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2	Synbiotic potential of colored rice flour as wall material and encapsulated with
3	Lactobacillus plantarum TISTR 1465:Its impacts on gut bacterial population and metabolic
4	activities
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25 Abstract

Black waxy and red jasmine rice flour (6.5 and 18.7% amylose content, respectively) 26 were modified using pullulanase followed by heat-moisture treatment (hydrolyzed-HMT) to 27 produce potential synbiotic microencapsulation with Lactobacillus plantarum TISTR 1465. 28 Hydrolyzed-HMT of colored rice flours showed restricted pasting properties, lower breakdown 29 and higher thermal properties than native flour (P < 0.05). Hydrolysis treatment was able to 30 promote a low molecular weight starch that easily formed a crystalline structure after HMT. As a 31 consequence, a significant increase in slowly digestible starch (23.65% to 36.96%) and resistant 32 33 starch (11.41% to 14.36%) and a decrease in rapidly digestible starch (47.70% to 40.30%) were 34 more noticeable in waxy flour than native flour. Microcapsules made from black waxy rice flour hydrolyzed for 36 h followed by HMT (hydrolyzed 36h-HMT) obtained the highest survival rate 35 of L. plantarum (89.56%), even after longer storage time (90 days, 4°C). In the stage of 36 37 simulated gastric fluid, the survival rate of L. plantarum in hydrolyzed 36h-HMT microcapsules was much higher (88.02%; 8.07 Log CFU/g) than gum arabic (75.38%; 6.12 Log CFU/g) and no 38 carrier (36.86%; 3.34 Log CFU/g). At the end of simulated intestinal fluid, hydrolyzed 36h-HMT 39 40 showed much higher survival (81.56%; 7.48 Log CFU/g) than gum arabic (58.1%; 4.72 Log CFU/g) and no carrier (0%). Under scanning electron microscopy, starch granules of the 41 hydrolyzed 36h-HMT were seen as polyhedral shapes in the spherical aggregates that carried the 42 microorganisms and reduced their injury and mortality. Short-chain fatty acids of the hydrolyzed 43 36h-HMT were much higher than positive control at every fermentation time (P < 0.01). The 44 fluorescence in situ hybridization data showed that the prebiotic property of hydrolyzed 36h-45 HMT can better aid the beneficial probiotic Lactobacillus spp. growth after 24 h fermentation 46 than the negative control (from Log 8.40 ± 0.50 to 7.03 ± 0.20 , P < 0.05) and commercial prebiotic 47 Orafti®Synergy1 (Log 8.40 \pm 0.50 to 7.47 \pm 0.08, P < 0.01). Microencapsulation of hydrolyzed 48 black waxy rice flour followed by HMT is proposed as a potential synbiotic ingredient to apply 49 50 in functional foods, further studies of these novel formulations are needed to determine in vivo 51 cell delivery performance and efficacy.

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53	Keywords: black wa	axy rice; pullulanase; heat mositure treatment; synbiotic; fluorescence in situ					
54	hybridization						
55							
56	Chemical compounds: 1,1-diphenyl-2-picrylhydrazyl (PubChem CID:74358); 2,2'-azino-bis-3-						
57	ethylbenzothiazoline	-6-sulfonic acid (PubChem CID: 16240279); Trolox (PubChem CID:					
58	40634)						
59							
60	Abbreviations						
61	Hydrolyzed-HMT	: Hydrolyzed heat-moisture-treated rice flour					
62	FISH	: Fluorescence in situ hybridization					
63	SCFAs	: Short-chain fatty acids					
64	SDS	: Slowly digestible starch					
65	RS	: Resistant starch					
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72 **1. Introduction**

Slowly digestible starch (SDS) and resistant starch (RS) are defined as the slow- to non-73 digestible portion of starch that cannot be digested after ingestion for 20-120 min and longer than 74 120 min, respectively. Indigestible carbohydrates, which can pass through the upper part of the 75 gastrointestinal tract into the large intestine, are subsequently fermented by gut microbes (Zhang 76 & Hamaker, 2009; Wang et al., 2002; Casterline et al., 1997). These prebiotic carbohydrates 77 should be able to provide an absorption site for probiotic adherence and, at the same time, a 78 79 carbon source for probiotic bacteria. These results in favorable metabolites such as short-chain fatty acids (SCFAs) in the human colon (Brouns et al., 2002; Crittenden al., 2001; Johnson & 80 Gee, 1996). SDS and RS are known to aid human health benefits, such as stable glucose 81 82 metabolism, reduced risk of diabetes, obesity, cardiovascular disease and colonic cancer (He, Liu, & Zhang, 2008; Lehmann & Robin, 2007). RS content of native colored rice from 83 indigenous Thai rice varieties is about 5.33-5.69% (Pongjanta, Chomsri, & Meechoui, 2016). 84 Factors affecting SDS and RS formation are crystallinity, chain length distribution, amylose and 85 amylopectin ratios, and retrogradation (Shi & Gao, 2011; Miao, Jiang, & Zhang, 2009). SDS and 86 RS can be produced by physical, chemical and enzymatic modifications. Physical modification is 87 88 considered safe for human consumption.

