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**Title:** Regulation of pH attenuates toxicity of a byproduct produced by an ethanogenic strain of *Sphingomonas* sp. A1 during ethanol fermentation from alginate

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21 **ABSTRACT**

22 Marine macroalgae is a promising carbon source that contains alginate and mannitol as  
23 major carbohydrates. A bioengineered ethanogenic strain of the bacterium  
24 *Sphingomonas* sp. A1 can produce ethanol from alginate, but not mannitol, whereas the  
25 yeast *Saccharomyces paradoxus* NBRC 0259-3 can produce ethanol from mannitol, but  
26 not alginate. Thus, one practical approach for converting both alginate and mannitol into  
27 ethanol would involve two-step fermentation, in which the ethanogenic bacterium  
28 initially converts alginate into ethanol, and then the yeast produces ethanol from  
29 mannitol. In this study, we found that during fermentation from alginate, the  
30 ethanogenic bacterium lost viability and secreted toxic byproducts into the medium.  
31 These toxic byproducts inhibited bacterial growth and killed bacterial cells, and also  
32 inhibited growth of *S. paradoxus* NBRC 0259-3. We discovered that adjusting the pH of  
33 the culture supernatant or the culture medium containing the toxic byproducts to 6.0  
34 attenuated the toxicity toward both bacteria and yeast, and also extended the period of  
35 viability of the bacterium. Although continuous adjustment of pH to 6.0 failed to  
36 improve the ethanol productivity of this ethanogenic bacterium, this pH adjustment  
37 worked very well in the two-step fermentation due to the attenuation of toxicity toward  
38 *S. paradoxus* NBRC 0259-3. These findings provide information critical for  
39 establishment of a practical system for ethanol production from brown macroalgae.

40

41 **INTRODUCTION**

42 Macroalgae has several advantages as a promising source of biofuels: it is more  
43 productive than land crops; its cultivation requires no arable land, irrigation water, or  
44 fertilizer; and it contains no lignin.<sup>1-5</sup> As a group, the macroalgae include the red, green,  
45 and brown macroalgae. One of the major carbohydrates in brown algae is alginate, a  
46 linear polysaccharide consisting of  $\beta$ -D-mannuronate (M) and its C5 epimer  
47  $\alpha$ -L-guluronate (G), in which the two monosaccharides are arranged as polyM, polyG,  
48 and heteropolymeric random sequences (polyMG).<sup>6</sup> The brown algae *Laminaria*  
49 *japonica* and genera *Sargassum* and *Turbinaria* contain up to 40% w/v alginate.<sup>7,8</sup> The  
50 other major carbohydrate in brown algae is mannitol, a sugar alcohol corresponding to  
51 mannose.<sup>9</sup> *L. japonica* contains up to 30% w/v mannitol.<sup>8</sup> In a review, Zubia *et al.*  
52 reported mannitol content up to 33% w/v in several brown algae of genera *Sargassum*  
53 and *Turbinaria*,<sup>7</sup> and Horn *et al.* reported that the brown alga *Laminaria hyperborea*  
54 contains 25% w/v mannitol.<sup>10</sup>

55 Two systems for producing ethanol from alginate have been established using  
56 bacteria.<sup>11,12</sup> One system utilizes a bioengineered ethanologenic strain of *Sphingomonas*  
57 sp. A1 (ethanologenic strain MK3353), which carries genes for pyruvate decarboxylase  
58 (PDC) and alcohol dehydrogenase (ADH) from *Zymomonas mobilis* and has acquired  
59 the ability to produce ethanol from alginate<sup>11</sup>; however, *Sphingomonas* sp. A1 is unable  
60 to assimilate mannitol.<sup>13</sup> The other system is a bioengineered ethanologenic *Escherichia*  
61 *coli* strain that carries genes for alginate utilization; this strain can produce ethanol from  
62 a mixture of mannitol and alginate derived from brown algae (kombu; *Saccharina*  
63 *japonica*).<sup>12</sup> In addition, we have recently established a system for ethanol production  
64 from mannitol that utilizes the yeast *Saccharomyces paradoxus* NBRC 0259-3, which

65 unlike *Saccharomyces cerevisiae* can naturally assimilate mannitol.<sup>14</sup> Thus, one practical  
66 approach for converting both alginate and mannitol into ethanol would involve two-step  
67 fermentation, in which the ethanogenic strain MK3353 initially converts alginate into  
68 ethanol, and then the yeast produces ethanol from mannitol.

69 In order to establish a practical system for production of ethanol from alginate and  
70 mannitol, it is necessary to achieve greater understanding of ethanol production by the  
71 ethanogenic strain MK3353. In this study, we found that this bacterium secretes toxic  
72 byproducts that inhibit the growth of both itself and *S. paradoxus* NBRC 0259-3, and  
73 kill the bacterium itself, during ethanol fermentation from alginate. We also discovered  
74 that adjusting the media pH to 6.0 attenuates this toxicity and enhances the two-step  
75 fermentation process.

76

## 77 RESULTS

78 **Process of ethanol production from alginate.** In order to understand the process of  
79 ethanol production from alginate, the ethanogenic strain MK3353 (*Sphingomonas* sp.  
80 A1 lacking in lactate dehydrogenase [LDH] gene and carrying genes for PDC and ADH  
81 in the broad-host range plasmid pKS13)<sup>11</sup> and the control strain MK3567  
82 (*Sphingomonas* sp. A1 lacking in LDH gene and carrying pKS13 alone) were cultivated  
83 in liquid alginate medium containing 5% w/v alginate at 30°C and 95 strokes per  
84 minutes (spm). Growth ( $A_{600}$  of the culture), viability (cfu), ethanol concentration in the  
85 supernatant (hereafter, we refer to supernatant from a culture of *Sphingomonas* sp. A1 as  
86 “A1-supernatant”), and alginate concentration in the culture were determined (Fig. 1).

