

Bioprospecting *Aspergillus* section *Nigri* in Atlantic Forest soil and plant litter

Bioprospecção de Aspergillus seção Nigri em solo e serrapilheira da Mata Atlântica

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ABSTRACT: The use of fungi as a source of enzymes has become widespread in various industrial and commercial areas, and *Aspergillus* section *Nigri* has significant potential for producing enzymes. The aim of this study was to isolate *Aspergillus* section *Nigri* from plant litter and soil from the Atlantic Forest biome and evaluate it with regards to hydrolytic enzyme production. The trials for producing the enzymes were carried out in Petri dishes, using different culture mediums adapted for microbial growth and with the respective substrates for inducing enzyme production — cellulase (carboxymethyl cellulose), protease (skimmed milk), amylase (soluble starch), pectinase (citrus pectin), and phytase (Pikovskaya medium). Forty-two fungi were isolated, 16.7% derived from the plant litter layer and 83.3% derived from soil at a depth of 0 to 5 cm and 5 to 10 cm. All of the isolated lineages presented amylase, protease, and phytase production, with 90.4% producing cellulase and no lineage producing pectinase. From the results, the significant potential for Atlantic Forest fungi as hydrolytic enzyme producers could be perceived. The enzymatic activity evaluations presented a satisfactory result when compared with the scientific literature.

KEYWORDS: filamentous fungi; enzymatic index; hydrolytic enzymes.

RESUMO: A utilização dos fungos como fonte de enzimas vem adquirindo *status* de destaque nas mais variadas áreas industriais e comerciais, e os *Aspergillus* membros da seção *Nigri* possuem significativo potencial para produção de enzimas. Os objetivos deste estudo foram isolar e avaliar *Aspergillus* da seção *Nigri* de serrapilheira e solos do bioma Mata Atlântica quanto à produção de enzimas hidrolíticas. Os ensaios para produção das enzimas foram realizados em placas de Petri, utilizando diferentes meios de cultivo, adequados ao crescimento microbiano e com a presença dos respectivos substratos indutores à produção das enzimas — celulasas (carboximetilcelulose), proteases (leite desnatado), amilases (amido solúvel), pectinases (pectina cítrica) e fitase (meio Pikovskaya). Foram isolados 42 fungos, sendo desse total 16,7% provenientes da camada de serrapilheira e 83,3% provenientes do solo na profundidade de 0 a 5 cm e 5 a 10 cm. Todas as linhagens isoladas apresentaram produção de amilases, protease e fitase, 90,4% produziram celulase, e nenhuma linhagem produziu pectinase. Com esses resultados, percebeu-se significativo potencial dos fungos da Mata Atlântica como produtores de enzimas hidrolíticas. As avaliações da atividade enzimática apresentaram resultado satisfatório quando comparados à literatura científica.

PALAVRAS-CHAVE: fungos filamentosos; índice enzimático; enzimas hidrolíticas.

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INTRODUCTION

Brazil contains the greatest biodiversity on the planet. Among the country's various biomes, the Atlantic Forest features as one of the 34 global "hotspots". Its climatic conditions enable the existence of microorganisms that decompose and degrade forest biomass, making the soil in this biome very rich in nutrients. Among these microorganisms, fungi are of crucial importance to the functioning of this ecosystem.

Fungi of the *Aspergillus* genus are filamentous, cosmopolitan, saprophytes, and are mainly found in regions with tropical and subtropical climates (KLICH, 2002). Some species of this genus are used in biotechnology, in which they produce a diversity of biocompounds such as antibiotics, microtoxins, enzymes, organic acids, and phenolic compounds, among other substances (DAGENAIS; KELLER, 2009).

Aspergillus stands out as being the most important genus for the commercial production of extracellular enzymes (NOVAKI, 2009), the most widely used species being *Aspergillus flavus*, *A. niger*, *A. awamori*, *A. oryzae*, *A. nidulans*, *A. fumigatus*, *A. clavatus*, *A. glaucus*, *A. ustus*, and *A. versicolor* (SLIVINSKI, 2007). Also in this genus, the species from the *Nigri* section have been extensively studied, as they are good producers of enzymes such as amylase, cellulose, pectinase, protease, and phytase (SALES et al., 2010; RODRÍGUEZ et al., 2011; REZENDE et al., 2012; MACIEL, 2013).

