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1 **NANO-/MICROSTRUCTURE OF EXTRUDED SPIRULINA/STARCH FOAMS IN**
2 **RELATION TO THEIR TEXTURAL PROPERTIES**

3
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20 **Abstract**

21 This work reports on an in-depth characterization of the nano- and microstructure of
22 extruded starch foams loaded with the microalga *Spirulina* (1, 5 and 10 wt.%), as well as the
23 implications of *Spirulina* incorporation on the textural properties of the foams. Due to the
24 gelatinization process occurring during extrusion, the crystalline and lamellar structures
25 originally present in the starch granule were disrupted, resulting in very amorphous foams.
26 Moreover, the crystalline structure of the fatty acids present in the raw microalga was lost
27 during processing. The presence of *Spirulina* intracellular components induced the formation
28 of more thermally-stable V-type crystallites through complexation with amylose, hence
29 producing slightly more crystalline foams ($X_c \sim 5-9\%$) than the pure extruded starch (X_c
30 $\sim 3\%$). This affected the microstructure of the hybrid foams, which showed more densely
31 packed and well-connected porous structures. Microstructural changes had an impact on the
32 texture of the foams, which became harder with greater *Spirulina* loadings. The foams
33 underwent very limited re-crystallization upon storage, which was further reduced by the
34 presence of *Spirulina*. Interestingly, the free fatty acids from *Spirulina* re-crystallized and
35 the resistant starch content in the 10% *Spirulina* foam increased, which could potentially be
36 interesting from a nutritional perspective. These results show the potential of extrusion
37 cooking to produce healthier snack foods and highlight the suitability of advanced
38 characterization tools such as neutron tomography and small angle X-ray scattering to
39 investigate food structure.

40

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43 **Keywords:** microalgae; SAXS; WAXS; neutron tomography; gelatinization

44

45 **1. Introduction**

46 Extrusion cooking is one of the processing methods used in the food industry to produce
47 healthier snack foods, which is increasingly being used to replace conventional technologies
48 such as frying. The process is based on the application of high temperature, pressure and
49 shear stress, resulting in major structural changes (such as starch gelatinization, protein
50 denaturation, formation of amylose-lipid complexes and degradation of bioactive
51 substances) in the raw materials added to the extruder (Camire, Camire, & Krumhar, 1990).
52 After the cooking step, the melt fluid is shaped into the desired form in the die, thus yielding
53 the final product. The properties of the manufactured products can be adjusted by modifying
54 the processing parameters (such as feed moisture, temperature profile, screw speed and feed
55 rate) (Ding, Ainsworth, Plunkett, Tucker, & Marson, 2006; Ding, Ainsworth, Tucker, &
56 Marson, 2005; Hashimoto & Grossmann, 2003) and by selecting suitable raw materials.
57 Cereal flours, rich in starch, are typically the main component in the formulations used to
58 produce extruded snacks, since they present good processability and enable the production
59 of snacks with acceptable physical characteristics, i.e. an adequate expansion coefficient,
60 hardness and density (Guy, 2001; Launay & Lisch, 1983). However, they present low
61 nutritional value due to their relatively limited protein content and deficiency in certain
62 essential amino acids (Bressani, Wilson, Béhar, & Scrimshaw, 1960; Howe, Jansen, &
63 Gilfillan, 1965). In order to improve their nutritional profile, starch-based snack foods have
64 been enriched with several protein-rich ingredients such as legumes and dairy (Anton, Gary
65 Fulcher, & Arntfield, 2009; Onwulata, Konstance, Smith, & Holsinger, 2001; Rampersad,
66 Badrie, & Comissiong, 2003; Veronica, Olusola, & Adebowale, 2006). However, the
67 incorporation of these additional ingredients into the extrusion formulations does not only
68 have an impact from the nutritional perspective, but also has technological implications,
69 since the processing parameters need to be adjusted and the structural, textural and sensory

70 properties of the extruded snacks may be affected (Lazou & Krokida, 2010; Pastor-Cavada,
71 et al., 2011).

72
73 Besides demanding products with added nutritional value and health benefits, consumers
74 are increasingly searching for more natural ingredients, such as algae-based products. In
75 this context, microalgae are gaining a great interest in the food industry for their use as
76 nutritional supplements and food additives (Chacón-Lee & González-Mariño, 2010;
77 Vigani, et al., 2015). Some microalgae species have been used in human nutrition for many
78 centuries and the interest lays in their high protein content, lipid profile, with relatively high
79 amounts of $\omega 3$ and $\omega 6$ fatty acids, and the presence of bioactive compounds such as
80 carotenoids, phycobilins, polysaccharides, vitamins, and sterols (Chacón-Lee, et al., 2010;
81 Spolaore, Joannis-Cassan, Duran, & Isambert, 2006). In particular, *Spirulina*, filamentous
82 blue-green microalgae or cyanobacteria, are one of the most widely used species in the
83 healthy-food market due to their large content in proteins, γ -linolenic acid and other
84 compounds such as vitamins (mainly B vitamins), minerals, pigments (phycocyanin, β -
85 carotene and xanthophyll), γ -tocopherol and phenolic compounds, which confer these
86 microalgae high antioxidant activities (Khan, Bhadouria, & Bisen, 2005; Miranda, Cintra,
87 Barros, & Mancini-Filho, 1998). Since these microalgae have received the GRAS
88 (Generally Recognized As Safe) certification, they have already been incorporated into
89 different types of food products, such as shakes for elderly (Santos, Freitas, Moreira,
90 Zanfonato, & Costa, 2016), pasta (De Marco, Steffolani, Martínez, & León, 2014; Fradique,
91 et al., 2010; Özyurt, et al., 2015) and extruded snacks (Joshi, Bera, & Panesar, 2014;
92 Bárbara Franco Lucas, de Morais, Santos, & Costa, 2018; Barbara Franco Lucas, Morais,
93 Santos, & Costa, 2017; Morsy, Sharoba, & Hem, 2014). Although most of the existing
94 literature has focused on the nutritional aspects, the incorporation of *Spirulina* may have an

95 impact on the structural changes undergone by food during processing. This may be
96 especially important in the case of starchy foods, whose processing is based on the
97 occurrence of a phenomenon known as gelatinization. Native starch granules display a
98 hierarchical structure consisting of alternating amorphous and semi-crystalline growth rings
99 of 120–400 nm thickness. At the same time, the semi-crystalline rings are formed by a
100 lamellar structure of alternating crystalline and amorphous regions with a repeat distance
101 of 9–10 nm (Blazek & Gilbert, 2011). When processed at high temperature and moisture
102 conditions, starch is known to undergo a gelatinization process in which its hierarchical
103 structure is disrupted. The gelatinization process is initiated by the access of water towards
104 the interior of starch granules, which swell and absorb water, followed by the disruption of
105 the crystalline and lamellar structure and the leakage of amylose from the starch granules
106 to form a continuous gel. In fact, it has been recently demonstrated that *Spirulina*
107 microalgae interfere with the gelatinization process of starch and as a result, lipid-amylose
108 complexes can be formed, promoting the crystallisation of V-type structures (Martínez-
109 Sanz, Fabra, G. Gómez-Mascaraque, & López-Rubio, 2018). Thus, the incorporation of
110 *Spirulina* into starch-based snack foods is not only expected to impact the nutritional value,
111 but also to affect the structural properties of the obtained products. In this study we have
112 performed a detailed investigation of the structural effects of the addition of *Spirulina* into
113 corn starch foams produced by means of extrusion cooking. For the first time a combination
114 of advanced characterisation techniques (ranging from the nanometre to micrometre length
115 scale), such as X-ray scattering (SAXS/WAXS) and neutron tomography, together with
116 scanning electron microscopy (SEM), were used to understand the structural changes
117 occurring in extruded starch foams as result of the addition of *Spirulina* microalgae. Some
118 of the main advantages of using scattering and tomography techniques are related to their
119 non-destructive character and their ability able to provide structural information on native

120 samples, since they minimise experimental artefacts derived from sample treatment.
121 Furthermore, the use of neutrons is particularly interesting for the analysis of food systems
122 by means tomography due to the higher contrast of neutrons with hydrogens (abundant in
123 food components), as compared with X-rays. These results are highly relevant in the
124 development of new food products, since structural changes are expected to affect the
125 sensory properties and digestibility of the produced snacks.

126

127 **2. Materials and methods**

128 **2.1 Materials**

129 Commercial corn starch flour (19% amylose) was purchased locally (Sweden). *Spirulina*
130 *platensis*, in the form of dry powder, was kindly donated by Dr. Acién from the University
131 of Almeria (Spain). The composition of the raw *Spirulina* in terms of total carbohydrate,
132 protein, ash, lipids and polyphenols was determined using standard methods. The total
133 carbohydrate content was determined after sulphuric acid hydrolysis, following the method
134 described in previous work (Martínez-Sanz, et al., 2019). The protein content was
135 determined by the Kjeldahl method, using a factor of 6.25 to convert the nitrogen to protein
136 content (Boussiba & Richmond, 1979). The ash content was determined by the standard
137 method TAPPI T211 om-07. Lipids were estimated by performing a Soxhlet extraction of
138 ca. 1g of dry *Spirulina* with 800 ml of toluene:ethanol 2:1 (v/v) for 24 h. The total phenolic
139 content was estimated by the Folin-Ciocalteu colorimetric assay (Singleton, Orthofer, &
140 Lamuela-Raventós, 1999). The gross chemical composition of the microalgal biomass was
141 determined as: 40.6% ash, 26.7% protein, 15.1% lipid; 8.0% carbohydrate and 0.9%
142 polyphenols.

143

144 **2.2 Preparation of starch and starch-*Spirulina* foams**

145 Pure starch and hybrid foams with three different loadings of *Spirulina* (1, 5 and 10 wt.%
146 dry weight) were prepared by extrusion cooking. The samples were processed with 30 wt.%
147 moisture content to facilitate starch gelatinization during processing and let to equilibrate for
148 24 hours before extrusion. A single-screw extruder (Brabender Duisburg DCE-330,
149 Germany) equipped with a variable speed D-C drive unit was used. The extruder barrel had
150 a 25 mm diameter with length to diameter ratio (L: D) of 20:1. The screw was a 2:1
151 compression ratio with a linearly tapered rod and 20 equidistantly positioned flights. A round
152 circular die with a 2 mm diameter and a length of 90 mm was used. Extrusion temperature
153 was kept constant at 120 °C, 150 °C, 150 °C and 140 °C for zones 1, 2, 3 and 4 respectively.
154 A screw speed of 150 rpm was used for all extrusion processes. The extrudates were
155 collected when the operation conditions were at constant steady state. Each formulation was
156 processed twice. The obtained samples were stored in an equilibrated relative humidity
157 cabinet at 30% RH and 14 °C for two-three days (T0) and one month (TF) prior to their
158 characterization. Expansion index was calculated as the ratio of the diameter of the extruded
159 sample to the diameter of the die.

160

161 **2.3 Attenuated total reflectance (ATR) FT-IR analysis**

162 Small pieces from the produced foams were grounded into powder and the obtained
163 materials were analysed by FT-IR in attenuated total reflectance (ATR) mode using a
164 Thermo Nicolet Nexus (GMI, USA) equipment. The spectra were taken at 4 cm⁻¹ resolution
165 in a wavelength range between 400-4000 cm⁻¹ and averaging a minimum of 32 scans.

166

167 **2.4 Determination of resistant starch content**

168 Resistant starch content was measured using the Megazyme Resistant Starch Assay Kit
169 (Megazyme International, Wicklow, Ireland), based on the AACC method 32-40.01 (AACC,

170 2009). Extruded samples were grounded to a fine powder in a blender. The samples (100
171 mg) were incubated with 4 mL mixture of pancreatic α -amylase and amyloglucosidase for
172 16 h (at 37 °C, 200 strokes/min) in a shaking water bath (Julabo, Seelbach, Germany) to
173 hydrolyze digestible starch to glucose. The reaction was terminated with 4 mL of 99 %
174 ethanol and the indigested resistant starch fraction was obtained after centrifugation (1500
175 \times g, 10 min). The residue was washed twice with 50 % ethanol and solubilized with 2 mL of
176 KOH solution 2 M in an ice bath. The solution was neutralized after 20 min with 8 mL
177 sodium acetate buffer (1.2 M, pH 3.8), and incubated with amyloglucosidase (0.1 mL, 3300
178 U/mL) at 50 °C for 30 min. Aliquots (0.1 mL) of the diluent were taken and mixed with 3
179 mL of glucose oxidase/oxidase reagent (GOPOD) and the mixture was incubated at 50°C
180 for 20 min. Absorbance at 510 nm was measured using a spectrophotometer (Ultrospec,
181 Biochrom, Cambridge, Germany). The samples were measured in duplicates and two
182 different batches of extruded samples were assessed. Resistant starch was calculated as g per
183 100 g of dry sample.

