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1 **Mashing with unmalted sorghum using a novel low**
2 **temperature enzyme system: impacts of sorghum**
3 **grain composition and microstructure**

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13 TITLE RUNNING HEAD: A low temperature mash schedule for unmalted sorghum

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23 **Abstract**

24 Brewing lager beers from unmalted sorghum traditionally requires the use of high
25 temperature mashing and exogenous enzymes to ensure adequate starch conversion.
26 Here, a novel low-temperature mashing system is compared to a more traditional
27 mash in terms of the wort quality produced (laboratory scale) from five unmalted
28 sorghums (2 brewing and 3 non-brewing varieties). The low temperature mash
29 generated worts of comparable quality to those resulting from a traditional energy
30 intensive mash protocol. Furthermore, its performance was less dependant on
31 sorghum raw material quality, such that it may facilitate the use of what were
32 previously considered non-brewing varieties. Whilst brewing sorghums were of lower
33 protein content, protein *per se* did not correlate with mashing performance. Rather, it
34 was the way in which protein was structured (particularly the strength of protein-
35 starch interactions) which most influenced brewing performance. RVA profile was
36 the easiest way of identifying this characteristic as potentially problematic.

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40 **Keywords**

41 Sorghum brewing, exogenous mash enzymes, mashing, sorghum starch

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43 **1. Introduction**

44 Brewing lager style beers using predominantly unmalted sorghum requires the use of
45 exogenous mash enzymes and specific mashing schedules tailored to the conversion of
46 sorghum starch. This is in part due to the high gelatinisation temperatures of sorghum
47 starch (Espinosa-Ramírez, Pérez-Carillo & Serna-Salvídor, 2014) which typically
48 require that mash is first heated to a high temperature (e.g. 95 °C) in order to fully
49 gelatinise the starch, followed by cooling and addition of exogenous mash enzymes to
50 assure breakdown of starch to sugars. Production of Western-style beers with
51 sorghum is currently limited to the use of light-skinned, low polyphenol sorghum
52 cultivars. Traditional sorghum beers are usually produced using brown or red skinned
53 sorghum cultivars (Lyumugabe, Gros, Nzungize, Bajyana, & Thonart, 2012). Use of
54 darker skinned, high tannin cultivars in brewing is thought to result in inhibition of
55 mash enzymes and an objectionable increase in product bitterness (Kobue-Lekalake,
56 Taylor, & de Kock, 2007; Novellie, 1981). However, some workers have suggested
57 that the use of high-tannin sorghum cultivars is responsible for only a minor increase
58 in bitterness, not detectable by all panellists (Daiber, 1975). In addition, the impact of
59 polyphenols on saccharification has also been disputed in mashing using sorghum
60 malts (Dufour, Melotte, & Srebrnik, 1992). It has been suggested that the reduced
61 saccharification of some sorghum malts is due not to polyphenols, but to starch
62 characteristics and poor diastatic potential (Dufour, Melotte, & Srebrnik, 1992).

63 As new enzyme blends become available which enable lower temperature mashing
64 conditions to be employed, it is of interest to study how this impacts on the brewing
65 performance of different sorghum cultivars. Furthermore, in some regions the
66 objective to brew with locally produced raw materials can make it of interest to use
67 varieties previously considered as sub-optimal for brewing, but which show good

68 agronomic performance. In the present study, five sorghum samples were sourced:
69 two brewing cultivars (yellow (Nigeria) and yellow (Cameroon)) and three forage
70 cultivars (red (Mexico), white (Nigeria) and white (Ghana)). Laboratory scale
71 brewing trials were conducted with each of the cultivars, comparing the performance
72 of a traditional (high temperature) sorghum mashing schedule with a novel low
73 temperature schedule utilising an exogenous enzyme blend (Figure 1). The latter was
74 developed to enable the digestion of sorghum starch without a high temperature
75 gelatinisation stand prior to saccharification. One objective of the trials was to
76 determine whether the low temperature mashing system could produce worts of
77 comparable brewing quality to those brewed using the traditional mash schedule. A
78 further objective was to study the impacts of cultivar on mashing performance and to
79 try to better understand the interactions between kernel structure and composition and
80 mashing performance.

81 **2. Materials and Methods.**

82 *2.1 Sorghum grain samples*

83 Five samples of sorghum grain were sourced by Diageo and Kerry Enzymes. As
84 cultivar identities were unavailable each sorghum variety was identified by colour and
85 country of origin. Two brewing cultivars were received: yellow (Nigeria) and yellow
86 (Cameroon). Three forage cultivars were received: red (Mexico), white (Nigeria) and
87 white (Ghana). Upon arrival, samples were stored at 4°C in plastic bins (as advised
88 by Kerry Enzymes).

89 *2.2 Sorghum grain compositional analysis*

90 *2.2.1 Moisture content*

91 Milled samples (5 g; 0.2 mm EBC fine grind) were weighed into pre-weighed foil
92 trays and placed into a convection oven at 130°C for 90 min. Samples were removed
93 into a desiccator, allowed to cool for 30 min and re-weighed to calculate moisture
94 content through weight loss. Five replicate measurements were taken.

95 *2.2.2 Starch content*

96 Starch content was determined using a Starch (GO/P) Assay Kit (Sigma STA20).
97 Whole grain samples (10 mg and 50 mg respectively) were finely milled in a coffee
98 grinder and used with the kit alongside a wheat starch standard (10 mg). To remove
99 non-starch sugars from the sample before analysis the samples were incubated at
100 85°C for 5 min then washed twice in 80 % (v/v) aqueous ethanol solution. To allow
101 the kit to act effectively upon sorghum's resistant starch, 2 mL dimethyl sulphoxide
102 (DMSO) was added to each sample prior to analysis; these were then incubated in
103 boiling water for 5 min.

104 *2.2.3 Cellulose and hemicellulose*

105 Ion chromatography was used to determine monomeric sugars in acid hydrolysed
106 samples, whilst HPLC was used to determine sugar degradation products produced.
107 Cellulose was estimated as the sum of glucose and hydroxymethylfurfural (HMF)
108 minus determined starch. Hemicellulose was estimated as the summed concentrations
109 of xylose, arabinose and furfural. Samples (60 mg) were weighed into heat resistant
110 screw-capped (with PTFE seal) glass tubes. 12 M H₂SO₄ (2 mL) was added and the
111 contents incubated at 37°C for 1 h. Water (22 mL) was added and the sample was
112 further incubated at 100°C for 2 h. The samples were filtered over glass microfibre
113 syringe filters (Whatman 25 mm 0.45 µm GD/X glass microfibre) and 1 mL was
114 transferred to HPLC vials and analysed via ion chromatography and HPLC.

115 *2.2.3.1 Ion Chromatography Analysis*

116 Sample (10µL) was injected onto a Dionex CarboPac20 column (3 mm x 150 mm)
117 coupled to a Dionex ICS 3000 with an electrochemical detector (Dionex, California,
118 USA). The samples were eluted isocratically with degassed 10 mM NaOH at a flow
119 rate of 0.5 mL/min running at around 3000 psi. Compounds were detected using an
120 electrochemical cell over a 30 min run time. The column was regenerated after each
121 sample run by flushing with 200 mM NaOH at 0.5 mL/min for 10 min.

122 *2.2.3.2 HPLC Analysis*

123 10 µL was injected onto a C18 Techsphere column (250 x 4.6 mm ID; HPLC
124 Technology, Macclesfield, UK) using a Waters 2695 liquid chromatograph (Waters,
125 Massachusetts, USA). Gradient elution was used to separate the analytes, using a
126 solvent mixture of 1 % (v/v) acetic acid (aq): methanol (80:20) ramped to 50:50 over
127 30 min period with a total flow rate of 1 mL/min at a pressure of approx. 2950 psi.
128 Compounds were detected using a Waters 996 Photodiode-Array detector using UV

129 detection at 270 nm. After 30 min the methanol was increased to 100 % over 1 min,
130 held for 2 min before returning to initial solvent conditions for the next run. External
131 standards of hydroxymethylfurfural (HMF) and furfural (0.1 g/L) were used for
132 calibration. Samples were analysed in triplicate.

