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Production of β -(1,3)-glucanases by *Trichoderma harzianum* Rifai: Optimization and Application to Produce Gluco-oligosaccharides from Paramylon and Pustulan

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Abstract

 β -(1 \rightarrow 3)-Glucanases were produced by *Trichoderma harzianum* Rifai PAMB-86 cultivated on botryosphaeran in a bench-fermenter and optimised by the response surface method. Maximal enzyme titres occurred at 5 days, initial pH 5.5 and aeration of 1.5vvm. β -(1 \rightarrow 3)-The β -glucanolytic enzyme complex produced by *T. harzianum* Rifai PAMB-86 was fractionated by gel filtration into 2 fractions (F-I, F-II), and employed to produce gluco-oligosaccharides from algal paramylon ((1 \rightarrow 3)- β -D-glucan) and lichen pustulan ((1 \rightarrow 6)- β -D-glucan). Both enzymes attacked paramylon to the extent of ~15-20% in 30 min releasing glucose and laminaribiose as major end-products, and laminarioligosaccharides of degree of polymerization (DP) ≥3. Only F-I degraded pustulan resulting in ~2% degradation at 30 min, with glucose, gentiobiose and gentio-oligosaccharides of DP ≥4 as major products. The difference in the nature of the hydrolysis products can be explained by the substrate specificities of each enzyme fraction, and the structural differences of the β -D-glucans attacked.

Keywords: Trichoderma harzianum PAMB-86; β -(1 \rightarrow 3)-Glucanases; Response surface method; Paramylon; Lichen Pustulan; Gluco-oligosaccharides

Introduction

Many fungal β -D-glucans have been described that possess biological activities including immunomodulation [1]. Modification of polysaccharides has contributed to the development of new industrial applications for these biopolymers in a variety of commercial sectors such as pharmaceuticals, cosmetics and foods [2,3]. Modification through hydrolysis produces gluco-oligosaccharides which have demonstrated health benefits that influence the immune system, and are recognized as having biological response modifying activities [4,5].

Paramylon, a storage (13)- β -D-glucan derived from the alga, *Euglena gracilis*, consists of a non-ramified chain constituted of D-glucose units [6]. Pustulan is a linear (1 \rightarrow 6)- β -D-Glucan usually isolated from lichens [7], but these β -D-glucans are also present in fungi [8], where they exist as a cell wall component. More recently, water-soluble exocellular (1 \rightarrow 6)- β -D-glucans have been reported for several *Botryosphaeria rhodina* strains [9], and *Guignardia citricarpa* [8]. Pustulan is commonly used as a substrate for assaying β -1(\rightarrow 6)glucanase activity [10], and as carbon source in screening for this type of enzyme activity [11].

 β -(1→3)-Glucanases have been used to depolymerize (1→3) (1→6)-β-D-glucans to obtain degraded fragments that possess biological and functional properties for use as pharmaceuticals, and inclusions as prebiotics in foods and beverages [4,12,13]. The β-(1→3)glucanases are predominantly of two enzyme types: *exo*- (EC 3.2.1.58) and *endo*- (EC 3.2.1.39) acting that specifically attack β-(1→3)-linked D-glucans [14]. Accessory enzymes that complement the key enzymes involved in the degradation of the (1→3)(1→6)-β-D-glucans include the β-(1→6)-glucanases (EC 3.2.1.75) and β-glucosidases (EC 3.2.1.21); each produced by several fungal species [15,16]. *Trichoderma* species have been widely investigated as bio-control agents and are renowned to produce hydrolytic enzymes that act synergistically on plant and fungal cell wall polysaccharides. Enzymes from *Trichoderma* species, and esp. *T. harzianum*, have been used to degrade extracellular (1 \rightarrow 3) (1 \rightarrow 6)- β -D-glucans to produce gluco-oligosaccharides [10,17].

In this work, we report on the production of β -(1 \rightarrow 3)-glucanases by *Trichoderma harzianum* Rifai PAMB-86 cultivated in a fermenter on (1 \rightarrow 3)(1 \rightarrow 6)- β -D-glucan (botryosphaeran) as sole carbon source, and their optimization by the response surface method (RSM). The β -(1 \rightarrow 3)-glucanases produced were partially fractionated and employed to enzymatically hydrolyze paramylon and pustulan. Studies involving specific β -glucanases in the hydrolysis of algal and lichen β -glucans are important to understand the nature of the hydrolysis reactions involved, and to develop new strategies to obtain glucooligosaccharides on a large scale for biotechnological applications.

