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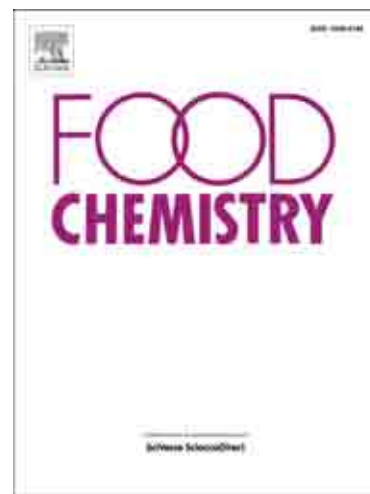
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1 Water-in-oil-in-water double emulsion for the delivery
2 of starter cultures in reduced-salt moromi fermentation
3 of soy sauce

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9

10 **Abstract**

11 This study investigated the application of water-oil-water ($W_1/O/W_2$) double emulsions (DE)
12 for yeast encapsulation and sequential inoculation of *Zygosaccharomyces rouxii* and
13 *Tetragenococcus halophilus* in moromi stage of soy sauce fermentation with reduced NaCl
14 and/or substitution with KCl. *Z. rouxii* and *T. halophilus* were incorporated in the internal W_1
15 and external W_2 phase of DE, respectively. NaCl reduction and substitution promoted *T.*
16 *halophilus* growth to 8.88 log CFU/mL, accompanied with faster sugar depletion and
17 enhanced lactic acid production. Reducing NaCl without substitution increased the final pH
18 (5.49) and decreased alcohols, acids, esters, furan and phenol content. However, the
19 application of DE resulted in moromi with similar microbiological and physicochemical
20 characteristics to that of high-salt. . Principal component analysis of GC-MS data

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21 demonstrated that the reduced-salt moromi had identical aroma profile to that obtained in the
22 standard one, indicating the feasibility of producing low-salt soy sauce without compromising
23 its quality.

24 Keywords: Soy sauce; Moromi fermentation; Salt reduction; W₁/O/W₂ double emulsion;
25 Yeast encapsulation; Sequential inoculation; Aroma compounds; GC-MS.

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26 1. Introduction

27 Soy sauce is a traditional fermented seasoning that is popular in Asia and throughout the
28 world, due to its intense umami taste and characteristic flavor. Soy sauce production process
29 involves a 2-step fermentation process, called *koji* and *moromi*. Koji fermentation begins by
30 mixing steam-cooked soybeans and roasted wheat flour with spores of mold, such as
31 *Aspergillus oryzae* or *Aspergillus sojae*, and, after 3 days of incubation, a compact mass is
32 formed due to mycelium growth (Zhu & Tramper, 2013). The resulting koji is then immersed
33 in brine solution, typically containing 18–22% NaCl to initiate the second step of
34 fermentation and produce moromi mash, and left to ferment for approximately 6 months.
35 During this stage, a wide range of microbial species grow spontaneously and produce
36 numerous flavor compounds, which are essential to the organoleptic properties of the final
37 product. *Tetragenococcus halophilus* and *Zygosaccharomyces rouxii* have been considered as
38 the most predominant osmophilic lactic acid bacteria (LAB) and yeast, respectively, and play
39 major roles in the aroma formation (Wilfred F.M. Röling, Timotius, Prasetyo, Stouthamer, &
40 Van Verseveld, 1994). The use of brine with high NaCl concentration in moromi
41 fermentation is important to control undesirable microorganisms and improve the flavor
42 profile and texture of the final product (Song, Jeong, & Baik, 2015a). However, high NaCl
43 content contributes to excessive sodium intake, which has been reported to increase risk of
44 hypertension, cardiovascular disease, and renal dysfunction (Kremer, Mojet, & Shimojo,
45 2009). Furthermore, the World Health Organization (WHO) recommends a limitation of
46 average daily intake of sodium to 2 g, which is equivalent to 5 g of salts (WHO, 2012). As a
47 consequence, producing soy sauce with low NaCl content without compromising its quality
48 and consumer acceptability is a challenge, and low salt soy sauce products are now available.
49 Soy sauce production with reduced NaCl has been investigated by different approaches.
50 Moromi fermentation in the absence of NaCl was possible by autolyzing koji under high

51 temperature prior to fermentation (Muramatsu, Sano, Uzuka, & Company, 1993).
52 Nevertheless, the absence of salt during fermentation may result in the growth of spoilage
53 microorganisms and the quality of final product can differ from the original. Salt reduction
54 during moromi fermentation could result in lower content of essential acids, alcohols, and
55 esters, and higher acidity content (Song et al., 2015a). Such problems could be counteracted
56 by the addition of mixed cultures of indigenous yeast species (Song et al., 2015a) as well as
57 combining LAB and yeasts (Singracha et al., 2017).

58 However, a recent study showed that the final aroma profile in moromi fermentation was
59 compromised due to antagonism between co-inoculated *Tetragenococcus halophilus* and
60 *Zygosaccharomyces rouxii* while their sequential inoculation could improve the aroma
61 complexity (Devanthi, Linforth, Onyeaka, & Gkatzionis, 2018). The application of sequential
62 inoculation of mixed cultures has been reported to improve flavor quality of fermented foods
63 and beverages. Modulation of the inoculation time was found to be key in achieving the
64 desired quality of apple cider (Ye, Yue, & Yuan, 2014). Furthermore, in whey fermentation,
65 sequential inoculation of *Kluyveromyces lactis* B10 and *Torulaspora delbrueckii* B14 after 48
66 h improved volatile compounds production (e.g. alcohols and esters) (Andrade, Melo,
67 Genisheva, Schwan, & Duarte, 2017). Higher production of 3-sulfanylhhexyl acetate (3SHA)
68 and 3SH (3-sulfanyl-1-hexanol), which are the most important volatiles in Sauvignon blanc
69 aroma, has been achieved with sequential culture of *T. delbrueckii* and *S. cerevisiae*.

70 A formulation is needed to control the sequential delivery and activity of microbial cultures
71 in soy sauce fermentation. Water-in-oil-in-water ($W_1/O/W_2$) double emulsions (DE) have
72 been studied in recent years for their ability to encapsulate hydrophilic substances, including
73 bacteria for protection and controlled release. Their multi-compartmentalized structure is
74 created by dispersing a water-in-oil (W_1/O) emulsion in another aqueous phase (W_2). Recent
75 studies have focused on probiotic bacteria encapsulation in DE for enhancing survival during

76 digestion (Eslami, Davarpanah, & Vahabzadeh, 2016; Shima, Morita, Yamashita, & Adachi,
77 2006). The instability of DE structure can be used to modulate the release of bacterial cells by
78 utilizing changes in osmotic balance (El Kadri, Gun, Overton, Bakalis, & Gkatzionis, 2016;
79 El Kadri, Overton, Bakalis, & Gkatzionis, 2015) as they would occur during fermentation. A
80 previous study demonstrated that the inherent DE instability acted as a mechanism for
81 gradual release of *Z. rouxii*, which could be linked to changes in glucose concentration in the
82 medium (Devanthi, El Kadri, Bowden, Spyropoulos, & Gkatzionis, 2018).

83 In this study, the application of DE for the encapsulation and sequential delivery of *T.*
84 *halophilus* and *Z. rouxii* cultures was tested in conditions reflecting moromi fermentation
85 with reduced NaCl content and/or substitution with KCl. The stability of DE in moromi was
86 examined by monitoring its microstructure, oil globules size, and distribution. Furthermore,
87 microbial population and physicochemical changes as well as volatile compounds formation
88 were monitored.

89 **2. Materials and Methods**

90 *2.1 Materials, chemicals, and microorganisms*

91 Soy and wheat flour were purchased from a local retailer (UK). *Aspergillus oryzae* 126842
92 was purchased from Centre for Agriculture and Biosciences International (Egham, UK).
93 *Tetragenococcus halophilus* 9477 and *Zygosaccharomyces rouxii* 1682 were purchased from
94 National Collection of Industrial Food and Marine Bacteria Ltd. (Aberdeen, UK) and
95 National Collection of Yeast Cultures (Norwich, UK), respectively. Sodium chloride (NaCl,
96 extra pure) was purchased from Acros Organics (Fairlawn, NJ). Microbiological growth
97 media used were Czapek Dox Agar (CDA; Oxoid Ltd., Basingstoke, UK), Brain Heart
98 Infusion agar (BHI, Oxoid Ltd., UK), de Man, Rogosa, and Sharpe broth (MRS broth, Oxoid
99 Ltd., UK), Yeast Malt agar (YM agar, Sigma-Aldrich, Gillingham, UK), Yeast Malt broth

100 (YM broth, Sigma-Aldrich, UK). Bacteria and yeast growth were controlled using
101 chloramphenicol (Oxoid Ltd., UK) and natamycin (Sigma-Aldrich, UK), respectively. 1-
102 Octen-3-ol (purity $\geq 98\%$) was purchased from Sigma-Aldrich. Soybean oil (Alfa Aesar,
103 Heysham, UK) was used as the oil phase of the DE. Polysorbate 80 (Tween 80, Sigma-
104 Aldrich, United Kingdom) and polyglycerol polyricinoleate (PGPR, Danisco A/S,
105 Copenhagen, Denmark) were used as water and oil soluble emulsifiers, respectively.

