

STATE UNIVERSITY OF MARINGA
AGRICULTURAL SCIENCES CENTER

EFFECTS OF LIGNOCELLULOLYTIC ENZYMES PRODUCED
BY *PLEUROTUS OSTREATUS* ON THE NUTRITIVE VALUE OF
WHOLE-PLANT CORN SILAGE

Author: Bruna Calvo Agostinho
Advisor: Prof. Dr. João Luiz Pratti Daniel
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State of Parana
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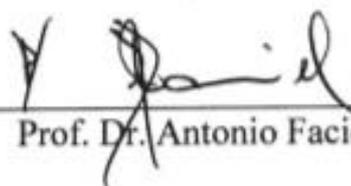
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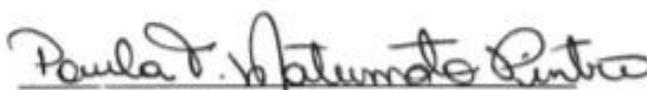
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NUTRITIVE VALUE OF WHOLE-PLANT CORN SILAGE

Autora: Bruna Calvo Agostinho
Orientador: Prof. Dr. João Luiz Pratti Daniel

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Prof. Dr. Antonio Faciola


Profª Drª Paula Toshimi
Matumoto Pinto


Prof. Dr. Luiz Felipe Ferraretto


Profª Drª Kathy Gisela Arriola


Prof. Dr. João Luiz Pratti Daniel
Orientador

“The size of your dreams must always exceed your current capacity to achieve them.

If your dreams do not scare you, they are not big enough.”

Ellen Johnson Sirleaf

*To my mother, Marta Calvo,
who always support me and believe in my dreams.*

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BIOGRAFY

BRUNA CALVO AGUSTINHO, daughter of Marta Calvo and Messias Antonio Agustinho, was born in Lupionopolis, Parana, on August 26th, 1992.

In January of 2015, she earned her bachelor's degree in Animal Sciences.

In March of 2015, she joined the Master's Degree in Graduate Program in Animal Science at the State University of Maringá.

In February of 2017, she submitted to the examining board to receive the title of Masters in Animal Sciences.

In March of 2017, she joined the Ph.D. Degree in Graduate Program in Animal Science at the State University of Maringa.

In February of 2020, she was approved in the Qualify Exams.

In December of 2020, she submitted her Ph.D. dissertation.

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ABSTRACT

1 Three experiments were carried out to examine the effect of lignocellulolytic enzymes
2 on the nutritive value of whole-plant corn silage (WPCS). In **experiment 1**, we
3 examined the effect of spent substrate from *Pleurotus ostreatus* cultivation (SSPO) on
4 the chemical composition, antioxidant capacity, lignin monomers, and *in vitro*
5 digestibility of WPCS. In **experiment 2**, we evaluated the performance of lactating
6 goats fed WPCS treated with different levels of SSPO. In **experiment 3**, we verified the
7 activity of an enzymatic complex at different pH and the effects of adding increasing
8 levels of that enzymatic complex produced by *Pleurotus ostreatus* on the fermentative
9 profile, chemical composition, and ruminal digestibility along of the days on the onset
10 of fermentation of WPCS. In **experiment 1**, four levels of lignocellulolytic enzymes
11 from spent substrate of *Pleurotus ostreatus* were tested in a completely randomized
12 design: 0, 10, 20, and 30 mg of lignocellulosic enzymes/kg of fresh matter, and four
13 replicates per treatment (vacuum-sealed bags). The bags were opened 60 d after
14 ensiling. The NDF, ADF, lignin, and cellulose concentration decreased quadratically. At
15 the nadir point, SSPO decreased NDF by 14.1%, ADF by 19.5%, lignin by 9.07%, and
16 cellulose by 22.1% compared to the untreated silage. Therefore, SSPO led to a quadratic
17 increase in IVDMD of WPCS (+10.3 % at vertex). In **experiment 2**, WPCS treated with
18 three enzyme levels (0, 10, or 30 mg/kg fresh matter, chosen based on experiment 1)
19 were fed to lactating goats as part of TMR. Nine lactating Saanen goats (62.68 ± 7.62 kg
20 BW; 44 ± 8 days in milk; 2.91 ± 0.81 kg of milk/day, mean \pm SD) were assigned to three 3×3
21 Latin squares. Intake and digestibility of dry matter and nutrients, microbial protein
22 syntheses, and milk production and composition were examined. The SSPO increase

23 the *in vivo* total-tract ADF digestibility quadratically. Additionally, the concentration of
24 polyphenols in milk increased linearly with the addition of SSPO in WPCS; however,
25 no other differences were detected among treatments. In **experiment 3**, the
26 lignocellulolytic enzymatic complex was obtained through *in vitro* cultivation of
27 *Pleurotus ostreatus*, and the activities of laccase, lignin peroxidase, manganese
28 peroxidase, endo and exoglucanase, xylanase, and mannanase were determined at pH 3,
29 4, 5, and 6. Following, five different enzymatic complex levels were tested in a
30 completely randomized block design: 0; 9; 18; 27, and 36 mg of lignocellulosic
31 enzymes/kg of fresh matter (FM) of whole-plant corn, with four replicates per treatment
32 (vacuum-sealed bags). The bags were opened after 1, 2, 3, and 7 d of ensiling to
33 evaluate the onset of fermentation and after 30 d of storage to evaluate the fermentation,
34 chemical composition, and *in situ* digestibility of WPCS. Laccase showed highest
35 activity at pH 5 ($P < 0.01$), whereas manganese peroxidase and lignin peroxidase had a
36 higher activity at pH 4 ($P < 0.01$; < 0.01 , respectively). There was no interaction
37 between the enzymatic complex and days of fermentation ($P > 0.11$). The concentration
38 of WSC decreased quadratically at the onset of fermentation ($P = 0.02$) due to its
39 consumption that led to a quadratic increase of lactic acid ($P = 0.01$) and a linear
40 increase of acetic acid ($P = 0.02$). As a result of increasing those organic acid
41 concentrations, pH decreased quadratically ($P = 0.01$). Lignin concentration decreased
42 linearly ($P = 0.04$) with the enzymatic complex at 30 d of storage. The collective
43 interpretation of these results leads to the conclusion that 10 mg of lignocellulolytic
44 enzymes from SSPO per kg of FM of WPC presented the best effect in silage
45 production due to more evident reduction in NDF, ADF, and lignin concentration and
46 increased ADF digestibility of lactation goats.

47 **Keywords:** fiber, fibrolytic enzyme, laccase, lignin, white-rot fungi

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I. INTRODUCTION

62 Fiber is one of the most predominant fractions that constitute ruminant diets (Van
63 Soest, 1994). Its concentration can vary according to the ingredients that compose the
64 diet, and its main function is related to provide effective physical capacity and energy
65 through its digestion (NRC, 2001).

66 However, the extension of fiber digestion has a huge variation between the forage
67 sources, and it is strongly related to three factors: total cell wall content, variation in
68 structural arrangement associated with plant anatomy, and cell wall composition
69 (Wilson, 1994), as reviewed below.

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71

1. Literature Review

72

1.1. Plant cell wall

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Plant cell walls are complex structures vital to plant survival, providing structural integrity and flexibility (Houston et al., 2016) and defense against pathogens and stress as a physical barrier (Underwood, 2012). The type of polysaccharides in the cell wall varies according to plant species, location in the wall (primary or secondary), plant tissue, maturity, and environment (Doblin et al., 2010).

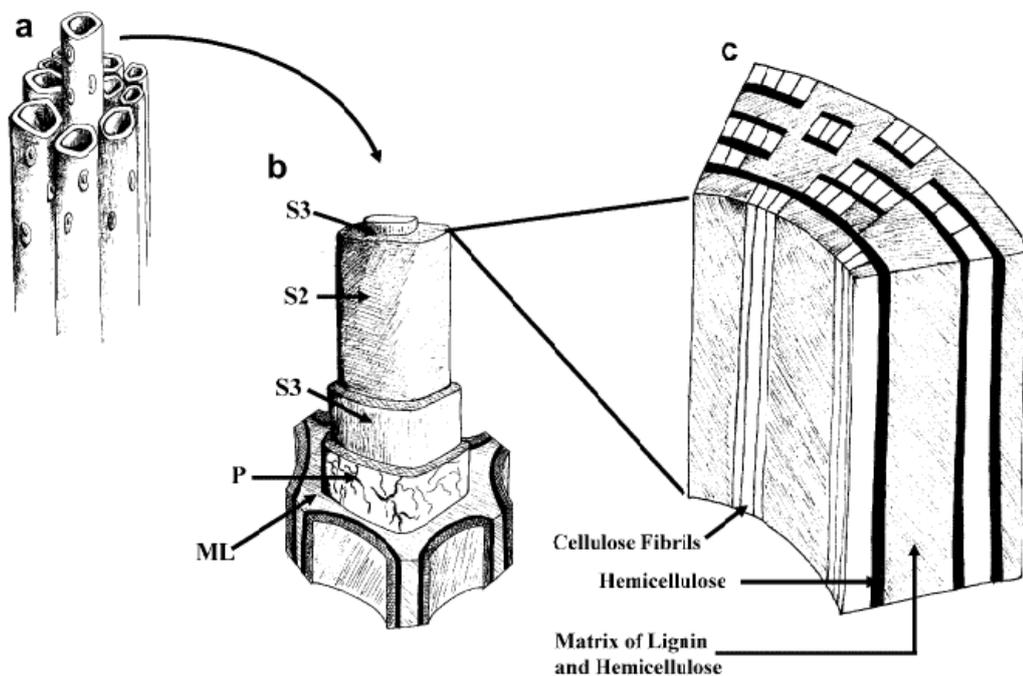
Plant cell walls are divided into primary and secondary walls and middle lamella (Figure 1). The primary wall consists of a cellulosic network encapsulated in a matrix of other polysaccharides, such as xylan and pectins (O'Neill and York, 2003).

Polysaccharides (cellulose and hemicellulose) represent approximately 90% of the primary wall, whereas structural glycoproteins (2-10%), phenolic esters (<2%), and minerals (1-5%) comprise a minor proportion (O'Neill and York, 2003).

84 The secondary wall contains celluloses, hemicelluloses (xylan and glucomannan),
 85 and lignin, with a low proportion of structural protein amount. Three layers compose the
 86 secondary wall (S1, S2, and S3). In S1, the cellulose microfibrils present crossed in
 87 each other; in the S2, they are oriented nearly parallel to the cell elongation axis; in the
 88 S3, they are oriented in a flat helix (Zhong et al., 2019). Between cell walls are present
 89 the middle lamella, which cements two cells together and is constituted of pectin
 90 (Srivastava, 2001).

91

92



93

94 Figure 1. Configuration of the plant cell wall. a) adjacent cells, b) cell wall layers, c)
 95 distribution of lignin and polysaccharides (from Pérez et al., 2002, adapted from Kirk and
 96 Cullen (1998)).

97 Abbreviation: ML: middle lamella; P: primary cell wall; S3, S2, and S3: layers in the secondary
 98 cell wall.

99

100 Cellulose is the most abundant carbohydrate in the cell wall, and it is formed by
 101 glucose molecules linked by β -1,4 bonds, whose chain length can present variation
 102 between 100 and 14000 residues (Béguin and Unitd, 1994). The chains are highly
 103 ordered, presenting a crystalline structure with amorphous regions. The degree of
 104 crystallinity varies according to the material, between 0% and 100% (Béguin and Unitd,
 105 1994), and the crystallinity degree of cellulose is associated with a linear reduction of
 106 degradation (Hall et al., 2010).

107 On the other hand, hemicellulose is a heterogeneous group composed of polymers
108 of pentoses (xylose, arabinose), hexoses (mannose, glucose, galactose), and sugar acids.
109 Hemicelluloses are strongly associated with lignin in the secondary wall and are
110 classified according to the main sugar residue in the backbone (Wyman et al., 2005).
111 The main residues are xylose, mannose, galactose, glucose, arabinose, and glucuronic
112 acid (Zhou et al., 2017). The linkage usually is β -1,4- and occasionally β -1,3-glycosidic
113 bonds (Pérez et al., 2002). Xylans are the most common hemicellulose group, with the
114 backbone of β -1,4-linkage xylose residues (Scheller and Ulvskov, 2010), and it has a
115 critical function to link lignin to hemicellulose (Wyman et al., 2005). Mannan is another
116 relevant group; its structure presents β -1,4-linkage between mannose residue and
117 backbone. The mannan is divided into glucomannan, galactomannan, and
118 galactoglucomanan, according to the composition of the residues (Ebringerova et al.,
119 2005).

120 Another important cell wall component is the lignin. Unlike cellulose and
121 hemicellulose, lignin is not a polysaccharide. Lignin is a complex structure formed from
122 oxidative-coupling of monolignols that are *p*-coumaryl, coniferyl, and sinapyl alcohols,
123 where the respective phenylpropanoid units are *p*-hydrophenyl (H), guaiacyl (G), and
124 syringyl (S) (Wong, 2009). Lignin is a heterogeneous group due to the extensive
125 possibility of monolignols combination (Hatakeyama and Hatakeyama, 2010). For
126 instance, grasses present G:S:H ratio, with proportions around 22:44:34, respectively
127 (Van Soest, 1994).

128 Lignin affects forage digestibility because it acts as a physical barrier. Cell wall
129 digestibility and plant maturity are negatively related since the plant lignification
130 increases with maturity. Its functions are related to the rigidity of the cell wall, reducing
131 water loss, and avoiding plant disease (Moore and Jung, 2001).

132

133 1.2. *Fiber in ruminant nutrition*

134 Forage is commonly used in ruminants diets, and in general, has high fiber
135 concentration. Fiber plays an essential role in maintaining ruminal function by
136 stimulating chewing and salivation, mat formation, and rumen motility. Therefore, a
137 minimal inclusion of roughage in ruminant diets is vital to avoid acidosis (Allen, 1997).

138 The minimum required level of physically effective neutral detergent fiber
139 (peNDF) to maintain animal health corresponds to approximately 31.2% peNDF

140 inclusive of particles >1.18 mm or 18.5% peNDF inclusive of particles >8 mm (Zebeli
141 et al., 2012). The peNDF is provided through forage addition in the diets and is altered
142 by forage proportion or particle length of forages (Yang and Beauchemin, 2007).

143 On the other hand, fiber is a significant energy source by microbial degradation
144 that produces volatile fatty acids (Tamminga, 1993). Fiber usually is provided to the
145 animal as grass pasture and conserved forage.

146

147 *1.3.Silage in ruminant nutrition*

148 Forage conservation is the process of preserving the forage to feed the animals
149 posteriorly. It is the key to feed cattle when there is no forage enough due to
150 temperature and rainfall (Gallaher and Pitman, 2000), or the animals stay the most of
151 the life inside the barn (Mayne et al., 2011). Therefore, forage conservation is a
152 technique essential to produce dairy cows and beef cattle efficiently. The two
153 techniques most used in the world to conserve forage are silage and hay. These
154 techniques offer benefits over the rest of the techniques, such as minimizing losses and
155 keeping forage with high quality during a long period, preserving the original feed
156 composition (Muck and Shinnars, 2001). Moreover, the ensilage process has the
157 advantage over hay production because it is less dependent on climate conditions
158 (Wilkinson and Rinne, 2017).

