



Antimicrobial Activity of Extract from the Abdominal Gland of African Weaver Ants (*Oecophylla longinoda*) on the Microorganisms Isolated from Cocoa Pods

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Abstract

This research was conducted to investigate into the antimicrobial potency of the abdominal gland extract of African Weaver Ant (AWA) against microorganisms isolated from Cocoa pods. Six (6) fresh samples of both infected and uninfected Cocoa pods gotten from a village farm in Ibule-soro, Ondo State were transferred to the laboratory under aseptic conditions. One gram from the pod of each sample was subjected to microbial analysis for isolation and the antimicrobial effects of the AWA abdominal extract was subsequently evaluated. Selected conventional antibiotics were used as controls. Bacterial isolates obtained were *Bacillus subtilis*, *Pseudomonas aeruginosa*, *Serratia fonticola*, *Proteus penneri*, *Enterobacter cloacae*, *Citrobacter freundii*, *Edwardsiella tarda*, *Klebsiella oxytoca* and *Proteus mirabilis*, while fungal isolates were *Aspergillus aculeatinus*, *Alternaria* sp., *Cloridium chlamydosporis*, *Aspergillus niger*, *Phytophthora palmivora*, *Aspergillus flavus*, *Aspergillus tamari*, *Fusarium* sp., *Penicillium chrysogenum*, *Penicillium camemberti*, *Saccharomyces cerevisiae*, *Saccharomyces ludwigii*. Four of all the bacterial isolates (*Bacillus subtilis*, *Edwardsiella tarda*, *Proteus penneri*, *Proteus mirabilis*) used for this study exhibited high levels of sensitivity to the abdominal gland of African weaver ants with the following zones of inhibition: 19 mm, 19 mm, 20 mm and 21 mm respectively. Similarly, fungal isolates were observed to be sensitive to the *Oecophylla longinoda* abdominal extract, with *Aspergillus flavus* having the highest zone of inhibition (14.89 mm). This research reveals the potentials of abdominal extracts from African weaver ants (*Oecophylla longinoda*) in antimicrobial capacities against most aetiological agents of Cocoa diseases.

Keywords: Cocoa; *Oecophylla Longinoda*; Antibacterial; Antifungal

Introduction

Cacao, a species in the genus *Theobroma*, is the most economically important species in this genus (Acebo-Guerrero 2012). It is an upright, evergreen tree that grows up to 15 metres, but it is usually kept below 5 metres to make for easy harvesting. The tree has a polymorphic characteristic - its seedling grows to between 1 to 2 m in height as a single stem after which it separates into the jorquette (a whorl of lateral-growing 3-5 branches). Two to three years after planting, flowers begin to grow from pads or cushions beneath the bark. Under optimal conditions these pads will continue to produce flowers for 60 to 100 years. Some self-incompatible cacao varieties usually require cross-pollination, while others that are completely self-compatible do not need to be cross-pollinated. The cacao fruit (pod) is cut from the tree without impairing the flower pad below the surface because impaired flower pads will no longer bear fruit. The pods are harvested for the seeds (beans) usually found inside them. Each pod contains about 30 to 60 beans, these are enclosed with white mucilage that is high in sugar content. Pods are picked at maturity for their primary use which is the production of chocolate. Cacao can be grown only within the tropical belt [1]. Ivory Coast on the west coast of Africa, is reputed to be the largest cacao-producing region where 68% of the world's cacao is being harvested. *Theobroma cacao* is believed to have its

origin in The Amazon Basin which makes up only a portion of the 15% of world production from the Americas [2]. *Phytophthora* pod rot (PPR or black pod) and Cocoa swollen shoot virus (CSSV) are two diseases considered to be highly important in the production of cocoa, these pathogens have consequently received great attention in Nigeria. *Phytophthora* pod rot caused by *P. palmivora* and *P. megakarya*, was acclaimed the most important yield-limiting factor in the Nigerian cocoa industry, with total annual losses estimated at up to 30 to 35% of the nation's crop [3], while 30 - 90% of the total global crop loss is caused by this disease [4]. However, severity of this disease varies depending on climatic factors like rainfall and humidity and the variety of cocoa concerned. Seedling diseases, cankers of stems, trunks and crowns of cocoa trees could be resultant from *Phytophthora* spp. infestations. The sources of inoculum for PPR are soil, ant tents, infected flower cushions, dry pods on the cocoa trees or ground, debris formed from heaping of cocoa pod husks. Spread from infected pods is by water splash and drips during rains, living vectors, wind-blown sprays and wounds or non-intact surfaces on pods usually caused by knives, rodents

Cocoa is a very important cash crop but the advent of the black pod disease caused by *Phytophthora megakarya* has greatly reduced the revenue generated by farmers. *P. megakarya* black pod is the single most important fungal disease of cocoa which has

been described by some farmers as a strange disease [8], this is due to the extreme virulence and the heavy losses associated with the pathogen. Many highly or extremely toxic pesticides (Class I) are being used in developing countries [9]. Spilling of pesticides over the back of the operator owing to a faulty locking cap of the container is a major cause of poisoning when using knapsack or trombone sprayer [10].

Various species of Weaver ants (*Oecophylla* spp.) have been used for pest management in tropical agro-ecosystems across Australia, Asia and Africa [11,12]. They have proven to be highly effective biological control agents being capable of controlling over 50 species of insect pests in at least 12 different tropical crops [11,13]. The use of conventional synthetic insecticide control was substituted with weaver ants and this led to increased net incomes of more than 70% in Australian cashew and mango plantations [14,15]. The peculiarities of extremely organized predatory behaviour, ability to forage the entire areas occupied by colonies and their potential to expand into new areas are responsible for the success of weaver ants in killing or driving out many potential pests [16,17]. In the year 1991, Dejean demonstrated the predatory efficiency of the African weaver ant *Oecophylla longinoda* and estimated that a colony of 12 nests captured approximately 45,000 prey individuals per year [18].