133 *2.2.4 Lipid*

134 Solvent-extractable lipid was determined via an adapted Folch determination
135 (Cequier-Sanchez, Rodriguez, Ravelo, & Zarate, 2008). Sample (400 mg) was added
136 to a capped glass test tube with 12 mL dichloromethane/methanol (2:1; v/v). The
137 samples were left for 2 h at room temperature with occasional hand agitation before
138 filtering through a Whatman GD/X glass microfiber filter (0.45 µm pore size). To the
139 filtrate 2.5 mL KCl (0.88 %; v/v) was added and after vigorous agitation the samples
140 were centrifuged at 380 x g at 4°C for 5 min. The aqueous upper layer was discarded
141 and the lower phase was dried over nitrogen gas. The remaining lipid was weighed.
142 Four replicate analyses were performed.

143 *2.2.5 Protein*

144 A Thermo Flash Nitrogen Analyser (ThermoFisher Scientific, Waltham,
145 Massachusetts, USA) was used to determine protein content of the samples. Sample
146 (50 mg) was sealed in a tin capsule and combusted at approximately 1800°C.
147 Quantitation was achieved with Eager 300 software using an L-aspartic acid standard.
148 Protein was determined using the N x 6.25 conversion factor. Samples were analysed
149 in triplicate.

150 *2.2.6 Ash*

151 Ash content was determined according to the method proposed by Santos, Jimenez,
152 Bartolome, Gomez-Cordoves, & del Nozal (2003). Sorghum, wheat or spent grain
153 sample (1 g) was accurately weighed into crucibles of known mass (ashed to constant

154 mass); these were placed into a muffle furnace at 580°C for 24 h. After ashing the
155 samples were placed directly into a desiccator for 30 min. The sample was then
156 accurately weighed to 3 decimal places. Samples were analysed in triplicate.

157 2.2.7 Lignin

158 Determination of sorghum lignin was achieved via an adapted version of the acetyl
159 bromide method (Iiyama & Wallis, 1990), as it was necessary to firstly remove tannin
160 from the grain. Tannin was washed from the milled sorghum grain using a method
161 adapted from Morrison, Asiedu, Stuchbury, & Powell (1995). Milled sorghum (100
162 mg) was weighed into a polypropylene tube and mixed with 10 mL acetone:water
163 (70:30; v/v). Samples were incubated at 30°C in a water bath for 30 min. After
164 incubation, samples were centrifuged at 500 g for 5 min. Extraction was repeated
165 twice with acetone:water (70:30; v/v) before a final wash with acetone. Water was
166 used to quantitatively transfer the samples into thick-walled glass tubes. The samples
167 were dried at 50°C for 48 h. To the dried samples 4 ml acetyl bromide reagent (25 %
168 acetyl bromide in glacial acetic acid) was added. The tubes were capped and
169 incubated in a water bath for 2 h at 50°C then allowed to cool for 5 min. To prepare
170 standards 10 mg lignin (Sigma 471003) was added to 4.5 mL dioxane and 1.5 mL
171 water and incubated at 50°C for 30 min (along with a dioxane/water blank) then
172 allowed to cool for 5 min. Aliquots of the incubated standard solution (0.2, 0.3, 0.4,
173 0.5 and 0.6 mL) and the blank (0.6 mL) were added to separate glass test tubes and
174 0.5 mL of the acetyl bromide reagent was added to each. Samples, standards and the
175 blank were made up to 16 mL with glacial acetic acid and 0.5 mL of this solution was
176 transferred to a glass test-tube. To each test-tube, 2.5 mL glacial acetic acid, 1.5 mL
177 sodium hydroxide (0.3 M) and 0.5 ml hydroxylamine hydrochloride (0.5 M) were
178 added. Sample volume was adjusted to 10 mL with glacial acetic acid and then

179 transferred to quartz cuvettes for analysis at 280 nm using a spectrophotometer.

180 Lignin content was calculated using the standard curve detailed. Four analyses were
181 performed for each sample.

182 *2.2.8 Tannin*

183 Sorghum grain tannin content was determined using the Vanillin-HCl method (Price,
184 Vanscoyoc, & Butler, 1978). Milled Sample (200 mg) was weighed into a
185 polypropylene tube, to this, 10 mL of 1 % (v/v) HCl was added; samples were
186 agitated on a roller bed for 20 min. Samples were centrifuged at 3000 x g to clear the
187 supernatant of particulate matter. A 1 mL aliquot of sample was added to a glass test
188 tube, to this 5 mL working vanillin reagent was added (using a blank of 1 mL sample
189 and 5 mL 4 % (v/v) HCl). Samples were incubated at 30°C for 20 min and measured
190 immediately at 500 nm. Five replicate samples were analysed.

191 *2.3 Determination of α -amylase and β -amylase in sorghum flours*

192 *2.3.1 Enzyme extraction*

193 Enzyme extracts for both assays were produced using a Megazyme Betamyl-3 kit (K-
194 BETA3; Megazyme, Co. Wicklow, Ireland). Grain sample was milled using a DLFU
195 laboratory disc mill using the EBC fine setting. Milled grain (0.5 g) was weighed into
196 a 15 mL polypropylene tube, to this, 5 mL extraction buffer (1 M Tris/HCL, 20 mM
197 disodium EDTA solution) was added. Extractions proceeded on a Stuart SRT60
198 roller bed (Bibby Scientific) for 1 h and were then centrifuged for 10 min at 2000 x g.
199 Kit efficacy was monitored using wheat flour controls of known α -amylase and β -
200 amylase activity.

201 *2.3.2 Determination of α -amylase and β -amylase*

202 Amylase activities in sorghum flour extracts were assayed using Megazyme test kits
203 (Megazyme, Bray, Ireland) and standard methodologies. α -amylase was determined

204 using the Ceralpha kit (K-CERA) whilst β -amylase was determined using the
205 Betamyl-3 kit. Five replicate samples were analysed in each case.

206 *2.4 Measurement of sorghum starch amylose content*

207 Amylose content was determined using a Megazyme Amylose/Amylopectin kit
208 (K-AMYL). Starch was precipitated from milled sample (25 mg) using 95 % (v/v)
209 ethanol. Starch samples were dissolved in ConA solvent and filtered through
210 Fisherbrand QL100 filter papers (Fisher Scientific, Loughborough, Leicestershire,
211 UK). Upon addition of ConA solution, amylopectin was precipitated from solution
212 and removed by centrifugation (14000 x g for 10 min). Amylose supernatant and
213 total starch samples were hydrolysed to glucose with a mixture of amyloglucosidase
214 and fungal α -amylase. Liberated glucose was treated with GOPOD reagent (glucose
215 oxidase, peroxidase and 4-aminoantipyrine) and GOPOD buffer (*p*-hydroxybenzoic
216 acid).

217 Absorbance was monitored at 510 nm for amylose and total starch samples allowing
218 percentage amylose to be calculated. Amylopectin was calculated by subtraction of
219 amylose from total starch. Samples were analysed in triplicate.

220 *2.5 Estimation of grain hardness and 100 grain weight*

221 Grain hardness was indirectly determined according to the sodium nitrate method of
222 grain floatation (Hallgren & Murty, 1983). Sodium nitrate was dissolved in RO
223 water to yield a solution of SG 1.300. A sample of 100 sorghum grains were weighed
224 to give 100 grain weight, the same samples were then used for grain floatation. Grain
225 samples were placed into the sodium nitrate solution and stirred for 30 seconds;
226 floating kernels were removed from the solution and counted. Five replicate readings
227 were taken for each sorghum sample.

228 *2.6 Imaging of sorghum grain ultrastructure using Scanning Electron microscopy*

229 *(SEM)*

230 Sorghum samples were deposited onto a conductive carbon pad and then mounted on
231 a standard 12 mm SEM stub and transferred directly to the SEM. All samples were
232 imaged at an accelerating voltage of 5-10 kV. The microscope used was an FEI
233 Quanta 3D 200 (FEI, Hillsboro, Oregon, USA).

234 *2.7 Starch extraction and purification from sorghum grain samples*

235 Sorghum starch was extracted according to the method of Beta, Corke, Rooney, &
236 Taylor (2001). Whole sorghum grain was steeped in 0.25 % (w/v) sodium hydroxide
237 (200 mL) for 24 h at 5°C. Steeped grains were drained and washed with 200 mL RO
238 water, then milled in a Waring blender. Sorghum slurry was passed through a 75 µm
239 pore size sieve; materials left on the sieve were milled again until they could pass
240 through the sieve. The filtrate was collected in polypropylene tubes and allowed to
241 settle for 1 h. Tubes were centrifuged at 760 x g for 10 min. The supernatant was
242 discarded and protein (grey material) was scraped from the top of the pellet using a
243 metal spatula; samples were washed with excess water until the pellet was white.
244 Recovered starch was dried at 40°C for 24 h.

