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Size-fractionated dissolved primary production and
carbohydrate composition of the coccolithophore *Emiliana huxleyi*

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20 **Abstract**

21 Extracellular release (ER) by phytoplankton is the major source of fresh dissolved organic
22 carbon (DOC) in marine ecosystems and accompanies primary production during all growth
23 phases. Little is known, so far, on size and composition of released molecules, and to which
24 extent ER occurs passively, by leakage, or actively, by exudation. Here, we report on ER by
25 the widespread and bloom-forming coccolithophore *Emiliana huxleyi* grown under steady
26 state conditions in phosphorus controlled chemostats (N:P=29, growth rate of $\mu=0.2\text{ d}^{-1}$) at
27 present day and high CO₂ concentrations. ¹⁴C incubations were accomplished to determine
28 primary production (PP), comprised by particulate (PO¹⁴C) and dissolved organic carbon
29 (DO¹⁴C). Concentration and composition of particulate combined carbohydrates (pCCHO),
30 and of high molecular weight (>1 kDa, HMW) dissolved combined carbohydrates (dCCHO)
31 were determined by ion chromatography. Information on size distribution of ER products
32 was obtained by investigating distinct size classes (<0.40 μm (DO¹⁴C), <0.45 μm (HMW-
33 dCCHO), <1000 kDa, <100 kDa and <10 kDa) of DO¹⁴C and HMW-dCCHO. Our results
34 revealed relatively low ER during steady state growth, corresponding to ~4.5% of primary
35 production, and similar ER rates for all size classes. Acidic sugars had a significant share on
36 freshly produced pCCHO as well as on HMW-dCCHO. While pCCHO and the smallest size
37 fraction (<10 kDa) of HMW-dCCHO exhibited a similar sugar composition, dominated by
38 high percentages of glucose (74-80 Mol %), the composition of HMW-dCCHO size-classes
39 >10 kDa was significantly different with higher Mol % of arabinose. Mol % of acidic sugars
40 increased and Mol % glucose decreased with increasing size of HMW-dCCHO. We conclude
41 that larger polysaccharides follow different production and release pathways than smaller
42 molecules, potentially serving distinct ecological and biogeochemical functions.

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48 **Keywords:** Exudation, carbohydrates, DOC, primary production, coccolithophores

49 **1. Introduction**

50 The global ocean inventory of dissolved organic carbon (DOC) is estimated to be in a range
51 of 662 – 700 Gt (Hansell and Carlson 1998, Ogawa and Tanoue 2003). A common
52 classification of marine DOC relies on its reactivity and discriminates between labile
53 (LDOC), semi-labile (SLDOC), semi-refractory (SRDOC), refractory (RDOC) and ultra-
54 refractory (URDOC) DOC with lifetimes of hours to days, weeks to months, month to years,
55 centuries or even millennia (Kirchman 1993, Carlson & Ducklow 1995, Anderson &
56 Williams 1999, Hansell, 2013). Only a small fraction of marine DOC is considered reactive;
57 LDOC (< 0.2 Gt) and SLDOC (6 ± 2 Gt) (Hansell, 2013). In general, these compounds are
58 freshly produced by plankton and represent the major nutritional resource for heterotrophic
59 microorganisms (Cherrier et al., 1996; Amon & Benner 1996, Amon et al., 2001, Benner,
60 2002, Azam & Malfatti 2007, Davis et al., 2009). Especially during the summer season,
61 SLDOC can accumulate in temperate waters, and becomes available for deep convective
62 mixing, contributing to the biological carbon pump (Hopkinson and Vallino, 2005; Hansell et
63 al., 2009). Microbial assimilation of DOC as well as the formation of gel particles, such as
64 transparent exopolymer particles (TEP), lead to a repartitioning of DOC into the particulate
65 organic carbon (POC) pool (Alldredge et al., 1993; Chin et al., 1998; Engel et al., 2004), the
66 sinking of which represents another pathway for carbon export and storage in the ocean. In
67 addition, microbial processing of fresh DOC may result in formation of recalcitrant
68 compounds with longer residence time, also increasing the carbon dioxide (CO₂) storage
69 potential in the ocean (Jiao and Zheng 2011). Thus, deeper insights to the origin and quality
70 of DOC in the ocean can greatly abet our ability to quantify carbon and nutrient cycling in
71 the ocean.

72

73 The ultimate source of organic carbon in the ocean is primary production, and extracellular
74 release (ER, also referred to as dissolved primary production) of organic carbon is the
75 primary source of fresh DOC, followed by cell lysis (Fuhrman 1999), grazing (Møller 2005),
76 enzymatic particle solubilisation (Cho and Azam, 1988; Smith et al., 1992) and sloppy
77 feeding (Copping and Lorenzen, 1980; Nagata, 2000). The major components of
78 phytoplankton ER are high molecular weight (HMW, >1 kDa) dissolved combined
79 carbohydrates (dCCHO), representing also the largest characterizable fraction of marine
80 dissolved organic matter (DOM); 15-35 % DOC in the surface ocean, 5-10 % DOC in the
81 deep ocean (Benner et al., 1992 Pakulski & Benner 1994, Biddanda and Benner, 1997,
82 Ogawa and Tanoue, 2003). Composition of HMW-dCCHO in seawater is usually determined
83 on the basis of monomeric sugars after hydrolysis of the polymer chains, and resembles

84 either phytoplankton biomass itself (Pakulski & Benner 1994, Børsheim et al. 1999) or
85 extracellular CCHO from phytoplankton cultures (Biersmith & Benner 1998, Aluwihare &
86 Repeta 1999, Aluwihare et al. 2002). The latter are usually comprised by neutral hexoses,
87 pentoses and deoxysugars like glucose, galactose and mannose, by amino sugars like
88 glucosamine and galactosamine and by uronic acids e.g. galacturonic acid and glucuronic
89 acid (Aluwihare et al., 1997; Biersmith & Benner, 1998; Aluwihare & Repeta, 1999; Engel et
90 al. 2010, Borchard & Engel, 2012).

91

92 ER is a normal function of healthy algae cells during all stages of growth (Fogg 1966, Mague
93 et al. 1980, Bjørnsen 1988, Borchard and Engel 2012, Lopez-Sandoval et al. 2011) and can
94 comprise up to 80 % of primary production (Sharp, 1977, Mague, 1980, Fogg, 1983,
95 Bjørnsen, 1988). Two conceptual models have been proposed for phytoplankton ER: *i*) the
96 *passive diffusion model* that describes the leakage of smaller molecules from inside the cell to
97 its surrounding environment (Fogg, 1983, Bjørnsen, 1988), and *ii*) the *overflow model* that
98 assumes an energy consuming exudation of HMW compounds (Fogg 1983, Nagata 2000,
99 Schartau et al. 2007). According to the *passive diffusion model*, DOC crosses the cell
100 membrane independently from primary production (PP) during day and night, and ER
101 correlates to phytoplankton biomass and cell size. A higher relative contribution of ER to
102 total PP would therefore be expected in communities dominated by small cells due to their
103 higher surface to volume ratio (Bjørnsen 1988, Kiørboe & Hansen 1993, Marañón et al.
104 1996).

105 Central aspects of the *overflow model* are a dependence of ER on PP rates, the absence of ER
106 at night and a high share of HMW substances (Williams 1990 and references therein, Nagata
107 2000). Fogg (1966) proposed that photosynthesis and build-up of organic carbon is primarily
108 regulated by irradiance, while cell growth is controlled by the availability of inorganic
109 nutrients. The discharge of photosynthates, not utilized for cell growth, was suggested to be
110 more energy-efficient than intracellular storage (Wangersky 1978, Wood & van Valen 1990).
111 In accordance with the *overflow model*, data from coastal, marine and estuarine systems
112 revealed a linear relationship between PP and ER, and factors influencing PP were suggested
113 to also affect ER (Baines and Pace 1991). Such effects were shown for light (Zlotnik &
114 Dubinsky 1989) and later suggested also for CO₂ (Engel 2002) and temperature (Moran et al.
115 2006). Under nutrient limitation, however, substantial ER was observed when PP was
116 reduced, leading to higher percentages of extracellular release (PER) (Myklestad et al. 1989,
117 Goldman et al. 1992, Obernosterer & Herndl 1995, Halewood et al. 2012). Under such
118 conditions, decoupled from PP and biomass, ER becomes difficult to estimate, both in terms

119 of quantity and quality. Moreover, phytoplankton cells display a large physiological plasticity
120 for nutrient requirements, i.e. the nutrient cell quota, which varies with environmental
121 conditions or among different taxonomic groups (Geider and LaRoche, 2002).

