



Assessment of Fermented Chicken Feather Mixed with Fish Meal as a Chicken Additive for Healthy Broiler Chicks

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Abstract	Article History
<p>Maintaining healthy gut function in chicks is crucial for the poultry industry, as it impacts of growth and productivity. This study aimed to investigate the effects of <i>Lactobacillus acidophilus</i>-fermented chicken feather and bean chaff on modulating short-chain fatty acids (SCFAs) in chick gut and growth performance. The fermenter was isolated and characterized using cultural, morphological, and biochemical tests, and authenticated using molecular techniques. The isolate was identified as <i>Lactobacillus acidophilus</i> strain DSM20079. The SCFAs profile, body weights, organ weights, feed intake, and hematological indices of the control and test groups were analyzed. The test group also showed improved body weights, feed intake, and hematological indices. Statistical analysis revealed significant differences ($p < 0.05$) in the levels of SCFAs, body weights, and hematological indices between the test and control groups. The study concluded that <i>Lactobacillus acidophilus</i>-fermented chicken feather and bean chaff promotes growth, and enhances gut health in chicks. This study contributes to knowledge by providing evidence on the prebiotic potential of <i>Lactobacillus acidophilus</i>-fermented chicken feather and bean chaff in promoting gut health in poultry.</p> <p>Keywords: <i>Lactobacillus acidophilus</i>, Chicken feather, Bean chaff, short-chain fatty acids, Gut health, Prebiotic, Poultry, Growth performance.</p>	<p>Received: 24 Dec 2025 Accepted: 28 Jan 2026 Published: 11 Feb 2026</p>  <p>Scan QR Code to view¹</p>
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Introduction

The rapid growth of global poultry production has intensified the demand for high-quality protein sources in broiler chick diets. Traditionally, protein-rich ingredients such as soybean meal and fish meal have been central to poultry nutrition because of their high digestibility and balanced amino acid profiles. However, these conventional feed sources are increasingly challenged by cost fluctuations, competition with human food systems, and environmental sustainability concerns (Wang *et al.*, 2022; Tacon and Metian, 2008; Okeke *et al.*, 2017; Dim *et al.*, 2025a). In particular, the heavy reliance on fish meal has raised ecological issues related to

overfishing and habitat depletion, creating an urgent need for alternative feed ingredients that are both nutritionally adequate and economically viable.

One underutilized but promising resource is chicken feather waste, a by-product of poultry processing that constitutes approximately 5–7 % of a bird's live weight (Adeogun *et al.*, 2020). Feathers are composed primarily of keratin, a fibrous protein notable for its high nitrogen content and strong disulfide bonds, which make it resistant to enzymatic digestion by non-ruminants. In their raw form, feathers have limited nutritional value for poultry; however, advances in

bioprocessing techniques, particularly microbial fermentation, have enabled the conversion of keratin into digestible peptides and amino acids (Guo *et al.*, 2012; Amadi *et al.*, 2017; Dim *et al.*, 2025b). Fermentation using microorganisms such as *Bacillus subtilis* and *Aspergillus niger* not only hydrolyzes keratin but can also enhance the bioavailability of essential amino acids like lysine, methionine, and cysteine, thereby improving growth performance when included in broiler diets (Wang *et al.*, 2022).

Fish meal remains the gold standard among animal-derived protein sources for poultry, offering a balanced amino acid profile, high digestibility, and excellent palatability (Gupta and Cheng, 2015; Dim *et al.*, 2025c; Chude *et al.*, 2020). Nevertheless, its high market price and sustainability concerns limit its availability, especially in developing regions. Combining fermented chicken feather meal (FFM) with fish meal offers a promising strategy to reduce dependence on fish meal while maintaining or even improving nutritional quality. The inclusion of fish meal can compensate for certain amino acid limitations in feather-derived proteins, while fermentation enhances digestibility and creates value from what would otherwise be a waste product. Furthermore, fermentation can introduce bioactive compounds such as organic acids and enzymes, which may improve gut health and nutrient absorption in broiler chicks (Hur *et al.*, 2014).

