

# Biodegradation and Sugar Release from Canola Plant Biomass by Selected White Rot Fungi

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## Abstract

Canola crop is rich in plant biomass. It is considered a major cash crop in North America and a potential source for biofuel. We evaluated six strains of white rot basidiomycetes under solid state fermentation (SSF) for their potentials to secrete oxidative and hydrolytic enzymes to biodegrade canola plant biomass (CPB), and release sugars. *Fuscoporia gilva* and *Pleurotus tuberregium* produced high amount of laccase (440.86 U/L and 480.63 U/L at day 7), as well as carboxymethylcellulase (CMCase) (4.78 U/mL at day 21 and 3.13 U/mL at day 14) and xylanase (4.48 U/mL and 7.8 U/mL at day 21), respectively. *Bjerkandera adusta* showed high amount of MnP (50.4 U/L) and peroxidase (64.5 U/L), relative to the other strains. Loss of organic matter peaked after 21 days of incubation in all the tested strains; however, the best result (34.0%) was shown in *P. tuberregium*. The highest lignin loss was observed in *Coriolopsis caperata* strains. Among the sugar polymers, hemicellulose was highly degraded by *P. tuberregium* and *P. pulmonarius* (4.1% - 4.6%), while cellulose (3.3% - 4.3%) was mainly degraded by *F. gilva* and *B. adusta*. Glucose was the dominant sugar released by all the fungi tested, with the highest concentration of 1.25 mg/mL produced by *B. adusta* at day 14 of incubation. Results indicate that selected white rot fungi can achieve significant delignification of CPB within 14 days of solid state fermentation. Their importance in low cost pretreatment of lignocellulosic biomass prior to conversion into biofuels and bio-products of economic importance is discussed.

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## Keywords

### Canola Plant Biomass, Bio-Delignification, White Rot Fungi, Lignocellulose

## 1. Introduction

Canola is a lignocellulosic rich plant biomass and bioenergy crop in North America [1]. The plant biomass is usually left as trash in fields or ploughed back into the soil after harvest. Its basal stems and roots are woody, and slower to decompose relative to the stem portions [2]. Lignin (14.2%), cellulose (42.4%), and hemicellulose (16.4%) are major components of the canola plant material [3]. Among these fractions, lignin consists of a heterogeneous and highly cross-linked polymer [4], while cellulose is organized as glucose polymeric long linear chains and is arranged in various levels of fibrils [5] [6]. Hemicelluloses, in contrast, are ramified polymers composed of various types of carbohydrates and phenolic compounds, among which there are xylan, glucan and arabinoglucan that form the sugar backbone in its structure [7].

White rot fungi (WRF) have been studied extensively due to their potential value in biopulping, bleaching, and other lignin removal schemes [8]. They can be cultivated on different lignocellulosic materials: agricultural wastes, sawdust, paper products, etc. [9] [10]. Some WRF have selective delignification properties, by which they preferentially degrade lignin over cellulose [11]. There are several reports on the application of WRF in the delignification of plant residues to improve digestibility [12]-[15]. The pretreatment of biomass for sugar release and delignification to enhance their use in bio-energy applications are also reported [16] [17]. Ohgren *et al.* [18] reported that sugar yields from both hemicellulose and cellulose are critical parameters for an economically-feasible ethanol production process; and that partial delignification of pretreated biomass increases the overall yield of sugar.

Oxidative and hydrolytic extracellular enzymes (manganese peroxidase, laccase, cellulases and xylanases) from these white rot fungi are able to depolymerize lignocellulosics [19] [20]. There is evidence that the extracellular polysaccharides (EPS) produced by lignocellulolytic fungi play an important role in the degradation process [21]. These EPS can immobilize the exocellular enzymes. According to Catley [22], the gel formed by these biopolymers prevents the hyphal dehydration, permits cell adherence to other cells or surfaces and could possibly select molecules that have positive influence on environmental degradation.

The conversion of lignocellulosics into fermentable sugars is a key limiting step in industrial processes of cellulosic ethanol production from biomass. Industrially-produced biomass sugars require harsh physical-chemical pretreatments (such as steam explosion, with or without diluted sulfuric acid) to loosen lignin and release fibrils of cellulose and monomers from lignocellulosic components [5].

The increasing demand for canola oil and press cake for food, feed and bioenergy applications has led to increased acreage used for canola cultivation, and the production of enormous amounts of biomass as waste. There is the lack of information or studies regarding the biodegradation or bioconversion of canola biomass using white rot fungi. Our objective was to determine the enzymatic potentials of selected WRF on the biodegradation and release of sugar from canola plant biomass under solid state fermentation (SSF) conditions.

## 2. Materials and Methods

### 2.1. Organisms and Cultivation

The white rot fungi used in this study included two strains of *Coriolopsis caperata* (Berk.) Murrill, and one each of *Bjerkandera adusta* (Willd.) P. Karst, *Fuscoporia gilva* (Schwein.) T. Wagner and M. Fisch., *Pleurotus tuberregium* (Fr.) Sing., and *Pleurotus pulmonarius* (Fr.) Qué. They were obtained from culture collections at the Mushroom Biology and Fungal Biotechnology Laboratory (MBFBL), North Carolina Agricultural & Technical State University, Greensboro, NC, USA.