106 2.2 Culture preparation

107 *Aspergillus oryzae* was maintained on CDA at 25 °C. The spore suspension of *A. oryzae* was
108 prepared according to the method described by Chou and Ling (1998) with slight
109 modification. Briefly, spores were obtained by growing *A. oryzae* on CDA at 25 °C for 7
110 days. NaCl solution (0.85%, w/v) solution containing 0.01% of Tween 80 (Sigma-Aldrich,
111 UK) was added into the agar slant bottle followed by vigorous mixing to collect the spores.
112 The number of spores were counted using an improved Neubauer hemocytometer and
113 adjusted to 10^6 spores/mL. *Tetragenococcus halophilus* was maintained on BHI with 10%
114 (w/v) NaCl and incubated at 37 °C. *T. halophilus* was grown in MRS broth with 7% NaCl for
115 36 h and the cell concentration was adjusted to a final concentration of 10^6 cells/mL.
116 *Zygosaccharomyces rouxii* was maintained on YM agar with 5% (w/v) NaCl and incubated at
117 25 °C. The inoculum was prepared by growing *Z. rouxii* in YM broth containing 5% (w/v)
118 NaCl in a 30 °C shaker incubator for 24 h and cell concentration was adjusted to 10^6
119 cells/mL.

120 2.3 DE preparation

121 The DEs were prepared using a 2-step emulsification method at ambient temperature by
122 using a high shear mixer (Silverson L5M). In the first step, W₁/O primary emulsion was
123 prepared by mixing sterile 6% (w/v) NaCl solution into the oil phase (soybean oil with 2% wt

124 PGPR) at W_1 :oil phase ratio of 20:80 at 1700 rpm for 2 min. For yeast encapsulation, *Z.*
125 *rouxii* suspension in 6% (w/v) NaCl solution (10^7 cells/mL) was used as W_1 .

126 In the second stage, W_1/O was re-emulsified in the continuous phase (W_2 ;sterile 6% (w/v)
127 NaCl in water with 1% wt Tween 80) at 2000 rpm for 1 min ($W_1/O:W_2$ ratio of 20:80). The
128 final concentration of encapsulated *Z. rouxii* cells was $\sim 10^5$ cells/mL. DEs containing *T.*
129 *halophilus* in the W_2 were prepared by directly adding 2 mL of *T. halophilus* (10^6 cells/mL)
130 into the W_2 after the mixing process.

131 2.4 Soy sauce fermentation

132 **Koji preparation:** Koji was prepared using the modified method of Su et al. (2005). Soy and
133 wheat flour were sterilized at 121 °C for 15 min in an LTE Series 300 autoclave (LTE
134 Scientific Ltd, Oldham, UK). Soy flour moisture was maintained by mixing 100 g of soy
135 flour with 120 mL of sterile distilled water. The cooked soy flour was cooled to room
136 temperature and then mixed thoroughly with the wheat flour (1:1 w/w). The mixture was
137 inoculated with *A. oryzae* spore to a final concentration of 10^5 spores/g substrate (Chou &
138 Ling, 1998). The inoculated substrates were transferred into sterile Petri dishes (d:140 mm)
139 and incubated at 30 °C for 3 days.

140 **Moromi preparation:** Different types of brine (18% w/v NaCl; 6% w/v NaCl and 12% w/v
141 KCl; 6% w/v NaCl) were added to the koji with ratio of 1:5 (koji:brine) to create moromi
142 $A_{[18\%]}$, $B_{[6:12\%]}$, and $C_{[6\%]}$ respectively, followed by inoculation as shown in Figure 1. Moromi
143 $A_{[18\%]}$ and $B_{[6:12\%]}$ were simultaneously inoculated with *T. halophilus* and *Z. rouxii*. Three
144 different moromi C were prepared according to the inoculation method of *Z. rouxii*. Moromi
145 $C1_{[6\%]}$ was simultaneously inoculated with *T. halophilus* and *Z. rouxii*, while moromi $C2_{[6\%]}$
146 and $C3_{[6\%]}$ were inoculated with *Z. rouxii* after 1 week and 2 weeks, respectively. Moromi
147 $C4_{[6\%]}$ was inoculated with DE (10% v/v) containing *T. halophilus* and *Z. rouxii*, which had

148 been incorporated in its W_2 and W_1 phase, respectively, prior to inoculation. The inoculated
149 moromi mashes were then incubated at 30 °C for 4 weeks and samples were taken at Week 0,
150 1, 2, 3, and 4. *T. halophilus* was grown on BHI agar supplemented with 7% (w/v) NaCl and
151 natamycin while the cell count of *Z. rouxii* was done on YM agar with the addition of 5%
152 (w/v) NaCl, and 100 mg/L chloramphenicol. In order to study the effect of koji:brine ratio on
153 DE stability, koji was mixed with 18% w/v NaCl solution with koji:brine ratio of 1:3, 1:5, and
154 1:7 followed by incubation at 30 °C for 7 days.

155 2.5 Rheological measurements

156 Rheological characterization of moromi was done by measuring the viscosity of koji mixed
157 with varying concentrations of brine solution (18% NaCl w/v). The viscosity was measured
158 for moromi containing koji:brine ratio of 1:3, 1:5, 1:7 and brine only at 30 °C using AR-G2
159 rheometer (TA instruments, New Castle, DE) on a parallel plate geometry (d: 40 mm). The
160 apparent viscosity was measured over a shear rate range of 0.1–100 s^{-1} . Briefly, 1 mL of
161 sample was placed between the cone and the plate, and measurement was started
162 immediately. In total, 30 data points were recorded at 10-s intervals during the shearing.
163 Shear stress was determined as a function of shear rate. Data were fitted to power-law model
164 (Barnes et al., 1989):

$$165 \quad \eta = K \cdot \gamma^{n-1} \quad (1)$$

166 where; η refers to viscosity (Pa s), K to consistency coefficient (Pa sⁿ), γ to shear rate (s^{-1}),
167 and n to flow behavior index (dimensionless).

168 2.6 Physicochemical analysis

169 Soy mash samples were centrifuged at 15000 g for 15 min at ambient temperature. The
170 supernatant regarded as raw soy sauce was transferred to microtubes and kept at –20 °C until

171 analysis. Total reducing sugar (D-glucose and D-fructose), total lactic acid (L-lactic acid and
172 D-lactic acid), ethanol, and L-glutamic acid were analyzed using enzymatic assay kit
173 (Megazyme, International Ireland Ltd., Bray, Ireland) according to the manufacturer's
174 instructions. Changes in pH were monitored using a pH meter (SevenCompact S220, Mettler
175 Toledo, Germany).

176 2.7 Volatile compound analysis (SPME GC-MS)

177 An automated headspace solid-phase microextraction method (SPME) followed by GC-MS
178 analysis was used for evaluating the *in vitro* production of microbial volatile organic
179 compounds. Soy sauce mash samples (1.5 g) were transferred into 20-mL headspace vials
180 (22.5 mm × 75.5 mm, Grace Alltech, Thermo Fisher UK) and the vials were sealed with
181 magnetic cap (20 mm diameter, 5 mm center, PTFE / Silicone Liner; Grace Alltech). Samples
182 were allowed to equilibrate at 22 °C for 30 min before analysis. Three replicates were
183 prepared for all samples.

184 The volatiles extraction was performed using a 1-cm Stableflex fiber coated with 50/30 µm
185 divinylbenzene-Carboxen on polydimethylsiloxane bonded to a flexible fused silica core
186 (Supelco, Bellefonte, PA). It was conditioned for 90 min at 300 °C in the injection port. The
187 fiber was pushed out of the housing and inserted into the vials through the center of the vial
188 cap. The penetration depth was fixed at 22 mm. The extraction was carried out by exposing
189 the fiber to the headspace for 10 min at 40 °C. For all analyses, desorption time was set to 10
190 min at 230 °C.

