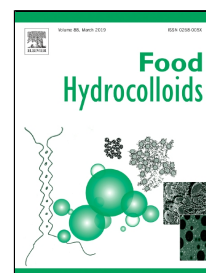


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Emulsifying properties of hemp proteins: effect of isolation technique

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1 Emulsifying properties of hemp proteins: effect of isolation technique

2
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14

15 Abstract

16 Hemp protein was isolated from hemp seed meal using two different isolation procedures: alkali
17 extraction/isoelectric precipitation (HPI) and micellization (HMI). The ability of these proteins to form
18 and stabilize 10% (w/w) sunflower oil-in-water emulsions (at pH = 3.0) was studied at three different
19 concentrations, 0.25, 0.75 and 1.5% (w/w), by monitoring emulsion droplet size distribution,
20 microstructural and morphological properties, rheological behaviour and stability against flocculation,
21 coalescence and creaming. In addition, hemp proteins were analysed for water solubility, denaturation
22 degree and surface/interfacial activity. HMI protein, which was found to be less denatured after
23 isolation, exhibited higher solubility and slightly higher surface/interfacial activity than HPI protein. HMI
24 emulsions possessed a smaller volume mean droplet diameter ($d_{4,3} = 1.92 - 3.42 \mu\text{m}$ in 2% SDS) than HPI
25 emulsions ($d_{4,3} = 2.25 - 15.77 \mu\text{m}$ in 2% SDS). While HMI stabilized emulsions were characterized with
26 individual droplets covered by protein film, both confocal laser scanning microscopy and flocculation
27 indices indicated occurrence of bridging flocculation in HPI stabilized emulsions. Protein aggregation,
28 which induced flocculation of the droplets, contributed to higher apparent viscosity of HPI stabilized
29 emulsions compared to HMI stabilized emulsions. Interestingly, emulsions stabilized with 1.5% (w/w)
30 HPI exhibited much better creaming and coalescence stability than other emulsions due to the
31 formation of a weak transient network of floccules and higher continuous phase viscosity which both
32 suppressed the movement of the droplets.

33 **Key words:** *hemp protein, isolation technique, emulsion stability, droplet size distribution, bridging*
34 *flocculation*

35

36

37

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38 1 Introduction

39

40 Proteins commonly used for food and pharmaceutical applications are whey protein, casein,
41 soy, egg white proteins, gelatin, bovine serum albumin, collagen, etc. (Lam & Nickerson, 2013).
42 However, over the past few years, food and cosmetics industries have been testing the
43 potential use of vegetable proteins from alternative sources due to increased interest.

44 The use of plant-based proteins as an alternative to animal derived proteins is desired due to
45 the higher sustainability, abundancy, and wide variety of sources (Bučko et al., 2015; Hadnađev
46 et al., 2017). Oilseeds are rich source of proteins, which can be extracted from the oil seed
47 meal, a by-product of the oil processing.

48 Hemp (*Cannabis sativa* L.) is a widely cultivated plant of industrial importance. In the past, it
49 was grown to produce hemp fibre for durable fabrics, while today it is becoming increasingly
50 utilized for edible oil extraction (Pojić et al., 2014). Hemp seed contains over 30% oil and 25%
51 protein, with a considerable amount of dietary fibre, vitamins and minerals. Hemp seed main
52 proteins are albumin and edestin. Hempseed proteins are an excellent source of digestible
53 amino acids when compared to other vegetable proteins like borage meal, canola meal
54 (Callaway, 2004). A direct comparison of the amino acid profiles from egg white, hempseed and
55 soy bean has shown that hemp seed is comparable to both egg white and soybean, which are
56 both considered to be high quality proteins (Tang, Ten, Wang, & Yang, 2006). In order to isolate
57 the proteins from hempseed meal (HMP), a by-product obtained after processing the seeds into
58 edible oil, alkali extraction is usually employed, followed by isoelectric protein precipitation or
59 micellization techniques (Hadnađev et al., 2018; Malomo & Aluko, 2015; Tang et al., 2006).
60 Malomo, He, & Aluko (2014) have reported that after protein alkali extraction, the protein
61 content in the obtained isolate (84.15%) was significantly higher than the determined value in
62 the starting material - HMP (44.32%). Comparing two isolation techniques, Hadnađev et al.
63 (2018) concluded that micellization (a "salting in-salting out" process) resulted in higher hemp
64 protein purity (98.87%) than the alkaline extraction/isoelectric precipitation technique
65 (91.44%).

66 Proteins generally exhibit a wide range of techno-functionalities such as foaming, emulsifying,
67 gelling and film-forming ability, fat absorption capacity, etc. (Pojić, Mišan, & Tiwari, 2018). Due
68 to their amphiphilic nature and film forming abilities, proteins are widely used as functional
69 ingredients for the formation and stabilisation of emulsions (Lam & Nickerson, 2013). While
70 small molecular weight emulsifiers diffuse rapidly to the interface during emulsion formation,
71 proteins diffuse at a much slower rate due to their higher molecular weight (McClements,
72 1999). Once adhering to the interface, proteins form viscoelastic films at the surface of the oil
73 droplets providing electrostatic and steric stabilization which are largely influenced by protein
74 type and environmental conditions (Tcholakova, Denkov, Ivanov, & Campbell, 2006a). At the
75 isoelectric pH value, where the net charge of the protein molecules is neutral, protein

76 aggregation occurs, leading to emulsion instability (Foegeding & Davis, 2011; Guzey &
77 McClements, 2007). Beyond the isoelectric point, pH conditions influence the overall charge
78 and functional behaviour of the protein. For example, improved emulsion stability and
79 viscoelastic properties of interfaces of amaranth proteins were observed at pH 2.0, in contrast
80 to pH 8.0; this was related to the denatured state of the proteins at pH 2.0, where proteins
81 formed a harder interfacial film, as compared to pH 8.0 (Ventureira et al., 2012). Moreover,
82 additional emulsion stabilization is achieved by proteins present within the continuous phase
83 which act as bulking agents, increasing emulsion viscosity, and thus reducing the mobility and
84 coalescence of oil droplets within the emulsion (Jafari, Beheshti, & Assadpoor, 2012).

85 The physico-chemical properties of proteins play an important role in determining their ability
86 to form and stabilize the emulsion (Papalamprou, Doxastakis, & Kiosseoglou, 2010). Solubility is
87 one of the most important features of proteins, since many functional properties of protein
88 depend upon their capacity to initially go into solution (Radha & Prakash, 2009). In general,
89 hemp protein exhibits low solubility in comparison to other vegetable proteins, which is
90 attributed to edestin (11S globulin) aggregation at pH below 7.0 (Malomo et al., 2014). The
91 employed protein isolation technique can influence protein solubility. It was shown that alkali
92 extracted hemp protein (HPI) has minimum solubility at pH 5.0, which increased as pH was
93 decreased or increased. In contrast, salt extracted hemp protein (HMI) has minimum solubility
94 at pH 6.0. The higher solubility of HMI at lower pHs suggests that HMI may be in a more native
95 state than HPI (Hadnađev et al., 2018).