Material and Methods

Materials

Paramylon (an unbranched $(1\rightarrow 3)$ - β -D-glucan from *Euglena* gracilis) was purchased from Fluka. Pustulan $((1\rightarrow 6)$ - β -D-glucan), a kind gift from Professor Philip Gorin (Universidade Federal do Paraná, Curitiba-PR, Brazil), was isolated from the lichen, *Actinogyra* muehlenbergii [7]. Laminarin (*Laminaria digitata*), laminaribiose,

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laminaritriose, gentiobiose and *p*-nitrophenyl- β -D-glucopyranoside (*p*-NPG) were purchased from Sigma-Aldrich. Botryosphaeran (a (1 \rightarrow 3)(1 \rightarrow 6)- β -D-glucan) was obtained from *Botryosphaeria rhodina* MAMB-05 according to Barbosa et al. [18].

Microorganism and culture conditions

Trichoderma harzianum Rifai (isolate PAMB-86) was obtained from decaying peroba rosa wood (Aspidosperma spp.) and maintained on xylose-agar at 4°C [11] incorporating Vogel minimal salts medium (VMSM) [19]. T. harzianum Rifai PAMB-86 conidia were used to inoculate (1×108 spores/flask) three 125 ml Erlenmeyer flasks containing 25 ml liquid medium comprising VMSM and glucose (10 g/l) during 72 h, 180 rpm at 28°C. The mycelium resulting was used as inoculum in fermenter-based experiments. The fungal isolate was cultivated in a laboratory fermenter of 2.5 l capacity equipped with a pH electrode, and facilities for control of agitation and aeration (Technal, Brazil), and was operated in a batch continuous mode. The fermenter was equipped with four removable standard baffles, and a top driven agitator shaft mounted with three standard blades (Rushton impellers). Fermentation was carried out using 600 ml medium comprising VMSM and botryosphaeran (1.5 g/l) as sole carbon source at 28°C and 120 rpm. Aeration, initial pH and time of growth for the production of β -(1 \rightarrow 3)-glucanases by *T. harzianum* Rifai PAMB-86, were used as variables in the statistical experimental design outlined below. Extracellular fluid (ECF) was recovered following centrifugation (7000xg/10 min), exhaustively dialyzed against deionised water at 4°C, and used as the source of enzyme.

Enzyme assays

β-Glucanase activity was determined using the polysaccharides: laminarin (β-(1→3)-glucanase) and pustulan (β-(1→6)-glucanase) according to Giese et al. [10]. The hydrolysis products released were measured as reducing sugars [20]. The unit of each β-glucanase activity was defined as the number of µmol reducing sugars produced/ min. β-Glucosidase activity was measured against *p*-nitrophenyl-βglucopyranoside (15 mM) as substrate, and incubated at 55°C for 20 min [10]. The unit of β-glucosidase activity was defined as the number of µmol *p*-nitrophenol liberated/min.

Analytical procedures

Hydrolysis products arising from various glucans were measured by the reducing sugars method [20]. Glucose from gluco-oligosaccharides was measured by the glucose oxidase method using a kit (Glicose Enz Color reagent kit; Bio Diagnóstica, Curitiba, Brazil).

HPAEC/PAD analysis of enzymatic hydrolysates

Following enzymatic hydrolysis, undigested polymeric material remaining in the hydrolysates was removed by precipitation with 3 volumes of ethanol [10]. The supernatant was recovered by centrifugation (7000x g/10min), and the ethanol evaporated under vacuum. The resulting syrup was re-solubilized in water, and used for determination of reducing sugars, and aliquots of 0.025 ml taken for sugar analysis by High Performance Anionic Exchange Chromatography with Pulsed Amperometric Detection (HPAEC/PAD) on a Dionex Chromatograph DX 500. Mono- and oligo- saccharides were separated on a CarboPac PA100 (Dionex Chromatography) column (4×250 mm) equipped with a PA-100 guard column at a flow rate of 1.0 ml/min. The column was equilibrated in 0.1 M NaOH (97%) and 0.5 M sodium acetate (3%). After 15 min, a linear 0-0.25 M concentration gradient of sodium acetate was applied over a 70 min interval while

the concentration of NaOH remained at 0.1 M. Sugar quantification was carried out from peak area measurements using response factors obtained from authentic standard sugars, and identified by their retention times (T_R ± SD min): glucose (2.93 ± 0.06 min), gentiobiose (4.57 ± 0.21 min), laminaribiose (7.12 ± 0.80 min) and laminaritriose (17.34 ± 0.20 min). Peaks exiting the chromatography column after 20 min were considered oligosaccharides of DP 4 and greater than 4.

