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# Assessment of microbiological quality and nutritional values of a processed edible weevil caterpillar (*Rhynchophorus phoenicis*) in Port Harcourt, southern Nigeria

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# ABSTRACT

Investigation into the microbiological quality of processed *Rhynchophorus* species revealed the presence of bacteria and fungi. Total bacterial and fungal counts were  $1.08 \times 10^5$  CFU/g and  $9.2 \times 10^2$  CFU/g respectively. Bacterial isolates identified included, *Lactobacillus plantarum*, *Staphylococcus aureus*, *Bacillus subtilis*, *Pseudomonas aeruginosa* and *Proteus vulgaris*, while the fungal species included *Cladosporum* sp, *Penicillium verrecosum*, *Aspergilus flavus and Fusarum poae* and one species of yeast. *Rhynchophorus* sp is rich in protein (71.63%), indicating high nutritional quality. The crude fibre (1.59%) and ash (11.97%) are high and could aid bowel movement and increase mineral contents respectively in the human body. High moisture (12.14%) and protein contents could encourage spoilage by proteophilic and hydrophilic bacteria and fungi. Poor sanitation and inadequate storage and marketing conditions may contribute to contamination and recontamination of the products. Intensive education on pre and post-harvest (handling) activities, including proper disposal of intestinal contents, preservation with spices and the application of atmospheric packaging system could reduce contamination with microbes.

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Keywords: Bayelsa suya (Rhynchophorus sp) larvae, nutritional and Microbiological qualities.

### **INTRODUCTION**

Global demand for food security and safety coupled with increased cases of malnutrition resulting from inadequate and high cost of protein diets, especially in Africa and developing countries necessitated the consumption of insects as a new food alternative that enrich the basic diet of man. The rise in the consumption of insects as food has become inevitable in part of sub-Saharan Africa in particular and Africa in general. Though the use of insects as food had been indigenous to Africa, other continents like Asia, Australia and Latin America have utilized them either as food directly or as food source for many years (Allotey and Mpuchane, 2003).

A large variety of insects including, winged termites, beetles, grubs, ants, locusts, crickets and grasshoppers are edible. These insects are eaten roasted, dried, fried or added to other foods. The larvae of an emperor moth, *Imbresia belina* are presently the most commonly utilized insects in southern Africa

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(Allotey and Mpuchane, 2003; Gashe et al., 1997). The caterpillar of a lepidopteran, *Bunaea alcinoe* is edible and is rich in amino acids and proteins (Amadi et al., 2005; Braide et al., 2009; Braide et al., 2010). The Pedi people of southern Africa prefer insects to meat (Quin, 1959), so do the Yukpa people of Venezuela and Colombia prefer traditional insect food to fresh meat (Ruddle, 1973). In Zambia, nutrient analyses of foods consumed by the Bemba revealed 65 g protein/100 g of insects for dried caterpillars compared to 32 g for dried fish, 30 g for roasted venison and lesser amount for other foods (DeFoliart, 2002).

The rise in the world population has placed greater demand on food security. Insects are abundant and contain useful nutrients like protein, vitamins and minerals (Siame et al., 1996; Motshegwe et al., 1998; Oliviera et al., 1976; Braide et al., 2009; Braide et al., 2010). Africa, a developing continent is facing numerous food problems, and a search for new food alternatives that enrich the basic diet and also supply the traditional African food habits and culture has become inevitable. General studies on the traditional food life style of southern Africa suggest that insects may have supplied a substantial amount of animal protein in their diet (Gelfand, 1971; Quin, 1959). People of the tropical areas of Africa, Asia and Neotropics have utilized the mature larvae of the palm weevil, Rhynchophorus species as food sources for centuries (Allotey & Mpuchane, 2003).

Presently in Nigeria, numerous edible insects and caterpillar are known, consumed and even preserved, but basic information and data on their proximate composition and microbiological qualities are not sufficient. Oliviera et al. (1976) reported that *Rhynchophorus sp* found in parts of Nigeria and Zaire is rich in Zinc, protein, thiamine and riboflavin. Wachukwu et al. (2002) had reported on the microorganisms associated with *Rhynchophorus phoenicis*.

The caterpillars are called "itun" in Yoruba and "Eruru Ngwo" in Igbo all in Nigeria. Rhynchophorus usually breeds inside palm tree under stress and fallen palms (Fasoranti & Ajiboye, 1993). The method of harvesting, preparation, storage and marketing unhygienic and susceptible is to contamination. During processing the larvae may come in contact with soil and other rotting materials and may become recontaminated by microorganisms that can cause spoilage during drying and storage. It is therefore imperative to determine the sanitary quality of this edible insect.

The study reports on the microbiological quality and the nutritional values of an edible caterpillar of a *Rhynchophorus* species.

