

# Inherent Microorganisms Affects the Quality of a Nigerian Fermented Beverage “Agadagidi” During Production

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## Authors' contributions

*This work was carried out in collaboration among all authors. The protocol and first draft of the manuscript was written by authors KJM and DJA designed this research. Second draft and corrections were jointly performed by authors OM and AOA Literature searches were managed by author ESO, HJA and OLO. Authors POG and IMA managed the analyses of the study. Authors OBAM and JOA performed the statistical analyses. All authors read and approved the final manuscript.*

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## ABSTRACT

This research produced wine from unripe plantain fruits using spontaneous fermentation method. The period of production was five days. The purpose of this study was to isolate and identify microorganisms associated with unripe plantain fruits, determine the physicochemical parameters of the fermenting medium. Microbial count, foaming density, alcohol content, enzymes assay, sensory analysis, physicochemical properties, mineral content, antioxidants properties, and antinutrient content of the sample were investigated every 24 h for 5 days. A total of seven bacteria and four fungi consisting of yeasts and moulds were isolated during the study. The microbial loads of bacteria and fungi of the unfermented plantain fruits were  $8.0 \times 10^6 \pm 0.01$  cfu/mL and  $14.2 \times 10^4$

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$\pm 0.01$  sfu/mL respectively. The temperature ( $^{\circ}\text{C}$ ), pH and titratable acidity (%) ranged from 25.55-32.50, 4.51-5.50 and 0.99-3.50 respectively. The sample was observed to be colourless all through the fermentation periods. Turbidity of the samples increased during fermentation from 43.50 to 111.00. The data obtained from this work has shown the importance of unripe plantain micro-flora in the production of "Agadagidi". It is also concluded that consortium of microorganisms inherent within the plantain fruits are involved in plantain fruits fermentation. This information can contribute to a better understanding of the "Agadagidi" production process for a consistent quality beverage.

**Keywords:** "Agadagidi"; local alcoholic beverages; ripe and unripe plantain; plantain wine; microbial loads; microflora.

## 1. INTRODUCTION

Plantain (*Musa paradisiaca* Linn) is one of the most important staple food crops for millions of people both in developed and developing countries. It is one of the foods commonly consumed in the West Africa sub-region, North America and the Caribbean [1]. In West and central Africa, more than 10 million tons are produced annually and are traded locally [2]. In Nigeria, its consumption cuts across the indigenous groups and the numerous socio-economic classes because of the ease of preparation and consumption [3]. "Agadagidi," a cloudy effervescent sweet-sour taste typical African traditional alcoholic beverage is made from overripe bananas and plantains through fermentation. It is common in south-western part of Nigeria [4].

Fermentation is a metabolic process that converts sugar to acid and alcohol through the actions of molds, yeasts and bacteria [5]. Fermentation is an age long method of processing cereals and legumes. Food fermentation, especially lactic acid fermentation is an important technology in Africa [6]. The technology is indigenous and is adaptable to the culture of the people [7]. The fermentation process meets the requirements of being low-cost, preventing food spoilage and food-borne diseases with respect to consumers living in a climate that aids the rapid deterioration of food [8-9]. In addition, fermented foods are of particular importance in ensuring adequate intake of protein and/or calories in the diet [10-11]. Fermentation actually holds promise as a food processing method that can be used to diversify the food uses of some under exploited plant foods like plantain [12]. The traditional processing of these foods needs to be changed or modified to improve their nutritional status [13].

The techniques used in the production of wines from tropical fruits are similar to those of grape

wine production which include pressing out the juice, fermenting, maturing and bottling [14]. Fermentation leads to changes in appearance of food which is characterized by different reactions of microorganisms [15]. Fermentation of foods covering a wide range of microbial and enzymatic processing of foods and ingredients is used to achieve desirable characteristics such as prolonged shelf-life, improved safety, attractive flavour, nutritional enrichment and promotion of health [16,17].

## 2. MATERIALS AND METHODS

### 2.1 Collection of Sample

Unripe plantain fruits (*Musa paradisiaca*, Linn) were obtained from a farmland in Arigidi-Akoko, Ondo State, Nigeria. The plantain fruits were kept in a sterile air tight polythene bags and transported to the Microbiology Postgraduate Laboratory, Federal University of Technology, Akure for further analysis.

