 Focus on the composition of cell wall related polysaccharides Tom M.M. Bernaerts *, Lore Gheysen *, Clare Kyomugasho *, Zahra Jamsazzadeh Kermani *, Stéphanie Vandionant *, Imogen Foubert *, Marc E. Hendricks *, Ann M. Van Loey *.* Laboratory of Food Technology, Leuven Food Science and Nutrition Research Centre (LFoRCe), Department of Microbial and Molecular Systems (M*S), Katholicke Universiteit Leuven, Kasteelpark Arenberg 22, Box 2457, 3001 Heverlee, Belgium Authors are affiliated to: Laboratory of Food Technology (member of Leuven Food Science and Nutrition Research Center, LFoRCe), Department of Microbial and Molecular Systems (M*S), KU Leuven, Kasteelpark Arenberg 22 box 2457, 3001 Heverlee, Belgium Laboratory Food and Lipids (member of Leuven Food Science and Nutrition Research Center, LFoRCe), Department of Microbial and Molecular Systems (M*S), KU Leuven, Kasteelpark Arenberg 22 box 2457, 3001 Heverlee, Belgium b Laboratory Food and Lipids (member of Leuven Food Science and Nutrition Research Center, LFoRCe), Department of Microbial and Molecular Systems (M*S), KU Leuven Kulak, E. Sabbelaan 53, 8500 Kortrijk, Belgium Fax: +32 16 321 960 Telephone (+32 16 321 960 Telephone (+32 16 321 967 E-mail: ann.value (@kuleuven.be 	1	Comparison of microalgal biomasses as functional food ingredients:
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24 Abstract

25 Microalgae are rich in several nutritional and health-beneficial components, showing great potential as 26 functional food ingredients. To this extent, knowledge of the biomass composition is essential in the selection 27 of suitable microalgae species for specific food applications. Surprisingly, although cell wall polysaccharides 28 are generally reported to play a role in functionality, limited attention has been given to the cell wall related polysaccharides of microalgae so far. Therefore, this study aimed to characterize dry biomasses of ten 29 30 microalgae species with potential as functional food ingredients, with a particular focus on the composition of cell wall related polysaccharides. The investigated species were Arthrospira platensis, Chlorella vulgaris, 31 Diacronema lutheri, Tisochrysis lutea, Nannochloropsis sp., Odontella aurita, Phaeodactylum tricornutum, 32 Porphyridium cruentum, Schizochytrium sp. and Tetraselmis chuii. Lipids, proteins and ash made up a large 33 fraction of the biomasses, except for the freshwater algae C. vulgaris and A. platensis which were mainly 34 composed of proteins and polysaccharides. Generally, low amounts of storage polysaccharides (2 - 8%) were 35 observed in the investigated microalgae species, while extracellular polymeric substances were only present in 36 P. cruentum, O. aurita, C. vulgaris and A. platensis. Cell wall polysaccharides contributed approximately 10% 37 38 of the biomass and were composed of heteropolysaccharides, showing at least five different monosaccharides. 39 Moreover, the presence of uronic acids and sulfate groups provides anionic characteristics to the cell wall related polysaccharides of several microalgae. As a result, these polysaccharides show potential to display interesting 40 functionalities as bioactive or technological substances. 41

- 42 Keywords
- Biomass composition cell wall polysaccharides exopolysaccharides monosaccharide uronic acid sulfate
 Highlights
- 45 Carbohydrates separately quantified as storage-, cell wall- and exo-polysaccharides
- 46 Exopolysaccharides were observed in P. cruentum, C. vulgaris, O. aurita, A. platensis
- 47 Glucose, galactose, xylose and mannose are dominant sugars in the polysaccharides
- 48 Uronic acid and sulfate groups confer anionic characteristics to the polysaccharides

49 Abbreviations

- 50 ω 3-LC-PUFA omega-3 long chain polyunsaturated fatty acids
- 51 CWPS cell wall polysaccharides
- 52 EPMS extracellular polymeric substances
- 53 EPS extracellular polysaccharides
- 54 SPS storage polysaccharides
- 55 UHPH ultra high pressure homogenization

56

57 1. Introduction

Microalgae are a promising source of several nutritional and health-beneficial components, including 58 59 omega-3 long chain polyunsaturated fatty acids (ω 3-LC-PUFA), proteins, minerals and antioxidants. In 60 recent decades, research has been exploring their potential as a functional food ingredient, to enhance the nutritional value of food products [1-3]. Since there is a large number of microalgae species available and 61 62 the composition of microalgal biomasses largely varies among different species, knowledge on the biochemical composition is required for the selection of suitable microalgae towards specific food 63 applications. However, even for a specific microalga species, variable biomass profiles are reported in 64 different studies. Part of this variability is attributed to differences in cultivation conditions, since many 65 environmental factors such as temperature, salinity and nutrient availability can strongly affect the chemical 66 composition of microalgae [4]. While this allows the optimization of cultivation conditions to maximize the 67 production of specific biomolecules, it also results in an increased complexity in comparing different 68 microalgal biomass profiles. On the other hand, the diverse biomass compositions found in literature can 69 also be attributed to distinct analytical approaches used in different studies. For instance, protein contents 70 can be determined by colorimetric assays or elemental analysis of nitrogen. While the former methods are 71 sensitive to interferences and require pretreatments to completely release intracellular proteins, the latter 72 relies on the use of nitrogen-to-protein conversion factors, but different conversion factors have been used 73 by different authors (ranging from 4.44 to 6.25) [5,6]. As a consequence, there is still a demand for studies 74 comparing microalgal biomass profiles using standardized protocols. 75

The nutritional valuable components of microalgae are stored inside the microalgal cell, which is protected 76 by a cell wall (except for a few species). As a consequence, the cell wall plays an important role as a natural 77 barrier, limiting extraction yields of high-value products or resulting in a low bioavailability of intracellular 78 components [7,8]. In this context, extensive research has been performed on the disruption of microalgal 79 cells, including chemical modifications and mechanical, thermal or ultrasonication processes [9]. Although 80 several treatments proved successful for many microalgae, optimization is still required for species 81 82 possessing a very rigid cell wall. In recent decades, the use of cell wall degrading enzymes has gained 83 interest as this shows some advantages, such as a minimal impact on the desired nutrients and low energy

requirements [7,10,11]. However, this approach requires the precise knowledge of the cell wall composition,
for the appropriate selection of specific cell wall degrading enzymes.

Insight into the composition of the cell wall is not only desired in terms of process optimization, but also 86 because distinct cell wall related polysaccharides might show potential for several biotechnological 87 purposes. To date, commercialization of high value products from microalgae is mainly targeted to ω 3-LC-88 89 PUFA, antioxidants or pigments, while microalgal polysaccharides are receiving limited attention. This might be due to the lack of knowledge on the composition and structure of cell wall-related polysaccharides, 90 with only few studies suggesting the potential of cell wall related polysaccharides for several applications. 91 According to de Jesus Raposo et al. (2015) [12], sulfated polysaccharides of microalgae display various 92 bioactivities, such as antiviral, antioxidant and anti-inflammatory activities. Moreover, exopolysaccharides 93 of the red microalga Porphyridium sp. show unique rheological properties and might therefore be used as 94 thickening agents in food products [13]. Thus, establishing the composition of cell wall related 95 polysaccharides, such as the monosaccharide profile or the degree of sulfation, could increase the 96 functionality of microalgal sources towards several applications. 97

Cell wall related polysaccharides comprise different types of polymers, including cell wall polysaccharides 98 (CWPS) and extracellular polymers. The latter are generally described as polymers that can be secreted into 99 the surrounding environment, such as the cultivation medium. Since the amount of secreted material 100 101 depends on the growth conditions and time of harvesting, the extracellular polymers can be both found as solubilized polymers in the aqueous phase as well as an external layer still surrounding the microalgal cell. 102 Although many microalgae species and cyanobacteria secrete extracellular polymers into the cultivation 103 medium, the type of secreted material is often unclear in literature, primarily due to distinct terminology. 104 The secreted material is often called EPS, referring to either extracellular polymeric substances, 105 extracellular polysaccharides or exopolysaccharides, although terms as released polysaccharides (RPS), 106 extracellular organic matter (EOM) or algogenic organic matter (AOM) are also commonly used [14,15]. 107 Depending on the definition, different classes of organic macromolecules are included, such as 108 109 polysaccharides, proteins, nucleic acids, phospholipids and smaller molecules [15]. In this study, a 110 distinction will be made between all polymeric material that can be secreted into the environment (referred

to as extracellular polymeric substances, EPMS) and polysaccharides that can be secreted (referred to asextracellular polysaccharides, EPS).