191 Chromatography was carried out using a Trace GC Ultra gas chromatography (Thermo
192 Electron Corporation, Hemel Hempstead, UK) equipped with a polar column ZB-Wax (30 m
193 × 0.25 mm I.D.; film thickness: 1 µm) from Phenomenex (Torrance, CA). Mass spectrometry

194 (MS) was performed with a DSQ mass spectrometer (Thermo Electron Corporation, Hemel
195 Hempstead, UK)). GC-MS parameters were set according to a previous study (Gkatzionis,
196 Linforth, & Dodd, 2009): The temperature of the injection port was 230 °C. Helium was
197 employed as the carrier gas, at a constant pressure of 17 psi. The oven temperature program
198 was as follows: an initial temperature of 40 °C was maintained for 2 min, increasing at a rate
199 of 8 °C /min to a final temperature of 220 °C. The transfer line from the gas chromatograph
200 to the mass spectrometer was held at 250 °C. The mass spectrometer was operated in positive
201 ionization electron impact mode (EI+) at 70 eV. The detector was operated in scan mode (2
202 scans/s) scanning from m/z 20 to 250. Source temperature was 200 °C.

203 Compounds were identified by comparing their retention times and mass spectra with those
204 of standards or their retention indices (RI) with those published in databases and their mass
205 spectra with the National Institute of Standards and Technology (NIST) mass spectral library
206 using XCalibur Software (Thermo Electron Corporation, UK). The signal intensity for each
207 compound was expressed relative to the signal observed when the headspace above a 0.1
208 $\mu\text{g/mL}$ 1-octen-3-ol solution was sampled.

209 2.8 *DE stability characterization*

210 DE samples were placed onto the microscope slides and the microstructure was observed
211 under a light microscope (Olympus BX50) with a 10 \times objective lens. Images were taken
212 using a Moticam 10 camera *via* Motic Images Plus video acquisition software at 17fps. The
213 oil droplets size distribution of DE was determined from microscopic images using image
214 analysis software (ImageJ), by measuring the diameter of at least 500 oil droplets from 3
215 different samples of DE.

216 2.9 Statistical analysis

217 Microbial cell enumeration, physicochemical tests, and volatile compounds analysis were
218 conducted in triplicate and repeated in two independent experiments. The results were
219 presented as mean \pm standard deviation. Significant differences among means were tested by
220 one-way analysis of variances (ANOVA) using IBM SPSS Statistics Software Version 21 at
221 $p < 0.05$ and Tukey's test was applied for means comparison. Principal component analysis
222 (PCA) was performed using XLSTATTM version 2015.6.01.24027 (Addinsoft, New York,
223 NY) to reduce the dimensionality of the dataset and show the differences in volatile
224 compounds among the soy sauce samples. Observations/variables were chosen as data format
225 and Pearson's correlation matrix was used as PCA type.

226 3. Results and Discussion

227 3.1 The effect of viscosity on the stability of DE in moromi

228 DEs were formulated using ingredients relevant to moromi constituents and soybean oil was
229 used as the oil phase. Since the reduced-salt moromi contained 6% NaCl, the internal W_1 and
230 external W_2 phase of DE also contained 6% NaCl. This aimed to balance the osmotic
231 pressure between the two phases, thus reducing instability of DE due to water movement
232 across the oil phase (Mezzenga, Folmer, & Hughes, 2004).

233 In order to describe the relationship between the viscosity of moromi and DE stability,
234 moromi formulations with different viscosities were tested by varying the ratio of koji:brine
235 (1:3, 1:5, and 1:7). The Power-Law model was used to describe the flow curves of the
236 moromi. The rheological parameters of this model are presented in Table S1. All the moromi
237 formulations exhibited non-Newtonian behavior at shear rates ranging between 0.1 and 100 s⁻¹
238 at 30 °C (Figure S1a). Moreover, the plot of the viscosity against shear rate of the koji and
239 brine mixtures yielded a flow index (n) of less than 1 (shear thinning), indicating that their

240 flow behavior had a non-Newtonian profile. Similar non-Newtonian behavior has been
241 reported for semi-solids of similar composition to koji which could be attributed to the
242 presence of high molecular weight components, such as proteins or dextrin (Manohar,
243 Manohar, & Rao, 1998).

244 DE maintained its microstructure after 4 weeks of fermentation (Figure 2a). However, the oil
245 globule size significantly decreased from 27.88 μm to 11.40 μm (Figure 2b and 2c). This
246 could be attributed to the high viscosity of the moromi system. The viscosity increased when
247 the amount of brine added was decreased (Figure S1a). After incorporation into the moromi
248 system, the DE stability was determined by observing its microstructure (i.e. inner W_1 phase)
249 using microscopy and monitoring the oil globule size. The initial oil globule size (31.84 μm)
250 decreased immediately after incorporation into the moromi slurry and during storage (Figure
251 S1b and S1c). However, the decrease in koji:brine 1:3 was more noticeable compared to
252 those with higher fractions of brine. By the end of storage, the oil globule size of DE in koji :
253 brine 1:3, 1:5, and 1:7 was 6.84 μm , 18.02 μm , and 15.29 μm , respectively. Moreover, all the
254 oil globules in koji:brine 1:3 completely lost their inner phase, while in koji:brine 1:5 and 1:7,
255 the DE structure was maintained (Figure S1d). These data indicate that DEs were destabilized
256 in the moromi system; however, the destabilization was not proportional to the viscosity of
257 the moromi.

258 3.2 *The effect of salt reduction and inoculation sequence on the growth of *T. halophilus** 259 *and *Z. rouxii**

260 Salt concentration is a significant parameter that determines soy sauce fermentation process
261 by affecting microbial growth. High salt concentration is typically used in soy sauce
262 fermentation, in order to suppress the growth of undesirable microorganism as well as
263 improving the organoleptic properties of the final product. *T. halophilus* growth was

264 suppressed during the first 2 weeks of fermentation (from 6.30 log CFU/mL to 4.17 log
265 CFU/mL) when 18% NaCl (A_[18%]) was present in moromi (Figure 3a). Meanwhile, its
266 growth was significantly enhanced when part of the NaCl was replaced with KCl (B_[6:12%])
267 and maintained high viability, reaching 7.88 log CFU/mL. Interestingly, the growth of *T.*
268 *halophilus* in A_[18%] recovered after 2 weeks and exceeded B_[6:12%] by the end of incubation.
269 In any case, the growth was higher at the lowest salt concentration (C1_[6%], C2_[6%], C3_[6%])
270 throughout the fermentation, where the cell count sharply increased to 8.49 log CFU/mL
271 within the first week and remained stable throughout the incubation period. Although *T.*
272 *halophilus* is an osmophilic LAB that can tolerate up to 26% NaCl, it grows best at 5 to 10%
273 w/v (Taniguchi et al., 1988). Therefore, raising the NaCl concentration can increase the
274 osmotic stress, reducing the ability of *T. halophilus* to grow (Kobayashi et al., 2004). This
275 indicated that *T. halophilus* could not grow immediately after inoculation in the presence of
276 high NaCl concentration, as previously described by Taniguchi et al. (1988).

277 The growth of *T. halophilus* under reduced-salt environment was enhanced when it was
278 simultaneously inoculated with *Z. rouxii* (C1_[6%]) compared to sequential inoculation (C2_[6%],
279 C3_[6%]) and gradual release in DE (C4_[6%]). The addition of *Z. rouxii* from the early stage of
280 fermentation might have supplied a variety of metabolites such as pyruvate, amino acids, and
281 vitamins, which are essential for the early stage of bacterial growth (Devanthi et al., 2018;
282 Sudun, Wulijideligen, Arakawa, Miyamoto, & Miyamoto, 2013).

283 *Z. rouxii* was not affected significantly by salt concentration during the first 3 weeks of
284 fermentation. However, low-salt moromi (C1_[6%]) suffered a decrease in its population at
285 Week 4, in contrast to the enhanced growth of *T. halophilus*. *Z. rouxii* is typically added to
286 enhance flavor and aroma formation in soy sauce production through alcoholic fermentation
287 (Van Der Sluis, Tramper, & Wijffels, 2001; Wah, Walaisri, Assavanig, Niamsiri, & Lertsiri,
288 2013). In a previous study by Singracha et al. (2017), the addition of *Z. rouxii* in combination

289 with *T. halophilus* and *Pichia guilliermondii* was shown to increase the total population of
290 lactic acid bacteria and yeast in reduced-salt moromi fermentation. Since *Z. rouxii* grows
291 optimally at low pH, *Z. rouxii* would be better added at the later stage of fermentation, once
292 moromi is acidified due to organic acids production by *T. halophilus*. In the present study, *Z.*
293 *rouxii* sequential inoculation (C2_[6%] and C3_[6%]) and gradual release in DE (C4_[6%]) did not
294 have significant effect on growth, as this seemed to depend primarily on the salt formulation
295 and less on inoculation sequence (Figure 3b).