Factorial design and analysis

Fermentation conditions to optimise β -(1- \exists 3)-glucanase production by *T. harzianum* Rifai PAMB-86 were studied as variables in a factorial design and analysis by the response-surface method (Table 1). The independent variables were x_1 = aeration (vvm), x_2 = initial pH, and x_3 = time of growth. The level of these variables and the variation levels for experimental studies on β -(1 \Rightarrow 3)-glucanase production ($Y_1 = U/$ ml) are shown in Table 1. Analysis of variance (ANOVA) and multiple regression analysis were performed using STATISTICA Version 6.

Fractionation of β -glucanolytic complex from T. harzianum Rifai PAMB-86

Extracellular fluid (600 ml) containing the enzyme was lyophilized following centrifugation to remove the mycelium. Lyophilized ECF was dissolved in 20 mM sodium acetate buffer (pH 5.0) and applied to a column of Sephadex G-100 (2.5×90 cm, Pharmacia Biotech, Sweden), and eluted with 20 mM sodium acetate buffer (pH 5.0) at a flow rate of 15 ml/h. Fractions of 2.5 ml were collected and analyzed for β -(1 \rightarrow 3)-glucanase, β -(1 \rightarrow 6)-glucanase and β -glucosidase activities. Two fractions were isolated (designated F-I and F-II), and used as the source of enzyme to hydrolyse paramylon and pustulan. All fractionation steps were performed at 4°C.

Results and Discussion

Production of β -(1+3)-glucanases by Trichoderma harzianum Rifai PAMB-86

Production of microbial β -(1 \rightarrow 3)-glucanases are strongly

Experimental run	Variables in coded levels			β -(1 \rightarrow 3)-Glucanase activity (U/ml)			
	X 1	X ₂	X ₃	Experime	ental	Predicted	
1	-1	-1	-1	0.69		0.62	
2	-1	-1	+1	1.88		1.88	
3	-1	+1	-1	0.92		0.93	
4	-1	+1	+1	1.52		1.58	
5	+1	-1	-1	0.94		0.95	
6	+1	-1	+1	1.96		2.02	
7	+1	+1	-1	0.60		0.66	
8	+1	+1	+1	0.72		1.11	
9	+1	0	+1	3.71		3.46	
10	0	+1	+1	1.63		1.37	
11	+1	+1	0	1.72		1.46	
12	0	0	0	3.65		3.71	
13	0	0	0	3.52		3.71	
—	Real levels						
Factor	-1			0	+1		
x_1 , aeration (vvm)	1.0			1.5		2.0	
x ₂ , initial pH	3.0 5.5			8.0			
x_{3} , time of growth (days)		2		5		8	

Table 1: A central composite design [30] defining conditions for aeration, initial pH and time of growth on the production of β -(1 \rightarrow 3)-glucanases by *Trichoderma harzianum* Rifai PAMB-86.

influenced by the level of β -D-glucan present as inducer, and the types of glucosidic linkages present in the β -glucans used as carbon sources [21-24]. In a previous study, production of β -(1 \rightarrow 3)-glucanases by *T*. *harzianum* Rifai grown on botryosphaeran (a $(1\rightarrow 3)(1\rightarrow 6)$ - β -D-glucan from B. rhodina MAMB-05 [23] was optimized in shake-flasks. The results showed that production of β -(1 \rightarrow 3)-glucanases was dependent upon the concentration of botryosphaeran and the time of growth, and highest enzyme titres obtained were 1.2 U/ml. A statistical mixturedesign indicated a synergistic effect of botryosphaeran on β -(1 \rightarrow 3)glucanases production by T. harzianum Rifai in combination to glucose and lactose as a mixture of carbon sources [24]. It was impractical to increase the botryosphaeran concentration in the nutrient medium in attempts to enhance enzyme titres as this exopolysaccharide is sparingly soluble in water (limit of 3 g/l), and aqueous solutions formed are rather viscous and difficult to manage. Instead, fermentation variables such as initial pH and aeration rate were evaluated in a 2.5 l fermenter (600 ml) to improve β -(1 \rightarrow 3)-glucanase titres by *T. harzianum* Rifai PAMB-86 when cultivated on botryosphaeran. Enzyme production was optimised by the response surface method.

