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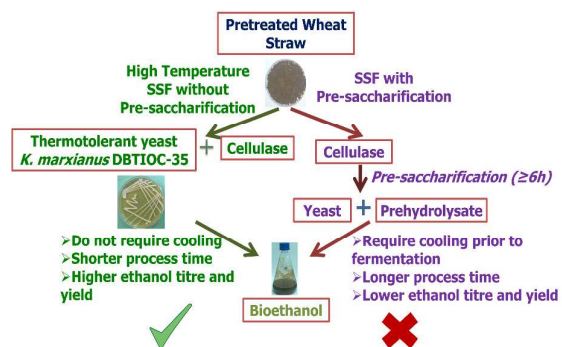
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Application of thermotolerant yeast *Kluyveromyces marxianus* DBTIOC-35 in SSF decreases overall process time, and increases yield by allowing elimination of presaccharification step and use of high biomass concentration.



1           **Second generation bioethanol production at high gravity of pilot-scale**  
2           **pretreated wheat straw employing newly isolated thermotolerant yeast**

3                           *Kluyveromyces marxianus* DBTIOC-35

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13

14 **Abstract**

15 Second-generation bioethanol production was studied by a newly isolated thermotolerant  
16 yeast strain at 42 °C and above using pilot-scale dilute acid pretreated wheat straw (WS) as  
17 feedstock. This strain was identified as *Kluyveromyces marxianus* DBTIOC-35 by  
18 biochemical characterization as well as molecular phylogenetic analysis of ITS-5.8S rRNA  
19 gene and D1/D2 domain of the 26S rRNA gene after PCR amplification and sequencing.  
20 Simultaneous saccharification and fermentation (SSF) at 42 and 45°C using 10 % biomass  
21 loading resulted in ethanol titer of 29.0 and 16.1 g/L, respectively. At 42°C ethanol  
22 productivity was higher during SSF (0.92 g/L/h) than separate hydrolysis and fermentation  
23 (SHF) (0.49 g/L/h) at 20% biomass loading. Results indicated that at 20% biomass  
24 loading, SSF without pre-saccharification led to more ethanol production (66.2 g/L with  
25 83.3% yield) at a faster rate than SSF with pre-saccharification (PSSF) which produced  
26 ethanol titer of 61.8 g/L, 77.7% yield and productivity of 0.86 g/L/h. Based on these  
27 findings, application of newly isolated yeast *K. marxianus* DBTIOC-35 in SSF of  
28 lignocellulosic biomass can eliminate pre-saccharification step which is a novel advantage  
29 of thermotolerant yeasts in terms of cutting down the overall biomass to bioethanol process  
30 time and enhancing bioethanol titer, yields and productivities.

31 **Keywords:** SSF; thermotolerant yeast; *Kluyveromyces*; bioethanol; wheat straw; dilute-  
32 acid pilot scale pretreatment

33

## 34 **Introduction**

35 Bioethanol production is considered a green technology for mitigating greenhouse  
36 gas emission and partial replacement of fossil transportation fuels (such as gasoline).<sup>1, 2</sup>  
37 Bioethanol production has increased enormously during the past decade. Bioethanol made  
38 from lignocellulosic biomass (such as corn stover, rice straw, sugarcane bagasse, wheat  
39 straw etc.) is termed second generation (2G) bioethanol in contrast to first-generation  
40 bioethanol that is derived from sugar and starch based materials (such as sugarcane and  
41 grains).<sup>3</sup> Feedstock for 2G bioethanol are abundant, renewable, cheaper and do not  
42 compete with food resources. Of the various agricultural lignocellulosic wastes wheat  
43 straw has the second largest availability of approximately 354.34 million tons with an  
44 estimated bioethanol production potential of at least 104 Giga litres.<sup>4</sup> Due to food versus  
45 fuel concerns developing nations like India cannot utilize food resources (such as grains or  
46 sugar) for biofuel production and must rely upon renewable lignocellulosic biomass.

47 2G bioethanol production from lignocellulosic feedstock involves pre-treatment,  
48 saccharification, fermentation and ethanol recovery. Pre-treatment helps in decreasing the  
49 crystallinity of cellulose, increasing biomass surface area, removing hemicellulose, and  
50 breaking lignin seal to make cellulose more accessible to cellulase enzymes. The  
51 biochemical route for bioethanol generation relies upon conversion of cellulose (and  
52 hemicelluloses) into fermentable sugars using (hemi-) cellulolytic enzymes followed by  
53 fermentation of sugars into ethanol via separate hydrolysis and fermentation (SHF) or  
54 simultaneous saccharification and fermentation (SSF).<sup>5</sup> Some bottlenecks to economically  
55 viable biochemical conversion processes include better pre-treatment process to increase  
56 enzyme accessibility, the cost and efficiency of cellulase enzymes, blending of cellulase  
57 enzymes to achieve better hydrolysis,<sup>6</sup> process and strain engineering to improve xylose  
58 and glucose co-fermentation<sup>5</sup> etc. In its original form, the batch SSF reactor initially

59 contains substrate, enzymes as well as yeast cells at the intended concentrations.<sup>5</sup> Major  
60 drawback of SSF process is mis-match of optimum temperatures of enzymes and most of  
61 fermenting microorganisms (e.g. the optimum temperatures are approximately 50 °C and  
62 30-35 °C, respectively). Therefore, several modifications in SSF have been suggested,  
63 such as inclusion of a pre-saccharification step at optimum enzyme temperature to allow  
64 sugar hydrolysis before cooling down to yeast optimum fermentation temperature,<sup>1,7</sup>  
65 varying the temperature during SSF and use of fed-batch approach for biomass addition.<sup>8-9</sup>  
66 Though these alterations can enhance the ethanol titer, the overall fermentation time is also  
67 increased resulting in low productivities that are undesirable for commercially viable  
68 process. Thermotolerance of the fermenting microorganisms is vital aspect in this regard.<sup>10</sup>  
69 Thermotolerant yeasts strains that have been evaluated in SSF process  
70 include *Kluyveromyces marxianus*, *Fabospora fragilis*, *Saccharomyces uvarum*, *Candida*  
71 *brassicae*, *C. lusitaniae*, etc. Use of thermotolerant yeasts can allow fermentation at  
72 temperatures closer to the optimal range (around 50°C) of the enzymes. Therefore, high  
73 degree of hydrolysis and subsequent higher ethanol yields can be obtained within lower  
74 time making SSF process more efficient. Other benefits of bioethanol fermentations at  
75 high temperature ( $\geq 40^{\circ}\text{C}$ ) are faster fermentation rates, reduction in cooling costs and  
76 prevention of contamination.<sup>11</sup> Therefore, application of thermotolerant microorganisms  
77 that can ferment high ethanol concentration can be a major development in SSF process.  
78 Higher temperatures are reported to negatively affect yeast viability and growth<sup>12</sup> which  
79 makes the search for better thermotolerant microorganisms even more necessary.

80 The aim of the current study was to enhance bioethanol production using dilute  
81 acid pilot scale pretreated wheat straw. To achieve this, indigenous strain of thermotolerant  
82 yeast was evaluated for fermentation at  $\geq 42^{\circ}\text{C}$  under different process configurations.  
83 Though there are few reports on use of thermotolerant yeasts, to the best of our knowledge

84 no comparable literature has highlighted the importance of thermotolerant yeasts in  
85 decreasing the lignocellulose to ethanol bioconversion process time by complete  
86 elimination of presaccharification step and enhancement of productivities during SSF and  
87 only fewer studies have utilized pilot scale pretreated biomass in SSF for ethanol  
88 production.