122 Despite their role in marine carbon cycling, processes involved in the production,
123 consumption and remineralisation of extracellular organic matter are little understood and
124 have largely been neglected in biogeochemical models (Flynn et al., 2008, Repeta &
125 Aluwihare, 2006; Hansell et al., 2009, Hansell, 2013). So far, it is not known if extracellular
126 products are mainly released by leakage or by exudation processes, or how much leakage and
127 exudation products differ. We also don't know if and how the physiological status of the cell
128 influences the composition of extracellular products, and whether or not such differences in
129 chemical signatures subsequently affect their microbial cycling, remineralisation rate, or
130 affinity to form gel particles.

131

132 In order to improve our understanding on ER, we conducted a chemostat experiment with *E.*
133 *huxleyi* under fully controlled nutrient supply and growth rate. *Emiliania huxleyi* is a bloom
134 forming cosmopolitan coccolithophore species, and known to produce a methylated, acidic
135 polysaccharide that plays a central role in coccolith formation and agglutination (Fichtinger-
136 Schepmann, 1979, De Jong, 1979). ER by *E. huxleyi* cells was reported earlier (Aluwihare
137 and Repeta 1999, Biddanda and Benner 1997, Borchard and Engel 2012) and carbohydrates
138 were shown to provide a substantial fraction of freshly produced HMW-DOC (35-94 %)
139 (Aluwihare and Repeta 1999, Biddanda and Benner 1997).

140

141 This study was part of a larger experiment investigating carbon and nutrient cycling under
142 different $p\text{CO}_2$ conditions at steady state growth in *E. huxleyi*. No effect of the CO_2 treatment
143 was observed for elemental stoichiometry of cells as well as for TEP production (Engel et al.
144 2014). This study focusses on primary production of POC and DOC by *E. huxleyi*, the
145 carbohydrate composition of cells and for the first time on different size fractions of released
146 compounds.

147

148 With our study we wanted (i) to determine ER of DOC and carbohydrates by combining rate
149 measurements for particulate and dissolved primary production with analyses of carbohydrate
150 concentration, and (ii) to characterize monomeric carbohydrate composition in different size
151 classes of DOC in order to elucidate mechanisms of ER. We chose the continuous culture
152 approach, because here cells can be grown under nutrient limitation at steady state biomass.
153 Thus, in a chemostat the increase in extracellular organic matter can primarily be attributed to

154 growing phytoplankton cells and not to cell lysis and decay, processes that co-occur with ER
155 when batch cultures or natural populations become nutrient depleted.
156

157 **2. Methods**

158 **2.1 Experimental setup**

159 A calcifying strain of *E. huxleyi* (PML B92/11) was grown as continuous culture in two
160 chemostats (~9.2 L each) at a constant dilution rate of $D = 0.2 \text{ d}^{-1}$. A more detailed
161 description of the chemostat principle and the experimental set-up are given by Borchard et
162 al. (2011), Borchard and Engel (2012), and Engel et al. (2014), respectively. Temperature
163 was set to $14.0 \pm 0.1^\circ\text{C}$. Irradiance was provided at a 16h:8h light:dark cycle with a photon
164 flux density of $190 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$ (TL-D Delux Pro, Philips; QSL 100, Biospherical
165 Instruments, Inc.). Nutrient medium was prepared from sterile-filtered (Sartobran P, 0.2 μm
166 capsule, Sartorius) aged natural seawater (NSW) with a salinity of 33, total alkalinity (TA) of
167 $2250 \mu\text{mol kg}^{-1}$ seawater and a pH of 8.24. The seawater was enriched with nutrients
168 according to the f/2 recipe of Guillard and Ryther (Guillard & Ryther 1962) with final
169 concentrations of $43 \mu\text{mol L}^{-1} \text{NO}_3^-$ and $1.5 \mu\text{mol L}^{-1} \text{PO}_4^{3-}$. The nutrient medium was treated
170 for 3 h with UV irradiation (Microfloat 1/0, a.c.k. aqua concept GmbH) for sterilisation
171 before the addition of sterile-filtered (0.2 μm , Minisart, Sartorius) f/2-vitamins. Axenic
172 conditions, however, could not be maintained in the 9.2 L chemostats over the long period of
173 time.

174 Equilibration of the medium with CO_2 was obtained by constant aeration with 380 and 750
175 $\mu\text{atm CO}_2$, respectively. To minimize effects of calcification by *E. huxleyi* on carbonate
176 chemistry in the incubators, TA in the reservoir tank was increased by addition of
177 bicarbonate (LaRoche et al. 2010) resulting in $2460 \mu\text{mol kg}^{-1}$ seawater. *E. huxleyi* cells
178 were pre-cultured for 30 d at prescribed CO_2 concentrations and temperature conditions in f/2
179 media in order to avoid short term stress effects on cell physiology. Each chemostat incubator
180 was then inoculated to a final density of $\sim 5000 \text{ cells ml}^{-1}$. Cultures were grown in batch mode
181 for 5 d until the constant medium supply was applied at a dilution rate (D) of $D=0.2 \text{ d}^{-1}$. Cells
182 were kept in suspension by gentle mixing at $50 \text{ rotations min}^{-1}$. Here, we report data derived
183 from samplings during steady state growth on experimental day 30, 34, 38, 42 and 44 for ^{14}C
184 rate measurements and on day 38, 42 and 44 for carbohydrate analyses and size
185 fractionations of those and ^{14}C exudation. All samples were taken 3 hours after lights on to
186 avoid biases due to physiological variations during the day-night cycle.

187

188 **2.2 Cell density and chemical analysis**

189 **2.2.1 Cell density** was determined daily as the mean of three consecutive measurements of
190 $500 \mu\text{l}$ by an electronic particle counter (Coulter Multisizer III, Beckman Coulter) equipped
191 with a $100 \mu\text{m}$ aperture. $0.2 \mu\text{m}$ pre-filtered (Minisart 2000, Sartorius) NSW with a salinity

192 of 33 was used to dilute the samples 1:100. After microscopic inspection, particles with an
193 equivalent spherical diameter in a range of 3.2 μm to 8.0 μm were identified as *E. huxleyi*
194 cells.

195 **2.2.2 Nutrient samples** were filtered through 0.2 μm syringe filters (Minisart, Sartorius) and
196 stored frozen at -20°C until analysis. Measurements of NO_3^- , NO_2^- , NH_4^+ and PO_4^{3-} were
197 made spectrophotometrically after Grasshof et al. (1999) using an Evolution 3 autoanalyzer
198 (Alliance Instruments). Detection limits were 0.3 $\mu\text{mol L}^{-1}$ for N and 0.01 $\mu\text{mol L}^{-1}$ for P.

199 **2.2.3 Primary production and exudation** were measured by applying the ^{14}C incubation
200 method according to Steemann Nielsen (Steemann Nielsen 1952) and Gargas (Gargas 1975).
201 Triplicate samples (75 ml each) were taken from each chemostat, transferred into cell culture
202 flasks (25 cm^2 , Corning[®]) and spiked with approximately 5 $\mu\text{Ci NaHCO}_3^-$ (Hartmann
203 Analytics, specific activity 40-60 mCi/mmole). Each triplicate set was incubated for about
204 4 h at original experimental light and temperature settings, but without aeration.
205 Simultaneously, dark uptake was measured in triplicate from 75 ml samples incubated in the
206 dark. Added activity in the samples was determined by removing a 100 μl aliquot from three
207 dark bottles prior to incubation and transferred to 6 ml liquid scintillation vials in which
208 200 μl of 2N NaOH were placed. 4 ml liquid scintillation cocktail (Ultima Gold AB) were
209 added before counting. Incubations were stopped by gentle filtration on 0.40 μm
210 polycarbonate filters (Nucleopore) at low vacuum (<150 mbar) to avoid cell breakage. The
211 filters (PO^{14}C) were covered with 250 μl 1 M HCl in order to remove inorganic ^{14}C . After a
212 few seconds they were rinsed with 10 ml filtered seawater. Filters were transferred to 6 ml
213 scintillation vials, 4 ml liquid scintillation cocktail (Ultima Gold AB) were added and
214 samples were stored overnight before being counted in a Packard Tri Carb Liquid
215 Scintillation Counter. Carbon incorporation rates were calculated in accordance to Borchard
216 and Engel (2012).

217 For the determination of released dissolved organic carbon (DO^{14}C), 4 ml of the filtrate were
218 transferred into 20 ml scintillation vials and acidified to $\text{pH} < 2$ by the addition of 100 μl 1 M
219 HCl and left open under the fume hood for 24 hours. For size fractionation of DO^{14}C ,
220 triplicate sets of 10 ml sample were transferred into Macrosep[®] centrifugal devices with
221 membrane cut off of <1000 kDa, <100 kDa and <10 kDa, respectively. After centrifugation
222 (Heraeus, Megafuge[®] 1.0 R) for 15 min at 4000 rounds per minute, 4 ml sample were
223 transferred into 20 ml liquid scintillation vials. In the following, samples were treated as the
224 whole DO^{14}C samples and after the outgassing of inorganic ^{14}C , 15 ml liquid scintillation
225 cocktail were added. Counting and calculations were accomplished after Borchard and Engel
226 (2012).

