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1	NANO-/MICROSTRUCTURE OF EXTRUDED SPIRULINA/STARCH FOAMS IN
2	RELATION TO THEIR TEXTURAL PROPERTIES
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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. Moreover, the crystalline structure of the fatty acids present in the raw microalga was lost 26 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 ( $X_{C}$ ~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.

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

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

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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 nutritional supplements and food additives (Chacón-Lee & González-Mariño, 2010; 76 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, y-linolenic acid and other 84 compounds such as vitamins (mainly B vitamins), minerals, pigments (phycocyanin, β-85 carotene and xanthophyll), y-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 especially important in the case of starchy foods, whose processing is based on the 96 97 occurrence of a phenomenon known as gelatinization. Native starch granules display a hierarchical structure consisting of alternating amorphous and semi-crystalline growth rings 98 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 of 9-10 nm (Blazek & Gilbert, 2011). When processed at high temperature and moisture 101 102 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 method TAPPI T211 om-07. Lipids were estimated by performing a Soxhlet extraction of 137 138 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-Ciocalteau colorimetric assay (Singleton, Orthofer, & 140 Lamuela-Raventós, 1999). The gross chemical composition of the microalgal biomass was determined as: 40.6% ash, 26.7% protein, 15.1% lipid; 8.0% carbohydrate and 0.9% 141 142 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.% dry weight) were prepared by extrusion cooking. The samples were processed with 30 wt.% 146 147 moisture content to facilitate starch gelatinization during processing and let to equilibrate for 24 hours before extrusion. A single-screw extruder (Brabender Duisburg DCE-330, 148 149 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 compression ratio with a linearly tapered rod and 20 equidistantly positioned flights. A round 151 152 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 Thermo Nicolet Nexus (GMI, USA) equipment. The spectra were taken at 4 cm<sup>-1</sup> resolution 164 in a wavelength range between 400-4000 cm<sup>-1</sup> and averaging a minimum of 32 scans. 165

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#### 2.4 Determination of resistant starch content

168 Resistant starch content was measured using the Megazyme Resistant Starch Assay Kit

(Megazyme International, Wicklow, Ireland), based on the AACC method 32-40.01 (AACC, 169

170 2009). Extruded samples were grounded to a fine powder in a blender. The samples (100 mg) were incubated with 4 mL mixture of pancreatic α-amylase and amyloglucosidase for 171 172 16 h (at 37 °C, 200 strokes/min) in a shaking water bath (Julabo, Seelbach, Germany) to hydrolyze digestible starch to glucose. The reaction was terminated with 4 mL of 99 % 173 174 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 KOH solution 2 M in an ice bath. The solution was neutralized after 20 min with 8 mL 176 177 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/peroxidase 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 100 g of dry sample. 183

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#### 185

# 2.5 Scanning electron microscopy (SEM)

SEM measurements were conducted on a Hitachi microscope (Hitachi S-4800) at an accelerating voltage of 10 kV and a working distance of 8-16 mm. The pure and composite starch films were cryo-fractured after immersion in liquid nitrogen to observe the crosssections. The samples were then sputtered with a gold–palladium mixture under vacuum 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 characterized shortly after being processed ( $t_0$ ) and after prolonged storage at 30% RH ( $t_f$ ), 196 197 both in their native dry state and after being hydrated in excess water for at least 24 h. The energy of the incident photons was 12.4 KeV (an equivalent wavelength,  $\lambda$ , of 1 Å). The 198 199 SAXS diffraction patterns were collected by means of a single-photon counting detector, Pilatus 1 M, with an active area of 168.7×179.4mm<sup>2</sup>, an effective pixel size of 172×172 µm<sup>2</sup> 200 201 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 Å<sup>-1</sup>. The WAXS diffraction patterns 203 were collected by means of a 3 CCD detectors, Rayonix LX255-HS, with an active area of  $85 \times 255$  mm<sup>2</sup>, an effective pixel size of  $44 \times 44 \ \mu$ m<sup>2</sup> and a dynamic range of 16 bits. In this 204 205 case, the sample-to-detector distance was set to 144.3 mm, corresponding to a maximum q value of 8.023 Å<sup>-1</sup>. This detector was tilted with a pitch of 29.38°. The exposure time, 206 207 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** SA

