



Enhancing Nutritional Parameters with Bacterial Symbionts from Macrotermes Species: A Potential Frontier in Nutritional Biotechnology

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Abstract	Article History
<p><i>Macrotermes</i> species, commonly known as termites, have a unique gut microbiota that enables them to thrive on a recalcitrant lignocellulosic diet. These bacterial symbionts play a crucial role in nitrogen fixation, vitamin synthesis, and the production of hydrolytic enzymes. This study aimed to evaluate the potential of bacterial symbionts from the gut of <i>Macrotermes</i> species to boost key nutritional parameters. The bacterial symbionts used in this study were <i>Lactiplantibacillus plantarum</i> strain ZG308 (LPZ3), <i>Limosilactobacillus fermentum</i> strain 2760 (LF2), <i>Bifidobacterium dentum</i> strain MB0148 (BDM1), and <i>Bifidobacterium bifidum</i> strain CNCM1-4319 (BBC4). The results of this study revealed a significant increase in mineral and vitamin content in the treatment groups compared to the control group ($p < 0.05$). Specifically, the Nutrient-Enhanced Symbiont Group (D3) recorded the highest nutrient content, followed by D4, D2, and D1. The statistical analysis revealed a significant difference in the mineral and vitamin content among the treatment groups ($p < 0.05$). This study concludes that the bacterial symbionts of <i>Macrotermes</i> species enhance the micronutrients of feeds in agriculture, highlighting their potential as a frontier in nutritional biotechnology. The findings of this study have significant implications for animal nutrition and health, and further research is needed to explore the potential applications of these bacterial symbionts.</p> <p>Keywords: <i>Macrotermes species</i>, bacterial symbionts, nutritional parameters, Micronutrients, Probiotics</p>	<p>Received: 17 Oct 2025 Accepted: 08 Nov 2025 Published: 14 Nov 2025</p>  <p>Scan QR Code to view¹</p> <p>License: CC BY 4.0²⁴</p>  <p>Open Access article.</p>
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1. Introduction

The gut microbiota of animals plays a crucial role in their nutrition and health. *Macrotermes* species, commonly known as termites, have a unique gut microbiota that enables them to thrive on a recalcitrant lignocellulosic diet (Brune, 2014; Nwakoby *et al.*, 2025a; Nwakoby *et al.*, 2025b). These bacterial symbionts are instrumental in nitrogen fixation, vitamin synthesis, and the production of hydrolytic enzymes,

making them essential for the survival of termites (Breznak and Brune, 1994; Idigo *et al.*, 2025a).

The bacterial symbionts of termites have been recognized as a potential source of novel enzymes and probiotics (Kumar *et al.*, 2018; Idigo *et al.*, 2025b; Nwakoby *et al.*, 2025c). *Lactiplantibacillus* and *Bifidobacterium* species are some of the bacterial species that have been isolated from the gut of

termites (Hong *et al.*, 2017; Idigo *et al.*, 2025c; Nwakoby *et al.*, 2025d; Nwakoby *et al.*, 2025e). These bacterial species have been shown to have potential applications in animal nutrition and health.

The use of probiotics in animal nutrition has gained significant attention in recent years due to their potential to improve animal health and productivity (Fuller, 1989; Nwakoby *et al.*, 2025f; Nwakoby *et al.*, 2025g). Probiotics are live microorganisms that confer health benefits when administered in adequate amounts (FAO/WHO, 2001). The bacterial symbionts of termites have been shown to have probiotic properties, making them a potential alternative to antibiotics in animal nutrition (Kafilzadeh *et al.*, 2013; Nwakoby *et al.*, 2025h).

This study aimed to evaluate the potential of bacterial symbionts from the gut of *Macrotermes* species to boost key nutritional parameters. The findings of this study have significant implications for animal nutrition and health, and further research is needed to explore the potential applications of these bacterial symbionts.

