

**Original Article** 

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# **Microbiological, nutritional quality and antioxidant activity of fermented Delonix regia seeds**

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## **1. Introduction**

Fermented products via fermentation combined with other food processing techniques constituted integral part of diets in many homes around the world mostly in African and Asian countries. Different traditional food processing techniques such as cooking, smoking, sprouting/germination and less often toasting are commonly practiced in most of the African countries. However, fermented foods have gained popularities and well accepted by all in these countries due to lots of nutritional benefits accrued to them (1,2). They are produced by the use of microorganisms and hydrolytic enzymes with lead to desirable quality that are quite different from raw or unfermented organic

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substrates (3). Their beneficial effects from the nutraceutical point of view include immunemodulatory, antiallergenic, antihypertensive and antitumourigenic (4,5,6). Some metabolites released on them during fermentation by consortium of microorganisms involved in the process also improved their nutritional status and extended their shelf life (7). Their thiamine, nicotinic acid, riboflavin and perhaps, protein content are enhanced as a result of microbial activity. Other dietary benefits of fermented foods include improvement of food safety, minerals bioavailability and vitamins, toxins reduction and improved digestibility and industrial utility (Jimoh et al., 2012).

The principal key player in food fermentation process is lactic acid bacteria (LAB); they help in the production of antimicrobial metabolites such as

bacteriocins, reuterins, hydrogen peroxide, diacetyls and organic acids. The presences of these antimicrobial compounds in fermented foods either inhibit or kill unwanted food borne pathogens and spoilage microorganisms. A number of LAB genera to be precised *Enterococcus spp.*, *Leuconostoc spp.*, *Lactobacillus spp.*, *Pediococcus spp.*, *Lactococcus spp.* and *Streptococcus spp.* have been screened and confirmed to produce these inhibitory substances during the course of fermentation (8). Some LAB are known as good producers of antioxidants such as flavanoids, benzoquinones, alkaloids, terpenoides, xanthones, phenols, tetralones and steroids (9). Fermented foods enriched with antioxidants protect consumers' body against oxidative damage (9).

Dietary supplements that contain antioxidants help in disease prevention and maintenance of good health. Antioxidants prevent human body from diseases such as cancer and coronary heart disease. Carcinogenic and toxic nature of synthetic or commercially available antioxidants has led to search for natural antioxidants (10). Previous investigations have revealed that plant foods such as vegetables, fruits, spices, tea and wines are proven sources of biological antioxidants (11). Many plant parts such as flowers, root, seed, stem and leaves contain arrays of antioxidant compounds which are considered as safe and inexpensive source of antioxidant from natural origin (12). However, antioxidants within the plant tissues are either lockup or available in inactive forms. Thus, microbial activities and their enzymes are required to make these compounds free in utilizable forms for use in traditional and modern medicine (13).

In folk medicine, *D. regia*, an average tall flowering plant has been used effectively for the treatment of ailments such as rheumatism, hemiplagia, leucorrhoea, inflammation and constipation (14). Its therapeutic potential is attributed to the fact that it has antioxidant, antimicrobial, anti-diarrhoeal, anti-ulcer and antiinflammatory activity (15). In search of underutilized plant material with desirable nutritional and pharmacokinetic properties, *D. regia* seed was therefore considered a plant part of choice and hence subjected to submerged state fermentation with a view to unlock its phyto-constituents. This research assessed the microbiological, nutritional quality and antioxidant profile of fermented *D. regia* seeds.