To date, information in literature on the amount of cell wall related polysaccharides in microalgae is scarce. 113 114 In fact, quantification of microalgal polysaccharides is usually done by analyzing the total carbohydrate content, thus including both storage polysaccharides (SPS) and cell wall related polysaccharides. However, 115 116 these two types of polysaccharides exhibit different functions in the microalgal cell. The main function of SPS is the storage of energy, providing substrates for metabolic processes and allowing survival of the 117 organism during dark periods. In contrast, cell wall related polysaccharides, comprising CWPS and EPMS, 118 play an important structural role in the microalgal cell. Whereas CWPS provide resistance to turgor pressure, 119 interactions between EPMS of different cells allow the creation of multicellular structures [16]. As a 120 consequence, both types of polysaccharides can contribute to different functional properties of the biomass, 121 depending on their structure and composition. Total carbohydrate contents (in fact expressed as glucose-122 equivalents due to the use of non-specific colorimetric assays) do therefore not allow the prediction of the 123 124 potential of microalgal polysaccharides.

Although several authors have reported monosaccharide profiles of microalgae, the composition of the cell 125 wall related polysaccharides of many microalgae is still unknown. On the one hand, some studies presented 126 monosaccharide profiles after hydrolysis of the total biomass [17,18]. However, due to possible interference 127 of other components, such as SPS and glycolipids, these results provide only limited information on the cell 128 wall composition. On the other hand, some authors have described the composition of cell wall related 129 polysaccharides, but the results were mostly concerning specific polysaccharide fractions obtained by a 130 selective extraction procedure [19,20]. Studies focusing on the polysaccharide composition of the whole 131 cell wall are therefore very limited. Moreover, large variability in cell wall composition has been reported 132 within a genus, a species and even within a strain, which can be due to differences in cultivation conditions 133 or depending on the life stage of the cell [7], further limiting the comparison among the studies available. 134 135 Therefore, the aim of this study is to apply a universal procedure for extraction of the total cell wall related 136 polysaccharides, including CWPS and EPMS, of commercially available microalgae species followed by a 137 detailed characterization.

138 The microalgae species used in this study were selected for their potential as functional food ingredients: 139 Arthrospira platensis, Chlorella vulgaris, Diacronema lutheri, Tisochrysis lutea (formerly listed as Isochrysis galbana), Nannochloropsis sp., Odontella aurita, Phaeodactylum tricornutum, Porphyridium 140 141 cruentum, Schizochytrium sp. and Tetraselmis chuii. Most of them show interesting nutritional profiles, e.g. containing ω 3-LC-PUFA, proteins rich in essential amino acids and antioxidants. In addition, some of these 142 biomasses have been accepted or authorized under the European novel food regulation, or applications are 143 ongoing. Finally, by selecting these microalgae a diverse taxonomic spectrum was obtained, composed of 144 photoautotrophic eukaryotic species classified as Chlorophyta (C. vulgaris, T. chui), Rhodophyta 145 (P. cruentum), Haptophyta (D. lutheri, T. lutea), Eustigmatophyta (Nannochloropsis sp.), Bacillariophyta 146 or diatoms (O. aurita, P. tricornutum), one heterotrophic species belonging to Labyrinthulomyceta 147 (Schizochytrium sp.) and one prokaryotic cyanobacterium (A. platensis) [21] 148

149 The objective of this study is to provide a fair comparison of ten microalgae species that are of interest for use as functional food ingredients. On the one hand, the microalgae were characterized in terms of biomass 150 composition. On the other hand, the composition of the cell wall related polysaccharides CWPS and EPMS 151 were established, by determining the monosaccharide profile, uronic acid content and sulfate content. All 152 analyses were performed on commercially available dry biomasses, with regard to the application of dried 153 microalgal biomass as a functional ingredient in food products, as food ingredients are generally delivered 154 155 in a dry form to guarantee long term storage stability. The insights provided by this study can facilitate an appropriate selection of microalgae species for enhancing the nutritional value of food products, as well as 156 revealing their potential as bioactive or biotechnological substances. 157

- 158
- 159 2. Materials and Methods
- 160 **2.1. Microalgal biomass**

161 Commercially available microalgal biomass was obtained from different companies. Lyophilized 162 biomasses of *Nannochloropsis* sp. and *Tisochrysis lutea* were obtained from Proviron (Hemiksem, 163 Belgium). *Odontella aurita* was purchased from Innovalg (Bouin, France), *Tetraselmis chuii* from 164 Fitoplancton Marino (Cádiz, Spain) and the cyanobacterium *Arthrospira platensis* from Earthrise (Irvine,

165 CA, USA). Lyophilized biomass of *Schizochytrium* sp. was kindly donated by Mara Renewables
166 Corporation (Dartmouth, Canada). Spray-dried biomass of *Chlorella vulgaris* was obtained from Allma
167 (Lisbon, Portugal). Biomasses of *Phaeodactylum tricornutum* and *Porphyridium cruentum* were obtained
168 as a wet paste from Necton (Olhão, Portugal) and immediately lyophilized. All biomasses were stored in
169 closed containers at -80 °C until use.

Biomass of *Diacronema lutheri* (CCAP 931/1) was produced in-house. This species was cultured in Wright's Cryptophyte medium in 125 L pilot-scale tubular airlift photobioreactors. Prior to use, the medium was sterilized by membrane filtration (0.2 μ m pore size). The photobioreactors were continuously illuminated (125 μ mol photons/m²·s) and the culture was maintained at pH 7.5 by automated CO₂ injection. The biomass was harvested at the end of the exponential growth phase by centrifugation, lyophilized and stored at -80 °C.

- Even though no full details on cultivation and harvesting conditions could be provided due to restrictions imposed by some companies, the data obtained will be indicative for dry biomasses towards food applications. Possible deviations from optimal cultivation conditions would generally be noticed from the amount of storage components, such as SPS and lipids, rather than structural components such as cell wall related polysaccharides [4]. However, the obtained biomass profiles correspond well with those reported in literature for the investigated microalgae species grown under standard conditions, as extensively described in **Section 3.1**.
- 183 **2.2.** Characterization of the biomass composition
- 184 **2.2.1.Moisture**

Moisture content of the biomass was determined in triplicate by vacuum-drying as described by Nguyen *et al.* (2016) [22]. Briefly, 20 mg of microalgal biomass was dried using a vacuum oven (UNIEQUIP 1445-2, Planegg, Germany), with sequential drying steps for 1 h at 0.8, 0.6 and 0.4 bar and for 30 min at 0.2 bar. The difference in weight before and after drying was expressed as a percentage, representing the moisture content. The average moisture content was used in the calculation of the chemical components, being expressed in percentage of total dry matter.

191 **2.2.2.Lipids**

192 Lipid content was determined according to the method optimized by Ryckebosch, Muylaert & Foubert 193 (2012) [23]. Briefly, 4 mL methanol, 2 mL chloroform and 0.4 mL water were added to 100 mg of dry biomass and the samples were homogenized. Then, 2 mL chloroform and 2 mL water were added and the 194 samples were again homogenized. After centrifugation (10 min, 750g, 25 °C), the upper aqueous layer was 195 discarded, while the lower solvent layer was collected. The remaining pellet was re-extracted with 4 mL of 196 chloroform:methanol (1:1 v/v) and centrifuged (10 min, 750g, 25 °C). The solvent phase was collected and 197 the extraction procedure was repeated on the pellet. All solvent layers were combined and filtered through 198 a layer of sodium sulphate to remove remaining water. The solvent was removed by rotary evaporation and 199 200 lipids were quantified gravimetrically. Lipid content was determined in triplicate.

201 **2.2.3.Proteins**

Protein content was determined by the Dumas method. Approximately 1-2 mg of biomass was transferred
to tin capsules and total nitrogen content was analyzed using an elemental analyzer (Carlo-Erba EA1108,
Thermo Scientific, Waltham, MA, USA). Protein content was estimated from total nitrogen content,
multiplied by an overall conversion factor of 4.78, as proposed by Lourenço *et al.* (2004) [5]. The analysis
was performed in triplicate.

