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1 Co-production of ethanol-hydrogen by genetically engineered *Escherichia coli* in  
2 sustainable biorefineries for lignocellulosic ethanol production

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23

24 **Abstract**

25 This work shows the impact of the hydrogen and ethanol co-production –via dark  
26 fermentation– using a genetically modified *Escherichia coli* in the environmental and  
27 economic sustainability of a lignocellulose-based biorefinery. Wheat straw (WS) and corn  
28 stover (CS) were used as feedstock pretreated with either a dilute acid pretreatment (DAP)  
29 or an autohydrolysis followed by very-dilute acid pretreatment (AH-VDAP) to compare the  
30 effect of the lignocellulosic matrix and the pretreatment as strategies for obtaining rich  
31 hemicellulosic hydrolysates, which were used as substrate in the dark fermentation  
32 experiments. Further, their impact on the profitability was determined on biorefinery  
33 conceptual designs incorporating the experimental results of the pretreatment and dark  
34 fermentation stages. The dark fermentation stage contributed with 20% to 30% of the total  
35 ethanol production in the lignocellulose-based biorefinery designs proposed in this work.  
36 Techno economic and sustainability analyses established that the biorefinery design using  
37 WS as feedstock and employing AH-VDAP presented the lowest negative environmental  
38 impact with the lowest Total Production Cost. The results show that co-production schemes  
39 could be an alternative for lignocellulosic ethanol biorefineries.

40

41 **Keywords:** lignocellulosic biomass; biorefineries; dark fermentation; metabolic  
42 engineering; techno economic analysis; sustainability analysis

## 43 **1 Introduction**

44 One of the most urgent and important challenges of this century is averting global warming  
45 whilst satisfying the growing energy demands of humankind. Renewable energies (*e.g.*  
46 solar, wind and biofuels) seems to be the most promising alternatives to address this  
47 challenge [1]. Therefore, the design, development, and optimisation of sustainable  
48 biorefineries for the efficient production of biofuels are needed to duly provide society with  
49 this renewable energy source [2,3]. A biorefinery is a facility in which biomass is converted  
50 into marketable products and energy using multistep processing approaches [4].  
51 Lignocellulose is a sustainable and world-wide available biomass, thus a suitable feedstock  
52 for biorefining purposes [5,6]. Wheat straw (WS) and corn stover (CS) are lignocellulosic  
53 biomasses (LCB) with potential as feed-stocks for producing bioenergy and high value  
54 added products [7] since they are the most abundant agricultural residues worldwide. The  
55 annual global production of these residuals is around 0.5 and 1 billion tons, respectively  
56 [8,9]. Their glucan and xylan pools represents a significant potential source of glucose and  
57 xylose [10,11]. These carbohydrates, after being pretreated and hydrolysed [12], produce  
58 biofuels (*e.g.*, alcohols or hydrogen) by different fermentation strategies. Among them,  
59 dark fermentation has been extensively studied, concluding that high yields and  
60 productivities as well as low production costs are required to achieve profitable industrial-  
61 scale production [13–15]. Genetically modified microorganisms with redesigned metabolic  
62 systems have been hailed as a possible solution since they can improve yield and  
63 productivity, thus reducing CAPEX and OPEX [16–18]. In particular, *Escherichia coli* is  
64 the most convenient onset to engineering microbial catalysts for biofuel production owing  
65 to extensive knowledge of its genetic and metabolism [19,20]. Among those biofuels  
66 produced using microbial fermentation, hydrogen has gained interest because its eco-

67 friendly nature and energy content (120 kJ/g), as well as ethanol due to mature production  
68 technology and its well established existing fuel market [16,21]. *E. coli* is capable to co-  
69 produce hydrogen and ethanol through dark fermentation from pentoses and hexoses  
70 (analytical grade), as well as from hemicellulose hydrolysates? [22,23]. This is because  
71 oxidative decarboxylation of pyruvate to produce acetyl-CoA and formate. Therefore, the  
72 co-production of hydrogen and ethanol can be more profitable than their production in  
73 separate fermentation stages [24]. Moreover, co-production schemes can improve the  
74 energy balance of the biorefinery designs [25,26].

75 This work studies the impact of hydrogen and ethanol co-production from hemicellulose –  
76 via dark fermentation– using a genetically modified *E. coli* strain in the sustainability of a  
77 biorefinery producing ethanol and hydrogen using lignocellulosic biomass. Two types of  
78 lignocellulosic biomass, which were subject to two different pretreatment methods to obtain  
79 hemicellulose-rich hydrolysates, and then used in the dark fermentation as substrate.  
80 Results were used for designing of biorefineries employing the dark-fermentation stage for  
81 improving the biorefinery energy balance. This stage provided part of the ethanol produced  
82 in the biorefinery, as well as the hydrogen used together with the biogas and solid residues  
83 from the wastewater treatment stage for cogenerating energy. The dark fermentation  
84 batches were carried out with hemicellulosic hydrolysates from WS and CS obtained from  
85 dilute acid pretreatment (DAP) or autohydrolysis followed by very-dilute acid pretreatment  
86 (AH-VDAP). These pretreatment methods were selected based on their glucose and  
87 hemicellulose high yields obtained in post-pretreatment stages from previous experiences  
88 on pilot-scale continuous mode pretreatment strategies [10,11,27] and their high yield in  
89 hydrogen production [28,29]. Techno economic and sustainability analyses of the resulting  
90 biorefinery designs was carried out considering the environmental and economic domains

91 [30] to identified the role of the co-production of hydrogen and ethanol strategy on the  
92 proposed biorefinery designs.

93

## 94 **2 Experimental setup and procedures**

### 95 **2.1 Feedstocks**

96 Wheat straw (WS) and corn stover (CS) were harvested in the spring of 2017 in La Barca  
97 (Jalisco, Mexico). Both feedstocks were milled with a hammer mill (Azteca 301012) using  
98 a 1.27 cm screen. LCB composition was determined according to NREL analytical  
99 procedures [31]. Cellulose, hemicellulose, and lignin content in LCB (dry basis) were  
100 48.88, 17.83 and 6.51% for WS; and 43.00, 22.11 and 18.00% for CS, respectively.

101

### 102 **2.2 LCB pretreatment methods**

103 Hemicellulosic hydrolysates were produced by two methods: a) dilute acid pretreatment  
104 (DAP) and b) autohydrolysis followed by a very-dilute acid pretreatment (AH-VDAP).  
105 DAP was carried out in an autoclave at 121°C for 1 hour with a 15% (w/v) solid loading  
106 and 1.5% (v/v) H<sub>2</sub>SO<sub>4</sub>. Liquid fractions from DAP using WS and CS as feedstock were  
107 identified as WSC and CSC, respectively. Autohydrolysis (AH) was carried out in a semi-  
108 pilot scale pretreatment continuous tubular reactor (PCTR) at 1034 kPa (185°C) with a  
109 mean residence time of 18 min [11]. Pretreated biomass from AH was further hydrolysed in  
110 an autoclave at 121°C for 60 min using H<sub>2</sub>SO<sub>4</sub> 0.25% (v/v) with a 1:2 (w/v) ratio solids  
111 loading. Liquid fractions from AH-VDAP using wheat straw and corn stover as feedstock  
112 were identified as WSP and CSP, respectively. The hydrolysates composition is shown in  
113 Table 1. Further hydrolysates dilutions were made to obtain concentration between 10-15

114 g/L of total reducing sugars, which were used in the experiments of co-production of  
115 hydrogen and ethanol via dark fermentation described in the following subsection.

116

### 117 **2.3 Hydrogen and ethanol co-production via dark fermentation by the genetically** 118 **modified *E. coli* strain**

119 The genetically modified *E. coli* strain used in this work has a genotype that corresponds to  
120 absence of *hycA*, *ldhA* and *frdD* genes. These genes were deleted as described elsewhere  
121 [32]. This genotype confers the strain the ability to overproduce ethanol (EtOH) and  
122 hydrogen (H<sub>2</sub>), as well as decreasing the amounts of lactic and succinic acids produced.  
123 From now on, this strain will be referred as EtOH-H<sub>2</sub>-coproducing *E. coli*. The co-  
124 production of hydrogen and ethanol was performed using hemicellulosic hydrolysates as  
125 substrates, at 31°C and initial pH of 8.2. Diluted WSC, CSC, WSP, and CSP were used to  
126 determine the effect of LCB pretreatment method on the co-production of hydrogen and  
127 ethanol by the coproducing *E. coli*. These experiments were carried out in anaerobic  
128 serological bottles (0.01 L working volume) containing 10-15 g/L of total reducing sugars,  
129 B buffer [33], 1 mL/L trace elements solution [34], 0.01 g/L MgSO<sub>4</sub> and 1 g/L yeast  
130 extract. Cultures were started with an optical density of 0.2 measured at a wavelength of  
131 600 nm and were shaken at 200 rpm until no generation of hydrogen was observed. The  
132 experiments were carried out in quadruplicate. Production of hydrogen and ethanol was  
133 measured as it is indicated in Section 2.4.

134

### 135 **2.4 Analytical methods**

136 Total reducing sugars (TRS) was determined by the dinitrosalicylic acid (DNS) method  
137 [35], with some modifications as follows: 250 µL of the diluted sample with 750 µL of

138 DNS reagent (10 g/L NaOH, 200 g/L KNaC<sub>4</sub>H<sub>4</sub>O<sub>6</sub>·4H<sub>2</sub>O, 0.5 g/L Na<sub>2</sub>S<sub>2</sub>O<sub>5</sub>, 2 g/l C<sub>6</sub>H<sub>6</sub>O, 10  
139 g/L 3,5-Dinitrosalicylic acid) were heated for 5 minutes at 100°C and then cooled down to  
140 room temperature. Once tempered, 400 µL of distilled water were added. Xylose (0.1 to 1.0  
141 g/L) was used as the reference standard. The absorbance was measured at 595 nm (iMark™  
142 Microplate Absorbance Reader).

143 Simple sugars and metabolites were quantified by an Agilent HPLC equipped with a  
144 refractive index detector (Agilent Technologies 1220 Infinity LC), using a Rezex™ ROA-  
145 Organic Acid H<sup>+</sup> (Phenomenex) column, operated at 60°C with H<sub>2</sub>SO<sub>4</sub> 0.0025 M as a  
146 mobile phase (0.50 mL/min). Furfural was analysed by gas chromatography (Agilent  
147 Technologies 6890N Network GC Systems) using a capillary column HP-Innowax (30 m  
148 length × 0.25 mm inner diameter × 0.25 µm film thickness; Agilent Technologies). Injector  
149 and flame ionization detector (FID) temperatures were 220 and 250 °C, respectively.  
150 Helium was used as carrier gas at a flow rate of 25 cm<sup>3</sup>/min. Analyses were performed with  
151 a split ratio of 10:1 and a temperature program of 35 °C for 2 min, then 10°C/min to 210°C  
152 for 1 min.

153 Gas production was measured by acidified water (pH ≤ 2) displacement in an inverted  
154 burette connected to serological bottles with rubber tubing and a needle. The hydrogen  
155 concentration (% , v/v) in the gas phase was determined by gas chromatography with a  
156 thermal conductivity detector (Agilent Technologies 6890N Network GC Systems) using  
157 an Agilent J&W HP-PLOT Molesieve column (30 m length × 0.32 mm inner diameter × 12  
158 µm film thickness) under the following conditions: 200°C, injector temperature; 280°C,  
159 detector temperature; 300°C, oven temperature. Helium was used as carrier gas. Hydrogen  
160 volume was corrected to standard conditions of temperature and pressure (298.15K and 10<sup>5</sup>  
161 Pa).



