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1 **Supramolecular structure of jackfruit seed starch and its relationship with digestibility and**
2 **physicochemical properties**

3

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13 **ABSTRACT**

14 The influence of supramolecular structure on the physicochemical properties and digestibility
15 of jackfruit seed starch (JSS) were investigated. Compared with maize and cassava starches (MS
16 and CS), JSS had smaller granules and higher amylose content (JSS: 24.90%; CS: 16.68%; and MS:
17 22.42%), which contributed to higher gelatinization temperature (T_0 : 81.11°C) and setback viscosity
18 (548.9 mPa·s). From scanning electron microscopy, the digestion of JSS was observed mainly at the
19 granule surface. Due to its higher crystallinity (JSS: 30.6%; CS: 30.3%; and MS: 27.4%) and more
20 ordered semi-crystalline lamellae, JSS had a high RS content (74.26%) and melting enthalpy (19.61
21 J/g). In other words, the supramolecular structure of JSS extensively determined its digestibility and
22 resistance to heat and mechanical shear treatment.

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24 *Keywords:*

25 Jackfruit seed starch; Supramolecular structure; Resistant starch; Digestibility; Thermal properties

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38 **Highlights:**

39 ✓ Jackfruit seed starch (JSS) had higher resistant starch content than other starches

40 ✓ High crystallinity and ordered semi-crystalline lamellae were the major reasons for the
41 higher enzyme-resistance of JSS

42 ✓ Digestion of JSS occurred mainly at the granule surface

43 ✓ Digestion caused slight decrease in crystallinity and lamellar regularity of JSS

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58 Chemical compounds studied in this article

59 Starch (PubChem CID: 24836924); Sodium hydroxide (PubChem CID: 14798); Water (PubChem

60 CID: 962); Hydrochloric acid (PubChem CID: 313); Ethanol (PubChem CID: 702); Acetic acid

61 (PubChem CID: 176); Iodine (PubChem CID: 807); Potassium iodine (PubChem CID: 4875);

62 Sodium acetate (PubChem CID: 517045).

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65 **1. Introduction**

66 Starch is one of the most important carbohydrates in human diets and has been extensively used

67 as a food ingredient. Understanding starch digestibility is of great interest to food industry and

68 importance for diet-related disorders such as obesity, diabetes, and cardiovascular diseases. Not all

69 starch can be digested in the small intestine, where the portion of starch that is not digested is

70 termed resistant starch (RS) (Asp & Björck, 1992). Physiological benefits have been correlated to

71 the RS consumption (Englyst & Hudson, 1996; Jenkins et al., 1998), which notably alters fecal bulk

72 and short-chain fatty acid metabolism, thus promoting the colonic health (Jenkins et al., 1998).

73 Because hydrolysis influences all level of food processing and nutrition, several arguments

74 prevail for a closer examination of the effects of hydrolytic enzymes on native starch granules. The

75 hydrolysis process of starches includes the diffusion of enzymes to the granule surface, followed by

76 the adsorption and subsequent catalytic events (Colonna, Leloup & Buleon, 1992). Previous studies

77 have shown that the action of α -amylase on starches from different botanical origins results in

78 varied digestion kinetics and degradation patterns (Fuwa, Takaya, Sugimoto & Marshall, 1980;

79 Sarikaya, Higasa, Adachi & Mikami, 2000). Generally, starch is a mixture of two types of
80 macromolecules, amylose and amylopectin (Hizukuri, 1985). Double or single helices of amylose
81 and amylopectin can be packed to form amorphous and crystalline regions (Oates, 1997), which is
82 the basis of the supramolecular structure (granule morphology, fractal structure, lamellar structure,
83 and crystalline structure) of starch. There are many structural factors of starch that affect the pattern
84 and rate of enzymatic hydrolysis, such as the size and shape of granules, granule integrity, porosity
85 of granules, crystallinity, amylose/amylopectin ratio, phosphate content, proteins, and lipids on the
86 granule surface (Copeland, Blazek, Salman & Tang, 2009; Dona, Pages, Gilbert & Kuchel, 2010;
87 Planchot, Colonna, Gallant & Bouchet, 1995; Robertson, Wong, Lee, Wagschal, Smith & Orts,
88 2006; Tester, Qi & Karkalas, 2006). The features of native starch granules that control the site, rate
89 and extent of hydrolysis by α -amylase are interrelated and not easily definable. Thus, studying the
90 changes of supramolecular structure would help to build the ability to manipulate and understand
91 the hydrolysis of starch granules.

92 Jackfruit is one of the most popular tropical fruits grown in Asia especially in Thailand. Its
93 seeds take up 10–15% of the whole fruits and contain abundant starch and proteins. With the rapid
94 development of the cultivating and processing industry of jackfruit, however, most seeds are
95 discarded, which causes a huge waste of starch resource. Jackfruit seed starch has not been
96 considered and exploited as a potent source of starch. To solve this problem, there have been studies
97 on the isolation and the properties of starch extracted from jackfruit seeds to verify its applicability
98 in food, pharmaceuticals and other uses. Jackfruit seed starch has the Type-A crystallinity pattern and
99 a high amylose content (Madruga, de Albuquerque, Silva, do Amaral, Magnani & Neto, 2014).

100 Compared with other starches, jackfruit seed starch has significantly higher gelatinization

101 temperature and lower breakdown viscosity, suggesting that this starch can be used to products
102 where a high level of gelatinization is not desirable during cooking (Bobbio, EI-Dash, Bobbio &
103 Rodrigues, 1978; Kittipongpatana & Kittipongpatana, 2011; Rengsutthi & Charoenrein, 2011;
104 Theivasanthi & Alagar, 2011; Tulyathan, Tananuwong, Songjinda & Jaiboon, 2002; Yi &
105 Shenghong, 2006). However, the literature provides little information about the structural features
106 of jackfruit seed starch and its effects on different properties. In particular, while the supramolecular
107 structure and its effect on the hydrolysis of native jackfruit seed starch are essential to ensure the
108 nutritional value and a diverse range of food industry uses, this information has not been reported so
109 far.

