.30 ^a	1.02 ^b	0.01	0.00
22–42 d	1.91 ^a	1.79 ^{ab}	1.68 ^b	1.61 ^b	1.23 ^c	0.02	0.01
1–42 d	1.67 ^a	1.57 ^a	1.51 ^a	1.48 ^{ab}	1.14 ^b	0.01	0.01

Notes: ^{a,b,c}Row means with common superscripts do not differ significantly. SEM = Standard error of the mean. BWG = Body weight gain. FI = Feed intake. FCR = Feed conversion ratio. PROTO (only basal diet; BD), PROT10 (BD + 1 g protease), PROT15 (BD + 1.5 g protease), PROT20 (BD + 2 g protease) and PROT25 (BD + 2.5 g protease). Three hundred birds were used for the analysis.

gut development, health, and functionality, influencing the digestion and absorption of nutrients (Biasato et al. 2018). The results obtained from the present study showed that dietary *Bacillus* protease at a 2.5 g/kg diet (PROT25) enhanced the growth and development of the gut. Generally, birds fed PROT25 recorded an increased villi height in the duodenum, jejunum, and ileum sections of the gut than the control group. The results of the present study showed that *Bacillus* protease in caecum could improve the development of the small intestine by reducing the crypt depth of ileum and increasing the villus height of ileum and duodenum (Wexler 2007). In the previous studies, *Bacillus* protease could produce volatile fatty acid (VFAs) through carbohydrate fermentation (Louis and Flint 2009), which could maintain gut health and served as the major source of energy for the gut mucosa to promote the growth of gut villus (Louis and Flint 2009). According to Vu et al. (2021), *Bacillus* protease could actually be beneficial for the development of intestinal morphology through the regulation and stabilization of gastrointestinal microbiota, the improvement of the gut morphology

and mucosal immunity and the elevation of beneficial bacteria to maintain the overall growth and development of gut morphology in broilers. As protease increases protein digestibility, the amino acids and polypeptides absorbed by the chicken would increase. In this scenario, protease help to create the availability of more amino acids for birds' utilization. It also influences the production of short-chain fatty acid concentrations (Fan et al. 2015; Apajalahti and Vienola 2016). The short-chain fatty acids (SCFAs) are the primary products of the breakdown of non-digestible carbohydrates and amino acids by gut bacteria. Collectively, they are a major source of energy for colon cells (Peng et al. 2009). SCFAs improve the gut health through a number of local effects, ranging from maintenance of intestinal barrier integrity, mucus production, and protection against inflammation to reduction of the risk of colorectal cancer (Gaudier et al. 2009; Lewis et al. 2010; O'Keefe 2016). The height of the villi is one of the morphological traits, and an increase in the villi height will result in an enhanced absorption ability (Markovic et al. 2009; Parviz et al. 2018). The small intestinal villi structure mainly utilizes

Table 5. The effect of dietary protease supplementation on protein intake and protein efficiency ratio of broiler birds.

Treatments	PROTO	PROT10	PROT15	PROT20	PROT25	p-value	SEM
Items							
DPI (g)	20.54 ^a	19.38 ^{ab}	19.86 ^{ab}	19.30 ^{ab}	17.71 ^b	0.04	0.84
DPER (g)	0.07 ^b	0.07 ^b	0.08 ^b	0.07 ^b	0.10 ^a	0.01	0.00
DMEI (kcal/bird)	312.61 ^a	295.12 ^a	287.33 ^{ab}	293.56 ^a	254.28 ^c	0.05	3.85
DEER(kcal/100 kcal)	0.49 ^b	0.50 ^b	0.50 ^b	0.50 ^b	0.68 ^a	0.02	0.07
Starter phase							
PI (g)	301.31 ^a	288.21 ^a	285.49 ^{ab}	298.50 ^a	258.91 ^b	0.05	2.87
PER (g)	3.14 ^b	3.27 ^b	3.33 ^b	3.21 ^b	4.11 ^a	0.03	0.01
MEI (kcal/bird)	3784.07 ^a	3614.84 ^a	3582.90 ^{ab}	3743.67 ^a	3248.17 ^c	0.05	13.78
EER(gain/100 kcal)	24.93 ^{bc}	26.04 ^b	26.56 ^b	25.54 ^b	32.79 ^a	0.03	0.81
Finisher phase							
PI (g)	606.83 ^a	569.31 ^{ab}	559.70 ^{ab}	557.29 ^{ab}	482.08 ^b	0.05	3.10
PER (g)	2.82 ^b	2.84 ^b	2.87 ^b	2.83 ^b	4.03 ^a	0.01	0.01
MEI (kcal/bird)	9345.44 ^a	8780.26 ^a	8585.01 ^{ab}	8585.70 ^{ab}	7431.56 ^b	0.05	12.89
EER(gain/100 kcal)	18.36 ^b	18.38 ^b	18.51 ^b	18.27 ^b	26.10 ^a	0.02	0.93
Overall performance							
PI (g)	908.14 ^a	857.52 ^{ab}	855.19 ^{ab}	855.79 ^{ab}	740.99 ^b	0.04	4.19
PER (g)	2.93 ^b	2.98 ^b	3.02 ^b	2.96 ^b	4.06 ^a	0.02	0.01
MEI (kcal/bird)	13129.50 ^a	12395.10 ^{ab}	12267.90 ^{ab}	12329.40 ^{ab}	10679.70 ^b	0.05	39.67
EER(gain/100 kcal)	20.59 ^{bc}	20.97 ^b	21.19 ^b	20.84 ^b	28.56 ^a	0.03	0.90

