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**Pre-fermentation of malt whisky wort using *Lactobacillus plantarum* and its influence on new-
make spirit character**

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22 **Abstract**

23 Distillery fermentations are non-sterile, which allow bacterial communities to flourish, typically
24 towards the end of fermentation. The effect of beginning the bacterial fermentation at the start of
25 fermentation was investigated. Wort was treated for 48 hours using a commercial strain of
26 *Lactobacillus plantarum* followed by fermentation using a distilling strain of *Saccharomyces cerevisiae*.

27 The treated wash showed a substantial increase in lactic, acetic and succinic acids Sensory analysis
28 determined that the spirit produced with bacterial treatment were significantly different ($p < 0.05$) and
29 chemical analysis demonstrated an increase in the production of ethyl acetate.

30 These results show that pre-treatment using species of *Lactobacillus* could be utilised to alter the
31 quality of new-make spirit in a distillery. By using bacterial cultures present in the surroundings or raw
32 materials, distillers could allow naturally occurring or commercially available microflora to be added
33 thus enhancing flavour development during fermentation and producing different spirit characters.

34 **Words: 150 (limit 150)**

35

36 **Keywords**

37 Whisky, fermentation, organic acids, bacteria, *Lactobacillus plantarum*

38 1. Introduction

39 During the production of malt whisky, there is no boiling stage during wort production, resulting in
40 'non-sterile' conditions for fermentation. Therefore, bacteria and other microflora, derived from the
41 local environment (dust, mash tun, washbacks (fermentation vessel)) or from the raw materials (malt,
42 cereals, yeast) (O'Sullivan, Walsh, O'Mahony, Fitzgerald, & Sinderen, 1999), can develop throughout
43 fermentation (van Beek & Priest, 2002, 2003). These microorganisms, along with the desired strain of
44 *Saccharomyces* yeast, influence the flavour of the fermented wort – known as wash once fermentation
45 is complete. In a typical malt whisky distillery, this wash will be distilled twice in copper pot stills to
46 produce new-make spirit.

47 Despite bacterial cultures and wild yeasts typically being associated with beer spoilage (Hammond,
48 Brennan, & Price, 1999; Suzuki, 2015; Vriesekoop, Krahl, Hucker, & Menz, 2012), the growth of
49 bacteria and their fermentation in Scotch whisky fermentations is often anecdotally linked with
50 enhanced new-make spirit quality and the taxonomy of the various bacterial cultures present has been
51 well studied in the industry (Cachat, 2005; Mekanjuola & Springham, 1984; Simpson, Pettersson, &
52 Priest, 2001; van Beek et al., 2002, 2003; Wilson, 2008). It is not common for Scottish distilleries to
53 encourage the presence of such cultures as they are enabled through the absence of wort boiling. Also
54 the use of wooden washbacks which harbour higher levels of bacteria are commonplace.
55 Furthermore, distillers will often increase fermentation time to encourage a secondary, lactic,
56 fermentation (Lee, Paterson, Piggott, & Richardson, 2001). Elsewhere in the world, more typically in
57 the United States, a portion of spent mash (processed grain, low in carbohydrates) from the previous
58 fermentation may be included in the fermenter, reducing the pH of the wort which will encourage the
59 growth of lactic acid bacteria during fermentation (Buglass, 2011). Although the growth of bacteria
60 may not be entirely eradicated, it is generally maintained at low levels so not to impact the spirit yield
61 from a given fermentation either through competition for fermentable sugar (Barbour & Priest, 1988)
62 or by initiating yeast flocculation (Mekanjuola, Tymon, & Springham, 1992).

63 Detailed studies of bacterial communities present in distillery environments have been carried out
64 (Simpson et al., 2001; van Beek et al., 2003), with species of *Lactobacillus*, found to be the dominating
65 genus due to its ability to survive the high alcohol content of late whisky fermentations (7.5 – 8.5 %
66 v/v ethanol), low pH (typically 3.5-4.0) and anaerobic conditions (Piggott, Sharp, & Duncan, 1989).
67 Despite this field being well studied, the specific impact that these cultures have on new-make spirit
68 is poorly understood. This could be due to several different factors. It is challenging to determine
69 differences between spirit quality from distillery to distillery as a myriad of other factors also influence
70 the perceived quality of new-make spirit. This includes, but is not limited to, mashing regime, yeast
71 pitching (addition) rate and strain, fermentation volume, time and temperature as well as still design
72 and cut points during distillation. Furthermore, researchers have established that bacterial cultures
73 typically thrive towards the end of fermentation, when the specific gravity (SG) is low (below 1.000)
74 (Rose, 1977; Wilson, 2014). Late in fermentation there are limited nutrients to be metabolised, these
75 are typically derived from the autolysis of dead yeast cells. Traditionally it has been understood by
76 distillers, and confirmed by research, that if the bacterial growth phase occurred too early in
77 fermentation it would be detrimental to ethanol production and therefore, final spirit yield (Barbour
78 et al., 1988; Dolan, 1976). The duration of fermentations carried out by several distillers is often linked,
79 specifically in marketing materials, with fruity, estery and waxy character in the new-make spirit. This
80 concept is loosely backed up by literature and argues that bacteria may increase the ester content of
81 the distilled spirit (Geddes & Rifkin, 1989; Priest, van Beek, & Cachat, 2002). The specific influence of
82 the bacterial species is difficult to prove both in the lab (due to a lack of indigenous microflora) or at
83 the industrial scale (due to variances between distilleries). By moving the bacterial fermentation prior
84 to the pitching of *S. cerevisiae* the impact of *Lactobacillus* on new-make spirit character can be
85 determined.

86 Due to process intensification, the replacement of wooden washbacks with stainless steel vessels and
87 more robust cleaning practices in many whisky distilleries for product control purposes, bacterial
88 growth and subsequent impact on new-make spirit quality may be limited. Links between washback

89 material and spirit quality are often made, although lacking in scientific evidence, linkages have been
90 postulated between wooden washbacks and higher content of fruity esters due to the prevalence of
91 bacteria within the more highly porous wooden washbacks (Yonezawa & Stewart, 2004).

92 Strains of distillery microflora, surviving fermentation conditions, can be categorised as
93 homofermenting (*Lactococcus*, *Pediococcus* and *Streptococcus*), heterofermenting (*Leuconostoc* and
94 *Lactobacillus*) (Von Wright & Axelsson, 2004; Wilson, 2014). Homofermentative species use the
95 glycolytic (or Embden–Meyerhof–Parnas) pathway and produce 2 mol lactic acid from 1 mol glucose
96 (Burgé, Saulou-Bérion, Moussa, Allais, Athes, & Spinnler, 2015, Wilson, 2014). Facultative
97 heterofermenting strains can metabolise glucose via the 6-phosphogluconate/phosphoketolase (6-
98 PG/PK) pathway which produce either 2 mol of lactic acid or 1 mol ethanol and carbon dioxide per
99 mol of glucose depending on prevailing conditions.

100 Other metabolism pathways of *Lactobacillus* have also been shown to form succinic acid from lactic
101 or citric acid (Dudley & Steele, 2005; Zalán, Hudáček, Štětina, Chumchalová, & Halász, 2010) and also
102 play a role in the decarboxylation of cinnamic acids (van Beek & Priest, 2000) all of which may influence
103 new make spirit character

104 Production of metabolism by-products, particularly, organic acids could have a substantial impact on
105 the flavour and aroma characteristics of the distilled wash. Due to the presence of alcohols such as
106 ethanol and higher alcohols (butanol, isoamyl alcohol) and low pH conditions, condensation reactions
107 between these alcohols and acids can result in esterification (Eqn 1). This effect has been shown in
108 laboratory experiments (Swan & Burtles, 1978).



110 Equation 1 – Generalised condensation reaction for ester production in acidic conditions

111 Furthermore, the volatility of esters has been shown to align well with the distillation profile of ethanol
112 in both experimental and model systems, increasing the likelihood that these compounds will have an
113 impact on flavour in the spirit cut of a distillation run (Ikari & Kubo, 1975).

114 Esters typically have a lower vapour pressure than their corresponding alcohols and acids and
115 therefore, these products will be more readily distilled, thus driving the equilibrium of this reaction to
116 the right by Le Chatelier's principle. Esters have a large impact on new-make spirit quality due to their
117 low volatility and low aroma thresholds contributing fruity, sometimes floral aromas to new-make
118 spirit (Buxton & Hughes, 2013; Christoph & Bauer-Christoph, 2007). Thus, an understanding of the
119 production of such congeners could lead to a greater understanding of how the behaviour of bacteria
120 species in a malt whisky distillery affect new-make spirit character.