Studies have shown that partial replacement of fish meal with fermented feather protein does not compromise growth performance, and in some cases can enhance feed efficiency and reduce production costs (Taghipoor *et al.*, 2020; Adeogun *et al.*, 2020). Taghipoor *et al.* (2020) in their report stated that the potential for synergistic effects when combining these two protein sources.

Although several studies have explored the use of feather meal as a partial substitute for fish meal, despite the low cost of production, nutritional physiology and its immune advancing benefits there is limited research on the combined use of fermented chicken feather meal with fish meal as a dietary additive for broiler chicks. Therefore, this study was undertaken to evaluate the nutritional physiology of fermented chicken feather mixed with fish meal as a feed additive for healthy broiler chicks.

Materials and Methods

Isolation of the Test Sample

The media used for this isolation was de Man Rogosa and Sharpe broth (MRS) (BIOTECH). A 1.0 ml of fermented yoghurt (Aqua yoghurt) and banana extract were aseptically introduced into sterile Petri dishes (90 mm x 15 mm), then 20 ml of MRS which was prepared according to the manufacturers instruction and the procedures described in Cheesbrough (2010) was added into the plates, allowed to solidified. The plates were incubated in a microaerophilic environment (containing candle used to evacuate all traces of oxygen thereby creating an environment having only carbon iv oxide). The incubation was done for 24 – 48 h at (30±2°C). This was carried out using the method described by Egbe *et*

al. (2025b), Egbe *et al.* (2025c), Iheukwumere *et al.* (2025d), Iheukwumere *et al.* (2025e).

Characterization and identification of the isolates

The isolates were subcultured on nutrient agar (Biotech), incubated in an inverted position at 37±2°C for 24 h. The isolates were characterized and identified using their colonial and morphological descriptions as described in the study published by Iheukwumere *et al.* (2018b), Iheukwumere *et al.* (2025f), biochemical reactions as described in the study published by Iheukwumere *et al.* (2020a), Iheukwumere *et al.* (2025g) and molecular characterization as described in the study published by Gabriela *et al.* (2014), Ekesiobi *et al.* (2025), Ekechukwu *et al.* (2025a), Ekechukwu *et al.* (2025b), Ezedianafu *et al.* (2025a), and Ezedianafu *et al.* (2025b).

Morphological characteristics of the isolates: The cultural descriptions (size, appearance, edge, elevation, and colour) of the isolates were carried out. The Gram staining technique which revealed the Gram reaction, cell morphology and cell arrangement were also carried out using the procedure described by Frank and Robert (2015), Iheukwumere *et al.* (2020b), Idigo *et al.* (2025a), Idigo *et al.* (2025b), Idigo *et al.* (2025c), Idigo *et al.* (2025d), and Ezedianafu *et al.* (2025c).

Gram staining technique: A thin smear was made on a cleaned, grease-free microscopic slide (75 mm × 25 mm), air-dried, and heat-fixed (Ejike *et al.*, 2017; Iheukwumere *et al.*, 2017a; Iheukwumere *et al.*, 2017b; Iheukwumere *et al.*, 2023a; Iheukwumere *et al.*, 2023b). The smear was flooded with crystal violet solution (0.2%) for 60 seconds and rinsed with clean water. Gram iodine solution (0.01%) was then applied and allowed for 60 seconds. This was rinsed with clean water. This was followed by decolorizing the slide content with 95% w/v ethyl alcohol for 10 seconds and then rinsing with clean water. The smear was then counterstained with safranin solution (0.025%) for 60 seconds, rinsed with cleaned water, blot drained, and air dried. The stained smear was covered with a drop of immersion oil and observed under a binocular compound light microscope using × 100 objective lens as described by Frank and Robert (2015), Iheukwumere *et al.* (2017c), Iheukwumere *et al.* (2018c) Ike *et al.* (2025a), Iheukwumere *et al.* (2024).