162

### 163 **3 Modelling and simulation**

#### 164 **3.1 Process description of the biorefinery design**

165 Biorefining schemes were designed to produce ethanol from lignocellulosic biomass. All  
166 schemes are similar, differing only in the feedstock (WS or CS) and pretreatment method  
167 (DAP or AH-VDAP). Fig. 1 shows a block diagram of the proposed biorefinery conceptual  
168 design co-producing ethanol and hydrogen using WS or CS as feedstocks with an installed  
169 capacity of 500-ton biomass/day. Biogas produced from biorefining residues, hydrogen,  
170 lignin, and fermentation residues are used in a co-generation stage for steam and electricity  
171 production. The biorefineries designs were based on the models previously described  
172 elsewhere [30,36], with the following particularities:

- 173 a) DAP or AH-VDAP were applied as pretreatment methods of LCB to obtain  
174 hemicellulosic hydrolysates (see description on Section 2.2).
- 175 b) Hemicellulosic hydrolysates from the pretreatment stage (see description on Section 2.3)  
176 were used as substrate by the EtOH-H<sub>2</sub>-coproducing *E. coli* strain. In the biorefinery  
177 designs, this stage provided part of the total production of ethanol, as well as the  
178 hydrogen used in the cogeneration stage.

179 The biorefinery designs evaluated were termed WSB1, WSB2, CSB1 and CSB2 (WSB:  
180 biorefineries using wheat straw as feedstock; CSB: biorefineries with corn stover as  
181 feedstock; 1: DAP as pretreatment method; 2: AH-VDAP as pretreatment method). Each  
182 design was composed by a traditional lignocellulosic ethanol production train [feedstock  
183 conditioning (feedstock cleaning and size reduction), pretreatment (DAP or AH-VDAP),  
184 enzymatic saccharification, alcoholic fermentation, separation (azeotropic distillation and  
185 molecular sieving)], a dark fermentation (hydrogen and ethanol co-production) stage, a

186 wastewater treatment (anaerobic treatment, aerobic treatment and clarification) plant, and a  
187 cogeneration (steam and electricity) stage. The process inputs for each designs were raw  
188 materials ( $H_2SO_4$ ,  $Ca(OH)_2$ , enzymes, yeasts, bacteria, *E. coli* WDH-LF, flocculants, and  
189 antifoams), utilities (fresh water, pressurized air, electricity, steam-generator fuel) and the  
190 feedstock (WS or CS). The outputs were energy (electricity), steam, wastes (water,  $CO_2$ ,  
191 ashes, cake, and other solid wastes) and ethanol as product. Biorefineries mass and energy  
192 steady state balances were implemented in continuous mode and solved using the SuperPro  
193 Designer v8.5 (SPD) simulator [37]. Process conditions and reactions rates for pretreatment  
194 and dark fermentation stages correspond to the experimental data obtained as described in  
195 Sections 2.2 and 2.3. Integration of energy and 20% of water recirculation to the process  
196 were considered. Process details are provided in the Supporting Information.

197

### 198 **3.2 Techno economic analysis**

199 The profitability of each biorefinery design proposed in the section above was analysed  
200 with techno economic analysis tools previously implemented and tested with similar  
201 designs to those proposed in this work [38,39]. The analysis was based on the Discounted  
202 Cash Flow Analysis (DCFA) method for Net Present Value ( $NPV$ ) = 0 [40], calculating  
203 total capital investment, total production cost ( $TPC$ ), and their contributions for all  
204 biorefinery designs. The biorefinery energy integration was carried out using the Pinch  
205 Point Analysis technique for maximum energy recovery [40]. The End Use Energy Ratio  
206 ( $EER$ ) was employed to evaluate the energy efficiency of each design.  $EER$  was defined as  
207 ratio of energy produced (steam, electricity, and chemical energy from ethanol) to the total  
208 energy consumed in the process (heating/cooling requirements and electricity) [41].  
209 Equipment size and cost were calculated based on plant capacity using the SPD economic

210 data-base and its construction material and capacity-based correlations. All costs and  
211 financial parameters corresponded to conditions (c. 2018) of the Mexican economy.  
212 Commissioning and plant life periods were fixed at 3 and 15 years, respectively, with 330  
213 operating days/year. The interest rate was set at 6% and equity at 30%. Full production was  
214 assumed to begin by the end biorefinery's commissioning.

215

### 216 **3.3 Sustainability analysis**

217 A sustainability analysis method –previously developed for assessing prospective  
218 biorefining technologies [30]– was employed to quantify the impact of the biorefinery  
219 designs in the environmental and economic domains. Where, those impacts **are** calculated  
220 with quantitative indicators for each domain. Each indicator **is** integrated by one or more  
221 metrics related to design and/or process variables of the biorefinery design in question.  
222 Table 2 shows the indicators and metrics **to be** evaluated for each domain (based on the  
223 sustainability framework previously used for similar evaluations [36,38]). Six indicators **are**  
224 part of the environmental domain whilst two indicators were defined for the economic  
225 counterpart. All metrics (environmental and economic) **are** translated to the same functional  
226 unit using ad-hoc dimensional functions and conversion coefficients that **are** defined based  
227 on regulatory frameworks where the biorefinery facilities would be located. In this study,  
228 the functional unit chosen was USD/MJ<sub>out</sub> to monetize the impacts per of unit of energy  
229 delivered by the biorefinery. The translation of metric values to the same functional unit  
230 was obtained applying the following equation

$$231 \eta_i = M_i \cdot C_i,$$

232 Where  $\eta_i$  is the monetized metric  $i$ ;  $M_i$  the metric value, and  $C_i$  the monetizing coefficient  
233 (*i.e.* conversion). In addition, signs were assigned to each metric according to its positive or

234 negative impact to the corresponding domain. A positive value **is** associated to a benefit  
235 received by the stakeholders, whilst a negative value **might** be interpreted as a cost that  
236 stakeholders must cover. Table 3 contains the metric values, as well as monetizing  
237 coefficients and their corresponding monetized values (USD/MJ<sub>out</sub>). One of them, the  
238 *Emitted GHG* indicator, or “carbon intensity” with its associated metric **was** calculated as  
239 the carbon dioxide produced during the fermentation processes and electricity cogeneration  
240 stages. The monetization **considers** an impact cost of \$123 USD per metric ton of CO<sub>2</sub> [42].  
241 This cost **includes** the damage caused to water resources, land and biodiversity, agriculture  
242 and forestry, ecosystems, and human health. The *Emitted non-GHG* indicator, whose metric  
243 **is** composed by SO<sub>2</sub> emissions in the electricity generation stage, **was** monetized using the  
244 trading value of SO<sub>2</sub> in the US Acid Rain Program [43]. *Water consumption* and  
245 *Wastewater quality* indicators were monetized according to current Mexican environmental  
246 regulations [44,45]. These indicators **are** formed by more than one metric (Table 2).  
247 Therefore, their monetized value **is** the sum of their monetized metrics. The *Amount of*  
248 *produced solid wastes* indicator was monetized using the cost of solid waste management  
249 services in Mexico, which is \$76.89 USD per ton of waste transferred and disposed [46].  
250 As was described in Section 3.2, the *EER* indicator **is** the ratio of energy produced to the  
251 total energy consumed in the process. In the economic domain, two indicators **were**  
252 considered: *TPC* and *Electrical productivity*. The *TPC* indicator was monetized by  
253 translating all products to their energy equivalents. Finally, the *Electrical productivity*  
254 indicator **was** monetizing using the cost per MJ of electricity produced as a fraction of the  
255 total energy generated by the biorefinery [37].  
256 Sustainability indicators per domain **are** calculated by adding their corresponding  
257 indicators. The “global sustainability value” **is** the sum of all the environmental and

258 economic indicators once they are monetized. For a comparative analysis of the impact of  
259 each indicator in the biorefinery designs, all metric values were normalized with respect to  
260 WSB1 design considered the base case.

261

## 262 **4 Results and discussion**

### 263 **4.1 LCB pretreatment methods**

264 The characterization of hemicellulosic hydrolysates (hemicellulose to pentoses conversion,  
265 composition of simple sugars, degradation compounds) from pretreatment is shown in  
266 Table 1. As expected, xylose was the highest concentration sugar monomer in the  
267 hydrolysates obtained with both LCB pretreatment methods. DAP decomposed  
268 hemicellulose while maintained cellulose and lignin almost intact [27,47]. WSC and CSC  
269 produced 29 and 34.3 g/L of xylose, respectively. Regarding AH-VDAP, the xylan  
270 backbone was selectively depolymerized during autohydrolysis, resulting into  
271 xylooligosaccharides (XOS) as main reaction products [48]. These XOS were  
272 depolymerized during the subsequent very-diluted acid pretreatment stage, resulting in  
273 large concentrations of xylose at the end of the pretreatment. WSP and CSP generated 39.8  
274 and 41.1 g/L of xylose, respectively. Glucose concentrations in WSP and CSP were 3-fold  
275 less than in WSC and CSC hydrolysates (Table 1) since the glucan of LCB biomass was  
276 not depolymerized during autohydrolysis as reported previously [11], and the sulphuric acid  
277 concentration of very-diluted acid pretreatment stage was chosen for depolymerizing XOS.  
278 The aims of pretreatment are to disrupt the crystalline cellulose structure and to fractionate  
279 the main components of the feedstock [49]. However, during pretreatment of LCB, by-  
280 products are often produced that can inhibit downstream biochemical processes. These  
281 inhibitors are formed when hemicellulose, cellulose and/or lignin are solubilized and

282 degraded [50,51]. Acetate was the main pretreatment by-product found in all hydrolysates,  
283 followed by formate and furfural. Acetate results from the hydrolysis of acetyl groups of  
284 hemicelluloses [52], and it was detected in concentrations higher than 4 g/L (Table 1). Even  
285 though acetate, formate and furfural have relatively low toxicity [50], to avoid their  
286 possible inhibiting effects in dark fermentation experiments, the hydrolysates were diluted  
287 with water. After dilution, the concentration of total reducing sugars was 15.1, 9.7, 10.2 and  
288 11.1 g/L in WSC, WSP, CSC and CSP, respectively.