121 Organic acids may also be produced from the tricarboxylic acid (TCA) cycle. Typically, this occurs early
122 through the action of yeast during the lag phase of fermentation. Changes in the organic content of a
123 rice and malted barley based beer have been studied, showing that acetic, lactic and succinic acids all
124 increase largely during fermentation (Li & Liu, 2015). Generally, the release of organic acids only occurs
125 when the TCA cycle is not completed and therefore, could also serve as an indicator of poor cell growth
126 or heightened yeast stress. The organic acid content of beers has been studied and shown that levels
127 of acetic, lactic and succinic acids all increase during fermentation with malic and fumaric acids
128 remaining stable however, there is limited information on the organic acids formed during malt whisky
129 fermentations (Li et al., 2015).

130 As detailed in the next section of this paper, a high-performance liquid chromatography (HPLC)
131 method was developed to investigate the concentration of organic acids in fermenting wort. Initially,
132 a commercial strain of *Lactobacillus plantarum* was inoculated into distillers' wort. Following three
133 bacterial treatments, a control (CP), where no bacterial treatment occurred, a "sterile" treatment (SS)
134 where bacteria was pitched into boiled wort and "non-sterile" treatment (US) where bacteria was
135 pitched into unboiled wort. A commercially available distilling strain of *Saccharomyces cerevisiae* was

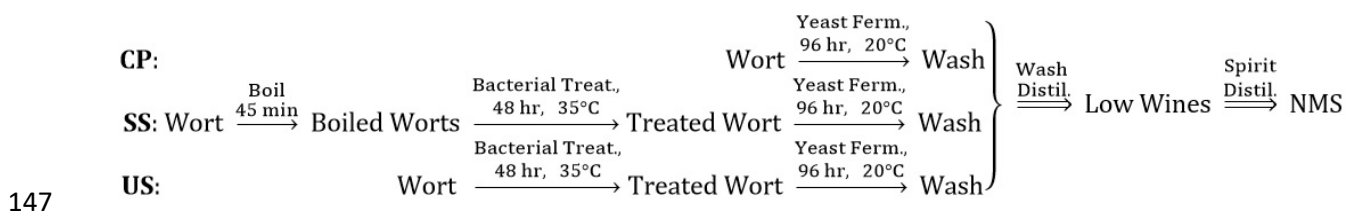
136 then pitched into the wort. Fermentation data was collected and modelled using a 4-parameter
137 logistic equation to determine any differences in fermentation profiles. The fermented wash was then
138 first distilled to produce low-wines and then subsequently to produce a new-make spirit which was
139 subjected to chemical and sensory analysis.

140 **2. Materials and Methods**

141 All specific gravity readings were measured by filtering and degassing the sample followed by
 142 measurement using an Anton-Paar DMA 35 (Anton-Paar, St Albans, GBR). The pH was recorded using
 143 a VWR pH110 meter (VWR, Lutterworth, GBR) and all fermentation samples (pre-distillation aliquots)
 144 were frozen prior to analysis.

145 *2.1 Experimental Plan*

146 An outline of the experimental procedure is shown in figure 1.



148 Figure 1 – Experimental outline, showing the three spirit producing regimes employed. All wash
 149 treatments were subjected to the same experimental procedures to produce low wines and new make
 150 spirit (NMS).

151 *2.2 Wort Production*

152 Wort (200 L) of specific gravity 1.066 was mashed using 55 kg distillers malted Concerto barley
 153 (Simpsons Malt, Berwick-upon-Tweed, GBR) in the International Centre for Brewing and Distilling
 154 (ICBD) pilot brewery using a two-roller mill (Fraser Agricultural, Inverurie, GBR). The grain was milled
 155 and then mashed with 150 L of water at 64 °C for 1 hour prior to separation using a lauter tun. The
 156 grain bed was then sparged continuously with 81 °C water to obtain the desired volume and gravity.
 157 The produced wort was frozen in 20 L aliquots for future use. All wort was thawed prior to use.

158

159

160 *2.3 Wort Treatment*

161 Two types of treated wort were produced in triplicate – a “sterile sour” wort (SS), produced by boiling
162 the mixture for 45 min and siphoned into a fermentation vessel to remove any hot break formed. An
163 “unsterile sour” wort (US) which was not boiled prior to pitching the bacteria. The bacterial culture
164 was a commercially available freeze-dried strain of *Lactobacillus plantarum* (Lallemand Inc.,
165 Felixstowe, GBR). Bacteria were pitched according to the manufacturer’s instructions at a rate of 1 g
166 hL⁻¹ and wort was soured for 48 hrs at 35 °C. The control wort used (CP) was treated without bacteria
167 and was not boiled prior to use.

168 *2.4 Fermentation*

169 A commercially available strain of dried *Saccharomyces cerevisiae* (N379) distiller’s yeast (Lallemand
170 Inc., Felixstowe, GBR) was pitched directly into the untreated control (“control pitch”, CP) and two
171 treated worts (i.e., an unsterile (US) and sterile (SS)). All fermentations were pitched with yeast at a
172 rate of 1 g L⁻¹ (approximately 2 x10⁷ cells mL⁻¹). The yeast was pitched directly into the wort at 20 °C
173 and was allowed to ferment for 96 hours. This temperature inhibited further bacterial fermentation
174 and encouraged the domination of *S. cerevisiae*. The fermented wash was filtered and analysed using
175 an Anton-Paar DMA 4500 coupled to an AlcoLyzer ME module (Anton-Paar, St Albans, GBR) to
176 determine final gravity and alcohol by volume (%v/v ethanol).

177 *2.5 Distillation*

178 Fermented wash was distilled on ICBD’s pilot distillery. During all distillation runs the % v/v ethanol of
179 the distillate was recorded to ensure consistent profiles were achieved to avoid alteration of flavour.
180 Stills were charged with 20 L (or 18 L with SS boiled wort) wash. Wash was heated using a steam coil
181 and a steady run off of low-wines was achieved. The wash distillation was ceased when the ethanol
182 level of the distillate reached 1 %v/v (an economic cut point).

183 Spirit distillation was carried out in 5 L copper Al-Ambiq stills (Al-Ambiq, Gandra, PRT) using a worm-
184 tub style condenser and heated using a 5 L isomantle (Electrothermal, Staffordshire, GBR) at a

185 constant heat setting. At the start of this second distillation foreshots (50 ml) were collected and then
186 the spirit cut was collected by allowing the %v/v ethanol of the distillate to fall by 12 % before halting
187 the collection of new make spirit.

188 *2.6 Chemicals*

189 All chemicals were supplied at purity $\geq 98\%$ (Sigma-Aldrich, Dorset, GBR). HPLC eluent was produced
190 using distilled water which was further purified and degassed by vacuum filtration (0.45 μm).

191 *2.7 Organic Acids Analysis*

192 Methods were adapted for wort and wash analysis from those found in the literature (Li & Liu, 2015;
193 Lopez & Gomez, 1996; Park, Shin, Lee, & Lee, 2017). Organic acids were analysed using a Shimadzu
194 LC-20AD (Shimadzu, Tokyo, JPN) chromatogram fitted with SIL-20A HT autosampler and a CTO-10AS
195 column oven coupled to a SPD-M20A diode array detector. Two Phenomenex Kinetex[®] (Phenomenex,
196 Macclesfield, GBR) 150 x 4.6 mm C18 columns were used as the stationary phase and a H₃PO₄ (5 mM,
197 pH 2.1) solution was used as the mobile phase. A flow rate of 0.65 ml min⁻¹ was achieved with a column
198 oven temperature of 60 °C. A 10 μL sample was injected and the run time was 30 minutes. All
199 compounds were detected at wavelength of 210 nm.

200 Formic, malic, lactic, acetic, fumaric and succinic acid were used as external standards. Gallic acid (10
201 g L⁻¹) was used as an internal standard by addition of 10 μL to 1 ml of sample, giving a working
202 concentration of 100 mg L⁻¹ and was undetectable in representative samples of wort and wash.
203 However, the compound may be found in low quantities (far below the instruments limit of detection)
204 in beer, potentially due to the addition of stabilisers which help to precipitate polyphenols and
205 proteins (Gerhäuser, 2005; Leiper, Stewart, McKeown, Nock, & Thompson, 2005). Standards were
206 made in ranges specific to their typical concentration found in wort and wash and were reflective of
207 the amount found in beer samples based on peer-review reports (Li & Liu, 2015). All standard curves

208 had a coefficient of determination (r^2) ≥ 0.999 and repeatability in wort, wash and beer of relative
209 standard deviation $<10\%$.