### 2.2 Traditional Preparation of "Agadagidi" Sample

The plantain fruits were divided into two portions; A and B. Portion A was left unfermented (control sample), while Portion B was fermented for 5 days.

The unripe plantain fruits were washed under running water to remove dirt, peeled and cut into small pieces. Sliced plantain of 28.6 grams were soaked in 500 mL of sterile water in a sterilized earthenware pot for five (5) days and were mashed. At the expiration of fermentation, the plantain fruit was then filtered through two layers of muslin cloth to remove the plantain pulp. The fermented alcoholic drink was then dispensed into sterile containers. The liquid then served as "Agadagidi".

The unfermented sample was withdrawn immediately after water was added into the plantain fruits.

## 2.3 Isolation and Identification of Microorganisms

Nutrient agar (NA), nutrient broth (NB), Potato dextrose agar (PDA), Potato dextrose broth (PDB) were prepared according to manufacturer's specification for the isolation of bacteria and fungi. At every 24 h for 5 days, samples were aseptically withdrawn and serially diluted for the isolation of bacteria and fungi. Microorganisms were incubated at 37°C and 27°C for bacteria and fungi respectively. Colonies and spore forming units formed on the media were counted and identified [18].

## 2.4 Determination of Colour

Colour determination was done using the method of "The Standard Lovibond Nessleriser Disc". Each sample of 5 mL was put in one tube of Lovibond Comparator and equal volume of distilled water was also put in the second tube. The colour was then read on the disc. The colours were expressed in Hazen Units (H.U) [19].

## 2.5 Determination of Total Titratable Acidity (TTA)

The total titratable acidity of the fermenting broth was determined every day as described by [20]. Exactly 10 mL of the sample was pipetted into a conical flask with 3 drops of phenolphthalein as an indicator was titrated against 0.1 M NaOH. The end point was noticed with the pink colour appearance due to phenolphthalein indicator. The volume of the end point was recorded when the pink colour was noted [21].

## 2.6 Determination of pH

The pH value of each sample were determined using Extech Instrument pH 100 after standardization with appropriate buffers (4 and 7 solutions). The electrode sensor of the pH meter was inserted directly into 20 mL sample in a clean 50 mL glass beaker. The electrode was then left in the media for 3 min to stabilize after which the pH were read.

## 2.7 Determination of Temperature

The temperature of fermenting sample was determined. The temperature was determined every 24 h using a portable temperature meter.

## 2.8 Determination of Turbidity

Turbidity was determined using the cell riser which was installed in the cell holder of model 2100A Turbidimeter in one of the two highest ranges (100 Or 1000 NTU). A clean sample cell of turbidimeter was filled with 25 mL of each sample and covered with light shield and turbidity read in Nephelometric Turbidity Units (NTU).

## 2.9 Statistical Analysis

The result of the experiment was in triplicate. The data obtained were subjected to one way analysis of variance (ANOVA) SPSS version 2.0. Differences were considered significant at  $P < 0.05$ .

## 3. RESULTS

### 3.1 Isolation and Enumeration of Microorganisms Associated with Plantain Fruits

#### 3.1.1 Total viable microbial counts from the samples

The total viable bacterial counts ( $10^6$  cfu/mL) and fungal counts ( $10^4$  sfu/mL) are shown in Figs 1 and 2 respectively. The total viable bacterial count of unfermented sample was  $8.01 \times 10^6$  cfu/mL while that of fungi was  $1.42 \times 10^5$  sfu/mL. Day 5 had the highest viable bacterial count  $3.20 \times 10^7$  cfu/mL while Day 4 had the highest viable fungal count  $3.00 \times 10^5$  sfu/mL.

