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Extensive cross disciplinary analysis of biological and chemical control of *Calanus finmarchicus* reproduction during an aldehyde forming diatom bloom in mesocosms

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ABSTRACT

Egg and faecal pellet production and egg hatching success of the calanoid copepod *Calanus finmarchicus* was monitored over a period of 14 days (14–28 April, 2008) while fed water from 4 differently treated mesocosms and ambient water. Two of the mesocosms used were inoculated with the polyunsaturated aldehyde (PUA) producing diatom *Skeletonema marinoi* while 2 received only nutrient additions with or without silica. The mesocosms developed blooms of *S. marinoi*, mixed diatoms or the haptophyte *Phaeocystis pouchetii*, respectively. Faecal pellet production of *C. finmarchicus* increased with increasing food availability. Egg production increased with time in all mesocosms to a maximum single female production of 232 eggs female⁻¹ day⁻¹ (average of 90 eggs female⁻¹ day⁻¹) and followed the development of ciliates and *P. pouchetii*, but was not affected by the observed high (up to 15 nmol L⁻¹) PUA production potential of the phytoplankton. The hatching success of the eggs produced on the mesocosm diets was high (78–96%) and was not affected by either aldehydes in the maternal diet or exposure to the dissolved aldehydes in the water.

Keywords: Mesocosm, *Calanus finmarchicus*, egg production, plankton interactions, *Skeletonema marinoi*

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40 **Introduction:**

41 The effect diatoms have on copepod reproduction, hatching, growth and survival has been a topic of
42 many papers for more than a decade. The studies have generally agreed that pure diatom diets are
43 not always the best food for copepods (Kleppel et al. 1991; Poulet et al. 1994; Ianora et al. 1995;
44 Jónasdóttir and Kiørboe 1996; Ianora et al. 2004). Pure diatom diets are found to have no, some,
45 and up to detrimental effects on egg production rates and especially egg hatching (Ban et al. 1997)
46 and survival of nauplii (Poulet et al. 1994; Koski et al. 2008). The effect can be diatom species
47 specific (Ban et al. 1997) but also strain dependent differences induce different responses in
48 feeding-, egg production, and hatching rates (Ask et al. 2006; Dutz et al. 2008). Different copepod
49 species also react differently in their reproductive output to specific diatom diets (Ianora et al. 2003;
50 Md Amin et al. 2011). Differences in feeding rates can though also be related to the size of the
51 diatom (Rey et al. 2001). The negative effect of diatoms on copepod reproduction and survival has
52 been explained by the differential ability of diatom species/strains to generate polyunsaturated
53 aldehydes (PUA) while mechanically handled by copepods (Miralto et al. 1999; Pohnert et al. 2002;
54 Wichard et al. 2007). Also differential nutritional quality of diatoms (Jónasdóttir and Kiørboe
55 1996), poor nutrient uptake in the copepod gut due to rapid gut passage time when feeding diatom
56 diets (Dutz et al. 2008) have been brought forward as possible causes for the observed detrimental
57 effects.

58
59 Protein, fatty acid, sterol and other biochemical components have been shown to change
60 significantly with the growth stage of diatoms and have, for example, been related to nutrient
61 limitation and light levels (Harrison et al. 1977; Claustre et al. 1989; Jónasdóttir 1994; Brown et al.
62 1996; Klein Breteler et al. 2005). The production of polyunsaturated aldehydes from some diatoms
63 is also shown to increase with senescing of a culture or a bloom (Ribalet et al. 2007; Vidoudez and
64 Pohnert 2008). *Skeletonema marinoi* (Sarno and Singone) (previously referred to as *S. costatum*,
65 Sarno et al. 2005) is one of the diatom species that produces heptadienal, octadienal and octatrienal
66 PUAs upon cell wounding and senescence in high concentrations (Vidoudez and Pohnert 2008).
67 This diatom is a common spring blooming species in the northern waters and occurs naturally in
68 high abundances (over 500 µg C L⁻¹) in the Raunefjord and other Norwegian fjords and often
69 dominates the diatom and total algae biomass during the spring bloom (Erga and Heimdal 1984,
70 Erga 1989).

71
72 The copepod *Calanus finmarchicus* (Gunnerus) is one of the most important copepod species in the
73 boreal north-east Atlantic waters where it constitutes up to 70% of the mesozooplankton biomass
74 during summer (Fransz et al., 1991). *C. finmarchicus* ascends as copepodite stage 5 and adult from

75 diapause from great depths in the Southern Norwegian Sea (Heath 1999) and starts reproduction
76 before the initiation of the spring bloom (Richardson et al. 1999; Niehoff et al. 1999; Jónasdóttir et
77 al. 2008). However, the major egg production usually occurs during the diatom bloom (Niehoff et
78 al. 1999; Debes et al. 2008). Diatoms are important food items for *C. finmarchicus* nauplii (Irigoien
79 et al. 2003; Castellani et al. 2008) while the adult females seem to have broader selection of food
80 types. Studies often demonstrate *C. finmarchicus* females having a positive selection for diatoms
81 (Meyer-Harms et al. 1999; Koski and Riser 2006; Koski 2007), but they also select alternative food
82 such as ciliates and larger prey when available (Mayor et al. 2006; Koski 2007). Still, when diatoms
83 are not eaten in proportion to their availability they often can make up a significant part of the diet
84 for *C. finmarchicus* females and late copepodites (Nejstgaard et al. 1997; 2001a, b). Therefore the
85 diatom spring bloom is of importance for initial start of the *C. finmarchicus* population growth
86 (Debes et al. 2008; Jónasdóttir et al. 2008).

87
88 In the present study we used large mesocosms to investigate the biological and chemical effect of
89 PUA producing diatom blooms on *C. finmarchicus* growth and reproduction. Large mesocosms are
90 an ideal experimental system to investigate the simultaneous chemical and biological effects on
91 plankton communities, including copepods. Such systems allow large volume sampling that is
92 essential for chemical analyses and multiple bottle incubations without greatly affecting the
93 plankton community in the mesocosms. With different manipulation of series of mesocosm bags
94 diverse natural plankton communities can develop in parallel throughout their respective bloom
95 cycles. A number of previous mesocosm studies using similar set-ups at the site of the present study
96 the Raunefjord, the Norwegian National Mesocosm Centre, Espegrend, University of Bergen, have
97 been used to successfully provoke blooms of naturally occurring plankton in near natural conditions
98 (see discussions in Williams and Egge 1998; Nejstgaard et al. 1997, 2006).

99
100 The goal of this study was to test if the PUA producing diatom *S. marinoi* affected the reproductive
101 success, and growth of *C. finmarchicus* over a period of a bloom. *S. marinoi/costatum* has been a
102 focus of several studies showing that when used as a long term single diet negative effects were
103 observed for *Calanus helgolandicus* on egg production and hatching (Ban et al. 1997; Ianora et al.
104 2003). However, no negative effects were observed on *C. helgolandicus* egg production and
105 hatching when *S. marinoi* was mixed in the diet in a field (Irigoien et al. 2000). *C. finmarchicus*
106 reproduction was not affected by pure *S. marinoi* diets (Ban et al. 1997). These differential effects
107 could have been *Skeletonema* strain dependent. Therefore, the idea with the present study was to
108 use a *S. marinoi* strain that is known to produce high amounts of polyunsaturated aldehydes. The
109 intent was to closely follow the chemical changes associated with the respective developing blooms

110 and to compare it simultaneously with the copepod growth, egg production, hatching success and
111 naupliar survival, in as close as possible natural setting, mesocosms. This would allow us to directly
112 assess the results for effects of specific food composition such as diatom rich diets. The fatty acid
113 content of females and eggs was followed over the mesocosm blooms and those reported in another
114 manuscript along with naupliar survival (Koski et al. unpublished). Due to the vast amount of
115 different chemical and biological measurements conducted on the mesocosm water, this is one of
116 the most extensive studies to date on the biological and chemical control on egg production rates of
117 this species.

118

119 **Material and Methods**

120 *Mesocosms*

121 The experiments were conducted over a 14 day period between the 14th and 28th of May 2008 at the
122 Espegrend marine biological field station by Raunefjord, Norway (latitude 60.16 °N; longitude:
123 5.14 °E). On April 14th 6 transparent polyethylene enclosures (each 11 m³, 2 m diameter, non
124 covered) were filled with unfiltered sea water from 4 m depth just outside the mesocosms, and the
125 following day April 15th (day 1) the mesocosms received different treatments. In the present study
126 we sampled 4 of those mesocosms, labelled B, C, E and F. All mesocosms were fertilized with 0.4
127 $\mu\text{mol L}^{-1}$ phosphate and 4.24 $\mu\text{mol L}^{-1}$ nitrate. Mesocosm B had no more treatment, mesocosm C, E
128 and F had additional 3.61 $\mu\text{mol L}^{-1}$ silicate fertilization and mesocosms E and F were inoculated
129 with cultured *Skeletonema marinoi* (Strain G4, University of Bergen) corresponding to an increase
130 of the low *in situ* concentrations with ca. 400 and 1000 cells mL⁻¹ respectively (Table 1). Water
131 temperature was measured daily throughout the water column in the mesocosms with a SAIV
132 SD204 CTD.

133

134 Water was sampled from each of the mesocosms every morning. The water was immediately
135 brought to the laboratory in 5 and 20 L carboys where filtrations and incubations took place.

136

137 *Chlorophyll filtration, extraction and analysis*

138 Chlorophyll *a* (Chl *a*) concentrations were determined daily in the mesocosm bags by filtering
139 duplicate water samples onto 0.2- μm Gelman 47 mm diameter polycarbonate filters, and a single
140 serial fractionation series onto 10, 5, 1 and 0.2 μm Gelman 47 mm diameter polycarbonate filters.
141 The filters were extracted immediately in 90% acetone overnight at 4°C and analyzed using a
142 Turner Designs 10-AU fluorometer (Turner Designs, Sunnyvale, CA, USA).

143

144 *Protist plankton*

145 Protists were counted *in vivo* by a Cytobuoy™ scanning flowcytometer (CytoBuoy b.v. Woerden,
146 The Netherlands) and by a FlowCAM II™ (Fluid Imaging Technologies, Yarmouth, ME USA)
147 fitted with black and white camera. *S. marinoi* were counted in triplicates by the Cytobuoy
148 flowcytometer following the procedures outlined by Takabayashi et al. (2006). The flowcytometer
149 was unable to identify the colonial form of *Phaeocystis pouchetii* so cell concentrations were
150 counted on colony images of the FlowCAM. The FlowCAM was set in auto-image mode using a
151 4X objective and a total sample volume of ca 4.2 mL (run time 20 min) were analysed daily from
152 each mesocosm. Each sample was fed the FlowCAM through a vertical tube from a beaker mounted
153 above the vertical mounted flow-cell, in a fashion that prevented particle sedimentation in the flow
154 system. The sample was stirred by a small lab stirrer set at the lowest speed (ca 5 rpm). The
155 dimension of the flow cell, determined by the objective was 2 mm × 0.1 mm (10X) (Vitrocom,
156 Mountain Lakes, NJ, USA). *Phaeocystis* colonies were counted and the numbers of cells were
157 estimated by use of a calibrated regression between manually counted number of colonial cells per
158 colony and colony grey scale area of FlowCAM images. Major protozoan groups >15µm were
159 enumerated by the FlowCAM. Images were manually sorted into species classes such as ciliates and
160 major heterotrophic dinoflagellates. The biomass of the individual groups such as dinoflagellates,
161 ciliates, and total were converted to carbon biomass using the generic carbon-to-volume
162 relationship given by Menden-Deuer and Lessard (2000).

