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Title: Effects of Ball Milling on the Structural, Thermal, and Rheological Properties of Oat Bran Protein Flour

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Keywords: oat protein; FTIR spectroscopy; DSC; amylose-lipid complex; dispersion.

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Abstract: Oat bran protein flour (OBPF), containing protein, starch, and lipid as major constituents, was ball milled and subsequently evaluated on structural conformation, thermal properties, particle size distributions, and rheological properties. Prior to ball milling, characterisation of OBPF were conducted by means of Fourier transform infrared (FTIR) spectroscopy and differential scanning calorimetry (DSC) showing the existence of aggregated protein and starch-lipid complexes as predominant constituents of OBPF. Ball milling altered structural conformations of both protein and starch. Moreover, increase of ball milling time gradually decreased the transition enthalpy changes of amylose-lipid complexes upon heating which can be related to disruption of amylose-lipid complexes helical structure. Ball milling at higher speed resulted to smaller average particle size distributions of OBPF. Dynamic mechanical spectra of concentrated dispersions containing ball milled OBPF exhibited lower storage (G') and loss (G") moduli compared to control sample due to reduced particles volume packing. Moduli-frequency sweep data satisfactory fitted the Power Law's model.

22nd of October 2017

Journal of Food Engineering

Resubmission of manuscript JFOODENG-D-17-01095R1

Dear Editor,

Thank you for the opportunity to revise our manuscript. We appreciate the careful review and constructive suggestions. The manuscript has been rewritten base on the suggestions, with a new title "Effects of Ball Milling on the Structural, Thermal, and Rheological Properties of Oat Bran Protein Flour". Following this letter are our responses to the reviewers' comments, including how and where the text was modified.

Thank you for your considerations.

Most sincerely,

Kurnia Ramadhan

Detailed Response to Reviewers

Reviewer #1

Comments	Responses				
The manuscript reports a study on structural	The manuscript has been rewritten and				
issues of an oats bran protein concentrate	added with more quantitative data presented				
using spectroscopy, calorimetry and	in tables.				
rheology. Standard procedures were used,					
but for a publication in Journal of Food					
Engineering, I was expecting a better					
quantitative analysis than presented. For					
example, the rheograms (moduli-frequency)					
could be modelled to obtain parameters than					
can be statistically compared. The					
manuscript is weak in statistics, even though					
the effects of milling and fractionation or					
defatting of the concentrate are discussed. A					
properly revised manuscript would meet the					
quality required for the journal.					
Lines 223-225: Consider including the	Revised as suggested. These lines have been				
statistics (+- standard deviation, SD or	rewritten and can be found in revised				
standard error of mean, SEM) of the particle	manuscript starting from the Line 255.				
sizes.					
Lines 243-252: It would have been better to	Revised as suggested. These lines have been				
model the moduli-frequency relationships to	rewritten and can be found in revised				
obtain quantitative parameters for a deeper	manuscript starting from the Line 284.				

analysis of the results, trends and effects of	
milling. A sentence like "However, the	
slope of G' and G" versus frequency curves	
seems to remain unchanged of OBPC	
dispersion", is too qualitative and eye-	
balling for the journal.	
Figure 3: Is it not possible to define some	Revised as suggested.
quantitative parameters from the spectra to	
characterize the samples? I was thinking	
absorbance units at those specified	
wavenumbers for protein structures could be	
used. Absorbance units at certain	
wavenumbers are used in FTIR studies on	
starches to properly characterize amorphous	
and crystalline structures in them.	
Figure 7: Consider modelling the rheograms	Frequency sweep data have been fitted to
for quantifiable characterization parameters	the Power Law's model and presented in
to deeply discuss the rheological behaviours	table. The rheogram remain unchanged.
of the samples.	
Table 1: Better to include the statistics to	The words "significantly" have been
justify using 'significant', 'significantly	removed. Fitting parameters of frequency
different' or 'not significant' in the text (lines	sweep tests have been added.
205, 233 and 247, for example). This table	
could also contain the characterization	
parameters for the rheological behaviours of	

the samples.	

