

Xylanases and cellulases biosynthesis by selected fungi in a simple and economic bio system using sugarcane straw

Tania Sila Campioni*, Ana Flávia de Azevedo Carvalho, Franciane Cristina de Figueiredo, Douglas Fernandes da Silva, Pedro de Oliva Neto.

Associated Laboratory of Bioenergy Research Institute (IPBEN), Bioprocess Unit, São Paulo State University (UNESP), Av. Dom Antonio, 2100, Zip code 19806-380, Assis, SP, Brazil.

Abstract— Sugarcane straw (SS) was used in an economic biosystem to evaluate the production of xylanases and cellulases in submerged fermentation (SmF) by axenic and mixed mode from *Trichoderma* and *Aspergillus* species. *T. reesei* QM9414 axenic culture reached the highest xylanase production (90.2 U/mL) and 0.5 FPU/mL of cellulase activity. The evaluation of agro-industrial residues on fibrolytic enzymes production was performed by a D-optimal design, and revealed the best supplementation of 100% SS, while wheat bran and citric pulp showed lower inductive effects on enzymes production. Also, the scale-up in a stirred tank showed the same yield production profile (xylanase ~ 90 U/mL and cellulase 0.6 FPU/mL). Xylanase was characterized by an optimum pH of 5-6 and temperature at 50 °C, and thermal stability was below 50 °C. The ion Mn²⁺ (5 and 10 mM) had a stimulatory effect on xylanase activity. The biobleaching application showed that 30 U/g of xylanases during 15 min decreased Kappa number in 9.37. These results indicate SS as an alternative substrate for fungi fibrolytic enzymes production and the xylanase with low cellulase extract as a potential biobleaching application.

Keywords— sugarcane straw, xylanase, cellulase, axenic and mixed cultures, fibrolytic enzymes.

I. INTRODUCTION

Due to broadened use of renewable energy sources for biofuels and high-value products production worldwide, including organic wastes mainly produced by agricultural countries, demand for green technologies has increased replacing the extensive usages of fossil fuels (Ferreira-Leitão et al., 2010; Carpio et al., 2019). Sugarcane cultivation is one of the major agricultural activities in Brazil which produced 620.4 million tons in 2018-2019 (Conab, 2019). During the sugarcane burning harvest system, almost 27 kg of carbon dioxide is released into the atmosphere per ton of sugarcane processed, related to burn (40%), fertilizers (20%) and fossil fuels use (18%), thus this quantity can decrease using no-burning system (Figueiredo et al., 2010). The São Paulo state law number 11.241/ 2002 established that, after 2017, 80% of sugarcane, harvesting should be mechanized and after 2021, no more burning will be permitted in mechanized areas. As a consequence of this new system implantation, almost 15 Mg ha⁻¹ dry biomass has been left in the field yearly, mainly SS (sugarcane straw) residue (Hassuani et al., 2005).

Straw represents around one-third of the total primary energy of the sugarcane crop, with a composition very

similar to the widely used bagasse, mainly cellulose, hemicellulose and lignin, 30, 30 and 25%, respectively (Leal et al., 2013). The straw residue in the soil range from positive impacts, such as increase in the macrofauna (mainly worms and ants), nutrients recycling, water storage, carbon accumulation, control of soil erosion and weed infestation, to negative impacts, such as increase in pest populations and biomass loss production (Leal et al., 2013; Carvalho et al., 2017). In fact, a research showed that 50% of SS residue in the soil is necessary to improve the yield of sugarcane crop but the other 50% should be recovered to be used in eco-friendly processes (Aquino et al., 2017). Depending on the amount and characteristics, that residue could be collected to produce energy or co-products such as enzymes (Carvalho et al., 2016; Silva et al., 2018), xylitol (Hernández-Pérez et al 2016), and biodegradable products such as cups, and straws (Gankin, 2019).

In addition, the enzyme technology has continuously replaced the traditional chemical processes in many areas, especially fine chemical and pharmaceutical industries (Choi et al., 2015). The global market for industrial enzymes expects to increase from nearly \$5.5 billion in 2018 to \$7.0 billion in 2023 with a compound annual

growth rate (CAGR) of 4.9% for 2018-2023 (Dewan et al., 2017). The importance of enzyme technology includes the knowledge of fermentation and downstream process, and a high number of available enzymes and applications are developed by the improvement of these technologies (Li et al., 2012). In this sense, the use of agro-industrial residues as carbon source for enzyme biosynthesis by microorganisms, which have potential to decrease the production costs and the final price of enzymes (Salmon et al., 2016; Abdullah et al., 2015). Currently, cellulases represent the third higher industrial enzyme production, and their applications are in cotton, paper recycling, juice extraction, detergent and feed industry (Acharya and Chaudhary, 2012). Other important fibrolytic enzymes are the xylanases, responsible for the hemicellulose hydrolysis. Filamentous fungi produce xylan-degrading enzymes, which is the main interest to industrial purposes due to its low-cost production and the final price of the product as well (Abdullah et al., 2015). Mesophilic fungus as the genera *Aspergillus* and *Trichoderma* have a remarkable importance on xylanases and cellulases improvement production, since they can be cultivated in mixed culture (Ahamed and Vermette 2008; Wen et al., 2005; Dhillon et al., 2011).

Although the efficiency of SS as a feedstock and inducer for cellulase production by some microorganisms were reported to *Streptomyces sp* SLBA-08 (Macedo et al., 2013) and *Trichoderma citrinoviride* (Guerra et al., 2006), in literature there is a lack studies of SS as feedstock for xylanase and cellulase production by *T. reesei*, *Trichoderma harzianum* and *Aspergillus fumigatus* in SmF (submerged fermentation).

In the present study, fibrolytic enzymes production was conducted considering the formulation of the culture medium with SS agro-residue and fungi from *Trichoderma* and *Aspergillus* genera, in axenic and mixed cultures. In addition, the biochemical characterization of the xylanases produced in the best conditions was performed considering biobleaching and future application.

II. MATERIAL AND METHODS

2.1 Microorganisms and substrates

The microorganisms tested in axenic cultures were: *Trichoderma reesei* (Tropical Culture Collection of André Tosello Foundation CCT -2768), *T. reesei* QM9414, *Trichoderma harzianum* N51, *T. harzianum* FS09, *Aspergillus fumigatus* M51 and *A. fumigatus* U2370. These cultures were selected in a previous study as the best producers of fibrolytic enzymes (Carvalho et al., 2015). They were cultured in plates containing 3.9% (w/v) Potato Dextrose Agar (PDA) medium for 7 days at 28 °C and

stored at 4 °C. Lignocellulosic substrates were used as carbon source in the culture medium. The SS was obtained from Água Bonita Mill, Tarumã-SP, Brazil, pretreated (autoclave at 121 °C, 15 min, 1 atm), and milled (14 mesh). The citrus pulp (CP) (from Citrovita, Catanduva-SP, Brazil) was milled (14 mesh), and wheat bran (WB) was used without any previous treatment (from Moinho Nacional, Assis-SP, Brazil).