The fungal inocula were prepared by cultivating each fungus in basal medium of glucose (10 g·L<sup>-1</sup>); KH<sub>2</sub>PO<sub>4</sub> (0.8 g·L<sup>-1</sup>); NH<sub>4</sub>NO<sub>3</sub> (2 g·L<sup>-1</sup>); Na<sub>2</sub>HPO<sub>4</sub> (0.4 g·L<sup>-1</sup>); MgSO<sub>4</sub>·7H<sub>2</sub>O (0.5 g·L<sup>-1</sup>); and yeast extract (2 g·L<sup>-1</sup>). The following microelements were added to the basal medium: ZnSO<sub>4</sub>·7H<sub>2</sub>O (0.001 g·L<sup>-1</sup>); FeSO<sub>4</sub>·7H<sub>2</sub>O (0.005 g·L<sup>-1</sup>); CaCl<sub>2</sub>·2H<sub>2</sub>O (0.06 g·L<sup>-1</sup>); CuSO<sub>4</sub>·7H<sub>2</sub>O (0.005 g·L<sup>-1</sup>); MnSO<sub>4</sub>·H<sub>2</sub>O (0.005 g·L<sup>-1</sup>). The medium

was adjusted to pH 6.0. The fungal mycelia cultivation was performed in 100 mL of the above medium in 500 mL flasks sterilized at 121°C for 15 minutes, then inoculated with 5 agar blocks (4 mm diameter) containing actively growing mycelia (7-day-old) of test fungi. The flasks inoculated with *C. caperata*, *B. adusta*, or *F. gilva* were incubated at 25°C, while those inoculated with *P. tuberregium* and *P. pulmonarius* were incubated at 30°C.

## 2.2. Solid State Fermentation

The solid state fermentation was conducted in 250 mL Erlenmeyer flasks, containing 5 g (dry weight) of milled canola plant materials (substrate) mixed with water (ratio 1:4) and sterilized at 121°C for 15 min. Flasks containing the sterilized substrates were each inoculated with 2 mL homogenate of inoculum (43 - 52 mg of mycelia dry weight) prepared as described above, and incubated at temperatures designated for the various strains. Three replicate sample flasks per strain were withdrawn at 7, 14 and 21 days of incubation. Each sample was extracted with 40 mL sodium acetate buffer (100 mM, pH 5.0), for 2 h at 4°C and filtered through Whatman 1 paper. The filtrate was centrifuged at 6000 rpm for 15 min to obtain the crude extract, polysaccharide and water soluble sugar content. The leftover solid materials from each flask were dried at 60°C to reach a constant weight and values obtained was used to calculate lost organic matter (LOM), *i.e.*, the percent difference in dry weight between the test substrate and the control (uninoculated substrate) [16] [23]. The leftover solid materials were also analyzed for lignin, cellulose, and hemicellulose components using the method for dietary fiber [24].

## 2.3. Enzyme Activity Assays

Laccase activity was determined using 2,2'-azino-bis (3-ethylbenzthiazoline-6-sulphonic acid (ABTS) as substrate and absorbance measured at 420 nm [25]. The reaction mixture contained 50 mM acetate buffer (pH 3.8), 1 mM ABTS, and 100 µL diluted culture filtrate. Manganese depended peroxidase (MnP) activity was assayed by the oxidation of phenol red. The reaction buffer (1 mL) contained sodium lactate-succinate buffer (25 mM, pH 4.5), 2 mM H<sub>2</sub>O<sub>2</sub>, 3 mM Phenol Red, and culture filtrate. The reaction was terminated by the addition of 2 M NaOH and absorbance recorded at 610 nm. All enzyme assays were carried out at room temperature (25°C). One unit of enzyme activity was defined as an amount of enzyme that transformed 1 µmol substrate per minute [26].

Carboxymethylcellulase (CMCase) activity was assayed with 1% solution of carboxymethyl cellulose as enzyme substrate in 0.05 M citrate buffer (pH 5.0) according to IUPAC recommendations [27]. The reaction mixture was incubated for 10 minutes in a buffer at a temperature of 50°C. Xylanase activity was assayed using a 1% solution of xylan from birch wood (Roth 7500) as substrate in 0.05 M citrate buffer (pH 5.0). The release of glucose and xylose, respectively, in 10 minutes at 50°C was measured using the dinitrosalicylic acid method [28]. One unit of enzyme activity is defined as 1 µmol of glucose or xylose equivalents released per minute under the given conditions.

The 1,4-β-glucosidase (EC 3.2.1.2) and 1,4-β-xylosidase (EC 3.2.1.37) activities were determined by measuring the rate of hydrolysis of, *p*-nitrophenyl-β-D-glucopyranoside and *p*-nitrophenyl-β-D-xylopyranoside respectively, as described by Poutanen and Puls [29]. The reaction mixture containing 1.8 mL of 2.5 mM substrate and 0.2 mL of the culture extract filtrate were incubated at 50°C for 10 minutes. The reaction was stopped by adding 1 mL of 1 M Na<sub>2</sub>CO<sub>3</sub> into the reaction mixture. The enzyme quantity catalyzing the release of 1 µmol of *p*-nitrophenol per min under these conditions was considered to be a unit of enzyme activity.