The use of these fungi as enzyme producers is recognized by the Food and Drug Administration (FDA), from the United States, and the products derived can be used for safe human consumption (SAID; PIETRO, 2002). Thus, some species of the *Aspergillus* genus have a Generally Recognized as Safe (GRAS) status, due to their low toxicity and historic use in the preparation and production of foods and drinks (ABARCA et al., 2004).

Enzymatic technology is being used to obtain advantages in establishing a technologically clean process, in which hydrolytics are produced by different species of *Aspergillus*, with a broad diversity regarding the characteristics of these enzymes, enhancing their application in different industry sectors, such as the textile, pulp and paper, pharmaceutical, and especially food industries (BARATTO et al., 2011).

The main sectors that use a large number of these enzymes are the food and drink sectors, in which amylase and cellulase enzymes are used in preparing the mash for producing beer. In baking, amylase and peptidase enzymes are used to facilitate the manipulation of dough. Peptidase is mainly used in processing fruit juices and wines, and in the treatment of silage along to phytase, as it improves nutritional quality and facilitates digestion by ruminant and monogastric animals (PANDEY et al., 2005; UENOJO; PASTORE, 2007; BECKER et al., 2009).

Therefore, studies have focused on selecting new fungi as a source of new enzymes, exploring the biotechnological potential

of global biodiversity (ABBASI et al., 2011; RODRÍGUES et al., 2011; MACIEL, 2013). The inoculation technique is used as a way of selecting, with Petri dishes and a solid medium containing a source of enzymatic activity induction, such as starch, pectin, skimmed milk, calcium phosphate, and carboxymethyl cellulose (GOPINATH et al., 2005).

Considering the little knowledge regarding the fungi found in areas of the Atlantic Forest, the importance of studies that contribute to expanding the data on fungi diversity in this threatened biome is clear, as well as the use of these fungi in biotechnological processes.

MATERIAL AND METHODS

Plant Litter and Soil Sampling

The plant litter and soil sample collection areas are located in the central west of Minas Gerais state, in the municipality of Sete Lagoas.

The plant litter and soil samples were collected in two periods: the first wet, at the end of January, and the second dry, at the beginning of September. Nine forest fragments were sampled, with the sampling carried out at three depths (plant litter, 0 to 5 cm, and 5 to 10 cm), with three repetitions for each depth, thus totaling 162 samples. After collection, the packages with the samples were stored at 4°C until analysis.

Isolating the fungal lineages

The isolations were carried out using five grams of soil and plant litter resuspended in a Erlenmeyer flask with 45 mL of 0.1% peptone containing 0.1% Tween 80, then shaken for 30 min in a vertical rod homogenizer at 150 rpm. The culture medium used was Dicloran Rosa de Bengal Cloranfenicol (DRBC), with 0.1 mL doses of the respective dilutions (10^2 to 10^6), incubated in ovens at 25°C for five to seven days (FRAGA; PEREIRA, 2012).

Morphological analyses of the isolated lineages

The morphological identification was carried out based on a suspension of *Aspergillus* section *Nigri* spores in a solution containing 0.2% agar and 0.05% Tween 80. From the suspension, 0.2 mL was used for inoculation at three equidistant points in the Petri dish containing the Czapek Yeast Agar (CYA) and Malt Extract Agar (MEA) culture mediums, in triplicate for each medium.

The incubation was carried out at 25 and 37°C for the dishes containing the CYA medium and 25°C for the MEA medium. After an inoculation period of seven days, their

macroscopic (color of surface and back of the colony, colony diameter, presence or absence and color of sclerotia) and microscopic (vesicle diameter and form, phialide length and width, color, size, form and texture of the conidia, and color of the hyphae) characteristics were analyzed. Inoculations were also carried out in Petri dishes containing MEA medium for 10 days to verify growth at temperatures of 15, 33, 36, and 40°C (KLICH, 2002; SAMSON et al., 2004; SAMSON et al., 2007; VARGA et al., 2011; MEIJER et al., 2011).