2.2 Selection of microorganisms in axenic and mixed cultures

The axenic and mixed strains were cultivated in Erlenmeyer flasks (250 mL) by SmF containing 80 mL medium (m/v): 3.0% pretreated SS, 0.1% (NH₄)₂SO₄, 0.0017% MgSO₄·7H₂O, 0.1% K₂HPO₄, 0.0028% ZnSO₄, 0.1% NH₄H₂PO₄, 0.06% KCl, 0.1% yeast extract and 0.1% sucrose at pH 4.5 (Silva et al., 2013). Each fungus spores suspension was prepared by incubating the cultures on PDA plates at 28 °C for about 10 days, until sufficient sporulation was observed. The spores were harvested using 0.1% Tween 80 solution (v/v) for inoculation purposes (about 1x10⁶ cells/mL). Flasks were inoculated and incubated at 28 °C, in an orbital shaker at 180 rpm for 360 h. The biomass was separated by 15 min centrifugation at 4 °C and 2900 x g. The liquid fraction was used as a crude enzymes extract. The binary mixtures of *T. harzianum* FS09 and *A. fumigatus* M51; *T. harzianum* FS09 and *T. reesei* QM9414; *T. reesei* QM9414 and *A. fumigatus* M51; as well as the ternary mixture of *T. harzianum* FS09, *T. reesei* QM9414 and *A. fumigatus* M51; in concentration of spores at 1x10⁶ cells/mL for each one, were combined since they are considered the best xylanase and cellulase producers of this study.

2.3 Formulation of culture medium with mixtures of agro-industrial residues for fibrolytic enzymes production

The SmF of selected microorganism was performed in Erlenmeyer flasks (250 ml, with 80 mL of medium described previously (section 2.2) during 288 h of incubation in a shaker at 28 °C and 180 rpm. D-Optimal mixture design was performed in order to evaluate the effect of individual substrates and the interactions among them in ternary mixtures on xylanase and cellulase production (Fernández-Núñez et al., 2016; Nunes et al., 2017). The number of experimental combinations in each experimental design was enough to fit special cubic models for response variables. The parameters and restrictions of the mixtures were: SS (60–100% w/w range), CP (0–40% w/w range) and WB (0–20% w/w range). A control experiment using 100% (w/v) of each substrate was performed at the same conditions. The D-

optimal experimental design was set up with restrictions and analyzed using Design-Expert software (Design-Expert® software, version 10, Stat-Ease, Inc., Minneapolis, MN, USA). The statistical results were made considering a significance level of 0.05. The strength of linear relationships between actual and predicted values by different models was assessed using the linear correlation coefficient (R^2). The xylanolytic activity in ternary mixtures of agro-industrial residues D-Optimal experimental design was optimized using a desirability function. The optimization criterion was to maximize xylanase activity according to a fitted polynomial for this variable.

2.4 Stirred tank bioreactor culture

The enzyme production by selected microorganism was scaled-up in 2 L BioFlo 115 fermenter (New Brunswick, New Jersey, USA) using medium and inoculation as previously described (section 2.2), working volume of 1.5 L, and Rushton impeller. The culture conditions were 28 °C, 1.7 volume of air per volume of medium per minute (vvm), pH 4.5 for 288 h. Dissolved oxygen was measured by an oxygen electrode and pH was measured and controlled with 1.0% (v/v) H_2SO_4 and 1.0 M NaOH.

2.5 Biochemical characterization of fungal xylanase

The biochemical characterization of xylanases produced from selected microorganism in SmF using selected substrate as described in the following protocols (Carvalho et al., 2006; Carvalho et al., 2010).

2.5.1 Optimum pH and stability

Optimum pH was evaluated by measuring enzyme activity at 50 °C using different buffers: sodium acetate (pH 3.0-6.0), sodium phosphate (pH 6.0-8.0), Tris-HCl (pH 8.0-9.0), and glycine-NaOH (pH 9.0-11.0) and a reaction mixture containing 0.65 mL 0.5% (w/v) xylan in 0.25 M buffer and 0.10 mL crude enzyme. For pH stability, crude enzyme extract was diluted (1:1) in buffers and maintained at 25 °C for 20, 40 and 60 min. An aliquot was used to determine the remaining activity (section 2.6).

2.5.2 Optimum temperature and thermostability

The optimal temperature was determined by incubating the reaction mixture at 20-70 °C (10 min) and assaying the enzyme activity at the optimum pH, in the same reaction mixture (2.6). For thermostability assay, the enzyme solution was incubated at various temperatures (20-70 °C) for 20, 40 and 60 min at pH 5.0 in sealed tubes

to prevent evaporation. The enzyme solution was maintained at these temperatures and times. Aliquots were removed and placed on ice before assaying for residual enzyme activity at optimum pH and temperature.

2.5.3 Effect of ions and EDTA

The effects of ions (Cu^{+2} , Mg^{2+} , Mn^{2+} , Zn^{2+} , Fe^{3+} , Ag^+) and EDTA (Ethylenediamine tetra-acetic acid) on xylanase activity were evaluated. Solutions concentrations of 5 and 10 mM were added to the reaction mixture at the concentration of 0.2% (v/v). The calculation of the percentage of enzyme activity was performed based on the reference sample without addition of any ion.

2.6 Enzymes activity assay

Xylanase activity was assayed at 50 °C in a reaction with 0.1 mL raw enzyme extract and 0.65 mL of 0.5% (m/v) xylan Birchwood solution (Sigma-Aldrich) in 250 mM sodium acetate buffer, at pH 5 for 10 min (Bailey et al., 1993). The reducing sugar concentration was quantified by the dinitrosalicylic acid (DNS) method (Miller, 1960). One unit (U) of xylanase activity was defined as the amount of enzyme to release 1 μ mol of reducing sugar per minute per mL of reaction. The cellulase activity was determined by Ghose (1987). One FPU here is defined as μ moles glucose equivalents released from Whatman n°. 1 per min averaged over 60 min, considering the low enzyme concentration in the raw enzymatic extract.

2.7 Biobleaching

Xylanase from *T. reesei* QM9414 was studied for biobleaching process of Kraft pulp as well as to evaluate its potential use as biobleaching agent. The amount of enzyme used for hydrolysis was 30 units of enzyme per gram of pulp samples. Test conditions were performed in a sealed polyethylene bags with sodium acetate buffer (pH 5.0), at 50 °C for 15 min (soaking stage). Treatment started by diluting the enzyme in the same buffer (pre-heated at 50 °C), adding the solution on pulp samples and then mixed by kneading the bags during 30 s. The final pulp content in the reaction mixture was 3%. Controls were prepared by adding distilled water instead of enzyme. After the enzymatic hydrolysis, the bags were boiled at 100 °C for 5 min to disable the enzymes, cooled and filtered on a Büchner funnel to form paper sheets, used for kappa number analysis.

III. RESULTS AND DISCUSSION

3.1 Selection of fungi for fibrolytic enzymes production in axenic and mixed cultures using SS as a carbon source

3.1.1 Axenic fungal cultures

All tested microorganisms showed xylanases and cellulases production using SS substrate as the sole carbon source in SmF (Fig. 1A and 1B). *T. reesei* QM9414 strain stood out compared to other fungi tested, reaching the highest production of 90 U/mL for xylanase and 0.56 FPU/mL for cellulase at 288 h of fermentation. Nevertheless, the fungi *A. fumigatus* M51 and *A. fumigatus* U2370 also showed good results for xylanases production, approximately 70 U/mL (Fig. 1A). However, after 288 h the enzymes activities decreased, probably due to protease presence in SmF (Silva et al., 2016; Haab et al., 1999). In literature, a higher concentration of xylanase was obtained when compared to 3.38 U/mL at 120 h of cultivation by *Trichoderma inhamatum* (Silva et al., 2015). Also, xylanase activity achieved 43.7 U/mL at 144 h of cultivation by *T. reesei* CCT 2768, 35 U/mL by *A. fumigatus* M51 and 28 U/mL by *A. fumigatus* U2370, using sugarcane bagasse in culture medium (Carvalho et al., 2010).