Notes: ^{a,b,c}Row means with different superscripts differ significantly. SEM = Standard error of the mean. DPI = daily protein intake. DPER = daily protein efficiency ratio. DMEI = daily metabolizable energy intake. DEER = daily energy efficiency ratio. PI = protein intake. PER = protein efficiency ratio. MEI = metabolizable energy intake. EER = energy efficiency ratio. PROTO (only basal diet; BD), PROT10 (BD + 1 g protease), PROT15 (BD + 1.5 g protease), PROT20 (BD + 2 g protease) and PROT25 (BD + 2.5 g protease). Three hundred birds were used for the analysis.

Table 6. Effects of dietary protease supplementation on apparent digestibility (%) of dry matter, crude protein, crude fat, and crude fibre in broiler finisher diets.

Parameters	Diets					SEM	<i>p</i> -value
	PROTO	PROT10	PROT15	PROT20	PROT25		
Crude fibre	11.99 ^b	12.29 ^b	16.56 ^{ab}	17.29 ^a	18.55 ^a	0.07	0.04
Crude protein	78.42 ^c	79.55 ^c	84.04 ^b	87.99 ^a	89.97 ^a	0.18	0.02
Crude Fat	76.85	77.79	78.04	78.56	77.26	0.16	0.08
Dry Matter	75.00 ^b	74.11 ^b	77.01 ^{ab}	81.90 ^a	82.83 ^a	0.19	0.03

Notes: ^{a,b,c}Row means with different superscripts differ significantly. SEM = Standard error of the mean. PROTO (only basal diet; BD), PROT10 (BD + 1 g protease), PROT15 (BD + 1.5 g protease), PROT20 (BD + 2 g protease) and PROT25 (BD + 2.5 g protease). Two birds from each cell/replicate were used for the analysis.

nutrients (Fuller 2004; Chwen et al. 2013) and the increase in villi height increases nutrient absorption capacity in the gut due to increased absorptive area, expression of brush border enzyme, and nutrient transport system, leading to a higher body weight gain of chicken (Parviz et al. 2018). *Bacillus* protease in PROT25 group may have reduced the intestinal colonization and infections process, hence reducing intestinal mucosa inflammation, which improved the height of villi and function of secretion, digestion, and absorption of nutrients (Khan and Iqbal 2016). According to Sobolewska et al. (2017), higher villi height and their increased absorbent surface area translate into improved feed utilization and chicken health. The villi height was lowest for the duodenum, jejunum, and ileum of PROTO birds, which may be the reason for their poor performance. The condition of the crypt dept explains the state of health and functional status of the chicken intestine. Their sizes determine the rate of the renewal process of intestinal epithelium (Haihan et al. 2019; Ivana et al. 2019). The proliferation of the intestinal cells takes place, particularly in the crypts (Kaunitz and Akiba 2019), and a large deeper crypt shows increased nutrient needs for intestinal care and low efficiency of chicken. Lower (shallow) crypt depth observed in our study for chickens fed *Bacillus* protease inclusion at 2.5 g/kg diet (PROT25) is a gainful approach to reduce the cost of intestinal care in chickens. According to Xu et al. (2003) and Ivana et al. (2019), the crypt is the villi factory, and a large crypt indicates a faster tissue turnover and an increased request for new tissue. Again, the decreased crypt depth in our study may indicate a proficient tissue turnover, good gut condition, and improved chicken body weight. Moreover, some results showed that the villi and microvilli of intestinal mucosa could affect the production of the digestive enzymes (Wu et al. 2013). Our study showed an increase in villi height and villus height: crypt ratio for PROT25 chickens. The improved mucosal morphology was attributed to increased enzymatic bio-activities of gastrointestinal contents. Enhanced intestinal morphology results in better nutrient absorption, decreased secretion in the gastrointestinal tract, increased disease resistance, and improved overall performance (Ebrahimzadeh et al. 2018). The lowest villi height: crypt depth ratio was recorded for chickens fed the control diet (PROTO). According to Parsaie et al. (2007) and Ensari and Marsh (2018), higher villi is related with the efficient absorption capacity of the enterocytes, while short villi decrease the surface area for nutrient absorption. However, the epithelial cells of the villi emanate from the crypt, and a large crypt shows fast enterocyte turnover and increased need for mucosal tissue maintenance requirements (Ayoola et al. 2015). The rapid enterocyte production

and the epithelial cell renewal rate impact the small intestinal mucosa's protein and energy requirements, reducing the animal's feed efficiency (Luo et al. 2009; Liisa et al. 2020). Reduced villi height and deepened crypts increased secretion in the GI tract, resulting in birds' poor growth performance (Parsaie et al. 2007; Ivana et al. 2019). According to Kim and Khan (2013), the mucus deposit in the small intestine is from goblet cells, which permits gastric secretion and forms a shielding obstruction against mechanical and chemical damages such as ingested feed, microorganisms, and pathogens (Hollingsworth and Swanson 2004; Sobolewska et al. 2017). Meanwhile, goblet cells are modified epithelial cells that produce mucus on the mucous membrane surface (epithelium layer and an underlying lamina propria of loose connective tissue) of the small intestine. The mucous membrane secretes mucous, a thick protective fluid that prevents pathogens and dirt from entering the body and prevents bodily tissue from becoming dehydrated. Muscularis is a thin layer of smooth muscle that supports mucous and provides it to move and fold. It is responsible for the segmental contractions and peristaltic movement in the GIT. These muscles cause feed to move and mix together with the digestive enzyme down the GIT. In the present study, only the muscularis thickness for the duodenum and ileum was affected by the dietary *Bacillus* protease. The study of Ebrahimzadeh et al. (2018) contradicts the results of the present study. They reported a decreased muscularis thickness with the inclusion of enzyme fortified grape pomace.