296 3.3 Physicochemical changes during fermentation

297 The changes in pH, reducing sugar, lactic acid, ethanol, and glutamic acid were measured to
298 monitor the fermentation progress, as they are associated with the growth of microorganisms
299 (Figure 4). Besides increasing in population during soy sauce fermentation, LAB also utilize
300 and convert carbohydrates into organic acids, which can bring the pH down. Reduction in pH
301 can also occur due to the accumulation of free fatty acids, amino acids, and peptides
302 containing carboxylic side chains, resulting from other microbial activities and raw materials
303 hydrolysis (Hoang et al., 2016; Van Der Sluis et al., 2001; Yanfang & Wenyi, 2009). As
304 shown in Figure 4a, pH of all moromi samples decreased from ~5.3 to final pH of ~4.8,
305 which was similar to values reported in previous studies of traditional Korean (Song, Jeong,
306 & Baik, 2015b) and reduced-salt soy sauce (Singracha et al., 2017). The pH decreased within
307 two weeks and then remained constant throughout the fermentation period, except for C1_[6%],
308 where pH increased to 5.49. The reduction in pH was associated with the increase in the
309 lactic acid amount produced by *T. halophilus* (Figure 4c). Although lactic acid production
310 was greatly suppressed by 18% NaCl, the reduction in pH was unaffected, which could be
311 due to production of other organic acids. Although *T. halophilus* is known as
312 homofermentative, some strains are regarded as heterofermentative and they are able to
313 produce acetic acid (Justé et al., 2012). Moreover, homofermentative strains of *T. halophilus*

314 are reported to undergo mixed acid fermentation under certain growth conditions (Wilred F.
315 M. Röling & van Verseveld, 1997).

316 The production of lactic acid was significantly lower in the presence of high salt
317 concentration, and high sodium content had a greater impact on the suppression (Figure 4c).
318 In low salt concentration, microorganisms are able to perform faster metabolic activity,
319 therefore producing higher amount of acids (Hoang et al., 2016). In the present study, lactic
320 acid production in reduced-salt moromi was enhanced when the inoculation of *Z. rouxii* was
321 modulated, sequentially or gradually by using DE. In co-inoculation, *Z. rouxii* might have
322 changed the physicochemical properties of the substrate, which could suppress the
323 fermentation of lactic acid by *T. halophilus*, as reported in a previous study (Devanthi et al.,
324 2018).

325 Reducing sugar is important during fermentation as it serves as a carbon source for microbial
326 growth as well as flavor and aroma formation. The initial content of total reducing sugar in
327 all moromi samples ranged from 2.68 to 3.49 g/L and it constantly decreased throughout the
328 incubation period (Figure 4b), which was in agreement with the previous study by Zhang,
329 Zhou, Cui, Huang, and Wu (2016). The reduction patterns were comparable regardless of salt
330 concentration and sequence of inoculation. During fermentation, reducing sugar is consumed
331 by microbes or possibly reacts with free amino acids during the Maillard reaction (Kim &
332 Lee, 2008). Since the fungal amylase, which breaks down the polysaccharide into simple
333 sugars, was heat-inactivated prior to the moromi stage, the amount of reducing sugar was
334 expected to decrease over time. The reducing sugar content in moromi decreased faster when
335 low salt concentration (B_[6:12%], C1_[6%], and C4_[6%]) was used. This could be attributed to
336 faster metabolic activity of the microbes, which also corresponded to higher *T. halophilus*
337 population and lactic acid production (Hoang et al., 2016). Furthermore, the reducing sugar
338 content decreased at a slower rate when *Z. rouxii* was inoculated sequentially after 1 or 2

339 weeks of fermentation, but not when DE was used. This was expected since *Z. rouxii* is the
340 main user of sugar for biomass and ethanol production (Devanthi et al., 2018). The activity of
341 the released *Z. rouxii* cells might have caused faster sugar depletion in DE (C4_[6%]).

342 Ethanol production was highly affected by variation in salt concentration and sequence of
343 inoculation (Figure 4d). In low-salt moromi (C1_[6%]), the amount of ethanol constantly
344 decreased after 2 weeks of fermentation compared to a high concentration of salt (A_[18%] and
345 B_[6.12%]). However, the decrease in ethanol production was compensated when *Z. rouxii* was
346 added simultaneously (C2_[6%] and C3_[6%]) or using DE (C4_[6%]). Interestingly, ethanol
347 production with a similar pattern to A_[18%] and B and at highest concentration was achieved
348 when *Z. rouxii* was encapsulated in DE.

349 *Z. rouxii* is known to produce extracellular glutaminase, which is a proteolytic enzyme that
350 converts L-glutamine derived from soy protein to L-glutamic acid (Iyer & Singhal, 2008;
351 Kashyap, Sabu, Pandey, Szakacs, & Soccol, 2002). Unlike the glutaminase produced by koji
352 mold, *Z. rouxii* glutaminase is more tolerant against high salinity. L-Glutamic acid is essential
353 for improving the flavor of the final product since it contributes to the “umami” taste of the
354 soy sauce. Therefore, high activity of glutaminase is desirable, in order to increase the
355 production of L-glutamic acid. As shown in Figure 4e, the amount of glutamic acid increased
356 after the fermentation process and the final concentration of glutamic acid between samples
357 did not differ significantly ($p > 0.05$).

358 3.4 Formation of volatile compounds

359 A total of 38 volatile compounds was detected in the moromi samples using SPME-GC/MS,
360 including 15 alcohols, 5 acids, 8 aldehydes, 4 esters, 1 furan, 1 phenol, 3 ketones, and 1
361 alkene (Table 1). Alcohol was found to be the most abundant compound in all samples,

362 comprising more than 90% of the total volatiles, as previously found in high-salt liquid state
363 fermentation, low-salt solid-state fermentation, and Koikuchi soy sauce (Feng et al., 2015).

364 Salt reduction (C1_[6%]) was shown to have a great influence on the volatiles production in
365 moromi, especially alcohols (Table 1). Yeasts contribute to the formation of alcohols through
366 the reduction of related aldehydes (Sun, Jiang, & Zhao, 2010; Van Der Sluis et al., 2001).

367 Lowering salt concentration to 6% w/v (C1_[6%]) significantly ($p < 0.05$) enhanced the
368 production of 2,4-dimethyl-3-pentanol, 2,6-dimethyl-4-heptanol, 3-methyl-1-butanol, 5-
369 nonanol, and phenylethyl alcohol. On the other hand, the production of ethanol and propanol
370 was reduced in low salt concentration (C1_[6%]), which was in agreement with the previously
371 studied reduced-salt Korean soy sauce (Song, Jeong, & Baik, 2015a). Partial salt substitution
372 with KCl (B_[6:12%]) did not affect the production of most volatile compounds, except for 2-
373 furanmethanol, 2-methoxy-5-methylphenol, and 2-methyl-1-propanol which were
374 significantly ($p < 0.05$) lower compared to sample A_[18%]. In previous studies reported by
375 Sasaki (1996) and Jansen, Veurink, Euverink, & Dijkhuizen (2003), the production of higher
376 alcohols, including phenylethyl alcohol, 3-methyl-1-butanol, 1-propanol, and 2-methyl-1-
377 propanol, was found to decrease with an increase of NaCl concentration. However, the
378 amounts of 1-propanol and 2-methyl-1-propanol decreased under reduced NaCl conditions
379 (B_[6:12%]). This might have arisen from decreasing uptake of the related amino acid by yeast,
380 since these compounds are mainly produced by *Z. rouxii* from their corresponding branched-
381 chain amino acids *via* the Ehrlich pathway (Van Der Sluis et al., 2001). The method of
382 inoculation was found to affect the production of most alcohols in the reduced-salt moromi.

383 Moromi with similar flavor pattern to those containing high salt concentrations (A_[18%] and
384 B_[6:12%]) was achieved when *Z. rouxii* was added sequentially at Week 1 (C2_[6%]) or using DE
385 (C4_[6%]). The addition of *Z. rouxii* at Week 2 resulted in significantly ($p < 0.05$) lower
386 amounts of 2-furanmethanol, 3-methyl-1-butanol, ethanol, 1-heptanol, 1-hexanol, and 1-

387 propanol. This result corresponds to the ethanol measurement during fermentation by using
388 enzymatic reaction (Figure 4d).

