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Sylvester Owusu Boakye \*

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# Isolation, Screening and Identification of Pectinolytic Fungi from the Soil of Decomposed Plant Materials

Sylvester Owusu Boakye 1\*, Hilary Domakyaara Zakpaa 1, Malik Borigu 2 and Abudu Ballu Duwiejuah 3

<sup>1</sup>Department of Biochemistry and Biotechnology, College of Science, Faculty of Biosciences, Kwame Nkrumah University of Science and Technology, Kumasi-Ghana.

<sup>2</sup>Department of Crop and Soil Sciences, Kwame Nkrumah University of Science and Technology, Kumasi-Ghana.

<sup>3</sup>Department of Biotechnology and Molecular Biology, Faculty of Biosciences, University for Development Studies, Tamale-Ghana.

\*Corresponding Author: Sylvester Owusu Boakye. Department of Biochemistry and Biotechnology, College of Science, Faculty of Biosciences, Kwame Nkrumah University of Science and Technology, Kumasi-Ghana.

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## Abstract

Pectinases are essential enzymes and command a quarter of all food enzymes sold globally, and are recognised as excellent workhorses in plethora of industries. These enzymes are mainly sourced from animals, microorganisms and plants, with microbes serving as the main sources owing to the quality, quantity, ease of extraction and economic considerations. This research study isolated, screened, and identified five pectinase-producing fungi from the soil of decaying fruits and vegetables. The isolated fungal species were cultured and screened for highest pectinase production using pectinase screening agar medium containing 1% citrus pectin. Four isolates identified as Aspergillus niger, Fusarium species, Trichoderma species and Aspergillus flavus, exhibited very high values of pectinase hydrolysis based on clear zone method, which were 25, 23, 20, and 20 mm for the respective isolates. A secondary pectin hydrolysis screening exercise was conducted afterwards to obtain the highest level of reducing sugars and pectinase released by the isolates. Aspergillus niger, once again, proved to be the best as it recorded maximum reducing sugar of 3.92 mg/mL and pectinase enzyme activity of 36.23 U/mL. Soil have the potential to serve as the primary source of fungal species capable of degrading pectin to release pectinase enzymes, which have numerous applications in the production of pharmaceuticals, food, beverages, animal feed, textiles, detergents, protoplast fusion, pulp and paper and biofuels.

**Keywords:** pectinase; fungal species; Aspergillus niger; pectin; screening

#### 1.Introduction

Pectin is a hecteropolysaccharide that serves as an important component of the cell wall and other internal components of higher plants. It confers mechanical support and flexibility to plants by acting as a cementing material that holds the cell walls of adjacent cells together [1,2]. Pectin is made up of numerous units that consists of D-galacturonic acids with carboxyl groups mostly esterified with methanol, with the main chain being made up of  $\alpha$ -(1, 4) glycosidic bonds. Pectin with its methyl groups removed is called pectate or pectic acid, which is also called polygalacturonic acid [3]. The structure of pectin contains homogalacturonan which appears as linear biopolymer and is made up of methyl esterified or acetylated D-galacturonic acid residues or monomers containing  $\alpha$ -(1, 4) glycosidic linkages. Pectin is formed when about 75% of carboxylic acids found in homogalacturonans become

methylated. Pectic acid is also formed when less than 75% of the carboxylic acids are methylated. Polygalacturonic acid results when there is no methyl esterification [4]. Also, protopectin represents the pectic substances that are not soluble in water [5]. Soluble pectic substances can be formed after limited hydrolysis of protopectin [6]. Pectinic substances, which act as the methylated form of galacturonic acid are formed when there is enzymatic hydrolysis of protopectin in ripe fruits [7].

Pectin is non-toxic, biocompatible and biodegradable and can dissolve in water. It has crucial stabilizing, hydration, gelling, thickening and swelling properties, which makes it very useful in many commercial applications. A few of the commercial applications include food designs, production of fruits juices, drinks, and fermented, confectionary pharmaceutical and health products. Pectin is also utilized as nano-encapsulator, drug delivery agent,

constituent of pro-health formulations, dietary fibre, anti-cancer agent, and hydrogel formulation material. It is also used to lower blood cholesterol and low-density lipoprotein levels as well as being used in the treatment of diarrhoea and constipation [8–10]. The primary sources of pectin have been citrus fruits, apple pomace, mango peels and sugar beet pulp [11–13].