207 2.2.4.Storage polysaccharides

208

2.2.4.1. Starch, floridean starch and glycogen

According to literature, starch is present in C. vulgaris, T. chuii and Schizochytrium sp. [24–26], whereas 209 R. cruentum contains floridean starch as SPS [27]. Lastly, the cyanobacterium A. platensis contains 210 glycogen [28]. Since all these SPS are polyglucans with α -1,4 and α -1,6-linkages, they were quantified 211 212 using the same procedure, based on the method of Pleissner & Eriksen (2012) [29]. Briefly, 50-100 mg 213 biomass was washed in MOPS buffer (3-(N-morpholino)propanesulfonic acid, 55 mM, pH 7) and cells were subsequently disrupted using ultra high pressure homogenization (UHPH) at 250 MPa (Stansted Fluid 214 Power SPCH-10, Harlow, United Kingdom). A single pass was applied for most microalgae, except for the 215 216 more rigid C. vulgaris requiring two passes for full cell disruption. After addition of ethanol and dimethyl

sulfoxide, the mixtures were boiled for 15 min at 100 °C to gelatinize (floridean) starch. Samples were then 217 incubated with thermostable α -amylase from *Bacillus licheniformis* (519 U/mg, Sigma-Aldrich) for 10 min 218 at 90 °C, followed by incubation with amyloglucosidase from Aspergillus niger (70 U/mg, Sigma-Aldrich) 219 220 for 1 h at 50 °C. Both enzymes were added to achieve a ratio of 10 U/mg biomass. The mixtures were subsequently centrifuged (15 min, 5000g, 25 °C) and the glucose content in the supernatant was measured 221 using the glucose oxidase method [30]. SPS content was finally calculated multiplying the glucose content 222 by 0.9, taking into account the addition of a water molecule during enzymatic hydrolysis of the polyglucans. 223 The analysis was done in triplicate. 224

225

2.2.4.2. Chrysolaminarin

Chrysolaminarin content was determined for Nannochloropsis sp., T. lutea, P. tricornutum, O. aurita and 226 D. lutheri, using a modified version of the method described by Granum & Myklestad (2002) [31]. Dry 227 biomass was suspended in demineralized water, after which the cells were disrupted using UHPH at 228 250 MPa. Whereas two passes were used for full disruption of Nannochloropsis sp. cells, a single pass at 229 250 MPa was sufficient for the other microalgae species. The disrupted microalgae were lyophilized and 230 β -1,3-glucans were subsequently extracted with 0.05 M sulfuric acid at 60 °C for 10 min. The extract was 231 filtered using GF/C glass fiber filters and the glucose content was determined using the phenol-sulfuric 232 acid method [32]. Glucose content was multiplied by 0.9 to quantify the chrysolaminarin content. 233 Extraction and quantification of chrysolaminarin was performed in triplicate. 234

235 **2.2.5.** Ash an

2.2.5.Ash and minerals

Approximately 20 mg of biomass was ashed in a muffle furnace (Nabertherm Controller P330, Lilienthal,
Germany), operating for 24 h at 550 °C. The ashes were weighed and dissolved in 10 mL of ultrapure water
(organic free, 18 MΩ cm resistance). These solutions were then acidified with 0.1 mL of 65% HNO₃ and
filtered through a 0.45 µm syringe filter (Chromafil[®] A-45/25, Macherey-Nagel, Duren, Germany).
Mineral composition was determined using inductively coupled plasma optical emission spectrometry
(ICP-OES, Perkin-Elmer Optima 3300 DV, Norwalk, CT, USA). Ash content and mineral composition
were determined in triplicate.

243 2.2.6.Cell wall related polysaccharides

- The extraction and characterization of CWPS and EPMS is described later in section 2.3. The total content of CWPS and EPMS was calculated from the monosaccharide and uronic acid content using **Eq. 2**, taking into account the addition of a water molecule to the dehydrosugar in a polysaccharide chain (**Eq. 1**):
- 247

 $(dehydrosugar)_n + n \cdot H_2 0 \rightarrow n \cdot monosaccharide$ (Eq. 1)

248 Polysaccharide (%) = $\sum \left(\frac{MM_{dehydrosugar}}{MM_{monosaccharide}} \times monosaccharide (%) \right) + \sum \left(\frac{MM_{dehydro\,uronic\,acid}}{MM_{uronic\,acid}} \times uronic\,acid (%) \right)$ (Eq. 2)

249 2.3. Composition of cell wall related polysaccharides

250 **2.3.1.Extraction**

A schematic representation of the procedure for extraction of EPMS and CWPS is shown in **Fig. 1**.

252 **2.3.1.1.** Extraction of extracellular polymeric substances

In the current study, cell wall related polysaccharides are extracted from dry biomass. Therefore, our definition of EPMS only includes the polymeric material of the external layer that is still attached to the dried cells, that can be secreted when resuspending the biomass in a simulated growth medium. The presence of an extracellular layer surrounding the cells in the dry biomass was visualized for *P. cruentum* (as indicated by the arrows in **Fig. A-1**, supplementary material).

A procedure was implemented for extraction of EPMS based on methods of Hanlon et al. (2006) and Patel 258 et al. (2013) [33,34]. Dry biomass (1.5 g) was suspended in 30 mL of saline solution at pH 7.5 to mimic 259 cultivation conditions of the microalgae. All marine microalgae were suspended in 2.5% w/v NaCl, 260 whereas the freshwater species C. vulgaris was suspended in 0.02% w/v NaCl. All suspensions were 261 incubated for 16 h at 25 °C, allowing the EPMS to dissolve into the medium. Subsequently, suspensions 262 263 were centrifuged (10 min, 10000g, 4 °C), followed by a second centrifugation step of the supernatants (30 min, 17000g, 4 °C) to completely separate the biomass from the medium. The integrity of the cells in 264 the residual biomass was confirmed using light microscopy (Olympus BX-51, Optical Co.Ltd, Tokyo, 265 Japan). The successful removal of EPMS was clearly observed for P. cruentum, since the extracellular 266

layers observed in the dry biomass had disappeared after the extraction of EPMS (Fig. A-2, supplementary
information). Cold ethanol (95% v/v) was added to the resulting supernatant to precipitate EPMS, ensuring
a final ethanol concentration higher than 70% v/v. The solution was vacuum filtered using MN 615 filter
paper and the insoluble residue was extensively dialyzed against demineralized water for 48 h
(Spectra/Por[®], MWCO 3.5 kDa, Spectrum Laboratories, CA, USA). Finally, the dialyzed EPMS extracts
were lyophilized (Alpha 2-4 LSC plus, Christ, Osterode, Germany).

273

2.3.1.2. Extraction of cell wall polysaccharides

In the present study, CWPS are considered the structural polysaccharides that are strongly bound in the cell wall. In contrast with EPMS, CWPS are not secreted into the surrounding environment, but play an important structural role in microalgal cells such as providing resistance to turgor pressure. CWPS were extracted by isolating an alcohol insoluble residue according to McFeeters & Armstrong (1984) [35], with the additional removal of lipids, SPS and proteins as demonstrated by several authors [20,36,37].

The residual biomass, i.e. the pellets obtained in the two centrifugation steps in section 2.3.1.1, was 279 resuspended in 100 mL MOPS buffer (55 mM, pH 7). Cells were disrupted using UHPH at 250 MPa, 280 applying four passes for the rigid microalgae *Nanhochloropsis* sp. and *C. vulgaris*, while a single pass was 281 used for the other microalgae species. Cold ethanol was added to the suspensions (> 70% v/v), the mixtures 282 were centrifuged (10 min, 10000g, 4 °C) and the pellet was recovered. Lipids were removed by a 283 hexane:isopropanol (3:2 v/v) extraction, based on the method of Ryckebosch et al. (2013) [38]. 284 Hexane: isopropanol (3:2) was chosen as extraction solvent instead of chloroform: methanol (1:1), as the 285 latter solvent mixture can also remove some non-lipid substances from the microalgal biomass [39]. In 286 short, 30 mL of hexane: isopropanol (3:2) was added to the pellet, mixed and centrifuged (10 min, 900g, 287 20 °C) to remove the upper solvent layer. This lipid extraction was repeated to obtain a defatted pellet. 288 289 Afterwards, SPS were enzymatically removed using either endo- β -1,3-glucanase or a combination of α -290 amylase and amyloglucosidase, depending on the type of SPS [37,40]. For removal of (floridean) starch or 291 glycogen, the defatted pellet was resuspended in sodium acetate buffer (140 mM, pH 4.5) and heated for 292 10 min at 100 °C to gelatinize the (floridean) starch. The low pH of 4.5 was used to avoid possible

293 degradation reactions of CWPS occurring at high temperatures, such as beta-elimination of pectic 294 polysaccharides [41]. After gelatinization, the pH was adjusted to 5.2 and α -amylase from *Bacillus* sp. 295 (839 U/mg, Sigma-Aldrich) and amyloglucosidase from A. niger (300 U/mL, Sigma-Aldrich) were both 296 added in a ratio of 1 U/mg initial biomass. The mixtures were incubated for 4 h at 60 °C. To remove chrysolaminarin, the defatted pellet was resuspended in sodium acetate buffer (100 mM, pH 4.5) and 297 heated for 10 min at 100 °C. Afterwards, pH was adjusted to 5 and endo-β-1,3-glucanase of barley 298 (2500 U/mL, Megazyme) was added to achieve 400 U/g initial biomass and the mixture was incubated for 299 4 h at 40 °C. After enzymatic incubations, cold ethanol was added to all samples (>70% v/v), the 300 suspensions were centrifuged (10 min, 10000g, 4 °C) and the pellets were recovered. Lastly, proteins were 301 removed enzymatically. Therefore, pellets were resuspended in phosphate buffer (80 mM, pH 7.5) and 302 Subtilisin A protease from B. licheniformis (12 U/mg, Sigma-Aldrich) was added to achieve 50 U/g initial 303 biomass. The mixtures were incubated for 1 h at 60 °C and after addition of cold ethanol (> 70% v/v), they 304 were centrifuged (10 min, 10000g, 4 °C). The pellet was finally washed in acetone, vacuum filtered and 305 306 dried overnight at 40 °C. This residue was considered as CWPS.