389 Salt reduction was also found to affect the production of several acids. The amount of 4-
390 methyl-2-oxovaleric acid was enhanced in reduced-salt moromi, only when *Z. rouxii* was
391 added simultaneously. Meanwhile, 2-methylpropanoic acid, which contributes to cheese/fatty
392 odor, was found to be significantly lower in all reduced-salt moromi samples. However,
393 noticeably higher amount of 2-methylpropanoic acid was detected when *Z. rouxii* was added
394 at Week 2. The production of some acids, including 3-methylbutanoic acid (cheese/sweet)
395 and acetic acid (sour/vinegar-like odor), was found to be enhanced when *Z. rouxii* inoculation
396 was delayed for 2 weeks. Acetic acid production was also similar when DE was used. These
397 acids have been reported as the highest odor-active compounds in Chinese soy sauce (Feng et
398 al., 2014). Among these acids, 2-methylpropanoic acid and 3-methylbutanoic acid are formed
399 *via* branched-chain α -keto acid catabolism (Song et al., 2015b)

400 Aldehydes contribute to nutty and malty aroma in soy sauce (Feng et al., 2015). In the present
401 study, most aldehyde compounds were not affected by salt reduction, except for 2-
402 methylpropanal which was significantly enhanced in reduced-salt moromi when mixed
403 cultures were added simultaneously (C1_[6%]). This branched-chain aldehyde is considered as
404 an important flavor compound, perceived as malty, chocolate-like, with low taste threshold
405 (Smit, Engels, & Smit, 2009). It is generated from branched-chain amino acid valine *via*
406 Strecker degradation or microbial activity, which then can be converted to its corresponding
407 alcohol (2-methyl-1-propanol) and/or acid (2-methylpropanoic acid) (Ardö, 2006; Song et al.,
408 2015a). The effect of modulating the inoculation time of *Z. rouxii* on aldehydes formation
409 was hardly seen, except for benzaldehyde (burnt sugar/sweet) and furfural (bread/sweet),
410 which were significantly enhanced in C2_[6%] and C4_[6%], respectively.

411 Replacing NaCl with KCl decreased the amount of 2-phenylethyl acetate, which contributes
412 to honey, rosy odor. However, this could be compensated for by adding *Z. rouxii* at Week 1
413 of the fermentation process. In our previous study, it was found that the production of 2-
414 phenylethyl acetate could be enhanced by adding *Z. rouxii* sequentially rather than
415 simultaneously (Devanthi et al., 2018). *Z. rouxii* enhances the production of esters (Van Der
416 Sluis et al., 2001), although the production of isoamyl acetate (banana aroma) was
417 significantly enhanced in reduced-salt moromi. This was only observed in C1_[6%], while the
418 amount of isoamyl acetate in C2_[6%], C3_[6%], and C4_[6%] was similar to that at high salt
419 concentration (A_[18%] and B_[6:12%]).

420 The only furan and phenol compounds detected in all moromi samples were 3-acetyl-2,5-
421 dimethylfuran and 2-methoxy-5-methylphenol, respectively. These were produced in
422 negligible amount when either salt or NaCl were reduced, except when *Z. rouxii* was added at
423 Week 1. Several ketones, such as 3-methyl-2-pentanone and acetoin, were produced at
424 significantly higher concentrations in reduced-salt moromi. The amounts of these compounds
425 were similar to moromi containing high salt (A_[18%] and B_[6:12%]) when *Z. rouxii* was added
426 sequentially, with or without DE.

427 3.5 Principal component analysis

428 PCA analysis was conducted, in order to gain more understanding on the relationship
429 between the fermentation conditions and profiles of volatile compounds. The first (PC1) and
430 second principal component (PC2) accounted for 30.60% and 21.43% of the total variance,
431 respectively (Figure 5a-b). The PCA score plot demonstrates distinct separation of some
432 moromi samples (Figure 5a). In the case of co-inoculated samples, low salt moromi sample
433 (C1_[6%]) was differentiated from high salt moromi sample (A_[18%]) while reduced NaCl sample
434 (substituted with 12% KCl; B) was positioned in the middle of PC1. This indicates that salt

435 reduction affected the aroma profile of moromi. Replacing part of NaCl with KCl (B_[6:12%])
436 was associated with lower content of 2-furanmethanol, 2-methyl-1-propanol, 2-phenylethyl
437 acetate, 3-acetyl-2,5-dimethylfuran, and 2-methoxy-5-methylphenol. C1_[6%] was associated
438 with high amounts of 3-methyl-1-butanol, phenylethyl alcohol, 2,4-dimethyl-3-pentanol, 2,6-
439 dimethyl-4-heptanol, isoamyl acetate, 2-methylpropanal, 4-methyl-2-oxovaleric acid, 3-
440 methyl-2-pentanone, and acetoin (Figure 5b).

441 The method of inoculation was found to affect the aroma profiles, and adding *Z. rouxii*
442 encapsulated in DE or sequentially after 1 week matched the aroma profile obtained with
443 high salt concentration. This was not the case when *Z. rouxii* was added sequentially after 2
444 weeks of fermentation. Clustering of samples A_[18%], C2_[6%], and C4_[6%] was influenced by
445 compounds such as 2-furanmethanol, 2,4-dimethyl-3-pentanol, 2,6-dimethyl-4-heptanol, 3-
446 methyl-1-butanol, 5-nonanol, ethanol, 1-hexanol, methanol, phenylethyl alcohol, 4-methyl-2-
447 oxovaleric acid, 3-methylbutanoic acid, acetic acid, propionic acid, 2-methylbutanal, 2-
448 methylpropanal, 3-methylbutanal, furfural, hexanal, pentanal, propanal, ethyl acetate, ethyl
449 propionate, isoamyl acetate, 3-methyl-2-pentanone, acetoin, acetone, and D-limonene.

450 4. Conclusion

451 Salt reduction could affect microbial growth and physicochemical changes during moromi
452 fermentation. Low salt concentrations could promote *T. halophilus* growth and enhance lactic
453 acid production. However, the final overall aroma balance differed from the original soy
454 sauce fermented with high salt concentration, indicated by lower content of some alcohols,
455 acids, esters, furan, and phenol. The use of DE for delivering the mixed cultures of *T.*
456 *halophilus* and *Z. rouxii* in reduced-salt moromi could compensate for such changes by
457 promoting the formation of some essential volatile compounds, including alcohols (e.g., 2-
458 furanmethanol and ethanol) and esters (e.g., 2-phenylethyl acetate). This indicates the

459 possibility of producing soy sauce in a low salt environment with a volatile profile pattern
460 identical to the original high-salt soy sauce. The results obtained in this study provide the soy
461 sauce industry with a new technique for standardizing the microbial activity and aroma
462 development, which also offers health benefits to the consumers, due to low salt content in
463 the final product. However, since modulating the release has a great impact on the aroma
464 formation, further study is needed in order to tailor the physicochemical properties of DE,
465 therefore enabling the cell release in a more controlled manner.

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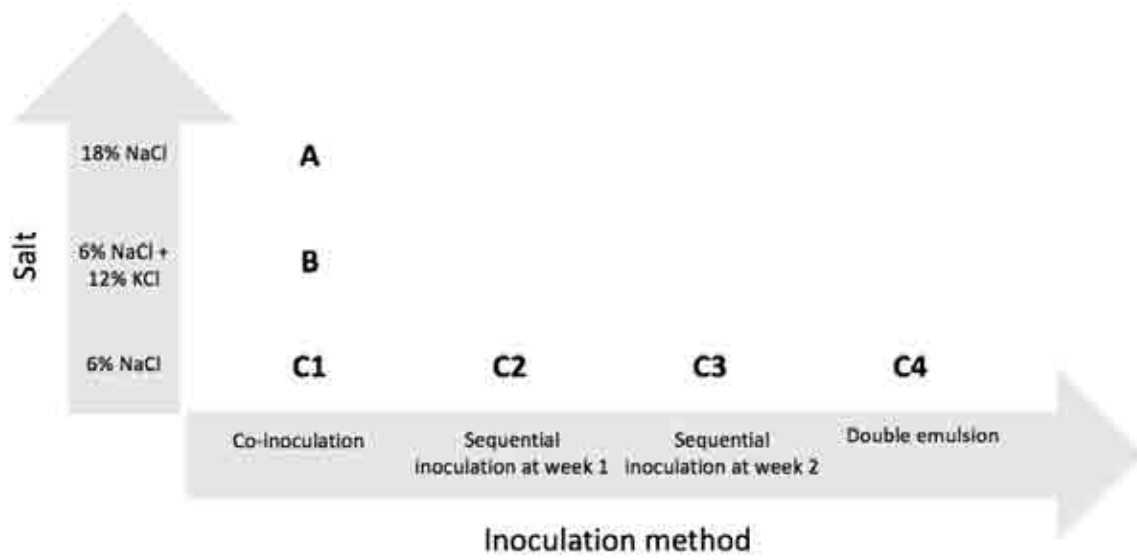
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593 Figure 1. Set of moromi samples varying in salt composition and inoculation method