227 Primary Production (PP) was derived from the sum of $PO^{14}C$ and $DO^{14}C$. The percentage of
228 extracellular release (PER) was calculated as $(DO^{14}C/PP)*100$.

229

230 **2.2.4. Total combined carbohydrates (tCCHO)** and high molecular weight (HMW; >1 kDa)
231 **dissolved combined carbohydrates (dCCHO)** were determined by ion chromatography after
232 Engel and Händel (2011). Duplicate samples for HMW-dCCHO were filtered through 0.45
233 μm syringe-filters (GHP membrane, Acrodisk, Pall Corporation) and stored in combusted (8
234 h at 500°C) glass vials at -20°C. Samples for tCCHO remained unfiltered and were stored
235 identically.

236 For size fractionation of HMW-dCCHO, 10 ml sample were transferred into Macrosep®
237 centrifugal devices with a molecular weight cut-off (MWCO) of 1000 kDa, 100 kDa and 10
238 kDa, respectively. After centrifugation (Heraeus, Megafuge® 1.0 R) for 15 min at 4000
239 rounds per minute, samples were transferred into combusted (8 h at 500°C) glass vials and
240 stored at -20°C. Before usage, Macrosep® devices were rinsed twice by centrifugation with
241 ultrapure water to avoid any contamination with carbohydrate compounds in the membrane.
242 Concentrations of CCHO in these blanks were tested to be below the detection limit and did
243 therefore not affect analyses.

244 Prior to analysis, samples were desalinated by membrane dialysis (1 kDa MWCO, Spectra
245 Por) for 6 h at 0°C and thereafter hydrolyzed with HCl at a final concentration of 0.8 M for
246 20 h at 100°C to yield monomeric CHO. Samples were stored at -20°C over night and then
247 neutralized by acid evaporation (N_2) at 50°C. Dried samples were solubilised in ultra pure
248 water before determination of CHO monomers by high performance anion exchange
249 chromatography (HPAEC) coupled with pulsed amperometric detection (PAD) on a Dionex
250 ICS 3000 (Engel & Händel 2011). A Dionex CarboPac PA10 guard column (2x50 mm)
251 coupled to a Dionex CarboPac PA10 analytical column (2x250 mm) was applied for
252 separation of fucose (Fuc), rhamnose (Rha), arabinose (Ara), galactosamine (GalN),
253 glucoseamine (GlcN), galactose (Gal), glucose (Glc), mannose/xylose (Man/Xyl) (quantified
254 together due to co-elution), galacturonic acid (Gal-URA) and glucuronic acid (Glc-URA).
255 Detection limits are 10 nM. **Particulate CCHO (pCCHO)** were derived from subtraction of
256 HMW-dCCHO from tCCHO and thus represent carbohydrates in the size fraction > 0.45 μm .
257 Concentrations of CHO are given as μmol carbon per volume of seawater ($\mu mol C L^{-1}$) and
258 composition of CCHO is expressed as Mol % CCHO.

259 **Size fractions** of $DO^{14}C$ and HMW-dCCHO obtained by Macrosep® centrifugal devices
260 were subtracted from each other in order to present data for each size class. Definitions for
261 size classes are given in table 1.

262

263 **2.2.5** For **total alkalinity** (TA), 25 ml of each sample were measured by titrating with 0.05 M
264 HCl until the buffering capacity of the water samples was consumed and all bases of interest
265 were protonated to zero level species. Analysis was accomplished with an automatic titrator
266 (TitroLine® *alpha plus*, SI Analytics) equipped with a sample changer (TW *alpha plus*, SI
267 Analytics) and a piston burette (Titronic®110 *plus*, SI Analytics). The pH was monitored by
268 a two-point calibrated (buffer solution pH 4.006 and pH 6.865; Applichem, standardised
269 according to DIN 19266) electrode (Schott® Instruments IoLine). The concentration of TA in
270 $\mu\text{mol kg}^{-1}$ seawater was calculated from linear regression of the absolute numbers of protons
271 in solution and the total volume (sample plus HCl) in the range of pH 4 and 3. Determination
272 of the seawater carbonate chemistry was conducted by using the program co2sys (Lewis &
273 Wallace 1998) with pH (calibrated by the use of reference materials provided by A. Dickson)
274 and TA being the input parameters.

275

276 **2.3 Data treatment**

277 All samplings were accomplished during the steady state period of the experiment when the
278 growth rate (μ) was equal to the dilution rate (D). The samplings over time thus represent
279 replicates of the same physiological state and values of the respective parameters are given as
280 average \pm standard deviation. Since CO_2 induced no differences between the *present day* and
281 the *high CO₂* chemostat, they were used as replicate treatments and values are given as mean
282 values with single standard deviation if not stated otherwise.

283 In order to relate daily rates ($\mu\text{mol L}^{-1} \text{d}^{-1}$) directly to concentrations ($\mu\text{mol L}^{-1}$), data were
284 converted into each other by applying a growth rate of 0.2 d^{-1} . For cell normalized carbon
285 values concentrations and rates were divided by the cell number.

286 Differences in carbohydrate composition for the different size fractions were tested by means
287 of analysis of co-variance (two-way ANOVA). Differences as response to CO_2 conditions
288 were tested by means of a *t*-test. Statistical significance was accepted for $p < 0.05$. All
289 calculations were performed using the software package Sigma Plot 10.01 (SysStat).

290

291 **3. Results**

292 **3.1 Growth, nutrients and carbonate chemistry**

293 Growth and biogeochemical composition of *Emiliana huxleyi* as well as carbonate and
294 nutrient chemistry during this chemostat experiment are described in more detail in Engel et
295 al. (2014). Briefly, on day 28 of the experiment, the steady state was reached with the
296 dilution rate (D) being equal to the growth rate (μ) of *E. huxleyi*. Cell abundances and basic
297 parameters such as particulate organic carbon (POC), nitrogen (PON), phosphorus (POP) and
298 chlorophyll *a* (chl *a*) remained constant until the end of the experiment proving the constant
299 physiological state of *E. huxleyi* (Engel et al., 2014).

300 During the steady state period, cell densities were similar in the *present day* and *high CO₂*
301 treatment and averaged $5.2 \cdot 10^8 \pm 18.6$ % cells L⁻¹ and $5.1 \cdot 10^8 \pm 19.7$ % cells L⁻¹,
302 respectively. High variations resulted exclusively from intensive sampling between days 42
303 and 44. Until day 42 variations did not exceed 11.6 % and biomass production was accepted
304 as balanced growth as a result of controlled nutrient supply. During steady state (days 30-44),
305 both, NO₃⁻ and PO₄³⁻ concentrations were below the detection limit in both treatments. P-
306 limitation was likely more severe than N-limitation, given a nutrient supply N:P ratio of ~29
307 and indicated also by PON:POP ratios clearly >16 (Engel et al., 2014). $p\text{CO}_2$ was calculated
308 from pH and TA and yielded significantly different values between treatments of 337 ± 94
309 (*present day*) and 623 ± 139 (*high CO₂*) μatm . Time averaged values given here differ
310 slightly from those given by Engel et al. (2014) as the latter used data from replicate
311 chemostats per CO₂ treatment, while only one chemostat per treatment was sampled for the
312 purpose of this study.

313

314 **3.2 Primary production and exudation**

315 As determined for cell densities, PO¹⁴C and DO¹⁴C production rates derived from replicate
316 sampling during steady state growth varied <11 % confirming the physiological steady state
317 of *E. huxleyi* grown in the chemostats. PO¹⁴C production of 173 ± 17 and 168 ± 16 $\mu\text{mol C L}^{-1}$
318 d^{-1} and DO¹⁴C production of 8.0 ± 0.7 and 8.2 ± 1.1 $\mu\text{mol C L}^{-1} \text{d}^{-1}$ were determined for the
319 *present day* and *high CO₂* μatm treatment, respectively (Fig. 1). Production rates of PO¹⁴C
320 and DO¹⁴C were not significantly different between the CO₂ treatments (Mann-Whitney
321 Rank sum tests and t-tests, $n=5$, $p>0.69$) and were thus averaged for both treatments: $171 \pm$
322 16 $\mu\text{mol C L}^{-1} \text{d}^{-1}$ (PO¹⁴C) and 8.1 ± 0.9 $\mu\text{mol C L}^{-1} \text{d}^{-1}$ (DO¹⁴C).