Reviewer #2

Comments	Responses
Line 15: Instead of "Fourier Transform	Revised as suggested. This line has been
Infrared", "Fourier transform infrared" is	rewritten and can be found in revised
more appropriate.	manuscript starting from the Line 14.
Lines 25-26, 44-45, 58-63, 82-83, 185-187,	Revised as suggested. These lines have been
192-193, 206-207, 215-216: These	rewritten.
sentences should be revised.	
In the manuscript, 35 references were used.	References have been updated.
From these references, only 9 references	
were published within the last 8 years (since	
2010). This means that the authors have not	
mentioned from recent studies sufficiently.	
Lines 30-46: "Introduction" section should	Revised as suggested. These lines have been
be re-written. Literature studies given in this	rewritten and can be found in revised
section is not sufficient.	manuscript starting from the Line 27.
Line 63: Please use small letters in the	Revised as suggested. This line has been
spelling of "Propanol".	rewritten and can be found in revised
	manuscript starting from the Line 65.
Line 111: The abbreviation of second is not	Revised as suggested. This line has been
"sec".	rewritten and can be found in revised
	manuscript starting from the Line 128.

Line 138: Please use subscript for "2" in "-	Revised as suggested. This line has been
СН2-".	rewritten and can be found in revised
	manuscript starting from the Line 158
Lines 254-259: Important numerical results	Revised as suggested. These lines have been
should be indicated in "Conclusions"	rewritten and can be found in revised
section.	manuscript starting from the Line 296.

Reviewer #3

Comments	Responses
This manuscript deals with the influence of	The manuscript has been rewritten and
chemical features and ball milling on the	added with more quantitative data presented
structuring characteristics of oat bran	in tables.
protein. For this purpose, several	
experimental tests were performed. The	
research topic falls within the scope of the	
Journal of Food Engineering and could be	
of the scientific interest. Nevertheless, the	
content of the manuscript evidence some	
lacks that must be improved. Please	
consider the following comments:	
Keywords: Personal consideration: Words	Revised as suggested.
included in the title should be removed from	
keywords section.	
Introduction: This section should be	Revised as suggested.
extended to include updated references in	

the field (last 5 years).	
Materials and methods: Replicates should	Revised as suggested.
be included in all sections.	
Further detail why experimental conditions	Revised as suggested.
were selected or some reference should be	
included in some methods section (see as	
e.g. ball milling procedure, preparation of	
OBPC dispersions, among others).	
Figure 6. Log representation is commonly	Revised as suggested.
used for rheological measurements.	
References should be updated: Only 3	References have been updated.
references from last 5 years.	
Tbale 1. Standard deviations should be	Revised as suggested.
included.	

Highlights

- Ball milling alters structural conformations of aggregated proteins and starch-lipid complexes
- Mechanical impact caused by ball milling disrupted the ordered structure of amyloselipid complexes
- Changes in mechanical moduli of dispersions are mainly contributed by differences in particle size distributions after ball milling

1	Effects of Ball Milling on the Structural, Thermal, and Rheological Properties of Oat
2	Bran Protein Flour
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Abstract

Oat bran protein flour (OBPF), containing protein, starch, and lipid as major constituents, 11 12 was ball milled and subsequently evaluated on structural conformation, thermal properties, 13 particle size distributions, and rheological properties. Prior to ball milling, characterisation of 14 OBPF were conducted by means of Fourier transform infrared (FTIR) spectroscopy and 15 differential scanning calorimetry (DSC) showing the existence of aggregated protein and 16 starch-lipid complexes as predominant constituents of OBPF. Ball milling altered structural 17 conformations of both protein and starch. Moreover, increase of ball milling time gradually decreased the transition enthalpy changes of amylose-lipid complexes upon heating which 18 19 can be related to disruption of amylose-lipid complexes helical structure. Ball milling at 20 higher speed resulted to smaller average particle size distributions of OBPF. Dynamic 21 mechanical spectra of concentrated dispersions containing ball milled OBPF exhibited lower 22 storage (G') and loss (G") moduli compared to control sample due to reduced particles volume packing. Moduli-frequency sweep data satisfactory fitted the Power Law's model. 23

24

25 Keywords: oat protein, FTIR spectroscopy, DSC, amylose-lipid complex, dispersion.

26

1. Introduction

Interests in protein-rich foods are rising favoured by its health-promoting effects. The benefits of dietary protein intake has been associated with body weight management and maintenance of muscle mass and function (Deutz et al., 2014; Westerterp-Plantenga et al., 2017). The urge to search for alternative food protein sources have led to valorisation of underexploited plants and side stream agricultural products.

32 Oat bran is a by-product of oat flour milling and have been used as a fibre-enriching 33 ingredient. Although oat bran contains abundance of protein, its full utilisation remains 34 challenging. Enzymatic pre-treatment of oat bran thick cell walls is necessary to assist 35 protein extraction, otherwise a highly alkaline pH is required to release protein that 36 adversely affect the nutritional quality (Jodayree et al., 2012). Low solubility of oat 37 protein in slightly acidic to neutral pH range also restrict its applicability for foods 38 (Konak et al., 2014). In order to bridge the gap between the limitations and advantages of oat bran protein for utilisation in foods, a feasible separation technique has been 39 40 introduced to obtain protein-enriched flour with low degree of purity (Sozer et al., 2017). The presence of protein, starch and lipid, as constituents of composite flour, are known 41 42 contributing to the physicochemical properties and functionality of flour (Puncha-Arnon 43 and Uttapap, 2013; Saleh, 2017).