The dual modification proposed in this study was a combination of two safe processes:
heat-moisture treatment (HMT) and enzymatic modification. Significant increase in SDS (40.8
%) was found after HMT of native brown rice (Chung et al., 2012). Enzymes such as α amylase,
β amylase, amylo-sucrase, pullulanase and iso-amylase were often used to cleave the outer chain
of amylopectin, hence increasing starch chain mobility which led to a highly crystalline

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structure. Shorter amylose chains were usually required, accounting for its ability to be readily 94 re-associated into a more orderly crystalline structure (Shi & Gao, 2011). For SDS and RS 95 production, amylo-sucrase was reported to be able to hydrolyze starch chains in waxy/normal 96 rice and potato starch into DP 25-36 that aided crystalline perfection (Shin et al., 2010). 97 Pullulanase treatment followed by repeated retrogradation was reported to increase SDS and RS 98 content (25.4% and 50.1% respectively) in waxy maize starch (Miao, Jiang, & Zhang, 2009). 99 The most SDS was reported to be produced by debranching waxy starch with pullulanase for 4 h 100 and subsequent storage at 1°C. Waxy starch is more suitable to make SDS (Guraya, James, & 101 102 Champagne, 2001). Starch that was hydrolyzed and allowed to crystallize showed higher RS 103 than those only debranching (Cai & Shi, 2010). The co-process of hydrolysis followed by HMT was reported to increase the ratio of linear glucan α -D-(1, 4), which supported crystalline 104 105 structure and resulted in a more enzymatic-resistant starch (Mutungi et al., 2010; Lin et al., 2009). 106

Various polysaccharides that have been used as encapsulating materials include gum 107 arabic, inulin, oligosaccharides, maltodextrin and resistant starch (Perdana et al., 2014; Soukoulis 108 et al., 2014a; Fritzen-Freire et al, 2013; Fahimdanesh et al., 2012; Desmond et al., 2002). These 109 polysaccharides have different degrees of prebiotic effects, depending on its structure and 110 composition that favor probiotic growth. The carrier matrix of gum arabic and sodium caseinate 111 in low-melting-point fat microparticles was reported to enhance probiotic survival after spray 112 113 drying, storage and *in vitro* digestion (from 1.20 Log CFU/g to 2.55 Log CFU/g) (Liu et al., 114 2016). Prebiotic edible films made from native rice and corn starch, mixed with gelatine, sodium 115 caseinate and soy protein concentrate were used to encapsulate L. rhamnosus GG in a bread 116 coating. The viability of L. rhamnosus GG is increased by 3 to 7-fold under simulated gastro-

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intestinal conditions (Soukoulis et al., 2014b; Soukoulis et al., 2016). Glucose-oligosaccharides
and polydextrose were also reported to enhance *L. rhamnosus* GG viability in prebiotic edible
films during air drying. Inulin was the most effective material to maintain sub-lethal amounts of *L. rhamnosus* GG during storage (Soukoulis et al., 2014a).

The nutritious colored rice of interest in this study accounted for high antioxidant 121 activity. The anthocyanin content in black/purple, red and wild rice were 3276.0, 93.5 and 27.2 122 μ g/g, respectively (Abdel-Aal et al., 2006). The synergistic effect of malvidin-3-glucoside mixed 123 with other anthocyanins was reported to improve the growth of the good bacteria (Hidalgo et al., 124 125 2012). The colored rice flour was modified by the dual process of using HMT and enzymatic modification to obtain appropriate degrees of slow to indigestible prebiotic starch. Highly 126 indigestible starch aids probiotic survival but lowers its utilization as a carbon source. On the 127 128 contrary, rapidly digestible starch lowers probiotic protection in the gut system but serves as a good carbon source. Therefore, this study aimed to evaluate this trade-off effect using *in vitro* 129 human faecal batch fermentation that closely mimics the real human gut system. 130

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132 **2. Materials and Methods**

133 2.1. Preparation of hydrolyzed-HMT rice flour

Paddy rice of the black waxy "kam leum pua variety" (Phrae Rice Research Center in Phrae, Thailand) and the "red jasmine variety" (Khonkaen Rice Research Center in Khonkaen, Thailand) were dehulled to obtain brown rice grains. The brown rice was steeped in water for 3 h and then wet-milled using a double-disk stone mill to produce a 10% (w/v) rice flour slurry. The flour slurry was adjusted to pH 4.5 with a 0.1 M sodium acetate buffer. The enzyme pullulanase (OPTIMAX® L-1000, 1000 ASPU/g, Siam Victory Chemicals, Thailand) (0.2 g) was added into

