

1 **Efficacy of a novel sequential enzymatic hydrolysis of lignocellulosic biomass and**
2 **inhibition characteristics of monosugars**

3 *Sibashish Baksi¹, Akash K Ball¹, Ujjaini Sarkar^{*1}, Debopam Banerjee¹, Alexander Wentzel³,*
4 *Heinz A Preisig⁴, Jagdish Chandra Kuniyal², Cansu Birgen⁴, Sudeshna Saha¹, Bernd*
5 *Wittgens³, Sidsel Markussen³*

6

7 **Abstract**

8 Efficient production of sugar monomers from lignocellulose is often hampered by serious
9 bottle-necks in biomass hydrolysis. The present study reveals that ultra-sonication assisted
10 pretreatment following autoclaving, termed as combined pretreatment, can lead to more
11 efficient delignification of lignocellulosic biomass and an open, deformed polysaccharide
12 matrix, found favorable for subsequent enzymatic hydrolysis, is formed. The pattern of
13 inhibition for the enzymatic hydrolysis reaction on combined-pretreated saw dust is
14 identified. Two main inhibition models (competitive and noncompetitive) are proposed and a
15 better fit of experimental values with the theoretical values for the competitive inhibition
16 model validates the proposition that in the present experiment, glucose inhibits the enzymes
17 competitively. Additionally, accuracy of the inhibitory kinetics based models is estimated
18 over a series of enzyme and substrate concentrations. A prominent departure in the range of
19 residual concentrations from the competitive model supports the same proposition, in
20 comparison to the non-competitive model.

21 **Key words:** enzymatic hydrolysis; enzyme inhibition; lignocellulose, cellulase; xylanase; β -
22 glucosidase.

23 **1. Introduction**

24 With increasing global population and industrialization, the energy demand and consumption
25 rates have been increasing progressively for the last few decades. Although fossil fuels are
26 still the most convenient and promising source to meet this elevated energy demands, its
27 sources are slowly depleting. Moreover, its exploitation is directly linked to global warming,
28 environmental pollution, climate change and health hazards(Lavoine, Desloges, Dufresne, &
29 Bras, 2012). Thus, there is an urgent need to find alternative energy resources. Along with
30 nuclear-, solar-, wind-and hydro-power, biofuel represents an important future energy carrier
31 and a sustainable substitute for fossil fuels. In this context, the biofuel production from
32 Lignocellulosic Biomass (LB) is gaining importance due to its abundance and sustainable
33 production process. Lignocellulosic biomass, essentially consisting of cellulose,
34 hemicelluloses and lignin are considered as the primary feedstock for production of biofuels.
35 Lignin forms a hetero-matrix with sugar polymers by cross-linking via hydrogen bonding,
36 covalent bonding such as ester linkages and physical encrustation. In the hetero-matrix,
37 cellulose molecules form the inner core of the structure, which is surrounded by a lignin-
38 hemicellulose based cross-linked matrix. It is essential to remove lignin from lignocellulose
39 feedstock in order to efficiently access the reducible sugar polymers for hydrolysis using
40 enzymes (Agbor, Cicek, Sparling, Berlin, & Levin, 2011; Betts, Dart, Ball, & Pedlar, 1991;
41 Hu & Ragauskas, 2012).

42 Over the years, many approaches are developed and implemented to generate biofuels from
43 LB. Some of the most popular methods adopted for initial pretreatment of biomass include
44 conventional alkaline peroxide-based pretreatment, acid pretreatment, steam explosion,
45 ammonia fiber expansion (AFEX), hot liquid water pretreatment, etc. While several processes
46 remove lignin efficiently from LB, it is found that a large portion of hemicelluloses and
47 celluloses degrade due to the high temperature often applied (Y. Sun & Cheng, 2002).

48 Consequently, a number of soluble toxic chemicals like furfural and HMF are generated
49 along with the degradation of hemicelluloses (W.-H. Chen, Hsu, Lu, Lee, & Lin, 2011).
50 AFEX pretreatment generates much smaller amount of inhibitors and increases the surface
51 area considerably, but it is not effective in removing lignin from biomass with high lignin
52 content. In spite of being very efficient, acid pretreatment is environmentally incompatible
53 since significant amounts of toxic chemicals and inhibitors are formed. Additionally,
54 corrosion is also a major concern of acid pretreatment (Wu, Yu, Chan, Kim, & Mai, 2000;
55 Yoon, Wu, & Lee, 1995; Zhu, Pan, & Zalesny, 2010). Considering the advantages,
56 disadvantages and cost of all pretreatment procedures, conventional alkaline peroxide
57 pretreatment is found to be a promising cost-effective option for pretreatment with much less
58 production of inhibitors.

59 Enzymatic hydrolysis of delignified biomass liberates fermentable sugars from pretreated LB,
60 with commercial enzymes, for decades. However, improved hydrolysis protocols are still
61 needed in order to overcome various problems arising due to the presence of inhibitors during
62 hydrolysis. Typically, two types of inhibition are observed during enzymatic hydrolysis of
63 LB - *product inhibition* and *substrate inhibition*. A couple of research groups tried to
64 overcome inhibition by using enzymes extracted from different genetically modified
65 organisms, significant progress is yet to be made (Fenila & Shastri, 2016; Sternberg,
66 Vuayakumar, & Reese, 1977).

67 *Morus serrata* (MS, mulberry), popularly known as Himalayan Mulberry, is a promising LB
68 source as this species can be used to generate monomeric sugars. Its optimum temperature for
69 growth ranges from 18°C to 30°C (Arya, Kalia, & Arya, 2000; Sargent, 1896). MS is widely
70 used in wood technology-based industries. A huge amount of wood powder is generated from
71 MS, which is generally discarded as waste. However, this powder can be used as a
72 prospective source of LB for generation of monomeric sugars. Other than lignin and

73 carbohydrate, various components present in other different types of biomass accounting for
74 the undetermined portion are enlisted below (Table 1).

75 Table 1 Components present in various other types of biomass accounting for the
76 undetermined portion (*other than lignin and carbohydrate*).

Biomass	Components	References
Switch grass	proteins, waxes, resins, gums, chlorophyll	(Xu, Cheng, Sharma- Shivappa, & Burns, 2010)
Bermuda Grass	uronic acids, acetyl groups, minerals, waxes, resins, gums	(Wang, Keshwani, Redding, & Cheng, 2010)
Hybrid poplar	ash, acetyl, anhydroglucuronic acid	(Luo, Brink, & Blanch, 2002)

77

78 In the present study, LB from MS is first de-lignified using a novel two-step pretreatment
79 process and subsequently de-lignified biomass is hydrolyzed using a specific enzyme
80 protocol to maximize both C₆ and C₅ sugars. Two inhibition models, competitive and non-
81 competitive, are validated for various ranges of substrate and enzyme concentrations in order
82 to generate primarily glucose and xylose.

83 **2. Materials and Methods**

84 *2.1. Sample preparation*

85 After harvesting branches from MS trees by using a hand saw, Himalayan Mulberry chips
86 and sawdust are prepared by using a chain saw (Make: STIHL, Germany; Model: Cast Iron
87 Chain Saw; Material: MS-180) and a circular saw (Make: BOSCH; Model: GKS190). Finely

88 ground mulberry powder is then passed through a 85 mesh screen size sieve prior to be
89 further used in order to remove impurities and larger particles.

90 2.2. *Pretreatment*

91 A range of different combination of pretreatment protocol is applied to maximize the removal
92 of lignin from mulberry samples while minimizing biomass denaturation and formation of
93 inhibitors for subsequent hydrolysis.

94 2.2.1. *Alkaline peroxide pretreatment*

95 Conventional alkaline-peroxide pretreatment (AP) is performed using 25 g moisture free
96 mulberry powder with a solid to alkaline-peroxide solution ratio of 1:40 (w/v) for 5h at 50°C.
97 The alkaline peroxide solution consists of 2% H₂O₂ [v/v] where the pH is maintained at 11.5
98 with NaOH. Alkaline peroxide pretreatment is performed with agitation at 150rpm (Banerjee,
99 Car, Scott-Craig, Hodge, & Walton, 2011).

100 2.2.2. *Combined pretreatment*

101 A novel combined pretreatment (CP) procedure is introduced to maximize removal of lignin
102 from the wood sample. 25 g of moisture free wood sample is submerged in alkaline solution
103 (pH 11.5), prepared by dissolving NaOH pellets into distilled water, followed by autoclaving
104 at 121°C for 60 min. Successively, the solid fraction is isolated, washed and dried at 40°C.
105 Next, the dried sample is submerged in 2%alkaline-peroxide solution [v/v] (pH 11.5) and
106 sonicated 20 times using a probe sonicator [Make: PCI analytics, India; power used: 300W;
107 probe diameter: 9mm]. After sonication, the solid fraction is separated by filtration using
108 Whatman no. 1 filter papers (Make: Sartorius, Germany), neutralized with de-ionized water
109 and dried overnight in a hot air oven at 40°C. The dried, sonicated mulberry powder is then
110 again treated with alkaline peroxide solution [pH 11.5, 2% H₂O₂(v/v)] for 5h at 50°C.
111 Subsequently, the insoluble solid material is separated and washed with de-ionized water

112 until pH of the solution is at neutral. Finally, the sample is again dried overnight in a hot air
113 oven at 40°C.