4.2. Growth performance traits

The inclusion of *Bacillus* protease impacted the BWG, FI, and FCR of broilers at all growth phases (starter phase, finisher phase, and overall phase). Chickens fed with the PROT25 diet had a higher BWG and improved FCR with less FI compared to those provided with the other dietary treatments. Our research findings align with Ghazi et al. (2002) and Cerrate et al. (2019). They attributed a better body weight and feed conversion ratio to improved true metabolizable energy and true nitrogen digestibility. Dietary fibre is implicated in the fast passage rate of the digesta (Wilfarta et al. 2007; Sayehban et al. 2015; Al-harhi 2016), impairing broiler growth. The better performance of birds fed with PROT25 diet in our study may be due to the slow digesta passage rate, which offers maximum opportunity for increased bio-activities of *Bacillus* protease in the gut. As the slow passage rate of fibre digesta persists in the intestine, it provides greater chances for improved feed digestion and growth performance (Sayehban et al. 2016).

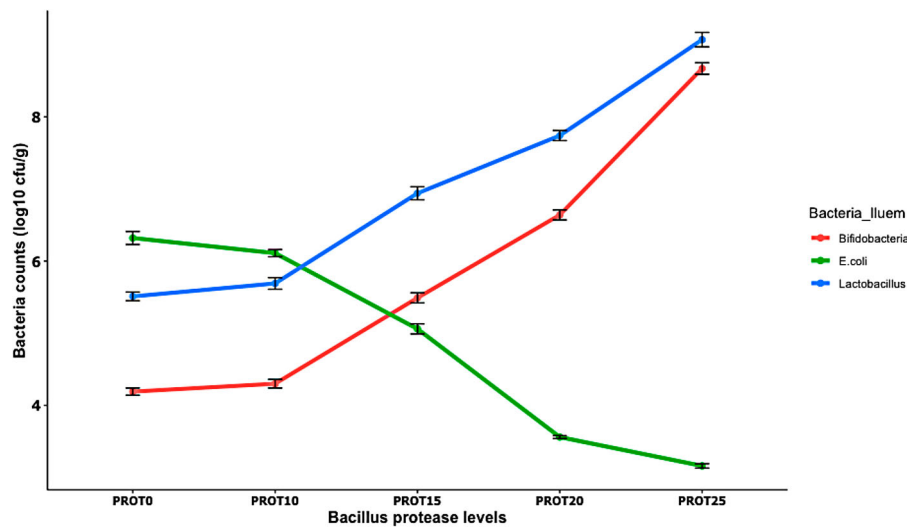


Figure 3. The effect of dietary *Bacillus* protease on *Lactobacillus*, *Bifidobacteria* and *E. coli* counts (\log_{10} cfu/g) in the ileum section of the intestine. Two birds from each cell/replicate were used for the analysis.

Dietary whole wheat (high fibre) decreases the bio-activities of amylase in the pancreatic tissue, while the inclusion of protease increases the bio-activities of chymotrypsin and lipase (Engberg et al. 2005; Selle et al. 2010). The present findings are in line with the study of Dosković et al. (2013), who suggested that exogenous protease complemented endogenous protease to increase intestinal and pancreatic protease production in young birds. Essentially, this phenomenon clearly suggests that the production of endogenous protease may be inadequate in the initial post-hatch period in poultry. Our results are in accordance with that of Odetallah et al. (2003) and Mahmood et al. (2017), who reported a significant increase in BWG with a decreased FI after supplying protease fortified standard protein diet to broilers for 21 days. Similar results (Odetallah et al. 2005) recorded an improved BWG and FCR in broilers fed dietary protease (versazyme) in high and normal protein diets. Improved BWG and FCR of chickens fed dietary protease may be attributed to efficient nutrient digestibility

and its utilization, mostly when feed ingredients are of inferior/low quality and low bioavailability (Kocher et al. 2002; Cowieson and Adeola 2008; Gervais et al. 2019). According to Freitas et al. (2011), improvement in BWG and FCR with protease supplementation is due to its direct effect in protein digestibility, liberating more amino acids for protein synthesis. Gao et al. (2008) reported that supplementation of protease could change the nutritional and physiological status of the broilers and improve growth rate. The studies of Freitas et al. (2011) and Law et al. (2018) showed a general decrease in FI of chickens fed dietary protease, similar to the results generated in our study. The fact that birds consumed less on a nutrient-dense diet and satisfied their nutrient requirements may explain the decreased FI reported for birds fed PROT25 in the present study. The ingredients of each diet determine whether the animal will consume more or less before satisfying its nutritional requirement (Ani and Oyeagu 2012). Several authors also share the same view (Alam et al. 2003; Oyeagu