110 The aim of the present study was to investigate the functional properties and enzyme digestion
111 of jackfruit seed starch, as well as the related hierarchical structure changes in the native starch
112 granule that control the susceptibility of starch to enzymatic hydrolysis. The results of jackfruit seed
113 starch were compared with cassava starch and maize starch, which are two of the most popular
114 starches used in food industry. This would provide us with nutritional implications which are
115 instrumental for practical applications.

116

117 **2. Materials and methods**

118 *2.1. Materials*

119 Jackfruit Seed Starch (JSS) was isolated from jackfruit seeds using a modified method of
120 (Bobbio, EI-Dash, Bobbio & Rodrigues, 1978). The seeds were manually separated from the
121 mucilage, and then the aril and spermoderm were peeled off. The peeled seeds were slurried in a
122 Waring Blender (HR 1727 Philips, Zhuhai, China) with an equal weight of a 0.1% sodium

123 hydroxide solution for approximately 10 min. Then, the slurry was pressed through multiple gauzes
124 to remove seed fibers. The resulting milking suspension was allowed to decant at 4–5°C and
125 rewashed with distilled water to eliminate soluble sugars. The supernatant was drained, and the
126 upper brown sediment was scraped. The remaining sediment was mixed with 0.1% sodium
127 hydroxide solution and filtered through a sieve (0.058 mm mesh size) to eliminate fibers. When the
128 supernatant became clear, the filtrate was neutralized with 0.1M hydrochloric acid to pH 7.0, and
129 the slurry was centrifuged at 3,000 g for 20 min. The starch was dried at 40°C for 24 h. The starch
130 was grounded with a mortar, passed through a sieve (0.15 mm mesh size), packed in a plastic bag
131 and kept at room temperature until further use. The yield of JSS from Jackfruit seed was
132 25.45–27.34 g/100 g (dry basis).

133 Cassava starch (CS) was purchased from Vietnamese Food and Investment Co., Ltd. (Nanning,
134 China). Maize starch (MS) was from Inner Mongolia Wang Yu Biotechnology Co., Ltd. (Inner
135 Mongolia, China). The moisture contents of JSS, CS, and MS, determined using a moisture
136 analyzer (DHS20-1, Sartorius Stedim Biotech GmbH, Germany), were 13.03%, 13.44%, and
137 13.25%, respectively. Porcine pancreatic α -amylase and amyloglucosidase were purchased from
138 Sigma-Aldrich (St. Louis, MO, USA). A glucose-oxidase peroxidase (GOPOD) assay kit was from
139 Megazyme International Ireland, Ltd. (Wicklow, Ireland). Potato amylose was purchased from
140 Heilongjiang Academy of Agricultural Sciences (Harbin, China).

141

142 2.2. Starch characterization

143 2.2.1. Amylose content analysis

144 The RS content of each sample (JSS, CS, and MS) was determined using a modified method of

145 ISO 6647-2:2007, of the International Standardization Organization (ISO, 2007).

146 0.1 g of the starch (dry basis) was accurately weighed and dissolved in 1 ml of ethanol and 9 ml
147 of sodium hydroxide solution (1 M), then heated in boiling water for 10 min. After cooling off, this
148 solution was then diluted to 100 mL in a volumetric flask with deionized water. An aliquot (2.50
149 mL) of this solution was then diluted with 25.00 mL of water, 0.50 mL of acetic acid solution (1 M),
150 0.50 mL of I₂/KI solution (0.0025 M I₂, and 0.0065 M KI), and the absorbance of this solution was
151 read in a 1cm path length quartz cell at 620 nm using an Evolution UV/Visible spectrophotometer
152 (Thermo Scientific, Waltham, USA). The amylose from potato (amylose content: 97.0%) was used
153 for the calibration curve ($R^2=0.9962$).

154

155 2.2.2. *Differential scanning calorimetry (DSC)*

156 Thermal behaviors of JSS, CS, and MS were studied using a PerkinElmer DSC 8000
157 (PerkinElmer, Waltham, America) with an internal coolant (Intercooler 2P) and nitrogen purge gas.
158 A high-pressure stainless steel pan (PerkinElmer No. B0182901) with a gold-plated copper seal
159 (PerkinElmer No. 042-191758) was used to achieve a constant moisture content (MC) during DSC
160 measurements. The sample, with about 70% MC, was prepared by premixing the starch with added
161 water in a sealed glass vial, which was kept at 20°C for 24 h before measurement. About a 4 mg
162 (dry basis) sample, scanned from 40 to 120°C, was used in this study. A slow heating rate of
163 5°C/min was used. The onset temperature (T_o), peak temperature (T_p), conclusion temperature (T_c),
164 and enthalpy (ΔH) of starch gelatinization were calculated. The enthalpy was calculated based on
165 the weight of dry basis starch.

166

167 2.2.3. *Pasting properties*

168 Pasting properties were studied using an Anton Paar MCR302 (Anton Paar China, Shanghai,
169 China). The sample slurry (6% concentration, starch on dry basis), after 1 min pre-shearing, was
170 heated from 30°C to 95°C at a heating rate of 5°C/min, held at 95°C for 15 min, and cooled to 50°C
171 at 5°C/min. Then the sample was held at 50°C for 15 min. The changes of viscosity were recorded.

172

173 2.3. *Enzyme digestion of starches*

174 2.3.1. *In vitro digestibility of native starches*

175 For native JSS, CS, and MS, the starch digestibility was determined following the modified
176 method of Englyst (Englyst, Kingman & Cummings, 1992). Based on the rate of hydrolysis, starch
177 was defined as rapidly-digestible starch (RDS, digested within 20 min), slowly-digestible starch
178 (SDS, digested between 20 min and 120 min), and resistant starch (RS, undigested within 120 min).