Physical modification of food materials by means of ball milling has gained attention due to ability in changing functionality. The modified functionality can be attributed to alteration in structural conformation as found in ball milled starch and starch-enriched food materials (Dhital et al., 2011; Liu et al., 2011; Roa et al., 2014a). However, there are limited studies reporting the effects of ball milling on structural conformation changes in food proteins including whey and soy protein (Liu et al., 2017; Sun et al., 2015). 50 This present study aims to evaluate the alteration on structural conformation of 51 chemical constituents within ball milled oat bran protein flour, and subsequently to 52 extrapolate the conformational changes into the thermal and rheological properties of its 53 dispersion. This study would provide useful insights to extend the utilisation of oat bran 54 protein flour in food formulations.

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2. Materials and methods

57 2.1. Materials

58 Oat bran protein flour (OBPF) was supplied by Tate & Lyle Oat Ingredients 59 (Kimstad, Sweden) as a brand of PrOatein®. This material was in the form of coarse 60 powder and declared to have ca. 94% dry matter containing 54% protein, 17% fat, 18% 61 carbohydrate, and 2% fibre on a dry basis. This material was prepared from oat bran that 62 have gone through enzymatic, thermal, and physical separation processes.

63

64 2.2. Elimination of lipid and starch

Elimination of lipid and reduction of starch were intended to provide a correct 65 66 assignment of FTIR spectra bands and interpretation of thermal properties in the later 67 analyses. Lipid were extracted using solvent in soxhlet apparatus as described by Pandey 68 and Shrivastava (Pandey and Shrivastava, 2017) with modifications. Two solvents with 69 different polarity were used, i.e. hexane for the non-polar solvent, and isopropanol for the 70 polar solvent. Defatted samples are addressed as NPDO for the hexane-defatted OBPF 71 and PDO for the isopropanol-defatted OBPF. Protein extraction from defatted samples, 72 i.e. NPDO and PDO, did not lead to reasonable protein yields for further analysis. Therefore, in this study, starch reduction was chosen to obtain a higher protein 73 concentration. Starch reduction was conducted by dispersing and stirring the defatted 74

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samples into alkaline solution for 30 min as described by De Souza et al. (De Souza et al.,

76 2016). The sample was then lyophilised to obtain dry powder and addressed as APDO.

77

78 2.3. Ball milling procedure

Ball milling was carried out by using Planetary Micro Mill (Pulverisette 7 classic line,
FritschGmbH, Germany) as described by Abbaszadeh et. al. (2014) with modifications.
Two and a half grams of OBPF was put into 12 ml grinding bowl containing 6 Zirconium
dioxide grinding balls. High speed milling was run at 800 revolutions per minute (rpm) for
15 min. The low speed milling was carried out at 200 rpm for 240 min. The applied
milling cycle configuration was 5 min milling followed by 5 min rest to dissipate heat.
Reversed milling directions were set to prevent ball slippage at both milling speeds.

86

87 2.4. Preparation of OBPF dispersions

OBPF dispersions (5%, 20%, and 30% wt) were prepared by suspending the flour into distilled water as described by Roa et al. (Roa et al., 2015) with modifications. These concentrations were selected according to previous trials conducted with OBPF dispersions from 5% to 30% wt. A dilute system (5% wt) was selected to evaluate the particle size distributions, while the 20% wt dispersion was subjected to thermal analysis for its flow-ability reason, and the 30% wt dispersion was selected for rheological analysis to avoid sedimentation.