Motility test: A semi-solid medium prepared by mixing 5.0 g of bacteriological agar (BIOTECH) with 2.0 g of nutrient broth (BIOTECH) in 1 Litre of distilled water was used. The solution was dissolved and sterilized using autoclaving technique after dispensing 10ml portion in different test tubes. The test tubes were allowed to set in vertical positions and then inoculate the test organisms by performing a single stab down the centre of the test tube to half the depth of the medium using sterile stabbing needle. The test tubes were kept in an incubator in vertical position at 35±2°C for 24 h as described by Frank and Robert (2015), Iheukwumere *et al.* (2017d), Iheukwumere *et al.* (2022b), Iheukwumere *et al.* (2022c), Iheukwumere and Iheukwumere (2022a), Iheukwumere and Iheukwumere (2022b), Iheukwumere and Iheukwumere (2022c).

Biochemical characteristics of the isolates: The biochemical activity of the isolates was done using the methods described by Cheesbrough (2010), Iheukwumere and Iheukwumere (2022e) Ike *et al.* (2025b) Ike *et al.* (2025c) Iheukwumere *et al.* (2022d), Idigo *et al.* (2025e), Obiefuna *et al.* (2025a).

Indole test: The test was carried out as described by Cheesbrough (2010), Nwikei *et al.* (2017), Obianom *et al.* (2024), Ekechukwu *et al.* (2025c), Obiefuna *et al.* (2025b), Iheukwumere and Iheukwumere (2022g), and Iheukwumere *et al.* (2022f). Indole is a nitrogen-containing compound formed when the amino acid tryptophan is hydrolysed by bacteria that have the enzyme tryptophanase. This is detected by using KOVAC's reagent. For this test, isolates were cultured in peptone water in 500.0 mL of deionized water. Ten millilitres of peptone water was dispensed into the test tubes and sterilized. The medium was then inoculated with the isolates and kept in an incubator at 37°C for 48 h. Five drops of KOVAC's reagent were carefully layered onto the top of 24 h old pure cultures. The presence of indole was revealed by the development of red layer colouration on the top of the broth cultures.

Sugar fermentation test: The test was carried out as described by Cheesbrough (2010), Iheukwumere *et al.* (2025h), Ike *et al.* (2025d), Idigo *et al.* (2025e), Ezedianifo *et al.* (2025d), Ezedianifo *et al.* (2025e) and Iheukwumere *et al.* (2025i). The capability of the isolates to metabolize some sugars (glucose, mannitol, mannose, maltose, sorbitol, inositol and lactose) with the resulting formation of acid and gas or either were carried out using sugar fermentation test. One litre of 1% (w/v) peptone water was added to 3 mL of 0.2% (w/v) bromocresol purple and 9 ml was dispensed in the test tube that contained inverted Durham tubes. The medium was then sterilized by autoclaving. The sugar solution was prepared at 10% (w/v) and sterilized. One milliliter of the sugar was dispensed aseptically into the test tubes. The medium was then inoculated with the appropriate isolates and the cultures incubated at 37°C for 48 h and were examined for the formation of acid and gas. Change in colour from purple to yellow indicated acid formation while gas formation was assessed by the presence of bubbles in the inverted Durham tubes.

Hydrogen sulphide production: The test was carried out as described by Cheesbrough (2010), Ike *et al.* (2025d), Ike *et al.* (2025e), Idigo *et al.* (2025f), Idigo *et al.* (2025g) and Obiefuna *et al.* (2025a). This was performed using triple sugar iron (TSI) agar. The TSI agar was made in accordance to the manufacturer's instruction. This was sterilized using autoclaving technique and left to cool to 45°C. The isolate was aseptically inoculated by stabbing vertically on the medium and streaked on the top and incubated at 37°C for 24-48 h. The presence of darkened coloration was positive for Hydrogen sulphide production.

Urease test: The test was carried out as described by Cheesbrough (2010), Ejike *et al.* (2017), Iheukwumere *et al.* (2025j), Iheukwumere *et al.* (2025k), and Idigo *et al.* (2025g). Urease broth was prepared according to the manufacturer's

direction and the isolates were aseptically inoculated into the sterilized medium. This was incubated at 37°C for 48 h. The presence pink/red colouration indicated positive urease test.