163

164 *Filtration for lipids and PUA and chemical analyses*

165 For PUA sampling and analysis, see Vidoudez et al. (2011b). For the determination of the fatty acid
166 and sterol profile of the plankton, 0.5-5 L were filtered on GF/C filters under moderate vacuum.
167 The filters were transferred into 4 mL brown glass vials and covered with 1 mL of methanol and 2
168 µL of internal standard (myristic acid-d27; Sigma-Aldrich, Deisenhofen, Germany), 10 mg mL⁻¹ in
169 methanol) and the samples were kept at -20°C until the end of the experiment. The samples were
170 transported frozen and stored there at -80°C until analysis. The fatty acid and sterol profiles were
171 determined simultaneously by gas chromatography-mass spectrometer (GC-MS) after
172 transesterification (modified from Lepage and Roy, 1984) and silylation: samples were cooled to 77
173 K in a liquid nitrogen bath prior freeze-drying. Five hundred µL of methanol:acetyl chloride 9:1 (>
174 99.0%, Sigma-Aldrich), 300 µL of hexane, and two glass beads were added to the freeze-dried
175 samples before tightly closing the vials. Samples were sonicated in an ultrasound bath for 10 min
176 before vortexing for 1 min and subsequently incubated at 100°C for 10 min. Samples were cooled
177 on an ice bath and 1 mL of deionised water was added and the samples were vortexed for another
178 minute. Phase separation was achieved by centrifugation. The hexane phase was then transferred
179 into 1.5 mL glass vials (Macherey-Nagel, Düren, Germany). Five hundred µL of chloroform were

180 then added to the remaining aqueous sample and the vials were vortexed for 5 min. After
181 centrifugation, the chloroform phase was combined with the hexane phase. One mL of chloroform
182 was then added to the samples and the vortex-centrifugation cycle was repeated as described above.
183 The combined organic phases were dried over anhydrous sodium sulphate and the supernatant was
184 removed from the drying agent by pipetting. The solvent was evaporated under vacuum and the
185 samples were re-suspended in 100 μ L of hexane-chloroform (1:1). After transfer into 200 μ L glass
186 inserts (Macherey-Nagel, Düren, Germany), 50 μ L of N-nethyl-N-(trimethylsilyl)trifluoroacetamide
187 (Macherey-Nagel, Düren, Germany) was added, the vials were closed with caps fitted with PTFE-
188 butyl-PTFE septa and incubated at 60°C for 20 min to allow derivatisation of the sterols. The
189 samples were then stored at -80°C until analyses by GC-MS. A GCT premier time-of-flight mass
190 spectrometer (Waters, Manchester, UK) coupled with an Agilent 6890N gas chromatograph
191 equipped with a DB-5ms 30 m column (0.25 mm internal diameter, 0.25 μ m film thickness, with 10
192 m Duraguard pre-column, Agilent, Waldbronn, Germany) was used with the following parameters:
193 GC program: 60°C for 5 min, then ramping to 300°C at 10°C min⁻¹. This last temperature was held
194 for 10 min. The injector, transfer line and ion source temperature was set to 300°C. Electron Impact
195 ionisation (EI 70 eV) was used. The samples were injected in Split 1-15 depending on the
196 concentration. Fatty acid and sterols were identified by comparison of the retention times and mass
197 spectra with authentic standards. The PUA analysed include 2,4-heptadienal, 2,4-octadienal, 2,4,7-
198 octatrienal, 2,4-decadienal and 2,4,7-decatrienal (these metabolites will be referred to in the
199 following by their names not including the numbering of the double bonds). Quantification of fatty
200 acids, sterols and PUA was validated using established analytical criteria (data not shown).

201

202 *Egg production*

203 Sampling of copepods

204 *Calanus finmarchicus* for incubations were sampled from a station in the Raunefjord 0.5 nautical
205 miles North West of the field station, by using a WP-2 net with a 200 μ m mesh size and a non-
206 filtering cod end, hauled obliquely from ca 20-m to the surface at ca 0.5 m s⁻¹. Several tows were
207 taken to obtain sufficient number of individuals. On deck, the content of the cod end was
208 immediately diluted with ambient water and the sample was brought back in thermo boxes to the
209 laboratory within 2 hours from the first net tow, and was kept cool with cooling elements. All
210 handling of copepods, incubations and counting of eggs and faecal pellets was carried out in a dim
211 lighted 7°C temperature controlled walk-in room. At day 7 the temperature in the mesocosms had
212 increased to 8°C and the temperature the walk-in room was changed accordingly to 7.5°C.
213 Photoperiod in the walk-in room was kept at 16:8 h, dim light: dark, and all incubations described
214 below were covered.

215

216 Egg production experiments

217 Incubations started on day 2 after the inoculation of the mesocosms, with first results on day 3. For
218 the egg production experiments, 20 replicates of 600 ml beakers were set up for each mesocosm
219 treatment and an additional natural seawater control with water from outside the mesocosms. Each
220 beaker contained a Plexiglas insert with a 300 μm false bottom mesh which kept the female away
221 from her eggs and faecal pellets that in turn passed the mesh and remained on the bottom of the
222 beaker. One *C. finmarchicus* female was placed in each of the beaker inserts. Initially, the filtered
223 seawater control was set up with 8 replicate females (as above). These controls were run for only 5
224 days. On day 6, new controls were set up with 10 new females and ambient water and run for the
225 remaining experimental period. These new females came from the net tow sampled at day 1 and
226 had received ambient food (replaced daily), but their egg production had not been monitored prior
227 to day 5. Each day females were transferred to a new food suspension from the newly sampled
228 mesocosms and ambient sea water. Eggs and faecal pellets were concentrated on a 50 μm sieve and
229 gently flushed into a small Petri dish and immediately counted. Every other day the Petri dish
230 containing eggs were set aside for 3 days allowing the eggs to hatch, after which the nauplii and
231 remaining eggs were counted. The other days eggs were sampled for lipid analyses (Koski et al.
232 unpublished). Daily, 500 ml of water from each mesocosm were filtered through 50 μm sieve to
233 account for background eggs and faecal pellets.

234

235 *Faecal pellet measurements and calculations*

236 Faecal pellets were sampled every other day after counting and fixed in 4% formalin for later size
237 measurements. Pellets were measured by using an inverted microscope at 100 x magnification.
238 Length and width of at least 30 pellets were measured for each of the sampling dates. Volume was
239 calculated as the volume of a cylinder.

240

241 *Statistics*

242 Comparisons between mesocosms were done with One-way ANOVA, but with Kruskal Wallis
243 ANOVA on Ranks if data could not be normalized. General Linear Method 2-way ANOVA was
244 used to compare faecal pellet sizes. Holm-Sidak or Dunn's *post hoc* pair-wise comparisons were
245 carried out when the ANOVAs gave significant differences. Data reduction was conducted on fatty
246 acid, sterol, microplankton and chlorophyll by use of principle component analysis (PCA). A proxy
247 variable, representing the group of significantly correlated variables on each different principle
248 components, was tested against egg production, faecal pellet production and hatching success using
249 stepwise multiple regression analysis. Hatching values were arcsine transformed and egg and faecal

pellet production data were square-root transformed, to meet normality and equal variance requirements. The statistical packages Sigma Stat and PASW were used for the analyses.

Results

Mesocosm development, chlorophyll a, phyto- and microplankton

The temperature increased in the mesocosms and in the ambient water from 6.3°C on the first day to 8.7°C at the last day (data not shown). The temperature of the incubations was kept constant at 7 °C for the first 6 days and increased to 7.5°C at day 7 till the end of the experiment. The temperature change in the mesocosms did therefore not transfer to the incubations.

The total Chl *a* concentrations showed that all manipulated mesocosms developed phytoplankton blooms, while the ambient water remained at low post-bloom concentrations throughout the experimental period (Fig. 1a). An un-manipulated mesocosm, mesocosm A, not presented in this study, developed the same way as the ambient outside the mesocosms indicating that the mesocosms enclosure did not generate artificial growth conditions (H.H. Jakobsen, J.C. Nejstgaard and G. Pohnert pers. com). Mesocosm F inoculated with the highest amount of *Skeletonema marinoi* both reached the highest total Chl *a* concentration and culminated at day 7, one day before all other mesocosms. Mesocosm B and C showed a less pronounced peak with the largest fraction of the Chl *a* (> 10 µm) culminating around day 12 (Fig. 1b). Mesocosm E inoculated with a lower amount of *S. marinoi* showed an intermediate development in Chl *a*. *S. marinoi* were a major component of the phytoplankton in both mesocosms E and F, and reached peak concentrations of 133 and 480 µg C L⁻¹, respectively (Fig. 2a). *S. marinoi* were also detected in the other mesocosms (B and C) but at extreme low concentrations (< 6 µgC L⁻¹). Colonies of *Phaeocystis pouchetii* (Hariot) Lagerheim followed the decline of the *S. marinoi* bloom in mesocosms E with maximum concentrations ranging between 400 and 670 µgC L⁻¹ and coincided with the *S. marinoi* bloom in mesocosms F maintaining concentrations about 300 µgC L⁻¹ during days 8-10. In mesocosms B and C *P. pouchetii* colonies were the dominant phytoplankton biomass >15 µm (peak 1300 and 1200 µgC L⁻¹ respectively, Fig. 2). Heterotrophic dinoflagellates (sum of *Gyrodinium spirale* (Bergh) Kofoid and Swezy, *G. dominans* Hulbert and *Protoperidinium bipes* Poulsen) were found in similar concentrations in all mesocosms (100-150 µgC L⁻¹; Fig. 2a). While the concentrations of ciliates were below 100 µgC L⁻¹ in both the ambient water and the mesocosm B, the ciliates reached higher concentrations in mesocosms with *S. marinoi* (C, E, F), with the highest concentration in mesocosm E (400 µgC L⁻¹, Fig 2). The relative contribution of the protist types from day 4 and onward is presented in Fig. 2b.

285 *Chemical composition*

286 Fatty acids

287 The polyunsaturated fatty acids (PUFA) in mesocosms B and C were dominated by 18:3(n-6) and
288 22:6(n-3) (Table 2). In mesocosm C that had received the additional Si addition the fatty acid
289 16:1(n-7) which is typical for diatoms could be detected in higher amounts than in mesocosm B.
290 Algae from mesocosm E and F had considerably higher proportion of 16:1(n-7) and 20:5(n-3) than
291 those of mesocosms B and C. High levels of 22:6(n-3) were also detected in these mesocosms but
292 highest in mesocosm F (Table 2). The monounsaturated fatty acids (MUFA) and PUFA followed
293 the same pattern as the bloom development (Fig. 3).

294

295 Sterols

296 The main particulate sterols were cholesterol, brassicasterol and sitosterol (Fig. 3). The sterol
297 content (total sterol) was significantly correlated with *P. pouchetii* ($R^2 = 0.65$, $p < 0.001$). In
298 mesocosm F particularly high concentrations of cholesterol were detected.

299

300 PUA

301 The dynamics of the polyunsaturated aldehydes is described elsewhere (Vidoudez et al. 2011).
302 Heptadienal and octadienal production was correlated to *S. marinoi* only, while decadienal
303 concentration was correlated to the combined densities of *P. pouchetii* and *S. marinoi*.