245 *2.8 Starch pasting properties: Rapid Visco Analyser measurements*

246 Pasting profiles were established for sorghum grains and extracted sorghum starches
247 with the use of an RVA super 4 (Newport Scientific, Jessup, Maryland) using
248 ThermoLine for Windows software. Milled sorghum grain (3 g) was weighed into an
249 aluminium beaker; to this either 25 mL RO water or 25 mL 10 mM silver nitrate (to
250 inhibit native amylases) was added (Batey, Hayden, Cai, Sharp, Cornish, Morell, et
251 al., 2001). Samples were stirred at 960 rpm for the first 10 seconds and 160 rpm for
252 the remainder of the test. Samples were heated with the following temperature

253 profile: hold at 50°C for 2 min, heat to 95°C at 7.15°C/min, hold at 95°C for 12 min,
254 cool to 50°C at 9°C/min, hold at 50°C for 4 min. Samples were analysed in triplicate.

255 *2.9 Differential scanning calorimetry*

256 Samples (approximately 5 mg) were weighed into aluminium pans and dispersed in
257 15 mg RO water. To ensure homogenous sample dispersion the aluminium pans were
258 mixed overnight on a roller bed. Mixed samples were analysed using a DSC823e
259 differential scanning calorimeter (Mettler-Toldeo, Greifensee, Switzerland). Samples
260 were measured between 10°C and 95°C (temperature ramp of 10°C/min).

261 *2.10 Mashing schedules*

262 Brewing liquor (reverse osmosis water; RO) supplemented with potassium
263 metabisulphite (1 g/kg) and calcium chloride dihydrate (2 g/kg) was heated to 50°C
264 using a water bath. Sorghum grain was milled to EBC fine grade (0.2 mm gap
265 setting) using a DFLU laboratory disc mill (Bühler Group, Uzwil, Switzerland). Grist
266 (100 g) was weighed into a metal mashing beaker and mixed with the atempered
267 brewing liquor (300 mL). Mash pH was adjusted to pH 5.5 by addition of 10 % (w/v)
268 aqueous lactic acid. Enzymes were added as per either the low or high temperature
269 mashing regimes (Figure 1) and the mashing beakers were added to a bench top mash
270 bath (1-cube R12, Havlickuv Brod, Czech Replublic). The 1-cube mash bath was
271 preheated to 50°C prior to mashing; upon sample addition a temperature profile was
272 selected according to the enzyme system being used (Figure 1). Mash was stirred at
273 the Hartong speed setting as the Congress setting was insufficient to stir the mash.
274 Beakers were covered with aluminium foil for the duration of the mash to minimise
275 evaporation.

276 After mashing, samples were placed immediately into a 20°C water bath and allowed
277 to cool for 20 min. Cooled samples were made up to a standard weight of 700 g with
278 RO water.

279 *2.11 Standard wort (and fermented wort) analyses*

280 *2.11.1 Wort run-off volume after 10 minutes*

281 Samples were filtered through pleated filter papers (Whatman 2555 1/2 320 mm) into
282 individual Erlenmeyer flasks. After 100 mL wort had passed through the filter the
283 funnel was moved into a clean 500 mL flask and the initial 100 mL filtrate was
284 replaced into the funnel. After 10 min the funnel was moved into a 1 L Erlenmeyer
285 flask and allowed to completely drain. The volume of wort collected during those 10
286 min of filtration was measured as an index of speed of filtration.

287 *2.11.2 Analysis of wort turbidity*

288 Wort haze was measured using a Vos Rota turbidity meter (Haffmans, Venlo,
289 Netherlands). The Vos Rota chamber was rinsed and filled with RO water. Glass
290 cuvettes (60 mm diameter) were filled with filtered wort sample, capped and placed
291 into the Vos Rota chamber. Scattered light was measured at angles of 90° and 25°
292 using a wavelength of 650 nm. The turbidity meter was calibrated up to 20 EBC
293 units; samples exceeding this value were diluted appropriately to fit within the
294 calibrated range of the device.

295 *2.11.3 Wort Colour*

296 Wort colour was determined according to Analytica-EBC method 4.7.1.
297 (<http://www.analytica-ebc.com/>).

298 *2.11.4 Specific gravity and percentage alcohol of samples*

299 Density, specific gravity (SG) and alcohol content of wort and fermented samples
300 were determined using an Anton Paar DMA 4500 and AlcoLyzer Plus (Anton Paar,

301 Graz, Austria). Sample was passed through a Minisart cellulose acetate 0.45 µm
302 syringe filter (Sartorius, Göttingen, Germany) into a 50 mL polypropylene tube.
303 Sample (30 mL) was passed through both the DMA 4500 and AlcoLyzer Plus and was
304 equilibrated to 20.00°C before measurement..

305 *2.11.5 Free amino nitrogen determination*

306 The free amino nitrogen (FAN) content of samples was determined according to
307 Analytica-EBC method 8.10 (ninhydrin method; <http://www.analytica-ebc.com/>).
308 Samples absorbance values (570 nm) were compared against a glycine standard
309 solution (2 mg/L). Samples were analysed in triplicate.

310 *2.12 Small scale fermentation of wort*

311 Small scale fermentations (100 mL) were conducted on worts produced using both the
312 high and low temperature mashing regimes.

313 *2.12.1 Yeast propagation*

314 A metal loop was used to transfer *Saccharomyces cerevisiae* strain Bry 96 ale yeast
315 (Siebel Institute, Chicago, Illinois, USA) from an agar slope into 10 mL autoclaved
316 YPD media (1 % (w/w) yeast extract, 2 % peptone, 2 % glucose in RO water). The
317 culture was incubated in a Ceromat BS-1 incubator (Sartorius) heated to 25°C and
318 shaking at 120 rpm. After 4 days the culture was transferred to a 250 mL Erlenmeyer
319 flask containing 90 mL YPD media. After a further 3 days the culture was transferred
320 to a 2 L flask containing 900 mL YPD. Finally, after 4 more days the cells were
321 harvested. Yeast slurry was centrifuged at 1,370 g in a J2-21 centrifuge (Beckman
322 Coulter Inc, Brea, California); the supernatant was discarded and the pellet
323 resuspended in RO water. A total yeast cell count was performed.

324 *2.12.2 Simulated wort boiling and wort aeration*

325 Wort was placed uncovered onto a Stuart SB162 stirring hot plate (Bibby Scientific;
326 preheated to 300°C) and allowed to heat for 55 min, samples were then capped and
327 heated for an additional 5 min before being removed from the heat. Samples were
328 immediately plunged into iced water for 30 min to cool. Cooled wort (100 mL) was
329 transferred aseptically into autoclaved 125 mL Wheaton serum bottles (containing a
330 12 x 4.5 mm stirrer bar) that were then sealed with a foam bung. Vessels were placed
331 onto magnetic stirrer plates inside a cooled incubator (LMS Ltd, Sevenoaks, United
332 Kingdom) set to 4°C and left to aerate overnight. Incubator temperature was
333 increased to 18°C two hours before pitching.

334 *2.12.3 Fermentation conditions*

335 Yeast cells were pitched into wort at a rate of 1×10^6 cells/mL/°Plato (Casey &
336 Bamforth, 2010; Fix, 1999) before vessels were sealed with butyl rubber bungs and
337 crimp caps. The butyl rubber bungs were then pierced with a Bunsen valve to allow
338 CO₂ formed during fermentation to exit the vessel whilst preventing the entrance of
339 potential contaminants. Finally, fermentation vessels were placed onto stirrer plates
340 (300 rpm) and incubated at 18°C for 236 h. Fermentation progress was monitored
341 regularly by measuring the weight of the vessel.