#### 3.1.2 Cultural, morphological, and biochemical characteristics of microorganisms isolated from the plantain fruits

The cultural, morphological, and biochemical characteristics of bacteria, yeasts, and moulds isolated from the plantain fruits are shown in Tables 1, 2 and 3 respectively. A total of 7 bacteria, 2 yeasts, and 2 moulds were isolated from the fermenting sample. The identified bacteria isolate from unripe plantain fruits during fermentation include; *Escherichia coli*, *Staphylococcus aureus*, *Bacillus subtilis*, *Bacillus cereus*, *Lactobacillus plantarum*, *Streptococcus faecalis* and *Leuconostoc mesenteriodes*.

The Moulds identified from unripe plantain fruits during fermentation include; *Aspergillus niger* and *Aspergillus flavus*. *Saccharomyces*

*cerevisiae* and *Candida utilis* were the yeasts isolated during fermentation.

### 3.1.3 Microbial succession during fermentation

Table 4 shows the bacterial and fungal succession during fermentation of plantain

unripe. *Bacillus* spp., *A. niger* and *A. flavus* were predominant at the beginning of the fermentation process. Day 2 to day 5 of the fermentation process, Lactic acid bacteria were the predominant bacteria involved in the fermentation while *Sacharomyces cerevisiae* and *Candida utilis* were the predominant fungi involved in the fermentation process.

