



Microbiology and Chemical Preservation of Eggplants Sold in Five Popular Markets in Owerri, Imo State, Nigeria



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Abstract

Preservatives have been frequently used to improve the shelf life of food. The preservation of eggplants was determined using three chemical preservatives; ascorbic acid, sodium benzoate and sodium metabisulphite applied at different concentrations. The microbial load was determined to ascertain the efficacy of the preservatives. The microbiological population and diversity of the eggplant was assessed before and after treatment using standard methods. The plate count techniques were adopted to determine the total colony forming unit of the sample. Antimicrobial activities revealed that sodium benzoate inhibits the growth of bacterial and fungal species even at low concentration. Sodium metabisulphite was effective on bacteria while ascorbic acid showed a stimulatory response on the growth of fungi and bacteria.

Four bacteria species namely, *Staphylococcus aureus*, *Bacillus subtilis*, *Micrococcus luteus* and *Enterococcus faecalis* and four fungal species, namely, *Saccharomyces*, *Fusarium*, *Aspergillus* and *Rhizopus* were isolated. Most of the isolates are soil and water borne which could serve as vehicle for contamination of the eggplants. Some of the isolates like *Staphylococcus* and *Bacillus* are involved in food spoilage and borne infections while *Aspergillus* and *Fusarium* species are notable causes of mycotoxicosis in man and animals. This study shows that preservatives applied at appropriate concentrations can create adequate protection from spoilage microorganisms, and therefore recommends the use of sodium benzoate to enhance the keeping quality of eggplants

Keywords: Eggplants; Microbiology; Preservatives

Abbreviations: GDP: Gross Domestic Product; NA: Nutrient Agar; PDA: Potato Dextrose Agar; MCA: MacConkey Agar

Introduction

The huge loss of fruits and vegetables to spoilage is becoming alarming. The enormous economic loss to farmers and reduction in gross domestic product (GDP) to producing nations has been the discourse in many international conferences. The high moisture contents and nutritional components of fruits and vegetables predispose it to both chemical and microbial spoilage [1-8]. Fruits, apart from its nutritional values have so many therapeutic potentials. Some contain phytochemical and antioxidant agents [9-11,5], while some have been employed in production of beverages, probiotics and prebiotics. Garden egg contains flavonoids, tropine, glycoalkaloids, arginine, lanosterol, gramisterol, aspartic acid as important constituents [9]. The plant is reported to have analgesia, antipyretic, antioxidant, anti-inflammatory, antiasthmatic,

hypolipidemic, hypotensive, antiplatelet, intraocular pressure reducing and anaphylactic reaction inhibitory activities [12].

The demand for garden egg is on the increase as it has gone beyond being a staple fruit to a more commercial commodity that is now a major source of income for producing households and marketers in the forest zone [13]. Since fruits are generally acidic, they are naturally amenable to preservation. Secondly, increased acidity can activate chemical reactions such as pectin which lowers water activity and reduces the possibility of microbial growth. Dehydration is among the oldest and most common forms of fruit preservation as moisture in the fruit is driven off; leaving a stable food that has moisture content below that at which microorganisms

can grow. This research reports the microbiological quality of garden eggs and different chemical preservation methods.

Materials and Methods

Sample Collection and Preparation

Four hundred fresh garden eggs were collected randomly from five markets (Ekeonunwa, Ihiagwa, Relief, Ezeobodo and Nekede) in Imo state, Nigeria. Two hundred samples were subjected to microbiological analysis without treatment with preservatives, while the remaining two hundred were treated with different concentrations of preservatives before subjecting to routine microbiological analysis.

Microbiological Analyses of Samples

Ten grams of fresh eggplants were rinsed in ninety milliliters of sterile distilled water to obtain 10¹ dilution. Further dilutions were made decimally until 10⁵ dilution was obtained. One-tenth milliliter (0.1ml) of the fifth dilution was plated in duplicates onto freshly prepared surface dried Nutrient agar (NA), Potato Dextrose Agar (PDA) and MacConkey agar (MCA). The inocula were spread evenly before incubated at ambient temperature (28±02°C) [14-16].

Preparation of Preservatives and Treatment of Samples

Two hundred milliliters of distilled water were sterilized in a 500mls conical flask. This was repeated in four sets. Different concentrations (10g, 5g, 3g, and 1g) of ascorbic acid was dissolved in the sterile distilled water and shaken vigorously. The procedure was repeated with sodium metabisulphite and sodium benzoate.

Twenty grams (20g) of fresh eggplants were suspended on the prepared preservatives and left to stand on the bench for 24h. The microbiological procedure was repeated for the samples treated with the preservatives.

Enumeration and Characterization of Microbial Isolates

The number of viable colonies developed were enumerated and express as colony forming unit per grams [15,16]. The characterization and identification of bacteria and fungi isolates were based on the colonial morphology, microscopic and standard biochemical test [17-21].