## 2.4. Exopolysaccharides Assay

Crude exopolysaccharides (EPS) was determined as previously reported, briefly, 1 volume of crude extract from the biomass was mixed with 4 volumes of absolute ethanol, stirred and left for 24 hours at 4°C. Precipitated EPS were collected by centrifugation at 6000 g for 20 min, dried and weighed [30].

## 2.5. Sugar Assay

The sugar content of the crude extract filtrate (pH 5.5) was quantitatively determined as reported previously, on a HPLC system utilizing Shodex KC-811 column [16].

## 2.6. Statistical Analysis

SPSS software version 17 (SPSS Inc., Chicago) was used for statistical analysis. A one-way analysis of variance

was carried out at  $\alpha = 0.05$ , and Duncan's multiple range test was used to compare the enzyme activities, polysaccharide production and biomass utilization, while Dunnett's statistics was used to compare the mean concentrations at the various sampling days with the un-inoculated as control.

### 3. Results and Discussions

#### 3.1. Macromolecules Degradation and Polysaccharide Production

The results indicate that among the three macromolecules, only lignin was significantly degraded (**Table 1**). After seven days only *B. adusta* showed significant degradation in lignin content. However, after 14 days, the extent of degradation by all the organisms was significantly different ( $P < 0.05$ ) from the control. Although the extent of the degradation differed among the tested organisms, *P. pulmonarius* had the highest degradation rates (residual lignin content had decreased from 15.24% to 10.9%), which was significantly different ( $P < 0.05$ ) from the other organisms (**Table 1**). However, after 21 days of incubation, all the organisms degraded lignin significantly ( $P < 0.05$ ), with the least residual lignin at 8.6% and 9.2% on substrate incubated with *C. caperata* 1 and 2, respectively, and followed by *P. pulmonarius* (10%). Significant degradation of hemicellulose was observed in *B. adusta* and *P. pulmonarius* and *C. caperata* 2 after seven days and *P. tuberregium* after 14 days. Cellulose degradation was negligible in all strains tested except in *B. adusta*, where significant degradation was observed after 14 days. All macromolecule degradation data are shown in **Table 1**.

Analysis of canola plant materials by the selected white rot fungi strains indicate selective delignification, since only the lignin component was significantly degraded, while the hemicelluloses and cellulose remained relatively intact. The selective biodelignification of lignocellulosic materials has high biotechnological importance in biofuels production, bioenergy applications [17] [18], feed and forage digestibility [6], as well as other environmental applications especially in the areas of waste conversion and recycling [31] [32].

In the early stages of fermentation, WRF have the added advantage of selective lignin biodegradation while conserving the cellulose in various plant biomass. In this study, the highest lignin loss was recorded in the two *C. caperata* isolates (38.7% and 42.9%), followed by *P. pulmonarius* (33.3%) and the rest of the fungi (14.7% - 24.0%) after 21 days of incubation. Akin *et al.* [12] showed lignin degradation of *Ceriporiopsis subvermispora* (29% - 33%), and *Cyathus stercoreus* (63% - 77%) after 6 weeks of incubation with Bermuda grass. Rois and Eyzaguirre [32] reported the selective degradation of lignin by the fungus *Ganoderma australe* and similar reports exist for various WRF tested for selective biomass delignification with *Pleurotus*, *Pycnoporus*, *Ischnoderma*, *Phlebia* spp [4] [33] [34]. Hatakka [35] studied the pretreatment of wheat straw by 19 WRF and found that 35% of the straw was converted to sugar by *Pleurotus ostreatus* in 5 weeks.

In all the WRF tested, the loss of organic matter (LOM) was highest after 21 days in all strains tested (**Table 2**). It is also visible that the organisms that showed highest degradation at the beginning did not turn out to be those with the highest degradation after three weeks, indicating the variability in degradation capacity versus fermentation time. In *P. tuberregium*, lignin correlated inversely with LOM ( $r = 0.999$ ;  $P < 0.05$ ), which clearly indicated that lignin biodegradation is associated with LOM. A loss in organic matter of 34% was recorded for *P. tuberregium*, which was significantly different from the other tested WRF (23.6% - 25.8%). This result recorded during this study is within the range (12.8% - 34.1%) recorded for *Grifola frondosa* during the biodegradation of oak sawdust in a 55-day fermentation experiment [36] and was lower than 55.8% recorded for *Lentinus squarrosulus* with cornstalk after 30 days of incubation [16]. However, the fact that this level of degradation happened in *P. tuberregium* and other tested strains within three weeks indicates their rapid degradation capacity, which may be useful in the pretreatment/delignification of biomass prior to use in producing cellulosic ethanol and other biomass to bio-products applications.