304

305 *Data reduction*

306 Data reduction (PCA) sorted the food environment into 4 major components that explained 87% of
307 the variance of the variables in the analysis (Table 3). Component 1 was composed of the
308 introduced diatom *S. marinoi*, hepta-, octa- and decadienal and Chl *a* >10µm, and to a lesser degree
309 cholesterol and monounsaturated fatty acids (MUFA). The second component was composed of *P.*
310 *pouchetii*, the brassica-, campe- and sitosterols. The third component was based on the ciliates, the
311 heterotrophic dinoflagellates, *G. spirale* and *P. bipes* and to a lesser degree on *G. dominans*.
312 Component 4 was composed of the PUFAs 22:6(n-3), 18:3(n-3) and 18:3(n-6) and to a lesser degree
313 20:5(n-3).

314

315 *Egg production*

316 After initial drop from the first to the second day of incubation (between days 3 and 4) average egg
317 production rates (EPR: eggs female⁻¹ day⁻¹) in all mesocosms increased gradually with time from ca
318 10 eggs female⁻¹ day⁻¹ till ca 80 eggs female⁻¹ day⁻¹ in the end of the experiment (Fig. 4a). EPR in
319 filtered seawater (FSW) dropped to 0 at the third measurement (day 5) while in the ambient

seawater (day 7-14) the average EPR measured 12 ± 7 eggs female⁻¹ d⁻¹ (\pm SE) and did not change significantly over time (Kruskal Wallis one-way ANOVA, $H_6 = 7.1$, $p = 0.314$). There were no significant differences in egg production rates between mesocosms when compared over the whole period (days 5-14), but the egg production rates at the final day (day 14) differed significantly between the ambient water and each of the mesocosms C, E and F ($p < 0.05$, Dunn's *post hoc* comparison).

The maximum daily egg number was highest (232 eggs) for a female incubated in mesocosm E water, but similarly high maximum numbers were also in the other mesocosm treatments (167, 173 and 202 eggs from water from mesocosm F, B and C respectively, Table 1).

The fraction of spawning females increased with time (Fig. 4b) but was significantly lower in the ambient food treatment compared to all the mesocosm treatments (Holm-Sidak *post hoc* comparison of means, $p > 0.05$). Additionally, there was a significant difference between the fraction of females producing eggs in waters from mesocosm B and C (Holm-Sidak *post hoc* comparison of means, $p > 0.05$).

The cumulative egg production was highest in the treatment with water from mesocosm F but lowest and almost half the value in the treatment from mesocosm B (Table 1). Female mortality was highest in the incubations from mesocosms B and E (35 and 40% respectively). No female died after the 12 day incubations from mesocosm F, which also had highest percent of producing females (88%, Table 1). The cumulative egg production does not correct for loss of females in the incubations as time went on, and reflects therefore the potential production of a population receiving the different treatments.

Due to the observed 2 day adjustment in EPR to the food availability (Fig. 4a), stepwise multiple regression was run on EPR excluding the EPR on the first 2 days of the experiment (days 3 and 4) and excluding the FSW treatment. Hence, due to this 2 day delay in EPR in response to the food environment, all correlations were done on the food available 2 days earlier. The environmental variables used in the analysis were based on the PCA data reduction, but using the measures of *S. marinoi* as a representative for the highly correlated variables on principal component (PC) 1, *P. pouchetii* for PC2, ciliates for PC3 and 22:6(n-3) for PC4. Additionally the variables that had lower but significant loading on the different PCs were added separately to the regression analyses. These were cholesterol (for MUFA on PC1), *G. dominans* (PC3) and 20:5(n-3) (PC4, but also high on PC3; Table 3). Egg production rates of *C. finmarchicus* were best explained by the 22:6(n-3) fatty

acid (51% of the total variation in EPR explained) but the combination of the protoplankton types *P. pouchetii* and ciliates contributed additional 27% to the model explanation. Finally by adding *S. marinoi* to the model improvement the explanation power by extra 3% (Stepwise Multiple Regression, $R^2 = 0.81$, $F_4 = 41.7$, $p < 0.001$, Table 4, Fig. 5). To further investigate the effects of specific food types and PUA on the observed egg production rates we used the method described in detail in Irigoien et al. (2005) based on the mixing model by Jónasdóttir et al. (1998). The relation between the residuals of the regression between EPR and total protist carbon (shown in Fig 5e) were tested against the fraction of relative abundance of *S. marinoi* and *P. pouchetii* in the food environment (Fig. 6a, b). The regressions were not significant which means that none of these food types were specifically important for *C. finmarchicus* egg production rates. The same was apparent with the relative abundance of ciliates ($R^2 = 0.00$; $P = 0.87$) and dinoflagellates ($R^2 = 0.02$; $P = 0.32$; plots not shown). Similarly, no negative effects were apparent when the concentration of dissolved and particulate PUA were related to the residuals of EPR (Fig. 6c,d) ($R^2 = 0.00$ and $R^2 = 0.00$; $P = 0.87$ and $P = 0.87$ respectively).

Hatching

Hatching success of eggs increased from ca 60% on day one (where eggs made from diet previous to the incubations) to 83-89% hatching (Fig. 4c, Table 1). No difference was found in hatching success between any of the mesocosm treatments (Kruskal Wallis ANOVA on arcsine transformed hatching data: $H_4 = 4.5$, $p = 0.35$) when day 1 is excluded, and no significant difference within the treatments during the incubations. Koski et al. (unpublished) presents in further details the hatching and the eggs and naupliar development in the present study.

Faecal pellet production

Number of faecal pellets produced per female per day (Table 1) differed significantly between the mesocosm treatments (One way ANOVA, $F_4 = 6.97$, $p < 0.001$). Significant differences were observed between pellet production of copepods feeding on water from mesocosm C produced significantly more faecal pellets compared to copepods feeding on mesocosm B. The copepods in mesocosm C, E and F produced significantly higher number of pellets compared to the copepods feeding in the ambient water treatment (Holm-Sidak *post hoc* comparisons at $p = 0.05$). In all but the ambient water the faecal pellet production increased for the first 7-11 days after which the production decreased again.

Faecal pellet length and consequently their volume (Fig. 4d) increased significantly with time in all mesocosm treatments (linear regression for all mesocosms $p < 0.001$ with $R^2 = 0.54$, 0.51 , 0.19 and

0.12 for mesocosms B, C, E and F, respectively on volume, data not shown). There was a significant difference in faecal pellet volume size between the mesocosms, where pellets from mesocosm B were significantly larger than the pellets from all the other mesocosms (Holm Sidak *post hoc* pairwise comparison $p = 0.05$).

Faecal pellet production rate as faecal volume (FPR: $\mu\text{m}^3 \text{ female}^{-1} \text{ day}^{-1}$) of *C. finmarchicus* were best explained with the combination of the concentrations of ciliates and *P. pouchetii*, and to a lesser degree the 22:6(n-3) fatty acid (Stepwise Multiple Regression $R^2 = 0.71$, $F_3 = 44.0$, $p < 0.001$, Table 4). The model included the Chl *a* in the size fraction 5-10 μm as more realistic representative of PC 1 for faecal pellet production than cholesterol. The individual correlations are demonstrated in Fig. 7.

There was a significant positive relationship between FPR and egg production rates ($R^2 = 0.79$; $p >$ Fig. 8).

Discussion/Conclusion

The development of the plankton communities in the 4 mesocosms and the ambient water were well suited for a comparative study as it generated a wide variety of food types and chemical compositions at broad concentration ranges. There were mainly 2 types of systems developing, one dominated with *Phaeocystis pouchetii*, the fatty acids 18:3(n-6) and 22:6(n-3) and cholesterol, and medium levels of decadienal and low levels of heptadienal, and the other with *Skeletonema marinoi*, the fatty acids 16:1(n-7), 20:5(n-3) and 22:6(n-3) increasing with time and high levels of hepta- and decadienal PUAs. Ciliates and heterotrophic dinoflagellates increased with time and peaked at different times and different concentrations in all mesocosms.

From day 7 and onwards in mesocosms F and from day 9 and 10 the mesocosms B, C and E, food availability was very abundant ($> 800 \mu\text{g C L}^{-1}$, Fig. 2); a concentration by which *Calanus finmarchicus* should not have experienced food limitation (Båmsted et al. 1999). The maximum faecal pellet production rate was achieved between 600-1000 $\mu\text{gC L}^{-1}$ (Fig 7e), while maximum egg production was achieved between 400 and 700 $\mu\text{gC L}^{-1}$ (Fig. 5e). Similar egg production in all mesocosms (B-F) could imply that individual food types dominating the different mesocosms (Fig. 2) were not important in the determination of reproductive success. However, copepods do not necessarily have fed on them in proportion. According to the optimal foraging theory *C. finmarchicus* could have allowed itself to be selective with the high concentration of mixed diet available (Fig. 2) and avoid the least favourable food items. Based on the high PUA content of *S.*

425 *marinoi* (1.2 - 4.2 fmol cell⁻¹, Ribalet et al. 2007) we could expect avoidance of this potentially
 426 harmful diatom. There are, however, several indicators that *C. finmarchicus* in our study did not
 427 specifically avoid *S. marinoi*. The only direct measure of ingestion is from the parallel study of
 428 Barofsky et al. (2010) that report that ingestion of *S. marinoi* was highest 1-3 days after the peak of
 429 the *S. marinoi* blooms in mesocosm E and F (their mesocosm 1 and 2 respectively; measured every
 430 other day). This is during the decline of the bloom when PUAs were expected to be at its highest
 431 according to the study of Ribalet et al. (2007). The second indication of *C. finmarchicus* actively
 432 feeding on *S. marinoi* is based on fatty acid composition of the females (presented in Koski et al.
 433 unpublished). The diatom fatty acid 16:1(n-7) was found in significantly higher concentrations in
 434 females from the *S. marinoi* impregnated mesocosms E and F compared to mesocosms B and C.
 435 Koski et al. (unpublished) additionally show that the fatty acids in the eggs did not differ from
 436 seston fatty acids, which indicates no selective feeding, and no selective control of fatty acids from
 437 females to the eggs. Therefore, even though we cannot state that no selection took place we can say
 438 with certainty that *C. finmarchicus* did not avoid eating *S. marinoi*.

439
 440 Faecal pellet volume production (FPR) is a good indicator of *Calanus* feeding (e.g. Nejstgaard et al.
 441 2001b). The faecal pellet production was high (30-40 pellets female⁻¹ d⁻¹) and similar in most
 442 mesocosms (Table 1) and comparable to e.g. measurements conducted by Rey et al. (1999) on *C.*
 443 *finmarchicus*. In our study FPR (volume based) was highest in mesocosms E, F and C where
 444 *Phaeocystis* was abundant, but FPR was lower in mesocosms B (Fig. 4d, Table 1) where there was
 445 less food diversity, and *P. pouchetii* dominated most of the period with 85-95 % of the biomass.
 446 Faecal pellet production is also dependent on the diet type (Besiktepe and Dam 2002, Dutz et al.
 447 2008), but in the present study it followed food abundance fairly well up to approximately 1000µg
 448 C⁻¹ (Fig7e). On the whole (all mesocosms) the pellet production was best explained with *P.*
 449 *pouchetii*, the PUFA proxy (22:6(n-3)) and ciliate concentrations but not *S. marinoi* (Table 4).
 450 However, as argued above and by the study of Barofsky et al. (2010) it does not imply that *C.*
 451 *finmarchicus* did not eat *S. marinoi*. Our measures also show that the pellet production in the water
 452 from these *S. marinoi* inoculated mesocosms was highest one or two days after the peaks of the
 453 relevant bloom (Fig. 4d). The peak of the particulate PUA coincided with the *S. marinoi* peaks and
 454 lasted couple of days after the blooms. Therefore, ingestion of potentially deterrent PUA was thus
 455 during and right after the peak of the bloom, when Barofsky et al. (2010) recorded the main feeding
 456 on *S. marinoi*. The decrease in faecal pellet production was directly related to decrease in the food
 457 availability and coincides with the carbon concentration of protoplankton (*P. pouchetii*,
 458 dinoflagellates and ciliates, Table 4).