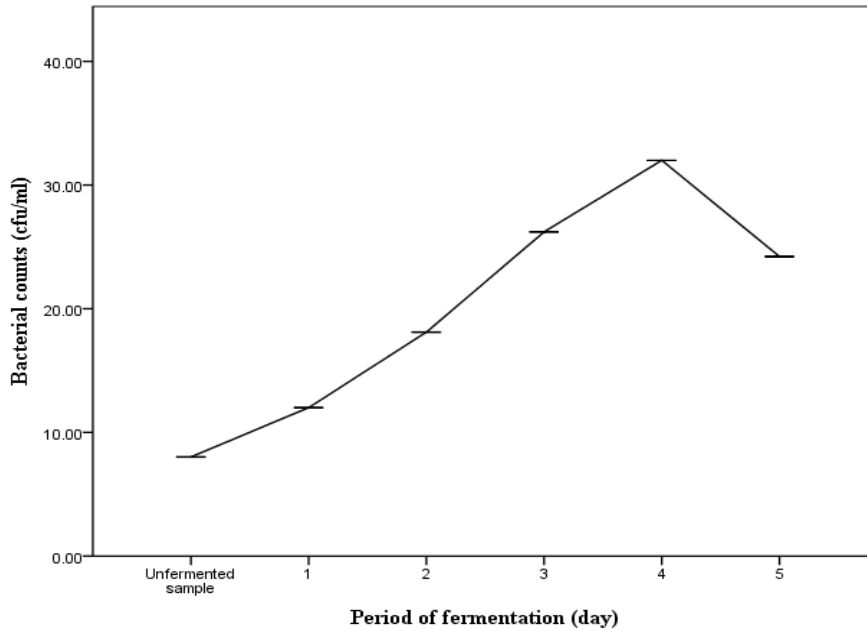


Fig. 1. Bacterial counts of unfermented and fermented unripe plantain samples

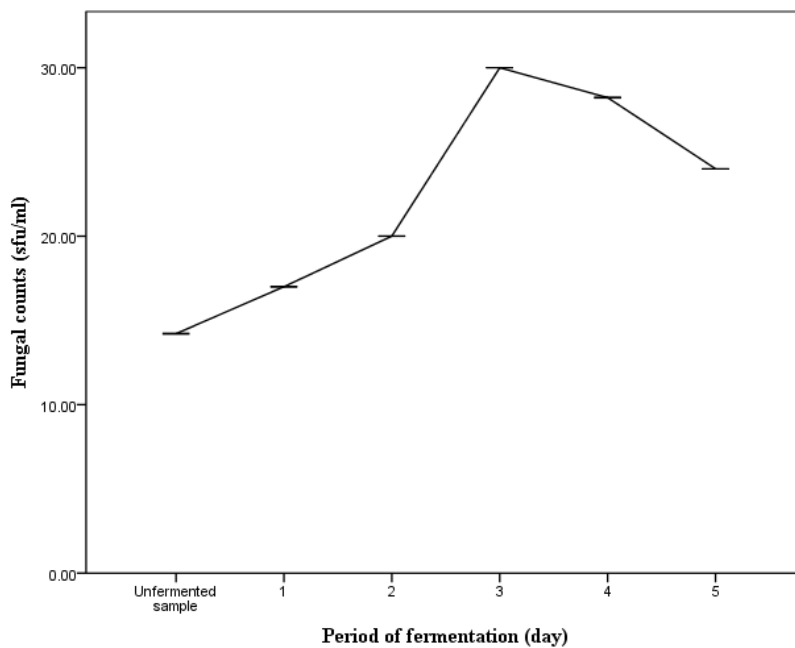


Fig. 2. Fungal counts of unfermented and fermented unripe plantain samples

**Table 1. Morphological and biochemical characteristics of bacteria isolated from Samples**

	A	B	C	D	E	F	G
Colour	Cream	Pale yellow	White	White	Yellow	Cream	White
Shape	Rod	Cocci	Rod	Rod	Cocci	Rod	Cocci
Edge	Entire	Entire	Irregular	Irregular	Entire	Irregular	Entire
Elevation	Flat	Raised	Flat	Flat	Raised	Circular	Circular
Surface	Rough	Smooth	Smooth	Rough	Smooth	Smooth	Smooth
Gram	-ve	+ve	+ve	+ve	+ve	+ve	+ve
Catalase	+ve	+ve	+ve	+ve	-ve	-ve	-ve
Coagulase	+ve	-ve	-ve	-ve	-ve	-ve	-ve
Spore	-ve	-ve	+ve	+ve	-ve	-ve	+ve
Motility	+ve	+ve	+ve	+ve	-ve	-ve	-ve
Oxidase	-ve	-ve	-ve	+ve	-ve	-ve	-ve
Indole	+ve	+ve	-ve	-ve	-ve	-ve	-ve
MethylRed	+ve	+ve	-ve	-ve	-ve	-ve	-ve
Urease	-ve	-ve	-ve	-ve	-ve	-ve	-ve
Starch	-ve	-ve	+ve	+ve	-ve	-ve	+ve
Glucose	AG	AG	A	A	-	AG	AG
Sucrose	-	A	A	A	-	-	AG
Maltose	A	A	A	A	-	A	AG
Lactose	AG	A	A	A	A	AG	AG
Fructose	-	A	AG	A	A	A	AG
Galactose	AG	AG	AG	-	-	AG	A

Key: A – *Escherichia coli* B - *Staphylococcus aureus* C – *Bacillus subtilis*

D – *Bacillus cereus*

E - *Streptococcus faecalis* F–*Lactobacillus Plantarum*

G– *Leuconostoc mesenteriodes*, A= Acid present and gas absent, AG= Acid and gas present, -ve = Negative, +ve = Positive=Absent

**Table 2. Morphological and biochemical characteristics of yeast isolated from samples**

	A	B
<b>Cultural Characteristics</b>		
Colour	Cream	Brown-black
Size	Medium	Medium
Surface	Smooth	Smooth
Shape	Oval	Cylindrical
Spore	+	+
Mycelium	-	-
<b>Sugar Fermentation</b>		
Glucose	FA	FA
Maltose	FA	FA
Lactose	A	-
Fructose	FA	FA
Sucrose	FA	FA
Galactose	FA	-

Key: A – *Saccharomyces cerevisiae* B - *Candida utilis*

A = Acid, FA = Fermentation and Assimilation, - = Absent, + = Present

### 3.1.4 Physicochemical properties of samples during fermentation

There was no visible colour observed during fermentation periods.

Fig. 3 shows steady increment in temperature (°C) in both plantain fruit samples. The highest

temperature value was on day 5 (32.50) of the fermentation period. Temperature value of unfermented sample was (25.54) which differed significantly ( $P < 0.05$ ) from the value obtained for day 1 to day 5 of fermentation. The highest pH value was obtained in unfermented sample (6.07), while the lowest pH was on day 5 (4.71).

The pH values of the unfermented sample and the fermented sample differed significantly ( $P < 0.05$ ) from each other (Fig. 4).

Fig. 5 shows that day 0 (unfermented sample) has the lowest titratable acidity (%) with a value of (0.99) while the highest was on day 5 with value of (3.50). There was significant difference ( $P < 0.05$ ) among the values of days; 1 to day 5 of the sample.