Oecophylla worker ants are utilised directly as a commercial product, as they are reported to be a source of traditional Chinese and Indian medicines against arthritis and some other abnormal health conditions [19,20]. Cocoa as a cash crop has over the years been faced with various diseases which have resulted into reduced yield and decrease in the total revenue of cocoa farmers. Consequently, some cocoa farms have been abandoned while other farmers have switched over to the cultivation of other crops [8]. This research is therefore designed to investigate into the antimicrobial properties of extracts from the abdominal gland of *O. longinoda* in the prevention and treatment of diseases affecting Cocoa. This research aims to isolate and characterize the microorganisms from the infected and uninfected Cocoa pods, to evaluate the antimicrobial properties of extract from the abdominal gland of the African Weaver Ants (*Oecophylla longinoda*) using conventional antibacterial and antifungal agents as a control study group to the research.

Materials and Methods

Collection of Samples

Six samples, (three Infected and three uninfected) Cocoa samples were collected from a selected farm in Ibule – soro, Ondo State, aseptically into already sterilized containers and were transported to the Microbiology research laboratory of the Federal University of Technology, Akure for microbial analysis.

Sterilization of Materials and Media

All glassware used were properly washed with detergent and rinsed thoroughly with water, allowed to dry and sterilized in the hot air oven at 160°C for 2 hours. The culture media used were prepared in the conical flasks following the manufacturer's standard. Test tubes used were plugged with cotton wool and wrapped with aluminium foil. Inoculating loops and needles were sterilized using a Bunsen burner till red heat was acquired. Work surfaces were swabbed with cotton wool soaked in 70% alcohol before and after each analysis. Hands were washed with an antiseptic soap before and after each experiment to avoid contamination and infection.

Isolation of Bacteria from Cocoa pods

The surface of the pod was washed properly with sterile distilled water [21-23]. One gram (1g) of the Cocoa pod was cut from each of the infected and uninfected pods and ground using mortar and pestle. This was transferred into the first test tube (containing

10 ml of sterile distilled water) and diluted serially in a stepwise manner (in test tubes with 9 ml of sterile distilled water) to 10⁻⁷ dilution test tube. One millilitre (1.0 ml) each of the dilution 10⁻⁵, 10⁻⁶ and 10⁻⁷ was plated on Nutrient agar, the plates were inverted and incubated at 37°C for 24h after which the plates were examined for growth.

Isolation and Characterization of Bacterial Isolates

The bacterial growth on nutrient agar was sub-cultured by streaking until pure colonies were obtained according to the conventional procedure as highlighted by [24]. Cultural characteristics of the discrete bacterial colonies such as colour, shape, pigmentation and opacity on nutrient agar were observed and noted. This was followed by characterization of the isolates [25] that includes microscopic examination for cell morphology, gram reaction and biochemical tests on 24h old cultures. Discrete bacterial colonies were cultured on double strength nutrient agar slant and incubated at 37°C for 24h, growth was observed and the slants were stored in the refrigerator for preservation of bacterial isolates.

Morphological and Biochemical Characterization of Bacterial Isolates

Different tests were carried out on isolates which include; Gram reaction, Urease, Oxidase, Voges Proskauer, Catalase test [24], Indole production, Motility and Citrate utilization test [25]. Bacterial isolates were identified using standard cultural, morphological, and biochemical characteristics as described by Cheesbrough [25].

Hydrogen Sulphide and Gas Production on Triple Sugar Iron (TSI) agar

The medium was prepared according to the manufacturer's specification. Five milliliter (5 ml) was dispensed into sterilized test tube and autoclaved at 121°C for 15 minutes. The medium was allowed to solidify in a slanting position to give a butt 25 - 30 mm deep and a slope 20 - 25 mm long. A straight wire was used to inoculate TSI agar slope medium with the inoculum of the test isolate by stabbing the butt and then streaking the surface of the slope in a zig-zag pattern and incubate at 37°C for 24h. TSI reactions are based on the fermentation of lactose and glucose (dextrose) and the production of hydrogen sulphide. Presence of cracks in the tube indicates gas production and black colouration indicates hydrogen sulphide production [25].

McFarland Turbidity Standard

Standardization of inoculum was done using WPA colorimeter. Five milliliter (5 ml) of sterile normal saline in test tubes of equal sizes was inoculated with 18 - 24h old culture. An uninoculated test tube serves as a control to calibrate the colorimeter. The suspension of each isolate was adjusted using 0.5 McFarland standard at 590 nm to obtain about 10⁶ - 10⁸ cfu/ml for the isolates

Antibacterial Susceptibility Test

Antibiotic resistance of bacteria was determined by the single disc diffusion method with the use of Mueller-Hinton agar, according to the Kirby-Bauer's method. The medium used was Mueller-Hinton agar (MHA) in petri dishes. The surface of the agar was aseptically seeded with a standardized saline suspension of the isolate using sterile swab sticks. Antibiotic discs consisting Pefloxacin (10 µg), Gentamycin (10 µg), Ampiclox (30 µg), Zinacef (20 µg), Amoxicillin (30 µg), Rocephin (25 µg), Ciprofloxacin (10 µg), Streptomycin (30 µg), Septrin (30 µg), Erythromycin (10 µg), Chloramphenicol (30 µg), Sparfloxacin (10 µg), Tarivid (30 µg), Augmentin (25 µg) were aseptically placed on the surface of the MHA and the plates were incubated at 37°C for 18 - 24 hours. The diameters of the zones of inhibition were then measured using a transparent meter rule to the nearest millimeter.