95

96 2.5. Attenuated total reflectance – Fourier transform infrared (ATR-FTIR) spectroscopy

97 The infrared spectra were obtained by using a Bruker Tensor 27 System (Bruker 98 Optik GmbH, Ettlingen, Germany) equipped with diamond Attenuated Total Reflection 99 (ATR) crystal and operated using OPUS software version 7.2.139.1294. The tests were 100 performed at ambient temperature. The open-air background test was run prior to sample 101 measurement. Approximately 50 mg of dry sample was placed into contact with the 102 diamond ATR crystal and sample holder. Each absorbance measurement was conducted under 128 scans with 4 cm^{-1} resolutions. The presented infrared spectra were obtained as 103 104 a mean from three measurements of duplicate samples that have undergone normalisation 105 and baseline correction. Height of peaks were determined by calculating the absorption 106 intensity differences between peak and its own baseline. Peak height ratio of selected 107 absorption bands was used to quantify the infrared spectral changes. The amide I band, i.e. $\sim 1630 \text{ cm}^{-1}$, was selected as reference band in evaluating spectral changes due to 108 109 elimination of lipids and starch. Calculation of absorbance intensity ratio ~997 to ~1022 cm⁻¹ was used to assess structural conformation changes on starch as described by Liu et 110 al. (Liu et al., 2011). The use of second derivative spectra was intended to resolve the 111 overlapping bands both in amide and saccharide regions (Figure 3). The assignment of 112 113 protein secondary structure within amide I band referred to a study reported by Tang and 114 Ma (2009). Further processing of infrared spectra was conducted using Spectragryph 115 optical spectroscopy software version 1.0.2 (https://www.effemm2.de/spectragryph/).

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- 2.6. Micro-differential scanning calorimetry

Scanning calorimetry was performed in a Micro-Differential Scanning Calorimeter III (Setaram Instrumentation, France). About 800 mg of dispersions containing 20% wt solid sample was transferred into hermetically sealed sample vial made from hastelloy, while similar weight of deionized water was used as reference. Heating-cooling processes were performed in two cycles from 20 to 120 °C at a rate of 1 °C min⁻¹. Measurement was run in duplicate for each sample. The heat flow curves were processed using Calisto Processing 124

software v1.43 (AKTS, Switzerland) to obtain the maximum heat absorption temperatures (T_m) and the transition enthalpy changes (Δ H).

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127 2.7. Particle size distributions

128 Particle size and size distributions were measured based on the principles of light 129 scattering using LS13320 Laser Diffraction Particle Size Analyzer (Beckman Coulter, 130 High Wycombe, UK). A dispersion of 5% OBPF in distilled water was pipetted into a 131 liquid module sample cell. The measurement was run for 60 s under following conditions: 132 Fraunhofer theory was used as optical model treating particles in spherical approximation, 133 refractive indexes of dispersant and sample were 1.33 and 1.6 respectively. The particle 134 size was determined by the volume-weighted mean diameter size. Each measurement was 135 run in duplicate.

136

137 2.8. Rheology

138 Structuring properties of OBPF dispersion was evaluated using rheological analysis. 139 Small deformation tests of oat dispersion were conducted using a stress-controlled 140 rheometer (Physica MCR 301, Anton Paar, Austria) with a serrated parallel-plate 141 geometry of 50 mm diameter and 1 mm gap. The linear viscoelastic region (LVR) was determined by strain amplitude sweep tests at a fixed angular frequency of 10 rad s^{-1} . The 142 temperature sweep tests were performed at a constant angular frequency of 10 rad s^{-1} and 143 144 a strain amplitude of 0.1%. A programmed heating-cooling process was used, i.e. heating from 20 to 100 °C, temperature hold at 100 °C for 10 min, and cooling from 100 to 20 °C 145 with heating and cooling rate of 5 °C min⁻¹. Prevention of evaporation and temperature 146 147 gradient were taken into account by using low-viscosity mineral oil and a peltier-148 controlled hood. Subsequently, frequency sweep tests were carried out from 0.5 to 50 rad

 s^{-1} at constant strain amplitude of 0.1%. Frequency sweep data were correlated by means 149 150 of the Power Law model using the following equations:

151
$$G' = a'\omega^{b'} \tag{1}$$

$$152 \qquad \qquad G'' = a'' \omega^{b''} \tag{2}$$

153 where ω , G', G", and are angular frequency, storage, and loss modulus respectively, and a', a", b', and b" intercepts and slope of corresponding fitting parameters. Each test was 154 155 run in duplicate.

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- 157

3. Results and discussion

158 3.1. Assignment of OBPF chemical constituent in infrared spectra

159 Infrared spectra of OBPF are illustrated in Figure 1 showing chemical-bond 160 vibrations of mixture system. Protein as the primary constituent can be traced by the 161 presence of amide A, I, and II absorption bands. Amide A vibrations occurred at higher 162 wavenumber and is contributed by N-H groups stretching vibrations. Amide I and II are predominantly induced by C=O and C-N stretching vibrations, and appear as two 163 consecutive absorption signals between 1700 and 1500 cm⁻¹ (Barth, 2007). The Amide A, 164 I, and II absorption bands of OBPF appear at wavenumbers 3271, 1630, and 1517 cm⁻¹ 165 166 respectively. Evaluation of amide I region can provide useful information on protein 167 secondary structure conformation which will be discussed in the later section.