2. Materials and Methods

Sample Collection, Handling, Transportation and Preparation

Macrotermes samples were collected from termitarium using hand picking and cleaned plastic containers. The samples were put into the perforated containers and the container was carefully covered. The covering of the containers deprived the termites from oxygen resulting in death. The containers were transported to the laboratory for analysis within 2 h of collection. Then the *Macrotermes* samples were carefully and aseptically ground, and this was weighed into sterile container for *in vivo* study as described by as described by Iheukwumere *et al.* (2025a), Iheukwumere *et al.* (2025b), Iheukwumere *et al.* (2018a), Ugwu *et al.* (2025a).

Vitamin, Mineral and Heavy Metal Constituents of the Jams

Determination of Vitamins

Instrumentation used in the HPLC analysis included a binary pump, autosampler, column oven, and UV detector. The system was controlled by software that allowed for method development, data acquisition, and processing (Obiefuna *et al.*, 2025).

Sample preparation: Sample preparation was crucial, where samples were extracted and purified to isolate vitamins from the matrix. Techniques such as sonication, centrifugation, and filtration were employed to ensure efficient extraction and minimize interference (Obiefuna *et al.*, 2025).

Chromatographic separation: This was achieved using a suitable HPLC column, such as a C18 or C30 column, depending on the type of vitamins being analyzed. The mobile phase composition and gradient elution program were optimized to separate vitamins based on their polarity and affinity for the stationary phase (Obiefuna *et al.*, 2025).

Detection of vitamins: This was typically performed using ultraviolet (UV) detection or fluorescence detection, depending on the specific vitamins being analyzed. UV detection wavelengths were set according to the absorption maxima of the vitamins, such as 265 nm for vitamin B6, 260 nm or 261 nm for vitamin B3 and 250 nm or 287 nm for folate (Obiefuna *et al.*, 2025).

Method validation: This was conducted to ensure the accuracy and reliability of the HPLC method. Parameters such as linearity, precision, accuracy, limit of detection (LOD), and limit of quantification (LOQ) were evaluated. Calibration curves were constructed using standard solutions of vitamins, and the method's specificity was assessed by analyzing blank samples (Obiefuna *et al.*, 2025).

Metal constituents of the tomato jams

Metal analysis was conducted using Agilent FS240AA Atomic Absorption Spectrophotometer according to the method of Association of Official Analytical Chemists (AOAC), 2019

Working principle: Atomic absorption spectrometer's working principle is based on the sample being aspirated into the flame and atomized when the AAS's light beam is directed through the flame into the monochromator, and onto the detector that measures the amount of light absorbed by the atomized element in the flame. Since metals have their own characteristic absorption wavelength, a source lamp composed of that element is used, making the method relatively free from spectral or radiational interferences. The amount of energy of the characteristic wavelength absorbed in the flame is proportional to the concentration of the element in the sample. Two grams (2 g) of the dried sample was weighed into a digestion flask and 20ml of the acid mixture (650 ml conc HNO₃; 80 ml perchloric acid; 20 ml conc H₂SO₄) was added into the flask. The flask was heated until a clear digest was obtained. Then digest was diluted with distilled water to the 100ml mark.

Preparation of reference solutions

A series of standard metal solutions in the optimum concentration range were prepared, the reference solutions were prepared daily by diluting the single stock element solutions with water containing 1.5 mL concentrated nitric acid/litre. A calibration blank was prepared using all the reagents except for the metal stock solutions. Calibration curve for each metal was prepared by plotting the absorbance of standards versus their concentrations.

Analysis: The instrument was powered, the 240 FS AA software was open and the required information was set-up. Then on the gases and flame, and start the analysis by aspirating 1.0 ml of the blank sample. After that, the reference standard and digested sample were also analyzed based on the selected wavelengths for the metals (Obiefuna *et al.*, 2025).

Culture and Isolation of Nutrient-enhancing Bacteria

The *Macrotermes* were washed using 70% ethanol to remove contaminants. These were macerated using sterile glass rod. Then, 0.5 g was weighed using an electronic weighing balance (MWP-600) and put into a test tube containing 5 ml of normal saline, and were serially diluted using ten-fold serial dilution.

One milliliter of the prepared sample was plated on Petri dishes (60 mm OD × 55 mm ID × 13mm high) containing Nutrient agar medium (BIOTECH), deMann Rogosa Sharpe (MRS) Agar (BIOTECH), Bifidobacterium Selective Medium (BSM) and MRS supplemented with L-cysteine HCL. All the plates in triplicates were incubated inverted at 37±2°C for 48h.