210 Samples were centrifuged at 4500 rpm for 15 min and 1 ml of the centrifuged media was transferred
211 into a sample vial containing 100 μl of gallic acid internal standard.

212 *2.8 GC-FID Analysis*

213 Higher alcohols were analysed using an Agilent 7820 A operating in liquid injection mode (Agilent,
214 California, USA). Separation was achieved using an Agilent DB-WAX 30 m x 0.25 mm x 0.25 μm column.
215 The oven started at 35 $^{\circ}\text{C}$ with a 6 min hold, temperature was ramped up to 60 $^{\circ}\text{C}$ at a rate of 5 $^{\circ}\text{C}$
216 min^{-1} with a 2 min hold followed by a ramp to 210 $^{\circ}\text{C}$ at a rate of 20 $^{\circ}\text{C min}^{-1}$ and a final ramp to 250
217 $^{\circ}\text{C}$ at a rate of 70 $^{\circ}\text{C min}^{-1}$ with a hold of 1 min. Compounds were detected using an FID detector at
218 270 $^{\circ}\text{C}$.

219 A 1 ml sample aliquot was added to a vial, followed by 100 μl of pentan-1-ol to give a final
220 concentration in vial of 500 mg L^{-1} . From this vial, 1 μl was injected on to the GC column. Congeners
221 analysed were acetaldehyde, ethyl acetate, acetal, methanol, n-propanol, isobutanol, isoamyl acetate,
222 n-butanol, isoamyl alcohol and furfural. Calibration ranges were in the range of 5 – 1250 mg L^{-1} with
223 the exception of isoamyl alcohol which was measured in a range of 10 – 2500 mg L^{-1} . All calibrations
224 had $r^2 > 0.99$.

225 *2.9 GC-MS Analysis*

226 Ester analysis was conducted on a Shimdadzu QP2010 Ultra GC/MS with AOC 5000 autosampler
227 (Shimdadzu, Tokyo, JPN) using a 65 μm PDMS/DVB solid phase microextraction (SPME) fibre (Supleco,
228 Pennsylvania, USA). A sample (1 ml) was added to a headspace flask and mixed with distilled water (5
229 ml). Methyl heptanoate (10 μL , 25 mg L^{-1}) was used as an internal standard. The sample was incubated
230 for 5 min, followed by 5 min of adsorption of analyte by exposing the fibre to the headspace of the
231 sample vial. Sample was then desorbed onto a GC column for 1 min at 200 $^{\circ}\text{C}$.

232 Separation was achieved on a DB-WAX 30 m x 0.25 mm x 0.25 μm column (Agilent, California, USA)
233 the oven program ran 40 $^{\circ}\text{C}$ for 3 min, followed by an initial ramp to 100 $^{\circ}\text{C}$ at 10 $^{\circ}\text{C min}^{-1}$ and a second
234 ramp to 160 $^{\circ}\text{C}$ at 4 $^{\circ}\text{C min}^{-1}$. The final ramp raised the oven temperature to 220 $^{\circ}\text{C}$ at a rate of 10 $^{\circ}\text{C}$
235 min^{-1} and held for 10 min. The mass spectrometer was operated in both SIM and SCAN mode. SIM ions
236 used were m/z 45, 55, 74, 88, 101, 104, 129. The SCAN mode ranged from m/z 42 – 400.

237 *2.10 Sensory Analysis*

238 An untrained panel ($n=12$) was used to evaluate the olfactory aspects of the spirit samples produced.
239 All samples were presented blind to the panellists after dilution to 20 % v/v ethanol.

240 A triangle test was used to determine if a significant statistical difference existed between the
241 produced spirits. Three different sets consisting of various combinations of the new-make spirit
242 samples were presented to the panellists. Each set contained two spirits – two of the same new-make
243 sample and one different. Candidates were asked to choose which sample they believed to be
244 different. The test was carried out as stated in the guidelines of BS EN ISO 4120 (British Standard,
245 2004). From the results, the statistical confidence interval was determined to prove that a difference
246 between samples could be determined by the panel.

247 Consumer panel testing was then carried out using the same sensory analysts. Ten sensory
248 characteristics were selected using the revised Scotch whisky flavour wheel that were deemed
249 appropriate for new-make spirit and recognised by the Scotch Whisky Research Institute for both
250 trained panel testing and consumer trials (Lee et al., 2001). Candidates were asked to rank each flavour
251 component on a scale from 0 to 5, where 0 was undetectable and 5 was strongly detectable. Finally,
252 candidates were presented with three anonymous samples and asked to rank their preference.

253

254 *2.11 Data Analysis*

255 To enhance understanding of the fermentations produced, the specific gravity (SG) was modelled in
256 RStudio (RStudio, Boston, USA) using the 4-parameter logistic equation (Eqn 2) as this has been shown
257 to effectively model fermentations in beer to prevent premature yeast flocculation (ASBC, 2012;
258 MacIntosh, 2015; MacIntosh, Adler, Eck, Speers, & Speers, 2012; MacIntosh, Josey, & Speers, 2016;
259 Speers, Rogers, & Smith, 2003).

$$260 \quad P_{(t)} = P_e + \frac{P_i - P_e}{1 + \exp(-B(t-M))} \quad [2]$$

261 Equation 2 – The 4-parameter logistic equations used to model the specific gravity decline during
262 fermentation

263 Where: $P_{(t)}$ is the specific gravity at given time t ,

264 P_i is the initial asymptotic gravity,

265 P_e is the final asymptotic gravity (FG),

266 B is a function of the gradient at the point of inflection,

267 M is the point of inflection with respect to time.

268

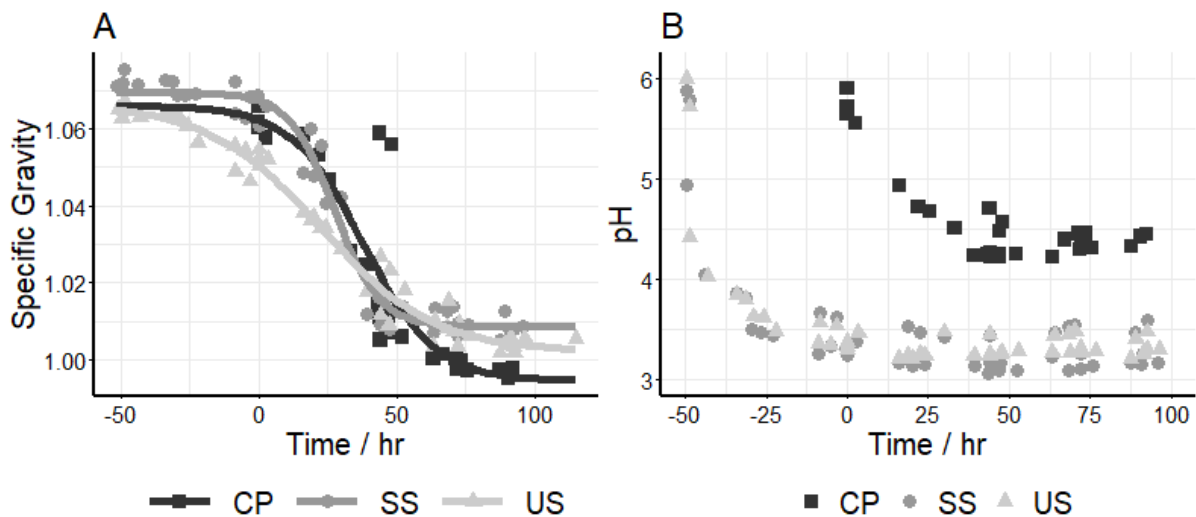
269 Data analysis was carried out in Microsoft Excel (Microsoft, Redmond, WA, USA) and OriginPro
270 (Originlab, MA, USA). All significance tested was conducted in RStudio (RStudio, Boston, MA, USA,
271 2019) with significance determined at the 95 % confidence interval.

272 3. Results and Discussion

273 3.1 Fermentation Data

274 The fermentation profiles are shown in Figure 2, and the coefficients to the applied logistic functions
275 (Supplementary 2) demonstrate that each fermentation followed a consistent sigmoidal pattern.
276 Fermentation of the unsterile wort appeared to have a shortened lag phase, with very little effect seen
277 with respect to the change in specific gravity upon pitching *S. cerevisiae*, however, the overall rate of
278 the exponential phase was the lowest ($B = 0.049 \pm 0.005$). Both treated worts did not fully attenuate
279 (Figure 2a), with the sterile treated wash fermenting 84.4 ± 2.0 % of the gravity (carbohydrates),
280 compared with the control wort and the unsterile treated fermenting to 94.4 ± 2.3 % of the control
281 wort. Part of this loss will be through the formation of biomass and consumption of carbohydrates by
282 the bacterial population which was to be expected.