## 89 **Experimental**

### 90 **Materials**

91 Peptone, yeast extract, D-glucose and all other medium components used in this  
92 study were procured from HiMedia, India. Antibiotics ampicillin and streptomycin were  
93 purchased from MP Biomedicals, USA and Sigma-Aldrich, USA, respectively.  
94 SacchariSeb C6, a commercial cellulase powder, was purchased from Advanced Enzymes  
95 Inc, India. Wheat straw was obtained from local farmers.

### 96 **Fermenting microorganism and culture conditions**

97 Whey samples collected in sterile bottles from local dairies at Noida, India were  
98 used for isolation of thermotolerant yeasts by enrichment culture technique using YPD  
99 medium (yeast extract - 1%, peptone - 2%, glucose - 5%, filter sterilized ampicillin and  
100 streptomycin - 50 mg/L each). After enrichment for 48 h, the developed yeast colonies  
101 were evaluated qualitatively and quantitatively for bioethanol production at  $\geq 40^{\circ}\text{C}$ . The  
102 most promising isolate DBT-IOC-35 was selected for this study on basis of maximum  
103 ethanol titer, yield and ethanol efficiency during fermentation on glucose at  $42^{\circ}\text{C}$  (not  
104 shown). This strain was maintained at  $4^{\circ}\text{C}$  on YPD agar slants and at  $-20^{\circ}\text{C}$  as glycerol  
105 stock for short and long term storage, respectively. Yeast fermentation media (YFM) used  
106 for bioethanol production contained (%):  $\text{KH}_2\text{PO}_4$ , 0.1;  $\text{NH}_4\text{Cl}$ , 0.03;  $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ , 0.2

107 Yeast extract, 0.5 and Peptone, 0.5; pH 5.0. For inoculum development, yeast cells were  
108 grown at 42°C for 24 h in 150 mL YPD medium. Yeast cells were recovered after  
109 centrifugation at 5000 rpm for 20 min, washed twice in saline and used as inoculum  
110 (approximately 1 g/L, dry wt.) for bioethanol production.

#### 111 **Biochemical characterization and molecular phylogenetic analysis**

112 The yeast cell morphology was examined by bright field microscopy (Nikon  
113 eclipse Ni, Tokyo, Japan). Biochemical profile of the strain was studied on basis of  
114 carbohydrate utilization characteristics using a HiCarbohydrate™ Kit (HiMedia Lab. Pvt.  
115 Ltd, India) by following manufacturer's instructions. For molecular phylogenetic analysis,  
116 the genomic DNA was extracted and purified from overnight grown culture of strain DBT-  
117 IOC-35 using Rapid Yeast Genomic DNA Extraction Kit (Bio Basic Inc., Ontario,  
118 Canada). The internal transcribed spacer region (ITS) 1 - 5.8S rRNA gene – ITS 2 region  
119 and D1/D2 domain of the 26S rRNA genes were amplified by polymerase chain reaction  
120 (PCR) using the primer pairs ITS1, ITS4 and NL1, NL4 respectively.<sup>13, 14</sup> The PCR  
121 products were purified using HiPurA PCR Product Purification Kit (HiMedia, India) and  
122 then sequenced by Macrogen Inc. (Korea) using an automated Applied Biosystems®  
123 3730/3730xl DNA analyzer. The phylogenetic analyses were performed using the program  
124 CLUSTALW. Phylogenetic tree were constructed using the neighbour-joining method  
125 with the program MEGA4 and boot-strap analysis based on 1000 replicates.<sup>15</sup>

#### 126 **Pilot scale dilute acid pretreatment and composition analysis of biomass**

127 Pretreatment of WS was carried out in 250 kg biomass/day capacity continuous  
128 pilot-scale pretreatment reactor. This multipurpose pilot plant is capable of pre-treatment  
129 operation for multiple feed (wheat straw, rice straw, sugarcane bagasse, cotton stalk, corn  
130 stover, woody biomass, etc.) under a wide range of operating conditions and pre-treatment



131 chemicals (acid and alkali). Wheat straw was milled to 4-5 mm; air dried and then soaked  
132 in the acid solution (2.5%, w/w) for 30 min in a soaking chamber specially equipped with  
133 spray and circulation of acid solution. The wet biomass, after soaking was hung for 2 h and  
134 further pressed for 15 min at a pressure of up to 100 Bar in a hydraulic filter press to  
135 remove water. Then biomass was fed into the reactor at a rate of 10 kg/h and treated at  
136 160°C and 6 bar with a residence time of 15 min. Residence time was controlled by the  
137 screw speed of the reactor. The pre-treated biomass slurry was collected in the slurry tank,  
138 cooled and then transferred through a peristaltic pump to a high speed centrifuge for  
139 separating solids (cellulose and lignin) and liquid (hemicelluloses).<sup>16</sup> Composition analysis  
140 of untreated (UWS) and pretreated wheat straw (PWS) was carried out by following the  
141 procedure of Sluiter *et al.*<sup>17</sup> After thorough washing and removal of extra water by  
142 pressing, moisture content of PWS was determined using MA150 electronic moisture  
143 analyzer (Sartorius weighing technology GmbH, Gottingen, Germany).

#### 144 **Enzymatic saccharification of biomass**

145 Enzymatic saccharification of PWS was carried out in 500 mL capped flasks  
146 containing 200 mL reaction volume at 150 rpm. Influence of various parameters such as  
147 different biomass loadings (5 to 20%, wt.), temperature (30 to 50°C) and enzyme dosage  
148 (10 to 60 filter paper units (FPU)/g biomass) on enzyme hydrolysis was studied. The initial  
149 pH of hydrolysis slurry was maintained at 5.0 using 1 M citrate buffer.

#### 150 **Separate hydrolysis and fermentation (SHF)**

151 For separate hydrolysis and fermentation studies, enzyme hydrolysis was carried  
152 out for 72 h at 50°C using different loadings of PWS (10, 15 and 20%, wt.) as described  
153 earlier. Sterilized YFM and yeast inoculum were added after enzyme hydrolysis and flasks

154 were incubated further at 42°C for 72 h. Samples were withdrawn every 24 h for  
155 estimation of ethanol and sugars.

### 156 **Simultaneous saccharification and fermentation (SSF)**

157 SSF experiments were performed in 500 mL capped Erlenmeyer flasks and the  
158 process was optimized for most favourable conditions of temperature (42 and 45°C),  
159 substrate concentration (10, 15, 20 and 25 % wt.) and enzyme dosage (10, 20, 30, 40, 50,  
160 60 FPU/g biomass). After addition of YFM and yeast inoculum, flasks were incubated at  
161 42°C and 150 rpm for 72 h. Samples were withdrawn at regular intervals for estimation of  
162 bioethanol and sugars. SSF ethanol results were reported as percentage of theoretical yield,  
163 considering the availability of all potential glucose in biomass for fermentation and a  
164 theoretical yield of 0.51 g ethanol/g glucose.<sup>16</sup>

### 165 **Pre-saccharification and simultaneous saccharification and fermentation (PSSF)**

166 The PSSF experiments were carried out in 500 mL flasks and included 6 h  
167 enzymatic pre-saccharification step at 50°C followed by SSF for 72 h. Prior to the start of  
168 the fermentation, temperature was lowered to 42°C and nutrients and yeasts were added.  
169 Three different biomass loadings (10, 15 and 20%, wt) were investigated. Samples were  
170 withdrawn after 6 h pre-saccharification (0 h fermentation) and then every 24 h during  
171 SSF, and were analysed for ethanol and glucose content.