159 Silage is produced on spontaneous fermentation of water-soluble carbohydrates
160 (WSC) in compounds, such as organic acids and alcohols by epiphytic lactic acid
161 bacteria from the plant, under anaerobic conditions (Kung et al., 2017). Organic acids
162 are responsible for decreasing pH, where lactic and acetic acid are those organic acids
163 with higher concentrations in most silages. Lactic acid is the strongest organic acid
164 produced during the fermentation process; consequently, it is the one that contributes
165 the most to decreasing and maintaining acidic pH (Kung et al., 2017). While, acetic acid
166 works to improve aerobic stability because it inhibits molds and yeast growth at opening
167 time, whose silage is exposed to oxygen (Danner et al., 2003).

168 Corn (*Zea mays L.*) is the most common feedstuff used to produce silage and is
169 highly used as a forage source in dairy diets. Corn has a high yield per area and
170 provides fiber and energy in dairy diets (Grant and Ferraretto, 2018). To produce high-
171 quality corn silage is essential to pay attention to some factors, such as hybrid choice,
172 dry matter control, speed of packing, particle size, covering, and utilization of additives

173 (Allen et al., 2003). The hybrid choice is the first point to consider in silage production.
174 Among the various types of hybrids, there are conventional or dual-purpose hybrids;
175 silage-specific hybrids, which produce silage with better nutritive value; brown midrib
176 mutant (BMR) hybrids, which present less lignin and a higher fiber digestibility; and
177 leafy hybrids, which produce more leaves above the ear, and consequently more yield
178 per area (Ferraretto and Shaver, 2015). Dry matter content is another crucial factor, and,
179 ideally, the whole-plant corn (WPC) must have approximately 35% of dry matter. When
180 the dry matter is higher than 40%, the material could have a problem during compaction
181 that hampers removing oxygen, leading to heat damage by Maillard Reaction, and
182 spoilage. However, a low dry matter content in the WPC (less than 30%) also can lead
183 to losses of nutrients due to the leaching process that impair the nutritional value (Muck
184 and Shinnors, 2001). Packing speed must be done as quickly as possible to avoid losses
185 by fermentation of the water-soluble carbohydrates in aerobic conditions and cell
186 respiration (Moser, 1995). The recommendation to chop length of the WPC must
187 provide enough physically effective fiber to the animal and good silage compaction to
188 avoid problems in the storage process. Regarding silage covering film, it is essential to
189 prevent damage from precipitation, animal attack, sunlight exposure, and the most
190 crucial point, to ensure anaerobiosis conditions (Bernardes, 2016). The utilization of
191 additives at ensiling is a key factor in producing good silage.

192 The additives can improve corn silage fermentation, avoid nutrient losses, and
193 increase digestibility. Among those additives are inoculants, chemical additives, and
194 enzymes, such as proteolytic, cellulolytic, and hemicellulolytic (Muck et al., 2018).
195 Enzymatic additives is a class of additive which can improve silage composition and
196 digestibility. Different enzymatic additives have been tested and used in silages,
197 including fibrolytic and proteolytic enzymes.

198 The proteases improve starch digestibility in whole-plant corn silage since they
199 can hydrolyze prolamin in the matrix covering the starch (Young et al., 2012; Der
200 Bedrosian and Kung, 2019). Fibrolytic enzymes aim to break down the fibrous
201 carbohydrates and consequently increase dry matter digestibility. Fibrolytic enzymes
202 usually are used as a mix of cellulase and hemicellulase, and have been applied in
203 combination with bacterial inoculants (Muck et al., 2018).

204 1.4. Fibrolytic enzymes

205 Enzymes are complex three-dimensional proteins whose structure comprises
206 amino acids linked via an amide bond (Blanco and Blanco, 2017). Enzyme names are
207 based on the substrate that it acts, followed by the suffix –ase. For instance, cellulase is
208 the name of the enzyme group that degrades cellulose (Blanco and Blanco, 2017).
209 Cellulases hydrolyze β -1,4 linkages in celluloses chains, and they are classified in
210 endoglucanases, exoglucanases, and β -glucosidases (Gupta et al., 2013).
211 Endoglucanases hydrolyze internal cellulose bonds at the amorphous regions in
212 oligomers releasing new terminal ends, while exoglucanases, also known as
213 cellobiohydrolases, hydrolyze long-chain oligosaccharides (Pérez et al., 2002).
214 Exoglucanases are divided into two types, and they both act unidirectionally on the long
215 oligomers chain, one from reducing and the other one from nonreducing ends, releasing
216 cellobiose (Liu and Kokare, 2017), while β -glucosidases release D-glucose from
217 cellobiose and soluble cellodextrins (Wyman et al., 2005).

218 Hemicellulase is an extensive group of enzymes that hydrolyze hemicellulose,
219 whose main enzymes are xylanase, mannanase, α -arabinofuranosidase (Shallom and
220 Shoham, 2003). Xylanases degrade β -1,4 linkages, converting linear polysaccharide
221 xylan into xylose (Liu and Kokare, 2017), and it is the most studied hemicellulase given
222 that xylan is the predominant hemicellulose (Kormelink and Voragen, 1993). Another
223 important hemicellulose is endo- β -mannanases that release mannose from
224 polysaccharides (Freiesleben et al., 2018).

225 In general, commercial enzymes are complex and are comprised of a combination
226 of various enzymes. A commonly used enzyme combination is cellulase and
227 hemicellulase (McAllister et al., 2009), which act synergistically to intensify the
228 substrate degradation (Song et al., 2016). Most commercial products are derived from
229 fungi cultivation, mainly from *Trichoderma longibrachiatum*, *Aspergillus niger* and *A.*
230 *oryzae* (Pendleton 2000, cited in Beauchemin et al., 2004). Although there are many
231 cellulolytic and hemicellulolytic enzymes, there are no commercial enzymes that
232 breakdown lignin in animal nutrition. The development of a commercial enzyme that
233 prioritizes lignin breakdown can become a key method to increasing fiber digestibility.

234 Enzymes capable of lignin breakdown have the ability to oxidize a huge variety of
235 organic and inorganic compounds (Wong, 2009), which act by cross-linking
236 phenylpropanoid units with the aromatic ring, cleaving aromatic compounds (Liu and
237 Kokare, 2017); this effect has shown to work synergistically (Wong, 2009).

238 Laccases are multicopper enzymes consisting of four copper atoms per molecule
239 in the catalytic center that are one Cu atom type 1 (T1), one Cu atom type 2 (T2), and
240 two Cu atoms type 3 (T3) (Wong, 2009). In the enzymatic resting form, all atoms of
241 copper are Cu^{2+} , which corresponds to the oxidated state. The first step of catalysis is
242 reducing the Cu^{2+} T1 to Cu^{1+} as an electron acceptor from the substrate. After this step,
243 the next electrons extracted from the reducing substrate are transferred to T2/T3 site,
244 and in this time, the enzyme is transformed to the fully reduced (Giardina et al., 2010).
245 The cycle of laccase is O_2 dependent, which due to the high molecular mass (~60-90
246 kDa) (Shekher et al., 2011) and the low redox potential; this enzyme requires a mediator
247 to oxidize non-phenolic compounds (Pollegioni et al., 2015a).

248 Lignin peroxidase (LiP) and manganese peroxidase (MnP) are also enzymes that
249 break down lignin. These enzymes are dependent on H_2O_2 or another organic peroxide
250 to act as an electron acceptor during catalysis (Pollegioni et al., 2015b). LiP is a
251 glycoprotein with a heme group in the active center, is nonspecific to the substrate, and
252 is known to oxidize phenolic and non-phenolic compounds (Falade et al., 2017). The
253 LiP-I (first active form) is formed from the reaction between LiP and H_2O_2 , and it is
254 pH-independent. However, the reaction of LiP-I to oxidate the substrate and form LiP-II
255 is pH-dependent, and it usually decreases when the pH increases. The optimum pH in
256 this part of the cycle is around 3 (Tien and Kirk, 1984). The next reaction where LiP-II
257 is converted to LiP-III (inactive form) is pH-dependent and occurs when there is excess
258 of H_2O_2 (Wong, 2009).

259 The enzyme MnP catalyzes the oxidation of Mn (II) to Mn (III). This Mn (III) is
260 chelated into organic acids, as oxalate, malonate, and fumarate, and it favors the
261 capacity to penetrate small molecular pores between cellulose microfibrils to breakdown
262 lignin (Makela et al., 2015). The MnP oxidizes a massive variety of monomeric phenols
263 during the Mn oxidation, catalyzing a molecule of H_2O_2 to form MnP-I and the release
264 of H_2O . Following, Mg^{2+} is oxidized, forming MnP-II and Mg^{3+} (Vrsanska et al., 2016).
265 The Mn-P-II is oxidized and produces MnP-III, which in turn oxidizes another Mg^{2+} to
266 Mg^{3+} . In the trivalent state, magnesium can oxidize and break down phenolic
267 compounds in the presence of a second mediator, such as glutathione. Mg^{3+} is highly
268 reactive and subsequently oxidizes non-phenolic substrates, attributing to electron
269 abstraction from aromatic rings (Wong, 2009).

270 1.5. *Fibrolytic enzymes in ruminant nutrition*

271 Ruminants can use fiber as a source of energy due to their mutualism with ruminal
272 microorganisms. Fiber degradation yields volatile fatty acids, the main source of
273 metabolizable energy for ruminants (Bergman, 1990). However, part of the fiber cannot
274 be degraded by the microorganisms due to its recalcitrance, which is caused by its
275 organization as well as the association between lignin and carbohydrates. Thus, some
276 alternatives to increase fiber utilization have been developed, such as mechanical
277 processing, genetic improvement, alkali and acid treatments, bacterial inoculants, and
278 exogenous fibrolytic enzymes (Adesogan et al., 2019).

279 Exogenous fibrolytic enzymes release sugars and oligomers when applied to the
280 substrate before feeding animals, thus improving de fiber digestibility. Some studies
281 with these types of enzymes in dairy cow nutrition have shown an increase in fiber
282 digestibility and milk production (Arriola et al., 2017). However, the results are
283 inconsistent (Beauchemin et al., 2003) since there were improvements in animal
284 performance in some experiments, while there were no differences between the
285 treatments in other researches. These variations can be attributed to variable factors
286 among the experiments, such as diet chemical composition, doses, pH and temperature,
287 cofactors, animal category, and application methods (Adesogan et al., 2019).

289 1.6. *Pleurotus ostreatus*

290 The white-rot fungi belong to the group basidiomycetous fungi, and it is known to
291 produce enzymes that can hydrolyze cellulose and hemicellulose, and break down lignin
292 (Chandra and Madakka, 2019). *Pleurotus ostreatus* is one of those white-rot fungi
293 species and produces carboxymethylcellulose, xylanase, laccase, manganese peroxidase,
294 and lignin peroxidase (Elisashvili et al., 2003; Membrillo et al., 2008).

295 Researchers have been developing with *Pleurotus ostreatus* in agriculture wastes,
296 such as rice straw, wheat straw, with the goal to enhance digestibility and viability for
297 animal nutrition. Fazaeli et al. (2004) observed that diets with wheat straw previously
298 treated with *Pleurotus ostreatus* presented greater dry matter digestibility compared to
299 untreated straw. Kholif et al. (2014) observed that rice crop residue treated with
300 *Pleurotus ostreatus* allowed to clover replacement (*Trifolium alexandrinum*) by 50%
301 without reducing the digestibility of neutral fiber detergent.

302 Recently, Machado et al. (2020) evaluated an enzymatic complex produced by
 303 *Pleurotus ostreatus* and used it to treat whole-plant corn and sugarcane at ensiling. They
 304 reported a reduction in the concentration of lignin, cellulose, and hemicellulose, as well
 305 as greater *in vitro* digestibility, and antioxidant capacity in both corn and sugarcane
 306 silages treated with enzyme complex.

307 Moreover, *Pleurotus ostreatus* is one of the most cultivated fungi to produce
 308 mushrooms in the world. The cultivation is performed in lignocellulosic material, as
 309 palm heart waste, sugarcane bagasse, and hay. The fungi produce lignocellulolytic
 310 enzymes that can degrade cellulose and hemicellulose, leading to the release of glucose
 311 (Mikiashvili et al., 2006). After mushroom harvest, the cultivation medium becomes
 312 low in nutrients and is named as spent mushroom substrate. The spent mushroom
 313 substrate of *Pleurotus ostreatus* (SSPO) has been used to fertilize the soil, fed animals,
 314 or burned; however, it presents a high concentration of lignocellulolytic enzymes and
 315 can be used as an alternative enzymatic source (Phan and Sabaratnam, 2012). Although
 316 the SSPO has the potential to improve the digestibility of feedstuffs through the
 317 enzymatic action. Therefore, it has just been tested as a feed and not as an enzyme
 318 source to improve the chemical composition of the feeds. Therefore, this dissertation
 319 aimed to evaluate the addition of sources of lignocellulolytic enzymes from *Pleurotus*
 320 *ostreatus* (spent substrate and enzymatic complex produced in laboratory) in whole-
 321 plant corn at ensiling, and performance of goats fed with silage treated with
 322 lignocellulolytic enzymes.

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324 2. Reference

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II. OBJECTIVE

The objective of this study was to examine the effect of two sources of lignocellulolytic enzymes from *Pleurotus ostreatus* (spent mushroom substrate and enzymatic complex) in whole-plant corn silage and the performance of lactating goats fed corn silage treated with different levels of spent mushroom substrate.

ABSTRACT

557
558 The objectives of this study were to examine the effect of the spent substrate from
559 *Pleurotus ostreatus* cultivation (SSPO) on corn silage chemical composition,
560 antioxidant capacity, lignin monomers, and *in vitro* digestibility, as well as the
561 performance of lactating goats fed corn silage treated with different levels of SSPO. In
562 experiment 1, four levels of lignocellulolytic enzymes were tested in a completely
563 randomized design: 0, 10, 20, and 30 mg of lignocellulosic enzymes/kg of fresh matter,
564 and four replicates per treatment (vacuum-sealed bags). The bags were opened 60 d
565 after ensiling. In experiment 2, corn silage treated with three enzyme levels (0, 10, or 30
566 mg/kg fresh matter) was fed to lactating goats as part of TMR. Nine lactating Saanen
567 goats (62.68 ± 7.62 kg BW; 44 ± 8 days in milk; 2.91 ± 0.81 kg of milk/day, mean \pm SD)
568 were assigned to three 3×3 Latin squares. Data were analyzed using the GLIMMIX
569 procedure of SAS, and the means were compared by linear and quadratic orthogonal
570 contrast. In experiment 1, NDF, ADF, lignin, and cellulose decreased quadratically. At
571 the nadir point, SSPO decreased NDF by 14.1%, ADF by 19.5%, lignin by 9.07%, and
572 cellulose by 22.1% compared to the untreated silage. Therefore, SSPO led to a quadratic
573 increase in IVDMD of corn silage (+10.3 % at the vertex). In experiment 2, SSPO
574 increased quadratically the *in vivo* total-tract ADF digestibility. Also, the concentration
575 of polyphenols in the milk of goats increased linearly with the addition of SSPO in corn
576 silage, but no other differences were detected among treatments. In summary, adding 10
577 mg of lignocellulolytic enzymes from SSPO per kg fresh matter of corn at ensiling
578 presented the best effect in silage production, with a more evident reduction in ADF and
579 NDF concentration and increased ADF digestibility *in vivo*.