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140 the flour suspension (110 g) to obtain a concentration of 20 ASPU/g of flour (dry basis). The solution was incubated at 55°C for 8, 24 and 36 h in a shaking water bath. The solution was 141 142 centrifuged (3000 g) for 10 min, the precipitate was washed twice with distilled water and 143 collected by centrifugation (Miao, Jiang, & Zhang, 2009). The precipitate was oven dried at 40°C until the target HMT moisture content (25%) was obtained. The rice flour sample was then 144 put in a sealed screw-cap container and equilibrated at room temperature for 24 h. The 145 equilibrated containers were then placed in a hot air oven (100°C) for 1 h. After that, the treated 146 flour was taken out and dried in a hot air oven (40 °C) until a 12% moisture content was 147 obtained. The obtained sample or "hydrolyzed-HMT" rice flour was milled (ultra-centrifugal 148 149 mill type ZM1, Retsch GmbH, Germany) and sieved to a particle size of 100 mesh, put in sealed plastic bag and kept at 4°C. 150

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152 2.2. Physico-chemical properties of hydrolyzed-HMT rice flour

153 2.2.1. Pasting properties

The pasting property of the hydrolyzed-HMT rice flour was determined by the Rapid Visco Analyzer (model RVA3D; Newport Scientific, NSW, Australia). The hydrolyzed-HMT rice flour (3.00±0.01 g) was mixed with distilled water (25 mL) in a metal RVA canister (AACC Method 61-02, 2000). The sample suspension was heated in the RVA using the heating profile for rice flour. The sample was heated from 50°C to 95°C at rate 12°C/min and held at 95°C for 2.5 min, cooled down to 50°C at a similar rate and held at 50°C for 2 min. The total running time for each sample was 13 min.

161 2.2.2. Thermal properties

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The thermal properties of all samples were determined by a differential scanning calorimeter (DSC Star[®] System; Mettler Toledo AG, Switzerland). Approximately 12 mg of flour was put directly into an aluminum pan using a flour to water ratio of 1:3 (w/w). The pan was sealed and equilibrated 1 h at room temperature before the analysis. The DSC scanning temperature range was set at 25-95°C using a heating rate of 10°C/min. The DSC was calibrated using indium as a standard and an empty aluminium pan as reference (Cham & Suwannaporn,

168 2010). The parameters were analyzed using STARe evaluation software v12.10 (Mettler Toledo169 AG, Switzerland).

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171 2.3. In vitro digestibility of hydrolyzed-HMT rice flour

172 2.3.1 Rapid digestible starch (RDS) and slow digestible starch (SDS) content

An enzyme solution was freshly prepared by adding porcine pancreatic α -amylase 173 (Sigma A-3176, Sigma-Aldrich, UK; 16 U/mg) (1.5 g) in a sodium acetate buffer (pH 5.2) (10 174 mL). The mixture was incubated at 37°C for 10 min and centrifuged (1500 g) for 10 min. The 175 supernatant was transferred into a beaker and mixed with amylo-glucosidase (Sigma A-7095, 176 177 Sigma-Aldrich, UK; 300 U/mL) at 8:1 (v/v) (Mutungi et al., 2011). The modified rice flour (100 mg) was suspended in a 0.1 M sodium acetate buffer (21 mL) and incubated at 37°C with 178 continuous shaking (200 strokes/min) for 15 min. The freshly prepared enzyme solution (1.6 179 mL) was added to the suspension, mixed for 1 min and incubated at 37 °C in a shaking water 180 bath (200 strokes/min). After incubation for 20 and 120 min, an aliquot (0.2 mL) was taken and 181 added into absolute ethanol (4 mL), mixed well and centrifuged (5000 g) for 10 min. The 182 supernatant was then collected for RDS and SDS determination. The glucose content was 183 184 measured by adding a glucose oxidase-peroxidase assay kit (GOPOD, Megazyme International,

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Ireland) into the aliquot and incubated at 50°C for 20 and 120 min. The aliquot was then measured in a spectrophotometer at 510 nm absorbance. Starch fractions that were digested (measured as % glucose) within 20 min and 20–120 min were calculated as RDS and SDS (Lin et al., 2009; Englyst et al., 1992).