### 2.7 SAXS/WAXS data fitting

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

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

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

222

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

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

where y is a fitted value for a given point,  $y_i$  is the measured data value for the point and  $\sigma_i$  is an estimate of the standard deviation for  $y_i$ . The curve fitting operation is carried out iteratively and for each iteration, the fitting coefficients are refined to minimize  $\chi^2$ . The crystallinity index X<sub>c</sub> was determined from the obtained fitting results by applying the following equation:

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

where  $A_{Total}$  is the sum of the areas under all the diffraction peaks and  $\Sigma A_{Crystal}$  is the sum of 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, Hélary, & 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% Spirulina (n=3) and 10% Spirulina (n=3), where n equals the number of scanned samples. 243 244 The scanning parameters were the following: exposure time, 60s and number of projections, 180; number of dark and flat-field projections, 10; sample-to-detector distance, 50 mm; field 245 246 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 with a sCMOS sensor (effective number of pixels, 2560 x 2160 pixels<sup>2</sup>; physical pixel size: 248 249  $6.5 \times 6.5 \ \mu\text{m}^2$ ) coupled to a 50  $\mu\text{m}$  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). Furthermore, the camera was cooled down to -30°C, to reduce noise. Pieces of few 251 252 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.

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### 5 **2.8.2 Image pre-processing and 3D reconstruction**

An in house-developed macro from the IMAGINE imaging station and implemented in ImageJ (Rueden, et al., 2017) was used to normalize the projection images, remove noise and correct the tilt-angle. Hereinafter, the SYRMEP Tomo Project (STP) (Brun, et al., 2017) was used to reconstruct a 3D-map of the linear attenuation coefficient (LAC) [cm<sup>-1</sup>] of the 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, 2011) was applied. One slice from each type of sample group was used for combined 269 270 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**

Quantitative 3D image analysis was performed using the Pore3D software library (Brun, et al., 2010) on Representative Volume of Interest (RVI) of the respective data sets, following an RVI-test, as further described in the Supplementary Material section 'Representative Volume of Interest (RVI) test'. The following image parameters were quantified: 1) Volume 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 of the VOI as further described by (Parfitt, et al., 1987), 3) Structure Thickness (St.Th.) of 289 290 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: Borgefors(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

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

305

# 306 **2.9 Uniaxial Compression testing**

Uniaxial compression tests were performed on an Instron material testing device (model 5542, Instron, Norwood, MA, USA). Extruded samples were carefully cut to cylindrical shapes with a height of ca. 1 cm with the help of a sharp scalpel and placed between flat metal surfaces covered with emery paper to avoid slippage. A 3 cm cylindrical probe was used to compress the samples until fracture, using a 500 N load cell (load values during compression were in the range of 125-200 N) at a crosshead speed of 1 mm/s. At least 5 replicates of each type of extruded sample were measured. Force (N) and distance (mm) were converted to true stress ( $\sigma_T$ ) and true strain ( $\varepsilon_T$ ) using equations 1 and 2 by the instrument software Blue Hill.

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

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

Where  $h_0$  is the initial height of the sample,  $\Delta h(t)$  is the change in height during compression and  $r_0$  is the initial radius of the sample. The compression modulus was obtained as the slope of the initial linear zone of the true stress vs. true strain plots. Mean values of the critical stress and the compression modulus were compared by Tukey and 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**