87 The ethanogenic strain MK3353 started to produce a large quantity of ethanol  
88 after 1 day of cultivation, as reported,<sup>11</sup> but the control strain MK3567 not. Ethanol

89 concentration reached a maximum after 3 days of cultivation, and did not increase  
90 thereafter (Fig. 1B). Accordingly, a larger amount of alginate was consumed by the  
91 ethanogenic strain MK3353 than the control strain MK3567, indicating that  
92 introduction of the genes encoding PDC and ADH improved alginate metabolism. The  
93 ethanogenic strain MK3353 consumed almost all alginate after 3 days of cultivation  
94 (Fig. 1C). The pH of the culture of the ethanogenic strain MK3353 became slightly  
95 alkaline (from pH 8.0 at day 0 to pH 8.7 at day 3 and pH 9.0 at day 4), whereas that of  
96 the control MK3567 strain was not (from pH 8.0 at day 0 to pH 6.3 at day 1 and pH 7.2  
97 at day 4) (data not shown). Notably, the ethanogenic strain MK3353 started to lose  
98 viability after 2 days of cultivation, and completely lost viability at day 5. We attributed  
99 the loss of ethanol-production capacity to this loss of cell viability. No remarkable  
100 morphological change was observed by transmission electron microscopy (TEM) or  
101 scanning electron microscopy (SEM) analyses of the ethanogenic strain MK3353 or  
102 the control strain MK3567 after 1, 2, or 3 days of cultivation (Supplementary Fig. S1).

103

104 **Toxic byproducts produced by the ethanogenic strain MK3353.** The data described  
105 above suggested that the loss of ethanol-production ability and the decrease in viability  
106 were caused by complete consumption of alginate. However, this possibility was ruled  
107 out by the observation that addition of alginate (1 g/day) each day after 2 days of  
108 cultivation had no effect on ethanol concentration or viability of the ethanogenic strain  
109 MK3353 (data not shown).

110 Another possibility is that ethanol itself caused the loss of cell viability. However,  
111 when the ethanogenic strain MK3353 was cultivated in liquid alginate medium  
112 containing 5% w/v alginate and an initial concentration of 1.0% w/v (10 g/l) ethanol,

113 there was no loss of viability, and ethanol was still produced from alginate, although the  
114 added ethanol did delay cell growth (Supplementary Fig. S2). Thus, the reduction in  
115 viability could not be attributed to the ethanol produced by the ethanogenic strain  
116 MK3353.

117 We then hypothesized that some toxic byproduct other than ethanol might be  
118 produced by the ethanogenic strain MK3353, but not the control strain MK3567. To  
119 test this idea, we examined the growth of the ethanogenic strain MK3353 in liquid  
120 alginate media containing 0.4% w/v alginate and 0, 10, 25, or 50% v/v of  
121 A1-supernatant from 1-, 2-, 3-, or 4-day culture of the ethanogenic strain MK3353 or  
122 the control strain MK3567 (Fig. 2A). Growth inhibition was observed in the presence of  
123 A1-supernatants from 2-, 3-, and 4-day cultures of the ethanogenic strain MK3353,  
124 but not A1-supernatant from 1-day culture of the ethanogenic strain MK3353 strain or  
125 A1-supernatants from any cultures of the control strain MK3567. Moreover, the  
126 inhibitory effect was dependent on the concentration of A1-supernatant. Collectively,  
127 these data indicate that toxic byproducts were produced specifically by the  
128 ethanogenic strain MK3353 (i.e., in a manner dependent on the presence of the genes  
129 encoding PDC and ADH), and that the concentration of these byproducts increased over  
130 the cultivation period.

131 A1-supernatant from 4-day culture of the ethanogenic strain MK3353 killed the  
132 bacterial cells, whereas A1-supernatant from 4-day culture of the control strain MK3567  
133 did not (Fig. 2B). Inhibition of the growth of *S. paradoxus* NBRC 0259-3 was also  
134 observed in the presence of A1-supernatant from 3-day culture of the ethanogenic  
135 strain MK3353, but not in the presence of A1-supernatant from 3-day culture of the  
136 control strain MK3567 (Fig. 2C); however, A1-supernatant of 3-day culture of the

137 ethanogenic strain MK3353 did not kill *S. paradoxus* NBRC 0259-3 cells (Fig. 2D).

138

139 **Attenuation of the toxic effects of byproducts secreted by the ethanogenic strain**

140 **MK3353.** We examined the effect of pH on the toxic effects of A1-supernatant from

141 cultures of the ethanogenic strain MK3353. The pH of A1-supernatant from 3-day

142 culture was approximately 8.7. We adjusted the pH to 4.0, 5.0, 6.0, 7.0, and 8.0; all

143 samples were adjusted to the same final volume by addition of sterilized water, as

144 necessary, to rule out effects of dilution. As shown in Fig. 3A, growth inhibition of the

145 ethanogenic strain MK3353 was not observed in the presence of A1-supernatant at pH

146 of 5.0 or 6.0. Moreover, the killing effect of A1-supernatant at pH 6.0 was much weaker

147 than that of A1-supernatant at pH 5.0 (Fig. 3B). Growth inhibition of *S. paradoxus*

148 NBRC 0259-3 was also not detected in the presence of A1-supernatants at pH of 6.0 (or

149 at 4.0 and 5.0) (Fig. 3C). These data indicate that adjustment of pH to 6.0 attenuated the

150 toxicity of the byproducts in A1-supernatants.

151

152 **Ethanol production by the ethanogenic strain MK3353 with continuous**

153 **adjustment of pH to 6.0.** The data described above led us to predict that continuous

154 adjustment of culture pH to 6.0 would improve ethanol productivity from alginate. To

155 test this prediction, we performed ethanol fermentation in liquid alginate medium

156 containing 5% w/v alginate (25 ml) using the ethanogenic strain MK3353. The pH of

157 the culture was adjusted to pH 4.0, 5.0, 6.0, 7.0, or 8.0 every day (Fig. 4A). As expected,

158 adjustment of pH to 6.0 improved cell viability (Fig. 4B); however, ethanol production

159 was not improved by this adjustment (Fig. 4C), indicating that the cells were viable for

160 a longer period at pH 6.0, but still lost the ability to produce ethanol from alginate.

161 Addition of alginate (0.625 g or 1.25 g) after 3 days of cultivation had no effect on  
162 ethanol concentration or cell viability of the ethanologenic strain MK3353 (data not  
163 shown). Therefore, we speculate that although the toxic effects of byproducts in the  
164 culture were attenuated by pH adjustment, these byproducts could still inhibit the  
165 reactions involved in production of ethanol from alginate.