### 172 **Analytical methods**

173 Total reducing sugars during enzyme hydrolysis were estimated by DNS method<sup>18</sup>  
174 by reading absorbance at 540 nm using UV-VIS Spectrophotometer (UV 1800  
175 Spectrophotometer, Shimadzu, Japan). Sugars (glucose, xylose, cellobiose, galactose,  
176 arabinose, and mannose), glycerol and inhibitors in pretreatment slurry were analyzed by

177 high performance liquid chromatography (HPLC, Waters, USA) equipped with a refractive  
178 index detector and Aminex HPX-87H column 300 mm×7.8 mm ID (Bio-Rad Labs,  
179 Hercules, CA) . For ethanol estimation Clarus-680 Gas Chromatograph (Perkin-Elmer,  
180 USA) fitted with a 30 m long Carbowax-PEG column (Perkin-Elmer, USA) having inner  
181 diameter of 0.32 mm was used. The conditions used for HPLC and GC analysis and  
182 cellulase enzyme assays were same as described earlier.<sup>16</sup> All samples were appropriately  
183 diluted and filtered through a 0.22 µm disc-filter prior to analysis. All experiments were  
184 conducted in triplicates and average values are shown.

### 185 **Calculations**

186 Hydrolysis yield (%) =

$$187 \quad [\text{Total Sugars (g/L)} / (0.511 * f * \{\text{Biomass}\}_0 * 1.111)] * 100 \quad (1)$$

$$188 \quad \text{Ethanol titer (g/L)} = \text{EtOH}_t - \text{EtOH}_0 \quad (2)$$

$$189 \quad \text{Ethanol yield (\%)} = [\text{Ethanol titer} / (0.511 * f * \{\text{Biomass}\}_0 * 1.111)] * 100\% \quad (3)$$

$$190 \quad \text{Ethanol productivity (g/L/h)} = \text{Ethanol titer} / t \quad (4)$$

191 Where, (EtOH<sub>t</sub> - EtOH<sub>0</sub>) indicates total ethanol produced during fermentation (g/L)  
192 run obtained by calculating the difference between ethanol produced initially (0 h  
193 fermentation) and after time 't', {Biomass}<sub>0</sub> is the initial dry biomass concentration (g/L), f  
194 is cellulose fraction of dry biomass, 0.511 is the conversion factor for glucose to ethanol  
195 based on stoichiometric biochemistry of yeast and 1.111 is the conversion factor for  
196 cellulose to equivalent glucose.<sup>16</sup>

### 197 **Results and Discussion**

#### 198 **Identification and characterization of strain DBTIOC-35**

199 One of the desired characteristics for development of an efficient bioethanol  
200 fermentation process is the application of yeast strains that can endure high stress of  
201 temperature or ethanol during fermentation process. Thus, isolation and characterization of  
202 thermotolerant and stress tolerant yeasts is critical in improving the bioethanol production  
203 processes via SSF.<sup>10</sup> For this study, thermotolerant yeasts were isolated and qualitatively  
204 screened for growth and fermentation at 42°C and 45°C. Strain DBT-IOC-35 exhibited  
205 best ethanol fermentation characteristics (0.49 g ethanol/g glucose) under shake flask  
206 conditions (data not shown). This isolate had typical yeast-like colonies on YPD agar  
207 plates and reproduced by budding. The colonies were observed as single, or in pair. It  
208 fermented dextrose, lactose, fructose, galactose, and raffinose efficiently whereas  
209 utilization of xylose, arabinose, mannose, glycerol,  $\alpha$ -methyl-d-glucoside, xylitol showed  
210 variable pattern during carbohydrate utilization test. This strain could not utilize citrate and  
211 malonate as sole carbon sources, but hydrolysed esculin and metabolized ONPG, thus  
212 indicating  $\beta$ -galactosidase activity. Biochemical profile of the strain matched closely to the  
213 reference strain *K. marxianus* MTCC-4136. Molecular phylogenetic analysis of ITS-5.8S  
214 rRNA gene and D1/D2 domain of the 26S rRNA gene of DBTIOC-35 revealed maximum  
215 similarities to *K. marxianus* Kw1696 (HE650694) (Fig. S1) and *K. marxianus* DX3-3  
216 (GU565206) (Fig. S2). Therefore, isolate DBTIOC-35 was finally identified as *K.*  
217 *marxianus*. ITS-5.8S and 26S rRNA gene sequences were submitted to NCBI GenBank  
218 under accession numbers KP192932 and KP192933, respectively. Many studies have  
219 reported the abilities of *K. marxianus* strains to grow and ferment rapidly at temperatures  $\geq$   
220 40 °C<sup>10</sup> but the reported ethanol yields or concentrations are mostly low. Therefore, to  
221 explore full potential of this strain in bioethanol applications, it was evaluated in SHF and  
222 SSF.

### 223 **Biomass pre-treatment and composition analysis**

224 Composition analysis (cellulose, hemicelluloses and lignin) of extractive free UWS  
225 and PWS indicated substantial removal of hemicelluloses (from 26.3% in UWS to 3.8% in  
226 PWS) and relative increase of 3.9% in the lignin content. Cellulose content also increased  
227 from 51.2% to 69.8% after pre-treatment. These findings are in accordance with that of  
228 Chen *et al.*<sup>19</sup> The dilute sulfuric acid pretreatment at high temperature hydrolyzes  
229 hemicellulose into monomeric sugars (xylose, arabinose, galactose, glucose, and mannose)  
230 and oligomers. Small amount of lignin is also depolymerized during acid pretreatment and  
231 might re-condense to form an altered lignin polymer. Removal of hemicellulose increases  
232 surface area and pore volume of the substrate which enhances the yield and rate of  
233 enzymatic hydrolysis of the cellulose rich biomass.<sup>20</sup> After pretreatment, most of the  
234 soluble monomeric and/or oligomeric sugars and inhibitors like furfural, hydroxyl-methyl  
235 furfurals and acetic acid remained in the liquid supernatant obtained after centrifugation. In  
236 most of the studies on bioethanol production, dilute acid pretreatment of biomass is  
237 generally carried out at a lab scale (mostly autoclaving the biomass containing shake flasks  
238 or bottles) with lower volume of pretreated biomass. In this study, dilute acid based  
239 pretreatment was carried out in a continuous pilot scale pretreatment reactor and this  
240 pretreated wheat straw was further used for enzymatic saccharification and bioethanol  
241 production studies.

#### 242 **Effect of enzyme dosage, biomass loading and temperature on enzymatic hydrolysis** 243 **of PWS**

244 Fig. 1a and 1b show the effect of enzyme dosage on hydrolysis of PWS at 50 °C  
245 and 42 °C, respectively. Pretreated WS was made free of sugars and inhibitors (phenolics,  
246 acetic acid, furfural etc.) due to their known detrimental effects on cellulase enzyme.<sup>21</sup>  
247 Hydrolysis yield increased with increase in the enzyme dosage at both temperatures. The  
248 yields at 42°C were lower than that at 50°C. Hydrolysis yield of 85% was attained by

249 using enzyme dosage of 45 FPU/g at 42°C in comparison to 30 FPU/g at 50°C. Therefore,  
250 45 FPU/g PWS was chosen for further investigation. Karthika *et al.*<sup>22</sup> previously reported  
251 an enzyme dose of 30 FPU/g to obtain 80% hydrolysis within 48 h. PWS hydrolysis at  
252 different biomass loadings at two different temperatures is depicted in Fig 1c and 1d. At  
253 both temperatures, little or no difference in hydrolysis yield was observed at 5 and 10%  
254 biomass loading (dry wt). But there was reduction in hydrolysis yield at higher solid  
255 loadings (15% to 20%) which was in accordance with Oloffsson *et al.*<sup>5</sup> This reduction in  
256 hydrolysis yield may be either due to increased viscosity resulting in mass transfer  
257 limitations or mixing difficulties or due to feedback inhibition on enzyme. However, use of  
258 high solids can significantly improve ethanol concentration in fermentation broth and is  
259 desirable for reducing ethanol recovery cost.<sup>23</sup> Influence of temperature on hydrolysis of  
260 PWS at 10% solid loading and enzyme dose of 45 FPU/g is shown in Fig. 2. As much as  
261 83% and 90% hydrolysis yields were obtained at 24 h at 42°C and 45°C, respectively.  
262 These conditions were chosen for SSF experiments. The enzymatic hydrolysis at higher  
263 temperatures would potentially reduce the reaction time but cellulase system is rapidly  
264 inactivated above 45°C. Therefore, to increase saccharification and subsequent ethanol  
265 fermentation, SSF can be adopted since the sugars released by enzymatic hydrolysis are  
266 immediately consumed by the yeast and therefore feedback inhibition on enzyme is  
267 potentially avoided. These points were taken into consideration during SSF experiments in  
268 this study.