Enzymatic index of the lineages

The enzymatic index (EI) is one of the parameters used in order to evaluate the production of enzymes by microorganisms in a solid medium. For the microorganisms considered enzyme producers, there is a direct correlation between the diameter of the halo of degradation and their degradative ability (LIN et al., 1991). In this context, the *Aspergillus* section *Nigri* isolates were evaluated as producers of cellulose, peptidase, amylase, phytase, and pectinase. Inoculation of the lineages was carried out in dishes containing the mediums, in a 10^4 spores/mL concentration, and incubated for five days at 25°C.

To detect the enzymes, the cellulase activity was evaluated using the synthetic medium containing carboxymethyl cellulose (CMC) as the only source of carbon, the composition being in accordance with RUEGGER; TAU-K-TORNISIELO (2004). After the incubation period, the revelation of the halo of hydrolysis was carried out by adding 10 mL of Red Congo coloring solution (2.5 mL in a Tris buffer 0.1 M, pH 8) over each colony, removing the solution after 30 min and washing the cultures in 5 mL of NaCl 0.5 M solution in this same buffer, for 15 min. Carboxymethyl cellulose degradation around the fungal colony is detected via the formation of a yellow halo around the colony (POINTING, 1999).

The degradation capacity of the amylase and pectinase was verified in accordance with HANKIN; ANAGNOSTAKIS (1975). The amylolytic activity was detected after the addition of 5 mL of an iodine solution over each colony. Formation of a yellow zone around the fungal colony and the development of petrol blue coloring on the remaining surface of the Petri dish indicate the occurrence of starch hydrolysis. The pectinolytic activity was verified after adding the chloridric acid solution (HCL) 5 N, for 5 min.

The degradation capacity of the peptidase was verified in accordance with GOPINATH et al. (2005).

Pikovskaya Medium (PIKOVSKAYA, 1948) was used to evaluate the productive capacity of the phytase. The halo indicating the production of peptidase and phytase enzymes can be seen without a revealing solution, with a clear zone being detected around the colonies.

The colony diameters and halos produced were measured with a pachymeter (NOGUEIRA; CAVALCANTI, 1996). The enzymatic index (EI) was calculated by the ratio between the

average diameter of the halo of degradation and the average diameter of the colony, as proposed by STAMFORD et al. (1998).

Statistical Analysis

The experiment was conducted in an entirely randomized design (ERD). All of the tests were carried out with three repetitions, in accordance with the number of positive treatments for each enzyme. The data obtained were submitted for analysis of variance, using the Sisvar software (FERREIRA, 2000). The averages of the treatments were compared using the Scott-Knott test at a 5% level of probability.

RESULTS AND DISCUSSION

In this perspective, 1,854 fungal lineages from the plant litter and soil samples were isolated, 907 lineages in the wet period and 947 lineages in the dry period. Of this total number of isolates, 42 were *Aspergillus* section *Nigri*, 16.7% derived from the plant litter layer and 83.3% derived from soil at depths of 0 to 5 cm and 5 to 10 cm.

In order to evaluate the cellulase production potential, it was observed that, of the lineages isolated from *Aspergillus* section *Nigri*, 38 of them produced cellulase; that is, 90.4% presented a halo of hydrolysis. The statistical analysis of the cellulase production showed that there was a difference between the enzymatic indices of the isolates analyzed, forming three different groups (a-c). Four isolates stood out from the rest in relation to the enzymatic index: *A. aculeatus*, 42 with 1.36; *A. aculeatus*, 13 with 1.37; *A. aculeatinus*, 9; and *A. aculeatus*, 36 with 1.38, and those that presented a negative result were the *A. niger* aggregates (Table 1).

According to RUEGGER; TAU-K-TORNISIELO (2004), enzymatic indices of around 1.6 are values found in fungi of the *Aspergillus* sp. genus, in which this index can be used as a useful measure for selecting lineages within the same species. These same authors evaluated the cellulase activity of fungi isolated from soil and observed that 45% of the isolates presented positive activity, since there was formation of a halo, indicating degradation of the medium; however, the *A. niger* species obtained an EI value of 1.4, close to the values found in our study for species of the *Nigri* section.

