



1 **The Arctic picoeukaryote *Micromonas pusilla* benefits**
2 **synergistically from warming and ocean acidification**

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15 **Abstract**

16 In the Arctic Ocean, climate change effects such as warming and ocean acidification (OA) are
17 manifesting faster than in other regions. Yet, we are lacking a mechanistic understanding of the
18 interactive effects of these drivers on Arctic primary producers. In the current study, one of the
19 most abundant species of the Arctic Ocean, the prasinophyte *Micromonas pusilla*, was exposed
20 to a range of different pCO₂ levels at two temperatures representing realistic scenarios for
21 current and future conditions. We observed that warming and OA synergistically increased
22 growth rates at intermediate to high pCO₂ levels. Furthermore, elevated temperatures shifted
23 the pCO₂-optimum of biomass production to higher levels. Based on changes in cellular
24 composition and photophysiology, we hypothesise that the observed synergies can be explained
25 by beneficial effects of warming on carbon fixation in combination with facilitated carbon
26 acquisition under OA. Our findings help to understand the higher abundances of picoeukaryotes
27 such as *M. pusilla* under OA, as has been observed in many mesocosm studies.



28 **1 Introduction**

29 With the progress in using molecular tools to describe marine biodiversity in the past decades,
30 the scientific community has become increasingly aware of the underestimated importance of
31 picoeukaryotes, for both primary and export production of the world's oceans (Richardson and
32 Jackson, 2007; Worden and Not, 2008). Larger phytoplankton such as diatoms are efficient
33 vectors for carbon export due to aggregate formation and ingestion by large zooplankton
34 leading to the production of fast-settling faecal pellets (Sherr et al., 2003). In contrast,
35 picoeukaryotes are mainly grazed by smaller heterotrophic protists such as ciliates, which have
36 a low carbon retention, excrete relatively more dissolved material, and thus fuel recycled
37 production (Sherr and Sherr, 2002). Hence, changes in the relative abundance of pico- and
38 nanoeukaryotes can have large implications for food webs and biogeochemistry (Worden et al.,
39 2015).

40 Picoeukaryotes tend to dominate low nutrient environments, which is often attributed to
41 their high surface:volume ratios and mixotrophic capacities (Raven, 1998; McKie-Krisberg and
42 Sanders, 2014). The low nutrient concentrations in the Arctic surface ocean, for example, cause
43 picoeukaryotes to be particularly successful in this region. In fact, the globally occurring
44 prasinophyte *Micromonas pusilla* is considered the most abundant species in the Arctic ocean
45 (Šlapeta et al., 2006; Lovejoy et al., 2007; Marquardt et al., 2016). In this environment, strong
46 stratification causes nutrient limitation throughout the summer and autumn months (Tremblay
47 et al., 2015), and the occurrence of the polar night requires organisms to either form resting
48 stages or to have heterotrophic capacities (Tremblay et al., 2009; Lovejoy, 2014; Berge et al.,
49 2015; Vader et al., 2015).

50 Climate change effects manifest faster in the Arctic than anywhere else on the planet
51 (Stocker, 2014). In this region, temperatures rise more than twice as fast as the rest of the globe
52 (Miller et al., 2010). The concurrent rapid reduction in ice cover allows for more light
53 penetration and longer growing seasons, while increased stratification due to ice melt and
54 warming constrain nutrient supply to surface waters, both of which will change the dynamics
55 of primary production (Arrigo et al., 2008; Wassmann and Reigstad, 2011). Ocean acidification
56 (OA) is also especially pronounced in the Arctic Ocean, because low temperatures and
57 alkalinity make the system sensitive to anthropogenic CO₂ loading (AMAP, 2013; Qi et al.,
58 2017). Picoeukaryotes such as *M. pusilla* may benefit from these changes and are considered
59 potential winners of climate change. In the Canadian Arctic, for example, picoeukaryote
60 abundances are increasing as surface waters get warmer, fresher and more oligotrophic (Li et
61 al., 2009). Regarding OA effects, the majority of studies on natural phytoplankton assemblages



62 have shown picoeukaryotes, particularly *M. pusilla*, to increase in relative abundance with
63 increasing pCO₂ levels (Engel et al., 2008; Meakin and Wyman, 2011; Newbold et al., 2012;
64 Brussaard et al., 2013; Schulz et al., 2017). Despite the evident sensitivity of *M. pusilla* to
65 changes in pCO₂ levels, a detailed assessment of the OA effects, their interaction with warming
66 as well as the underlying mechanisms in this important species is still missing.