184

185 **2.5 Scanning electron microscopy (SEM)**

186 SEM measurements were conducted on a Hitachi microscope (Hitachi S-4800) at an
187 accelerating voltage of 10 kV and a working distance of 8-16 mm. The pure and composite
188 starch films were cryo-fractured after immersion in liquid nitrogen to observe the cross-
189 sections. The samples were then sputtered with a gold-palladium mixture under vacuum
190 during 3 minutes before their morphology was examined.

191

192 **2.6 Simultaneous small and wide angle X-ray scattering (SAXS/WAXS) experiments**

193 Combined small and wide angle X-ray scattering (SAXS and WAXS, respectively)
194 experiments were carried out in the Non Crystalline Diffraction beamline, BL-11, at the

195 ALBA synchrotron light source ("ALBA synchrotron"). The obtained foams were
 196 characterized shortly after being processed (t_0) and after prolonged storage at 30% RH (t_f),
 197 both in their native dry state and after being hydrated in excess water for at least 24 h. The
 198 energy of the incident photons was 12.4 KeV (an equivalent wavelength, λ , of 1 Å). The
 199 SAXS diffraction patterns were collected by means of a single-photon counting detector,
 200 Pilatus 1 M, with an active area of 168.7×179.4mm², an effective pixel size of 172×172 μm²
 201 and a dynamic range of 20 bits. The sample-to-detector distance was set to 7740 mm,
 202 resulting in a q range with a maximum value of $q=0.189 \text{ Å}^{-1}$. The WAXS diffraction patterns
 203 were collected by means of a 3 CCD detectors, Rayonix LX255-HS, with an active area of
 204 85×255mm², an effective pixel size of 44×44 μm² and a dynamic range of 16 bits. In this
 205 case, the sample-to-detector distance was set to 144.3 mm, corresponding to a maximum q
 206 value of 8.023 Å⁻¹. This detector was tilted with a pitch of 29.38°. The exposure time,
 207 common to both detectors, was 2 s. This time was optimized in preliminary measurements,
 208 in order to maximize the signal to noise ratio while avoiding detector saturation.

209 2.7 SAXS/WAXS data fitting

210 SAXS data were fitted using the Igor NIST analysis macro suite (Kline, 2006) and applying
 211 a mathematical function consisting of a power-law term plus one Gaussian-Lorentzian
 212 peak, similar to that previously reported for several starch samples (A. Lopez-Rubio, et al.,
 213 2007; Salman, et al., 2009):

$$\begin{aligned}
 214 \quad I(q) = & A \cdot q^{-m} + \left[R \cdot \left[I_{max} \cdot \left(1 + \left(\frac{2 \cdot (q - q_{max})}{\Delta q} \right)^2 \right)^{-1} \right] + \left[(1 - R) \cdot \left[I_{max} \cdot \exp \left[-\frac{1}{2} \cdot \right. \right. \right. \right. \\
 215 \quad & \left. \left. \left. \left(\frac{q - q_{max}}{\Delta q} \right)^2 \right] \right] \right] + bkg \quad (1)
 \end{aligned}$$

216 The first term in equation (1) corresponds to the power-law function (where A is a prefactor
217 and m is the power-law exponent) to account for the underlying diffuse scattering, the
218 second and third terms correspond to the Lorentzian and Gaussian functions used to
219 describe the starch lamellar peak (where q_{max} is the peak position, I_{max} is the intensity of
220 the peak, Δq is the full width at half maximum and R is the Lorentzian to Gaussian ratio of
221 the peak shape) and the fourth term accounts for the incoherent background signal.

222

223 WAXS peak fitting was performed in Igor, following the procedure described in a previous
224 work (Amparo Lopez-Rubio, Flanagan, Gilbert, & Gidley, 2008). The obtained values
225 from the fitting coefficients are those that minimize the value of Chi-squared, which is
226 defined as:

$$227 \chi^2 = \sum \left(\frac{y - y_i}{\sigma_i} \right)^2 \quad (2)$$

228 where y is a fitted value for a given point, y_i is the measured data value for the point and
229 σ_i is an estimate of the standard deviation for y_i . The curve fitting operation is carried out
230 iteratively and for each iteration, the fitting coefficients are refined to minimize χ^2 . The
231 crystallinity index X_C was determined from the obtained fitting results by applying the
232 following equation:

$$233 X_C(\%) = \frac{\sum A_{Crystal}}{A_{Total}} \times 100 \quad (3)$$

234 where A_{Total} is the sum of the areas under all the diffraction peaks and $\sum A_{Crystal}$ is the sum of
235 the areas corresponding to the crystalline peaks.

236

237 **2.8 Neutron tomography**

238 **2.8.1 Data acquisition**

239 Neutron tomography experiments were performed at the cold neutron imaging station,
240 IMAGINE (Ott, Loupiac, Désert, Hélyary, & Lavie, 2015) at the ORPHEE reactor of the
241 French national neutron facility, the *Léon Brillouin Laboratory* (Saclay, France). The 4 types
242 of foams produced by extrusion were scanned: pure starch (n=2), 1% *Spirulina* (n=3), 5%
243 *Spirulina* (n=3) and 10% *Spirulina* (n=3), where n equals the number of scanned samples.
244 The scanning parameters were the following: exposure time, 60s and number of projections,
245 180; number of dark and flat-field projections, 10; sample-to-detector distance, 50 mm; field
246 of view (FOV), 47mm × 40mm; effective isotropic pixel size, 18.5 μm. The characteristics
247 of the utilized detector setup were the following; a Neo sCMOS (ANDOR) camera equipped
248 with a sCMOS sensor (effective number of pixels, 2560 x 2160 pixels²; physical pixel size:
249 6.5 x 6.5 μm²) coupled to a 50μm thick gadolinium scintillator (RC Tritec, 2014), followed
250 by optimal light magnification using a 35 mm objective (Canon EF 35 mm f/2.0 IS USM).
251 Furthermore, the camera was cooled down to -30°C, to reduce noise. Pieces of few
252 centimeters of the extruded samples (T0) were placed in an aluminum film inside a
253 cylindrical aluminum cell placed in front of the scintillator on the rotating table.

254

255 **2.8.2 Image pre-processing and 3D reconstruction**

256 An in house-developed macro from the IMAGINE imaging station and implemented in
257 ImageJ (Rueden, et al., 2017) was used to normalize the projection images, remove noise
258 and correct the tilt-angle. Hereinafter, the SYRMEP Tomo Project (STP) (Brun, et al., 2017)
259 was used to reconstruct a 3D-map of the linear attenuation coefficient (LAC) [cm⁻¹] of the
260 neutrons for the scanned samples. The LAC was reconstructed on 32-bit grey-level scale.

261

262 **2.8.3 Image post processing**

263 The reconstructed slices contained two well-separated peaks, associated to air and
264 starch/*Spirulina*. Standard grey-level thresholding was shown to be unable to accurately
265 segment the foams, due to the large variance in grey-scale values between the thinner and
266 thicker cell wall structures, as shown in Figure S1A. Therefore, an automated (supervised)
267 pixel-level classification method (based on intensity, edge and texture with $\sigma = 0.7$)
268 implemented in the Ilastik software framework (Sommer, Straehle, Koethe, & Hamprecht,
269 2011) was applied. One slice from each type of sample group was used for combined
270 training, wherein after the automated segmentation method was applied to all respective
271 slices of the given data sets, as shown in Figure S1B.

272

273 In extrusion, both the material composition and the utilized processing parameters determine
274 the pore size and cell wall thickness. Consequently, samples with a very thin outer cell wall
275 thickness tend to rupture easily, as shown in (Bárbara Franco Lucas, et al., 2018). The open
276 path between the air outside and inside the sample respectively hampers more advanced
277 image analysis, especially so called skeletonization, followed by the calculation of the
278 Connectivity Density (Conn.D). Therefore, an image processing protocol implemented in
279 ImageJ, which successfully closes the open path at broken cell walls, was developed and
280 applied to all of the hereby addressed data sets, as shown in Figure S1.

281

282 **2.8.4 Quantitative 3D image analysis**

283 Quantitative 3D image analysis was performed using the Pore3D software library (Brun, et
284 al., 2010) on Representative Volume of Interest (RVI) of the respective data sets, following
285 an RVI-test, as further described in the Supplementary Material section ‘Representative
286 Volume of Interest (RVI) test’. The following image parameters were quantified: 1) Volume
287 percentage (Vol. Perc.) of the air part, also referred to as ‘porosity’, 2) Specific Surface Area,

288 computed as the quotient between the surface area of the object divided with the total volume
289 of the VOI as further described by (Parfitt, et al., 1987), 3) Structure Thickness (St.Th.) of
290 the extruded foams, hereinafter referred to as ‘cell wall thickness’ and 4) Conn.D, as well as
291 5) pore and 6) throat sizes, based on skeletonization, as further described in (Furlan, et al.,
292 2018), hereinafter referred to as Method 1. For objective comparison, the pore size was also
293 estimated on the individually separated macro pores, by applying a ‘Distance Transform
294 Watershed 3D’ algorithm in the MorphLibJ package (Legland, Arganda-Carreras, &
295 Andrey, 2016) in ImageJ, using the following parameters: Borgfors(3,4,5) and
296 ‘normalization dynamic’: 6.0, followed by quantification using Pore3D, hereinafter referred
297 to as Method 2. However, only Method 1 is capable of estimating both the pore and the
298 throat size.

299

300 **2.8.5 3D Visualization**

301 The 3D volume renderings were performed using the Drishti package for visualization of
302 volumetric datasets (Limaye, 2012) and using ‘Volume viewer’ in ImageJ (Rueden, et al.,
303 2017) for visualization of the ‘local thickness’ (Hildebrand & Rüeggsegger, 1997), as well as
304 the individually separated macro pores, as separated by ‘Wathershedding’.

305

306 **2.9 Uniaxial Compression testing**

307 Uniaxial compression tests were performed on an Instron material testing device (model
308 5542, Instron, Norwood, MA, USA). Extruded samples were carefully cut to cylindrical
309 shapes with a height of ca. 1 cm with the help of a sharp scalpel and placed between flat
310 metal surfaces covered with emery paper to avoid slippage. A 3 cm cylindrical probe was
311 used to compress the samples until fracture, using a 500 N load cell (load values during
312 compression were in the range of 125-200 N) at a crosshead speed of 1 mm/s. At least 5

313 replicates of each type of extruded sample were measured. Force (N) and distance (mm)
314 were converted to true stress (σ_T) and true strain (ε_T) using equations 1 and 2 by the
315 instrument software Blue Hill.

$$316 \quad \sigma_T = \frac{F(t)(h_0 - \Delta h(t))}{\pi r_0^2 h_0} \quad (4)$$

$$317 \quad \varepsilon_T = \ln\left(\frac{h_0}{h_0 - \Delta h(t)}\right) \quad (5)$$

318 Where h_0 is the initial height of the sample, $\Delta h(t)$ is the change in height during
319 compression and r_0 is the initial radius of the sample. The compression modulus was
320 obtained as the slope of the initial linear zone of the true stress vs. true strain plots. Mean
321 values of the critical stress and the compression modulus were compared by Tukey and
322 Fisher tests with a confidence interval of 95% using Excel Stats software.

323

324 **3. Results and Discussion**

325 **3.1 Nanostructure of the as-processed foams**

326 In this work, different contents of *Spirulina* (1, 5 and 10 wt.%) were incorporated into corn
327 starch formulations to produce foams by means of extrusion cooking. To investigate the
328 effects of the added microalgae on the structure of the foams, the samples were characterized
329 by wide angle and small angle X-ray scattering (WAXS and SAXS). These techniques are
330 very powerful tools to extract information on the lamellar and crystalline arrangement of
331 starch, as widely reported in the literature (Blazek, et al., 2011; P. Jenkins, et al., 1994).