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

332

The WAXS patterns of the raw materials and the as-processed foams (T0) are shown in Figure 1A. As observed the raw corn starch presented WAXS patterns typical from the Atype crystalline allomorph, characterized by the main diffraction peaks located at  $2\theta = 15.1^{\circ}$ , 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 20 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 of lipids determined in the sample (ca. 15%), thus supporting the hypothesis of crystallized 342 343 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 treatments is known to induce the formation of V-type crystalline structures in corn starch 353 354 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 very easily, even with only mild stirring (Fabra, et al., 2017; Martínez-Sanz, et al., 2018), 358 359 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 high362 humidity thermal treatment in the presence of Spirulina (Martínez-Sanz, et al., 2018). Thus, it can be assumed that the shearing effect during the extrusion cooking process promoted 363 364 cell wall disruption and facilitated the release of intracellular components which were able to form complexes with amylose. Moreover, the WAXS results suggest that the formed V-365 366 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 cooking process. This is in agreement with previous work which reported that a small 368 369 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

While WAXS is able to provide information on the crystalline structure of starch, SAXS is 378 379 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 starch presented a weak and broad lamellar peak centred at ca. 0.071 Å<sup>-1</sup> in the native dry 381 382 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 more intense and sharp in the hydrated state and its position shifted towards ca. 0.063 Å<sup>-1</sup> 391 392 (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 confirmed by the change in the power-law exponent from the low q region (0.003-0.005  $\text{\AA}^-$ 398 <sup>1</sup>), which decreased from ca. 4 for the raw starch (corresponding to the Porod's ideal two-399 400 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 higher exponents, similar to the values here reported, have been estimated for melt-404 405 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

412 process, both for the pure starch and the blends with the added *Spirulina*.







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

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

Power-law	Lamellar	Width of	Lamellar	
exponent	peak position	the lamellar	repeat	X <sub>C</sub> (%)
	(Å <sup>-1</sup> )	peak (Å <sup>-1</sup> )	distance (Å)	

Raw Spirulina	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% Spirulina	2.34 (5)				4.59
1					
5% Spirulina	2.52(1)				6.62
1					
10% Spirulina	2.73 (1)				8.72
1					

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

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

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

445 An in-depth 3D characterization of the microstructure from the different foams was carried out by means of neutron tomography. The porous structure of the extruded foams could be 446 447 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 distributions, calculated with method 1 and 2, as described in the Supplementary Material 457 458 section (Figures S4, S5 and S6), indicating that the pore size (mean and standard deviation) 459 decreased with increasing concentration of Spirulina.





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 described by 'Method 1'. The height of the samples were 11.2 mm and 7.1 mm for the pure 467 468 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.

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

	Porosity (%)	Specific Surface Area (mm <sup>-1</sup> )	Cell wall thickness (µm)	Conn.D (mm <sup>-3</sup> )	Pore Size (µm) Method 1 (Method 2)	Throat Size (µm) Method 1
Starch	67.6 ± 3.1	$1.7 \pm 0.1$	234.0 ± 18.4	$0.06 \pm 0.00$	$1109.0 \pm 1068.9$ $(1210.9 \pm 922.1)$	$161.2 \pm 114.9$
1% Spirulina	64.8 ± 1.2	2.5 ± 0.1	199.1 ± 0.2	$0.07 \pm 0.00$	$707.7 \pm 527.4$ (859.9 ± 600.6)	$178.0\pm133.2$
5% Spirulina	57.2 ± 2.4	3.8 ± 0.1	165.1 ± 1.7	$0.34 \pm 0.05$	$652.8 \pm 279.1$ $(671.3 \pm 247.9)$	$152.9 \pm 91.9$
10% Spirulina	50.3 ± 5.0	$4.2 \pm 0.2$	180.1 ± 25.2	$0.79 \pm 0.09$	$528.5 \pm 198.4$ (574.4 $\pm 196.3$ )	$159.0\pm92.5$



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

The structural changes undergone by starch during the extrusion cooking process were 485 486 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  $3.1 \pm 0.2$  for the corn starch to  $2.8 \pm 0.1$ ,  $2.3 \pm 0.1$  and  $2.2 \pm$ 493 0.1 for the hybrid samples containing 1%, 5% and 10% Spirulina, respectively. These results 494 are in agreeement 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 extrussion 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.