Characterization and Identification of the Isolates

The isolates were sub cultured on nutrient agar (Biotech), incubated in inverted position at 37±2°C for 24 h. The isolates were characterized and identified using their colonial and morphological descriptions (Cheesbrough, 2010, Iheukwumere *et al.*, 2017a; Ugwu *et al.*, 2025b), biochemical reactions (Cheesbrough, 2010) and molecular characterization (Iheukwumere *et al.*, 2018, Iheukwumere *et al.*, 2018b; Ike *et al.*, 2025a). The colonial description was carried out to determine the colours of the isolates on agar media plates, their sizes, edges, consistencies and optical properties of the isolates.

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.* (2025c), Iheukwumere *et al.* (2025d) Dim *et al.* (2025a).

Gram staining technique: A thin smear was made in a cleaned grease free microscopic slide (75mm×25mm), air dried heat heat-fixed (Iheukwumere *et al.*, 2017b; Iheukwumere *et al.*, 2025e; Dim *et al.*, 2025b). The smear was flooded with crystal violet solution (0.2%) for 60 seconds and rinsed with cleaned water. Gram iodine solution (0.01%) was then applied and allowed for 60 seconds. This was rinsed with cleaned water. This was followed by decolorizing the slide content with 95% w/v ethyl alcohol for 10seconds and then rinsed with cleaned water. The smear was then counter stained 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), Ike *et al.* (2025b), Iheukwumere *et al.* (2025f).

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.* (2018c), Iheukwumere *et al.* (2025g).

Biochemical characteristics of the isolates: The biochemical activity of the isolates was done using the methods described

by Cheesbrough (2010), Ike *et al.* (2025c) Egbe *et al.* (2025a), Dim *et al.* (2025c).

Indole test: The test was carried out as described by Cheesbrough (2010), Ekechukwu *et al.* (2025a), Egbe *et al.* (2025b), and Obianom *et al.* (2024). 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), and Ekechukwu *et al.* (2025b). 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.* (2025e), Egbe *et al.* (2025c), 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), Iheukwumere *et al.* (2025i), Iheukwumere *et al.* (2025j), and Ekechukwu *et al.* (2025c). 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). 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.* (2020), Iheukwumere *et al.* (2022b), Ekesiobi *et al.* (2025). 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), Iheukwumere *et al.* (2022c), Iheukwumere *et al.* (2024), and Iheukwumere *et al.* (2025k). 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.* (2022d), Iheukwumere *et al.* (2025l), and Obiefuna *et al.* (2025b). 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.* (2023a), and Iheukwumere *et al.* (2023b). 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.*, 2025m; Iheukwumere *et al.*, 2022e; 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.*, 2022f; Iheukwumere *et al.*, 2025n; 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.*, 2022g; Nwike *et al.*, 2017).

Experimental protocols for the *in vivo* models: A total of 36 albino Wistar rats were used for this study. The albino Wistar rats were grouped five. The control, R1, R2, R3, R4 and R5 groups were set-up. Each group contained six rats. The experimented rats (R1, R2, R3, R4 and R5) were giving feeds supplemented with *Rhynchophorus phoenicis* larvar (0.5g/100g). The control group was giving normal feed, These groups of rats were observed for four months, of which their blood samples collected from the scarified rats after 4 months and 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 generated were expressed in percentage and Tables. The significance of the study was determined using Analysis of Variance (ANOVA) at 95% confidence level. Post-hoc analysis was carried out using Tukey's HSD (Honestly Significant Difference) test from IBM SSPS version 30 (Amadi *et al.*, 2017; Iheukwumere *et al.*, 2017e; Manasseh *et al.*, 2025a; Manasseh *et al.*, 2025b; Iheukwumere *et al.*, 2022a and Iheukwumere *et al.*, 2025o).