289

#### 290 **4.2 Dark fermentation by the EtOH-H<sub>2</sub>-coproducing *E. coli* strain**

291 The effect of DAP and AH-VDAP hydrolysates on the EtOH-H<sub>2</sub>-coproducing *E. coli* are  
292 shown in Fig. 2. Using WSC as substrate up to 1.3-fold more hydrogen was obtained than  
293 with WSP (Fig. 2A). This is because of the difference of TRS concentration among  
294 hydrolysates after dilution, with WSC having 1.6-fold more TRS than WSP. Similar event  
295 was observed using CSC or CSP as substrates. With a TRS concentration in CSP 1.1-fold  
296 higher than in CSC, CSP produced  $2,930.3 \pm 189.4$  mL H<sub>2</sub>/L whilst CSC reached  $2,576.4 \pm$   
297  $220.4$  mL H<sub>2</sub>/L. Regarding hydrogen production rate,  $22.1 \pm 1.1$ ,  $18.3 \pm 1.0$ ,  $20.2 \pm 1.7$  and  
298  $18.5 \pm 1.3$  mL H<sub>2</sub>/L·h were obtained using WSC, WSP, CSC and CSP as substrate,  
299 respectively. Note that production rates were higher with DAP hydrolysates than with the  
300 AH-VDAP counterparts. Hydrogen production kinetics and the percentage of hydrogen in  
301 the gas attained by the EtOH-H<sub>2</sub>-coproducing *E. coli* are showed in Fig. 3. None of the  
302 batches presented lag phase since hydrogen was found from the first gas sampling (17 h).  
303 The maximum concentration of hydrogen (% , v/v) in the gas phase detected was 56% (at 17  
304 h, WSC), 45% (at 40 h, WSP), 46% (at 40 h, CSC) and 50.2% (at 40 h, CSP). Hydrogen  
305 production declined after 120 h of fermentation, regardless of the kind of feedstock and

306 type of pretreatment used. Fig. 2B shows the hydrogen yield and TRS consumption by the  
307 EtOH-H<sub>2</sub>-coproducing *E. coli* strain. The TRS consumption was 10% higher using WSP as  
308 substrate than with WSC. However, this was not observed with CSC or CSP. The yield of  
309 hydrogen achieved by EtOH-H<sub>2</sub>-coproducing strain was  $311.5 \pm 30.7$ ,  $323.1 \pm 6.6$ ,  $312.3 \pm$   
310  $26.7$  and  $337.1 \pm 21.8$  mL H<sub>2</sub>/g TRS using WSC, WSP, CSC and CSP as substrate,  
311 respectively. Regarding ethanol production (Fig. 2C), up to  $3.54 \pm 0.27$  g/L were produced  
312 at the end of fermentation, achieving yields in the range of  $0.32 \pm 0.01$  to  $0.34 \pm 0.06$  g  
313 EtOH/g TRS (for all hydrolysates). Therefore, amount of ethanol produced per TRS unit  
314 seems to be not affected by feedstock or pretreatment method.

315 The co-production of hydrogen and ethanol by microorganisms had been studied using  
316 genetically engineered *E. coli* strains. Different molecular strategies had been tested to  
317 enhance the fermentation efficiency of *E. coli* strains, such as deletion of genes including  
318 those to produce hydrogenases, negative regulator of the formate regulon, lactate  
319 dehydrogenase, fumarate reductase and phosphoglucose isomerase, among other  
320 [20,25,26,53], as well as heterologous gene expression [54]. Many of these studies used  
321 glucose as carbon source instead of LCB hydrolysates. Interestingly, reported hydrogen and  
322 ethanol yields are lower than those obtained in this work (Table 4). In previous studies,  
323 whet straw hydrolysate was used as substrate for co-production hydrogen and ethanol by  
324 metabolic engineered *E. coli* strains, WDHL [22] and WDHGFA [23]. Reported yields of  
325 hydrogen and ethanol obtained by WDHL strain were 159 mL H<sub>2</sub>/g sugar and 0.32 g  
326 EtOH/g sugar, while WDHGFA strain reached 160 mL H<sub>2</sub>/g sugar and 0.26 g EtOH/g  
327 sugar. These amounts are either similar or lower up to 47% than those obtained here as seen  
328 in Table 4.

329 Since the aim of this work was to improve the lignocellulosic ethanol biorefinery  
330 performance by co-producing hydrogen and ethanol by genetically modified *E. coli*, the  
331 experimental data provided above was included in the conceptual design of (environmental  
332 and economic) sustainable biorefineries, as described in the following subsections.

333

### 334 **4.3 Mass balances**

335 As mentioned previously, DAP and AH-VDAP of WS and CS were included as  
336 pretreatment methods to compare their effect on the dark fermentation performance and  
337 therefore on the biorefinery economics. The process schemes evaluated were WSB1,  
338 WSB2, CSB1 and CSB2 (see Section 3.1 for a detail description). The mass balances for  
339 the stages involved in ethanol production for all biorefineries are shown in Figs. 4 and 5.  
340 For mass conversion ( $X_{A \rightarrow B}$ ) data see Tables A1, A2 and A3 in the Supporting Information.  
341 Table 5 shows output flowrates from each of the biorefining stages. CSB1 and CSB2  
342 produced 4,825 and 4,912 kg/h of pentoses in the pretreatment stage, 24% more than  
343 WSB1 and WSB2. This is because hemicellulose fractions in CS are 1.2-fold higher than in  
344 WS, as well as the  $X_{Hemic. \rightarrow Pentoses}$  during the pretreatment stage in CSB1 and CSB2 is 7%  
345 and 5% higher than in WSB1 and WSB2, respectively.

346 In the dark fermentation stage, the hemicellulosic hydrolysates from the pretreatment stage  
347 were used by the EtOH-H<sub>2</sub>-coproducing *E. coli* strain to obtain hydrogen and ethanol.  
348 WSB1, WSB2, CSB1 and CSB2 produced 26.3, 25.1, 44.1 and 45.8 kg/h of hydrogen,  
349 respectively, which were fed to the cogeneration stage for electricity production. The  
350 difference in hydrogen production among schemes is due to a smaller  $X_{Sugars \rightarrow H_2}$  (15%) and  
351 the lower hemicellulose fraction for WS (Table 5). Regarding ethanol, WSB1, WSB2,



352 CSB1 and CSB2 produced 6,093, 6,126, 5,742 and 5,796 kg/h of ethanol, respectively.  
353 Around 20-30% of this comes from dark fermentation, and the rest from alcoholic  
354 fermentation (Table 5). Since WS presented the highest cellulose content, WS-based  
355 biorefineries produces 6% more ethanol than CS-based counterparts.

356

#### 357 **4.4 Techno economic analysis results**

358 After establishing the contribution of dark fermentation over ethanol production in the  
359 proposed conceptual designs, their profitability was determined by a techno economic  
360 analysis considering Mexican economic conditions (c. 2018). The total equipment cost is  
361 10.1%, 46.8% and 22.2% higher in WSB1, CSB1 and CSB2 compare to WSB2,  
362 respectively (Fig. 6A). The equipment cost per stage, as shown in Fig 6B, is similar in all  
363 cases, except for the pretreatment and dark fermentation stages. On one hand, CS-based  
364 biorefineries have the most expensive dark fermentation stage (\$57-58 USD millions) due  
365 to the higher amount of pentoses obtained from the pretreatment stage than with WS.  
366 Therefore, a larger amount of water is needed to achieve the sugars concentration required  
367 in the dark fermentation stage. As a consequence, higher volume reactors must be  
368 employed with larger costs. On the other hand, the pretreatment stage equipment cost of  
369 AH-VDAP biorefineries (WSB2 and CSB2) is about 60 % lower than their DAP  
370 counterparts since a continuous reactor was considered for this case.

371 Fig. 7 shows the TPC calculated per litre of ethanol for each biorefining design, as well as  
372 their ethanol production. The lowest *TPC* (\$1.37 UDS/L EtOH, Fig. 7A) was obtained by  
373 WSB2, which is 17.9, 43.1 and 15.9% lower than those obtained for WSB1, CSB1 and  
374 CSB2, respectively. Even when WS is 2.5-fold more expensive than CS (Table 1),  
375 feedstock cost seems not to contribute to *TPC* in that proportion. WS seems to be a better

376 feedstock for ethanol production compared to CS due to its higher cellulose content. The  
377 most important contributors to *TPC*, as shown in Fig. 7B, are operation cost followed by  
378 services (cooling and heating) and total capital investment, with values around 29.8-34.7%,  
379 14.5-22.4% and 15.4-18.3%, respectively. Operating cost include maintenance, operating  
380 supplies, labour, and direct supervision, laboratory charges, patents, and royalties.  
381 Regarding the services contribution to *TPC*, the highest values were corresponding to those  
382 designs using DAP pretreatment since a higher amount of cooling water is required by the  
383 process during pretreatment stage. Electricity consumption is not a relevant contributor to  
384 *TPC*. Electricity demand (*Electricity<sub>in</sub>*) of all biorefineries is more than 5,800 kWh of  
385 electricity (Fig. 8). However, they produce (*Electricity<sub>out</sub>*) just around 20-30% of this  
386 demand, thus *EER* is lower than 0.50 for all designs. WSB2 is the most economical option,  
387 because it produces the largest amount of ethanol with the lowest equipment cost.

388

## 389 **4.5 Sustainability analysis results**

390

### 391 **4.5.1 Environmental Sustainability Analysis Results**

392 The results of the sustainability analysis for the environmental domain are summarized in  
393 Table 3. Once monetized, all indicators were of negative value, with the *EER* indicator as  
394 the main contributor in this domain, with a contribution of more than 67% of the *Total*  
395 *Environmental Indicator* for all designs. The electricity dependence has been observed in  
396 the analysis of other biorefinery designs producing biofuels as the main product [30]. Other  
397 important indicator is *EGHG* with a contribution of around 15-21%. *WCo* and *WWQ*  
398 indicators provide a contribution about  $\leq 7\%$  for all cases because the biorefinery was  
399 designed for water recirculation and for complying with the Mexican regulatory framework

400 for discharges to water bodies [44]. *SW* indicator is the lowest contributor with  $\leq 3.7\%$  for  
401 all cases.

402 To compare the environmental indicators performance among biorefineries, metric values  
403 were normalized with respect to WSB1 (base case) as shown in Fig. 9. *T* and *pH* metrics  
404 were not included because they are similar in all cases. For  $M_{CO_2}$  –which is related to GHG  
405 emission generated during dark fermentation, alcoholic fermentation and cogeneration  
406 stages–, CS-based biorefineries produced around 22-25% more g CO<sub>2eq</sub> per MJ than the  
407 base case since a higher amount of lignin, H<sub>2</sub> and biogas are fed to the cogeneration stage  
408 which is the main contributor to this metric and therefore to GHG emissions indicator. In  
409 the case of  $M_{SO_2}$ , the only non-greenhouse gas considered in this work is SO<sub>2</sub>, produced  
410 during the cogeneration stage due to sulphur contained in LCB. Corn stover (CSB1 and  
411 CSB2) biorefineries emitted 20% and 28% less g SO<sub>2eq</sub> by MJ produced than the base case,  
412 respectively. This is principally due to differences in feedstocks composition. Regarding  
413 water consumption ( $M_{fw}$ ), CS-based biorefineries employ around 26-39% more water than  
414 their WS-based counterparts. This is because the water required adjusting TRS in the  
415 pretreatment output stream feeding the dark fermentation stage. Discharged water ( $M_{dw}$ ,  
416  $L_{discharged\ water}/MJ_{out}$ ) by WSB1 was 36, 30 and 73% lower than WSB2, CSB1 and CSB2,  
417 respectively. For the *WWQ* indicator, the metrics  $M_{COD}$  and  $M_{dp}$  –which are related to  
418 organic material and other pollutants content in wastewater treated– are lower in CSB1 and  
419 CSB2 than in WS-based biorefineries, because CSB1 and CSB2 streams are more diluted  
420 than those for the other two designs. The metric ( $M_{sw}$ ) of *SW* indicator is directly related to  
421 solids generated by *pH* adjustment, as well as ash production in the cogeneration stage and  
422 dust from the conditioning stage. The *pH* in the dark fermentation stage by EtOH-H<sub>2</sub>-

423 coproducing *E. coli* is 8.2. Therefore, the hydrolysates coming from the DAP-based  
424 biorefineries (WSB1 and CSB1) demand larger Ca(OH)<sub>2</sub> amounts than their counterparts,  
425 thus producing 89 and 77% more solid wastes than WSB2 and CSB2, respectively.  
426 Regarding energy self-sufficiency, WSB2 is the biorefinery with the highest *EER* value  
427 (0.49) because is the biorefinery with the largest ethanol production, surpassing in 12% the  
428 base scheme (Fig. 9). However, none of the biorefinery designs was energetically self-  
429 sufficient.