459

460 The egg production rate of *Calanus finmarchicus* was relatively high in most of mesocosms, both in
461 the beginning and the end of our study. We measured one of the highest egg number ever recorded
462 of a single female over 24 hr period for this species: 200 and 230 eggs for individual females in
463 mesocosms C and E (for comparison see overview in Jónasdóttir et al. 2005, 2008). Even though
464 we compared the EPR with food condition 2 days earlier, and therefore did not use the 2 last days of
465 the experiment in the analyses, the rise, peak and the fall of both *P. pouchetii* and the *S. marinoi*
466 blooms were included in the diet for *C. finmarchicus* in the analyses (Fig. 2). Food availability was
467 the main determinant for the observed EPR as strongly indicated by the close correlation between
468 FPR and EPR (Fig. 8) and the highly significant correlation between protist carbon and EPR (Fig
469 5e). The increase in spawning frequency (Fig. 4b) also follows the food availability. This
470 correlation is based on a combination of different food types that build up the best explainable
471 model for the variation in the egg production rate. Total protist carbon explained 71% of the
472 variation in EPR (non linear model; Fig 5e). A more detailed analysis of the specific food
473 environment including food quality indicators (PUFA and sterols) could however, explain 81% of
474 the variation in the linear stepwise multiple regression model. The breakdown of the analysis
475 revealed that seston PUFA (22:6(n-3)) and *P. pouchetii* together explained 71% of the variation in
476 the observed EPR. Ciliates (as a proxy for flagellates, *Gyrodinium spirale* and *Protoperidinium*
477 *bipebs*) and *Skeletonema* (as proxy for octa-, deca-, and heptadienal) made up the rest leaving only
478 19% of the variation unexplained. Interestingly, *S. marinoi* and accordingly the potentially
479 detrimental PUAs had positive, rather than negative effect on the EPR, in contrast to what was
480 expected.

481

482 The source of the PUFAs in the mesocosms is a combination from the diatoms high in the fatty acid
483 20:5(n-3) (Viso and Marty 1993) and dinoflagellates that are high in C18 and 22:6(n-3) PUFAs
484 (e.g. Jónasdóttir 1994). *P. pouchetii* does on the other hand contain very low levels of PUFA
485 (Nichols et al. 1991), but may be rich in camper-, brassica- and sitosterols as they were highly
486 correlated with *P. pouchetii* in the present experiment (Table 3). PUFAs and sterols have been
487 shown to be essential for copepod egg production and growth (e.g. Jónasdóttir and Kiørboe 1996;
488 Hassett 2004; Klein Breteler, 2005), but do not seem to have been limiting the EPR in the present
489 study. The different protist types therefore, seem to have complemented each other to generate ideal
490 food condition for *C. finmarchicus* during the experiment.

491

492 The only difference in the reproduction between mesocosms observed in this study was in the
493 fraction of spawning females in mesocosms B and C (Fig 4b, Table 1). EPR in mesocosms B water
494 was also lower than in mesocosms C although the difference was not significant. We cannot explain

the difference in the spawning performance by the quality of the seston measured as polyunsaturated fatty acids normalized to carbon, but a significant positive correlation with low regression fit was found between seston cholesterol (carbon normalized) in mesocosms B and C and the fraction of spawning females (arcsine transformed, $R^2 = 0.17$, $p = 0.049$, $n = 18$). However, this relationship did not hold when all mesocosms were included in the analysis. The best relationship in spawning performance including all mesocosms and ambient water was with carbon content of the seston ($R^2 = 0.49$, $p < 0.001$, $n = 55$; data not shown). In any case, on the final day of the experiment there were no differences in the spawning performance of the females feeding on the different mesocosm water.

In this and most studies the PUA concentrations documented are the potential concentrations, i.e. the concentration that could be released from the algae, when wounded either when eaten or during senescence. The parameter dissolved PUA represents concentration of these metabolites in the water and thus the concentrations copepods and their eggs are exposed to. In addition to the PUA transferred through the diet of the female these dissolved PUA have the potential to directly affect hatching and naupliar survival. There are many studies that demonstrate inhibition of egg, embryo and naupliar development by direct exposure to dissolved PUA. Most of the studies that give PUA concentrations or enough information to calculate the PUA concentrations are summarised in Table 5. Most reports of detrimental effects of PUA are found with the commercially available aldehyde decadial, and inhibition of copepod egg hatching, cell division of embryos and starfish sperm motility were found at direct exposure at concentrations between 3,300-10,000 nmol L⁻¹ of dissolved decadial. Studies on the effects of post ingestion of PUA producing diatoms (maternal effects) have given more variable results; from no effects to poor hatching and/or naupliar survival (see Table 5). Therefore the question arises if the PUA concentration in our mesocosm experiment was just too low to cause failure in hatching and development. The potential PUA concentration (bound in phytoplankton) in this study was between 0 - 17 nmol L⁻¹ where the dissolved PUA was somewhat lower (0 - 4.9 nM). These concentrations are orders of magnitude lower than those used in the laboratory studies that demonstrate detrimental damage on eggs and nauplii (Table 5).

While there are remarkably few records of natural concentrations of PUA – particulate or dissolved, the few available studies show very similar concentrations as we measured in our mesocosms (0 - 28 nmol L⁻¹ particulate PUA and 0 - 0.1 nM dissolved PUA, see Table 5). Our particulate concentrations are therefore close to the highest reported concentrations in the Adriatic Sea (28 nmol L⁻¹) with no apparent effect on the eggs and egg production. The dissolved fraction in the mesocosms was however 50 times higher than the ones measured at the same station as the high

530 particulate PUA in the Adriatic (Table 5) but still with no measurable effect on the egg hatching.
531 The comparison summarised in Table 5 shows that most studies reporting detrimental effects of
532 dissolved aldehydes used concentrations 2 to 7 orders of magnitude higher than ever measured in
533 the natural environment and are therefore not useful, if not directly misleading for our
534 understanding of how the chemical and biological environment affects impact copepod
535 reproduction and growth. Those studies only demonstrate that the most active PUA, decadienal
536 does not have any effect on egg hatching and naupliar development at highest natural
537 concentrations and the natural dissolved concentrations can well increase over 1000 fold before
538 they directly affect egg and naupliar development. Indeed, Wichard et al. (2007) who determined
539 particulate PUA in the field did not find any correlation to the copepod egg production hatching or
540 naupliar abnormality. Similarly, in a laboratory study Dutz et al. (2008) demonstrated that PUA was
541 not the determining factor in hatching failure of the copepod eggs using potential concentrations 2-
542 100 times higher than the highest potential natural concentrations and ingestion rates of PUA of 0.1
543 nmol female⁻¹ d⁻¹. Therefore presence of potentially PUA producing diatoms does not seem to be
544 the exclusive factor controlling reproductive success of copepods in nature.

545

546 In summary, this study shows that *C. finmarchicus* egg production depends primarily on food
547 availability as a combination of ciliate, *P. pouchetii* and the availability of the essential PUFAs
548 representing range of diets of complimenting nutritional quality. Egg quality was high on the
549 mesocosm diets and was not inhibited by any means. This study further demonstrates that one of the
550 highest measured natural PUA concentrations both as potential PUA and dissolved PUA, associated
551 with *S. marinoi* blooms did not affect *C. finmarchicus* reproduction success.

552

553

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567

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 735

736 **Figure Legends:**

737 **Fig. 1** Daily development of a) total Chl *a* and b) Chl *a* in size fraction >10µm (µg L⁻¹) in
738 mesocosms: B (added phosphate and nitrate), C (added phosphate, nitrate and silicate), E (added
739 phosphate, nitrate, silicate and 400 *S. marinoi* L⁻¹) and F (added phosphate, nitrate, silicate and 400
740 *S. marinoi* L⁻¹) and the untreated water outside the mesocosms (Ambient). For detailed description
741 of the mesocosm treatments see Materials and Methods. Shaded area shows the days not used in the
742 analysis for EPR.

743

744 **Fig. 2** Daily development of A) protoplankton biomass (µg C L⁻¹) and B) % composition from day
745 4 in the ambient seawater and in mesocosm B, C, E and F. Black fill: the diatom *Skeletonema*
746 *marinoi*, hatched fill: ciliates, white fill: total heterotrophic dinoflagellates (sum of *Gyrodinium*
747 *spirale*, *G. dominans* and *Protoperdinium bipebs*) and stippled fill: the cryptophyte *Phaeocystis*
748 *pouchetii*

749

750 **Fig. 3** Daily development of total fatty acids, monounsaturated fatty acids (MUFA),
751 polyunsaturated fatty acids (PUFA), and total sterols, chole-, brassica-, sitosterols (all µg L⁻¹) in the
752 mesocosms B, C, E and F and ambient water. Symbols as in figure 1.

753

754 **Fig. 4** Daily development of *Calanus finmarchicus* a) egg production rate (EPR: eggs female⁻¹ day⁻¹)
755 ¹), b) fraction of females producing eggs, c) egg hatching success (%), and d) faecal pellet
756 production (FPR: µm³ female⁻¹ day⁻¹ x 10⁶) from mesocosms B, C, E and F and the ambient water
757 (from day 7). EPR shows additionally EPR from filtered seawater incubation (day 3-6).

758

759 **Fig. 5** *Calanus finmarchicus* egg production rate (eggs female⁻¹ day⁻¹) as a function of
760 protoplankton biomass a) *P. pouchetii*, b) ciliates, c) *S. marinoi*, and d) *G. dominans*, e) total carbon
761 concentration of protoplankton (all in µgC L⁻¹) and f) concentration of the fatty acid 22:6(n-3) (µg
762 L⁻¹). Symbols as in Figure 1.

763

764 **Fig. 6** Relation between the residuals of the regression presented in Figure 5e and the carbon based
765 proportion of a) *S. marinoi* and b) *P. pouchetii* and the concentrations of c) particulate PUA (µg L⁻¹)
766 and d) dissolved PUA (nM).

767

768 **Fig. 7** *Calanus finmarchicus* faecal pellet production (µm³ female⁻¹ day⁻¹ x 10⁶) as a function of
769 protoplankton biomass a) *P. pouchetii*, b) ciliates, c) *S. marinoi*, and d) *G. dominans*, e) total carbon

770 concentration of protoplankton (all in $\mu\text{gC L}^{-1}$) and f) concentration of the fatty acid 22:6(n-3) (μg
771 L^{-1}). Symbols as in Figure 1.

772
773 **Fig. 8** Correlation between *Calanus finmarchicus* egg production rate (eggs female⁻¹ day⁻¹) and
774 faecal pellet production (μm^3 female⁻¹ day⁻¹ x 10⁶). Symbols as in Fig 1.

