



M. Jalil, A. M., Edwards, C. A., Combet Aspray, E., Ibrahim, M., and Garcia, A. L. (2015) Combined effects of added beta glucan and black tea in breads on starch functionality. *International Journal of Food Sciences and Nutrition*, 66(2). 159-165.

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Combined effects of added beta glucan and black tea in breads on starch functionality

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This manuscript is published in *International Journal of Food Sciences and Nutrition* (2015) (doi:10.3109/09637486.2014.971225)

1 **ABSTRACT**

2 Bread and tea are usually consumed separately but there may be different food matrix
3 interactions and changes in starch characteristics when they are combined in bread. This
4 study developed breads (white bread, WF; black tea, BT; beta glucan, β G; beta glucan plus
5 black tea, β GBT) and determined their starch functionalities. Breads were developed using a
6 standard baking recipe and determined their starch characteristics. There was no significant
7 difference in starch hydrolysis between BT and WF but β GBT reduced early (10 min) starch
8 hydrolysis compared with β G. The starch granules in β G and β GBT were elliptical and
9 closely packed together. These results suggest that the addition of beta glucan and black tea
10 to bread preserved the elliptical starch granules and lowered short-term starch hydrolysis.

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14 Keywords: beta glucan bread, starch granules, starch hydrolysis, food matrix

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21 **Introduction**

22 Tea is one of the most common beverages consumed around the world. Long-term
23 intake of tea has been associated with risk reduction for cardiovascular diseases, cancers,
24 hypertension and diabetes mellitus (Gardner et al., 2007; Khan & Mukhtar, 2012). The health
25 effects of tea have been attributed to the presence of bioactive polyphenols (Rothwell et al.,
26 2012). Acute intake of tea also reduced postprandial glycaemia in humans (Bryans et al.,
27 2007).

28 Beta glucan is a soluble dietary fibre with mixed β -(1 \rightarrow 3) and β -(1 \rightarrow 4) linkages. The
29 effects of beta glucans on post-prandial glycaemic and insulinaemic responses in humans
30 have been well established (Ostman et al., 2006). The European Food Safety Authority
31 (ESFA) approved a health claim in which 4 g of beta glucan from either oats or barley per 30
32 g available carbohydrate is recommended to reduce glycaemic response without
33 disproportionately increasing postprandial insulinaemic response (Agostoni et al., 2011).
34 Hence, beta glucan can be used as an active ingredient in formulating products aiming at
35 reducing postprandial blood glucose.

36 Bread is the most popular starchy food in Europe with an average intake of 50 kg bread
37 per person per year (Bakers, 2014). Bread contains gluten and starch whose properties are
38 directly influenced by different stages of bread making (mixing ingredients, proofing and
39 baking) (Rosell, 2011). Breads are a good target for further development to improve their
40 functional properties (Hayta & Gamze, 2011). In addition, they have been used as a model to
41 study food matrix interactions (Juntunen et al., 2002; Juntunen et al., 2003). Food
42 constituents can interact in several ways when eaten separately but if cooked together in a
43 product such as bread, these interactions may be more complex and influence other
44 components during food processing. The addition of polyphenols and/or dietary fibres to

45 foods may directly or indirectly modify the properties of the starch components, their
46 digestibility, and antioxidant properties (Juntunen et al., 2002; Juntunen et al., 2003; Rosen et
47 al., 2011; Gawlik-Dziki et al., 2013). The addition of guar gum (guar galactomannan) to
48 bread reduced *in vitro* starch hydrolysis by forming a physical ‘barrier’ to starch-alpha-
49 amylase interactions (Brennan et al., 1996). The mechanism was due to the layer of
50 galactomannan mucilage that coated the starch granules and bread matrix. Tea polyphenols
51 (epigallocatechin gallate and epigallocatechin) remain stable during bread making (Wang &
52 Zhou, 2004) and thus may confer further health benefits when incorporated into bread.
53 Studies have shown that starch hydrolysis was directly related to glycemia and insulinaemic
54 response in acute human feeding trials (Ekstrom et al., 2013). In this study we hypothesized
55 that the addition of black tea and/or beta glucan will change the characteristics and
56 functionality of wheat bread.