179 In brief, porcine pancreatic α -amylase (3 g) was dispersed in water (20 mL), stirred for 10 min
180 and centrifuged at 3000 g for 15 min. The supernatant (13.5 mL) was transferred to a beaker, and
181 225 U of amyloglucosidase and 1 mL of deionized water were added to the solution. The enzymatic
182 solution should be freshly prepared for each digestion. Duplicate samples (one named Sample A,
183 the other Sample B) of each starch (JSS, CS, and MS) (1 g, dry basis) were dispersed in 20 mL of
184 0.1 M sodium acetate buffer (pH = 5.2) and then mixed with an enzyme solution (5 mL) consisting
185 of the pancreatic extract and amyloglucosidase. The dispersion was incubated in a 37 °C shaking
186 water-bath at 180 strokes/min. An aliquot (0.5 mL) of Sample A was taken at interval of 20 min and
187 mixed with 20 ml of 70% ethanol. The mixed solution of Sample A was centrifuged at 3000 g for 10
188 min, and then the supernatant was used for hydrolyzing the glucose content, measured by the

189 glucose oxidase-peroxidase reagent. Sample B was mixed with ethanol to eliminate the activities of
190 enzyme, and then the dispersion was centrifuged at 3,000 g for 20 min. After three times of mixing
191 with ethanol and centrifugation, the sediments of Sample B were dried at 40°C for 12 h, named
192 JSS-20, CS-20, and MS-20 (“20” means the time interval (min) for which the three starches were
193 hydrolyzed), respectively. When the time interval reached 120 min, another aliquot (0.5 mL) of
194 Sample A was taken and mixed with 20 ml of 70% ethanol, centrifuged to analyze the hydrolyzed
195 glucose content. The sediments were treated using the same method of Sample B. These sediments
196 were JSS-120, CS-120, and MS-120, respectively.

197 2.3.2. *Scanning electron microscopy (SEM)*

198 Granule morphology was studied using an EVO18 scanning electron microscope (ZEISS,
199 Germany) operated at a high voltage of 10.0 kV. Before the SEM examination, the samples were
200 coated with a gold thin film.

201

202 2.3.3. *Small-angle X-ray scattering (SAXS)*

203 A SAXSess small angle X-ray scattering system (Anton Paar, Austria), operated at 50 mA and
204 40 kV, using Cu K α radiation with a wavelength of 0.1542 nm as the X-ray source, was applied to
205 perform the SAXS measurements according to our previously method (Zhu, Li, Chen & Li, 2012)
206 with proper modification. Each sample was placed in a paste sample cell and exposed at the
207 incident X-ray monochromatic beam for 10 min. The data, recorded using an image plate, were
208 collected by the IP Reader software with a PerkinElmer storage phosphor system.

209 The samples used for the SAXS measurement were prepared by premixing the starch with
210 added water in glass vials and were equilibrated at 20°C for 24 h before the analysis. The total MC

211 of each sample was 65%. All data were normalized, and the background intensity and smeared
212 intensity were removed using the SAXSquant 3.0 software for further analysis.

213

214 2.3.4. *Polarized light microscopy*

215 Polarized light microscopy was performed using a polarized light microscope (PLM)
216 (Axioskop 40 Pol/40A Pol, ZEISS, Oberkochen, Germany) equipped with a 35mm SLA camera
217 (Power Shot G5, Canon, Tokyo, Japan). The magnification was 500 (50×10). Each sample was
218 dispersed as 10 mg (wet basis) of starch in 1 mL of distilled water in a glass vial. Then, a drop of
219 the starch suspension was transferred onto a slide and covered by a coverslip. Polarized light was
220 used for observation.

221

222 2.3.5. *X-ray diffraction (XRD)*

223 XRD analysis was performed with an Xpert PRO diffractometer (Panalytical, Netherlands),
224 operated at 40 mA and 40 kV, using Cu K α radiation with a wavelength of 0.1542 nm as the X-ray
225 source. The scanning of diffraction angle (2θ) was from 5° to 40° with a scanning speed of 10°/min
226 and scanning step of 0.033°. The MC of each sample was about 10%. The relative crystallinity of
227 each sample was calculated using a previous method (Hermans & Weidinger, 1948).

228

229 2.4 *Statistical analysis*

230 The mean values and differences were analyzed using Duncan's multiple-range test. Analysis of
231 variance (ANOVA), followed by the least significant difference test (LSD-test), was performed
232 using the software SPSS (Version 22.0). The significance level was set at $p < 0.05$.

233

234 3. Results and discussion

235 3.1. Amylose contents and *in vitro* enzyme digestion analysis of native starches

236 The amylose/amylopectin ratio is an important index of starch and it can influence digestion
237 and swelling properties through the way of amylose and amylopectin packed. As seen from Table 1,
238 compared to CS and MS, the amylose content of JSS was higher (24.90%), which was similar to a
239 previous finding (Li & Zhong, 2004). CS had the lowest amylose content, only 16.68%. Based on
240 the Englyst test, the percentages of RDS, SDS, and RS in JSS were 5.92%, 19.82%, and 74.26%,
241 respectively. The RS content of JSS was much higher than CS and MS while RDS and SDS were
242 lower, indicating that JSS had strong anti-enzymatic capability. Interestingly, MS had the lowest RS
243 content but the highest SDS content, suggesting that it is a good material of SDS. The
244 slow-digestion property of MS is more likely to be controlled by its inherent structure (perhaps
245 amylopectin chain length distribution) although the existence of surface porous channels might
246 contribute to a high rate of starch hydrolysis (Zhang, Ao & Hamaker, 2006).

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250 3.2. Supramolecular structure characteristics of native and hydrolyzed starches

251 3.2.1. Granule morphology

252 Fig. 1 shows the SEM images of JSS, CS and MS in their native states and after 20min and
253 120min enzyme hydrolysis. The JSS and CS granules had round to bell shapes with a smooth
254 surface. Unlike the other two starches, the MS granules were irregular in shape with small pores and

255 pits randomly distributed on a rough surface. The JSS granules were less irregular in shape, being
256 smaller than the CS and MS granules.