430

#### 431 **4.5.2 Economic Sustainability Analysis Results**

432 The results of the sustainability analysis for the economic domain are presented in Table 3.  
433 After monetization, *TPC* is the most relevant indicator, with a 99% contribution for all  
434 cases. Therefore, WSB2 is the best alternative in the economic domain due to its lowest  
435 *TPC*. The indicator normalization using WSB1 as base case is shown in Fig. 10. WSB2  
436 exhibited the lowest *TPC* (Fig. 7A), due to the lowest total equipment investment and  
437 highest ethanol production (Figs. 6A, 7A) as explained in Section 4.4. CSB1 and CSB2  
438 exhibited the highest electrical productivity, surpassing in 60% and 46% the base case,  
439 respectively. Since the contribution of this indicator to the economic sustainability indicator  
440 is  $\leq 1\%$ , its impact is not relevant in the *Total Economic Indicator*.

441

#### 442 **4.5.3 Global Sustainability Analysis Results**

443 The indicator values for each domain together with the global sustainability indicator are  
444 shown in Fig. 11. These values represent what stakeholders should pay per each MJ  
445 produced for either fines, and environmental damages caused by the biorefinery regarding  
446 the environmental domain or production costs considering the economic domain. From an

447 environmental point of view, the lowest impact is associated with WSB2, (-0.047  
448 USD/MJ<sub>out</sub>) since its *EER* indicator (positive) was the highest, as well as *EGHG* and *SW*  
449 (negative) indicators were the lowest values of all designs. The absolute value of this  
450 indicator is 26, 69 and 48% lower than those calculated for WSB1, CSB1 and CSB2,  
451 respectively. From an economic perspective, WSB2, again, achieved the lowest value of the  
452 four proposed schemes, with -0.064 USD per MJ produced, mainly due to its lowest *TPC*.  
453 Further, considering the *Global Sustainability Indicator*, the smallest value of all  
454 biorefinery scenarios is -0.111 USD per MJ produced, associated to WSB2. 43% of its  
455 value corresponds to the *Total Environmental Indicator*, and the rest to the *Total Economic*  
456 *Indicator*. The second-best option is WSB1, with a global impact value 23% higher than  
457 that for WSB2.

458

## 459 **5 Conclusions**

460 The dark fermentation stage –by EtOH-H<sub>2</sub>-coproducing *E. coli*– contributes with 20 to  
461 30% of the total ethanol production in the lignocellulose-based biorefinery designs is  
462 proposed in this work. From all designs, WSB2 (wheat straw as feedstock and AH-VDAP  
463 as LCB pretreatment method) could generate the smallest environmental impact with the  
464 lowest *TPC*, which is up to 43% lower than its counterparts. The sustainability analysis  
465 shows the importance of environmental issues compared against economic aspects, fact that  
466 is not evident using conventional techno economic analysis tools. Based on the regulatory  
467 framework employed, the environmental monetized impact of the most sustainable design  
468 resulted almost as important as the economic aspects of it. Therefore, the results show that  
469 co-production schemes are an alternative for ethanol biorefineries that must be explored  
470 further.

471

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477

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680 **Figure captions**

681 **Graphical abstract**

682

683 **Fig. 1 Biorefinery block diagram.** The process inputs are indicated in blue arrows, and the outputs  
684 are marked in green arrows

685

686 **Fig. 2 Effect of the pretreatment method of lignocellulosic biomass in co-production of**  
687 **hydrogen and ethanol by EtOH-H<sub>2</sub>-coproducing *E. coli*.** Batch cultures of 0.01L were performed  
688 at 31°C and initial pH 8.2 using WSP, WSC, CSP and CSC as substrates. Production and  
689 production rate of hydrogen (A), hydrogen yield and TRS consumption (B), production and yield of  
690 ethanol (C). Data are presented as mean ± standard deviation

691

692 **Fig. 3 Kinetics of hydrogen production.** Batch cultures (0.01 L) at 31°C and initial pH 8.2 using  
693 WSP (A), WSC (B), CSP (C) and CSC (D) as substrates. Data are presented as mean ± standard  
694 deviation

695

696 **Fig. 4 Mass balance for biorefining stages in DAP biorefineries (WSB1 and CSB1)**

697

698 **Fig. 5 Mass balance for biorefining stages in AH-VDAP biorefineries (WSB2 and CSB2)**

699

700 **Fig. 6 Total equipment investment (A) and equipment investment contributions by stage (B)**  
701 **for all biorefineries schemes**

702

703 **Fig. 7 Technoeconomic analysis results.** *TPC* and ethanol production (A), *TPC* contributions (B)

704

705 **Fig. 8 Electricity in-out, electrical productivity and *ERR* for all biorefinery designs**

706

707 **Fig. 9 Indicator analysis for the environmental domain**

708

709 **Fig. 10 Indicator analysis for the economic domain**

710

711 **Fig. 11 Sustainability global values for each biorefinery design**

712 **Table 1** Characterization of the hemicellulosic hydrolysates

Hydrolysate	Feedstock	Feedstock cost (USD/kg)	Pretreatment	$X_{Hemic. \rightarrow Pentoses}$ (%)	Composition (g/L)					
					Glucose	Xylose	Arabinose	Formate	Acetate	Furfural
WSC	Wheat straw	\$0.08	DAP	88	5.5	29	6.8	1.8	4.2	ND
WSP			AH-VDAP	90	1.8	39.8	7.9	2.6	7.8	1.2
CSC	Corn stover	\$0.03	DAP	95	4.7	34.3	9.4	1.2	4.1	ND
CSP			AH-VDAP	95	1.5	41.1	11.3	3.3	4.2	0.8

713 DAP: Diluted acid pretreatment; AH-VDAP: Autohydrolysis followed by very-diluted acid pretreatment;  $X_{Hemic. \rightarrow Pentoses}$ : Hemicellulose to pentoses mass-  
 714 conversion; ND: No determinate

715



716 **Table 2** Sustainability framework

Domain	Indicator	Metric, units	Dimensional function	Reference
Environmental	<i>Emitted GHG (EGHG)</i>	$M_{CO_2}, gCO_{2eq}/MJ_{out}$	$M_{CO_2} \cdot C_{CO_2}$	[42]
	<i>Emitted non-GHG (NGHG)</i>	$M_{SO_2}, gSO_{2eq}/MJ_{out}$	$M_{SO_2} \cdot C_{SO_2}$	[43]
	<i>Water consumption (WCo)</i>	$M_{fw}, L_{fresh\ water}/MJ_{out}$	$M_{fw} \cdot C_{fw}$	[44,45]
		$M_{dw}, L_{discharged\ water}/MJ_{out}$	$M_{dw} \cdot C_{dw}$	
	<i>Wastewater quality (WWQ)</i>	$M_{COD}, mgCOD/L_{water}$	$M_{COD} \cdot C_{COD}$	
		$M_{dp}, kg\ dissolved\ pollutants/MJ_{out}$	$M_{dp} \cdot C_{dp}$	
		T, °C pH	- -	- -
<i>Amount of produced solid wastes (SW)</i>	$M_{sw}, kg\ disposable\ wastes/MJ_{out}$	$M_{sw} \cdot C_{ws}$	[46]	
<i>End Use Energy ratio (EER)</i>	$M_{EER}, MJ_{out}/MJ_{in}$	$(M_{EER}-1) \cdot C_{TPC}$	-	
Economic	<i>Total production cost (TPC)</i>	$M_{TPC}, USD/L_{EtOH}$	$M_{TPC} \cdot C_{TEP}$	-
	<i>Electrical productivity (E)</i>	$M_E,$ $Electricity_{out}/Electricity_{in}$	$(M_E-1) \cdot C_E$	[37]

717  $C_{TEP}$ : Total energy produced;  $C_E$ : Cost per MJ of electricity produced as a fraction of total energy produced by  
718 the biorefinery

719

720 **Table 3** Metric values for WSB1, WSB2, CSB1 and CSB2 biorefineries and monetizing coefficients for translating metric units to monetized  
 721 indicators (USD/MJ<sub>out</sub>)

Indicator	Metric	Metric value ( $M_i$ )				Monetizing coefficient ( $C_i$ )				Monetized metric value ( $\eta_i = M_i \cdot C_i$ ; USD/MJ <sub>out</sub> )				Metric contributions (%)				
		WSB1	WSB2	CSB1	CSB2	WSB1	WSB2	CSB1	CSB2	WSB1	WSB2	CSB1	CSB2	WSB1	WSB2	CSB1	CSB2	
<i>EGHG</i>	$M_{CO_2}$	80.70	80.16	101.11	98.76					-1.23E-04								
<i>NGHG</i>	$M_{SO_2}$	9.14	8.40	7.29	6.55					-6.00E-08								
<i>WCo</i>	$M_{fw}$	1.40	1.57	1.77	1.95					-1.01E-03								
	$M_{dw}$	0.84	1.14	1.09	1.45					-2.38E-04								
<i>WWQ</i>	$M_{COD}$	129.99	115.45	99.46	91.63	-2.06E-05	-2.78E-05	-2.79E-05	-3.68E-05	-2.68E-03	-3.21E-03	-2.77E-03	-3.37E-03	4.52	6.80	3.47	4.82	
	$M_{dp}$	5.16E-10	4.53E-10	4.31E-10	3.92E-10					-1.40E-04								
	T	32.58	31.56	33.01	31.95					0	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
	pH	7	7	7	7					0	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
<i>SW</i>	$M_{sw}$	3.00E-02	3.31E-03	2.78E-02	6.54E-03					-7.40E-02								
<i>ERR</i>	$M_{EER}$	0.44	0.49	0.33	0.29	-7.61E-02	-6.30E-02	-9.00E-02	-7.29E-02	-4.29E-02	-3.21E-02	-6.07E-02	-5.16E-02	72.29	67.90	75.85	73.81	
		<i>Total Environmental Indicator</i>								-0.059	-0.047	-0.080	-0.070					
<i>TPC</i>	$M_{TPC}$	1.61	1.37	1.96	1.59	-4.72E-02	-4.61E-02	-4.60E-02	-4.60E-02	-7.61E-02	-6.299E-02	-9.01E-02	-7.29E-02	99.26	99.16	98.97	98.81	
<i>E</i>	$M_E$	0.21	0.17	0.33	0.30	-2.75E-03	-3.07E-03	-2.81E-03	-2.93E-03	-5.69E-04	-5.32E-04	-9.33E-04	-8.81E-04	0.74	0.84	1.03	1.19	
		<i>Total Economic Indicator</i>								-0.077	-0.064	-0.091	-0.074					
		<i>Global Sustainability Indicator</i>								-0.136	-0.111	-0.171	-0.144					