57 **2. Materials and methods**

58 *2.1 Materials*

59 White wheat flour and easy bake yeast were purchased from Allinson (Peterborough,
60 United Kingdom), unsalted butter from Morning Fresh (Caerphilly, United Kingdom), dried
61 skimmed milk powder from WM Morrisons Supermarkets PLC (Bradford, United Kingdom),
62 barley beta glucan concentrate (GlucageTM) from DKSH (containing 17.9% carbohydrate,
63 9.9% moisture, 4.9% protein and other nitrogenous compounds, 1.9% lipid and 2% ash
64 (according to manufacturer data) (Quai du Rhône, France) and freeze-dried pure tea granules
65 from Tata Global Beverages GB LTD (containing 452.17 ± 18.93 mg gallic acid
66 equivalents/100 ml) (Greenford, United Kingdom).

67 2.2 *Bread making*

68 Breads were prepared using standard bread making techniques (Burton-Freeman,
69 2010). All ingredients (**Table 1**) were weighed (in triplicate) on a digital kitchen scale
70 (Brabanita, UK) into a baking pan and mixed manually before being placed in a domestic
71 bread maker (Morphy Richards Ltd, South Yorkshire, UK). White bread (WF) and black tea
72 bread (BT) were prepared using standard program 8 (bread mixes, 150 min) while beta
73 glucan (β G) and beta glucan plus tea (β GBT) bread using program 5 (French bread, 210 min).
74 The addition of beta glucan competed with gluten for water; therefore, more water was
75 needed to compensate for water uptake by beta glucan (Jacobs et al., 2008). Additional
76 proofing time allows extended fermentation by yeast to take place and this improved dough
77 quality. The latter program allows extended proofing time and is crucial for the development
78 of both β G and β GBT breads. Fully developed bread was allowed to cool for a total of 60 min
79 which included 30 min in bread pan and 30 min at room temperature. The breads were
80 individually weighed and sliced into 1.5 cm thickness. All breads were analysed in triplicate
81 and used for each analysis. The structure of the breads was visualised using digital still
82 camera at 4x power (Sony Cybershot, Sony Corp, Japan).

83 2.3 *Proximate analysis*

84 2.3.1 *Protein content (Kjedahl method)*

85 One gram of bread was weighed in a digestion flask for protein analysis. The sample
86 was digested at 450°C for 1 h using Foss Tecator Digestor (Foss Tecator AB, Höganäs,
87 Sweden), then distilled and titrated in an automated Kjeltac 2300 Analyzer (Foss Tecator AB,

88 Höganäs, Sweden). A specific Jones factor for bread of 5.70 was used for calculation of
89 protein content.

90 2.3.2 *Fat content (solvent extraction, Soxtec method)*

91 One gram of bread sample was used for the determination of fat content. Seventy ml of
92 petroleum ether (Fisher Scientific, USA) was added to the sample in an aluminium thimble
93 which was carefully loaded into a Soxtec 2050 Automatic System (Foss Analytical AB,
94 Höganäs, Sweden).

95 All analyses were done in triplicate and nutrient contents are expressed as g/100 g fresh
96 weight.

97 2.4 *Resistant starch (RS)*

98 2.4.1 *Hydrolysis and solubilisation of non-resistant starch*

99 Resistant starch (RS) was determined using a Resistant Starch kit (Megazyme
100 International Wicklow, AOAC Method 2002.02 and AACC Method 32-40). Samples (100
101 mg) were incubated with pancreatic α -amylase (EC 3.2.1.1) containing amyloglucosidase
102 (AMG) (EC 3.2.1.3) at 37°C with continuous shaking (200 strokes /min) for 16 h. The tubes
103 were centrifuged at 3,000 rpm for 10 min, the supernatant was carefully decanted and the
104 pellets re-suspended in 2 ml of 50% ethanol with vigorous stirring on a vortex mixer. A
105 further 6 ml 50% ethanol was added, mixed and centrifuged again at 3,000 rpm for 10 min.
106 The supernatant was decanted and the extraction repeated followed by centrifugation.