342 **3. Results & Discussion**

343 *3.1 Characterisation and analysis of sorghum samples*

344 Measurement of 100 grain weight for each sample (Table 1) confirmed the visual
345 observation that the two brewing sorghum cultivars (the yellow sorghums from
346 Nigeria and Cameroon) were larger in size than the agricultural cultivars. Looking at
347 the grain compositional analysis (Table 1), the brewing cultivars were notably lower
348 in protein and higher in starch than the forage sorghums, confirming their value as
349 brewing raw materials. The starch contents reported are within the broad range
350 expected for sorghum grain (55.6-75.2 % db; Jambunathan & Subramanian, 1988),
351 whilst the range of protein contents reported (8.5-10.6 % db) falls in a tight band
352 relative to the overall range for sorghum cultivars (4.4-21.1 % db) suggested by
353 Jambunathan & Subramanian (1988). The Ghanaian white sorghum had the lowest
354 starch content of the varieties tested and a surprisingly high cellulose content (22.4%
355 db, versus 3.6-15.2% db for the remaining samples).

356 Tannins are usually associated with the pigmented seed coat of the sorghum grain
357 (Dlamini, Taylor, & Rooney, 2007). Thus, it was not surprising that the highly
358 pigmented, red sorghum had the highest concentration of condensed tannins
359 (measured in catechin equivalents, Table 1). However, it is interesting to note that,
360 apart from the yellow (Nigeria) sample, all of the sorghum cultivars contained
361 significant amounts of tannin. The tannin contents reported here are within the ranges
362 typically quoted for sorghum cultivars (Earp, Akingbala, Ring, & Rooney, 1981).
363 Increased tannin content in sorghum has been linked to a number of issues during
364 brewing, mostly attributed to the ability of tannins to bind proteinaceous material.
365 Tannins have been found to negatively impact the diastatic power of sorghum malts
366 through amylase binding (Beta, Rooney, Marovatsanga, & Taylor, 2000).

367 Furthermore, tannins have been implicated in inhibition of protease activity (Elmaki,
368 Babiker, & El Tinay, 1999); this is usually associated with poor digestibility in human
369 or livestock diet, but could likewise result in reduced proteolysis during brewery
370 mashing.

371 Based on the amount of amylose (Table 1) in the sorghum starches, all of the cultivars
372 investigated here fell into the heterowaxy classification (Sang, Bean, Seib, Pedersen,
373 & Shi, 2008). Waxy sorghum starch contains very little amylose (<3.5 %) compared
374 to normal sorghum starch (>23.6 %), heterowaxy starch amylose content is
375 intermediary between these two categories. The yellow (Nigeria) sorghum was
376 highest in amylose content (21.4%) whilst the Mexican red sorghum had the lowest
377 amylose content (13.0%).

378 α -amylase activity was only detectable at low levels in the white sorghum from
379 Nigeria (Table 1). This is not surprising as α -amylase is mainly produced 24-36 h
380 after the onset of germination and is not thought to be present in the grain before this
381 (Aisien & Palmer, 1983). The activity of β -amylase was either not detectable, or
382 present at very low level (Table 1). This finding is in agreement with the current
383 literature which suggests β -amylase in sorghum grain is either not present or is
384 present with limiting quantities (Taylor, Dlamini, & Kruger, 2013). In spite of the low
385 diastatic activities identified, it was important to complete this analysis by way of
386 context for the RVA and brewing experiments.

387 *3.2 SEM imaging of sorghum grain samples*

388 Scanning electron microscopy (SEM) allowed for high resolution imaging of the
389 interior of each grain sample (e.g. Figures 2A & B). Cursory investigation of the
390 samples by SEM showed the grains to be relatively similar (excluding overall size and
391 shape), with all samples displaying the characteristic sorghum grain features of an

392 embryo, an endosperm and a pericarp-testa (the outer-coat of the grain). However,
393 use of higher magnification SEM enabled a closer look at the detailed structures of the
394 different cultivars. The endosperm tissue of the grains all displayed areas of tightly
395 packed and loosely packed starch granules, defined as corneous and floury endosperm
396 tissue respectively (Hoseney, Davis, & Harbers, 1974). However, within these
397 structures there was noticeable variation between the grains. The two brewing
398 cultivars possessed a clear delineation between the corneous and floury endosperm
399 (e.g. Figure 2C), this was not evident in the other varieties. The border between
400 corneous and floury endosperm was not clear in the red variety, with tightly packed
401 granules transitioning gradually to a looser structure toward the centre of the
402 caryopsis. In addition, the floury region of the red cultivar was not as loosely packed
403 as the brewing varieties. A feature unique to the white variety from Nigeria was the
404 presence of extensive regions of loosely packed starch granules at the periphery of the
405 endosperm. This was interesting as floury (loosely packed) endosperm tissue is
406 usually associated with the centre of the sorghum caryopsis (Rooney & Miller, 1981).
407 The central region of this cultivar possessed very little observable floury endosperm
408 tissue. The other white cultivar, from Ghana possessed little observable floury
409 endosperm with corneous endosperm extending throughout the grain (Figure 2B).
410 Spherical structures were observed between the starch granules of sorghum samples
411 (e.g. Figures 2D & E). Confocal laser scanning microscopy and fluorescent staining
412 with Rhodamine B was used to confirm the identity of these structures as protein (data
413 not shown). These are probably prolamins, the storage protein that accounts for 60-70
414 % of sorghum protein (Duodu, Taylor, Belton, & Hamaker, 2003).
415 In agreement with prior literature (Seckinger & Wolf, 1973), protein bodies were
416 abundant towards the endosperm periphery, becoming less so in the corneous

417 endosperm and floury endosperm. In the corneous endosperm, spherical protein
418 bodies were concentrated between starch granules (e.g. Figure 2E). Starch granules in
419 corneous endosperm were less spherical and irregularly shaped (Figure 2E).
420 Polygonal starch granules are thought to be formed by constriction by storage proteins
421 caused by water loss during maturation of the caryopsis (Hoseney, Davis, & Harbers,
422 1974). As the starch granules become packed together, protein bodies are compacted
423 and concentrated between starch granules.

424 Imaging of crudely purified sorghum starch further illustrated the close interaction
425 between protein matrix and starch granule (Figure 2F). Many starch granules had
426 clear indentations, with some containing protein that survived purification. The white
427 sorghum from Ghana displayed the greatest degree of protein surviving crude starch
428 isolation, indicating a particularly strong protein-starch interaction in this cultivar.
429 Such interactions have the potential to hinder starch swelling and hydration during
430 brewery mashing (Almeida-Dominguez, Suhendro, & Rooney, 1997).

431 *3.3 Thermophysical properties of sorghum flours and extracted/purified starches*

432 Pasting profiles of sorghum flours in water revealed key differences between the
433 sorghum varieties investigated (Figure 3A). The pasting profile of the yellow cultivar
434 from Nigeria closely resembled that of a barley control (not shown) and displayed the
435 highest peak viscosity and final viscosity. Both yellow (Cameroon) and white
436 (Nigeria) displayed low peak and final viscosities (Table 2), this was hypothesised to
437 be due to enzyme activity within the sorghum flours, although only the White
438 (Nigeria) sorghum contained detectable α -amylase activity (Table 1). Use of silver
439 nitrate (10 mM) to inhibit enzymes during Rapid Visco Analyser (RVA) testing
440 revealed a pasting profile markedly different to that obtained with water (Figure 3B &
441 Table 2). During enzyme-inhibited RVA all sorghum flours displayed an increase in

442 viscosity as compared to RVA using water. This suggested the presence of enzyme
443 activity within the sorghum flours. For silver nitrate RVA, white (Nigeria) and yellow
444 (Cameroon) displayed pasting profiles similar to the other sorghums with the
445 exception of the white sorghum cultivar from Ghana, which displayed a unique
446 pasting profile (Figures 3A and 3B), with neither a clear viscosity peak nor viscosity
447 trough being observed. The characteristic lack of a viscosity peak was observed with
448 or without silver nitrate addition, suggesting that enzyme activity was not the cause of
449 this feature. Lack of a clear viscosity peak in maize has been linked to poor starch
450 granule hydration and swelling as a result of protein-starch interactions (Almeida-
451 Dominguez, Suhendro, & Rooney, 1997). The hypothesis that protein starch
452 interactions inhibited starch granule swelling in the white sorghum from Ghana is
453 supported by the SEM imaging results (Figure 2F, Section 3.2). The impact of
454 protein-starch interaction on starch granule swelling is thought to be exacerbated in
455 material originating from the corneous endosperm due to the tightly packed condition
456 of the starch (Almeida-Dominguez, Suhendro, & Rooney, 1997). In agreement with
457 findings from SEM imaging, a simple floaters test for grain hardness (Table 1)
458 suggested the white sorghum from Ghana contained the highest proportion of
459 corneous endosperm as compared to the other sorghum samples (since increased
460 endosperm density, reflecting a higher proportion of corneous material, will cause the
461 grains to sink rather than float)Furthermore, RVA analysis of starch isolated from the
462 white sorghum (Ghana) revealed a pasting profile similar to the other sorghums
463 analysed (Figure 3C and Table 2). This suggests that poor swelling was not an
464 indigenous characteristic of the starch in that cultivar and was instead mediated by a
465 component removed during purification.