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260 The susceptibility of starch granules can be classified by the degree and manner by which the
261 granules are eroded and corroded. As seen from SEM, the degree of digestion of starch followed the
262 order: MS > CS > JSS, contrary to the trend of RS (Table 1), which is as expected. Besides, the
263 observed levels of digestion were comparable between large and small granules for all three raw
264 starches. Some small granules in JSS-20 and CS-20 even became hollow with only a thin external
265 shell structure. This suggests a fundamental difference in the mode of α -amylase and
266 amyloglucosidase action, according to the granule size. Smaller granules, by virtue of their higher
267 available surface area per unit mass, facilitate the diffusion and adsorption of enzymes (Colonna,
268 Leloup & Buleon, 1992).

269 Digestion of JSS was not clearly apparent; the main indication was a less smooth and more
270 rugged granule surface with a few pits (JSS-20 and JSS-120, in Fig.1). Enzymatic digestion of CS
271 was apparent from the increased surface roughness and formation of deep cracks and large holes in
272 many granules (CS-20 and CS-120 in Fig.1). After 20min of enzymatic digestion, some CS granules
273 were in a truncated form (CS-20 in Fig.1). Truncatures are weak points in the granule structure that
274 lead to increased susceptibility, resulting in enhanced hydrolysis of CS. (Valetudie, Colonna,
275 Bouchet & Gallant, 1993). Because of no pores and smooth surfaces, SEM micrographs for JSS and
276 CS showed that enzymatic erosion occurred mainly at the surface. The MS granules showed

277 extensive corrosion, mainly in the direction of the radial axis and only a few granules remained
278 intact. The surface pores of hydrolyzed MS became larger and deeper into granules because of the
279 more extensive hydrolysis (MS-20 in Fig.1). After 120min hydrolysis, some granules were split,
280 exposing their layered internal structure (MS-120 in Fig.1). The layered internal structure showed
281 different susceptibility of the semi-crystalline structure and amorphous growth rings toward
282 digestion (Zhang, Ao & Hamaker, 2006).

283

284 3.2.2. Lamellar structure characteristics

285 The double-logarithmic SAXS patterns of native and hydrolyzed starch residues are shown in
286 Fig. 2. From this figure, we can obtain some parameters of a theoretical model for the lamellar
287 structure in starch (Cameron & Donald, 1993a, b), including d , the average thickness of the
288 semi-crystalline lamellae; $\Delta\rho = \rho_c - \rho_a$ (where ρ_c and ρ_a are the electron densities of the crystalline
289 regions and the amorphous regions in the semi-crystalline lamellae), the difference in electron
290 density between the crystalline lamellae and the amorphous lamellae; $\Delta\rho_u = \rho_u - \rho_a$ (where ρ_u is the
291 electron density of the amorphous background), the difference in electron density between the value
292 of q of the peak at *ca.* 0.6 nm^{-1} can be used to calculate the average repeat distance (d) of the
293 semi-crystalline lamellae in starch granules according to the Woolf-Bragg's equation $d = 2\pi/q$
294 (Blazek & Gilbert, 2010; Vermeylen, Goderis & Delcour, 2006). Table 2 shows the SAXS
295 parameters from the peaks of native and hydrolyzed starches. It can be seen from Table 2 that the
296 average thickness of the semi-crystalline lamellae of JSS and CS were thinner than that of MS (JSS:
297 9.06 nm; CS: 9.14 nm; and MS: 9.42 nm) and the peak areas of JSS and CS were larger than MS
298 (JSS: 0.1288; CS: 0.1248; and MS: 0.0800). This indicates JSS and CS may have more ordered

299 semi-crystalline lamellae than MS.

300

301

302

303 The $\log I \sim \log q$ SAXS patterns of JSS, CS, and MS and their hydrolyzed residues are
304 presented in Fig. 2a, b and c. The scattering intensity changed slightly for JSS (JSS-20 and JSS-120
305 in Fig. 2a) during the whole enzymatic hydrolysis. After 120min hydrolysis, the scattering intensity
306 at the low q region showed an increasing trend (JSS-120 in Fig. 2a) and the definition of the peak of
307 JSS-120 was lower than those of JSS and JSS-20. This can be explained by the easier disturbance of
308 starch molecular arrangement in the amorphous background than in the amorphous lamellae by
309 α -amylase, thus resulting in an increase in $\Delta\rho_u$ (Cameron & Donald, 1992). All the analysis of JSS
310 showed that most of the semi-crystalline lamellae of JSS remained intact even after 120min
311 hydrolysis. And the slight changes in the scattering intensity of JSS, JSS-20, and JSS-120 explained
312 a high RS content of JSS and less obvious surface erosion. However for CS and MS (CS-20 in Fig.
313 2b and MS-20 in Fig. 2c), the q region around the peak showed a decreasing trend, suggesting the
314 crystalline regions in the semi-crystalline lamellae were disturbed after 20min hydrolysis. And the
315 scattering intensity at the low q region showed an increasing trend, due to more destruction to the
316 amorphous background than to the amorphous lamellae. After 120min hydrolysis (CS-120 in Fig.
317 2b and MS-120 in Fig. 2c), the scattering intensity decreased to an extensive degree. It is noted that
318 the decrease of scattering intensity in MS was faster during the first 20 min of enzymatic hydrolysis
319 and slower from 20 min to 120 min than in CS. This could be an excellent explanation for the
320 higher SDS content of MS. Based on the above discussion, a conclusion can be made that the

321 semi-crystalline lamellae of JSS were more ordered and thus more resistant to the hydrolysis than
322 those of CS and MS.