307

- 308 Fig. 1 Schematic overview of the extraction of extracellular polymeric substances (EPMS) and cell wall polysaccharides 309 (CWPS).
- 310

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2.3.2.Quantification 311

Quantification of monosaccharides and uronic acids 312 2.3.2.1.

Monosaccharide and uronic acid composition of CWPS and EPMS were determined according to De Ruiter 313 314 et al. (1992) [42]. Polysaccharides were first hydrolyzed using methanolysis combined with trifluoroacetic acid (TFA) hydrolysis. The resulting monosaccharides and uronic acids were then quantified by high 315 performance anion exchange chromatography combined with pulsed amperometric detection (HPAEC-316 317 PAD) Methanolysis and TFA hydrolysis was performed as described by De Ruiter et al. (1992) [42]. Briefly, 318 sample was dissolved in ultrapure water (organic free, 18 M Ω cm resistance) in a concentration of 2 mg/mL,

- of which 40 µL was transferred to a pyrex tube and dried by N2 evaporation at 45 °C. Then, 2 mL of 2 M 320
- 321 methanolic HCl was added to the sample and the mixtures were incubated for 16 h at 80 °C. The solvent

322 was removed by N2 evaporation at 30 °C. Subsequently, 2 mL of 2 M TFA was added to the tubes and 323 incubated for 1 h at 121 °C. After removing the solvent by N₂ evaporation at 45 °C, the hydrolysate was dissolved in ultrapure water (organic free, 18 M Ω cm resistance) and filtered through a 0.45 µm syringe 324 filter (Chromafil[®] A-45/25, Macherey-Nagel, Duren, Germany). The hydrolysis was performed in triplicate. 325 326 Monosaccharides and uronic acids in the hydrolysates were identified and quantified by HPAEC-PAD, as described by Jamsazzadeh Kermani et al. (2014) [43]. A Dionex system (DX600) equipped with a GS50 327 gradient pump, a CarboPacTM PA20 column (150×3 mm), a CarboPacTM PA20 guard column (30×3 mm) 328 and an ED50 electrochemical detector were used (Dionex, Sunnyvale, CA, USA). The detector was 329 equipped with a reference pH electrode (Ag/AgCl) and a gold electrode. Equilibration was performed for 330 10 min with 100 mM NaOH, after which 10 µL of hydrolysate was injected. First, isocratic elution of 331 monosaccharides was performed for 20 min. In order to achieve a complete chromatographic separation of 332 all monosaccharides analyzed, each sample was eluted using 0.5 mM NaOH as well as 15 mM NaOH. 333 Secondly, uronic acids were isocratically eluted with 500 mM NaOH for 10 min. 334

Mixtures of commercial sugar standards (D-glucose, D-galactose, D-xylose, D-mannose, L-rhamnose, Larabinose, L-fucose, D-ribose, D-glucosamine, D-galacturonic acid and D-glucuronic acid) at varying concentrations (1-10 ppm) were used as external standards for identification and quantification. These standards were also subjected to the above-mentioned hydrolysis in order to correct for monosaccharide degradation during methanolysis and TFA hydrolysis.

340

2.3.2.2. Quantification of sulfate groups

Sulfate groups in the polysaccharide samples were quantified using the barium chloride-gelatin method of Dodgson & Price (1962) [44]. Briefly, 2-5 mg of sample was hydrolyzed with 1 M HCl for 5 h at 100 °C. The hydrolyzed sample was then incubated with 3% trichloroacetic acid and barium chloride-gelatin reagent for 15 min at 25 °C, and the absorbance was measured at 360 nm. A standard curve was prepared with K₂SO₄. Samples were corrected for UV-absorbing materials formed during hydrolysis, which were determined in absence of barium as described by Dodgson & Price (1962) [44]. The analysis was performed in triplicate.

2.4. Statistical analysis

- 349 The data obtained are presented as the average of three measurements \pm standard error. Differences in 350 biomass composition of different microalgae species were statistically analyzed using one-way ANOVA combined with Tukey's test for multiple comparison (P < 0.05) with JMP statistical software (JMP Pro 12, 351
- 352 Cary, NC, USA).
- 353

3. Results and discussion 354

- 355 3.1. Characterization of the biomass composition
- Results obtained from the biochemical characterization of the microalgal biomasses are presented in 356 Table 1. These results are expressed as a percentage of dry matter, as they were corrected for the moisture 357 contents (1.5 - 8.2%). Generally, substantial differences were found in the biomass composition amongst 358 different species, with each microalga showing a distinct biomass nutrient profile. 359
- **Table 1** Biochemical composition of microalgal biomasses, expressed as the average percentage of dry matter (%) \pm standard deviation of triplicate measurements. (SPS: storage polysaccharides; CWPS: cell wall polysaccharides; EPS: extracellular polysaccharides; n.d.: not detected). The results were compared statistically (one-way ANOVA). Significant differences (Tukey 360 361 362 363 test, P < 0.05) within each column are indicated with different letters (n = 3).

	Lipids	Proteins	SPS	CWPS	EPS	Ash
P. cruentum	11.5 ± 1.0 e	28.2 ± 1.2 e	2.1 ± 0.1 e	$9.6\pm0.3~\textbf{b}$	$2.6\pm0.2~a$	17.9 ± 0.6 a
C. vulgaris	6.6 ± 1.5 f	39.4 ± 0.3 b	1.8 ± 0.1 e	$9.3\pm0.7~\textbf{b}$	$0.5\pm0.1~c$	$6.7\pm0.1~\textbf{d}$
T. chuii	11.9 ± 1.1 e	31.1 \pm 0.3 d	7.8 ± 0.2 bc	17.0 ± 1.0 a	n.d.	$14.5\pm0.3~\text{bc}$
P. tricornutum	17.1 ± 0.9 d	29.4 ± 0.4 de	$5.2\pm0.7~\textbf{d}$	$5.3\pm0.2~\textbf{d}$	n.d.	$15.9\pm0.9~\boldsymbol{b}$
O. aurita	12.8 ± 1.1 e	$20.2\pm0.1~\mathbf{f}$	$21.7\pm0.6~a$	7.4 ± 0.3 c	$1.2\pm0.1~\textbf{b}$	$14.1\pm0.6~c$
Nannochloropsis sp.	31.7 ± 1.3 b	35.1 ± 0.8 c	$5.9\pm0.5~\text{cd}$	3.8 ± 0.1 e	n.d.	$6.1\pm0.9~\textbf{d}$
Schizochytrium sp.	73.9 ± 1.7 a	$10.2\pm0.1~{\rm g}$	6.3 ± 1.1 bcd	6.9 ± 0.4 c	n.d.	3.4 ± 0.3 e
T. lutea	$27.2 \pm 1.6 $ c	$29.8\pm0.7~\text{de}$	7.3 ± 0.4 bc	$3.5\pm0.1~f$	n.d.	$13.5\pm0.6~c$
D. lutheri	31.5 ± 1.4 b	$21.9\pm0.3~{\rm f}$	7.2 ± 0.8 bc	$10.2\pm0.5~\textbf{b}$	n.d.	13.0 ± 0.4 c
A. platensis	5.5 ± 0.5 f	47.4 ± 0.1 a	$8.1\pm0.6~\textbf{b}$	$9.7\pm0.7~\boldsymbol{b}$	0.7 ± 0.1 c	$4.5\pm0.2~e$

364

A large diversity is observed in terms of lipid content among all microalgae. A. platensis and C. vulgaris 365 presented low lipid contents, making up 5.5% and 6.6% of the biomass, respectively. Similar lipid contents 366 367 were found in literature for biomasses of Arthrospira sp. (3.6 - 7.5%) [1,3,45,46], while a wider range of fat contents was observed for C. vulgaris. Although some authors reported similar values of 5% oil for 368 C. vulgaris [1,45], higher lipid contents (13 - 20%) have also been found [3]. This could be related to 369