### 269 **Bioethanol production employing thermotolerant yeast DBTIOC-35**

270 Both SHF and SSF are established processes for ethanol production from  
271 lignocellulosic biomass. Major advantage of SHF is that both hydrolysis and fermentation  
272 steps can be run separately under their optimal conditions at 50°C and 30°C, respectively.  
273 SSF on the other hand is usually carried out at 35°C at pH 5. Based on previous reports on

274 SSF, biomass loading of 10% wt., yeast inoculums concentration of 1 g/L in SSF and  
275 YFM as fermentation medium were used during bioethanol fermentation in this study.<sup>5, 24-  
276 27</sup>

### 277 **Separate hydrolysis and fermentation**

278 For SHF experiments PWS was hydrolysed by the cellulase enzyme at three  
279 different loadings 10, 15, 20 % (wt.) for 72 h and the hydrolysate was subjected to  
280 fermentation after supplementation with YFM nutrients and yeast cells. Initial sugar  
281 concentrations, cellulose to glucose conversion after hydrolysis, ethanol yield (g ethanol/g  
282 glucose) and % yield (of theoretical maximum) obtained during SHF are presented in  
283 Table 1. Glucose in the SHF flasks (taking into account the dilution due to addition of  
284 medium and inoculum) ranged from 65.5 to 99.9 g/L, which corresponded to respective  
285 biomass concentrations. Increase in substrate concentration resulted in decreased sugar  
286 conversion and glucose yield. The ethanol yields (g/g) obtained during SHF were close to  
287 those of control fermentation experiments (0.49 g ethanol/g glucose) using synthetic media  
288 (data not shown), which indicated overall good fermentation performance of strain  
289 DBTIOC-35. In SHF10 flasks almost complete utilization of glucose along with ethanol  
290 concentration of 26.8 g/L was observed after 24 h. Residual glucose in SHF15 and SHF20  
291 flasks indicated incomplete sugar utilization and decreased the ethanol yield. Ethanol  
292 productivity (initial 24 h) in all flasks was proportional to biomass concentration. In  
293 SHF20 flasks ethanol concentration increased, however, some of the glucose released  
294 during saccharification remained unutilized even up to 72 h indicating the decrease in cell  
295 viability. During SHF of lignocellulosic biomass, amount of ethanol produced in  
296 fermentation step mainly depends upon sugar concentration obtained after saccharification  
297 step. Therefore, to achieve higher ethanol concentrations, and reduce fermentation vessel

298 size and wastewater streams higher initial substrate concentrations are preferred during  
299 SHF.<sup>7</sup>

### 300 **Simultaneous saccharification and fermentation (SSF)**

301 SSF experiments were aimed at increasing the final ethanol concentration. The  
302 frequently reported main advantages associated with this process are prevention of end-  
303 product inhibition during enzymatic hydrolysis resulting in comparatively higher  
304 saccharification and ethanol production, use of a single vessel for saccharification and  
305 fermentation step, elimination of separation of residual biomass prior to fermentation.<sup>26</sup>

### 306 *Effect of temperature on SSF*

307 Fig. 3a and 3b depict ethanol concentrations and yields (% of theoretical ethanol)  
308 estimated at 16, 24, 48, 72, 96 and 120 h during SSF at 42°C and 45°C, respectively.  
309 Tolerance to high temperatures besides high ethanol and sugar tolerance is an important  
310 aspect for selecting robust bioethanol producing microorganisms. Therefore, bioethanol  
311 production via SSF was studied at higher temperatures (> 40°C) at 10% initial biomass  
312 loading. Maximum ethanol concentration of 29.0 and 16.1 g/L were achieved,  
313 corresponding to ethanol yields (percentage of theoretical) of 73% and 40.5% at 42 and  
314 45°C, respectively. SSF at 45°C resulted in lower ethanol production than at 42°C.  
315 However, the observed ethanol concentration was much higher than 6.18 g/L from rice  
316 cellulignin<sup>11</sup> and 10 g/L from wheat straw.<sup>25</sup> At this temperature, accumulation of sugars  
317 (glucose as well as some xylose) was noticeable and was mainly due to inhibitory or lethal  
318 effect of this temperature on yeast growth and metabolism. Lin *et al.*,<sup>28</sup> have reported 80%  
319 decrease in viability of *K. marxianus* as a reason for decreased ethanol fermentation at  
320 45°C. At both the temperatures used in this study, maximum ethanol production was  
321 observed between 72-96 h, 42°C being the more favourable temperature in terms of



322 bioethanol titer, yield as well as productivity and was used further. Though there are a few  
323 reports on bioethanol fermentation at temperatures as high as 45°C, but the obtained  
324 ethanol yields are inferior to that obtained at lower temperatures (42°C or 37°C). Due to  
325 decrease in cell viability, ethanol production decreases considerably at higher  
326 temperatures.<sup>10</sup>

### 327 *Effect of biomass loading on SSF*

328 To further increase ethanol concentration in the fermentation broth, SSF process  
329 was carried out at high substrate loading. Fig. 3c, 3d and 3e show ethanol production  
330 during SSF at 42°C using different biomass loadings. After 72 h, highest ethanol yields at  
331 20 and 25% biomass loadings were 83.3% and 67.9%, respectively. Substrate  
332 concentration of 25% hampered the ethanol yield. With increase in solid content, increased  
333 viscosity creates difficulty in stirring and limits the enzymatic hydrolysis of biomass  
334 possibly due to inhibition of binding of cellulase to substrate under such conditions.<sup>7, 23</sup>  
335 However, in present study, ethanol yield increased with increase in biomass loading from  
336 10 to 20%, wt. in contrast to reports indicating decrease in ethanol yield above 10-15%.<sup>5</sup>  
337 One probable reason might be release of sugars in controlled manner (as no residual  
338 glucose was observed in the SSF medium) and immediate utilization and fermentation to  
339 ethanol by yeast. Detailed studies are needed to determine exact reasons for improvements  
340 in ethanol yields with increasing biomass loading up to 20%. Ethanol concentrations of  
341 66.2 and 67.4 g/L were the maximum observed at 20 and 25% biomass loadings,  
342 respectively. Higher ethanol concentration obtained during SSF experiments might have  
343 been lethal under the process conditions (e.g. higher temperature) and possibly inhibited  
344 further ethanol production. This was supported by the accumulation of residual glucose  
345 when biomass loading of 25% was used. Moreover, very high dry matter content in the  
346 SSF process increased the viscosity, at least initially, and might inhibited even mixing of

347 slurry. Though fed-batch as well as pre-saccharification strategies can alleviate some of the  
348 problems associated with high DM concentrations, use of good bioethanol fermenting  
349 thermotolerant yeast strains such as *K. marxianus* DBTIOC-35 could provide a better  
350 alternate solution.