Through multiple regression analysis of the experimental data, a second-order polynomial equation was obtained for β -(1 \rightarrow 3)-glucanase production (Equation 1).

$$\hat{Y}_{1} = 3,71294 + 0,42793x_{3} - 1,89259x_{2}^{2}$$
⁽¹⁾

Linear-effect terms of the variables x_1 and x_2 and squared-effect terms of the variables x_1 and x_2 were discarded as being non-significant. According to the results, an intercept was significant indicating that the central-points (aeration, 1.5 vvm; initial pH, 5.5; and 5 d growth) were correctly chosen. The variable more important for β -(1 \rightarrow 3)-glucanase production by *T. harzianum* Rifai PAMB-86 was the initial pH followed by the time of growth. The analysis of variance (ANOVA) showed the lack-of-fit (p > 0.05) was not significant, indicating that the model was predictive (Table 2). The experimental and predicted β -(1 \rightarrow 3)-glucanase activities were in agreement (Table 1). The R-squared value implied 97% of the variability in the observed response values could be explained by the model, or by experimental factors and their interactions. The pure error was low, indicating good reproducibility of the experimental data.

Statistical analysis by RSM has also been successfully employed by Donzelli et al. [25] to optimise the production of β -(1 \rightarrow 3)-glucanases by *Trichoderma atroviride* in shake-flask cultivation using

Source of variation	Sum of squares	Degrees of freedom	Mean square	<i>F-</i> test	p
Aeration, X ₁ (L)	0.0106	1	0.0106	1.1745	0.4744
Aeration, X ₁ (Q)	0.0007	1	0.0007	0.0805	0.8240
Initial pH, X ₂ (L)	0.1969	1	0.1969	21.8204	0.1342
Initial pH, X ₂ (Q)	3.9728	1	3.9728	440.2251	0.0303
Time of growth, X ₃ (L)	1.6013	1	1.6013	177.4382	0.0477
Time of growth, X ₃ (Q)	0.3674	1	0.3675	40.7178	0.0989
X ₁ X ₂	0.1866	1	0.1866	20.6795	0.1378
X ₁ X ₃	0.0197	1	0.0197	2.1837	0.3787
X ₂ X ₃	0.1935	1	0.1935	21.4390	0.1354
Lack of fit	0.3990	2	0.1995	22.1092	0.1487
Pure Error	0.0090	1	0.0090		
Total	15.5154	12			

L: linear effect; Q: quadratic effect; % of explained variance = 97.37; % of maximum explained variance = 88.479; X₁, X₂, X₃ are variables described in Table 1.

Table 2: ANOVA data for the production of β -1,3-glucanases by Trichoderma harzianum Rifai PAMB-86, and test of significance for the regression coefficients.

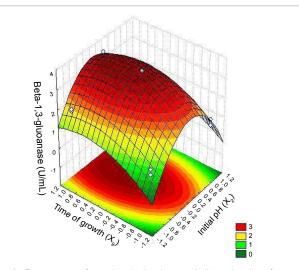


Figure 1: Response surface plot depicted as a 3-dimensional surface and contour showing the effect of β -(1 \rightarrow 3)-glucanase production (Y₁) by *Trichoderma harzianum* Rifai PAMB-86 as a function of initial pH (X₂) and time of growth (X₃), at fixed aeration (X₄ = 1.5 vvm) according to central composite design.

different $(1\rightarrow 3)$ - β -D-glucans as sole carbon sources. According to our experimental data, maximum β - $(1\rightarrow 3)$ -glucanase production (3.7 U/ml) by *T. harzianum* Rifai PAMB-86 occurred within 5 d of growth, an initial pH of 5.5 and aeration of 1.5 vvm. This is a 3-fold improvement over shake flask-grown cultures at the same concentration of botryosphaeran. The 3-dimensional surface and contour plot showing the effect of β - $(1\rightarrow 3)$ -glucanase production as a function of initial pH and time of growth is shown in Figure 1.

There were no significant observed alterations (p < 0.05) between the initial and final pH in the experimental runs. This behavior was also reported by Théodore & Panda [22] with cultures of a *T. harzianum* strain using glucose as sole carbon source, and an initial pH of 4.7 was chosen as the best pH for optimal β -(1 \rightarrow 3)-glucanase production. Donzelli et al. [25] evaluated growth and β -(1 \rightarrow 3)-glucanase production by *T. atroviride* grown on scleroglucan (a (1 \rightarrow 3)(1 \rightarrow 6)- β -D-glucan from *Sclerotium glucanicum*) and glucose as carbon sources, and verified a relationship between β -(1 \rightarrow 3)-glucanase activities and initial pH. An increase in initial pH from 5.5 to 6.5 increased enzyme production 16fold depending on other factors that were applied.