114 2.3. *Isolation of hemicellulose from pretreatment solutions*

115 Hemicelluloses have a tendency to dissolve in acid or concentrated alkaline solution and can
116 be isolated by precipitation. After pretreatments followed by separation (*see* section 2.2), the
117 alkaline peroxide solution is acidified to a pH of 4.4 using 10% HCl [v/v] solution. Next, three
118 volumes of 95% ethanol is added to the acidified solution and incubated overnight, allowing
119 the dissolved hemicelluloses to precipitate. Hemicelluloses thus precipitated are recovered
120 afterwards by filtration and then washed with 72% ethanol followed by drying (Subhedar &
121 Gogate, 2014).

122 2.4. *Estimation of lignin content in the biomass*

123 The amount of lignin present in untreated as well as in the pretreated mulberry wood powder
124 is determined quantitatively following TAPPI T222 method (Tappi, 2002). Specifically, 1g
125 (± 0.1 g) of each sample (test specimen) is dissolved in 15ml of 72% sulfuric acid and
126 incubated at 20°C for 2h. Then, the final volume is adjusted to 575ml with distilled water and
127 the mixture is boiled for 4h and left overnight for precipitation of lignin. The precipitated
128 lignin is isolated using a glass crucible and quantified along with acid-soluble lignin
129 (Standard, 2002).

$$130 \quad \text{Lignin}(\%) = \frac{\text{Weight of lignin (g)}}{\text{Weight of the test specimen(g)}} \times 100 \quad (1)$$

131 Where weight of the lignin (g) = [weight of the crucible with lignin (g) - weight of the
132 crucible (g)].

133 2.5. *Estimation of total carbohydrates in crude and pretreated mulberry wood powder*
134 Total reducing sugar in crude and pretreated sample are determined with 3,5-dinitrosalicylic
135 acid (DNS method) using glucose as the standard (Miller, 1959; Sluiter et al., 2008; Y. Sun &
136 Cheng, 2005; Van Wychen & Laurens, 2017). 0.3 g each of the samples is hydrolyzed with
137 3ml of 72% H₂SO₄ and the mixture is kept at 30°C for 1h. Afterwards, the solution is diluted
138 to 4% H₂SO₄ by adding 84 ml of distilled water followed by autoclaving for 1h. Next the
139 mixture is centrifuged and the clear supernatant is collected. 3 ml of clear supernatant is
140 mixed with 3 ml of DNS reagent and boiled for 5 min at 100°C. Then the mixture is cooled
141 down and the color intensities are recorded in a UV/Vis spectrophotometer [Make:
142 PerkinElmer, USA; Model: Lambda 365] at 575nm. Quantitative estimation of total reducing
143 sugar is based on the standard curve generated using glucose equivalent.

144

145 2.6. *Crystallinity of biomass using X-Ray Diffraction (XRD) method*

146 Changes in the crystallinity of biomass depend largely on the specific method of
147 delignification as crystalline and amorphous cellulose along with lignin and hemicelluloses
148 largely constitute the structure of the biomass matrix. The degree of crystallinity of crude and
149 pretreated biomass is determined using an X-ray diffractometer [Make: Rigaku, Japan;
150 Model: Giegerflex D/Max B], employing a Cu K α radiation source functioning at 40kV and
151 30 mA. Scan is performed in the range of 2 θ [Bragg angle=5°- 40°] at a scanning rate of
152 0.05° per second. The crystallinity index (CrI) of various samples is determined using the
153 scattered data in equation (2) (Baksi et al., 2018; Zhang et al., 2014):

154
$$CrI (\%) = \frac{I_{002} - I_{am}}{I_{002}} \times 100 \quad (2)$$

155 I_{002} and I_{am} represent the scattered intensity of the crystalline portion of biomass at about
156 $2\theta = 22.5^\circ$ and the amorphous portion of biomass at about $2\theta = 16.8^\circ$, respectively.

157 2.7. *Enzymatic hydrolysis*

158 Enzymatic hydrolysis is carried out using a cellulase blend (SAE0020, 1000U/g, 1.2g/ml)
159 along with cellulase [C1184, 1.3U/mg, derived from *A. niger*], hemicellulase
160 [H2125, 1.5U/mg, derived from *A. niger*] and β -glucosidase [49290, 7.7 U/mg, derived from
161 almonds]. Using these enzymes, a cocktail is prepared with a ratio of cellulase mix:
162 hemicellulase : β -glucosidase = 1:1:2 [unit basis], whereas the cellulase mix is prepared using
163 cellulase blend [SAE0020] and cellulase [C1184] with a ratio of 1:1.8 [unit basis].

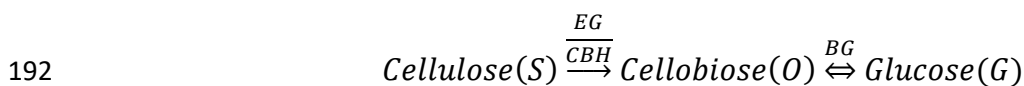
164 Enzymatic hydrolysis is performed in a stoppered conical flask in 50 ml solution of citrate
165 buffer (pH 4.8) along with sodium azide [0.1% (w/v)] in order to inhibit microbial
166 contamination. A series of hydrolysis experiments is performed, using different enzyme
167 cocktail (mix) concentrations (1.28, 6.66, 11.23 and 17.8g/L) at three different
168 substrate [pretreated mulberry powder] loadings (25, 50 and 125 g dry biomass/L). In each
169 hydrolysis experiment, every substrate (25, 50 and 125 g dry biomass/L) is supplemented with
170 0.2 g of hemicelluloses, taken from pretreatment liquor precipitate. Initially, the reaction
171 mixture, comprising of substrate, buffer and sodium azide, is agitated at 115 rpm. When the
172 temperature reached 50°C , enzymes are added into the reaction mixture to initiate
173 hydrolysis. Samples are withdrawn from the reaction mixture at different time intervals and
174 placed in a boiling water bath for 10 minutes to terminate the reaction by deactivating the
175 enzymes. Thereafter, free sugars are estimated using High Performance Liquid
176 Chromatography (HPLC).

177 2.8. *Analysis of sugars by HPLC*

178 Quantification of hydrolyzed sugars is carried out using HPLC (Make: Waters; Model: 2489)
179 fitted with an RI detector (Make: Waters; Model: 2414) for measuring the change in
180 refractive index of the column effluent passing through the flow-cell. A Brownlee amino
181 column (Make: PerkinElmer, USA; Material: N9303501) is used to separate sugars at
182 ambient temperature. The mobile phase used consists of HPLC grade acetonitrile and
183 ultrapure water [70:30 (v/v)] at a constant flow rate of 0.6 ml/min. The temperature of the RI
184 detector is maintained at 45°C with a sensitivity of 16, while the column temperature is kept
185 at 30°C. Sugars are finally quantified using standard curves generated with standard grade
186 glucose (CAS No.: 50-99-7) and xylose (CAS No.: 58-86-6).

187 2.9. *Proposed enzyme kinetics*

188 Enzymatic hydrolysis is a complicated process as it comprises of heterogeneous substrate and
189 an enzyme blend composed of endoglucanase (EG), cellobiohydrolase (CBH), β -glucosidase
190 (BG) and hemicellulase (XY). The overall phenomena of hydrolysis of biomass can be
191 described with the help of two heterogeneous reactions as follows:



193



195 The activity of an enzyme is directly correlated with its initial hydrolysis rate as well as the
196 maximum velocity of hydrolysis.

197 It is well established that the enzyme activity is inhibited with an increase in product
198 (*cellobiose* and *glucose*) concentration during enzymatic hydrolysis (Bezerra, Dias, Fraga, &
199 Pereira, 2006). Two types of inhibition, namely, competitive and noncompetitive, are

200 predominately present in hydrolysis systems and responsible for inhibiting enzyme activity.
 201 In order to assess the type of inhibition present, time-integrated expressions of competitive
 202 and noncompetitive models are considered(Andrić, Meyer, Jensen, & Dam-Johansen, 2010),
 203 *see* Table 2:

204 Table 2 Equations of competitive and noncompetitive inhibition.

Inhibition model	Equations
Competitive	$t = \frac{1}{k_{cat}E_0} \left(- \left(K_M + \frac{K_M}{K_I c} (cG_0 + S_0) \right) \ln \left(1 - \frac{c(G - G_0)}{S_0} \right) + \left(1 - \frac{K_M}{K_I c} \right) c(G - G_0) \right) \quad (3)$
Non-competitive	$t = \frac{1}{k_{cat}E_0} \left(- \left(K_M + \frac{K_M}{K_I c} (cG_0 + S_0) \right) \ln \left(1 - \frac{c(G - G_0)}{S_0} \right) + \left(1 - \frac{K_M}{K_I c} \right) c(G - G_0) + \frac{c}{2K_I} (G^2 - G_0^2) \right) \quad (4)$

205
 206 Here, k_{cat} is the apparent cellulase turn-over number, E_0 is the initial enzyme concentration
 207 (g/L), K_I is the enzyme–glucose complex dissociation constant or inhibition constant,
 208 whereas K_M represents the apparent Michaelis constant, which corresponds to the affinity
 209 between cellulose and cellulase. S_0 and G_0 stand for initial concentration of cellulose from
 210 pre-treated saw dust (g/L) and initial glucose concentration (g/L), respectively. c (=0.9)
 211 represents the reciprocal value of the number of glucose units present in a molecule of
 212 cellulose obtained from pre-treated saw dust. In order to evaluate the proposed models with
 213 experimental data, it is necessary to find out the values of the apparent cellulase turn-over

214 number k_{cat} and the inhibition constant K_I for each of the substrate loadings (25, 50, and
 215 125g/L), hydrolyzed with predetermined enzyme concentrations (1.28, 6.66, 11.23, 17.8g/L).
 216 The apparent kinetic parameters of the inhibition models are evaluated using a nonlinear
 217 least-square method. Each model is regressed on all available data. The quality assessment of
 218 the model prediction of net glucose concentration is based on the R^2 value of each of the
 219 nonlinear fits. The solution algorithm is developed with the help of MATLAB R2017a using
 220 the ode45¹ function. Based on a suitable fit of the kinetic model with the experimental data,
 221 the mode of inhibition executed by the end product (glucose) is estimated.