Exopolysaccharides (EPS) produced by all the test organisms at the various incubation periods, were all significantly higher ( $P < 0.01$ ) than the control (1.4 mg/mL) (**Table 2**). Among the different incubation times, polysaccharide production was highest after 14 days, followed by after 21 days and least after 7 days. The highest polysaccharide (5.3 mg/mL) was produced by *F. gilva* and *C. caperata* 2 after 14 days of incubation. Other investigators have similarly reported the production of polysaccharide during the fungal degradation of wood biomass. *L. squarrosulus* produced polysaccharides attaining peak concentration of 4.0 and 5.13 mg/mL after 18 and 24 days of incubation [16]. Mikiashvili *et al.* [36] found that *G. frondosa* produced polysaccharide ranging 3.2 - 3.5 mg/mL under SSF after 55 days of incubation. The roles of EPS secreted by fungi include mycelia adhesion to the substrate, protection against dehydration; and storage as a carbon source [21]. Protection against

**Table 1.** Lignin, cellulose and hemicellulose content in canola plant biomass incubated with different species of Basidiomycetes.

Species	Days	Hemicellulose (%)	Cellulose (%)	Lignin (%)
<i>C. caperata</i> 1	7	16.98 ± 0.60 def	45.95 ± 0.90 cd	14.92 ± 0.33 ijk
	14	16.81 ± 0.45 def	45.55 ± 0.42 bcd	11.87 ± 0.40 def <sup>*</sup>
	21	16.41 ± 0.40 cde	46.51 ± 0.38 cd	8.68 ± 0.83 a <sup>*</sup>
<i>C. caperata</i> 2	7	15.88 ± 0.57 abcd	44.20 ± 0.46 abc	15.18 ± 0.64 ijk
	14	17.09 ± 0.34 def	45.32 ± 1.54 bcd	12.96 ± 1.42 fgh <sup>*</sup>
	21	17.08 ± 0.55 def	45.96 ± 0.67 cd	9.24 ± 1.25 ab <sup>*</sup>
<i>B. adusta</i>	7	15.61 ± 0.44 abc <sup>*</sup>	44.28 ± 0.63 abc	13.83 ± 0.37 ghi <sup>*</sup>
	14	16.18 ± 0.47 bcde	43.34 ± 1.52 a <sup>*</sup>	11.88 ± 0.54 def <sup>*</sup>
	21	16.93 ± 0.45 def	44.51 ± 1.92 abc	11.66 ± 1.19 def <sup>*</sup>
<i>F. gilva</i>	7	16.99 ± 0.84 def	46.04 ± 1.39 cd	15.75 ± 0.58 ijk
	14	16.84 ± 0.31 def	44.74 ± 0.39 abcd	12.52 ± 0.68 ef <sup>*</sup>
	21	17.51 ± 0.32 def	43.83 ± 0.39 ab	12.76 ± 0.60 g <sup>*</sup>
<i>P. tuberregium</i>	7	16.59 ± 1.20 cdef	44.99 ± 0.45 abcd	15.03 ± 0.29 ijk
	14	14.95 ± 0.17 a <sup>*</sup>	45.32 ± 2.08 bcd	14.26 ± 1.63 ij <sup>*</sup>
	21	16.06 ± 0.49 bcde	45.88 ± 0.87 cd	11.39 ± 0.36 de <sup>*</sup>
<i>P. pulmonarius</i>	7	15.38 ± 0.04 ab <sup>*</sup>	44.87 ± 1.39 abcd	14.06 ± 0.49 hij <sup>*</sup>
	14	15.36 ± 0.22 ab <sup>*</sup>	44.54 ± 0.37 abc	10.92 ± 0.62 cd <sup>*</sup>
	21	16.14 ± 0.16 def	44.77 ± 0.07 abcd	10.05 ± 0.20 bc
Uninoculate substrate		16.84 ± 0.82 def	45.33 ± 0.46 bcd	15.24 ± 0.28 ijk

Each value is expressed as mean ± SE (n = 3). <sup>\*</sup>Measurements that is significantly different from the control at  $\alpha < 0.05$ . Means along the column with the same alphabetic numbers are not significantly different at  $\alpha > 0.05$ .

**Table 2.** Loss of organic matter and exopolysaccharides content in canola plant biomass during solid state fermentation with selected white rot fungi.

Species	Days	LOM (%)	Exopolysaccharides (mg/ml)
<i>C. caperata</i> 1	7	17.47 ± 1.81 e	2.20 ± 0.17 a
	14	20.0 ± 0.87 cd	4.47 ± 0.03 bcd
	21	23.93 ± 1.40 b	4.10 ± 0.12 bc
<i>C. caperata</i> 2	7	18.20 ± 1.83 de	2.37 ± 0.32 a
	14	18.93 ± 2.04 de	5.33 ± 0.12 f
	21	24.93 ± 2.21 b	3.97 ± 0.09 b
<i>B. adusta</i>	7	17.20 ± 1.40 e	2.33 ± 0.24 a
	14	17.20 ± 0.72 e	4.83 ± 0.09 de
	21	25.47 ± 0.31 b	4.70 ± 0.31 cde
<i>F. gilva</i>	7	17.20 ± 1.40 e	4.83 ± 0.15 de
	14	17.20 ± 0.72 e	5.37 ± 0.29 f
	21	23.60 ± 1.74 b	4.53 ± 0.35 bcd
<i>P. tuberregium</i>	7	12.87 ± 0.90 f	4.20 ± 0.15 bcd
	14	16.53 ± 1.30 e	4.30 ± 0.20 bcd
	21	34.07 ± 1.10 a	4.33 ± 0.18 bcd
<i>P. pulmonarius</i>	7	18.00 ± 0.92 e	2.40 ± 0.30 a
	14	21.33 ± 0.90 c	4.40 ± 0.31 bcd
	21	25.80 ± 1.31 b	4.80 ± 0.10 cde