323

324 *3.2.3 Crystalline characteristics*

325 Normally, a birefringence cross can be observed when the starch granule is exposed under
326 polarized light, due to orderly-arranged starch molecules of crystalline regions and
327 disorderly-arranged starch molecules of amorphous regions. Therefore, information about the
328 crystalline structure of starch can be reflected by the birefringence pattern when starch granules
329 suffered from hydrolysis or external attack. The polarized light microscope images of JSS, CS, and
330 MS and their hydrolyzed residues are shown in Fig. 3. Given the different sizes of JSS, CS, and MS
331 granules, native JSS showed weaker birefringence intensity than CS and MS, while CS showed the
332 strongest intensity. It is noted that the birefringence intensity remained almost the same for JSS after
333 enzyme hydrolysis for 120 min, suggesting most of crystalline structure of JSS was retained.
334 Nevertheless, the birefringence intensity decreased significantly for CS and MS (especially for MS),
335 and the birefringence crosses became less apparent, owing to the disturbance of double helices in
336 their crystallites during enzyme digestion. This result is consistent with the analysis of SAXS.

337

338

339

340 Fig. 4 shows the XRD patterns of JSS, CS, and MS, and their hydrolyzed residues. It is seen
341 that JSS and MS displayed a typical A-type crystalline structure with main diffraction peaks at *ca.*
342 15, 17, 18 and 23° (2θ) (Tulyathan, Tananuwong, Songjinda & Jariboon, 2002; Zobel, 1964). CS

343 exhibited a weak diffraction maximum at $5.6^\circ(2\theta)$, and the $17^\circ(2\theta)$ peak was somewhat more
344 intense than its $18^\circ(2\theta)$ neighbor (Chrastil, 1987). Both features indicated CS contained some
345 B-type crystalline structure but the main structure was still A-type. The degree of relative
346 crystallinity of starch followed the order: $JSS \approx CS > MS$. According to the XRD patterns of
347 partly-digested starches of JSS, CS and MS, the crystalline types of all three starches remained
348 essentially unchanged after digestion. However, after enzyme treatment, decreased diffraction
349 intensities were observed (Figure 4a, b, and c). The relative crystallinity of JSS changed moderately,
350 decreased from 30.6% to 27.6% (Table 2) after 20min digestion, while CS and MS decreased more
351 sharply from 30.3% to 23.6% and 27.4% to 19.4%, respectively. These results suggest that
352 hydrolysis did occur in the crystalline regions despite that most of crystalline structure of JSS was
353 retained after 120min hydrolysis.

354

355 It is noted that although JSS and CS both had a smooth surface and similar relative crystallinity
356 (Table 2), the RS content of JSS was higher than CS. This can be demonstrated by the observation
357 that the degree of the ordered structure in semi-crystalline lamellae was in the order $JSS \cdot CS \cdot MS$
358 in the SAXS, suggesting not only the crystallinity but the way how molecules are ordered play a
359 key role in the enzyme digestion of JSS. Another reason could be due to their amylose/amylopectin
360 ratio. Specifically, a higher amylose content may mean an increased number of long chains and
361 facilitate the amylose-lipid complex formation on the granule surface, leading to an increased
362 content of enzyme-resistant starch (Crowe, Seligman & Copeland, 2000; Cui & Oates, 1999;
363 Tufvesson, Skrabanja, Björck, Elmståhl & Eliasson, 2001). The surface pores and low relative
364 crystallinity of MS could contribute to its high RDS and low RS contents.

365 When the α -amylase attacks starch granules, the double helices must first be unwound, as
366 single-stranded helices are the polymeric substrates for the enzyme (Larson, Day & McPherson,
367 2010). The amylopectin double helices can only be unwound if they are dissociated from their
368 crystallites. However, the amylopectin side chains of starch strongly interact, not only with their
369 helical duplex partners, but also with other neighboring helices. Thus, more ordered crystalline
370 structure leads to a lower rate of enzymatic hydrolysis because of stronger interactions between
371 neighboring helices. Normally, higher crystallinity is in consistent with more ordered arrangement
372 of amylopectin double helices in the semi-crystalline lamellae, since the crystallinity reflects the
373 long range order of starch. In the light of these principles, the more ordered crystalline structure
374 (corresponding to more ordered semi-crystalline lamellae and high relative crystallinity) was the
375 main reason for the strong anti-enzymatic capability of JSS.

376

377 3.3. *Thermal behavior*

378 Fig.5a shows the DSC thermograms of JSS, CS and MS in excess water (70 wt.%) and the
379 related thermal parameters were shown in Table 3. From Fig.5a and Table 3, it was obvious that JSS
380 had the highest gelatinization temperature (T_0 : 81.11°C), followed by MS (T_0 : 65.58°C) and CS (T_0 :
381 60.47°C). The higher T_0 , T_p , and T_c of JSS could be due to a higher content of amylose-lipid
382 complexes with an increased amylose content, resulting in reduced swelling of the granule
383 (Karkalas & Raphaelides, 1986; Pycia, Juszczak, Galkowska & Witczak, 2012; Svihus, Uhlen &
384 Harstad, 2005; Tester & Morrison, 1990). The higher gelatinization temperature of JSS may also
385 reflect its much longer amylopectin chains, as there is a significant positive correlation between the
386 DSC gelatinization parameters and the amylopectin unit-chain length distribution of starches (Jane

387 et al., 1999; Noda et al., 1998; Shi & Seib, 1995; Srichuwong, Sunarti, Mishima, Isono &
388 Hisamatsu, 2005a). Since the granule size followed the order CS • MS • JSS (Fig.1), another reason
389 could be related to the size of starch granules since larger granules might be more vulnerable during
390 heating (Chiotelli & Le Meste, 2002; Kaur, Singh & Sodhi, 2002; Vasanthan & Bhatta, 1996). JSS
391 and MS showed rather symmetric peaks and had similar ΔT , which was narrower than that of CS.
392 This indicates that the crystalline structure of JSS and MS are more unified and consistent than that
393 of CS, resulting in more homogeneous heat conductivity. Higher ΔT of CS was proposed to arise
394 from the inconsistency of crystalline structure corresponding to the melting of B-type in CS
395 although the main structure in CS was A-type. JSS and CS had similar ΔH (Table 3), due to their
396 similar relative crystallinity, which were higher than that of MS. The higher ΔH values suggested
397 that the interactions (via hydrogen bonding) between double helices (which were packed in clusters)
398 forming the crystalline regions of JSS and CS were probably more extensive than in MS (Cooke &
399 Gidley, 1992; Zhou, Hoover & Liu, 2004).