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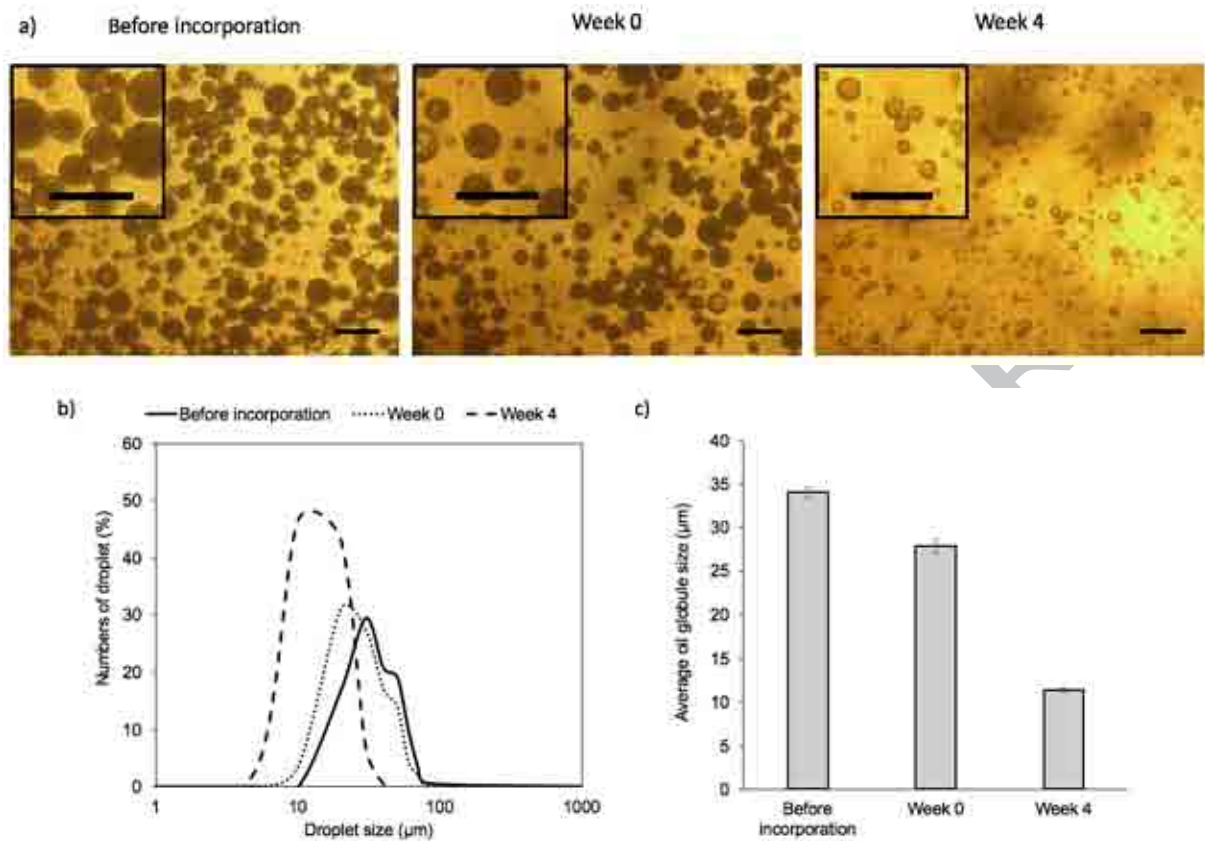
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606 Figure 2. (a) Optical micrograph of W₁/O/W₂ DE before and after incorporation into moromi,

607 and after 4 weeks of fermentation. Scale bar: 100 μm. (b) Oil globule size distribution before

608 and after fermentation. (c) Average oil globule size before and after fermentation.