323 Cell normalized production of PO¹⁴C and DO¹⁴C during the steady state period were on
324 average 0.33 ± 0.04 and 0.015 ± 0.002 $\text{pmol C cell}^{-1} \text{d}^{-1}$ for both treatments. Similar PO¹⁴C
325 and DO¹⁴C production rates in both chemostats are reflected in comparable percentages of

326 extracellular release (PER) of 4.42 ± 0.22 % (*present day*) and 4.70 ± 0.92 % (*high CO₂*) and
327 also for the size classes of DO¹⁴C no CO₂ effect was determined (Fig. 1).

328 Averaged for both treatments, size fractionated (see table 1 for definition) DO¹⁴C production
329 ranged between 1.27 ± 0.53 (*medium*) and 2.74 ± 0.88 (*very large*) $\mu\text{mol C L}^{-1} \text{d}^{-1}$. Relative
330 contribution of different DO¹⁴C size classes to total DO¹⁴C was 33.6 ± 9.31 %, (*very large*),
331 24.6 ± 7.90 % (*large*), 15.9 ± 7.15 % (*medium*) and 25.8 ± 3.55 % (*small*). Thus, total DO¹⁴C
332 was comprised by comparable shares of DO¹⁴C in these size classes with slightly higher
333 proportions in the *very large* fraction.

334

335 **3.3 Combined carbohydrates**

336 Initial HMW-dCCHO concentrations of 7.02 ± 0.15 $\mu\text{mol C L}^{-1}$ were determined in the
337 natural seawater (NSW) media. Corrected for NSW values, carbohydrate concentration
338 during steady state growth of *Emiliania huxleyi* was 103 ± 28 (*present day*) and 104 ± 31
339 $\mu\text{mol C L}^{-1}$ (*high CO₂*) for pCCHO, and 15.2 ± 2.1 (*present day*) and 15.8 ± 2.4 (*high CO₂*)
340 $\mu\text{mol C L}^{-1}$ for fresh HMW-dCCHO, and hence very similar between the two CO₂ treatments
341 (Fig. 2). Averaged for both treatments, 87 ± 3 % of tCCHO were present in the particulate
342 fraction (pCCHO). *E. huxleyi* produced pCCHO in order of 104 ± 27 $\mu\text{mol C L}^{-1}$ (0.20 ± 0.02
343 pmol C cell^{-1}) equivalent to 20.7 ± 5.3 $\mu\text{mol C L}^{-1} \text{d}^{-1}$ at a growth rate of 0.2d^{-1} , representing
344 about 12.5 % of the daily produced PO¹⁴C (Table 2). Freshly produced HMW-dCCHO was
345 15 $\mu\text{mol C L}^{-1}$ (0.043 ± 0.004 pmol C cell^{-1}), equivalent to about 40 % of freshly produced
346 DO¹⁴C. Fresh carbohydrate concentrations in various size classes (see table 1 for definition)
347 also revealed a strong similarity between the *present day* and the *high CO₂* treatment (Fig. 2,
348 t-tests, $n=6$, $p>0.269$) and are therefore given as average values in the following. In the
349 different size classes, HMW-dCCHO comprised between 29.5 ± 9.3 % (*small*) and $59.7 \pm$
350 17.2 % (*Very large*) of DO¹⁴C (Table 2).

351 HMW-dCCHO yielded 7.07 ± 1.06 (*very large*), 3.57 ± 1.21 (*large*), 3.08 ± 1.26 (*medium*)
352 and 3.09 ± 0.92 $\mu\text{mol C L}^{-1}$ (*small*), suggesting that freshly released HMW-dCCHO were
353 primarily comprised by *very large* HMW-dCCHO ($46 \pm 3\%$ C). HMW-dCCHO in *large*,
354 *medium*, and *small* contributed 23 ± 6 , 20 ± 4 and 20 ± 3 % C, respectively, to total HMW-
355 dCCHO.

356 Size fractionation of HMW-dCCHO in NSW yielded concentrations of 2.39 ± 0.21 (*very*
357 *large*), 1.31 ± 0.09 (*large*), 1.28 ± 0.09 (*medium*) and 1.95 ± 0.10 (*small*) $\mu\text{mol C L}^{-1}$ (Fig. 3a,
358 left panel). During the experiment, extracellular release by *E. huxleyi* enriched the NSW to
359 HMW-dCCHO concentrations of 22.2 ± 2.1 $\mu\text{mol C L}^{-1}$ with size fractions of 8.55 ± 1.08

360 (very large), 4.64 ± 1.33 (large), 4.09 ± 1.21 (medium) and 5.18 ± 0.72 $\mu\text{mol C L}^{-1}$ (small)
361 (Fig. 3a, right panel).

362 **3.4 Carbohydrate composition of exudates**

363 Sugar monomers of three different types comprised the combined carbohydrates (CCHO)
364 determined during the present experiment: Neutral sugars (Fuc, Rha, Ara, Gal, Glc and
365 coeluting Man/Xyl), amino sugars (GalN and GlcN) and uronic acids (Gal-URA and Glc-
366 URA). Various amounts of these monomers were detected in HMW-dCCHO of the initial
367 NSW used for the present experiment (Fig. 3b, left panel). Size fractions of HMW-dCCHO
368 in NSW did not show any significant variation in monomeric composition ($p > 0.462$).
369 However, relative to the other size fractions Man/Xyl was slightly enriched in *small*, while a
370 smaller proportion of Fuc was detected in this fraction. No significant differences in
371 monomeric composition of CCHO produced by *Emiliana huxleyi* were determined between
372 the *present day* and *high CO₂* treatment ($p > 0.881$). Therefore, average values are given for
373 replicate sampling during steady state growth and both treatments in the following (Fig. 3
374 and Table 3).

375 ER by *E. huxleyi* led to a clear change in HMW-dCCHO composition of the NSW (Fig. 3b);
376 primarily caused by Ara being significantly higher ($p < 0.001$) in all size classes, except for
377 *small* (Fig. 3b). In HMW-dCCHO derived from *E. huxleyi*, Ara and Glc were the dominant
378 sugars with 46 and 21 Mol %, respectively followed by Gal-URA (11 Mol %), Man/Xyl (10
379 Mol %), Glc-URA (5.7 Mol %), Gal (4.6 Mol %) and Rha (2.4 Mol %) (Table 3 and Fig. 3b,
380 right panel). Proportions of other monomers comprised less than 0.5 mol % and declined in
381 the following order: Fuc > GalN > GlcN (Table 3). In pCCHO, Glc was the most abundant
382 sugar (74 ± 4 Mol %), followed by Rha (6.6 ± 0.8 Mol %) and Gal-URA (5.2 ± 0.7 Mol %).
383 Man/Xyl, Ara and Glc-URA ranged between 3.2 ± 1.7 and 4.2 ± 1.0 Mol % while Fuc and the
384 amino sugars GalN and GlcN contributed only a minor fraction (< 0.5 Mol %) to pCCHO
385 (Table 3).

386 Hence, composition of pCCHO was substantially different from the composition determined
387 for freshly produced HMW-dCCHO ($p < 0.002$), except for the proportions of Gal and Glc-
388 URA. This difference is mainly attributed to a smaller proportion of Glc in the dissolved
389 fraction along with a more than 10 fold higher share of Ara and also higher proportions of
390 Man/Xyl and Gal-URA (Table 3).

391

392 Carbohydrate composition of the investigated dCCHO size fractions was significantly
393 different also ($p < 0.002$). Ara was dominant in *very large*, *large*, and *medium* HMW-dCCHO,

394 but not in *the small* fraction in which its contribution was significantly smaller than in all
395 other size classes ($p < 0.01$). Most interestingly, the proportion of Glc increased with
396 decreasing size class, while the proportion of Gal-URA clearly decreased (Fig. 4). In *small*,
397 Glc contribution was 80 ± 12 Mol % and significantly higher than in all other size classes of
398 HMW-dCCHO ($p < 0.002$) (Table 3). Contribution of Glc to *very large* dCCHO was
399 negligible (< 0.5 Mol %). In contrast, Gal-URA contributed 18 Mol % to *very large*, but only
400 1 Mol % to *small*. Proportions of Gal were also decreasing the smaller the HMW-dCCHO
401 size classes, albeit not as clearly as for Gal-URA. Gal ranged from 6 (*very large*) to < 0.5 Mol
402 % (*small*). Contributions of Rha and Man/Xyl varied among size classes. Mol % of Fuc and
403 both amino sugars, GalN and GlcN were negligible.