Pectin can be hydrolyzed by a group of enzymes called pectinases by splitting the glycosidic bonds existing between pectin molecules [14]. Pectinases are able to break down pectin oligo-D- galacturonate present in plant cell walls into monogalacturonic acids. The hydrolysis is achieved when pectinolytic enzymes cleave the α-1, 4 glycosidic linkages of polygalacturonic acid [15]. Pectinases can be classified as extracellular and intracellular. Extracellular pectinase enzymes represent those secreted outside of the cell into the surrounding media in which the microbes live, whereas intracellular pectinase enzymes are those secreted and act within the cell membrane [16]. Nevertheless, extracellular and intracellular pectinases are also classified into three broader groups based on their mode of action on pectic substances [16,17], which include Protopectinases, Esterases, and Depolymerases [18-21]. Pectin esterases are able to catalyze the deesterification of methyl groups in pectin to form pectic acids. Pectin depolymerases, depending on their mechanism of action, are able to cleave the glycosidic bonds found in pectic acids. Also, protopectinases are able to break down protopectin [22]. In terms of their nature, pectinases can be categorized as acidic and alkaline. Acidic pectinases usually enjoy their applications in wine, beverage and fruit juice industries for clarification and purification of fruit juices and wines [13,23]. Alkaline pectinases on the other hand, find their applications in textile industries and wastewater treatments, [4,24,25].

Among the various pectinases, extracellular pectinases derived from fungal species have been known to possess excellent abilities compared to pectinases from other sources such as plants, animals and bacterial [26] and their method of extraction is very simple as the integrity of the cells remains intact. Other added advantages include high yields, economic feasibility, stability and easy genetic manipulation. Also, fungal-derived pectinolytic enzymes have generally regarded as safe (GRAS) status [22]. For these and many more reasons, extracellular pectinases isolated from fungal species especially that of Aspergillus spp. are in high demand in various applications [27,28]. The applications extend across food/feed, beverage, pulp and paper, biofuel, textile, oil extraction and waste and environmental remediation [13,21,29,30]. Pectinases employed in fruit and wine industries can enhance taste and clarity and overall chromatic appearance compared to untreated ones [24].

The soil, as part of the natural environment, serves as a huge reservoir for fungal species that can be isolated and employed in the production of pectinase enzymes. Microorganisms isolated from the soil are mesophilic, can tolerate stress and have the ability to synthesize pectinase enzymes in large quantities in wider spectrum of growth parameters [31]. Soils particularly found in plant food processing locations, markets and organic waste dumpsites have been proven to be very much dominant with pectin-degrading fungus [5]. Recent advances indicate that the trend to identify novel enzymes from microorganisms from such environments is accelerating due to their inherent stability and efficiency [32,33]. Hence, pectinolytic microbes were obtained from the soils exposed to high and low temperatures, and subsequent identification was carried out in this current investigation.

### 2. Materials and Methods

## 2.1. Collection of Soil Sample

Soil composed of plant-decayed materials were collected from the dumpsites of fruits, vegetables and plant materials in some markets in Kumasi as these places contained a lot of spoiled fruits and vegetable wastes and decaying plant materials. All the samples were pooled together and kept in sterile plastic containers and carried to the microbiology laboratories of Kwame

Nkrumah University of Science and Technology where fungal species were isolated from them.

#### 2.2. Preparation of Medium for Isolation of Fungal Species

A sterile nutrient medium containing 1% citrus pectin; FeSO4.7H2O (5 g/L), MgSO4.7H2O (0.02%), (NH4)2SO4 (0.14%), K2HPO4 (0.2%) as well as nutrient solution (0.1%) containing: ZnSO4.7H2O (1.4 mg/L), MnSO4.H2O (1.6 mg/L), CoCl2 (2.0 mg/L) was prepared, and mixed with 3% agar-agar (the gelling agent) and autoclaved at 121oC for 20 minutes [20]. It was then allowed to cool to around 45°C before being placed into Petri dishes to gel. To ensure sterility, the plates were incubated overnight in an incubator at 37oC.