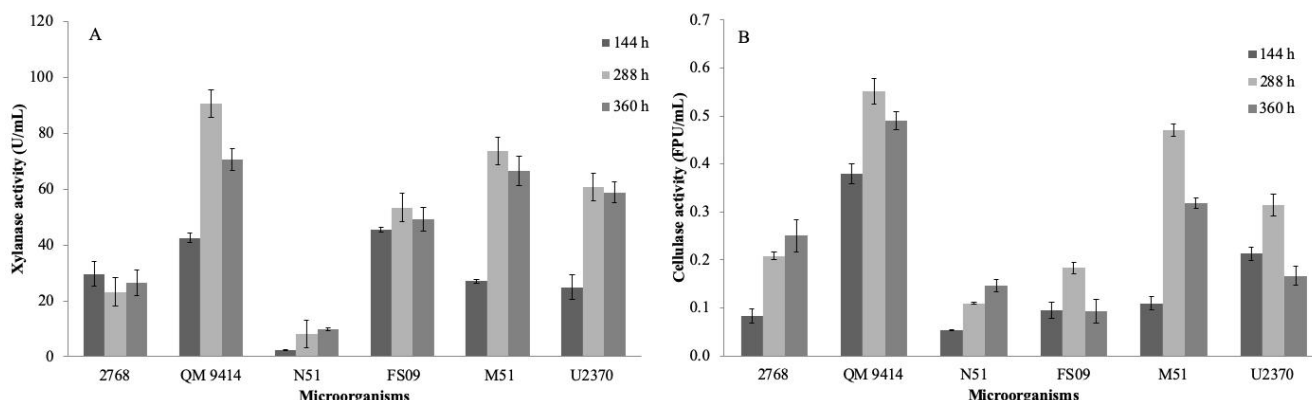


Fig. 1: A) Profile of xylanase production by fungi: *T. reesei* 2768 (2768), *T. reesei* QM9414 (QM9414), *T. harzianum* N51 (N51), *T. harzianum* FS09 (FS09), *A. fumigatus* M51 (M51) and *A. fumigatus* U2370 (U2370), in SmF using SS as substrate (28 °C, pH 4.5, 180 rpm). Each bar value was the average of three replicate experiments, and the error bars show the data ranges. B) Profile of cellulase production by fungi: *T. reesei* 2768 (2768), *T. reesei* QM9414 (QM9414), *T. harzianum* N51 (N51), *T. harzianum* FS09 (FS09), *A. fumigatus* M51 (M51) and *A. fumigatus* U2370 (U2370), in SmF using SS as substrate (28 °C, pH 4.5, 180 rpm). Each bar value was the average of three replicate experiments, and the error bars show the data ranges.

When fibrolytic enzymes biosynthesis from these mixed cultures were compared to axenic culture (Fig. 1-2), the enzyme activities of mixed cultures were lower. However, this result was not expected according to literature (Ahamed and Vermette, 2008; Wen et al., 2005; Dhillon et al., 2011), since the mixed cultures with *Trichoderma* and *Aspergillus* genera resulted in a

The fungi *T. harzianum* FS09, *A. fumigatus* M51 and *T. reesei* QM 9414 were the best cellulases producers, 0.2, 0.4 and 0.6 FPU/mL at 288 h, respectively (Fig. 1B). However, these results obtained to cellulases were lower compared to those found in other studies such as Zhang et al (2014) (0.93 FPU/mL, 96 h) and Xiong et al (2016) (2.33 FPU/mL, 144 h) also produced by *Trichoderma* species, although in these studies were used different substrates as pretreated corn stover and a synthetic medium, respectively. The fact that *T. reesei* QM 9414 produced low cellulases is important for pulp biobleaching application of xylanases for reducing the chlorinated compounds in the paper mills.

3.1.2 Mixed fungal cultures

The mixed fungal and axenic cultures were compared in the present study. Since the *Trichoderma* and *Aspergillus* co-culture system has been reported in literature (Ahamed and Vermette, 2008; Wen et al., 2005), the followed mixtures were proposed: *T. reesei* QM 9414, *A. fumigatus* M51 and *T. harzianum* FS09. Xylanase and cellulase production profile by mixed cultures during 360 h of cultivation were evaluated (Fig. 2A and 2B).

complete enzymatic pool that acts synergistically better in substrate degradation compared to respective axenic culture. According to Duff et al. (1987), fungi species started a substrate competition between them, consequently blocking the enzyme production. The fibrolytic enzymes biosynthesis by *Aspergillus* inhibited the enzymes biosynthesis of *Trichoderma*, probably due

to the catalysis of those enzymes already produced. Proteases or endotoxins biosynthesis could degrade or inhibit the cellulases. In addition, a competition between these microorganisms for the same nutrients in the medium is another hypothesis. The carbon source is reported an important parameter to a successful mixed culture (Dhillon et al., 2011).

Although the results were lower than axenic cultivation for xylanase production, the mixed culture *T. reesei* QM 9414 and *A. fumigatus* M51 reached the maximum value of 60 U/mL (Fig. 2A). On the other hand, it was better than produced by Zhang et al. (2014) (2.5 U/mL), but with another strain (*T. reesei* Rut C-30). These authors also reported a slightly improvement on cellulase production (22.89 - 24.17 U/g) respectively from axenic to mixed cultures, in solid state fermentation (SSF), while the substrate consumption was better in mixed culture. *T. reesei* mutant and *A. niger* in mixed culture resulted in an improvement on enzymes production comparing to single culture by non-mutant strain (Gutierrez-Correa et al., 1999). A synergy in mixed culture of *Trichoderma* and *Aspergillus* was also verified for substrate degradation and consequently a higher

enzyme synthesis (Ahamed and Vermette, 2008). However, the culture of *T. reesei* and *A. phoenicis* ATCC329 xylanase was worse compared to axenic culture in the present study (Wen et al., 2005). Enzymes production by a single culture is preferred to achieve the better substrate degradation from its synergic effect, despite the mixed culture improves cellulases and β -glucosidases production by *T. reesei* QM9414 and *A. terreus* SUK-1 (Wen et al., 2005). In fact, other authors reported the competition by *Trichoderma* and *Aspergillus* to the same nutrients in the medium in a mixed culture (Ahamed and Vermette, 2008; Duff et al., 1987; Anthony et al., 2016). As *T. reesei* showed a great production of xylanases, this strain was selected for the next steps of this work with emphasis for xylanases.

For fibrolytic enzymes production, 3% (m/v) of the substrates SS, CP and WB were evaluated isolated by *T. reesei* QM9414 in SmF medium (Table 1). The culture medium formulated by SS only as substrate showed a higher performance for xylanases biosynthesis (90 U/mL) than other residues. For cellulases production, the cultures of *T. reesei* QM9414 also showed a highest preference for SS (0.6 FPU/mL) (Table 1).

Table 1: Fibrolytic enzymes production by *T. reesei* QM 9414 using agro-industrial residues and its respective chemical composition.

Substrate**	Xylanase activity (U/mL)	Cellulase activity (FPU/mL)	Cellulose (%) w/w	Hemicellulose (%) w/w	Lignin (%) w/w	Reference
Sugarcane Straw	90.6±7.04	0.56±<0.1	33.77	27.38	21.28	Szczerbowski, et al., 2014
Wheat bran	37.7±4.23	<0.10±<0.1	22.3	32	4	Marín et al., 2007
Citrus pulp	31.0±5.87	0.10±<0.1	24.52	7.57	7.51	Rahman et al., 2017

*The results are related with the average and standard deviation of three experiments. **(3% w/v).