67 Like all photosynthetic organisms, cells of *M. pusilla* need to maintain a balance
68 between energy sources (i.e. light harvesting by the photosynthetic apparatus) and sinks (most
69 importantly carbon fixation in the Calvin cycle) to prevent harmful levels of excitation pressure
70 on the photosynthetic electron transport chain (Behrenfeld et al., 2008). Light harvesting and
71 electron transport in the photosystems are largely independent of changes in temperature and
72 pCO₂ (Mock and Hoch, 2005; Hoppe et al., 2015), but the impact of these drivers on energy
73 sinks can potentially affect the energy balance of the cell: The beneficial effects of elevated
74 pCO₂ observed in phytoplankton are thought to be caused by increased diffusive CO₂ supply,
75 reduced CO₂ leakage, or by lowered costs to operate their CO₂ concentrating mechanisms (Rost
76 et al., 2008; Bach et al., 2013). Elevated temperatures, on the other hand, can change enzyme
77 kinetics including those involved in the Calvin cycle, thus leading to a larger sink of excitation
78 energy (Maxwell et al., 1994; Toseland et al., 2013). Hence, both ocean warming and
79 acidification potentially increase the efficiency of photosynthesis and biomass production, at
80 least up to the organisms' respective optimum levels. Above these levels, temperatures and
81 proton concentrations start to disrupt enzymatic processes, increase the need for pH
82 homeostasis, and impair the delicate regulation of cellular processes (Levitt, 1980; Taylor et
83 al., 2001; Flynn et al., 2012). Thus, the complex balance between beneficial and detrimental
84 effects will determine whether the combination of warming and OA will synergistically
85 promote or deteriorate phytoplankton growth and biomass build-up.

86 In the current study, we aim to investigate the responses of an Arctic *M. pusilla* strain
87 to warming and OA. To this end, *M. pusilla* was grown at four pCO₂ levels ranging from
88 preindustrial to future scenarios (180-1400 µatm) under 2°C and 6°C, which represent the
89 magnitude of the projected future temperature increase in this region (Collins et al., 2013), but
90 also the current spring and summer temperatures in the environment where the strain was
91 isolated (Hegseth et al., in press).



92 **2 Material & Methods**

93

94 **2.1 Culture conditions**

95 Monoclonal cultures of the picoeukaryote *Micromonas pusilla* (Butcher) I. Manton & M. Parke
96 (isolated in 2014 by K. Wolf in Kongsfjorden, Svalbard, 79°N; taxonomic identification
97 confirmed by rDNA sequencing of SSU, LSU and ITS sequences) were grown in 1-L glass
98 bottles in semi-continuous dilute-batch cultures (max 129,000 cells mL⁻¹; diluted every 3-4
99 days) under constant irradiances of $150 \pm 26 \mu\text{mol photons m}^{-2} \text{s}^{-1}$. Media consisted of 0.2 μm
100 sterile-filtered Arctic seawater with a salinity of 32.7 enriched with macronutrients, trace metals
101 and vitamins according to F/2_R medium (Guillard and Ryther, 1962). Light intensities were
102 provided by daylight lamps (Philips Master TL-D 18W; emission peaks at wavelength of 440,
103 560 and 635 nm), adjusted by neutral density screens and monitored using a LI-1400 data logger
104 (Li-Cor) equipped with a 4 π -sensor (Walz). Cells were growing at four different CO₂ partial
105 pressures (pCO₂; 180, 380, 1000, and 1400 μatm) and two temperatures ($2.2 \pm 0.3^\circ\text{C}$ and $6.3 \pm$
106 0.2°C). Cultures were acclimated to these conditions for at least 7 generations prior to sampling.