466 One of the primary issues associated with sorghum brewing is a high starch
467 gelatinisation temperature. Use of differential scanning calorimetry revealed that all
468 of the sorghum cultivars studied here had a gelatinisation temperature (Table 2) in
469 excess of that expected for barley malt (62-63°C; Palmer, Etokakpan, & Igyor, 1989).
470 The red sorghum sourced from Mexico had the lowest gelatinisation peak temperature
471 (68.9°C) whilst the other sorghums gelatinised at higher temperatures (peak
472 temperature 72.9-74.5°C) Interestingly, an association was observed between starch
473 amylose content and peak gelatinisation temperature (Tables 1 and 2). This is in
474 agreement with the findings for rice and maize previously determined by other
475 researchers (Knutson, 1990; Varavinit, Shobsngob, Varanyanond, Chinachoti, &
476 Naivikul, 2003). The complex nature of starch gelatinisation is highlighted by
477 comparison of DSC analysis of sorghum flour and sorghum starch in Table 2.
478 Gelatinisation of isolated sorghum starches was achieved at a lower value than their
479 counterpart sorghum flours. Swelling of starch granules is required for efficient
480 gelatinisation, this process has been found to be restricted by interactions of starch
481 with lipids and proteins (Debet & Gidley, 2006). The lower gelatinisation
482 temperatures observed in isolated sorghum starches can probably be accounted for by
483 the removal of lipids and proteins that could inhibit granule swelling.

484 *3.4 Laboratory mashing of unmalted sorghum samples*

485 Each of the five sorghum samples were mashed using both the high and low
486 temperature mash schedules depicted in Figure 1. Analytical data for the resulting
487 wort samples is presented in Table 3, alongside post-fermentation data indicating
488 ethanol yield and fermentability when each wort was fermented at laboratory scale.
489 Together these data enable the brewing value of the worts to be appraised, with

490 reference both to the efficacy of the novel low temperature mashing schedule and also
491 to the impacts of sorghum grain composition and structure on the mashing process.

492 Hot water extract (HWE) is a key indicator of brewing efficiency. It represents the
493 proportion of grist material solubilised during mashing and is calculated based on the
494 extract content of wort (expressed in °Plato) and the amount of dry matter in the grist.

495 The yellow (Nigerian) brewing sorghum had the highest HWE (82.6%; Table 3) using
496 the high temperature (conventional) mash schedule. Surprisingly the other brewing
497 cultivar from Cameroon had a lower HWE (78.6%) than two of the forage cultivars
498 using this mash schedule. Most interestingly, the low temperature mashing schedule
499 evened out the differences between cultivars, yielding HWE values ranging between
500 81.1-82.7% for all samples bar the Ghanaian white sorghum (72.9%). This probably
501 reflects the activity of the Promalt S-LTP enzyme blend which was apparently able to
502 convert starch to sugars at low temperature consistently and irrespective of grain
503 protein content. The white sorghum from Ghana performed worst in terms of HWE
504 with either mashing schedule and has previously (Section 3.2) been noted to exhibit a
505 high proportion of corneous endosperm and strong starch-protein interactions. This
506 presumably caused problems with starch swelling and conversion using either
507 brewing schedule. Increased corneous endosperm has been associated with reduced
508 saccharification during mashing as a result of strong starch-protein interactions
509 causing inferior amylase access (Espinosa-Ramirez, Perez-Carrillo, & Serna-Saldivar,
510 2014). This hypothesis is corroborated by the RVA results (Table 2). Furthermore the
511 Ghanaian sorghum had the lowest starch content of all of the samples (49.3% db;
512 Table 1).

513 Whilst extract is an important economic consideration, the brewer also needs to
514 understand the value of that extract for alcohol production through fermentation. This

515 is appraised here in terms of the individual and total amounts of fermentable sugars
516 generated in wort. Whilst some of the forage sorghums performed reasonably well in
517 terms of extract potential, the known brewing cultivars resulted in significantly higher
518 total fermentable sugars using either mashing schedule (Table 3). Interestingly, the
519 yellow Nigerian brewing cultivar gave the highest fermentable sugars yield using the
520 high temperature mash schedule, but was exceeded in this regard by the other
521 (Cameroonian) brewing variety when mashed using the low temperature regime.
522 Furthermore, all cultivars yielded higher amounts of fermentable sugars using the low
523 temperature mash schedule relative to equivalent data for the high temperature mash.
524 The profile of fermentable sugars in wort is principally determined by the enzymes
525 present and their interaction with the mash time-temperature schedule. Thus, radically
526 different profiles were obtained when comparing the two mash schedules, but
527 comparing within each schedule, there was minimal impact of cultivar on fermentable
528 sugar spectrum (Table 3). The main feature of this data set is thus the very high
529 glucose concentrations (36.8-45.5 g/L) in low temperature mashed worts, due to the
530 inclusion of an amyloglucosidase enzyme in the formulation (Amylo 300). In
531 comparison, for the high temperature mashed worts, glucose concentrations ranged
532 from 9-12.5 g/L and maltose was the major wort fermentable sugar (30.9-47.3 g/L).
533 It has been suggested that tannins can be involved in amylase binding and inactivation
534 (Okolo & Ezeogu, 1996). Review of the present data set fails to support this
535 hypothesis, with analysed tannin levels (Table 1) showing no obvious association with
536 fermentable sugars yield (Table 3). We conclude that other factors were more
537 significant in determining the yield of sugars and that tannins were not limiting on
538 amylase activity at the concentrations noted (35-74 mg/g db catechin equivalents) and
539 with the concentrations of exogenous enzymes used.

540 Mashing with the white variety from Nigeria produced wort comparable to the
541 brewing cultivars in both high- and low-temperature mashing systems in terms of
542 extract. Despite this, worts of the white sorghum from Nigeria were lower in glucose,
543 maltose and maltotriose content. This probably resulted from incomplete hydrolysis
544 of soluble, yet unfermentable dextrins in the wort.

545 Based on the current results, the high-temperature system performed optimally with
546 the yellow cultivar from Nigeria but with reduced efficiency when acting upon the
547 other varieties. The low-temperature enzyme system is assumed to act on
548 ungelatinised starch, without the need for efficient starch dissolution, and it is likely
549 that starch characteristics had a lesser impact on mashing efficiency in this case.

550 The Free amino nitrogen (FAN) content of worts produced (44-94 mg/L; Table 3)
551 were comparable to published data for worts produced from 100 % unmalted sorghum
552 grain (e.g. 51 mg/L; (Bajomo & Young, 1993)). For all cultivars the low temperature
553 mash schedule gave marginally higher FAN contents relative to those from the high
554 temperature mashes. However, all of these worts would likely require
555 supplementation with additional nitrogen sources prior to fermentation as they would
556 not provide the minimum of 100-230 mg/L FAN (dependent on wort gravity) thought
557 to be required for efficient yeast cell fermentation (Pierce, 1987). Worts produced
558 from the Mexican red sorghum and the white variety from Nigeria gave higher FAN
559 worts than did the brewing cultivars. However, they would still be considered FAN
560 deficient relative to a barley malt wort (e.g. 158 mg/L; Bajomo & Young, 1993).
561 Worts produced using the Ghanaian sorghum had significantly lower FAN contents as
562 compared to other worts when using either enzyme system. Since this variety had a
563 similar protein content to the other agricultural varieties (Table 1) a reduced wort

564 FAN content implies issues with proteolysis during mashing, which might again
565 reflect the impacts of strong starch granule-protein interactions.