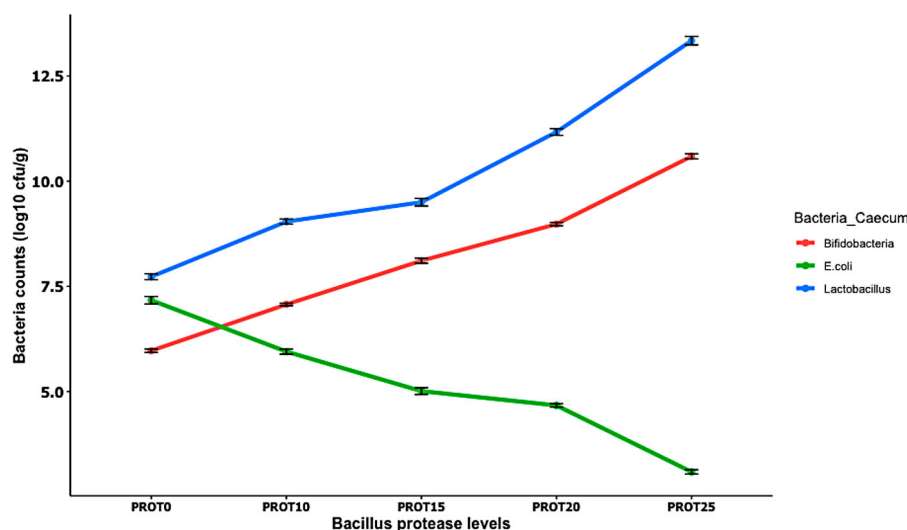


Figure 4. The effect of dietary *Bacillus* protease on *Lactobacillus*, *Bifidobacteria*, and *E. coli* counts (\log_{10} cfu/g) in the caecum section of the intestine. Two birds from each cell/replicate were used for the analysis.

Table 7. The effect of dietary *Bacillus* protease supplementation on carcass characteristics of broiler birds.

Treatment	PROTO	PROT10	PROT15	PROT20	PROT25	SEM	<i>p</i> -value
Slaughter weight (g)	2556.00 ^c	2569.20 ^c	2599.80 ^c	2802.80 ^b	3050.00 ^a	20.01	0.04
Carcass weight (g)	2006.80 ^{cd}	2014.05 ^{cd}	2050.09 ^c	2220.90 ^b	2508.30 ^a	18.97	0.02
Neck weight (g)	85.02	80.00	80.02	80.77	83.33	2.32	0.11
Wing weight (g)	83.05	77.07	80.01	78.33	82.65	2.06	0.09
Drumstick weight (g)	81.00 ^{bc}	85.67 ^b	88.27 ^b	89.11 ^b	108.47 ^a	2.57	0.01
Thigh weight (g)	80.33 ^{bc}	81.93 ^{bc}	88.33 ^b	89.67 ^b	110.47 ^a	2.98	0.04
Breast weight (g)	379.66 ^c	390.48 ^b	400.20 ^b	515.00 ^a	535.60 ^a	5.76	0.02
Shank weight (g)	38.20	34.80	35.00	35.27	39.66	0.98	0.17
Carcass yield (%)	77.26 ^b	78.19 ^b	79.78 ^b	79.79 ^b	84.24 ^a	1.97	0.01
Neck yield (%)	3.83	3.98	3.97	3.94	3.88	0.07	0.07
Wing yield (%)	3.74	3.84	3.89	3.72	3.78	0.06	0.10
Drumstick yield (%)	3.83 ^b	3.93 ^b	3.89 ^b	3.96 ^b	4.32 ^a	0.08	0.02
Thigh yield (%)	3.97 ^c	4.03 ^c	4.04 ^c	4.18 ^b	4.40 ^a	0.09	0.02
Breast yield (%)	18.30 ^b	17.44 ^b	17.93 ^b	20.24 ^a	21.35 ^a	0.76	0.05
Shank yield (%)	1.72	1.73	1.74	1.72	1.70	0.02	0.15

Notes: ^{a,b,c}Row means with different superscripts differ significantly. SEM = Standard error of the mean. PROTO = Basal Diet: BD (without PROT; protease). PROT10 = BD + 1 g PROT. PROT15 = BD + 1.5 g PROT. PROT20 = BD + 2 g PROT. PROT25 = BD + 2.5 g PROT. Five birds from each cell/replicate were used for the analysis.

et al. 2016; Oyeagu et al. 2019a). Indeed, Hajati et al. (2009) and Hajati (2010) suggested that exogenous enzyme inclusion may improve broiler performance by improving nutrient digestibility. The reduction of the digesta passage rate in the gut, which activates viscosity reduction is one of the mechanisms of achieving nutrient digestibility (Oyeagu et al. 2015). However, Fru-Nji et al. (2011) reported no change on the BWG of broilers fed dietary inclusion of protease at 3% during the finisher phase, which contradicts the present results. Such discrepancies may be due to variations in feed ingredient contents, levels of exogenous enzymes inclusion, or breeds of chicken used (Angel et al. 2011).