189 2.3.2 Resistant starch (RS) content

190 A flour sample (100 mg) was weighed into a centrifugal tube and a KCl-HCl buffer (pH 191 1.5) (10 mL) was added. Pepsin solution (0.2 mL) was added into the mixture in order to remove 192 protein. The pepsin solution was prepared by adding pepsin (P-7125, Sigma-Aldrich, UK) (1 g) in the KCl-HCl buffer (10 mL). The solution was mixed well and incubated in a shaking water 193 194 bath (40°C) for 60 min and cooled down to room temperature. The solution was then added with a 0.1 M Trismaleate buffer (pH 6.9) (9 mL) and mixed with a pancreatic α -amylase solution 195 (1mL). The pancreatic α -amylase solution was prepared by adding pancreatic α -amylase (A-196 3176, Sigma-Aldrich, UK) (40 mg) into a tris-maleate buffer (1 mL). The solution was incubated 197 at 37°C for 16 h. The digested sample was centrifuged and the sediment was washed with 198 distilled water (10 mL) and centrifuged. After that, the precipitate was mixed with distilled water 199 200 (3 mL) and 4 M KOH (pH 4.75) (3 mL) for 30 min at room temperature. Then, 2 M HCl (5.5 mL) and 0.4 M sodium acetate buffer (pH 4.75) (3 mL) were added. Next, amyloglucosidase (A-201 7095, Sigma-Aldrich, UK) (80 μ L) was incubated (60°C) for 45 min and centrifuged, and the 202 203 supernatant was collected. The residue was washed with distilled water and centrifuged, and the supernatant was combined to make 100 mL. Sample solutions (0.1 mL) were pipetted into a test 204 tube and added with glucose oxidase-peroxidase kit reagent (GOPOD, Megazyme International, 205 206 Ireland) (3 mL), mixed well, and incubated (50° C) for 20 min. The solution was measured by a 207 spectrophotometer using absorbance 510 nm against a blank reagent. The resistant starch content 208 was calculated as mg of glucose x 0.9. The standard curve was determined using known glucose209 concentrations (Goni et al., 1996).

210 2.4. Antioxidative activity of hydrolyzed-HMT rice flour

A flour sample (1 g) was extracted using an extraction solvent (methanol:water, 80:20 v/v) (25 mL), mixed for 24 h and centrifuged (2500 g) for 10 min. The collected supernatant was analyzed for antioxidant activity using the following assays:

214 2.4.1. ABTS radical cation decolorization assay

For scavenging activity determination, a 2,2'-azino-bis-3-ethylbenzothiazoline-6-sulfonic
acid radical (ABTS) (Sigma-Aldrich, UK) solution was prepared by reacting a 7 mM aqueous
solution of ABTS with potassium persulfate in the dark at room temperature for 16 h before use.
The ABTS solution (1.5 mL) was added into the extracted sample (1.5 mL), mixed well and
measured at 734 nm absorbance using a spectrophotometer. The result was expressed as Trolox
equivalent antioxidant capacity (TEAC) in µM of Trolox/g flour (Jeng et al., 2012).

221 2.4.2. DPPH radical scavenging assay

The scavenging effect on the 1,1-diphenyl-2-picrylhydrazyl (DPPH) free radical was measured according to Brand-Williams et al. (1995). The working solution of 0.2 mM DPPH was prepared by dissolving DPPH (3.94 mg) in methanol (50 mL). The extracted sample (1.5 mL) was added to the DPPH working solution (1.5 mL). The mixture was left at room temperature for 30 min and was measured at 517 nm absorbance using a spectrophotometer. The scavenging effect was expressed as Trolox equivalent antioxidant capacity in μM of Trolox/g flour.

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230 2.5.

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2.5. In vitro batch culture fermentation

In vitro fermentation describes the method used by Costabile et al. (2015). Experiments 232 were carried out using fresh faecal samples from three healthy donors (1 female aged 26 and 2 233 234 males aged 25 and 31). Donors never had any metabolic or gastrointestinal diseases, did not take any probiotic/prebiotic supplements or antibiotics within the last 6 months. Each donor provided 235 written consent and filled in a standard questionnaire regarding their health status, drug use, 236 clinical anamnesis and lifestyles. This study was approved by the University of Reading 237 Research Ethics Committee (UREC 15/20). Each faecal sample was diluted to 1:10 (w/w) with 238 239 0.1 mol/L of phosphate buffer solution (PBS) at pH 7.4 and then homogenized (240 paddle beats/min) (Stomacher 400, Seward, West Sussex, UK) for 2 min. The solution was placed in an 240 anaerobic jar (AnaeroJarTM 2.5 L, Oxoid Ltd) equipped with a gas generating kit 241 242 (AnaeroGenTM, Oxoid). A flour sample (0.5 g) was added into an anaerobic fermenter within 15 min of voiding (Rycroft et al. 2001). The experiment was run in triplicate (3 vessels per donor). 243 Oligofructose-enriched inulin (Orafti[®]Synergy1, Belgium) (0.5 g) was used as a positive control 244 and a vessel without any sample was used as a negative control. The fermentation system was 245 run for 24 h. The sample solution was collected at 0, 4, 8, and 24 h to analyse for bacterial 246 247 populations and short-chain fatty acids.