404 **4. Discussion**

405 *4.1 Particulate and dissolved primary production*

406 Nutrient limitation and low growth rate did not hamper organic carbon production of
407 *Emiliana huxleyi* during the present study. Cell normalized production of PO^{14}C was on
408 average $\sim 0.33 \text{ pmol C cell}^{-1} \text{ d}^{-1}$ and well within the range of published values (0.12 – 0.64
409 $\text{pmol C cell}^{-1} \text{ d}^{-1}$; Biddanda and Benner 1997, Borchard and Engel, 2012). The partitioning of
410 organic carbon between dissolved and particulate pool was shown earlier to be highly
411 influenced by environmental conditions such as light, temperature and nutrient supply
412 (Myklestad and Haug, 1972, Zlotnik and Dubinsky 1989, Staats et al., 2000, Wetz and
413 Wheeler, 2007). Nutrient depletion, however, seems to be the major factor leading to excess
414 DOC excretion from algae cells to the surrounding environment and was reported from a
415 variety of field and lab experiments (Fogg, 1983, Wood and VanValen 1990, Smith and
416 Underwood, 2000, Lopez Sandoval 2010, 2011). Extracellular release (ER) in the range from
417 0-80 % was reported over the past decades and only after a long lasting debate primarily
418 concerning methodological constraints (Sharp, 1977, Mague, 1980, Fogg, 1983, Bjørnsen,
419 1988), it is nowadays accepted, that ER is a normal function of healthy algae cells occurring
420 during all stages of growth. In exponentially growing cells in culture, ER typically ranges
421 between 2 and 10 %, while in natural marine environments ER is generally higher by 10-20
422 % (see Nagata, 2000 and references therein). A relatively constant percentage of extracellular
423 release (PER) of 20 % was reported for field samples over different ecosystems covering
424 oligotrophic and eutrophic regions (Marañón et al., 2005). Increased PER (up to 37 %)
425 however, were observed for nutrient limited algae, during the transition period of exponential
426 to stationary growth and during senescence of natural phytoplankton communities (Lopez-
427 Sandoval, 2010, 2011, Engel et al. 2013). In chemostats, despite the strict control of nutrient
428 supply and growth rate, cells still grow exponentially. A decoupling of carbon to nutrient
429 metabolism in continuous cultures can occur due to a change in growth rate (e.g. change the
430 inflow of nutrient media) and results in changes in the partitioning between dissolved and
431 particulate carbon pools, as shown with the same *E. huxleyi* strain (B 92/11) by Borchard and
432 Engel (2012). In their study, down-regulation of the growth rate from $\mu=0.3 \text{ d}^{-1}$ to $\mu=0.1 \text{ d}^{-1}$
433 induced a slight increase in DO^{14}C production, while the PO^{14}C production was significantly
434 minimized, resulting in higher PER. Cells then adapted to the steady state and high PER
435 remained constant. During the present study, growth of *E. huxleyi* was also balanced to the
436 nutrient supply but cells were not exposed to any stress due to nutritional changes. Thus,
437 production of DO^{14}C was not explicitly stimulated by changing experimental conditions, and,
438 albeit constantly P-limited, the cell normalized DO^{14}C production of $\sim 0.015 \text{ pmol C cell}^{-1} \text{ d}^{-1}$

439 represented an ER of ~4.5 %, well within the above mentioned range for non-stressed algae.
440 Full acclimation to environmental conditions during steady state growth may also explain the
441 absence of a CO₂ effect on primary production and exudation during this study, and shows
442 that *E. huxleyi* is in principle capable of acclimating to different CO₂ concentrations. Engel et
443 al. (2014) suggested that exudation may be more sensitive to changes in *p*CO₂ during
444 transient growth phase, such as towards the end of phytoplankton blooms, when cells become
445 nutrient limited. Indeed, significant responses of ER to changes in *p*CO₂ have mainly been
446 reported for phytoplankton blooms (Engel et al. 2013), batch and semi-continuous cultures
447 (Thornton 2009, Barcelos e Ramos 2014), or when growing conditions changed during
448 chemostat studies (Borchard and Engel 2012).

449

450 4.2 Combined carbohydrate production

451 HMW-dCCHO freshly produced by *C. huxleyi* during steady state growth represented about
452 40 % of freshly produced DO¹⁴C (Table 2). This is a lower estimate because low molecular
453 weight-DOC (<1 kDa, LMW) would be detected by the ¹⁴C-incubation method (Stemann
454 Nielsen, 1952) during the determination of DO¹⁴C, but would escape the analysis of HMW-
455 dCCHO due to the molecular cut off >1 kDa during desalinization of seawater samples
456 (Engel & Händel 2011). In the surface ocean, HMW compounds of dissolved organic matter
457 (DOM) were found to be more abundant (30-35 %) compared to deeper waters (20-25 %) and
458 it was concluded that HMW-DOM inherits a higher reactivity and shorter lifetimes,
459 while LMW-DOM is rather refractory (Amon and Benner, 1996, Ogawa and Tanoue, 2003).
460 Major reaction processes of HMW compounds are heterotrophic degradation (Amon and
461 Benner, 1996, Guo et al., 2002, Aluwihare and Repeta, 1999) and gel particle formation
462 (Mari & Burd, 1998, Leppard, 1995, Passow, 2000, Passow 2002 and references therein).
463 Thus, the HMW-DOM pool is directly linked to processes significant for organic carbon
464 dynamics, nutrient cycling and oxygen consumption in the ocean. Assembly and coagulation
465 of polymeric precursors has been proposed as mechanism leading to the formation of marine
466 gel particles, such as TEP. Specifically, divalent cation bridging of acidic sugars, such as
467 uronic acids is assumed to be involved in bonding between polysaccharide chains. The
468 release of larger polysaccharides with relatively high Mol % Gal-URA as observed for *E.*
469 *huxleyi* in this study may be an important first step for high TEP concentrations, observed
470 previously (Engel et al. 2004, Harlay et al. 2009). However, absolute rates of ER were
471 relatively low and apparently insufficient to induce TEP formation during this study. Engel et
472 al. (2014) suggested that responses to variations in environmental factors, specifically to
473 changes in nutrient supply, are responsible for excess carbon accumulation inside the cell and

474 for exudation of carbohydrates. Sampling during this study was conducted during the period
475 of steady state growth. This may explain the observed relatively low rates of ER, including
476 potential TEP precursors.

477

478 4.2.1 Monomeric composition of CCHO

479 Natural seawater (NSW) used in the present study to prepare the nutrient media was collected
480 from the North Sea and kept under dark and cool conditions for several months before usage.
481 HMW-dCCHO monosaccharide composition of NSW was dominated by Glc (24 Mol %) and
482 Man/Xyl (24 Mol %). Also high Mol % (~10) of Fuc, Gal, Gal-URA and GlcURA
483 were determined, while other monomers were of minor importance (Fig. 3b, left panel). The
484 composition of the aged NSW used here differs from those obtained from the Northwest
485 Atlantic, the Sargasso Sea and the Gulf of Mexico (Aluwihare et al., 1997 and references
486 therein), especially concerning comparably low proportions of Rha and Gal (Fig. 3b, left
487 panel). Differences in carbohydrate composition of the seawater can be explained by seasonal
488 or geographical divergences as well as by storage time of NSW. Monomeric composition of
489 HMW-dCCHO released by *E. huxleyi* during the present experiment was substantially
490 different from the initial NSW composition (Fig. 3b) and the compositional shift was
491 primarily induced by a profound relative increase in Ara. The HMW-dCCHO and pCCHO
492 derived from *E. huxleyi* during this experiment contained a similar composition as
493 determined earlier for cellular and extracellular carbohydrates derived from this species (De
494 Jong et al. 1979, Fichtinger Schepman et al. 1979, Nanninga et al. 1996, Bilan & Usov 2001).
495 Cellular pCCHO of *E. huxleyi* differed clearly not only from NSW but also from HMW-
496 dCCHO (Fig. 3b, right panel). This is in accordance with previous studies showing
497 differences between intracellular and extracellular CCHO compositions for various algae
498 (Mague, 1980, Aluwihare, 1999, 2002).