FERNANDES (2009) tested 24 fungal lineages isolated from various sources in order to detect cellulase, among which nine belonged to the *Aspergillus* section *Nigri* genus: *A. carbonarius*, EI = 1.4 and 1.14; *A. niger*, EI = 1.10 and 1.27; *A. foetidus*, EI = 1.13 and 1.16; *A. niger* aggregates, EI = 1.11, where these last EI values were close to those ones found in this study. AGUSTINI et al. (2012) also carried out a qualitative test of cellulase production, using the CMC medium in 553 isolates, containing fungi and bacteria, and found the

Table 1. Evaluation of the lineages of *Aspergillus* section *Nigri* isolated from the plant litter and soil, by their respective Enzymatic Indices.

<i>Aspergillus</i> Isolates	Cellulase			Peptidase (Protease)			Amylase		
	Øc	Øh	E.I.*	Øc	Øh	E.I.*	Øc	Øh	E.I.*
02 <i>A. aculeatus</i>	2.57	3.30	1.29 ^b	6.5	7.83	1.20 ^d	4.1	4.8	1.16 ^b
03 <i>A. aculeatus</i>	2.90	3.63	1.25 ^b	5.57	7.5	1.25 ^c	3.2	3.8	1.22 ^a
04 <i>A. aculeatinus</i>	3.63	4.10	1.13 ^c	6.87	8.33	1.21 ^d	4.1	4.8	1.16 ^b
05 <i>A. aculeatinus</i>	3.60	4.27	1.18 ^c	6.77	8.7	1.28 ^c	4.1	5.2	1.28 ^a
06 <i>A. aculeatinus</i>	3.63	4.30	1.18 ^c	7.5	8.83	1.17 ^e	4.8	6.1	1.26 ^a
07 <i>A. aculeatinus</i>	3.80	4.47	1.17 ^c	7.2	8.73	1.21 ^d	4.5	5.3	1.16 ^b
08 <i>A. aculeatus</i>	3.13	3.87	1.24 ^b	6.17	7.3	1.18 ^e	3.9	4.8	1.22 ^a
09 <i>A. aculeatinus</i>	2.63	3.57	1.38 ^a	7.03	8.5	1.21 ^d	4.1	4.9	1.21 ^a
10 <i>A. aculeatus</i>	3.27	3.93	1.20 ^c	6.47	8	1.23 ^c	3.6	4.5	1.24 ^a
11 <i>A. aculeatus</i>	3.33	4.27	1.28 ^b	5.93	7.27	1.22 ^d	3.8	4.4	1.17 ^b
12 <i>A. aculeatus</i>	3.00	3.53	1.18 ^c	6.83	8.1	1.18 ^e	4.5	5.3	1.18 ^b
13 <i>A. aculeatus</i>	2.57	3.50	1.37 ^a	5.53	7.1	1.29 ^c	4.0	4.7	1.16 ^b
14 <i>A. aculeatus</i>	3.23	3.77	1.17 ^c	5.63	7.1	1.26 ^c	3.8	4.7	1.23 ^a
15 <i>A. aculeatinus</i>	3.83	4.30	1.12 ^c	6.97	8.5	1.22 ^d	4.2	5.3	1.25 ^a
16 <i>A. aculeatus</i>	2.80	3.67	1.31 ^b	6.07	7.33	1.21 ^d	4.2	4.9	1.18 ^b