351 ***Effect of Pre-saccharification and simultaneous saccharification and fermentation***  
352 ***(PSSF)***

353 Ethanol production rate in SSF is governed more by cellulose hydrolysis rather  
354 than glucose fermentation.<sup>23</sup> But when the biomass concentration is increased in SSF,  
355 mixing of the solids is decreased. Therefore, one approach repeatedly employed is to carry  
356 out prehydrolysis or pre-saccharification of the biomass at optimum temperature of  
357 enzyme for some time prior to yeast inoculation. This makes the SSF slurry more fluid and  
358 easy to handle. Time course of ethanol production during PSSF at 42°C with different  
359 biomass loading has been shown in Fig. 4a-4c. Pre-saccharification step of 6 h at 50°C at  
360 different WS concentrations was evaluated during this study. The conditions for PSSF  
361 were chosen based on the obtained hydrolysis efficiency of more than 54% during  
362 enzymatic hydrolysis experiments (Fig. 1). The main aim of PSSF was to partially  
363 hydrolyse the cellulose to sugars prior to yeast addition so that ethanol production during  
364 initial phase of PSSF could be increased. The positive effect of PSSF on ethanol  
365 production during initial 24 h was evident in experiments at 15 and 20 % biomass loading.  
366 However, after 48 h both SSF and PSSH resulted in almost similar ethanol concentration  
367 and yields. After 72 h ethanol production was lower than SSF performed without any pre-  
368 saccharification step. Thus, the pre-saccharification step in comparison to SSF did not  
369 result in enhanced ethanol production by *K. marxianus* DBTIOC-35 in this study. Similar  
370 findings have been reported earlier for olive pruning biomass, corn stover and barley straw  
371 due to more enzyme deactivation at high temperature pre-saccharification step.<sup>1, 23</sup> This

372 might have decreased the ethanol yield by lowering the enzyme hydrolysis in the later  
373 stages of PSSF. Lower ethanol yields can also be a consequence of decreased water  
374 activity in the fermentation broth, especially at higher biomass loading or due to sudden  
375 osmotic changes in extracellular environment of the yeast which leads to osmostress  
376 response as a defense mechanism by making intracellular physiological changes such as  
377 glycerol formation.<sup>29</sup> Glycerol synthesis is an alternate means for NAD<sup>+</sup> replenishment  
378 during SSF process. Glycerol concentration in SSF broths were in the range of 2.1 to 3.56  
379 g/L (not shown) and are comparable to 2.62 g/L reported previously during SSF of  
380 sunflower meal.<sup>30</sup> However, low extracellular glycerol may not rule out the formation of  
381 intracellular glycerol.

### 382 *Effect of enzyme dosage on SSF*

383 Total rate of ethanol production during SSF process is largely controlled by the  
384 enzymatic hydrolysis of the solid biomass. Previous studies on SSF at different enzyme  
385 loadings have exhibited strong positive correlation between enzyme loading and the  
386 overall ethanol yield.<sup>5</sup> However, SSF process can be made more cost-effective by  
387 decreasing the amount of cellulase needed to a level that doesn't compromise the ethanol  
388 production significantly. Therefore, SSF experiments were performed using different  
389 cellulase dosages at 20% biomass concentration (Fig. 5). At low enzyme loadings of 20  
390 FPU/g, the overall ethanol productivity was not much affected and only a slight decrease  
391 from 0.87 g/L/h (at 40 FPU/g) to 0.82 g/L/h was observed. Similar previous findings have  
392 been attributed to better conversion of cellulose to glucose at higher cellulase loadings.<sup>5</sup>  
393 Application of higher enzyme dosages may seem uneconomical as such, but recycling of  
394 cellulase enzymes can be a potential strategy that can cut down the cost of enzymes used  
395 for biomass bioconversion. This will require more efforts in effective separation of  
396 enzymes dissolved in enzymatic hydrolysate and/or bound to left over solid residuals

397 mainly lignin, especially when using acid pretreated biomass.<sup>31</sup> Further decrease in  
398 enzyme loading to 10 FPU/g decreased ethanol productivity to 0.67 g/L/h which was much  
399 lower than at higher enzyme dosages. From process economy point of view, application of  
400 lower enzyme dose will definitely be more important in future in terms of cost savings and  
401 can make up the loss due to decreased productivity to some extent. However, choice of  
402 optimal enzyme dose will depend upon balanced trade-off between both the process  
403 economy and productivity.

#### 404 **Comparison of SHF, SSF and PSSF**

405 Table 2 compares the ethanol production by thermotolerant yeast *K. marxianus*  
406 DBTIOC-35 using different process configurations SHF, SSF and PSSF at different  
407 loadings of PWS. In SHF with 10% biomass loading, faster initial (24 h) rate of ethanol  
408 production was observed than SSF or PSSF due to more availability of glucose initially. At  
409 solids loading above 10%, glucose availability increased in SSF but enzyme inhibition  
410 possibly became more dominant in SHF due to increased sugar release. As a result, initial  
411 rates of ethanol production for SHF and SSF differed at 20% biomass loading. The  
412 removal of sugar inhibition on enzyme in SSF enabled more sugar to be continuously  
413 released which in turn continuously increased ethanol production and resulted in higher  
414 ethanol titers. These findings are very well supported by the results of previous study.<sup>32</sup>  
415 Highest productivity were obtained during PSSF of PWS at biomass loadings  $\geq 15\%$  (wt.),  
416 however, these were only marginally better than that obtained during SSF without 6 h long  
417 pre-saccharification step. The overall productivities, as well as titer and yields during SSF  
418 process at different biomass loadings were better than other process configurations. During  
419 SSF and PSSF cellobiose and xylose concentration never exceeded 0.4 and 4.0 g/L,  
420 respectively and galactose, mannose and arabinose were not detectable.

421 Ethanol production by *K. marxianus* DBTIOC-35 during SSF has been compared  
422 with previous studies in Table 3. As can be seen, maximum overall ethanol productivity of  
423 0.92 g/L/h obtained in this study at 20 % biomass loading was better than 0.39 g/L/h  
424 obtained by both Mohagheghi *et al.*<sup>21</sup> and Bollók *et al.*,<sup>33</sup> and 0.885 g/L/h reported by  
425 Jain *et al.*<sup>34</sup> Though there are some previous reports on SSF without pre-saccharification step,  
426 simple SSF process i.e. without any presaccharification or any other modifications (such as  
427 delayed temperature SSF or non-isothermal SSF) have not resulted in such high ethanol titres of  
428 66.2 g/L (ethanol yield of 83.3%) as have been obtained in this study under high gravity SSF of  
429 pilot scale dilute acid pretreated wheat straw. Zhang *et al.*,<sup>8</sup> reported higher ethanol titer (84.7 g/L)  
430 than this study; however the process took longer (96 h) with comparatively lower ethanol yields  
431 and the biomass was added in fed batch process to increase biomass loading. Moreover, their  
432 process involved serial acid and alkali pretreatment which removes both hemicellulose and lignin  
433 but at the same time increases cost and environmental concerns.

434 Success of any SSF process depends upon balanced trade-off between optimal  
435 temperatures of enzymes and yeast which can be achieved by carrying out SSF process at  
436 temperatures above 40°C. Besides high fermentation rates, high temperature SSF may  
437 reduce the possible contamination by mesophilic microorganisms and enhance the  
438 hydrolytic enzyme activities.<sup>35</sup> Therefore, thermotolerant yeasts are highly desirable for  
439 cost-effective commercial production of second generation bioethanol as these can  
440 withstand higher temperatures encountered due to heat generation during large-scale  
441 industrial fermentations. The SSF process without pre-saccharification (as used in this study) has  
442 certain advantages such as, saving at least 4 h to 24 h of the overall process time, the energy used  
443 in heating and cooling during and after pre-saccharification, respectively, increased stability of the  
444 used enzyme during the process (in comparison to when higher temperature is used) etc. Ethanol  
445 concentrations obtained in the present study can be improved further by employing fed-  
446 batch SSF or temperature shift SSF process.