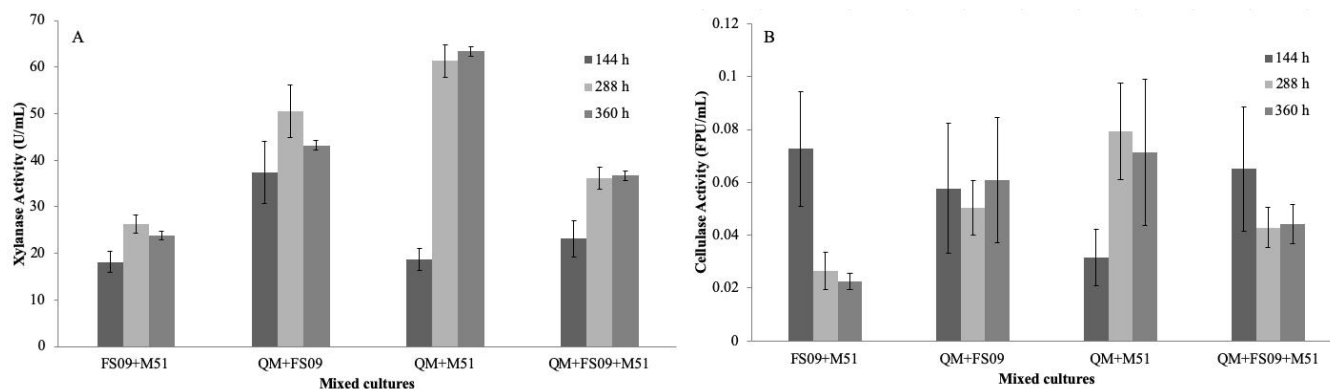


Fig. 2: A) Profile of xylanase production in mixed cultures: *T. harzianum* FS09 + *A. fumigatus* M51 (FS09+M51); *T. harzianum* FS09 + *T. reesei* QM 9414 (QM+FS09); *T. reesei* QM 9414+ *A. fumigatus* M51 (QM+M51); *T. harzianum* FS09 + *T. reesei* QM 9414+ *A. fumigatus* M51 (QM+FS09+M51), in SmF using SS as substrate (28 °C, pH 4.5, 180 rpm). B) Profile of cellulase production in mixed cultures: *T. harzianum* FS09 + *A. fumigatus* M51 (FS09+M51); *T. harzianum* FS09 + *T. reesei* QM 9414 (QM+FS09); *T. reesei* QM 9414+ *A. fumigatus* M51 (QM+M51); *T. harzianum* FS09 + *T. reesei* QM 9414+ *A. fumigatus* M51 (QM+FS09+M51), in SmF using SS as substrate (28 °C, pH 4.5, 180 rpm). Each bar value was the average of three replicate experiments, and the error bars show the data ranges.

3.2 The effect of the mixture of agro-industrial residues in formulated media for fibrolytic enzyme production by *T. reesei* QM 9414

The use of WB as substrate was proposed since in literature was observed higher xylanase production in SSF culture (Dhillon et al., 2011; Guimarães et al., 2013) for this residue. The substrates compositions (Table 1) suggest that CP and WB should be more easily hydrolyzed due to their low lignin content. In addition, this fact is responsible for a better xylanases production in SS, since SS residue has high level of lignin makes the degradation of the fiber more difficult and it demands more fibrolytic enzymes.

In the second set of experiments, a D-Optimal mixture experimental design was used to evaluate the synergistic or antagonistic effects of the mixed carbon sources in SmF to produce fibrolytic enzymes by *T. reesei* QM 9414 in 12 days (Table 2). When xylanase and cellulase activities were evaluated, for ternary mixtures of these substrates, were modeled in D-optimal design, cubic models were satisfactorily fitted to the experimental data (model significance tests, $p < 0.05$ and lack of fit tests, $p > 0.05$).

$$\begin{aligned} \text{Xylanase activity (U/mL)} &= 89.18*A+80.18*B+1408.6*C-3.97*AB-2716.56*AC- \\ &2693.27*BC+3926.94*ABC+269.98*AB(A- \\ &B)+1683.22*AC(A-C)+1.798*BC(B-C) \text{ Eq. (1)} \\ \text{Cellulase activity (U/mL)} &= 0.52*A+0.43*B+10.87*C- \\ &0.11*AB-21.73*AC- \\ &21.4734*BC+30.88*ABC+1.88*AB(A-B)+11.89*AC(A- \\ &C)+13.45*BC(B-C). \text{ Eq.(2)} \end{aligned}$$

The equations for xylanase and cellulase activities (Equations 1-2 for actual values) in conjunction with contour Graphs (Fig. 3A and 3B) showed the major contribution of SS for higher values of fibrolytic enzymes activities.

The SS influence on xylanase activity was noticed that activity increased with higher substrate concentration, while for CP residue a slight increment on xylanase activity was observed. The substrate WB was not interesting for this purpose since the results were not satisfactory.

Table 2: Results derived from D-optimal experimental design for ternary mixtures of SS, CP and WB as carbon sources in SmF by *T. reesei* QM9414 (pH 4.5, 28 °C, 288 h).

Experiment	Sugarcane Straw (% m/m)	Citrus Pulp (% m/m)	Wheat Bran (% m/m)	Xylanase Activity (U/mL)	Cellulase Activity (FPU/mL)
1	80.0	0.0	20.0	69.4	0.3
2	75.0	15.0	10.0	83.9	0.4
3	60.0	20.0	20.0	70.0	0.3

4	60.0	40.0	0.0	81.8	0.4
5	60.0	20.0	20.0	71.8	0.3
6	90.0	0.0	10.0	67.3	0.2
7	82.5	7.5	10.0	93.3	0.4
8	66.67	20.0	13.33	61.2	0.2
9	60.0	40.0	0.0	78.4	0.4
10 (C)	100.0	0.0	0.0	90.2	0.5
11 (C)	100.0	0.0	0.0	88.1	0.5
12	80.0	20.0	0.0	83.9	0.4
13	70.0	10.0	20.0	84.7	0.4
14	67.5	27.5	5.0	88.2	0.4
15	80.0	20.0	0.0	83.5	0.5

The math models are expressed in Eq. 1-2, with coded variables showing the enzymatic activities as function of: A = SS (w/w), B = CP (w/w), and C = WB (w/w). According to ANOVA, each activity response desired, xylanase and cellulase activities produced were statistically significant ($p < 0.05$), respectively, for the cubic math models with high Regression coefficient ($R^2_{adj} = 0.95, 0.93$).

Regarding the cellulase production, SS in a relatively higher concentration presented great activities. However, WB did not represent any synergic effect with other substrates. CP presented a positive effect on cellulase activity within the range interactions. On the other hand, these results are in disagreement with some authors that found an improvement on enzymes production in optimization studies of mixed substrates. Das et al. (2013) showed cellulase production increased 1.3-fold after the medium optimization, containing WB and rice straw by *A. fumigatus* ABK9. WB also performed a positive effect (21%) in the xylanase production by *A. flavus* (Guimarães et al., 2013).

Considering the final purpose of the use of crude enzymatic extract rich in xylanases and poor in cellulases, which are an important characteristic for biobleaching of kraft pulp (Guimarães et al., 2013; Nagar et al., 2010), the optimization of parameters was adjusted to reach a maximum of xylanases and low cellulases production. The optimal set of factors to maximize xylanase production by *T. reesei* was 100% SS, which the experiment 10 reached 90.2 U/mL (Table 2). The most significant results were achieved with 100% SS with desirability predicted for the model was 0.92. The result was validated (in triplicate) in the same conditions (100% SS). The predicted result from the desirability function was 89.2 U/mL and the result obtained, 90.2 U/mL,

presented no significant difference (Anova+Tukey, $p > 0.05$). The crude enzymes extract under this condition was rich in xylanases and poor in cellulases, a ratio of 1:0.005 U/mL, respectively.