Each value is expressed as mean ± SE (n = 3). Means along the column with the same alphabetic numbers are not significantly different at  $\alpha > 0.05$ .

dehydration and environmental injuries can be considered a general function of extracellular polysaccharides. The EPS by nature of their glucan content that is composed mainly of glucose, mannose, galactose, xylose and fucose [37] [38], could also be considered as storage compounds, which are consumed by the fungus when exogenous carbon sources are limited [21]. Therefore, the observed decrease in EPS seen in most tested WRF after two weeks of incubation, is a result of degradation and utilization of the sugar monomers in the EPS. Also, some of the sugars detected could actually be by-products from the degradation of the EPS formed in the earlier stages of growth of the fungus on the substrate. Other roles of the extracellular polysaccharides are to facilitate the diffusion and concentration of biodegradation enzymes, and also provide a medium with suitable ionic and pH conditions for degradation activities [21]. McCue and Shetty [39] investigated the involvement of lignin degradation activities in SSB of soybean by *Lentinus edodes* and they observed that decreased antioxidant activity was associated with total peroxidase and laccase activity. There are many reports on studies of EPS production by white rot fungi in submerged fermentation [40]-[42]. Their application is in nutraceuticals, especially  $\beta$ -glucans [43] and environmental remediation [44]. However, studies on exopolysaccharide production in lignocellulosic substrates under SSF have been few [16] [36].

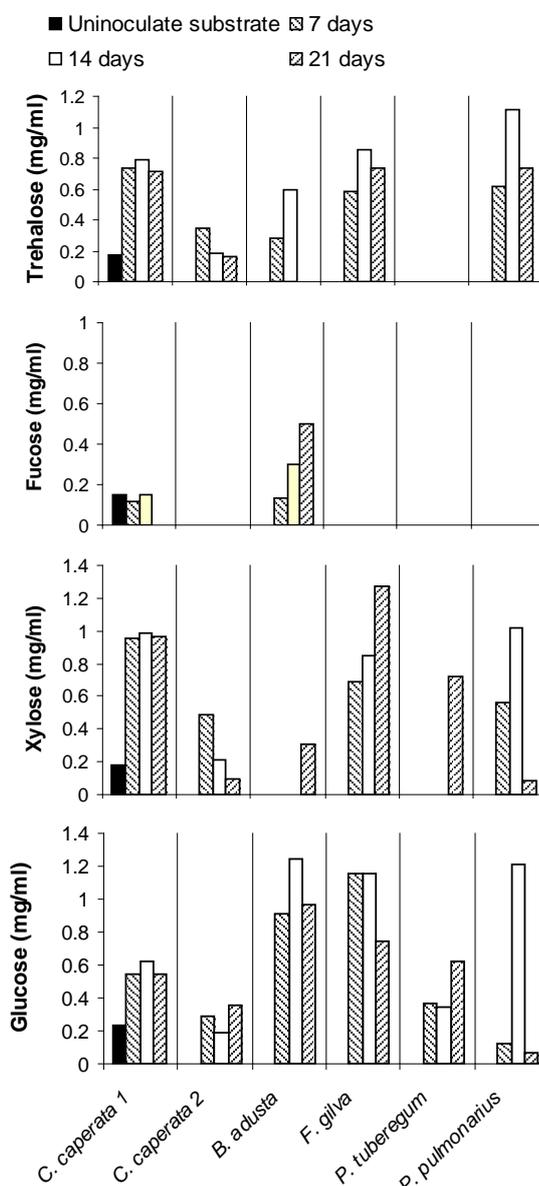
### 3.2. Soluble Sugar Content

Sugars were released to varying extents by the different test organisms during the biodegradation of plant materials (Figure 1). All the organisms released sugars throughout the three weeks incubation period. The highest glucose concentration (1.25 mg/mL) was determined for *B. adusta* after 14 days of incubation. The highest glucose concentration after 14 days of incubation was consistent among test strains. The trend was that glucose increased after one week, attained peak concentration after two weeks and declined after three weeks except for *C. caperata* 2 and *P. tuberregium*, which had the highest glucose concentration after three weeks. A similar trend was exhibited in trehalose levels except for *C. caperata* 2, which had the highest trehalose released after one week; however, this sugar was not detected with *P. tuberregium* at any time during the fermentation period. It is possible that released trehalose is rapidly metabolized to levels beyond detection limits used in this experiment. Another possibility is that it is never produced in the degradation/metabolic pathways for canola plants by *P. tuberregium*. The highest concentration of xylose, 1.27 mg/mL and 1.02 mg/mL was recorded for *F. gilva* after three weeks and *P. pulmonarius* after two weeks respectively. Significant quantities of fructose were only released by *C. caperata* 1 and *B. adusta*. It is worthy to mention that *C. caperata* 1 and *C. caperata* 2, though same species, exhibited different behavior on sugar release from the plant materials used in our study (Figure 1).