## 2.3. Preparation of Potato Dextrose Agar for Isolation of Fungal Species

Potato Dextrose Agar (PDA) was made from a commercial dehydrated OXOID product. Thirty- nine PDA powder (39.0 g) was dissolved in 1 L of distilled water, and sterilized at 121oC for 20 minutes and at pressure of 1.2 g/cm2 according to the product recommendations.

## 2.4. Isolation and Sub-Culturing of Fungal Species

The simple dilution plate method was adopted here. One (1) g of soil sample was diluted ten- fold in 9 mL of sterile distilled water. One (1) mL of the 6th dilution (10-6) was flooded onto three replicates of the previously prepared sterile Potato Dextrose Agar (PDA) nutrient medium, which had been supplemented with chloramphenicol antibiotic to inhibit bacterial growth. The plates were then incubated at room temperature (25oC - 28oC) for 7 days, until visible fungal colonies appeared. All morphologically distinct colonies were purified by sub-culturing on different plates and streaking with a sterile inoculation loop. This process proceeded until pure fungal isolates were obtained. Pure cultures were maintained on PDA slants, refrigerated and sub-cultured at regular intervals throughout the study and were used as stock cultures. All pure fungal growths were counted with a colony counter, and identified appropriately. Their frequency of occurrence were subsequently established as well.

#### 2.5. Identification of Fungal Isolates

The identification procedure for all the fungal isolates was based on macromorphological and micromorphological characteristics. Macromorphological identification was done by observing colony characteristics such as colour, texture, and spore structure, the form of mycelia and/or pattern of growth. The micromorphological characteristics like separation, spore shapes, among others were determined using the usual lactophenol cotton blue procedure (LPCB). All the fungal isolates had been defined in accordance with the manual for determining the identity of fungi [34] and Illustrated Genera of Imperfect Fungi by Barnett and Hunter, 1972 [35]. In summary, a three-day-old pure culture was utilized to prepare slides for microscopic examination. A small amount of mycelia was placed on the slide, along with a drop of lactophenol cotton blue reagent. A cover slip was laid over it, and the specimen was examined under a light microscope at X400 magnification. Identification of the species was accomplished by comparing features to the micrographs, and the photographs taken included at the end of this section.

## 2.6. Qualitative (Primary) Screening for Pectinolytic Fungal Species

The process of screening for pectinase synthesis was carried out with the use of the prepared cultivation medium (in the previous section) by applying the spot inoculation method. In brief, pure cultures of the various fungal species were introduced into the already prepared sterile and solidified pectin agar medium. The plates that were inoculated then incubated at a temperature of 30°C for four days (96 hours), and following that, clear observable colony of fungi could be seen on every petri dish. Thereafter, the petri dishes were soaked with Iodine-Potassium iodide solutions (a mixture of 1.0g of iodine

and 5.0g of potassium iodide dissolved in 330 mL distilled water) for about 10 minutes in order to boost visibility for measuring the zone of clearance / carbohydrate hydrolysis [20]. A distinct zone formed surrounding the pectinase-producing colonies was observed and subsequently measured. The fungal isolates with the most promising clear zone of pectinase hydrolysis were chosen and preserved under 4oC in the refrigerator on pectin agar and later used in the next step.