107 Different pCO₂ conditions were achieved by aeration of the incubation bottles with air
108 of the respective pCO₂ levels delivered through sterile 0.2- μm air-filters (Midisart 2000,
109 Sartorius stedim) for 24 h prior to inoculation. Gas mixtures were generated using a gas flow
110 controller (CGM 2000 MCZ Umwelttechnik), in which CO₂-free air (<1 ppmv CO₂; Dominick
111 Hunter) was mixed with pure CO₂ (Air Liquide Deutschland). The pCO₂ levels in the gas
112 mixtures were regularly monitored with a non-dispersive infrared analyzer system (LI6252, LI-
113 COR Biosciences), calibrated with CO₂-free air and purchased gas mixtures of 150 ± 10 and
114 1000 ± 20 ppmv CO₂ (Air Liquide Deutschland).

115

116 **2.2 Carbonate chemistry**

117 Samples for total alkalinity (A_T) were filtered through 0.7- μm glass fibre filters (GF/F,
118 Whatman) and stored in borosilicate bottles at 3°C. A_T was estimated from duplicate
119 potentiometric titration (Brewer et al., 1986) using a TitroLine alpha plus (Schott Instruments).
120 A_T values were corrected for systematic errors based on measurements of certified reference
121 materials (CRMs provided by Prof. A. Dickson, Scripps, USA; batch #111; reproducibility ± 5
122 $\mu\text{mol kg}^{-1}$). Total dissolved inorganic carbon (C_T) samples were filtered through 0.2- μm
123 cellulose-acetate filters (Sartorius stedim) and stored in gas-tight borosilicate bottles at 3°C. C_T
124 was measured colorimetrically in triplicates with a QuAAtro autoanalyzer (Seal; Stoll et al.
125 2001). The analyser was calibrated with NaHCO₃ solutions (with a salinity of 35, achieved by



126 addition of NaCl) to achieve concentrations ranging from 1800 to 2300 $\mu\text{mol C}_T \text{ kg}^{-1}$. CRMs
127 were used for corrections of errors in instrument performance such as baseline drifts
128 (reproducibility $\pm 8 \mu\text{mol kg}^{-1}$). Seawater pH_{total} was measured potentiometrically with a two-
129 point calibrated glass reference electrode (IOline, Schott Instruments). An internal TRIS-based
130 reference standard (Dickson et al., 2007) was used to correct for variability on electrode
131 performance (reproducibility ± 0.015 pH units). Following recommendations by Hoppe et al.
132 (2012), seawater carbonate chemistry including pCO_2 was calculated from A_T and pH using
133 $\text{CO}_{2\text{SYS}}$ (Pierrot et al., 2006). The dissociation constants of carbonic acid of Mehrbach et al.
134 (1973), as refitted by Dickson and Millero (1987), were used for calculations. Dissociation
135 constants for KHSO_4 were taken from Dickson (1990).

136

137 **2.3 Growth, elemental composition and production rates**

138 Samples for cell counts were fixed with glutaraldehyde (0.5% final concentration). After gentle
139 mixing, samples were stored at room temperature in the dark for 15 min, and subsequently
140 frozen in liquid nitrogen and stored at -80°C . Prior to analysis, samples were thawed on ice and
141 mixed thoroughly. After addition of 10 μL SybrGreen working solution (dissolved in DMSO)
142 and 10 μL YG beads working solution (1 μm -Flouresbrite calibration beads grade YG,
143 Polyscience), samples were counted on an Accuri C6 flow cytometer (BD Biosciences)
144 equipped with a blue solid-state laser (488 nm excitation wavelength) run on medium fluidics
145 settings (35 $\mu\text{L min}^{-1}$; 16 μm core size) with a limit of 50,000 events or 250 μL . Analysis was
146 performed based on red (FL3 channel, >670 nm) and green (FL1 channel, 533 ± 30 nm)
147 fluorescence, as well as sideward and forward light scattering. Specific growth rates constants
148 (μ) were determined from exponential fits of cell counts over 4 consecutive days.