17 <i>A. aculeatus</i>	3.37	4.03	1.20 ^c	5.7	6.77	1.18 ^e	3.2	3.9	1.23 ^a
18 <i>A. aculeatus</i>	3.23	4.17	1.29 ^b	5.37	6.77	1.26 ^e	4.0	4.8	1.18 ^b
20 <i>A. aculeatinus</i>	3.30	4.17	1.26 ^b	7.03	8.77	1.24 ^c	4.2	5.3	1.25 ^a
21 <i>A. aculeatinus</i>	3.30	4.17	1.26 ^b	7.03	8.77	1.25 ^c	3.6	4.5	1.26 ^a
22 <i>A. aculeatinus</i>	3.60	4.07	1.13 ^c	6.9	8.83	1.28 ^c	4.6	5.6	1.23 ^a
23 <i>A. aculeatus</i>	3.07	3.80	1.24 ^b	5.6	6.73	1.20 ^d	3.5	4.4	1.27 ^a
24 <i>A. aculeatinus</i>	3.03	3.50	1.15 ^c	6.67	8	1.20 ^d	3.5	4.1	1.19 ^b
25 <i>A. aculeatus</i>	3.53	4.27	1.21 ^c	5.43	6.77	1.24 ^c	3.9	4.6	1.19 ^b
26 <i>A. aculeatus</i>	3.57	4.10	1.15 ^c	6.47	7.53	1.16 ^e	4.6	5.2	1.14 ^b
28 <i>A. aculeatinus</i>	2.77	3.50	1.28 ^b	7.07	8.73	1.23 ^c	4.6	5.7	1.24 ^a
29 <i>A. aculeatus</i>	3.57	4.43	1.24 ^b	6.6	7.67	1.16 ^e	4.2	4.9	1.18 ^b
31 <i>A. aculeatinus</i>	3.53	4.27	1.21 ^c	6.73	8.27	1.22 ^d	3.5	4.3	1.25 ^a
32 <i>A. aculeatus</i>	3.63	4.10	1.13 ^c	5.57	6.83	1.23 ^d	4.0	4.7	1.17 ^b
33 <i>A. aculeatus</i>	3.17	3.90	1.23 ^b	5.8	7.27	1.25 ^c	3.9	4.6	1.19 ^b
34 <i>A. aculeatus</i>	3.40	4.27	1.25 ^b	6.83	8.1	1.18 ^c	4.0	4.9	1.23 ^a
35 <i>A. aculeatus</i>	3.57	4.23	1.19 ^c	6.97	8.17	1.17 ^e	3.3	3.9	1.20 ^b
36 <i>A. aculeatus</i>	2.67	3.67	1.38 ^a	6.33	8	1.26 ^c	4.4	5.3	1.19 ^b
37 <i>A. aculeatus</i>	3.33	4.07	1.22 ^c	5.43	6.77	1.24 ^c	3.5	4.4	1.25 ^a
38 <i>A. aculeatus</i>	3.57	4.33	1.21 ^c	6	7.27	1.21 ^d	4.0	4.7	1.17 ^b
39 <i>A. aculeatus</i>	3.63	4.30	1.18 ^c	5.63	7.1	1.26 ^c	3.6	4.5	1.24 ^a
40 <i>A. aculeatus</i>	3.70	4.37	1.18 ^c	6.67	7.73	1.15 ^e	4.2	4.8	1.16 ^b
41 <i>A. aculeatus</i>	3.43	4.37	1.27 ^b	6.07	7.07	1.16 ^e	3.8	4.7	1.23 ^a
42 <i>A. aculeatus</i>	3.10	4.23	1.36 ^a	5.37	6.77	1.26 ^c	4.7	5.8	1.22 ^a
Aggregates									
01 <i>A. niger</i>	-	-	-	5.3	7.43	1.40 ^a	3.7	4.3	1.18 ^b
19 <i>A. niger</i>	-	-	-	6.1	8.17	1.34 ^b	4.2	5.2	1.26 ^a
27 <i>A. niger</i>	-	-	-	6	8.35	1.40 ^a	3.9	5.0	1.28 ^a
30 <i>A. niger</i>	-	-	-	6.2	8.33	1.34 ^a	4.5	5.6	1.25 ^a

