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Capability of *Trichoderma viride* to Produce Cellulolytic and Pectolytic Enzymes

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Abstract

Background: Species of the genus *Trichoderma* have been used in the food and textile industries to produce cellulases and other enzymes that degrade complex polysaccharide structures. *Trichoderma* species have been utilized to make cellulases and other enzymes that break down intricate polysaccharide structures in the food and textile industries. The study investigated the nutritional requirements and the production of enzymes by the fungus *Trichoderma viride* fungus' ability to produce enzymes).

Methods: We used a medium supplemented with cellulolytic and pectic substances for enzyme production.

Results: The results of the study proved that. Methionine, Glutamic acid, and leucine effectively enhanced mycelial growth. Findings regarding impact of pH level on the development of the fungus *T. viride* indicated that the maximum growth was at pH 5.0. However, growth decreased dramatically with increasing pH values. We examined the power of the fungus *T. viride* to produce cellulolytic and pectolytic enzymes in various substrates in the current study.

Conclusion: The findings demonstrated that Pectin was the best substrate for pectolytic enzyme synthesis, whereas Carboxymethyl cellulose (CMC) was the best substrate for the cellulolytic enzyme.

Introduction

Trichoderma filamentous fungi of the rhizosphere constitute a significant component of diverse soil ecosystem microbiomes and are distinguished by their capacity to penetrate plant roots. Trichoderma's employment in agriculture can be effectively ensured by thoroughly understanding its characteristics, including metabolic activity and the nature of interactions with plants and other microorganisms [1]. *T. harzianum* and *T. viride* are the two most significant species in the genus. The genus species are adaptable fungi used in various commercial and industrial applications. In addition to many bioactive metabolites against NTCB, fungi also produce colors, antioxidants, polysaccharides, and enzymes that may be investigated as natural alternatives to synthetic food additives. Through better fermentation conditions, metabolic engineering techniques, and fungi genetic manipulation, research in this area has enhanced the yields of metabolites for industrial purposes [2].

Trichoderma is frequently found as an indoor pollutant and has been shown to make up a sizeable portion of the fungal biomass in soil [3, 4, 5]. Trichoderma is frequently found in soil and woody materials. Due to its potent disintegrating function as spoilage organisms and for use in various plans for degrading cellulose waste, it has recently attracted significant economic interest [6].

Fungi produce cellulases and other enzymes that break down complex polysaccharide structures, and they have been exploited in the food and textile industries. Many extracellular enzymes are produced by them quite effectively. They are employed commercially in synthesizing enzymes like cellulases that can digest tough polysaccharides. These applications are typical in the businesses related to both food and textiles. The enzymes are also used in poultry feed to increase the digestion of hemicelluloses from barley or other crops [7].

According to Wang et al. [8], During their aggressive action against plant pathogens, Trichoderma species exhibit five mechanisms: antibiosis via secondary metabolite production; myco-parasitism via hydrolytic enzyme secretion; competition for space and nutrients; promotion of plant growth; and induction of systemic resistance in the host plant.

It is possible to add additional enzymatic capabilities and expand the applications of a selective consortium of fungal strains to deconstruct biomass into sugar [9, 10]. Co-cultivating diverse species of fungi enabled the production of enzymatic extracts abundant in different kinds of enzymes [11, 12]. In their investigation, Romero-Peláez et al. [13] shown that co-cultures of the ascomycete *T. reesei* RUT-C30 and the basidiomycete *P. lecontei* resulted in enzymatic extracts high in

cellulases, ligninases, and hemicellulases, allowing for an increase in the hydrolysis yield of pretreated sugarcane bagasse.

Pectin depolymerization is frequently related to the ripening process of fruit. A particular group of pectic enzymes breaks down pectic enzymes produced by many plant pathogens. Due to their importance, these enzymes affect how fruits and vegetables are stored after harvest [14]. Numerous studies have been conducted on these enzymes.

Although many fungi can live on cellulose or create enzymes that break down amorphous cellulose, it should be stressed that only a few fungi can fully develop extracellular cellulase systems that can break down crystalline cellulose [15]. The most well-known species are *Trichoderma* and *Phanerochaete* [16].

Cellulosic enzyme production has been attributed to a variety of microorganisms. *Trichoderma viride*, *Verticillium albo-atrum*, *Fusarium oxysporum*, *F. solani*, and *Stemphylium botryosum* are among them [17]. *Trichoderma* spp. were thought to be essential contributors to the lyses of *Rhizoctonia solani* cells by producing cellulase and chitinase enzymes [18].