96 An emulsifying activity index (EAI) and emulsifying stability index (ESI) of hemp stabilized
97 emulsions has been reported by Tang et al. (2006), which has shown that EAI profiles of HPI at
98 different pH values have the similar pattern to the protein solubility (PS) profiles, suggesting a
99 possible relation between those two characteristics. Lower EAI and ESI of hemp protein
100 stabilized emulsions, when compared with those of soy protein, were attributed to the
101 propensity of hemp protein to form covalent disulfide bonds between individual proteins and
102 its subsequent aggregation at neutral or acidic pH (Tang et al., 2006). The decreased solubility
103 of canola (CaPI) and flax seed (FIPI) protein isolates, relative to whey protein isolate (WPI), were
104 responsible for their lower EAI and ESI, although emulsion capacity of CaPI was comparable to
105 WPI (Karaca Low, & Nickerson, 2011b). Karaca et al. (2011b) also reported that CaPI and FIPI
106 produced by salt extraction showed improved solubility and interfacial activity compared to
107 those produced by isoelectric precipitation. Although salt extracted CaPI exhibited better EAI
108 than isoelectric precipitated CaPI, the former was characterized with a higher mean emulsion
109 droplet diameter and rapid creaming. On the contrary, the isolation method did not influence
110 significantly the EAI and creaming stability of FIPI. The different response of CaPI and FIPI to
111 isolation technique, however, remains unclear.

112 In order to corroborate deeply the effect of protein isolation technique on emulsion formation
113 as well as the ability of hemp protein to stabilize emulsions, the aim of this work was to

114 compare the emulsifying properties of alkali extracted/isoelectric precipitated hemp protein
115 (HPI) and micellar (salt extracted) hemp protein (HMI).

116 In general, emulsifying properties of hemp proteins are poorly investigated, mostly due to their
117 low solubility. Therefore, enzymatically hydrolysed version of hemp protein was mostly
118 investigated for its emulsifying ability (Yin et al., 2008), while hemp protein isolate stabilized
119 emulsions were only characterized in terms of emulsifying activity index (EAI) and emulsifying
120 stability index (ESI) in comparison to soy protein isolate. However, there is a lack of knowledge
121 in the hemp protein emulsions morphology and structure, rheological behaviour, stability
122 mechanism, effect of solubility and denaturation degree on emulsifying ability and stability, etc.
123 In this paper, emulsifying properties of two hemp proteins (HPI and HMI), isolated using
124 different techniques, were compared at different protein concentrations and pH condition far
125 from the isoelectric point ($\text{pH} = 3.0$). These conditions were chosen based on previous studies
126 investigating the influence of pH on protein emulsifying properties. These studies confirmed
127 improved emulsification and stability against coalescence with protein-stabilized emulsions at
128 pHs below the isoelectric point (pI), due to extensive protein unfolding induced by the extreme
129 pH conditions (Liang & Tang, 2013; Ventureira et al., 2012). According to Gharsallaoui, Cases,
130 Chambin, & Saurel (2009) better stability of pea protein-based emulsions at acidic conditions in
131 comparison to neutral or alkali pH, was influenced by the ability of pea protein to form a
132 stronger and denser viscoelastic network when adsorbed at the interface.

133 The emulsions prepared in this paper were characterized in terms of their droplet size
134 distribution, microstructural and morphological properties, rheological behaviour and stability
135 and related to protein solubility, denaturation degree and surface/interfacial activity.

136

137 **2 Materials and methods**

138

139 **2.1 Materials**

140 Hemp pellets obtained as a by-product in cold oil processing were provided by the local
141 company Svet Konoplje (Kisač, Serbia). Refined sunflower oil was purchased in local market.
142 Nile blue was procured from Sigma-Aldrich. All chemicals used were of reagent grade and
143 deionized water was used to prepare all solutions used in the experimental work.

144

145 **2.2 Preparation of protein isolates**

146 Hemp pellets were ground in a Foss Knifetec 1095 laboratory mill (Foss, Hillerød, Denmark)
147 equipped with water circulation, to ensure sample cooling and avoid overheating during milling
148 procedure. The obtained ground hemp meal was separated into two fractions using a universal
149 laboratory sifter (Bühler AG, Uzwil, Switzerland): a coarse fraction (particle size $\geq 250\mu\text{m}$) and
150 fine fraction (particle size $< 250\mu\text{m}$). The fine hemp meal fraction contained significantly higher

151 protein content than the course fraction, and thus, was employed for further protein extraction
152 (Pojić et al., 2014).

153 Prior to the hemp protein extraction process, fine hemp meal was defatted using a triple
154 hexane extraction at 1:3 hemp meal to hexane ratio for 2 h each. Subsequently, the defatted
155 hempseed meal was air dried in fume hood at room temperature.

156 After drying, defatted hempseed meal was used, as a starting material, to isolate proteins with
157 the aid of two different extraction techniques: alkali extraction/isoelectric precipitation and
158 micellization, both techniques are described in detailed by Hadnađev et al. (2018). Both alkali
159 extracted hemp protein, assigned as HPI ($91.44 \pm 0.34\%$ protein content on dry matter basis),
160 and micelle protein referred to as HMI ($98.87 \pm 0.41\%$ protein content on dry matter basis) were
161 prepared in triplicates and compared for their emulsifying ability.

162

163 **2.3 Protein solubility determination (at pH 3.0)**

164 Under constant stirring, 1% (w/w) suspensions of hemp protein isolates were adjusted to pH
165 3.0 with the aid of 1 M HCl followed by further stirring for 1 h at room temperature (23 °C) with
166 pH maintenance throughout. Afterwards, centrifugation was applied to the protein solution for
167 30 min at 7500xg. The obtained supernatant was filtered, diluted accordingly and the protein
168 content was determined by the Lowry method (Lowry, Rosenbrough, Fair, & Randall, 1951).
169 Protein solubility was determined as the ratio of protein content in supernatant of the
170 suspension as compared to the total protein content. The measurements were performed in
171 triplicates.

172

173 **2.4 Protein suspension/solution preparation**

174 HMI and HPI protein solutions were prepared by dispersing 0.25, 0.75 and 1.5% (w/w) of
175 proteins isolates (calculated on protein content) in deionized water. Under constant stirring, pH
176 was adjusted to 3.0 with 1.0 M HCl and stirred for additional 2 h at room temperature.
177 Subsequently, sodium azide (0.02% w/v), which was used as an antimicrobial agent, was added
178 and solutions were stored overnight at 4 °C in order to allow complete protein hydration. Two
179 separate batches of suspensions were prepared for analysis.