The conversion of woody biomass to fermentable sugars is challenging and needs to be overcome in cellulosic ethanol production as well as delignification to improve its digestibility when used as feed. Isikhuemhen *et al.* [16] reported the release of glucose (0.64 - 2.96 mg/mL), galactose (0.62 - 2.02 mg/mL) and fructose (0.18 - 0.38 mg/mL) during SSF of cornstalk using *Lentinus squarrosulus*. Ohgren *et al.* [18] reported 96% - 104% glucose yield and 70% - 74% xylose yield during the delignification of corn stover using enzymes. In our study, sugar appeared to be partially released, since the hemicelluloses and cellulose were barely degraded by the tested fungi. However, it appears that the period of sugar release peaked after two weeks, after which, there is increased utilization/assimilation of the sugars for fungi growth and metabolism. Our results suggest that to conserve sugars, and enhance pretreatment that achieves biodelignification using WRF should terminate after two weeks in most of the tested fungi. The pretreated material can then be subjected to cellulolytic enzyme activities or other forms of hydrolysis to release sugars, which can be used as feedstock for cellulosic ethanol production. Szczodrak and Fiedurek [14] reported that the production of lignocellulolytic enzymes appears to be the most expensive process, accounting to 43.7% of the total cost of cellulosic ethanol production. This cost consideration underscores the importance of this study, which used low cost SSF for the selective biodelignification of the plant materials. According to Sun and Cheng [17], the advantages of biological treatment of biomass over conventional chemicals include low energy requirements and mild environmental conditions though the rate of biological pretreatment process is slow.

### 3.3. Oxidative Enzyme Activities

Laccase activities were detected in all species tested (Table 3). In *B. adusta*, polysaccharide directly correlated with laccase ( $r = 0.998$ ;  $P < 0.05$ ), which suggests that lignin degradation by this fungus could contribute to the synthesis of polysaccharide. Generally, laccase increased after day 7 and attained a peak concentration on day



**Figure 1.** Sugar released during the biodegradation of CPB by white rot fungi. Uninoculated substrate contains: 0.23 mg/ml glucose, 0.17 mg/ml trehalose, 0.18 mg/ml xylose and 0.15 mg/ml fucose.

14 but declined to lower level on day 21 ( $P < 0.05$ ) except for *F. gilva* and *P. tuberregium*, which expressed peak enzyme concentration of 440.86 U/L and 480.63 U/L respectively after seven days. There was no correlation between laccase and lignin degradation among the fungi tested in this study. Hammel [31] reported that most ligninolytic fungi produce laccase. *L. squarrosulus* had a peak laccase activity of 154.5 U/L at day 6 of SSF with cornstalk [16]. Mikiashvilli *et al.* [36] reported peak laccase activity of 703.3 U/L by *G. frondosa* under SSF after 55 days of incubation. Laccase has been described as blue copper oxidases that catalyze the one-electron oxidation of lignin, phenolics and other electron rich compounds [31].

MnP also exhibited a pattern similar to laccase, increasing after seven days and attaining a peak activity after 14 days (Table 3). *B. adusta* and *P. pulmonarius* had the highest MnP concentration of 50 U/L after two weeks, while the least (0.23 U/L) was recorded for *C. caperata* 1 after 21 days. Unlike laccase and MnP, peroxidase exhibited different pattern among the different organisms tested. For *C. caperata* 1 and *C. caperata* 2, peroxidase had

**Table 3.** Lignocellulose degrading enzyme activities of different white rot fungi during solid state fermentation on canola plant material.