149 Particulate organic carbon (POC) and nitrogen (PON) were measured after filtration
150 onto precombusted (15h, 500°C) GF/F filters (Whatman). Filters were stored at -20°C and
151 dried for at least 12 h at 60°C prior to sample preparation. Analysis was performed using a
152 CHNS-O elemental analyser (Euro EA 3000, HEKAtech). Contents of POC and PON were
153 corrected for blank measurements and normalised to filtered volume and cell densities to yield
154 cellular quotas. Production rates of POC were calculated by multiplying the cellular quota with
155 the specific growth rate constant of the respective incubation. Samples for determination of
156 chlorophyll *a* (Chl *a*) were filtered onto GF/F filters (Whatman), immediately placed into liquid
157 nitrogen and stored at -80°C until analysis. Chl *a* was subsequently extracted in 8 mL 90%
158 acetone at 4°C over night. Chl *a* concentrations were determined on a fluorometer (TD-700,



159 Turner Designs), using an acidification step (1M HCl) to determine phaeopigments (Knap et
160 al., 1996).

161

162 **2.4 Variable Chl fluorescence**

163 Photophysiological characteristics, based on photosystem II (PSII) variable Chl fluorescence,
164 were measured using a fast repetition rate fluorometer (FRRf; FastOcean PTX, Chelsea
165 Technologies) in combination with a FastAct Laboratory system (Chelsea Technologies). The
166 excitation wavelength of the fluorometer's light-emitting diodes (LEDs) was 450 nm, and the
167 applied light intensity was 1.3×10^{22} photons $m^{-2} s^{-1}$. The FRRf was used in single turnover
168 mode, with a saturation phase comprising 100 flashlets on a 2 μs pitch and a relaxation phase
169 comprising 40 flashlets on a 50 μs pitch. Measurements from all replicates ($n=3$) were
170 conducted in a temperature-controlled chamber ($\pm 0.2^\circ C$) at the respective treatment
171 temperature.

172 After subtraction of a blank value, the minimum (F_0 and F_0' for light- and dark-
173 acclimated measurements, respectively) and maximum Chl fluorescence (F_m and F_m' for light-
174 and dark-acclimated measurements, respectively) were estimated from iterative algorithms for
175 induction (Kolber et al., 1998) and relaxation phase (Oxborough, 2012) after 15 min of dark
176 acclimation, which was sufficient to achieve a dark-acclimated state (data not shown). All
177 fluorescence parameters were calculated by standard equations (Genty et al., 1989; Maxwell
178 and Johnson, 2000). Maximum quantum yields of PSII (apparent PSII photochemical quantum
179 efficiency; F_v/F_m) were calculated as

$$180 \quad F_v/F_m = (F_m - F_0)/F_m \quad (1)$$

181 Fluorescence based photosynthesis-irradiance curves (PI) were conducted at six irradiances (I)
182 between 33 and 672 μmol photons $m^{-2} s^{-1}$, with an acclimation time of 10 min per light step.
183 Relative electron transfer rate (rETR) through PSII for each light step was calculated as:

$$184 \quad rETR = ((F_m' - F_0')/F_m') * I \quad (2)$$

185 Following the suggestion by Silsbe and Kromkamp (2012), the light-use efficiency was
186 estimated by fitting the data to the model by (Webb et al., 1974):

$$187 \quad rETR = rETR_{max} * [1 - e^{-(\alpha * I)/rETR_{max}}] \quad (3)$$

188 The light saturation index (E_k) was then calculated as $rETR_{max}/\alpha$. Maximum non-photochemical
189 quenching of Chl fluorescence (NPQ) at irradiances of 672 μmol photons $m^{-2} s^{-1}$ (i.e. the highest
190 irradiance step of the PI curve) were calculated using the normalized Stern-Volmer coefficient,
191 also termed NSV, as described in McKew et al. (2013):



192 $(F_q'/F_v')-1 = F_0'/F_v'$ (4)

193 where F_q' is the differences between measured and maximal fluorescence (Suggett et al., 2010).

194 F_0' was measured after each light step (with a duration of 90 s).

195

196 **2.5 Statistics**

197 All data is given as the mean of three biological replicates with \pm one standard deviation. To

198 test for significant differences between the treatments, two-way analyses of variance (ANOVA)

199 with additional normality (Kolmogorov-Smirnov) and Post Hoc (Holm-Sidak) tests were

200 performed. The significance level was set to 0.05. Statistical analyses were performed with the

201 program SigmaPlot (SysStat Software Inc, Version 12.5).