180

181 **2.5 Protein solutions surface and interfacial tension measurements**

182 The surface and interfacial tension measurements were carried out on a digital Krüss Easy Dyne
183 tensiometer (Hamburg, Germany) using a du Noüy ring method. The surface tension values
184 were corrected by Harkins and Jordan method, which is integrated into the electronic
185 Krüss tensiometer (Milanović, Krstonošić, Dokić, Hadnađev, & Dapčević Hadnađev, 2015). Prior
186 to surface tension (air-protein solution) measurements, the ring was immersed in the liquid and
187 left for 10 min in order to allow the surface to equilibrate. For interfacial tension measurements
188 (oil-protein solution), the ring was firstly immersed in the solution, and afterwards, sunflower

189 oil was poured on the top of the solution surface followed by system equilibration for 10 min.
 190 The reported surface/interfacial tension values were the average of five measurements per
 191 batch at constant temperature of 25 ± 0.1 °C. Surface tension of the solvent (water adjusted to
 192 pH = 3.0) was recorded as 72.3 ± 0.09 mN/m, while its interfacial tension was 23.3 ± 0.17 mN/m.
 193 Surface activity was expressed as surface/interfacial pressure:

$$194 \quad \pi = \sigma_0 - \sigma \quad (1)$$

196 where σ_0 and σ are surface/interfacial tensions of solvent and protein solution, respectively
 197 (Bučko et al., 2015).
 198

200 2.6 Emulsion preparation

201 Six different hemp protein isolate solutions were prepared (0.25, 0.75 and 1.5% (w/w) HPI and
 202 HMI), as described in section 2.4, to act as the continuous phases in emulsion preparation. Oil-
 203 in-water emulsions were prepared by blending 90% (w/w) hemp protein isolate solution (0.25 -
 204 1.5% protein content per continuous phase) with 10% (w/w) sunflower oil. The emulsions were
 205 pre-homogenized using an Ultraturrax T-25 (IKA® Werke GmbH & Co. KG, Germany) under
 206 continuous mixing at a speed of 12000 rpm for 60 s. Afterwards, pre-homogenized emulsions
 207 were homogenized at 400 bar using an APV Lab 1000 homogenizer (SPX flow, Germany). Two
 208 separate batches of emulsion were prepared for analysis.
 209

210 2.7 Determination of droplet-size distribution

211 Particle size distribution of the freshly prepared and stored (24 h) emulsions was measured at
 212 room temperature utilizing a Malvern Mastersizer 3000 equipped with a Hydro R cell (Malvern
 213 Instruments Ltd, Worcestershire, UK). An obscuration between 4-10% was employed and
 214 deionized water or 2.0% (w/v) SDS solution was used as the dispersant. Refractive index values
 215 of 1.33 for continuous phase and 1.471 for sunflower oil were used as the optical parameters,
 216 while emulsion droplet absorbance was set to 0.1. Relative refractive index, which represents
 217 the ratio of the dispersed refractive index to that of the dispersant, was set at 1.106. Droplet
 218 size distribution was reported as distribution profile as well as the mean volume diameter ($d_{4,3}$):
 219

$$220 \quad d_{4,3} = \frac{\sum n_i d_i^4}{\sum n_i d_i^3} \quad (2)$$

221 where n_i is the number of droplets with diameter d_i .

222 Droplet size distribution parameters were presented as the mean value of three measurements
 223 of each batch.
 224

225

226 **2.8 Flocculation and coalescence quantification**

227 Flocculation and coalescence indices were evaluated as described by Liang & Tang (2013).
 228 Mean volume diameter ($d_{4,3}$) determined as described above for freshly prepared (0 h) and
 229 stored (24 h) emulsions diluted in deionized water with and without 2.0% (w/v) SDS was used
 230 to calculate the percentage of flocculation index (%FI) and percentage of coalescence index
 231 (%CI):

232

$$233 \quad Fl(\%) = \left[\frac{d_{4,3} \text{ in water}}{d_{4,3} \text{ in 2\% SDS}} - 1 \right] \cdot 100 \quad (3)$$

234

$$235 \quad Cl(\%) = \left[\frac{d_{4,3} (24 \text{ h})}{d_{4,3} (0 \text{ h})} - 1 \right] \cdot 100 \quad (4)$$

236

237 **2.9 Confocal laser scanning microscopy**

238 Confocal laser scanning microscopy (CLSM) was performed using a Leica TCS SP5 Confocal Laser
 239 Scanning Microscope (CLSM; Leica Microsystems CMS GmbH, Wetzlar, Germany) using 63x oil
 240 immersion objective at 3 times zoom factor. Emulsions were imaged used dual confocal
 241 illumination through the use of an Argon laser at 488 nm and a Helium/Neon laser at 633 nm to
 242 show lipids and proteins, respectively. To prepare the samples for imaging 10 μ L of Nile Blue
 243 (0.1 g/100 μ L) was added into 1 mL of emulsion and vortexed for 10 s. After vortexing,
 244 approximately 50 μ L of labelled emulsion was pipetted onto a microscope slide and a coverslip
 245 was placed on top. Images were taken using simultaneous dual-channel imaging and were
 246 pseudo-coloured to show protein (red) and lipids (green). Emulsions were evaluated 24 h after
 247 preparation to qualify the microstructure of the droplets after equilibration.

248

249 **2.10 Rheological properties of emulsions**

250 Rheological measurements were carried out using a Haake MARS rheometer (Thermo Scientific,
 251 Germany) equipped with DG41 Ti cylinder measuring geometry. For each emulsion, 6.3 ml of
 252 liquid was carefully loaded into measuring cup. The steady-state flow measurements were
 253 recorded at 25 ± 0.1 °C in the range of 0-100 1/s. Shear rate was increased linearly for 3 min,
 254 then it was held on maximum shear rate of 100 1/s for 2 min, afterwards the shear rate was
 255 decreased linearly from 100 to 0 1/s for 3 min. The obtained data were described using power
 256 law model:

257

$$258 \quad \tau = K\dot{\gamma}^n \quad (5)$$

259

260 where τ is the shear stress (Pa), K is the consistency index (Pa s^n), and n is the flow behaviour
261 index. All the rheological measurements were performed in triplicates for each batch 24 h after
262 preparation.

263

264 **2.11 Emulsion stability measurements**

265 Stability of previously prepared emulsions was estimated using both simple (under gravitational
266 forces) and rapid (under centrifugal forces) creaming methods. Gravitational separation in
267 emulsions was followed by visual observation. The tested emulsions were placed in 10 ml
268 sealed transparent glass-cylinders, gently agitated to ensure initial homogeneity and left for 14
269 days at room temperature. The extent of creaming was characterized as the bottom (serum)
270 layer height to the total height of the emulsion.

271 Rapid stability method was conducted using a LUMiSizer Dispersion Analyser (L.U.M. GmbH,
272 Germany). The measurements were performed by applying centrifugal sedimentation in order
273 to accelerate the instability phenomena of tested emulsions such as sedimentation, creaming
274 or flocculation (Petzold, Goltzsche, Mende, Schwarz, & Jaeger, 2009; Sobisch & Lerche, 2008).
275 The obtained recordings refer to intensities of transmitted light as a function of time i.e.
276 percentage of light absorbance per hour which is related to "creaming rate". Creaming rate is
277 inversely related to emulsion stability, and thus, the LUMiSizer can be employed to judge the
278 performance of an emulsion system. The parameters on the LUMiSizer were as follows: 600 μL
279 of emulsion, 1500 rpm, time 3600 s, time interval 60 s and temperature 25 °C. Measurements
280 were conducted in duplicates for each batch 24 h after preparation.