Antibacterial assay of the Ants

The abdomen of the *Oecophylla longinoda* ant was cut and crushed using mortar and pestle and then homogenized with Dimethyl sulfoxide (DMSO) under aseptic conditions and diluted into three different concentrations in such a way that 100 µl homogenate contained glands of 3, 6 and 12 the assay was conducted using agar well diffusion method [26].

Isolation of Fungi from Cocoa Pods

The surface of the pod was washed properly with sterile distilled water [21-23]. One gram (1g) of the Cocoa pod was cut from each of the infected and uninfected pods and ground using mortar and pestle. This was transferred into the first test tube (containing 10 ml of sterile distilled water) and diluted serially in a stepwise manner (in test tubes with 9ml of sterile distilled water) to 10⁻⁷ dilution test tube. One millilitre (1.0 ml) each of the dilution 10⁻³, 10⁻⁴ and 10⁻⁵ was plated on Potato dextrose agar, the plates were incubated aerobically at 25 ± 2°C for 3 - 7 days after which the plates were examined for growth.

Fungal mycelia were teased with sterile inoculating needle. With two drops of Lactophenol - cotton blue dye were added to the sample on a clean glass slide and covered with slip and then viewed under the light microscope using medium and high power objective lenses. The Fungal characterization was based on micro morphological features which include the following; nature of conidia shape, sporangiophore, arthrospores, spore head, rhizoid and hyphae (septate or non-septate). Isolated fungal colonies were identified for both microscopic and macroscopic examinations [24].

Antifungal Susceptibility Test of Fungi isolated from Cocoa pods

This test was used to determine the sensitivity of fungal isolates to selected antifungal agents. Four commercial antifungal agents (Griseofulvin, Ketoconazole, Itraconazole and Clotrimazole) were used in this assay. These were reconstituted to a concentration of 40 mg/ml each and agar well method was adopted for the test. The medium used was Mueller-Hinton agar. The surface of the agar was aseptically seeded with a standardized saline suspension of the isolate using sterile swab sticks and the plates were incubated at 37°C for 24 hours. The diameter of the zone of inhibition was then measured using a transparent meter rule to the nearest millimeter.

Antifungal assay of the Ants

The abdomen of the *Oecophylla longinoda* ant was cut and crushed using mortar and pestle and then homogenized with DMSO under aseptic conditions and diluted into three different concentrations in such a way that 100 µl homogenate contained glands of 3, 6 and 12 using agar well diffusion method [26].

Results

Bacterial load on the Cocoa Pods

Mean total bacterial counts were shown in table 1. The result revealed there was significant difference (p ≤ 0.05) in the bacte-

rial load of the Cocoa pods. The bacterial load on the Cocoa pods showed a range of 15.00 ± 0.29 x 10⁶ cfu/g to 20.00 ± 1.15 x 10⁶ cfu/g in infected pods and 5.00 ± 0.57 x 10⁶ cfu/g to 10.00 ± 0.98 x 10⁶ cfu/g in uninfected pods. Bacterial load of infected sample 1 was significantly (p ≤ 0.05) higher than other samples.

Samples	Infected Cocoa pods (cfu/g x 10 ⁶)	Uninfected Cocoa pods (cfu/g x 10 ⁶)
1	20.00 ± 1.15 ^b	10.00 ± 0.98 ^b
2	15.00 ± 0.29 ^a	7.00 ± 0.34 ^a
3	19.00 ± 0.69 ^b	5.00 ± 0.57 ^a

Table 1: Bacterial Load on the Cocoa Pods.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at p ≤ 0.05

Key: SE: Standard Error; Cfu: Colony Forming Unit

Fungal load on the Cocoa Pods

Mean total fungal spore counts were shown in table 2. The result revealed there was significant difference (p ≤ 0.05) in total fungal spores count load of the Cocoa pods. The fungal load of Cocoa pods showed a range of 2.67 ± 0.33 x 10⁶ sfu/g to 11.33 ± 0.88 x 10⁶ sfu/g in infected pods and 1.33 ± 0.33 x 10⁶ sfu/g to 2.33 ± 0.33 x 10⁶ sfu/g in uninfected pods. Fungal load of infected sample 1 was significantly (p ≤ 0.05) higher than other samples.

Samples	Infected (sfu/g x 10 ³)	Uninfected (sfu/g x 10 ³)
1	11.33 ± 0.88 ^c	1.33 ± 0.33 ^a
2	6.00 ± 0.58 ^b	1.66 ± 0.33 ^a
3	2.67 ± 0.33 ^a	2.33 ± 0.33 ^a

Table 2: Fungal Load on Cocoa Pods.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at p ≤ 0.05

Key: SE: Standard Error; Sfu: Spore Forming Unit

Morphological, Biochemical Characteristics and Identification of isolates from Cocoa Pods

Table 3 shows biochemical tests that were carried out on the bacterial isolates; Gram stain, Cell shape, motility, catalase, citrate, oxidase, indole, reaction on Triple Sugar Iron. All the bacterial isolates show different biochemical reactions and were characterized morphologically based on microscopy into cocci and bacilli. The isolates were identified as *Bacillus subtilis*, *Pseudomonas aeruginosa*, *Serratia fonticola*, *Proteus penneri*, *Enterobacter cloacae*, *Citrobacter freundii*, *Edwardsiella tarda*, *Klebsiella oxytoca* and *Proteus mirabilis*.