332

333 The WAXS patterns of the raw materials and the as-processed foams (T0) are shown in
334 Figure 1A. As observed the raw corn starch presented WAXS patterns typical from the A-
335 type crystalline allomorph, characterized by the main diffraction peaks located at $2\theta = 15.1^\circ$,
336 17.0° , 18.1° and 23.1° . On the other hand, the raw *Spirulina* presented several sharp peaks,

337 of which the most intense ones were located at 2θ values of ca. 20.6° , 29.1° , 29.6° , 30.9° ,
338 31.7° , 38.0° , 41.8° and 47.1° the most intense ones. The appearance of these peaks has been
339 previously associated to the presence of crystallized fatty acids in this microalga (Martínez-
340 Sanz, et al., 2018), although they have not been assigned to any particular crystalline
341 allomorph yet. In fact, the crystallinity of the raw *Spirulina* is very close to the total amount
342 of lipids determined in the sample (ca. 15%), thus supporting the hypothesis of crystallized
343 fatty acids present in the native microalga. After the extrusion cooking process, a very
344 amorphous structure, showing a WAXS pattern resembling the characteristic pattern of the
345 V-type crystalline allomorph, was observed for the pure starch foam. Interestingly, the
346 position and relative intensity of the scattering peaks did not correspond to any of the
347 reported pure V-type crystalline structures. This is in agreement with a previous study which
348 demonstrated that corn starch samples shortly after being processed by extrusion presented
349 a non-stable structure consisting of a mixture of different crystalline allomorphs (Benito-
350 González, López-Rubio, & Martínez-Sanz, 2019). The samples containing *Spirulina* showed
351 scattering patterns similar to those reported for V-type starches and displayed greater
352 crystallinity values than the pure starch foam (cf. Table 1). The application of heat/moisture
353 treatments is known to induce the formation of V-type crystalline structures in corn starch
354 due to the complexation of the fatty acids and phospholipids naturally present in the raw
355 cereal with the amylose chains from starch (Kugimiya, Donovan, & Wong, 1980; Zobel,
356 1988). Our results indicate that the formation of V-type crystallinity was clearly favoured by
357 the presence of *Spirulina*. The weak cell walls from *Spirulina* have been seen to break down
358 very easily, even with only mild stirring (Fabra, et al., 2017; Martínez-Sanz, et al., 2018),
359 thereby enabling the release of the components contained within the cells, such as lipids and
360 proteins. Furthermore, a previous study showed that the formation of V-type crystallites as
361 the result of lipid-amylose complexation takes place when subjecting corn starch to a high-

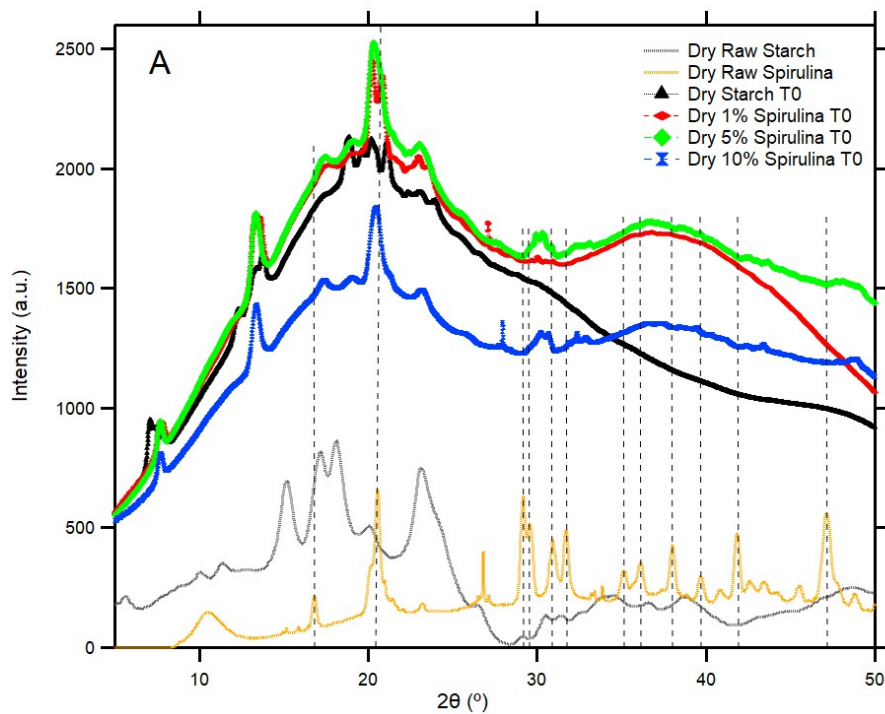
362 humidity thermal treatment in the presence of *Spirulina* (Martínez-Sanz, et al., 2018). Thus,
363 it can be assumed that the shearing effect during the extrusion cooking process promoted
364 cell wall disruption and facilitated the release of intracellular components which were able
365 to form complexes with amylose. Moreover, the WAXS results suggest that the formed V-
366 type complexes were more thermally stable than the original A-type crystallites present in
367 the raw corn starch, hence producing slightly more crystalline samples after the extrusion
368 cooking process. This is in agreement with previous work which reported that a small
369 fraction of these V-type crystallites was resistant at 110 °C (Martínez-Sanz, et al., 2018).
370 Accordingly, the WAXS results suggest that, in parallel with the gelatinization process, the
371 presence of *Spirulina* cell wall components favoured the formation of V-type complexes,
372 which were more thermally stable than the original A-type crystallites present in the raw
373 corn starch. This resulted in the production of slightly more crystalline samples after the
374 extrusion cooking process. It should be noted that none of the peaks detected in the hybrid
375 foams corresponded to those assigned to the raw microalgae, suggesting that the fatty acid
376 crystals were melted upon processing.

377
378 While WAXS is able to provide information on the crystalline structure of starch, SAXS is
379 mostly used to understand the lamellar arrangement. The SAXS patterns from the raw
380 materials and the obtained foams (T0) are shown in Figure 1B. As observed, only the native
381 starch presented a weak and broad lamellar peak centred at ca. 0.071 \AA^{-1} in the native dry
382 state, which corresponds to a lamellar repeat distance of ca. 8.8 nm. The identification of the
383 starch lamellar peaks is typically much more evident in hydrated samples. This may be due
384 to the plasticization of amylopectin branch points, increasing the mobility of the amylopectin
385 double helices, which are then able to align in layers (i.e. lamellae), or simply to increased
386 contrast between the amorphous and the semicrystalline lamellae as a result of the easier

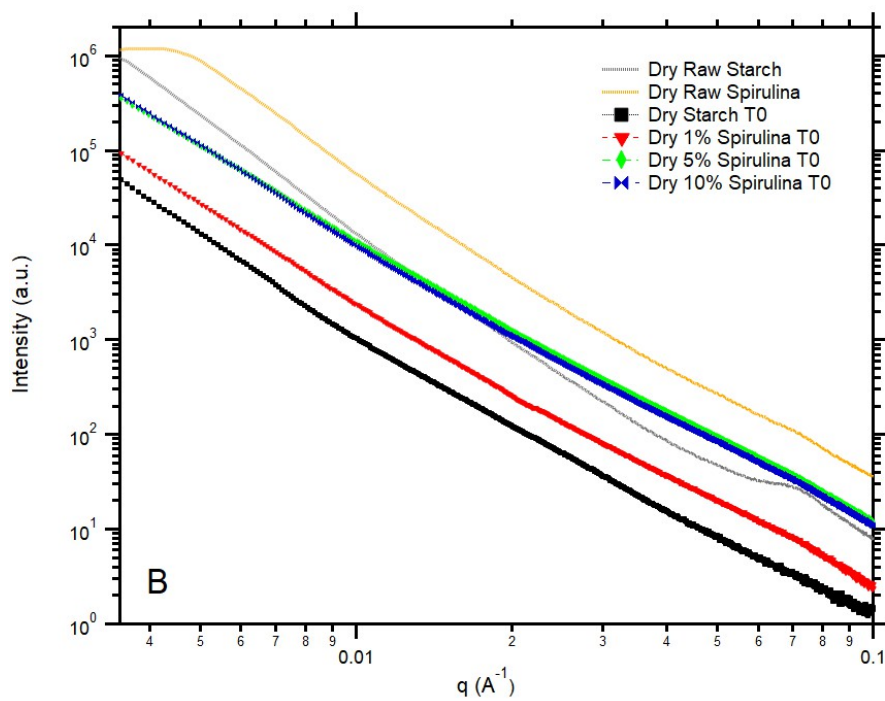
387 access of water towards the amorphous part of the granules. On the contrary, in the dry state
388 the amylopectin branch points become less flexible and as a result, their arrangement into
389 lamellae is more limited, although the correlations within the double helices persist (giving
390 rise to the WAXS peaks). Accordingly, the lamellar peak detected in the raw starch became
391 more intense and sharp in the hydrated state and its position shifted towards ca. 0.063 \AA^{-1}
392 (cf. Figure 1C), corresponding to a lamellar repeat distance of ca. 9.9 nm, in agreement with
393 the lamellar distances previously reported for other corn starches (P. J. Jenkins, Cameron, &
394 Donald, 1993; Kuang, et al., 2017; Martínez-Sanz, et al., 2018). This peak was absent in the
395 patterns from the foams, pointing out towards a disruption of the lamellar structure during
396 the extrusion cooking process, in line with the WAXS results and in agreement with previous
397 works (Benito-González, Martínez-Sanz, & Lopez-Rubio, 2018). This was further
398 confirmed by the change in the power-law exponent from the low q region ($0.003\text{-}0.005 \text{ \AA}^{-1}$
399 1), which decreased from ca. 4 for the raw starch (corresponding to the Porod's ideal two-
400 phase system and typically reported for native starches (Chanvrier, et al., 2007; Lopez-
401 Rubio, Htoon, & Gilbert, 2007)) to 2.3-2.7 for the foams (corresponding to mass fractal
402 structures). Power-law exponents close to 2 have been reported for gelatinized starches
403 where the lamellar order was completely lost (Lopez-Rubio, et al., 2007), while slightly
404 higher exponents, similar to the values here reported, have been estimated for melt-
405 compounded cellulose-starch samples (Benito-González, et al., 2018). Extrusion of starch
406 samples in excess water has been reported to completely destroy the lamellar structure
407 (Waigh, Gidley, Komanshek & Donald, 2000), whereas a weak 9-10 nm lamellar peak has
408 been seen to remain when the extrusion process was carried out at relatively low water
409 contents (Chanvrier, Uthayakumaran, Appelqvist, Gidley, Gilbert & López-Rubio, 2007).
410 Our results support the fact that the lamellar structure of starch was extensively disrupted

411 with the addition of 30 wt.% moisture and the conditions used for the extrusion cooking
412 process, both for the pure starch and the blends with the added *Spirulina*.

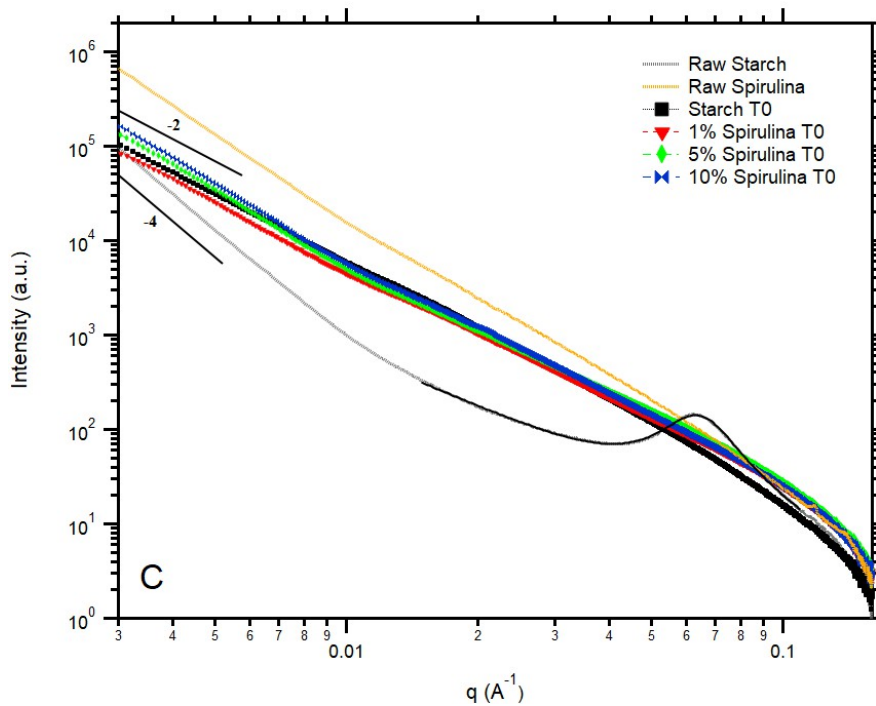
413



414



415



416

417 **Figure 1.** (A) WAXS patterns from the raw starch and *Spirulina* and from the as-processed
 418 foams (T0). Curves from raw starch and *Spirulina* have been offset for clarity. (B) SAXS
 419 patterns of the raw starch and *Spirulina* and the foams after being processed (T0) in their
 420 native dry state and (C) SAXS patterns from the fully hydrated samples. Dotted lines
 421 correspond to the experimental data, whereas solid lines correspond to the fitting curves.