281

282 **2.12 Statistical analysis**

283 The obtained results were expressed as mean values of replicate analyses. Analysis of variance
284 (ANOVA) and Tukey's test were used to assess significant differences among the mean values at
285 significance level of 0.05. The statistical analyses were performed by software TIBCO Statistica™
286 13.3.0 (TIBCO Software Inc., Palo Alto, CA, USA).

287

288 **3 Results and discussion**

289

290 **3.1 Protein isolates characterization**

291 Protein isolates from hemp seed meal were prepared using two different isolation techniques:
292 alkaline extraction/isoelectric precipitation (HPI) and micellization (a "salting in - salting out"
293 process; HMI). Both techniques resulted in the extraction of the same protein fraction, i.e.
294 edestin (11S globulin) as identified by protein electrophoretic profiles (Hadnađev et al., 2018).
295 However, according to FTIR spectra and differential scanning calorimetry, HPI and HMI differ in
296 their secondary structure, since highly alkaline conditions during HPI extraction leads to partial

297 protein denaturation, thus HMI is generally found in a more preserved native state (Hadnađev
298 et al., 2018).

299 The influence of isolation technique on the solubility of two hemp proteins, prepared in this
300 study (HPI and HMI), is presented in Table 1. In general, under the chosen conditions (pH = 3.0),
301 both protein isolates had solubility higher than 70%, which is an important prerequisite for a
302 protein to be an effective emulsifier (Liang & Tang, 2013). Comparing the isolates prepared in
303 this study, it can be noticed that HMI possessed significantly better solubility at pH 3.0 than HPI,
304 which is in agreement with previous studies using the employed isolation techniques on canola
305 and flaxseed protein isolates (Karaca et al., 2011b; Krause, Schultz, & Dudek, 2002). The
306 observed lower solubility of HPI could be the consequence of partial denaturation during its
307 preparation (Hadnađev et al., 2018). **On the contrary, due to mild isolation conditions,
308 hydrophobic part of HMI was placed in the core region of the molecule.**

309 The effect of protein preparation procedure as well as proteins solution concentration on
310 surface/interfacial pressure is summarized in Table 1.

311 By increasing the protein concentration in solution an increase in surface/interfacial pressure
312 was observed, i.e. more proteins were adsorbed at the interface. Niño, Sánchez, Fernández, &
313 Patino (2001) reported that the protein concentration dependence on surface pressure
314 expresses classical sigmoidal behaviour, i.e. increase in surface pressure with protein
315 concentration and reaching of a plateau at the maximum protein concentration in the bulk
316 phase. Only a slight increase in the surface pressure was observed with higher concentrations
317 for both HPI and HMI, suggesting that the examined concentration range (0.25 – 1.50% w/w)
318 was close to the plateau value. In general, the decrease in interfacial tension between air/water
319 and oil/water due to hemp protein adsorption was 24.93 – 32.61 mN/m and 9.13 – 15.73
320 mN/m, respectively, which is comparable to the values recorded for other plant protein
321 isolates, such as pumpkinseed protein (Bučko et al., 2015), lentil protein isolate (Joshi et al.,
322 2012), amaranth protein (Ventureira et al., 2012).

323 Comparing the two proteins, HMI expressed slightly higher surface/interfacial activity than HPI.
324 This was probably influenced by higher solubility of HMI since higher protein solubility led to
325 increased biopolymer mobility and greater diffusion rates to the interface (Lam & Nickerson,
326 2013; Karaca Low, & Nickerson, 2011a). Liang & Tang (2013) also reported that the pea protein
327 adsorption at the interface at pH 3.0 was closely associated with its solubility. Onsaard,
328 Vittayanont, Srigam, & McClements (2006) also noticed that freeze–thaw treated coconut
329 cream protein (CCP2), which was characterized with more than two times higher solubility than
330 isoelectric precipitated protein (CCP1), attained higher surface pressure at the air–water
331 interface.

332 According to the solubility values (at pH = 3.0) of hemp proteins prepared in this study, more
333 pronounced differences in their surface activities could be expected.

334 The results of Liang & Tang (2013) indicated that protein behaviour at the interface is also
335 highly correlated with its surface hydrophobicity. Although it could be expected that increase in
336 protein denaturation degree due to isolation conditions would lead to increase in its
337 hydrophobicity, study on the effect of extraction pH on the properties of the pigeonpea and
338 cowpea protein isolates has demonstrated that increase in the degree of protein denaturation
339 resulted in an increase in the % of hydrophobic amino acids, while the exposed
340 hydrophobicities decreased (Mwasaru et al., 1999).

341 The results obtained in this study indicated that, besides solubility and surface hydrophobicity,
342 the amount of protein adsorption at the surface/interface is also affected by the unfolding and
343 conformational flexibility of the protein at the interface (Liang & Tang, 2013), i.e. changes in the
344 protein secondary structure and agglomeration. According to FTIR spectra HPI protein
345 secondary structure was characterized with lower intensity of the peaks corresponding to
346 native protein structural elements such as intramolecular β -sheets and α -helices, as well as
347 higher intensities of "aggregation peaks" indicating enhanced protein aggregation compared to
348 HMI (Hadnađev et al., 2018). The results of the protein dispersions (30% by weight)
349 microstructure have also revealed that HMI behaviour was governed by electrostatic
350 repulsions, while hydrophobic interactions were noted between the HPI molecules (Dapčević-
351 Hadnađev, Hadnađev, Lazaridou, Moschakis, & Biliaderis, 2018). Therefore, increased
352 hydrophobic interactions forced the HPI protein molecules to the interface in order to minimize
353 unfavorable interactions (Papalamprou et al., 2010), although it was characterized with lower
354 solubility in comparison to HMI. Bučko et al. (2015) have reported that the largest surface
355 pressure of pumpkin protein solution was detected at pH = 5.0 where it exhibited the lowest
356 solubility and ascribed that to lower hydration of proteins and the increased hydrophobic
357 interactions.

358

359 **3.2 Protein isolates stabilized emulsions characterization**

360 Emulsifying properties of hemp proteins (HPI and HMI) at pH 3.0 were evaluated at various
361 protein concentrations (0.25, 0.75 and 1.5%, w/v) via emulsion droplet-size analysis,
362 microscopy, rheological and stability measurements.

363

364 **3.2.1 Emulsifying ability**

365 The droplet size distributions of emulsions in the absence of SDS are shown in Figure 1 and
366 were found to be mostly bimodal. Emulsions containing HPI were characterized with two
367 droplets populations: i) smaller droplets (diameter 0.5 – 1 μ m) which size did not change with
368 protein concentration and ii) larger droplets (diameters 10 – 50 μ m, 4 – 20 μ m and 1 – 5 μ m for
369 0.25, 0.75 and 1.5%, w/v HPI, respectively) which size decreased with increase in the protein
370 concentration. Emulsions stabilized with HMI had one distribution peak at $d = 0.4 - 1 \mu$ m which

371 height increased with increase in protein concentration and the second peak at $d = 1 - 5 \mu\text{m}$
372 which slightly shifted toward larger sizes with the decrease in the protein concentration.