Variations in turbidity during fermentation of unripe and ripe plantain fruits are shown in Fig. 6. Sample had the turbidity ranging from  $50.01 \pm 0.01$  to  $111.00 \pm 0.00$  from day 1 to day 5. Turbidity gradually increased significantly ( $p \leq 0.05$ ) in the sample till the end of fermentation (Fig. 6).

#### 4. DISCUSSION

The microbiological changes observed during fermentation could be as a result of diverse group of microorganisms present during fermentation process [22]. The bacterial count in the plantain fruit was higher compared to fungal count. This is in agreement with the report of Akinyosoye and Arotupin [23] stating that bacterial are most numerous organism in the soil. The high counts of bacteria, mould and yeast in the plantain fruits during fermentation may be due to the high concentration of nutrients which agrees with the findings of Arotupin et al. [24] that arrays of microorganisms could increase in population due to the nutrient rich nature of the substrate, thus, supporting the growth and proliferation of microorganisms. Low pH could be responsible for the increased fungal population [25]. The lignocellulosic structure of plantain fruits may be a factor responsible for the low microbial population in the unfermented sample [26].

The presence of bacteria such as *B. cereus*, and *B. subtilis* in the unfermented sample agrees with the findings of Oriola et al. [27]. The presence moulds such as *A. niger*, and *A. flavus* in the unfermented sample concurs with the finding of Oriola et al. [27] when *A. flavus*, *A. fumigatus* and *P. notatum* were isolated from uncrushed plantain fruits. The dominant population of *A. flavus* particularly in the unfermented sample may be due to the presence of nutrients available within the plantain fruits for utilization. Nasrin et al. [28] attested to this when combination of molasses and jackfruit were used as a substrate for mutant strain of *A. niger* for

citric acid production. *Saccharomyces cerevisiae* and *Aspergillus* spp. may as well be responsible for lactic acid production which concurs with the documentation of Ogbonnaya and Chukwu, [29] that some yeasts (*Saccharomyces*) and moulds (*Penicillium*, *Aspergillus* and *Botrytis*) produce lactic acid.

The decrease observed in bacterial load during plantain fruits fermentation on day 5 may be due to inhibitory effect of bioactive substances [30,31]. The higher prevalence of rod-shaped lactic acid bacteria in this study corroborated the study of Nwokoro and Chukwu [32] who reported that lactic acid bacteria commonly predominates during food fermentation. This is because they are aciduric in nature [32,33].

The appearance and increase in the yeast count after 24 h of fermentation is attributed to the decrease in the pH that creates conditions ideal for yeast growth. This is similar with the finding reported for other fermented beverages [34]. Roles of yeasts have also been reported to include improving the organoleptic qualities by producing different flavours and aroma in different foods [35]. The moulds (*Aspergillus flavus* and *Aspergillus niger*) isolated in the study are commonly present as contaminants in fruits and legumes and do not appear to play any significant important role in the fermentation [36,37].

The subsequent disappearance of moulds after 24 h observed in this study as well as previous studies [37,32,33] was probably due to the low oxygen tension in the fermenting matrix. It could also be due to the presence of organic acids especially lactic acid as it could be seen that the growth of lactic bacteria and yeasts increased gradually throughout fermentation while the numbers of moulds decreased. The growth of lactic acid bacteria and yeasts can inhibit the growth of moulds [38].