## 447 **Conclusions**

448 Thermotolerant yeast *K. marxianus* DBTIOC-35 capable of producing high titer  
449 ethanol above 40°C with a maximum fermentation temperature of 45°C has been  
450 successfully isolated. SSF of dilute acid pretreated WS at higher solids loadings at 42°C  
451 using an efficient thermotolerant yeast can improve bioethanol titer and yield even without  
452 the pre-saccharification step and the overall biomass to ethanol conversion process time  
453 can be reduced. As far as our knowledge is concerned, ethanol titers and yields reported in  
454 this study are the highest when using dilute acid pretreated biomass in unmodified SSF  
455 process. Further studies on scale-up are underway and will certainly prove fruitful for  
456 development of a more sustainable and greener process for biomass to ethanol  
457 bioconversion.

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522

523 **Legends to Figures**

524 **Fig. 1. Enzymatic hydrolysis of PWS at different enzyme dose and 10% biomass**  
525 **loading at (a) 50°C and (b) 42°C; and at different biomass loadings and 45 FPU/g at**  
526 **(c) 50°C and (d) 42°C.**

527 **Fig. 2. Influence of temperature on biomass hydrolysis at 10% solid loading and**  
528 **enzyme dosage of 45 FPU/g.**

529 **Fig. 3. Ethanol production during SSF of WS at different temperatures and biomass**  
530 **loadings. SSF at (a) 42°C using 10% biomass loading; (b) SSF at 45°C using 10%**  
531 **biomass loading; SSF at 42°C using (c) 15% biomass loading; (d) 20% biomass**  
532 **loading; and (e) 25% biomass loading.**

533 **Fig. 4. Time course of ethanol production during PSSF at 42°C using (a) 10%**  
534 **biomass loading, (b) 15% biomass loading, and (c) 20% biomass loading.**

535 **Fig. 5. Ethanol production during SSF at different enzyme dosages. SSF was carried**  
536 **out at 42°C for 72 h using 20% biomass loading.**

537

538

539 **Table 1. Initial glucose concentration, glucose conversion and ethanol yield during**  
 540 **SHF at different loadings of PWS\*.**

Experiment ID	Substrate Concentration (wt.)	Initial glucose, g/L	Glucose Conversion	Time	Yield, g/g	Yield, % of theoretical (On basis of sugar utilized)
<b>SHF10</b>	10%	62.5	96.9%	24 h	0.47	91.5
				48 h	0.43	84.8
				72 h	0.43	83.5
<b>SHF15</b>	15%	87.4	90.0%	24 h	0.48	94.2
				48 h	0.43	84.3
				72 h	0.41	80.7
<b>SHF20</b>	20%	99.9	82.2%	24 h	0.49	95.4
				48 h	0.42	79.2
				72 h	0.40	78.0

541

542 \*Enzymatic hydrolysis was done using enzyme dose of 45 FPU/g at 50°C and different  
 543 biomass loadings for 72 h and glucose conversion (after HPLC analysis) was calculated  
 544 from cellulose content of PWS on dry matter basis

545

546 **Table 2. Comparison of ethanol concentrations and yields (% of theoretical) obtained**  
 547 **during SHF, SSF and PSSF at different loadings of PWS.**

Biomass Loadings (% wt.)	Final Ethanol concentration (g/L)			Yield (%)			Productivity (g/L/h)		
	SHF	SSF	PSSF	SHF	SSF	PSSF	SHF	SSF	PSSF
10%	26.3	29.0	27.6	67.8	73.0	66.5	1.11 (0.36)	1.06 (0.4)	1.06 (0.38)
15%	32.3	45.9	44.2	51.6	77.0	72.5	1.28 (0.45)	1.61 (0.64)	1.69 (0.61)
20%	35.2	66.2	61.8	39.8	83.3	77.7	1.31 (0.49)	2.17 (0.92)	2.27 (0.86)

548 Initial and overall ethanol productivities were calculated respectively, for 24 h and 72 h  
 549 (Values in parenthesis) fermentation period i.e after addition of yeast cells and does not  
 550 take into account the hydrolysis time (as in SHF or PSSF). SHF (72 h hydrolysis and 72 h  
 551 fermentation), SSF (72 h) and PSSF (6 h hydrolysis and 72 h fermentation) were carried  
 552 out at 42°C.

553 **Table 3. Comparison of bioethanol production via SSF process by *K. marxianus* DBTIOC-35 with previous studies.**

Microorganism	Substrate, Pre-treatment and solids loading (% , wt.)*	Enzyme Dosage	Fermentation Temperature and Time	Ethanol titer, g/L; yield, % (or g/100g); productivity (g/L/h)	Ref.
<i>S. cerevisiae</i> DQ1	A, CS, 30%	10-15 FPU/g DM Commercial cellulase	40°C	48 g/L; 65.6 %	<b>Chu et al.,<sup>9</sup></b>
<i>S. cerevisiae</i>	A, WS, 20%	0.24 g Cellulase 150L/g cellulose	37 °C, 144 h	57 g/L; 80%; 0.39 g/L/h	<b>Mohagheghi et al.,<sup>21</sup></b>
<i>K. marxianus</i> Y.00243	SE spruce, 5%	37 FPU/g+38 BGLU/g of commercial enzymes	42°C, 23 h	9.1 g/L; 14 g/100 g DM; 0.39g/L/h	<b>Bollók et al.,<sup>33</sup></b>
<i>K. marxianus</i> CECT 10875	SE WS, 10% WIS	15 FPU/g Celluclast 1.5 L + 12.6 IU/g Novozyme 188	42°C, 72 h	70%; 12 g ethanol/100 DM	<b>Ballesteros et al.,<sup>25</sup></b>
Dry yeast ( <i>S. cerevisiae</i> )	A+B, CC, 19%	30 FPU/g glucan Commercial cellulase	37°C, 96 h	62.7 g/L; 81.2; 0.65 g/L/h	<b>Zhang et al.,<sup>8</sup></b>
Dry yeast ( <i>S. cerevisiae</i> )	A+B treated CC, (19% initial + 6% after 4 h)	30 FPU/g glucan Commercial cellulase	37°C, 96 h	84.7 g/L; 79.6%; 0.88 g/L/h	<b>Zhang et al.,<sup>8</sup></b>
<i>K. marxianus</i> DBTIOC-35	A, WS, 20%	SaccariSeb C6 45 FPU/g	42°C, 72 h	66.2 g/L; 83.3%; (33.1g/100 g DM); 0.92 g/L/h	<b>This study</b>
<i>K. marxianus</i> DBTIOC-35	A, WS, 25%	SaccariSeb C6 45 FPU/g	42°C, 72 h	67.4 g/L; 67.9% (33.7g/100 DM); 0.94 g/L/h	<b>This study</b>
<i>K. marxianus</i> NRRL Y-6860	A, RS cellulignin	25 FPU/g Cellubrix and 25 IU Novozyme 188	45°C, 12 h	6.18 g/L; 47%; 24 g/100g	<b>Castro &amp; Roberto,<sup>11</sup></b>
<i>Kluyveromyces</i> sp. IIPE453	A, RS , 10%	1% v/w SacchariSeb C6	45°C,	17.7 g/L; 0.885 g/L/h	<b>Jain et al.,<sup>35</sup></b>
<i>K. marxianus</i> IMB3	B, WS, 6%	2% (v/v) commercial cellulase	45°C, 60 h	3.6 g/L; 48%; 20g/100g WS	<b>Boyle et al.,<sup>36</sup></b>
<i>K. marxianus</i> DBTIOC-35	A, WS, 10%	SacchariSeb C6 45 FPU/g	45°C, 72 h	16.1 g/L; 31.5%; 0.52g/L/h	<b>This study</b>