373 In order to measure the effect of droplet flocculation, droplet-size distribution was also
374 determined in emulsions diluted in 2% SDS. The volume mean droplet diameter ($d_{4,3}$) of
375 emulsions with deionized water and 2% SDS as dispersant was calculated and summarized in
376 Table 2.

377 As the concentration of the HPI protein increased, the volume mean droplet diameter of the
378 emulsion significantly decreased (Table 2). The same increase in the concentration of HMI
379 protein was found to have less progressive influence on the $d_{4,3}$ as compared to HPI stabilized
380 emulsions. Thus HMI was deemed to have better emulsifying properties than the HPI protein,
381 since addition of HPI in lower concentrations was not enough to completely saturate the total
382 surface of emulsion droplets, which resulted in larger droplets and lower emulsifying activity.
383 The better emulsifying ability of HMI might be associated with its higher solubility at pH 3.0
384 and, consequently, faster adsorption rate at the interface. A similar relationship between
385 protein solubility and emulsifying ability at pH 3.0 was observed by Liang & Tang (2013).
386 Moreover, for HPI emulsions $d_{4,3}$ in water was significantly higher than $d_{4,3}$ in 2% SDS indicating
387 flocculation of oil droplets. Oil droplets flocculation was quantified by calculating flocculation
388 index (FI) which is also listed in Table 2. For HPI, the FI progressively increased with the increase
389 in protein concentration from 0.25 to 0.75 %, while further increase in protein concentration to
390 1.5% led to decline in FI value. On the contrary, HMI emulsions were characterized with small
391 flocculation indexes.

392

393 **3.2.2 Microstructure**

394 As both HPI and HMI were characterized with a similar surface activity but created emulsions
395 with different droplet sizes, especially at lower protein concentrations, and exhibited
396 differences in flocculation behaviour, confocal laser scanning microscopy (CLSM) was employed
397 to understand the possible mechanism involved (Fig. 2). Images capture of the HPI emulsions
398 demonstrated a visible decrease in the average droplet size with a subsequent increase in
399 protein concentration (Fig. 2 a-c), as confirmed by particle size measurements (Fig. 1 and Table
400 2).

401 As expected from FI index values (Table 2), CLSM images of the HPI stabilized emulsions showed
402 flocculation of the droplets, which appeared to be connected by protein aggregates (Fig. 2a-c).

403 In general, in biopolymer stabilized emulsions, two types of droplet-droplet interactions could
404 be distinguished: i) depletion and ii) bridging flocculation (Dickinson, 2003). Depletion
405 flocculation occurs when the biopolymer in the continuous phase of the emulsion is
406 unadsorbed or poorly adsorbed and is driven by an osmotic pressure gradient associated with
407 the exclusion of biopolymer from a narrow region surrounding the droplet, which results in
408 droplets attraction toward each other (McClements, 2000). On the contrary, bridging

409 flocculation occurs when biopolymers, added at lower levels, adsorbs onto the surface of more
410 than one droplet leading to droplets connection via bridges and subsequently their flocculation
411 (Dickinson, 2003). The interactions between the interfacial layer and the emulsion droplet
412 influence the prevailing flocculation mechanism.

413 According to Fig. 2 (a-c) bridging flocculation prevailed over depletion flocculation since the
414 droplets (green coloured) were surrounded and interconnected with protein molecules (red
415 coloured).

416 Formation of droplets aggregates was also reported in other hemp protein stabilized emulsions.
417 In one study, hemp protein was enzymatically hydrolysed; while this increased the solubility of
418 the protein, the emulsifying properties of the protein decreased; this was also hypothesized to
419 be due to presence of aggregates within the continuous aqueous phase which inhibited the
420 formation of viscoelastic film on the droplet surface (Yin et al., 2008). Furthermore, in a study
421 by Wang, Jiang, & Xiong (2018), hemp milk emulsions were studied for its stability using a
422 native and alkali adjusted version of the system. As seen within this study, the protein present
423 in its native state was able to create emulsions with enough droplet-droplet static repulsion to
424 generate a homogenous distribution of oil droplets. However, in the alkali adjusted solution
425 large aggregates of oil droplets were found and the droplet size of the emulsion subsequently
426 increased.

427 The differences in emulsifying ability of HPI and HMI could have been because they were
428 characterized with different denaturation degrees prior to emulsion preparation. According to
429 Fig. 2 (d-f), HMI had the ability to generate repulsive interactions (steric and/or electrostatic)
430 between the oil droplets which took prevalence over hydrophobic or covalent interactions
431 between proteins. Some protein-protein aggregates observed in Fig. 2 (e-f), might originate
432 from HMI exposure to extreme pH (3.0) and subsequent high pressure homogenization process
433 which led to protein structural unfolding and exposure of its hydrophobic core (Tcholakova,
434 Denkov, Sidzhakova, & Campbell, 2006b; Wang et al., 2018). On the contrary, the isolation
435 process of HPI provoked earlier protein conformational changes which allowed the formation of
436 hydrophobic bonds between proteins at different droplet interfaces, thus causing droplets
437 aggregation. Moreover, since hemp proteins possess a high concentration of sulfur-containing
438 amino acids (Hadnadev et al., 2018), exposure of previously hidden sulfhydryl groups upon
439 protein unravelling, and thus, consequent disulfide bond formation between protein molecules
440 must be considered. Different HPI and HMI emulsifying activities were probably the result of
441 different ratio of bonds formed at the interface and those formed between two droplet
442 interfaces. Similarly, Wang et al. (2012) reported that temperature (90 °C) induced
443 denaturation of soy proteins resulted in enhanced emulsion stability due to formation of
444 disulfide bonds with neighbouring proteins. However, increased heat treatment (120 °C) caused
445 a reduction in the emulsifying capabilities of the soy proteins due to excessive bonding among
446 the proteins and formation of aggregates.

447 Moreover, flocculation of the droplets covered with HPI could also be related to the fact that,
448 at pH 3.0, HPI contained higher amount of non-soluble proteins (Table 1). According to
449 Ventureira et al. (2012) the emulsions made with total proteins show more flocculation than
450 those made with soluble protein, since non-soluble proteins could be adsorbed in two different
451 interfaces at the same time thus leading to bridging flocculation.