775 **Table 1.** *Calanus finmarchicus* biological rates in the different experimental treatments. Average egg production rate (EPR: eggs female⁻¹
776 day⁻¹), cumulative egg production (eggs), non producing females (% Fem w/o eggs), female mortality (%), hatching success (H%), faecal
777 volume production (FPR: μm^3 female⁻¹ day⁻¹ x 10⁶) and faecal pellet production (# pellets: no. pellets female⁻¹ day⁻¹). FSW: 0.2 μm
778 filtered seawater control during days 3-6. Amb: ambient seawater control during days 7-14. All averages list \pm SE.
779

Mesocos m	Treatment additions				EPR ± SE Day 5-14	EPR range Day 5-14	Cum EPR Day 5-14	% Fem w/o eggs	% Mortality	H% ± SE Day 5-14	FPR x 10 ⁶ Day 3-14	# Pellets Day 3-14
	μmol L ⁻¹	cells L ⁻¹										
	N	P	Si	<i>S. marinoi</i>								
B	4.24	0.4	-	-	32 ± 6	0-173	4497	23	35	83 ± 2	25 ± 2	27 ± 2
C	4.24	0.4	3.61	-	46 ± 8	0-202	7959	15	20	88 ± 2	54 ± 4	43 ± 2
E	4.24	0.4	3.61	400	45 ± 7	0-232	5863	23	40	86 ± 3	45 ± 4	40 ± 3
F	4.24	0.4	3.61	1000	42 ± 8	0-167	8039	12	0	89 ± 3	43 ± 2	42 ± 2
FSW _{d 3-6}	-	-	-	-	6 ± 3	0- 38	-	47	-	-	-	9 ± 3
Amb _{d 7-14}	-	-	-	-	9 ± 5	0 - 90	-	50	20	89 ± 2	4 ± 0.5	11 ± 2

780

781

782

783 **Table 2.** Development of the major fatty acids in mesocosms B, C, E and F. The fatty acids 20:3(n-
784 3) and 20:3(n-6) were either not detected (n.d.) or found in trace amounts (T). nm: Not measured.
785

Day	16:1 (n-7)	18:1 (n-9)	18:2 (n-6)	18:3 (n-3)	18:3 (n-6)	20:1 (n-9)	20:2 (n-6)	20:4 (n-6)	20:5 (n-3)	22:1 (n-9)	22:2 (n-6)	22:6 (n-3)
µg (cells in 1L) ⁻¹												
Mesocosm B												
0	T	2.6	1.2	1.5	7.0	0.3	T	T	3.2	0.6	0.7	4.7
1	T	1.9	0.8	1.1	9.6	0.2	T	T	3.0	0.5	0.5	4.6
2	T	T	0.5	0.6	6.2	T	T	T	1.9	0.3	0.3	3.2
3	T	1.9	1.3	1.4	14.9	n.d.	n.d.	n.d.	2.3	0.3	T	4.4
4	9.8	4.0	2.9	4.9	51.3	0.3	n.d.	0.7	7.3	0.3	0.4	20.3
5	10.5	3.5	2.8	4.4	59.8	0.3	n.d.	1.3	12.4	0.3	0.4	32.4
6	9.0	5.2	3.1	4.2	59.6	0.3	n.d.	1.1	10.7	0.4	n.d.	56.3
7	8.4	6.0	3.3	3.6	44.9	0.4	n.d.	T	6.3	T	0.4	34.4
8	15.8	9.8	5.7	7.1	91.6	0.9	n.d.	2.5	15.5	T	0.9	81.1
9	n.d.	15.1	6.9	6.5	60.1	1.1	n.d.	3.0	11.2	T	1.0	55.6
10	T	23.4	15.3	11.8	132.8	1.9	n.d.	T	12.7	T	2.4	72.7
11	n.d.	19.7	13.0	9.4	83.4	1.8	n.d.	T	13.1	T	1.6	84.9
12	17.6	26.6	22.7	22.2	164.9	2.0	4.6	T	25.0	n.d.	1.7	196.4
13	17.3	25.7	21.2	14.7	124.3	1.9	4.4	T	14.1	2.2	2.1	129.4
Mesocosm C												
0	T	1.7	1.0	1.0	5.9	0.3	n.d.	n.d.	n.d.	0.7	0.7	3.6
1	T	1.7	0.8	1.3	14.3	0.3	n.d.	T	n.d.	0.8	0.7	6.5
2	T	1.3	0.5	0.7	8.5	0.3	n.d.	T	T	0.4	0.3	4.1
3	T	1.7	0.8	1.3	9.8	0.2	n.d.	T	1.8	0.3	0.2	5.6
4	11.3	3.5	1.8	3.6	40.9	0.2	T	0.8	7.7	0.3	0.6	27.6
5	15.9	4.3	2.8	5.4	57.5	0.2	T	1.2	8.7	0.3	0.6	44.2
6	21.6	5.4	3.5	7.4	77.2	0.3	T	1.1	14.8	0.5	1.1	59.5
7	21.6	5.7	2.6	5.2	68.6	0.5	T	1.6	13.3	0.6	1.0	50.2
8	16.5	7.4	3.9	5.2	51.0	0.9	T	2.6	13.7	1.2	1.1	47.5
9	15.6	8.0	3.7	4.1	50.1	0.8	T	T	14.0	1.2	1.0	37.6
10	9.1	9.8	n.d.	5.5	57.4	1.1	T	T	11.1	1.5	1.3	42.7
11	31.8	11.2	8.2	7.1	n.d.	1.2	n.d.	T	16.2	1.5	1.4	63.9
12	26.9	10.9	7.2	5.3	50.3	1.2	n.d.	T	9.1	1.5	1.3	43.9
13	92.4	22.5	23.2	15.6	165.4	2.1	4.6	5.5	27.5	2.3	2.6	226.2
Mesocosm E												
0	n.d.	T	0.6	0.9	4.9	n.d.	n.d.	n.d.	n.d.	n.d.	0.4	2.4
1	n.d.	T	n.d.	0.6	4.8	n.d.	n.d.	n.d.	T	n.d.	0.2	1.4
2	n.d.	n.d.	0.5	0.5	3.8	T	n.d.	n.d.	T	T	T	0.9
3	10.7	1.8	1.3	3.1	22.0	0.2	n.d.	0.6	6.3	0.3	0.3	16.8
4	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.	n.m.
5	76.0	n.d.	1.8	n.d.	95.7	n.d.	n.d.	n.d.	T	n.d.	n.d.	93.0
6	n.d.	n.d.	2.2	n.d.	134.8	n.d.	n.d.	n.d.	T	n.d.	n.d.	79.5
7	108.4	n.d.	2.4	n.d.	74.9	n.d.	n.d.	n.d.	T	n.d.	n.d.	57.1
8	159.5	T	2.9	7.6	64.2	n.d.	n.d.	n.d.	21.6	n.d.	n.d.	67.7
9	191.9	7.2	4.6	9.2	96.5	n.d.	n.d.	n.d.	22.2	n.d.	T	86.8
10	145.4	13.1	8.1	9.0	87.0	1.0	n.d.	3.4	36.1	1.1	1.1	80.4
11	67.9	12.5	7.1	8.5	53.5	1.0	n.d.	T	23.9	1.1	0.9	75.1
12	147.0	18.1	12.7	13.0	96.1	T	2.3	T	49.7	1.2	1.1	142.4
13	81.0	12.5	7.7	8.3	56.1	0.8	1.4	T	24.4	0.8	0.9	87.1
Mesocosm F												
0	T	1.8	0.6	0.9	5.4	T	n.d.	n.d.	n.d.	0.5	0.6	1.9
1	2.9	n.d.	0.5	0.5	4.5	0.2	n.d.	n.d.	T	0.3	0.4	2.5
2	3.4	1.2	n.d.	0.5	4.7	0.2	n.d.	T	T	0.3	0.3	2.3
3	9.9	1.4	0.7	1.3	10.0	0.2	n.d.	T	3.6	0.3	0.2	4.7
4	22.9	n.d.	0.6	0.8	16.3	n.d.	n.d.	n.d.	T	n.d.	T	10.8
5	108.3	2.9	1.8	5.8	82.8	n.d.	n.d.	1.3	28.7	n.d.	n.d.	80.2
6	171.2	3.7	2.4	7.8	92.9	n.d.	n.d.	T	52.6	n.d.	0.4	98.9
7	253.9	7.9	7.3	16.4	188.9	n.d.	n.d.	2.4	71.6	n.d.	T	140.6
8	223.2	9.0	7.0	15.7	110.2	T	T	3.8	57.1	n.d.	n.d.	135.2
9	284.1	5.5	n.d.	8.7	95.6	n.d.	n.d.	n.d.	41.0	n.d.	n.d.	106.8
10	105.6	7.3	3.6	8.1	74.5	T	n.d.	T	38.1	n.d.	n.d.	111.4
11	123.2	8.8	4.5	8.3	59.6	T	n.d.	T	42.4	0.6	0.9	104.4
12	183.2	14.0	4.1	11.8	79.6	n.d.	n.d.	n.d.	49.7	n.d.	n.d.	356.3
13	120.3	8.8	3.6	7.4	39.6	0.4	n.d.	n.d.	39.0	n.d.	T	136.4

786

787 **Table: 3**

788 Results from the data reduction by the Principal Component Analysis. The principal components
 789 and their % contribution to explaining the variance in the whole dataset (combined = 87 %). The
 790 scores show the rotated components acquired by using the Equamax method with Kaiser
 791 Normalization in 10 iterations. Bold letters show highly correlated variables on each component,
 792 bold italic variables that are moderately correlated on that variable. MUFA: mono unsaturated fatty
 793 acids.

794
 795

	Principal Component			
	1	2	3	4
	54 %	17 %	10 %	6 %
Octadienal	0.92		0.17	0.14
Heptadienal	0.92		0.17	0.17
<i>Skeletonema marinoi</i>	0.85			0.39
Decadienal	0.81	0.30	0.31	0.13
Chlorophyll <i>a</i> > 10 µm	0.71	0.54		0.26
MUFA	0.62		0.57	0.48
Cholesterol	0.64	0.36	0.20	0.57
<i>Phaeocystis pouchetii</i>	0.17	0.91	0.17	
Brassicasterol		0.87		0.38
Campesterol		0.80	0.35	
Sitosterol		0.80	0.23	0.40
Cilates	0.31	0.17	0.86	
<i>Gyrodinium spirale</i>		0.24	0.84	0.37
<i>Protoperidinium bipes</i>		0.28	0.81	0.33
<i>Gyrodinium dominans</i>	0.47	0.23	0.61	0.34
Docosahexaenoic acid; 22:6(n-3)		0.14	0.49	0.79
Linoleic acid; 18:3(n-3)	0.20	0.41	0.33	0.77
Gamma Linolenic acid 18:3(n-6)	0.36	0.42	0.17	0.73
Eicosapentaenoic acid; 20:5(n-3)	0.42		0.51	0.66

796
 797

Table 4

Results from multiple stepwise regression on *Calanus finmarchicus* egg production rates (EPR: eggs female⁻¹ d⁻¹) and faecal pellet production (FPR: μm³ female⁻¹ d⁻¹) and the food environment, *S. marinoi*, *G. dominans*, *P. pouchetii*, ciliates, the fatty acids 22:6(n-3), 20:5(n-3) and cholesterol (or Chl *a* in the size fraction 5-10 μm). The stepwise process selects only those variables (labelled “in”) that contribute to the best regression. Variables not in the regression are labelled “out”. Model statistics show R²: coefficient of determination of the multiple regression, *F*_{df}: the F ratio with *df*: degrees of freedom, significance value *p* of the multiple regression and number of observations *N* in the regression. For each variable in the model, "F-to-remove" is the F statistic for its coefficient within the regression; for each variable not in the model, "F-to-enter" is the F statistic that its coefficient would have if it were the next variable added in the regression. *p*_{var}: significance of the variable within the regression where *: *p* < 0.05, **: *p* < 0.01, ***: *p* < 0.001.