566 Another characteristic of note during mashing was turbidity in worts of the
567 Cameroonian and Mexican cultivars. During high-temperature mashing of both
568 cultivars high turbidity wort was produced (Table 3); this was not observed with use
569 of the low-temperature mashing system. Wort haze can be attributed to a number of
570 causative factors, including lipid content, polyphenol-protein interactions and the
571 survival of β -glucan in the wort (Steiner, Becker, & Gastl, 2010). Interestingly, these
572 two varieties were both of characteristically low amylose content (Table 1); perhaps
573 poor amylopectin hydrolysis could have contributed to haze formation. Wort samples
574 in this research were only run through a filter paper, it is possible that turbidity may
575 not be an issue in at industrial scale using a mash filter.

576 *3.5 Laboratory scale fermentation trials*

577 Worts produced from five different sorghum cultivars were fermented at small scale
578 (100 mL). The fermentations of the low-temperature mashed worts displayed higher
579 final alcohol contents (% ABV) as compared to those of the high-temperature system
580 (Table 3) although they took significantly longer to reach attenuation (final gravity).
581 In addition, fermentations of low-temperature mashing were lower in residual extract
582 and FAN content, suggesting a proportionately greater utilisation of wort components.
583 Despite the fact that worts produced using the low-temperature system contained
584 higher amounts of fermentable sugars and FAN as compared to high-temperature
585 mashed worts, fermentation profiles showed that they fermented relatively slowly by
586 comparison (data not shown). Fermentation of worts produced from the high-
587 temperature system were mostly complete within 120 h. For low temperature mashed
588 worts fermentation was not fully attenuated even after 236 h. This was most likely

589 due to the sugar profiles of the worts. Worts produced by low-temperature mashing
590 were rich in glucose (due to the amyloglucosidase enzyme addition), which has
591 previously been linked to inhibited glucose uptake, yeast growth and slow
592 fermentation (MacGregor, Bazin, Macri, & Babb, 1999; Phaweni, O'Connor-Cox,
593 Pickerell, & Axcell, 1993). The results illustrate that simply providing a greater
594 content of fermentable sugar and FAN does not guarantee an efficient fermentation.

595 The worts of the Mexican sorghum and agricultural white sorghum (Nigeria) from
596 low-temperature mashing were of comparable fermentability and final alcohol yield to
597 those produced using brewing cultivars. This was despite them having a lower starch
598 content in the original grist (Table 1).

599 The results obtained here suggest that worts produced using the low-temperature
600 mashing system can result in fermentation alcohol yields comparable to the high-
601 temperature mashing system. In addition, the low-temperature system appeared less
602 dependant on the raw materials used. However, fermentation of the low-temperature
603 mashed worts was relatively slow, indicating a deficiency in a component required for
604 efficient fermentation or the presence of a component at inhibitory concentrations.

605 **4. Conclusions**

606 A novel low-temperature mashing system was shown to produce worts of comparable
607 brewing value to those resulting from a more traditional, energy intensive, high-
608 temperature mash. The energy savings of operating with the low temperature system
609 would be substantial at industrial scale because i) T_{\max} for the schedule was reduced
610 from 95°C to 78°C, ii) the energy requirements of heating a mash to 95°C and then
611 cooling it back to 65°C to saccharify the mash are removed and iii) the overall mash
612 schedule is shorter by approximately 2 hours. Furthermore, our results offer
613 preliminary encouragement that the novel low-temperature mashing regime
614 compensates for some raw material quality differences and narrowed the gap in
615 brewing performance between the use of brewing and non-brewing sorghum cultivars.
616 It thus has the potential to facilitate broader use of locally produced sorghum varieties
617 in brewing, although full substantiation of this is beyond the scope of the present
618 paper. The noted issue with long, sluggish fermentation times for the low temperature
619 mashed worts is readily solvable in brewing practice. The excellent apparent
620 fermentability results confirm that the worts had the required alcohol yield potential,
621 albeit that the fermentations took a long time to attenuate. Fermentation vigour would
622 most likely be improved by i) substituting different diastatic enzyme blends for the
623 Amylo300 (amyloglucosidase) used here. This enzyme is not the component which
624 confers the low temperature gelatinisation property and it generates high
625 concentrations of glucose in worts which subsequently can slow yeast glucose uptake
626 (Phaweni, O'Connor-Cox, Pickerell, & Axcell, 1993), or ii) the use of supplementary
627 yeast nutrients (nitrogen source, Zn^{2+} , etc.).
628 With regard to the impacts of cultivar composition, starch properties and
629 ultrastructure on brewing performance it was interesting to note that with either

630 mashing schedule the impacts of kernel structure, and in particular evidence of strong
631 starch-protein interactions had a far greater influence than did starch gelatinisation
632 temperature – although the latter is more frequently used to assess likely brewing
633 performance. Thus the noted lower gelatinisation temperature range for the red
634 sorghum from Mexico did not offer a significant advantage in terms of extract or
635 fermentable sugars yield. Whilst the brewing varieties were of lower protein content,
636 protein *per se* did not correlate with mashing performance. Thus, the red sorghum
637 contained the highest amount of protein (and tannins) but yielded respectable brewing
638 performance, particularly when mashed using the low temperature regime. Hence our
639 work suggests that it is the way in which protein is structured and in particular the
640 strength of protein-starch granule interactions which most influenced brewing
641 performance. Thus the white (Ghana) sorghum performed poorly using either mash
642 schedule. The RVA profile represented the easiest way of identifying this sorghum as
643 potentially problematic for brewing use.

644 In the present work there was no support for the hypothesis that tannin levels
645 negatively impact on brewing performance (with the levels of exogenous enzymes
646 used here), although this was not the main focus of the study and no sensory tests
647 were performed on beers to evaluate the levels of astringency conferred.

648

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655 **Conflict of Interest**

656 The authors are not aware of any conflict of interest relating to publication of the

657 enclosed material.

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Table 1: Analytical data for the five sorghum cultivars used in the trial.

	Yellow (Nigeria)	Yellow (Cameroon)	Red (Mexico)	White (Nigeria)	White (Ghana)
Ash (g/100g db)	1.8 ±0.19	1.8 ±0.16	2.3 ±0.18	2.2 ±0.35	2.5 ±0.18
Cellulose (g/100g db)	11.7 ±3.1	3.6 ±2.39	13.1 ±3.28	15.2 ±4.22	22.4 ±0.95
Hemicellulose (g/100g db)	7.6 ±0.60	6.6 ±2.11	6.9 ±2.67	5.6 ±1.79	4.1 ±0.31
Lignin (g/100g db)	5.3 ±1.09	7.4 ±1.66	6.7 ±1.8	5.0 ±0.95	4.9 ±1.03
Lipid (g/100g db)	3.1 ±0.2	3.2 ±0.16	2.7 ±0.15	3.2 ±0.23	3.9 ±0.53
Protein (g/100g db)	9.4 ±0.06	8.5 ±0.59	10.6 ±0.2	9.8 ±0.36	10.2 ±0.30
Starch (g/100g db)	61.7 ±6.04	64.4 ±2.33	58.1 ±2.39	55.8 ±1.75	49.3 ±0.62
TOTAL	100.49	95.53	100.32	96.7	97.19
Moisture content (%)	11.2 ±0.16	10.4 ±0.04	14.7 ±0.13	11.5 ±0.05	11.4 ±0.21
Amylose (%)	21.4 ±2.60	14.1 ±0.00	13.0 ±1.20	18.4 ±3.10	18.9 ±1.80
Floater (%)	94.0 ±1.00	99.7 ±0.60	90.3 ±3.20	95.3 ±0.60	32.0 ±1.00
100 grain weight	3.94 ±0.31	5.24 ±0.43	2.26 ±0.07	3.22 ±0.28	2.55 ±0.28
α-amylase (DU/g)	nd	nd	nd	1.66±0.24	nd
β-amylase (betamyl-3-units)	nd	0.19±0.096	0.21±0.058	0.23±0.081	nd
Catechin equivalents mg/ g (dry weight basis)					
Tannins (Vanillin-HCl method)	nd	48 ±27	74 ±6	43 ±7	35 ±6

Results are the mean of at least triplicate independent analyses ± standard deviation

Table 2: Thermophysical properties of sorghum flours and starches according to Differential Scanning Calorimetry (DSC) and Rapid Visco Analysis (RVA) in the presence of 10 mM silver nitrate.