#### 2.7 Quantitative (Secondary) Screening for Pectinolytic Fungal Species

This experiment utilized the protocol established by Kaur and Kaur [36] with some modifications. The four isolates with higher degrees of pectinolytic activity showing larger zones of pectin hydrolysis obtained in the previous experiment were used in a submerged fermentation process. 50 mL of the fermentation medium (without the 3% agar agar) prepared earlier underwent inoculation with 1mL of spore suspension of each of the culture isolates. It was made to incubate at 30°C for 7 days on a rotary shaker at 120 revolutions per minute. Throughout the experiment, triplicate Erlenmeyer flasks (250mL) were used for each species of fungus. Once the incubation period elapsed, the crude pectinolytic enzyme underwent extraction. The extraction was done by mixing 50mLof sodium acetate buffer with the fermented broth cultures. The mixture was centrifuged for 15 minutes at 4000 revolutions per minute to remove the debris and cells. The supernatants were labeled as crude enzymes. The pectinolytic activities and reducing sugars of the enzyme were measured by estimating the reducing sugars generated from pectin using the 3, 5-dinitrosalicylic acid (DNS) technique. Glucose was employed as the benchmark. Each sample's pectinase activity was defined as the quantity of enzyme required to produce 1 µmol reducing groups/minute.

#### 3. Results and Discussion

#### 3.1. Isolation and Sub-Culturing of Fungal Species

Table 1 presents the results of identified species of fungus from the soil samples sourced from the various locations. In all, 5 fungal isolates were obtained from the pooled soil samples. The microbes were designated as FS1, FS2, FS3, FS4 and FS5 based on the petri dishes they were cultured on. This result is clearly in line with recent developments which have involved screening a significant number of microbial species from diverse soil resources, agrowastes and environments. Such developments are aimed at finding microbial species capable of degrading polysaccharides present in plant biomass that would illicit synthesis of pectinases [5,37–39]. These microbial creatures play crucial roles in the maintenance of nutrient cycles, biodiversity and degradation of organic matter. Microbes have also been shown to be very useful for industrial processes. The fact that, all the 5

isolates being fungal species perfectly agrees with Meyer et al., 2018. According to them, fungal species are the most dominant microbial consortia present in the environment. The dominance is as a result of their robustness in their lifecycle and activities [31].

The wide range of fungal species in the natural environment, particularly in plant food processing locations, markets and organic waste dumpsites, is also in strong alignment with Thakur et al., 2021 [5]. It is of no doubt that, soil serves as a reservoir for plant-degrading microorganisms that can be employed to carry out many processes. Previous investigations established by Ezenwelu et al., 2022 [41] demonstrated the existence of Aspergillus niger as well as Rhizopus spp. within the environments composed of decaying plant materials. Ametefe et al., 2021 [42], recoded Aspergillus niger and Penicillium spp as part of the fungi they isolated from the soil. Aside the soil, decaying fruits have also been implicated to harbour many microbial consortia especially fungal species. For instance, Okonji et al., 2019 [31] isolated Aspergillus fumigatus from decaying citrus. Mat Jalil et al., 2021[43] also isolated Aspergillus niger (LFP-1 strain) from decomposed citrus fruits. Budu et al., 2023 [39], showed the abundance of fungal spp. in decaying cassava peels. Plant-decaying soils (especially that of fruits and vegetables) are attractive to the growth and survival of fungi. This is because, they serve as a rich source of nutrients and the soil creates conducive environment that is highly favourable to fungal growth and survival [40].

As a result, such soils have the potential to draw fungi to colonize and flourish on them. Secondly, the physical structure of fruits and vegetables, which is usually characterized by their fibrous and porous nature provide a generous surface area and soil environment for fungal spores to land and establish their growth and thrive successfully. The physical structure in addition to texture, pH, oxygen content, cation exchange capacity and redox reactions of the soil provide synergistic conditions for microbe wellbeing. As such, their presence on organic matter in the soil, as a natural consequence, underscores the roles fungi play as the agents involved in breaking down and recycling of organic components and nutrients.

#### 3.2. Identification of Fungal Isolates

Fungal isolates FS1, FS2, FS3, FS4 and FS5 were thoroughly identified in accordance with their macromorphological and micromorphological features [34,35]. At the end of this section, Aspergillus flavus, Fusarium spp., Aspergillus niger, Trichoderma spp. and saccharomyces spp. were realized. Table 1 below demonstrates a summary of the various fungal species successfully obtained and identified and their characteristics.