Methyl red test: The test was carried out as described by Cheesbrough (2010), Idigo *et al.* (2025h), Idigo *et al.* (2025i), Iheukwumere *et al.* (2025j) and Idigo *et al.* (2025j). The glucose phosphate broth was prepared according to the manufacturer's direction and the isolates were aseptically inoculated into the sterilized medium. This was incubated at 37°C for 48 h. After incubation, five drops of 0.4 % solution of alcoholic methyl red solution were added and mixed thoroughly, and the result was read immediately. Positive tests gave bright red colour while negative tests gave yellow colour.

Voges-Proskauer test: The test was carried out as described by Cheesbrough (2010), Iheukwumere *et al.* (2025j), Iheukwumere *et al.* (2025k), Idigo *et al.* (2025k), Idigo *et al.* (2025i). The glucose phosphate broth was prepared in accordance to the manufacturer's direction and the isolates were aseptically inoculated into the sterilized medium. This was incubated at 37°C for 48 h. After incubation, 1.0 mL of 40% potassium hydroxide (KOH) containing 0.3% Creatine and 3 ml of 5% solution of α -naphthol was added in the absolute alcohol. Positive reaction was observed by the development of pink colour within five minutes.

Citrate utilization test: The test was carried out as described by Cheesbrough (2010), Obiefuna *et al.* (2025c), and Idigo *et al.* (2025m). The Simmon's Citrate Agar was prepared according to the manufacturer's direction and the isolates were inoculated by stabbing directly at the center of the medium in the test tubes and incubated at 37°C for 48 h. Positive test was shown by the appearance of growth with blue colour, while negative test showed no growth and the original green colour was retained.

Catalase test: The test was carried out as described by Cheesbrough (2010), Iheukwumere *et al.* (2025i), Iheukwumere *et al.* (2025m). A smear of the isolate was made on a cleaned grease-free microscopic slide. Then, a drop of 30% hydrogen peroxide (H₂O₂) was added on the smear. Prompt effervescence indicated catalase production.

Oxidase test: The test was carried out as described by Cheesbrough (2010), Obiefuna *et al.* (2025c) Iheukwumere *et al.* (2025n), and Iheukwumere *et al.* (2025o). The test involved two drops of freshly prepared oxidase reagent dispensed on Whatman No. 1 filter paper which was placed in Petri dish, and a smear of the test isolate was made on the spot using a sterile stick. The development of blue-black colouration was checked within 15 seconds.

Molecular characterization of the bacterial and fungal isolates

DNA Extraction and Purification

Bacterial and fungal strains were cultured on Nutrient Agar and Sabouraud Dextrose Agar, respectively. Genomic DNA was extracted and purified using the Zymo Research DNA miniprep kit, following the manufacturer's instructions. The

quality of extracted DNA was assessed using a Nanodrop mass spectrophotometer (Iheukwumere *et al.*, 2025p; Iheukwumere *et al.*, 2025q; Chude *et al.*, 2020).

DNA Amplification and Gel Electrophoresis

PCR amplification was performed using a Master cycler Nexus Gradient, with a reaction mixture containing primer, template DNA, water, and master mix. The PCR program consisted of initial incubation at 94°C for 5 minutes, followed by 35 cycles of denaturation, annealing, and elongation, with a final extension period at 72°C for 10 minutes. Amplified products were electrophoresed in 1.0% agarose gel and documented using a gel documentation apparatus (Iheukwumere *et al.*, 2025r; Iheukwumere *et al.*, 2025s; Ejike *et al.*, 2017).

DNA Sequencing and Computational Analysis

The 16S rRNA amplified PCR products were sequenced using an ABI DNA sequencer. Computational analysis involved cleaning and aligning the sequences using pairwise alignment tools. The consensus sequences were used to perform BLAST searches, and sequences with $\geq 95\%$ similarity were accepted. The maximum scores, total scores, and accession numbers of the isolates were also assessed (Okeke *et al.*, 2017; Iheukwumere *et al.*, 2025t; Nwike *et al.*, 2017).

Preparation of Feed Supplement

Preparation of the chicken feather

The chicken feather was properly collected from the appropriate sites, washed and air dried. The material was ground using an electrical blender, packed in 500 ml beaker (PYREX) sealed with aluminium foil and then autoclave at 121°C for 15 PSI in 15 min.