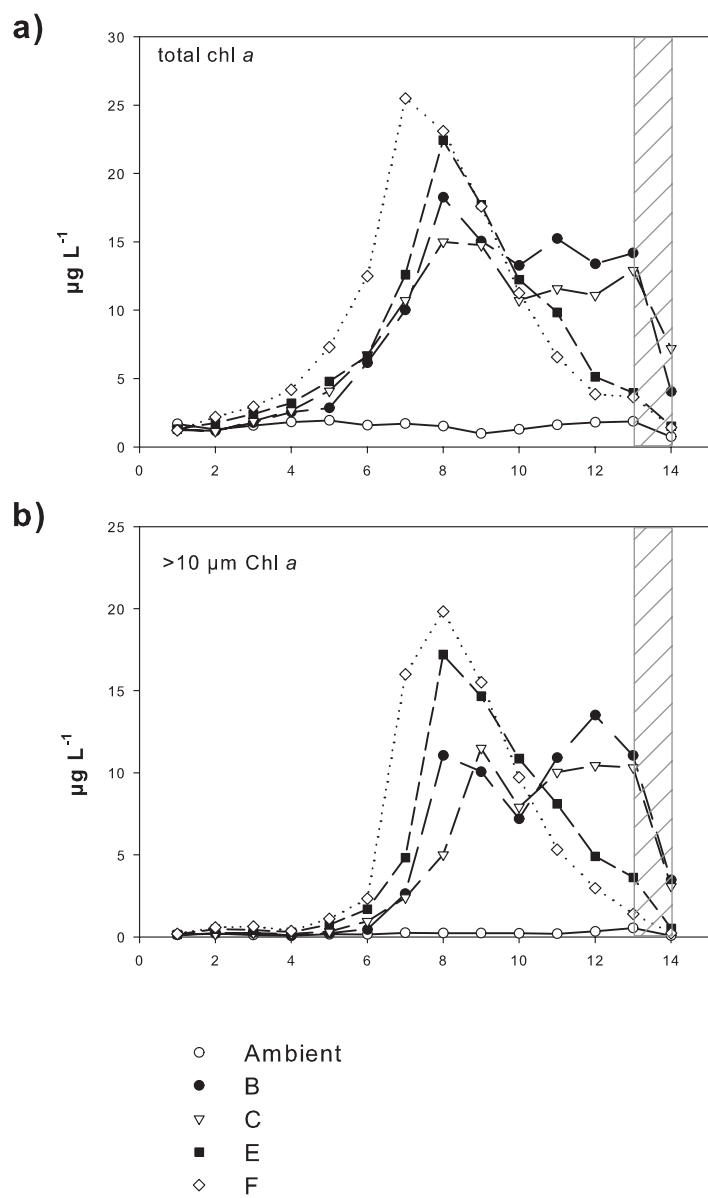
	Variable		F-to-remove	F-to-enter	p var
EPR	<i>Skeletonema marinoi</i>	in	6.3		0.016*
	ciliates	in	15.0		<0.001***
R ² = 0.81	<i>Gyrodinium dominans</i>	out		1.1	0.292
F ₄ = 41.7	<i>Phaeocystis pouchetii</i>	in	20.6		<0.001***
p < 0.001	22:6(n-3)	in	27.4		0.001***
N = 45	Cholesterol	out		0.04	0.814
	20:5(n-3)	out		0.04	0.961
FPR	<i>Skeletonema marinoi</i>	out		1.1	0.610
	ciliates	in	8.4		<0.006**
R ² = 0.78	<i>Gyrodinium dominans</i>	in	4.5		0.038*
F ₃ = 43.3	<i>Phaeocystis pouchetii</i>	in	43.6		<0.001***
p < 0.001	22:6(n-3)	in	10.9		0.002**
N = 54	Chlorophyll <i>a</i> 5-10μm	out		2.7	0.107
	20:5(n-3)	out		0.9	0.339

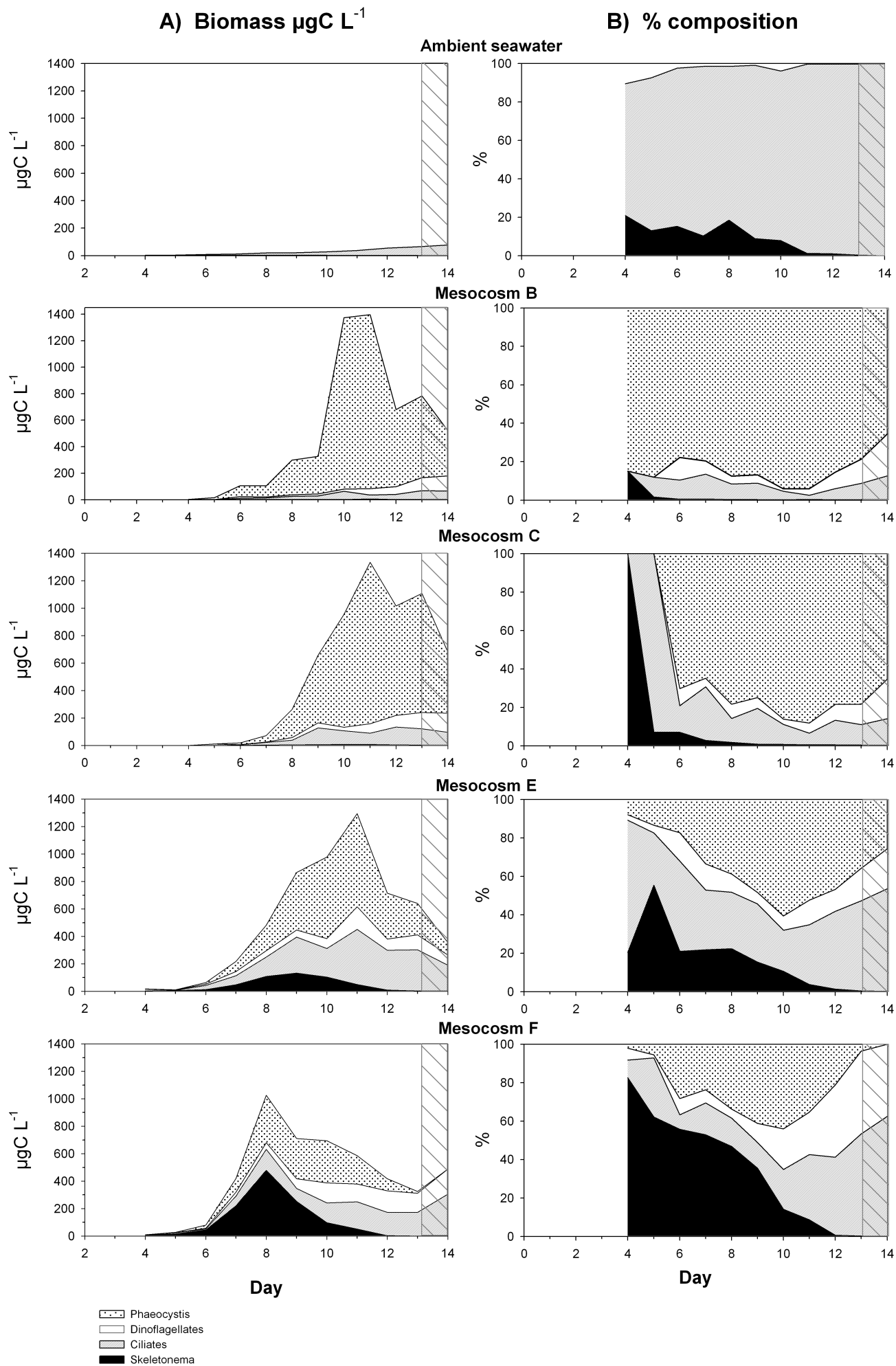
Table 5.

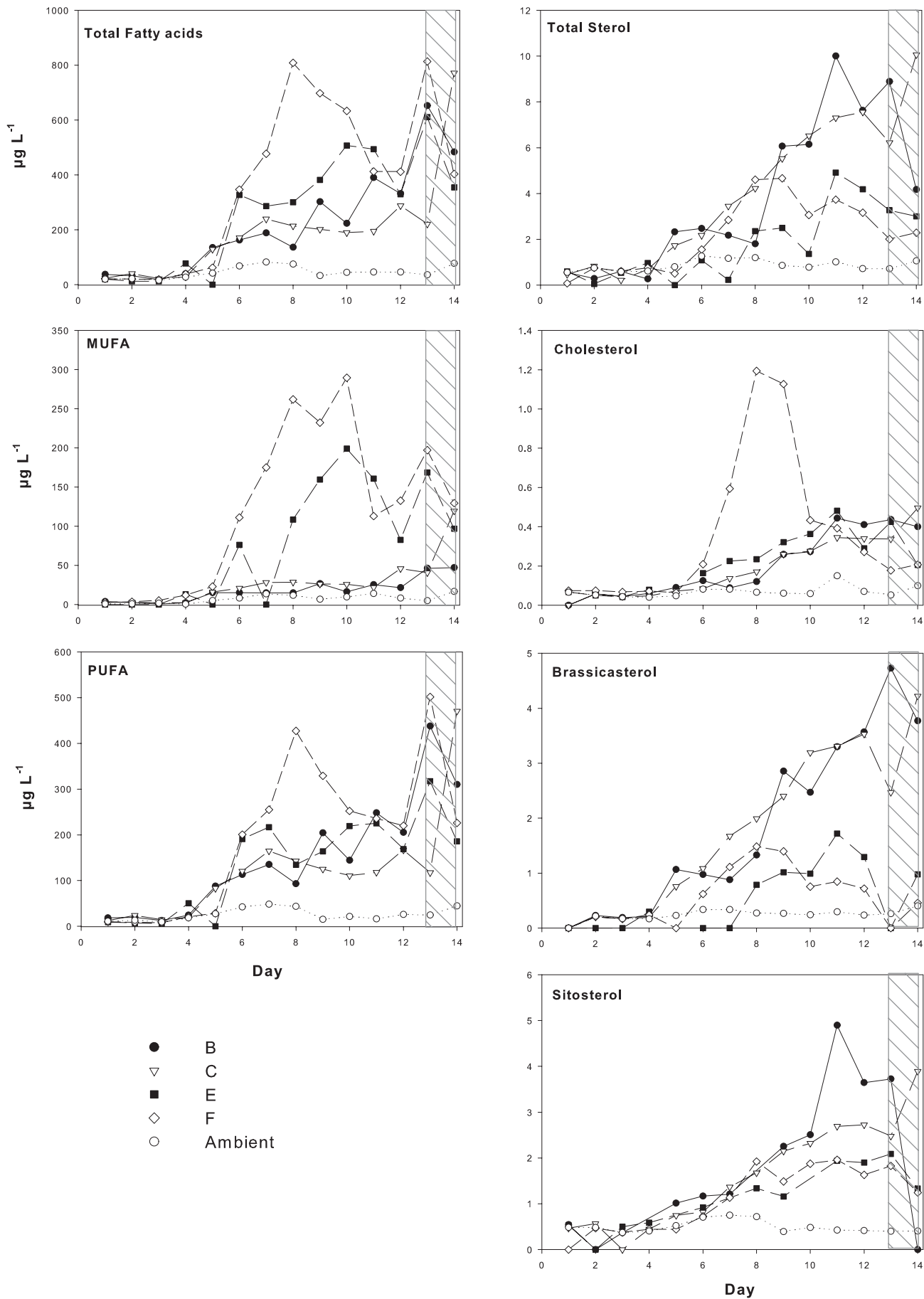
Review of concentrations used for studies testing polyunsaturated aldehydes on various biological activities of copepods, cladoceran and echinoderms. All units transferred to nmol L⁻¹ (shaded column). Units reported in the respective publications are shown in bold; otherwise calculated from cell numbers and PUA content given in the respective publications. The average molecular weight of 150 g mol⁻¹ was used as representative for the most commonly tested decadi- and decatrienal aldehydes. Other aldehyde types have lower molecular weight and therefore transfer to nmol is conservative. Aldehyde types are: HD: Heptadienal, OD: Octadienal, OT: Octatrienal, DE: Decenal, DD: Decadienal, DT: Decatrienal, PUA: Polyunsaturated aldehydes