222 3. Results and Discussion

223 3.1. Chemical composition of saw dust

224 The chemical composition of crude Himalayan mulberry wood dust and its modifications
 225 during various stages of pretreatment is estimated (refer Table 3).

226

227 Table 3 Modification of chemical composition at different stages of combined pretreatment
 228 (CP) of Himalayan mulberry wood powder.

Various Stages of Pretreatment	Composition (Dry wt %)					
	Lignin	Glucan	Xylan	Mannan	Galactan	Other (Ash, Wax, Extractives) (wt %)
Crude	32.1±0.03	44.8±0.4	6.24±0.2	8.5±0.06	4.3±0.2	4.06±0.1

¹ode45 is a very useful function for solving non-stiff ordinary differential equation and it is a medium order method.

Autoclave (at 121°C for 1h)	29.77±0.3	52.27±0.2	17.96±0.4	-	-	-
Probe sonication (for 1h)	25.83±0.2	56.44±0.2	17.73±0.3	-	-	-
Alkaline-peroxide pretreatment (5h at 50°C)	23.97±0.3	58.38±0.89	17.65±0.2	-	-	-

229

230

231 It is observed from the table that crude Himalayan mulberry wood powder contains 44.8%
232 glucan whereas three different hemicellulosic sugars like xylan (6.24%), Mannan (8.5%) and
233 Galactan (4.3%) are also found in mulberry wood powder. Glucan is found as the corner-stone
234 of the lignocellulosic biomass and hemicellulosic sugars like mannan and xylan often exist in
235 a complex with Glucan and galactan, termed as glucomannan, (galacto) glucomannan and
236 glucoxylan (Geng, Sun, Sun, & Lu, 2003). Following Combined Pretreatment (CP), a
237 considerable depletion of lignin is observed along with elevation of glucan and xylose
238 content (Table 3) in relative percentages. However, no traces of mannan and galactan are
239 found in the biomass following combined pretreatment. Alkaline-peroxide pretreatment is
240 strongly correlated to primary solubilization and partial degradation of the macromolecular
241 hemicelluloses (J. Sun, Mao, Sun, & Sun, 2005). This significant increase in the xylose
242 content, following combined pretreatment, provides an evidence that in the mulberry wood
243 cell walls, xylose resides in the main chain of hemicelluloses while galactose and mannose
244 are probably present in side chains and are thus released relatively easily after pretreatment of

245 moderate severity. Dissolution of all galactose from Douglas fir (*Pseudotsuga menziesii*)
246 wood chips after only 30 min of pretreatment with 2% alkaline-peroxide pretreatment is also
247 reported (Alvarez-Vasco & Zhang, 2013). Removal of galactose strongly assists in
248 solubilizing the glucomannan structure. Even at low temperature (93°C), as soon as the wood
249 powder comes in contact with alkaline-peroxide solution, significant loss of glucomannan
250 (=75%) is reported (Wigell, Brelid, & Theliander, 2007). Moreover, presence of hydrogen
251 peroxide also facilitates removal of mannan moieties significantly from biomass (Alvarez-
252 Vasco & Zhang, 2013). On the other hand, xylan has a removal pattern that differs
253 significantly from that of glucomannan. A combination of high temperature and higher
254 peroxide concentration is needed to dissolve and remove xylose moieties present in the major
255 backbone chain of hemicelluloses. A limited loss of xylose is reported at temperatures below
256 139°C (Fang, Sun, & Tomkinson, 2000; Wigell et al., 2007).

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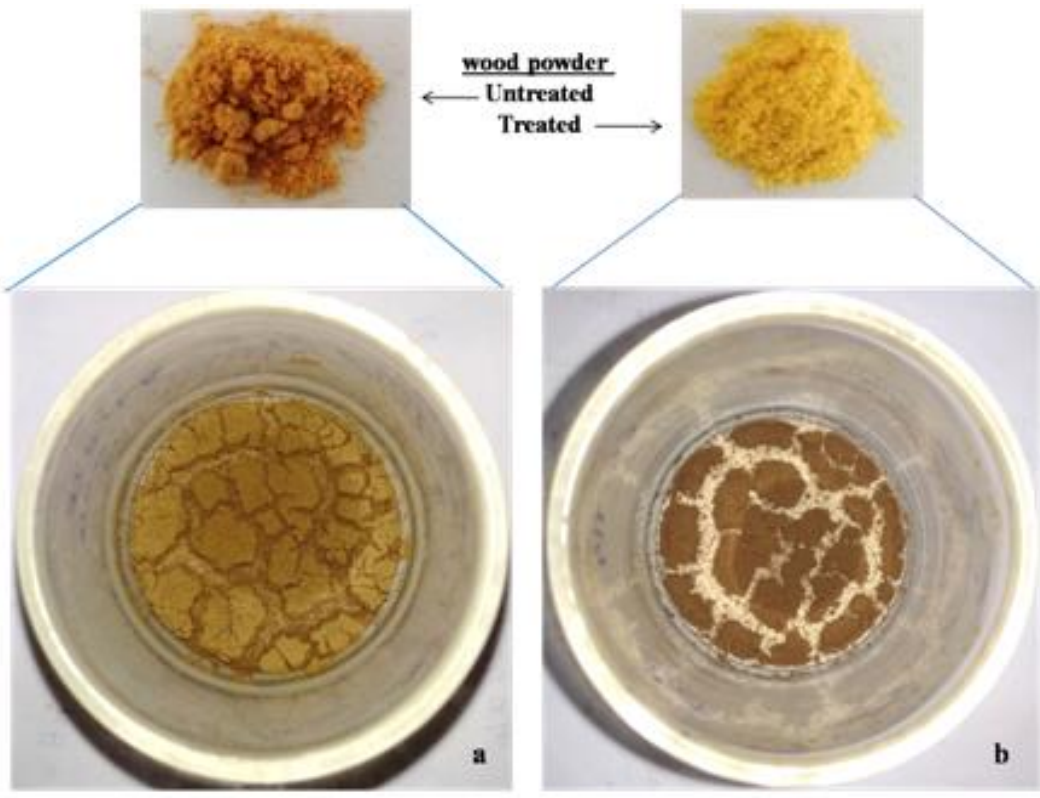
258 3.2. *Effect of various pretreatment procedures on mulberry powder*

259 Starting with an untreated biomass with a lignin content of 32.1 % (dry weight), CP is found
260 to remove lignin from untreated biomass more efficiently (23.97 % of dry weight remaining)
261 than alkaline peroxide (AP) based pretreatment (28.61% of dry weight remaining). Amount of
262 total carbohydrates increases from 63.84 wt % to 68.75 wt % following AP whereas to 74.03
263 wt % after CP. From this scenario, it can be inferred that combined pretreatment is
264 substantially more efficient as compared to AP for optimum delignification. This is crucially
265 beneficial for downstream enzymatic hydrolysis of the pretreated biomass.

266 There are various types of covalent and non-covalent linkages present in the lignocellulosic
267 matrix that make the matrix stable. Two important chemicals used in AP are NaOH and
268 H₂O₂. In alkaline conditions, H₂O₂ is readily decomposed into hydroxyl radicals and
269 superoxide anions. These radicals cleave several inter-unit bonds, introducing hydrophilic

270 carboxyl groups into the lignin structure and eventually dissolve lignin and hemicelluloses
271 into the pretreatment solution(Betts et al., 1991). While several ester and ether bonds are
272 easily broken during alkaline pre-treatment, most of these covalent bonds remain intact. In
273 CP, the mulberry powder is treated in an autoclave [pressure=15psi (*gauge*)], where high
274 pressure and temperature substantially help deforming the LB heteromatrix.Subsequently,
275 ultrasonic waves generate bubbles, which on collapsing due to wave compression, form
276 micro-jets enabling the cell walls to break, thereby deforming the matrix to a great extent
277 with an effective increase in the available surface area. This likely allows the radicals formed
278 from H₂O₂ to access a larger surface area of the lignocellulosic matrix at a time. Eventually
279 these radicals break and saponify more bonds present in the matrix. The dark brown color of
280 untreated biomass is transformed to light yellow after delignification followed by washing
281 [see Figure 1].

282



283

284 Figure 1 Acid insoluble lignin, on crucible bed, obtained from (a) untreated material and
285 (b) pretreated material (CP).