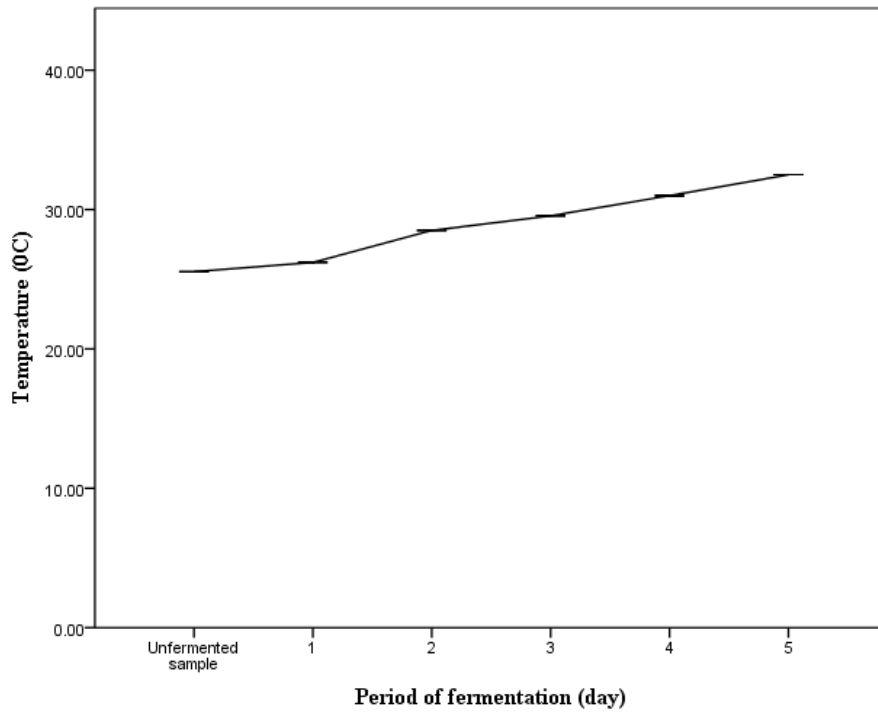
The *Bacillus* species isolated in this study which was *Bacillus cereus* and *Bacillus subtilis*. Previous reports [36,32] have also isolated *Bacillus* species in the fermentation and they have been reported to show saccharolytic activities [32]. These microorganisms persisted towards the end of the fermentation indicating that they continued the saccharification of starch to release sugars. The presence of lactic acid bacteria as dominant organisms may be due to their aciduric nature [39].

**Table 3. Morphology characteristics of mould isolated from samples**

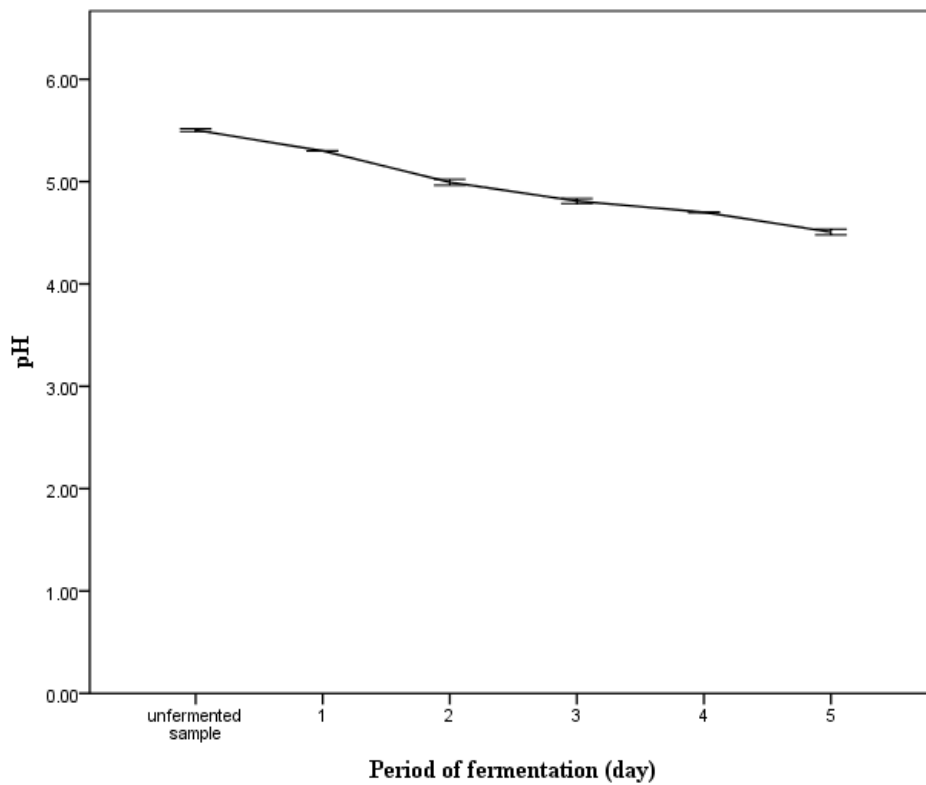
Isolates	Cultural characteristics	Spores/conidia arrangement under the microscope	Identity of isolates
A1	The surface is rough. Spores are granular flat, often with radial grooves yellow at first but quickly become bright yellow green with age. Sizes of colonies are medium.	Conidia are globose to sub globose with pale green colour. Septate hyphae with long conidiophore bearing the conidia	<i>Aspergillus flavus</i>
A2	The surface is smooth with brown mycelia growth. With whitish colonies large in size.	An upright conidiopore that terminates in a swelling, bearing phialides at the apex radiating from the entire surface. Conidial are one celled and densely packed. Spores are black.	<i>Aspergillus niger</i>