202



203 **3 Results**

204

205 **3.1 Carbonate Chemistry**

206 Regular dilution of cultures with pre-aerated seawater medium kept carbonate chemistry stable
207 over the course of the experiment. More specifically, in each bottle the drift in A_T and C_T
208 compared to initial values was $\leq 3\%$ and $\leq 4\%$, respectively (data not shown). Final carbonate
209 chemistry in the 2°C treatments yielded $p\text{CO}_2$ levels of 197 ± 3 , 323 ± 12 , 959 ± 22 and 1380
210 $\pm 53 \mu\text{atm}$ (Table 1). In the 6°C treatments, $p\text{CO}_2$ levels were 198 ± 6 , 394 ± 10 , 1036 ± 31 and
211 $1449 \pm 18 \mu\text{atm}$. Please note that the same $p\text{CO}_2$ level translates into differing dissolved CO_2
212 concentrations at different temperatures due to the temperature dependency of the carbonate
213 system. Specifically, the treatment $p\text{CO}_2$ values translated into up to 13% lower dissolved CO_2
214 concentrations in the 6°C compared to the 2°C treatment (Table 1; cf. Figure S11).
215 Concurrently, the $p\text{CO}_2$ levels at 2°C corresponded to pH_{total} values of 8.30 ± 0.01 , 8.11 ± 0.01 ,
216 7.68 ± 0.01 and 7.52 ± 0.02 , respectively. In the 6°C treatment, pH_{total} values of the four $p\text{CO}_2$
217 treatments were 8.30 ± 0.01 , 8.04 ± 0.01 , 7.65 ± 0.01 and 7.52 ± 0.01 , respectively.

218

219 **3.2 Growth and biomass build-up**

220 Growth rates constants of exponentially growing *M. pusilla* cultures were significantly affected
221 by the applied treatments (Figure 1, Table 2, S11). Depending on the $p\text{CO}_2$ level, temperature
222 increased growth by 20 to 60% with an average of 0.80 d^{-1} under low and 1.10 d^{-1} under high
223 temperature conditions (two-way ANOVA, $F = 328$, $p < 0.001$). Overall, there was also a
224 positive $p\text{CO}_2$ effect on growth (two-way ANOVA, $F = 9$, $p = 0.001$), even though no linear
225 trends with either $p\text{CO}_2$ or $[\text{CO}_2]$ were observed (Figure 1, S11). The observed $p\text{CO}_2$ responses
226 also differed between temperature levels, indicating a significant interaction between both
227 drivers (two-way ANOVA, $F = 12$, $p < 0.001$): Under low temperature, growth increased
228 significantly from 180 to 380 $\mu\text{atm } p\text{CO}_2$ (post-hoc, $t = 3.1$, $p = 0.04$), while there was a
229 declining, yet insignificant trend in growth with further increases in $p\text{CO}_2$. Under high
230 temperature, growth increased from 180 to 1000 $\mu\text{atm } p\text{CO}_2$ (post-hoc, $t = 5.6$, $p < 0.001$), and
231 declined from 1000 to 1400 $\mu\text{atm } p\text{CO}_2$ (post-hoc, $t = 5.9$, $p < 0.001$). Thus, warming shifted
232 the optimum range for growth to higher $p\text{CO}_2$ levels (Figure 1A).

233 This trend was also observed in terms of POC production rates (Figure 1B, Table 2),
234 with significant effects of temperature (Table S11; two-way ANOVA, $F = 356$, $p < 0.001$), $p\text{CO}_2$
235 (two-way ANOVA, $F = 7$, $p = 0.003$), and their interaction (two-way ANOVA, $F = 29$, p
236 < 0.001). At low temperatures, higher production rates were observed at 180 and 380 μatm



237 compared to those at 1000 and 1400 μatm pCO_2 (post-hoc tests, $t = 3.5$, $p = 0.016$ and $t = 3.0$,
238 $p = 0.046$, respectively). At high temperatures, POC production rates were significantly higher
239 at 1000 μatm than at all other pCO_2 levels (post-hoc tests, e.g. $t = 9.1$, $p < 0.001$ for 380 vs.
240 1000 μatm and $t = 7.4$, $p < 0.001$ for 1000 vs. 1400 μatm), again indicating an upward shift in
241 the pCO_2 optimum with warming.