283 Limited utilisation of all the nutrients in the wort is likely to be due to the prevailing conditions at the
284 end of fermentation. The average final pH measured in SS and US was 3.35 ± 0.22 and 3.34 ± 0.13
285 respectively. It is commonplace for distillery yeast strains to have a pH tolerance down to *ca.* 3.5 and
286 so it is likely that the pH limit of the yeast has been reached. Although treatment brought the pH below
287 this value for much of fermentation, it is likely that a loss of cell viability, caused by the combination
288 of increasing levels of alcohol and low pH, causes the yeast to cease activity. Higher ethanol
289 concentrations will increase the re-entry of protons into the cell membrane, reducing the pH within
290 the cytoplasm and inhibiting membrane function (Walker, 1999). It has been reported that *S.*
291 *cerevisiae* has various mechanisms to adapt to organic acid stress, however, it is likely that the
292 accumulation of such acids has exceeded the tolerance of this strain (Sugiyama, Sasano, & Harashima,
293 2015). A drop in yeast viability is commonly observed and may also be due to a rise in bacterial number
294 towards the end of fermentation such as *Lactobacillus delbreuckii* and *Lactobacillus paracasei* (Wilson,
295 2014).



296 Figure 2a & 2b – 2). a. (left) Trend in specific gravity during fermentation modelled by the 4-parameter
 297 logistic function (coefficients in Supplementary 2). b. (Right) Trend in pH during fermentation.

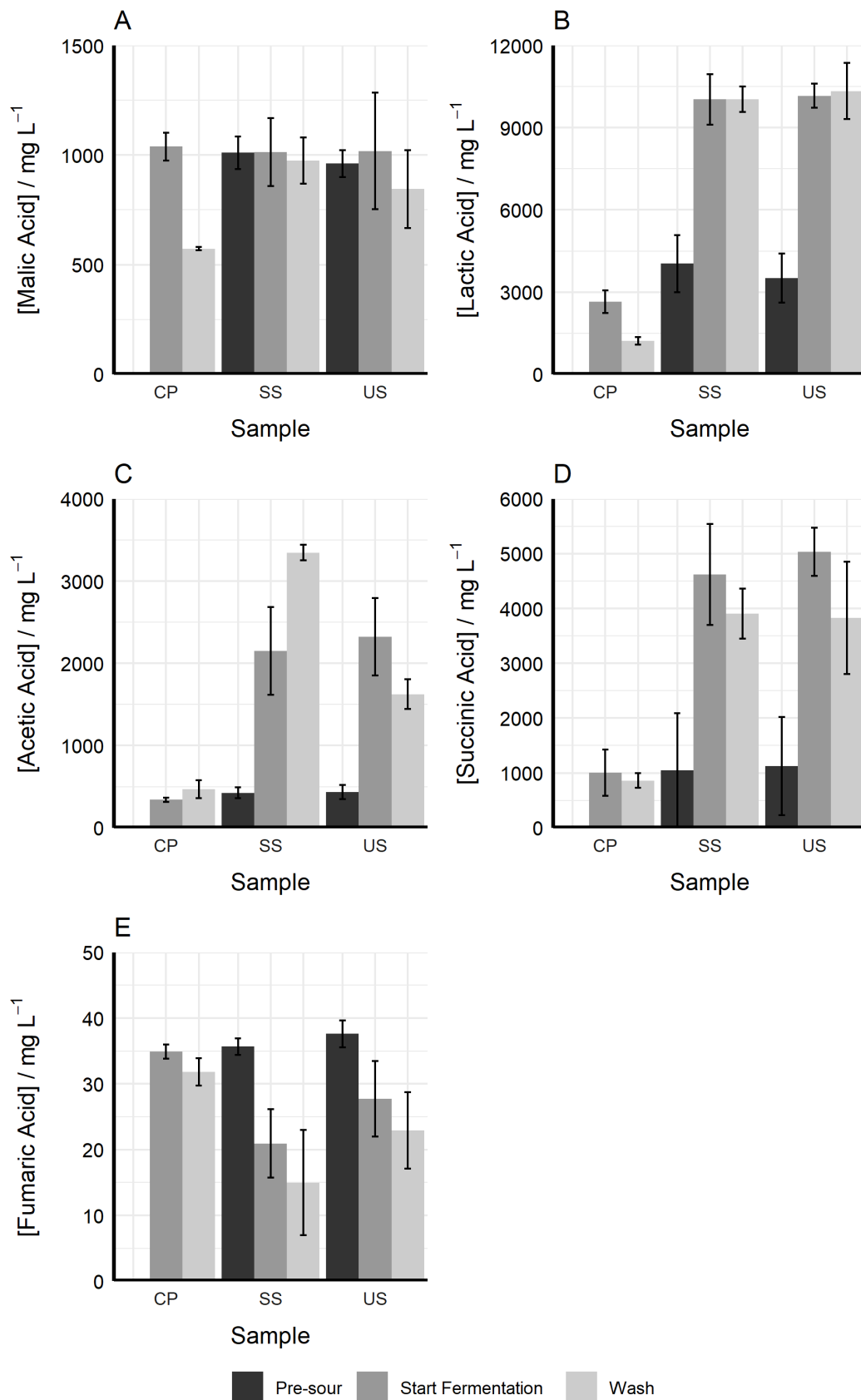
298 The presence of bacteria will also reduce the opportunity for yeast cell growth at the start of
 299 fermentation through the consumption of both dissolved O₂ and nutrients in the wort prior to yeast
 300 addition. As dried yeast was used, the impact of nutrient depletion will be less, however, as the yeast
 301 will have reduced viability in its dry form, this may have reduced cell growth after pitching. Cheung et
 302 al., have previously shown that yeast format has an impact on the yeasts ability to handle stressful
 303 conditions (Cheung, Brosnan, Phister, & Smart, 2012). This will increase the likelihood of stuck
 304 fermentations that do not fully attenuate. Bacteria (specifically *L. plantarum*) has been shown to
 305 increase the amount of flocculation occurring during the early stages of fermentations when both
 306 bacteria and yeast are pitched together under laboratory conditions (Makanjuola et al., 1992).
 307 Increased flocculation early in fermentation would increase the likelihood of an aberrant
 308 fermentation.

309 Fermentations with a low pH may be further limited due to a decline in enzymatic activity. Enzymes
 310 have been shown to continue to break down complex wort sugars throughout fermentation in whisky
 311 fermentations (Makanjuola et al., 1992; Vriesekoop, Rathband, MacKinlay, & Bryce, 2010). In the
 312 sterile (SS) sample, mashing enzymes will have been completely denatured during the boiling stage.
 313 Whereas, of the unsterile sour (US) fermentation, shifting the pH during fermentation (Figure 2b) may

314 have resulted in dextrin-degrading enzymes becoming inactive ceasing the breakdown wort sugars,
315 resulting in the observed final gravity (Bryce, McCaffery, Cooper, & Brosnan, 2002; Vriesekoop et al.,
316 2010).

317 3.2 Analytical Data - Fermentation

318 The organic acid profile during both treatment and fermentation is consistent among similar batches
319 of wort (Supplementary 3). Large differences, particularly during treatment, show that the levels of
320 lactic, acetic and succinic acids to increase in concentration up to five times (Figure 3, Table 1). Acetic
321 acid had the greatest percentage increase in both sterile and unsterile worts but lactic acid had the
322 highest final concentration of the acids measured (Figure 3). These concentrations of organic acids are
323 then carried forward into the subsequent fermentation with *S. cerevisiae*, however, levels of acetic
324 were found to be reduced at the start of the fermentation with yeast, particularly those in the US
325 sample – where a drop of around 1000 mg L⁻¹ during the first 24 hours of fermentation was observed
326 (Supplementary 3). *S. cerevisiae* has the ability to metabolise acetic acid and has been shown to do so
327 in lychee wine, particularly at 20 °C (Shang, Zeng, Zhu, & Zhong, 2016). Acetic acid may be taken up
328 by the cell, reacting with acetyl-CoA where it could form secondary metabolites, contributing to spirit
329 character such as higher alcohols, esters, fatty acids and other lipids. The trend of organic acids
330 throughout treatment and fermentation showed that acetic acid declines rapidly during the first 40
331 hours of fermentation when yeast is producing ethanol and other higher alcohols which could react
332 with the acetic acid, further supporting this postulation. The levels of acetic acid were observed to
333 increase towards the end of the SS fermentation (after 30 hours) however, it is suggested that this is
334 due to yeast autolysis and secretion of cellular compounds and organic acids into the wort.