Isolate code	Gram's reaction/shape	Motility test	Catalase test	Citrate test	Oxidase test	Indole Test	Top	Butt	H ₂ S	Gas production	Methyl red	Voges-Proskauer	Urease test	Probable Organisms
I ₁ a	+ /rod	+	+	+	-	-	Y	R	-	-	-	+	-	<i>Bacillus subtilis</i>
I ₁ b	- /rod	+	+	+	+	-	R	R	-	-	-	-	-	<i>Pseudomonas aeruginosa</i>
I ₂ a	- /rod	+	+	-	-	-	Y	Y	-	+	-	+	-	<i>Serratia fonticola</i>
I ₂ b	+ /rod	+	+	-	-	-	R	Y	-	-	-	+	-	<i>Bacillus subtilis</i>
I ₂ c	- /rod	+	+	-	-	-	R	Y	-	-	+	-	+	<i>Proteus penneri</i>
I ₃ a	- /rod	+	+	-	-	-	Y	Y	-	+	-	+	-	<i>Enterobacter cloacae</i>
I ₃ b	- /rod	+	+	-	-	-	Y	Y	-	+	+	+	-	<i>Citrobacter freundii</i>
I ₃ c	- /rod	+	+	-	-	-	R	Y	+	+	-	-	-	<i>Edwardsiella tarda</i>
I ₃ d	- /rod	+	+	-	-	-	Y	Y	-	+	+	+	-	<i>Citrobacter freundii</i>
UI ₁ a	- /rod	+	+	+	-	-	Y	Y	-	+	-	+	-	<i>Klebsiella oxytoca</i>
UI ₁ b	- /rod	+	+	-	-	-	R	Y	-	+	+	-	+	<i>Proteus penneri</i>
UI ₂ a	- /rod	+	+	-	-	-	R	Y	+	+	+	-	+	<i>Proteus mirabilis</i>
UI ₃ a	- /rod	+	+	-	+	-	Y	R	-	+	-	-	-	<i>Pseudomonas aeruginosa</i>

Table 3: Morphological, Biochemical Characteristics and identification of Bacteria isolated from Cocoa pods.

Key: +: Positive; -: Negative; Y: Yellow; R: Red

Identification and cultural characteristics of fungi isolated from Cocoa pods

Table 4 revealed the fungal characterization which was based on micro and macro morphological features that include the following; nature of conidia shape, sporangiophore, arthrospores, spores head, rhizoid and hyphae (septate or non-septate). Isolated fungal colonies were identified from both microscopic and macroscopic examinations and they were identified to be *Aspergillus aculeatinus*, *Penicillium chrysogenum*, *Cloridium chlamydo sporis*, *Aspergillus niger*, *Saccharomyces ludwigii*, *Phytophthora palmivora*, *Aspergillus flavus*, *Saccharomyces cerevisiae*, *Aspergillus tamari*, *Fusarium sp.*, *Penicillium camemberti*, *Alternaria species*.

Fungal isolates	Morphological and microscopy characterization
<i>Aspergillus aculeatinus</i>	Dark green colouration on agar plate, clavate swelling, conidiophores not too long, upright, conidia globose or ovoid, conidophores standing inform of mortar pistle, thick with swollen head.
<i>Penicillium chrysogenum</i>	Green colonies on culture. Conidia are unicellular round to ovoid, hyaline or pigmented. Phialides grouped in brush-like cluster at the end of the conidiophores.
<i>Penicillium camemberti</i>	White to pink or orange colonies. Mycelium consists of a highly branched network of multinucleate septate.
<i>Alternaria sp.</i>	Colonies are fast growing, black or greyish. Conidia are obclavate, sometimes ovoid or ellipsoidal, often with a short conical break smooth-walled or verrucose.
<i>Cloridium chlamydo sporis</i>	Black on culture. Conidiophores slender and swollen, lightly pigmented, simple or septate, conidia subhyaline, 1-celled globose or oblong.
<i>Aspergillus niger</i>	Black colouration with black conidia with white lining on the culture. Conidiophores are very thick and long.
<i>Saccharomyces ludwigii</i>	Cream to yellow colonies with budding cells.
<i>Phytophthora palmivora</i>	White colonies with uneven edges. From microscopic observation, there was presence of hyphae hyaline, branches and non-partition.
<i>Aspergillus flavus</i>	Green colonies on culture, globose conidia. The hyphae is branched and the septate conidiophore are long, rough, septate and granular.
<i>Saccharomyces cerevisiae</i>	White to cream, smooth, gliborous yeast-like colonies. Large globose to ellipsoidal cells with multilateral budding.
<i>Aspergillus tamari</i>	Dirty green colouration on the culture, dense spotting on culture
<i>Fusarium culmorum</i>	White cotton wool-like mould growth with no distinct aerial mycelium. Conidiophore are short, single, lateral monophialides

Table 4: Morphological characteristics of fungal isolates.

Comparative distribution and occurrence of bacteria isolated from Cocoa pods

The total number of bacteria isolated from each of the samples collected were presented in table 5a, the occurrence of each bacterial isolates from both the infected and uninfected pods as well as the percentage occurrences were also presented.