242

243 3.3 Cellular composition

244 Overall, POC quota (Figure 2 a, Table 2, Table S11) were significantly higher under elevated
245 compared to low temperature (two-way ANOVA, $F = 24$, $p < 0.001$), but no overarching trend
246 with pCO_2 was observed. Under low temperature, cells had significantly higher POC quota at
247 low pCO_2 levels (180 and 380 μatm) compared to high pCO_2 levels (1000 and 1400 μatm ; all
248 four post-hoc tests significant, e.g. 380 vs 1000 μatm : $t = 2.8$, $p = 0.033$). This trend reversed
249 under high temperature, where POC quota were highest under 1000 and 1400 μatm (post-hoc
250 test, $t = 3.5$, $p = 0.024$). Thus, temperature and pCO_2 levels exhibited a significant interactive
251 effect on POC quota (two-way ANOVA, $F = 10$, $p < 0.001$).

252 Similar trends were observed in terms of cellular PON quota, where temperature (two-
253 way ANOVA, $F = 5$, $p = 0.045$) and its interaction with pCO_2 (two-way ANOVA, $F = 10$, p
254 < 0.001) significantly affected the results. Here, opposing pCO_2 effects under different
255 temperatures were more subtle, with PON quota under low temperatures only being
256 significantly decreased between 380 and 1400 μatm (post-hoc test, $t = 3.3$, $p = 0.027$), while
257 under high temperature PON quota significantly increased from 180 and 380 to 1000 μatm
258 pCO_2 (post-hoc tests, $t = 3.7$, $p = 0.012$ and $t = 2.8$, $p = 0.028$, respectively).

259 Regarding cellular Chl *a* quota, there were no significant effects of temperature or pCO_2
260 alone (Figure 2 b, Table 2, Table S11), but a significant interaction between the two drivers
261 (two-way ANOVA, $F = 18$, $p < 0.001$): Under low temperature, Chl *a* quota decreased from low
262 (180 μatm) to high pCO_2 levels (1000 and 1400 μatm ; post-hoc tests, $t = 5.0$, $p < 0.001$ and $t =$
263 3.9 , $p = 0.006$, respectively). Under high temperature, the opposite trend was observed, where
264 Chl *a* quota increased from low (180 and 380 μatm) to high pCO_2 levels (1000 and 1400 μatm ;
265 all four post-hoc tests significant, e.g. 380 vs 1000 μatm : $t = 3.0$, $p = 0.027$).

266 Molar C:N ratios of the biomass (Figure 2 c, Table 2, Table S11) increased with
267 temperature (two-way ANOVA, $F = 14$, $p = 0.002$), yet this overall difference was mainly
268 driven by results at low pCO_2 levels (180 and 380 μatm ; post-hoc tests, $t = 2.7$, $p = 0.017$ and t
269 $= 3.5$, $p = 0.003$, respectively). By itself, pCO_2 did not significantly affect C:N ratios.

270 The ratios of C:Chl *a* (Figure 2 d, Table 2, Table S11) were elevated under high



271 compared to low temperature conditions (two-way ANOVA, $F = 14$, $p = 0.002$), an effect that
272 was most pronounced at $p\text{CO}_2$ levels of $180 \mu\text{atm}$ (post-hoc test, $t = 5.5$, $p < 0.001$). While there
273 was no effect of $p\text{CO}_2$ on C:Chl α at low temperature, C:Chl α decreased with increasing $p\text{CO}_2$
274 at high temperature (two-way ANOVA, interaction term, $F = 6$, $p = 0.007$; 180 vs. $1400 \mu\text{atm}$
275 at 6°C post-hoc test, $t = 3.9$, $p = 0.008$).

276

277 **3.4 Chl α fluorescence-based photophysiology**

278 The effects of the applied treatments on photophysiology were investigated by means of FRRf,
279 which investigates photochemistry at photosystem II (PSII). No effects of the applied
280 treatments were observed in most parameters investigated (Table 3, SI1). This was true for the
281 dark-acclimated quantum yield efficiency of PSII (F_v/F_m), which was similar in all treatments
282 with values of 0.45 ± 0.06 , as well as for absorption cross section of PSII light harvesting (σ_{PSII}).