# **2. Materials and methods**

2.1. *Delonix regia* seeds collection and preparation for microbiological and chemical analyses

Microorganisms associated with fermented *D. regia* seeds were isolated according to the method of Chelule et al. (16). Two hundred grams (200 g) milled seeds of *D. regia* was suspended in a clean transparent screw capped glass jar containing two liters of sterile distilled water. The mixture was incubated in static condition for three days at room temperature  $(28\pm2°C)$ , and at interval of 24 h, isolation of microorganisms were carried out using pour plating technique with suitable dilution factors and incubated at 30°C for 24 and 72 h for bacteria and fungi respectively. The resultant bacterial and fungal colonies were counted and expressed as CFU per millimeters and SFU/ millimeters respectively. Pure bacterial isolates were presumptively identified via morphological and some biochemical identification. Thereafter, Bergey's Manual of Determinative Bacteria was consulted as a standard reference to tentatively identify the bacterial isolates (17). Pure fungal isolates were identified on the basis of cultural characters as well as microscopic structure (18).

# 2.3. Proximate composition of fermented *D. regia* seeds

The method of AOAC (19) was adopted in the determination of the proximate composition of the fermented samples. The proximate parameters considered were moisture content, ash content, fat content, crude fiber and crude protein content, carbohydrates and moisture contents.

# 2.4. Total titratable acidity and pH determination

Total titratable acidity (TTA) and pH of the fermenting substrate were determined as described by Akharayi and Omoya (20). TTA was determined by diluting fermented slurry with sterile distilled water, and the mixture with the addition of (few drops) 1% phenolphthalein was titrated against 0.1 M NaOH. The pH was determined by dipping a calibrated pH meter (Crison Basic Model 20) into the fermented samples; the readings were taken and recorded.

# 2.5. Antioxidant profile of fermented *D. regia* seeds

The antioxidant parameters evaluated from the fermented *D. regia* seeds were total flavonoid, ferric reduction property (FRP), free radicals scavenging (FRS) ability and total antioxidant activity.

## 2.5.1. Total flavonoid determination

The colorimeter assay was used to determine the total flavonoid content in the fermented *D. regia* seeds

and the absorbance was read at 510 nm against the reagent blank. Total flavonoid content in the samples was expressed as  $mg/g(21)$ .

## 2.5.2. Determination of FRP

FRP of the fermented *D. regia* seeds was determined according to Pulido et al. (22). Fermented *D. regia* seeds (0.3 ml) was diluted with equal volume of 200 mM sodium phosphate buffer pH 6.6 and 0.25 ml of 1% KFC. The mixture in a test tube was incubated at 50°C for 20 min, followed by the addition of 0.25 ml of 10% TCA and centrifuged at 2000 rpm for 10 min. After centrifugation, 1.0 ml of the clear supernatant was diluted with 1 ml distilled water and  $0.1\%$  FeCl<sub>3</sub> and the reading was taken at 700 nm.

#### 2.5.3. Determination of FRS

The FRS of the fermented *D. regia* seeds was determined against DPPH (1,1-diphenyl-2 picryhydrazyl) as described by Gyamfi et al. (23). One ml from the fermentation medium was mixed with 1 ml of 0.4 mM methanolic solution of the DPPH, the mixture was left in the dark for 30 min and the absorbance was measured at 516 nm.

#### 2.5.4. Total antioxidant activity determination

The determination of total antioxidant activity (TAA) was based on the reduction of Mo (VI) - Mo (V) by the fermented *D. regia* seeds and the subsequent formation of a green phosphate/Mo (V) complex at acidic pH (24). TAA of the fermented samples was determined by adding different concentrations of samples with 3 ml of the reagent solution (0.6 M sulphuric acid, 28 mM sodium phosphate and 4 mM ammonium molybdate) followed by the incubation of the reaction mixture in water bath at 95°C for 1 h 30 min. The mixture was cooled to room temperature against blank and the absorbance was measured at 695 nm. The TAA was expressed as garlic acid equivalent.

## 2.6. Statistical analysis

 **Table 3.** Morphological and biochemical characterization of bacterial isolates from fermented *D. regia* seeds

Data obtained were analyzed by analysis of variance and significant differences between means were compared using Duncan multiple range test.