Isolates	Infected			Uninfected			Total	Occurrence (%)
	A	B	C	A	B	C		
<i>Bacillus subtilis</i>	+	+	-	-	-	-	2	9.1
<i>Pseudomonas aeruginosa</i>	+	-	-	-	+	-	2	9.1
<i>Serratia fonticola</i>		+	-	-	-	-	1	4.5
<i>Proteus penneri</i>		+	+	-	-	-	2	9.1
<i>Enterobacter cloacae</i>	+	-	+	-	-	-	2	9.1
<i>Citrobacter freundii</i>		-	+	-	-	-	1	4.5
<i>Edwardsiella tarda</i>		+	-	+	-	-	2	9.1
<i>Klebsiella oxytoca</i>		-	-	+	-	+	2	9.1
<i>Proteus penneri</i>	+	+	+	+	+	+	6	27.3
<i>Proteus mirabilis</i>	-	+	-	-	+	-	2	9.1
Total	4	6	4	2	3	3	22	100

Table 5a: Percentage occurrence of bacteria isolated from Cocoa pods.

Key: +: Present; -: Absent.

Comparative distribution and occurrence of fungi isolated from Cocoa pods

The total number of fungal species isolated from each of the samples collected were presented in table 5b, the occurrence of each of the fungal isolates from both the infected and the uninfected as well as the percentage occurrences were also presented.

Susceptibility pattern of Gram negative bacteria

All the isolated bacteria were resistant to Septrin, Chloramphenicol, Amoxicillin, Augmentin with zone (0 mm -7 mm) and *Pseudomonas* species gave intermediate resistance to Streptomycin and Tarivid with zone (8 mm -10 mm) as revealed in table 6. On the contrary, *Pseudomonas aeruginosa*, *Citrobacter freundii*, *Klebsiella oxytoca*, *Proteus mirabilis*, *Pseudomonas* spp. were all susceptible to Ciprofloxacin. *Citrobacter freundii*, *Klebsiella oxytoca*, *Proteus mirabilis* and *Pseudomonas* spp. also displayed susceptibility to Pefloxacin. Then, *Citrobacter freundii*, *Klebsiella oxytoca*, *Proteus mirabilis* showed susceptibility to Tarivid whereas only, *Pseudomonas aeruginosa*, *Klebsiella oxytoca*, *Proteus mirabilis* were susceptible to Sparfloxacin.

Isolates	Infected			Uninfected			Total	Occurrence (%)
	A	B	C	A	B	C		
<i>Aspergillus aculeatinus</i>	+	-	-	-	-	-	1	5
<i>Penicillium chrysogenum</i>	+	-	-	-	-	-	1	5
<i>Chloridium chlamyosporis</i>		+	-	-	-	-	1	5
<i>Aspergillus niger</i>		-	+	-	-	-	1	5
<i>Saccharomyces ludwigii</i>		+	+	-	-	-	2	10
<i>Phytophthora palmivora</i>	+	-	-	-	-	-	1	5
<i>Aspergillus flavus</i>		-	-	+	-	-	1	5
<i>Saccharomyces cerevisiae</i>		+	-	+	-	-	2	10
<i>Aspergillus tamari</i>		-	-	-	+	-	1	5
<i>Fusarium sp.</i>	+	+	+	+	+	+	6	30
<i>Penicillium camemberti</i>	+	-	+	-	-	-	2	10
<i>Alternaria sp.</i>	-	+	-	-	-	-	1	5
Total	5	5	4	2	2	2	20	100

Table 5b: Percentage occurrence of fungi isolated from Cocoa pods.

Key: +: Present; -: Absent

Isolate code	OFX (µg)	S (µg)	SXT (µg)	CH (µg)	SP (µg)	CPX (10 µg)	AM (30 µg)	AU (25 µg)	CN (10 µg)	PEF (10 µg)
I ₁ b	10.00 ± 0.03 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	6.03 ± 0.41 ^b	12.03 ± 0.32 ^b	12.17 ± 0.32 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	7.00 ± 0.33 ^b
I ₂ a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
I ₂ c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
I ₃ a	12.00 ± 0.12 ^c	10.01 ± 0.45 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	15.00 ± 0.45 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	7.00 ± 0.94 ^b	14.00 ± 0.35 ^c
I ₃ b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
I ₃ c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
UI ₁ a	12.00 ± 0.23 ^c	12.00 ± 0.21 ^c	7.02 ± 0.01 ^b	0.00 ± 0.00 ^a	12.00 ± 0.44 ^b	14.00 ± 0.43 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	12.02 ± 0.36 ^b
UI ₂ a	12.00 ± 0.11 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	6.03 ± 0.22 ^b	12.00 ± 0.16 ^b	12.00 ± 0.35 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	7.00 ± 0.52 ^b	12.00 ± 0.01 ^b
UI ₃ a	10.00 ± 0.15 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	7.00 ± 0.18 ^b	0.00 ± 0.00 ^a	12.00 ± 0.19 ^b	7.20 ± 0.22 ^b	6.00 ± 0.91 ^b	7.10 ± 0.58 ^b	12.00 ± 0.00 ^b

Table 6: Susceptibility pattern of Gram negative bacteria.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at $p \leq 0.05$

Key: I₁b: *Pseudomonas aeruginosa*; I₂a: *Serratia fonticola*; I₂c: *Proteus penneri*; I₃a: *Enterobacter cloacae*; I₃b: *Citrobacter freundii*; I₃c: *Edwardsiella tarda*; UI₁a: *Klebsiella oxytoca*; UI₂a: *Proteus mirabilis*; UI₃a: *Pseudomonas* species; OFX: Tarivid; S: Streptomycin; SXT: Septrin; CH: Chloramphenicol; SP: Sparfloxacin; CPX: Ciprofloxacin; AM: Amoxicillin; AU: Augmentin; CN: Gentamycin; PEF: Pefloxacin

Susceptibility pattern of Gram positive bacteria

Bacillus subtilis revealed high susceptibility to Ciprofloxacin, Rocephin, Pefloxacin and Erythromycin, while *Bacillus* species showed high susceptibility to Rocephin only as seen in table 7. On the other hand, *Bacillus* species were resistant to the remaining antibiotics.