286

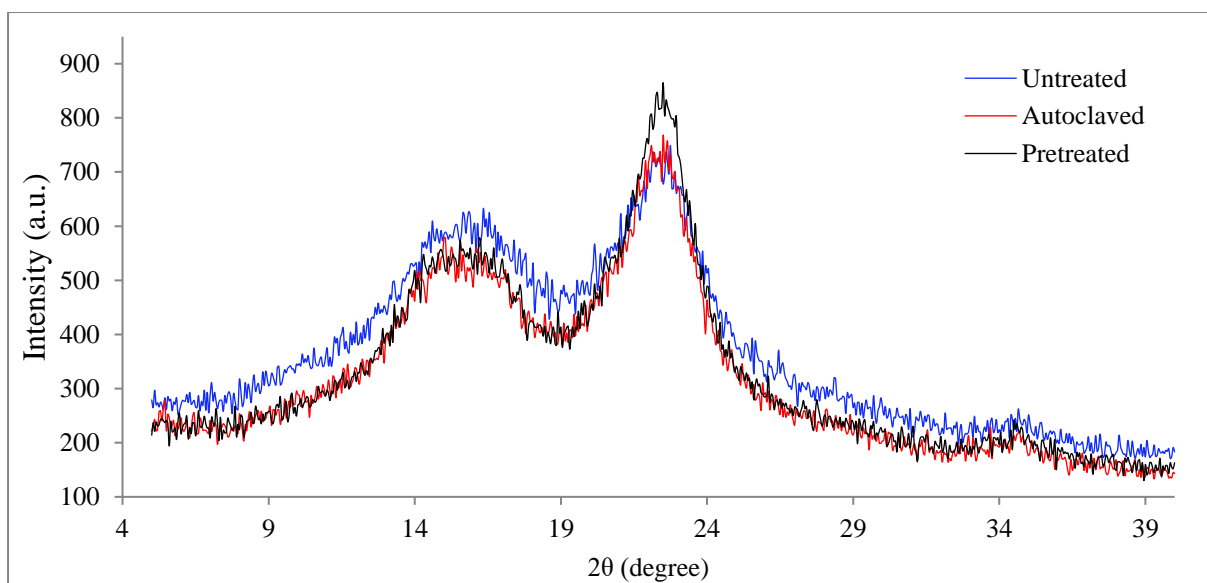
287 Obviously, removal of lignin (brown colour) from the wood powder causes discoloration of
288 the material. Additionally, being an efficient procedure for optimum delignification of
289 mulberry wood powder, pretreatment liquors acquired during various steps of combined
290 pretreatment (step I: autoclave, step II: probe sonication and step III: 5h long pretreatment
291 with alkaline-peroxide solution) are collected and then dissolved fractions of hemicelluloses
292 are precipitated from the liquor. A cumulative amount of 10.5% of total hemicelluloses is
293 finally recovered following CP and the same is further hydrolyzed along with pretreated
294 biomass (CP), in the downstream enzymatic hydrolysis.

295

296 3.3. Crystalline structure of biomass

297 The crystalline fingerprints of untreated and combined pretreated mulberry powder are
298 determined using XRD analysis as represented in Figure 2.

299



300

301 Figure 2 X-Ray diffraction patterns of untreated, autoclaved and pretreated (CP) material.

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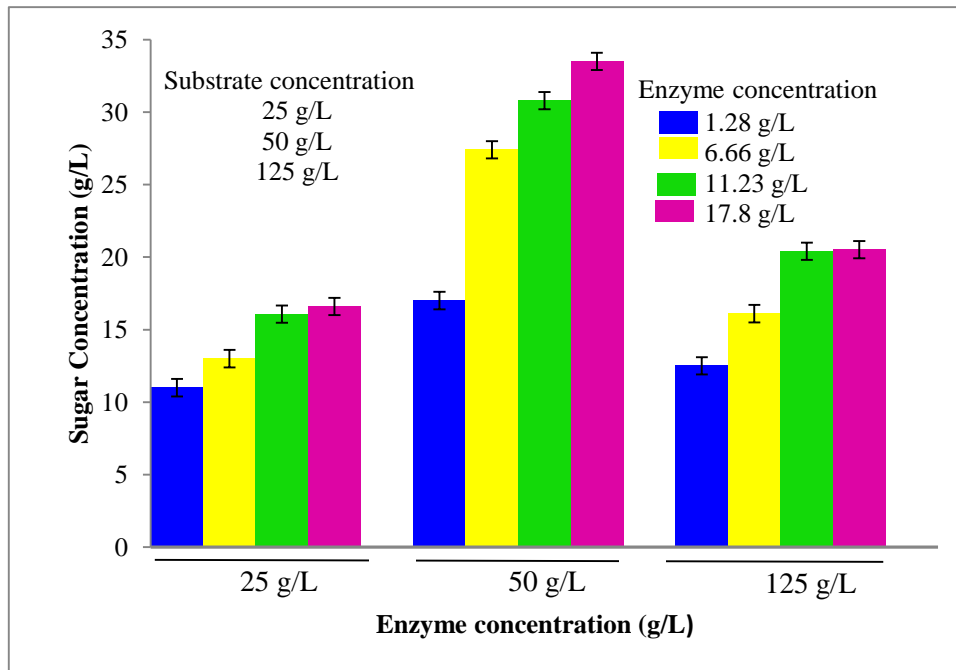
303 The XRD diffraction pattern of each sample exhibited two sharp peaks at around $2\theta=22.5^\circ$
304 and 16.8° which correspond to the (2,0) and (1,0) lattice planes of crystalline cellulose I,
305 respectively(S. Chen, Ling, Zhang, Kim, & Xu, 2018). Crystallinity of the mulberry powder
306 significantly increase after combined pretreatment (see Figure 2).Crystallinity of the
307 untreated materials is calculated as 19.91% whereas after autoclaving, the crystallinity
308 increases to 33.98% and eventually to 41.39% after completion of combined pretreatment.
309 This finding indicates removal of amorphous portion from the solid biomass material with
310 subsequent liberation of crystalline cellulose. Due to the exposure of untreated material to
311 high temperature and pressure during autoclaving, most of the bonds present in the
312 amorphous region of the lignocellulosic matrix might have broken. Further exposure of the
313 autoclaved material to ultra-sonication and downstream 5h long alkaline-peroxide treatment
314 might have helped in breaking the additional covalent and non-covalent bonds in the
315 amorphous region. Amorphous regions of lignocelluloses are composed of lignin and
316 hemicelluloses. It is therefore evident that combined pretreatment leads to cleavage of bonds
317 in this region and eventually solubilize hemicelluloses and lignin in the pretreatment liquor
318 and finally increases the global crystallinity of biomass.

319

320 3.4. *Enzymatic saccharification of combined pretreated (CP) mulberry powder*

321 The results of enzymatic hydrolysis of pretreated mulberry sawdust are shown in Fig.3.
322 It is evident from Fig.3 that, for a particular substrate concentration, yield of total reducing
323 sugar increases with an escalated enzyme concentration, varying in the range of 1.28 g/L to
324 17.8 g/L. Additionally, the effect of substrate loading for a particular enzyme concentration
325 can also be explained from the figure. An increased amount of total reducing sugar can be
326 achieved with a particular enzyme concentration by increasing substrate loading from 25g/L

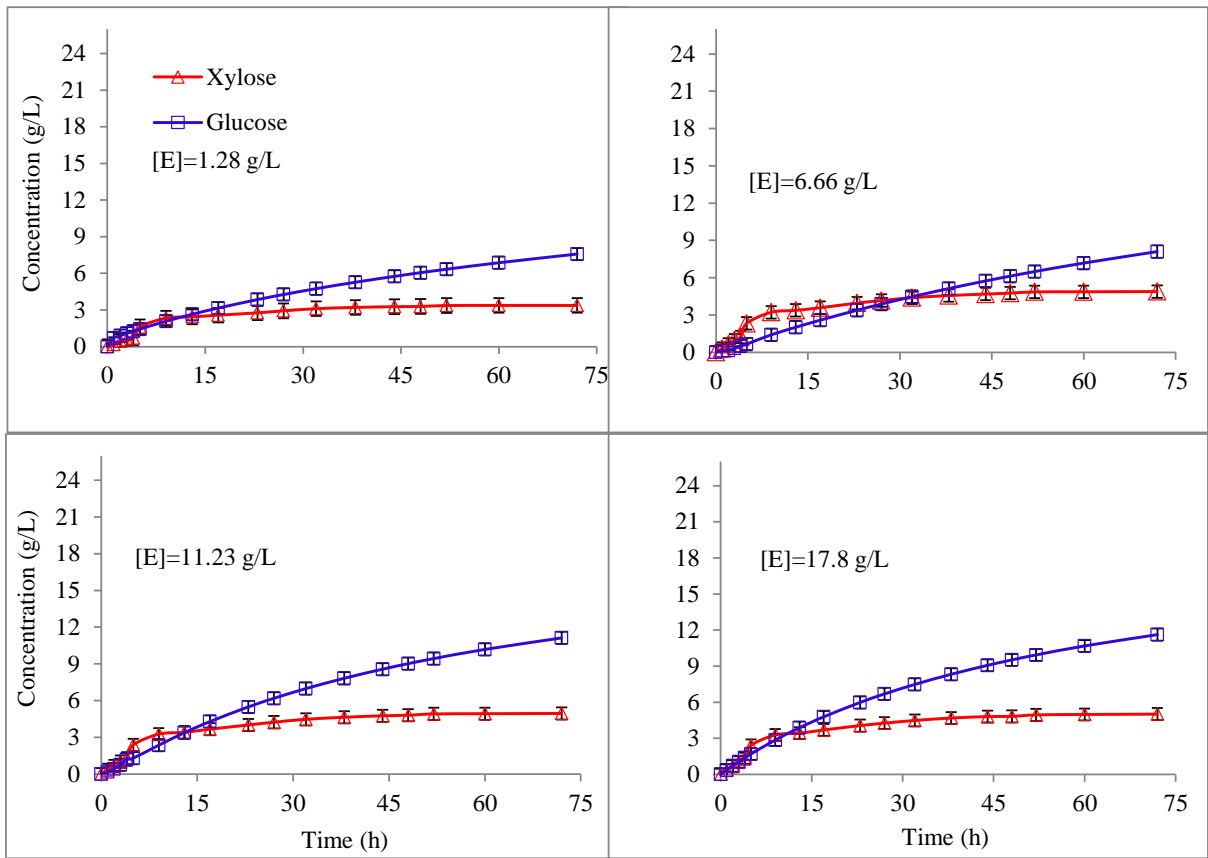
327 to 50 g/L. Surprisingly, further increment of substrate loading till 125 g/L resulted in a sugar
 328 yield lower than the same achieved with 50 g/L substrate concentration. The concentrated
 329 (125 g/L substrate) reaction mixture forms a thick slurry that induces mass transfer limitation,
 330 thereby apparently reducing the sugar yield (O'Dwyer, Zhu, Granda, & Holtzapple, 2007).
 331



332
 333 Figure 3 Yield of total reducing sugar (g/L) using various substrate (25, 50, 125 g/L) and
 334 enzyme (1.28, 6.66, 11.23, 17.8 g/L) concentrations.