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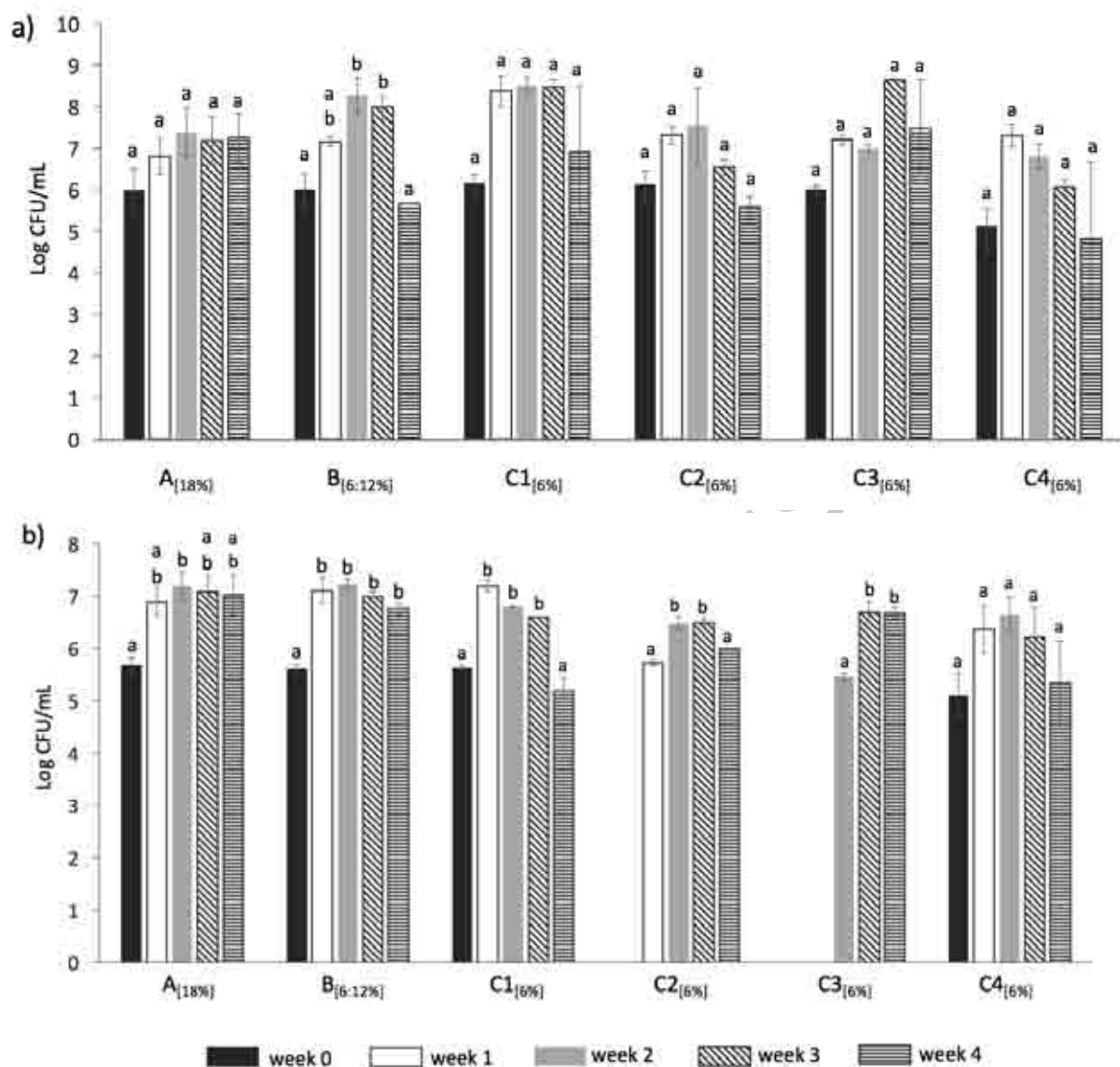
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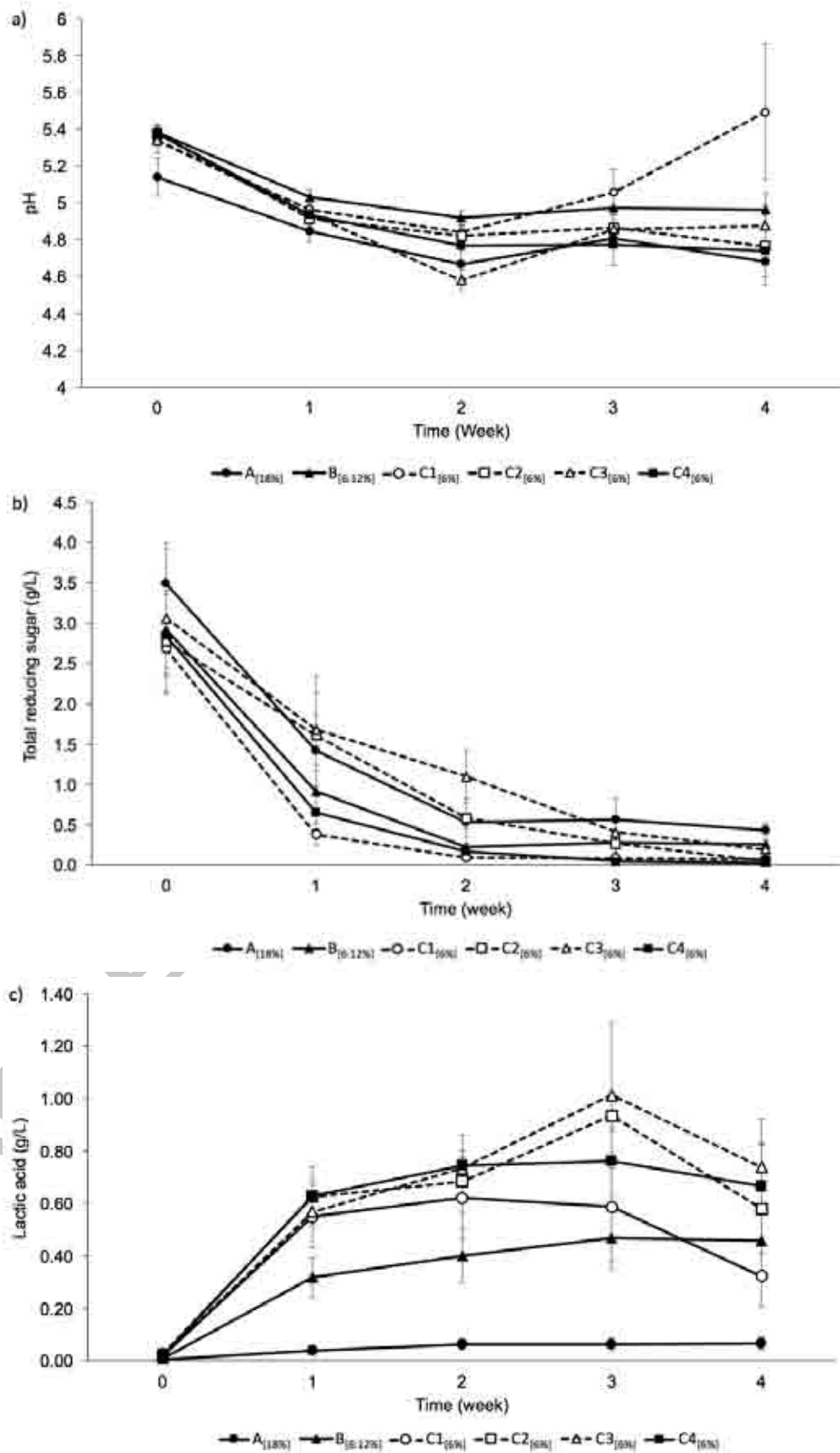
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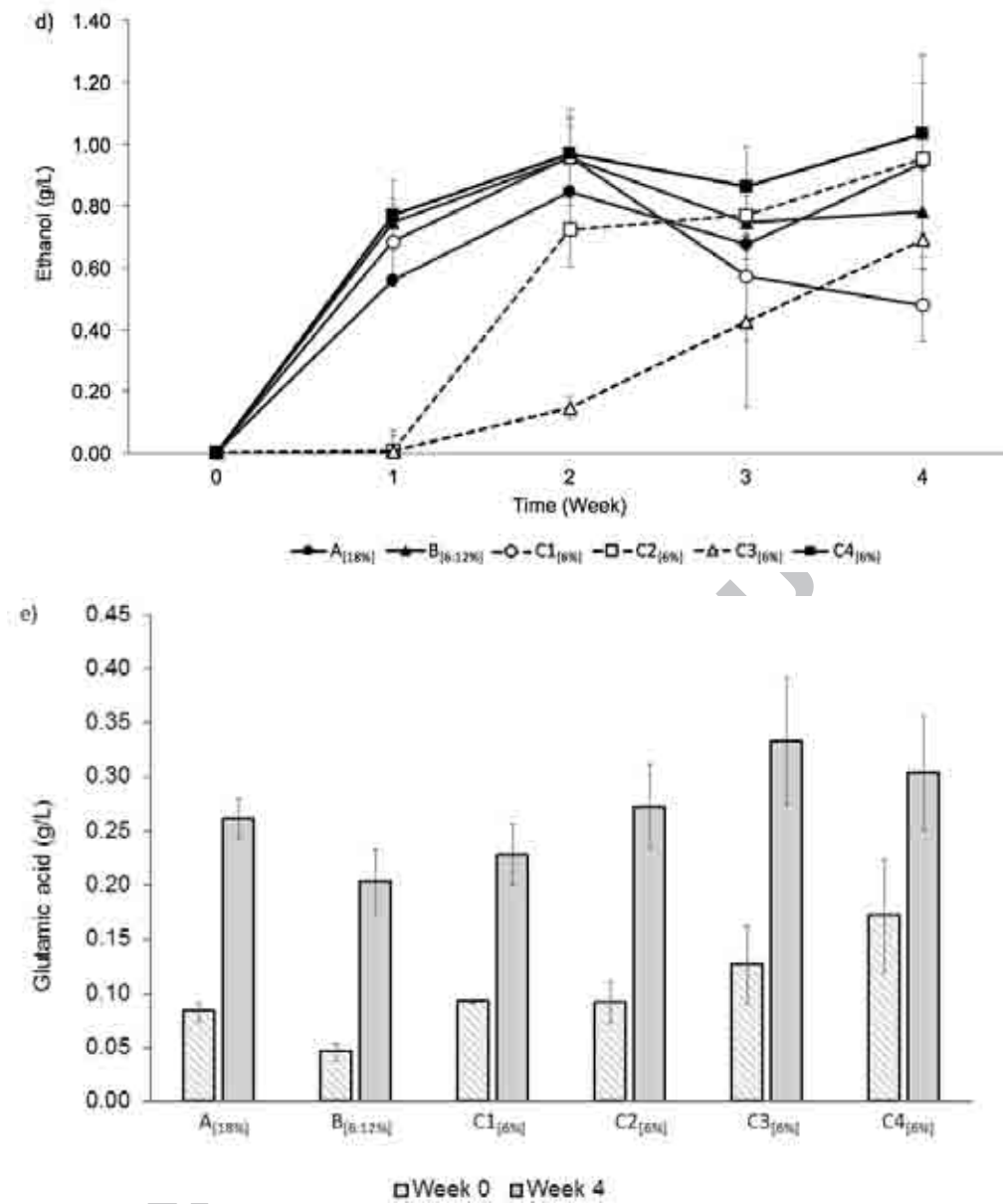
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617 Figure 3. Changes in population of (a) *T. halophilus* and (b) *Z. rouxii* during fermentation of
 618 low and high salt moromi at 30 °C. The samples contained co-inoculated *T. halophilus* and *Z.*
 619 *rouxii* in 18% NaCl (A_[18%]), 6% NaCl and 12% KCl (B_[6:12%]), 6% NaCl (C1_[6%]), and
 620 sequentially inoculated *T. halophilus* and *Z. rouxii* at week 1 (C2_[6%]), week 2 (C3_[6%]), or
 621 with DE (C4_[6%]). The addition time of *Z. rouxii* cells for sequential inoculation is indicated
 622 by the asterisk mark (*). Means within the same group with different letters (a, b, c) are
 623 significantly different (p<0.05).