4.3. Protein and energy utilization efficiency

Bacillus protease supplementation increased energy and protein utilization efficiency with less protein and energy intake for birds fed with the PROT25 diet throughout the feeding period. The efficient utilization of energy and protein recorded for PROT25 birds may be the reason for their improved BWG and FCR (Table 4), as supported by Kamran et al. (2008) findings. This study, however, contradicts the findings of Moosavi et al. (2012), who recorded no effect of dietary protease on both PER and EER of broiler chickens. According to Olukosi et al. (2015), NSPase and protease inclusion improved the metabolizable energy. It is important to note that different ingredients of broiler diets and the differences within them affect nutrient digestibility (Jha and Mishra 2021; Tejeda and Kim 2021). For instance, protein digestibility is high in corn-soybean meal-based broiler diets but lower in animal-derived meals. The disparity may result from excessive thermal processing and the type of contained animal by-products (Douglas et al. 2000; Goldflus et al. 2006; de Coca-Sinova et al. 2008; Oyeagu et al. 2016). Variation in protein and energy digestibility could be observed within the same feed ingredient (Lemme et al. 2004; Rojas and Stein 2017). Considering the diet composition used in this study, birds fed with the PROT25 diet showed improved BWG (Table 4) due to efficient utilization of protein and energy. Also, it may be that the level of *Bacillus* protease in PROT25 diet improved protein availability as it broke down protein–starch bonds within feed ingredients (Selle et al. 2013). The effective

separation of protein–starch bond due to protease inclusion increases digestion and facilitates the utilization of extra energy by the animal (du Plessis and van Rensburg 2014). The diets contain amino acids complexes with reducing sugars that could potentially make them impossible to digest. The possibility of beneficial impact of exogenous protease in liberating amino acids from reducing complex sugars cannot be ignored. It has been suggested that exogenous protease acts by releasing peptides from anti-nutritional factors present in the feed ingredients, cleaving linkage between amino acids–starch complexes, supplementing the endogenous production of peptidases, and reducing enzymatic secretions and protein turnover; thereby providing amino acids for protein synthesis and deposition (Cowieson and Roos 2013). It was reported that age of a bird and diet are known to influence pancreatic output and enzyme composition (Isaksen et al. 2010). Although more research is needed, in line with our results, there is evidence that the addition of exogenous protease to the diet increases pancreatic production and secretion of endogenous proteolytic enzymes (Jeferson et al. 2020), an effect that result in higher protein digestibility (Yuan et al. 2017). It is important to note that amino acid deficiency is known to reduce growth rate and feed efficiency in broilers (Freitas et al. 2011). The better performance of birds fed the highest *Bacillus* protease inclusion level (PROT25) than other groups in our study may be due to the maximum pancreatic enzyme secretion that increased protein digestibility. According to Lemme et al. (2004) and Freitas et al. (2011), significant amounts of protein and energy escape via the GI tract without being completely digested; thus, the addition of exogenous protease optimized feed protein and energy digestibility and utilization (Mohammadigheisar and Kim 2018; Ndazigaruye et al. 2019). Dietary protease enhanced BWG and the efficiency of feed digestion by improving amino acid content and energy utilization (Kaczmarek et al. 2014). It also reduces proteolytic fermentation, bacterial toxins, and the number of nutrients excreted in faeces, reducing potential pollutants in the environment (Mahmood et al. 2017). Improved digestion, utilization, and bio-availability of amino acids were reported when protease cleaved trypsin inhibitor (anti-nutrient) (Ndazigaruye et al. 2019). Better growth performance reported for broilers fed PROT25 may

Table 8. The effect of dietary protease supplementation on meat quality attributes of broiler birds.

Treatment	PROT0	PROT10	PROT15	PROT20	PROT25	SEM	p-value
Colour							
a*(redness)	1.32	1.08	1.44	1.14	1.62	0.00	0.11
b*(yellowness)	15.92	15.64	14.98	16.81	14.15	0.13	0.31
L*(lightness)	49.87	46.96	46.97	49.41	49.27	0.67	0.20
Cooking loss (%)	17.53 ^a	16.96 ^a	14.83 ^b	13.98 ^b	13.26 ^{bc}	0.13	0.03
Shear force (N)	8.32	9.37	9.60	9.81	9.87	0.07	0.09
Drip loss (%)	8.12	7.89	6.74	5.63	5.10	0.02	0.07
WHC (%)	18.17 ^b	19.10 ^{ab}	19.81 ^{ab}	20.29 ^{ab}	21.92 ^a	0.25	0.02
pH ₀	6.04	6.05	6.16	6.24	6.30	0.02	0.23
pH _u	5.64	5.72	5.58	5.73	5.67	0.02	0.15
Temp1 (°C)	34.75	34.54	34.55	35.19	34.67	0.46	0.22
Temp2 (°C)	11.24	11.56	11.64	11.79	11.69	0.11	0.10

Notes: ^{a,b,c} Row means with different superscripts differ significantly. SEM = Standard error of the mean. WHC = water holding capacity. PROT0 = Basal Diet: BD (without PROT; protease). PROT10 = BD + 1 g PROT. PROT15 = BD + 1.5 g PROT. PROT20 = BD + 2 g PROT. PROT25 = BD + 2.5 g PROT. Five birds from each cell/replicate were used for the analysis.

result from increased protein digestibility and capacity for energy utilization. Also, it may be due to alterations in intestinal morphological traits (Cowieson, Lu, et al. 2017) or other gut health indicators, such as volatile fatty acids (VFA) and secretory immunoglobulin A (IgA).