168 Infrared spectral changes as affected by lipids and starch eliminations are presented in Table 1. Removal of neutral lipids by non-polar solvent drastically decreased the 169 absorption signal at ~2924, ~2854, and ~1744 cm⁻¹. However, the use of polar solvent 170 diminished the second peaks and left small signal of the third peak. This might indicate 171 the asymmetric stretching vibrations of C-H at ~ 2924 cm⁻¹ is not solely contributed by of 172 lipid alone, but also from other major chemical constituents. The peak at ~2854 cm⁻¹ 173

174 could be uniquely assigned to $-CH_2$ - symmetric stretching of fatty acid chains. The peak 175 ~1744 cm⁻¹ corresponds to carbonyl (C=O) stretching vibration identifying the ester 176 functional group of lipid (Cerqueira et al., 2012). Oats contain higher amounts of polar 177 lipids compared to other cereals, such as glycolipids and monoacyl lipids (Doehlert et al., 178 2010).

Saccharide region of infrared spectra ranges from 1200 to 900 cm⁻¹. Three intense 179 absorption bands at 1150, 1078, and 1022 cm⁻¹ confirming typical vibration 180 181 characteristics of C-O, C-C, O-H bonds and asymmetric stretching of C-O-C glycosidic 182 bonds within the starch (Liu et al., 2011; Roa et al., 2014b). The strongest absorption band at 1022 cm⁻¹ indicated predominant disordered conformational structure of OBPF 183 184 starch that have gone through thermal treatment during production. This peak contains 185 shoulder at the lower wavenumber that possibly indicate small amount of crystalline conformation. Alkaline exposure treatment dramatically decreased absorbance intensity 186 of the three above mentioned peaks, indicating irreversible solubilisation of starch. This 187 188 confirms the correct assignment for starch within infrared spectra and can be used to 189 evaluate structural conformations in the later section.

190

191 3.2. Thermal properties of OBPF dispersions

192 The DSC thermograms and the corresponding thermal transition peak temperature and 193 enthalpy changes of OBPF dispersions are presented in Figure 2 and Table 2 respectively. 194 Upon the first heating, a broad endothermic peak appeared at c.a. 90 °C and consequently 195 shifted to a higher temperature when reheated. Whereas upon cooling, a sharp exothermic 196 peak was observed at c.a. 74 °C in either cycles. The Δ H values was 1.7 J g⁻¹ on the first 197 heating and subsequently decrease over the following cycles. The use of polar solvent 198 greatly reduced Δ H values due to elimination of higher amounts of lipids as confirmed by 199 FTIR study. However, calorimetry scanning of extracted lipids did not show any thermal 200 transition events (data not shown). Starch solubilisation by alkaline exposure further 201 decreased the Δ H values. It shows that the Δ H values are more dependent to the starch 202 and lipids rather than the protein.

203 The reversible thermal events at very high temperature upon heating, i.e. 90 °C, can 204 be associated to the characteristics of starch-lipid complexes in the presence of protein contained in thermally treated foods as found in previous studies (De Pilli et al., 2015; 205 206 Moisio et al., 2015; Wang et al., 2017). When the heated starch is cooled back in the 207 presence of lipid, the amylose recrystallize and form complexes with lipids by the 208 inclusion of fatty acid chain into helical cavity (Tufvesson et al., 2003). The amount of 209 amylose-lipid complexes is the major factor determining the ΔH value as proposed by 210 Kawai et al. (2012). Interestingly, melting temperature of starch-lipid complexes slightly 211 increased upon second heating hence indicating a more thermally-stable structure as a 212 result of complexes reformation in more ordered structure.

213

3.3.Ball milling effects on structural conformation of OBPF chemical constituents

215 Ball milling affected structural conformation changes on both protein and starch in 216 solid-state as illustrated in Figure 3 and Table 3. Prior to ball milling, secondary structure 217 of protein prominently presented as β -sheets structure and some minor helical structures. 218 This result demonstrated a higher composition in β -sheets compared to previous study on 219 oat protein isolate secondary structure at low moisture (Liu et al., 2009). The explanation 220 for this condition is the protein contained in OBPF has been thermally treated during the 221 production, hence most likely in denatured and aggregated form. The β -sheet structures 222 have been associated with hydrophobic and aggregated protein (Maltais et al., 2008; Nikolaus Wellner et al., 2004; Tang and Ma, 2009). However, other study on oat protein 223

224 conformation in aqueous dispersion showed lower relative amount of β -sheets as the major 225 structure (Jing et al., 2016). Secondary structure of protein in dispersion is a result from 226 peptide-solvent interactions in aqueous environment, whereas at low moisture the peptide-227 peptide interactions are predominant factor due to dense packing in a solid environment.