Isolate	Characteristics (Macro-and Micromorphological)	Possible Identity
FS <sub>1</sub>	Appeared white initially and later turned grey or yellowish brown whilstpale yellow appeared as the reverse; colonies were fast growing with the texture appearing velvety or powdery, with a cottony appearance; conidial heads appeared as compact clusters, typically yellowish-green to yellow in color; large spore-bearing heads, spherical, and densely packed; clear, thin, and short hyphae; greenish mycelia; branched greenish conidia and septate hyphae; dense unbranched conidiophores with rounded or flask- shaped vesicle; elongated Phialides with constricted neck.	Aspergillus flavus
FS <sub>2</sub>	Colony had fast and aggressive growthand looked fluffy white initially, turned black later; generated enormous black spores; reverse appeared pale yellow; wrinkled mycelia with visible dense and lengthy aerial hyphae; conidia were unbranched; texture looked velvety orpowdery and the conidial produced heads with compact clusters, typically dark green to black in colour; hyphae looked septate, dense and branched; long, unbranched, and terminate conidiophores and gave rise to the	Aspergillus niger.

## 3.3. Qualitative (primary) screening for pectinolytic fungal species

One of the numerous ways to show pectinase production ability of microorganisms especially fungal species is the method popularly known as pectin/carbohydrate hydrolysis or zone of clearance. Such procedures have been known to be very effective in determining and selecting the best fungal species needed to carry out pectinase production [19,31,37]. By adopting similar paths, this current study conducted a protocol showing carbohydrate hydrolysis. The aim was to select the isolates with the highest pectin

degradation potential, after 5 days of incubation. After subjecting the isolates to pectin hydrolysis, the result showed that, all the fungal species possessed various degrees of pectinolytic activity (Table 2 and Figure 1). Aspergillus niger (FS2) emerged slightly on top as the most efficient microbial species (25.0 mm) among the tested species. Fusarium species (FS4) followed second with (23.0 mm), then Trichoderma species (FS3) and Aspergillus flavus (FS1) produced (20.0 mm each). Saccharomyces species (FS5) gave 7.0 mm, hence it was left out in subsequent experiment.

Microorganism	Zone of Pectinolytic activity/mm
Aspergillus niger	25.00
Fusarium spp.	23.00
Trichoderma spp.	20.00
Aspergillus flavus	20.00
Saccharomyces spp.	7.00

Table 2. Qualitative analysis of fungal species based on zone of pectin hydrolysis achieved.



A: Fusarium species

B: Aspergillus niger

C: Aspergillus flavus



D: Trichoderma species

E: Saccharomyces species

Figure 1. Qualitative analysis of fungal species based on zone of pectin hydrolysis. (A) Fusarium spp.;

(B) Aspergillus niger; (C) Aspergillus flavus; (D) Trichoderma spp.; (E) Saccharomyces spp.

It is well noted that, the strains of Aspergillus spp. have been known to dominate pectinase enzyme manufacturing in various studies and industrial applications [31]. The fact that, fungal species dominated the synthesis of pectinase in terms of zone of clearance in current study is favourably supported by El-Ghomary et al., 2021 [38]. In their work, they opined that, majority of commercial pectinase enzymes employed in industries are generated from species of fungal sources including Rhizopus stolonifer, Aspergillus and Penicillium spp. Olivad and Dawit, 2018 [27] and Adedayo et al., 2021 [37], also asserted that, commercial pectinase enzyme production has recognized Aspergillus and Penicillium spp. as the main sources of pectinase enzymes. El-Ghomary et al., 2021 [38] isolated four strains of Aspergillus spp. that showed clear zone of pectin hydrolysis of 3.62 mm, 3.70 mm, 3.91 mm, and 4.75 mm for the respective strains, with Aspergillus flavus emerging as the species with highest zone of clearance. Current result is also, in agreement with Okonji et al., 2019 [31] when fungal isolate identified as Aspergillus fumigatus showed the highest pectin hydrolysis. Other numerous earlier investigations have also indicated the vital role played by Aspergillus spp. [13,19,29,43].