Results

Table 1 shows the total microbial populations prior to the addition of preservatives. Counts ranged between 3.3 x 10⁵ - 1.43 x 10⁷ for heterotrophic bacteria, 1.0 x 10⁵ - 1.45 x 10⁷ for heterotrophic fungi and 1.0 x 10⁴ - 3.2 x 10⁴ for coliform bacteria. Table 2 shows the biochemical and carbohydrate characterization of bacteria isolated from the untreated samples. Four species of bacteria namely, *Staphylococcus aureus*, *Bacillus subtilis*, and *Micrococcus luteus* and *Enterococcus faecalis* were isolated. Four fungal species namely *Rhizopus nigrican*, *Saccharomyces cerevisiae*, *Aspergillus* and *Fusarium* species were isolated from the sample as shown in Table 3. Table 4 shows the total count and colony characterization of bacterial isolates after treatment with different concentrations of preservatives. Samples treated with ascorbic acid did not show any effect, especially at concentrations 1g and 3g, as the organisms grew luxuriantly. Remarkable result was evident at concentrations 5g and 10g as there was reduction on both microbial population and diversity.

Table 1: Total microbial population on garden eggs before treatment with preservatives.

Sample Code	Total Heterotrophic Bacterial Counts on Nutrient Agar	Total Heterotrophic Fungal Counts on Potato Dextrose Agar	Total Coliform Counts on Macconkey Agar
EKM1	4.5 x 10 ⁵	1.2 x 10 ⁴	1.0 x 10 ⁴
EKM2	5.6 x 10 ⁵	2.9 x 10 ⁴	1.2 x 10 ⁴
EKM3	6.2 x 10 ⁵	2.6 x 10 ⁴	1.5 x 10 ⁴
EKM4	3.3 x 10 ⁵	1.1 x 10 ⁴	1.2 x 10 ⁴
EKM5	3.8 x 10 ⁵	7.1 x 10 ⁴	1.0 x 10 ⁴
IHM1	5.5 x 10 ⁵	2.9 x 10 ⁴	1.9 x 10 ⁴
IHM2	6.0 x 10 ⁵	3.3 x 10 ⁵	1.5 x 10 ⁴
IHM3	4.1 x 10 ⁵	3.9 x 10 ⁵	1.8 x 10 ⁴
IHM4	4.4 x 10 ⁵	1.0 x 10 ⁵	1.0 x 10 ⁴
IHM5	5.0 x 10 ⁵	2.1 x 10 ⁵	1.0 x 10 ⁴
RFM1	1.11 x 10 ⁷	1.22 x 10 ⁵	1.1 x 10 ⁴
RFM2	1.43 x 10 ⁷	1.11 x 10 ⁷	2.2 x 10 ⁴
RFM3	1.23 x 10 ⁷	1.28 x 10 ⁷	1.4 x 10 ⁴
RFM4	8.9 x 10 ⁵	1.05 x 10 ⁷	1.0 x 10 ⁴ 10 ⁴
RFM5	6.5 x 10 ⁵	1.45 x 10 ⁷	1.0 x 10 ⁴
EZM1	6.9 x 10 ⁵	3.4 x 10 ⁵	1.2 x 10 ⁴
EZM2	5.5 x 10 ⁵	3.0 x 10 ⁵	1.9 x 10 ⁴
EZM3	3.6 x 10 ⁵	6.8 x 10 ⁵	1.6 x 10 ⁴
EZM4	4.0 x 10 ⁵	4.5 x 10 ⁵	1.0 x 10 ⁴

EZM5	6.1×10^5	5.5×10^5	2.8×10^4
NKM1	3.9×10^5	2.1×10^5	2.2×10^4
NKM2	3.3×10^5	3.5×10^5	2.0×10^4
NKM3	5.2×10^5	1.2×10^5	1.2×10^4
NKM4	3.9×10^5	1.0×10^5	2.2×10^4
NKM5	5.5×10^5	1.6×10^5	3.2×10^4

Note: EKM, Ekeonuwa market; IHM, Ihiagwa market; RFM, Relief market; EZM, Ezeobodo market; NKM, Nekede market.

Table 2: Biochemical characterization and carbohydrate formation test of bacteria isolated from samples.

Gram Morphology	Cat	Oxi	In	MR	Cit	NO ₃	Urease	Carbohydrate Fermentation						Identity of Isolate
								Glu	Suc	Mal	Lac	Mann	Xyl	
+S	+	-	+	-	-	+	+	+	+	+	+	+	-	<i>Staphylococcus aureus</i>
+R	+	-	-	-	+	+	-	+	-	-	-	+	+	<i>Bacillus subtilis</i>
+S	+	-	-	+	+	+	-	-	-	-	-	-	-	<i>Micrococcus Luteus</i>
+S	+	-	-	+	+	-	-	+	+	-	+	+	-	<i>Enterococcus faecalis</i>

Note: Cat, catalase; Oxi, oxidase; In, indole; MR, methyl red; Cit, citrate; NO₃, Nitrate reduction; Glu, glucose; Suc, sucrose; Mal, maltose; Mann, Mannose; Xyl, xylose; Lac, lactase: +S, gram positive spherical (round) shape: +R, gram positive rod shape.