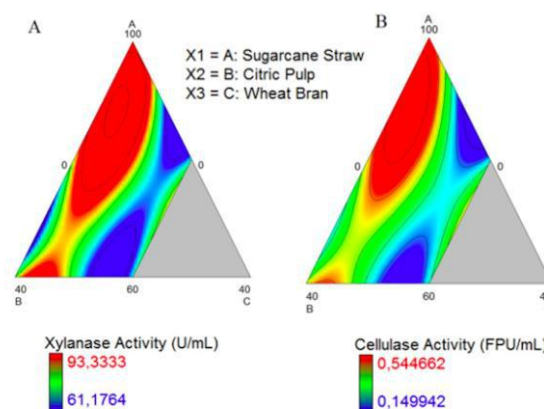


Fig. 3: Contour plots of responses generated by the interactions of the A = SS (w/w); B = CP (w/w); C = WB (w/w), on fibrolytic activities. A) Xylanase activity and B) Cellulase activity produced in SmF by *T. reesei* QM9414 using SS, CP and WB as substrates (28 °C, pH 4.5, 180 rpm).

Regarding the xylanase application on kraft pulp biobleaching, Campioni et al. (2019) studied xylanase extract produced by *T. reesei* QM9414 in SmF with SS and optimized the biobleaching parameters. The best conditions were 30 U/g of xylanase, at pH 5, at 50 °C during 30 min and resulted a 12.5% of Kappa number reduction. After the xylanase biobleaching, the final chlorine dioxide consumption reduced to 10%, maintaining the same brightness compared to control on the subsequent chemical process. In addition, an important parameter for biobleaching application is the xylanase combined with low cellulase concentration or

even no cellulase activity, otherwise, higher amount of this enzyme could degrade the pulp.

It is known about the successful application of enzymes depends not only on the substrate choice but a simple bioprocess and mainly a low-cost production as well. As mentioned previously (section 1), regarding the transition of no sugarcane burning on harvest system (São Paulo State No. 11.241/2002), the SS residue has been left large amounts on fields, which influenced the dynamics of sugarcane production in several aspects (Carvalho et al., 2017). Additionally, SS has been considered a low-cost residue, which the average of value of US \$9.38/ton (Carpio et al., 2019). In this sense, several lignocellulosic agro-industrial residues have been widely evaluated as substrate for xylanase production, such as sugarcane bagasse, WB, sawdust, soy flour, maize straw and others (Knob et al., 2013). Although the use of agro-industrial residues has been extensively described in literature, there is the concern about multiple and complex process steps, consequently become more expensive and difficult to scale up. For example, the substrate pretreatment procedures, waste of extensive washing with distilled water (Knob et al., 2014), chemical pretreatments and in some cases they can generate other toxic compounds for microorganisms and become difficult to find an appropriate destination (Robl et al., 2015). Therefore, this study is a cost effective and simple using SS as a potential substrate for fibrolytic production by *T. reesei* QM9414 and its biobleaching application. After the selection of microorganism and agro-industrial residue used as carbon sources, the enzymatic production was scaled up in bioreactor using 1.5 L working volume and controlled conditions, resulting in 88.02 ± 4.54 U/mL and 0.41 ± 0.1 FPU/mL, for xylanase and cellulase respectively, proving a high level of xylanase production using 100% of SS by *T. reesei* QM9414 can be obtained by this simple and economical bioprocess. On the other hand, the enzyme production losses were detected in scaling-up of *T. harzianum* P49P11 in SmF using sugarcane bagasse in stirred tank bioreactor (Haab et al., 1990).

3.3 Xylanases biochemical characterization

The enzymatic extract produced by *T. reesei* QM9414 cultivated in SS medium (12 culture days)

showed the highest xylanase activity at pH 5 (100 U/mL) (Fig. 4A). The lower range (pH 3-4) and basic pH (pH 8-11) strongly decreased the enzymatic activity. In spite of this, when basic pH was performed the Tris-HCl buffer was chosen than sodium phosphate due to higher enzyme activity in the same pH 8, respectively 65 and 20 U/mL. Xylanase residual activities linearly decreased after the incubation time (20, 40 and 60 min) for all pH ranges (Fig. 4B). The loss of activity varied from 20-95% compared to control, and a higher loss was at pH 8, after 60 min of incubation. In the range of pH 5-6, the enzyme remained 80% active after all incubation times. Xylanases from other *Trichoderma* species was also found in literature with optimum pH 5-6, but with broader pH ranges (Table 3).

Considering pH close to 5.0 as xylanase optimum pH, some applications were found in literature. Zhang et al (2014) proposed the use of xylanases as an additive in bird feed, due to pH range used in this feed was 5.5-6.5. Other sectors are possible such as juice mills (Nagar et al., 2010) and bioethanol (Ferreira-Leitão et al., 2010; Carpio et al., 2019).

In this study, xylanase *T. reesei* QM9414 optimum activity was observed at 50 °C (Fig. 5A). This temperature is commonly reported by *Trichoderma* SC9 and *T. inhamatum* (Tab. 3), beyond microorganisms from other genera: *Paenibacillus macquariensis* (Terrasan et al., 2013) and *Penicillium janczewskii* (Jänis et al., 2001).

The Fig. 5B depicts the thermostability. In temperatures of 20-30 °C, and after 20, 40 and 60 min, xylanase retained almost 80% of its activity. On the other hand, in temperatures higher than 50 °C, a linear decrease in enzymatic activity was observed, except at the point at 50 °C for 20 min, which the activity just improved slightly and then decreased again. Xylanase produced by *T. reesei* QM9414 showed optimum at 50 °C temperature of incubation. The low thermostability of xylanase by other species of *Trichoderma* was also observed in literature.

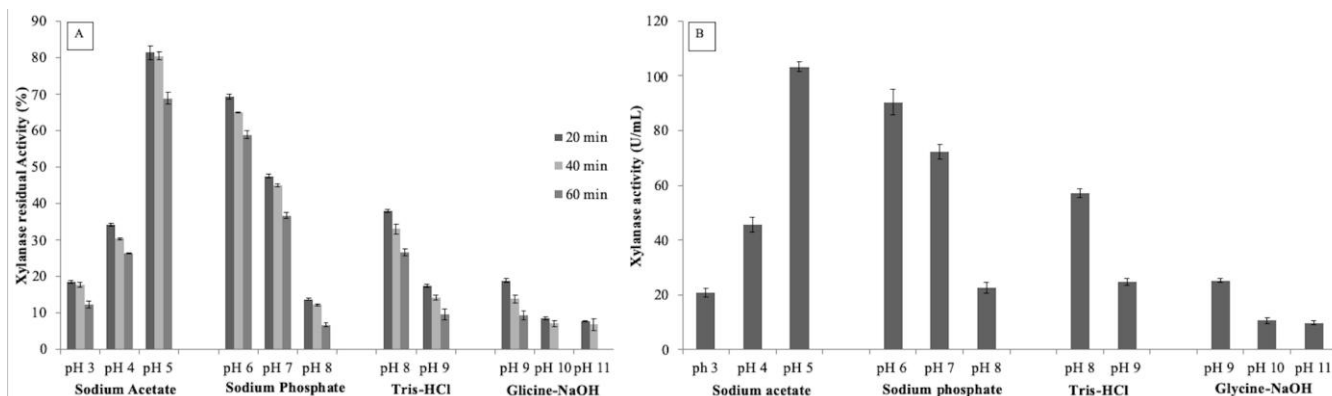


Fig. 4: A) Effect of pH on xylanase activity of the crude extract produced by *T. reesei* QM9414 cultivated with SS (pH 4.5, 28 °C, 288 h). Each bar value was the average of three replicate experiments, and the error bars show the data ranges. B) Effect of pH on xylanase activity stability of the crude extract produced by *T. reesei* QM9414 cultivated with SS (pH 4.5, 28 °C, 288 h). Each bar value was the average of three replicate experiments, and the error bars show the data ranges.