107 *2.4.2 Measurement of resistant starch*

108 2 M KOH (2 ml) was added to each tube to re-suspend the pellets and stirred using a
109 magnetic stirrer for 20 min in an ice water bath. Thereafter, sodium acetate buffer (pH 3.8)
110 was added to each tube and incubated with 0.1 ml of AMG at 50°C for 30 min. An aliquot of
111 the solution was transferred into glass test tubes and 3.0 ml of glucose oxidase/peroxidase
112 (GOPOD) reagent added, incubated at 50°C for 20 min and absorbance measured at 510 nm
113 against the reagent blank (BioMate 3, Thermo Electron Corporation, Madison, USA).

114 *2.4.3 Measurement of digestible (solubilised) starch*

115 The supernatant obtained above was pooled in a volumetric flask and 0.1 ml incubated
116 with 10 µl of dilute AMG solution for 20 min at 50°C. Finally, 3.0 ml of GOPOD reagent
117 was added, incubated at 50°C for 20 min and absorbance read at 510 nm against a reagent
118 blank.

119 All analyses were done in triplicate and nutrient contents are expressed as g/100 g fresh
120 weight.

121 *2.5 Beta glucan content*

122 A Mixed-Linkage Beta-Glucan kit (McCleary method) from Megazyme International
123 Ireland (Wicklow, Ireland) was used for the determination of beta glucan. Sodium phosphate
124 buffer (4.0 ml, 20 mM, pH 6.5) was added to 1 g bread samples and stirred on a vortex mixer.
125 Lichenase (EC 3.2.1.73) was added and incubated for 1 h at 50°C. Sodium acetate buffer (5.0
126 ml, 200 mM, pH 4.0) was added and mixed on a vortex mixer. The tubes were centrifuged at
127 3,500 g for 10 min. Aliquots (0.1 ml) were incubated with β-glucosidase (EC 3.2.1.21) in 50

128 mM sodium acetate buffer (pH 4.0) at 50°C for 10 min. Finally, 3.0 ml of GOPOD reagent
129 was added, incubated at 50°C for 20 min and absorbance measured at 510 nm against a
130 reagent blank. The glucose content was calculated using Megazyme-Calc, a calculation sheet
131 provided by the manufacturer. All analyses were done in triplicate and nutrient contents are
132 expressed as g/100 g fresh weight.

133 2.6 *Starch hydrolysis of breads*

134 Starch hydrolysis was determined using a commercially available assay (Megazyme
135 International, Wicklow, Ireland). Breads containing 50 mg available carbohydrate were
136 added to 4 ml pancreatic α -amylase (5 mg/ml). Tubes were incubated at 37°C and the
137 reaction was stopped by adding 0.7 ml of ethanol (99% v/v) at different time points (0, 10,
138 30, 60, 90, 120, 150 and 180 min). Sodium acetate buffer (100 mM, pH 4.5) was added and
139 incubated with 10 μ l of dilute AMG solution for 20 min at 50°C. Finally, 3 ml of GOPOD
140 reagent was added, incubated for 20 min at 50°C and absorbance read at 510 nm against a
141 reagent blank. The glucose content was calculated using Megazyme-Calc, a calculation sheet
142 provided by the manufacturer. Glucose content was converted to starch using a 0.9
143 multiplying factor. Results are expressed as percentage (%) of total hydrolysed starch at
144 different time.