# MATERIALS AND METHODS

### **Description and sources of sample**

Samples of processed Rhynchophorus larvae were bought from hawkers at Rumuokoro Motor Park in Port Harcourt, Rivers State, Nigeria. The sample was identified by Professor B.A.Okwapkpam, an entomologist in the Department of Applied and Environmental Biology, Rivers State University of Science and Technology, Nkpolu, Harcourt. Processed Port Rhynchophorus larvae were sold in sticks in an open market and along major motor parks. Samples were taken immediately to the National Root Crops Research Institute (NRCRI) Laboratory, Umudike, Abia State, Nigeria and stored at 4 °C prior to analysis.

### Preparation of sample for analysis

Prior to analysis, samples were examined and extraneous materials removed. Samples were divided into two parts, "A" and "B". Part "A" was u sed to determine the proximate composition, while part "B" was used to study the microbiological quality.

Sample "A" was spread out on a laboratory tray and dried in an oven at 65  $^{\circ}$ C, until a constant weight was obtained. The sample was ground in an Author Thomas

laboratory mill and sieved through 1 mm sieve to obtain a powder of the sample used for proximate analysis (AOAC, 2000).

Twenty grams of sample "B" was measured and blended in 180 ml sterile distilled water. One-milliliter portion was aseptically transferred into 9 ml distilled water and serial dilutions continued until a concentration of  $10^{-6}$  was obtained.

## **Determination of microbial counts**

One-tenth milliliter (0.1 ml) of the 4<sup>th</sup> and 6<sup>th</sup> dilutions obtained from sample "B" was aseptically inoculated onto freshly prepared surface-dried potato dextrose agar (PDA) and nutrient agar (NA) plates respectively. Inoculums were spread evenly over the surface of the plate with a sterile hockey stick-like spreader. The experiment was done in triplicate for each medium. PDA culture plates were incubated at room temperature for 5 days, while NA culture plates were incubated at 37 °C for 48 h. On establishment of growth, the number of bacteria colonies formed on each culture plate was counted using the Gallenkamp electronic colony counter. Fungi counts were obtained with a hand lens. The mean counts from the triplicate plate was obtained and multiplied with the appropriate dilution factor to obtain the total viable cells per unit weight of the sample expressed as the colony forming unit per grams (CFU/g) of the sample (Harrigan and McCance, 1990; ICMSF, 1978; Pelczar and Chan, 1977; Cheesbrough, 2000).

# Characterization and identification of isolates

Visible colonies of bacterial isolates were subjected to colonial characterization described in Cheesbrough (2000); Pelczar et al. (1993) and Pelczar and Chan (1977). Pure cultures of isolates were further characterized by microscopic and biochemical methods (Cheesbrough, 2000; Pelczar et al., 1993). The identities of the isolate was matched against those is Buchanan and Gibbon (1974).

Cultures of fungi were identified based on macro-and micromorphology, reverse and surface colouration of colonies grown on PDA (Pelczar and Chan, 1977; Harrigan and 1990: McCance. Abbey, 2007). Mycotoxigenic fungi of Aspergillus flavus and Aspergillus parasiticus were distinguished from the other Aspergillus species by a bright orange-yellow reverse colour using A. flavus and A. parasiticus medium prepared according to the recipe of Pitt and Hocking (1994) and Pitt et al. (1983). Penicillium identified using colony species were diameters, macro-and micromorphology according to the standardized conditions of Pitt (1979) and Pitt and Hocking (1985). Penicillium species were grown on three different media, namely, Czepak yeast extract agar, malt extract agar and 25% glycerol nitrate agar for 7 days in the dark at 5 °C, 25 °C and 37 °C respectively (Pitt and Hocking, 1985; Pitt and Hocking, 1994). Colony diameters were measured after 7 days. Other fungal isolates were matched against those in Barnnet and Hunters (1987).

# Determination of proximate composition of sample

The protein content of the sample was determined by the Kjedahl method adopted by James (1995) and Chang (2003) in which nitrogen content was determined and multiplied by 6.25. Ash content was determined by the furnace gravimetric method (James, 1995). The soxhlet solvent extraction method of James (1995) was employed in the determination of the fat content. Moisture content was determined by the gravimetric method of A.O.A.C. (2000). The Weenden gravimetric method of Kirk and Sawyer (1998) was used in the determination of crude fiber. Analysis was based on dry matter and expressed in percentage.

### RESULTS

The results obtained from the microbiological analysis of the larvae of

*Rhynchophorus* species revealed that both bacteria and fungi were involved in the contamination of the processed samples (Tables 1 and 2). The total bacterial and fungal counts were  $1.68 \times 10^5$  CFU/g and  $9.2 \times 10^2$  CFU/g respectively. The Tables also revealed the colonial characteristics of the isolates.