Table 3: Comparative xylanase characteristics produced by different *Trichoderma* species in literature.

Microorganism	Optimum pH	Stability range pH	Optimum temperature (°C)	Reference
<i>T. reesei</i> QM9414	5.0	5.0-6.0	50	This work
<i>T. inhamatum</i>	Xyl I: 5-5.5 Xyl II: 5	Xyl I: 4.5-6.5 Xyl II: 5.0	50 (both)	Silva et al., 2015
<i>Trichoderma</i> sp SC9	6.0	3.5-9.0	42.5	Zhou et al., 2011
<i>T. harzianum</i> 1073 D3	5.0	3.0-7.0	60	Isil and Nilufer, 2005
<i>T. reesei</i>	6.0	3.0-8.0	-	He et al., 2009

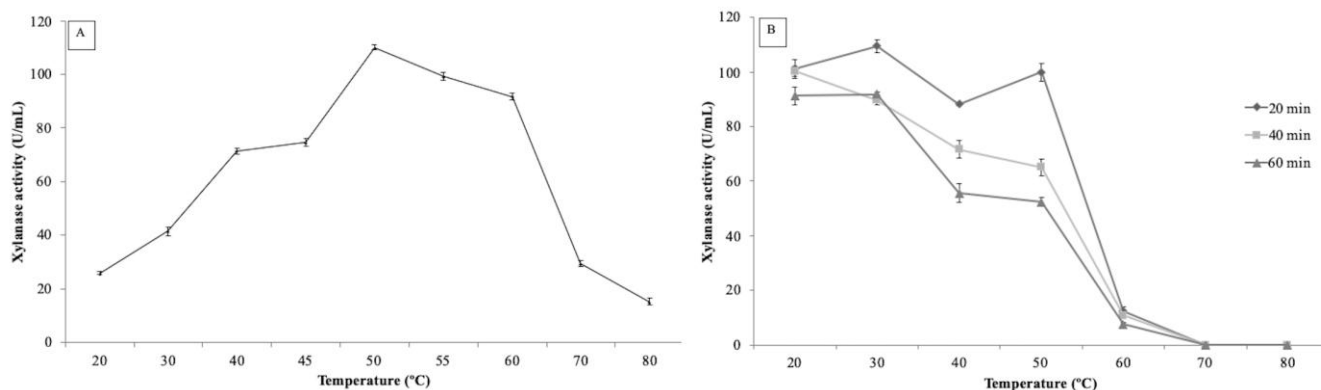


Fig. 5: A) Effect of temperature on xylanase activity produced from *T. reesei* QM9414. B) Thermostability of xylanase produced by *T. reesei* QM9414 (pH 4.5, 28 °C, 288 h). Each bar was the average of three replicate experiments, and the error bars show the data ranges.

The thermostability of *T. inhamatum* xylanase presented a half-life of 2.2 h at 40 °C, and subsequently when the temperature reached 50 °C this time dropped drastically to 2 min (Silva et al., 2015). Another work showed the stability of *T. reesei* RUT C-30 xylanase was 94% at 50 °C after 30 min of incubation (He et al., 2009). The thermostability loss of xylanase from *Trichoderma* genus in temperatures higher than 50 °C can be explained

by a conformational structure change (López and Estrada, 2014), as well as the loss of secondary structure at 58.8 °C and tertiary one in 56.3 °C, reflecting in decrease of activity (Cobos and Estrada, 2003). Some additives in xylanases can be applied to solve the thermostability loss, such as polyhydroxylic co-solvents addiction (Xiong et al., 2004) and mutations in bisulfide bounds (Blanco et al., 1995). The effect of activation or inhibition of ions

and EDTA on xylanases activities were evaluated and considering two ions solution concentrations, 5 and 10 mM. When the Cu^{2+} , Mg^{2+} , Mn^{2+} and Zn^{2+} ions were added, there was an increment on the enzymatic activity (Table 4). The most expressive result was the Mn^{2+} , 39 and 49%, for the respective concentrations. In contrast, 10 mM of ions Cu^{2+} and Ag^{+} resulted in a strong inhibition of xylanase, 21 and 18% respectively. In literature, the presence of Mn^{2+} and Zn^{2+} also increased xylanase activity produced by *T. harzianum* 1073 D3, whereas in the presence of Mg^{2+} and Cu^{2+} the activity was not affected (Isil and Nilufer, 2005). According to Blanco et al. (1995) Mn^{2+} and Cu^{2+} did not affect the xylanase activity, while Mg^{2+} had a stimulatory effect. In addition, Mn^{2+} also stimulated the enzymatic activity for xylanases from *Paenibacillus macquariensis* (Terrasan et al., 2013). In this last work Cu^{2+} and Fe^{3+} caused inhibition on the enzymatic activity, whereas Mn^{2+} and Mg^{2+} presented no difference compared to control. EDTA caused a slightly decreased on the xylanase activity at concentrations of 5 and 10 mM, 10 and 0.8%, respectively (Table 4). The explanation of the authors for this fact was that an enzyme needs divalent ions for catalysis. In other works, EDTA caused inhibition of the enzymatic activity of xylanases in the concentrations of 1, 2 and 10 mM (Silva et al., 2008).

3.4 Biobleaching

In order to evaluate the xylanase efficiency for cellulose pulp biobleaching, the pulp was clarified by *T. reesei* QM9414 crude extract and 30 Units of xylanase per gram of pulp in 15 min was successfully effective compared to controls. Xylanase reduced the kappa number in 9.37% (2.1 kappa points). In literature, xylanase produced by *A. caespitosus* reduced kappa number only in 1.7% (xyl I), and the conditions were 10 U/g dry pulp in 2 hours (Sandrim et al., 2005).

Table 4: Ions and EDTA effect on xylanase activity produced by *T. reesei* QM9414 in SS medium.

	Xylanase Activity (%)	
	5 mM	10 mM
Cu^{2+}	106.2	78.7
Mg^{2+}	106.5	108.7
Mn^{2+}	138.8	148.7
Zn^{2+}	104.4	111.9
Fe^{3+}	89.8	93.6
Ag^{1+}	98.9	82.0
EDTA	89.9	99.2

IV. CONCLUSIONS

Sugarcane straw was evaluated as the main carbon source in axenic SmF of *T. reesei* QM 9414 to produce fibrolytic enzymes in a simple and economical bioprocess. Also, the xylanase production was successfully scaled-up from shaker flasks to bioreactor, maintaining the same culture conditions, without loss of enzyme production. This enzyme was characterized, accordingly interesting conditions for some industrial applications, mainly potential on biobleaching of kraft pulp proposes.

ACKNOWLEDGEMENTS

The authors wish to thank Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (Capes), Brazil, and Função de Amparo à Pesquisa (Fapesp - 2014/24188-1), São Paulo, Brazil, for financial support.