4.4. Apparent nutrient digestibility

Improved feed efficiency is of vital importance in animal nutrition because it reduces environmental pollution from poultry and production costs. Poultry cannot digest certain compounds in the feedstuffs, and these compounds may interfere with the animal's digestive system (Rada et al. 2016). These problems were made evident or noticed because animals cannot endogenously produce the necessary enzymes to degrade these compounds (Khattak et al. 2006), restricting the full release of nutrients to the animal. However, *Bacillus* protease was used in this study to reduce the total nutrient excretion, which results from incomplete digestion. In the present study, *Bacillus* protease inclusion improved the digestibility of crude fibre, crude protein, and dry matter but not crude fat. Improved nutrient digestibility traits enhance growth performance, reduce nitrogen excretion, and improve overall gut health due to reduced putrefaction in the distal intestine (Walk et al. 2019). The higher nutrient digestibility values recorded in this study for birds fed PROT25 may be due to the positive effects of the gastrointestinal tract fermentation by lowering gastric acids, decreasing the pathogenic microbial activity, and enhancing mucosal architecture or structure (Scott 2002; Akinola et al. 2015). Furthermore, fermentation results to the catabolism of feed constituents, reducing the viscous nature of grain-legumes, synthesizing B-vitamins, and increasing mineral extractability (Badau et al. 2005). According to Angel et al. (2011); and Liu et al. (2013), the inclusion of protease in poultry diets improved amino acid digestibility. They reported beneficial impacts of exogenous protease on amino acid and crude protein digestibility which might be attributed to their capacity to target protease inhibitors and/or target the cereal portion of the diet (Liu et al. 2013). It has been suggested that protease supplementation may help neutralize anti-nutritional factors such as antigenic proteins, trypsin inhibitors, and lectins; improving protein digestibility (Douglas et al. 2000). Previous studies (Cowieson and Ravindran 2008;

Amerah et al. 2017) also found that supplementary protease improved protein digestibility and nitrogen retention. Nevertheless, Campasino et al. (2015) reported that protease supplementation did not improve protein digestibility at 10% inclusion level. Surprisingly, Crude Protein digestibility was improved at 15% protease inclusion level. According to Kim et al. (2020) Supplementation of protease has been employed in diets to reduce crude protein levels in order to minimize dietary protein waste, nitrogen excretion, and nitrogen emission into the environment without compromising bird performance. In the past, (Olukosi et al. 2015; Opoku et al. 2015; Hussain et al. 2019) protease addition has led to improved nitrogen digestibility when an alternate raw material with more indigestible fraction was added. Walk et al. (2019) and Brown et al. (2020) recorded improvements on BWG and FCR of chicken and attributed the results to improved digestibility of protein caused by dietary protease. It improved the crude protein and amino acid digestibility (Cowieson and Roos 2013) and enhanced the ileal digestibility of energy (Kalmendal and Tauson 2012). According to Cowieson, Zaefarian, et al. (2017), a beneficial effect of exogenous protease on mucin secretion, intestinal integrity, and immunity was recorded. Freitas et al. (2011) and Akinola et al. (2015) found a significant effect of protease supplementation only in the dry matter, while Kamel et al. (2015) and Zuo et al. (2015) recorded an improved crude protein digestibility. This study showed that *Bacillus* protease could increase dry matter, crude fibre, and crude protein digestibility, thus affecting growth performance (Table 4). Dietary protease increased the digestion of many essential amino acids in chicken (Allouche et al. 2015; Kamel et al. 2015).

4.5. Gut micro examination

There is a synergy between gut micro-flora that colonized the gastrointestinal tract and their host at the early post-hatch period. The gut micro-flora performs an essential task in the health and nutrition of the host by encouraging digestion and absorption of nutrients, preventing the migration of the pathogen, and sustaining normal mucosal immunity. The significant traits to explain the microbial structure and diversity are the richness and evenness of the bacteria. According to Kiarie et al. (2014) and Leeming et al. (2019), there is a general acceptance that diet manipulation can influence the