452

453 3.2.3 Rheological behaviour

454 The rheological properties of emulsions were also monitored since they represent a useful
455 analytical measurement to provide insights into the structural organization and interactions of
456 the components within emulsions. Moreover, they influence emulsion shelf life and sensory
457 properties (McClements, 1999). According to flow behaviour indexes (n) obtained from power
458 law model (Eq. 5) all the emulsions showed near-Newtonian behaviour ($n=0.9004-1.0130$), i.e.
459 their viscosities were nearly independent on applied shear rates. The similar behaviour was
460 noticed by Demetriades, Coupland & McClements (1997) for 20 wt% corn oil-in-water
461 emulsions stabilized by 2 wt% whey protein isolate. In their study flow behaviour index was
462 close to 1 at high and low pH values. Emulsions (20 vol. %) stabilized with 2 wt% soy 11S
463 globulin obtained at low homogenizing pressure have also shown Newtonian flow behaviour
464 with quite low viscosities (Floury, Desrumaux & Legrand, 2002).

465 In general, rheological measurements (Fig. 3) revealed that HPI stabilized emulsions had slightly
466 higher viscosities than HMI stabilized emulsions.

467 The higher apparent viscosity of the emulsion prepared with HPI can be explained by the
468 occurrence of the bridging flocculation and enlargements of apparent droplet size. These
469 findings were in accordance with other studies which also reported that flocculated emulsions
470 were more viscous (Sun, Gunasekaran, & Richards, 2007; Wang et al., 2018; Xu, Wang, Jiang,
471 Yuan, & Gao, 2012). **According to McClements (1999), an emulsion containing flocculated
472 droplets has a higher viscosity than an emulsion containing the same number of isolated
473 droplets due to ability of the flocs to trap some of the continuous phase within their structure
474 thus exhibiting a higher effective volume fraction than the actual volume fraction of the
475 individual droplets.** Moreover, differences in the viscosities of continuous phase also
476 contributed to differences in the viscosity of final emulsion. As it can be noticed in Fig. 4, HPI
477 dispersion (1.5% w/w) expressed a higher viscosity than HMI dispersion of the same
478 concentration due to more intensive HPI protein-protein interactions and aggregation and
479 consequent increase in hydrodynamic volume size (Wang et al., 2018).

480 Moreover, rheological behaviour of HMI stabilized emulsions was not influenced with increase
481 in protein concentration, indicating that emulsion viscosity was insensitive to the amount of
482 unadsorbed HMI in the continuous phase. On the contrary, HPI stabilized emulsions have
483 exhibited an increase in apparent viscosity with the decrease in emulsifier concentration.
484 Higher viscosity of 0.25% HPI emulsions could be ascribed to the presence of large flocs

485 composed of large droplets (Fig. 2a) that were characterized with increased hydrodynamic
486 volume in comparison to 0.75% and 1.50% HPI emulsions, which consequently led to the
487 increase in resistance to flow and then of emulsion viscosity.

488

489 **3.2.4 Emulsion stability**

490 The stability of HPI and HMI emulsions upon storage of 24 h was evaluated in terms of
491 flocculation (FI) and coalescence (CI) indexes.

492 The coalescence index of droplets was determined as the relative change in $d_{4,3}$ (in 2% SDS) of
493 droplets, before and after storage for 24 h and summarized in Table 2. The flocculation stability
494 of the fresh and stored (24 h) emulsions was also evaluated as the relative change in $d_{4,3}$
495 determined in water and 2% SDS as dispersants and the results are also included in Table 2.

496 As expected, the CI for both HPI and HMI was much lower at higher protein concentrations than
497 that at lower concentration values (Table 2), confirming the results of other studies (Liang &
498 Tang, 2014). The highest coalescence stability was noticed in 1.5% HPI emulsions, which was
499 also characterized by the highest content of non-soluble proteins (Table 1). According to Wang
500 et al. (2018), interactions between the adsorbed and unadsorbed proteins at droplet interface
501 could impart a steric stabilization against further coalescence during storage, contributing to
502 the enhancement of emulsion stability. HPI emulsions prepared with low protein
503 concentrations (0.25 – 0.75%) were unstable against coalescence, as well as susceptible to
504 droplet flocculation. Moreover, in these emulsions FI increased after the storage of 24 h. On the
505 contrary, 1.5% HPI emulsions were more stable against flocculation in comparison to 0.25% and
506 0.75% HPI emulsions.

507 Liang & Tang (2014) also reported progressive increase in the flocculation index of pea protein
508 stabilized emulsions, as the concentration increased to 1.0 g/100 ml, followed by gradual
509 decrease in flocculation index with further increase in protein concentration. This was
510 explained by the dominance of the inter-droplet attractive interactions between the adsorbed
511 proteins on individual droplets at low protein concentrations and the prevalence of the
512 interactions between the adsorbed and unadsorbed proteins over the inter-droplet interactions
513 at high protein concentrations (Liang & Tang, 2014).

514 The similar phenomenon was observed in emulsions stabilized with protein-polysaccharide
515 mixtures, where low concentration of polysaccharide caused polymeric linkages between
516 proteins adsorbed on emulsion droplets leading to bridging flocculation. Higher concentrations
517 of the same polysaccharide were sufficient to completely cover the droplet interface and
518 stabilize the emulsion via steric and electrostatic forces (Dickinson & Eriksson, 1991; Dickinson
519 & Galazka, 1991).

520 Comparing the flocculation stability of emulsions prepared in this study, HMI emulsions were
521 characterized with significantly lower FI in comparison to HPI.

522 Thus it can be concluded that the mechanism for the emulsion instability vary with the
523 technique used to isolate the hemp protein and consequently protein denaturation degree.
524 **Droplet coalescence was the underlying mechanism of the HMI emulsion instability, while**
525 **instability of HPI emulsions was governed by bridging flocculation.**

526 The creaming stability of all the tested emulsions was also evaluated. In general, all emulsions
527 are thermodynamically unstable systems and if stored long enough, they will separate into an
528 opaque layer consisting of oil droplets ("cream") at the top and a slightly turbid or transparent
529 aqueous layer ("serum") at the bottom (Liang & Tang, 2013).

530 In this study, creaming stability was monitored under gravitational and centrifugal forces. The
531 gravitational separation of 0.25% HPI emulsion was the largest among all the tested emulsions
532 since this emulsion developed into two layers after 1 day of storage (Figure 4). The creaming
533 index at the first day of 0.25% HPI emulsion storage was 78% and it slightly increased over the
534 storage period. The other tested emulsions did not exhibited visible creaming under
535 gravitational forces during the tested storage period (Figure 4).

536 Therefore, accelerated destabilization test performed with the aid of Lumisizer measurements
537 was also used to determine the effect of HPI and HMI concentration on emulsions stability. The
538 measurement is based on the monitoring of changes in intensity of the parallel transmitted NIR
539 light over the entire sample length during centrifugation at different time intervals (Lim & Roos,
540 2015). At the beginning of the centrifugation, only a small amount of light is transmitted along
541 the sample cell, while, during centrifugation, the heavier and more transparent aqueous phase
542 moves to the bottom. This accelerated creaming process promotes transmission as the less
543 transparent oil phase moves to the top (Xiang, Liu, Fan, & Gao, 2015). The larger changes in the
544 transmission with centrifugation, i.e. higher slope of the integral transmission vs. time, indicate
545 emulsions with lower stability (Lim & Roos, 2015; Xiang, Liu, Fan, & Gao, 2015).