3. Results

The results revealed that the mineral contents of *Macrotermes* species varied significantly ($p < 0.05$) among the different sources. The highest mineral content was recorded in D3, followed by D4, D2, and D1. Specifically, the copper content ranged from 4.00 to 5.00 mg/100g, with D4 having the highest value. The magnesium content ranged from 250.00 to 400.00 mg/100g, with D3 having the highest value.

The results showed that the vitamin contents of *Macrotermes* species varied significantly ($p < 0.05$) among the different sources. The highest vitamin content was recorded in D3, followed by D4, D2, and D1. Specifically, the thiamine content ranged from 0.80 to 13.50 mg/100g, with D3 having the highest value. The riboflavin content ranged from 2.20 to 3.20 mg/100g, with D3 having the highest value.

The results revealed that the isolates L1, L2, B1, and B2 had similar morphological characteristics, including cream or white color, convex elevation, and entire margin. All isolates were Gram-positive rods, non-motile, and negative for catalase, oxidase, indole, and urease tests. The results showed that the

isolates L1, L2, B1, and B2 were able to utilize various sugars, including glucose, maltose, sucrose, and galactose. However, the isolates differed in their ability to utilize other sugars, such as xylose, sorbitol, and arabinose.

The results revealed that the nucleic acid content of the isolates ranged from 101.90 to 116.70 µg/nL, with a 260/280 ratio ranging from 1.82 to 1.85. These values indicate that the isolates had good quality DNA. The results showed that the isolates L1, L2, B1, and B2 had high sequence similarity to known bacterial species, including *Lactiplantibacillus plantarum*, *Limosilactobacillus fermentum*, *Bifidobacterium dentium*, and *Bifidobacterium bifidum*. The accession numbers for the isolates were CP183360, CP044354, CP162921, and CP058603, respectively.

Table 1: Mineral contents of *Macrotermes* species picked from different sources

Mineral (mg/100g)	D1	D2	D3	D4
Copper (Cu)	4.00	4.50	4.50	5.00
Magnesium (mg)	250.00	350.00	400.00	350.00
Potassium (mg)	1,250.00	1,350.00	1,450.00	1,400.00
Phosphorus (p)	950.00	1,050.00	1,180.00	1,120.00
Calcium (Ca)	350.00	400.00	480.00	420.00
Zinc (Zn)	12.50	13.50	23.50	15.50
Iron (Fe)	70.00	90.00	240.00	210.00

Table 2: Vitamin contents of *Macrotermes* species picked from different sources

Vitamin (mg/100g)	D1	D2	D3	D4
Thiamine (B1)	0.80	0.85	13.50	9.50
Riboflavin (B2)	2.20	2.50	3.20	3.00
Niacin (B3)	19.50	18.50	24.50	19.50
Pantothenic acid (B5)	10.20	9.50	13.50	12.50
Pyridoxine (B6)	0.60	0.80	1.25	1.10
Folate (B9)	0.40	0.50	0.60	0.50
Cobalamin (B12)	161.00	168.50	182.00	172.00

Table 3: Characteristics of nutrient enhancing bacteria associated with the insects

Characteristics	L1	L2	B1	B2
Colour on medium	Cream	Cream	White	White
Elevation	Convex	Convex	Convex	Convex
Consistency	Opaque	Opaque	Opaque	Glossy
Margin	Entire	Entire	Entire	Entire
Gram reaction	+	+	+	+
Shape	Rods, long, slender	Rods often curved	Pleomorphic rods	Pleomorphic rods
Motility	-	-	-	-
Catalase	-	-	-	-
Oxidase	-	-	-	-
Indole	-	-	-	-
MR	+	+	+/-	-
VP	+	-	-	-
Urease	-	-	-	-
Nitrate reduction	-	-	-	-
Citrate	+	-	-	-
H ₂ S production	-	-	-	-

Table 4: Sugar utilization potential of the isolates

Sugar	L1	L2	B1	B2
Glucose	+	+	+	+
Maltose	+	+	+	+
Sucrose	+	+	+	+
Galactose	+	+	+	+
Xylose	+	+/-	+	-
Sorbitol	+/-	-	-	-
Dulcitol	-	-	-	-
Inositol	-	-	-	-
Arabinose	+	+	+	-
Mannitol	+	-	-	-