Cellulose, in its native form, is insoluble and partly crystalline. Cellulose conversion to glucose needs a complex of enzymes, initially designated as; C1, Cx, and cellobiase [19]. The Cx (endo B- 1,4 glucanase) or the carboxymethyl cellulase (CMCase) splits soluble cellulose (carboxy methyl cellulose) to produce glucose from cellobiose.

The pectolytic enzymes include three main groups of enzymes; the esterases (Pectinmethylesterases PME), the polygalacturonases (P.G.), and the lyases (pectin methyl trans-eliminates PTE). The pectin methyl esterases (PME) from bacteria (*Clostridium multi-fermentans*), and fungi (*Botrytis cinerea*, *Corticium rolfisii*, *Fusarium oxysporum*, and *Phytophthora infestans*) as was reported by Forster [20]. The lyases (pectin methyl trans-eliminates PTE) break the α -1,4 linkage through a trans-eliminate cleavage, accompanied by a simultaneous elimination of H from C-5. Microorganisms in natural settings are responsible for regulating enzyme production to ensure the best effective utilization of the carbon supply that is present [21]. The ultimate objective of the industry is to discover and make use of naturally occurring microorganisms that are capable of manufacturing the desired enzymes in large yields and with a high level of productivity. The purpose of this study was to evaluate several elements of the capacity of the fungus *T. viride* to produce the virus.

Methods

The isolates of the fungus *T. viride* used in the present study were obtained from the Plant Pathology

Laboratory, Agriculture Research Corporation, Wad Medani, Sudan. Five different carbohydrates (glucose, sucrose fructose, galactose, and mannose) and five amino acids (glutamic acid, aniline, Arginine, leucine, and methionine) were obtained from the Food Science and Technology Department Lab. The fungus was grown on PDA slants and stored at -200 C before being.

Media used: Potato Dextrose Agar (PDA)

This study used Oxoid compounds to make PDA media locally. Fungal growth was sustained on this media. After boiling 200 grams of peeled and diced potato in one liter of distilled water for an hour, the extract was filtered and made up to 1 liter. Steaming and guiding dissolved D-glucose and agar. Then 100 ml samples of the medium were dispensed in conical flasks with cotton plugs and aluminum foil and sterilized in the autoclave at 121°C (15 lb/in²) for 15 minutes. It was refrigerated at 40°C. When needed, the medium was melted in a water bath and put into sanitized Petri dishes. For cellulase and pectic enzyme synthesis, the fungus was cultivated on basal salt medium with the following.

K₂HPO₄ (1.0 g)
KH₂PO₄ (0.5 g)
(NH₄)₂SO₄ (1.0 g)
Substrate (10.0 g)
Distilled water 1000ml

Three 5.0 mm fungal discs from an actively growing fungus were injected into the medium after 15 minutes of autoclaving at 121°C, cooling, and incubating on a rotary incubator at room temperature. The following substrates produced the cellulase enzyme; Carboxymethylcellulose (CMC), cotton fibers, filter paper, and glucose. Nevertheless, for the production of the pectic enzyme, the following were used; Pectin, glucose, and orange peels.

Effects of amino acids on fungal radial growth

PDA was utilized for fungal radial growth [22]. Substituting medium D-glucose yielded five amino acids at varied amounts. Sterilized media was placed into Petri dishes and solidified at room temperature (28-30°C). A fungal growth disc from an actively developing *T. viride* culture on PDA was injected into each cemented Petri plate. Petri dishes were cultured at room temperature for eight days after inoculation. All treatments were tripled. Each disc in a Petri dish was averaged two crossed dimensions to measure growth diameter every 48 hours. Radial growth was calculated as a percentage of Petri dish diameter.

Effects of the amino acids on mycelial dry weight

PDB was made and distributed in 100 ml conical flasks (45 ml each). For each amino acid, 0.0, 25.0, 50.0, 75.0, and 100.0 concentrations were synthesized. Each

solution was sterilized in an autoclave at 121°C (15 lb/in²) for 15 minutes and cooled to room temperature. A sterile cork-borer (5.0 mm diameter) formed three discs from an edge of an actively developing *T. viride* culture on a solidified PDA medium to inoculate each flask. For 12 days, inoculated plates were incubated at 28-30°C. Following incubation, the culture was filtered through Wittman's No. 1 filter papers and dried at 800 C for 24 hours before weighing. All treatments were tripled [23].