The microscopic and biochemical characterization of the bacteria isolated are presented in Tables 3 and 4 respectively,

while Table 5 shows the microscopic characteristics of the fungal isolates.

The identities of the bacteria and fungi isolated from the processed larvae are shown in Tables 4 and 5 respectively.

The percentage means value of the total energy and proximate composition of the sample is shown in Tables 6.

Table 1: Total heterotrophic counts and colonial characteristics of bacteria isolated on
Nutrient agar.

Colony code	Total counts	Colonial morphology
	(CFU/g)	
RNY 1		Rough creamy colonies with grey background
RNY 2		Smooth circular, cream low convex colonies
RNY 3		Large irregular cream colonies
RNY 4		Smooth and slimy with green pigment
RNY 5	$(1.68 \times 10^5)$	Large swarmy colonies
Number in parenth	pesis total counts of ha	cteria isolated

Number in parenthesis, total counts of bacteria isolated

Colony code	Total counts (CFU/g)	Colonial characteristics	Most probable identity
RNY A		Tall dark and upright colonies, branched	Cladosporum sp
		near the apex in clusters	
RNY B		Black-dark brown colonies with visible	Aspergillus sp
		tiny spores above the mycelia	
RNY C		Dark green rough surfaced velvety mass	Penicillium sp
		without clearly visible spores	
RNY D		Extensive Cotton-like mass of mycelia	<i>Fusarium</i> sp
		growth with slight pigmentation	
RNY E	$(9.2 \times 10^2)$	Whitish to cream coloured, round and	Saccharomyces
		butyrous colonies with limited growth	sp

**Table 2:** Total heterotrophic count and colonial characteristics of fungal isolates.

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Colony code	Mot	Gram rnx	Spore	Flagella	Capsule	Cell morphology	Most probable identity		
RNY 1	-	+	-	-	-	Long slender rod	Lactobacillus sp		
RNY 2	-	+	-	-	-	Oval cells in clusters	Staphylococcus sp		
RNY 3	-	+	+	-	-	Large short single rods	Bacillus sp		
RNY 4	+	-	-	+	-	Short small single rods	Pseudomonas sp		
RNY 5	+	-	-	+	-	Coccal to long filament	Proteus sp		

 Table 3: Microscopic characteristics of bacterial isolates.

Mot, motility; Gram rnx, Gram reaction

 Table 4: Biochemical characteristics of bacterial isolates.

Cat	Oxi	Coag	In	MR	VP	Cit	NO <sub>3</sub>	H <sub>2</sub> S	Ure	Glu	Suc	Ma	Lac	Man	Xyl	Identity of isolate
												1		n		
-	-	-	-	+	-	-	-	-	-	+	+	-	+	-	+	Lactobacillus plantarum
+	-	+	-	+	-	+	+	-	-	+	+	+	+	+	+	Staphylococcus aureus
+	-	-	-	+	-	+	+	+	-	+	+	+	+	+	+	Bacillus subtilis
+	+	-	-	+	-	+	+	+	-	+	+	+	-	-	-	Pseudomonas aeruginosa
+	-	-	+	+	-	-	+	+	+	+	+	+	-	-	+	Proteus vulgaris

Cat, catalase; Oxi, oxidase; Coag, coagulase; In, indole; MR, methyl red; VP, Voges Proskaeur; Cit, citrate; NO<sub>3</sub>, nitrate reduction; H<sub>2</sub>S, hydrogen sulphite production; Ure, urease production Glu, glucose; Suc, sucrose; Mal, maltose; Lac, lactose; Mann, mannitol; Xly, xylose

Colony code	Cell morphology	Identity of isolates
RNY A	Conidia seen as few cells with varying sizes and shapes.	Cladosporum sp
	Conidia borne of short-branched clusters and the	
	conidiophores had no swelling.	
RNY B	Simple upright unbranched conidiophores with tappers	Aspergillus flavus
	developing into an enlarge globase swelling at the apex.	
	Conidiophore was hyaline and conidia were seen borne on	
	short chain of sterigma.	
RNY C	Upright unbranched conidiophores which are septate	Penicillium
	giving a general picture of clustered fingers near the apex.	verrecosum
	Conidia were seen borne on philiade.	
RNY D	Short slender stocky irregular branched conidiophore.	Fusarium poae
	There appeared to be larger conidia on which smaller ones	
	were borne. Large conidia were seen as almost boat shaped	
	and roundish with thickened walls.	
RNY E	Oval cells, with larger cells attached with small round cells	Saccharomyces
	(small buds).	cerevisiae

**Table 5:** Microscopic characteristics of fungal isolates.

Table 6: Total energy and proximate composition of *Rhynchophorus* sp.