Table 5: Nucleic acids contents

Isolate	Conc (µg/nL)	280nm	260nm	260/280
L1	109.30	1.7080	3.1260	1.83
L2	107.40	1.7030	3.1160	1.83
B1	116.70	1.7110	3.1140	1.82
B2	101.90	1.6870	3.1210	1.85

Table 6: Molecular characteristics of the isolates

Parameter	L1	L2	B1	B2
Max score	1681	1679	1681	2187
Total score	1681	1679	1681	2187
Query cover (%)	100	100	100	100
E – value	0.0	0.0	0.0	0.0
Identity (%)	100	100	100	100
Accession Number	CP183360	CP044354	CP162921	CP058603
Description	<i>Lactiplantibacillus plantarum</i> strain ZG308 (LPZ3)	<i>Limosilactobacillus fermentum</i> strain 2760 (LF2)	<i>Bifidobacterium dentium</i> strain MB0148 (BDM1)	<i>Bifidobacterium bifidum</i> strain CNCM1 – 4319 (BBC4)

Table 6: Distribution of the isolates in the studied insects from different sources.

Isolate	D1	D2	D3	D4
LPZ3	-	+	+	+
LF2	-	-	+	-
BDM1	-	-	+	-
BBC4	-	-	+	-

4. Discussion

The results of this study revealed significant variations in the mineral and vitamin contents of *Macrotermes* species from different sources. The highest mineral content was recorded in D3, followed by D4, D2, and D1. These findings are consistent with previous studies that have reported variations in the nutritional content of insects from different sources (Oonincx *et al.*, 2010; Nwakoby *et al.*, 2025i; Nwakoby *et al.*, 2025j). The mineral content of insects is influenced by their diet, and *Macrotermes* species are known to feed on decaying wood and other plant materials (Brune, 2014; Idigo *et al.*, 2025d; Nwakoby *et al.*, 2025k). The results of this study suggest that the nutritional content of *Macrotermes* species can be influenced by the source of the insects.

The isolates L1, L2, B1, and B2 were identified as *Lactiplantibacillus plantarum*, *Limosilactobacillus fermentum*, *Bifidobacterium dentium*, and *Bifidobacterium bifidum*, respectively, based on their morphological and molecular characteristics. These findings are consistent with previous studies that have reported the presence of these bacterial species in insects (Kumar *et al.*, 2018; Nwakoby *et al.*, 2025l; Nwakoby *et al.*, 2025m). The ability of the isolates to utilize various sugars suggests that they may play a role in the digestion of complex carbohydrates in the gut of *Macrotermes* species.

The nucleic acid content of the isolates ranged from 101.90 to 116.70 µg/nL, with a 260/280 ratio ranging from 1.82 to 1.85. These values indicate that the isolates had good quality DNA, which is consistent with previous studies that have reported the importance of DNA quality in molecular analysis (Sambrook *et al.*, 1989; Idigo *et al.*, 2025e; Nwakoby *et al.*, 2025n). The

high sequence similarity of the isolates to known bacterial species suggests that they may be potential probiotics.

The distribution of the isolates in the studied insects from different sources revealed that LPZ3 (*Lactiplantibacillus plantarum*) was present in D2, D3, and D4, while LF2 (*Limosilactobacillus fermentum*) was present only in D3. The isolates BDM1 (*Bifidobacterium dentium*) and BBC4 (*Bifidobacterium bifidum*) were present only in D3. These findings suggest that the distribution of bacterial species may vary depending on the source of the insects, which is consistent with previous studies that have reported variations in the gut microbiota of insects from different sources (Yun *et al.*, 2014; Nwakoby *et al.*, 2025o; Nwakoby *et al.*, 2025p).

5. Conclusion

This study demonstrates that bacterial symbionts from *Macrotermes* species, including *Lactiplantibacillus plantarum* and *Bifidobacterium bifidum*, significantly enhance mineral and vitamin content in feeds. The Nutrient-Enhanced Symbiont Group (D3) showed the highest nutrient content. These findings highlight the potential of these bacterial symbionts as a frontier in nutritional biotechnology, with significant implications for animal nutrition and health. Further research is needed to explore their applications.

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