248 2.5.2. Short-chain fatty acids (SCFAs) analysis

A cell-free supernatant was obtained by centrifuging (13,000g) a sample solution (obtained from section 2.5.1) (1 mL) for 10 min and filtering it through a 0.22 μm membrane (Millipore, Cork, Ireland) directly into a vial. The SCFA content was measured by an ion exclusion high performance liquid chromatograph (LaChrom, Merck Hitachi, Dorset, UK) equipped with a pump (L-7100, Merck Hitachi, UK), RI detector (L-7490, Merck Hitachi, UK)

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with a wavelength of 210 nm and an auto-sampler (L-7200, Merck Hitachi, UK). The sample 254 solution (20 µL) was taken and injected into the HPLC with a prepacked Rezex ROA-organic 255 acid H+ 80% (300 x 7.8 mm) column (Phenomenex Inc, UK), at a flow rate of 0.5 mL/min at 256 257 84°C. H₂SO₄ (2.5 mM) was used as eluent. The amount of SCFAs (acetic, propionic and butyric) was calculated using standard samples at concentrations of 12.5, 25.0, 50.0, 75.0 and 100.0 mM. 258 2-ethylbutyric acid (20 mM) was used as an internal standard. All chemicals were provided by 259 260 Sigma-Aldrich (Dorset, UK). 2.5.3 Enumeration of faecal bacteria population by fluorescence *in situ* hybridization (FISH) 261 262 FISH was analyzed following the method of Costabile et al. (2015). The oligonucleotide probe (Sigma-Aldrich, UK) was designed to target specific regions of 16S rRNA. The probe was 263 commercially synthesized and labeled with fluorescent dye Cy3 (Sigma-Aldrich, UK). The 264 probes used in this study are detailed in Table 1. The bacterial groups to be enumerated were (1) 265 Eub338 I-II-III for the total number of bacteria, (2) Bif164 for Bifidobacterium spp., (3) Lab158 266 for Lactobacillus-Enterococcus spp., (4) Bac303 for the Bacteroides-Prevotella group and (5) 267 Chis150 for the *Clostridium histolyticum* subgroup. 268 Table 1 269

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271 **2.6** Preparation of potential synbiotic microencapsulation by spray drying

The stock cell solution of *L. plantarum* TISTR 1465 (Thailand Institute of Scientific and Technological Research, Thailand) was freshly prepared. A pure lyophilized culture was thawed, suspended in 0.85% sterile saline (2 mL) and incubated at 37°C for 24 h. When the liquid was turbid, *L. plantarum* (1 mL) was taken, activated in MRS broth (9 mL) and incubated at 37°C for 24 h. The culture was collected by centrifugation (6000 g) for 10 min in the early stationary

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phase. The bacterial pellets were washed and centrifuged 3 times with sterile saline (10 mL). The
final precipitate was suspended in sterile saline (5 mL) (Zhao et al., 2008). The cell concentration
used was approximately 10¹⁰ CFU/mL.

Native and all hydrolyzed-HMT rice flours (hydrolyzed for 8, 24, 36 h then HMT) were 280 used as wall materials for L. plantarum encapsulation. Gum arabic and maltodextrin were used 281 as control carriers. The carrier solution was prepared by mixing wall material in water to obtain a 282 final concentration of 20% (w/w). The mixture was then homogenized using a two-stage high 283 pressure homogenizer (5000 psi) (15MR-8TA, APV Gaulin, Inc., MA, USA). Previously 284 prepared stock cell solution (approximately 10¹⁰ CFU/mL) (5 mL) was mixed with Tween 80 285 (0.2 mL), followed by the carrier solution (94.8 mL). The mixture was stirred at 20°C for 20 min 286 287 and immediately fed into a spray dryer (GEA Niro, A/S, DK-2860, Soeborg, Denmark). The inlet temperature (140 \pm 5°C) and outlet temperature (70 \pm 5°C) were obtained from a previous 288 experiment with the highest survival rate (data not shown). The spray-dried powder was 289 collected from the base of the cyclone, put in a sealed plastic bag and stored at 4°C for the next 290 291 study.

292

293 **2.7. Enumeration of** *L. plantarum* in the microcapsules

294 2.7.1. Viability of *L. plantarum* after spray drying and storage

The viability of *L. plantarum* in the microcapsules was determined immediately after spray drying and after storage at 4°C for 30 and 90 days. Spray-dried powder (1 g) was suspended in 0.1% peptone water (w/w) (9 mL) and homogenized by vortex for 10 min at room temperature to ensure complete dissolution of the powder. A sample solution (1 mL) was plated on MRS agar and incubated anaerobically at 37°C for 48 h (Yonekura et al., 2014).