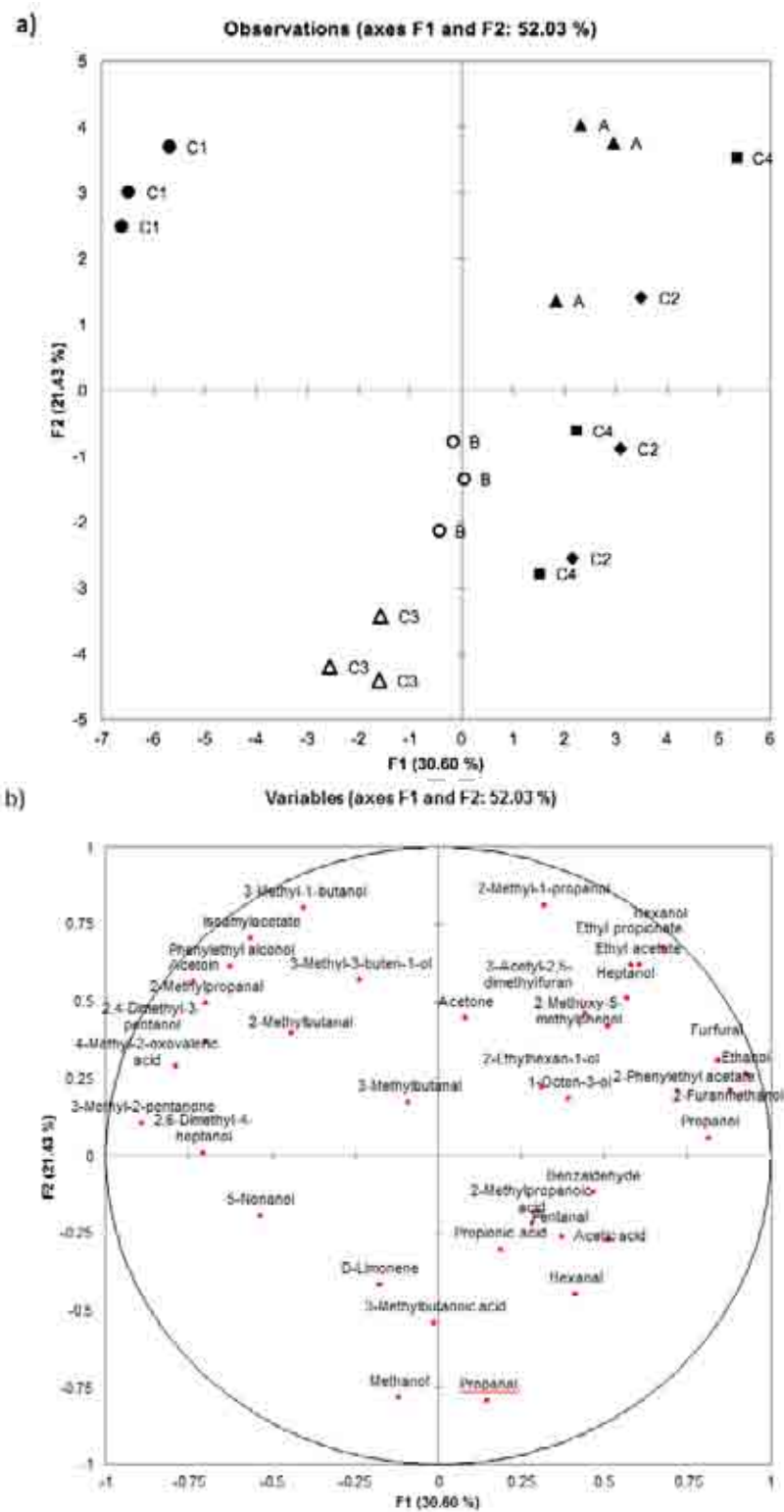




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626 Figure 4. Changes in (a) pH, (b) total reducing sugar, (c) lactic acid, (d) ethanol, and (e)

627 glutamic acid during fermentation of low and high salt moromi at 30 °C.



628 Figure 5. (a) PCA score plot of six moromi samples after 4-week fermentation. The scores
 629 are based on three replicates of each sample. The identical symbols represent triplicate
 630 measurements. (b) PCA loading plot of the aroma compounds detected in moromi after 4-
 631 week fermentation.

632

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635 **Table 1.** Aroma compounds found in moromi after 4 weeks of fermentation in low and high
 636 salt concentration. The values are relative to the peak area observed when the headspace
 637 above a 0.1 µg/mL 1-octen-3-ol solution was analyzed. Each value is based on three
 638 replicates.

Compound	LRI ¹	Day 30											
		A _[18%]		B _[6:12%]		C1 _[6%]		C2 _[6%]					
		mean	SD	mean	SD	mean	SD	mean	SD				
Alcohols													
1-octen-3-ol	1466	0.039	a	0.009	0.020	a	0.002	0.037	a	0.006	0.086	b	0.035
2-ethyl-1-hexanol	1508	0.029	a	0.003	0.028	ab	0.008	0.014	abc	0.007	0.025	abc	0.011
2-furanmethanol	1690	0.084	a	0.006	0.046	b	0.003	0.039	b	0.002	0.089	a	0.011
2-methyl-1-propanol	1115	0.385	a	0.055	0.186	bc	0.019	0.204	b	0.070	0.159	bc	0.030
2,4-dimethyl-3-pentanol	1395	0.013	a	0.001	0.019	a	0.001	0.298	b	0.203	0.021	a	0.004
2,6-dimethyl-4-heptanol	1506	0.031	ab	0.010	0.005	a	0.001	0.090	b	0.025	0.032	ab	0.006

3-methyl-3-buten-1-ol	1271	0.026	a	0.004	0.021	a	0.002	0.019	a	0.004	0.004	b	0.001
3-methyl-1-butanol	1225	25.061	a	3.033	18.089	ac	1.609	40.297	b	8.876	19.532	ac	3.675
5-nonanol	1473	0.002	a	0.000	0.001	a	0.000	0.006	b	0.003	0.005	ab	0.002
ethanol	950	33.709	a	2.159	27.538	ab	1.129	14.398	b	1.106	39.284	a	3.380
1-heptanol	1473	0.042	ab	0.006	0.016	a	0.002	0.016	a	0.002	0.015	a	0.006
1-hexanol	1371	0.071	a	0.012	0.033	ab	0.002	0.032	ab	0.010	0.046	ab	0.018
methanol	915	0.449	a	0.025	0.511	a	0.065	0.488	a	0.088	0.604	a	0.082
phenylethyl alcohol	1957	2.425	a	0.371	2.059	ac	0.121	3.801	b	0.352	2.010	ac	0.193
1-propanol	1057	0.376	ac	0.030	0.281	ab	0.023	0.180	b	0.030	0.468	cd	0.057
acids													
4-methyl-2-oxovaleric acid	1478	0.016	ac	0.005	0.006	a	0.000	0.040	b	0.006	0.012	ac	0.004
2-methylpropanoic acid	1596	0.113	a	0.046	0.057	ab	0.025	0.000	b	0.000	0.025	bc	0.007
3-methylbutanoic acid	1699	0.400	ab	0.153	0.205	a	0.081	0.008	a	0.012	0.116	a	0.103

acid

acetic acid	1481	0.161	ab	0.098	0.041	ab	0.040	0.000	a	0.000	0.212	ab	0.055
propionic acid	1565	0.010	a	0.010	0.018	a	0.023	0.015	a	0.025	0.061	a	0.090

aldehydes

2-methylbutanal	929	0.009	a	0.002	0.001	a	0.000	0.026	a	0.027	0.009	a	0.009
2-methylpropanal	824	0.026	a	0.009	0.009	a	0.002	0.127	b	0.057	0.027	a	0.022
3-methylbutanal	934	0.083	a	0.013	0.012	a	0.002	0.091	a	0.063	0.109	a	0.114
benzaldehyde	1568	0.021	a	0.004	0.034	ab	0.001	0.013	a	0.001	0.060	b	0.030
furfural	1500	0.014	ab	0.002	0.010	ab	0.001	0.009	a	0.000	0.015	b	0.003
hexanal	1104	0.001	a	0.001	0.002	a	0.001	0.001	a	0.000	0.003	a	0.001
pentanal	1001	0.014	a	0.007	0.015	a	0.007	0.015	a	0.004	0.029	a	0.010
propanal	807	0.004	a	0.000	0.005	a	0.001	0.003	a	0.001	0.004	a	0.002

esters

2-phenylethyl acetate	1860	0.316	ac	0.054	0.136	b	0.006	0.094	b	0.024	0.378	c	0.071
ethyl acetate	906	0.196	a	0.046	0.079	a	0.026	0.073	a	0.023	0.100	a	0.077

ethyl propionate	975	0.032	ab	0.008	0.006	a	0.000	0.010	ab	0.005	0.011	ab	0.009
isoamyl acetate	1141	0.256	a	0.074	0.029	a	0.006	0.878	b	0.280	0.115	a	0.119
furan													
3-acetyl-2,5-dimethylfuran	1450	0.186	a	0.061	0.001	bc	0.000	0.001	b	0.000	0.069	c	0.014
phenol													
2-methoxy-5-methylphenol	1614	0.258	a	0.081	0.001	b	0.000	0.001	b	0.000	0.169	a	0.035
ketone													
3-methyl-2-pentanone	1037	0.004	a	0.001	0.009	ac	0.001	0.032	b	0.011	0.000	a	0.000
acetoin	1318	0.656	a	0.039	0.451	a	0.043	1.624	b	0.572	0.066	a	0.013
acetone	829	0.671	a	0.088	0.334	a	0.082	0.507	a	0.084	0.405	a	0.107

Others

D-limonene	1217	0.006	a	0.003	0.001	a	0.001	0.004	a	0.004	0.002	a	0.001
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639 ¹LRI: linear retention indices of the compounds relative to an alkane series.

640 Means within the same row with different letters (a, b, c) are significantly different ($p < 0.05$)

641 A_[18%] : Co-inoculation; 18% NaCl

642 B_[6:12%] : Co-inoculation; 6% NaCl and 12% KCl

643 C1_[6%] : Co-inoculation; 6% NaCl

644 C2_[6%] : Sequential inoculation starting at Week 1; 6% NaCl

645 C3_[6%] : Sequential inoculation starting at Week 2; 6% NaCl

646 C4_[6%] : Inoculation with DE; 6% NaCl

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649 Highlights

- 650 • First study to utilize W₁/O/W₂ double emulsion (DE) in low-salt moromi
651 fermentation
- 652 • DE stability was dependent on but not proportional to moromi viscosity
- 653 • DE was utilized to control the inoculation of soy sauce starter cultures
- 654 • Volatile profile of low-salt moromi fermented with DE resembled that of high-salt
655 sample
- 656 • Sequential inoculation affected fermentation and volatile compounds formation

657

ACCEPTED MANUSCRIPT