335 Figure 3 – Measured concentrations of organic acids before treatment (pre-sour), in wort (start
 336 fermentation) and wash. Organic acids analysed were A) malic, B) lactic, C) acetic, D) succinic and E)
 337 fumaric. Error bars indicate the standard deviation on n = 3 samples.

338 Table 1 – Levels of lactic, acetic and succinic acid in wort and post treatment with *Lactobacillus* and
 339 differences in SS and US samples respectively.

Treatment	Acid	Average Concentration / mg L ⁻¹		% Difference in Concentration
		Wort	End of Treatment	
SS	Lactic	4039	10031	248
	Acetic	425	2150	505
	Succinic	1050	4621	440
US	Lactic	3507	10163	290
	Acetic	434	2325	536
	Succinic	1127	5033	446

340

341 The level of both lactic and succinic acid is consistently higher in the SS fermentation due to a reduced
 342 lag phase in the bacterial fermentation, a result of boiling the wort. The boiled wort will also have a
 343 large reduction in competing bacteria (certainly most if not all vegetative bacteria will be removed)
 344 and therefore pitched *L. plantarum* can be expected to dominate this phase of fermentation.

345 *3.3 Distillation Data*

346 The fermented wash was then distilled twice, the first distillation in a hybrid (copper and glass) still to
 347 produce low wines and then distilled in a copper pot still to produce new make spirit.

348 Low wines were found to demonstrate a significant decrease ($p < 0.05$) in litres of pure alcohol (LPA)
 349 between the control and treated samples, ranging from 1.79 LPA (control) to 1.46 LPA in the US wort.
 350 This demonstrates that during fermentation, less alcohol was produced from all wort samples pitched
 351 with bacteria, mainly due to the lack of fermentable sugars.

352 LPA of both foreshots and spirit cuts showed no significant difference ($p > 0.05$) between
 353 fermentations. This suggests that appropriate rectification was achieved on all distillation runs,

354 however, a reduction in spirit yield (LPA per kg malt) was observed from the fermentations using
355 treated (SS and US). This result was not unexpected as consumption of sugar by bacteria will lead to a
356 reduction of alcohol per molecule of sugar consumed (Wilson, 2014).

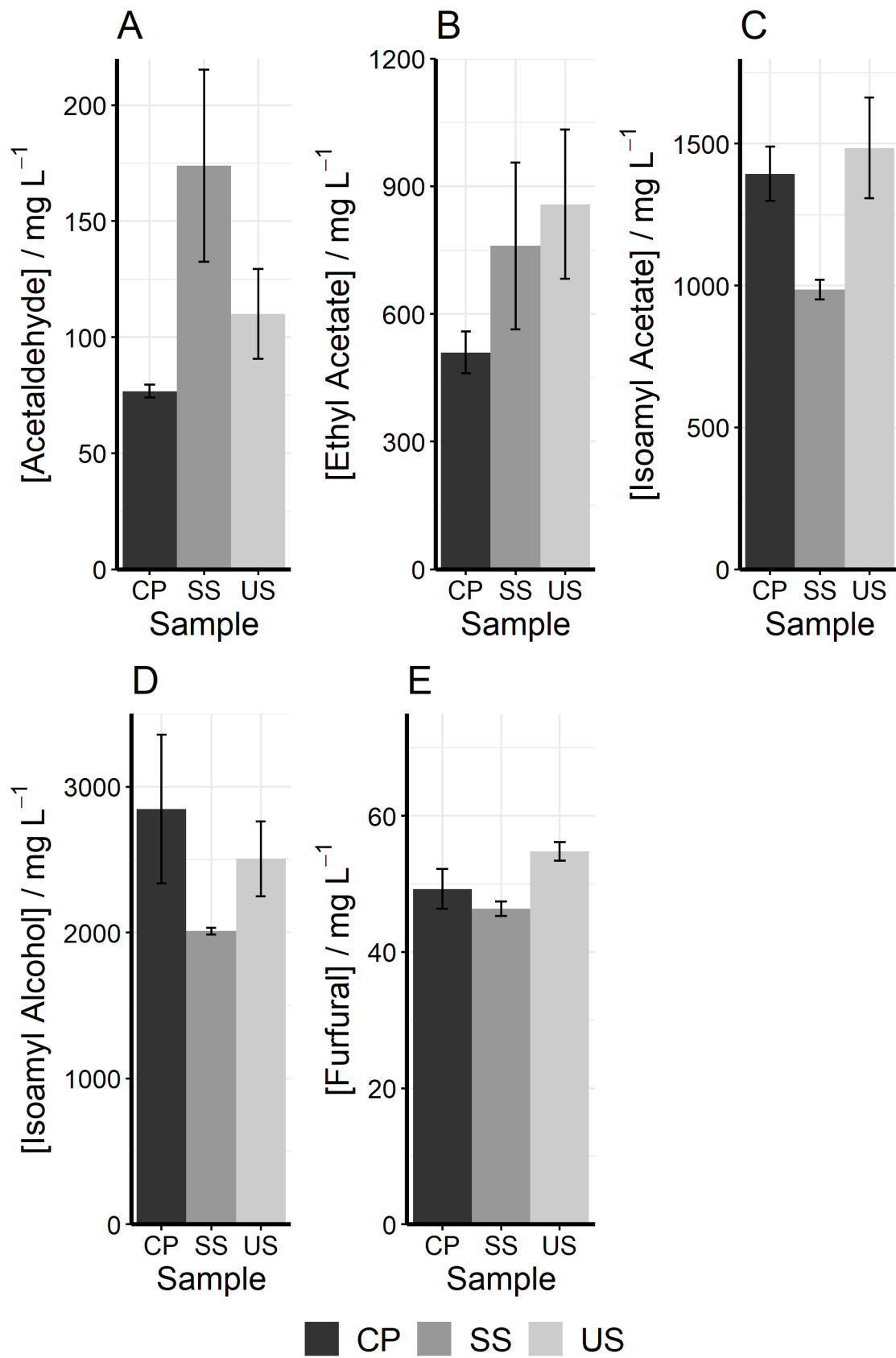
357 Chemical analysis of the distilled wash (low wines) showed an increase in concentration of ethyl
358 acetate, significantly ($p < 0.05$) in the US sample compared with the control ($103 \pm 23 \text{ mg L}^{-1}$ and $63 \pm$
359 13 mg L^{-1} respectively), with an increase in SS sample also observed ($79 \pm 22 \text{ mg L}^{-1}$). Furthermore, the
360 higher alcohols propanol, isobutanol and isoamyl alcohol were detected at lower concentrations in
361 soured low wines samples than in the control.

362 Chemical analysis from the second distillation (Figure 4) found that low levels of butanol were only
363 detected in the control sample at mean concentrations of 35.8 mg L^{-1} and 28.4 mg L^{-1} in the foreshots
364 and spirit cut respectively. Propanol was detected a concentration of 124.1 mg L^{-1} in the US spirit
365 sample (Figure 4).

366 In both soured fermentation regimes, levels of acetaldehyde and ethyl acetate were enhanced
367 substantially in comparison to those of the control sample. Acetaldehyde, associated an ethereal
368 aroma (Burdock, 2016) is not generally perceived as a positive attribute to new-make spirit.
369 Heightened levels of ethyl acetate in both sour samples are shown could be a result of initially
370 produced acetic acid (by *Lactobacillus*) becoming subsequently metabolised by yeast or may be
371 formed by esterification during one of the distillation stages. Typically, acetate esters confer fruity
372 notes in new-make spirit. These results concur with work by Ensor, Bryce, & Hill, (2015) who
373 demonstrated that higher concentrations of acetaldehyde and ethyl acetate are achieved when a
374 distillery fermentation is co-inoculated with *Lactobacillus* (Ensor, et al., 2015).

375 The reduction seen in higher alcohols may suggest that the organic acids produced during souring
376 treatment remove these higher alcohols in the form of esters, removing such analytes from the
377 samples produced. By changing the conditions during fermentation it is suggested that the metabolic

378 pathway of yeast changed and this led to a reduction in higher alcohols formed during treated
379 fermentations. Furthermore, reduced levels of amino acids (primarily due to metabolism by bacteria)
380 present in the wort may have influenced the amount of keto-acids produced within the yeast cell. This
381 would inhibit aldehyde formation, thus making producing less of the corresponding reduced product.
382 This would have had a positive influence on the aroma of the spirit.