580 **Keywords:** fibrolytic enzyme, laccase, lignin, white-rot mushroom

581

INTRODUCTION

582
583 Some technologies have been studied to increase fiber digestibility, such as
584 exogenous fibrolytic enzymes produced by fungi or bacteria (Beauchemin et al., 2003),
585 and these enzymes are directly provided to the animal or used to treat feedstuffs prior to
586 feeding (Adesogan et al., 2019).

587 Treating feedstuffs with fibrolytic enzymes at ensiling is an alternative to decrease
588 fiber concentration, change fermentation profile, and increase fiber digestibility, as
589 observed in some research with cellulolytic and hemicellulolytic enzymes (Colombatto
590 et al., 2004a; Dean et al., 2005). However, few studies aimed to degrade lignin present
591 in forage with exogenous enzymes (Machado et al., 2020; Lynch et al., 2014; Pech-
592 Cervantes et al., 2019).

593 Lignolytic enzymes, such as laccase, manganese peroxidase, and lignin peroxidase
594 can be used to break down lignin (Pérez et al., 2002) and possibly increase fiber
595 digestibility. These enzymes, as well as several cellulolytic and hemicellulolytic
596 enzymes, are produced by white-rot mushrooms, such as *Pleurotus ostreatus* (Bánfi et
597 al., 2015), as a mechanism to obtain energy and nutrients from substrates (Phan and
598 Sabaratnam, 2012).

599 Previous studies had used *Pleurotus ostreatus* to treat straws, increased their
600 digestibility (Fazaeli et al., 2004), and replaced conventional ingredients in ruminant
601 diets (Khattab et al., 2013; Kholif et al., 2014). Recently, Machado et al. (2020)
602 examined the effects of enzymes isolated from *Pleurotus ostreatus* on the conservation
603 and nutritive value of whole-plant corn and sugarcane silages. In these studies, the
604 enzymes reduced the concentration of lignin, cellulose, and hemicellulose and increased
605 *in vitro* digestibility and antioxidant capacity in corn and sugarcane silages (Machado et
606 al., 2020). However, there is no study on the effects of residue from mushroom
607 cultivation as a potential source of lignocellulolytic enzymes to treat feedstuffs.

608 Thus, the objectives of this study were to evaluate the effect of the spent substrate
609 from *Pleurotus ostreatus* cultivation (SSPO) on corn silage chemical composition,
610 antioxidant capacity, lignin monomers, and *in vitro* digestibility, as well as to evaluate
611 the intake, digestibility, microbial protein synthesis, and milk yield and composition of
612 lactating goats fed with corn silage treated with different levels of SSPO. We
613 hypothesized that the addition of SSPO at ensiling could decrease fiber concentration of
614 the corn plant through lignin degradation and consequently increase fiber digestibility in
615 lactating goats.

616

617 MATERIAL AND METHODS

618 This study consisted of two experiments. In experiment 1, we evaluated the
619 composition and *in vitro* digestibility of whole-plant corn silage treated with different
620 doses of SSPO. Based on the results of the first trial, in experiment 2, we examined the
621 effects of treating corn silage with different doses of SSPO on the performance of
622 lactating goats.

623

624 *Enzyme source characterization*

625 The spent substrate from *Pleurotus ostreatus* (Jacq.) P. Kumm cultivation (SSPO)
626 was obtained from a mushroom producer in Maringa, PR, Brazil, after the commercial
627 harvest. The spent substrate was originated from mushroom cultivation carried out in
628 plastic bags, in which the initial substrate consisted of a mix of tropical grass hay and
629 palm kernel waste.

630 The SSPO was freeze-dried, ground to pass a 1 mm screen (Marconi MA340,
631 Piracicaba, SP, Brazil), and homogenized. A subsample was collected for measuring the

632 activities of laccase, manganese peroxidase, lignin peroxidase, carboxymethylcellulase,
633 mannanase, and xylanase. The same batch of SSPO was used in both trials.

634 Laccase activity was determined in UV-Vis spectrophotometer (PC 300
635 ThermoScientific, Waltham, MA, USA) at 420 nm, through the 2,2'-azino-bis(3-
636 etilbenzotiazolina-6-sulfonato) (ABTS) oxidation, where we used 0.2 mL of ABTS
637 solution (20mM), 0.2 mL of enzymes solution, and 1.6 mL of McIlvaine Buffer (pH
638 4.0) with a reaction time of 6 min at 25°C. Laccase activity was expressed in units (U),
639 and one U was defined as μmol of ABTS oxidized per min (Li and Xu, 1999).

640 Manganese peroxidase activity was determined in UV-Vis spectrophotometer (PC
641 300 ThermoScientific, Waltham, MA, USA) at 270 nm according to Wariishi Method
642 (Wariishi et al., 1992) where 0.6 mL of sodium malonate buffer (50mM, pH 4.5), 1.2
643 mL of enzymes solution, 0.6 mL of MnSO_4 (4.5 mM), and 0.3mL of H_2O_2 (9 mM) with
644 a reaction time of 5 min at room temperature was used. Manganese peroxidase activity
645 was expressed in U, and one U was defined as 1 μmol MnSO_4 oxidized per min.

646 Lignin peroxidase activity was determined in UV-Vis spectrophotometer (PC 300
647 ThermoScientific, Waltham, MA, USA) at 310 nm according to Tien and Kirk (1984)
648 through veratryl alcohol oxidation to veratrilaldehyde (3,4 dimethoxybenzaldehyde),
649 where 0.75 mL of sodium tartrate buffer (10 mM, pH 3.0), 0.5 mL of enzyme solution,
650 0.25 mL of veratryl alcohol (3mM), and 0.10 mL of H_2O_2 (5 mM), with a reaction time
651 of 5 min at room temperature was used. Lignin peroxidase activity was expressed in U,
652 and one U was defined as 1 μmol veratryl alcohol oxidized per min.

653 Carboxymethylcellulase activity was determined in UV-Vis spectrophotometer (PC
654 300 ThermoScientific, Waltham, MA, USA) at 540 nm, according to Ghose (1987),
655 where 0.25 mL of carboxymethylcellulose (0.5% in 50 mL sodium acetate buffer, pH
656 5.8), as the substrate, and 0.25 mL of enzyme solution for the reaction was used. The

657 reaction time was 30 min at 50° C, and posteriorly the amount of released reducing
658 sugar was marked with dinitrosalicylic acid, according to Miller (1959). Glucose was
659 used to generate a standard curve. Carboxymethylcellulase activity was expressed in U,
660 and one U was defined as 1 μ mol of glucose released per min.

661 Xylanase activity was determined in UV-Vis spectrophotometer (PC 300
662 ThermoScientific, Waltham, MA, USA) at 540 nm, according to Damiano et al. (2003),
663 where 0.9 mL of Birchwood-Sigma xylan (1% in 50 mM sodium acetate buffer, pH
664 5.4), as the substrate, and 0.1 mL of enzyme solution for the reaction was used. The
665 reaction time was 30 min at 50° C, and posteriorly the amount of released reducing
666 sugar was marked with dinitrosalicylic acid, according to Miller (1959). Xylose was
667 used to generate a standard curve. Xylanase activity was expressed in U, and one U was
668 defined as 1 μ mol of xylose released per min.

669 Mannanase activity was determined in UV-Vis spectrophotometer (PC 300
670 ThermoScientific, Waltham, MA, USA) at 540 nm, according to Rättö and Poutanen
671 (1988), where 0.9 mL of galactoglucomannan (0.5% in 50 mM sodium acetate buffer,
672 pH 5.4), as the substrate, and 0.1 mL of enzyme solution for the reaction was used. The
673 reaction time was 30 min at 50° C, and posteriorly the amount of released reducing
674 sugar was marked with dinitrosalicylic acid, according to Miller (1959). Mannose was
675 used as standard. Mannanase activity was expressed in U, and one U was defined as 1
676 μ mol of mannose released per min.

677 Protein concentration was measured using the Bradford Protein Assay (Sigma-
678 Aldrich Co., St. Louis, MO), and BSA was used as standard, according to Bradford
679 (1976), and one mg of protein in this method was defined as one mg of the enzyme.

680

681 ***Experiment 1:***

682 ***Silage production and treatment***

683 Whole-plant corn hybrid Biomatrix BM 3061 (Biomatrix, Rio Claro, SP, Brazil)
684 was harvested with 29.3 % dry matter (DM) by manual cutting and chopped into
685 approximately 2 cm length particles in a stationary forage harvester. A sample was
686 collected to determine the plant chemical composition before ensiling. Approximately
687 500 g of forage was weighed, homogenized, treated with the respective levels of SSPO,
688 placed into nylon-polyethylene vacuum bags, and heat-sealed using a vacuum machine
689 (TecMaq, TM-250, Sao Paulo, Brazil) to ensure an anaerobic environment. The silage
690 bags were stored for 60 d after ensiling, in a dark environment at room temperature.

691 This trial consisted of a completely randomized design with four treatments (levels
692 of enzyme) and four replicates per treatment (vacuum-sealed bags). The tested levels
693 were 0, 10, 20, and 30 mg of lignocellulolytic enzymes/kg of fresh matter, which
694 correspond to 0; 1,100; 2,200, and 3,300 U/min/kg of fresh matter, respectively (sum of
695 enzymatic activities of laccase, manganese peroxidase, lignin peroxidase,
696 carboxymethylcellulase). The levels (mg/kg of fresh matter) were based on a previous
697 study carried out with lignocellulolytic enzyme from *Pleurotus ostreatus* produced in
698 our laboratory (Machado et al., 2020).

699

700 ***Chemical composition***

701 Mini silos were opened 60 d after ensiling, weighed to estimate the recovery of
702 nutrients, and samples of silages were collected to determine pH, chemical composition,
703 antioxidant capacity, and *in vitro* digestibility. Approximately 9 g of undried silage was
704 diluted with 60 mL of distilled water, manually mixed, and left to stand for 30 mins
705 before measuring pH, and it was measured in duplicate using a digital pH meter. Two

706 samples from each mini silo were dried in a forced-air oven at 55°C for 72 h and then
707 ground to pass through a 1 mm screen in a Wiley mill (Marconi MA340, Piracicaba, SP,
708 Brazil).

709 The absolute DM content was determined at 105°C using an oven according to
710 method No. 924.01 (AOAC, 1990). Ash was determined by combustion at 600°C for 6
711 h in a furnace, according to method No. 924.05 (AOAC, 1990). Crude protein was
712 determined by the Kjeldahl method (Method No. 990.03, AOAC, 1990). Neutral
713 detergent fiber (NDF) was determined, according to Mertens (2002), using thermostable
714 α -amylase without sodium sulfite. Acid detergent fiber (ADF) was determined
715 according to method No. 973.18 (AOAC, 1990). Ether extract (EE) was determined
716 according to method No. 7.060 (AOAC, 1990). The non-fibrous carbohydrates (NFC)
717 concentration was calculated, according to Weiss (1999), using the following equation:
718 $NFC = 100 - (\% NDF + \% CP + \% EE + \% \text{ash})$. The concentration of lignin was
719 determined using the acid detergent lignin methodology, according to Van Soest and
720 Wine (1968), by submitting the material to sulfuric acid (72/28, v/v in DI water)
721 sequentially following ADF analysis. The cellulose concentration was obtained by the
722 difference between the ADF and lignin, whereas hemicellulose concentration was
723 obtained by the difference between NDF and ADF. Acid detergent insoluble nitrogen
724 (ADIN) and neutral detergent insoluble nitrogen (NDIN) were obtained through
725 nitrogen determination according to AOAC (AOAC, 1990) method No. 990.03 using
726 the residue from ADF, and NDF analyses, respectively.

727 The DM recovery was estimated using the following equation: $DMR (\%) = 100 \times$
728 $((FMop \times DMop)/(FMen \times DMen))$, where FMop = fresh matter mass at opening;
729 DMop = dry matter content at opening; FMen = fresh matter mass at ensiling; DMen =

730 dry matter content at ensiling. Nutrient recovery was estimated as the mass of a given
731 nutrient at silo opening as a proportion of the mass of that nutrient at ensiling.

732 The fractionation of carbohydrates was carried out according to System CNCPS
733 equations (Sniffen et al., 1992): Fraction A + B1 = $100 - (C + B2)$; Fraction B2 = $100 \times$
734 $((\text{NDF}(\% \text{DM})) - \text{NDIN}(\% \text{CP}) \times 0.01 \times \text{CP}(\% \text{DM}) - \text{NDF}(\% \text{DM}) \times 0.01 \times$
735 $\text{Lignin}(\% \text{NDF}) \times (\text{Lignin} \times 2.4)) / \text{Total carbohydrates}(\% \text{DM})$; Fraction C = $(100 \times$
736 $\text{NDF}(\% \text{DM}) \times 0.01 \times \text{Lignin}(\% \text{NDF}) \times (\text{Lignin} \times 2.4)) / \text{Total carbohydrates}(\% \text{DM})$),
737 where: NDIN = neutral detergent insoluble nitrogen.

738 Lignin monomer composition was determined by oxidation with nitrobenzene,
739 according to Bubna et al. (2011). The material was purified to obtain a protein-free cell
740 wall. Approximately 0.3 g of sample was homogenized with sodium, potassium
741 phosphate buffer (7 mL, 50 mM, pH 7.0), centrifuged at 1,400 g for 2 min supernatant
742 was discarded. This process was repeated four more times with sodium and potassium
743 phosphate buffer, three times with Triton (7 mL, 1/99, v/v in phosphate buffer, pH 7.0),
744 and three times with NaCl (7 mL, 1 M in pH phosphate buffer, pH 7.0). Subsequently,
745 the sample was washed 3 times with distilled water (7 mL) and 2 times with acetone (5
746 mL). The precipitated material was kept in an oven at 60°C for 24 h for drying, cooled
747 in a desiccator, and weighed to determine the cell wall concentration.

748 Approximately 50 mg of the purified cell wall, 1 mL de NaOH 2 M e 100 µL de
749 nitrobenzeno were placed in ampoules and then sealed under temperature. The
750 ampoules were kept at 170°C for 150 min and stirred manually when they reached 75
751 min. The samples were cooled after the reaction period, washed with chloroform twice,
752 acidified with HCl 5M, and submitted again to chloroform twice to promote the
753 extraction. The extracted sample was dried in rotavapor (IKA rotary evaporator RV10
754 digital V, IKA, Wilmington, NC), diluted in methanol, filtered through a 0.45 µm filter,

755 and analyzed by High-performance liquid chromatography (HPLC) (Agilent 1100
756 Series, Massy, France), using column ACE C18-AR, 150 mm × 4.6 mm × 5µm,
757 Aberdeen, Scotland, using 4% methanol/acetic acid in water (20/80, v/v) as a mobile
758 phase and flow 1.2 mL/min for isocratic analysis and 20 mins for isocratic analysis.
759 Quantification of *p*-hydroxybenzaldehyde (H), guaiacyl (G), and syringaldehyde (S) at
760 290 nm. The results were expressed in mg/g of the cell wall. The S/G ratio was
761 calculated with the data obtained from the quantification.