microbial structure and diversity of the intestine. Exogenous enzymes can control the population of microbes due to the changes they impose in the gut content (Hubner et al. 2002; Masey O'Neill et al. 2014; Oyeagu et al. 2019b). The changes recorded in the present study for bacteria populations in the ileum and caecum sections of the gut are not consistent with the report of Gao et al. (2008), who recorded no effect for *Lactobacillus* and Coliform counts in the caecum of 21d old birds. Luo et al. (2009) reported no difference in the population of *E. coli*, *Lactobacillus*, and total aerobes in the ileum and caecum of chicken at 42d of the feeding trial. There were increased *Lactobacillus* and *Bifidobacteria* counts in both ileum and caecum parts of the gut as the level of *Bacillus* protease increased. Also, there was a drop/decrease in *E. coli* counts as the level of *Bacillus* protease increased in both the ileum and caecum parts of the intestine. These observations may be due to the ability of the enzyme to increase the levels of available substrate for microbial fermentation, which improves the digestibility of protein and the production of short-chain fatty acid (SCFA) in the GI tract. This process typically occurs due to the non-digested protein diet available to the intestinal microbes (Morrison and Preston 2016). A similar observation was reported by Nabizadeh et al. (2017). Dietary *Bacillus* protease influenced the proliferation of *Lactobacillus* and *Bifidobacteria* populations in the present study. These beneficial bacteria species are found throughout the digestive tract, predominantly in the small intestine. They are thought to contribute to nutrient absorption and are involved with bile salt hydrolysis (Xiao et al. 2017). *Lactobacillus* and *Bifidobacteria* are gram-positive and facultative anaerobic, and are fastidious with complex nutritional requirements, including fermentable carbohydrates, amino acids, peptides, vitamins, salts, and fatty acids (Staley et al. 2017). Apajalahti and Vienola (2016) hypothesized that protease would decrease *Lactobacilli* located in the small intestine; however, our data do not support this hypothesis, as *Bacillus* protease used in the present study increased the proliferation of *Lactobacillus* and *Bifidobacteria* in both caecum and ileum sections of the gut. Jozefiak et al. (2010) and Yadav and Jha (2019) reported that external protease reduced the viscosity in the intestine; it also helps developing competitive microbial communities and even higher intra-microbial competition. It diminishes microbial interference in nutrient absorption and contributes to the potential reduction of pathogenic microbes with improved growth traits. Dietary factors can influence the bacteria population. Some authors (Sugiharto and Ranjitkar 2019) reported a higher population of *E. coli* and *Lactobacillus* in the digesta of broilers that consumed wheat and barley than those that consumed corn diets. As the intestinal bacteria of chicken changes, it may alter the mucosal structure, which may subsequently influence the absorptive capacity of the nutrient (Munyaka et al. 2016). An increase in the SCFA density stimulates a gradual reduction of the proliferation rate of *Enterobacteria* but not that of *Lactobacillus* and *Bifidobacteria* (Van der Wielen et al. 2000; Nabizadeh et al. 2017). Inclusion of *Bacillus* protease influenced the production of SCFAs which are end products of bacterial fermentation of carbohydrates and amino acids (Jeferson et al. 2020). The SCFAs encourage the proliferation of beneficial microbes (such as *Lactobacillus*) and

prevent the inflammation of the gut (Morrison and Preston 2016). The majority of short-chain fatty acids are produced by the fermentation of carbohydrates and amino acids that escaped digestion in the small intestine. Some gastrointestinal microbes ferments amino acids and can deaminate them rapidly (Lee et al. 2018). Exogenous enzymes balance chicken intestinal bacteria, which may affect the host's health status and the extent of digestion (Bedford and Cowieson 2012; Alagawany et al. 2018; Sudhir and Rajesh 2019).

4.6. Carcass traits

Our study confirms that the *Bacillus* protease significantly affects the carcass, drumstick, thigh, and breast weights and carcass yield (dressing percentage), drumstick yield, thigh yield, and breast yield of broiler chickens. Improved carcass yield is as a result of adequate diet and chicken nutrition (Cardoso et al. 2011). For example, livestock will deposit more tissue (muscle) when they consume balanced diets. Dietary *Bacillus* protease at 2.5 g/kg offers improved nutrient digestibility and increases nutrients utilization for enhanced development of the muscle (tissue) and breast growth since breast cut yield accounts for approximately 40% of aggregate carcass yield (Oyeagu et al. 2019b). The addition of protease to feed improved poultry growth rates, carcass yield and caused some changes in the diversity of the broiler gut micro-biota. The presence of protease in the feed improved protein absorption, which increased the live body weight and cut yield of broiler chickens by 0.5 kg, and also reduced feed intake (Daria et al. 2020). Comparable to our results, Dalólio et al. (2015) confirmed the impact of dietary enzyme complexes on the carcass yield of chicken. According to Marapana (2016), dressing percentage and meat yield of different chicken body parts could be altered by diets. The increased yield of carcass, drumstick, thigh and breast meat of broilers fed with the PROT25 diet compared to other dietary treatments in our study may be due to the increased available nutrients for the development of tissue (retail cut yields). Dietary exogenous enzymes enhanced diet digestibility, substrates availability, and increased absorption of nutrients, which positively impacted muscle development (Oyeagu et al. 2019b). Dietary external enzymes such as amylase, xylanase, or protease boost the actions of the pancreatic and intestinal enzymes determined at 14 and 42-d-old chickens (Horvatovic et al. 2015). Intestinal enzymes play a crucial role in breaking down feeds, while pancreatic enzymes break down sugars, fats, and starches. These proteins speed up chemical reactions that turn nutrients into available substances for absorption. Dietary barley, maize, sorghum, and wheat without enzymes negatively impact intestinal enzyme activities, unlike enzyme supplemented diets (Shakouri et al. 2008). According to Yadav and Sah (2005) and Barletta (2011), microbial protease could break down stored proteins and proteinaceous anti-nutrients in diets. It was evident from the slaughter weight, carcass weight, and carcass yield (dressing percentage) that chickens with heavier muscle have a higher dressing percent and higher cut-yield than lighter chickens (Table 5). According to Wilfarta et al. (2007) and Sayehban et al. (2015), dietary fibre was implicated in the fast passage rate of the digesta which