## **3. Results**

Table 1 shows the total bacterial and lactic acid bacterial (LAB) counts from fermented *D. regia*. Total bacterial counts from fermented *D. regia* increased from  $1.3\times10^9$  cfu/ml at 0 h to  $4.4\times10^9$  cfu/ml at 48 h, and beyond, a reduced total bacterial load was recorded. LAB counts from fermented *D. regia* recorded an appreciable increase all through the fermentation time.

**Table 1.** Total bacterial and lactic acid bacterial count from fermented *D. regia*

Fermentation time (hour)	<b>Bacterial count</b> $(x109$ cfu/ml)	Lactic acid bacterial count $(x 10^9 \text{cfu/ml})$		
24	21	09		
48	4.4	10.9		
	31	189		

Table 2 shows the total fungal counts from the fermented *D. regia*. The fungal loads increased on PDA and SDA all through the fermentation time.





**Key:** PDA: Potato dextrose agar, SDA: Sabraud dextrose agar

Table 3 shows the morphological and biochemical characterization of bacteria isolated from the fermented Delonix regia seeds. The bacterial strains were identified as *Bacillus subtilis, Staphyloccocus aureus*, *Serratia marcescens*, *Streptomyces greceus*, *Micrococcu luteus*, *Lactobacillus plantarum* and *Lactobacillus casei* respectively.



+ = Positive, - = Negative, AG = Acid and Gas production; Ag = Acid production without Gas; LR = Long rod; SR = Short rod; R = Rod; C = Cocci

<b>Isolates</b>	Cultural characterization and microscopic observation	Probable organisms
code		
A11	Powdery, yellow to green colony and the reverse was yellowish. Radiated conidia head with	Aspergillus flavus
	hyaline, long, rough walled conidiophore and globose to subglobose finely roughened to	
	echinulate conidia	
A12	Powdery blue-green colonies with pale yellow reverse. Septate hyphae with smooth walled	
	conidiophores, ending in vesicle and subglobose shape.	Aspergillus Fumigates
A13	Sparse mycelium, whitish with a purple reverse. Septate hyphae with conidia moderately	Fusarium species
	curved and pointed at both end	
A14	Grayish green with reverse pale yellow color.	Penicillium notatum
A15	Colony whitish and dark brown to black-brown.	Rhizopus japonicum
A16	Bluish green with reverse yellow brown. Septate hyphae with smooth walled conidiophore	Penicillium italicum
	with ellipsoida to cylindrical conidia	
A17	Creamy color, oval shape with branched cell.	Saccharomyces cerevisiae

**Table 4.** Cultural and microscopic identification of fungi isolated from the fermented *D. regia* seeds

In Table 4, the cultural characterization and microscopic observation of fungal isolates associated with fermented *D. regia* seeds is revealed. The fungi consisted of *Aspergillus flavus*, *A. fumigatus, Fusarium spp., Penicillium notatum, Penicillium italicum, Rhizopus japonicum* and *Saccharomyces cerevisiae*, respectively.

Table 5 shows the TTA and pH value of the fermented *D. regia* seeds. The total titratable acidity increased with corresponding pH decrease with increase in the fermentation time from 24 to 72 h. The total titratable acidity of the fermented sample increased from 3.3 at 24 h to 18.1 at 72 h, while the pH decreased from 6.8 to 4.1.

**Table 5.** Total titratable acidity (TTA) and pH value of fermented *D. regia* seeds

<b>Fermentation time</b> 'days)	<b>TTA</b>	рH
24	3.3	6.8
48	14.1	4.9
77	18 1	

Table 6 shows proximate composition of the raw and fermented *D. regia* seeds. The moisture, fat and protein contents of the fermented samples increased from approximately 11, 2 and 20% in raw samples to 18, 5 and 32%, respectively in the fermented samples. There was remarkable reduction in the fiber and carbohydrate content by approximately 79 and 16% in the fermented samples when compared with unfermented samples.

The antioxidant component of the raw and fermented *D. regia* seeds is shown in Table 7. The FRAP, TAC, flavonoid and DPPH in the unfermented samples increased from approximately 26 mg/g, 9 mg/g, 1 mg/g and 10% to 44 mg/g, 32 mg/g, 3 mg/g and 10% respectively in the fermented samples.