228 Ball milling gradually increased relative amount of helical conformations in expense 229 of β -sheet structures. Longer ball milling time, regardless the applied speed, demonstrated 230 greater effect on protein conformational changes due to simultaneous mechanical impact. 231 Similar result of increasing α -helix and decreasing β -sheet structure after ball milling sov 232 protein isolate has been reported (Liu et al., 2017). Ball milling involves high pressure, 233 shear force, and air contact that might cause rearrangement of aggregated protein 234 secondary structure as suggested by Sun et al. (2015). Among the protein fractions and 235 sub-units, low molecular weight proteins are the most susceptible to disruptions caused by 236 ball milling (Thanatuksorn et al., 2009).

Evaluation of ball milling effects on starch conformation are presented in Table 3. A higher ratio of peak ~997 to ~1022 indicated small degree of ordered structure within starch conformation. The presence of amylose-lipid complexes has been reported to consists of some starch crystalline order (Kawai et al., 2012). In this study, longer ball milling time greatly reduced ordered structure of starch conformation. This ball milling effects on starch has been extensively studied in previous reports known by disruption effects on helical structures (Liu et al., 2011; Roa et al., 2015).

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245 3.4. Ball milling effects on thermal properties of OBPF dispersions

Thermal properties of OBPF prior to and after ball milling are presented in Table 4. Ball milling did not greatly affect the transition peak temperature, whereas the enthalpy values gradually decreased as the milling time increased, regardless the applied speed. Decreasing enthalpy values can be interpreted as the decrease on amounts of amyloselipid complexes. This is confirmed by further loss of crystallinity provided by FTIR spectroscopy study. Simultaneous mechanical impact during ball milling disrupted the helical structure of amylose, hence disintegrated the amylose-lipid complexes. Ball milled samples did not differ in transition temperature peak upon the second heating showing thermal behaviour remain unchanged.

255

256 3.5. Ball milling effects on particle size distribution

257 Particle size distributions of OBPF prior to and after ball milling are presented in 258 Figure 4 and Table 5. Prior to ball milling, wide range of particle size distributions were 259 observed. The D₁₀, D₅₀, and D₉₀ values represent 10, 50, and 90% particle size 260 distributions within the dispersed sample, whereas the $D_{(4,3)}$ show average particle sizes. 261 Ball milling at high speed reduced the particle size distributions to about quarter of initial values, e.g. from 116.45 \pm 21.85 to 27.67 \pm 0.93 $\mu m.$ However, low speed ball milling 262 263 could reduce to only about half of initial even after 360 min milling duration. This might 264 be explained that rotational speed is a major factor to cause particle breakage during ball 265 milling as suggested by Yin et al. (2015).

266

267 3.6. Ball milling effects on viscoelastic properties of OBPF dispersions

Figure 5 illustrated the viscoelastic properties of OBPF dispersions affected by ball milling and temperature changes. Ball milled samples at higher speed exhibited the lowest moduli at the same applied shear. Since the dispersions contain same weight fraction of samples, the volume packing fraction of HSST and LTST dispersions were lower, and hence decreased the storage moduli.

12

273 Temperature effects on moduli were observed in both heating and cooling cycles. 274 Moduli decreases occurred below 40 °C seems to be the effects of increasing particle mobility since there were no thermal events recorded by scanning calorimetry within this 275 276 temperature range. On the other hand, further decrease on moduli when temperature 277 approached 90 °C indicate the melting of starch-lipid complexes contained in dispersions. 278 Noticeable differences in melting behaviour of ball milled samples were observed using 279 the tangent of phase angle (tan δ). Dispersions containing ball milled samples showed 280 lower tan δ when reach the melting temperature that might due to the lower enthalpy 281 changes.

The moduli of dispersions increased steadily upon cooling back to room temperature due to the recrystallisation of melted particles. Moreover, cooled dispersions reached higher moduli compared to initial values prior to heating. This might indicate the connectivity between particles were built during cooling which most likely contributed by starch.