283 Furthermore, the fitted parameters of FRRf-based PI curves (α , $r\text{ETR}_{\text{max}}$ and E_K) were
284 independent of the experimental treatments (Table 3, SI1). In contrast, the rate constant of the
285 reopening of PSII reaction centres (τ_{ES} ; Table 3, SI1) was slightly yet significantly smaller
286 under high temperatures (two-way ANOVA, $F = 6$, $p = 0.029$), even though this overall
287 response also depended on the applied $p\text{CO}_2$ levels (two-way ANOVA, interaction term, $F = 4$,
288 $p = 0.033$).

289 Maximum non-photochemical quenching (NPQ_{max} ; Table 3, SI1) increased
290 significantly with $p\text{CO}_2$ (Table SI1; two-way ANOVA, $F = 0$, $p = 0.002$) while temperature had
291 no effect. Post-hoc tests revealed that this response was mainly driven by high NPQ_{max} values
292 at $1000 \mu\text{atm}$, which were significantly higher than in any other $p\text{CO}_2$ treatment (e.g. $t = 4.1$, p
293 $= 0.006$ for 380 vs. $1000 \mu\text{atm}$ and $t = 3.1$, $p = 0.030$ for 1000 vs. $1400 \mu\text{atm}$).

294 **4 Discussion**

295

296 **4.1 *Micromonas pusilla* benefits from warming**

297 We observed a strong stimulation of growth rates and biomass build-up with increasing
298 temperature (Figure 1, Table 2). Even though the isolate stems from 1.8°C water temperature,
299 the beneficial effects of warming from 2°C to 6°C are not surprising as *M. pusilla* is known to
300 dominate Arctic phytoplankton assemblages in the summer and autumn situations (Lovejoy et
301 al., 2007; Marquardt et al., 2016). Moreover, summer ocean surface temperatures at the location
302 of isolation usually reach 6°C or more (Hegseth et al., in press). Our results are also in line with
303 mesocosm experiments that indicate stimulatory effects of warming on picoplankton
304 abundances (Daufresne et al., 2009; Sommer et al., 2015) as well as with the temperature
305 optimum of 6-8°C observed for another Arctic strain of *M. pusilla* (Lovejoy et al., 2007).

306 Below the temperature optimum of a cell, warming causes an acceleration of the entire
307 metabolism, as enzymatic reactions run faster under these conditions (Eppley, 1972; Brown et
308 al., 2004). In this study, warming caused higher growth rates, POC quotas and biomass
309 production (Figure 2, Tables 2, SI1), indicating that particularly the fixation and storage of
310 carbon was facilitated by increasing temperature. Electron transport processes, on the other
311 hand, were largely independent of temperature (Tables 3, SI1). Thus, temperature affected the
312 balance between electron transport ('light reaction') and carbon fixation in the Calvin cycle
313 ('dark reactions'). Especially under relatively low temperatures, as investigated here, warming
314 can decrease the excitation pressure on the electron transport chain of the photosystems by
315 increasing the temperature-limited turnover rates of enzyme reaction such as RuBisCO (Mock
316 and Hoch, 2005). Thus, cells grown under low temperature need to invest relatively more
317 energy into biosynthesis than into photochemistry compared to cells grown under high
318 temperature (Toseland et al., 2013). While it has been shown that Antarctic diatoms can
319 compensate for slow RuBisCO kinetics by increasing the expression of this enzyme (Young et
320 al., 2014), it is unknown whether such acclimation responses also occur in prasinophytes.
321 Regarding the C:Chl *a* ratio, this can be taken as an indicator on how much resources the cell
322 retains as carbon biomass (e.g. structural and storage compounds) relative to how much is
323 invested into its light harvesting capacities (Halsey and Jones, 2015). In this study, the strong
324 temperature-dependent increase in C:Chl *a* (Figure 2, Table SI1) under potentially limiting
325 pCO₂ levels of 180 µatm suggests that under warming, the balance between light harvesting
326 and carbon fixation was indeed more beneficial for biomass build-up. Furthermore, elevated
327 temperature significantly decreased τ_{ES} (Table 3, SI1), which can serve as a proxy of the rate



328 at which down-stream processes can remove electrons from PSII (Kolber et al., 1998). Thus,
329 our results indicate that the drainage of electrons into carbon fixation was faster under warmer
330 conditions, explaining the higher growth and biomass production under these conditions.