335
 336 Apart from glucose, xylose is considered as the second most important monomeric sugar for
 337 downstream fermentation to generate second generation biofuel. Therefore, in the present
 338 study, individual C5 (xylose) and C6 (glucose) sugar concentrations generated under each
 339 hydrolysis condition are measured for about 3 days and analyzed using HPLC. Results are
 340 shown in Fig.4. During initial stages of hydrolysis, xylose liberates at a higher rate than
 341 glucose. At around 10 h of hydrolysis, xylose reaches an equilibrium concentration.
 342 Xylanase, present in the enzyme system obviously attacks the hemicellulose fraction of the

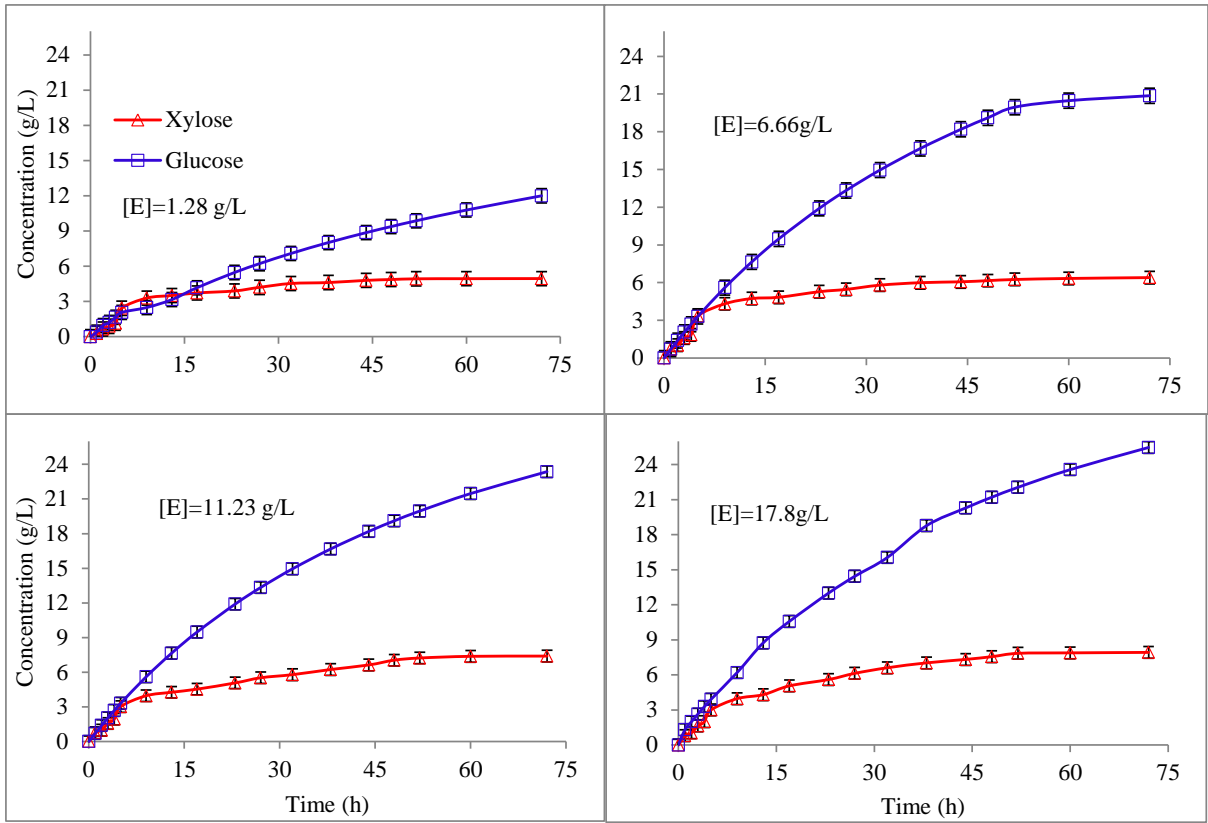
343 biomass efficiently and hydrolyzes it rapidly during the initial stages of hydrolysis, before
344 getting deactivated due to product (glucose) inhibition.



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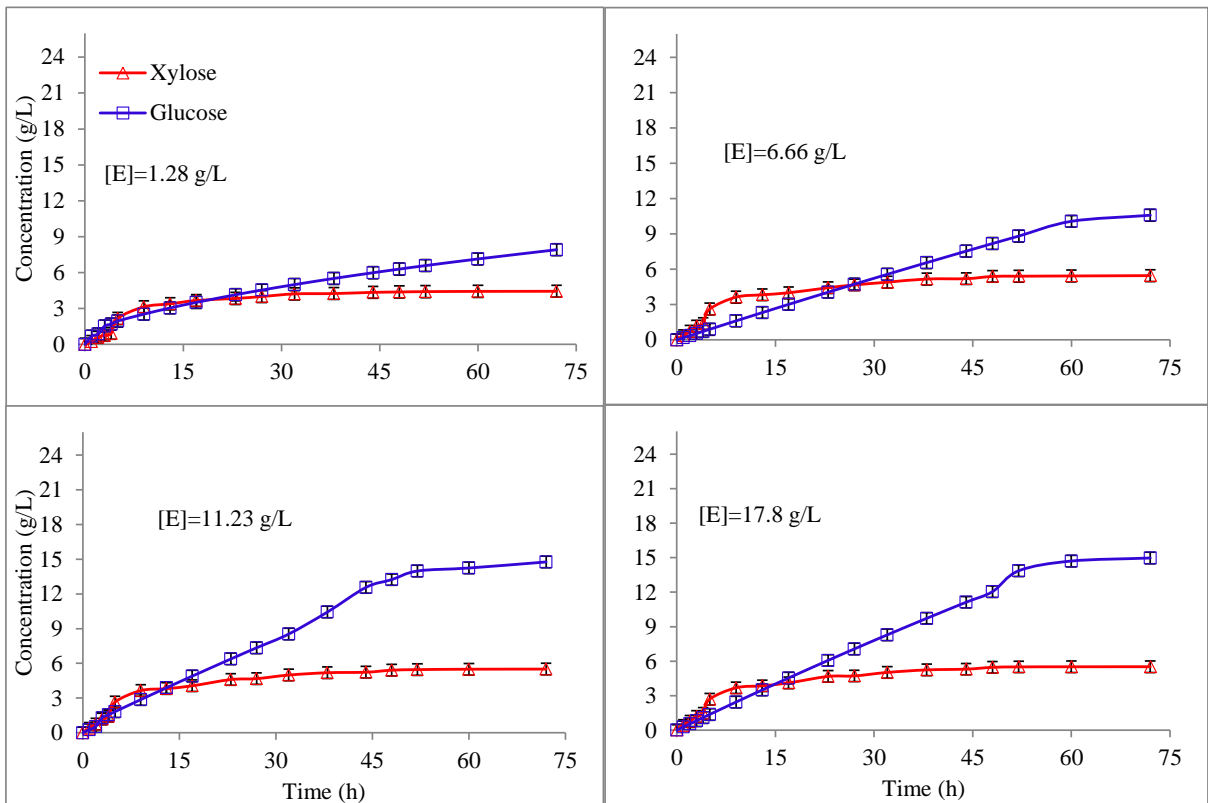
Figure 4(a)



347

348

Figure 4(b)



349

350

Figure 4(c)

351

352 Figure 4 Time dependent production of glucose and xylose from pretreated biomass with
353 different enzyme concentrations (a) 25g/L (b) 50g/L, (c) 125g/L.

354

355 Most likely product inhibition is not solely responsible for the rapid fall in xylose generation
356 rate. In fact, inhibitory effects exhibited by hemicellulose derived sugars are known to be
357 much less significant than the same by glucose (Xiao, Zhang, Gregg, & Saddler, 2004). It is
358 inferred that before 10 h of hydrolysis, all immediately accessible hemicellulose get
359 hydrolyzed. Also, any further increase in xylose concentration (at a much lower rate) is
360 connected to the hydrolysis of cellulose, proceeding, successively giving access to further
361 xylose moieties for hydrolysis (Lin, Yan, Liu, & Jiang, 2010).