Testing concentrations		Aldehyde types	Main effects	Source
$\mu\text{g mL}^{-1}$	nmol L ⁻¹			
0.5 - 2.0	3,300 – 13,300	DD DT-cis DT-trans	Hatching of <i>Temora longicornis</i> eggs reduced 10-15% at direct exposure of 3,300 nmol L ⁻¹ DD, 40-60% at 6,700 nmol L ⁻¹ and failed totally at 10,000 – 13,000 nmol L ⁻¹ . Sea urchin egg division failure at >3,300 nmol L ⁻¹ DD.	Miralto et al. 1999
0.6-600	4,000-4,000,000	DD OD DD	Cell division of echinoderm embryos first affected at direct exposure of 10,000 nmol L ⁻¹ DD (50% division). Post ingestion effect on <i>C. helgolandicus</i> eggs lowest pot. PUA 4,000 nmol L ⁻¹ with 20% reduction in hatching %. Remained 80% hatching till 400,000 nmol L ⁻¹ and had 70% reduction at 4 million nmol L ⁻¹ . NB: Eggs also exposed to increasing concentration of algae – can affect hatching (O ₂).	Adolph et al. 2004
0.05 - 50	333 - 333,000	DD	Motility of starfish sperm decreased with increased direct exposure to DD. 50% decrease in motility at 10,000 nmol L ⁻¹ .	Caldwell et al. 2004
0.5 - 1.5	3,300 – 10,000	DD	Post ingestion effect of the diatom <i>Skeletonema. C. helgolandicus</i> nauplii survive to CII after 3 & 5 days and to NII after 7 days of maternal and neonatal feeding. Ingestion ca 100 pg DD female ⁻¹ d ⁻¹ = 0.0007 [sic. 0.005] nmol female ⁻¹ d ⁻¹ . Addition of dissolved DD (direct exposure) 10,000 nmol L ⁻¹ to diets resulted in arrested development at CII while 3,300 and 6,700 nmol DD L ⁻¹ addition did not affect development.	Ianora et al. 2004
0.5 - 4.0	3,300 – 27,000	DD	Post ingestion effects of pure diatom diets on <i>Daphnia</i> embryos. First effect of 50% hatching after 5 clutches. Direct exposure of developing embryos to DD first affected at >10,000 nmol L ⁻¹ .	Carotenuto et al. 2005
1.4 – 15.0	9,300 – 100,000	DD OD HD	No effect on egg production of <i>Tisbe</i> or hatching after 3 day feeding on pure diatom diets with PUA potential. Direct exposure of nauplii to DD showed lethal dose (LD ₅₀ = 50% death) 9,300 nmol L ⁻¹ . Other PUA types less active. Nauplii most sensitive to direct exposure of DD, females LD ₅₀ > 40,000 nmol L ⁻¹ DD.	Taylor et al. 2007
0.1 - 0.6	53 - 638	OD OT DT	Post ingestion effect of PUA from var. diatom diets tested on copepod <i>T. longicornis</i> . Ingestion of 0.1 nmol PUA female ⁻¹ d ⁻¹ (should be 100pmol d ⁻¹ not 100 nmol d ⁻¹) did not negatively affect egg production or hatching success (10 days). PUA sum of OD, OT and DT.	Dutz et al. 2008
0.002	11.2	HD	Competition of fatty acids in the copepod gut (<i>T. longicornis</i>) when enzymes transform PUFA	Wichard et al. 2007

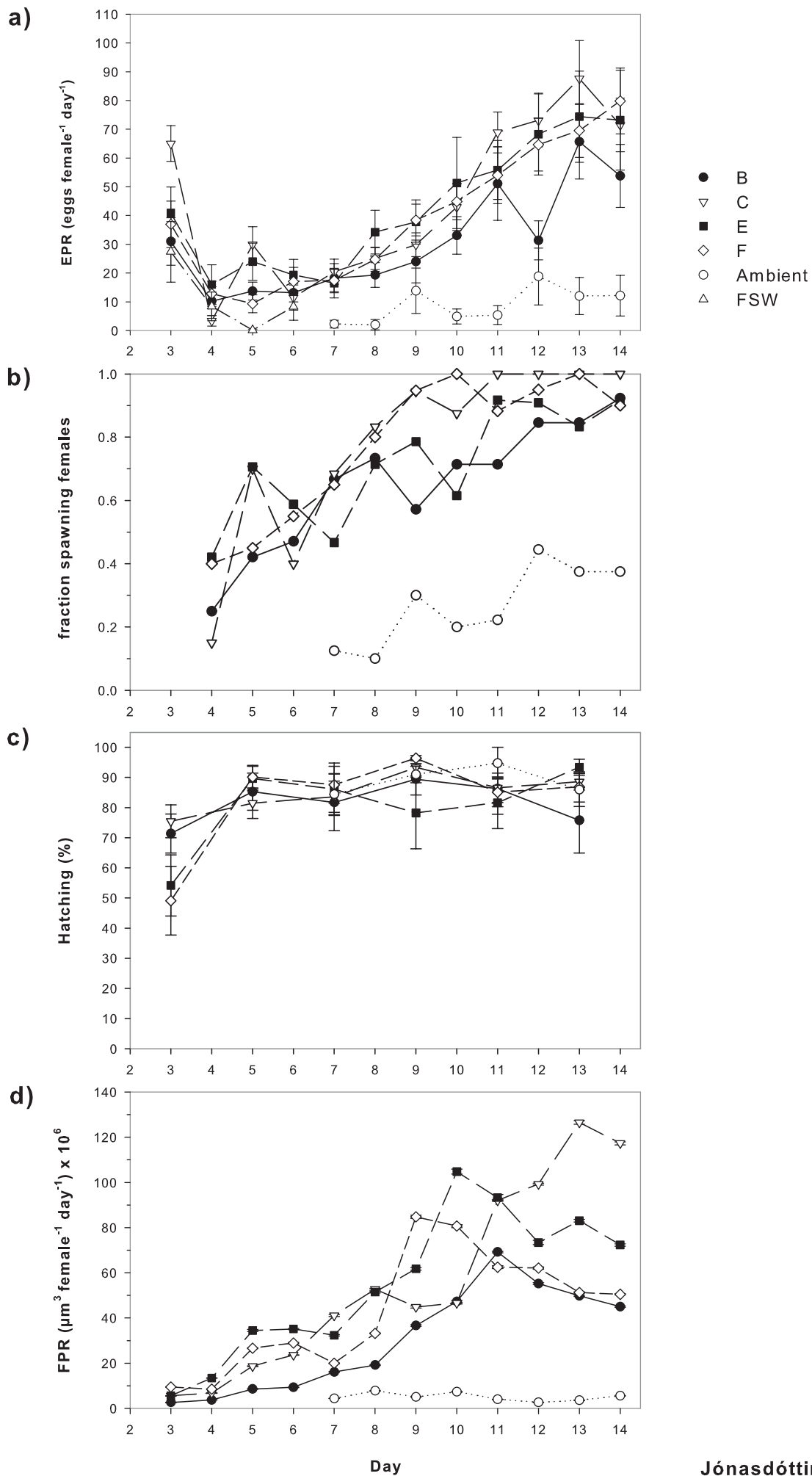
		OD OT DD DT	to PUA. Supplements of fatty acids improve hatching which is actually not very impaired by presence of high concentration of PUA.	
Measurements <i>in situ</i>				
1×10^{-4}	0.7	HD OD OT DD DT	English Channel natural diatom population during spring. Very high PUA per cell (47 fm cell ⁻¹) mainly DT and OD. Values are potential PUA.	Wichard et al. 2005
$2 \times 10^{-6} - 1 \times 10^{-4}$	0.01 - 1.3	HD OD OT	Seasonal PUA concentrations from Gullmarsfjorden, Sweden. Diatoms isolated and tested in the laboratory. Values are potential PUA. Higher PUA in senescing algae. Highest in senescing spring population.	Taylor et al. 2009
4×10^{-4}	2.7	HD OD OT	Natural concentration (sum potential PUA) during spring bloom offshore Dichanto, Chile. Variation in egg production, low hatching and high abnormal nauplii could not be explained with natural PUA or doubling of natural PUA concentrations.	Poulet et al. 2007
$1 \times 10^{-5} - 3 \times 10^{-5}$ 0.002 - 0.2	0.1 - 0.3 (<i>in situ</i>) 12-143 (exp)	OT OD DT HT	Natural concentration during natural diatom assembly in the English Channel and in feeding experiments using uni-algal isolates. Sum of potential PUA. Hatching and egg production of <i>C. helgolandicus</i> positively correlated with diatoms & PUA. Experiments with high PUA producers resulted in no effect of PUA on hatching or larval abnormality.	Wichard et al. 2008
$2 \times 10^{-5} - 0.004$	0.1 - 28 (part) 0 - 0.1 (dissolved)	HD OD	Natural concentrations in the Adriatic. Suggests that lysis of <i>Skeletonema marinoi</i> can contribute significantly to dissolved PUA. One station with the high concentration of both potential and dissolved PUA.	Vidoudez et al 2011a
0 - 0.003 0 - 7×10^{-4}	0 - 17 (part) 0 - 4.9 (dissolved)	HD OT OD DD DT	Range of the maximum values measured in mesocosms B, C, E and F. Sum of all measured dienal and trienal PUAs. Dominant PUA (peak concentration measured), HD (7 nmol L ⁻¹) and DD (9 nmol L ⁻¹) in mesocosms E& F.	This study Vidoudez et al. 2011b