145 2.7 *Microscopic study of breads structures*

146 Breads were sampled from the centre of the bread loaf and processed using a standard
147 protocol of 70% alcohol (1 h), 90% alcohol (1 h), absolute alcohol (6 h 30 min), xylene (2 h
148 30 min) cycles and fixed in paraffin. The sections were cut into 2.5 μ m thickness with a
149 microtome (Shandon Finesse E, Thermo Scientific, Runcorn, United Kingdom) and dried in

150 an oven for 1 h at 60°C. The bread sections were stained with Lugol's iodine solution
151 [(0.33% I₂, w/v) and 0.67% KI (w/v)] (Sigma Aldrich, Steinheim, Germany) for 2 min
152 followed by 0.1% Light Green (Gurr, BDH Ltd., Poole, United Kingdom) for another 2 min.
153 The slides were visualised under light microscopy at the magnification power of 40x. Under
154 light microscopy, amylopectin stains brown, amylose stains dark brown (appears in the centre
155 of starch) and gluten stains light green.

156 2.8 *Statistical analysis*

157 Data are presented as the mean ± standard deviation. One-way ANOVA (SPSS version
158 21.0, SPSS Inc., Chicago, USA) and *post hoc* LSD test were used to determine the mean
159 differences between groups. General Linear Model (Repeated Measures) was used to
160 determine the effect of time interactions. Values are considered significantly different at the
161 level of $p < 0.05$. Coefficient of variation (CV) was calculated as standard deviation/mean x
162 100.

163 3. Results

164 3.1 *Bread characteristics*

165 White bread (WF) and black tea bread (BT) had significantly ($p < 0.05$) lower bread
166 volumes than beta glucan (β G) and beta glucan plus black tea (β GBT) bread (**Table 2**). All
167 breads showed similar length ranging from 12.1 to 13.4 cm. β G and β GBT breads had
168 significantly lower height than WF and BT breads (8.1 - 8.7 cm and 13.0 - 13.6 cm,
169 respectively; $p < 0.05$). The cross-section of the bread structure is shown in **Figure 1**. Both
170 BT and β GBT appeared darker because of the black tea.

171 3.2 *Nutrient composition*

172 The total available carbohydrate of WF and BT was significantly ($p < 0.05$) higher than
173 that of β G and β GBT (**Table 2**). There was no significant difference in resistant starch
174 content. Digestible starch was lower ($p < 0.05$) in β G and β GBT than WF and BT. The
175 protein content ranged from 7.0 - 9.1 g/100 fresh weight and was similar between the
176 different types of bread. Fat content was significantly ($p < 0.05$) higher in WF bread than the
177 other breads. Moisture content of bread added with beta glucan was significantly ($p < 0.05$)
178 higher compared to WF and BT. The addition of beta glucan significantly ($p < 0.05$) reduced
179 total energy content of β G and β GBT.

180 3.3 *Microscopic study of bread structures*

181 The bread microstructures, starch granules and protein (gluten), were studied and
182 visualised under a light microscope (**Figure 2**). Amylopectin granules stained brown,
183 amylose dark brown (appears in the middle of starch granule) and gluten light green. Starch

184 granules in WF and BT were swollen and sheared into small circular structures. In contrast,
185 starch granules in β G and β GBT were elliptical and closely packed to each other. The green-
186 stained gluten area was embedded in between starch granules and the porous (irregular white
187 structures) area of the breads. The gluten networks were more prominent and appeared as a
188 more continuous matrix in β G and β GBT compared with WF and BT.

189 3.4 Starch hydrolysis of breads

190 Standard flour (SF) showed the lowest starch hydrolysis from 10 to 120 min (**Figure 3**).
191 There were significant ($p < 0.05$) time interactions for SF at 90, 120, 150 and 180 min
192 compared with 10 min. There was no significant difference in starch hydrolysis between WF
193 and BT from 0 to 180 min. Both WF and BT showed significantly ($p < 0.05$) higher starch
194 hydrolysis at 10 min compared with β G and β GBT. Combination of beta glucan and black tea
195 in β GBT significantly ($p < 0.05$) reduced early (10 min) starch hydrolysis by 25% compared
196 with β G. There were significant ($p < 0.05$) time interactions for β GBT at 60, 90 and 150 min
197 compared with 10 min.