300 2.7.2. Viability of *L. plantarum* under *in vitro* gastrointestinal condition

Simulated gastric fluid (SGF) was prepared by adding porcine pepsin (P-7125, Sigma-301 Aldrich, UK) (3 g) in salt solution (1 L). Salt solution was prepared by adding 125 mM NaCl 302 (7.305 g), 7 mM KCl (0.52 g) and 45 mM NaHCO₃ (3.78 g). Distilled water was added to make 303 a volume of 1 L and its pH adjusted to 2.5 with 0.1N HCl. Synbiotic powder (1 g) was added 304 into test tube that contained pre-warmed (37°C), freshly filtered, sterilized SGF (9 mL) and 305 mixed well. The aliquot was incubated in a water bath (37°C) for 90 min. Bacterial cells were 306 collected and washed with 0.85% NaCl by centrifugation (6000 g) for 10 min. After that, the cell 307 308 pellets were resuspended in simulated intestinal fluid (SIF). SIF was prepared by adding pancreatin (A-3176, Sigma-Aldrich, UK) (1 g) and bile extract (B-8631, Sigma-Aldrich, UK) 309 (1.5 g) in salt solution (1L) and its pH adjusted to 6.5 using 1 N NaOH. The aliquot was mixed 310 and incubated at 37°C. The digested aliquot (1 mL) was collected at 0, 90, 120 and 180 min and 311 added into normal saline (9 mL). The solution (1 mL) was then plated on MRS agar and 312 incubated at 37°C under anaerobic conditions for 48 h (Grimoud et al., 2010). 313

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315 **2.8. Scanning electron microscopy**

The morphology of the spray-dried microcapsules was determined by scanning electron microscopy (JM-560LV model, JEOL, Japan). Briefly, the encapsulated *L. plantarum* was fixed to the sample slide with 2.5% glutaraldehyde for 2 h, and washed with 0.1 M phosphate buffer (pH 7.2). The sample was fixed again with 2% tetroxide for 2 h, washed with deionized water and dehydrated by increasing the concentrations of ethanol solutions (50%, 70%, 80%, 90% and 100%). The dried powder was spread thinly onto a double-sided carbon adhesive disc and then anchored to the electron microscopy stub. The specimen was then coated with gold andexamined under scanning electron microscope.

324 **2.9. Data analysis**

Data was analyzed statistically using SPSS Version 16. The experimental data of hydrolyzed-HMT rice flour properties and enumeration of microencapsulated *L. plantarum* were analyzed by ANOVA and Duncan's multiple rank tests. The differences between the bacterial counts, substrates and SCFAs profiles at 0, 4, 8 and 24 h of fermentation were tested using ANOVA with Tukey's post-test (P < 0.05). All analysis were performed using a GraphPad Prism 5.0 (GraphPad Software, LaJolla, CA, USA).

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332 **3. Results and discussion**

333 **3.1. Pasting and thermal properties of hydrolyzed-HMT rice flour**

Black waxy and red jasmine rice flour (6.5 and 18.7% amylose content, respectively) 334 were modified using a debranching enzyme followed by heat-moisture treatment (HMT). A 335 decrease in pasting parameters was identified by the increase in starch crystallinity that limited 336 starch swelling (Figure 1a, 1b). Hydrolysis treatment is reported to promote an increase of low 337 molecular weight starch that easily formed a crystalline structure after HMT (Polesi & Sarmento, 338 2011). Higher paste viscosity increased noticeably in the hydrolyzed-HMT of red jasmine rice 339 flour (higher amylose content). Breakdown was observed to be substantially decreased, which 340 341 indicated more rigidity and resistance to shearing of the modified granules. The thermal 342 properties of gelatinization are used as thermal stability indices of starch granules. A higher gelatinization endotherm indicates more rigidity, more crystallinity and less swelling of starch 343 344 granules. All hydrolyzed-HMT rice flour had higher thermal properties than the native flour

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345	(p<0.05) (Figure 1c, 1d). Pullulanase, the debranching enzyme, attacked the 1, 6 glucosidic
346	linkages of amylopectin, generating a more linear chain (Reddy et al., 2015; Wong et al., 2007).
347	Later, HMT rearranged these linear chains into a more perfect crystal.
348	Figure 1
349	
350	3.2. In vitro digestibility of hydrolyzed-HMT rice flour
351	The enzyme digestibility of hydrolyzed-HMT rice flour was monitored by time needed to
352	convert starch to glucose in a simulated gastrointestinal environment. A significant increase in
353	SDS and RS and a decrease in RDS were noticeable in the hydrolyzed-HMT of black waxy rice
354	flour, while a much lower effect was found in the non-waxy type (Table 2). Waxy rice was more
355	susceptible to pullulanase, accounting for its higher α -1,6 branch point. Hydrolyzed 36 h-HMT
356	black waxy rice flour showed a desirable quality as it could produce high SDS and RS and low
357	RDS. SDS and RS that can pass through the stomach into the small and large intestines are
358	desirable, playing an important role in both prebiotic and protection wall materials in this study.
359	Table 2
360	
361	In this study, pullulanase was used to hydrolyze ungelatinized swollen starch granules
362	without liquefaction. Hydrolysis occurred mostly on a solid surface (Jung et al., 2013) that is not
363	strong enough to induce the formation of RS. RS required linear polymers of a minimum chain
364	length of approximately 10 DP to form double helices (Mutungi et al., 2010; Miao, Jiang, &
365	Zhang, 2009). Chain mobility improved during hydrothermal process that enabled crystalline
366	rearrangement, hence increasing indigestible starch formation. Limited enzyme reaction caused