370 differences in cultivation conditions, as it is known for Chlorophyceae (and oleaginous microalgae in 371 general) that biomass composition can be selectively modified by adapting the cultivation conditions, such 372 as nitrogen depletion and temperature [47]. In contrast, lipid content and composition of cvanobacteria are 373 less affected by changes in cultivation conditions [8], confirming the smaller range of lipid contents found 374 in literature for Arthrospira biomasses. In contrast to the former two species, Nannochloropsis sp., D. lutheri, T. lutea and P. tricornutum can be considered as lipid-rich microalgae, presenting lipid contents 375 of 17 - 32%. In fact, these microalgae species are considered among the most interesting sources of $\omega 3$ -LC-376 PUFA and similar values for total lipid content have been reported in literature [48]. In addition, a very high 377 lipid content was observed for Schizochytrium sp., making up 74% of the dry biomass. It was 378 microscopically confirmed that Schizochytrium sp. cells were completely filled with lipid bodies, as 379 previously reported by Morita et al. (2006) [49]. Generally, lipid contents of Schizochytrium sp. are 380 somewhat lower (32 - 50%) [50,51], although a similar content of 73% has been reported by Liang *et al.* 381 (2010) [52]. Finally, intermediate lipid contents (11-13%) were observed for P. cruentum, T. chuii and 382 O. aurita, corresponding to literature reports [48,53], 383

Proteins generally make up an important part of the biomass, being more than 20% for the investigated 384 microalgae (except for Schizochytrium sp.). The highest protein contents were observed for A. platensis and 385 C. vulgaris, comprising 47% and 39% of the dry biomass, respectively. These two species have been 386 387 commercialized as a source of proteins, containing high amounts of essential amino acids [2]. Although some authors presented similar protein contents, approximately 38% for C. vulgaris and 39-46% for 388 A. platensis [1,46], higher values (up to 70% for Arthrospira sp.) have also been reported [3,6,54,55]. 389 However, it should be noted that part of this variation can be ascribed to different analytical approaches, 390 such as the use of different correction factors for non-protein nitrogen in elemental analyses [5,6]. Large 391 amounts of proteins were also found for lipid-rich microalgae species (D. lutheri, T. lutea, 392 Nannochloropsis sp. and P. tricornutum), representing 22 - 35% of the biomass. Generally, the protein 393 394 values obtained are in agreement with those reported in literature [1,6,26,56–58].

In contrast to most studies on microalgal biomass profiles, different types of carbohydrates were quantified
 separately in this study. According to literature, four types of SPS are present in the selected microalgae,

397 being either starch, floridean starch, glycogen or chrysolaminarin. The three former SPS were analyzed by 398 enzymatic hydrolysis, while the latter was extracted chemically, followed by quantification of the 399 hydrolyzed glucose monomers. Relatively low SPS contents were observed for all microalgae species 400 (2-8%), with the exception of *O. aurita* biomass presenting 22% of chrysolaminarin. For biomasses of 401 P. cruentum, Schizochytrium sp. and A. platensis, the values obtained correspond with previously reported 402 SPS contents [26,28,59]. In contrast, the amount of SPS in C. vulgaris and P. tricornutum biomass differs from those in literature [24,60], although only one article was encountered for each microalga reporting SPS 403 404 contents grown under standard conditions. Typically, SPS are very sensitive to cultivation conditions and the time of harvesting. Whereas SPS contents are usually low in the exponential growth phase, they rapidly 405 increase when nutrients are exhausted in the stationary phase. Similarly, cultivation in a nutrient depleted 406 medium leads to accumulation of SPS [25,57,61]. The low amounts of SPS in these commercially obtained 407 408 biomasses therefore suggest that cells were harvested in the late exponential phase, preserving the nutritional quality of the biomass in terms of protein content and lipid composition. Biomass of O. aurita was cultivated 409 410 in open raceway ponds and could have experienced more stress in terms of nutrient heterogeneity or competition with other organisms, possibly resulting in higher chrysolaminarin contents compared to the 411 412 other microalgae species.

Two fractions of cell wall related polysaccharides were extracted from the microalgal biomass. On the one 413 414 hand, EPMS were comprised of polymers on the outer layer of the cell wall, which dissolved into the medium without disruption of the microalgal cells. EPMS were only obtained for four microalgae species 415 (P. cruentum, O. aurita, A. platensis and C. vulgaris), implying that the other microalgae do not have a layer 416 417 of extracellular polymers. These observations correspond to our previous study, in which an increased viscosity of the continuous serum phase was ascribed to the presence of extracellular polymers for these 418 four microalgae species. In addition, the absence of EPMS in *Nannochloropsis* sp. and *P. tricornutum* was 419 420 confirmed by that study, since no increased serum viscosity was observed for suspensions of these microalgae species [62]. The highest content of EPMS was found in *P. cruentum*, a red microalga which is 421 422 known for its sulfated extracellular polysaccharides [13,63]. In addition, EPMS were also obtained for 423 A. platensis, C. vulgaris and O. aurita. While EPMS have been previously reported for A. platensis [64,65]

424 and C. vulgaris [66,67], no information was found on the presence of EPMS in O. aurita. Furthermore, the 425 absence of EPMS in Nannochloropsis sp., T. lutea and the fusiform type of P. tricornutum as seen in 426 Table 1 was confirmed by other authors [68,69]. In contrast, although EPMS have been reported in literature 427 for *Tetraselmis* sp. and *Schizochytrium* sp. [68,70], they were not observed in this study. It should be noted that EPMS were extracted starting from dried biomass. Therefore, the yield obtained in this study is 428 probably lower than when EPMS are directly extracted from the cultivation medium, due to losses of EPMS 429 during harvesting steps. Although part of the extracellular polymers might have been secreted into the 430 cultivation medium, the presence of unsecreted extracellular layers surrounding the microalgal cells was 431 visualized using microscopy (Fig. A, supplementary material). Even if EPMS were partially lost during 432 harvesting of the microalgae, the data obtained are relevant in the context of food applications, as microalgal 433 biomass will be delivered in a dry form for use in food products to guarantee long term storage stability of 434 435 these functional ingredients. In addition, Table 1 shows the content of extracellular polysaccharides (EPS), quantified as the sum of monosaccharides and uronic acids in the extracts, without considering other 436 437 polymers such as proteins or co-extracted molecules like minerals and nucleic acids. However, when considering the weights of extracted EPMS, including other extracellular polymers, yields of 2.7 - 8.3%438 were obtained for these four microalgae species. Even though proteins were not quantified in EPMS samples 439 (since they are accounted for in the total protein content of the biomasses), it can be inferred that they are 440 441 present in all EPMS of the four microalgae species.

On the other hand, CWPS, which are structural polysaccharides constituting the cell wall, were extracted 442 after removal of EPMS followed by disruption of the cells. Generally, CWPS make up around 10% of the 443 microalgal biomass [54]. However, it is known that some microalgae contain a cell wall comprised of other 444 445 (non-polysaccharide) substances, resulting in a reduced amount of polysaccharides in the cell wall. For instance, several authors have reported the presence of algaenan in the Nannochloropsis sp. cell wall, a 446 resistant aliphatic biopolymer composed of ether-linked long alkyl chains of esterified monomers [71,72]. 447 448 This could explain the low CWPS content found in *Nannochloropsis* sp. compared to the other microalgae 449 species. Similarly, the cells of diatoms are surrounded by a silicified cell wall, called a frustule, which is 450 coated with a layer of organic material [73]. This structure is likely present in O. aurita and was visualized

451 by microscopic images in our previous study [62]. Although P. tricornutum is also classified as a diatom, 452 not all morphotypes contain this siliceous skeleton. In fact, the ovoid form is the only morphotype presenting 453 true silica valves, while the other morphotypes are poor in silica [73]. Thus, the lower amount of CWPS in 454 P. tricornutum cannot be attributed to the presence of a frustule. Finally, the lowest amount of CWPS was 455 found in T. lutea, representing only 3.5% of the total biomass. It is known that Isochrysis sp. cells are easily 456 disrupted, which is related to its weak cell wall structure [74]. Some authors even doubt the presence of a 457 cell wall in Isochrysis sp., claiming that this microalga species only contains a plasma membrane around 458 the cells [75]. However, the results obtained in this study imply that a cell wall layer of polysaccharides is present in T. lutea, but the low CWPS content suggests a rather small contribution, possibly related with its 459 460 low resistance to mechanical disruption.