198 **4. Discussion**

199 The aim of this study was to develop breads to determine food matrix interactions of two
200 food components with established health effects. We determined the effect of adding beta
201 glucan and/or black tea on starch structures and the impact of these interactions on starch
202 characteristics and functionality. This is the first study to our knowledge of the development
203 of a bread with combination of beta glucan and black tea on starch hydrolysis and
204 microscopic study of starch granules. The composition of protein (gluten), starch and water
205 play an important role in making good quality bread (Flander et al., 2007). Gluten is
206 responsible for dough formation while starch is important in textural properties and stability
207 of the bread. Water hydrates and expands gluten forming a viscoelastic protein network. The
208 added beta glucan competed with gluten for water and decreased rising of the dough during
209 proofing; therefore, more water was needed to compensate for water uptake by beta glucan;
210 this was necessary to improve dough quality (Hager et al., 2010; Jacobs et al., 2008). The
211 presence of mixed linkage (1→3)(1→4)-β-D-glucans stabilizes the air cells in bread dough
212 and improves coalescence (Wang et al., 1998). Others have demonstrated that the addition of
213 sorghum flour in flat bread lowers the proportion of rapidly digestible starch (RDS) and
214 starch digestibility by 29% and 25%, respectively (Yousif et al., 2012).

215

216 *In vitro* studies demonstrated that black teas reduced *in vitro* starch hydrolysis (Zhang
217 & Kashket, 1998; Guzar et al., 2012). Mechanistically, a structural relationship of flavonoids
218 and α-amylase activity has been described with hydrogen bonding as well as the formation of
219 a conjugate that stabilized the interaction with the active site of α-amylase (Lo Piparo et al.,
220 2008). This effect may have been mediated through inhibition of amylase activity. However,
221 in our study the addition of black tea in bread (BT) did not reduce starch hydrolysis. This

222 may be explained by differences in experimental design. In our study, tea was added to
223 breads as an ingredient during baking. Therefore, we would anticipate that this would lead to
224 a different effect than the one observed in previous studies because of complex food matrix-
225 interaction between gluten, black tea and starch in bread. In bread, the addition of apple
226 pectin and polyphenols extracts, from kiwi, blackcurrant or apple in dough development and
227 bread baking, directly influenced the cross-linking of gluten polymers which could lead to
228 more water holding and softer bread (Sivam et al., 2011). The type of polyphenols used may
229 have different mechanisms in forming complexes with bread protein (gluten). Highly polar
230 phenolic acids (*i.e.* the caffeic acid present in kiwi) are more mobile in bread than low
231 polarity polyphenols (*i.e.* anthocyanins and proanthocyanins in blackcurrant and apple).
232 Caffeic acid is attracted to charged components in protein and/or directly incorporates into
233 the protein mesh work with less steric hindrance (Sun-Waterhouse et al., 2009; Sivam et al.,
234 2011). Hence, we proposed that the presence of black tea polyphenols could form cross-
235 linking with gluten which leads to softer bread, exposing starch granules to α -amylase
236 activity and increased starch hydrolysis.

237 In guar gum wheat bread, the presence of galactomannan coated the starch granules and
238 protein matrix during bread making process and subsequently reduced starch hydrolysis
239 (Brennan et al., 1996). Our study showed that black tea had an additional effect when added
240 together with beta glucan. The incorporation of black tea and beta glucan significantly
241 reduced short-term (10 min) starch digestibility in β GBT compared with β G bread. The
242 microscopy suggested that both beta glucan and black tea preserved the starch granule
243 structure, which resulted in lower short-term starch hydrolysis. Black tea contains higher
244 levels of high molecular weight theaflavins and thearubigins compared to the other types of
245 teas (Shao et al., 1995). We propose that there is a food-matrix interaction between black tea

246 polyphenols, beta glucan and starch granules in the bread. The presence of black tea
247 polyphenols could form complexes with the gluten-network while beta glucan preserved the
248 starch structure and/or decreased the surface area for starch digestion by α -amylase and hence
249 reduced starch hydrolysis.