Furthermore, the relative dominance of Aspergillus niger, in current investigation, after exhibiting highest activity of pectinase in relation to zone of clearance (25 mm) is no fluke at all. This assertion is true as its ability to hydrolyze pectin-rich media has also been agreeably demonstrated in series of previous studies cited in Satapathy et al., 2019 [20]. In another closely

related study conducted by Adedayo et al. (2021) [37], Aspergillus niger showed a better zone of clearance of 7.0 mm whereas Aspergillus flavus produced 5.5 mm. In addition, Kamalambigeswari et al., 2018 [45], examined three distinct strains of Aspergillus niger (F-3, F-4, and F-P), which revealed isolate F-4 as one that exhibited the largest region of pectin breakdown (9 mm).

## 3.4. Quantitative screening of potent pectinolytic fungal species (secondary screening)

Due to their promising nature, all the four isolates with comparatively higher zones of pectin hydrolysis were selected for further work on pectinase production (secondary screening). The four isolates that were deemed the best at pectinolytic ability exhibiting a clear zone of clearance were scrutinized for quantitative evaluation of pectinase synthesis utilizing a selective enrichment technique to generate comparative assessments. The secondary pectin hydrolysis screening exercise was then performed to identify the greatest amount of reducing sugars generated by the isolates with their respective enzyme activity via enzymatic breakdown of pectin. The galacturonic acid released was measured spectrophotometrically. The result (Table 3) showed that among the four (4) fungal isolates, FS2, which was identified as Aspergillus niger, once again, emerged as the most potent fungal species for the production of pectinase after recording the highest value reducing sugar formation (3.92 mg/mL) and pectinase activity (36.23 U/mL) under submerged fermentation after seven (7) days of incubation.

Microorganism	Reducing sugar (mg/mL)	Enzyme activity (U/mL)
Aspergillus niger (FS <sub>2)</sub>	3.92	36.23
Aspergillus flavus (FS <sub>1</sub> )	3.68	32.52
Trichoderma spp. (FS <sub>3</sub> )	3.23	29.74
Fusarium spp. (FS <sub>4</sub> )	3.16	28.66

P-value = 0.0260.

Table 3. shows the figures obtained during the secondary screening.

This finding fits in with El-Ghomary et al., 2021 [38], who found that, fungal isolates tagged as F9 showed relatively the highest titre of pectinase (35.83 U/mL) than that of all the other isolates. Okonji et al. (2019) [31] also discovered that Aspergillus fumigatus yielded 42 U/mL of pectinase enzyme activity in submerged fermentation conditions. The results achieved in this study and

subsequent selection of fungal strains is also in line with Sudeep et al., 2020, who in their research, isolated 4 strains of fungal species with high pectinolytic activities from the soil. After thorough primary and secondary screening procedure, Aspergillus spp. was observed as the most potent fungus for pectinase enzyme production.

#### 4.Conclusions

Soils obtained from the selected markets in Kumasi were found to be great source of plethora of diverse pectin-degrading microorganisms especially fungal species. A. niger has proven to be very dominant probably as a result of its ubiquitous nature and its ability to colonize diverse environments and substrate as it thrives well in many growth media or substrates as proven by this and other studies. A successful collection of these industrially-relevant strains of microorganisms from the soil could very well contribute to the increasing demand and adequate supply of pectinase enzymes in commercial quantities by related industries. Pectinase enzymes from microbes would also be very useful in remediation of areas polluted with plant waste. Further researches should be directed towards the mechanisms involved in the molecular regulation of pectinase enzymes production among fungal species especially A. niger. With such knowledge, scientists would be able to optimize the production of pectinase for commercial purposes. Also, subsequent studies could be invested in screening of extreme environments with the aim of finding fungal species with enhanced capabilities that can synthesize pectinase with novel properties.

**Author Contributions:** Conceptualization and design of the study were done by all authors. Material preparation, data collection and analysis as well as the first draft of the manuscript were performed by Owusu Sylvester Boakye, Zakpaa Hilary Domakyaara reviewed, edited and provided valuable comments and guidance whilst Malik Borigu supervised the laboratory experiments. Abudu Ballu Duwiejuah reviewed and provided valuable comments. All authors read and approved the final manuscript.

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