400

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404 3.4. Pasting properties

405 Fig.5b shows the pasting properties of JSS, CS and MS. As seen from Table 3, the peak
406 viscosity (PV) of three starches followed the order JSS • CS • MS, which corresponded to the trend
407 of T_0 . The breakdown viscosity (BDV) of JSS (109.5 mPa·s) was lower than those of CS and MS

408 (473.2 mPa·s and 288.4 mPa·s, respectively). When viscosity reached PV, almost all of amylose
409 leached out and therefore BDV was less affected by amylose, but more by amylopectin fine
410 structure (Han & Hamaker, 2001). Lower BDV is another indicator that JSS may have much longer
411 amylopectin chains since dissociation of double helices of amylopectin leads to granule swelling
412 and affects pasting properties to some extent (Han & Hamaker, 2001; Srichuwong, Sunarti,
413 Mishima, Isono & Hisamatsu, 2005b). The final viscosity (FV) and setback viscosity (SBV)
414 indicate the re-association of the starch molecules involving amylose after gelatinization and a
415 formation of a gel network (Charles, Chang, Ko, Sriroth & Huang, 2004). JSS had higher FV and
416 SBV than CS and MS (Table 3), owing to a high amylose content (Sasaki, Yasui & Matsuki, 2000;
417 Vandeputte, Derycke, Geeroms & Delcour, 2003). The reason CS had less amylose content but
418 higher FV and SB than MS might be due to the finer amylopectin structure (enrichment in B2
419 chains) of CS (Srichuwong, Sunarti, Mishima, Isono & Hisamatsu, 2005b).

420

421 **4. Conclusion**

422 JSS granules were shown to be small, round to bell shapes, with a smooth surface and
423 displayed a typical A-type crystalline structure. Compared with MS and CS, JSS had higher
424 amylose content, higher RS content and more ordered semi-crystalline lamellae. According to the
425 DSC measurement, JSS had the highest T_0 . This might be because of the reduced swelling of the
426 granule, probably due to more amylose-lipid complexes with higher amylose content and to its
427 smaller granules which were more resistance to heat. JSS and CS had similar ΔH , due to their
428 similar relative crystallinity. From the pasting property study, the BDV of JSS was lower than those
429 of CS and MS while FV and SBV were higher. Lower BDV might indicate longer amylopectin

430 chains of JSS, which needs further investigation. As seen from SEM, the degree of digestion of
431 starch followed the order: MS > CS > JSS. Digestion of JSS only apparently occurred at the surface,
432 with a less smooth and more rugged granule surface with occasional pitting. In the course of
433 digestion, for JSS, the scattering intensity and the relative crystallinity were decreased slightly, and
434 the birefringence intensity remained almost the same. These observations indicate the more ordered
435 semi-crystalline lamellae and high relative crystallinity were the major factors for the stronger
436 anti-enzymatic capability of JSS than those of CS and MS. In conclusion, the results presented the
437 detailed related supramolecular structure changes (especially granular, crystalline, and lamellae
438 structure) of JSS granules that control the susceptibility of starch to enzymatic hydrolysis and the
439 physicochemical properties. The knowledge obtained from this work is expected to facilitate further
440 research on the nutritional and other properties of JSS for widening its industrial application.

441

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646 Figure Captions

647 Fig.1. SEM images of native and hydrolyzed starch residues at 1000 \times and 3000 \times magnification

648 Fig.2. Double-logarithmic SAXS patterns of native and hydrolyzed starch residues. (a) jackfruit

649 seed starch (JSS, JSS-20, and JSS-120); (b) cassava starch (CS, CS-20, and CS-120); (c) maize

650 starch (MS, MS-20, and MS-120).

651 Fig.3. Polarized light microscopic images of native and hydrolyzed starch residues

652 Fig.4. XRD patterns of native and hydrolyzed starch residues, (a) jackfruit seed starch (JSS,
653 JSS-20, and JSS-120); (b) cassava starch (CS, CS-20, and CS-120); (c) maize starch (MS, MS-20,
654 and MS-120).

655 Fig.5. Differential scanning calorimetry (DSC) thermographs (a), and viscosity curves (b) of
656 jackfruit seed starch, cassava starch and maize starch

657

658 Tables

659 Table 1 Amylose contents and *in vitro* enzyme digestion analysis of jackfruit seed starch (JSS), cassava starch (CS) and
660 maize starch (MS).

Raw starches	RDS (%)	SDS (%)	RS (%)	Amylose (%)
Jackfruit seed starch (JSS)	5.92±0.49 ^c	19.82±1.01 ^c	74.26±1.28 ^a	24.90±0.10 ^a
Cassava starch (CS)	10.50±0.04 ^b	38.43±0.03 ^b	51.07±0.08 ^b	16.68±0.54 ^c
Maize starch (MS)	12.04±0.04 ^a	69.73±1.14 ^a	18.23±1.18 ^c	22.42±0.19 ^b

661 Values are means of three determinations (±standard deviation); values followed by the different letters within a column
662 differ significantly ($p < 0.05$).