422

423 **Table 1.** Power-law exponents, lamellar peak parameters and crystallinity (X_C) for the raw
 424 materials and the foams after being processed (T0). The lamellar peak parameters were
 425 obtained by fitting the experimental SAXS data of the hydrated samples to the sum function
 426 of a power-law plus a Gaussian-Lorentzian peak and crystallinity was obtained from the
 427 peak fitting of the WAXS patterns from the dry samples. Standard deviation values on the
 428 last digit are shown in parentheses for the fitting parameters.

Power-law exponent	Lamellar peak position (\AA^{-1})	Width of the lamellar peak (\AA^{-1})	Lamellar repeat distance (\AA)	X_C (%)
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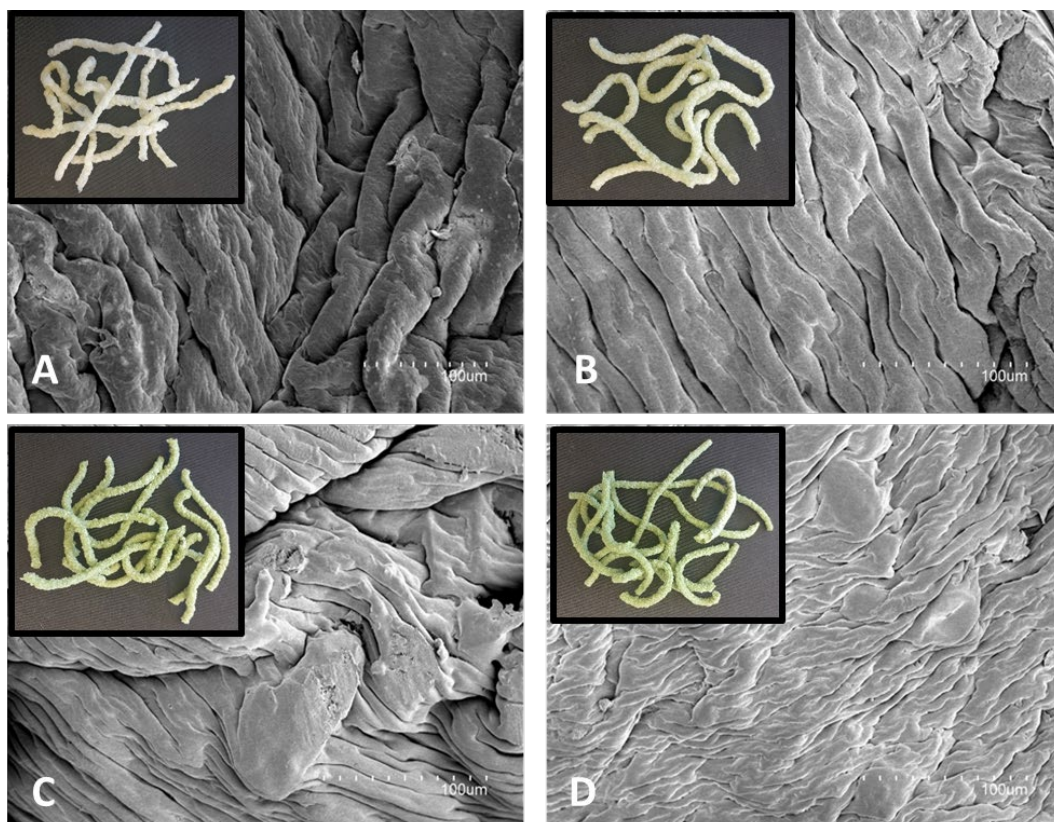
Raw <i>Spirulina</i>	3.24 (1)	---	---	---	14.98
Raw Starch	2.009 (3)	0.06335 (1)	0.02543 (5)	9.92	42.71
Starch	2.49 (1)	---	---	---	3.22
1% <i>Spirulina</i>	2.34 (5)	---	---	---	4.59
5% <i>Spirulina</i>	2.52 (1)	---	---	---	6.62
10% <i>Spirulina</i>	2.73 (1)	---	---	---	8.72

429

430 **3.2 Microstructure and mechanical properties of the as-processed foams**

431

432 To understand the effects of *Spirulina* addition at a larger size range, the microstructure of
433 the obtained foams was also investigated by SEM and neutron tomography. Representative
434 SEM images, together with pictures showing the visual appearance of the foams, are shown
435 in Figure 2. As observed, the addition of *Spirulina* produced foams with a greenish
436 coloration, which was more evident for loadings higher than 5 wt.%. Furthermore, as
437 evidenced by the SEM images, more densely packed structures were present in the foams
438 containing higher *Spirulina* loadings. Such effect that was also observed macroscopically.
439

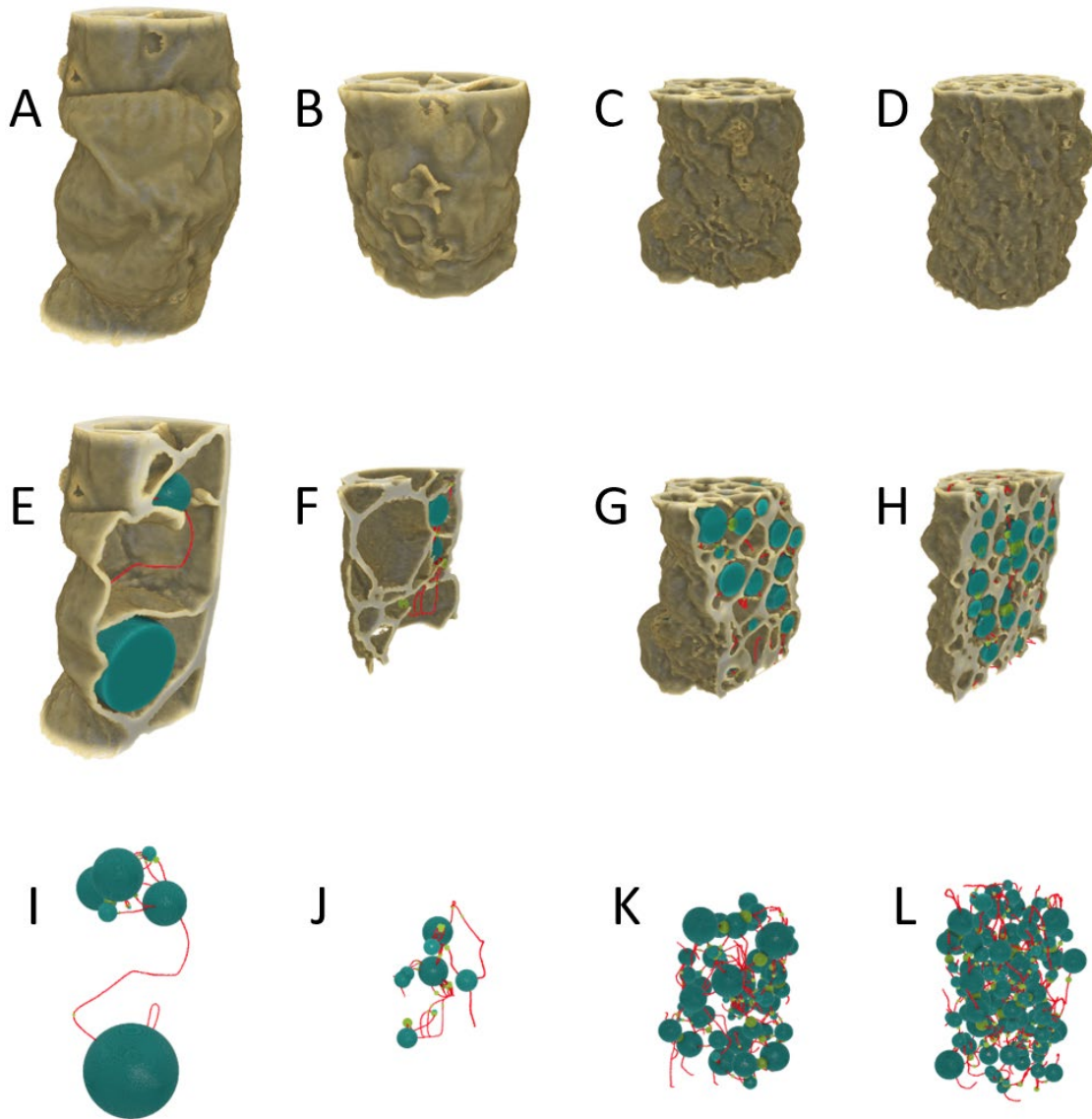


440
441 **Figure 2.** SEM micrographs from the as-processed foams (T0): (A) Pure starch, (B) 1%
442 *Spirulina*, (C) 5% *Spirulina* and (D) 10% *Spirulina*. Image inserts show the visual
443 appearance of the foams.

444

445 An in-depth 3D characterization of the microstructure from the different foams was carried
446 out by means of neutron tomography. The porous structure of the extruded foams could be
447 clearly resolved and visualized by this technique, as shown in Figure 3 and Figure 4. The
448 visual results were also supported with quantitative image analysis and the obtained
449 parameters are summarized in Table 2. As observed, loadings higher than 5 wt.% *Spirulina*
450 led to a reduction in the porosity. The specific surface area and connectivity density
451 (Conn.D) increased with increasing concentrations of *Spirulina*, indicating that more well-
452 connected foams were formed. In addition, the cell wall thickness decreased with the
453 addition of *Spirulina*, although there was no clear trend with the microalga concentration
454 (cf. Table 2 and Figure 4). However, no significant differences in pore size or throat size
455 could be measured, due to the large heterogeneity of the samples, which led to large standard
456 deviations. These data have thus been supported by pore and throat size probability
457 distributions, calculated with method 1 and 2, as described in the Supplementary Material
458 section (Figures S4, S5 and S6), indicating that the pore size (mean and standard deviation)
459 decreased with increasing concentration of *Spirulina*.