Effects of different pH levels on mycelial growth

Abubakar et al. [23] examined how pH affects mycelial growth. PDB medium was made and administered in 250 ml flask. Three 5.0 mm discs from an edge of an actively developing *T. viride* culture on solidified PDA media were injected into each flask. Samples pH was adjusted to be ranging from 4 – 9 pH (4, 5, and 6 were made by adding a concentrated HCl acid, while the pH values of 7, 8, and 9 were made by using NaOH solution). Three treatments were used for each pH level; the solutions were autoclaved at 121°C and cooled at room temperature (28 – 300°C). The flasks that were inoculated were placed in an incubator set at room temperature for a duration of 8 days. Following the incubation period, the cultures were subjected to filtration using a Whatman filter paper. The resulting mass was measured as the fresh weight (wt), after which it was subjected to a drying process at a temperature of 800°C for a duration of 24 hours. Subsequently, the dried mass was measured again as the dry weight (wt).

Production of cellulase and pectic enzymes

The fungus was grown on a basal salt medium containing the following constituents for cellulase and pectic enzyme production (prepared according to Nwodo-Chinedu et al [24]). The medium was autoclaved at 1210 C for 15 minutes, cooled, and inoculated by three fungal discs (5.0 mm diameter), cut from an actively growing fungus, and incubated on a rotary incubator at room temperature. The following substrates produced the cellulase enzyme; Carboxymethylcellulose (CMC), cotton fibers, filter paper, and glucose. However, Pectin, glucose, and orange peels were used for pectic enzyme production [25].

Enzyme assay: viscometrical method

Both the cellulase (Cx) and pectinase enzymes were assayed by the substrate viscosity reduction method using Viscometers (Cannon Fenke type) [26]. The viscometers were suspended in a water bath at 300 C. The reaction mixtures contained each of the followings:

Substrate (CMC or Pectin) 8.0 ml
 Water 1.0 ml
 CaCl₂ 0.01 ml
 Enzyme 1.0 ml

Before any viscometer was used, it was checked against a standard of water. A 100% reduction in viscosity was defined as the time it took to blow 10 ml of water.

Results

Effect of amino acids on radial growth

The effects of the different concentrations of the amino acids on the radial growth of the fungus were compared. The results showed that methionine was the most effective and gave significantly better results than all of the amino acids tested (Tables 1-3). It gave 100% radial growth at its higher concentration and the growth was also increased at its lower concentrations. Among the different amino acids tested, only Arginine was less effective in the radial growth of the fungus. Methionine and leucine were also significantly better, giving 99.4 and 99.45% radial growth at 50 mg/ml concentrations. The glutamic acid has seemlier significant to the leucine.

Conc (mg/ml)	% Radial growth					
	Incubation period days					
	2	4	2	8	2	x-coeff.
0	51.75	0	51.75	0	51.75	0
25	56.4	25	56.4	25	56.4	25
50	63.4	50	63.4	50	63.4	50
75	54.05	75	54.05	75	54.05	75
100	57.6	100	57.6	100	57.6	100
R ²	0.11	0.51	0.60	0.76	-	-
x-coeff.	0.33	0.71	0.77	0.87	-	-

Table 1: Effect of different concentrations of methionine on radial growth of *T. viride*.

Conc mg/ml	% Radial growth					
	Incubation period days					
	2	4	6	8	R ²	x-coeff.
0	60	73.9	83.9	85	0.89	0.94
25	48.05	54.2	62.55	64.2	0.94	0.97
50	65.6	92.25	98.9	99.45	0.76	0.87
75	52.8	76.7	83.3	85.6	0.81	0.90
100	40.55	63.9	71.1	72.75	0.81	0.90
R ²	0.302	0.0007	0.003	0.001		
x-coeff.	-0.54	0.027	-0.05	-0.036		

Table 2: Effect of different concentrations of DL- leucine on radial growth of *T. viride*.

The better growth of glutamic acid at its higher concentration gave 100% radial growth (Table 3). The other amino acids (alanine and Arginine) were ineffective in enhancing the radial growth of the fungus, even at their higher concentrations. Although alanine was slightly better than the control treatment at 50 mg/ml, it is far less than the other amino acids, while Arginine gave less radial growth than the control.