166

167 **Effects of pH adjustment on two-step fermentation.** Finally, we examined the effect  
168 of adjusting pH to 6.0 on the two-step fermentation that we demonstrated previously.<sup>14</sup>  
169 In our earlier study, we adjusted the pH of A1-supernatant of 3-day culture to 5.8, the  
170 same as that of the yeast extract/peptone (YP) that was added to A1-supernatant to  
171 support the growth of yeast.<sup>14</sup> In this study, we adjusted the pH of A1-supernatant of  
172 3-day culture to 6.0, based on the results described above, and prepared YPM-A1  
173 medium (final pH 6.1) by mixing 22.5 ml of A1-supernatant adjusted to pH 6.0, 2.5 ml  
174 of 10-fold concentrated YP (pH 5.6 in this study), and 0.5 g mannitol. We also prepared  
175 YPM-A1 medium (final pH 7.3) by mixing 22.5 ml of A1-supernatant without pH  
176 adjustment (pH 8.7), 2.5 ml of 10-fold concentrated YP (pH 5.6), and 0.5 g mannitol.  
177 The final mannitol concentrations in both media were 2% w/v. As controls, we also  
178 prepared YPM (pH 6.1) and YPM (pH 7.3).

179 We cultivated *S. paradoxus* NBRC 0259-3 in these four media and monitored  
180 ethanol production from mannitol (Fig. 5). As expected, *S. paradoxus* NBRC 0259-3  
181 grew and produced ethanol in YPM-A1 medium (pH 6.1), but not at all in YPM-A1  
182 medium (pH 7.3) (Fig. 5). Because *S. paradoxus* NBRC 0259-3 grew similarly in YPM  
183 (pH 6.1) and YPM (pH 7.3), and produced only slightly less ethanol in YPM (pH 7.3)  
184 than in YPM (pH 6.1) (Fig. 5), we concluded that the severe difference in growth and



185 ethanol production between YPM-A1 medium (pH 6.1) and YPM-A1 medium (pH 7.3)  
186 was not due to the pH difference alone, but was rather largely due to the attenuation of  
187 toxicity resulting from the pH adjustment. These observations demonstrate that  
188 adjustment of pH to 6.0 is also very effective in the two-step fermentation to produce  
189 ethanol from alginate and mannitol.

190

## 191 **DISCUSSION**

192 Due to the huge potential availability of marine macroalgae, and hence of alginate and  
193 mannitol, a system for production of ethanol from these carbohydrates would be of  
194 great value. Despite the importance of this goal, however, only two systems for  
195 production of ethanol from alginate have been established to date: one using engineered  
196 *Sphingomonas* sp. A1, and another using engineered *E. coli*.<sup>11, 12</sup>

197 The ethanogenic strain MK3353, a bioengineered strain of *Sphingomonas* sp.  
198 A1, carries genes for PDC and ADH from *Z. mobilis*.<sup>11</sup> In this strain, the PDC and ADH  
199 genes are controlled by a potent intrinsic promoter identified using DNA microarrays. In  
200 addition, the gene for LDH was deleted from the genome, because when this gene is  
201 present, lactate accumulates as the main byproduct of ethanol production.<sup>11</sup>  
202 Consequently, the ethanogenic strain MK3353 has acquired the capacity to produce as  
203 much as 13 g/L ethanol from 60 g/L alginate after a 72 h-fermentation.<sup>11</sup> However, as  
204 noted above, this bacterium is unable to assimilate mannitol.<sup>13</sup> To construct the  
205 engineered *E. coli* strain BAL1611, (i) an alginate-lyase secretion system, (ii) the genes  
206 for alginate degradation, transport, and metabolism, and (iii) the genes for PDC and  
207 ADH from *Z. mobilis* were introduced into an *E. coli* strain in which several genes  
208 (*pf1B-focA*, *frdABCD*, and *ldhA*) had been deleted from the genome. Due to the intrinsic

209 ability of *E. coli* to assimilate glucose and mannitol, the resulting engineered strain is  
210 able to produce 35–41 g/L ethanol after 150 h-fermentation from extracts (total mass,  
211 140 g) of *S. japonica* containing various carbohydrates such as alginate, mannitol, and  
212 glucose.<sup>12</sup>

213 Several bacterial species and strains are capable of producing ethanol from  
214 mannitol: the bacteria *Zymobacter palmae* (13 g/L ethanol from 38 g/L mannitol after  
215 70 h-fermentation), *E. coli* KO11 (25.8 g/L ethanol from 90 g/L mannitol after 120  
216 h-fermentation), and the aforementioned *E. coli* strain (BAL1611).<sup>9, 12, 15</sup> Ethanol has  
217 been produced from mannitol by some yeast strains, e.g., the *S. cerevisiae* polyploid  
218 strain BB1 (5 g/L ethanol from 50 g/L mannitol after 60 h-fermentation) and *Pichia*  
219 *angophorae* (10 g/L ethanol from 38 g/L mannitol after 75 h-fermentation).<sup>10, 16</sup> By  
220 contrast, however, other *S. cerevisiae* strains, e.g., polyploid BB2, haploid S288C, and  
221 haploid *Sc41 YJO*, are unable to assimilate mannitol for growth.<sup>16, 17</sup> Recently, we  
222 demonstrated that the *S. paradoxus* strain NBRC 0259-3 is more suitable for the  
223 production of ethanol from mannitol than *P. angophorae* and *E. coli* KO11.<sup>14</sup> *S.*  
224 *paradoxus* strain NBRC 0259-3 was derived from the original NBRC 0259 strain by  
225 cultivation for 3 days in medium containing mannitol, resulting in acquisition of higher  
226 capacity to produce ethanol from alginate: *S. paradoxus* NBRC 0259-3 strain produced  
227 40 g/L ethanol from 100 g/L mannitol after an 11-day fermentation.<sup>14</sup> We also  
228 succeeded in a two-step fermentation in which the ethanologenic strain MK3353  
229 initially converts alginate into ethanol, and then the yeast *S. paradoxus* NBRC 0259-3  
230 produces ethanol from mannitol. In that system, we adjusted the pH of A1-supernatant  
231 of 3-day culture to 5.8, the same as that of the YP added to A1-supernatant to support  
232 growth of the yeast.<sup>14</sup>

233 In order to establish a practical system for production of ethanol from alginate as  
234 well as mannitol, it is necessary to achieve greater understanding of ethanol production  
235 by these systems. In this study, we found that during fermentation of the ethanologenic  
236 *Sphingomonas* sp. A1 strain MK3353 from alginate, the culture became slightly alkaline,  
237 and the bacterium secreted toxic byproducts that inhibited the growth of both itself and  
238 *S. paradoxus* NBRC 0259-3, and also killed the bacterial cells. However, we discovered  
239 that this toxicity could be attenuated by adjusting the pH of toxic culture supernatant or  
240 culture medium to 6.0. This worked very well in the two-step fermentation, due to the  
241 reduction in toxicity toward *S. paradoxus* NBRC 0259-3. To our knowledge, this is the  
242 first report of the formation of toxic byproducts during ethanol fermentation from  
243 alginate.