762

763 *Antioxidants*

764 The silage extract to determine the total antioxidant capacity (TAC), polyphenols,
765 flavonoids, and reducing power were performed with a 0.5 g of sample and a 4.5 mL of
766 90% methanol-water (90/10, v/v). The extract was stirred on a rotary shaker overnight,
767 centrifuged at 2,500 g for 10 min, and the supernatant was used in the analyzes. The
768 reducing power was determined according to Zhu et al. (2002), with Santos et al. (2014)
769 modifications. The absorbance was determined in a UV-Vis spectrophotometer (PC 300
770 ThermoScientific, Waltham, MA, USA) and expressed as gallic acid equivalents (GAE;
771 mg/100 g of silage, in DM basis). TAC was determined as described by Rufino et al.
772 (2007) with the addition of radical ABTS^{•+} (2,2-azinobis-(3-ethyl-benzothiazolin-6-
773 sulfonic acid)) to the extract. The absorbance was determined in a UV-Vis
774 spectrophotometer (PC 300 ThermoScientific, Waltham, MA, USA) and expressed in
775 Trolox equivalent (mM Trolox/100 g of silage, in DM basis). Total polyphenols were
776 determined according to Singleton and Rossi (1965). The absorbance was determined in
777 a UV-Vis spectrophotometer (PC 300 ThermoScientific, Waltham, MA, USA) and
778 expressed as gallic acid equivalents (EAG; mg/100 g of silage in, DM basis). According
779 to Woisky and Salatino (1998), Flavonoids were determined with the modifications of

780 Sánchez et al. (2010). The absorbance was determined in a UV-Vis spectrophotometer
781 (PC 300 ThermoScientific, Waltham, MA, USA) and expressed as quercetin equivalent
782 (QE; mg/100 g of silage, in DM basis).

783

784 *In vitro Digestibility*

785 All animal procedures were approved by the Committee for Use of Animals in
786 Experimentation from the State University of Maringá. The *in vitro* dry matter
787 digestibility (IVDMD) was performed according to Tilley and Terry (1963). Ruminant
788 content was taken from two non-lactating water buffaloes through ruminal cannula.
789 Buffaloes were fed a diet (corn silage, ground corn, soybean meal, wheat meal, and a
790 vitamin-mineral supplement) formulated for the maintenance of nutritional requirement
791 (CP = 10%; TDN = 65%), according to Paul and Lal (2010). The liquid and solid phase
792 of the ruminal contents were collected manually prior to the morning feeding from the
793 ventral, central, and dorsal areas of the rumen. Ruminant contents were mixed in a
794 blender, filtered into four layers of cheesecloth, stored in thermo bottle. The bottle was
795 flushed with CO₂, and maintained at 39°C until incubations.

796 Corn silage sample (0.5 g), artificial saliva (40 mL), and ruminal fluid (10 mL)
797 were added to each tube, and tubes were saturated with CO₂ to maintain an anaerobic
798 environment. Incubations were performed in a water bath for 48 h at 39°C with constant
799 stirring. Duplicate samples were incubated in three separate *in vitro* runs. In each
800 incubation, two additional tubes containing only artificial saliva and ruminal fluid, and
801 another two tubes with a forage with a known IVDMD were included as blank and
802 standard, respectively. After incubation, residues were filtered in analytical filter paper
803 N° 40 and dried to calculate the DM disappearance.

804 The determination of *in vitro* NDF digestibility (IVNDFD) was performed
805 according to the methodology proposed by Tilley and Terry (1963) using the artificial
806 rumen Daisy II Fermenter® (Ankom Technology, Macedon, NY, USA), modified by
807 Holden (1999) (Holden, 1999). Approximately 0.5 g of silage was weighed into a non-
808 woven textile bag (100 g/m²). The ruminal fluid was collected as previously mentioned
809 in the methodology of IVDMD. The samples, 1600 mL of buffer solution, and 400 mL
810 of ruminal fluid were added to the jars. Subsequently, the jars were saturated with CO₂,
811 and constantly rotated at 39 °C for 48 h. The bags were drained, washed with water
812 until the water remained clean, and subsequently, the bags were frozen. The bags were
813 analyzed for NDF concentration in the fiber determiner (TE-149, Tecnal, Piracicaba,
814 SP, Brazil; Mertens et al., 2002). Duplicate samples were incubated in three separate *in*
815 *vitro* runs. In each incubation, two additional empties bags and another two bags with a
816 sample with known IVNDFD were included as blank and standard, respectively.

817

818 ***Experiment 2***

819 ***Ensiling and treatments***

820 The whole-plant corn hybrid DKB 290 Pro3 (Dekalb, Brazil) was harvested at a
821 half-milk line stage, chopped (theoretical length of cut of 10 mm), and the material was
822 manually ensiled in 200-L plastic drums (600 kg/m³ of FM). The corn was mixed with
823 SSPO at the compaction process. The silages were stored for 5 months before the
824 feeding trial. The tested levels were 0, 10, and 30 mg of lignocellulolytic enzymes/kg of
825 fresh matter, and those were chosen based on the results from experiment 1.

826

827 ***Animals and experimental diets***

828 The experiment was approved by the Committee for Use of Animals in
829 Experimentation of the State University of Maringá, PR (protocol 4361101018), and the
830 study fully complied with the ethical principles of animal experimentation prepared by
831 the Brazilian College of Animal Experimentation. The experiment was conducted at the
832 Iguatemi Experimental Farm, PR, Brazil.

833 Nine lactating Saanen goats (62.68 ± 7.62 kg of BW 44 \pm 8 days in milk; 2.91 ± 0.81
834 kg/d of milk mean \pm SD) were used in a triplicated 3×3 Latin square design, with three
835 periods and three treatments. The animals were maintained in individual stalls with free
836 access to water and diet.

837 Experimental diets were formulated to meet the nutritional requirements of
838 lactating goats weighing 60 kg and a milk yield of 3 kg/day of milk, according to NRC
839 (2007) (Table 5). The same concentrate was used in all the experimental diets, and those
840 diets differed according to the enzymatic levels added in the whole-plant corn at
841 ensiling.

842 Each experimental period lasted 22 d, with 17 d for adaptation to the treatments and
843 5 d of sampling. The animals were fed *ad libitum* with total mixed rations, and diets
844 were offered daily at 8:30 and 16:00 h, allowing 5 to 10% of refusal. The goats were
845 milked twice daily at 8:00 and 16:00 h.

846

847 ***Sampling***

848 The amount of feed provided and refusals from each animal were weighed and
849 recorded daily during the whole experiment. Feed was sampled from d 18 to d 21, and
850 refusals and feces from d 19 to d 22 of each experimental period. Fecal samples were
851 collected directly from the rectum at 8:30 and 16:30 hrs. The samples were stored at -

852 20°C, posteriorly thawed, dried in a convection oven at 55°C for 72 h, ground in Wiley
853 mill (Marconi MA340, Piracicaba, Brazil) to pass through a 2 mm sieve for indigestible
854 neutral detergent fiber (iNDF) determination and a 1 mm sieve for chemical
855 composition analysis. Fecal and refusal samples were pooled for each goat to obtain a
856 composite sample per animal per period.

857 Between d 19 and d 22 of each period, the milk yield was recorded daily, and the
858 energy-corrected (ECM) and fat-corrected milk (FCM) yield were calculated according
859 to Tyrrell and Reid (1965) and NRC (2001), respectively. Milk samples were obtained
860 from six consecutive milkings between d 20 and 22 of each period and divided into two
861 aliquots. The first aliquot, approximately 50 mL of the milk, was kept at room
862 temperature and preserved with 2-bromo-2-nitropropane-1,3-diol (Bronopol, San
863 Ramon, CA, USA) for determination of milk fat, protein, lactose, and milk urea
864 nitrogen (MUN). The second aliquot (2 mL of milk), without preservative, was frozen
865 at -80°C to determine antioxidants capacity.

866 Spot urine samples were collected on d 21 of each period 4 h after feeding to
867 estimate the efficiency of microbial protein synthesis, according to Chen and Gomes
868 (1992).

869

870 ***Chemical composition and antioxidants***

871 The chemical composition was determined on samples ground to pass a 1 mm
872 sieve, according to the methodologies previously described in experiment 1.

873 The iNDF was used as an intern marker to estimate the fecal production (Huhtanen
874 et al., 1994); thus the concentration of iNDF in feces, feed and refusals were performed
875 to determine the apparent total tract digestibility of DM, and nutrients. Samples were
876 incubated for 288 h in ruminal cannulated dairy cows fed a diet composed with 60% of

877 corn silage and 40% of grain mix (DM basis). After removal from the rumen, the bags
878 were drained, washed with water until the water remained clean, and analyzed for NDF.

879 The milk extract to determine TAC, polyphenols, and reducing power were
880 performed with the addition of 1 mL of milk and 9 ml of methanol. Posteriorly, samples
881 were vortexed for 5 min and centrifuged at $2,500\text{ g} \times 10\text{ min}$, and the supernatant was
882 used in the following analysis.

883 The reducing power was determined according to Zhu et al. (2002), with the
884 modifications of Santos et al. (2014). The absorbance was determined in a UV-Vis
885 spectrophotometer (PC 300 ThermoScientific, Waltham, MA, USA) and expressed as
886 gallic acid equivalents (GAE; mg/L of milk).

887 Total antioxidant capacity (TCA) was determined as described by Rufino et al.
888 (2007) with the addition of radical $\text{ABTS}^{+\cdot}$ (2,2-azinobis-(3-ethyl-benzothiazolin-6-
889 sulfonic acid)) to the extract. The absorbance was determined in a UV-Vis
890 spectrophotometer (PC 300 ThermoScientific, Waltham, MA, USA) and expressed in %
891 of $\text{ABTS}^{+\cdot}$ degradation (% of ABTS degradation).

892 Total polyphenols were determined according to Singleton and Rossi (1965). The
893 absorbance was determined in a UV-Vis spectrophotometer (PC 300 ThermoScientific,
894 Waltham, MA, USA) and expressed as gallic acid equivalents (EAG; g/L of milk).

895

896 *Milk and urine analyses*

897 The fat, protein, and lactose concentrations in milk were determined by infrared
898 spectroscopy (Bentley model 2000, Chaska, MN. USA) following procedure 972.16 of
899 AOAC (1990). MUN concentration was determined by a colorimetric method with the
900 Berthelot reaction (Chemspec 150, Chaska, MN, USA).

901 Creatinine concentration was analyzed in a semi-automatic biochemistry analyzer
 902 (Bio 2000–Bioplus, São Paulo, Brazil) using a commercial kit (Gold Analisa, Belo
 903 Horizonte, Brazil). Urine volume (L/d) was estimated by dividing the average daily
 904 creatinine excretion rate reported for goats of 26.05 mg/kg BW, according to Da
 905 Fonseca et al. (2006). Diluted urine was analyzed for allantoin, uric acid, xanthine and
 906 hypoxanthine concentrations according to methods of Chen and Gomes (1992). The
 907 purines derivatives (PD) absorbed (mmol/d) and intestinal flow of microbial nitrogen
 908 were calculated based on the following equations described by Chen and Gomes (1992):
 909 $PD = 0.84 \times + (0.150BW^{0.75}e^{-0.25X})$, and $MN (g/d) = 0.727 \times PD$, where X = purines
 910 derivatives excreted.

911

912 *Statistical analysis*

913 The outcomes from experiment 1 were analyzed using the GLIMMIX procedure of
 914 SAS (Statistical Analysis System, 9.4, Cary, NC, USA) with the following model: $Y_{ij} =$
 915 $\mu + E_i + e_{ij}$, where Y_{ij} = dependent variable; μ = overall mean; E_i = fixed effect of
 916 enzyme levels ($i = 0$ to 30 mg of enzyme/kg of FM); and e_{ij} = residue ($j = 1$ to 4). Data
 917 from IVDMD and IVNDFD were analyzed using the same model but including a
 918 random effect of the run (1 to 3). Means were compared by linear and quadratic
 919 orthogonal contrasts.

920 Data from experiment 2 were analyzed using the GLIMMIX procedure of SAS,
 921 with the following model: $Y_{ijk} = \mu + E_i + S_j + Ak(j) + Pl + e_{ijklm}$, where $Y_i =$
 922 dependent variable; μ = overall mean; E_i = fixed effect of enzyme levels ($i = 0$ to 30 mg
 923 of enzyme/kg of FM); S_j = fixed effect of square ($j = 1$ or 3); $Ak(j)$ random effect of
 924 goat nested within square ($j = 1$ to 9); Pl = fixed effect of period ($l = 1$ to 3); and e_{ijklm}
 925 = residue. Means were compared by linear and quadratic orthogonal contrasts. The

926 coefficients for orthogonal contrasts were determined using the IML procedure of SAS.
927 Differences were declared significant at $P \leq 0.05$ and tendencies if $P > 0.05$ and ≤ 0.10 .
928

929 RESULTS

930 *Experiment 1*

931 Spent substrate from *Pleurotus ostreatus* cultivation showed enzymatic activity for
932 three enzymes that degrade lignin. These enzymes were laccase (390 U/g/min), lignin
933 peroxidase (176 U/g/min), and manganese peroxidase (12.6 U/g/min; Table 1).
934 Enzymatic activity was also observed for carboxymethylcellulase (2,894 U/g/min);
935 however, no activity was observed for xylanase and mannanase, which are enzymes
936 capable of degrading hemicellulose.

937 The chemical composition of the whole-plant corn used in experiment 1 is
938 described in Table 2. There was no effect of SSPO ($P > 0.11$) on the recovery of
939 hemicellulose and lignin, while NFC recovery increased linearly ($P = 0.01$) (Table 3)
940 with the addition of SSPO. The NDF recovery tended to decrease quadratically ($P =$
941 0.09), which was associated with a quadratic reduction ($P < 0.01$) on cellulose recovery.

942 The addition of SSPO linearly decreased ($P = 0.02$) silage pH, while DM
943 concentration increased quadratically ($P = 0.03$; Table 4). The concentrations of NDF
944 and ADF decreased quadratically ($P = 0.02$; and < 0.01 , respectively). Due to the
945 enzymatic activities from SSPO, NDF decreased by 14.1%, and ADF decreased by
946 19.5% at the nadir point achieved with 10 mg/kg FM of enzymes compared to the
947 control treatment. Enzyme concentration also affected carbohydrate fractionation of the
948 corn silage. Fraction A+B1 increased quadratically ($P = 0.04$), whereas fraction C and
949 B2 decreased quadratically ($P < 0.01$; 0.01, respectively). The concentration of NFC
950 decreased linearly ($P = 0.01$) with the addition of SSPO at ensiling. The concentrations

951 of lignin ($P = 0.03$), cellulose ($P < 0.01$) and ADIN ($P < 0.02$) decreased quadratically
952 with higher doses of SSPO. Compared with the control, lignin concentration decreased
953 by 9.07% and cellulose by 22.1% at the nadir point achieved with 20 and 10 mg of
954 enzyme/kg FM respectively. Minimum values of ADIN were also achieved with 10 mg
955 of enzymes/kg of FM. The addition of SSPO at ensiling did not change ($P > 0.42$) the
956 concentrations of EE and CP in corn silage.

957 The addition of SSPO at ensiling increased IVDMD quadratically ($P < 0.01$), and
958 the IVNDFD increased linearly ($P < 0.01$). Regarding the antioxidants, the polyphenols
959 and reducing power were reduced linearly ($P < 0.01$; and < 0.01 respectively) as SSPO
960 levels increased. Other antioxidant parameters did not differ among SSPO levels ($P >$
961 0.05).