Antifungal susceptibility of fungal isolates

The result in table 8 revealed the susceptibility of fungal isolates to commercial antifungal agents. Only *Phytophthora palmivora* was susceptible to Griseofulvin while the other fungi were not susceptible; only *Penicillium camemberti*

was not susceptible to Ketoconazole, while all the other fungi were susceptible; *Penicillium chrysogenum*, *Penicillium camemberti*, *Alternaria* species were not susceptible to Itraconazole while other fungi were susceptible; *Alternaria* species, *Aspergillus niger*, *Saccharomyces cerevisiae* were not susceptible to Clotrimazole while others were susceptible; DMSO is the negative control.

Antibiotic activity of abdominal gland extract of *O. longinoda*

All the bacterial isolates were susceptible to both the extract and Ciprofloxacin except *Klebsiella oxytoca* which was not susceptible to Ciprofloxacin as revealed in table 9.

Isolate Code	S (30 µg)	CPX (10 µg)	R (25 µg)	AM (30 µg)	Z (20 µg)	APX (30 µg)	CN (10 µg)	PEF (10 µg)	E (10 µg)	SXT (30 µg)
I ₁ a	0.00 ± 0.00 ^a	17.00 ± 0.21 ^b	12.00 ± 0.21 ^b	7.01 ± 0.42 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	14.01 ± 0.31 ^b	12.20 ± 0.22 ^b	0.00 ± 0.00 ^a
I ₂ b	6.00 ± 0.21 ^b	7.02 ± 0.01 ^a	22.00 ± 0.00 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a

Table 7: Susceptibility pattern of Gram positive bacteria.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at $p \leq 0.05$.

Key: I₁a: *Bacillus subtilis*; I₂b: *Bacillus* species; S: Streptomycin; CPX: Ciprofloxacin; R: Rocephin; AM: Amoxicillin; Z: Zinacef; APX: Ampiclox; CN: Gentamycin; PEF: Pefloxacin; E: Erythromycin; SXT: Septrin.

Fungal Isolate	A (mm)	B (mm)	C (mm)	D (mm)	E (mm)
<i>Aspergillus aculeatinus</i>	0.00 ± 0.00 ^a	20.68 ± 0.17 ^e	12.03 ± 0.26 ^b	24.00 ± 0.35 ^f	0.00 ± 0.00 ^a
<i>Penicillium chrysogenum</i>	0.00 ± 0.00 ^a	11.97 ± 0.55 ^b	0.00 ± 0.00 ^a	24.00 ± 0.00 ^f	0.00 ± 0.00 ^a
<i>Penicillium camemberti</i>	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	24.03 ± 0.51 ^f	0.00 ± 0.00 ^a
<i>Alternaria sp</i>	0.00 ± 0.00 ^a	34.00 ± 0.23 ^h	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Chloridium chlamyosporis</i>	0.00 ± 0.00 ^a	24.00 ± 0.11 ^g	29.00 ± 0.05	9.00 ± 0.51 ^b	0.00 ± 0.00 ^a
<i>Aspergillus niger</i>	0.00 ± 0.00 ^a	16.03 ± 0.38 ^c	16.03 ± 0.54 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Saccharomyces ludwigii</i>	0.00 ± 0.00 ^a	24.00 ± 0.00 ^g	24.00 ± 0.03 ^g	19.03 ± 0.61 ^e	0.00 ± 0.00 ^a
<i>Phytophthora palmivora</i>	13.96 ± 0.54 ^b	24.06 ± 0.18 ^g	24.00 ± 0.05 ^g	34.00 ± 0.58 ^g	0.00 ± 0.00 ^a
<i>Aspergillus flavus</i>	0.00 ± 0.00 ^a	22.97 ± 0.43 ^f	23.00 ± 0.00 ^f	24.03 ± 0.19 ^f	0.00 ± 0.00 ^a
<i>Saccharomyces cerevisiae</i>	0.00 ± 0.00 ^a	20.00 ± 0.05 ^e	20.03 ± 0.54 ^d	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Aspergillus tamari</i>	0.00 ± 0.00 ^a	24.00 ± 0.58 ^g	23.96 ± 0.55 ^f	16.00 ± 0.41 ^d	0.00 ± 0.00 ^a
<i>Fusarium sp</i>	0.00 ± 0.00 ^a	19.00 ± 0.00 ^d	12.03 ± 0.04 ^b	12.00 ± 0.58 ^c	0.00 ± 0.00 ^a

Table 8: Antifungal Sensitivity Test.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at $p \leq 0.05$.