Finally, it was observed that all microalgal biomasses contain considerable amounts of minerals. Clear 461 differences are observed between marine and freshwater microalgae. Whereas most of the marine 462 microalgae presented high ash contents due to accumulation of minerals from the cultivation medium 463 464 (13-18% of the total biomass), the freshwater species C. vulgaris and A. platensis only presented ash contents of 6.7% and 4.5%, respectively. The lowest ash content was observed for the heterotrophic 465 Schizochytrium sp., representing only 3.4% of the total biomass, even though a higher value (11% ash) was 466 previously reported for biomass of S. limacinum [76]. Nevertheless, since Sun et al. (2014) stated that high 467 468 lipid contents coincide with decreased ash contents in Schizochytrium sp. [77], it is not surprising that a low ash content was observed in this study. In addition, a low ash content was also observed in 469 Nannochloropsis sp. biomass, even though this marine species was cultivated in a salt-rich medium. 470 Nevertheless, similar values (6 - 10%) have been previously reported by other authors [58,78]. In terms of 471 472 mineral composition, each microalga species presented a typical mineral profile, as shown in Table 2. Generally, marine microalgae were rich in the monovalent cations sodium and potassium. In fact, sodium 473 474 represented more than 60% of all minerals in biomasses of Nannochloropsis sp., Schizochytrium sp., 475 T. lutea and D. lutheri. In contrast, the freshwater species A. platensis and C. vulgaris contained low 476 amounts of sodium, representing less than 10% of the analyzed minerals in these biomasses. From a 477 nutritional point of view, P. tricornutum has the most interesting mineral profile, possessing very high

amounts of calcium and iron. Given that iron deficiency is considered as the most prevalent single nutritional 478 deficiency in the world [79], P. tricornutum biomass shows great potential as an iron source in the human 479 diet. In addition, this microalga presented substantial amounts of magnesium and manganese and a relatively 480 low amount of sodium, which was also observed by Rebolloso-Fuentes et al. (2001) [80]. Biomasses of 481 P. cruentum and T. chuii also contained considerable amounts of calcium and iron, however they may be of 482 483 lower interest for nutritional purposes due to higher amounts of sodium. Sodium consumption is after all associated with increased blood pressure and cardiovascular diseases and a reduced sodium intake is 484 therefore recommended [81]. Nevertheless, these three microalgae species, together with the freshwater 485 species A. platensis and C. vulgaris, can generally be considered as good sources of minerals for human 486 487 nutrition.

488Table 2 Mineral composition of microalgal biomasses, expressed as the average \pm standard deviation of triplicate measurements.489The results were compared statistically (one-way ANOVA). Significant differences (Tukey test, P < 0.05) within each column are490indicated with different letters (n = 3).

	Na	K	Ca	Mg	Fe	Zn	Mn	Cu
	(mg/g)	(mg/g)	(mg/g)	(mg/g)	(mg/100 g)	(mg/100 g)	(mg/100 g)	(mg/100 g)
P. cruentum	$27.1 \pm 1.4 \ \boldsymbol{a}$	17.5 ± 2.7 ab	13.3 ± 0.9 c	9.8 ± 0.5 a	131.1 ± 3.3 b	$6.9\pm0.5~a$	$8.7\pm0.4~\boldsymbol{b}$	0.9 ± 0.1 ab
C. vulgaris	$2.4\pm0.2~\textbf{e}$	$12.2\pm0.2~\textbf{cde}$	4.1 ± 0.6 e	1.1 ± 0.1 f	11.7 ± 6.6 ef	$1.8\pm0.8~\text{de}$	$1.6\pm0.8~\text{de}$	$0.6\pm0.1~\boldsymbol{b}$
T. chuii	$10.1\pm0.9~\textbf{cd}$	11.5 ± 0.7 cde	17.7 ± 0.8 b	3.6 ± 0.2 d	$49.0\pm6.4~\textbf{d}$	$1.3\pm0.1~\text{ef}$	$8.0\pm1.0~\boldsymbol{b}$	$0.9\pm0.2~~\textbf{ab}$
P. tricornutum	$7.1 \pm 1.8 ~\textbf{d}$	16.2 ± 1.3 abc	50.1 ± 3.2 a	3.7 ± 0.2 d	$240.6\pm14.0~\textbf{a}$	$2.9\pm0.3~\text{cd}$	$12.0\pm0.7~\textbf{a}$	$2.0\pm1.2~\textbf{a}$
O. aurita	$18.8\pm2.3~\textbf{b}$	19.1 ± 2.8 a	4.4 ± 0.5 de	3.5 ± 0.3 d	$19.2\pm2.7~e$	$0.3\pm0.01~{\rm f}$	11.9 ± 1.4 a	$0.2\pm0.02~\textbf{b}$
Nannochloropsis sp.	$12.8\pm0.9~c$	5.2 ±1.1 fg	1.1 ± 0.2 ef	$1.9\pm0.2~\textbf{e}$	7.3 ± 3.0 ef	$2.3\pm0.5~\text{de}$	$2.2\pm0.4~\text{cd}$	$0.5\pm0.2~\textbf{b}$
Schizochytrium sp.	$11.0\pm1.0~\textbf{c}$	4.3 ± 2.4 g	0.4 ± 0.2 f	$0.8\pm0.1~\boldsymbol{a}$	$1.5\pm0.7~\text{ef}$	$3.9\pm0.01~\text{bc}$	$0.1\pm0.02~\textbf{e}$	$0.1\pm0.01~\textbf{b}$
T. lutea	29.1 ± 0.9 a	10.0 ± 2.1 def	1.4 ± 0.1 ef	$1.9\pm0.1~\textbf{e}$	$1.0\pm0.4~\mathbf{f}$	$4.8\pm0.3~\textbf{b}$	$0.7\pm0.5~\text{de}$	$0.1\pm0.03~\textbf{b}$
D. lutheri	27.7 ± 1.7 a	7.6 ± 0.7 efg	3.7 ± 0.9 ef	$6.1\pm0.3~\textbf{b}$	$42.0\pm5.7~\textbf{d}$	$1.9\pm0.4~\text{de}$	$1.4\pm0.3~\text{de}$	$0.4\pm0.1~\textbf{b}$
A. platensis	2.8 ± 0.2 e	14.0 ± 0.9 bcd	$7.7\pm0.3~\textbf{d}$	$4.6\pm0.2~c$	93.3 ± 6.1 c	$1.9\pm0.5~\text{de}$	$3.8\pm0.1~c$	$0.5\pm0.03~\textbf{b}$
1		X						

492 **3.2. Composition of cell wall related polysaccharides**

The aim of this study was to apply a universal procedure for extraction of the total cell wall related polysaccharides followed by characterization of their monosaccharide and uronic acid composition. **Table 3** presents the monosaccharide and uronic acid profiles of CWPS of the investigated microalgae, with each sugar expressed as a percentage of the total amount of monosaccharides and uronic acids. Diverse monosaccharide profiles were observed for the different microalgae species. All microalgae species presented at least five different monosaccharides in the CWPS extracts, suggesting that their cell walls are composed of heteropolymers or multiple types of polysaccharides. Glucose, galactose, xylose and mannose

500 were generally the most abundant monosaccharides, while the ratio of galacturonic and glucuronic acid was 501 strongly dependent on the microalga species. The composition of the EPMS is shown in Table 4 for the four microalgae P. cruentum, C. vulgaris, O. aurita and A. platensis. As mentioned before, no EPMS could 502 be extracted for the other microalgae species. Similar to CWPS, EPMS are considered 503 heteropolysaccharides, containing both neutral monosaccharides and uronic acid residues. Finally, the 504 505 sulfate content of both polysaccharide fractions is given in Table 5. All microalgae contained sulfated polysaccharides, although the degree of sulfation was generally low (0.6 - 14%). However, it should be 506 noted that the obtained cell wall fractions were not further purified and still contained some impurities such 507 as ash and little amounts of proteins. As a consequence, sulfate contents of the purified cell wall related 508 polysaccharides might be somewhat higher than the data presented in Table 5. Even though no detailed 509 information was provided on the cultivation and harvesting conditions, the data obtained are expected to be 510 representative for the cell wall related polysaccharides of microalgae of the investigated species. Although 511 the amount of CWPS might be dependent on the cultivations conditions applied, it was previously shown 512 513 that the composition of neutral monosaccharides was constant for Chlorella species when cultivated under different growth conditions, such as elevated CO₂ concentration and nitrogen limitation [37,82]. The impact 514 of different cultivation conditions on uronic acids in cell wall related polysaccharides might be different, as 515 Cheng, Labavitch & VanderGheynst (2015) indicated an increased uronic acid content under elevated CO₂ 516 517 supply [82]. However, to the best of our knowledge, no reports were found on uronic acids in the microalgal cell wall being affected by other cultivation parameters. 518

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Table 3 Monosaccharide and uronic acid composition in cell wall polysaccharides (CWPS) of microalgae, expressed as the average percentage of total monosaccharides and uronic acids (%) \pm standard deviation of triplicate measurements (n = 3). (n.d.: not detected).