Table 3: Colonies and microscopic characterization of fungal isolates.

Colony Characterization	Microscopic Characterization	Identification of Isolate
Tall white filamentous hyphae with orange spores.	Non septate hyphae	<i>Rhizopus nigricans</i>
Moist and shiny cream colonies	Gram positive oval budding cells	<i>Saccharomyces spp.</i>
Dull and dry domed shape cream colonies	Gram positive ellipsoidal budding cells	<i>Saccharomyces ellipsoideus</i>
Yellow colonies	Gram positive oval budding cells	<i>Saccharomyces spp.</i>
Moist and shiny	Gram positive budding cells	<i>Saccharomyces spp.</i>
Cream colonies, short mycelium with black spores and yellow background	Septate hyphae hyphae. Conidial attached to vascular	<i>Aspergillus spp</i>
Black spores raised hyphae	Septate hyphae	<i>Aspergillus spp</i>
Short white mycelia with red background	Conidia sickle sloped	<i>Fusarium spp</i>

Table 4: Total count and colony characterization of Bacteria isolated on samples treated with preservatives.

Preservatives	Concentration of Preservatives (g)	Total Bacterial Counts (Cfu/ml)	Identities of Bacterial Isolates
Ascorbic Acid	1	8.1×10^3	<i>Bacillus cereus, Staphylococcus aureus, Micrococcus luteus, Enterococcus faecalis</i>
	3	7.2×10^3	<i>Bacillus cereus, Staphylococcus aureus, Enterococcus faecalis</i>
	5	2.1×10^2	<i>Bacillus cereus, Staphylococcus aureus</i>
	10	1.3×10^2	<i>Bacillus cereus, Staphylococcus aureus</i>
Sodium Benzoate	1	2.1×10^2	<i>Bacillus cereus, Staphylococcus aureus</i>
	3	1.0×10^2	
	5	0	<i>Bacillus cereus</i>
	10	0	

Sodium metabisulphite	1	1.04 x 10 ⁴	<i>Bacillus cereus</i> , <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Enterococcus faecalis</i>
	3	1.28 x 10 ⁴	<i>Bacillus cereus</i> , <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Enterococcus faecalis</i>
	5	1.49 x 10 ⁴	<i>Bacillus cereus</i> , <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Enterococcus faecalis</i>
	10	2.11 x 10 ⁴	<i>Bacillus cereus</i> , <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Enterococcus faecalis</i>

There was significant decrease in both cells counts and bacterial types in samples treated with sodium benzoate. The higher concentrations (5g and 10g) inhibited the growth of the organisms. Sodium metabisulphite was stimulatory and the bacterial populations increased with concentration of the preservatives. The effects of preservatives on the fungal isolates on the samples is shown in Table 5. The higher concentrations (5g and 10g) of sodium benzoate and sodium metabisulphite showed significant inhibition on the growth of the fungal isolates. Ascorbic

acid showed a stimulatory effect on the growth of bacteria and fungi. The percentage occurrence of bacteria and fungi isolated from the samples is shown in Table 6. The yeasts, *Saccharomyces species* and the bacterium, *Enterococcus faecalis* were dominant species isolated from the samples. The distribution of the bacteria and fungi isolated across the sample locations (markets) is shown in Table 7. *Saccharomyces*, *Bacillus* and *Enterococcus faecalis* were isolated in the sample across the markets.

Table 5: Total count and colony characterization of Fungi isolated on samples treated with preservatives.

Preservatives	Concentration of Preservatives (g)	Total Fungal Counts (Cfu/ml)	Identities of Fungal Isolates
Ascorbic Acid	1	1.96 x 10 ⁴	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i> , <i>Aspergillus sp</i> , <i>Fusarium sp</i>
	3	2.11 x 10 ⁴	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i> , <i>Aspergillus sp</i> , <i>Fusarium sp</i>
	5	2.49 x 10 ⁴	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i> , <i>Aspergillus sp</i> , <i>Fusarium sp</i>
	10	2.71 x 10 ⁴	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i> , <i>Aspergillus sp</i> , <i>Fusarium sp</i>
Sodium Benzoate	1	3.1 x 10 ³	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i>
	3	1.0 x 10 ²	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i>
	5	0	
	10	0	
Sodium metabisulphite	1	1.9 x 10 ³	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i>
	3	1.0 x 10 ²	<i>Saccharomyces cerevisiae</i> , <i>Saccharomyces ellipsoideus</i>
	5	0	
	10	0	<i>Bacillus cereus</i> , <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Enterococcus faecalis</i>

Table 6: Percentage distribution of Bacteria and Fungi in samples.