Species	Days	Laccase (U/L)	MnP (U/L)	Peroxidase (U/L)	CMCase (U/mL)	Glucosidase (U/mL)	Xylosidase (U/mL)	Xylanase (U/L)
<i>C. caperata</i> 1	7	40.52 ± 3.61 b	2.75 ± 0.98 a	3.08 ± 0.18 ab	0.32 ± 0.05 a	1.58 ± 0.05 bc	1.23 ± 0.04 efg	0.22 ± 0.13 a
	14	183.89 ± 9.90 g	0.96 ± 0.13 a	1.72 ± 0.26 ab	0.92 ± 0.09 abc	1.13 ± 0.09 b	1.26 ± 0.07 efg	2.46 ± 0.52 cd
	21	76.22 ± 2.76 c	0.23 ± 0.04 a	0.45 ± 0.09 a	1.10 ± 0.28 abcd	1.32 ± 0.16 b	0.97 ± 0.04 abcd	2.09 ± 0.49 bcd
<i>C. caperata</i> 2	7	56.06 ± 1.95 b	3.58 ± 0.27 a	4.24 ± 0.25 b	0.37 ± 0.18 a	1.91 ± 0.08 cd	1.15 ± 0.10 cdef	0.67 ± 0.13 ab
	14	221.08 ± 5.77 hi	1.79 ± 0.11 a	2.27 ± 0.27 ab	2.02 ± 0.24 bcde	1.14 ± 0.11 b	0.86 ± 0.06 a	2.54 ± 0.42 cd
	21	111.37 ± 4.65 d	1.26 ± 0.18 a	0.81 ± 0.20 a	1.20 ± 0.40 abcd	1.31 ± 0.07 b	0.92 ± 0.09 ab	2.09 ± 0.40 bcd
<i>B. adusta</i>	7	16.65 ± 3.20 a	1.74 ± 0.29 a	2.88 ± 0.13 ab	0.18 ± 0.05 a	1.33 ± 0.08 b	1.12 ± 0.05 bcde	0.30 ± 0.07 a
	14	189.63 ± 7.10 g	50.36 ± 3.03 f	65.35 ± 2.63 g	2.30 ± 0.33 cde	2.63 ± 0.17 f	0.85 ± 0.07 a	1.72 ± 0.33 abc
	21	169.65 ± 10.73 fg	18.38 ± 1.76 c	23.05 ± 0.64 e	1.47 ± 0.33 abcd	5.00 ± 0.35 h	1.30 ± 0.09 fg	2.54 ± 0.40 cd
<i>F. gilva</i>	7	440.86 ± 9.41 j	24.30 ± 2.89 de	24.21 ± 1.79 e	0.78 ± 0.20 ab	2.46 ± 0.07 ef	1.23 ± 0.05 efg	1.79 ± 0.13 abc
	14	160.03 ± 5.01 f	17.17 ± 0.79 c	10.71 ± 0.33 c	3.13 ± 0.56 e	2.31 ± 0.11 def	1.04 ± 0.02 abcd	4.48 ± 0.85 d
	21	137.46 ± 2.89 e	27.48 ± 0.22 e	30.60 ± 0.38 f	4.78 ± 0.24 f	2.15 ± 0.14 de	1.29 ± 0.10 fg	9.26 ± 0.64 g
<i>P. tuberregium</i>	7	480.63 ± 10.95 k	12.52 ± 1.13 b	11.54 ± 0.79 c	0.92 ± 0.24 abc	5.60 ± 0.26 i	1.27 ± 0.04 fg	3.58 ± 0.13 de
	14	133.76 ± 5.35 e	3.89 ± 0.88 a	2.75 ± 0.37 ab	3.13 ± 0.40 e	2.20 ± 0.12 def	0.95 ± 0.07 abc	4.33 ± 0.79 d
	21	231.25 ± 10.11 i	22.65 ± 0.60 d	14.67 ± 0.42 d	2.21 ± 0.48 cde	3.63 ± 0.07 g	1.40 ± 0.03 fg	6.42 ± 1.37 f
<i>P. pulmonarius</i>	7	8.14 ± 0.19 a	1.62 ± 0.28 a	2.75 ± 0.38 ab	0.37 ± 0.18 a	1.36 ± 0.06 b	1.15 ± 0.05 cde	0.60 ± 0.07 ab
	14	209.24 ± 4.98 h	49.75 ± 1.00 f	11.69 ± 0.81 c	2.41 ± 1.26 de	1.88 ± 0.14 cd	1.07 ± 0.10 abcde	1.49 ± 0.07 abc
	21	102.86 ± 6.80 d	25.99 ± 2.10 de	3.89 ± 0.66 b	1.29 ± 0.33 abcd	0.37 ± 0.02 a	1.19 ± 0.08 deg	0.75 ± 0.15 ab

Each value is expressed as mean ± SD (n = 3). Means along the column with the same alphabetic numbers are not significantly different at  $\alpha > 0.05$ .

peak concentration after seven days and decreased thereafter. For *B. adusta*, peroxidase activity of 2.88 U/L was detected after one week but attained peak activity of 63.3 U/L after 14 days and declined to 23 U/L after 21 days ( $P < 0.05$ ). For *F. gilva*, peroxidase had a peak concentration after one week, which decreased after two weeks, but increased again to reach high levels on day 21. For *P. tuberregium*, peroxidase had peak concentration after 21 days, although the concentration on day seven was significantly higher than after day 14 ( $P < 0.05$ ). *P. pulmonarius* had peak peroxidase concentration after 14 days. In *F. gilva*, polysaccharide was inversely correlated with MnP ( $r = 0.998$ ;  $P < 0.05$ ) and peroxidase ( $r = 0.999$ ;  $P < 0.01$ ) indicating that both enzymes, could enhance the biosynthesis of polysaccharide. In *C. caperata* 1, peroxidase directly correlated with residual lignin ( $r = 0.999$ ;  $P < 0.01$ ) and therefore indirectly with lignin loss, thus indicating that the enzyme is linked to lignin degradation. It is generally believed that WRF possess the ability to degrade lignin due to the secretion of peroxidases [45] [46]. Rios and Eyzaquirre [32] linked the selective degradation of lignin by *Ganoderma australis* to the production of MnP under low nitrogen content, low oxygen availability and high humidity. Mikiashvilli *et al.* [36] reported peak peroxidase activity of 22.6 U/L for *G. frondosa* grown on oak sawdust. Isikhuemhen *et al.* [16] reported a peak MnP activity of 13 U/L and peroxidase activity of 27.4 U/L for *L. squarrosulus* grown under SSF during six days of incubation.