244 In ethanol production from lignocellulosic biomass, inhibitory compounds  
245 including aldehyde inhibitors, ketone inhibitors, organic acid inhibitors, and  
246 phenol-based inhibitors are generated during the thermo-chemical pre-treatment of the  
247 biomass.<sup>18</sup> Several detoxification methods have been described, including physical  
248 treatments (evaporation and use of membranes), physicochemical treatments (ion  
249 exchange resins, neutralization, overliming, use of activated charcoal, and extraction  
250 with organic solvents), and biological treatments (use of enzymes and  
251 microorganisms).<sup>19</sup> In our ethanol fermentation from alginate, we used commercially  
252 supplied sodium alginate, and therefore did not need to pretreat brown macroalgal  
253 biomass. Because generation of toxic compounds was dependent on the presence of the  
254 PDC and ADH genes, it is possible that the toxic compounds were derived from the  
255 metabolic conversion of alginate to ethanol (Fig. 1).

256 Although the toxic compounds derived from ethanol fermentation from alginate

257 remain to be identified, our findings provide a rationale for the success of our previous  
258 two-step fermentation method, in which we adjusted the pH of A1-supernatant of 3-day  
259 culture to 5.8, the same as that of YP.<sup>14</sup> Furthermore, because yeasts generally prefer  
260 acidic conditions (as shown in Fig. 5), and acidic conditions generally prevent bacterial  
261 contamination, attenuation of the toxicity of A1-supernatant by adjustment of the pH to  
262 6.0 should be beneficial in the context of practical two-step fermentation. Therefore,  
263 such an approach should aid in the establishment of a practical system for ethanol  
264 production from brown macroalgae, a promising carbon source for bioethanol.

265

## 266 MATERIALS AND METHODS

267 **Strains and cultivation.** The ethanologenic *Sphingomonas* sp. A1 strain MK3353, that  
268 was previously called EPv104, lacks the LDH gene and carries the *Z. mobilis* genes for  
269 PDC and ADH on a broad-host range plasmid, pKS13.<sup>11</sup> The control *Sphingomonas* sp.  
270 A1 strain, MK3567, is also lacks in LDH gene, but carries pKS13 alone (i.e., without  
271 the *Z. mobilis* genes). *Sphingomonas* sp. A1 was transformed by triparental mating with  
272 *E. coli* DH5 $\alpha$  carrying pRK2013 as a helper.<sup>20</sup> *S. paradoxus* NBRC 0259-3 is a  
273 derivative of the original *S. paradoxus* strain NBRC 0259, which can naturally  
274 assimilate mannitol.<sup>14</sup> *S. paradoxus* NBRC 0259-3 has a higher capacity than the  
275 parental strain for production of ethanol from mannitol.<sup>14</sup>

276 Algininate medium consists of sodium alginate from brown algae [average molecular  
277 weight (MW), 300 kDa; ratio of mannuronate to guluronate, 3:1; Nacalai Tesque], 0.1%  
278 w/v (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, 0.1% w/v KH<sub>2</sub>PO<sub>4</sub>, 0.1% w/v Na<sub>2</sub>HPO<sub>4</sub>, 0.01% w/v MgSO<sub>4</sub>·7H<sub>2</sub>O, and  
279 0.01% w/v yeast extract (pH 8.0).<sup>11</sup> Antibiotics [20 mg/l tetracycline (Tet) and 25 mg/l  
280 kanamycin (Kan)] were included in media for cultivation of the ethanologenic strain

281 MK3353 and the control strain MK3567. For solid medium, 1.5% w/v agar and 0.5%  
282 w/v alginate were included. For liquid media, 0.4, 0.8, or 5% w/v alginate was included.  
283 For ethanol production, fresh cells of the ethanologenic strain MK3353 grown on  
284 alginate solid medium were inoculated into liquid medium containing 0.8% w/v alginate  
285 and precultured at 30°C for 24 h at 145 spm. Cells in the preculture were inoculated into  
286 liquid alginate medium containing 5% w/v alginate (100 and 25 ml media in a 300- and  
287 100-ml Erlenmeyer flasks, respectively), at an initial  $A_{600}$  of 0.1, and subsequently  
288 cultivated at 30°C and 95 spm.<sup>11</sup> After 3 days of cultivation, the culture was centrifuged  
289 at 20,000×g for 10 min, and the resultant supernatant was referred to as  
290 “A1-supernatant”. When necessary, the pH of A1-supernatant was adjusted.

291 YP medium consisted of 1% w/v yeast extract and 2% w/v tryptone (pH 5.6). For  
292 YPM medium, YP medium was supplemented with 2% w/v mannitol. For solid medium,  
293 2% w/v agar was included. YP medium was sterilized by autoclaving prior to addition  
294 of carbon sources. Ten-fold concentrated YP (pH 5.6) was sterilized by passage through  
295 a filter with 0.2- $\mu$ m pores. YPM-A1 medium consisted of 22.5 ml of A1-supernatant  
296 (pH-adjusted if necessary), 2.5 ml 10-fold concentrated YP (pH 5.6), and 0.5 g mannitol.  
297 For ethanol production, *S. paradoxus* NBRC 0259-3 was grown as reported  
298 previously.<sup>14</sup> Yeast supernatant was obtained by centrifugation of the culture at  
299 20,000×g at 4°C for 5 min.

300

301 **Analytical methods.** Concentration of ethanol in A1-supernatant or yeast supernatant  
302 was determined using assay kits (Roche) according to the manufacturer’s instructions.  
303 A standard curve was prepared for each assay using an ethanol standard solution.  
304 Alginate concentration in the culture was determined by the carbazole–sulfuric acid

305 method, using sodium alginate as a standard.<sup>21</sup> To determine colony-forming units (cfu),  
306 cultures were diluted in 1× P solution (1.1 mM KH<sub>2</sub>PO<sub>4</sub> plus 1.1 mM Na<sub>2</sub>HPO<sub>4</sub>), and  
307 streaked on solid alginate medium containing 20 mg/l Tet and 25 mg/l Kan.