546 Integral transmission (%) against time (s) for all systems is shown in Fig. 5. As expected, the
547 creaming behaviour of emulsions in accelerated test was highly related to the applied protein
548 concentration. Emulsions with lower protein concentrations creamed more readily as
549 compared to systems with 0.75 and 1.5% (w/w) emulsifier. For the 0.25% (w/w) HPI emulsion,
550 the creaming rate was high at the beginning, and after that, it gradually increased (Fig. 5). In
551 contrast, at the beginning, the creaming instability of the HMI emulsion at concentration of
552 0.25% (w/w) was lower than that of 0.25% (w/w) HPI emulsion, but it progressively increased
553 over a prolonged period and after 3000 s centrifugation it was even higher. This indicates that
554 creaming stability under accelerated mode is associated with the combined effect of droplet
555 size as well as the flocculation and coalescence stability.

556 In general, creaming stability of emulsions can be enhanced with decreased droplet size and
557 increased continuous phase viscosity (McClements, 1999). In this study, the creaming instability
558 of HMI emulsions was the consequence of decreased stability against coalescence (Table 2) and
559 low continuous phase viscosity (Fig. 3) which led to increased movement of droplets due to

560 gravity or Brownian motion; however, fast creaming in 0.25 and 0.75% (w/w) HPI emulsions
561 was influenced with large droplet sizes and protein induced droplet bridging flocculation.
562 Interestingly, emulsions with 1.5% (w/w) HPI exhibited the best stability against creaming,
563 despite the bridging flocculation of droplets observed (Fig. 2 f).

564 In contrast to 1.5% (w/w) HMI emulsions, HPI emulsion prepared with the same amount of
565 protein exhibited bridging flocculation (Fig. 2 c) and higher continuous phase viscosity (Fig. 3).

566 In general, at low or intermediate droplet concentrations, flocculation tends to increase the
567 creaming rate because the floccules have a larger effective size than the individual droplets,
568 while in concentrated emulsions flocculation retards creaming due to formation of a three-
569 dimensional network of aggregated floccules that prevents the individual droplets from moving
570 (McClements, 1999). However, according to McClements (1999), a network can form at lower
571 disperse phase volume fractions when the droplets in a floccule are more openly packed, and
572 therefore creaming is prevented at lower droplet concentrations. Since, in this paper, droplets
573 in floccules were openly packed (Fig. 2), the possible reason for higher creaming stability of
574 1.5% (w/w) HPI emulsion could be attribute to formation of a weak transient network of
575 floccules that suppressed the movements of the droplets. Ventureira, Martinez, & Anon, (2010)
576 also noticed that emulsion prepared at pH 2.0 with non-hydrolyzed amaranth proteins showed
577 lower creaming rate in comparison to other emulsions although it was characterized with the
578 highest degree of flocculation. Moreover, aggregation of HPI protein molecules in the
579 continuous phase influenced an increase in the viscosity (Fig. 3), which suppressed droplets
580 movement within the emulsion (Frelichowska, Bolzinger, & Chevalier, 2010; Liang & Tang,
581 2014).

582

583 **4 Conclusions**

584 Alkali extracted/isoelectric precipitated hemp protein (HPI) and micellar (salt extracted) hemp
585 protein (HMI) possess different emulsification and emulsion stabilization mechanisms. The two
586 proteins differed in their denaturation state which contributed to the observed changes in their
587 solubility and surface/interfacial activity. HMI protein, which had preserved native state after
588 isolation process, formed emulsions with small droplets ($d_{4,3} = 1.92 - 3.42 \mu\text{m}$ in 2% SDS) and
589 enough droplet-droplet static repulsion. However, low viscosity of HMI stabilized emulsions
590 enabled fast droplet movement and led to increased creaming and coalescence at lower
591 protein concentrations (0.25 – 0.75% w/w). In the HPI stabilized emulsions, isolation technique
592 favoured pH-induced structural unfolding of protein molecules, exposure of hydrophobic sites
593 and sulfhydryl groups and subsequent formation of protein connected droplet aggregates
594 during emulsification. At lower HPI concentrations (0.25 - 0.75 % w/w), the dominant type of
595 interactions were the ones between proteins adsorbed at the droplets which influenced
596 bridging flocculation, while at higher HPI concentrations (1.5% w/w), there was the prevalence
597 of the interactions between the adsorbed and unadsorbed proteins at droplet interface as well

598 as the protein-protein interactions in the continuous phase. While the latter influenced
599 formation of transient network of floccules and increased continuous phase viscosity thus
600 contributing to the emulsion stability, the former led to increase in emulsion droplet size and
601 instability.

602

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611

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740

741

742 **Figure captions**

743 **Figure 1.** The size distribution profiles of alkali extracted hemp protein (HPI) and micellar hemp
744 protein (HMI) stabilized emulsion (fresh emulsions, diluted in water) at different protein
745 concentrations (0.25-1.5% w/w)

746 **Figure 2.** Confocal laser scanning micrographs of hemp protein stabilized emulsions at pH 3.0.
747 Emulsions stabilized with alkali extracted hemp protein (HPI) are shown in images at (a) 0.25%,
748 (b) 0.75% and (c) 1.50% w/w in the aqueous phase. Native micellar hemp protein (HMI)
749 emulsions are also shown with various concentrations of HMI: (d) 0.25%; (e) 0.75% and (f)
750 1.50%. Sunflower oil acting as the oil phase in the emulsion is labelled as green with hemp
751 protein in red. Scale bar = 25 μ m

752 **Figure 3.** Flow curves of 10% O/W emulsions stabilized by alkali extracted hemp protein (HPI)
753 and micellar hemp protein (HMI) at various levels (0.25-1.5% w/w) and 1.5% w/w HPI and HMI
754 dispersions (continuous phases)

755 **Figure 4.** Images of the alkali extracted hemp protein (HPI) and micellar hemp protein (HMI)
756 stabilized emulsion at different protein concentrations (0.25-1.5% w/w) after storage for 14
757 days

758 **Figure 5.** Integral transmission profiles as a measure of stability of 10% O/W emulsions
759 stabilized by alkali extracted hemp protein (HPI) and micellar hemp protein (HMI) at various
760 concentrations (0.25-1.5% w/w)

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Table 1. Solubility at pH = 3.0, surface and interfacial pressure of hemp protein isolate dispersions of different concentrations prepared by different extraction techniques

Sample	Solubility at pH = 3.0 (%)	
HPI	71.2±3.56 ^a	
HMI	88.2±0.73 ^b	
Sample concentration	Surface pressure π (mN/m)	Interfacial pressure π (mN/m)
0% HPI/HMI	0 ^a	0 ^a
0.25% HPI	24.93±0.259 ^b	9.13±0.115 ^b
0.75% HPI	27.67±0.148 ^c	11.07±0.059 ^c
1.50% HPI	30.47±0.148 ^d	12.63±0.503 ^d
0.25% HMI	28.21±0.167 ^c	9.47±0.252 ^b
0.75% HMI	30.95±0.122 ^d	13.60±0.557 ^d
1.50% HMI	32.61±0.26 ^e	15.73±0.416 ^e