499

500 Neutral sugars generally dominated the HMW-dCCHO composition with ~83 mol %. These
501 results are consistent with findings by Aluwihare (1999), who report on HMW exudates from
502 *E. huxleyi* being mainly composed by neutral polysaccharides with Ara as the dominant
503 monomer (30 Mol %). However, the fraction of Ara observed during this study is
504 considerably higher than reported for ultrafiltered DOM (>1 kDa) by Biersmith and Benner
505 (1998) who also investigated non-axenic *E. huxleyi* as batch culture, and for HMW-dCCHO
506 sampled during a field study in the Bay of Biscay, when coccolithophores and presumably *E.*
507 *huxleyi* was the dominating phytoplankton organism (Engel et al. 2012); both studies
508 reported Ara of ~3 % Mol. Apart from well documented species specific differences in

509 CCHO composition (Aluwihare and Repeta, 1999, Mykkestad, 1974, Mykkestad et al., 1989),
510 variations in the composition of algal extracellular carbohydrates may be related to
511 physiological and ecological functions. Although freshly produced DOC is generally a
512 primary substrate for heterotrophic uptake, *E. huxleyi* exudates were shown to exhibit
513 recalcitrant features (Nanninga et al., 1996). Degradation experiments with the diatom
514 *Thalassiosira weissflogii* revealed a special role of Ara in carbohydrate accessibility, as it
515 escaped bacterial degradation over a period of two weeks (Aluwihare and Repeta, 1999).
516 Bacterial cell numbers during the present experiment were relatively high, between 2 and 3 x
517 10⁶ mL⁻¹, contributing ~2 % to particulate organic carbon (POC) and ~3 % to DOC (Engel et
518 al. 2014). Assuming a bacterial growth efficiency of 60 % (upper limit, Del Giorgio and
519 Cole, 1998), the bacterial carbon demand could have been about 2 % of POC and 5 % of
520 DOC. Relative to the freshly produced DO¹⁴C derived from rate measurements, however, a
521 share of up to 20 % may have been channeled into heterotrophic turn-over. This means that e
522 ER would be underestimated by 20% at most. The HMW-CCHO was thus to some extent
523 subject to bacterial reworking and the high proportions of Ara may be a result of the selective
524 removal of other monomers. In accordance with the findings of Aluwihare (1999),
525 concentration of Ara in dCCHO remained unchanged during a degradation experiment with
526 the same *E. huxleyi* strain investigated here, while dCCHO were reduced by ~60 % (Piontek
527 et al. 2010; J. Piontek pers. comm., 2014). However, we would expect that extensive
528 microbial degradation of larger dCCHO would lead to an increase of Ara Mol % in the *small*
529 size fraction. But this was not observed.

530

531 Alternatively, high Mol % Ara and low Mol % Glc may indeed be a characteristic of larger
532 carbohydrate molecules released by *E. huxleyi* that are recalcitrant to microbial
533 decomposition. Assuming these components are bad substrates for microbial utilization, their
534 controlled exudation, if physiologically necessary, may be ecologically advantageous for
535 algal cells that are competing with bacteria for nutrients such as phosphorus. This
536 corroborates earlier findings of DOM produced at P-depletion being more resistant to
537 bacterial degradation (Obernosterer and Herndl, 1995, Puddu, 2003). On the other hand
538 bacteria recycle organic phosphorus and a certain degree of bacterial activity will be
539 advantageous for regenerated productivity of algal cells. So far, little is known on how
540 nutrient limitation affects the composition of algal release products. We suggest that nutrient
541 availability may be one factor responsible for variability in carbohydrate composition
542 observed during various studies (Giroldo et al. 2005, Goldberg et al. 2010, Engel et al. 2013).
543 Assuming a certain degree of microbial modification, another explanation for the difference

544 of CCHO composition between culture studies, and those observed in natural seawater may
545 be the highly specific linkage between algal release and bacterial community response,
546 proposed by a series of recent studies (Teeling et al 2012, Taylor et al. 2014, Kabisch et al.
547 2014). These showed that the release of algal polysaccharides can induce a succession of
548 bacterial communities inhabiting different abilities for enzyme expression related to specific
549 carbohydrate degradation. Because the majority of marine bacteria cannot be kept in culture,
550 bacteria present in this chemostat study, and likely in all culture experiments, represent only a
551 small fraction of the natural diversity. The bacteria present in this study may have left a
552 different fingerprint on polysaccharide composition than natural communities. Short-term
553 incubation studies with natural bacterial communities may be required to better understand
554 the microbial fingerprint on DOM, specifically polysaccharide degradation. A better
555 understanding of the microbial fingerprint on DOM could also allow for tracing microbial
556 degradation activities in specific environments, such as the ocean's anoxic zones, or the
557 extreme oligotrophic seas.

558

559 4.2.2 Size fractionation of CCHO and DOC – Considerations on extracellular release

560 Quantitatively, each DO^{14}C size fraction contributed similar amounts to total DO^{14}C with
561 slightly higher proportions in the *very large* fraction (Fig. 1 and Table 2). Release rates of
562 HMW-dCCHO were similar for the different size fractions, but highest in the *very large*
563 fraction (Fig. 2, Table 2). On a total basis, ~40 % of produced DO^{14}C were characterized as
564 freshly produced HMW-dCCHO (Table 2). Contribution of dCCHO to fresh DOC was
565 lowest in the *small* size fraction (30 %) and highest in the *very large* (60 %) fraction (Table
566 2). Monomeric composition of different size classes of dCCHO enriched by *E. huxleyi*
567 exudates was profoundly different from those of the aged NSW used as culture media (Fig.
568 3). In aged NSW, monomers were more evenly distributed among size fractions (Fig. 3b, left
569 panel). In comparison, differences in monomeric composition of size classes in *E. huxleyi*
570 exudates were largely due to changes in Ara, Glc, and Gal-URA. Most remarkably, Ara the
571 dominant monomer in all larger dCCHO size classes, was of minor importance in the *small*
572 dCCHO size fraction and lowest in the particulate fraction (Fig. 3, right panel). This is in
573 accordance with the findings of Biersmith and Benner (1998), who also observed lower Mol
574 % Ara for particulate components of an *E. huxleyi* culture as well as for the cell lysate. In
575 contrast to Ara, Mol % Glc in our study was highest in the particulate and small fraction,
576 relatively small in the *medium to large*, and negligible in the *very large* fraction. This also
577 agrees well to earlier findings; Skoog et al. (2008) observed larger Mol % of Glc in LMW-
578 CCHO than in HMW-CCHO, while reporting less Mol % Ara in LMW- than in HMW-

579 CCHO. Thus, differences in size fractions of combined sugar molecules may be one factor
580 responsible for differences in CCHO composition of DOC between study sites.

581

582 In general, carbohydrate composition in the smallest size class was similar to cellular
583 *p*CCHO composition, while larger molecules were more distinct (Fig. 3, right panel). The ¹⁴C
584 method (Steemann Nielsen, 1952), applied here to measure primary production and ER of
585 organic carbon does not allow distinguishing if DOC is released from the cell passively, i.e.
586 by leakage, or actively by exudation. Leakage is hypothesised to be directly related to
587 biomass and cell size, suggesting a constant value of passive PER. The composition of the
588 *small* size class, and particularly the high share of Glc, resembled the cellular carbohydrate
589 composition (Fig. 3b, right panel). This finding suggests a non-selective, i.e. passive, release
590 of carbohydrates in the smallest size class determined here. Storage glucans in algae are
591 comprised exclusively by Glc in D formation and have a molecular weight of 5 – 10 kDa. D-
592 Glc was reported as major component of coccolith polysaccharide (CP) of *E. huxleyi*
593 (Fichtinger Schepman, 1979). For chloroplasts in higher plants, porins are described that
594 allow trans-membrane passage of hydrophilic molecules like sugars and amino acids up to a
595 molecular weight of 10 kDa without the use of energy (Flügge and Benz, 1984; Mohr and
596 Schopfer, 1992). The existence of porins in cell membranes of algae is likely but not
597 explicitly reported. If $DO^{14}C > 1$ and < 10 kDa and associated carbohydrates leak from the cell
598 in accordance to the passive diffusion model, this extracellular release is presumably linear
599 correlated to biomass (*property tax* – Sharp 1977). For molecules > 10 kDa, however,
600 different mechanisms for the extracellular release are to be expected, since larger molecules
601 cannot pass the membrane by diffusion, and CCHO composition clearly differs from
602 intracellular CCHO (Fig. 3b, right panel). If active release, i.e. exudation, follows the
603 overflow model, biomass growth and dissolved primary production might be strongly
604 decoupled (*income tax* – Sharp 1977). Moreover, exudation requires a series of physiological
605 processes involved in the synthesis, transport and trans-membrane release of exudates.
606 Hence, exudates likely vary in composition. Data obtained during the present study indicate,
607 that components > 10 kDa, rich in Ara and Gal-URA and poor in Glc, are transported actively
608 through the cell membrane.