554

555 \*CS = corn stover; WS= wheat straw; RS= Rice straw; WIS= water insoluble solids; A= acid pre-treatment; B= Alkali pre-treatment; A+B=Acid and alkali

556 combined pre-treatment; SE=steam explosion preachment

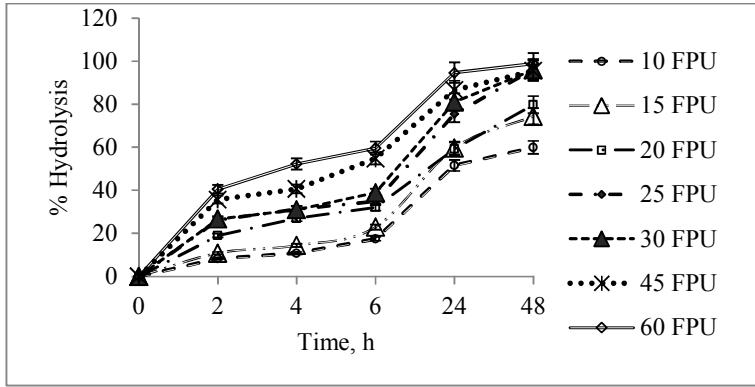


Fig. 1a.

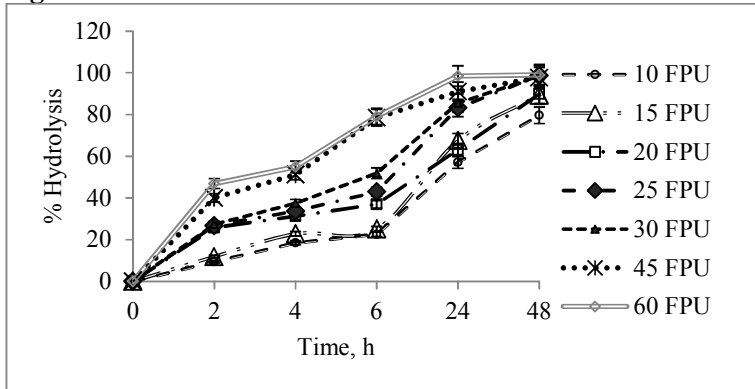


Fig. 1b.

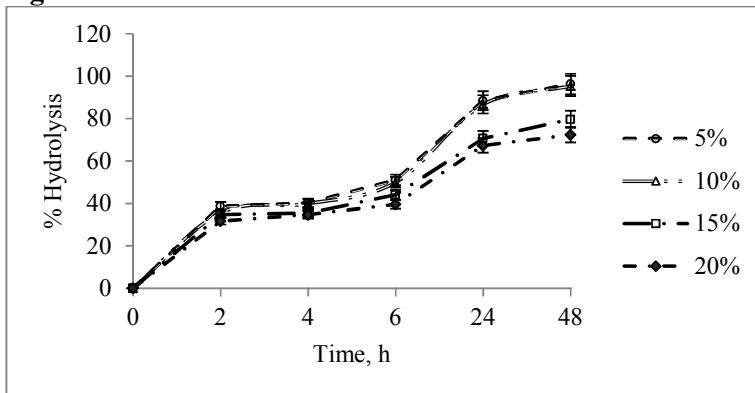


Fig. 1c.

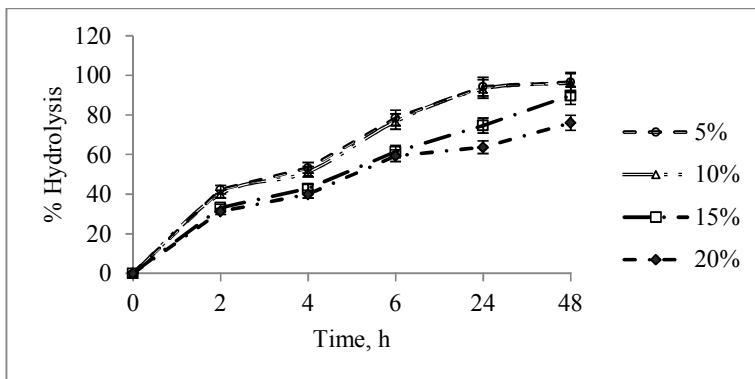


Fig. 1d.

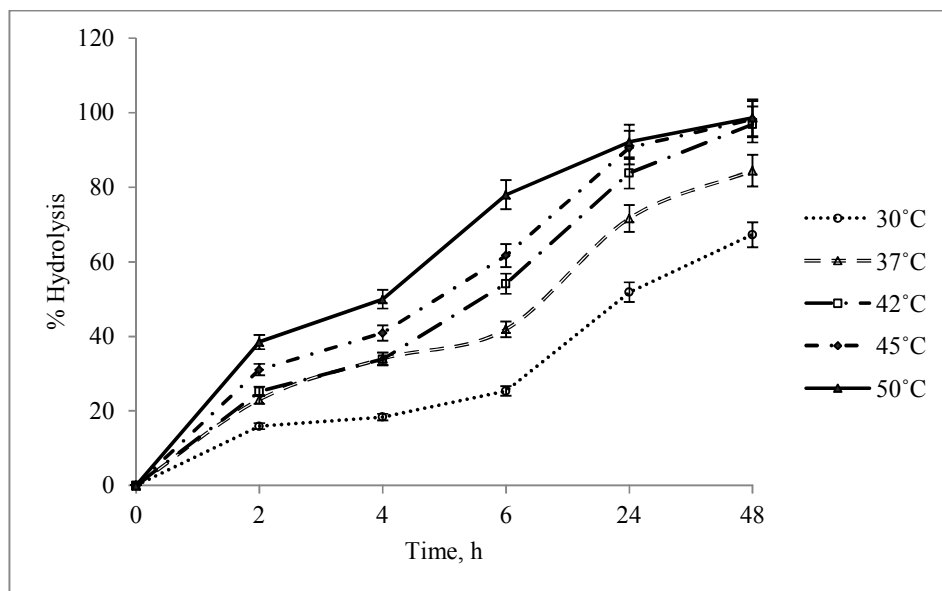


Fig. 2.

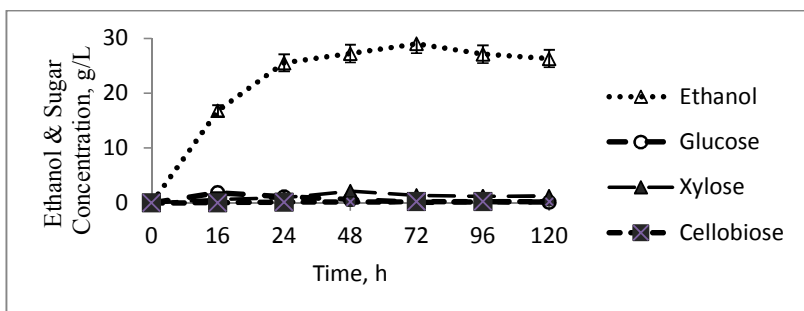


Fig. 3a.

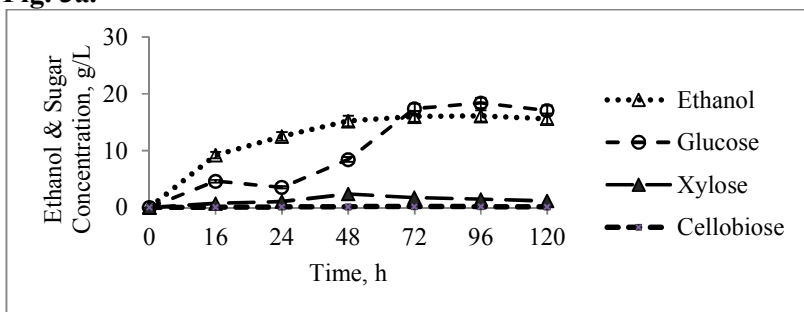


Fig. 3b.