Production of cellulosic enzymes

Culture media containing cellulose substrates (Carboxymethyl cellulose-CMC, cotton fibers, filter

paper, and L-glucose) were inoculated with the fungus *T. viride* and incubated at room temperature (28- 30°C) for 24 days. The enzyme activity was measured every eight days using the viscosity-reducing method [27]. The comparative activity of the cellulosic enzyme at different incubation intervals (8,16, and 24 days) is shown in Fig. (1). Activity was found to increase with time. Carboxymethyl cellulose (CMC) was the best substrate for cellulase enzyme production, giving 35% of the comparative Activity. Other materials include L-glucose (dextrose), which gave 28%, then cotton fibers which gave 21%, and the minor comparative Activity was recorded for the filter paper culture. *T. viride* was tested on cultures of three pectic substrates (Pectin, orange rind, and L-glucose). The enzyme activity was also measured by the viscosity-reducing method. The comparative Activity of the pectolytic enzyme at different incubation intervals (8, 16, and 24 days) and in different substrates is shown in Fig. (2). Pectin culture was found as the best one for pectolytic enzyme production, giving 52% comparative Activity. Followed by the orange rind, which gave 34%, while the lowest comparative Activity was recorded in the L-glucose cultures.

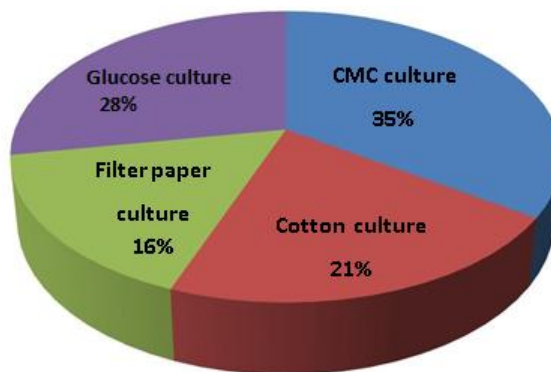


Figure 1: Comparative activity of cellulase enzymes produced by *T. viride* on different cultures after 8 days incubation time.

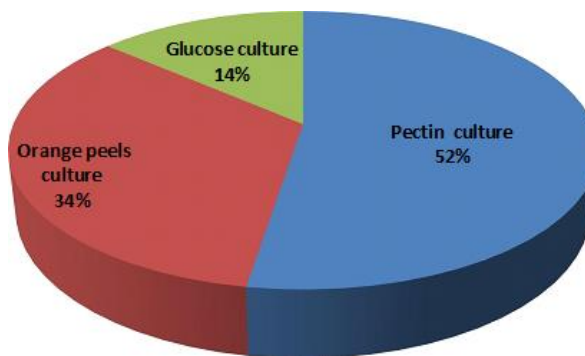


Figure 2: Comparative activity of pectolytic enzymes produced by *T. viride* on different cultures after 8 days incubation time.

Conc (mg/ ml)	% Radial growth					
	Incubation period days					
	2	4	2	8	2	x-coeff.
0	52.8	68.35	76.1	76.1	0.83	0.91
25	41.7	77.8	87.8	90.55	0.80	0.89
50	41.65	70	85.6	91.1	0.918	0.95
75	47.8	61.1	66.1	68.9	0.88	0.94
100	41.35	85.5	98.9	100	0.78	0.88
R ²	0.27	0.08	0.09	0.10	-	-
x-coeff.	-0.52	0.29	0.30	0.32	-	-

Table 3: Effect of different concentrations of Glutamic acid on radial growth of *T. viride*.

Discussion

The degradation and poor soil quality due to intensive agriculture, lousy land management techniques, and resource extraction from the soil are major global concerns. Contamination, erosion, salinization, flooding, and biodiversity loss are risks linked to soil deterioration [28]. Degraded soils are characterized by a lack of available organic matter in the soil as well as a depleted pool of organic carbon, with losses ranging from 25–75 percent of the initial soil organic matter and organic carbon pool [29]. Loss of biological variety and functioning occurs together with soil degradation. Recent research has demonstrated that anthropogenic activities and climate change impair soil's ability to carry out essential functions and processes such as primary production, nitrogen cycling, litter decomposition, and organic matter mineralization [30]. Most of the research under consideration reported that the rhizosphere had more microbial variety and richness than the control. The reason is that more nutrients and organic C substrates are available for mineralization [31].