367 SDS to increase and then become stagnant at 24-36 h because no liquefaction occurred. The

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368	result agreed well with Wong et al. (2007) in that linear long-chain dextrin did not increase much
369	in non-gelatinized sago starch treated with pullulanse for 24 h as compared to gelatinized starch.
370	
371	3.3. Antioxidant properties
372	The DPPH and ABTS scavenging activity of all samples decreased drastically after
373	modification (Figure 2). Soaking during hydrolysis incubation and spray drying deteriorated
374	some water-soluble pigments of colored rice flour. Modified black waxy rice flour drastically
375	decreased antioxidant activity in comparison with red jasmine rice flour. However, a higher level
376	of antioxidant activity in black waxy rice flour was still retained, accounting for its high initial
377	content in both free and bound forms. Protocatechuic acid, a phenolic compound in bound form,
378	was only found in black rice, whereas ferulic, p-coumaric and vanillic acid were found in both
379	red and black rice varieties (Sompong et al., 2011).
380	Figure 2
381	
382	3.4. Viability of <i>L. plantarum</i> after spray drying and storage
383	After spray drying, the viability of L. plantarum in all spray-dried samples was not much
384	different (9.36-9.71 log CFU/g) (Table 3). Yonekura et al. (2014) was also reported non-
385	significantly different in L. acidophilus survival after spray drying with sodium alginate,
386	hydroxypropylmethyl cellulose and chitosan (8.90-8.99 log CFU/g). However, a noticeable
387	difference in survival rates was observed in longer storage times (90 days). Maltodextrin showed

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the lowest survival rate (44.68%) while hydrolyzed 36h-HMT showed the highest survival rate

389 (89.56%). The modified black waxy rice flour protected *L. plantarum* better than red jasmine.

390 Protection often correlated with high glass transition temperature of wall materials which provide

stability for bacteria enclosed within a glassy matrix (Crowe et al., 1998; Leslie et al., 1995). The increase in glass transition temperatures can provide stabilization via the fixation of bacterial cells in a glassy state during spray drying (Perdana et al., 2014). Encapsulated microcapsules from hydrolyzed-HMT rice flour showed good protection properties supported by high bacterial survival even after a long storage time (90 days, 4°C). Hydrolyzed 36h-HMT of black waxy rice flour was selected to use as wall material throughout later experiments.

397

Table 3

398

399 **3.5.** Survival of *L. plantarum* in *in vitro* gastrointestinal environment

The sharp reduction in the viable cell count was observed in the stage of SGF (90 min). 400 At this stage, the survival rate of L. plantarum in encapsulated powder with hydrolyzed 36h-401 402 HMT, gum arabic and no carrier (free cell) were 88.02% (8.07 Log CFU/g), 75.38% (6.12 Log CFU/g) and 36.86% (3.34 Log CFU/g), respectively (Figure 3). At the end of SIF (180 min), the 403 hydrolyzed 36h-HMT showed much higher survival and was stabilized throughout the digestion 404 process. The survival rate of hydrolyzed 36h-HMT, gum arabic and no carrier (free cell) were 405 81.56% (7.48 Log CFU/g), 58.1% (4.72 Log CFU/g) and 0%, respectively. Without any carrier, 406 no viable cells of L. plantarum were found, as they are sensitive to the acid-bile condition. Gum 407 arabic, a frequently used commercial carrier, is a soluble fiber consisting mostly of the carboxyl 408 group. Spray-dried particles with gum arabic dissolved more easily in SGF and SIF solutions, 409 410 hence lowering its prebiotic property and probiotic protection. Xing et al. (2015) also suggested a 411 complex wall material that consisted of porous starch (10%), mannitol (3%) and glycerol (2%) to protect L. acidophilus against the intestinal system and refrigerated storage. The microparticles 412 413 made from hydrolyzed 36h-HMT showed SDS and RS properties as some particles remained

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after SGF and SIF conditions. These particles protect probiotic cells survival and help get thesecells through the lower part of the gastrointestinal tract to produce SCFAs.