362 At the same time, hydrolysis of cellulose to glucose proceeds at a lower rate than xylose
363 liberation but proceeds for much longer time and at a successively decreasing rate. Exo- and
364 endo-glucanase both attack and hydrolyze cellulose chains from cellulose fibrils attached to
365 the core matrix, followed by generation of cellobiose, while β -glucosidase immediately starts
366 hydrolyzing cellobiose molecules into glucose. Successive decrease in glucose production
367 rate is probably due to successive increase in feedback inhibition by the accumulating
368 glucose (Lai et al., 2014; Li et al., 2013). Total glucan and xylan recovery are estimated after
369 hydrolysis is complete (see Table 4).

370

371 Table 4 Carbohydrate recovery following completion of hydrolysis after combined
372 pretreatment (CP).

Substrate	Enzyme	Carbohydrate conversion and yield
-----------	--------	-----------------------------------

concentration (g/L)	Concentration (g/L)	Glucan conversion (%)	Glucose Yield (%)	Xylan conversion (%)	Xylose Yield(%)
25	1.28	51.07	45.963	39.95	35.156
	6.66	54.58	49.122	58.03	51.0664
	11.23	74.93	67.437	58.86	51.7968
	17.8	78.30	70.47	59.45	52.316
50	1.28	38.36	34.524	40.15	35.332
	6.66	71.50	64.35	51.91	45.6808
	11.23	80.06	72.054	60.01	52.8088
	17.8	87.26	78.534	64.23	56.5224
125	1.28	10.83	9.747	17.03	14.9864
	6.66	14.49	13.041	20.95	18.436
	11.23	20.23	18.207	21.14	18.6032
	17.8	20.50	18.45	21.18	18.6384

373

374 It can be seen from Table 4 that with a particular substrate-enzyme combination, a relatively
375 greater percentage of glucan is recovered with a relatively less amount of xylose. Similar
376 kind of outcome is reported where 85% glucan and 70% of xylose have been achieved with

377 an enzyme cocktail composed of cellulase, beta-glucosidase, xylanase and β -xylosidase (Qing
378 & Wyman, 2011). On the other hand, a similar trend is observed with a recovery of 40%
379 glucan and 27% xylan using an enzyme-cocktail, composed of endoglucanase,
380 cellobiohydrolase, β -glucosidase, endoxylanase, β -xylosidase and acetylxylan esterase (Barr,
381 Mertens, & Schall, 2012). These results elucidate that efficient and complete hydrolysis of
382 xylan is partially hindered due to the structural complexity of hemicellulose along with
383 specific requirements of hemi-cellulolytic enzymes (like formation of a precise transition
384 state in order to bind the substrate more efficiently and effectively). Additionally, the
385 competition between cellulase and hemicellulase for binding on the reactive sites of cellulose
386 make hemicellulase limited for xylan hydrolysis. Stronger binding of cellulase to xylan in
387 comparison to glucan, makes xylan occupied with cellulase faster. As a result, xylanase,
388 present in hemicellulase, cannot bind efficiently with xylan backbone and the same is another
389 potential reason for the incomplete hydrolysis of xylose. Chemical structure and
390 modifications also play a crucial role in the hydrolysis of xylan. Presence of substitutes like
391 4-O-Meglucuroic acid in xylan backbone can potentially deactivate endoxylanase (Barr et al.,
392 2012). Acetylation of xylan can also inhibit endoxylanase to a great extent. As these
393 substitutes are removed, during pretreatment and enzymatic hydrolysis, xylan becomes less
394 soluble and form aggregates that offers steric-hindrance, thereby retarding de-polymerization
395 (Wyman et al., 2005).

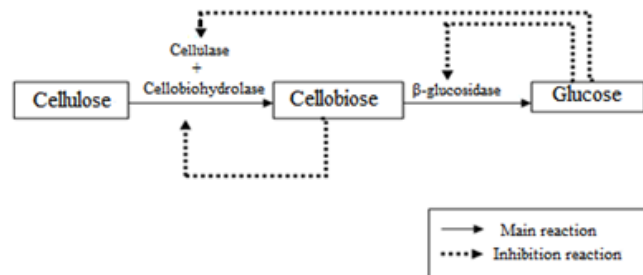
396

397 *3.5. Determination of Inhibition pattern and kinetic parameters*

398 In enzyme-substrate systems, two types of inhibition can be present: substrate inhibition and
399 product inhibition. Product inhibition has a serious influence on cellulose hydrolysis and
400 enzyme kinetics (Corazza, Calsavara, Moraes, Zanin, & Neitzel, 2005). Based on this and the
401 observation of successive decrease in glucose production rate in Fig.4., inhibition kinetics

402 associated with glucose production is investigated. A schematic diagram of inhibition
403 phenomena is presented in Fig.5. (Andrić et al., 2010).

404



405

406 Figure 5 Schematic representation of probable inhibition phenomena during enzymatic
407 hydrolysis of LB.

408

409 Though cellobiose and glucose both act as inhibitors during enzymatic hydrolysis of
410 lignocellulosic biomass, specific inhibition type(s), exerted only by glucose is considered
411 here. However, the biomass is a heterogeneous substrate and the commercial cellulolytic
412 enzyme system can be composed of different numbers and types of enzymes. It is
413 reasonably difficult to measure the initial intrinsic or apparent hydrolysis rate for this
414 cellulose-cellulase system. Thus, Dixon or Lineweaver-Burk plots cannot be applied for
415 evaluating the type of inhibition exhibited by glucose (Zhao, Wu, Yan, & Gao, 2004).
416 Competitive inhibition occurs in cellulose-cellulase systems and in between glucose and β -
417 glucosidase (Andrić et al., 2010; Lee & Fan, 1983). In competitive inhibition, the product
418 competes with the substrate for binding to the active site of the enzyme and inhibits it.
419 Increasing the substrate concentration is a way to overcome competitive inhibition exerted
420 by the key product. In the present study, increased product recovery is observed by
421 increasing the substrate concentration from 25 g/L to 50 g/L (see Fig. 3.). Following
422 hydrolysis of 25g/L concentrated substrate with predefined enzyme concentration, glucose

423 and xylose are produced as products and both compete for the active sites of the enzymes
 424 with the substrate because of their binding tendency to identical active sites. Therefore, those
 425 products act like inhibitors and displace the substrate from the active site of the enzymes and
 426 form an enzyme-inhibitor complex ultimately leading to loss of enzymes. A thermodynamic
 427 principle explains this inhibitory mechanism more accurately. Two equilibria, one between
 428 enzyme and inhibitors/products and the other between enzyme and substrate, exist in parallel
 429 and these two equilibria are not independent as the enzyme-substrate complex and enzyme-
 430 inhibitor complex equilibrate with the same free enzyme pool. Therefore, increasing enzyme-
 431 substrate concentration is the only way to eliminate the probability of enzyme-inhibitor
 432 complex formation. Increasing the substrate concentration from 25 g/L to 50 g/L offers more
 433 substrate for a particular enzyme concentration which in turn forms enzyme-substrate
 434 complex more frequently than enzyme-inhibitor complex. This supports the presence of
 435 competitive inhibition exerted by glucose. On the other hand, in noncompetitive inhibition,
 436 glucose would bind to the allosteric sites of the enzymes, thereby reducing its surface
 437 activity and inhibiting the enzyme non-competitively. Glucose has an equal binding
 438 affinity to both the free enzymes and the enzyme-substrate complex. In the present study,
 439 constants associated with competitive and noncompetitive inhibition are evaluated using
 440 equations (3) and (4) and presented in Table 5.

441 Table 5 Estimated parameters of inhibition kinetics.

Enzyme concentration (g/L)	Substrate concentration (g/L)	Competitive			Noncompetitive		
		k_m (g/L)	k_{cat} (sec ⁻¹)	K_i (g/L)	k_m (g/L)	k_{cat} (sec ⁻¹)	K_i (g/L)
1.28	25	2.288	0.142	1.594	0.057	.321	1.152

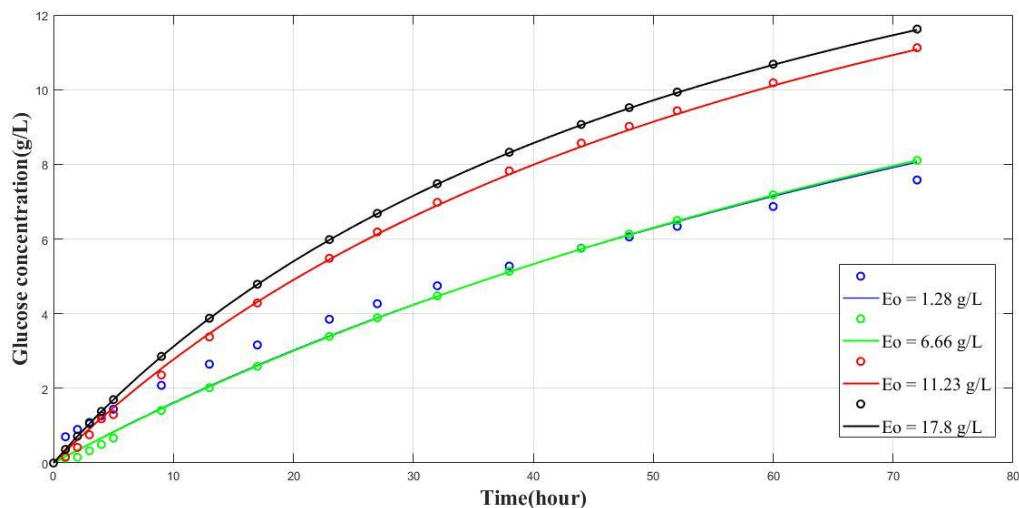
	50	2.274	.185	1.71	159.73	1.55	9.41
	125	2.227	.113	.658	-54.44	$.723 \times 10^6$.088
6.66	25	2.206	0.027	1.673	1.34	-12.29	-0.005
	50	3.922	0.111	2.213	1.326×10^6	6.02×10^3	14.39
	125	.865	0.024	.786	11.94	-20.715	-0.006
11.23	25	3.652	0.032	2.248	1.407	-6.2	-0.012
	50	4.222	0.065	2.816	432.89	0.91	48.34
	125	1.08	0.019	.938	16.19	-12.57	-0.012
17.8	25	6.162	0.027	3.196	1.33	-4.37	-0.012
	50	9.766	0.058	6.322	70.65	0.13	45.77
	125	1.140	.014	.992	5.99	-9.84	-0.009