According to Qiao et al.[32], the rhizosphere is a critical interface for interactions, signaling, and exchanging resources and energy between plants and soil microbes. This discovery demonstrates how these organisms often thrive in carbon-rich environments, which are prevalent in the rhizosphere, thanks to their high metabolic Activity, rapid development, and proliferative abilities. According to these results, O.A.s increase the supply of fresh organic materials, which widens niches, lessens competition, and fosters positive co-occurrence patterns with keystone saprotrophic species [8]. The rhizosphere also controls the dynamics of the C and N in the soil. The decomposition and transformation of organic compounds that were frequently more intense in the rhizosphere included cellulolysis, xylanolysis, ligninolytic, ureolysis, and chitinolysis, indicating the presence and Activity of more organisms that break down organic compounds [33].

Saprophytic fungi, including *Trichoderma* species significance in plant residue digestion and plant health, as well as the vast amount of microbial biomass they contribute to the soil, make them critical and typical

components of the soil rhizosphere. [34, 35]. *T. viride*, the subject of the present study, was very important. *Trichoderma* species, being saprophytic, are adapted to thrive in diverse situations, producing a wide array of enzymes [36, 37]. Microbial pectolytic enzymes, on the other hand, are known to play a vital role in many industrial processes such as; fiber extraction, clarification and pectination of fruit juices, and extraction of vegetable oils [38]. The present study tested the ability of the fungus *T. viride* to produce both cellulolytic and pectolytic enzymes in different substrates. The results showed that Carboxymethyl cellulose (CMC) was the best substrate for the cellulolytic enzyme, while Pectin was the best for the pectolytic enzyme production [39] in a similar study also reported that CMC was the best substrate for cellulolytic enzyme production. However, many fungi were reported to produce cellulolytic enzymes of high potency and stability; of these, *Trichoderma* species were found to be the most important [38]. Enzymatic hydrolysis of cellulose to glucose is simple, efficient, and cost-effective, reducing pollution and creating new food sources [39, 40].

The main polymer in plant cell walls, cellulose, is stable and recalcitrant [41]. The macromolecular structure of cellulose is more complicated than expected for a linear glucan chain polymer. Cellulose I is composed of β -(1,4)-d-glucan polymer fibrils. Supramolecular properties like lateral diameters vary by species in cellulose I fibrils. A fibril's polymers' order depends on their location, with surface polymers being the least restricted. Depending on the starting material and isolation method, cellulose I fibrils can form supramolecular structures, fibril aggregates, and the cellulose network, such as wood pulp fiber walls [42]. This complete porous network is space-oriented complexly. The surface area exposed to enzyme assault may limit material characteristics during hydrolysis [43]. Besides the complicated spatial orientation, cellulose structures have hydrophilic and hydrophobic faces, which exocellulase (CBHI) seems to prefer in cellulose hydrolysis [44].

The present study confirmed that *T. viride* needs some organic compounds. like nitrogenous compounds such as amino acids and others for their growth. Glutamic acid, methionine, and leucine were all influential in enhancing mycelial growth, but methionine is highly significant, while leucine and Glutamic acid gave a similar effect. The amino acid leucine gave the best mycelial growth at 50 % concentration. The effect of pH level on the fungus *T. viride* indicated that the maximum growth was at pH 5.0. However, growth decreased dramatically with increasing pH values.

The fungus *T. viride* had the ability to produce both cellulolytic and pectolytic enzyme production in some different substrates. Carboxymethylcellulose (CMC) was the best substrate for the cellulolytic enzyme. Furthermore, Pectin was the best for pectolytic enzyme production. For any industrial process or as a biocontrol agent, the growth of the fungus needs the addition of carbon and a nitrogen source to the medium. The best amino acids for culturing fungus are *T. viride*, leucine, and methionine, at a 50 mg/ ml concentration. Carboxymethylcellulose (CMC) was the best substrate for the cellulolytic enzyme. Pectin was the best for pectolytic enzyme production. The best incubation time for cellulolytic enzyme and pectolytic enzyme production was after 16 days. The present study confirmed that the fungus *T. viride* could produce cellulolytic and pectolytic enzymes in different substrates. Carboxymethyl cellulose (CMC) was the best substrate for cellulolytic enzymes. In addition, Pectin was the best for pectolytic enzyme production. Moreover, the best incubation time for cellulolytic enzyme and pectolytic enzyme production was after 16 days.

Competing Interest

The authors declare that there is no conflict of interest.

Author contribution

Conceptualization: A.S., W.A and S. H; methodology, A.S., S.H and N.A.; formal analysis, A.S., W.A and S. H; writing–original draft preparation, A.S and S.H.; writing–review and editing, A.S., S.H.; project administration, W.A. All authors have read and agreed to the published version of the final manuscript).

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