308

309 **TEM.** The culture was mixed with an equal volume of 100 mM sodium and potassium  
310 phosphate, pH 7.4 (PB) plus 4% paraformaldehyde and 4% glutaraldehyde, and then  
311 incubated at 4°C for 1 h. After centrifugation at 2,000×g for 2.5 min, the collected cells  
312 were again suspended in PB plus 2% glutaraldehyde and fixed overnight. The cells were  
313 rinsed three times with PB, followed by post-fixation with 2% osmium tetroxide in PB.  
314 The fixed cells were dehydrated with ethanol, infiltrated with propylene oxide, placed  
315 into a 7:3 mixture of propylene oxide and Quetol-812 (Nisshin EM, Tokyo, Japan), and  
316 incubated overnight with the lid open to volatilize propylene oxide. The cells were then  
317 transferred to 100% resin and polymerized at 60°C for 48 h. Ultra-thin sections  
318 (approximately 70 nm thick) were cut with a diamond knife using an Ultracut UCT  
319 ultramicrotome (Leica). Sections were placed on copper grids and stained with 2%  
320 uranyl acetate, followed by lead staining (Sigma). The sections were examined using a  
321 JEM-1200EX microscope (JEOL, Tokyo, Japan) at 80 kV.

322

323 **SEM.** Cells were fixed and dehydrated as for TEM. The dehydrated cells were  
324 substituted with tert-butyl alcohol and vacuum-dried using a DAP-6D dry vacuum  
325 pump (Ulvac Kiko) with slow decompression. After drying, the samples were coated  
326 with a thin layer (30 nm) of osmium using an NL-OPC80NS plasma coater (Nippon  
327 Laser & Electronics Laboratory). The samples were observed using a JSM-6340F  
328 scanning electron microscope (JEOL) at an electron voltage of 10.0 kV.

329

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333

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335   **REFERENCES**

- 336   1.     Huesemann M, Roesjadi G, Benemann J, Metting FB. Biofuels from microalgae  
337         and seaweeds. In: Vertès A, Qureshi N, Yukawa H, Blaschek HP, eds. Biomass to  
338         biofuels: strategies for global industries: Wiley, 2010:165-84.
- 339   2.     Adams JM, Gallagher JA, Donnison IS. Fermentation study on *Saccharina*  
340         *latisima* for bioethanol production considering variable pre-treatments. J Appl  
341         Phycol 2009; 21:569-74.
- 342   3.     Yoon JJ, Kim YJ, Kim SH, Ryu HJ, Choi JY, Kim GS, et al. Production of  
343         polysaccharides and corresponding sugars from red seaweed. Adv Mat Res  
344         2010; 93-94:463-6.
- 345   4.     John RP, Anisha GS, Nampoothiri KM, Pandey A. Micro and macroalgal  
346         biomass: a renewable source for bioethanol. Bioresour Technol 2011;  
347         102:186-93.
- 348   5.     Yanagisawa M, Kawai S, Murata K. Strategies for the production of high  
349         concentrations of bioethanol from seaweeds: production of high concentrations  
350         of bioethanol from seaweeds. Bioengineered 2013; 4:224-35.
- 351   6.     Gacesa P. Alginates. Carbohydr Polym 1988; 8:161-82.
- 352   7.     Zubia M, Payri C, Deslandes E. Alginate, mannitol, phenolic compounds and  
353         biological activities of two range-extending brown algae, *Sargassum*  
354         *mangarevense* and *Turbinaria ornata* (Phaeophyta: Fucales), from Tahiti (French  
355         Polynesia). J Appl Phycol 2008; 20:1033-43.
- 356   8.     Honya M, Kinoshita T, Ishikawa M, Mori H, Nisizawa K. Monthly  
357         determination of alginate, M/G ratio, mannitol, and minerals in cultivated  
358         *Laminaria Japonica*. Nippon Suisan Gakkaishi 1993; 59:295-9.



- 359 9. Horn SJ, Aasen IM, Østgaard K. Production of ethanol from mannitol by  
360 *Zymobacter palmae*. J Ind Microbiol Biotechnol 2000; 24:51-7.
- 361 10. Horn SJ, Aasen IM, Ostgaard K. Ethanol production from seaweed extract. J Ind  
362 Microbiol Biotechnol 2000; 25:249-54.
- 363 11. Takeda H, Yoneyama F, Kawai S, Hashimoto W, Murata K. Bioethanol  
364 production from marine biomass alginate by genetically engineered bacteria.  
365 Energy Environ Sci 2011; 4:2575-81
- 366 12. Wargacki AJ, Leonard E, Win MN, Regitsky DD, Santos CN, Kim PB, et al. An  
367 engineered microbial platform for direct biofuel production from brown  
368 macroalgae. Science 2012; 335:308-13.
- 369 13. Hisano T, Yonemoto Y, Yamashita T, Fukuda Y, Kimura A, Murata K. Direct  
370 uptake of alginate molecules through a pit on the bacterial-cell surface - a novel  
371 mechanism for the uptake of macromolecules. J Ferment Bioeng 1995;  
372 79:538-44.
- 373 14. Ota A, Kawai S, Oda H, Iohara K, Murata K. Production of ethanol from  
374 mannitol by the yeast strain *Saccharomyces paradoxus* NBRC 0259. J Biosci  
375 Bioeng 2013; 116:327-32.
- 376 15. Kim NJ, Li H, Jung K, Chang HN, Lee PC. Ethanol production from marine  
377 algal hydrolysates using *Escherichia coli* KO11. Bioresour Technol 2011;  
378 102:7466-9.
- 379 16. Quain DE, Boulton CA. Growth and metabolism of mannitol by strains of  
380 *Saccharomyces cerevisiae*. J Gen Microbiol 1987; 133:1675-84.
- 381 17. Perfect JR, Rude TH, Wong B, Flynn T, Chaturvedi V, Niehaus W. Identification  
382 of a *Cryptococcus neoformans* gene that directs expression of the cryptic