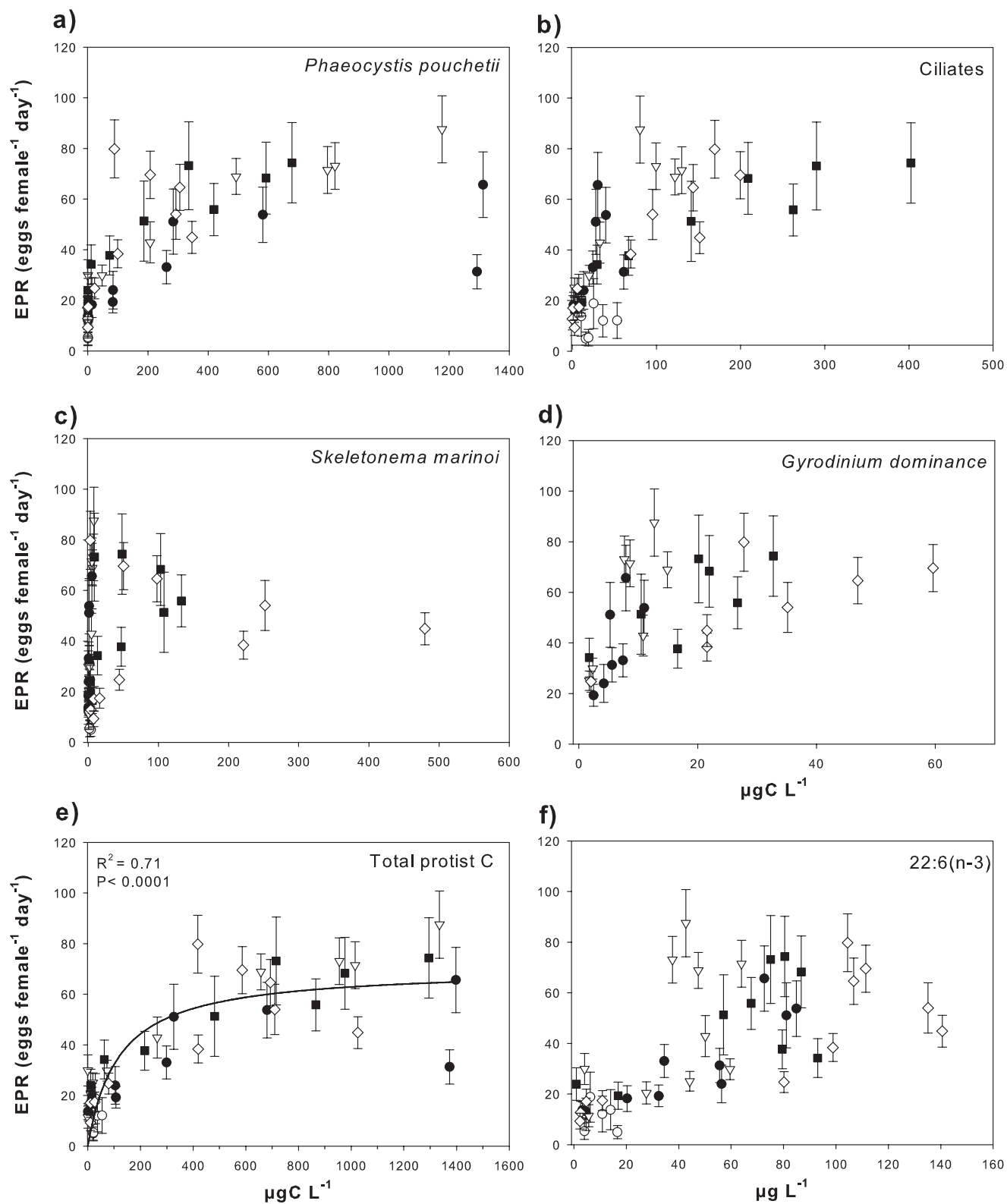




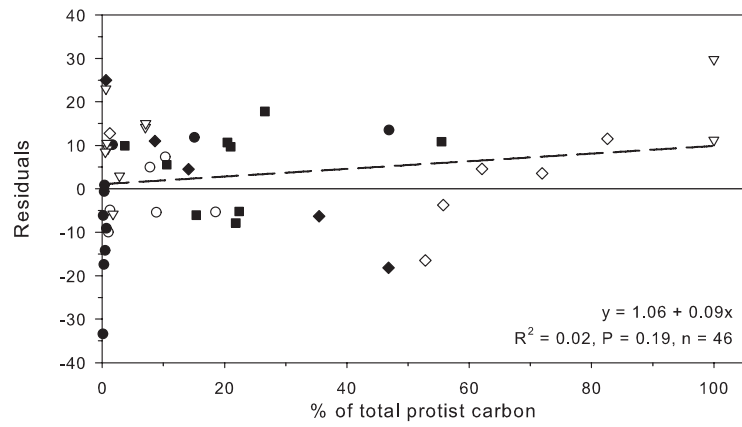
Jónasdóttir et al. Figure 3



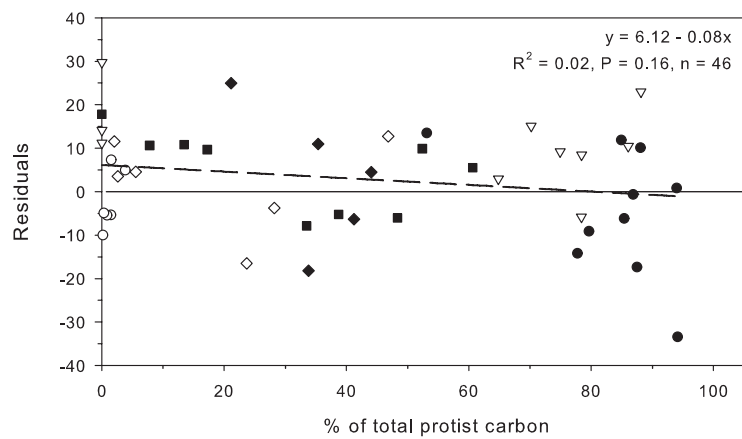
Jónasdóttir et al. Figure 4



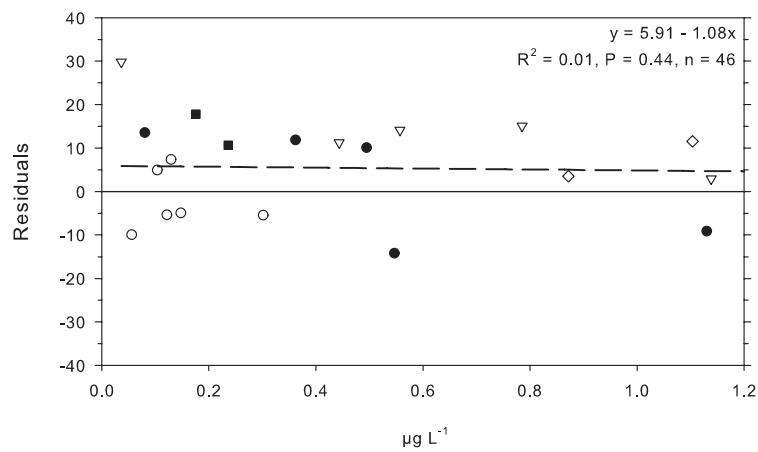
A) *Skeletonema marinoi*



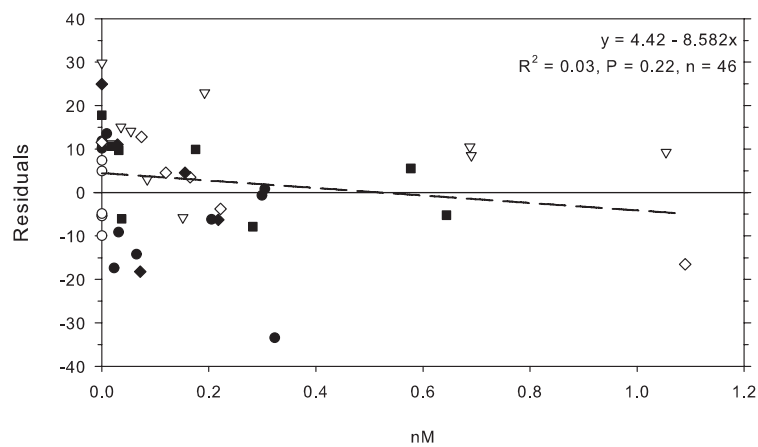
B) *Phaeocystis pouchetii*

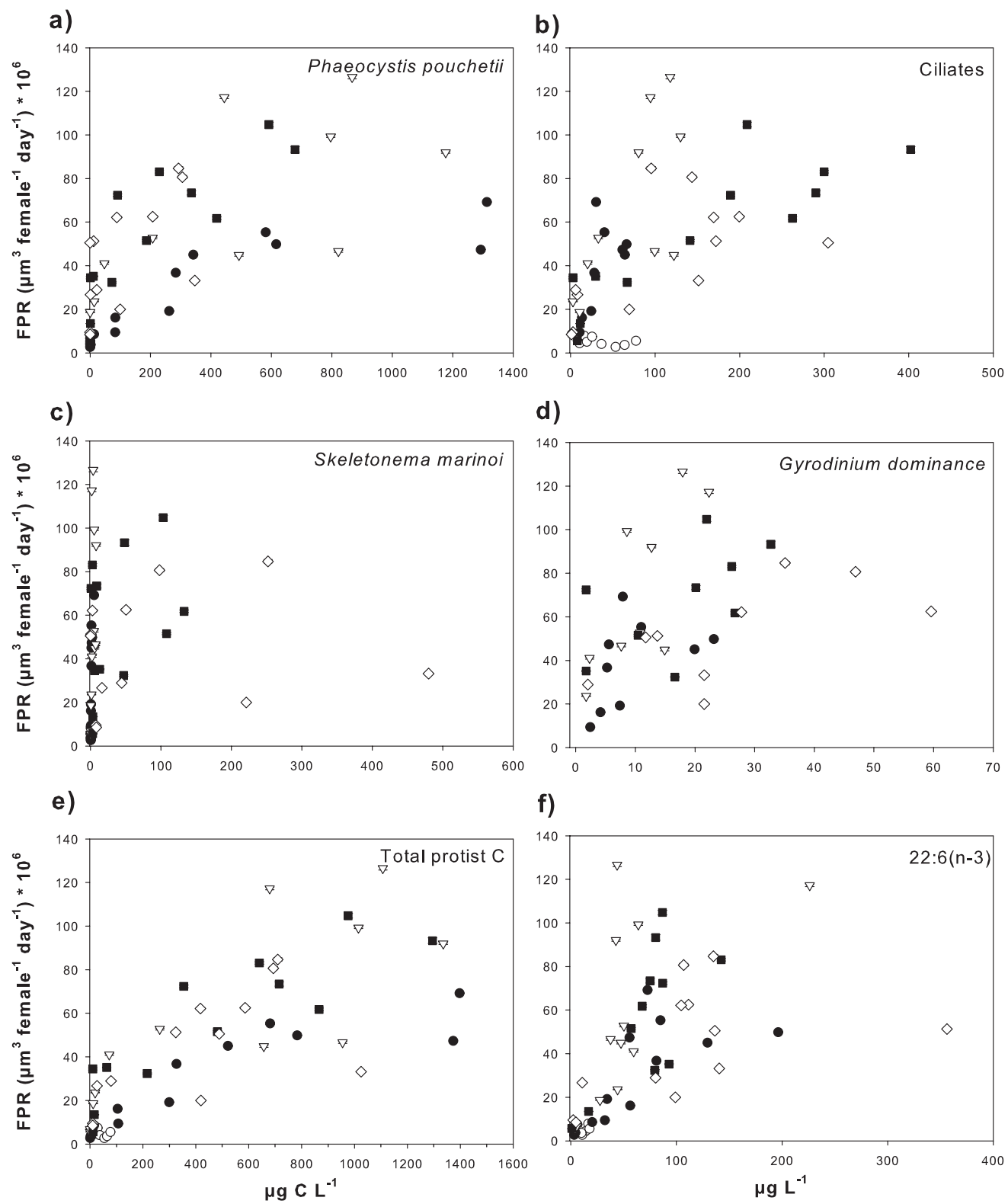


C) Particulate PUA



D) Dissolved PUA





Jónasdóttir et al Figure 7