460



461

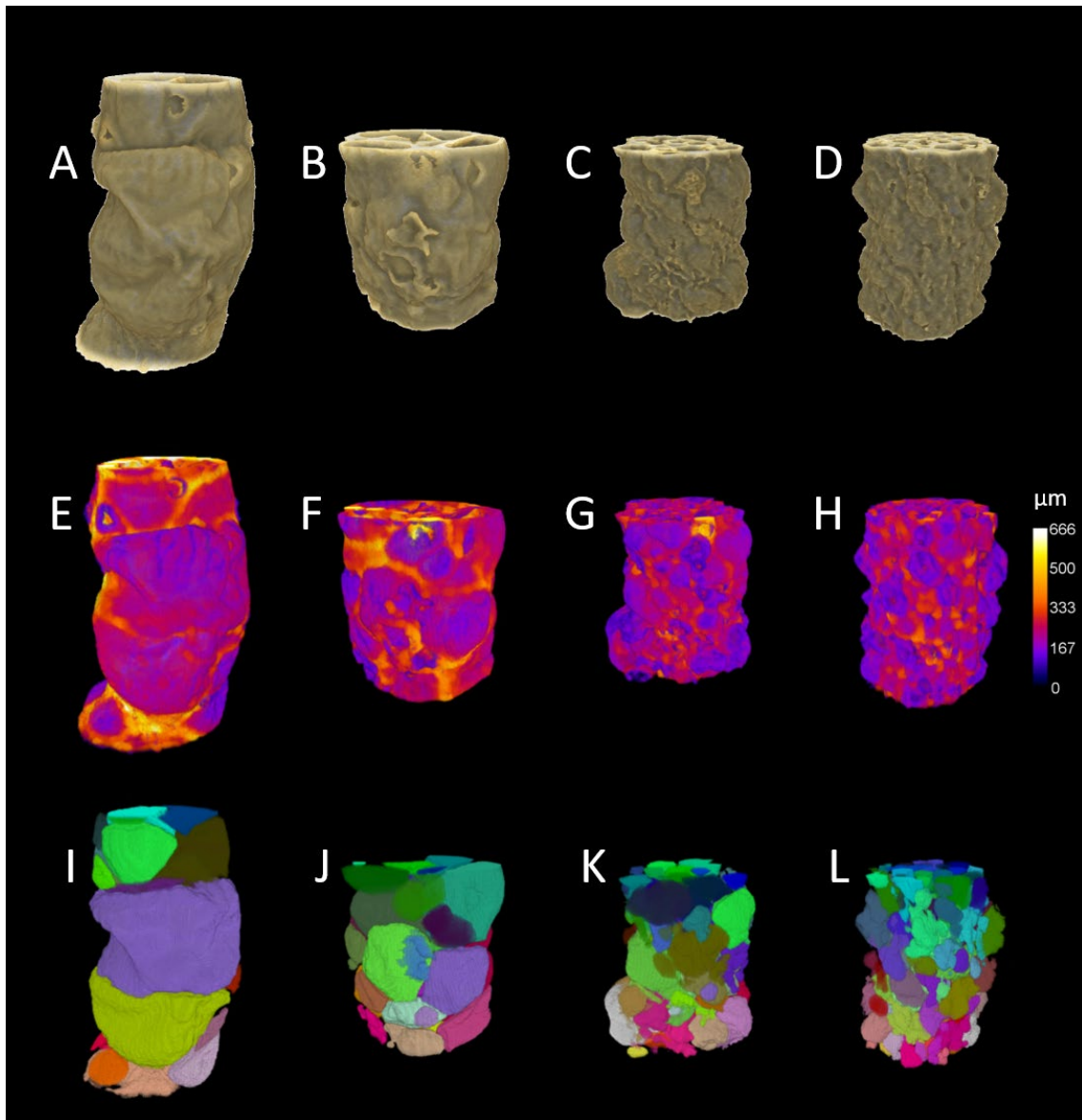
462 **Figure 3.** Visualization of the microstructure of the as-processed foams (T0) determined by
 463 neutron tomography. (A-D) show volume renderings of the extruded foams (in light brown),
 464 (E-H) with applied virtual cut planes. (E-H and I-L) show the longest connected image
 465 skeleton (in red) throughout the pore space, with maximal inscribed blobs in the porous part
 466 at the nodes (dark green) and at the throats (light green) along the longest skeleton, as
 467 described by ‘Method 1’. The height of the samples were 11.2 mm and 7.1 mm for the pure
 468 starch and the hybrid starch-*Spirulina* samples, respectively. (A, E, I) Pure starch, (B, F, J)
 469 1% *Spirulina*, (C, G, K) 5% *Spirulina* and (D, H, L) 10% *Spirulina*.

470

471 **Table 2.** Extracted image descriptors obtained by applying quantitative 3D image analysis
 472 on selected volumes of interests (VOIs), with a representative VOI height of 11.2 mm and
 473 7.1 mm for the pure starch and the hybrid starch-*Spirulina* samples, respectively. See Figures
 474 S2 and S3 for details of how the representative volume of interest was selected.

	Porosity (%)	Specific Surface Area (mm ⁻¹)	Cell wall thickness (μm)	Conn.D (mm ⁻³)	Pore Size (μm) Method 1 (Method 2)	Throat Size (μm) Method 1
Starch	67.6 ± 3.1	1.7 ± 0.1	234.0 ± 18.4	0.06 ± 0.00	1109.0 ± 1068.9 (1210.9 ± 922.1)	161.2 ± 114.9
1% <i>Spirulina</i>	64.8 ± 1.2	2.5 ± 0.1	199.1 ± 0.2	0.07 ± 0.00	707.7 ± 527.4 (859.9 ± 600.6)	178.0 ± 133.2
5% <i>Spirulina</i>	57.2 ± 2.4	3.8 ± 0.1	165.1 ± 1.7	0.34 ± 0.05	652.8 ± 279.1 (671.3 ± 247.9)	152.9 ± 91.9
10% <i>Spirulina</i>	50.3 ± 5.0	4.2 ± 0.2	180.1 ± 25.2	0.79 ± 0.09	528.5 ± 198.4 (574.4 ± 196.3)	159.0 ± 92.5

475



476

477 **Figure 4.** (A-D) Volume renderings of the as-processed foams (T0) (in light brown), as
 478 determined by neutron tomography. (E-H) Visualization of the cell wall thickness, which is
 479 colour coded, and given in μm . (I-L) Visualization of the individual pores, following 3D
 480 water shedding and colour labelling, as described by ‘Method 2’. The height of the samples
 481 were 11.2 mm and 7.1 mm for the starch and hybrid starch-*Spirulina* foams, respectively.
 482 (A, E, I) Pure starch, (B, F, J) 1% *Spirulina*, (C, G, K) 5% *Spirulina* and (D, H, L) 10%
 483 *Spirulina*.

484

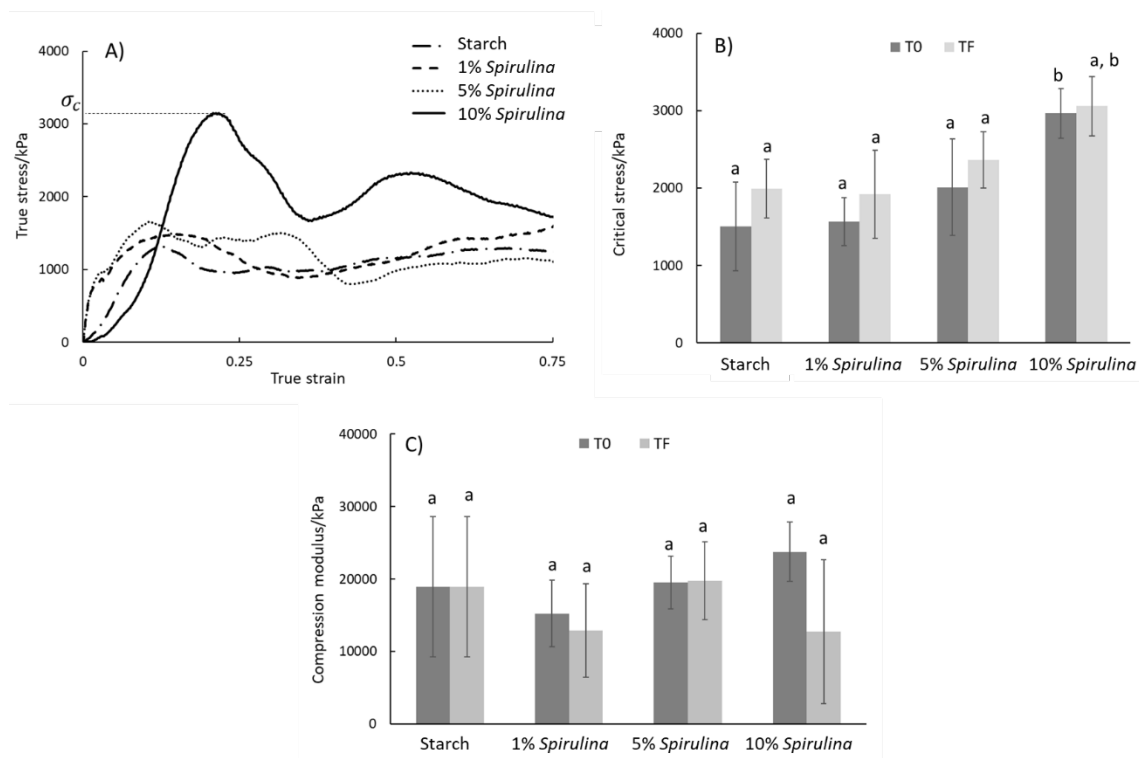
485 The structural changes undergone by starch during the extrusion cooking process were
486 expected to have a strong impact on the mechanical properties of the obtained foams, which
487 in turn are directly linked to their texture. Food texture is a very important attribute from the
488 perspective of consumers' acceptance and thus, it is crucial for the application of these foams
489 as food snacks. The expansion index of the extrudates is directly related to the microstructure
490 of extrudates, and has been shown to be a determining factor for the crunchiness (Joshi, et
491 al., 2014). The expansion index was calculated as described in the materials and methods
492 section and was reduced from 3.1 ± 0.2 for the corn starch to 2.8 ± 0.1 , 2.3 ± 0.1 and $2.2 \pm$
493 0.1 for the hybrid samples containing 1%, 5% and 10 % *Spirulina*, respectively. These results
494 are in agreement with previous works which showed no effect on the expansion index of
495 rice and corn flour mixtures (2:1) with the addition of 0.4-2.6 wt.% *Spirulina* (reporting
496 values which ranged from 2.5 to 4.5 depending on the extrusion conditions) (Bárbara
497 Franco Lucas, et al., 2018; Barbara Franco Lucas, et al., 2017; Morsy, et al., 2014), while
498 higher *Spirulina* loadings (6-8%) have been shown to reduce the expansion index (Tańska,
499 Konopka, & Ruszkowska, 2017). As suggested by the WAXS results, the *Spirulina* cell wall
500 components were able to form complexes with amylose during the extrusion cooking
501 process, leading to the formation of more thermally-stable V-type crystallites and, therefore,
502 producing slightly more crystalline foams with reduced elasticity and thus, limited ability to
503 expand. The reduced expansion index of the 5% and 10% *Spirulina* hybrid foams can be
504 directly linked to their more densely packed structures, as observed by SEM and neutron
505 tomography.

506

507 Compression experiments were also carried out to determine the mechanical properties of
508 the as-processed (T0) foams and typical compression true stress-strain curves are shown in
509 Figure 5A. A critical value of stress (σ_c), defined as the first maximum stress value, could

510 be identified for all the samples, but more clearly for the 10% *Spirulina* foam. After the
511 critical stress it was observed that the samples followed a brittle fracture as the compression
512 test progressed, indicated by the curve pattern with peaks up and down until the sample
513 collapses characteristic of this type of systems (Attenburrow, Goodband, Taylor, & Lillford,
514 1989). As deduced from Figure 5B, only the highest *Spirulina* loading resulted in a
515 significant increase (95% confidence interval) in the critical stress (2966 ± 322 kPa for 10%
516 *Spirulina*, as compared to 1505 ± 573 kPa, 1568 ± 310 kPa and 2011 ± 622 kPa for the pure
517 starch, 1% *Spirulina* and 5% *Spirulina*, respectively). A similar trend of increased hardness
518 has been previously reported with the addition of lower *Spirulina* contents into extruded
519 starch (Barbara Franco Lucas, et al., 2017) and was attributed to the effect of proteins in
520 restricting moisture accessibility towards starch. The slightly greater crystallinity of the
521 samples containing *Spirulina* can also explain the hardness increase. On the other hand, the
522 stiffness of the foams was not significantly affected, as suggested by their compression
523 moduli (15000-25000 kPa) (cf. Figure 5C), which were within the range of previously
524 reported values for extruded starch foams (Robin, et al., 2010). It should be noted that the
525 larger standard deviation values for the pure starch foam reflect a greater structural
526 heterogeneity. As previously discussed, the microstructure of the extruded samples consisted
527 of a porous network which may be described as a solid foam (cf. Figure 2 and Figure 3). The
528 micromechanics of foam deformation during compression starts by cell walls bending
529 elastically, and giving rise to the initial linear region in the stress-strain curves, followed by
530 structural collapse when a critical value of stress is reached (Ashby & Medalist, 1983). Ashby
531 and Gibson demonstrated that the ratio of the bulk density to the density of the cell wall
532 material can be correlated to the critical stress and modulus of foams (Ashby, et al., 1983).
533 The strongest sample (i.e. presenting the highest critical stress) would correspond to that
534 with the greater bulk density and containing thinner cell walls and a higher number of cells

535 (10% *Spirulina*), whilst coarse structures with a lower number of cells would be the weakest
 536 (pure starch and 1% *Spirulina*). The increased hardness of the 10% *Spirulina* foam is is
 537 expected to have some implications on the consumers' acceptance. This will be further
 538 evaluated by sensory analyses in the future.
 539



540
 541 **Figure 5.** (A) Representative true stress vs strain curves of starch-*Spirulina* foams
 542 conditioned at 30% RH. (B) Average critical stress for samples before (T0) and after storage
 543 (TF). (C) Average compression modulus of the extruded samples before (T0) and after
 544 storage (TF). Different letters indicate statistically significant differences between samples
 545 ($p \leq 0.05$).