**Table 7.** Antioxidant profile of the raw and fermented *D. regia* seeds

Antioxidant properties	Unfermented	Fermented	
Ferric reducing antioxidant	$26.10^{\circ} \pm 1.73$	$43.61b \pm 0.05$	
property (FRAP) $(mg/g)$			
Total antioxidant capacity	$9.06^a \pm 0.20$	$32.34b \pm 0.35$	
(TAC)(mg/g)			
Flavonoid $(mg/g)$	$0.95a \pm 0.02$	$2.98 + 0.04$	
DPPH(%)	$10.21a \pm 2.61$	$46.29b \pm 0.25$	

Data are represented as mean ± standard deviation. Means with the same superscript across the column are not significantly different from each other  $(p<0.05)$ 

#### **4. Discussion**

Array of microorganisms were isolated during the fermentation of *D. regia* seeds. Isolation of different microorganisms from fermented plant materials has been reported (25). The establishment of co-metabolism or microbial synergy between fungi and bacteria during the fermentation of plant materials is well documented (26,27). Lactic acid bacteria (LAB) create acid environment for the proliferation of fermentative fungi, and in return, vitamins, utilizable simple sugars and other growth factors are produced by fungi (28). The occurrence of different bacterial and fungal isolates such as LAB, *S. aureus*, *S. marcescens*, *B. subtilis, S.*

**Table 6.** Proximate composition of raw and fermented *D. regia* seeds (%)

Sample	<b>Moisture Content</b>	Ash Content	<b>Fat Content</b>	<b>Fiber Content</b>	Protein ∩ntent	Carbohvdrate ∩ontent⊂
Raw	$11.10a \pm 0.02$	$2.53 \pm 0.11$	$1.66a \pm 0.21$	$16.13 \pm 0.01$	$19.99a \pm 0.01$	$48.59b \pm 0.01$
DF	16.57b±0.01	$2.02b \pm 0.12$	$3.84b \pm 0.01$	$11.34b \pm 0.04$	$25.09b \pm 0.03$	$41.14a \pm 0.03$
AF	18.25c±0.03	$0.65$ a $\pm 0.11$	$4.67 \pm 0.02$	$3.4a \pm 0.02$	$32.16 \pm 0.15$	$40.87a \pm 0.02$

Data are represented as mean ± standard deviation. Means with the same superscript down the row are not significantly different from each other (p>0.05).Key: DF: During fermentation, AF: After fermentation.

*greceus*, *A. flavus*, *A. fumigatus*, *Fusarium spp.*, *P. notatum* , *R. japonicum*, *P. italicum* and *S. cerevisiae* during the fermentation of *D. regia* seeds suggest that they might be classified either as natural microflora or contaminating microorganisms emanated from the substrates meant for fermentation, soil, utensil used or fermentation vessels, aerosols, rodents and the personnel involved (29,30). Duangjitcharoen et al. (31) and Gunawan et al. (32) isolated LAB from fermented plant beverages and *Xylocarpus moluccensis* seeds respectively. The presence of *A. flavus*, *A. fumigatus* and *Fusarium spp*. signal great health concern since these fungi are known for mycotoxins production. Some strains of *A. flavus* produced carcinogenic toxic chemical called aflatoxins, while certain strains of *A. fumigatus* and *Fusarium spp*. secrete toxic fumonisins and zearalenone, respectively (30).

The fermentation of *D. regia* seeds was characterized by a decrease in pH and corresponding rise in TTA. Increase in TTA and decrease in pH during the spontaneous fermentation of plant seeds has been reported (33). The decrease in pH with a corresponding increase in TTA during fermentation has been reported by some researchers who had done similar investigation. Ayo (34), Ojokoh et al. (27), Amankwah et al. (36) and Ojokoh et al. (27) reported a decrease in the pH and a rise in the TTA of the fermenting medium for millet flour, millet-acha based kunun zaki (beverage), breadfruit-cowpea blend flours, maize flour and pearl millet and acha flour blends, respectively. Decrease in pH might be due to the presence of organic acids such as lactic acid by LAB involved in the fermentation of carbohydrate contents of the fermenting substrates (27,29).