Values followed by different letters indicate significant differences ($p \leq 0.05$)

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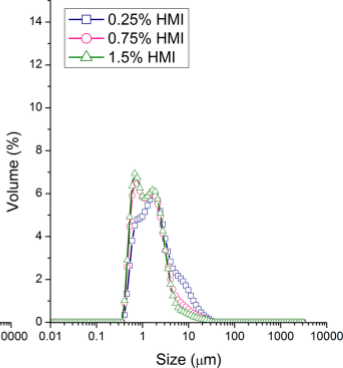
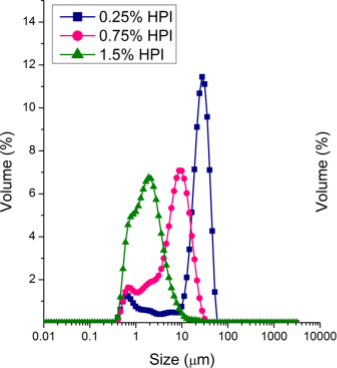
Table 2. Volume mean droplet diameter ($d_{4,3}$), flocculation (FI) and coalescence (CI) indices of HPI and HMI hemp protein stabilized emulsion, for different protein concentrations (0.25-1.5% w/w) immediately after preparation (0 h) and after 24 h of storage

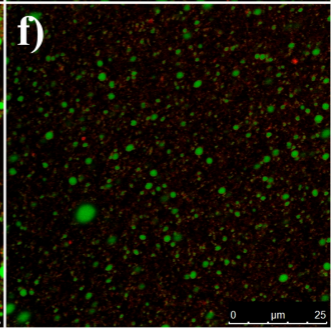
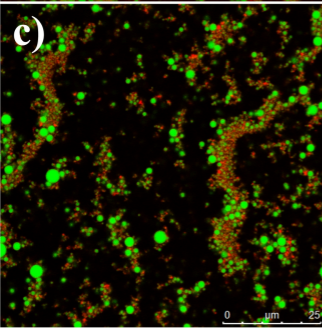
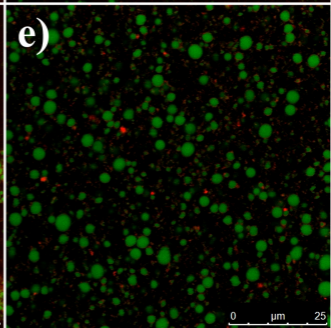
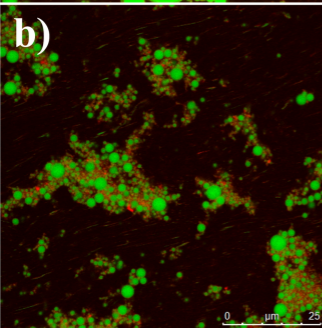
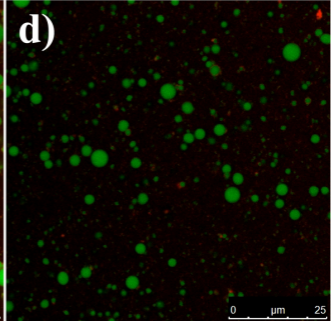
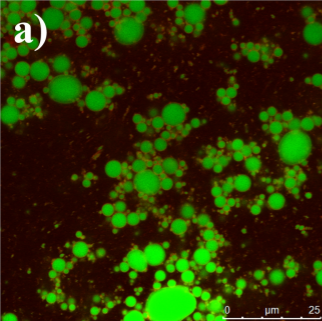
Sample	$d_{4,3}$ (μm)				FI (%)		CI (%)
	0 h		24 h		0 h	24 h	24 h
	Water	2% SDS	Water	2% SDS			2% SDS
0.25% HPI	24.17±0.814 ^c	15.77±0.404 ^c	39.80±1.670 ^d	24.13±1.320 ^c	53.3±2.56 ^c	65.0±3.46 ^c	53.1±7.43 ^d
0.75% HPI	8.76±0.431 ^b	3.80±0.141 ^b	12.15±0.495 ^c	5.10±0.283 ^b	130.6±2.77 ^d	138.3±3.51 ^d	34.2±2.45 ^c
1.5% HPI	2.71±0.260 ^a	2.25±0.160 ^a	2.86±0.290 ^{ab}	2.37±0.178 ^a	20.1±3.00 ^b	20.7±3.82 ^b	5.0±0.90 ^a
0.25% HMI	3.54±0.403 ^a	3.42±0.375 ^b	5.40±0.311 ^b	5.28±0.332 ^b	3.5±0.44 ^a	2.2±0.54 ^a	54.7±7.22 ^d
0.75% HMI	2.31±0.083 ^a	2.18±0.070 ^a	2.76±0.070 ^{ab}	2.63±0.090 ^a	5.8±0.56 ^a	5.0±0.93 ^a	20.5±1.31 ^{bc}
1.5% HMI	2.01±0.042 ^a	1.92±0.043 ^a	2.20±0.020 ^a	2.12±0.028 ^a	4.7±0.10 ^a	3.5±0.38 ^a	10.4±0.97 ^{ab}

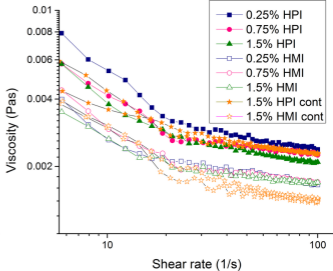
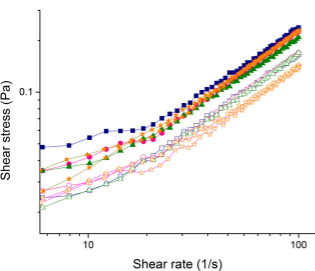
810 Values followed by different letters indicate significant differences ($p \leq 0.05$)

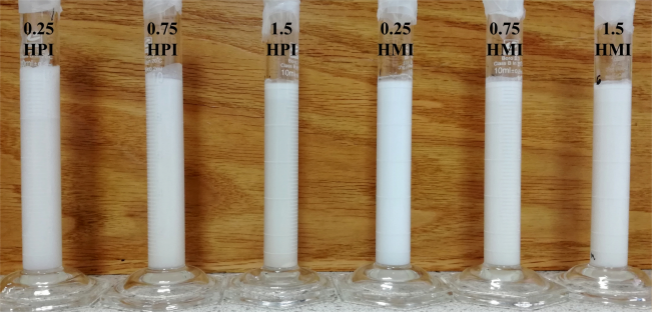
Highlights

- Hemp proteins were isolated using two different isolation procedures
- Alkali (HPI) and salt (HMI) extracted proteins differ in emulsifying ability
- HMI emulsions had smaller volume mean droplet diameter than HPI emulsions
- Microscopy indicated occurrence of bridging flocculation in HPI emulsions
- 1.5% HPI emulsions had the best stability due to formation of floccules network









0.25
HPI

0.75
HPI

1.5
HPI

0.25
HMI

0.75
HMI

1.5
HMI