383 Figure 4 - Concentration of key volatiles in spirit. Congeners analysed were A) acetaldehyde, B) ethyl
 384 acetate, C) isoamyl acetate, D) isoamyl alcohol and E) furfural. Error bars indicate the standard
 385 deviation on n = 3 samples.

386 Trace ester analysis of the main-cut fraction was also conducted using SPME-GC/MS. From the GC/MS
387 results, levels of all ethyl esters – excluding ethyl lactate and ethyl dodecanoate – were higher in the
388 control when compared with both soured samples (Supplementary 10).

389 This discrepancy somewhat disagrees with the literature on the subject (Geddes et al., 1989; Priest et
390 al., 2002), however, there are several aspects of the experimental set-up which may explain the
391 difference. Through the action of pre-souring the wort prior to fermentation by yeast, the yeast will
392 both produce less biomass due to a lack of nutrients available and, also due to the stressful
393 environment present from high levels of organic acids. Thus, levels of fatty acid esters such as ethyl
394 octanoate and ethyl decanoate – derived from the by-products of lipid synthesis and alcohols will be
395 reduced in the final distillate. Levels of ethanol are also higher in the control sample though this is
396 unlikely to have as great an impact as ethanol is the most abundant solvent in all samples.

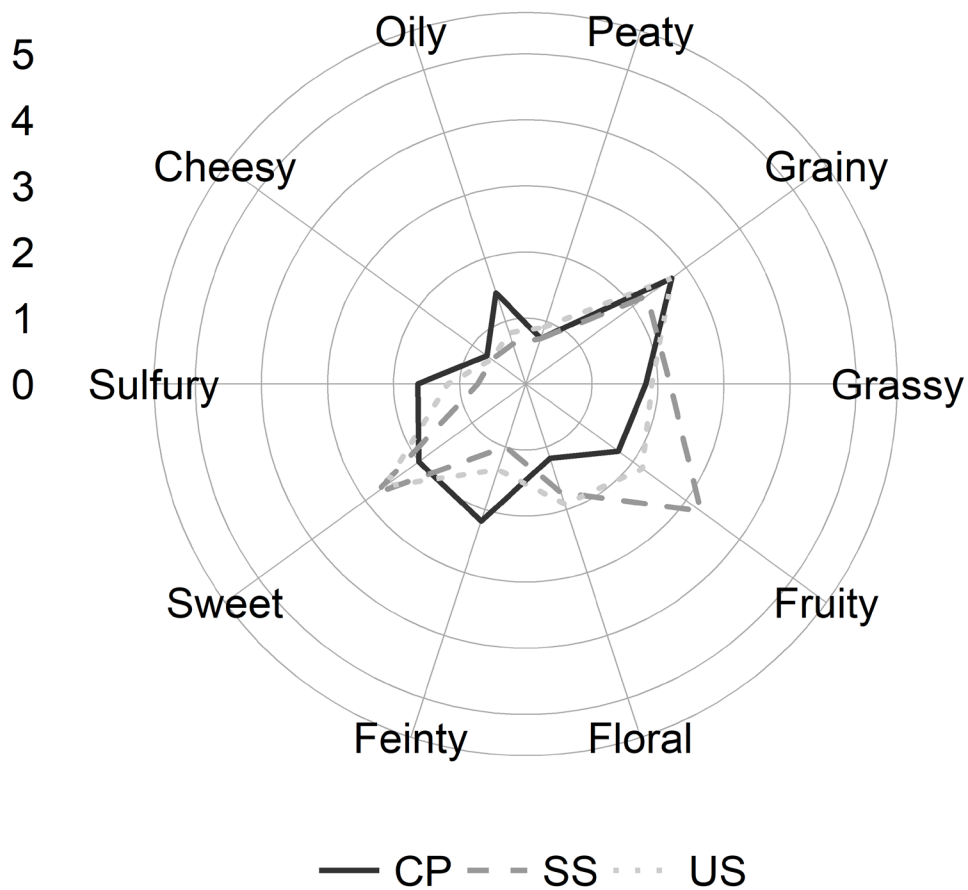
397 Levels of ethyl lactate were significantly ($p < 0.05$) higher in treated samples due to esterification of
398 the produced lactic acid with ethanol with levels in the control, SS and US samples being 3, 20 and 21
399 $\mu\text{g L}^{-1}$ respectively were observed. Similarly, phenylethyl acetate was shown to present at significantly
400 ($p < 0.05$) higher concentrations through similar reactions with phenylethyl alcohol this ester
401 contributes sweetness, floral and fruity characteristics to spirit (Burdock, 2016). Ethyl lactate confers
402 a waxy aroma, however, has not been seen to have a significant impact on the quality of new-make
403 spirit (Wilson, 2008).

404 Boiled wort will have a reduced level of proteins and other wort nutrients due to their removal during
405 the formation of hot break (precipitated proteins and other materials). Studies have shown that
406 reduced levels of such wort constituents increase levels of esters and promote the quality of the
407 subsequent spirit (Martin, 2002). Furthermore, production of esters and other metabolites derived
408 from yeast will be reduced in treated samples due to the reduction of fermentable sugar in comparison
409 with the control sample. Thus, the decrease in many levels of esters is to be expected. By boiling the
410 wort, wort sugars become more concentrated and the gravity (i.e., density) increases. Gravity

411 measurements in the SS sample at the end of treatment - t = 0 (Figure 2a) - is more similar to the
 412 original gravity of the control fermentation. As a result, the ester profile of the control and sample SS
 413 are similarly distributed.

414 *3.4 Sensory Data*

415 Figure 5 shows the average results of the quantitative descriptive analysis (QDA) conducted by the
 416 same panel.



417 Figure 5 – Spider diagram showing the mean result from the QDA of the new-make spirit samples
 418 produced, candidates ranked each component 0 - 5 on an arbitrary scale, n = 12.

419 Candidates selected spirit sample SS as their preference at 67 %, with a further 25 % selecting US as
 420 their preferred spirit. A final 8 % of candidates selected the control (CP) as their preference.

421 Sensory results depicted that all three spirits showed slight evidence of apparent differences between
422 samples when assessed by BS EN ISO 4120 (CI \geq 90 %, Supplementary 11) (British Standard, 2004).
423 Furthermore, evidence of a moderate apparent difference between the control sample and SS was
424 observed. It is postulated that a trained spirit sensory team is likely to have produced results with a
425 higher confidence interval, as random, untrained panels have been shown to lack consistency (Worch,
426 L , & Punter, 2010).

427 Further evidence for differences in spirit is shown by the quantitative descriptive analysis (Figure 5).
428 The panel detected greater fruity, floral and sweet aromas from both samples – attributes generally
429 associated with positive spirit character – with both soured samples. Typically, panellists ranked
430 sample SS higher in these categories compared with the US distillate. Studies involving *L. plantarum*
431 have shown previously that both the fruity and floral notes have been enhanced in new-make spirit
432 with inclusion of this species during fermentation (Takatani & Ikemoto, 2002). Heavier characteristics,
433 such as feinty, oily and sulfury were more generally associated with the control sample.

434 Following sensory testing 67 % of panellists preferred the sterile sample, with a further 25 % choosing
435 the unsterile sour sample and one panellist (8 %) choosing the control sample. Although the
436 preference test is subjective, as no instruction was given on what to base their opinion on, it does
437 demonstrate that preference correlated with positive attributes mentioned.

438 4. Conclusion

439 By use of a pre-fermentation treatment step to lower pH (souring), using a commercial strain of *L.*
440 *plantarum*, the organic acid profile of a typical Scotch whisky distillery wash was influenced. The
441 bacterial species produced lactic, acetic and succinic acids whose concentrations remained high
442 through subsequent fermentation using a known strain of *S. cerevisiae*. Souring significantly reduced
443 the pH of fermenting media and so, fermentability of the wort was reduced. Although the yields of
444 alcohol are significantly lower in the soured samples, a shorter souring period or earlier yeast pitching
445 time could be employed, which has the potential to reduce the final pH and allow the yeast to remain
446 viable and active until fermentation is completed.

447 The distillate of soured samples showed higher levels of acetate esters, particularly ethyl acetate and
448 phenylethyl acetate. Ethyl lactate – produced via the reaction of ethanol and lactic acid – was also
449 detected at elevated concentrations in treated distillates, depicting that organic acids produced by
450 the bacterial species can increase ester concentration in spirit. Levels of higher alcohols were also
451 reduced compared with a non-soured control.