546
 547 **3.2 Effect of retrogradation upon prolonged storage**

548 After being gelatinized, starch is capable of re-crystallizing upon storage at certain
 549 conditions. During this re-crystallization process, known as retrogradation, the amylose

550 chains gradually re-associate and re-organize to form a different ordered structure (Wang,
551 Li, Copeland, Niu, & Wang, 2015). The impact of the *Spirulina* microalga on the
552 retrogradation process was evaluated by characterizing the obtained foams after storage at
553 30% RH for one month (TF). This would be representative of the properties of the foams
554 during their shelf-life. The nanostructural changes undergone by the samples at TF were
555 investigated by means of WAXS and SAXS. The WAXS patterns from the stored foams are
556 shown in Figure 6A. As observed, the crystalline structure of the foams changed from the
557 non-stable V-type mixture observed shortly after processing the samples (T0) towards a V_A
558 type structure, in agreement with what has been previously reported for other extruded starch
559 samples (Benito-González, et al., 2019; J. J. Van Soest, Hulleman, De Wit, & Vliegenthart,
560 1996). Additionally, two peaks located at ca. 17° and 22°, which are characteristic from the
561 B-type allomorph, were detected, suggesting that a certain fraction of starch re-crystallized
562 into this type of structure. Corn starch has been reported to re-crystallize into a B-type
563 structure when stored at low temperature (Shamai et al. 2003) or high relative humidity
564 conditions (Fu et al. 2013) and similar results have also been reported for corn starch samples
565 processed by means of melt compounding in the presence of 50 wt.% moisture (Benito-
566 González, et al., 2019). The re-crystallization into B-type structures has been related to the
567 rearrangement and association of the outermost short branches of amylopectin (Hoover,
568 Hughes, Chung, & Liu, 2010). The crystallinity values calculated for the foams at TF (cf.
569 Table 3) were very far from the value corresponding to the native starch (cf. Table 1) and
570 indicate that only a small fraction of starch was able to re-crystallize upon storage at
571 30% RH. The pure extruded starch and the hybrid foams presented very similar
572 crystallinities at TF and only the highest *Spirulina* loading gave rise to a slight crystallinity
573 increase compared to the pure starch. Comparing with the values estimated just after
574 processing (T0), the retrogradation process gave rise to a crystallinity increase of 92% in the

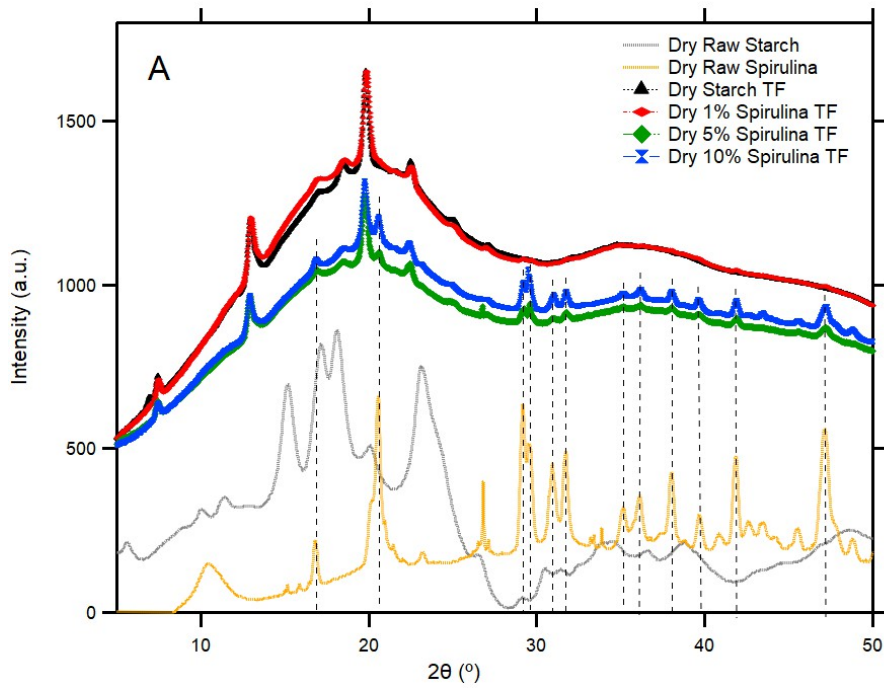
575 case of pure starch and 23% for the 1% *Spirulina* foam, whereas the crystallinity
576 corresponding to the starch fraction was slightly decreased for the 5% and 10% *Spirulina*
577 foams. It is known that greater moisture contents favour starch re-crystallization (J. Van
578 Soest & Knooren, 1997; Wang, et al., 2015) and in particular, the presence of moisture is
579 necessary for the transition of V-type complexes into B-type crystallites ((Kugimiya, et al.,
580 1980; Zobel, 1988). Previous studies have pointed out that the proteins present in *Spirulina*
581 may compete for water binding sites, hindering the hydration of starch granules (Batista, et
582 al., 2012; Barbara Franco Lucas, et al., 2017; Tańska, et al., 2017). These results confirm the
583 fact that the presence of the microalga limited the accessibility of moisture towards the starch
584 molecules and as a result, hindered the re-crystallization process. A similar effect has been
585 attributed to the presence of cellulose (Benito-González, et al., 2019) and other fillers which
586 interact with the hydroxyl groups from starch by the formation of hydrogen bonds (Ferrero,
587 Martino, & Zaritzky, 1994; Ghanbarzadeh, Almasi, & Entezami, 2011). It should be noted
588 that the 5% *Spirulina* and 10% *Spirulina* foams displayed additional weak peaks which
589 corresponded to the most intense scattering peaks from the raw *Spirulina*. This indicates that
590 the free fatty acids contained in the microalga were able to re-crystallize back to their native
591 form upon storage of the foams.

592
593 To confirm the effect of *Spirulina* on the crystallinity of the foams after storage, FT-IR
594 analyses were also carried out and the obtained spectra are shown in Figure S7. The spectral
595 region of 800–1200 cm^{-1} was used to investigate structural differences between samples, as
596 previous studies have suggested that bands in this ‘fingerprint’ region reflect changes in
597 polymer conformation and hydration of processed starches (Capron, Robert, Colonna,
598 Brogly, & Planchot, 2007; Htoon, et al., 2009; Sevenou, Hill, Farhat, & Mitchell, 2002; J. J.
599 G. van Soest & Vliegenthart, 1997). The band at 1022 cm^{-1} has been seen to increase in

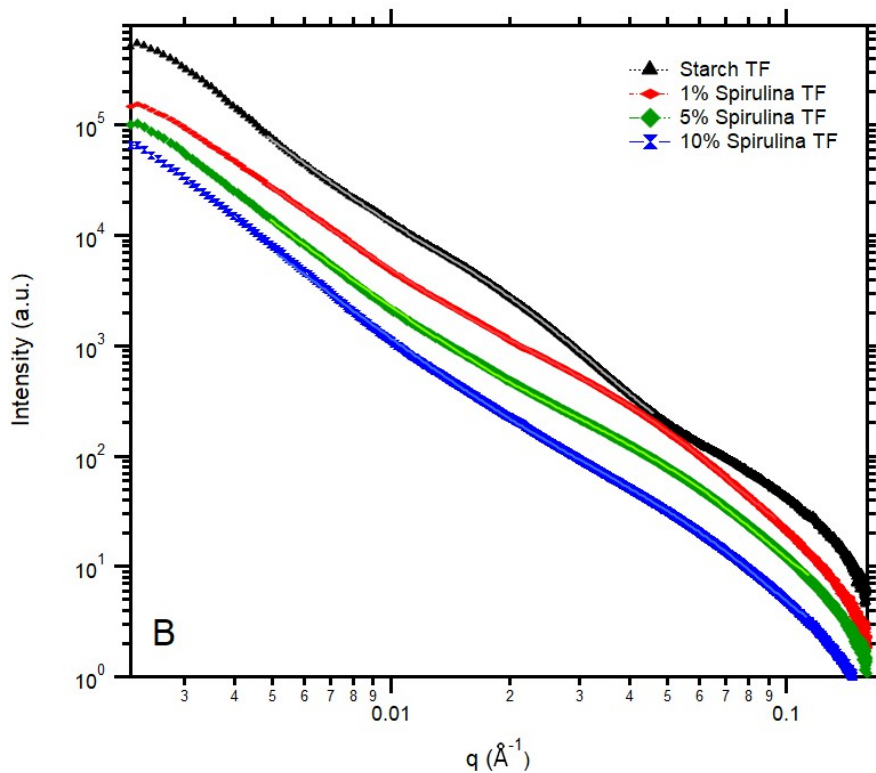
600 more amorphous samples, while the bands at 1000 and 1047 cm^{-1} become more defined in
601 more crystalline samples (Capron, et al., 2007; Sevenou, et al., 2002; J. J. G. van Soest, et
602 al., 1997). In particular, the trend in the intensity ratio for the 1022:1000 bands has been
603 reported to correlate reasonably with the crystallinity measured by X-ray diffraction
604 (Capron, et al., 2007). The values here obtained for the 1022:1000 intensity ratio were
605 approximately constant (ca. 0.96-0.94) for all the samples, confirming that starch presented
606 a similar degree of crystallinity in all the foams after storage for one month.

607
608 Very broad and weak shoulders were detected in the SAXS patterns from the foams after
609 storage (cf. Figure 6B). This type of scattering feature has been previously detected for
610 retrograded starches and has been assigned to the formation of heterogeneous semi-crystalline
611 structures (Shamai, Shimoni, & Bianco-Peled, 2004; Suzuki, Chiba, & Yarno, 1997). The
612 associated lamellar repeat distances ranged from ca. 51 nm for the pure starch to ca. 40-44
613 nm for the hybrid foams. Suzuki et al. reported lamellar distances of ca. 15-30 nm for potato
614 starch retrograded at low temperature (Suzuki, et al., 1997), whereas Shamai et al. noted the
615 appearance of a broad lamellar peak, characteristic of distances of ca. 20 nm, in high amylose
616 corn starch only when the samples were retrograded at low temperatures (Shamai, et al.,
617 2004). The broader shoulders observed for the hybrid foams indicate that the presence of
618 *Spirulina* promoted the re-crystallization of starch into more heterogeneous structures.
619 Furthermore, the power-law exponents in the low q region were 2.6-2.7 for all the samples,
620 confirming the existence of structures with low degree of order in the nanometre size range.
621 The slight increase in the power-law exponents of the foams at TF with respect to the
622 corresponding values at T0 confirm that the gelatinization process was irreversible and that
623 the neither the crystalline nor the lamellar structure of native starch were recovered upon
624 retrogradation. Presence of *Spirulina* and the effect of limited moisture accessibility

625 hindered the re-organization of amylose chains into crystalline domains and lamellae.
626 However, all the foams presented very similar crystallinity values after storage since the
627 mentioned effect upon storage was counteracted by the initially greater crystallinity of the
628 hybrid foams at T0.
629



630



631

632 **Figure 6.** (A) WAXS patterns from the raw starch and *Spirulina* and from the foams after
 633 storage for one month (TF) at 30% RH. Curves from raw starch and *Spirulina* have been
 634 offset for clarity. (B) SAXS patterns from the fully hydrated samples. Dotted lines
 635 correspond to the experimental data, whereas solid lines correspond to the fitting curves. The
 636 curves have been offset for clarity.

637

638 **Table 3.** Power-law exponents, lamellar peak parameters and crystallinity (X_C) for the foams
 639 after storage for one month (TF). The lamellar peak parameters were obtained by fitting the
 640 experimental SAXS data of the hydrated samples to the sum function of a power-law plus a
 641 Gaussian-Lorentzian peak and crystallinity was obtained from the peak fitting of the WAXS
 642 patterns from the dry samples. Standard deviation values on the last digit are shown in
 643 parentheses for the fitting parameters.

	Power-law exponent	Lamellar peak position (\AA^{-1})	Width of the lamellar peak (\AA^{-1})	Lamellar repeat distance (\AA)	X_c (%) ^(*)
Starch	2.678 (9)	0.0122 (1)	0.0141 (1)	51.40	6.19 [6.19]
1% <i>Spirulina</i>	2.561 (7)	0.0156 (4)	0.0346 (1)	40.28	5.65 [5.65]
5% <i>Spirulina</i>	2.680 (2)	0.0144 (4)	0.0441 (1)	43.72	6.24 [5.90]
10% <i>Spirulina</i>	2.716 (4)	0.0143 (6)	0.0454 (1)	43.87	8.57 [7.75]

644 ^(*)The crystallinity values calculated accounting only for the contribution of starch are shown
645 in brackets.