Key: A: Griseofulvin; B: Ketoconazole; C: Itraconazole; D: Clotrimazole; E: DMSO

Bacterial Isolate	A (mm)	B (mm)	C (mm)	D (mm)
<i>Pseudomonas aeruginosa</i>	14.00 ± 0.29 ^d	14.00 ± 0.00 ^{de}	14.00 ± 0.00 ^d	16.00 ± 0.46 ^d
<i>Serratia fonticola</i>	9.00 ± 0.23 ^b	12.00 ± 0.55 ^d	13.00 ± 0.00 ^c	16.00 ± 0.00 ^d
<i>Bacillus subtilis</i>	19.00 ± 0.06 ^e	13.03 ± 0.06 ^f	14.00 ± 0.29 ^d	19.00 ± 0.23 ^b
<i>Proteus penneri</i>	11.03 ± 0.04 ^c	10.00 ± 0.58 ^c	12.00 ± 0.00 ^c	20.00 ± 0.00 ^f
<i>Enterobacter cloacae</i>	9.03 ± 0.50 ^b	9.00 ± 0.58 ^{de}	11.00 ± 0.12 ^b	14.01 ± 0.46 ^b
<i>Citrobacter freundii</i>	14.00 ± 0.00	9.00 ± 0.17 ^{de}	12.03 ± 0.06 ^b	14.00 ± 0.00 ^b
<i>Edwardsiella tarda</i>	11.00 ± 0.01 ^c	14.33 ± 0.44 ^e	14.03 ± 0.08	19.00 ± 0.56
<i>Klebsiella oxytoca</i>	0.00 ± 0.00 ^a	10.00 ± 0.06 ^c	12.00 ± 0.00 ^c	15.00 ± 0.00 ^c
<i>Proteus mirabilis</i>	9.00 ± 0.00 ^b	5.00 ± 0.56 ^b	15.03 ± 0.00 ^e	21.03 ± 0.09 ^g

Table 9: Antibiotic activity of abdominal gland extract of *O. longinoda*.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at $p \leq 0.05$.

Antibacterial activity of abdominal gland extract of *O. longinoda*. B, C and D - varying concentrations of abdominal gland extract. Concentration B is gland from three ants; concentration C is gland from six ants; concentration D, gland from twelve ants. A is, Ciprofloxacin std 100 µl (40 mg/ml).

Antifungal activity of abdominal gland extract of *O. longinoda*

All the fungal isolates were susceptible to the extract except for *Penicillium chrysogenum*, *Penicillium camemberti*, *Saccharomyces ludwigii*, *Phytophthora palmivora* and *Fusarium* sp. which were not susceptible; of all the twelve isolated fungi tested with Ketoconazole, eleven (*Aspergillus aculeatinus*, *Alternaria species*, *Chloridium chlamydosporis*, *Aspergillus niger*, *Phytophthora palmivora*, *Aspergillus flavus*, *Aspergillus tamari*, *Fusarium* sp., *Saccharomyces cerevisiae*, *Saccharomyces ludwigii*, *Penicillium chrysogenum*) were susceptible while the only one that was not susceptible is *Penicillium camemberti*.

Fungal Isolate (code is to be removed)	A (mm)	B (mm)	C (mm)	D (mm)
<i>Aspergillus aculeatinus</i>	19.05 ± 0.14 ^b	8.95 ± 0.10 ^c	10.00 ± 0.01 ^b	11.00 ± 0.00 ^c
<i>Penicillium chrysogenum</i>	11.90 ± 0.15 ^d	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Penicillium camemberti</i>	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Alternaria</i> sp.	24.33 ± 0.30 ^c	8.00 ± 0.22 ^d	10.03 ± 0.00 ^b	12.51 ± 0.21 ^c
<i>Chloridium chlamydosporis</i>	24.36 ± 0.05 ^c	6.00 ± 0.32 ^b	7.90 ± 0.03 ^b	9.22 ± 0.25 ^b
<i>Aspergillus niger</i>	24.00 ± 0.03 ^c	8.34 ± 0.15 ^b	9.17 ± 0.33	10.12 ± 0.22 ^b
<i>Saccharomyces ludwigii</i>	24.00 ± 0.03 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Phytophthora palmivora</i>	24.33 ± 0.30 ^c	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a
<i>Aspergillus flavus</i>	20.02 ± 0.31 ^b	9.33 ± 0.00 ^d	12.11 ± 0.31 ^b	14.89 ± 0.06 ^e
<i>Saccharomyces cerevisiae</i>	20.00 ± 0.30 ^b	11.16 ± 0.02 ^c	11.17 ± 0.06 ^c	12.12 ± 0.16 ^d
<i>Aspergillus tamari</i>	24.01 ± 0.11 ^c	9.04 ± 0.06 ^c	11.00 ± 0.02 ^b	14.04 ± 0.01 ^e
<i>Fusarium</i> sp.	19.09 ± 0.15 ^b	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a	0.00 ± 0.00 ^a

Table 10: Antifungal activity of abdominal gland extract of *O. longinoda*.

Values are presented as mean ± SE and values in the same column carrying same superscript are not significantly different using Duncan's new multiple range test at $p \leq 0.05$.

Antifungal activity of abdominal gland extract of *O. longinoda*. B, C and D, Increasing concentrations of abdominal gland extract. Concentration B is gland from three ants; concentration C is glands from six ants; concentration D, gland from twelve ants. A is, Ketoconazole std 100 µl (40 mg/ml).

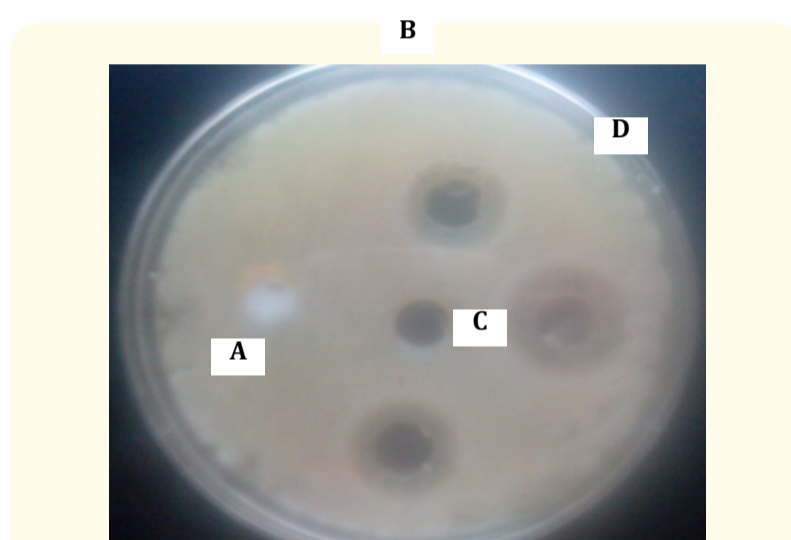


Plate 1: Antibacterial activity of *Oecophylla longinoda* abdominal gland extract against *Klebsiella oxytoca*.