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725 **Table 4** Comparison of hydrogen and ethanol yields obtained by genetically engineered *Escherichia coli* strains

Strain	Genotype description	Substrate	Hydrogen Yield (mL H <sub>2</sub> /g substrate)	Ethanol yield (g EtOH/g substrate)	Reference
SH9*_ <sub>ZG</sub>	<i>E. coli</i> BW25113 $\Delta hycA \Delta hyaAB \Delta hybBC \Delta ldhA \Delta frdAB$ $\Delta pfkA$ /pEcZG (pDK7 carrying <i>zwf</i> , and <i>gnd</i> )	Glucose (Glc)	265.6 <sup>‡</sup> (1.8 mol H <sub>2</sub> /mol Glc)	0.36 <sup>‡</sup> (1.4 mol EtOH/mol Glc)	[26]
SH5 $\Delta pgi$ _ <sub>ZLGG</sub>	<i>E. coli</i> BW25113 $\Delta hycA \Delta hyaAB \Delta hybBC \Delta ldhA \Delta frdAB$ $\Delta pgi$ /pLmZ-GoG (pDK7 carrying <i>zwf</i> of <i>E. coli</i> BW25113 and <i>gnd</i> of <i>G. oxydans</i> )		245.8 <sup>‡</sup> (1.74 mol H <sub>2</sub> /mol Glc)	0.41 <sup>‡</sup> (1.62 mol EtOH/mol Glc)	[25]
SS1- Recombinant <i>hybC</i>	<i>E. coli</i> SS1/pETDuet-1 (carrying <i>hybC</i> )		94.6 <sup>‡</sup> (0.67 mol H <sub>2</sub> /mol Glc)	0.15 <sup>‡</sup> (0.58 mol EtOH/mol Glc)	[53]
SH8*_ <sub>ZG</sub>	<i>E. coli</i> BW25113 $\Delta hycA \Delta hyaAB \Delta hybBC \Delta ldhA \Delta frdAB \Delta pfkA$ $\Delta pta$ - <i>ackA</i> -adaptive evolution /pEcG (pDK7 carrying <i>gnd</i> )		186.5 <sup>‡</sup> (1.32 mol H <sub>2</sub> /mol Glc)	0.35 <sup>‡</sup> (1.38 mol EtOH/mol Glc)	[54]
WDHL	<i>E. coli</i> W3110 $\Delta hycA \Delta lacI$		Wheat straw	159.3	0.32
WDHGFA	<i>E. coli</i> W3110 $\Delta hycA \Delta ptsG \Delta frdD \Delta ldhA$	hydrolysate	160 <sup>‡</sup> (0.24 mol H <sub>2</sub> /C-mol)	0.26 <sup>‡</sup> (0.195 mol EtOH/C-mol)	[23]
Ethanol- H <sub>2</sub> - coproducing <i>E. coli</i>	<i>E. coli</i> W3110 $\Delta hycA \Delta ldhA \Delta frdD$	WSC	311.5	0.33	This work
		WSP	323.1	0.32	
		CSC	312.3	0.34	
		CSP	337.1	0.34	

726 <sup>‡</sup>Converted units from the original data (reported units)

727

728 **Table 5** Pentoses, hydrogen and ethanol production during dark fermentation and alcoholic  
 729 fermentation stages in all biorefineries schemes

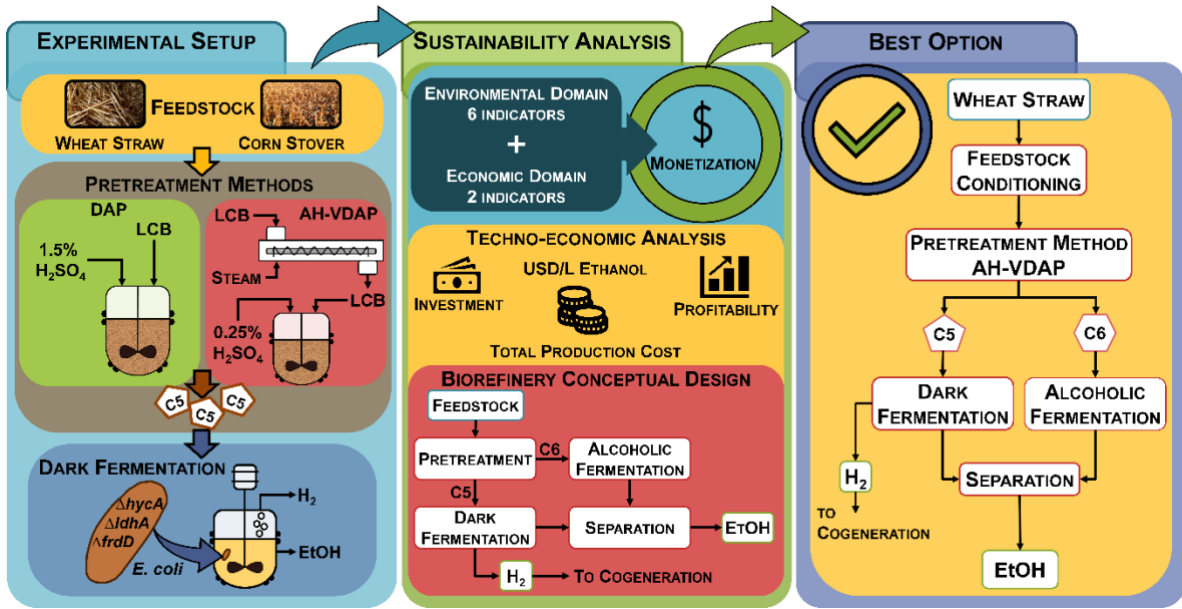
Biorefinery		WSB1	WSB2	CSB1	CSB2
Pretreatment stage	Pentoses (kg/h)	3,673	3,746	4,825	4,912
	$X_{Sugars \rightarrow H_2}$ (%)	14.6	15.1	19.9	21.5
Dark fermentation stage	H <sub>2</sub> (kg/h)	26.3	25.1	44.1	45.8
	$X_{Sugars \rightarrow EtOH}$ (%)	75.0	75.0	66.0	66.0
	EtOH (kg/h)	1,542	1,421	1,677	1,609
Alcoholic fermentation stage	EtOH (kg/h)	4,752	4,923	4,254	4,381
Ethanol production	EtOH (kg/h)	6,093	6,126	5,742	5,796

730  $X_{Sugars \rightarrow H_2}$ : Sugars (glucose and pentoses) to hydrogen mass-conversion during dark fermentation stage;

731  $X_{Sugars \rightarrow EtOH}$ : Sugars (glucose and pentoses) to ethanol mass-conversion during dark fermentation stage

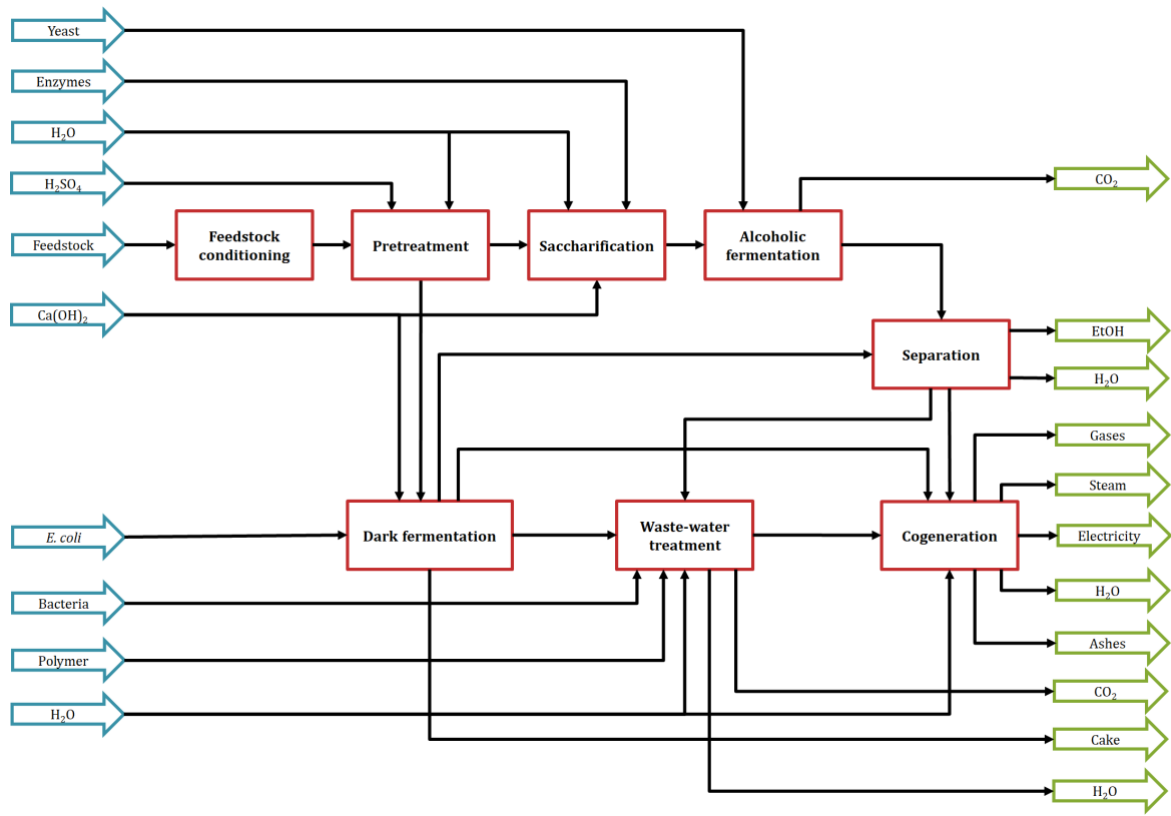
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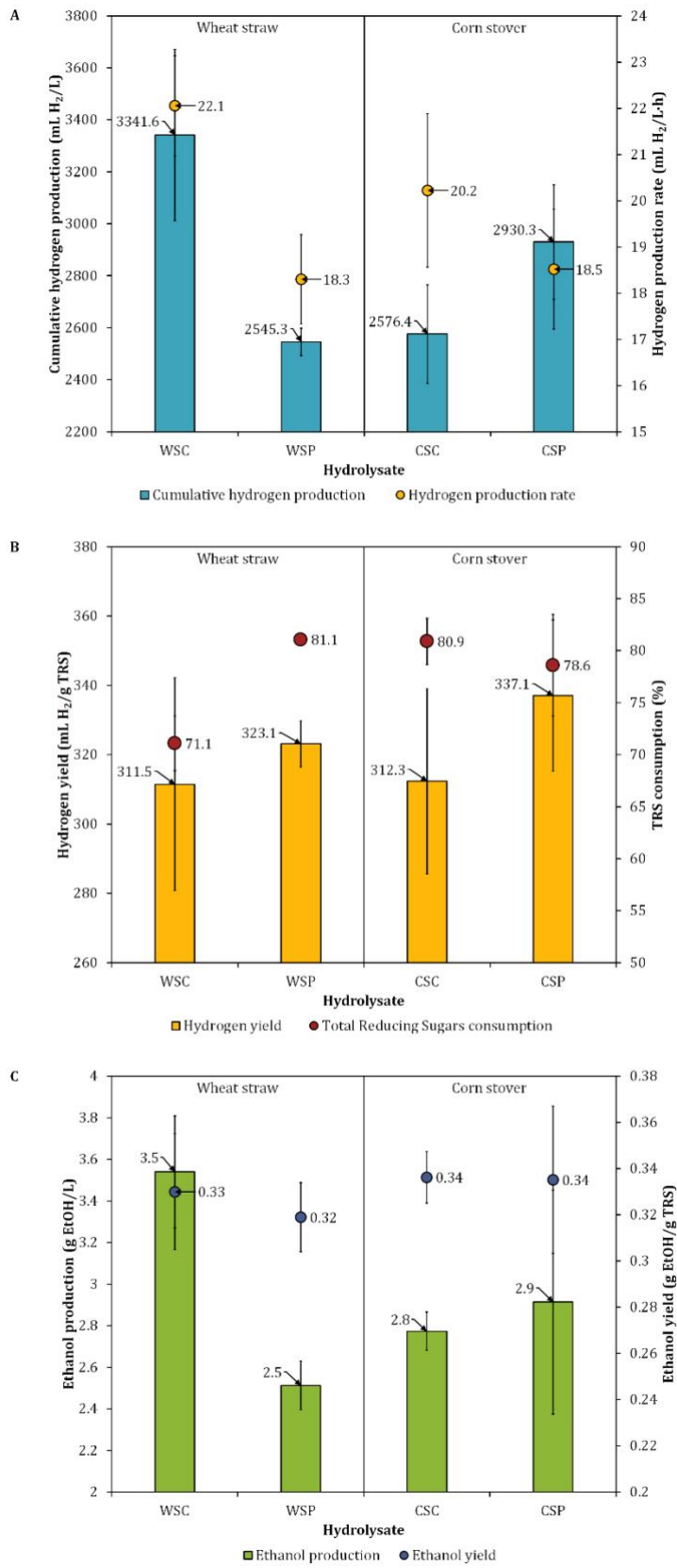
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735 Graphical abstract