Enzymatic hydrolysis of paramylon and pustulan

ECF containing β -(1 \rightarrow 3)-glucanase activity was concentrated by lyophyllization and the re-solubilised enzyme preparation submitted to gel filtration chromatography. Fractionation revealed the presence of two enzyme fractions (F-I and F-II) hydrolyzing laminarin (a low branched $(1\rightarrow 3)$ - β -D-glucan with $(1\rightarrow 6)$ - β -D-glucosidic linkages, ~5%). F-I contained mainly β -(1 \rightarrow 3)-glucanase activity (0.44 U/ml) as well as β -glucosidase (0.12 U/ml) and β -(1 \rightarrow 6)-glucanase activities (0.24 U/ml), while F-II contained lower β -(1 \rightarrow 3)-glucanase activity (0.30 U/ml), but no β -(1 \rightarrow 6)-glucanase activity. Enzymes F-I and F-II exhibited broad substrate specificity (Table 3; relative activity assigned 100% to laminarin). The enzyme fractions showed no activity towards carboxymethylcellulose and dextran. F-II was not able to hydrolyse lichen pustulan and the oligosaccharides cellotriose and cellotetraose. Both fractions preferred laminaribiose and laminaritriose as substrates to laminarin, while F-I showed highest activity towards cellobiose and gentiobiose.

The β -glucanase complex from *T. harzianum* Rifai PAMB-86 was fractionated in attempts to separate the enzyme components so as to minimize attack on the oligosaccharides liberated during the early stages of hydrolysis of β-D-glucans by enzyme. Earlier studies using a crude enzyme preparation had demonstrated that gluco-oligosaccharides of DP >3 were largely absent from hydrolysates of laminarin and botryosphaeran [10]. To resolve and avoid this situation, we examined each enzyme fraction on its ability to hydrolyze paramylon and pustulan for the production of fragments of DP >3. Enzyme F-I degraded both β -D-glucans differently as judged by the array of hydrolysis products determined by HPAEC/PAD analysis (Table 4), while F-II hydrolysed paramylon in a similar manner to F-I. Glucose was released in relatively large amounts early during hydrolysis by both enzymes attacking paramylon, and the amounts of reducing sugars liberated increased with time. By 30 min ~15-20% of paramylon had been degraded by both enzymes. Under similar conditions, hydrolysis of pustulan by F-I showed that glucose too was rapidly released, and the reducing sugar content in the hydrolysates increased with time, and after 30 min, ~2% of the pustulan was degraded. Enzymatic hydrolysates of paramylon by F-I and F-II contained laminaribiose and laminaritriose, but gentiobiose was absent, confirming that paramylon from Euglena *gracilis* did not carry glucose branches linked through β -(1 \rightarrow 6) bonds. Gluco-oligosaccharides of DP \geq 4 produced from paramylon decreased with time for both enzymes. F-I hydrolysates of pustulan contained gentiobiose and gluco-oligosaccharides of DP \geq 4 that increased with time of hydrolysis (Table 4). F-I thus appears to be an excellent enzyme preparation for the production of gluco-oligosaccharides of higher DP's from pustulan. The absence of laminaribiose and laminaritriose in F-I hydrolysates of pustulan confirmed that the pustulan of the lichen, Actinogyra muehlenbergii, did not carry any β -(1 \rightarrow 3)-linked glucose residues.

Curdlan, a bacterial linear $(1\rightarrow 3)$ - β -D-glucan, was reported to be hydrolyzed by a crude β - $(1\rightarrow 3)$ -glucanase preparation from *T*. *harzianum*, and under the conditions, 80% of this polysaccharide was hydrolyzed [13]. β -Glucanases from *Streptomyces* sp. were also reported to degrade curdlan to mono-, di- and various lower oligosaccharides [15]. The crude β -glucanolytic enzyme preparation from *T. harzianum* Rifai revealed that these enzymes were also able to produce a broad