	P. cruentum	C. vulgaris	T. chuii	P. tricornutum	O. aurita
Glucose	30.5 ± 1.1	41.5 ± 1.8	28.9 ± 4.1	4.4 ± 2.4	11.1 ± 0.8
Galactose	22.4 ± 0.3	8.6 ± 0.2	5.7 ± 0.9	3.8 ± 0.9	35.7 ± 0.6
Xylose	27.8 ± 2.4	n.d.	5.3 ± 0.7	14.3 ± 0.1	9.3 ± 0.6
Mannose	9.3 ± 0.5	34.8 ± 7.6	41.3 ± 3.4	46.4 ± 1.3	17.1 ± 1.2
Rhamnose	n.d.	2.7 ± 0.1	1.0 ± 0.2	8.9 ± 0.8	3.2 ± 1.1
Arabinose	n.d.	n.d.	0.8 ± 0.2	n.d.	n.d.
Fucose	n.d.	n.d.	0.7 ± 0.7	2.5 ± 0.3	12.4 ± 0.6
Ribose	n.d.	1.9 ± 0.1	n.d	2.9 ± 1.2	1.3 ± 0.6
Glucosamine	1.8 ± 1.3	2.9 ± 0.1	n.d	n.d.	n.d.
Galacturonic acid	3.9 ± 0.9	3.3 ± 0.2	15.1 ± 1.3	2.9 ± 0.1	4.0 ± 2.6
Glucuronic acid	4.3 ± 0.1	4.3 ± 0.1	1.2 ± 0.2	13.9 ± 0.1	5.9 ± 1.0

523

524 Table 3 (continued)

Table 3 (contin	ued)		. (-01	
	Nannochloropsis sp.	Schizochytrium sp.	T. lutea	D. lutheri	A. platensis
Glucose	75.8 ± 4.9	33.1 ± 1.6	22.7 ± 1.2	82.2 ± 3.3	49.8 ± 5.8
Galactose	6.4 ± 0.4	29.0 ± 3.5	14.8 ± 0.9	n.d.	3.8 ± 0.3
Xylose	3.5 ± 0.4	14.3 ± 0.2	9.6 ± 1.3	4.9 ± 2.7	n.d.
Mannose	4.7 ± 0.1	20.5 ± 3.4	16.4 ± 0.7	6.2 ± 0.3	29.8 ± 3.7
Rhamnose	3.0 ± 0.1	n.d.	1.3 ± 0.2	n.d.	6.7 ± 0.9
Arabinose	n.d.	n.d.	20.3 ± 0.9	3.4 ± 0.7	n.d.
Fucose	2.1 ± 0.2	n.d.	4.2 ± 0.4	0.8 ± 0.4	n.d.
Ribose	4.5 ± 0.1	n.d.	4.0 ± 0.5	1.3 ± 0.6	n.d.
Glucosamine	n.d.	3.1 ± 1.9	n.d.	n.d.	2.1 ± 0.3
Galacturonic acid	n.d.	n.d.	4.1 ± 3.3	n.d.	5.6 ± 2.9
Glucuronic acid	n.d.	n.d.	2.6 ± 0.4	1.2 ± 0.3	2.2 ± 0.3

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526 527 528 Table 4 Monosaccharide and uronic acid composition in extracellular polymeric substances (EPMS) of microalgae, expressed as the average percentage of total monosaccharides and uronic acids (%) \pm standard deviation of triplicate measurements (n = 3). (n.d.: not detected).

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	P. cruentum	C. vulgaris	O. aurita	A. platensis
Glucose	35.4 ± 5.6	24.2 ± 1.1	2.1 ± 0.3	11.8 ± 0.4
Galactose	21.3 ± 1.6	17.3 ± 0.5	60.8 ± 1.7	11.1 ± 0.6
Xylose	29.3 ± 1.8	6.3 ± 0.4	10.6 ± 0.2	2.7 ± 0.3
Mannose	3.8 ± 2.5	19.1 ± 0.5	3.3 ± 0.2	1.7 ± 0.1
Rhamnose	n.d.	11.0 ± 0.3	2.8 ± 0.2	29.1 ± 1.4
Arabinose	n.d.	1.9 ± 0.1	n.d.	n.d.
Fucose	1.3 ± 0.1	8.7 ± 0.1	15.9 ± 0.5	8.9 ± 0.2
Ribose	n.d.	n.d.	n.d.	25.8 ± 0.8
Glucosamine	n.d.	n.d.	n.d.	n.d.
Galacturonic acid	n.d.	3.3 ± 0.1	1.3 ± 0.1	n.d.
Glucuronic acid	8.9 ± 0.3	8.2 ± 0.4	3.2 ± 0.1	8.9 ± 0.3

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530 531 532 **Table 5** Sulfate content in cell wall polysaccharides (CWPS) and extracellular polymeric substances (EPMS) of microalgae, expressed as the average percentage of CWPS or EPMS (%) \pm standard deviation of triplicate measurements (n = 3). (n.a.: not analyzed)

	CWPS	EPMS
P. cruentum	4.24 ± 0.12	7.05 ± 0.18
C. vulgaris	1.14 ± 0.06	1.73 ± 0.21
T. chuii	1.18 ± 0.10	n.a.
P. tricornutum	6.89 ± 0.17	n.a.
O. aurita	10.89 ± 0.61	13.93 ± 0.52
Nannochloropsis sp.	6.43 ± 0.11	n.a.
Schizochytrium sp.	2.69 ± 0.08	n.a.
T. lutea	4.69 ± 0.20	n.a.
D. lutheri	5.99 ± 0.14	n.a.
A. platensis	0.64 ± 0.11	1.34 ± 0.13

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534 Cell wall related polysaccharides of *P. cruentum* have been extensively studied in literature, with major focus on its extracellular polysaccharides with unique rheological properties and bioactivities [13]. Cells of 535 Porphyridium sp. are encapsulated within a cell wall polysaccharide complex. The external part easily 536 dissolves into the cultivation medium and polymers therein are usually identified as soluble polysaccharides, 537 corresponding to EPMS in this study. On the other hand, the part that remains bound to the cell, usually 538 referred to as bound polysaccharides [83], corresponds to CWPS. The proposed cell wall structure, i.e. one 539 polysaccharide complex surrounding *P. cruentum* cells, can be confirmed based on our results, since CWPS 540 and EPMS showed comparable monosaccharide profiles, with glucose, galactose and xylose being the 541 dominant sugars. This was previously reported by several authors, but with variable molar ratios of these 542 monosaccharides [34,83,84]. In addition, uronic acids accounted for 8-9% of all monosaccharides in 543 CWPS and EPMS, in accordance with previous studies [34,85]. However, whereas EPMS consisted only 544 of glucuronic acid, CWPS presented both galacturonic and glucuronic acid. Another difference between 545 546 CWPS and EPMS was the degree of sulfation, which was lower in CWPS (4.2%) than in EPMS (7.1%). These sulfate contents fall in the range reported in literature for *Porphyridium* sp. (4 - 14%) [12,85]. 547 In CWPS of C. vulgaris, glucose and mannose were the most abundant monosaccharides, suggesting the 548

presence of glucomannans. Other authors also reported that these monosaccharides account for most of the cell wall of *Chlorella* sp. [86,87], although it should be noted that two other types of CWPS can also be found in chlorococcal algae [88]. However, glucomannans typically contain larger amounts of mannose than glucose, indicating that another source of glucose was present in the *C. vulgaris* cell wall. For instance,

553 glucose could also result from (partial) hydrolysis of cellulosic and hemicellulosic polymers. Chen et al. 554 (2013) described the cell wall of Chlorophyta consisting of an inner cell wall layer, mainly composed of 555 cellulose and hemicellulose, and an outer cell wall layer with a variable composition depending on the 556 species [89]. However, discrepancies in cell wall structures of *Chlorella* sp. have been described in literature. This can be partly attributed to taxonomic revisions, resulting in a new definition of the Chlorella 557 558 genus and therefore a transfer of previously assigned species to other genera [7]. Nevertheless, contrasting findings of several studies still result in a lack of clarity on the *Chlorella* sp. cell wall composition. Whereas 559 some authors ascribe the rigidity of the cell wall to cellulose microfibrils [89,90], others attribute it to a 560 chitin-like polysaccharide [7,11,91]. Given the low amount of glucosamine observed in this study, the latter 561 statement cannot be confirmed based on our results. Moreover, even though the presence of hemicelluloses 562 is mentioned by most authors, there is no consensus on the type of hemicelluloses in Chlorella sp. cell walls. 563 564 In fact, the presence of either arabinomannans, arabinogalactans, glucomannans or xyloglucans was reported in various studies [17,87,92]. Furthermore, CWPS of C. vulgaris contained low amounts of sulfate 565 566 groups. Although some authors reported substantial amounts of sulfate esters in certain species of the Chlorella genus [12,93], no reports were found on C. vulgaris. Finally, EPMS from C. vulgaris showed a 567 very heterogeneous composition, containing seven monosaccharides and two uronic acids, with a low 568 degree of sulfation. Whereas the values obtained are confirmed for EPMS of *Chlorella* sp. by some authors 569 [14,68], others show contrasting results [12,94]. It can thus be concluded that microalgae of the Chlorella 570 genus show a very large diversity in cell wall composition, presenting different cell wall structures 571 depending on the species and the strain. 572