- 383           *Saccharomyces cerevisiae* mannitol dehydrogenase gene. J Bacteriol 1996;  
384           178:5257-62.
- 385   18.   Liu ZL, Blaschek HP. Biomass conversion inhibitors and *in situ* detoxification.  
386           In: Vertès A, Qureshi N, Yukawa H, Blaschek HP, eds. Biomass to biofuels:  
387           strategies for global industries: Wiley, 2010:233-59.
- 388   19.   Canilha L, Chandel AK, Milessi TSD, Antunes FAF, Freitas WLD, Felipe MDA,  
389           et al. Bioconversion of sugarcane biomass into ethanol: an overview about  
390           composition, pretreatment methods, detoxification of hydrolysates, enzymatic  
391           saccharification, and ethanol fermentation. J Biomed Biotechnol 2012.
- 392   20.   Ditta G, Stanfield S, Corbin D, Helinski DR. Broad host range DNA cloning  
393           system for gram-negative bacteria: construction of a gene bank of *Rhizobium*  
394           *meliloti*. Proc Natl Acad Sci U S A 1980; 77:7347-51.
- 395   21.   Galambos JT. The reaction of carbazole with carbohydrates. I. Effect of borate  
396           and sulfamate on the carbazole color of sugars. Anal Biochem 1967; 19:119-32.
- 397
- 398
- 399

400 **FIGURE LEGENDS**

401 **Figure. 1.** Process of ethanol production from alginate. The ethanogenic strain  
402 MK3353 (closed symbol) and the control strain MK3567 (open symbol) were  
403 precultured, inoculated, and cultivated as described in MATERIALS AND METHODS  
404 in liquid alginate medium (100 ml) containing 5% w/v alginate at 30°C and 95 spm.  
405 Growth ( $A_{600}$  of the culture) (A), ethanol concentration in A1-supernatant (B), alginate  
406 concentration in the culture (C), and number of viable cells per 10  $\mu$ l culture (cfu) (D)  
407 were determined. Means and standard deviation (SD) of three independent experiments  
408 are shown.

409

410 **Figure. 2.** Growth-inhibitory and killing effects of A1-supernatants. (A)  
411 Growth-inhibitory effects of A1-supernatants on the ethanogenic strain MK3353.  
412 MK3353 was precultured as described in MATERIALS AND METHODS; inoculated  
413 into liquid alginate medium (1.0 ml) containing 0.4% w/v alginate plus 0, 10, 25, or  
414 50% v/v of A1-supernatant from 1- (black bar), 2- (hatched bar), 3- (gray bar), or 4-day  
415 (white bar) culture of MK3353; grown for 24 h at 145 spm and 30°C; and then the  $A_{600}$   
416 was measured. (B) Killing effects of A1-supernatant on MK3353. MK3353 was  
417 cultured as in (A), except that the bacteria were grown for 8 h in media containing 0%  
418 (open symbol) or 50% (closed symbol) v/v of A1-supernatant from 4-day culture of  
419 MK3353. During cultivation, viability was determined as described in MATERIALS  
420 AND METHODS. (C) Growth-inhibitory effect of A1-supernatant on *S. paradoxus*  
421 NBRC 0259-3. Fresh *S. paradoxus* NBRC 0259-3 cells on YPM solid medium were  
422 suspended in sterilized water (SDW) and inoculated into YPM (1.0 ml) containing 0, 10,  
423 25, 50, or 90% v/v of A1-supernatant from 3-day culture of MK3353 or MK3567. The

424 culture was grown for 24 h at 145 spm and 30°C, and then  $A_{600}$  was measured. (D)  
425 Effect of A1-supernatant on viability of *S. paradoxus* NBRC 0259-3. *S. paradoxus*  
426 NBRC 0259-3 was cultured as in (C), except that *S. paradoxus* NBRC 0259-3 was  
427 grown for 8 h in the media containing 0% (open symbol) or 50% (closed symbol) v/v of  
428 A1-supernatant from 3-day culture of MK3353. During cultivation, viability was  
429 determined as described in MATERIALS AND METHODS. (A–D) Means and SD of  
430 three independent experiments are shown.

431

432 **Figure 3.** pH adjustment attenuates the growth-inhibitory and killing effects of  
433 A1-supernatant. (A) Attenuation of growth-inhibitory effect of A1-supernatant on the  
434 ethanogenic strain MK3353. MK3353 was cultured as described for Fig. 2A, but in  
435 liquid alginate medium containing 0.4% w/v alginate plus 0% (cont.) or 50% v/v of  
436 A1-supernatant from 4-day culture of MK3353, with pH adjusted with HCl as indicated,  
437 and then  $A_{600}$  was measured. (B) Attenuation of the killing effect of A1-supernatant on  
438 MK3353. MK3353 was cultured as described in (A); but with pH adjusted to 5.0 or 6.0  
439 with HCl, or not adjusted (pH 8.7), in 25 ml liquid medium at 95 spm and 30°C. After  
440 24 h of cultivation, viability was determined. (C) Attenuation of growth-inhibitory  
441 effect of A1-supernatant on *S. paradoxus* NBRC 0259-3. Fresh *S. paradoxus* NBRC  
442 0259-3 cells on YPM solid medium were cultured as in Fig. 2C; but in YPM (1.0 ml)  
443 containing 0% (cont.) or 50% v/v of A1-supernatant from 3-day culture of MK3353,  
444 with pH adjusted with HCl as indicated, and then  $A_{600}$  was measured. (A–C) Means and  
445 SD of three independent experiments are shown.

446

447 **Figure 4.** Effect of continuous pH adjustment on ethanol production from alginate. (A)

448 pH profile of the culture. The ethanologenic strain MK3353 was grown in liquid  
449 alginate medium (25 ml) containing 5% w/v alginate at 95 spm and 30°C, as described  
450 in MATERIALS AND METHODS. The pH of the culture was adjusted with HCl or  
451 NaOH as indicated (open triangle, without pH adjustment; open square, pH 4.0; open  
452 diamond, pH 5.0; closed triangle, pH 6.0; closed square, pH 7.0; closed diamond, pH  
453 8.0) every 24 h. (B) Viability of the cells in the cultures. (C) Ethanol concentrations in  
454 the cultures. (A–C) Means and SD of three independent experiments are shown.