962 Lignin monomers were also affected by SSPO. *p*-Hydroxyphenyl tended to
963 decrease quadratically ($P = 0.06$), while syringyl tended to decrease linearly ($P = 0.07$)
964 and consequently decreased the syringyl:guaiacyl ratio linearly ($P = 0.01$) in the cell
965 wall of corn silage.

966

967 ***Experiment 2***

968 The intake of DM and nutrients were not affected ($P > 0.24$) by treatments (Table
969 6). Apparent total tract digestibility of ADF increased quadratically ($P < 0.01$), and
970 NDF digestibility tended to increase quadratically ($P = 0.10$) with SSPO levels. The
971 apparent digestibility of other nutrients was not affected by treatments ($P > 0.15$). The
972 absorption of purine derivatives and intestinal flow of microbial nitrogen were not
973 affected by SSPO levels ($P > 0.65$; Table 7).

974 The concentration of polyphenols in the milk increased linearly with SSPO levels
975 ($P < 0.01$; Table 8). Milk yield, composition, and the other antioxidants parameters did
976 not differ among SSPO levels ($P > 0.66$; 0.30; 0.13, respectively).

977

978

DISCUSSION

979 The fungi *Pleurotus ostreatus* produces enzymes capable of breaking down cell
980 wall polymers (Isikhuemhen and Mikiashvilli,2009), consequently reducing NDF and
981 ADF and increasing NFC concentration in silage, as shown in experiment 1.
982 Interestingly, ADF reduction was more pronounced than NDF because the fractions that
983 compose ADF were decreased (i.e., lignin and cellulose), while hemicellulose, which
984 represented approximately 43% in NDF, did not change. These results corroborate with
985 the observed by Colombatto et al. (2004), who reported a reduction in NDF and ADF
986 concentration in corn silage treated with a mixture of fibrolytic enzymes from different
987 fungi.

988 As expected, in this study, a decrease of cellulose was paired with a simultaneous
989 increase in NFC concentration. Cellulose degradation can explain an increase of NFC
990 concentration when forages are treated with fibrolytic enzymes to glucose and
991 oligosaccharides (that constitute NFC) through cellulase activity. The NFC can be
992 rapidly assimilated by ruminal microorganisms and positively correlates with volatile
993 fatty acid syntheses (Getachew et al., 2004). Carbohydrate A+B1 fraction corresponds
994 to the sugars and starch (Sniffen et al., 1992); therefore, the increase in this fraction by
995 SSPO is correlated to the increase in NFC concentration.

996 The reduction of lignin concentration in the corn silage observed in experiment 1,
997 was certainly a result of lignocellulolytic enzymes present in SSPO (laccase, lignin
998 peroxidase, and manganese peroxidase) capable of degrading phenolic and non-phenolic

999 compounds (Wong, 2009). Machado et al.(2020) observed a 33% reduction in lignin
1000 concentration in corn silage treated at ensiling with a complex of lignocellulolytic
1001 enzymes produced in the laboratory by *Pleurotus ostreatus*. However, this reduction
1002 observed by Machado et al. (2020) was approximately 3 times greater than in the
1003 current study. The difference could have potentially been caused by different
1004 proportions of enzymatic activities, lignin concentration, and arrangement of the plant
1005 cell wall components between studies. In the current study, the carboxymethylcellulase
1006 activity in SSPO was predominant, and consequently, there were a lower proportion of
1007 enzymes that degrade lignin.

1008 Khattab et al. (2013) also reported a reduction of cellulose and lignin when applied
1009 *Pleurotus ostreatus* on the rice harvest wastes. However, the proportion of reduction in
1010 lignin in this residue was around 50%, while in the present study, it was 11.3%. The
1011 reduction of cellulose concentration using SSPO in our study was 22.2%, whereas the
1012 authors reported a decrease of 3.7% when treating rice wastes. The different magnitude
1013 of reductions of fiber components may have occurred for several reasons, such as the
1014 anaerobic environment of silage production since oxygen is required for laccase cycling
1015 (Wong, 2009); the difference of *Pleurotus ostreatus* strains, as each strain can produce
1016 enzymes in different proportions and amounts (Membrillo et al., 2008); differences in
1017 chemical composition between the rice residue and the corn plant, due enzyme-feed
1018 specificity (Beauchemin et al., 2003); as well as, the way that the feeds were treated,
1019 since in the silage the enzymes were added through SSPO, and in the rice residue the
1020 fungus was placed to grow directly in the material.

1021 The reduction in ADIN concentration using the enzymes from SSPO follows the
1022 results observed with the reduction of lignin and might be likely through the cleavage of
1023 the linkage between N and lignin. The ADIN is recognized by its very low ruminal

1024 degradation and intestinal digestibility (Lanzas et al., 2008); thus, releasing nitrogen
1025 associated with lignin may increase the metabolizable protein for the animal. The
1026 reduction of carbohydrate fraction C was a consequence of lignin degradation.

1027 Although white-rot mushroom, as *Pleurotus ostreatus*, produces enzymes capable
1028 of degrading hemicellulose, such as xylanase and mannanase (Machado et al., 2020), in
1029 our study, the SSPO did not present hemicellulase activity (xylanase and mannanase).
1030 The absence of hemicellulase activity may be occurred due to culture conditions
1031 (Qinnghe et al., 2004; Membrillo et al., 2008) or be a characteristic of the strain used in
1032 our study (Membrillo et al., 2008).

1033 Applying SSPO to corn silage also altered lignin monomers. The effect observed in
1034 *p*-hydroxyphenyl and syringyl concentration is likely a result of cleavage provided by
1035 ligninolytic enzymes (Wong, 2009). Machado et al. (2020) reported the opposite for the
1036 concentration of polyphenols and lignin monomers in corn silage, since polyphenol
1037 concentration increased with the addition of lignocellulolytic enzymes, accompanied by
1038 an increase in the concentration of lignin monomers.

1039 The increase in IVDMD was likely due to a modification in corn silage chemical
1040 composition due to the reduction of lignin and cellulose concentration simultaneous to
1041 an increase of NFC. This increase in IVDMD corroborates with Fazaeli et al. (2004),
1042 who fed cows at the end of lactation with 30% wheat straw in the diet and observed that
1043 diet with wheat straw previously treated with *Pleurotus ostreatus* showed greater DM
1044 digestibility compared to the diet with untreated wheat straw.

1045 In experiment 2, alterations in corn silage composition were not consistent with
1046 experiment 1 that may be due to differences in corn hybrids and silo type. No difference
1047 was observed between the enzymatic levels in the silage chemical composition (data are
1048 not shown). However, the addition of SSPO increased quadratically total tract ADF and

1049 NDF digestibility that may be explained due by the loosening effect in the lignin-
1050 hemicellulose-cellulose complex. This loosening effect was previously reported by
1051 Taniguchi et al. (2005); it might favor the access to fiber carbohydrate and became
1052 easier the degradation of fiber by rumen microbial for the goats.

1053 Rojo et al. (2015) observed that goats fed with a similar concentration of CP, and
1054 NDF, and proportion of forage:concentrate in the diet presented a higher NDF
1055 digestibility when the TMR was treated with a cellulase, that corroborates with the
1056 tendency observed in our study. However, the authors also observed that the enzymes
1057 increased OM and DM digestibility of the diets, which was not observed in our study.

1058 The SSPO did not affect milk yield and nutrient concentration. However, the results
1059 of those variables in this study were similar to those related by Ferro et al. (2017) in
1060 Saanen goats milk, where the milk composition of goats was characterized.

1061 Despite that the concentration of total polyphenols in the silage was similar in both
1062 treatments (477 ± 6.09 mg/100 g of diet, DM basis), the concentration of polyphenols in
1063 milk increased with the addition of SSPO. The total polyphenol combines free, soluble
1064 conjugated, and insoluble-bound phenolic compounds (Madhujith and Shahidi, 2009).
1065 Therefore, the POSS may be released free phenolic compounds by the cleavage of
1066 insoluble-bound phenolic compounds present in the lignin, favoring the absorption and
1067 the transfer for the milk. The total polyphenols analysis determines polyphenols
1068 concentration. It does not distinguish the three categories in the quantification, and this
1069 justifies the absence of difference among polyphenols concentration in the diets, leading
1070 to a similar concentration.

1071

1072

CONCLUSION

1073
1074 Therefore, we concluded that adding 10 mg of lignocellulolytic enzymes from
1075 SSPO per kg fresh matter of whole-plant corn at ensiling presented the greatest effect on
1076 animal performance due to increasing total tract ADF digestibility. Thus, POSS may be
1077 an option at ensiling to improve fiber digestibility of lactating animals. Future studies
1078 should investigate the effect of those enzymes in WPCS on other lactating species, such
1079 as dairy cows, which may show an increase in milk yield.

1080

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1083 (Conceptualization, supervision, funding acquisition, writing - review, and editing);
1084 C.R.A. (Conceptualization, and investigation); E.M. (Conceptualization, methodology,
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TABLES

1299 Table 1. Characterization of enzymatic activity of spent substrate from *Pleurotus*
1300 *ostreatus* cultivation (SSPO)

Enzymes	Enzymatic activity (U/g/min) ¹
Laccase	390
Lignin peroxidase	176
Manganese peroxidase	12.6
Carboxymethylcellulase	2,894

1301 ¹ The enzymatic activities were identified based on the final product obtained in the reaction.

1302 Enzymatic activity was expressed in U/g of SSPO/min.

1303 Table 2. Chemical composition (% of dry matter) of whole-plant corn at ensiling
1304 (Experiment 1)

Chemical composition (%DM, unless otherwise stated)	
Dry matter (g/100 g)	29.3
Organic matter	96.4
NDF	47.9
ADF	27.4
Lignin	5.64
Crude protein	7.43
EE	2.40
NFC	38.7

1305 NDF= neutral detergent fiber; ADF= acid detergent fiber; Lignin = acid detergent lignin; EE=
1306 ether extract; NFC= non-fiber carbohydrates.

1307 Table 3. Recovery of fiber fractions (%) in corn silage treated with spent substrate from
 1308 *Pleurotus ostreatus* cultivation, after 60 d of storage (Experiment 1)

Recovery (%)	Treatments (mg enzymes/kg FM) ¹				SEM	P-value	
	0	10	20	30		Linear	Quadratic
Dry matter	86.7	93.3	88.6	92.7	1.83	0.14	0.51
NDF	90.1	83.0	84.7	86.8	2.50	0.48	0.09
NFC	78.4	103.3	89.1	98.4	3.56	0.01	0.05
Cellulose	98.9	81.1	85.4	86.0	2.21	0.01	<0.01
Hemicellulose	85.1	85.5	85.7	89.7	4.22	0.47	0.68
Lignin	83.1	81.2	75.5	79.2	2.27	0.11	0.24

1309 FM= fresh matter; NDF= neutral detergent fiber; EE= ether extract; NFC= non-fiber
 1310 carbohydrates ¹ = treatments corresponded to the following enzymatic activities: 0 (0 U/min/kg
 1311 FM), 10 (1,100 U/min/kg FM), 20 (2,200 U/min/kg FM), 30 (3,300 U/min/kg FM).

1312 Table 4. Chemical composition, carbohydrate fractionation (% of DM), antioxidant
 1313 capacity and lignin monomers (ug/mg protein-free cell wall) in corn silage treated with
 1314 spent substrate from *Pleurotus ostreatus* cultivation, after 60 d of storage (Experiment
 1315 1)

Parameters (%)	Treatments (mg enzymes/kg FM) ¹				SEM	P-value	
	0	10	20	30		Linear	Quadratic
pH	3.61	3.63	3.59	3.56	0.014	0.02	0.12
Chemical composition							
Dry matter	26.3	29.6	27.8	28.5	0.508	0.05	0.03
Organic matter	94.7	95.5	94.8	95.6	0.292	0.04	0.86
NDF	49.7	42.7	45.7	44.8	1.53	0.03	0.02
ADF	29.7	23.9	26.0	25.1	0.627	<0.01	<0.01
Lignin	5.40	4.91	4.79	4.82	0.105	<0.01	0.03
Hemicellulose	20.0	18.8	19.7	19.7	0.819	0.96	0.47
Cellulose	24.4	19.0	21.0	20.2	0.570	<0.01	<0.01
ADIN	5.25	4.54	4.92	4.96	0.113	0.38	<0.01
Crude protein	7.41	7.36	7.53	7.21	0.161	0.55	0.42
EE	2.61	2.65	2.68	2.57	0.087	0.78	0.43
NFC	34.9	40.6	38.9	41.2	0.871	<0.01	0.07
Fraction B2	50.54	46.0	47.5	46.0	0.932	0.01	0.13
Fraction C	8.11	6.37	6.44	6.03	0.274	<0.01	0.03
Fraction A+B1	41.3	47.6	46.1	48.0	0.954	<0.01	0.04
In vitro Digestibility (%)							
IVDMD	56.2	62.0	58.4	58.3	5.27	<0.01	<0.01
IVNDFD	48.8	55.1	48.2	53.4	4.66	<0.01	0.24
Antioxidant capacity							
TAC ²	758	757	756	756	0.479	0.05	0.74
Flavonoids ³	120	110	104	109	1.78	0.22	0.31
Reducing power ⁴	267	252	241	194	13.3	<0.01	0.25
Poliphenols ⁵	674	581	532	475	22.4	<0.01	0.45
Monomers lignin							
p-Hydroxyphenyl	12.8	6.23	12.8	20.1	3.11	0.07	0.06
Guaiacyl	118	92.1	116	118	9.81	0.59	0.18
Syringyl	1123	76.7	86.8	80.0	9.24	0.07	0.15
Syringyl:guaiacyl	0.958	0.832	0.754	0.677	0.063	0.01	0.71

1316 FM= fresh matter; NDF= neutral detergent fiber; ADF= acid detergent fiber; EE= ether extract;
 1317 NFC= non-fiber carbohydrates, TAC= total antioxidant capacity; ¹ = treatments corresponded to
 1318 the following enzymatic activities: 0 (0 U/min/kg FM), 10 (1,100 U/min/kg FM), 20 (2,200
 1319 U/min/kg FM), 30 (3,300 U/min/kg FM); ²= expressed in mM Trolox/100 g of silage (DM
 1320 basis); ³= expressed in mg of equivalent quercetin/100 g of silage (DM basis); ⁴= expressed in
 1321 mg of gallic acid/100 g of silage (DM basis); ⁵= expressed in mg of gallic acid/100 g of silage
 1322 (DM basis).

1323 Table 5. Ingredient and chemical composition of the experimental diets (% DM, unless
 1324 otherwise stated, Experiment 2)

Item	Treatments (mg enzymes/kg FM) ¹		
	0	10	30
Ingredients			
Corn silage	60.0	60.0	60.0
Ground corn	19.3	19.3	19.3
Soybean meal	19.1	19.1	19.1
Mineral and vitamin supplement	1.0	1.0	1.0
Dicalcium phosphate	0.6	0.6	0.6
Chemical composition			
DM (% of fresh matter)	53.0	53.0	53.0
Organic matter	94.4	94.7	95.1
Crude protein	15.3	15.3	15.3
Neutral detergent fiber	34.9	35.9	35.2
Acid detergent fiber	19.0	20.1	19.2
Lignin	3.16	3.05	2.78
Ether extract	2.76	2.72	2.64
ADIN, % of ADF	5.20	4.55	4.96

1325 ADIN = Acid detergent insoluble nitrogen; ¹ = treatments corresponded to the following
 1326 enzymatic activities: 0 (0 U/min/kg FM), 10 (1,100 U/min/kg FM), 30 (3,300 U/min/kg FM).