impairs broiler growth. The better performance of birds fed with the PROT25 diet in our study may be due to the slow digesta passage rate, which offers maximum opportunity for increased bio-activities of *Bacillus* protease in the gut. As the slow passage rate of fibre digesta persists in the intestine, it provides greater chances for improved feed digestion and absorption with higher muscle deposit which influences the cut yield percentage (Sayehban et al. 2016). It was reported that broilers with higher muscle deposit produces higher cut-yield percentage (Al-harathi 2016; Oyeagu et al. 2019b). The range of dressing percentage (77.19% to 82.24%) recorded in the present study is above the threshold reported by Schwehofer (2011). They noted that the average dressing percentage (carcass yield) of broiler chicken is 71%. Breast meat yield is one of the carcass components with the highest economic value. Breast meat continuously increases as a percentage of body weight in our study. A similar result was reported by Petracci et al. (2004) and Kanagaraju et al. (2019). The success of poultry meat production may be attributed to improvements in growth and carcass yield, mainly by increasing the proportion of chicken breast part. Some chicken parts such as the thigh, drumstick, and breast have higher commercial value (Marapana 2016). The sale of a whole chicken and its cut-up parts allows the consumers to have different cooking and taste choices (Cevger et al. 2004; Kaygisiz and Cevger 2010). The demand for high-quality cut-up parts at meat shelves in the shops has increased, and it has also influenced the poultry industry to change its marketing strategies. Today, the most marketed form of chicken is always the cut-up parts. However, the yield of breasts, thighs, and broiler drumsticks has become critical to processors (Young et al. 2001). Due to increased aggregate value, there is a higher tendency to sell cut-up parts more than the whole carcass. Silveira et al. (2010) reported a 25% leg quarter yield increase for birds fed dietary protease at 21 days of age than those fed the control diet. The authors reported no effects on breast meat yield. According to Cardoso et al. (2011), there was no difference in carcass yield at day 42 of chickens fed dietary multi-enzyme. Our results are similar to the findings of Alam et al. (2003), Wang et al. (2005), and Hajati (2010). They reported an increased carcass and breast yield of birds-fed dietary enzymes (xylanase + protease, phytase + protease, and xylanase + protease + amylase) due to improved digestion, higher nutrient availability, and absorption.

4.7. Meat quality traits

Chicken meat has specific sensory attributes that increase consumers' acceptance, such as appearance, texture, succulence, and flavour (Petracci and Baéza 2011). The most important factor affecting consumer choice is broiler meat colour (Werner et al. 2009). Our results showed no significant change in the chicken meat colour. Meat quality traits depend directly on environmental and management conditions, mainly the feed offered (Neves et al. 2014). Dietary enzymes influence efficient digestibility and promote muscle fibre deposit and whole carcass yield (Allouche et al. 2015). Increased protein deposition may adjust muscle fibre type and shape, altering meat quality parameters (Fatufe et al.

2004; Dalólio et al. 2016). Our study showed that the inclusion of *Bacillus* protease increased the values of WHC and decreased the CL of the breast meat. WHC concerns the ability of the meat to retain its water upon application of an external force (Omojola et al. 2014), and it is a primary indicator of the juiciness of the meat. Dietary *Bacillus* protease reduced CL which implies reduced water loss during cooking. Omojola et al. (2014) reported that meat with increased water loss (exudate) results in more nutrient loss and should be considered poor-quality meat. Dietary *Bacillus* protease used in our study reduced nutrient loss while also increasing the cooking yield of chicken. This is attributed to the specific hydrolyzing activities of *Bacillus* protease (from *Bacillus strain EA1*). It may be that microbial protease (modified or wild) from non-pathogenic microbes improved meat quality (Alaa et al. 2014). On the contrary, the traditional plant protease has broad specificities and breaks down connective tissue and myofibrillar proteins indiscriminately (Ashie et al. 2002; Madhusankha and Thilakarathna 2021). It leads to undesirable attributes in the tenderized meat (i. e., mushy texture, high water loss, bitterness, and off-flavour). The findings of our trial are in contrast to the studies of Werner et al. (2009), Zakaria et al. (2010), and Dalólio et al. (2015); they did not observe any effect of exogenous enzyme addition on meat quality parameters such as pH, WHC, colour, CL, shear force among others. According to Omojola et al. (2014), the cooking yield of meat is dependent on cooking loss, while decreased cooking loss results to higher WHC. The present study showed that supplementing *Bacillus* protease at a 2.5 g/kg diet improved performance compared to other treatment groups.

The dietary level (2.5 g PROT/kg feed) confirms the maximum/highest activities of *Bacillus* protease on the substrate, reflecting the efficient utilization of protein and energy and the overall performance of chickens. Since the peak growth performance of broiler birds was observed at the highest level of *Bacillus* protease supplementation (2.5 g PROT/kg feed), further study is required to evaluate the effects of *Bacillus* protease on the performance of broilers beyond 2.5 g PROT/kg feed (PROT25) inclusion. Nevertheless, our findings revealed that the inclusion of *Bacillus* protease in maize-soybean meal-based diets improved the health status, encouraged the development of gut histology, and provided an excellent gut microbiota environment for chickens (Khan and Naz 2013). We concluded that broiler chickens respond positively to supplementation with *Bacillus* protease concerning meat quality traits. The gut microflora and intestinal histological characteristics were improved with the highest inclusion of *Bacillus* protease (Abudabos et al. 2017). In other words, the highest inclusion of *Bacillus* protease improved nutrient digestion and utilization as it protects the gut's integrity, which helps stabilize the health status of chickens. Summarily, birds fed 2.5 g *Bacillus* protease/kg feed (PROT25) produced a better FCR and higher weight gain, retail cut yields, carcass yields, and improved gut integrity.

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