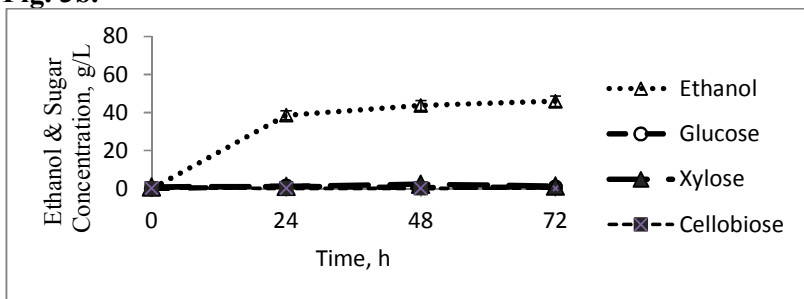


Fig. 3c.

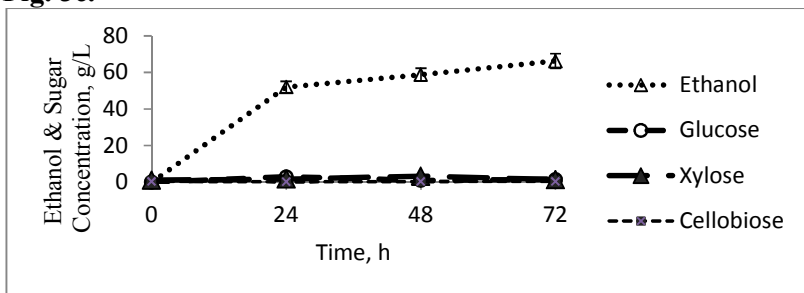


Fig. 3d.

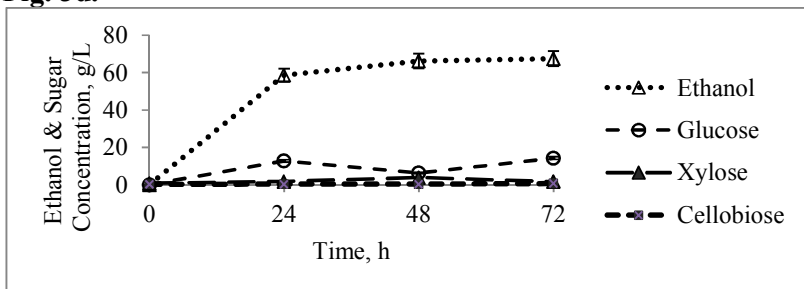


Fig. 3e.



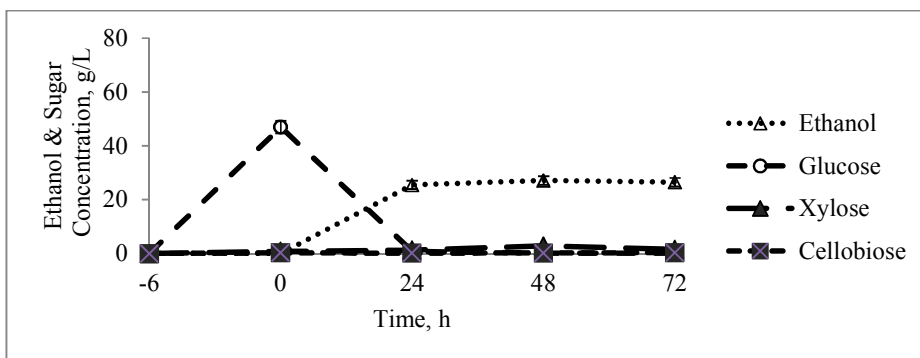


Fig. 4a.

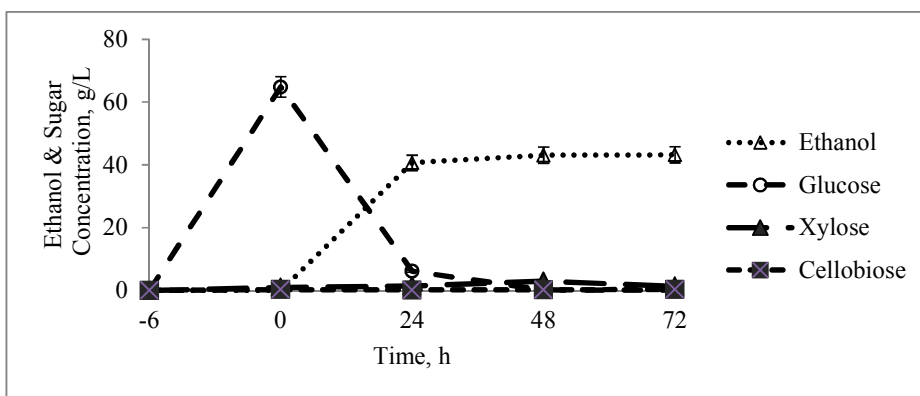


Fig. 4b.

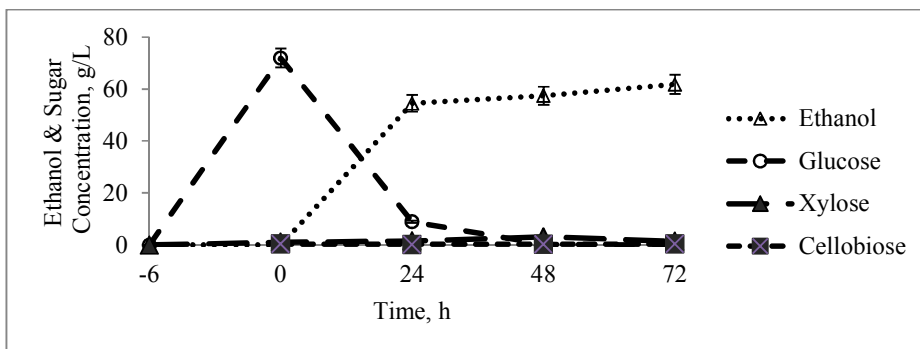
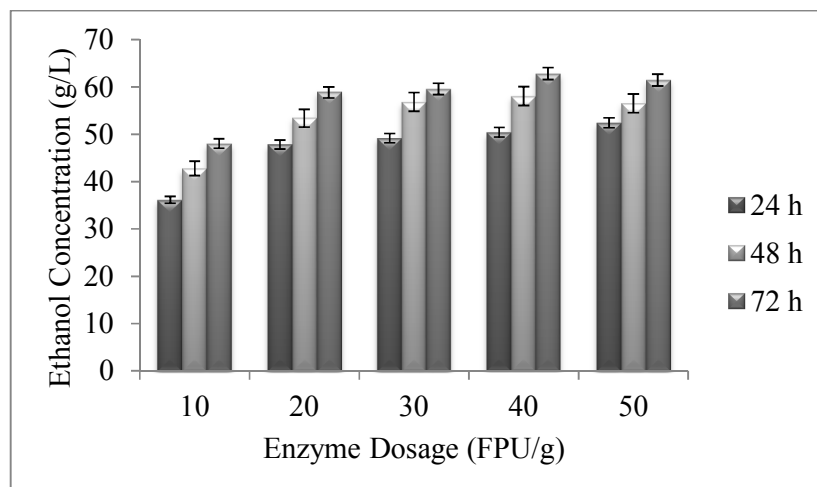


Fig. 4c.



**Fig. 5.**