1327 Table 6. Dry matter (DM) and nutrient intake (kg/d) and total tract apparent digestibility
 1328 (%) in lactating goats fed diets containing corn silage treated with spent substrate from
 1329 *Pleurotus ostreatus* cultivation (Experiment 2)

Items	Treatments (mg enzymes/kg FM) ¹			SEM	P-value	
	0	10	30		Linear	Quadratic
Intake						
Dry matter	2.09	2.04	2.06	0.106	0.76	0.62
Organic matter	1.93	1.92	2.01	0.103	0.30	0.57
Crude protein	0.322	0.315	0.330	0.0177	0.47	0.43
NDF	0.688	0.703	0.720	0.0353	0.31	0.87
ADF	0.372	0.394	0.391	0.0202	0.37	0.31
Ether extract	0.0574	0.0557	0.0573	0.00502	0.96	0.68
NFC	0.865	0.833	0.885	0.0511	0.43	0.24
TND	1.483	1.420	1.478	0.0732	0.89	0.28
Total-tract digestibility						
Dry matter	72.1	72.3	71.6	0.966	0.52	0.71
Organic matter	72.7	73.3	73.8	0.83	0.26	0.76
Crude protein	72.8	74.0	75.1	1.36	0.15	0.73
NDF	50.5	53.5	53.3	1.38	0.09	0.10
ADF	50.2	54.4	52.9	1.36	0.10	<0.01
Ether extract	82.1	81.5	82.1	1.66	0.95	0.77
NFC	89.6	88.7	88.8	0.815	0.50	0.53
TDN	73.1	73.5	73.8	0.853	0.42	0.84

1330 NDF= neutral detergent fiber; ADF= acid detergent fiber; NFC= non-fiber carbohydrates;
 1331 TDN= total digestible nutrient; ¹ = treatments corresponded to the following enzymatic
 1332 activities: 0 (0 U/min/kg FM), 10 (1,100 U/min/kg FM), 30 (3,300 U/min/kg FM).

1333 Table 7. Efficiency of protein synthesis in lactating goats fed diets containing corn
 1334 silage treated with spent substrate from *Pleurotus ostreatus* cultivation (Experiment 2)

Items	Treatments (mg enzymes/kg FM) ¹			SEM	P-value	
	0	10	30		Linear	Quadratic
Purine derivatives ²	17.4	18.2	17.6	1.55	0.97	0.65
Microbial nitrogen ³	12.6	13.2	12.8	1.13	0.97	0.65

1335 ²= purines derivatives absorbed was expressed in mmol/d; ³= microbial nitrogen was expressed
 1336 in g/d

1337 Table 8. Milk yield and composition and concentration of antioxidants in milk of goats
 1338 fed diets containing corn silage treated with spent substrate from *Pleurotus ostreatus*
 1339 cultivation (Experiment 2)

Items	Treatments (mg enzymes/kg FM) ¹			SEM	P-value	
	0	10	30		Linear	Quadratic
Milk Yield, kg/d						
Actual	2.92	2.88	2.92	0.380	0.95	0.76
Fat-corrected ²	2.54	2.49	2.52	0.350	0.94	0.66
Energy-corrected ³	2.76	2.70	2.75	0.380	0.97	0.61
Yield, g/d						
Fat	91.6	89.2	90.0	13.2	0.85	0.56
Protein	86.2	84.0	87.3	12.0	0.73	0.55
Lactose	124	123	123	16.7	0.99	0.85
Total Solids	324	316	319	49.9	0.81	0.67
Concentration, % <i>unless otherwise stated</i>						
Fat	3.09	3.09	3.06	0.136	0.76	0.94
Protein	2.95	2.93	3.00	0.073	0.31	0.39
Lactose	4.22	4.26	4.22	0.0654	0.86	0.30
Total Solids	11.0	10.9	10.9	0.386	0.44	0.72
MUN, mg/dL	12.2	12.6	11.9	1.24	0.41	0.37
Antioxidant						
TAC ⁴	19.7	19.2	21.5	3.20	0.13	0.36
Polyphenols ⁵	1.00	1.03	1.23	0.222	<0.01	0.51
Reducing power ⁶	145	145	154	17.9	0.48	0.86

1340 TAC = total antioxidant capacity; ¹ = treatments corresponded to the following
 1341 enzymatic activities: 0 (0 U/min/kg FM), 10 (1,100 U/min/kg FM), 30 (3,300 U/min/kg
 1342 FM); ² = FMC (kg/d) = 4% fat-corrected (kg/d) = (0.4 × milk yield (kg/d)) + (15.0 × fat
 1343 yield (kg/d)) [15]; ³ ECM (kg/d) = (0.327 × kg of milk) + (12.95 × kg of fat) + (7.2 × kg
 1344 of protein) [33]; ⁴ = expressed in % of ABTS degradation; ⁵ = expressed in g of
 1345 equivalent gallic acid/L of milk; ⁶ = expressed in mg of gallic acid/L of milk.

1346 **IV. Effects of lignocellulolytic enzymes on the fermentation profile,**
1347 **chemical composition, and in situ ruminal disappearance of whole-**
1348 **plant corn silage**

1349

1350 (Manuscript style and form consistent with the Instructions for Authors of the Journal

1351 Animal Feed Science and Technology)

1352

1353 **Abbreviations:** ADF, acid detergent fiber; aNDF, neutral detergent fiber assayed with a

1354 heat-stable alpha-amylase and inclusive of residual ash; DM, dry matter; FM, fresh

1355 matter; uNDF, undigested neutral detergent fiber; U, unit of enzyme activity; WPC,

1356 whole-plant corn; WPCS, whole-plant corn silage; WSC, water-soluble carbohydrates

1357

1358 **ABSTRACT:** The objective of this study was to examine the enzymatic activity of an
1359 enzymatic complex produced by *Pleurotus ostreatus* in different pH; and the effects of

1360 adding increasing levels of this enzymatic complex on the fermentation profile,

1361 chemical composition, and *in situ* ruminal disappearance of whole-plant corn silage

1362 (WPCS) at the onset of fermentation and after 30 d of ensiling. The lignocellulolytic

1363 enzymatic complex was obtained through *in vitro* cultivation of *Pleurotus ostreatus*.

1364 The activities of laccase, lignin peroxidase, manganese peroxidase, endo- and exo-

1365 glucanase, xylanase, and mannanase were determined at pH 3, 4, 5, and 6. Afterward,

1366 five enzymatic complex levels were tested in a completely randomized block design in

1367 the levels of 0; 9; 18; 27; and 36 mg of lignocellulosic enzymes/kg of fresh matter (FM)

1368 of whole-plant corn. There were four replicates per treatment (vacuum-sealed bags) per

1369 opening time. The bags were opened after 1, 2, 3, and 7 d of ensiling (onset of

1370 fermentation period) and after 30 d of storage to evaluate the fermentation, chemical

1371 composition, and *in situ* disappearance of WPCS. Laccase showed highest activity at
1372 pH 5 ($P < 0.01$), whereas manganese peroxidase and lignin peroxidase had a higher
1373 activity at pH 4 ($P < 0.01$; < 0.01 , respectively). There was no interaction between the
1374 enzymatic complex and days of fermentation ($P > 0.11$). Also, there was no effect of
1375 enzymatic complex addition ($P > 0.27$) on the fermentation profile and chemical
1376 composition at the onset of fermentation. The concentration of WSC decreased
1377 quadratically at the onset of fermentation ($P = 0.02$) through the fermentation, leading
1378 to a quadratic increase of lactic acid ($P = 0.01$) and a linear increase of acetic acid ($P =$
1379 0.02). As a consequence of increasing those organic acid concentrations, pH decreased
1380 quadratically ($P = 0.01$). Lignin concentration decreased linearly ($P = 0.04$) with the
1381 enzymatic complex levels at 30 d of storage; however, other nutrients ($P > 0.12$) and
1382 fermentation profile ($P > 0.11$, Table 5) did not change. The addition of
1383 lignocellulolytic enzymatic complex from *Pleurotus ostreatus* cultivation decreased
1384 lignin concentration in WPCS after 30 d of ensiling; however, it was not enough to
1385 affect the *in situ* disappearance.

1386 **Keywords:** fibrolytic enzyme, laccase, lignin, *Pleurotus ostreatus*, white-rot fungi

1387

1388 1. INTRODUCTION

1389 Whole-plant corn silage (WPCS) is the main forage source in dairy diets (Grant
1390 and Ferraretto, 2018). However, just approximately 560 g/kg of neutral fiber detergent
1391 (NDF) in the WPCS is digestible (Ferraretto and Shaver, 2015), and it is highly related
1392 to some factors, such as lignin. Lignin is a phenolic polymer that constitutes plant cell
1393 walls and negatively affects fiber degradability (Jung and Allen, 1995) due to the cross-
1394 linking of lignin to arabinoxylans (Hatfield et al., 2017). Meanwhile, reducing lignin

1395 concentration by changing genotypes (Oba and Allen, 2000) or cleaving lignin linkage
1396 are strategies to potentially increase fiber degradability (Machado et al., 2020).

1397 Some organisms, such as white-rot fungi, are known to produce lignocellulolytic
1398 enzymes as a mechanism to obtain energy and nutrients from fibrous substrates
1399 (Manavalan et al., 2015). *Pleurotus ostreatus*, a white-rot fungus, produces enzymes
1400 that degrade cellulose (endo- and exo-glucanase), hemicellulose (xylanase and
1401 mannanase), and lignin, such as laccase, lignin peroxidase, and manganese peroxidase
1402 (Leonowicz et al., 1999). Although these enzymes have the potential to break down
1403 lignin and possibly increase fiber degradability, few research trials have focused on
1404 evaluating the effects of adding these enzymes to whole-plant corn (WPC) at ensiling.

1405 Recently we reported the potential of lignocellulolytic enzymes produced by
1406 *Pleurotus ostreatus* in reducing the concentration of lignin, cellulose, and hemicellulose
1407 and increasing *in vitro* degradability and antioxidant capacity of WPCS (Machado et al.,
1408 2020). However, no study on the fermentation profile after 30 d of fermentation; and
1409 fermentation profile and chemical composition of the WPC at the onset of fermentation
1410 period with lignocellulolytic enzymatic complex from *Pleurotus ostreatus* cultivation.

1411 Therefore, the objective of this study was to evaluate the enzymatic activity of
1412 each fibrolytic enzyme present in the enzymatic complex at different pH and examine
1413 the effects of adding increasing levels of the enzymatic complex produced in the
1414 cultivation of *Pleurotus ostreatus* on the fermentative profile, chemical composition,
1415 and ruminal disappearance of WPCS along of the days of fermentation. We
1416 hypothesized that treating WPCS with the lignocellulolytic enzymatic complex from
1417 *Pleurotus ostreatus* would decrease the concentration of aNDF and lignin in the
1418 material and consequently increase *in situ* dry matter disappearance.

1419

1420 2. MATERIAL AND METHODS

1421

1422 2.1. Production of the enzymatic complex and their activities

1423 A sample of *Pleurotus ostreatus* (strain number 1833) was purchased from the
1424 company DSMZ[®] (Leibniz Institute DSMZ, Braunschweig, Germany) and propagated
1425 in Petri dishes using Potato Dextrose Agar (Sigma-Aldrich Co., St. Louis, MO) as
1426 culture medium and incubated for 10 d at room temperature. After grown, 10 mm-
1427 diameter disk of *Pleurotus ostreatus* was incubated in 125 mL flasks with 25 mL of
1428 liquid culture medium (0.5% sugarcane diluted in distilled water) and 0.5 g of Coastal
1429 bermudagrass hay (*Cynodon dactylon* [L.] Pers) ground at 2 mm as a carbon source.
1430 The flasks were placed in a platform shaker, incubated at 28°C in constant agitation for
1431 8 d. The liquid culture medium, proportion of carbon source, and incubation length
1432 were previously tested and chosen based on a pre-trial enzymatic activity. After each
1433 incubation, the material was frozen at -80° C, freeze-dried, and stored at 4°C to prevent
1434 possible enzymatic denaturation or degradation.

1435 The activities of lignocellulolytic enzymes were determined for laccase,
1436 manganese peroxidase, lignin peroxidase, endoglucanase, exoglucanase, xylanase, and
1437 mannanase at pH 3, 4, 5, and 6. The liquid extracts of the enzymatic complex to
1438 determine the enzymatic activities were carried out using the respective buffers adding 1
1439 g of the enzymatic complex to 19 mL of buffer in a 50 mL tube. The extracts were
1440 vortexed for 1 min, filtered in 2 layers of cheesecloth, centrifuged at 2,500 g for 10 min
1441 at 4°C, and the supernatants were used to determine the enzymatic activities.

1442 Laccase activity was determined in a spectrophotometer (Spectra Max 340 PC,
1443 Molecular Devices Corporation, Sunnyvale, CA, USA) at 420 nm, through the 2,2'-
1444 azino-bis(3- etilbenzotiazolina-6-sulfonato) (ABTS) oxidation, where 35 µL of ABTS
1445 solution (20mM), 35 µL of enzymes extract, and 280 µL of McIlvaine Buffer (corrected

1446 to each pH evaluated) were loaded in a 96-well UV microplate, incubated at 25°C for 5
1447 min. Laccase activity was expressed in units (U), and one U was defined as μmol of
1448 ABTS oxidized per min (Li and Xu, 1999).

1449 Manganese peroxidase activity was determined in a UV-Vis spectrophotometer
1450 (Jasco V-530, Jasco, Easton, MD, USA) at 270 nm according to Wariishi (1992), where
1451 0.6 mL of sodium malonate buffer (50mM, corrected to each pH evaluated), 1.2 mL of
1452 enzymes solution, 0.6 mL of MnSO_4 (4.5 mM), and 0.3 mL of H_2O_2 (9 mM) reacted for
1453 5 mins at room temperature. Manganese peroxidase activity was expressed in U, and
1454 one U was defined as 1 μmol MnSO_4 oxidized per min.

1455 Lignin peroxidase activity was determined in a UV-Vis spectrophotometer (Jasco
1456 V-530, Jasco, Easton, MD, USA) at 310 nm according to Tien and Kirk (1984) through
1457 veratryl alcohol oxidation to veratrilaldehyde (3,4 dimethoxybenzaldehyde).

1458 Specifically, 0.75 mL of sodium tartrate buffer (10 mM, corrected to each pH
1459 evaluated), 0.5 mL of enzyme solution, 0.25 mL of veratryl alcohol (3mM), and 0.10
1460 mL of H_2O_2 (5 mM), with a reaction time of 5 min at room temperature was used.

1461 Lignin peroxidase activity was expressed in U, and one U was defined as 1 μmol
1462 veratryl alcohol oxidized per min.

1463 Endoglucanase and exoglucanase activities were determined in a spectrophotometer
1464 (Spectra Max 340 PC, Molecular Devices Corporation, Sunnyvale, CA, USA) at 540
1465 nm, adapted from Wood and Bhat (1988). In the endoglucanase assay, 1 mL of
1466 carboxymethylcellulose (1%, w/v), as substrate, and 0.9 mL of citrate phosphate buffer
1467 (0.1 M, corrected to each pH evaluated) were added into a 19 mL borosilicate glass tube
1468 and incubated for 10 min at 39°C. Posteriorly, 0.1 mL of enzyme extract was added and
1469 incubated at 39° C for 5 min. Specifically, 3 mL of dinitrosalicylic acid was added into
1470 the tube and boiled for 5 min to mark the released reducing sugar, according to Miller

1471 (1959). Glucose was used as the standard. Endoglucanase activity was expressed in U,
1472 and one U was defined as 1 μmol of glucose released per min. The exoglucanase
1473 activity assay was similar, except the substrate was replaced with microcrystalline
1474 cellulose (1%, w/v).