609

610

611

612 **5. Conclusion**

613 Carbohydrates of high molecular weight (>1 kDa) as a product of primary production are
614 released from nutrient limited *E. huxleyi* during steady state growth. Compositional
615 difference between size fractions of combined carbohydrate suggest that dCCHO >10 kDa
616 are released by active exudation across the cell membrane whereas lower molecular weight
617 carbohydrates (<10 kDa) can pass the membrane passively by leakage. The underlying
618 mechanism of the release, however, needs to be further elucidated. If the presence of Ara is
619 indeed an indicator for less degradable exudates as suggested by this study or, if Ara
620 degradation requires activities of specific bacterial assemblages, needs further exploration,
621 i.e. by using axenic phytoplankton cultures combined with the addition of natural
622 bacterioplankton communities. At present our understanding of how microbial processes
623 shape the molecular composition of DOM, specifically of carbohydrates, is still at its infancy.
624 This study suggests that dCCHO composition and size may be valuable indicators of
625 processes related to autotrophy such as primary production and exudation, but may also keep
626 the fingerprint of heterotrophic degradation. A better understanding of compositional changes
627 in dCCHO, as major fraction of semi-labile DOC, may therefore help to unravel carbon
628 cycling and ecosystem dynamics in the ocean.

629

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Table 1: Definition of size classes for fractionated high molecular weight (HMW, >1 kDa) dissolved combined carbohydrates (HMW-dCCHO) and dissolved organic carbon (DO¹⁴C).

	HMW-dCCHO	DO¹⁴C
Total	1kDa < HMW-dCCHO < 0.45 μm	DO ¹⁴ C < 0.40μm
Very Large	1000 kDa < HMW-dCCHO < 0.45 μm	1000 kDa < DO ¹⁴ C < 0.40μm
Large	100 kDa < HMW-dCCHO < 1000 kDa	100 kDa < DO ¹⁴ C < 1000 kDa
medium	10 kDa < HMW-dCCHO < 100 kDa	10 kDa < DO ¹⁴ C < 100 kDa
Small	1kDa < HMW-dCCHO < 10 kDa	DO ¹⁴ C < 10 kDa

Table 2: Size class resolved production rates of high molecular weight (>1 kDa) carbohydrates (HMW-CCHO) and of fresh organic carbon (^{14}C) during the chemostat experiment, as well as contribution of carbon contained in HMW-CCHO to primary production (^{14}C) in particulate matter and in different size fractions of dissolved organic carbon . Values represent averages \pm standard deviation of replicate samplings and both treatments, $n=6$.

	HMW - CCHO $[\mu\text{mol C L}^{-1} \text{d}^{-1}]$ avg \pm sd	^{14}C $[\mu\text{mol C L}^{-1} \text{d}^{-1}]$ avg. \pm sd	HMW-CCHO : ^{14}C [%] avg. \pm sd
Particulate	20.7 \pm 5.34	171 \pm 15.9	12.5 \pm 3.54
Dissolved			
Total	3.10 \pm 0.41	8.11 \pm 0.88	40.0 \pm 5.37
Very Large	1.41 \pm 0.21	2.74 \pm 0.88	59.7 \pm 17.2
Large	0.71 \pm 0.24	2.01 \pm 0.72	44.6 \pm 24.7
Medium	0.62 \pm 0.25	1.27 \pm 0.53	52.9 \pm 33.7
Small	0.62 \pm 0.18	2.09 \pm 0.32	29.5 \pm 9.30

Table 3: Freshly produced combined carbohydrates (CCHO) in various size fractions. Average values (**bold**) and standard deviations (*italics*) in Mol % CCHO are given for replicate samplings and both treatments., n=6. Fuc, GalN and GlcN were always <0.5 Mol % and are not included.

CCHO [Mol %] Size fraction	Rha	Ara	Gal	Glc	Man/Xyl	Gal-URA	Glc-URA
pCCHO	6.56 <i>0.84</i>	3.69 <i>0.99</i>	3.09 <i>1.45</i>	74.0 <i>4.08</i>	3.22 <i>1.65</i>	5.18 <i>0.68</i>	4.23 <i>1.01</i>
HMW-dCCHO (total)	2.44 <i>0.70</i>	46.0 <i>3.0</i>	4.64 <i>1.95</i>	20.5 <i>7.48</i>	9.68 <i>2.37</i>	11.0 <i>4.40</i>	5.74 <i>2.99</i>
very large	3.46 <i>1.88</i>	54.2 <i>13.3</i>	6.34 <i>4.11</i>	<0.5 <i>-</i>	9.62 <i>6.35</i>	18.2 <i>5.35</i>	8.17 <i>6.11</i>
large	0.91 <i>0.85</i>	41.0 <i>24.2</i>	5.92 <i>5.10</i>	18.9 <i>11.2</i>	16.5 <i>18.9</i>	8.45 <i>13.3</i>	8.34 <i>13.41</i>
medium	1.71 <i>1.03</i>	48.8 <i>9.41</i>	3.54 <i>3.41</i>	34.9 <i>19.2</i>	8.20 <i>9.88</i>	2.96 <i>3.35</i>	<0.5 <i>-</i>
small	2.25 <i>1.54</i>	9.70 <i>6.17</i>	<0.5 <i>-</i>	79.8 <i>11.9</i>	4.03 <i>3.74</i>	1.13 <i>3.35</i>	2.64 <i>2.89</i>

pCCHO: particulate combined carbohydrates; HMW-dCCHO: high molecular weight (> 1 kDa, HMW) dissolved combined carbohydrates; Rha: rhamnose; Ara: arabinose; Gal: galactose; Glc: glucose; Man/Xyl: co-eluting mannose and xylose; Gal-URA: galacturonic acid; Glc-URA: glucuronic acid;

Figure 1

Dissolved (DO¹⁴C, left) and particulate (PO¹⁴C, right) primary production [$\mu\text{mol C L}^{-1} \text{d}^{-1}$] of *Emiliana huxleyi* at present day (filled bars) and high CO₂ (open bars) conditions. Daily rates are additionally given for each DO¹⁴C size fraction. Each bar corresponds to the average (\pm standard deviation) of replicate samplings (sampling 1-5, $n=5$) accomplished during the steady state period of the experiment.

Figure 2

Freshly produced high molecular weight (HMW > 1 kDa) dissolved combined carbohydrates (HMW-dCCHO, left), particulate CCHO (pCCHO, right) [$\mu\text{mol C L}^{-1}$] derived from *E. huxleyi* at present day (filled bars) and high CO₂ (open bars) conditions. Concentrations are additionally given for each size fraction of HMW-dCCHO. Each bar corresponds to the average (\pm standard deviation) of replicate samplings (samplings 3-5, $n=3$) accomplished during the steady state period of the experiment.

Figure 3

Concentration [$\mu\text{mol C L}^{-1}$] (a) and composition [Mol % CCHO] (b) of high molecular weight (>1 kDa) dissolved combined carbohydrates (HMW-dCCHO). Data are shown for natural seawater used to prepare the experimental culture media (left panels) and composition in natural seawater enriched with freshly produced HMW-dCCHO derived from *E. huxleyi* (rights panels) grown in chemostats. Due to the strong similarity between the present day and high CO₂ treatment, both were treated as replicates. Stacked bars show the average of replicate samplings (samplings 3-5, $n=6$) accomplished during the steady state period of the experiment.

*: Data for *HMW-dCCHO for Natural seawater* and *E. huxleyi* taken from Aluwihare (1999) for comparison. Here, only neutral carbohydrates are included, since amino- and acidic HMW-dCCHO were not analyzed.

Figure 4

Proportions of glucose (Glc) and galacturonic acid (Gal-URA) in high molecular weight (HMW > 1 kDa) dissolved combined carbohydrates (dCCHO) of different molecular weight size classes as defined in table 1. Due to the strong similarity between the present day and high CO₂ culture, both were treated as replicates. Bars show the average (\pm standard

deviation) of replicate samplings (sampling 3-5, $n=6$) accomplished during the steady state period of the experiment.