452 A sensory panel determined that the three regimes employed gave spirit that was statistically different
453 by use of a triangle test. Soured samples produced spirit of a sweet, floral and fruity characteristic.

454 Organic acids produced during fermentation may limit production of such esters during distillation. By
455 incorporating yeast or bacterial strains or fermentation conditions which encourage the production of
456 such acids (such as acetic) may produce distillate of a quality that is perceived as having increased
457 fruity and sweet notes. It is likely that the sensory contribution that bacteria has on new-make spirit
458 is one of both primary effects via metabolism and also secondary effects on the fermentation of yeast
459 (Takatani et al., 2002).

460 Although the Scotch Whisky Technical File (DEFRA, 2013) does not permit the specific addition of
461 bacterial species to wort directly, this process may have use in other markets where such restrictions

462 do not apply (CAN, IND, JPN or USA) thus giving enhanced control over the bacteria which may
463 inoculate their malt whisky production. Furthermore, organic acid producing yeasts are becoming
464 increasingly popular amongst craft brewers as an alternative to standard kettle souring bacteria. As
465 the Scotch whisky regulations do not specify which strain of yeast must be used to produce whisky
466 this work shows that yeast strains with the ability to produce organic acids may be of benefit to the
467 whisky new-make spirit quality.

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470 treatments were subjected to the same experimental procedures to produce low wines and
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486 differences in SS and US samples respectively.

487

488 **Conflict-of-interest statement**

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490 **References**

- 491 ASBC. (2012). Yeast-14 - Miniature fermentation assay. *ASBC Methods of Analysis, Yeast-14*(3), 1–4.
492 Retrieved from <http://methods.asbcnet.org/methods/yeast-14.pdf>
- 493 Barbour, E. A., & Priest, F. G. (1988). Some effects of *Lactobacillus* contamination in Scotch Whisky
494 Fermentations. *Journal of the Institute of Brewing*, *94*(2), 89–92.
495 <https://doi.org/10.1002/j.2050-0416.1988.tb04563.x>
- 496 British Standard. *Sensory analysis - Methodology - Triangle test (ISO 4120:2004)*. , (2004).
- 497 Bryce, J. H., McCaffery, C. A., Cooper, C. S., & Brosnan, J. M. (2002). Optimising the fermentability of
498 wort in a distillery – the role of limit dextrinase. In J. H. Bryce & G. G. Stewart (Eds.), *Worldwide*
499 *Distilled Spirits Conference - Tradition and Innovation* (pp. 69–78).
- 500 Buglass, A. J. (2011). Whiskeys. In *Handbook of Alcoholic Beverages* (pp. 515–534).
- 501 Burdock, G. A. (2016). *Fenaroli's Handbook of Flavor Ingredients*.
502 <https://doi.org/10.1201/9781439847503>
- 503 Burgé, G., Saulou-Bérion, C., Moussa, M., Allais, F., Athes, V., & Spinnler, H. E. (2015). Relationships
504 between the use of Embden Meyerhof pathway (EMP) or Phosphoketolase pathway (PKP) and
505 lactate production capabilities of diverse *Lactobacillus reuteri reuteri* strains. *Journal of*
506 *Microbiology*, *53*(10), 702–710. <https://doi.org/10.1007/s12275-015-5056-x>
- 507 Buxton, I., & Hughes, P. S. (2013). Crop-to-cask: Production of New Make Spirit. In *The Science and*
508 *Commerce of Whisky* (pp. 90–133). Retrieved from
509 <http://pubs.rsc.org/en/content/ebook/9781849731508#!divbookcontent>
- 510 Cachat, E. (2005). *Lactobacillus suntoryeus* sp. nov., isolated from malt whisky distilleries.
511 *International Journal of Systematic and Evolutionary Microbiology*, *55*(1), 31–34.
512 <https://doi.org/10.1099/ij.s.0.63266-0>

513 Cheung, A. W. Y., Brosnan, J. M., Phister, T., & Smart, K. A. (2012). Impact of dried, creamed and cake
514 supply formats on the genetic variation and ethanol tolerance of three *Saccharomyces*
515 *cerevisiae* distilling strains. *Journal of the Institute of Brewing*, 118(2), 152–162.
516 <https://doi.org/10.1002/jib.23>

517 Christoph, N., & Bauer-Christoph, C. (2007). Flavour of Spirit Drinks: Raw Materials, Fermentation,
518 Distillation, and Ageing. In *Flavour and Fragrances - Chemistry, Bioprocessing and Sustainability*
519 (pp. 219–240). <https://doi.org/978-3-540-49338-9>

520 DEFRA. *Technical File for Scotch Whisky*. , (2013).

521 Dolan, T. C. S. (1976). Some aspects of the impact of brewing science on Scotch malt whisky
522 production. *Journal of the Institute of Brewing*, 82(3), 177–181. [https://doi.org/10.1002/j.2050-](https://doi.org/10.1002/j.2050-0416.1976.tb03747.x)
523 [0416.1976.tb03747.x](https://doi.org/10.1002/j.2050-0416.1976.tb03747.x)

524 Dudley, E. G., & Steele, J. L. (2005). Succinate production and citrate catabolism by Cheddar cheese
525 nonstarter lactobacilli. *Journal of Applied Microbiology*, 98(1), 14–23.
526 <https://doi.org/10.1111/j.1365-2672.2004.02440.x>

527 Ensor, M., Bryce, J. H., & Hill, A. E. (2015). An investigation into the use of different yeast strains and
528 *Lactobacillus* on new make spirit. In I. Goodall, R. N. Fotheringham, D. Murray, R. A. Speers, & G.
529 Walker (Eds.), *Worldwide Distilled Spirits Conference - Future Challenges, New Solutions* (pp.
530 243–248). Context.

531 Geddes, P. A., & Rifkin, H. L. (1989). Influence of lactic acid bacteria on aldehyde, ester and higher
532 alcohol formation during Scotch whisky fermentations. In *Distilled Beverage Flavour* (pp. 193–
533 199).

534 Gerhäuser, C. (2005). Beer constituents as potential cancer chemopreventive agents. *European*
535 *Journal of Cancer*, 41(13), 1941–1954. <https://doi.org/10.1016/j.ejca.2005.04.012>

536 Hammond, J., Brennan, M., & Price, A. (1999). The Control of Microbial Spoilage of Beer. *Journal of*
537 *the Institute of Brewing*, 105(2), 113–120. <https://doi.org/10.1002/j.2050-0416.1999.tb00014.x>

538 Ikari, A., & Kubo, R. (1975). Behaviour of various impurities in simple distillation of aqueous solution
539 of ethanol. *Journal of Chemical Engineering of Japan*, 8(4), 294–299.
540 <https://doi.org/10.1252/jcej.8.294>

541 Lee, K.-Y. M., Paterson, A., Piggott, J. R., & Richardson, G. D. (2001). Origins of Flavour in Whiskies
542 and a Revised Flavour Wheel: a Review. *Journal of Institute of Brewing*, 107(5), 287–313.
543 <https://doi.org/10.1002/j.2050-0416.2001.tb00099.x>

544 Leiper, K. A., Stewart, G. G., McKeown, I. P., Nock, T., & Thompson, M. J. (2005). Optimising beer
545 stabilisation by the selective removal of tannoids and sensitive proteins. *Journal of the Institute*
546 *of Brewing*, 111(2), 118–127. <https://doi.org/10.1002/j.2050-0416.2005.tb00657.x>

547 Li, H., & Liu, F. (2015). Changes in Organic Acids during Beer Fermentation. *Journal of the American*
548 *Society of Brewing Chemists*, 73(3), 275–279. <https://doi.org/10.1094/ASBCJ-2015-0509-01>

549 Lopez, E. F., & Gomez, E. F. (1996). Simultaneous Determination of the Major Organic Acids, Sugars,
550 Glycerol, and Ethanol by HPLC in Grape Musts and White Wines. *Journal of Chromatographic*
551 *Science*, 34(5), 254–257. <https://doi.org/10.1093/chromsci/34.5.254>

552 MacIntosh, A. J. (2015). Modelling yeast growth and metabolism for optimum performance. In
553 *Brewing Microbiology*. <https://doi.org/10.1016/B978-1-78242-331-7.00003-4>

554 MacIntosh, A. J., Adler, J., Eck, E., Speers, R. A., & Speers. (2012). Suitability of the miniature
555 fermentability method to monitor industrial fermentations. *Journal of the American Society of*
556 *Brewing Chemists*, 70(3), 205–211. <https://doi.org/10.1094/ASBCJ-2012-0724-01>

557 MacIntosh, A. J., Josey, M., & Speers, R. A. (2016). *An Examination of Substrate and Product Kinetics*
558 *During Brewing Fermentations*. 74(14), 250–257. <https://doi.org/10.1094/ASBCJ-2016-4753-01>

559 Makanjuola, D. B., & Springham, D. G. (1984). Identification of lactic acid bacteria isolated from
560 different stages of malt whisky fermentations. *Journal of the Institute of Brewing*, 90(1), 13–19.
561 <https://doi.org/10.1002/j.2050-0416.1984.tb04226.x>

562 Makanjuola, D. B., Tymon, A., & Springham, D. G. (1992). Some effects of lactic acid bacteria on
563 laboratory-scale yeast fermentations. *Enzyme and Microbial Technology*, 14(5), 350–357.
564 [https://doi.org/10.1016/0141-0229\(92\)90002-6](https://doi.org/10.1016/0141-0229(92)90002-6)

565 Martin, S. A. (2002). *The influence of wort solids on the fermentation of all-malt scotch whisky wort*.
566 Heriot-Watt University.