**Table 4. Microbial succession during fermentation of plantain fruits**

Unfermented sample	Day 1	Day 2	Day 3	Day 4	Day 5
A <i>Bacillus cereus</i> , <i>Bacillus subtilis</i> , <i>Escherichia coli</i>	<i>Staphylococcus aureus</i> , <i>Streptococcus faecalis</i>	<i>Staphylococcus aureus</i> , <i>Lactobacillus plantarum</i> , <i>Leuconostoc mesenteriodes</i>	<i>Lactobacillus plantarum</i> , <i>Bacillus subtilis</i> . <i>Streptococcus faecalis</i>	<i>Lactobacillus plantarum</i> , <i>Streptococcus faecalis</i>	<i>Lactobacillus plantarum</i>
B <i>Aspergillus niger</i> , <i>Aspergillus flavus</i>	<i>Aspergillus flavus</i> , <i>Saccharomyces cerevisiae</i>	<i>Saccharomyces cerevisiae</i> , <i>Candida utilis</i>	<i>Candida utilis</i> , <i>Aspergillus niger</i> <i>Saccharomyces cerevisiae</i>	<i>Saccharomyces cerevisiae</i> , <i>Candida utilis</i>	<i>Saccharomyces cerevisiae</i> , <i>Candida utilis</i>