416

Figure 3

417 3.6. Scanning electron microscopy (SEM) of microencapsulated aggregates

Starch granules of the hydrolyzed 36h-HMT rice flour remained ungelatinized, which 418 were observed as polyhedral shapes formed in spherical aggregates. The spherical aggregates 419 were produced when starch granules of small size were rapidly dehydrated by spray drying with 420 low amounts of bonding agents (Zhao & Whistler, 1994) (Figure 4). The protein in the flour was 421 422 reported to promote the formation of these aggregates (Avila-Reyes et al., 2014). L. plantarum was scattered within the interstitial spaces and at the periphery of the aggregates (Figure 4). 423 These aggregates carried the microorganisms and reduced their injury and mortality. The 424 425 aggregates of microcapsules from hydrolyzed 36 h-HMT were similar to those encapsulated with various carbohydrates or colloids, such as inulin, gum arabic, maltodextrin and native starch 426 (Avila-Reves et al., 2014; Rodríguez-Huezo et al., 2007; Desmond et al., 2002) 427

428

Figure 4

429

430 **3.7. Short chain fatty acids (SCFAs)**

SCFAs were generated by the metabolism of faecal bacteria using wall materials as substrates. SCFAs were analyzed in comparison to a commercial prebiotic as a positive control (Table 4). Indigestible carbohydrates were consumed and converted into SCFAs by intestinal microbes (Casterline et al., 1997). SCFAs have an important role in maintaining gut health and host energy metabolism (den Besten et al., 2013; Dongowski et al., 2005). SCFAs of the hydrolyzed 36 h-HMT and the positive control were much higher than the negative control at

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437 every fermentation time (p<0.01). Hydrolyzed 36 h-HMT was able to ferment at a higher rate and produced higher amounts of SCFAs than the positive control. High amounts of acetate were 438 detected and rapidly increased after 4 h of fermentation. Acetate was reported as a main 439 440 metabolized product of starch and arabino-oligosaccharides by faecal bacteria (Vigsnaes et al., 2011). Acetic acid was the main metabolite produced by bifidobacteria in prebiotic stimulation 441 with oligofructose (Van der Meulen et al., 2006). Propionate and butyrate showed gradual 442 increases during faecal fermentation. Butvrate was formed by free acetate as a precursor of the 443 conversion by butyryl CoA from colon bacteria (Scott, Duncan, & Flint, 2008). 444

445

Table 4

446

447 **3.8. Modulation of bacterial populations by FISH**

In comparison with the Orafti[®]Synergy1 (a commercial prebiotic, inulin based), fermentaion of hydrolyzed 36 h-HMT sample caused significantly higher numbers of *Lactobacilli* (enumerated by Lab158 probe) at 8 h (from Log 7.91±0.02 to 7.50±0.05, P < 0.05) and 24 h (from Log 8.40±0.50 to 7.47±0.08, P < 0.01) (Figure 5). However, there was no significant difference between hydrolyzed-HMT and Orafti[®]Synergy1 in the other bacteria enumerations.

In comparison with the negative control, the fermentation of hydrolyzed 36h-HMT showed significantly higher numbers of lactobacilli (Lab158), *Bifidobacterium* ssp. (Bif164) and total bacteria (Eub338 I-II-III) (Figure 5). Numbers of lactobacilli in hydrolyzed 36 h-HMT fermentation was higher than the negative control for every fermentation time; 4 h (from Log 7.85±0.07 to 7.30±0.01, P < 0.05), 8 h (from Log 7.91±0.02 to 7.03±0.11, P < 0.01) and 24 h (from Log 8.40±0.50 to 7.03±0.20, P < 0.05). A significant increase in the Bif164 group was

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also found in hydrolyzed 36 h-HMT in comparison to the negative control after being fermented for 8 h (from Log 9.14±0.51 to 8.58±0.20, P < 0.05) and 24 h (from Log 9.50±0.11 to 8.10±0.20, P < 0.01). The total bacteria enumerated with Eub338 I-II-III showed significant increases at 8 h (from Log 9.43±0.40 to 8.63±0.21, P < 0.01) and 24 h (from Log 9.60±0.28 to 8.05±0.06, P <0.001). Only the numbers of the Chis150 group decreased at 24 h (from Log 7.22±0.20 to 6.65±0.31, P < 0.01). No significant changes in *Bacteroides-Prevotella* numbers (Bac303) were found for any fermentation time point or substrate.

The results indicate the prebiotic properties of hydrolyzed 36h-HMT that aid the growth 467 468 of beneficial probiotics (mainly *Lactobacillus* spp. and *Bifidobacterium* spp.) after fermentation, as quickly as 8 h earlier than the negative control. Moreover, a better or comparable prebiotic 469 quality of hydrolyzed 36 h-HMT was also found in comparison with commercial prebiotic of 470 471 inulin base. This result corresponds well with the SCFAs production (Table 4). The polysaccharide from wheat dextrin was also reported to enhance numbers of Lactobacillus in in 472 vitro batch fermentation (Noack et al., 2013). A significant increase of *Lactobacillus* in the mice 473 faeces was also found after applying amylomaize starch in the rat diet (Wang et al., 2002). 474

475

Figure 5

476

477 **4. Conclusion**

Encapsulated formulation with slow digestible colored rice flour could be used as a cheaper alternative functional food ingredient. Hydrolyzed-HMT of colored rice flour obtained both potential prebiotic and synbiotic properties due the antioxidative property. Moreover, its activity was stable throughout the 90-day studied shelf life further studies of these novel formulations are needed to determine in vivo cell delivery performance and efficacy.

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488	
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