1475 Xylanase activity was determined in a spectrophotometer (Spectra Max 340 PC,
1476 Molecular Devices Corporation, Sunnyvale, CA, USA) at 540 nm, adapted from Bailey
1477 et al. (1992). In the xylanase assay, 1 mL of xylan (1%, w/v) as substrate and 0.9 mL of
1478 citrate phosphate buffer (0.1 M, corrected to each pH evaluated) were added to a 19 mL
1479 borosilicate glass tube and incubated for 10 min at 39°C. Posteriorly, 0.1 mL of enzyme
1480 extract was added and incubated at 39° C for 5 min. Released reducing sugar was
1481 market as described previously, according to Miller (1959). Xylose was used as the
1482 standard. Xylanase activity was expressed in U, and one U was defined as 1 μmol of
1483 xylose released per min.

1484 Mannanase activity was determined in a spectrophotometer (Spectra Max 340 PC,
1485 Molecular Devices Corporation, Sunnyvale, CA, USA) at 540 nm, adapted from Rättö
1486 and Poutanen (1988), where 1 mL of galactoglucomannan (0.5%, w/v), as the substrate,
1487 as substrate, and 0.9 mL of citrate phosphate buffer (0.1 M, corrected to each pH
1488 evaluated) were added into a 19 mL borosilicate glass tube and incubated for 10 min at
1489 39°C. Posteriorly, 0.1 mL of enzyme extract was added and incubated at 39°C for 5
1490 min. Released reducing sugar was market as described previously, according to Miller
1491 (1959). Mannose was used as the standard. Mannanase activity was expressed in U, and
1492 one U was defined as 1 μmol of mannose released per min.

1493 Protein concentration was measured using the Bradford Protein Kit Assay (Sigma-
1494 Aldrich Co., St. Louis, MO) according to Bradford (1976). BSA was used as standard,
1495 and one mg of protein was determined as one mg of enzyme.

1496

1497 ***2.2. Silage preparation and experimental design***

1498 The WPC was harvested, and the ensiling process was carried out at the Plant
1499 Science Research and Education Unit (Citra, FL). The *in situ* incubation was carried out
1500 at the University of Florida Dairy Research Unit (Gainesville, FL). The procedures for
1501 animal care and handling required were approved by the Institutional Animal Use and
1502 Care Committee of the University of Florida.

1503 The WPC hybrid Syngenta NK1694-3111 (Syngenta International AG, Basel,
1504 Switzerland) was manually harvested at 378 g/mg of DM from 4 different locations in
1505 the same field. The average chemical composition is shown in Table 1.

1506 The experiment evaluated WPCS during the onset of fermentation stage and after
1507 30 d of ensiling. The onset of fermentation was carried out in a completely randomized
1508 block design using a 5 × 4 factorial split-plot arrangement of the treatments (5
1509 enzymatic levels and 4 opening times), totaling 20 treatments with 4 replicates per
1510 treatment. Each replicate originated from a different location in the field, named as plot.
1511 The enzymatic levels were 0, 9, 18, 27, and 36 mg of lignocellulosic enzymes/kg of
1512 WPC (FM basis), corresponding to 0, 0.587, 1.156, 1.734, and 2.312 g of enzymatic
1513 complex/kg of FM, and the opening times were 1, 2, 3, and 7 d after ensiling. Silage
1514 stored for 30 d of ensiling was performed in a completely randomized block design,
1515 with the same five enzymatic levels and four replicates. The enzymatic levels (mg of
1516 enzyme/kg FM) were based on a previous study carried out with lignocellulolytic
1517 enzyme from *Pleurotus ostreatus* produced in our research group (Machado et al.,
1518 2020).

1519 The WPC was chopped at 2 cm of the theoretical length of cut with a single line
1520 combine harvester Cibus S (Wintersteiger inc., Salt Lake City, USA) and immediately

1521 homogenized with the respective levels of lignocellulosic enzymes. Treated WPC (600
1522 g/bag) was immediately placed into nylon-polyethylene vacuum bags (89 μ m thickness,
1523 25.4 \times 35.6 cm; Doug Care Equipment Inc., Springville, USA) and heat-sealed using a
1524 vacuum machine (Bestvac; distributed by Doug Care Equipment Inc.). The silage bags
1525 were stored in a dark environment at room temperature.

1526

1527 ***2.3. Sample collection and analyses***

1528 The bags were weighted to estimate the dry matter (DM) losses and opened on the
1529 respective day of fermentation. Two sub-samples from each bag were dried in a
1530 forced-air oven at 55°C for 72 h. The two replicates were combined and ground in a
1531 Wiley mill to pass a 4 mm screen to determine the *in situ* DM and NDF disappearance
1532 and to pass a 1 mm screen to determine chemical composition and uNDF.

1533 An aqueous extract was prepared by mixing 20 g of the fresh sample plus 200 mL
1534 of double-distilled water in a Stomacher (Lab-Blender 400, Tekmar Company,
1535 Cincinnati, USA) at high speed for 30 seconds and filtered through two layers of
1536 cheesecloth. The pH was determined using a digital pH meter (Accumet XL25, Thermo
1537 Fisher Scientific Inc., Waltham, USA). Forty-mL aliquot of each extract was acidified
1538 with 0.4 mL of sulfuric acid in water (50/50, v/v), centrifuged at 7000 \times g for 15 min at
1539 4°C, and stored at -20°C to determine organic acids and NH₃-N.

1540 Organic acid concentrations were determined as described by Muck and
1541 Dickerson (1988) using HPLC (Merck Hitachi Elite La-Chrome; Hitachi L2400, Tokyo,
1542 Japan). A Bio-Rad Aminex HPX-87H ion exclusion column (300 \times 7.8-mm i.d.; Bio-
1543 Rad Laboratories, Hercules, CA) was used in an isocratic elution system containing
1544 0.015 M sulfuric acid in the mobile phase of HPLC with a UV detector (wavelength 210
1545 nm; L-2400, Hitachi), using a flow rate of 0.7 mL/min at 45°C.

1546 The NH₃-N concentration was determined according to Broderich and Kang
1547 (1980) and adapted to a plate reader by using 2 μL of the sample, 100 μL of phenol, and
1548 80 μL of hypochlorite in each well of the microplate. The plate was incubated at 95°C
1549 for 10 min and maintained at room temperature for 10 min for cooling. Absorbance
1550 readings were done utilizing a UV-Vis spectrophotometer at 620 nm (Spectra Max 340
1551 197 PC, Molecular Devices Corporation, Sunnyvale, USA).

1552 To determine the NDF *in situ* disappearance, approximately 5 g of ground
1553 material at 6 mm was weighed into filter bags (R1020, 10 x 20 cm, 50 ± 10-micron
1554 porosity; Ankom Technology, Macedon, USA) in duplicate for each sample. Bags were
1555 incubated for 30 h in two rumen-cannulated lactating Holstein cows (1 bag per sample
1556 per cow). Those animals used to the incubation were fed a diet consisting of 38% corn
1557 silage, 19% ground corn, 13% soybean meal, 11% cottonseed, 9% citrus pulp, 8.5%
1558 mineral premix, and 1.5% palmitic acid supplement.

1559 To determine the *in situ* undigested NDF (uNDF), approximately 0.5 g of
1560 ground material (1 mm) was weighed into fiber filter bags (F57, 25-micron porosity;
1561 Ankom,Technology, Macedon, USA) in duplicate for each sample. Bags were
1562 incubated for 240 h in two rumen-cannulated lactating Holstein cows (1 bag per sample
1563 per cow).

1564 All the bags were dried in a forced-air oven at 60°C for 48 h. The bags from 30
1565 h of incubation were weighed to determine the DM disappearance, and the replication
1566 was combined and ground to pass a 1 mm sieve, and samples were placed into fiber
1567 filter bags (F57, 25 micron porosity; Ankom,Technology, Macedon, USA) for aNDF
1568 analysis. The aNDF was determined according to Mertens et al. (2002), using
1569 thermostable α-amylase and sodium sulfite in Ankom 200 Fiber Analyzer (Ankom
1570 Technologies, Macedon, USA).

1571 The DM content was determined at 105°C using an oven according to method
1572 No. 924.01 (AOAC, 1990). Ash was determined by combustion at 600°C for 6 h in a
1573 furnace, according to method No. 924.05 (AOAC, 1990). Neutral detergent fiber
1574 (aNDF) was determined, according to Mertens (2002), using thermostable α -amylase
1575 and sodium sulfite. Acid detergent fiber (ADF) was determined according to method
1576 No. 973.18 (AOAC, 1990). The concentration of lignin was determined using the acid
1577 detergent lignin methodology, according to Van Soest and Wine (1968), by submitting
1578 the material to sulfuric acid (72/28, v/v in DI water) sequentially following ADF
1579 analysis. The cellulose concentration was obtained by the difference between the ADF
1580 and lignin, whereas hemicellulose concentration was obtained by the difference between
1581 aNDF and ADF. Water-soluble carbohydrate (WSC) was determined using the anthrone
1582 reaction test (Weiss et al., 1990) and the starch concentration, according to Hall et al.
1583 (2015), by colorimetric method.

1584

1585 *2.4. Statistical analyses*

1586 All analyses were carried out using the GLIMMIX procedure of SAS (version
1587 9.4, SAS Institute Inc., Cary, NC). Differences were declared significant at $P \leq 0.05$ and
1588 tendencies if $P > 0.05$ and ≤ 0.10 .

1589 For the **enzymatic activity**, data were analyzed in a completely randomized
1590 design using the following model:

$$1591 Y_{ij} = \mu + H_i + \varepsilon_{ij},$$

1592 where Y_{ij} = dependent variable; μ = overall mean; H_i = fixed effect of the pH (i
1593 = 3 to 6); and ε_{ij} = random error. Enzymatic activity means were compared using the
1594 Bonferroni t-test option whenever differences were observed.

1595 Data from the onset of fermentation stage were analyzed in a completely
 1596 randomized block design using a 5×4 factorial slip-plot arrangement of treatments (5
 1597 enzymatic levels and 4 ensiling time), using the following model:

$$1598 \quad Y_{ijkl} = \mu + P_i + E_j + P_i \times E_j + F_k + E_j \times F_k + \varepsilon_{ijkl},$$

1599 where Y_{ijkl} = dependent variable; μ = overall mean; P_i = random effect of plot (i
 1600 = 1 to 4); E_j = fixed effect of enzymatic level ($j = 0$ to 36 mg of enzymes/kg of FM); $P_i \times$
 1601 E_j = main-plot error; F_k = fixed effect of days of fermentation ($k = 1$ to 4); $E_j \times F_k$ =
 1602 fixed effect of interaction; and ε_{ijkl} = random error. Linear and quadratic effects were
 1603 tested for the enzymatic levels and fermentation length by using orthogonal contrasts.
 1604 The coefficients for orthogonal contrasts were determined using the IML procedure of
 1605 SAS.

1606 Data at 30 d of fermentation was analyzed in a completely randomized block
 1607 design, containing 5 enzymatic levels, using the following model:

$$1608 \quad Y_{ijk} = \mu + E_i + P_j + \varepsilon_{ijk},$$

1609 where Y_{ijk} = dependent variable; μ = overall mean; E_i = fixed effect of
 1610 enzymatic levels ($i = 0$ to 36 mg of enzymes/kg of FM); P_j = random effect of plot ($j = 1$
 1611 to 4); and ε_{ijk} = random error. Linear and quadratic effects were tested for the enzymatic
 1612 levels by using orthogonal contrasts.

1613

1614 3. RESULTS

1615 The characterization of the enzymatic complex produced by *Pleurotus ostreatus*
 1616 is presented in Table 2. Laccase showed the highest activity ($P < 0.01$) at pH 5, which
 1617 corresponded to 97.8 U/g of enzymatic complex per min. Laccase activity decreased at
 1618 pH 4 and 6 that corresponded to a reduction of approximately 23% when both were

1619 compared with the activity at pH 5. The lowest laccase activity was observed at pH 3,
1620 whose reduction was 40.2% compared with the activity observed at pH 5.

1621 Manganese peroxidase and lignin peroxidase had the highest activity at pH 4 (P
1622 < 0.01). Manganese peroxidase had no activity at pH 5 and 6 and low activity at pH 3.
1623 Lignin peroxidase activity decreased by approximately 68% at pH 3 and 98.7% at pH 5
1624 compared to the highest activity observed at pH 4. Mannanase had the highest activity
1625 at pH 5 ($P < 0.04$); however, it did not differ from the activities observed at pH 6 and 4.
1626 Activities of xylanase, endo- and exo-glucanase did not change from pH 3 to 6.

1627 The P -values for the effects of enzymatic complex, days on the onset of
1628 fermentation, and their interactions at the onset of fermentation are in Table 3. There
1629 was no interaction between enzymatic complex and days on the onset of fermentation
1630 ($P > 0.11$), also enzymatic complex did not affect fermentation profile and chemical
1631 composition at the onset of fermentation ($P > 0.21$). Differences observed for the main
1632 effect of days of fermentation are presented in Table 4. The pH decreased quadratically
1633 ($P = 0.01$), and the concentration of lactic acid increased quadratically ($P = 0.01$) as
1634 fermentation progressed. From 1 to 7 d, acetic acid ($P = 0.02$) and the total organic
1635 acids ($P < 0.01$) increased linearly. The concentrations of lignin, uNDF, and WSC
1636 decreased quadratically with days of fermentation ($P \leq 0.02$).

1637 After 30 d of storage, the fermentation profile was not affected by
1638 lignocellulolytic enzymes (Table 5). Ash concentration increased quadratically ($P =$
1639 0.05), whereas organic matter decreased quadratically ($P = 0.05$) with the addition of
1640 enzymes. The lignin concentration decreased linearly ($P = 0.04$) with the addition of the
1641 enzymatic complex (Table 6). Other nutrients were not affected by enzymatic levels (P
1642 > 0.11).

1643

1644 4. DISCUSSION

1645 White-rot fungi, such as *Pleurotus ostreatus*, have been reported in the literature
1646 to produce enzymes that break down lignin and degrade cellulose and hemicellulose
1647 (Bánfi et al., 2015). In this study, we observed that *Pleurotus ostreatus* produced three
1648 enzymes that break down lignin (laccase, lignin peroxidase, and manganese
1649 peroxidase), two types of cellulases (endoglucanase and exoglucanase), and two
1650 hemicellulases (xylanase and mannanase).