442

443 The value of inhibition constant increased gradually with escalated enzyme concentration
444 for a particular substrate loading during competitive inhibition (refer Table 5), indicating
445 reduced inhibition by the product formed. The same trend is also observed for a particular
446 enzyme concentration with escalated substrate concentration from 25 g/L to 50 g/L.
447 However, for a particular enzyme concentration, value of the inhibition constant decreased
448 when the substrate loading was increased to 125 g/L. As discussed above, this is due to the
449 film resistance created by the formation of thick slurry at such high substrate
450 concentrations and nonproductive irreversible adsorption of enzymes on surfaces of
451 substrate. Enzymatic hydrolysis with a loading of 50 g/L substrate, along with 17.8 g/L

452 enzyme concentration, is found optimum with an inhibition constant of 6.322 with
 453 minimum inhibition and maximum production of sugars. On the contrary, in case of
 454 noncompetitive inhibition, many of the kinetic constants appear with negative values
 455 which make this mode of inhibition unlikely for the current study. In early studies negative
 456 kinetic constants were also found for noncompetitive inhibition when pretreated cellulose
 457 was hydrolyzed with cellulase enzyme system [enzyme commission no.: 3.2.1.4](Caminal,
 458 Lopez-Santin, & Sola, 1985). It can be inferred that the product inhibition of the system
 459 studied here follows a competitive mode of inhibition. The theoretical data set, evaluated
 460 using the competitive kinetic constants in equation (3), is compared with experimental data
 461 and represented in Fig.6 with a value of $r^2 \geq 0.91$ except for the system composed of 125
 462 g/L substrate and 1.28 g/L enzyme ($r^2=0.89$).

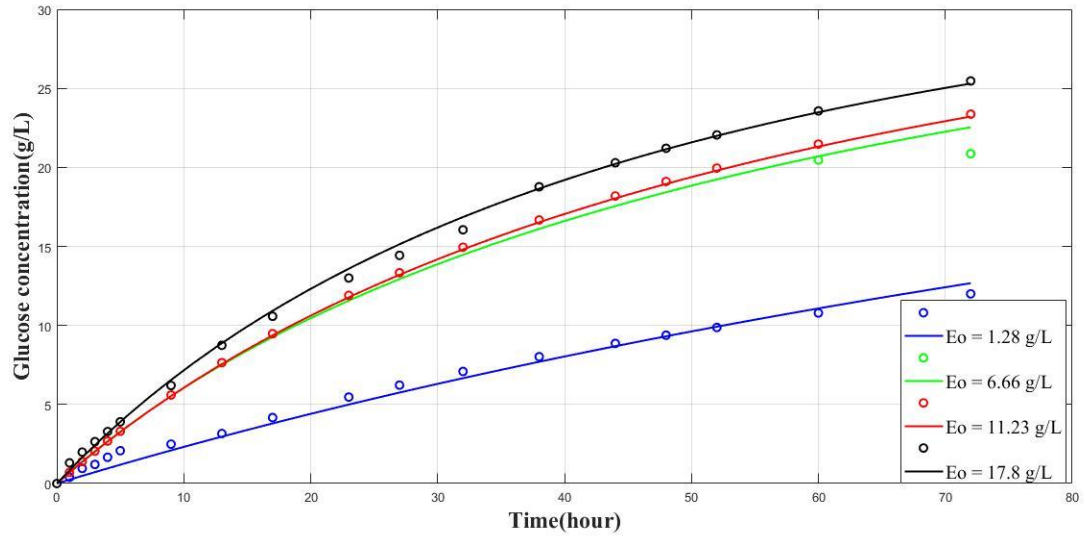
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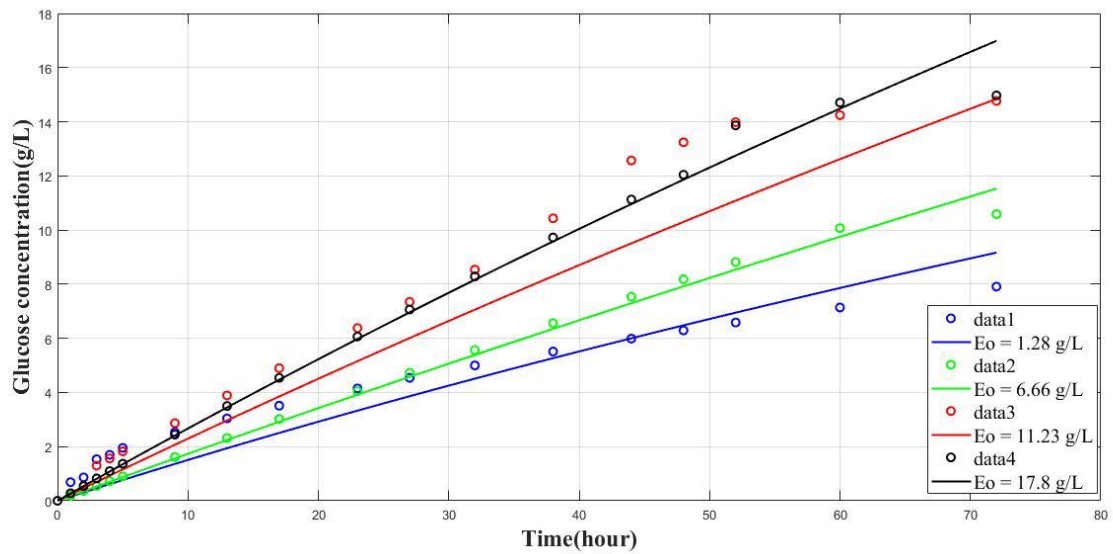
Figure 6(a)



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Figure 6(b)



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Figure 6(c)

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Figure 6 Predicted inhibition model outputs (curves with solid lines) along with

471

observed data (markers) for enzymatic hydrolysis of (a) 25g/L, (b) 50 g/L and (c) 125

472

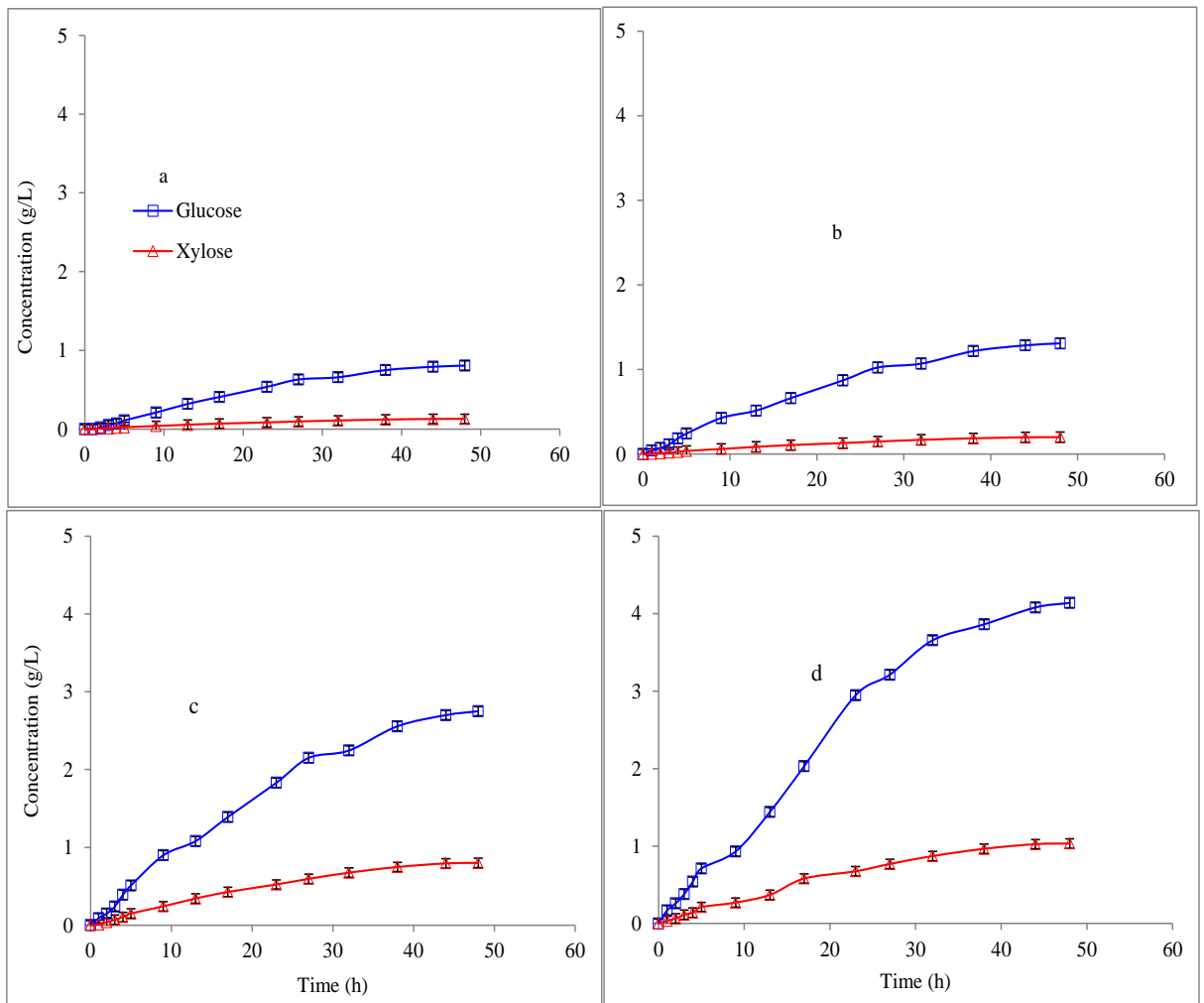
g/L pretreated substrate with various enzyme concentrations.