567 O’Sullivan, T. F., Walsh, Y., O’Mahony, A., Fitzgerald, G. F., & Sinderen, D. (1999). A Comparative
568 Study of Malthouse and Brewhouse Microflora. *Journal of the Institute of Brewing*, 105(1), 55–
569 61. <https://doi.org/10.1002/j.2050-0416.1999.tb00006.x>

570 Park, J.-M., Shin, J.-A., Lee, J. H., & Lee, K.-T. (2017). Development of a quantitative method for
571 organic acid in wine and beer using high performance liquid chromatography. *Food Science and*
572 *Biotechnology*, 26(2), 349–355. <https://doi.org/10.1007/s10068-017-0047-9>

573 Piggott, J. R., Sharp, R., & Duncan, R. E. B. (1989). Fermentation. In *The Science and Technology of*
574 *Whiskies* (pp. 89–117).

575 Priest, F. G., van Beek, S., & Cachat, E. (2002). Lactic acid bacteria and the Scotch whisky
576 fermentation. In J. H. Bryce & G. G. Stewart (Eds.), *Worldwide Distilled Spirits Conference -*
577 *Tradition and Innovation* (pp. 187–196).

578 Rose, A. H. (1977). Alcoholic Beverages. In *Economic Microbiology* (Vol. 1).
579 <https://doi.org/10.1002/jobm.19790190419>

580 Shang, Y., Zeng, Y., Zhu, P., & Zhong, Q. (2016). Acetate metabolism of *Saccharomyces cerevisiae* at
581 different temperatures during lychee wine fermentation. *Biotechnology & Biotechnological*

582 *Equipment*, 30(3), 512–520. <https://doi.org/10.1080/13102818.2016.1142831>

583 Simpson, K. L., Pettersson, B., & Priest, F. G. (2001). Characterization of *Lactobacilli* from Scotch malt
584 whisky distilleries and description of *Lactobacillus ferintoshensis* sp. nov., a new species
585 isolated from malt whisky fermentations. *Microbiology*, 147(4), 1007–1016.

586 Speers, R. A., Rogers, P., & Smith, B. (2003). Non-Linear Modelling of Industrial Brewing
587 Fermentations. *Journal of the Institute of Brewing*, 19(3), 229–235.
588 <https://doi.org/10.1002/j.2050-0416.2003.tb00163.x>

589 Sugiyama, M., Sasano, Y., & Harashima, S. (2015). Mechanism of Yeast Adaptation to Weak Organic
590 Acid Stress. In H. Takagi & H. Kitagaki (Eds.), *Stress Biology of Yeasts and Fungi* (pp. 107–122).
591 <https://doi.org/10.1007/978-4-431-55248-2>

592 Suzuki, K. (2015). Gram-positive spoilage bacteria in brewing. In *Brewing Microbiology - Managing*
593 *Microbes, Ensuring Quality and Valorising Waste* (pp. 141–169).

594 Swan, J. S., & Burtles, S. M. (1978). The development of flavour in potable spirits. *Chemical Society*
595 *Reviews*, 7(2), 201. <https://doi.org/10.1039/cs9780700201>

596 Takatani, T., & Ikemoto, H. (2002). Contribution of bacterial microflora in malt whisky quality. In J. H.
597 Bryce & G. G. Stewart (Eds.), *Distilled Spirits - Tradition and Innovation* (pp. 197–207).
598 Nottingham Press.

599 van Beek, S., & Priest, F. G. (2000). Decarboxylation of substituted cinnamic acids by lactic acid
600 bacteria isolated during malt whisky fermentation decarboxylation of substituted cinnamic
601 acids by lactic acid bacteria isolated during malt whisky fermentation. *Applied and*
602 *Environmental Microbiology*, 66(12), 1–8. [https://doi.org/10.1128/AEM.66.12.5322-](https://doi.org/10.1128/AEM.66.12.5322-5328.2000)
603 5328.2000.Updated

604 van Beek, S., & Priest, F. G. (2002). Evolution of the Lactic Acid Bacterial Community during Malt

605 Whisky Fermentation: a Polyphasic Study. *Applied and Environmental Microbiology*, 68(1), 297–
606 305. <https://doi.org/10.1128/AEM.68.1.297-305.2002>

607 van Beek, S., & Priest, F. G. (2003). Bacterial Diversity in Scotch Whisky Fermentations as Revealed
608 by Denaturing Gradient Gel Electrophoresis. *J Am Soc Brew Chem*, 61(1), 10–14.

609 Von Wright, A., & Axelsson, L. (2004). Lactic Acid Bacteria: An Introduction. In *Lactic Acid Bacteria:*
610 *Classification and Physiology* (pp. 1–66). <https://doi.org/10.1201/9780824752033.ch1>

611 Vriesekoop, F., Krahl, M., Hucker, B., & Menz, G. (2012). 125th Anniversary Review: Bacteria in
612 brewing: The good, the bad and the ugly. *Journal of the Institute of Brewing*, 118(4), 335–345.
613 <https://doi.org/10.1002/jib.49>

614 Vriesekoop, F., Rathband, A., MacKinlay, J., & Bryce, J. H. (2010). The Evolution of Dextrins During
615 the Mashing and Fermentation of All-malt Whisky Production. *Journal of the Institute of*
616 *Brewing*, 116(3), 230–238. <https://doi.org/10.1002/j.2050-0416.2010.tb00425.x>

617 Walker, G. (1999). Yeast Growth. In *Yeast Physiology and Biotechnology* (Vol. 1999, pp. 101–202).
618 [https://doi.org/10.1016/s0294-3506\(99\)80418-9](https://doi.org/10.1016/s0294-3506(99)80418-9)

619 Wilson, N. R. (2008). *The Effect of Lactic Acid Bacteria on Congener Composition and sensory*
620 *Characteristics of Scotch Malt Whisky* (Heriot-Watt University). [https://doi.org/10.1016/B978-](https://doi.org/10.1016/B978-0-408-04101-0.50004-8)
621 [0-408-04101-0.50004-8](https://doi.org/10.1016/B978-0-408-04101-0.50004-8)

622 Wilson, N. R. (2014). Contamination: bacteria and wild yeasts in a whisky fermentation. In *Whisky :*
623 *Technology, Production and Marketing* (pp. 147–154). [https://doi.org/10.1016/B978-0-12-](https://doi.org/10.1016/B978-0-12-4017435-1.00008-8)
624 [4017435-1.00008-8](https://doi.org/10.1016/B978-0-12-4017435-1.00008-8)

625 Worch, T., Lê, S., & Punter, P. (2010). How reliable are the consumers? Comparison of sensory
626 profiles from consumers and experts. *Food Quality and Preference*, 21(3), 309–318.
627 <https://doi.org/10.1016/j.foodqual.2009.06.001>

628 Yonezawa, T., & Stewart, G. G. (2004). Monitoring and controlling of whisky fermentation. In James
629 H. Bryce (Ed.), *Worldwide Distilled Spiritis Conference - Tradition and Innovation* (pp. 103–112).
630 Nottingham Press.

631 Zalán, Z., Hudáček, J., Štětina, J., Chumchalová, J., & Halász, A. (2010). Production of organic acids by
632 *Lactobacillus* strains in three different media. *European Food Research and Technology*, 230(3),
633 395–404. <https://doi.org/10.1007/s00217-009-1179-9>

634