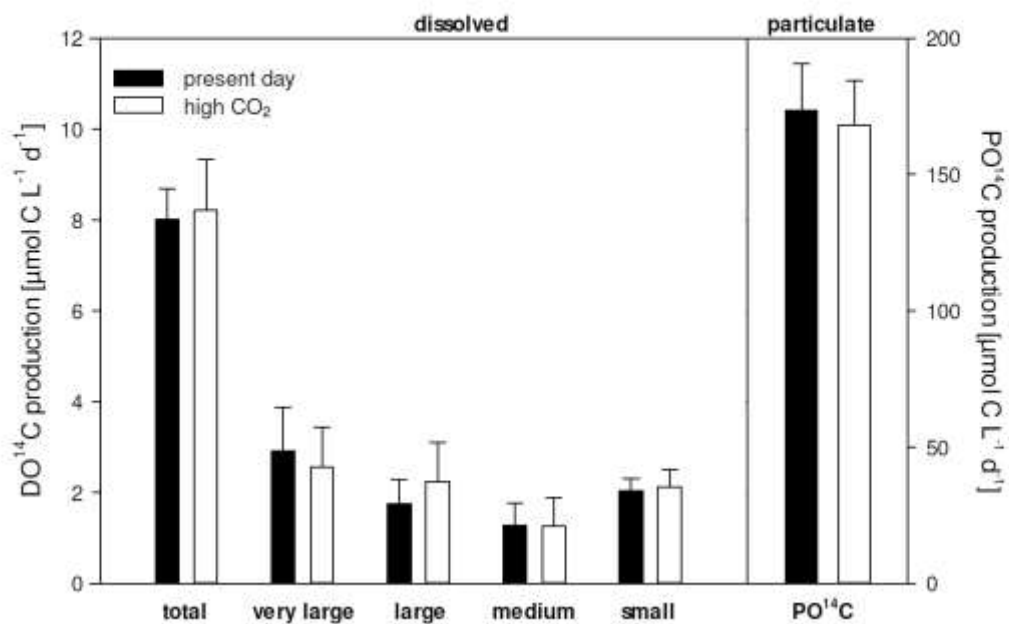


Figure 1

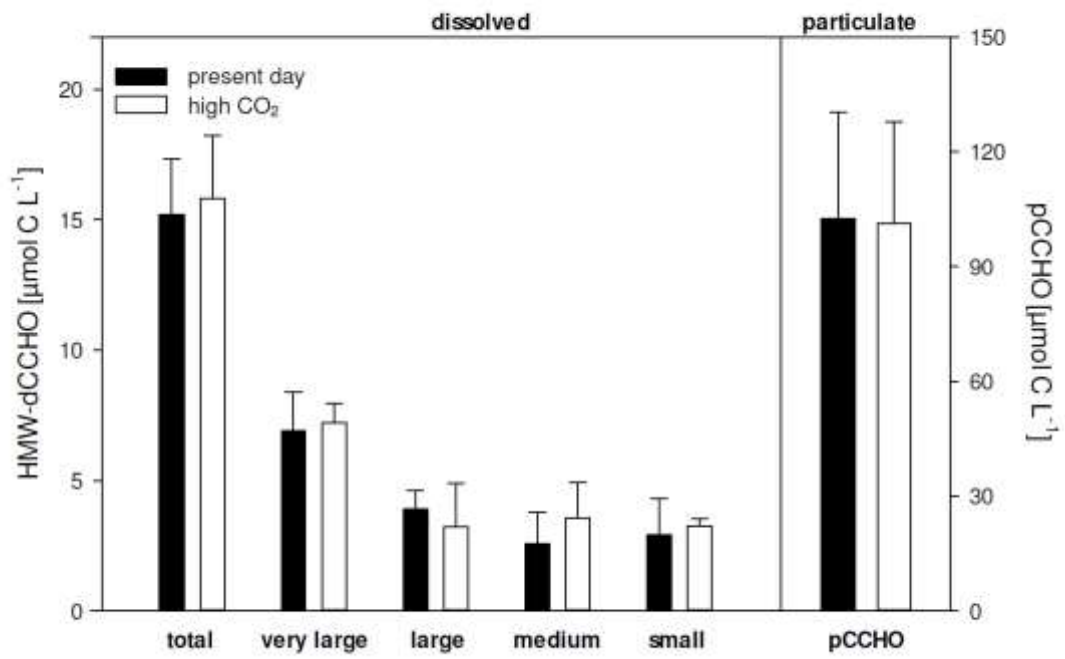


Figure 2

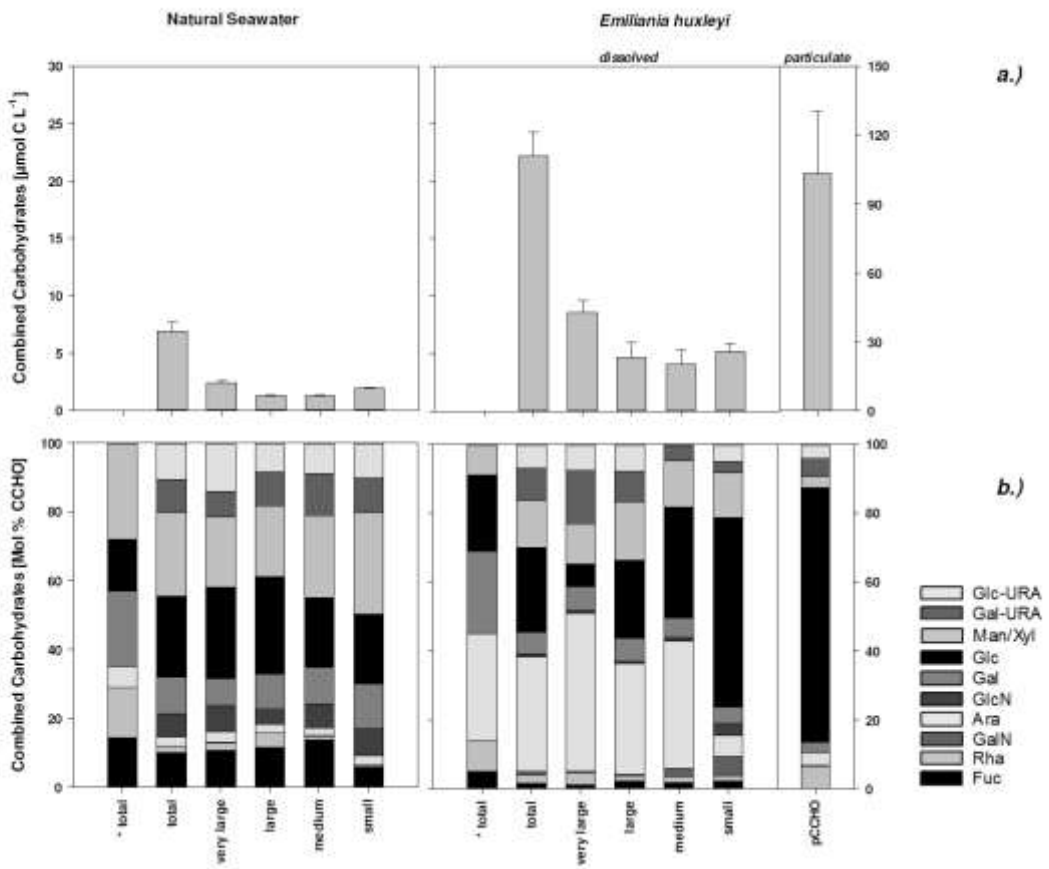


Figure 3

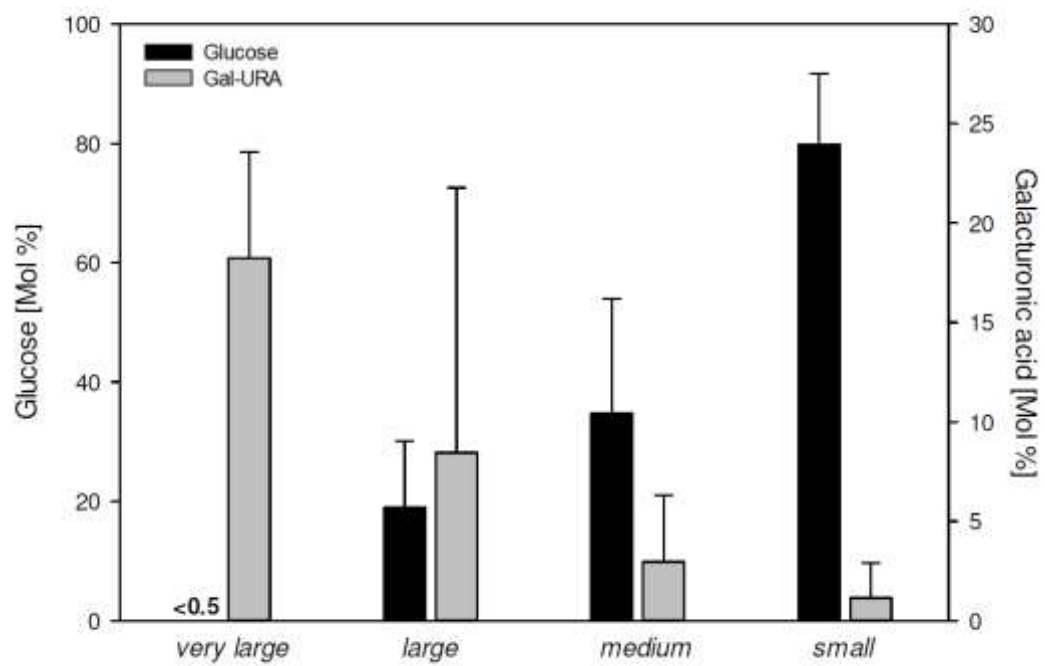


Figure 4