REFERENCES

- [1] Abdullah R., Nisar K., Aslam A., Iqtedar M. & Naz S. (2015). Enhanced production of xylanase from locally isolated fungal strain using agro-industrial residues under solid-state fermentation. *Nat Prod Res.* 29, 1006-1011.
- [2] Acharya S. & Chaudhary A. (2012). Bioprospecting thermophiles for cellulase production: a review. *Braz. J. of Microbiol.* 43, 844-856.
- [3] Ahamed A. & Vermette P. (2008). Enhanced enzyme production from mixed cultures of *Trichoderma reesei* RUT-C30 and *Aspergillus niger* LMA grown as fed batch in a stirred tank bioreactor. *Biochem. Eng. J.* 42, 41-46.
- [4] Anthony P., Harish B.S., Jampala P., Ramanujam S. & Uppuluri K.B. (2016). Statistical optimization, purification and applications of xylanase produced from mixed bacteria in a solid liquid fermentation using *Prosopis juliflora*. *Biocatal. Agric. Biotechnol.* 8, 213-220.
- [5] Aquino G.S., Conti Medina C., Costa D.C., Shahab M. & Santiago A.D. (2017). Sugarcane straw management and its impact on production and development of ratoons. (Supplement C), *Ind Crops Prod.* 102, 58-64.
- [6] Bailey M.J., Buchert J. & Viikari L. (1993). Effect of pH on production of xylanase by *Trichoderma reesei* on xylan- and cellulose-based media. *Appl Microbiol Biotechnol.* 40, 224-229.
- [7] Blanco A., Vidal T., Colom J.F. & Pastor F.I. (1995). Purification and properties of xylanase A from alkali-tolerant *Bacillus* sp. strain BP-23. *Appl Environ Microbiol.* 61, 4468-4470.
- [8] Campioni T.S., Moreira L.J., Moretto E., Nunes N.S.S. & Oliva-Neto P. (2019). Biobleaching of Kraft pulp using fungal xylanases produced from sugarcane straw and the subsequent decrease of chlorine consumption. *Biomass and Bioenergy*, 121, 22-27.
- [9] Carpio L.G.T. & Souza F.S. (2019). Competition between Second-Generation Ethanol and Bioelectricity using the

- Residual Biomass of Sugarcane: Effects of Uncertainty on the Production Mix. *Molecules*, 24, 369.
- [10] Carvalho A.F.A., Boscolo M., Silva R., Ferreira H. & Gomes E. (2010). Purification and characterization of the α -glucosidase produced by thermophilic fungus *Thermoascus aurantiacus* CBMAI 756. *J Microbiol.* 48, 452-459.
- [11] Carvalho A.F.A., Gonçalves A.Z., Silva R. & Gomes E. (2006). A specific short dextrin-hydrolyzing extracellular glucosidase from the thermophilic fungus *Thermoascus aurantiacus* 179-5. *J Microbiol.* 44, 276-283.
- [12] Carvalho A.F.A., Neto P.O., Almeida P.Z., Silva J.B., Escaramboni B. & Pastore G.M. (2015). Screening of xylanolytic *Aspergillus fumigatus* for prebiotic xylooligosaccharide production using bagasse. *Food Technol Biotechnol.* 53, 428-435.
- [13] Carvalho D.M., Queiroz J.H. & Colodette J.L. (2016). Assessment of alkaline pretreatment for the production of bioethanol from eucalyptus, sugarcane bagasse and sugarcane straw. *Ind Crops Prod.* 94, 932-941.
- [14] Carvalho J.L.N., Nogueiro R.C., Menandro L.M.S., et al. (2017). Agronomic and environmental implications of sugarcane straw removal: a major review. *GCB Bioenergy* 9, 1181-1195.
- [15] Choi J-M., Han S-S., Kim H-S. (2015). Industrial applications of enzyme biocatalysis: Current status and future aspects. *Biotechnol Adv.* 33, 1443-1454.
- [16] Cobos A. & Estrada P. (2003). Effect of polyhydroxylic cosolvents on the thermostability and activity of xylanase from *Trichoderma reesei* QM 9414. *Enzyme Microb Tech.* 33, 810-818.
- [17] Conab, Companhia Nacional de Abastecimento. 2019. Acompanhamento da safra brasileira de cana-de-açúcar. *Conab Safra 2018/19* 4, 1-75.
- [18] Das A., Paul T., Halder S.K., et al. (2013). Production of cellulolytic enzymes by *Aspergillus fumigatus* ABK9 in wheat bran-rice straw mixed substrate and use of cocktail enzymes for deinking of waste office paper pulp. *Bioresour Technol.* 128, 290-296.
- [19] Dewan S.S., 2018. (2019, October 02). Global Markets for Enzymes in Industrial Applications. Retrieved from: <https://www.bccresearch.com/marketresearch/biotechnology/global-markets-for-enzymes-in-industrial-applications.html>.
- [20] Dhillon G.S., Oberoi H.S., Kaur S., Bansal S. & Brar S.K. (2011). Value-addition of agricultural wastes for augmented cellulase and xylanase production through solid-state tray fermentation employing mixed-culture of fungi. *Ind Crops and Prod.* 34, 1160-1167.
- [21] Duff S.J.B., Cooper D.G. & Fuller O.M. (1987). Effect of media composition and growth conditions on production of cellulase and β -glucosidase by a mixed fungal fermentation. *Enzyme Microb Technol.* 9, 47-52.
- [22] Fernández-Núñez E.G., Barchi A.C., Ito S., et al. (2016). Artificial intelligence approach for high level production of amylase using *Rhizopus microsporus* var. oligosporus and different agro-industrial wastes. *J. Chem. Technol. Biot.* 92, 684-692.
- [23] Ferreira-Leitão V., Perrone C.C., Rodrigues J., Franke A.P.M., Macrelli S. & Zacchi G. (2010). An approach to the utilisation of CO₂ as impregnating agent in steam pretreatment of sugar cane bagasse and leaves for ethanol production, *Biotechnol. Biofuels* 7, 2-8.
- [24] Figueiredo E.B., Panosso A.R., Romão R. & La Scala N. (2010). Greenhouse gas emission associated with sugar production in southern Brazil. *Carbon Balance and Manag* 5, 3.
- Gankin E, (2019, October 18). Sustainability News. Retrieved from: <https://www.biofutura.com/en/materials/sugarcane>.
- [25] Ghose T.K. (1987). Measurement of cellulase activities. *Pure Appl Chem.* 59, 257-268.
- [26] Guerra G., Casado MR-LG., Arguelles J., Sánchez M.I., Manzano A.M. & Guzman T. (2006). Cellulase production with sugarcane straw by *Trichoderma citrinoviride* on solid bed. *Sugar Tech.* 8, 30-35.
- [27] Guimarães N.C.A., Sorgatto M., Peixoto-Nogueira S.C., et al. (2013). Bioprocess and biotechnology: effect of xylanase from *Aspergillus niger* and *Aspergillus flavus* on pulp biobleaching and enzyme production using agroindustrial residues as substrate. *Springer Plus* 2, 380.
- [28] Gutierrez-Correa M., Portal L., Moreno P., Tengerdy R.P. (1999). Mixed culture solid substrate fermentation of *Trichoderma reesei* with *Aspergillus niger* on sugar cane bagasse. *Bioresour Technol.* 68, 173-178.
- [29] Haab D., Hagspiel K., Szakmary K. & Kubicek C.P. (1990). Formation of the extracellular proteases from *Trichoderma reesei* QM 9414 involved in cellulase degradation. *J Biotechnol.* 16, 187-198.
- [30] Haab D., Hagspiel K., Szakmary K. & Kubicek C.P. (1990). Formation of the extracellular proteases from *Trichoderma reesei* QM 9414 involved in cellulase degradation. *J Biotechnol.* 16, 187-198.
- [31] Hassuani S.J., Leal M.R.L.V. & Macedo I.C. (2005). Biomass power generation: Sugar cane bagasse and trash. In: Série Caminhos para Sustentabilidade Piracicaba: PNUD-CTC, p.19-23.
- [31] He J., Yu B., Fau - Zhang K., Zhang K., Fau - Ding X., Ding X., Fau - Chen D. & Chen D. (2009). Expression of endo-1, 4-beta-xylanase from *Trichoderma reesei* in *Pichia pastoris* and functional characterization of the produced enzyme. *BMC Biotechnology* 16, 56-64.
- [32] Hernández-Pérez A.F., Costa I.A.L., Silva D.D.V. et al. (2016). Biochemical conversion of sugarcane straw hemicellulosic hydrolyzate supplemented with co-substrates for xylitol production. *Bioresour Technol.* 200, 1085-1088.
- [33] Isil S., Nilufer A. 2005. Investigation of factors affecting xylanase activity from *Trichoderma harzianum* 1073 D3. *Braz Arch Biol Technol.* 48, 187-193.
- [34] Jänis J., Rouvinen J., Leisola M., Turunen O. & Vainiotalo P. (2001). Thermostability of endo-1,4- β -xylanase II from *Trichoderma reesei* studied by electrospray ionization Fourier-transform ion cyclotron resonance MS,