The increase in the protein content of fermented *D. regia* seeds in this study agrees with the reports of Sade (37), Chukwu and Abdul-kadir (38) and Ojokoh et al. (27). For instance, Sade (37) reported an increase in the protein content of raw millet from 14.0 to 17.5% in the fermented millet, while Chukwu and Abdul-kadir (38) reported an increase in the protein content from 6.9 to 10.6% in acha. Similarly, Ojokoh et al. (27) reported an increase in the protein content of fermented pearl millet with acha by approximately 50% when compared with unfermented sample. The increase in the crude protein contents of the fermented *D. regia* seeds might be due to the secretion of hydrolase enzymes rich in protein and excess nitrogen entrapment by the consortium of microorganisms involved (39,40). The crude fiber decreased consistently all through the fermentation period. Many researchers have reported similar reduction in the amount of crude fiber in fermented

plant seeds or cereals. In their studies, Sade (37) and Amankwah et al. (36) reported a reduction in the crude fiber content of fermented millet and maize flours respectively. Similar result was reported by Ojokoh et al. (27) when pearl millet with acha was subjected to fermentation. The decrease in the crude fiber content of the fermented *D. regia* seeds might be connected with the ability of LAB to produce crude fiber-degrading enzymes in the course of fermentation (27,40). There was significant increase in the moisture content of fermented *D. regia* seeds when compared with the unfermented substrate. The finding from this study was in agreement with the reports of some researcher who had carried out similar experiment. There was an increase in the moisture content of fermented maize flour from 14.2 to approximately 20% (41). In another study by Abiola and Ekunrin (42), moisture content in the raw melon husk increased from 2.46 to 5.84% after 72 h of fermentation. Also, Zakari et al. (43) reported that the moisture content of pearl millet-bambara groundnut blend flours increased from 12.0 to 12.9% after fermentation. The reduced moisture content in the fermented samples might be due to the utilization of moisture by the microorganisms for various metabolic activities and this reduction might contribute to the longer shelf life and prevent microbial spoilage of the end product (44).

In this study, there was significant increase in the antioxidant profile of fermented *D. regia* seeds. Hunaefi et al. (45) reported an increase in the total flavonoids and flavonols of *orthosiphon aristatus* (Java Tea) when series of LAB fermentation methods was exploited. The increase in the DPPH content of the fermented *D. regia* seeds agrees with the report of Oliveira et al. (46) who reported an increase in the DPPH of fermented rice (*Oryza sativa*) bran from 32 to 59% after 96 h of solid state fermentation. Fermentation of black soybean by *Bacillus subtilis* was found to increase its total phenolic and flavonoid contents as well as its antioxidant activities of anti-DPPH radicals and ferric reducing antioxidant power (FRAP) (47). The significant increase in the antioxidant components evaluated might be attributed to the involvement of LAB in the fermentation (45). LAB have the ability to produce exopolysaccharides that power bank for antioxidant components. These exopolysaccharides produced by LAB demonstrated potential in vitro antioxidant properties (1,1 diphenyl-2- picrylhydrozyl radical scavenging activities, chelation of ferrous ion, inhibition of linoleic acid peroxidation and reducing power) during an oral administration of their strain (45).

## **5. Conclusion**

In conclusion, the fermented *D. regia* seeds had higher nutritive value and better antioxidant activity than the unfermented samples. The antioxidant properties of *D. regia* seeds could enhance through food grade LAB fermentation. Hence, fermented *D. regia* seeds can be exploited as feed ingredient and used as substitute to essential ingredients known to be expensive in infant food nutrition.

## **Conflict of interest**

There is no conflict of interest.

# **Acknowledgements**

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