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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 ( $\sigma_c$ ), defined as the first maximum stress value, could

be identified for all the samples, but more clearly for the 10% Spirulina foam. After the 510 511 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  $\pm$  322 kPa for 10% Spirulina, as compared to  $1505 \pm 573$  kPa,  $1568 \pm 310$  kPa and  $2011 \pm 622$  kPa for the pure 516 517 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 heterogenity. As previously discussed, the microstructure of the extruded samples consisted 526 527 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 colapse when a critical value of stress is reached (Ashby & Medalist, 1983). Ashby 531 and Gibson demostrated 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 with the greater bulk density and containing thinner cell walls and a higher number of cells 534

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

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

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# 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, Li, Copeland, Niu, & Wang, 2015). The impact of the Spirulina microalga on the 551 552 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 shown in Figure 6A. As observed, the crystalline structure of the foams changed from the 556 557 non-stable V-type mixture observed shortly after processing the samples (T0) towards a V<sub>A</sub> 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 processing (T0), the retrogradation process gave rise to a crystallinity increase of 92% in the 574

575 case of pure starch and 23% for the 1% Spirulina foam, whereas the crystallinity corresponding to the starch fraction was slightly decreased for the 5% and 10% Spirulina 576 577 foams. It is known that greater moisture contents favour starch re-crystallization (J. Van Soest & Knooren, 1997; Wang, et al., 2015) and in particular, the presence of moisture is 578 579 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.

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To confirm the effect of *Spirulina* on the crystallinity of the foams after storage, FT-IR analyses were also carried out and the obtained spectra are shown in Figure S7. The spectral region of 800–1200 cm<sup>-1</sup> was used to investigate structural differences between samples, as previous studies have suggested that bands in this 'fingerprint' region reflect changes in polymer conformation and hydration of processed starches (Capron, Robert, Colonna, Brogly, & Planchot, 2007; Htoon, et al., 2009; Sevenou, Hill, Farhat, & Mitchell, 2002; J. J. G. van Soest & Vliegenthart, 1997). The band at 1022 cm<sup>-1</sup> has been seen to increase in more amorphous samples, while the bands at 1000 and 1047 cm<sup>-1</sup> become more defined in more crystalline samples (Capron, et al., 2007; Sevenou, et al., 2002; J. J. G. van Soest, et al., 1997). In particular, the trend in the intensity ratio for the 1022:1000 bands has been reported to correlate reasonably with the crystallinity measured by X-ray diffraction (Capron, et al., 2007). The values here obtained for the 1022:1000 intensity ratio were approximately constant (ca. 0.96-0.94) for all the samples, confirming that starch presented a similar degree of crystallinity in all the foams after storage for one month.

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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 retrogaded 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 retrogradation. Presence of Spirulina and the effect of limited moisture accessibility 624

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





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

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

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

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<sup>(\*)</sup>The crystallinity values calculated accounting only for the contribution of starch are shown in brackets.

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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  $1.1 \pm 0.3$  g RS/100 g sample in the pure starch and the samples with 1%

663 and 5% Spirulina and a higher content of  $2.1 \pm 0.7$  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, the formation of V-type amylose-lipid complexes has been shown to produce a decrease in 668 669 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 design point of view, a higher content of resistant starch is desirable, since it is related to 673 lower calorie content and lower glycaemic loads (Fuentes-Zaragoza, Riquelme-Navarrete, 674 675 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.

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#### 681 **4. Conclusions**

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

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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.

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

708

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941 Supplementary Material





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

processing steps were performed in 3D. The results of these steps were used for bothquantitative 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.





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



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



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



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



Figure S6. Pore size distribution of the extruded foams (A) pure starch, (B) 1% *Spirulina*,
(C) 5% *Spirulina* and (D) 10% *Spirulina*, as quantified by Method 2. A Weibull distribution
(orange) was used to fit the data. The fitting works rather accurately for all pore sizes (A-D).



**Figure S7.** ATR-FTIR spectra from the raw starch and *Spirulina* and from the foams after

1007 storage. Curves have been offset for clarity.