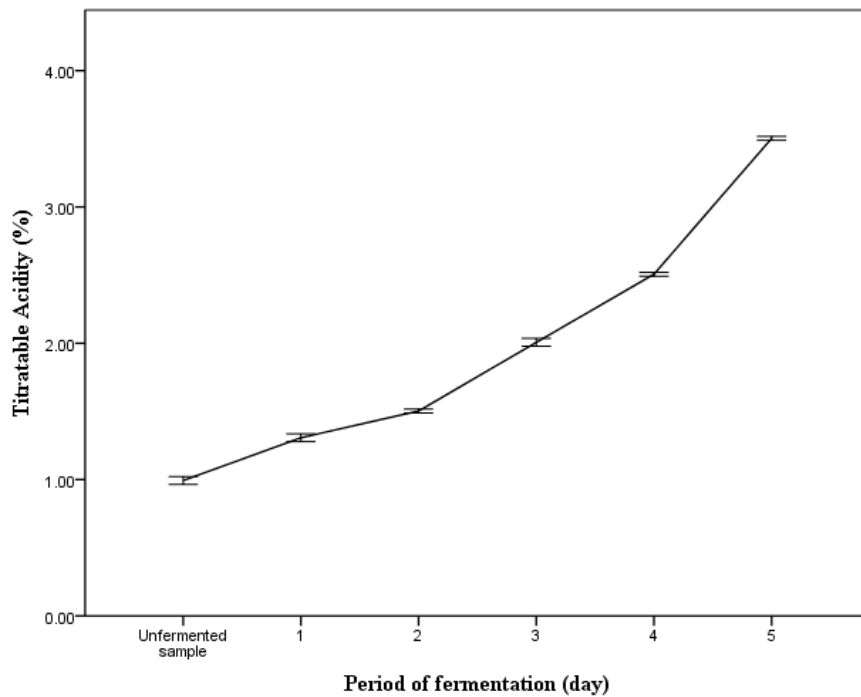


**Fig. 3. Temperature of unfermented and fermented unripe plantain samples**

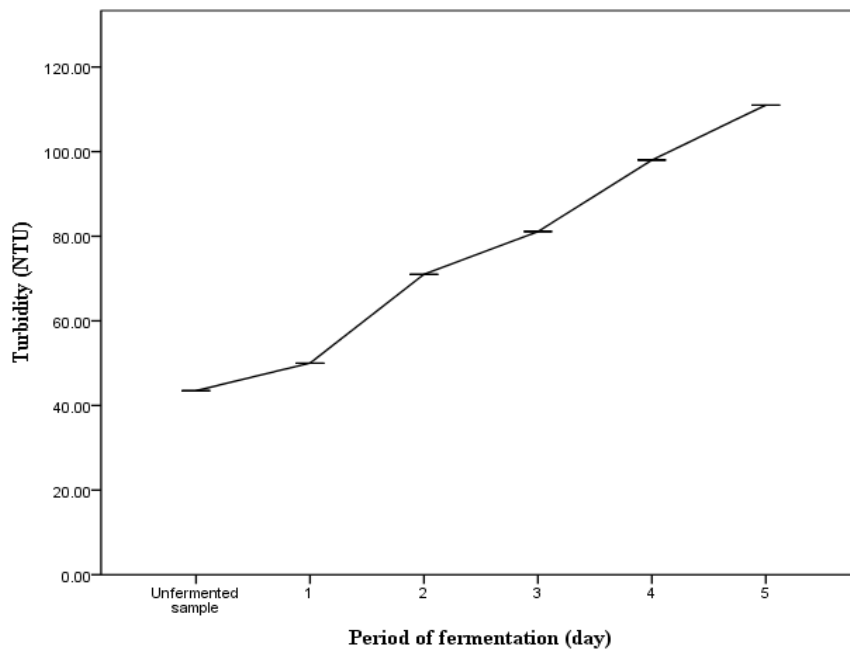


**Fig. 4. pH of unfermented and fermented unripe plantain samples**





**Fig. 5. Titratable acidity of unfermented and fermented unripe plantain samples**



**Fig. 6. Turbidity of unfermented and fermented unripe plantain samples**

This study revealed that unfermented and fermented unripe plantain fruits colour was observed to be colourless. This could be as a result of the environment and natural colour of the fruits. There was a steady decrease in the pH during fermentation. This might be as a result of

production of lactic acid by fermentative organisms (mainly lactic acid bacteria) responsible for the fermentation of “Agadagidi” [40]. A significant increase in the total titratable acidity (TTA) during fermentation was recorded in this study, and this could be as a result of

production of lactic acid and other organic acids by organisms responsible for the fermentation, as is the case with the decrease in pH. This observation of decrease in pH and increase in total titratable acidity (TTA) agrees with most fermentation studies including those of Wakil and Kazeem [41], Okwute and Olafiaji [42], and Modu et al. [43]. The early rise in titratable acidity and reduced pH is important to avoid proliferation of undesirable organisms resulting in poor fermentation [37]. During the process of fermentation, the temperature of samples was observed to increase as fermentation time progresses. The increases in temperature may be due to the activity of different microorganisms during fermentation process. This is similar to a result reported by [22], when he fermented ripe and unripe plantain flour. This supports the fact that fermentation is an exothermic process and that the heat generated was due to metabolic activities of microorganisms.

## 5. CONCLUSION

This study therefore showed that microorganisms such as *Escherichia coli*, *Staphylococcus aureus*, *Bacillus subtilis*, *Bacillus cereus*, *Lactobacillus plantarum*, *Streptococcus faecalis* and *Leuconostoc mesenteroides*, *Aspergillus niger* and *Aspergillus flavus*, *Saccharomyces cerevisiae* and *Candida utilis* were associated with plantain fruits. Thus, *Leuconostoc mesenteroides*, *Lactobacillus plantarum*, *Bacillus subtilis*, *Saccharomyces cerevisiae* and *Candida utilis* were found predominant in the production of plantain wine.

## COMPETING INTERESTS

Authors have declared that no competing interests exist.

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