Similar to *C. vulgaris*, mannose and glucose were the main monosaccharides in the CWPS of *T. chuii*.
Several authors reported these monosaccharides in *Tetraselmis* sp., but together with high amounts of galactose [95–97]. In addition, appreciable amounts of galacturonic acid in combination with minor amounts
of galactose, rhamnose, arabinose and fucose suggest the presence of pectic polysaccharides. In fact, Arora *et al.* (2012) have reported the presence of glucans, galactomannans and pectins in cell walls of *T. indica*[98]. It should be noted that the obtained monosaccharide composition probably does not provide the full
carbohydrate profile of the *T. chuii* cell wall. In fact, *Tetraselmis* sp. cell walls are built up of thecae which

- contain up to 80% of acidic polysaccharides, characterized as 3-deoxy-manno-2-octulosonic acid, 3-deoxy5-O-methyl-manno-2-octulosonic acid and 3-deoxy-lyxo-2-heptulosaric acid [7,99]. However, no attempts
 were made to quantify these keto-sugar acids in this study. Finally, a sulfate content of 1.2% was observed
 in CWPS of *T. chuii*, somewhat lower compared to previous studies [68,99].
- CWPS of the diatom P. tricornutum consisted mainly of mannose and glucuronic acid, representing 60% of 584 585 the total sugars. Several authors have previously identified a sulfated glucuronomannan as the most prominent polysaccharide present in *P. tricornutum* [97,100,101]. A sulfate content of 6.9% was obtained, 586 corresponding to previous reports [93,100]. In addition, minor amounts of other monosaccharides, mainly 587 xylose, rhamnose, glucose and galactose, suggest the presence of other hemicellulosic polymers in the cell 588 wall of *P. tricornutum* [17]. Although some authors suggest that sulfated glucuronomannans are a conserved 589 structural polymer in diatoms [101], this was not obvious from the CWPS composition of O. aurita, another 590 591 diatom. A heterogeneous monosaccharide profile was found, mainly containing galactose, mannose, fucose and glucose. EPMS of O. aurita also presented high amounts of galactose and fucose, while mannose and 592 593 glucose were only minor constituents of EPMS. Uronic acids accounted for 10% of the sugar composition in both fractions. In addition, high sulfate contents were found for CWPS and EPMS of O. aurita, being 594 11% and 14%, respectively. As a consequence, cell wall related polysaccharides of O. aurita possess anionic 595 characteristics and could therefore be an interesting source in terms of several functionalities. 596
- 597 CWPS of Nannochloropsis sp. consist of 75% glucose, together with minor amounts of other monosaccharides. This large fraction of glucose was previously observed by several authors and was 598 ascribed to cellulose polymers [71,97,102]. Scholz et al. (2014) described the cell wall of N. gaditana as a 599 bilayer structure, consisting of a cellulosic inner wall representing approximately 75% of the mass balance, 600 protected by an outer algaenan layer [71]. The latter layer was not identified nor quantified in this study. It 601 must be noted that microcrystalline cellulose is only partially hydrolyzed by methanolysis and TFA 602 hydrolysis, with a recovery of only 30 to 50%. This might be another explanation for the low amount of 603 604 CWPS of Nannochloropsis sp. presented in Table 1. No uronic acids were present in Nannochloropsis sp., 605 while a substantial amount of sulfate groups was observed. The latter results, in combination with low

amounts of other monosaccharides, suggest the presence of sulfated polysaccharides in the
 Nannochloropsis sp. cell wall in addition to cellulose and algaenan.

Only few studies investigated the cell wall composition of *Schizochytrium* sp., describing a thin noncellulosic cell wall with galactose as the principal monosaccharide [103,104]. In the current study, galactose
was also observed, however together with glucose, mannose and xylose. The latter two monosaccharides
have been previously reported in *Thraustochytrium* species. These fungi are phylogenetically related to *Schizochytrium* sp., based on the structure and formation of the cell wall in particular [103,104].
Furthermore, CWPS of *Schizochytirium* sp. presented a low degree of sulfation (2.7%) and uronic acids
were absent.

In CWPS of T. lutea, all monosaccharides were detected in substantial amounts, except for glucosamine. 615 The largest fraction of CWPS was composed of glucose, mannose, arabinose and galactose. These were 616 also the major monosaccharides observed in previous studies [95,97], although glucose represented 76% of 617 the total sugar composition in the study of Brown (1991) [97]. However, that study reported on the overall 618 polysaccharide composition, including the SPS chrysolaminarin. Since our results showed that T. lutea 619 contained a large fraction of chrysolaminarin in comparison with the fraction of CWPS (Table 1), it is 620 inferred that glucose is mainly attributed to chrysolaminarin, and only partially constituting the cell wall of 621 T. lutea. Finally, the CWPS also contained sulfated polysaccharides, with similar sulfate contents as 622 previously reported (1.1 - 5.5%) [20]. Although D. lutheri also belongs to the Haptophyta, i.e. the same 623 class as T. lutea, a different CWPS composition was observed for this microalga species. Glucose was found 624 as the principal monosaccharide, while mannose, xylose and arabinose were present in low amounts. Arnold 625 et al. (2015) reported that D. lutheri cell walls consist of small cellulose scales covered by an organic matrix 626 of hemicellulose, mainly xyloglucans [105]. Moreover, a sulfate content of 6% was determined for CWPS 627 of D. lutheri, indicating the presence of sulfated polysaccharides. 628

Finally, CWPS of the cyanobacterium *A. platensis* were mainly composed of glucose and mannose. In fact,
these monosaccharides are typically found in the cell walls of cyanobacteria, together with minor amounts
of galactose and xylose [106]. In addition, the observed amounts of rhamnose could result from spirulan
polymers, a sulfated polysaccharide which was previously extracted from *A. platensis* [107]. It should be

633 noted that polysaccharides probably represent a small fraction of the cell wall, since cyanobacterial cells are 634 typically covered by a peptidoglycan layer, a polymer composed of N-acetyl-glucosamine [106]. Surprisingly, only low amounts of glucosamine were detected in this study, which could be due to 635 636 incomplete hydrolysis of peptidoglycan by methanolysis and TFA hydrolysis. In fact, some authors hypothesized that a harsher HCl hydrolysis procedure would result in a more complete hydrolysis of N-637 638 acetyl-glucosamine containing polymers [17]. Furthermore, a heterogeneous monosaccharide profile was observed for EPMS of A. platensis, showing substantial amounts of rhamnose, ribose, glucose, galactose, 639 fucose and glucuronic acid. These sugars have been previously reported for EPMS of A. platensis, but in 640 different ratios [14,64]. Although several bioactivities are ascribed to specific sulfated polysaccharides of 641 A. platensis [14,107], it is assumed that these polymers only make up a small fraction of the cell wall related 642 polysaccharides, since low degrees of sulfation were observed in both CWPS and EPMS (0.6% and 1.4%). 643

644

645 **4.** Conclusions

This study compared the biomass profiles of ten microalgae species using a standardized protocol, mainly 646 focusing on the composition of their cell wall related polysaccharides. A large diversity in biomass 647 composition was demonstrated, allowing an appropriate selection of microalgae species towards specific 648 applications. Nevertheless, some taxonomic similarities were observed. Microalgae belonging to the 649 Haptophyta (T. lutea and D. lutheri) were characterized by high amounts of lipids and proteins, 650 651 intermediate SPS and ash contents, and the absence of EPMS. In contrast, C. vulgaris and T. chuii (Chlorophyta) were mainly composed of proteins and CWPS, while lipids represented a smaller fraction 652 of these biomasses. Less similarities were found among the two investigated diatoms, P. tricornutum and 653 O. aurita, which might be ascribed to distinct cultivation conditions resulting in a high SPS content in 654 655 O. aurita. Moreover, the presence of large amounts of structural components in most microalgae, such as EPS in *P. cruentum* or high protein contents in *C. vulgaris* and *A. platensis*, might explain the functionality 656 of microalgal biomasses as potential gelling or thickening agents. 657

658 Generally, CWPS made up 10% of the microalgal biomass, however the contribution of other non-659 polysaccharide substances to the cell wall could not be excluded. Extraction of EPMS was only successful

for four microalgae species, implying that the other microalgae do not possess extracellular polymers. Even 660 though CWPS and EPMS of all microalgae species were sulfated, their degree of sulfation was generally 661 low. In addition, some microalgae presented polysaccharides with substantial amounts of uronic acids and 662 sulfate groups, providing anionic characteristics to these polymers. The use of additional structural 663 analyses, including linkage analysis or the use of monoclonal antibodies for recognition of specific cell 664 665 wall polysaccharide epitopes, might provide additional insights into the molecular structure of the microalgal cell wall related polysaccharides. Nevertheless, based on the results obtained in this study, it 666 can be concluded that some microalgal cell wall related polysaccharides might have great potential and 667 should be further explored for use as bioactive or technological substances. 668

669

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