Øc: colony diameter (cm); Øh: halo diameter (cm); EI: enzymatic index; (-): did not grow and/or did not produce a halo; ^{a, b, c}: similarity of enzymatic index, in which same letters similar results.

A.s niger species to be a cellulase producer, in which the respective indices varied from 1 to 3.36 for all of the microorganisms.

MONTEIRO et al. (2012) selected cellulase producing isolates belonging to the *Aspergillus* and *Penicillium* genera, and, of the 135 isolates tested, only 20 were producers, and the five isolates that stood out with regards to qualitative production of cellulase belong to the *Aspergillus* genus.

With regards to protease production, a significant difference between the averages analyzed can be observed in Table 1, with the separation of five different groups (a-e) occurring with a variation in EI of between 1.15 and 1.40. The isolates that presented the highest indices were the fungi from the *A. niger* aggregate group, with EI = 1.40 and 1.34. None of the isolates presented a negative result.

In papers from FERNANDES (2009) and SCHUSTER et al. (2002), *A. niger* did not produce a halo of degradation indicative for proteolytic determination, unlike in this study, in which the *Nigri* section species produced a halo of degradation, although with $EI \leq 2.0$. PEREIRA (2012) also observed protease enzyme production when analyzing 40 isolates belonging to the EcoCenter Fungi Collection/Empresa de Pesquisa Agropecuária de Minas Gerais (EPAMIG) and the Microtoxins and Mycology Laboratory of Universidade Federal de Lavras (UFLA), of which six belonged to the *Aspergillus* genus and presented the same results, $EI \leq 2.0$, with there being no higher values for this genus. On the other hand, SOHAIL et al. (2009) obtained 81% protease producing fungi in 160 isolates, and among these the *A. niger* and *A. flavus* species were considered potential producers of proteolytic enzymes.

In the evaluation of the potential production of the amylase enzyme, 100% presented a halo of hydrolysis, the results of which are presented in Table 1. There was a small difference between the averages, forming two groups (a and b); however, there was no isolate that stood out. All the results indicated that the lineages tested are able to degrade the starch present in the culture medium, but no lineage presented enzymatic potential, since they did not reach $EI \geq 2.0$.

TEIXEIRA (1994) found that the greatest halos of enzymatic activity were produced by *A. niger* for amylases, cellulases, and pectinases. MORAES (2004) also observed that *A. niger* was able to degrade α -amylase. However, these results were not similar to those obtained in this study.

SILVA et al. (2006) did not find amylolytic activity in 29 *Aspergillus* isolates. When analyzing mutant lineages of the *Aspergillus nidulans* fungus, FLORES et al. (2010) observed little halos, smaller than 2.0. PEREIRA (2012) also studied the *Aspergillus* genus for amylase production; however, they were not producers of this enzyme using the methodologies tested. Thus, these papers partly support the results presented in this study, since they affirm that amylase production by *Aspergillus* in a solid culture medium may not be positive or involve little amylolytic activity.

All of the isolates in this paper presented growth in the selective medium for phytase, with an average colony diameter

of 70 mm; however, a halo of degradation was not formed, and it was, therefore, not possible to measure the enzymatic indices of the *Aspergillus* section *Nigri* isolates.

The non formation of a halo of degradation may be due to the extensive growth of the fungus in this culture medium, after five days of incubation, as the fungus presented a colony with a diameter close to the size of the Petri dish. However, according to CASTRO; PEREIRA (2010), visualization of a halo depends on various factors, besides the composition of the culture medium, and some chemical substances can interfere in the coloring, leading to false-positives or even causing its precipitation or inhibiting the link between it and the polysaccharides.

The 42 lineages did not grow and did not present a halo of hydrolysis indicating pectinase activity.

PEREIRA (2012) tested two methodologies for producing pectinase. For the first one, there was low sensitivity to detecting the enzyme, with low results occurring, in which the EI values of the *Aspergillus* fungi were between 1.3 and 1.6, compared with the second methodology, in which the EI was between 1.6 and 4.8. However, just as in the results obtained in this study, the mentioned papers showed little or no pectinase activity for fungi of the *Aspergillus* genus. This was unlike the studies carried out by SANDRI (2010), which evaluated 60 fungal isolates from *A. niger* and *A. oryzae* with regards to the formation of a halo of pectin degradation in a solid medium, a qualitative test, and subsequent production by fermentation. The data showed that of the 30 *Aspergillus niger* isolates only six did not form a halo of hydrolysis, and the EI value varied from 1.2 to 2.8.

It should be highlighted that the ability of a fungus to produce enzymes varies between species, as it also does between isolates from the same species, and it is quite variable. Innumerable factors of a biological and physical-chemical nature, such as pH and temperature, can influence the number and enzymatic activity of the microorganisms. The absence of activity, as occurred with the pectinase enzyme, may be attributed to the incapacity for secretion, to the direction of production to the intracellular metabolism, or to the insufficiency of the detection methods.

CONCLUSION

The results showed significant potential for Atlantic Forest fungi as producers of hydrolytic enzymes. All isolated lineages presented amylase, protease, and phytase production, with 90.4% producing cellulase and no lineage producing pectinase. The quantitative evaluations of enzymatic activity presented satisfactory results, and a quantitative evaluation is needed with regards to the potential of these fungi as producers of hydrolytic enzymes.

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