646

647 As observed in Figure 5B, although the differences were not significant, partially due to a
648 large variability between replicates, a trend of increased hardness (i.e. greater critical stress
649 values) was observed when comparing the samples at T0 and TF, which could be expected
650 from the structural changes occurring during storage (i.e. the amorphous as-processed starch
651 foam may re-organize into more ordered structures upon storage, leading to the formation of
652 more mechanically resistant foams). The weak effect is reasonable considering the very
653 limited re-crystallization of starch after storage. The same trend observed at T0 was detected
654 for the foams after storage, i.e. the presence of higher amounts of *Spirulina* led to the
655 formation of harder foams, although the high standard deviation values for the measured
656 samples preclude from extracting definitive conclusions. Regarding stiffness, no significant
657 differences were observed between samples before and after storage, partly due to the large
658 variability between samples.

659

660 Since the crystalline structure of starch is known to have a significant impact on its
661 digestibility, the content of resistant starch (RS) was determined in the foams at TF,
662 obtaining values of 1.1 ± 0.3 g RS/100 g sample in the pure starch and the samples with 1%

663 and 5% *Spirulina* and a higher content of 2.1 ± 0.7 g RS/100 g sample in the foam with 10%
664 *Spirulina*. This effect might be related to the formation of V-type amylose-lipid complexes,
665 which has been previously noted in starch-*Spirulina* blends (Martínez-Sanz, et al., 2018) and
666 as suggested by the WAXS patterns of the as-processed foams. These complexes are known
667 to form more stable crystalline structures than those found in native starches. For instance,
668 the formation of V-type amylose-lipid complexes has been shown to produce a decrease in
669 the amylose solubility, hence increasing the gelatinisation temperature (Eliasson, Carlson,
670 & Larsson, 1981). Furthermore, although amylose-lipid complexes are hydrolysed and
671 absorbed in the gastrointestinal tract to the same extent as free amylose, the hydrolysis
672 process has been seen to take place at a slower rate (J. Holm, et al., 1983). From a food
673 design point of view, a higher content of resistant starch is desirable, since it is related to
674 lower calorie content and lower glycaemic loads (Fuentes-Zaragoza, Riquelme-Navarrete,
675 Sánchez-Zapata, & Pérez-Álvarez, 2010). Thus, the addition of 10% *Spirulina* could be
676 interesting from a nutritional perspective, not only due to the increased protein content and
677 the antioxidant properties provided by bioactive compounds such as polyphenols (De Marco,
678 et al., 2014), but also due to the potential benefits associated to an increased content of
679 resistant starch in the foams.

680

681 **4. Conclusions**

682 Starch foams with different loadings (1, 5 and 10 wt.%) of the microalga *Spirulina* were
683 produced by means of extrusion cooking. The gelatinization process taking place during the
684 extrusion process led to the disruption of the crystalline and lamellar structure of native
685 starch, producing very amorphous foams. Furthermore, the crystallized fatty acids present
686 in the raw *Spirulina* were completely melted. The release of *Spirulina* intracellular

687 components promoted the formation of more stable V-type complexes with amylose, leaving
688 slightly more crystalline foams after processing. The microstructure of the foams was
689 strongly affected by the presence of the microalga and the hybrid foams exhibited more
690 densely packed and well-connected porous structures. This, in turn, affected the textural
691 properties of the foams, which became harder with higher *Spirulina* loadings.

692
693 All the foams presented a very limited re-crystallization upon storage, being this further
694 reduced in the case of the hybrid foams. As a result, all the foams presented similar
695 crystallinity values after storage. Furthermore, the starch lamellar arrangement was only
696 partially recovered, forming much more heterogeneous structures than those found in the
697 native starch. The formation of amylose complexes was translated into a higher resistant
698 starch content in the 10% *Spirulina* foam. This, together with the increased protein content
699 and the presence of bioactive compounds such as polyphenols, highlights the potential
700 benefits of adding *Spirulina* into starch extruded foams to produce snack foods with higher
701 nutritional value.

702
703 These results show that the combination of advanced characterization tools such as X-ray
704 scattering techniques and neutron tomography can provide very valuable information on the
705 nano- and microstructure of starch-based products, which will ultimately enable a rational
706 design of food products with the desired characteristics according to the consumers'
707 requirements.

708
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717

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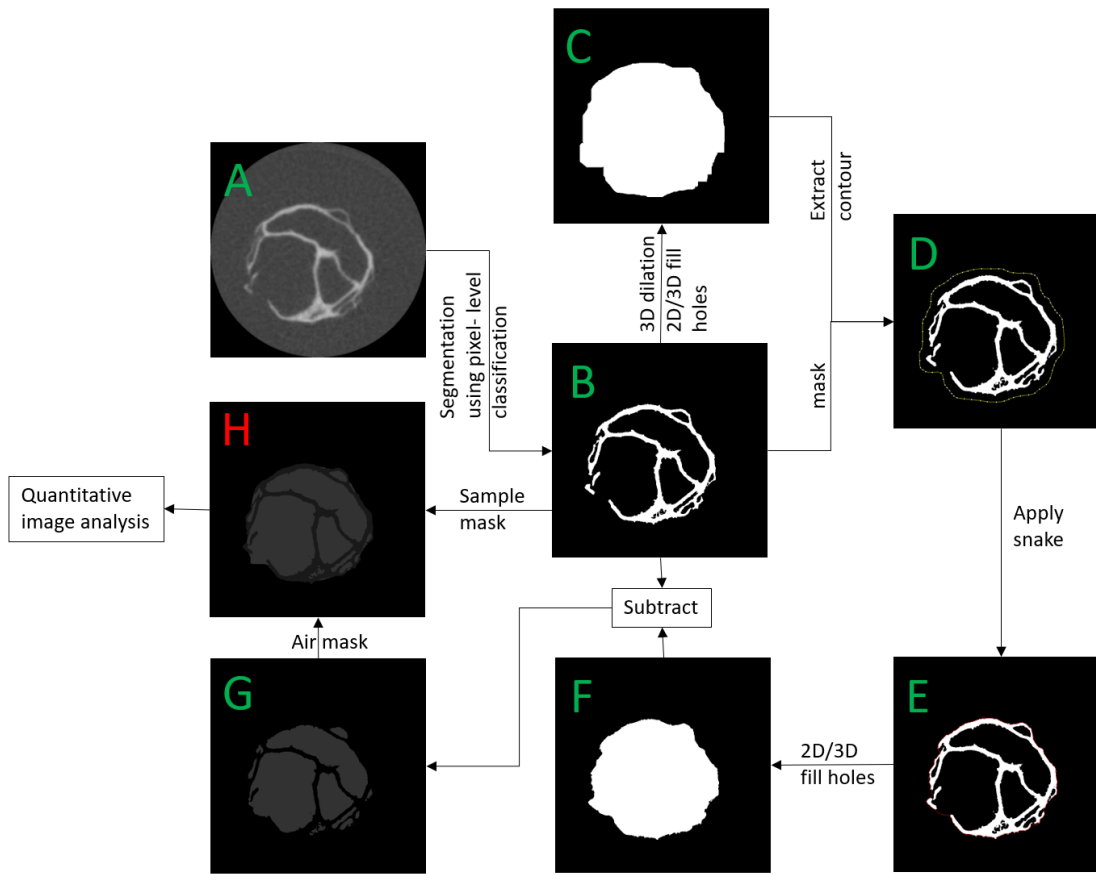
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940

941 **Supplementary Material**



942

943 **Figure S1.** Schematic drawing of the applied 2D/3D image processing protocol in order to
 944 segment the extruded samples with broken cell walls experiencing an open-air macro pore
 945 connection, thus allowing further quantitative 3D image analysis. The original image (A)
 946 with a varying grey-scale value for the thinner and thicker cell wall, as well as with
 947 physically broken cell wall, is segmented using a pixel-level classification method
 948 implemented in ‘Ilastik’. The resulting image (B) shows a pore connected to the outer air,
 949 via a thin broken cell wall, which impedes further 3D quantification of the air skeleton and
 950 quantification of both the connectivity density and pore size distribution. (C-G) shows the
 951 utilized image processing steps, finally resulting in image (H), where the extruded sample
 952 and the air pore space have been successfully segmented from the background. All image

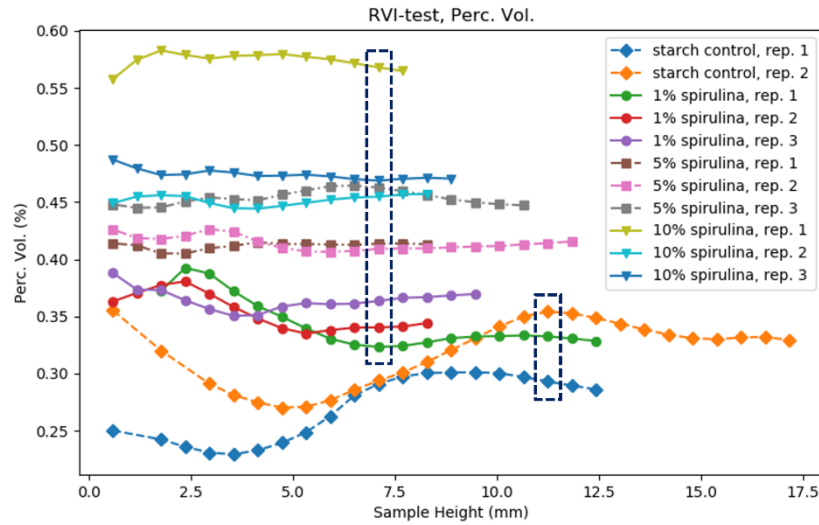
953 processing steps were performed in 3D. The results of these steps were used for both
954 quantitative image analysis and 3D-renderings.

955

956 **Representative Volume of Interest (RVI) test**

957 Quantitative 3D image analysis is commonly applied on so called volume of interests
958 (VOIs), which are often cubic, and extracted from different positions inside the sample, far
959 away from sample borders, which can often be non-representative to include in such an
960 analysis. The size of the considered VOI is determined by a prior Representative Volume of
961 Interest test (RVI-test) (Bear, 2013), where the quantified parameters, e.g. Volume
962 percentage (Vol.Perc.) and Connectivity Density (Conn.D) are plotted against the side length
963 of the VOI, as shown in Figures S2 and S3. The representative VOI-size is found when the
964 observed parameters ‘level out’ and are no longer affected by the varying size of the
965 considered VOI (Bear, 2013). Such an approach also works well on extruded samples with
966 a rather low porosity and a thick cell wall structure of the extrudate, as shown in (Karlsson,
967 Larsson, Lorén, Stading, & Rigdahl, 2019). However, for extruded samples with a very high
968 porosity, large pore size and very thin cell wall structure, as for the hereby described
969 extruded foams with a high starch content, quantitative analysis based on cubic VOIs is
970 challenging, due to the fact that the quantified parameters fail to ‘level out’, even for the
971 maximum possible applied VOI-size, following the RVI-test. Therefore, a modified RVI-
972 test, where the sample diameter is kept ‘as is’ and only the height of the VOI is allowed to
973 vary, was performed. For the given samples, a representative VOI height of 11.2 mm and
974 7.1 and was found for the pure starch and hybrid starch-*Spirulina* foams, respectively, as
975 shown in Figures S2 and S3.

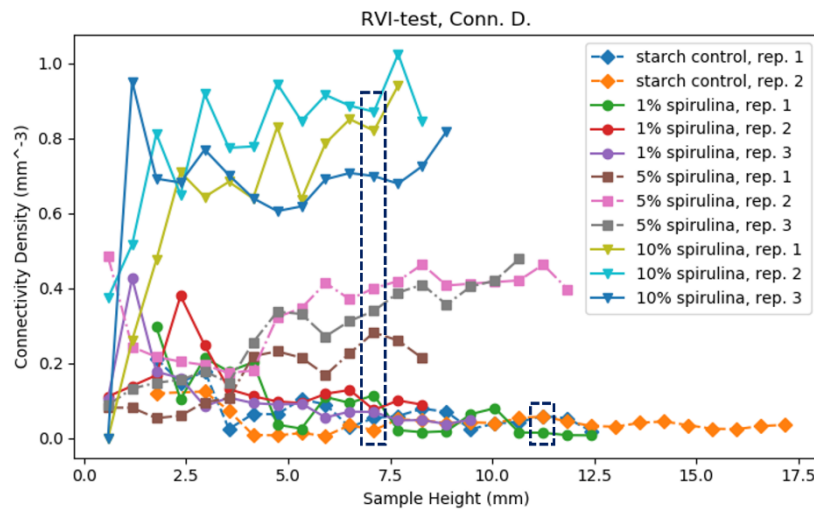
976



977

978 **Figure S2.** Test for finding the representative volume of interest (RVI-test), based on the
 979 Vol. Perc. as the evaluation parameter. The optimal height of the VOI was chosen at 11.2
 980 mm and 7.1 mm for the starch and hybrid starch-*Spirulina* foams, respectively.