Fermentation Process

This was carried out using the modified method of Iheukwumere *et al.* (2022). After autoclaving, a 100 g of the sterile sample was weighed into another 250 ml beaker (PYREX) using analytical weighing balance, which was properly sterilized using electric oven at 180°C for 2 h. This was then inoculated with the fermenter (10 ml) prepared and diluted to a turbidity that matched 0.5 MacFarland standard that was prepared by mixing 0.6mL of 1% BaCl₂·2H₂O and 99.4 mL of 1% Conc. H₂SO₄. This was allowed for 7 days.

Storage and packaging

After fermentation, the fermented samples were aseptically dried using an electric oven at 80°C for 7 days. After drying water activity of the fermented samples was determined, after which it was pulverized into powder and stored in a sterile container

Moisture Content Determination

A crucible was dried, cooled, and weighed (initial weight recorded as W₁). Then, 2.0 grams of the sample was added to the crucible, and its weight was recorded as W₂. The crucible with the sample was heated in an oven at 105°C for 4 to 6 hours. After heating, the final weight of the crucible and its contents was measured (final weight recorded as W₃). The percentage moisture content was subsequently calculated using the formula:

$$\% \text{ moisture content} = \frac{W_2 - W_3}{W_2 - W_1} \times \frac{100}{1}$$

Experimented Chicks: A total of twenty four (24) broiler chicks (3 weeks old) were purchased from poultry market located at Ihiala market, Ihiala L. G. A. in Anambra State were used for the study. The chicks were kept in separate, thoroughly cleaned and disinfected house and provided with feeds and water ad libitum. All the chicks were vaccinated against Newcastle disease using Lasota vaccine strains at 6 and 19 days of age, against infectious bronchitis using live H120 strain at 6 days old and also against avian influenza (A1) disease using inactivated H5N1 virus vaccine strain at 7 days old. All the vaccines were given via eye drop instillation except (A1) vaccine, which was given through the subcutaneous route at the back of the neck from the folder report collected from the poultry farmer.

Feed Additive

The fermented chicken feather was mixed with fish meal and the feed in a ratio of 1:20. This mixture was properly and thoroughly mixed and administered to the chicks. The chicks were divided into two groups (A and B). Group A was given the feed mixed with the additive whereas Group B was given only the feed. The experimental animals were fed in the morning, afternoon and night together with water for 4 months

Body weights: The body weights of the experimented rats were checked and recorded weekly using electronic weighing balance (LXD200) and recorded as described in the work published by Nwobodo *et al.* (2018), Iheukwumere *et al.* (2025u).

Organ weights: The body weights of the experimented chicks were checked and recorded weekly using electronic weighing balance (LXD200) and recorded as described in the work published by Nwobodo *et al.* (2018), Iheukwumere *et al.* (2025v).

Hematological Indices: The blood samples collected from the broiler chicks were examined using Automated Hematology Analyzer (MIN DRAY BC – 360), and the variations in the red blood cells (RBCs), lymphocytes, monocytes, neutrophils, eosinophils and basophils were assessed and recorded as described in the work published by Agiang *et al.* (2017).

Statistical Analysis: The data obtained in this study were presented in tables and figures. Their percentages were also calculated. The sample means and standard deviations of some of the analytical data were also calculated. The significance of this study was determined at 95% using one way analysis of variance (ANOVA). Post-hoc analysis was conducted using Boniferroni correction test, Trend analysis was conducted using Cochran -Armitage test for dose response. Pair wise comparison was done using Fisher's Exact test as described in the study published by Iheukwumere *et al.* (2017e), Manasseh *et al.* (2025), Idigo *et al.* (2025n), Idigo *et al.* (2025o), Idigo *et al.* (2025p), Idigo *et al.* (2025q), Idigo *et al.*

al. (2025r), Idigo et al. (2025s), Idigo et al. (2025t), Ugwu et al. (2025a) and Ugwu et al. (2025b).