1651 The enzymatic activity is recognized to be affected by temperature, substrate
1652 concentration, and pH (Beauchemin et al., 2003). Therefore, the activity of enzymes
1653 from *Pleurotus ostreatus* cultivation was tested at different pH. As expected, the
1654 optimum pH varied among enzymes. The optimum pH observed for laccase was at 5 in
1655 our study, previous studies (Manole et al., 2008; El-Batal et al., 2015). Laccase activity
1656 was likely impaired during the onset of fermentation due to the pH drop, once silage pH
1657 decreased to almost 4 on the first day of fermentation, and also to anaerobiosis
1658 establishment in silages. As demonstrated, at pH 4 the activity of this enzyme was
1659 suppressed by approximately 23% compared to the optimum pH (pH 5). Anaerobiosis
1660 also may affect laccase activity negatively, as O₂ is an essential cofactor for its catalytic
1661 cycle (Shekher et al., 2011). In the silo, the residual oxygen is consumed by plant cell
1662 respiration and aerobic or facultative microorganisms shortly after silo sealing (Elferink
1663 et al., 2000), allowing a short time of optimal conditions for proper enzyme function.

1664 The optimum pH to lignin peroxidase (LiP) was at 4, which is related to the pH
1665 required to LiP Compound I oxidize a non-phenolic aromatic and convert to LiP
1666 Compound II (Wong, 2009; Datta et al., 2017), being essential to lignin peroxidase
1667 cycle. The optimum pH favored the activity of this enzyme along the fermentation when
1668 the pH range was around 4.

1669 The higher activities of manganese peroxidase and lignin peroxidase at low pH
1670 may have allowed these enzymes to act during the whole fermentation. Also,
1671 manganese peroxidase and LiP do not require aerobic conditions for the cycle activation
1672 (Wong, 2009); however, manganese peroxidase does require Mn^{+2} , as a cofactor, to
1673 donate an electron converting to Mn^{+3} that oxidize and consequently cleave the lignin
1674 (Hofrichter et al., 2010). This requirement is convenient in materials as whole-plant
1675 corn because it has approximately 914 mg/kg DM of Mn (NRC, 2001), contributing to
1676 manganese peroxidase activity. Therefore, the reduction in lignin concentration in the
1677 corn silage at 30 d after ensiling with the addition of enzymatic complex may have
1678 resulted from lignin cleavage mainly by manganese and lignin peroxidases along silage
1679 fermentation. We expected to observe the effect of the enzymatic levels in the lignin
1680 concentration along the onset of fermentation, however, this was not observed. We
1681 attributed this lack of effect to not enough accumulation of lignin cleavage until 7 d of
1682 fermentation.

1683 Lignin works as a barrier that hinders the rumen microorganisms from degrading
1684 the fiber due to its high association with hemicellulose in the cell wall by cross-linking
1685 of lignin to arabinoxylans (Hatfield et al., 2017). Thus, degradation of lignin would be a
1686 way to increase the access of the microorganisms to hemicellulose and cellulose and
1687 possibly increase fiber degradability (van Kuijk et al., 2015). However, despite reducing
1688 the lignin concentration, *in situ* DM and NDF disappearance did not increase by adding
1689 the enzymatic complex.

1690 The reduction of lignin concentration with crescent levels of enzymatic complex
1691 was similar to that observed by Machado et al. (2020). However, in this previous study,
1692 the reduction in lignin led to an increase of *in vitro* DM and NDF digestibility that was
1693 not observed in the present study. Differences among studies may have occurred due to

1694 different lignin concentrations and the cell wall arrangement in the WPC, since in the
1695 current study, lignin concentration in WPC was quite lower than that observed by
1696 Machado et al. (2020) (16.3 vs. 66.4 g/kg of DM). Therefore, these factors related to
1697 lignin might favor the magnitude of its reduction by enzyme treatment (8.59% in the
1698 present study vs. 44.4% in Machado et al., 2020), and consequently, the enzymes
1699 benefited in NDF and DM degradability.

1700 Another factor that may be played a role in the lignin concentration and the
1701 degradability was the range in lignocellulolytic activities. Machado et al. (2020)
1702 observed higher activity of laccase, cellulases, and hemicellulases than in the present
1703 study. Differences in enzymatic activities are attributed to fungal strain, and culture
1704 medium conditions since Machado et al. (2020) used KIRK medium (Kirk et al., 1986),
1705 and the *Pleurotus ostreatus* was not described. In the current study, the culture medium
1706 was prepared with Bermuda grass (*Cynodon dactylon*) hay, sugarcane and water, chosen
1707 from a pre-trial to replace KIRK medium to decrease the production cost. Membrillo et
1708 al. (2008) also observed that two different strains of *Pleurotus ostreatus* produced
1709 different proportions of enzymes when the fungi were exposed to different medium
1710 conditions, which supported our results.

1711 Although the enzymatic complex presented activities of cellulases and
1712 hemicellulases, its addition onto WPC at ensiling did not affect the concentration of
1713 aNDF and ADF at the onset of fermentation or 30 d after ensiling. Colombatto et al.
1714 (2004) and Lynch et al. (2015) observed a reduction in the concentration of aNDF and
1715 ADF, that differed from the present study, using fibrolytic enzymes (a combination of
1716 cellulolytic and hemicellulolytic) in WPC at ensiling. This divergence may be attributed
1717 to the levels of enzymatic activities and/or the concentration and arrangement of NDF
1718 and ADF in the unfermented WSC. In those studies, the enzymatic activity per kg of

1719 WPC and the concentration of NDF and ADF in WPC was much higher than in our
1720 study.

1721 The concentrations of lactic acid, acetic acid, propionic acid, and NH₃-N, the
1722 pH, and the absence of butyric acid in the corn silage after 30 days of ensiling were
1723 similar to the values described in the literature to a good fermentation, evidencing that
1724 the WPCS showed a satisfactory fermentation (Kung et al., 2017).

1725 Our suggestion for future studies is to test the enzymatic complex from the
1726 *Pleurotus ostreatus* cultivation in WPC with high lignin and NDF concentration,
1727 usually observed in silage production in tropical areas (Correa et al., 2002), where the
1728 enzymes may express more effect.

1729

1730 5. CONCLUSIONS

1731 The addition of lignocellulolytic enzymatic complex from *Pleurotus ostreatus*
1732 cultivation decreased lignin concentration in the WPCS after 30 d of ensiling; however,
1733 this did not affect *in situ* disappearance of dry matter and neutral detergent fiber. This
1734 reduction is associated with the activities of lignin peroxidase and manganese
1735 peroxidase, which are capable of degrading lignin anaerobioses and low pH. No
1736 difference in the fermentation profile and chemical composition was observed at onset
1737 fermentation.

1738

1739 **CRedit authorship contribution statement:** Bruna C. Agostinho: Formal analysis,
1740 Investigation, Data curation, Writing - original draft, Writing - review & editing. Joao
1741 L. P. Daniel: Data curation, Supervision, Writing - review & editing, Supervision. Lucia
1742 M. Zeoula: Conceptualization, Investigation, Supervision. Luiz F. Ferraretto:
1743 Conceptualization, Investigation. Hugo F. Monteiro: Statistical analysis, Writing -

1744 review. Matheus R. Pupo: Formal analysis. Lucas G. Ghizzi: Formal analysis, Writing -
1745 review. Mariele C. N. Agarussi: Formal analysis. Celso Heinzen Junior: Formal
1746 analysis, Data curation. Richard R. Lobo: Formal analysis. Antonio P. Faciola:
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1751

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Tables

1903 Table 1. Chemical composition of unfermented whole-plant corn (WPC)

Item	Mean	SD
pH	5.84	0.16
DM, g/kg as-fed	378	27.7
aNDF, g/kg of DM	357	24.4
ADF, g/kg of DM	185	12.1
Cellulose, g/kg of DM	166	13.9
Hemicellulose, g/kg of DM	172	13.2
Lignin, g/kg of DM	18.5	2.35
Starch, g/kg of DM	325	20.7
WSC, g/kg of DM	75.4	2.06
Ash, g/kg of DM	24	0.89

1904 SD = standard deviation; DM = dry matter; aNDF = neutral detergent fiber determined with

1905 heat-stable alpha-amylase and inclusive of residual ash; ADF = acid detergent fiber; WSC =

1906 water soluble carbohydrates.

1907 Table 2. Characterization of enzymatic complex produced by *Pleurotus ostreatus*

Enzymatic activity, U/g/min ¹	pH				SEM	P-value
	3	4	5	6		
Laccase	58.5 ^c	72.9 ^b	97.8 ^a	78.6 ^b	2.01	<0.01
Mn Peroxidase ²	3.79 ^b	22.2 ^a	0.00 ^b	0.00 ^b	1.15	<0.01
Lignin Peroxidase	26.1 ^b	81.8 ^a	1.03 ^c	0.0 ^c	48.1	<0.01
Endoglucanase	16.3	14.3	19.3	14.1	0.17	0.61
Exoglucanase	23.4	27.9	35.4	28.8	1.82	0.64
Xylanase	26.6	28.2	28.6	27.1	0.07	0.42
Mannanase	17.0 ^b	18.5 ^{ab}	19.0 ^a	17.9 ^{ab}	0.03	0.04

1908 ^{a,b,c} Means with different superscript letters differed ($P \leq 0.05$).1909 ¹The enzymatic activities were identified based on the final product obtained in the reaction.

1910 Enzymatic activity was expressed in U/g of SSPO/min. Abbreviations: Mn peroxidase =

1911 Manganese peroxidase.

1912 Table 3. *P*-values for enzymatic complex addition, days of fermentation, and their
 1913 interaction at the onset of fermentation (1 to 7 d) in whole-plant corn silage treated with
 1914 lignocellulolytic enzymes from *Pleurotus ostreatus*

Item	<i>P</i> -value		
	Enzyme	Time	Enzyme*time
pH	0.62	<0.01	0.17
Total acids	0.26	<0.01	0.11
Lactic acid	0.43	<0.01	0.11
Acetic acid	0.21	<0.01	0.14
Propionic acid	0.57	0.02	0.46
N-NH ₃	0.29	<0.01	0.15
DM	0.36	0.11	0.70
Ash	0.96	0.41	0.99
OM	0.96	0.41	0.99
aNDF	0.57	0.48	0.82
ADF	0.52	0.19	0.71
Cellulose	0.43	0.22	0.66
Hemicellulose	0.58	0.13	0.57
Lignin	0.62	<0.01	0.30
WSC	0.37	<0.01	0.97
DMD	0.68	0.76	0.31
NDFD	0.57	0.34	0.12
uNDF	0.65	0.01	0.38
DM loss	0.91	0.64	0.31

1915 Abbreviations: DM = dry matter; aNDF = neutral detergent fiber determined with heat-stable
 1916 alpha-amylase and inclusive of residual ash; ADF = acid detergent fiber; WSC = water soluble
 1917 carbohydrates; uNDF = in situ undigested neutral detergent fiber; DMD = in situ dry matter
 1918 digestibility; NDFD = in situ neutral detergent fiber digestibility; DM loss = dry matter loss.

1919 Table 4. Fermentation profile and chemical composition of whole-plant corn silage at
 1920 the onset of fermentation, when averaged over lignocellulolytic enzymes from
 1921 *Pleurotus ostreatus*

Item	Day of ensiling				SEM	P-value	
	1	2	3	7		Linear	Quadratic
Fermentation profile							
pH	4.25	4.28	3.87	3.85	0.0413	<0.01	<0.01
Lactic acid ¹	18.2	21.8	33.7	44.3	2.53	<0.01	0.02
Acetic acid ¹	11.9	10.6	15.0	14.3	1.07	0.02	0.16
Propionic acid ¹	8.96	6.51	7.07	7.61	0.580	0.54	0.01
Total acids ¹	39.1	38.9	56.5	66.2	4.14	<0.01	0.09
NH ₃ -N ¹	29.0	33.4	35.8	40.9	1.77	<0.01	<0.01
Chemical composition, g/kg of dry matter							
Lignin	18.5	16.3	16.6	16.9	0.424	0.10	<0.01
WSC	42.2	36.3	28.6	17.1	2.09	<0.01	<0.01
uNDF	134	122	126	125	2.77	0.16	0.02

1922 Abbreviations: WSC = water soluble carbohydrates; uNDF = undigested neutral detergent fiber.

1923 ¹ = g/kg of dry matter.

1924 Table 5. Fermentation profile and dry matter loss of whole-plant corn silage treated with
 1925 lignocellulolytic enzymes from *Pleurotus ostreatus*, after 30 d of fermentation

Item	Treatments (mg enzymes/kg FM) ¹					SEM	P-value	
	0	9	18	27	36		Linear	Quadratic
pH	3.64	3.60	3.71	3.59	3.64	0.024	0.91	0.35
Total acids ²	71.84	69.1	64.4	83.6	64.6	7.05	0.99	0.67
Lactic acid ²	58.2	53.3	50.8	61.7	53.5	5.56	0.92	0.65
Acetic acid ²	8.38	8.44	8.19	12.4	7.76	1.77	0.53	0.34
Propionic acid ²	5.24	7.33	5.45	9.91	3.35	1.75	0.81	0.12
Butyric acid ²	ND	ND	ND	ND	ND	-	-	-
NH ₃ -N ²	48.9	47.9	50.8	45.4	46.7	3.18	0.16	0.52
DM Loss ³	133	144	131	125	127	19.5	0.33	0.74

1926 ND = not detected.; ¹ = treatments corresponded to the following enzymatic activities: 0
 1927 (0 g of enzymatic complex/kg of FM), 9 (0.587 g of enzymatic complex/kg of FM), 18
 1928 (1.156 g of enzymatic complex/kg of FM), 27 (1.734 g of enzymatic complex/kg of
 1929 FM), 36 (2.312 g of enzymatic complex/kg of FM); ² = g/kg of dry matter; ³ = DM loss =
 1930 dry matter loss.

1931 Table 6. Chemical composition and *in situ* disappearance of whole-plant corn silage
 1932 treated with lignocellulolytic enzymes from *Pleurotus ostreatus*, after 30 d of
 1933 fermentation

Item, g/kg of dry matter	Treatments (mg enzymes/kg FM) ¹					SEM	P-value	
	0	9	18	27	36		Linear	Quadratic
Dry matter, g/kg	354	349	354	358	357	12.8	0.28	0.63
Ash	23.7	26.6	27.0	25.0	25.6	0.874	0.44	0.05
Organic matter	976	973	973	975	974	0.874	0.44	0.05
NDF	364	391	392	371	358	20.6	0.57	0.15
ADF	190	209	211	205	196	11.5	0.78	0.12
Lignin	16.3	17.3	15.0	15.0	14.9	0.701	0.04	0.99
Cellulose	173	191	196	190	181	11.2	0.65	0.11
Hemicellulose	174	182	181	166	162	9.91	0.13	0.22
WSC	8.82	8.42	9.07	9.82	9.25	1.10	0.43	0.94
uNDF	116	126	129	123	121	5.33	0.73	0.12
DMD	661	667	663	679	692	14.2	0.15	0.55
NDFD	236	286	287	288	268	27.3	0.39	0.13

1934 ¹ = treatments corresponded to the following enzymatic activities: 0 (0 g of enzymatic
 1935 complex/kg of FM), 9 (0.587 g of enzymatic complex/kg of FM), 18 (1.156 g of
 1936 enzymatic complex/kg of FM), 27 (1.734 g of enzymatic complex/kg of FM), 36 (2.312
 1937 g of enzymatic complex/kg of FM); Abbreviations: DM = dry matter; aNDF = neutral
 1938 detergent fiber determined with heat-stable alpha-amylase and inclusive of residual ash; ADF =
 1939 acid detergent fiber; WSC = water soluble carbohydrates; uNDF = in situ undigested neutral
 1940 detergent fiber; DMD = in situ dry matter degradability; NDFD = in situ neutral detergent fiber
 1941 degradability.