663

664

665 Table 2. SAXS parameters and relative crystallinity of native and hydrolyzed starches.

Sample	q_{peak} (nm^{-1})	d (nm)	Peak Area	RC (%)
JSS	0.6934 ^{abc}	9.06 ^{de}	0.1288 ^a	30.6 ^{ab}
JSS-20	0.6868 ^{bcd}	9.15 ^{cd}	0.1240 ^a	28.5 ^{abcd}
JSS-120	0.7001 ^{ab}	8.98 ^{de}	0.0653 ^c	27.6 ^{bcd}
CS	0.6868 ^{bcd}	9.14 ^{de}	0.1248 ^a	30.3 ^{ab}
CS-20	0.6802 ^{cd}	9.24 ^c	0.0639 ^c	25.4 ^{def}
CS-120	0.6934 ^{abc}	9.01 ^{de}	0.0318 ^d	23.6 ^{efg}
MS	0.6670 ^e	9.42 ^b	0.0800 ^b	27.4 ^{cde}
MS-20	0.6604 ^e	9.51 ^b	0.0572 ^c	21.5 ^g
MS-120	0.6208 ^f	10.12 ^a	0.0213 ^d	19.4 ^g

666 Values are means of three determinations; values followed by the different letters within a column differ significantly (p
667 < 0.05).

668

669 Table 3 Gelatinization parameters and pasting properties of jackfruit seed starch (JSS), cassava starch (CS) and maize
 670 starch (MS)

Sample	JSS	CS	MS
T_o (°C)	81.11±0.53 ^a	60.47±1.00 ^c	65.58±0.45 ^b
T_p (°C)	85.39±0.64 ^a	65.88±0.78 ^c	69.43±0.15 ^b
T_c (°C)	91.70±1.12 ^a	79.32±0.84 ^b	75.48±0.38 ^c
ΔT ($T_c - T_o$)	10.59±0.65 ^b	18.85±1.85 ^a	9.90±0.09 ^b
ΔH (J/g)	19.61±0.76 ^a	19.67±0.41 ^a	15.86±0.32 ^b
PT (°C)	82.0±0.2 ^a	66.9±0.3 ^c	71.5±0.2 ^b
PV (mPa·s)	844.0±5.1 ^b	963.2±4.3 ^a	743.9±3.3 ^c
BDV (mPa·s)	109.5±2.4 ^c	473.2±1.5 ^a	288.4±1.8 ^b
FV (mPa·s)	1354.0±7.4 ^a	1044.0±6.3 ^b	827.9±5.2 ^c
SBV (mPa·s)	548.9±3.5 ^a	514.4±4.1 ^b	349.1±3.8 ^c

671 T_o , T_p and T_c correspond to onset, peak and conclusion gelatinization temperature (°C); whereas ΔH and ΔT represent
 672 melting enthalpy (J/g of starch) and gelatinization temperature range (°C) respectively. PT represents peak temperature
 673 (°C), whereas PV, BDV, FV, SBV correspond to peak viscosity, breakdown viscosity, final viscosity and setback
 674 viscosity (mPa·s) respectively.
 675 Values in the table are means of three determinations (\pm standard deviation); values followed by the different letters
 676 within a column differ significantly ($p < 0.05$).