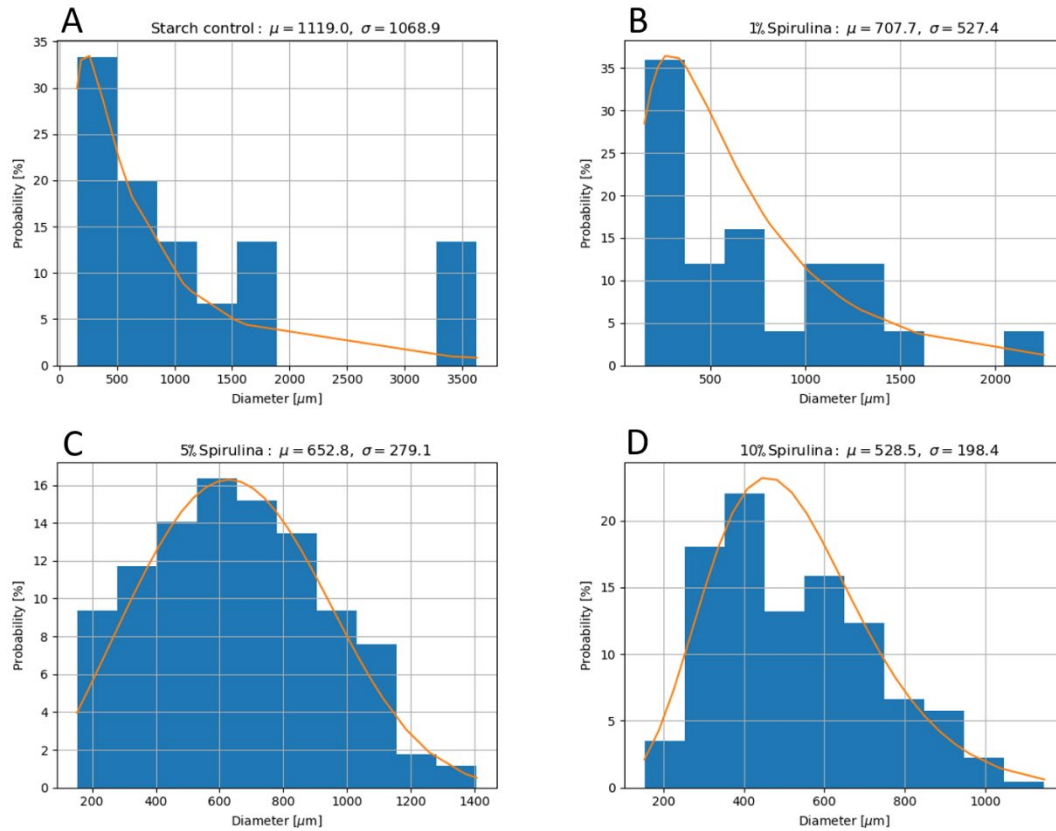
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982

983 **Figure S3.** Test for finding the representative volume of interest (RVI-test), based on the
 984 Conn. D. as the control parameter. The optimal height of the VOI was chosen at 11.2 mm
 985 and 7.1 mm for the starch and hybrid starch-*Spirulina* foams, respectively.

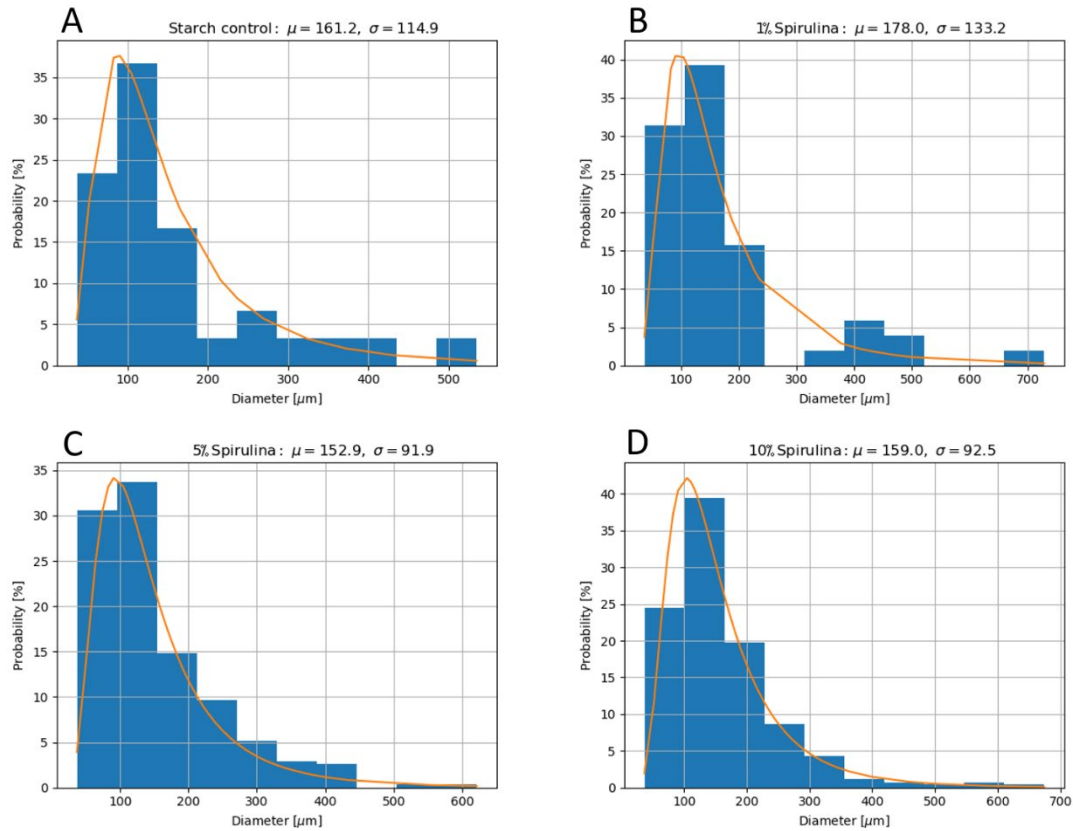
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987

988 **Figure S4.** Pore size distribution of the extruded foams (A) pure starch, (B) 1% *Spirulina*,
 989 (C) 5% *Spirulina* and (D) 10% *Spirulina*, as quantified by Method 1. A Weibull distribution
 990 (orange) was used to fit the data. The fitting works rather accurately for continuous pore
 991 sizes up to 1400 μm as in (C-D) but fails at pore sizes bigger than 2000 μm as in (A-B), due
 992 to the discrete number of available pores.

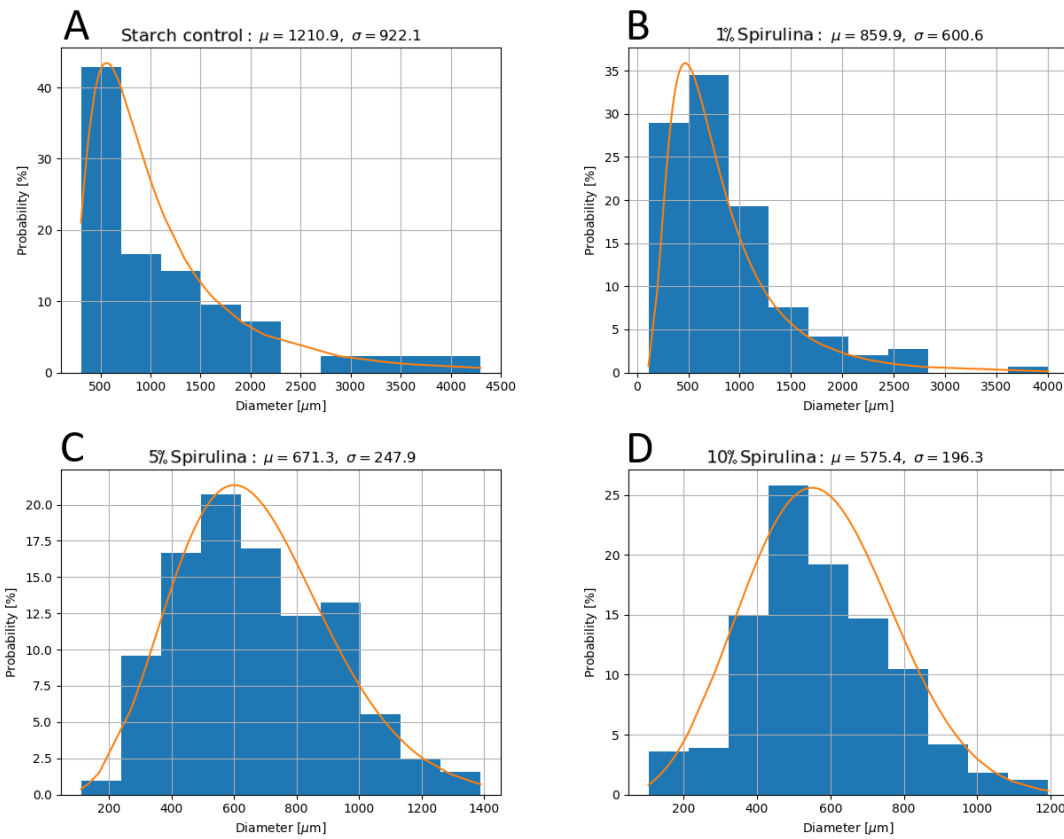
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994

995 **Figure S5.** Throat size distribution of of the extruded foams (A) pure starch, (B) 1%
 996 *Spirulina*, (C) 5% *Spirulina* and (D) 10% *Spirulina*, as quantified by Method 1. A Weibull
 997 distribution (orange) was used to fit the data. The fitting works rather accurately for
 998 continuous throat sizes in (C-D) and less well for the discrete throats sizes in (A-B)

999



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Figure S6. Pore size distribution of the extruded foams (A) pure starch, (B) 1% *Spirulina*,

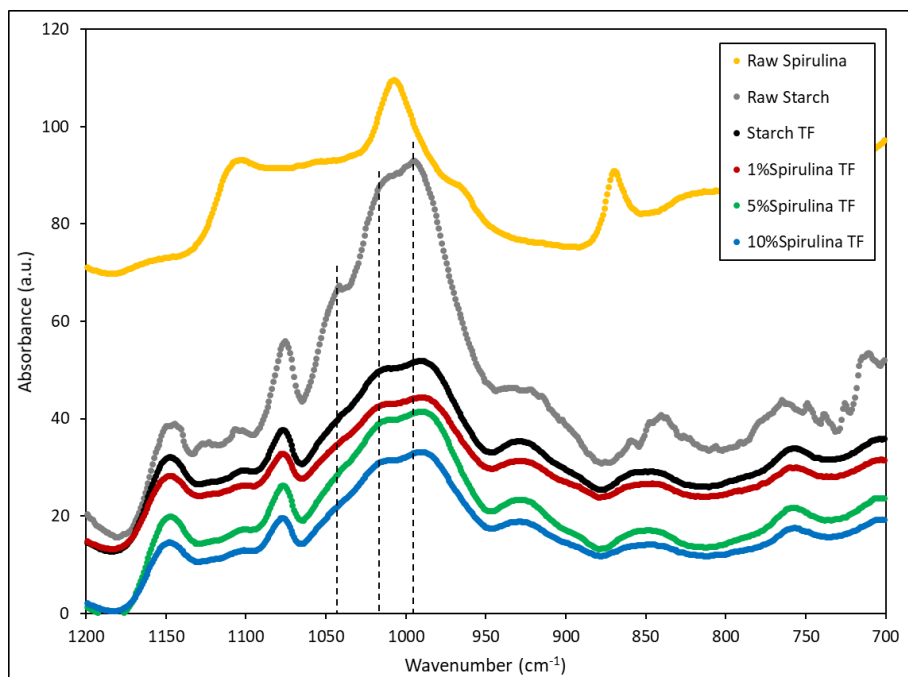
1002

(C) 5% *Spirulina* and (D) 10% *Spirulina*, as quantified by Method 2. A Weibull distribution

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(orange) was used to fit the data. The fitting works rather accurately for all pore sizes (A-D).

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Figure S7. ATR-FTIR spectra from the raw starch and *Spirulina* and from the foams after storage. Curves have been offset for clarity.