Key: A: Ciprofloxacin; B: 3 Abdominal Gland Extract of *O. longinoda*; C: 6 Abdominal Gland Extract of *O. longinoda*; D: 12 Abdominal Gland Extract of *O. longinoda*

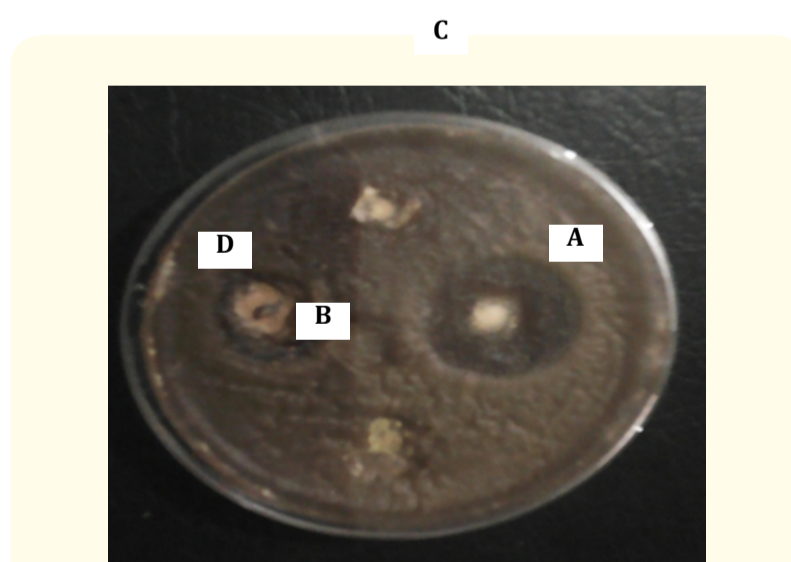


Plate 2: Antifungal activity of *Oecophylla longinoda* abdominal gland extract against *Aspergillus niger*.

Key: A: Ketoconazole; B: 3 Abdominal Gland Extract of *O. longinoda*; C: 6 Abdominal Gland Extract of *O. longinoda*; D: 12 Abdominal Gland Extract of *O. longinoda*

Discussion

These results revealed that microorganisms can be isolated from both infected and healthy Cocoa pods. The result showed that microorganisms (bacteria and fungi) isolated from Cocoa pods were more in infected pods than in healthy pods. The microorganisms isolated from the Cocoa pods is in agreement with the work of Adejumo and Adejoro [27] and Nasril [28].

Twenty-two (22) bacterial isolates were obtained of which *Proteus penneri* had the highest percentage occurrence of 27.3% while *Serratia* and *Citrobacter freundii* had the lowest percentage occurrence of 4.5%. 20 fungal isolates were obtained of which *Fusarium* sp. had the highest percentage occurrence of 30%, *Aspergillus aculeatinus*, *Penicillium chrysogenum*, *Cloridium chlamydosporis*, *Phytophthora palmivora*, *Aspergillus flavus*, *Aspergillus tamari*; *Alternaria* sp. had the lowest percentage occurrence of 5%.

All the tested bacterial isolates showed increase in susceptibility as the concentration of the extract increases this is in agreement with the work of Vidhu and Evans [26]; all the isolated bacteria were susceptible to the positive control (Ciprofloxacin) except *Klebsiella oxytoca* which was not susceptible to Ciprofloxacin.

Ketoconazole appeared to be the most effective antifungal agent in the control group. Of all the twelve fungal isolates tested with Ketoconazole, eleven (*Aspergillus aculeatinus*, *Alternaria* sp., *Cloridium chlamydosporis*, *Aspergillus niger*, *Phytophthora palmivora*, *Aspergillus flavus*, *Aspergillus tamari*, *Fusarium* sp., *Saccharomyces cerevisiae*, *Saccharomycodes ludwigii*, *Penicillium chrysogenum*) were susceptible while (*Penicillium camemberti*) was not susceptible.

All the fungal isolates showed increase in susceptibility as the concentration of the extract increases, this is in correspondence with the work of Vidhu and Evans [26] except for *Penicillium chrysogenum*, *Penicillium camemberti*, *Saccharomycodes ludwigii*, *Phytophthora palmivora* and *Fusarium* species which were not susceptible to all the concentrations; hence, in order to protect cocoa pods from infections, the population of *O. longinoda* must be very abundant, which was also suggested by Vanderplank [29].

Conclusion and Recommendation

The isolation of microorganisms from cocoa pods showed that both bacteria and fungi are present on both diseased and healthy pods and that the fact that microorganisms were isolated from both healthy and infected pods does not mean that they are both infected. On the other hand, the healthy pods might have been infected but have not started manifesting the symptoms of diseases. *Oecophylla longinoda* is a very good biocontrol agent as well as a potent antimicrobial agent which can be used by farmers in place of pesticides because it is relatively cheaper than pesticides and environmentally friendly as it is not harmful to human and the environment.

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