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679 Figures

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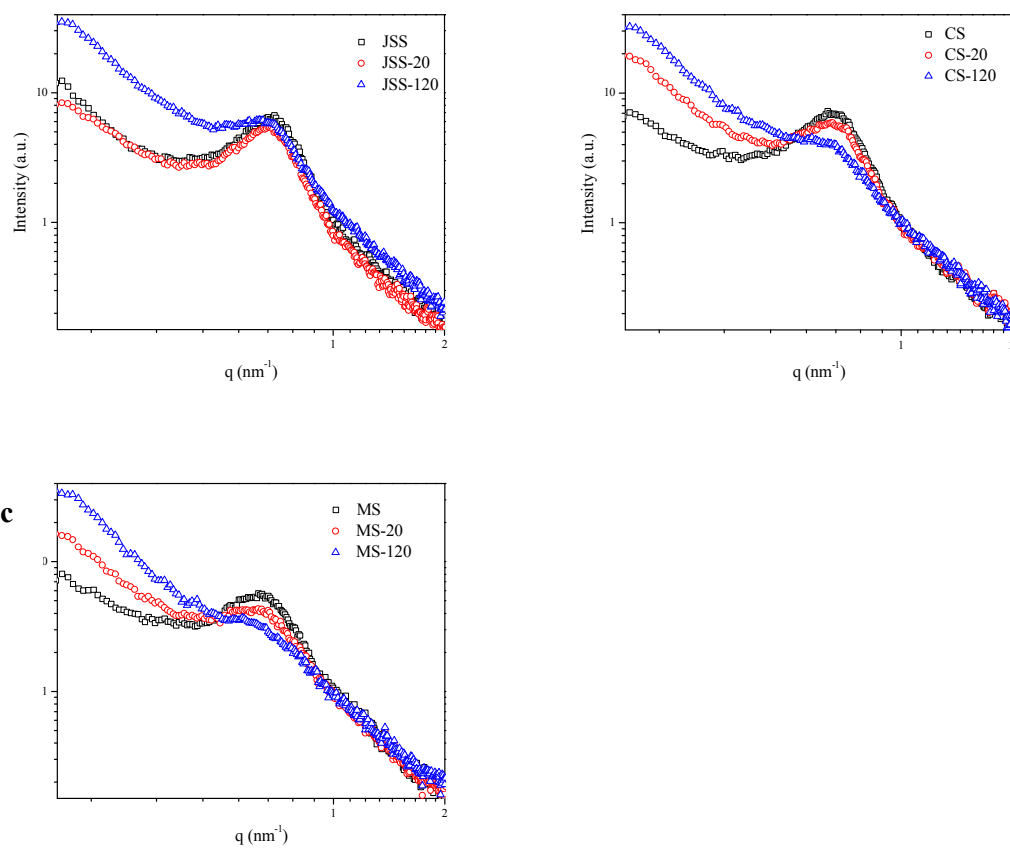
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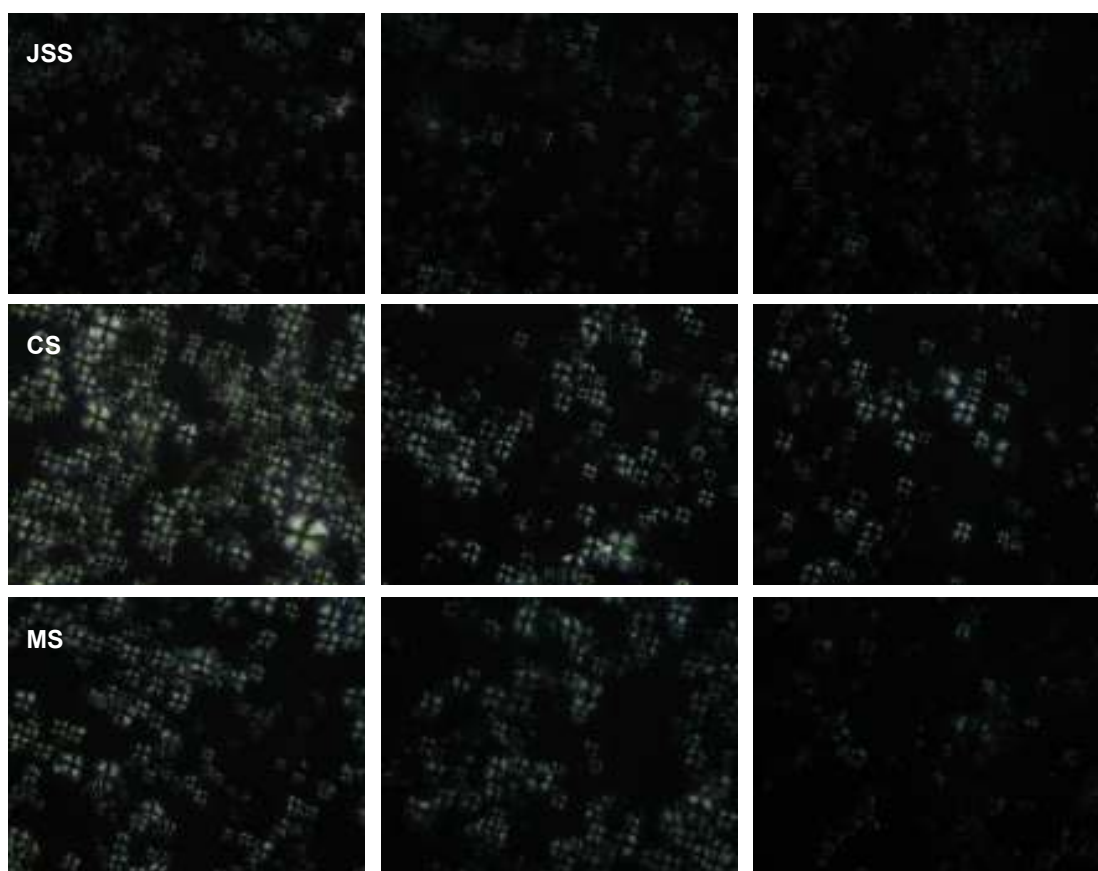
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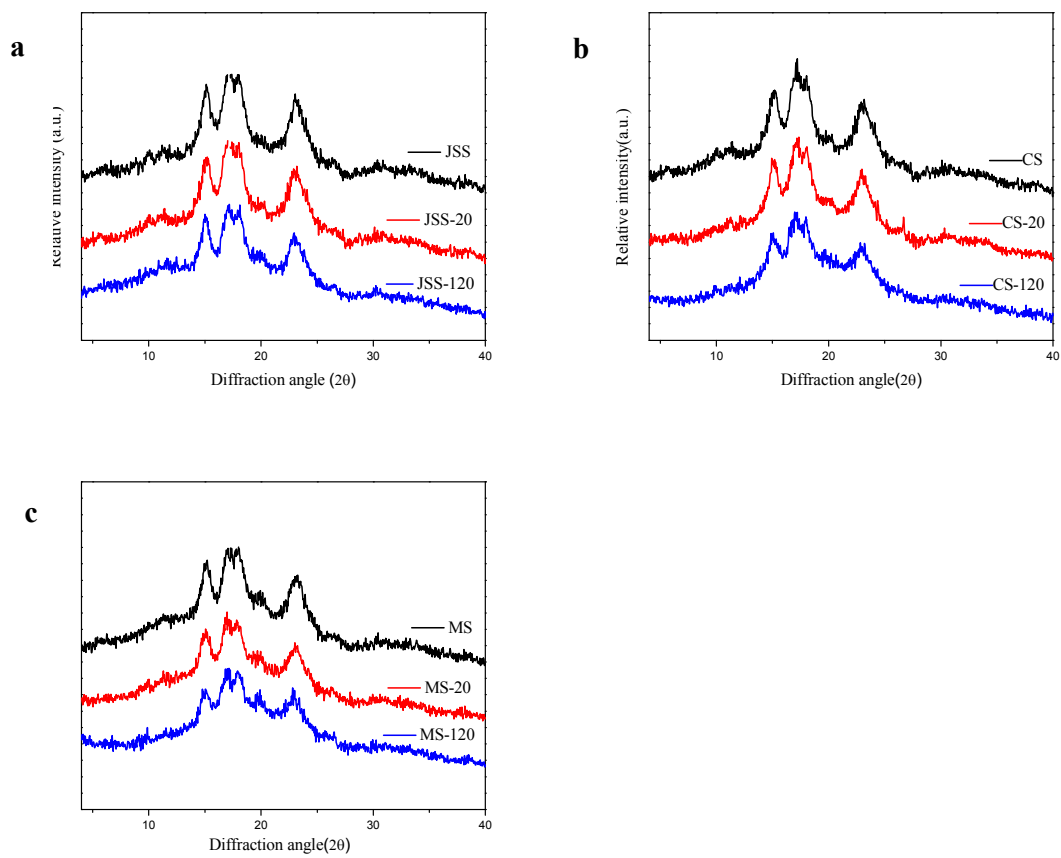
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711 Fig 5