287 Figure 6 illustrated the frequency-dependency of OBPF concentrated dispersions. 288 The moduli increased with increase of angular frequency. Effects of ball milling were 289 observed in the fitting parameters of frequency sweep data, presented in Table 6. 290 Decreases in intercepts are consistent with the decreases of G' and G" due to particle 291 volume packing effects. Ball milling also increased the slopes of moduli-frequency curves 292 which revealed a weaker gel-like structure. As the particle volume packing decreases, 293 solvent-particles interactions increase and inhibit the particle-particle interactions in 294 building the structure. This is agreed with previous study on other flour dispersions 295 (Ahmed et al., 2014). Both G' and G" of all samples were satisfactory fitted to Power Law's model, shown by the R^2 values close to 1. 296

297

298

4. Conclusions

299 Major constituent of OBPF were observed in the form of aggregated proteins and 300 starch-lipid complexes. Ball milling altered the structural conformations of protein and 301 starch, and reduced the enthalpy changes of starch-lipid complexes melting properties. 302 Higher speed ball milling reduced particle size distributions to about quarter of initial 303 values, e.g. from 116.45 \pm 21.85 to 27.67 \pm 0.93 µm, led to lower particle volume 304 packing, and hence decreased the mechanical moduli of dispersions. Subsequent heating-305 cooling processes of OBPF dispersions allowed the starch to build a stronger networked 306 structure. Therefore, the starch contained in OBPF plays as important constituent in 307 influencing the structure of dispersion.

308

309 Acknowledgments

310 The first author would like express gratitude to the Ministry of Research, Technology,

311 and Higher Education of the Republic of Indonesia for the financial support.

312

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Figure captions

Figure 1 Typical infrared spectra of oat bran protein flour

Figure 2 DSC thermogram of OBPF dispersions upon heating-cooling cycles

Figure 3 (A) Second derivative spectra at amide I region of OBPF prior to and after ball milling; (B) original and second derivative spectra of OBPF at saccharide region. For abbreviations see Table 3.

Figure 4 Volume-weighted particle size distributions of OBPC prior to and after ball milling. For abbreviations see Table 3.

Figure 5 Storage (filled) and loss (open) moduli (top), and tangent of phase angle (bottom) of OBPC concentrated dispersions prior to and after ball milling over changing temperatures. For abbreviations see Table 3.

Figure 6 Storage (filled) and loss (open) moduli of OBPC concentrated dispersions prior to and after ball milling over changing angular frequencies. For abbreviations see Table 3.

Sampla	Peak height ratio relative to peak at ~ 1630 cm ⁻¹					
Sample	~2924 cm ⁻¹	~2854 cm ⁻¹	~1744 cm ⁻¹	$\sim 1150 \text{ cm}^{-1}$	~1078 cm ⁻¹	~1022 cm ⁻¹
OBPF	0.55 ± 0.04	0.11 ± 0.02	0.32 ± 0.02	0.67 ± 0.05	0.49 ± 0.06	0.94 ± 0.14
NPDO	0.31 ± 0.02	0.02 ± 0.00	0.16 ± 0.08	0.69 ± 0.06	0.53 ± 0.06	0.92 ± 0.03
PDO	0.29 ± 0.01	N/A	0.08 ± 0.01	0.79 ± 0.08	0.54 ± 0.08	1.02 ± 0.15
APDO	0.21 ± 0.02	N/A	0.05 ± 0.01	0.48 ± 0.06	0.37 ± 0.04	0.61 ± 0.12

Table 1. Infrared spectral changes on selected bands of OBPF and its derived fractions

Abbreviations: OBPF (Oat bran protein flour), NPDO (Non polar solvent defatted OBPF),

PDO (Polar solvent defatted OBPF), APDO (Alkaline exposed PDO).

Comula	First heating		First cooling		Second heating		Second cooling	
Sample	$T_p (^{o}C)$	$\Delta H (J g^{-1} \text{ solid})$	$T_p (^{o}C)$	$\Delta H (J g^{-1} \text{ solid})$	$T_p (^{o}C)$	$\Delta H (J g^{-1} \text{ solid})$	$T_p (^{o}C)$	$\Delta H (J g^{-1} \text{ solid})$
OBPF	90.63 ± 0.32	1.70 ± 0.21	74.14 ± 0.04	-1.02 ± 0.04	94.42 ± 0.10	1.01 ± 0.02	73.97 ± 0.04	-0.94 ± 0.02
NPDO	91.12 ± 0.74	0.94 ± 0.18	74.49 ± 0.05	$\textbf{-0.84} \pm 0.02$	94.98 ± 0.08	0.72 ± 0.13	74.34 ± 0.03	$\textbf{-0.72} \pm 0.07$
PDO	90.41 ± 0.59	0.76 ± 0.02	73.80 ± 0.25	-0.64 ± 0.04	94.07 ± 0.16	0.61 ± 0.04	73.73 ± 0.25	$\textbf{-0.53} \pm 0.08$
APDO	92.07 ± 0.51	0.48 ± 0.01	72.28 ± 0.41	-0.32 ± 0.02	92.76 ± 0.16	0.26 ± 0.02	71.88 ± 0.61	-0.26 ± 0.05
For abbraviations see Table 1								

Table 2. Thermal transition peak temperatures (T_p) and enthalpy changes (ΔH) of OBPF dispersions and its derived fractions

For abbreviations see Table 1.