Bacterial Isolates	Percentage Occurrence (%)	Fungal Isolates	Percentage Occurrence (%)
<i>Staphylococcus aureus</i>	18 (16.7)	<i>Saccharomyces cerevisiae</i>	45 (38.8)
<i>Bacillus cereus</i>	21 (19.4)	<i>Saccharomyces ellipsoideus</i>	39 (33.6)
<i>Micrococcus luteus</i>	14 (13.0)	<i>Aspergillus sp</i>	12 (10.3)
<i>Enterococcus faecalis</i>	45 (41.7)	<i>Rhizopus nigricans</i>	4 (3.4)
<i>Bacillus subtilis</i>	10 (9.3)	<i>Fusarium sp</i>	16 (13.8)

Table 7:

Sample Locations	Bacteria Distribution	Fungi Distribution
EKM	<i>Bacillus cereus, Bacillus subtilis, Enterococcus faecalis, Staphylococcus aureus, Micrococcus luteus</i>	<i>Rhizopus nigricans, Saccharomyces cerevisiae, Saccharomyces ellipsoideus, Aspergillus sp, Fusarium sp</i>
EZM	<i>Bacillus cereus, Bacillus subtilis, Enterococcus faecalis, Staphylococcus aureus, Micrococcus luteus</i>	<i>Rhizopus nigricans, Saccharomyces cerevisiae, Saccharomyces ellipsoideus, Fusarium sp</i>
IHM	<i>Bacillus cereus, Bacillus subtilis, Enterococcus faecalis, Staphylococcus aureus, Micrococcus luteus</i>	<i>Rhizopus nigricans, Saccharomyces cerevisiae, Saccharomyces ellipsoideus, Aspergillus sp, Fusarium sp</i>
RFM	<i>Bacillus cereus, Enterococcus faecalis, Staphylococcus aureus, Micrococcus luteus</i>	<i>Saccharomyces cerevisiae, Saccharomyces ellipsoideus, Aspergillus sp, Fusarium sp</i>
NKM	<i>Bacillus cereus, Bacillus subtilis, Enterococcus faecalis</i>	<i>Rhizopus nigricans, Saccharomyces cerevisiae, Saccharomyces ellipsoideus,</i>

Discussion

The results obtained shows that the eggplants were contaminated with diverse microorganisms. Some of these isolates have been found to be associated with certain food borne illnesses, most of which arise from improper handling, preparation and poor food storage facilities [22,23], also can cause food spoilage [23,24]. The presence of some pathogenic bacteria such as *Bacillus subtilis*, *Enterococcus faecalis*, *Micrococcus luteus* and *Staphylococcus aureus* is worrisome because of their public health importance. *Bacillus subtilis* is a spore former that can withstand adverse effects of preservative(s). These structures pose resistant to environmental stress such as heat, ultraviolet radiation, chemical disinfectant and desiccation [23]. The presence of *Enterococcus faecalis* associated with faecal contamination is notable causative agent of food borne gastroenteritis and bacterial diarrhea disease [23,25-27].

Reported that some strains of *Aspergillus flavus* produce aflatoxins, which are cancerous in mammals including human. The presence of these microorganisms is an indication of the use of contaminated water by food handler and soil where they are cultivated [23,25,26]. *Saccharomyces cerevisiae* and *Bacillus species* are the major fermentative organisms and may influence microbial food deterioration and spoilage [26,28]. Results shows that the chemical preservatives used were effective against microbial growth

at different concentrations. This was evident in the reduction of bacterial and fungal count over time. Chemical preservatives have been used to improve the shelf lives of food substances as they inhibit, retard or arrest the growth of microorganisms [7,29-31]. Investigation of antimicrobial activities of the three preservatives used showed that sodium benzoate was more effective on both bacterial and fungal species.

This was also reported by [32]. This was evident as samples treated with sodium benzoate had a lower microbial count as well as decrease in microbial diversities. The low hydrogen ion concentration, (pH) could account for the reduction in both microbial populations and diversities [32-34]. The higher bacteria contamination shown in samples treated with sodium metabisulphite indicates a stimulatory response rather than antagonistic response. Sodium metabisulphite was very effective against fungi at higher concentrations, whereas ascorbic acid was stimulatory to the growth of fungi, especially the yeasts, *Saccharomyces*. This study has established microbiological safety of eggplants treated with chemical preservatives. The use of chemical preservatives in appropriate concentration in addition to other hurdles techniques recommended by [30] could serve as a check in the control of microbial contamination.

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