	Yellow (Nigeria)	Yellow (Cameroon)	Red (Mexico)	White (Nigeria)	White (Ghana)
Sorghum flour samples (DSC)					
Gelatinisation Onset (°C)	72.49	71.07	64.54	72.54	71.87
Gelatinisation Peak (°C)	77.34	76.48	71.47	76.83	76.87
Gelatinisation Endset (°C)	84.73	83.97	78.76	83.65	85.60
Sorghum flour samples (RVA)					
Time of gelatinization (s)	371	347	314	355	360
Peak Viscosity (cP)	733	1010	782	765	-
Trough Viscosity (cP)	661	786	605	723	-
Breakdown (cP)	72	224	177	41	-
Final Viscosity (cP)	1388	1617	1305	1469	1419
Total Setback (cP)	726	831	700	746	-
Sorghum starches (DSC)					
Gelatinisation Onset (°C)	70.32	68.43	63.36	69.83	69.95
Gelatinisation Peak (°C)	74.47	72.90	68.92	73.50	74.16
Gelatinisation Endset (°C)	80.28	79.14	75.58	79.35	80.79
Sorghum starches (RVA)					
Time of gelatinization (s)	325	315	274	319	322
Peak Viscosity (cP)	3928	4159	4893	3986	4068
Trough Viscosity (cP)	993	1044	1027	1229	1206
Breakdown (cP)	2935	3115	3866	2757	2862
Final Viscosity (cP)	2605	2718	2451	3137	3317
Total Setback (cP)	1612	1674	1424	1908	2111

Results are the mean of triplicate analyses.

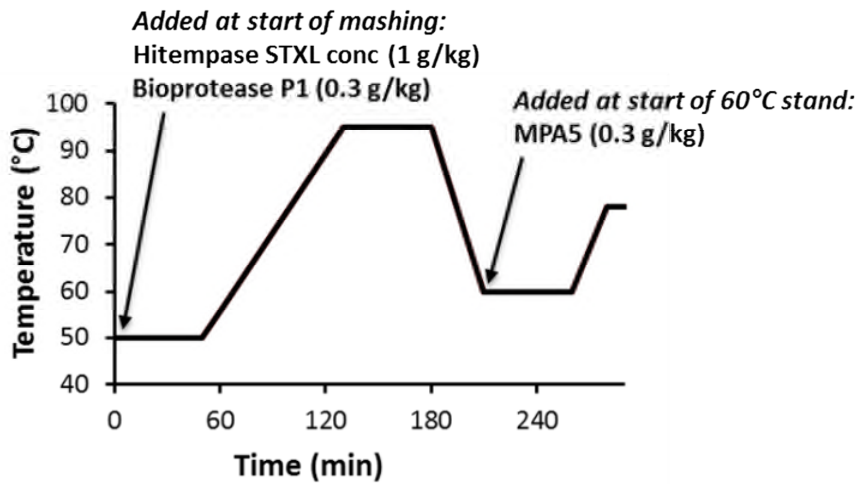
1 **Table 3:** Results to mashing and fermentation trials using five sorghum cultivars mashed using either the high temperature or low temperature
2 mashing schedule. Standard conditions: mashing-in pH 5.5, KMS 1 g/kg, CaCl₂·2H₂O 2 g/kg, enzymes.
3

	HT mashing system					LT mashing system				
Wort analyses	Yellow Nigeria	Yellow Cameroon	Red Mexico	White Nigeria	White Ghana	Yellow Nigeria	Yellow Cameroon	Red Mexico	White Nigeria	White Ghana
Extract (°P)	10.71 ±0.1	10.34 ±0.06	9.86 ±0.08	10.42 ±0.34	10.12 ±0.05	10.62 ±0.06	10.71 ±0.06	10.12 ±0.08	10.69 ±0.12	9.55 ±0.25
Hot Water Extract (% db)	82.6	78.6	78.8	80.4	77.7	81.8	81.7	81.1	82.7	72.9
Wort colour (EBC)	5 ±0	6 ±1	7 ±1	3 ±0	2 ±0	5 ±0	6 ±0	5 ±0	3 ±0	2 ±0
Wort pH	5.71 ±0.01	5.74 ±0.01	5.67 ±0.02	5.80 ±0.01	5.63 ±0.04	5.70 ±0.01	5.80 ±0.01	5.71 ±0.02	5.88 ±0.03	5.72 ±0.04
FAN (mg/L)	61 ±5	70 ±1	78 ±1	86 ±5	44 ±2	63 ±1	73 ±0	82 ±4	94 ±2	49 ±1
fructose (g/L)	0.7 ±0.1	1.3 ±0	0.3 ±0	0.7 ±0.1	0.5 ±0	0.6 ±0	1.1 ±0.1	nd ^a	0.7 ±0	0.3 ±0
glucose (g/L)	9.9 ±0.4	12.5 ±1.5	11.1 ±0.3	11 ±0.7	9 ±0.3	45.3 ±3.2	45.5 ±3.9	44 ±1.3	41.4 ±1.5	36.8 ±1.6
maltose (g/L)	47.3 ±2.8	41.1 ±3.4	37.5 ±1.4	32.9 ±2.1	30.9 ±1.9	26.7 ±3	27.6 ±1.8	24.9 ±0.9	26.6 ±0.9	22.3 ±2.4
maltotriose (g/L)	29.2 ±1	26.7 ±2.9	23.9 ±0.9	21.8 ±1.7	23.9 ±1.8	15.3 ±2.1	15.4 ±1.4	11.3 ±0.2	13 ±0.7	12.1 ±0.3
^a Fermentable sugars (TOTAL)	87.1	81.6	72.8	66.4	64.3	87.9	89.6	80.3	81.7	71.5
haze 25° (EBC)	1.86 ±0.01	8.78 ±0.19	7.79 ±2.13	1.9 ±0.47	0.13 ±0.04	1.7 ±0.2	1.08 ±0.06	0.63 ±0.24	2.18 ±0.07	0.22 ±0.07
haze 90° (EBC)	1.12 ±0.03	3.71 ±0.13	2.98 ±0.72	1.19 ±0.22	0.25 ±0.05	1.41 ±0.1	2.04 ±0.22	0.37 ±0.09	1.09 ±0.02	0.37 ±0.11
run-off volume (mL)	327 ±28	425 ±22	319 ±19	245 ±9	300 ±6	448 ±8	443 ±23	441 ±5	366 ±15	383 ±14
Post-fermentation analyses										
Alcohol content (% ABV)	4.68 ±0.16	4.37 ±0.59	4.40 ±0.09	4.17 ±0.41	4.11 ±0.15	4.69 ±0.25	5.13 ±0.61	4.71 ±0.25	5.11 ±0.17	4.36 ±0.12
Residual Extract (° Plato)	2.69 ±0.06	2.09 ±0.23	2.04 ±0.03	2.71 ±0.28	2.99 ±0.1	1.39 ±0.11	1.55 ±0.06	0.83 ±0.2	1.28 ±0.04	1.37 ±0.07
FAN (mg/L)	27 ±0	24 ±1	31 ±2	29 ±1	16 ±4	17 ±3	21 ±1	19 ±1	19 ±8	15 ±1
Apparent fermentability (%)	76.8	81.2	81.0	75.4	72.4	87.2	86.1	92.0	88.7	86.5

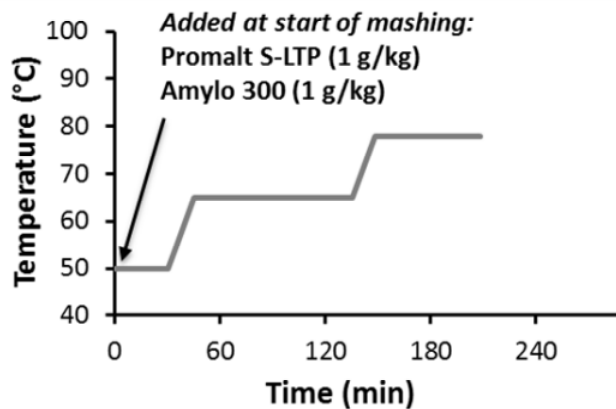
4 Results are the mean of triplicate independent mashes ± standard deviation. ^asum total of fructose, glucose, maltose and maltotriose.

Figure 1: Details of A) traditional high temperature and B) novel low temperature mashing regimes used in the research, together with details of the respective exogenous enzymes added.