Sample	Relative area of protein se	Starch		
Sample	Total β -sheets and β -turns (%)	Total α -helix and random coil (%)	~997/~1022 cm ⁻¹ ratio	
OBPF	88.03 ± 2.69	11.97 ± 2.69	0.63 ± 0.02	
HSST10	82.33 ± 2.68	17.67 ± 2.68	0.57 ± 0.06	
HSST20	82.53 ± 1.02	17.47 ± 1.02	0.56 ± 0.05	
HSST30	79.85 ± 0.61	20.15 ± 0.61	0.50 ± 0.02	
LSLT120	80.96 ± 4.39	19.04 ± 4.39	0.56 ± 0.04	
LSLT240	81.88 ± 1.38	18.12 ± 1.38	0.49 ± 0.04	
LSLT360	80.33 ± 6.95	19.67 ± 6.95	0.38 ± 0.04	

Table 3. Ball milling effects on protein and starch structural conformation

Abbreviations: OBPF (Oat bran protein flour), HSST10, 20, and 30 (Ball milled OBPF at high speed, i.e. 800 rpm for short time, i.e. 10, 20, or

30 min), LSLT120, 240, and 360 (Ball milled OBPF at low speed, i.e. 200 rpm for long time, i.e. 120, 240, or 360 min)

	First heating		First cooling		Second heating		Second cooling	
Sample	$T_n (^{\circ}C)$	$\Delta H (J g^{-1} \text{ solid})$	$T_n (^{\circ}C)$	$\Delta H (J g^{-1} solid)$	T_{p} (°C)	$\Delta H (J g^{-1} solid)$	T_{n} (°C)	$\Delta H (J g^{-1} \text{ solid})$
	r v /		r × ,		F ` '		F Ý	
OBPF	90.63 ± 0.32	1.70 ± 0.21	74.14 ± 0.04	-1.02 ± 0.04	94.42 ± 0.10	1.01 ± 0.02	73.97 ± 0.04	$\textbf{-0.94} \pm 0.02$
HSST30	90.50 ± 0.26	1.19 ± 0.14	73.72 ± 0.09	-0.78 ± 0.07	94.10 ± 0.01	0.65 ± 0.12	73.63 ± 0.10	-0.51 ± 0.01
LSLT360	91.20 ± 0.34	0.85 ± 0.07	74.04 ± 0.09	-0.77 ± 0.07	94.52 ± 0.07	0.66 ± 0.10	73.96 ± 0.07	-0.60 ± 0.04

Table 4. Thermal transition peak temperatures and enthalpy changes of OBPF dispersions affected by ball milling

For abbreviations see Table 3.

Sample	D _(4,3) (µm)	D ₁₀ (µm)	D ₅₀ (µm)	D ₉₀ (µm)	
OBPF	116.45 ± 21.85	27.33 ± 0.80	98.11 ± 5.79	203.50 ± 21.92	
HSST	27.67 ± 0.93	4.09 ± 0.10	23.22 ± 0.47	56.67 ± 1.66	
LSLT	49.33 ± 0.23	10.01 ± 0.86	46.88 ± 0.36	91.55 ± 0.05	

Table 5. Particle size distributions OBPI	F prior to and after ball milling
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For abbreviations see Table 3.

Table 6 . Parameters of moduli-frequency fitting based on the Power Law model	
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Sample	G'			G"		
	a' (kPa)	b'	R^2	<i>a'</i> (kPa)	b'	R^2
OBPF	7.94 ± 0.93	0.16 ± 0.00	0.986 ± 0.008	3.18 ± 0.17	0.21 ± 0.01	0.992 ± 0.004
HSST	2.44 ± 0.11	0.19 ± 0.05	0.985 ± 0.005	0.85 ± 0.35	0.24 ± 0.02	0.994 ± 0.002
LSLT	3.15 ± 0.67	0.19 ± 0.04	0.988 ± 0.002	1.11 ± 0.02	0.24 ± 0.02	0.994 ± 0.003

For abbreviations see Table 3.