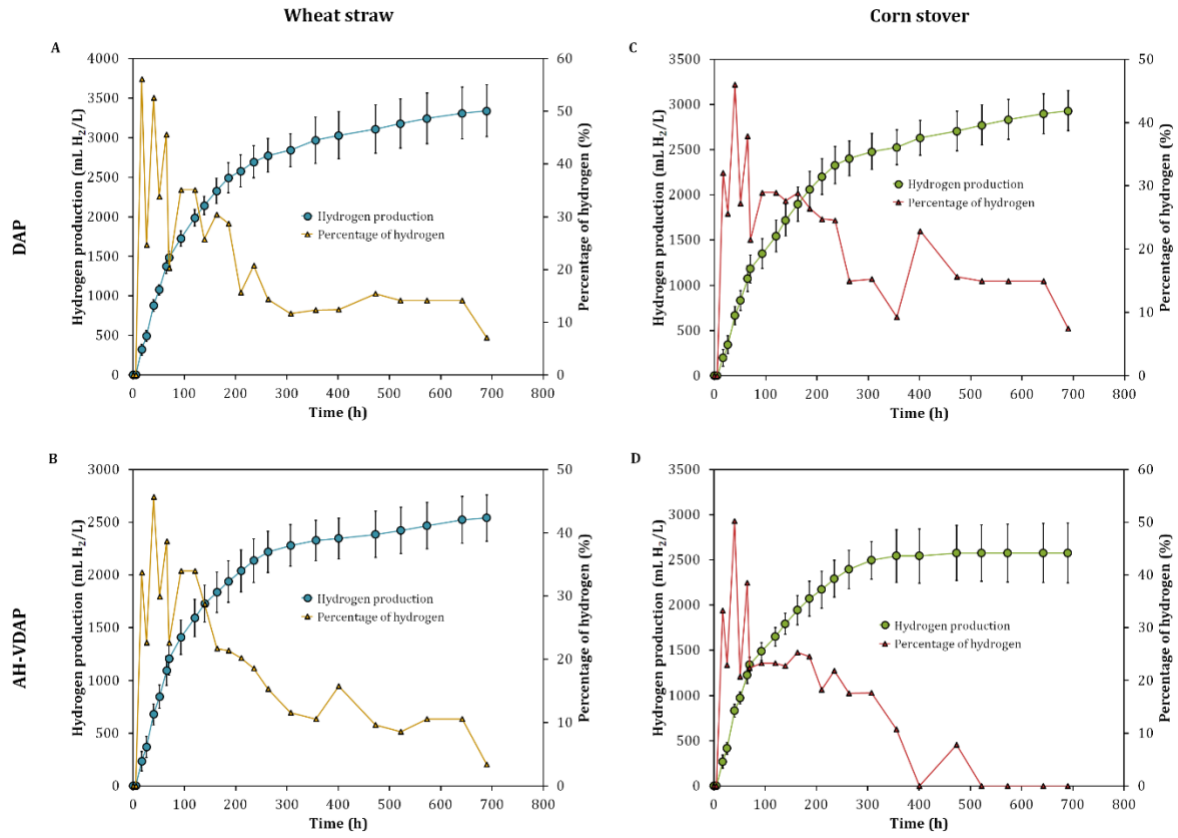
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737 **Fig. 1**



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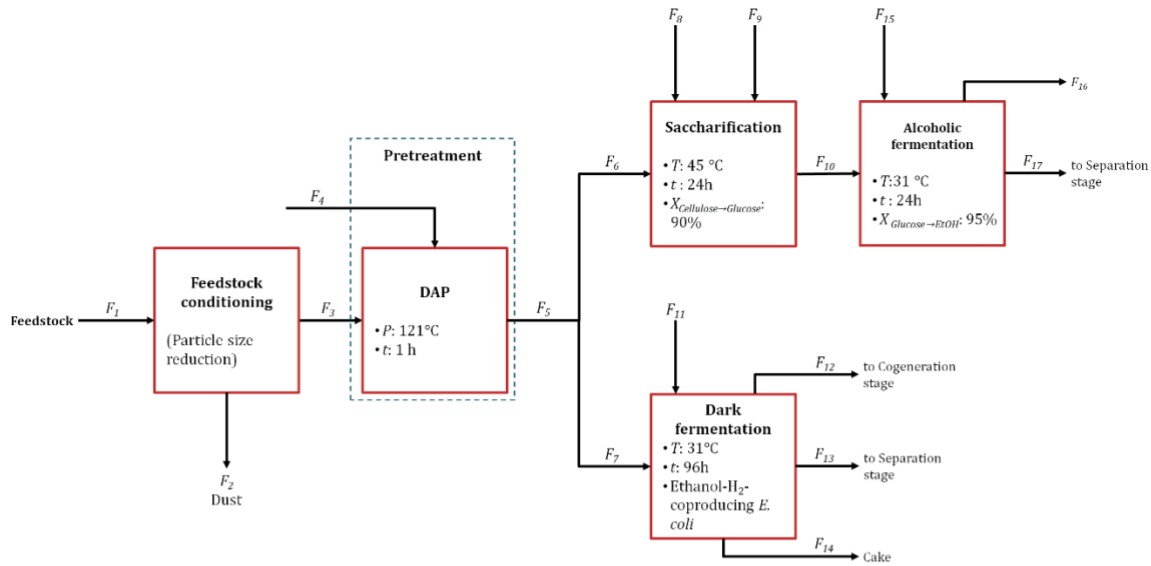
**Fig. 2**



**Fig. 3**

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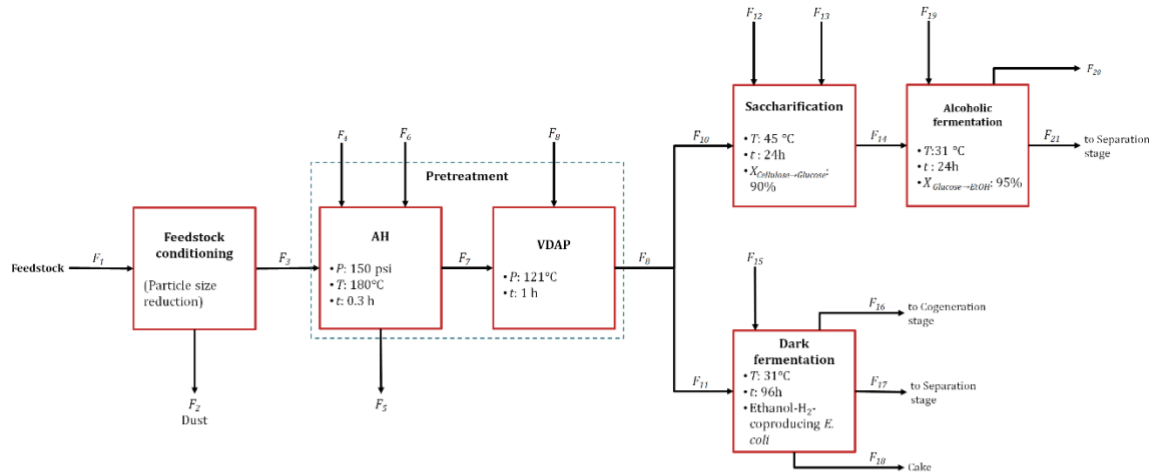




Biorefinery Feedstock Stream		$F_1$	$F_2$	$F_3$	$F_4$	$F_5$	$F_6$	$F_7$	$F_8$	$F_9$	$F_{10}$	$F_{11}$	$F_{12}$	$F_{13}$	$F_{14}$	$F_{15}$	$F_{16}$	$F_{17}$		
WSB1	Wheat straw	Flow (ton/day)	500.0	24.6	475.4	2315.2	2790.5	843.1	1949.9	5.4	1152.0	2003.7	2760.0	42.9	4149.0	471.8	4.4	109.0	1899.2	
		Composition (wt%)																		
		Cellulose	48.88	9.93	50.90		8.04	26.62					0.27							1.18
		Hemic.	17.83	3.62	18.57		0.38	1.27					0.53							0.56
		Lignig	6.51		6.85		1.17	3.86					1.62							1.71
		H <sub>2</sub> O				97.24	80.23	53.11	91.97		100.00		78.80	100.00		98.94	96.62			83.14
		Glucose					0.70	0.23	0.90				11.30			0.04	0.03			0.58
		Xylose					3.16	1.05	4.07				0.44			0.19	0.12			0.04
		EtOH														0.83	0.52			6.01
		H <sub>2</sub>																1.47		
		CO <sub>2</sub>														98.53				100.00
		H <sub>2</sub> SO <sub>4</sub>				2.76	2.29	0.51	3.06											
		Enzyme								100.00			0.27							
		Corn liquor																		87.57
		DAP																		12.43
NaOAc															2.72					
Other	26.78	86.45	23.68		4.04	13.36					6.77				0.01			6.78		
CSB1	Corn stover	Flow (ton/day)	500.0	3.3	496.8	2315.2	2811.9	842.2	1972.1	4.7	1200.0	2050.2	3960.0	51.1	5693.9	229.5	4.4	97.5	1957.1	
		Composition (wt%)																		
		Cellulose	43.00	66.04	42.85		7.02	23.44					0.96							1.01
		Hemic.	22.11	33.96	22.03		0.19	0.65					0.27							0.28
		Lignig	18.00		18.12		3.20	10.69					4.39							4.60
		H <sub>2</sub> O				97.24	79.49	53.08	90.79		100.00		79.45	100.00		99.06	51.21			83.23
		Glucose					0.61	0.20	0.78				9.71			0.04	0.06			0.49
		Xylose					4.12	1.37	5.28				0.56			0.24	0.42			0.06
		EtOH														0.66	1.16			5.22
		H <sub>2</sub>																2.07		
		CO <sub>2</sub>														97.93				100.00
		H <sub>2</sub> SO <sub>4</sub>				2.76	2.27	0.51	3.02											
		Enzyme								100.00										
		Corn liquor																		87.57
		DAP																		12.43
NaOAc															9.39					
Other	16.89		17.00		3.10	10.06	0.13				4.66				37.76			5.11		