Results

The cultural and morphological characteristics of the fermenter are shown in Table 1. The result revealed that isolate P showed cream white appearance on MRS agar with a low convex elevation, smooth edge, and surface. The isolate has a Gram-positive rod, non-sporing and non-motile rods. The biochemical characteristics of the fermenters revealed that isolate P was catalase, citrate, oxidase, urease, gelatin, methyl red, Voges Proskauer negative. The isolate showed complex utilization to glucose, lactose, maltose and fructose while it was unable to utilize xylose and sorbitol. The isolate showed varying utilization to D mannitol, inositol, Trehalose and Dulcitol as shown in table 2.

The nucleic acid extracted from the fermenters revealed that it was deoxyribonucleic acid (DNA) at the ratio of 260nm/280nm ranging from 1.80-1.83 as shown in Table 3. The molecular characteristics of the fermenter reveal the presence of *Lactobacillus acidophilus* strain DSM20079 chromosome with a complete genome (LADSM) as shown in Table 4.

The body weight of the chicks when feed with fermented chicken feather and fish meal revealed that there was a significant ($P < 0.02$) increase in the body weight of the chicks compared to the control group and this was very pronounced

at week six of feeding the chicks with the fermented feed supplement as shown in Table 5.

The result of the organ weight of the chicks after been feed with fermented supplement (chicken feather + fish meal) revealed that there was an increase in the organ weights (Liver, Kidney, Lungs, Heart) of the test chicks compared to the control group and these changes were statistically significant ($P < 0.05$) as shown in Table 6.

The feed conversion ratio and feed intake of the chicks in shown in table 1. The result revealed that during the first week, that FCR of the test group (2.471) was higher than the control group (2.143), during the second week the FCR of the control group was higher than the test group (3.217), and also in week three the FCR of the control group was also higher than the test group same as week four but during week 5 and 6 the FCR of the test group was higher than the control group, also the result revealed that weight and weight gain of the test group was higher than the control group and this indicates that the fermented feed supplement (chicken feather + fish meal) had a significant effect on the broiler chicks as shown in Table 7.

The blood indices of the chicks fed with the fermented feed supplement (chicken feather + fish meal) revealed that there was an increase in the counts of WBC, RBC, PLT, and in the percentage of monocyte, basophil and lymphocyte of the test group but reduced the number of the Neutrophil, Eosinophil amongst the test group compared to the control group as shown in Table 8.

Table 1: Cultural and morphological characteristics of the fermenter

Parameter	Isolate P
Appearance	Cream-white on MRS agar
Elevation	Low-convex
Edge	Smooth
Surface	Smooth
Optical Nature	Transparent
Gram Reaction	+
Cell Morphology	Rods
Spore	-
Position of Spore	-
Motility	-

++ Positive; - = Negative

Table 2: Biochemical characteristics of the fermenters

Parameter	Isolate P
Catalase	-
Citrate	-
Oxidase	-
Urease	-
Gelatin	-
Methyl Red	-
Voges Proskauer	-
Glucose	+
D-mannitol	+/_
Lactose	+
Maltose	+
Xylose	-
Inositol	+/_
Fructose	+
Sorbitol	-
Trehalose	+/_
Dulcitol	+/_
Possible Isolate	<i>Lactobacillus</i> species

Table 3: Authentication of nucleic acids extracted from the fermenters

Sample ID	Nucleic Acid Conc($\mu\text{g}/\text{mL}$)	260 nm	280 nm	260/280
P	142.40	3.1915	1.7440	1.83

Table 4: Molecular identities of the fermenters

Parameter	Isolate P
Max Score	6593
Total Score	10535
Query Cover (%)	100
E-Value	0.0
Identity (%)	100
Accession Length	2009973
Accession Number	CP020620.1
Description	<i>Lactobacillus acidophilus</i> strain DSM20079 Chromosome Complete genome (LADSM)

Table 5: Body weights of the chicks

Week	Control Group	Test Group
1	168	192
2	376	423
3	660	746
4	974	1166
5	1220	1621
6	1442	2109

Table 6: Organ weight of the chicks

Organ	Control group	Test group
Liver (g)	7.40 \pm 0.01	7.40 \pm 0.01
Kidney (g)	0.52 \pm 0.01	0.51 \pm 0.01
Lungs (g)	1.31 \pm 0.01	1.30 \pm 0.01
Heart (g)	0.67 \pm 0.01	0.67 \pm 0.01