Carbohydrate		Relative enzyme activity (%)			
substrates	Linkage type	Fraction-I	Fraction-II		
β-D-Glucans	'				
Laminarin	β-(1→3)(1→6)	100	100		
Botryosphaeran	β-(1→3)(1→6)	32	25		
Paramylon	β-(1→3)	27	31		
Pustulan	β-(1→6)	20	0		
CMC ¹	β-(1→4)	0	0		
β-Gluco-oligosacchari	ides				
Laminaribiose	β-(1→3)	140	138		
Laminaritriose	β-(1→3)	133	121		
laminaritetraose	β-(1→3)	98	90		
Gentiobiose	β-(1→6)	124	11		
Cellobiose	β-(1→4)	127	5		
Cellotriose	β-(1→4)	48	0		
Cellotetraose	β-(1→4)	40	0		

Substrates (1 mg/ml) were incubated in 25 mM sodium acetate buffer (pH 4.5) with each enzyme fraction (F-I or F-II; 1 unit of activity) for 10 min at 50°C. Aliquots were next withdrawn and assayed for reducing sugars and/or glucose.¹ carboxymethylcellulose

Table 3: Relative rates of hydrolysis of β -D-glucans and gluco-oligosaccharides by β -glucanase fractions from *Trichoderma harzianum* Rifai PAMB-86.

	β-Glucanase	Time of	Reducing	Hydrolysis products ¹ (Relative amounts, %)				
Substrate	fraction	hydrolysis (min)	sugars (µg/ml)	G,	$\mathbf{G}_{_{2G}}$	$\mathbf{G}_{_{2L}}$	G _{3L}	DP≥4
Paramylon _	Fraction I	5	110 ± 8	82	ND 2	16	0.2	2
		10	140 ± 11	82	ND	15	0.3	3
		20	142 ± 10	77	ND	20	0.3	3
		30	218 ± 6	67	ND	32	0.6	1
	Fraction II	5	82 ± 8	69	ND	22	0.5	9
		10	121 ± 11	73	ND	18	0.3	9
		20	131 ± 12	79	ND	19	ND	2
		30	168 ± 7	75	ND	24	ND	1
Pustulan	Fraction I	5	8 ± 0.5	67	22	ND	ND	11
		10	15 ± 0.9	65	25	ND	ND	10
		20	19 ± 0.9	62	7	ND	ND	31
		30	25 ± 0.8	50	23	ND	ND	27

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Enzymatic hydrolysis of paramylon and pustulan (1 g/l; solubilised by autoclaving) was conducted in 20 ml solution (25 mM sodium acetate buffer, pH 4.5) using 1 unit of each enzyme fraction (F-I, F-II; activity measured against laminarin) at 40°C. ¹G₁, glucose; G_{2G}, gentiobiose; G_{2L}, laminaribiose; G_{3L}, laminaritriose; DP, degree of polymerization; ²ND, not detectable

 Table 4: HPAEC-PAD analysis of enzymatic hydrolysates of paramylon and pustulan using enzyme fractions F-I and F-II from *Trichoderma harzianum* Rifai PAMB-86.

range of products from botryosphaeran in short incubation times [10] with only tetrasaccharides as the highest DP liberated. Grandpierre et al. [26] observed that the rapid release of glucose could inhibit enzyme action on curdlan.

Fungal β -(1 \rightarrow 6)-glucanases are associated with β -(1 \rightarrow 3)-glucanase activities in many microorganisms [16]. Reese et al. [27] observed gentiotetraose, gentiotriose, gentiobiose and glucose as hydrolysis products, and verified that these enzymes acted in an *endo*-glucanase manner. The β -(1 \rightarrow 6)-glucanases from *Acremonium* sp. IMI 383068 showed an improvement of hydrolysis products from pustulan when the incubation time was increased [28]. Wu et al. [29] used a β -(1 \rightarrow 6)-glucanase preparation from *Streptomyces rochei* to hydrolyze pustulan from the lichen, *Umbilicaria papullose*, and observed a range of oligosaccharides formed within 10 min of incubation.

Analysis of the products obtained after incubation of the β -Dglucans, paramylon and pustulan, with β -glucanases from *T. harzianum* Rifai PAMB-86 revealed more information about the specificity of the enzymes produced by this fungus, and this information can be useful in further studies to optimize gluco-oligosaccharide production.

In conclusion, the β -glucanases from *T. harzianum* Rifai PAMB-86 showed maximal enzyme titres at 5 days fermentation, initial pH 5.5 and aeration of 1.5 vvm. This enzyme complex was demonstrated to be useful in the production of gluco-oligosaccharides from algal paramylon and lichen pustulan, which may find applications as prebiotics (nutraceuticals) and modifiers of immunological activities.

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