Sample	Total energy (Kcal/100 g)	Moisture	Crude protein	Fat	СНО	Crude fibre	ash
RNY	387.01±0.9	$12.14\pm0.1$	71.63±0.7	8.25±0.03	$6.56 \pm 1.0$	$1.59\pm0.2$	11.97±0.4
(Means ± SD	))						

## DISCUSSION

The results obtained from the microbiological analysis of the processed Rhynchophorus species indicates the presence of five genera of bacteria: Lactobacillus plantarum, Staphylococcus aureus, Bacillus subtilis, Pseudomonas aeruginosa and Proteus vulgaris, and four genera of moulds, namely, Cladosporum sp, Aspergillus flavus, Penicillium verrucosum and Fusarium poae, as well as one yeast species. The total bacterial count of 1.68 x 10<sup>5</sup> CFU/g suggests contamination and poses health risk following consumption.

*Lactobacillus plantarum* are effective food spoilage organisms (Harrigan and McCance, 1990). Their presence indicates that the sample is highly susceptible to spoilage resulting from improper handling and processing. The sample is rich in protein and other essential nutrients required for the growth of Lactobacillus plantarum and other organisms involved in food spoilage. Staphylococcus aureus and Bacillus subtilis are major food are ubiquitous and contaminants. They cause food spoilage and usually give rise to food borne illnesses (Nester et al., 1998; Pelczar et al., 1986; Prescott et al., 2002). Staphylococcus aureus cause food intoxication, when allowed to incubate in certain foods and produce heat stable enterotoxins that renders food dangerous even though it appears normal (Prescott et al., 2002). Bacillus subtilis is an endospore former and predominates in soil, and are usually associated with spoilage of starchy foods (Prescott et al., 2002; Nesters et al., 1998; Pelczar et al., 1986).

Proteus vulgaris are also food contaminants that act on food with high protein content. They rarely give rise to food borne infections, but generally lower the nutritional quality of contaminated food (Prescott et al., 2002). Pseudomonas aeruginosa are food contaminants with high water activity (a<sub>w</sub>). They are moist spoilage bacteria characteristic of the processed Rhynohophorus species and are frequently associated with high level food contamination (Harrigan and McCance, 1990; Prescott et al., 2002; Nester et al., 1998). Pseudomonas aeruginosa produce protease and lipases that can catalyze reactions causing degradation of proteins and fats resulting in an undesirable flavour of the food products (Nester et al., 1998).

The fungi species, *Cladosporum* and *Fusarium poae* are food contaminants that have rapid growth rates. They cause deterioration and spoilage of stored products. *Fusarium poae* had been implicated in the production of a potent mycotoxin (Prescott et al., 2002; Frazier and Westhoff, 1978). Some of the isolates are soil borne microorganisms associated with metamorphosing moth larvae (Braide et al., 2008).

The ubiquity and the ability of Aspergillus and Penicillum species to grow at low water activity make them important in post harvest/processed contamination and decay (Blomberg and Alder, 1992; Pitt, 1995). Aspergillus flavus are well known producers of aflatoxins. Aflatoxins whose primary target is the liver are potent carcinogens, mutagens and teratogens and are acutely toxic to animals and human (Siame et al., 1996). Penicillium verrucosum produce citrinin and ochratoxins, and is rated the most important source of ochratoxin in food (Abramson, 1991). Ochratoxin A is a potent nephrotoxin and causes kidney damage in pigs and experimental animals (Pitt and Hocking, 1994; Abramson, 1991).

Results obtained from the proximate analysis showed that the processed larvae of

Rhynchophorus is rich in protein (71.63%), which indicates high nutritional quality as well as susceptibility to spoilage microorganisms. The presence of crude fibre and ash in high quantity is important as they aid bowel peristalsis and provide minerals to the body respectively (Wardlaw and Kessel, 2002; Delvin, 2006). The moisture content (12.14%) is high and is sufficient to encourage the growth and proliferation of food spoilage and disease causing bacteria and fungi.

Poor sanitation may be an important factor in the contamination of processed larvae sold and consumed. The method of collection and utensils used in the preparation together with inadequate storage and marketing may all contribute as sources of contamination. Therefore, use of clean water and utensils, adequate packaging and practice of good sanitary and hygiene within and around the environment could reduce recontamination of the ready-to-eat products.

Preparation of samples with spices containing antimicrobial compounds may discourage bacterial contamination. Spices such as garlic (*Allium sativum*) and Negro pepper (*Xylopia aethiopica*) had been reported to contain some antimicrobial compounds (Iwu et al., 1999). Furthermore, lowering the water activity level through curing or drying will reduce the growth of xerophilic and xeroduric organisms.

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