- hydrogen/deuterium-exchange reactions and dynamic light scattering. *Biochem J.* 356, 453.
- [35] Knob A., Beitel S.M., Fortkamp D., Terrasan C.R.F., Almeida A.F. 2013. Production, purification, and characterization of a major *Penicillium glabrum* xylanase using Brewer's spent grain as substrate. *BioMed Research International* 728735, 8.
- [36] Knob A., Fortkamp D., Prolo T., Izidoro S.C., Almeida J.M. 2014. Agro-residues as alternative for xylanase production by filamentous fungi. *BioResources* 9, 5738-5773.
- [37] Leal M.R.L.V., Galdos M.V., Scarpore F.V., Seabra J.E.A., Walter A., Oliveira C.O.F. (2013). Sugarcane straw availability, quality, recovery and energy use: A literature review. *Biomass and Bioenergy* 53, 11-19.
- [38] Li S., Yang X., Yang S., Zhu M., Wang X. 2012. Technology prospecting on enzymes: application, marketing and engineering. *Comput. Struct. Biotech. J.* 2, 1-11.
- [39] Lopéz G. & Estrada P. (2014). Effect of temperature on xylanase II from *Trichoderma reesei* QM 9414: A calorimetric, catalytic, and conformational study. *Enzyme Res.* 2014, 16.
- [40] Macedo E.P., Cerqueira C.L.O., Souza D.A.J., Bispo A.S.R., Coelho R.R.R. & Nascimento R.P. (2013). Production of cellulose-degrading enzyme on sisal and other agro-industrial residues using a new Brazilian actinobacteria strain *Streptomyces* sp. SLBA-08. *Braz J of Chem Eng.* 30, 729-735.
- [41] Marín, F.R. et al. (2007). By-products from different citrus processes as a source of customized functional fibres. *Food Chem*, 100, 736-741.
- [42] Miller G.L., Blum R., Glennon W.E. & Burton A.L. (1960). Measurement of carboxymethylcellulase activity. *Anal Biochem.* 2.
- [43] Nagar S., Gupta V.K., Kumar D., Kumar L., Kuhad R.C. (2010). Production and optimization of cellulase-free, alkali-stable xylanase by *Bacillus pumilus* SV-85S in submerged fermentation. *J Ind Microbiol Biotechnol.* 37, 71-83.
- [44] Nagar S., Mittal A., Gupta V.K. 2012. Enzymatic clarification of fruit juices (Apple, Pineapple, and Tomato) using purified *Bacillus pumilus* SV-85S xylanase. *Biotechnol Bioproc E.* 17, 1165-1175.
- [45] Nunes L.V., Correa F.F.B., Oliva-Neto P., et al. (2017). Lactic acid production from submerged fermentation of broken rice using undefined mixed culture. *World J Microbiol Biotechnol.* 33, 79.
- [46] Rahman, A. et al. (2017). Pretreatment of Wheat Bran for Suitable Reinforcement in Biocomposites. *J. Renew. Mater.* 5, n. Suppl 1, 62-73.
- [47] Robl D., Delabona P.S., Costa P.S., et al. (2015). Xylanase production by endophytic *Aspergillus niger* using pentose-rich hydrothermal liquor from sugarcane bagasse. *Biocatal Biotransfor.* 33, 175-187.
- [48] Salmon, D.N.X., Fendrich, R.C., Cruz, M.A., Montibeller, V.W., Vandenberghe, L.P.S., Soccol C.R. & Spier, M.R. (2016). Bioprocess for phytase production by *Ganoderma* sp. MR-56 in different types of bioreactors through submerged cultivation. *Biochem. Eng. J.* 114, 288-297.
- [49] Sandrim V.C., Rizzatti A.C.S., Terenzi H.F., Jorge J.A., Milagres A.M.F. & Polizeli M.L.T.M. (2005). Purification and biochemical characterization of two xylanases produced by *Aspergillus caespitosus* and their potential for kraft pulp bleaching. *Process Biochem.* 40, 1823-1828.
- [50] Sharma M., Mehta S. & Kumar A. (2013). Purification and characterization of alkaline xylanase secreted from *Paenibacillus macquariensis*. *Adv Microbiol.* 3, 32-41.
- [51] Silva D.F., Camargo R., Carvalho A.F.A. & Oliva-Neto P. (2013). Cellulolytic enzyme production by the fungi *Trichoderma reesei* QM9414 and *Trichoderma reesei* CCT2768 using citrus pulp. *Poster presented at the 6th Symposium of Applied Microbiology* Rio Claro, SP, Brazil.
- [52] Silva D.F., Hergesel L.M., Campioni T.S., Carvalho A.F.A. & Oliva-Neto P. (2018). Evaluation of different biological and chemical treatments in agroindustrial residues for the production of fungal glucanases and xylanases. *Process Biochem.* 67, 29-37.
- [53] Silva L.A.O., Carmona E.C. [2008]. Production and characterization of cellulase-free xylanase from *Trichoderma inhamatum*. *Appl Biochem Biotechnol.* 150, 117-125.
- [54] Silva, L.A.O., Terrasan C.R.F., Carmona E.C. (2015). Purification and characterization of xylanases from *Trichoderma inhamatum*. *Electron J Biotechnol.* 18, 307-313.
- [55] Szczerbowski, D. et al. (2014). Sugarcane biomass for biorefineries: Comparative composition of carbohydrate and non-carbohydrate components of bagasse and straw. *Carbohydr. Polym.* 114, 95-101.
- [56] Terrasan C.R.F., Temer B., Sarto C., Silva Júnior F.G., Carmona E.C. (2013). Xylanase and β -xylosidase from *Penicillium janczewskii*: production, physico-chemical properties, and application of the crude extract to pulp biobleaching. *BioResources* 8, 1292-1305.
- [57] Wen Z., Liao W. & Chen S. Production of cellulase/ β -glucosidase by the mixed fungi culture of *Trichoderma reesei* and *Aspergillus phoenicis* on dairy manure. 2005. In: Davison B.H., Evans B.R., Finkelstein M., McMillan J.D., (Eds). Twenty-Sixth Symposium on Biotechnology for Fuels and Chemicals, Humana Press, Totowa, pp. 93-104.
- [58] Xiong H., Fenel F., Leisola M., Turunen O. 2004. Engineering the thermostability of *Trichoderma reesei* endo-1,4- β -xylanase II by combination of disulphide bridges. *Extremophiles* 8, 393-400.
- [59] Xiong L., Kameshwar A.K.S., Chen X., et al. (2016). The ACEII recombinant *Trichoderma reesei* QM9414 strains with enhanced xylanase production and its applications in production of xylitol from tree barks. *Microb Cell Fact.* 15, 215.

- [60] Zhang L., Wang X., Ruan Z., et al. (2014). Fungal cellulase/xylanase production and corresponding hydrolysis using pretreated corn stover as Substrates. *Appl Biochem Biotechnol.* 172, 1045-1054.
- [61] Zhou P., Zhu H., Yan Q., Katrolia P., Jiang Z. 2011. Purification and properties of a psychrotrophic *Trichoderma* sp. xylanase and its gene sequence. *Appl Biochem Biotechnol.* 164, 944-956.