250 The blunted early starch hydrolysis could translate to benefits for human health by
251 various mechanisms. Firstly, it could reduce post-prandial glucose and insulin response *in*
252 *vivo*. It has been previously demonstrated that the supplementation of guar bread reduced
253 post-prandial insulin response in healthy individual (Ellis et al., 1991). Secondly, the
254 polyphenol-linked beta glucan will be passed undigested and metabolised by gut microbiota
255 in the large intestine to short chain fatty acids and phenolic acids. Tea has been metabolised
256 by microbiota in the large intestine into phenolic acids and short chain fatty acids (Stalmach
257 et al., 2009; Stalmach et al., 2010). Beta glucan was fermented by intestinal microbiota and
258 significantly increased propionate giving an acetate:propionate:butyrate production ratio of
259 51:32:17 which was considered as propionate-rich (Hughes et al., 2008). It has been
260 demonstrated that phenolic acid metabolites and short chain fatty acids decreased the levels
261 of reactive oxygen species, modified gut microbial balance and promote satiety
262 (Beloborodova et al., 2012; Parkar et al., 2013; Arora et al., 2011). The incorporation of
263 propionate in foods could induce hypophagic effects through inhibition of cholesterol
264 synthesis, reduced lipolysis and delayed gastric emptying rate (Arora et al., 2011). Hence,
265 thirdly, the addition of beta glucan in bread will increase the level of propionate by the action
266 of gut microbiota and promoting greater satiety. A further mechanistic study is warranted to
267 study the structural relationship between polyphenols and dietary fibre in reducing starch
268 hydrolysis and their fermentability by gut microbiota.

269 **5. Conclusions**

270 This study showed that the addition of beta glucan alone or combination of black tea
271 and beta glucan influenced starch functionality in bread. The combination of black tea and
272 beta glucan reduced the hydrolysis of rapidly digestible starch. This finding suggests a food
273 matrix effect of beta glucan and black tea resulting in preserved elliptical starch granules and
274 lowered short-term starch hydrolysis. The applicability of these newly developed breads as
275 functional foods is promising.

276 **Acknowledgements**

277 The authors would like to acknowledge Dr Matthis Riehl at the Centre for Cell
278 Engineering Lab, University of Glasgow for the microscope facility and Ministry of Higher
279 Education, Malaysia for the funding granted.

280 **Declaration of interest**

281 The authors would like to declare that there is no conflict of interest.

282

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Figure 1. Detail of bread structure. The structure of bread was visualised at 4x power. (a) WF, white bread; (b) BT, black tea bread; (c) β G, beta glucan bread; (d) β GBT, beta glucan plus tea bread.

Figure 2. Microscopic figures of bread structures. (a) WF, white bread; (b) BT, black tea bread; (c) β G, beta glucan bread; (d) β GBT, beta glucan plus tea bread. Amylopectin granule stains brown, amylose dark brown (appears in the middle of starch granule) and gluten light green.

Figure 3. Starch hydrolysis based on percentage (%) of total hydrolysed starch of different breads. WF, white bread; BT, black tea bread; β G, beta glucan bread; β GBT, beta glucan plus tea bread; SF, Standard flour (provided by supplier). Values are expressed as mean \pm standard deviation. Values with different letters are significantly different at the level of $p < 0.05$ within same time point. Asterisk (*) indicates significant ($p < 0.05$) time interactions vs 10 min. Coefficient of variation (CV) is less than 32.0%.