473

474 3.6. Effect of each pretreatment step on enzymatic hydrolysis outcome

475 In order to understand the effect of each pretreatment step on enzymatic hydrolysis,
476 substrate of a constant concentration (50 g/l) is withdrawn after each pretreatment process
477 and hydrolyzed with four known enzyme concentrations (1.28, 6.66, 11.23 and 17.8 g/l)
478 [See Figure 7 and Figure 8]. Additionally, recovery of glucan and xylan after hydrolysis of
479 the autoclaved and probe sonicated biomass is assessed in order to understand the effect
480 of each step of combined pretreatment on hydrolysis.

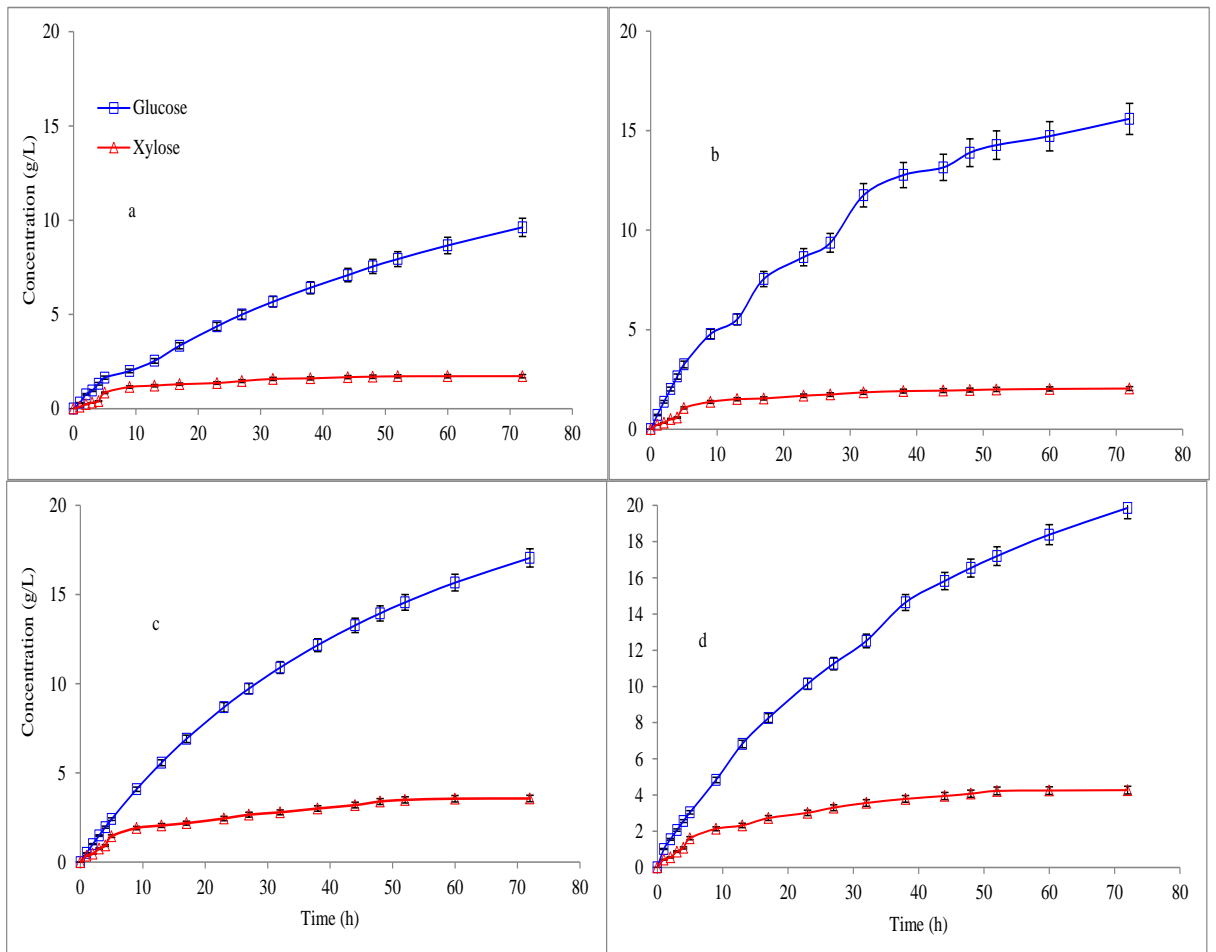
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482

483 Figure 7 Enzymatic hydrolysis of autoclaved substrate (50 g/L) with different known

484 enzyme concentrations [(a) 1.28 g/L, (b) 6.66 g/L, (c) 11.23 g/L, (d) 17.8 g/L].



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Figure 8 Enzymatic hydrolysis of probe sonicated substrate (50 g/L) with different known enzyme concentrations [(a) 1.28 g/L, (b) 6.66 g/L, (c) 11.23 g/L, (d) 17.8 g/L].

Table 6 Recovery of glucan and xylan (wt %) after hydrolysis using each step of combined pretreatment (CP) of the LB (substrate).

Pretreatment step	Substrate concentration (g/L)	Enzyme Concentration (g/L)	Carbohydrate conversion and yield			
			Glucan conversion (%)	Glucose Yield (%)	Xylan conversion (%)	Xylose Yield(%)

Autoclaved	50	1.28	3.06	2.754	1.44	1.2672
		6.66	4.97	4.473	2.23	1.9624
		11.23	10.50	9.45	8.90	7.832
		17.8	15.84	14.256	11.47	10.0936
Probe sonicated	50	1.28	34.09	30.681	19.01	16.7288
		6.66	55.24	49.716	23.11	20.3368
		11.23	60.42	54.378	40.29	35.4552
		17.8	70.38	63.342	48.3	42.504

492

493 It can be seen from the Figures (refer Figure 7 and Figure 8) that a minor amount of
494 glucose and xylose are generated following hydrolysis of autoclaved substrate. Using
495 lower concentration of enzymes like 1.28 g/L, only 3.06% glucan and 1.44% of xylan get
496 converted. Additionally, elevated yield of carbohydrate is accomplished with augmented
497 enzyme concentration with a corresponding yield of 14.256% glucose and 10.094% of
498 xylose (*enzyme concentration: 17.8g/L*). After autoclaving, the substrate still contains
499 considerable amount of lignin which is responsible for the irreversible and non-
500 productive binding of supplied enzymes on lignin surface. On the other hand, autoclaving
501 helps in the solubilization of lignin and hemicellulose side chains, keeping crystallinity of
502 cellulose unaffected. As a result, enzymes cannot substantially de-polymerize
503 lignocellulosic carbohydrates. On the contrary, better results are observed when probe
504 sonicated substrate (50g/L) is hydrolyzed with same enzyme concentrations [(a) 1.28 g/L,

505 (b) 6.66 g/L, (c) 11.23 g/L, (d) 17.8 g/L]. The highest level of glucose and xylose yield
506 are 63.342% and 42.504% respectively, following hydrolysis of probe sonicated substrate
507 with 17.8 g/L enzyme concentration. Probe sonication helps in the removal of amorphous
508 cellulosic materials along with considerable amount of lignin. Supplied enzymes can thus
509 access crystalline cellulose. Therefore, following hydrolysis of probe sonicated material,
510 an elevated concentration of glucose and xylose is observed.

511

512 **4. Conclusion**

513 By systematically comparing two different pretreatment methods (CP and AP), CP is found
514 to be a better option in terms of preparation of biomass for subsequent enzymatic hydrolysis
515 as it increases the accessible surface area of biomass for accommodating enzymes in the
516 active sites as well as increases the crystallinity. Appropriate enzyme loading per unit
517 substrate mass is found to be a crucial factor for optimal hydrolysis outcomes. Enzyme
518 cocktails used in the present study are found to be inhibited competitively with produced
519 glucose (*product inhibition*). Apart from enzyme and substrate loadings, the role of water is
520 also found to be crucial for retaining the optimum activity of enzymes and for efficient
521 hydrolysis of substrates. Lack of sufficient water in slurry (with large fraction of solids)
522 hinders the free movement and mixing of enzymes with its substrates, eventually leading to
523 inefficient hydrolysis. This condition even inhibits the enzymes at a relatively early stage of
524 hydrolysis.

525

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534

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