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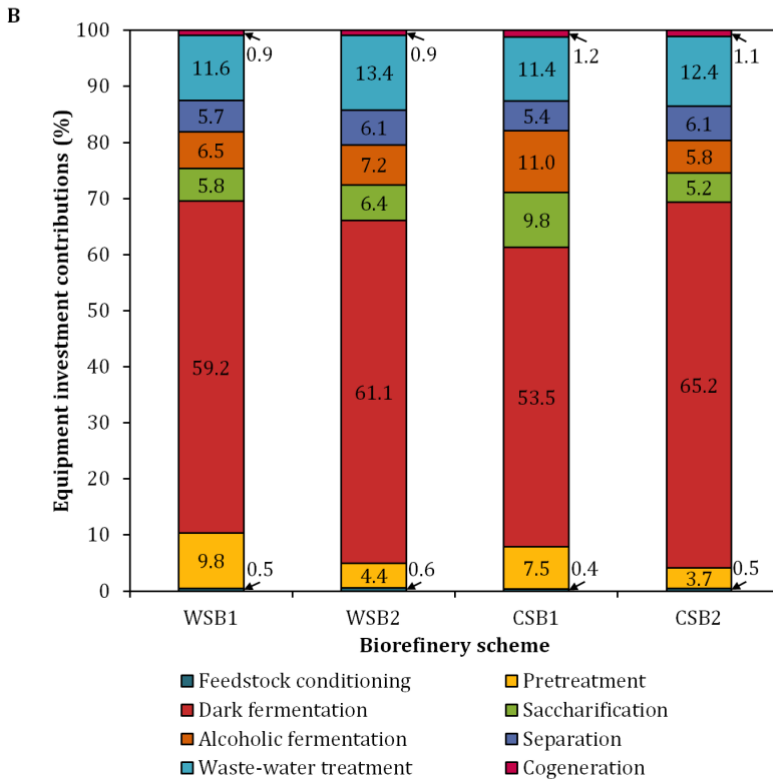
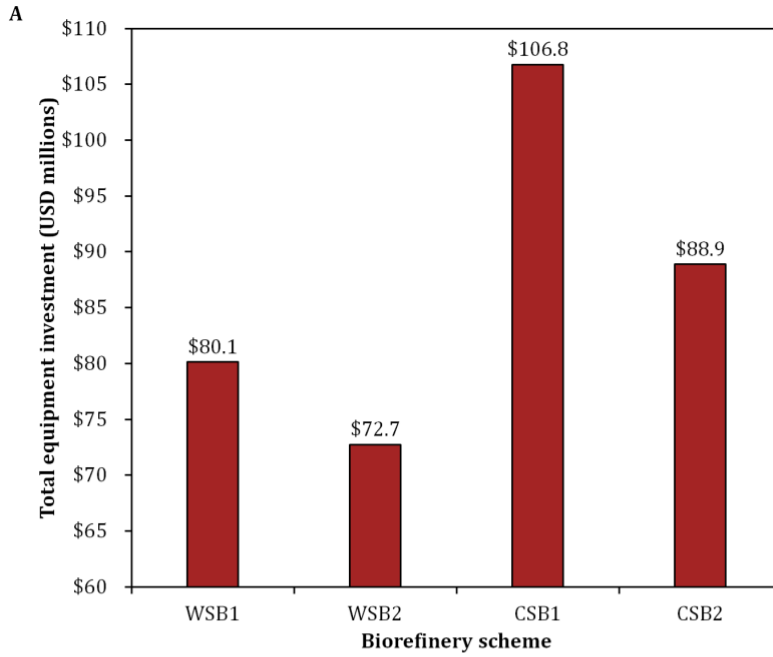
745 **Fig. 4**



Biorefinery Feedstock Stream		F <sub>1</sub>	F <sub>2</sub>	F <sub>3</sub>	F <sub>4</sub>	F <sub>5</sub>	F <sub>6</sub>	F <sub>7</sub>	F <sub>8</sub>	F <sub>9</sub>	F <sub>10</sub>	F <sub>11</sub>	F <sub>12</sub>	F <sub>13</sub>	F <sub>14</sub>	F <sub>15</sub>	F <sub>16</sub>	F <sub>17</sub>	F <sub>18</sub>	F <sub>19</sub>	F <sub>20</sub>	F <sub>21</sub>			
Flow (ton/day)		500.0	24.6	475.4	1920.0	1656.0	449.1	1188.4	16.9	1205.3	591.8	615.9	5.7	1397.2	1994.7	3840.0	39.8	3972.4	454.2	4.4	112.9	1886.3			
		Composition (wt%)																							
WSB2	Wheat straw	Cellulose	48.88	9.93	50.90				20.36		19.67	40.07			1.19									1.26	
		Hemic.	17.83	3.62	18.57				3.07		0.76	1.54			0.46										0.48
		Lignin	6.51		6.85				2.74		2.70	5.50			1.63										1.73
		H <sub>2</sub> O				100.00	100.00			59.96	99.54	59.61	32.78	85.55		100.00	78.58	100.00		98.99	96.68				83.10
		Steam						100.00																	
		XOS							4.08																
		Glucose										0.45	0.06	0.81		11.91				0.01					0.61
		Xylose										7.46	1.02	13.62		0.30				0.19					0.03
		EtOH																		0.80					6.26
		H <sub>2</sub>																	1.51						
		CO <sub>2</sub>																	98.49						100.00
		H <sub>2</sub> SO <sub>4</sub>								0.46															
		Enzyme													100.00										
		Corn liquor																							87.57
DAP																							12.43		
NaOAc																							2.70		
Other		26.78	86.45	23.69				9.48		9.35	19.03	0.01		5.93					0.62				6.53		
Flow (ton/day)		500.0	3.3	496.7	2123.8	1883.8	505.1	1241.9	16.9	1258.7	593.1	668.0	5.0	1365.7	1963.8	5040.0	50.0	5529.3	144.4	4.4	100.4	1867.8			
		Composition (wt%)																							
CSB2	Corn stover	Cellulose	43.00	66.04	42.85				17.14		16.57	35.17			1.06									1.12	
		Hemic.	22.11	33.96	22.03				3.97																4.82
		Lignin	18.00		18.12				7.25		7.15	15.17			4.58										82.78
		H <sub>2</sub> O				100.00	100.00			59.95	99.54	59.25	33.95	86.63		100.00	78.73	100.00		99.11	77.39				
		Steam						100.00																	
		XOS							4.46																
		Glucose										0.38	0.05	0.80		10.64				0.01					0.54
		Xylose										9.37	1.33	11.83		0.40				0.23					0.04
		EtOH																		0.65					5.63
		H <sub>2</sub>																			2.20				
		CO <sub>2</sub>																			97.80				100.00
		H <sub>2</sub> SO <sub>4</sub>								0.46															
		Enzyme													100.00										
		Corn liquor																							87.57
DAP																							12.43		
NaOAc																							15.48		
Other		16.89		17.00				6.81		7.28	14.33	0.74		4.59					7.13				5.07		

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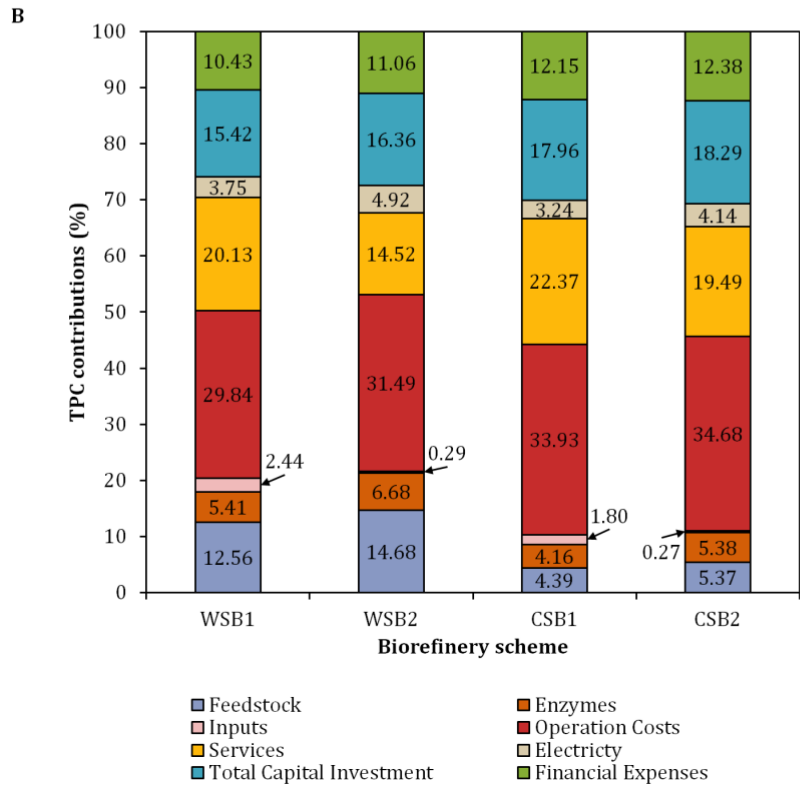
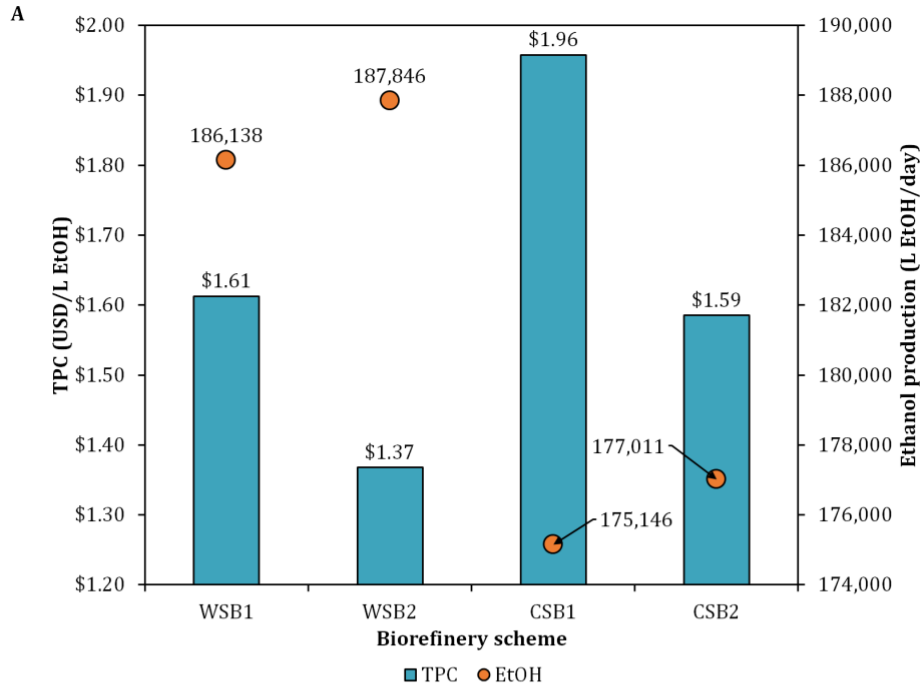
747 Fig. 5