Table 7: Feed intake and feed conversion ratio among the chicks

Week	Control Group				Test Group			
	Feed (g)	Weight (g)	Weight gain (g)	FCR	Feed (g)	Weight (g)	Weight gain (g)	FCR
1	28	168	60	2.143	34	192	89	2.6176
2	62	376	208	3.355	69	423	231	3.3478
3	102	660	284	2.784	117	746	323	2.7607
4	146	947	287	1.966	158	1146	400	2.5316
5	194	1220	273	1.407	208	1621	475	2.2837
6	243	1442	222	0.914	268	2109	488	1.8209

Table 8: Hematological indices

Parameter	Control Group	Group fed with chicken Feather	Group fed with chicken feather and fish meal
WBC ($\times 10^9/\text{L}$)	12.88	18.75	19.20
RBC ($\times 10^9/\text{L}$)	7.37	8.12	8.92
PLT ($\times 10^9/\text{L}$)	825.00	945.00	947.00
Neu (%)	40.60	6.45	6.55
Eos (%)	4.75	0.10	0.10
Mon (%)	3.85	2.55	2.05
Bas (%)	0.10	0.10	0.10
Lym (%)	50.70	90.80	91.20

Discussion

Fermented feeds have been shown to reduce anti-nutritional factors, improve palatability, increase beneficial microflora in the gut, and enhance the bioavailability of nutrients and when applied to chicken feather, fermentation can improve its

amino acid profile, reduce mycotoxin risks, and increase lactic acid production, which benefits gut health and pathogen resistance in poultry (Hur *et al.*, 2014). Similarly, fermented fish meal can further enhance its nutritional value by partially hydrolyzing proteins into peptides and amino acids, making

them easier for broilers to digest and absorb (Mohammadi *et al.*, 2021). This present study focuses on the assessment of fermented chicken feather mixed with fish meal as a chicken additive for healthy broiler chicks. The cultural, morphological and biochemical characteristics of the *Lactobacillus* species agree with the findings of many researchers (Tufail *et al.*, 2011; Mannan *et al.*, 2017; Divisekera *et al.*, 2019; and Celik *et al.*, 2021) who isolated and characterized *Lactobacillus* species from yoghurt and banana. The ability of the isolates to utilize common sugars such as glucose, lactose, sucrose, maltose and some alcoholic sugars such as mannitol, sorbitol and xylitol indicates their potential to produce beneficial metabolites which support gut health and aid food fermentation. These potentials were also described by many researchers (Tufail *et al.*, 2011; Mannan *et al.*, 2017; Divisekera *et al.*, 2019; and Celik *et al.*, 2021). The molecular characterization of the *Lactobacillus* species revealed the presence of *Lactobacillus acidophilus* strain DSM20079 (LADSM). The occurrence of *Lactobacillus acidophilus* in strains had been reported by Li *et al.* (2011), Anjum *et al.* (2014), and Mannan *et al.* (2017), who stated that the occurrence may be due to antagonism.

The results of fermented chicken feather showed an increased, weight gain, and this corroborated the findings of many researchers (Baluogu *et al.*, 2017; Onunkwo and Ekine, 2020; Yang *et al.*, 2020) who evaluated the nutritional quality of fermented chicken feather. Similar observations were found when the chicks were fed fermented fish meal. Compared to the combination of both fermented meals when fed to the chicks, the results revealed that there was a pronounced increase in the body weight gain, organ weight and immune response and these findings corroborated with the findings of many researchers (Sugiharto and Ranjekar, 2019; Zhang *et al.*, 2022 and Hong *et al.*, 2025) who evaluated the effect of fermented feed meals on the growth performance of broiler chicks and this could be attributed to lactic acid bacteria present in the feed samples.

Conclusion

This study has shown that *Lactobacillus acidophilus*-fermented chicken feather and bean chaff enhances gut health, promotes growth, and improves productivity in chicks. The fermented product improves body weights, feed intake, and hematological indices, offering a potential prebiotic solution for the poultry industry.

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