### 3.4. Hydrolytic Enzymes Activities

Four hydrolytic enzymes (CMCase, glucosidase, xylosidase, cellobiosidase and xylanase) were assayed during the study. CMCase concentrations were not significantly different ( $P < 0.05$ ) among the 6 tested strains after seven days (Table 3). However, the general trend exhibited by all the organisms is that CMCase concentration was at its least after seven days, increased to peak concentration after 14 days and declined thereafter, except for *F. gilva*, which produced the highest concentration of CMCase (4.78 U/mL) on day 21. Glucosidase exhibited a different pattern compared to the other enzymes previously presented (Table 3). The least glucosidase enzyme activity was recorded after 21 days for *P. pulmonarius* (0.33 U/mL), while the highest concentration (5.0 U/mL) was recorded after 21 for *B. adusta*. It has been demonstrated that during lignocellulosic biomass degradation, glucose yield correlated with the  $\beta$ -glucosidase activity from mutant *Trichoderma atroviride* TUB [47].

Among all the tested WRF, *C. caperata* 1, and *P. tuberregium* produced the least cellobiosidase activity (<1.0 U/mL) after 14 days, while the highest activity (2.17 U/mL) was recorded for *P. tuberregium* after 21 days (Table 3). With *C. caperata* 2, polysaccharide directly correlated with CMCase ( $r = 0.999$ ;  $P < 0.05$ ) while in *P. tuberregium*, CMCase inversely correlated with glucose ( $r = 1.000$ ;  $P < 0.01$ ). In *F. gilva*, cellulose inversely correlated with CMCase ( $r = 1.000$ ;  $P < 0.01$ ) while in *C. caperata* 2, cellulose inversely correlated with cellobiosidase ( $r = 1.000$ ;  $P < 0.01$ ) and in *F. gilva*, LOM directly correlated with cellobiosidase ( $r = 0.997$ ;  $P < 0.05$ ). These correlations suggest that these enzymes play some roles in the degradation of macromolecules present in canola plant materials and the release of fermentable sugars.

Xylanase activity exhibited different patterns among the different organisms (Table 3). Among all the organisms, xylanase had the least activity after seven days, but after 14 and 21 days, different patterns were observed among the tested organisms. In *C. caperata* 1, *C. caperata* 2 and *P. pulmonarius*, xylanase concentrations decreased after 14 or 21 days, whereas with *B. adusta*, *F. gilva* and *P. tuberregium*, it increased, and *P. tuberregium* expressing the highest xylanase activity of 9.2 U/mL on day 21. In *C. caperata* 1, xylanase activity inversely correlated with polysaccharide ( $r = 1.000$ ;  $P < 0.01$ ), while in *P. tuberregium*, lignin inversely correlated with xylanase ( $r = 0.998$ ;  $P < 0.05$ ). This may indicate that xylanase does play some significant roles in the degradation of ligninocellulose and the biosynthesis of polysaccharide. Boominathan and Reddy [48] reported that *P. chrysosporum* produces lignin degrading enzymes including lignin peroxidases and MnP during secondary metabolism in response to carbon or nitrogen limitations. Other investigators have reported similar hydrolytic enzymatic activities among wood degrading fungi. Isikhuemhen [16] reported the hydrolytic enzyme activities to be 6.0 U/mL CMCase) and 14.5 U/mL for xylanase during six days fermentation under of cornstark under SSF conditions. Kovacs *et al.* [47] demonstrated that xylose yield correlated with the  $\beta$ -xylosidase, using mutant, *Trichoderma atroviride* TUB. Other studies have shown the role of xylanase and xylosidase in the hydrolysis of hemicelluloses during enzymatic saccharification of lignocellulosics [18] [49].

## 4. Conclusion

The results presented indicated that the WRF tested have the ability to selectively degrade the lignin fraction of

canola plant biomass, and will produce both oxidative and hydrolytic signature enzymes; exopolysaccharides and fermentable sugars. The mode of biodegradation is considered selective delignification, because lignin was significantly degraded compared to hemicelluloses and cellulose components. The results indicated the lack of linear relationships between enzyme activities, macromolecules degradation, exopolysaccharide production and sugar release from plant biomass. This could indicate the various regulatory mechanisms that couple degradation/sugar release in such a way that maintains osmotic equilibrium for optimum physiological and metabolic activities within the different strains. However, selective biodelignification of lignocellulosic materials have several important industrial and biotechnological applications. It could become useful in the pretreatment of plant biomass to enhance feed and forage digestibility and plant biomass biodelignification prior to their use in cellulosic ethanol production, as well as waste conversion to other bioproducts of economic importance. An efficient bioconversion process by WRF could lead to plant biomass utilization in ways that improve the overall economy of canola production, as well as contributing to the reduction of wastes and pollution from canola production and applications.

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