A)



B)



Enzyme preparation	Principal Activities	Enzyme source	Temperature optimum	pH optimum
Amylo 300	amyloglucosidase	<i>A. niger</i>	75	4.0
Bioprotease P1	protease	<i>Bacillus spp.</i>	70	6.0
Hitempase STXL	α -amylase	<i>B. lichenformis</i>	90	6.0
MPA 5	α -amylase	<i>A. oryzae</i>	60	6.0
Promalt S-LTP	Amylolytic and proteolytic	GM and non-GM strains	50-70	5.0-7.0

Figure 2 Scanning electron micrographs showing: Longitudinal cross section through an entire caryopsis of A) yellow sorghum from Nigeria and B) white sorghum from Ghana. C) the border between floursy and corneous endosperm in the yellow (Nigeria) sample D) High magnification image of the floursy endosperm of yellow Nigerian sorghum E) corneous endosperm of the white Ghanaian sorghum and F) a starch granule isolated from the white sorghum originating in Ghana, labelled with (i) protein body and (ii) indentation.

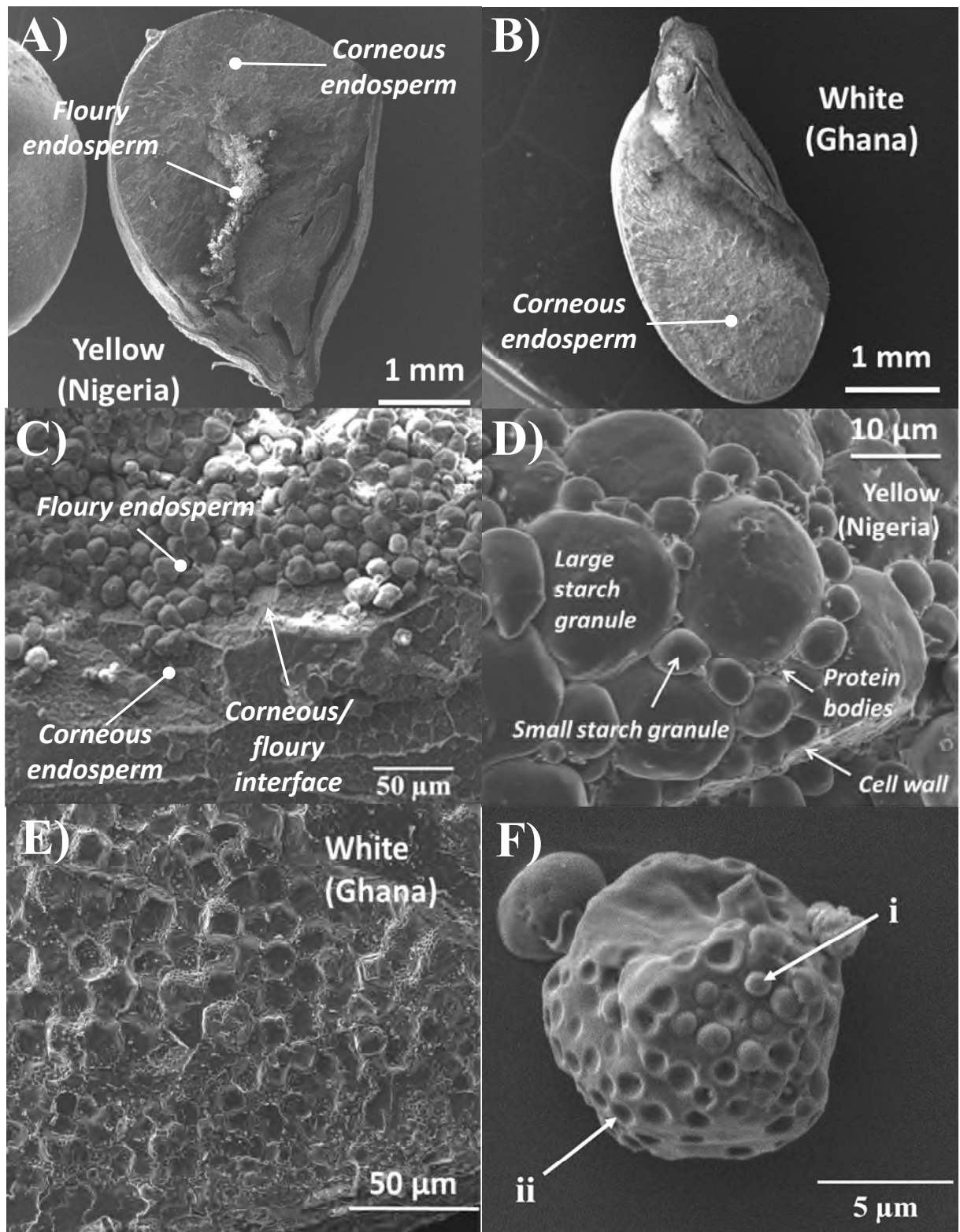
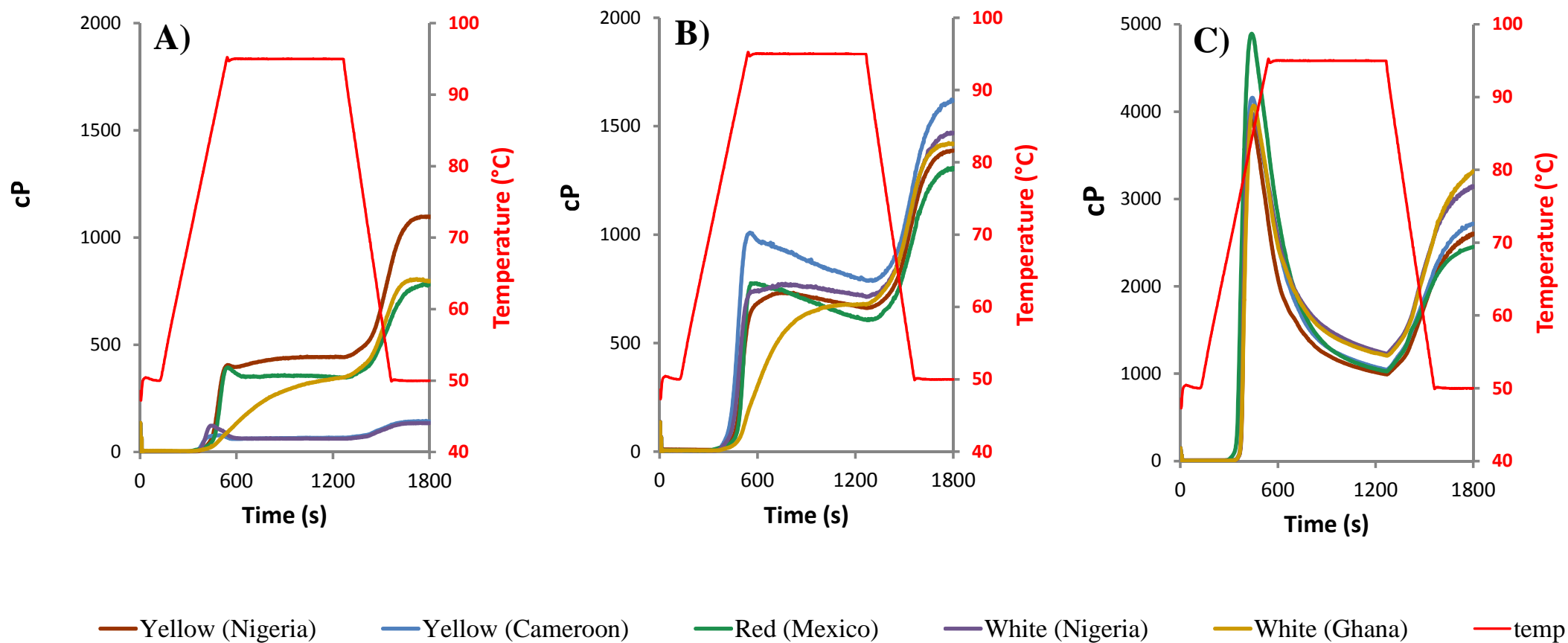


Figure 3: RVA pasting profiles of (A) sorghum flours tested in water (B) sorghum flours tested in 10 mM silver nitrate and (C) extracted and purified sorghum starches in 10 mM silver nitrate.



Results displayed are the mean of triplicate analyses.