455

456 **Figure 5.** Effect of pH adjustment on the two-step fermentation. *S. paradoxus* NBRC  
457 0259-3 maintained on solid YPM medium was precultured in 50 ml YPM liquid  
458 medium at 30°C in a 100-ml Erlenmeyer flask on a shaker (Personal Lt-10F) at 95  
459 spm.<sup>14</sup> After 1 day of cultivation, cells were collected, washed once with SDW,  
460 suspended in SDW, and added to 25 ml YPM (open triangles, pH 7.3; or open squares,  
461 pH 6.1) or 25 ml YPM-A1 medium (closed triangles, pH 7.3; or closed squares, pH 6.1)  
462 in a 50-ml Erlenmeyer flask to an initial  $A_{600}$  of 0.1; cultivation was continued at 30°C  
463 and 95 spm. YPM-A1 medium consists of 2.5 ml 10-fold concentrated YP, 0.5 g  
464 mannitol, and 22.5 ml of A1-supernatant from 3-day culture of the ethanologenic strain  
465 MK3353 without pH adjustment (resulting in a final pH of 7.3; closed triangles) or with  
466 adjustment to pH 6.0 (resulting in a final pH of 6.1; closed squares).  $A_{600}$  indicating cell  
467 growth (A) and ethanol concentration (B) were measured. (A, B) Means and maximum  
468 and minimum values of two independent experiments are shown.