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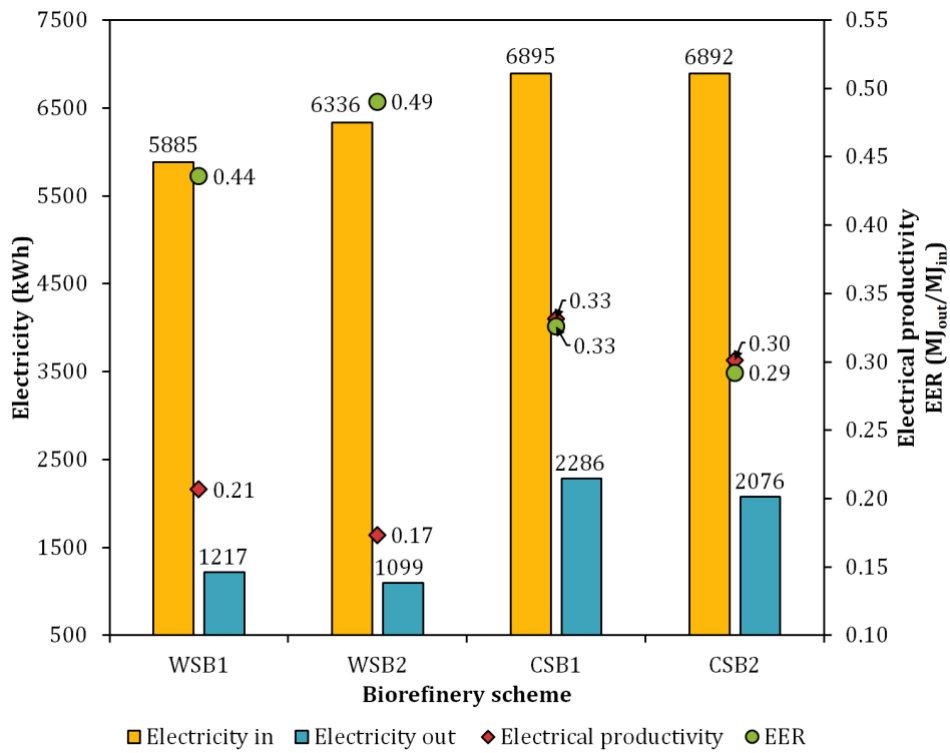
749 **Fig. 6**

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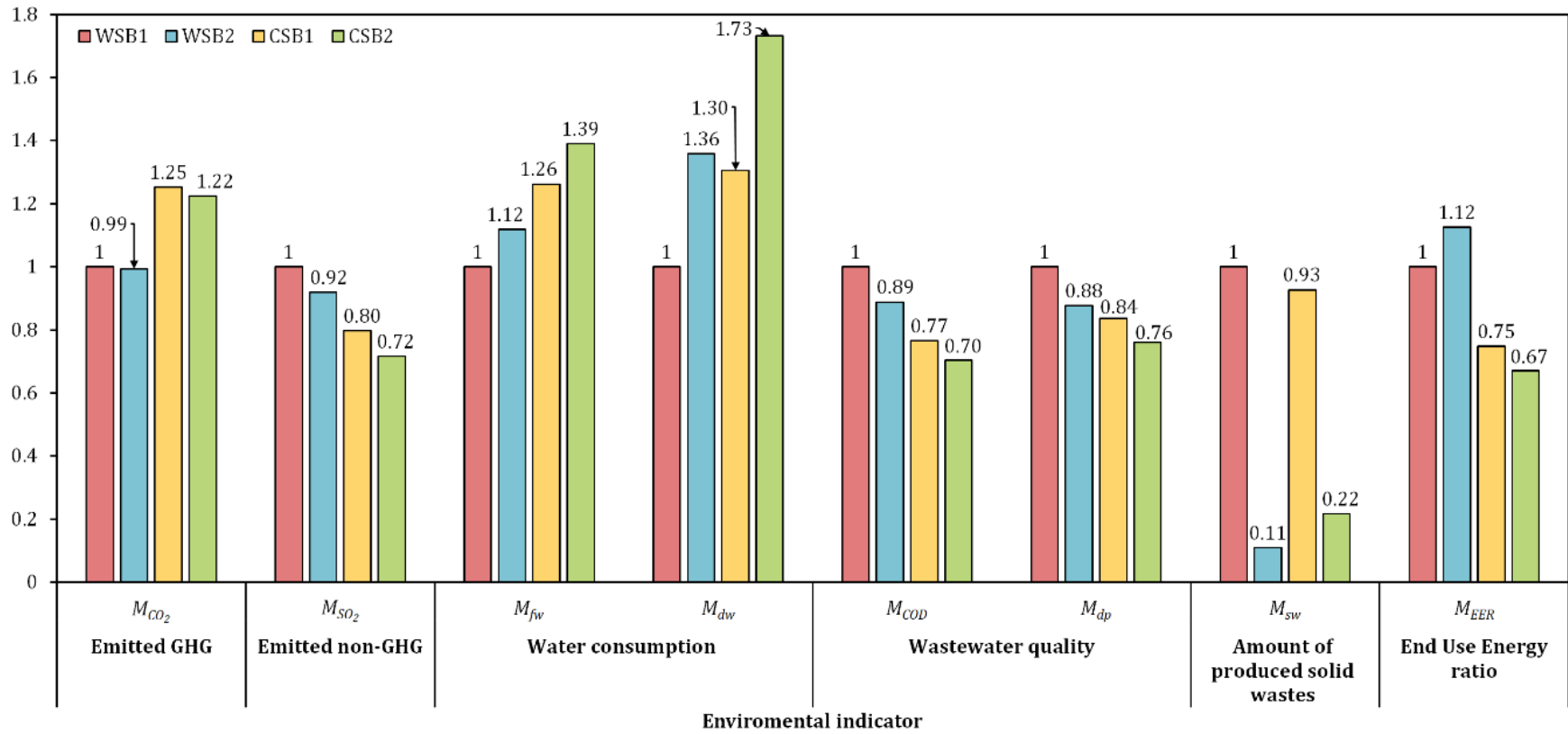
**Fig. 7**



754

755 **Fig. 8**

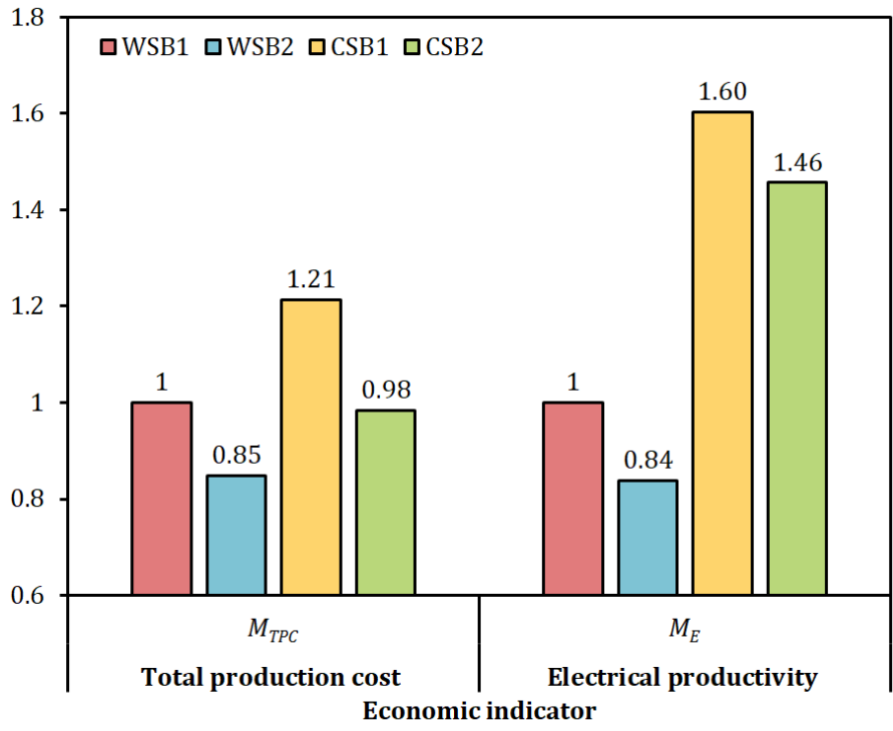
756



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758 **Fig. 9**

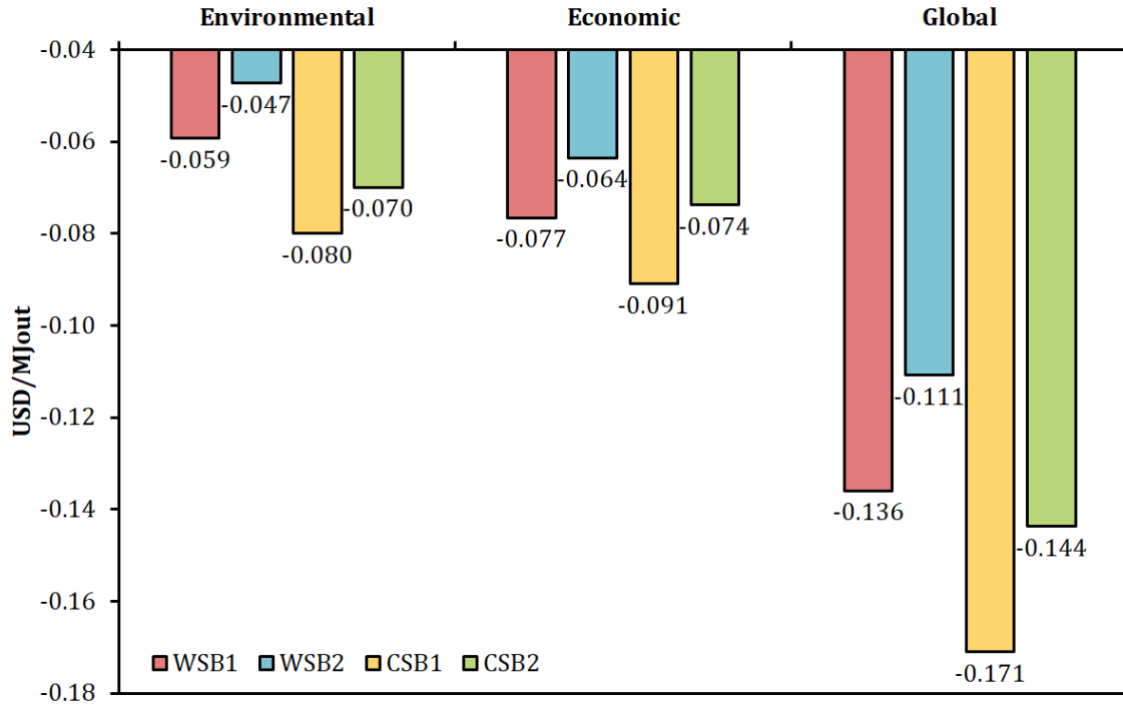
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761 **Fig. 10**

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764 **Fig. 11**