469

470

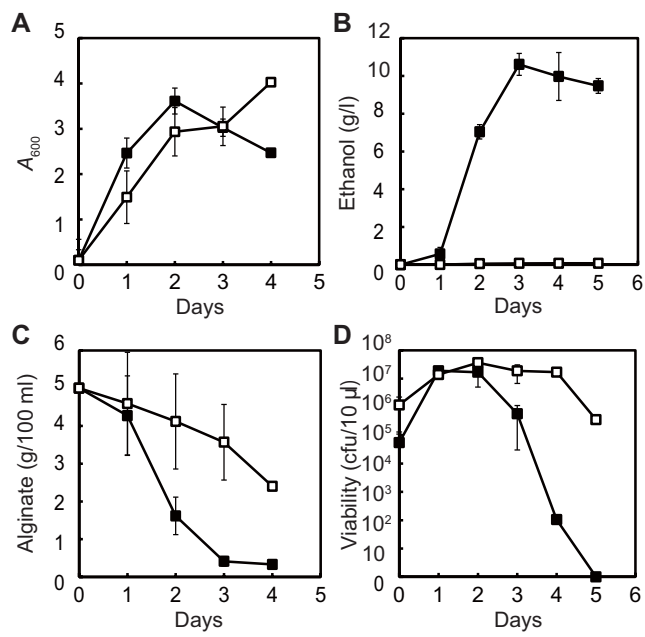


Fig. 1

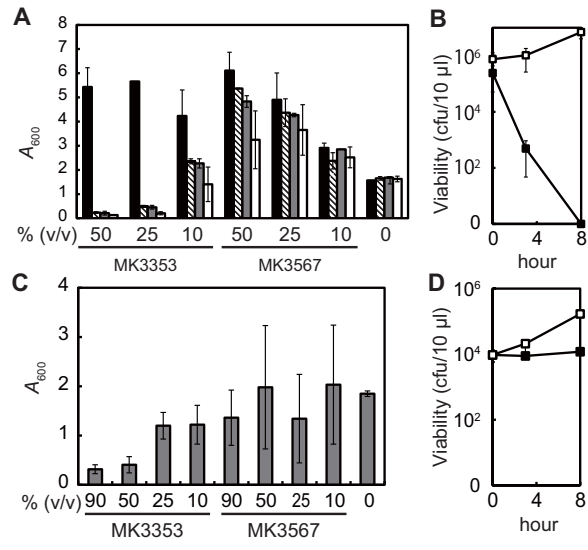


Fig. 2

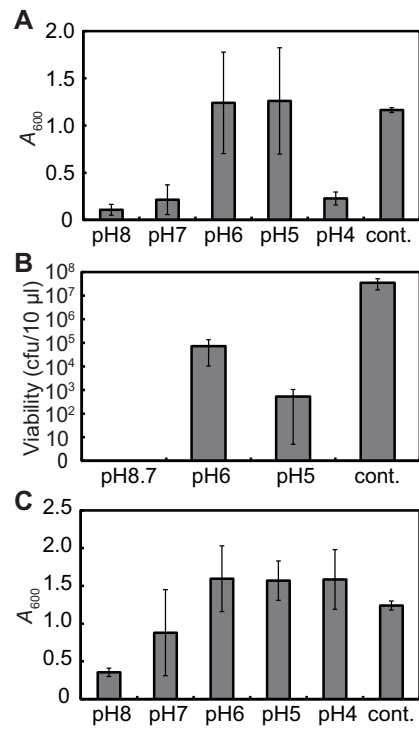


Fig. 3



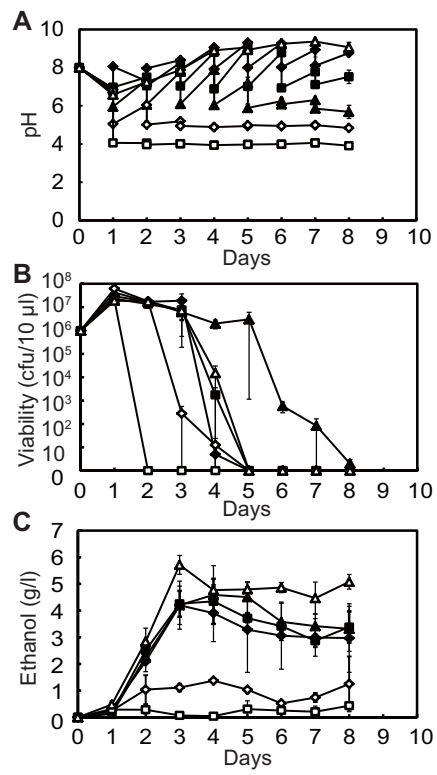


Fig. 4

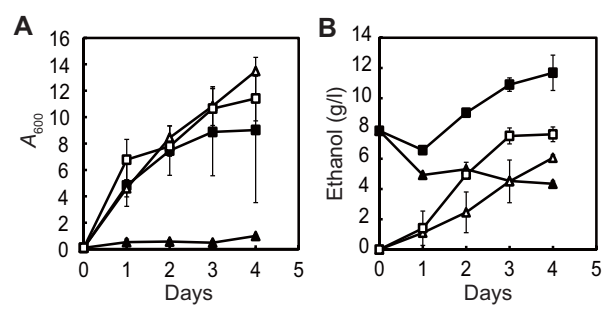
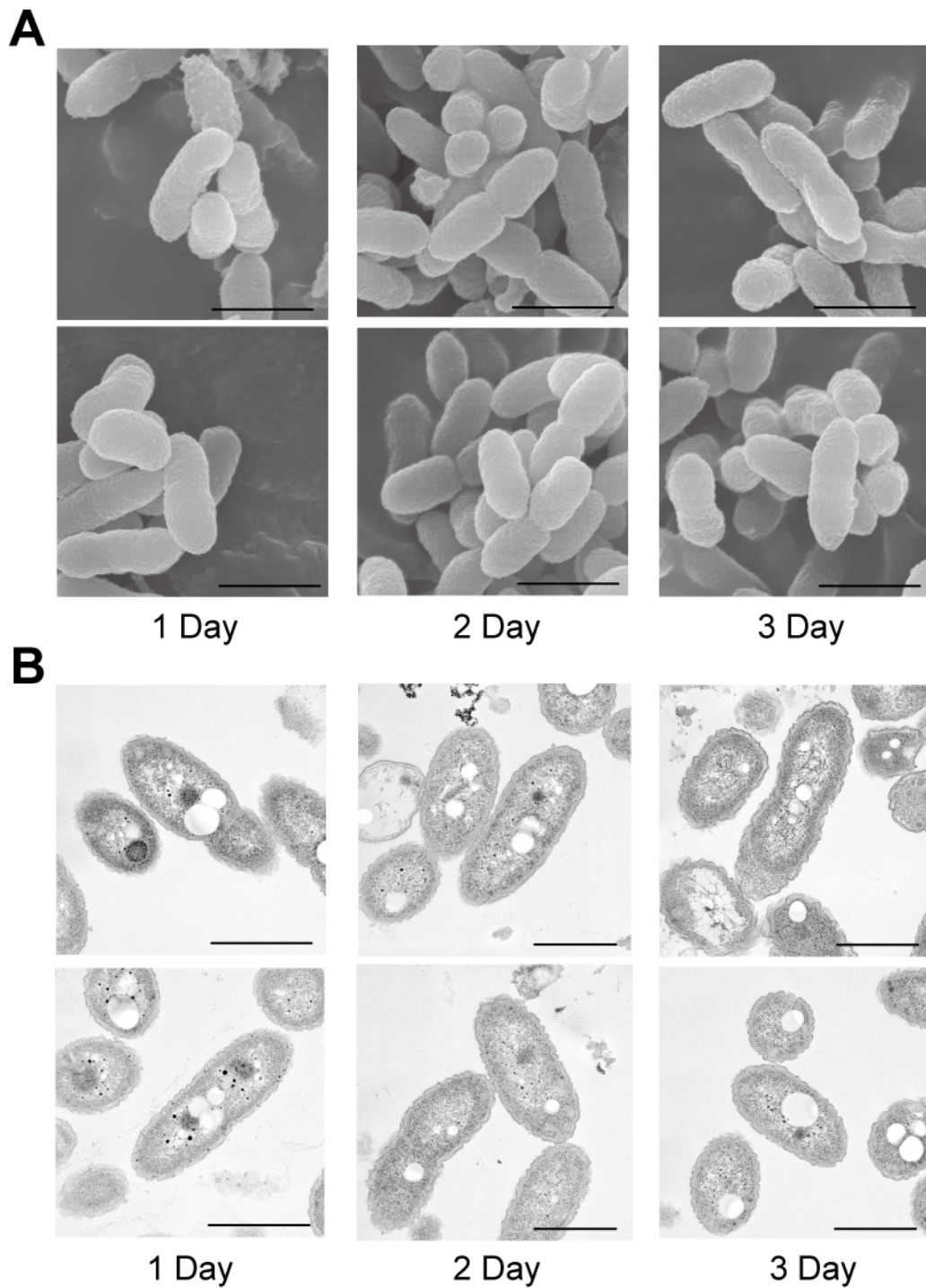
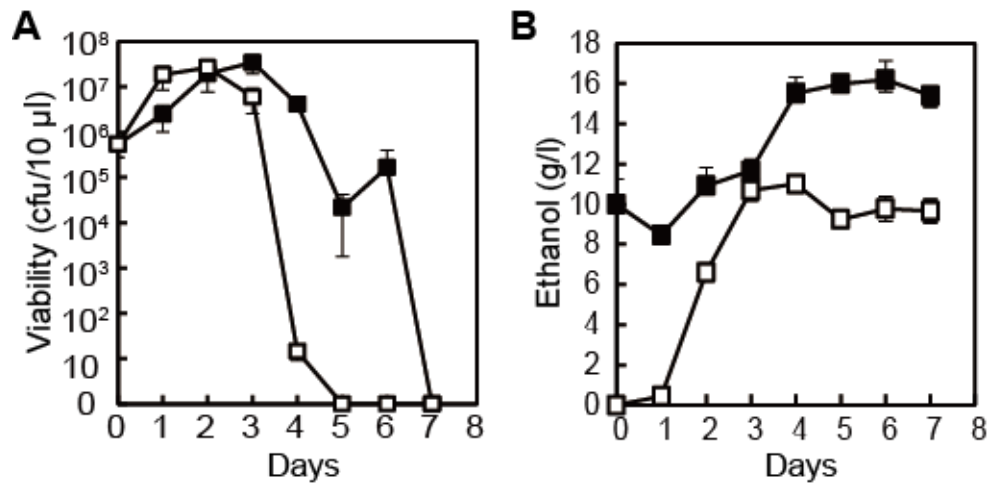


Fig. 5



**Supplementary Fig. S1.** SEM (A) and TEM (B) observations of cells of the ethanologenic strain MK3353 (upper panels) and the control strain MK3567 (lower panels). Cells were precultured, inoculated, and cultivated as described in MATERIALS AND METHODS in liquid alginate medium (100 ml) containing 5% w/v alginate at 30°C and 95 spm for the indicated periods. SEM and TEM observations were conducted

as described in MATERIALS AND METHODS. Scale bars: 1  $\mu\text{m}$  for SEM, 0.5  $\mu\text{m}$  for TEM.



**Supplementary Fig. S2.** Initial presence of ethanol causes no reduction in cell viability of ethanologenic strain MK3353. The ethanologenic strain MK3353 was cultivated as in Fig. S1 in the presence of an initial concentration of 0% w/v (open symbols) or 1.0% w/v [10 g/l] (closed symbols) ethanol. Number of viable cells per 10 µl culture (viability; cfu) (A) and ethanol concentration in A1-supernatant (B) were determined. Means and standard deviation (SD) of three independent experiments are shown.