331

332 **4.2 Warming shifts CO₂ optima towards higher pCO₂ levels**

333 Overall, also OA had a significantly positive effect on growth and biomass build-up (Figure 1).
334 This finding is in line with previous studies, which have shown that picoeukaryotes can benefit
335 strongly from OA in both laboratory and mesocosm studies (Meakin and Wyman, 2011;
336 Newbold et al., 2012; Schaum et al., 2012; Brussaard et al., 2013; Maat et al., 2014; Schulz et
337 al., 2017). Such positive response to OA could indicate that picoeukaryotes such as *M. pusilla*
338 are mainly dependent on diffusive CO₂ supply and thus directly benefit from higher CO₂
339 concentrations (Brussaard et al., 2013; Schulz et al., 2013; Schulz et al., 2017).

340 Despite this overall effect, growth rates of *M. pusilla* tended to follow an optimum
341 response curve over the tested range of glacial to elevated future pCO₂ levels (i.e. 180 to 1400
342 μ atm), i.e. growth increased with increasing pCO₂ from low to intermediate, but decreased
343 again under higher pCO₂ levels (Figure 1). Such an optimum behaviour can be expected for
344 most environmental drivers (Harley et al., 2017) and has previously been observed in response
345 to OA (Sett et al., 2014; Wolf et al., 2017). The response patterns in these studies were attributed
346 to a combination of beneficial effects of rising pCO₂ under potentially carbon-limiting
347 conditions for photosynthesis, and negative effects of declining pH on cellular homeostasis and
348 enzyme performance, which manifest mainly at high pCO₂ (Bach et al., 2013).

349 On a more general level, apparent discrepancies between OA studies can be attributed
350 to actual differences in the environmental settings and their interactive effects with pCO₂
351 (Riebesell and Gattuso, 2015). When comparing the two most commonly applied pCO₂ levels,
352 i.e. the present-day and the anticipated end-of-century situation, the effects of OA on most of
353 the investigated physiological parameters are reversed under 6°C compared to 2°C (Figure 3).
354 This illustrates how difficult it is to infer responses to OA from experiments applying only one
355 set of environmental conditions. It is also noteworthy that the combination of OA and warming
356 led to more densely packed cells (no change in cell size based on flow cytometric
357 measurements; data not shown) with similar stoichiometry compared to the control treatment
358 (Table 2). This indicates that cells managed to cope well with the experienced future conditions.
359 Furthermore, warming altered the OA-dependent change in most of the investigated parameters
360 in a direction that indicates higher fitness compared to low temperatures (e.g. higher growth
361 rates and higher elemental quota; Figure 3). Thus, the increase in growth under future compared



362 to ambient conditions was larger than what would be expected by the respective responses to
363 warming and OA in isolation, indicating synergistic beneficial effects of both drivers.

364

365 **4.3 Potential mechanism underlying the interaction between warming and OA**

366 The observed synergistic effects could be explained by their specific impacts on carbon
367 acquisition and fixation. As outlined in the introduction, light and dark reaction of
368 photosynthesis need to be balanced to achieve high biomass production while avoiding
369 photodamage (Behrenfeld et al., 2008). According to our data, this balance is shifted towards
370 higher biomass production rates under warming and OA.

371 At higher temperatures, seawater CO₂ concentrations were lower than under colder
372 conditions (Table 1; Zeebe and Wolf-Gladrow, 2001). At the same time, warming from 2°C to
373 6°C caused up to 60% higher growth and 110% higher biomass build-up rates (Figure 1, Table
374 2). Furthermore, the decrease in τ_{ES} indicates a faster transfer of photochemical energy into
375 downstream processes such as RuBisCO activity (Table 3). Increased carbon demand in concert
376 with lower carbon supply at higher temperatures thus increases the risk of CO₂ shortage in the
377 cell, which in turn causes OA to have larger effects than at colder temperatures. Moreover,
378 warming changes the kinetics of carbon fixation, with RuBisCO increasing its maximum
379 turnover rates but decreasing its affinity for CO₂ (Young et al., 2014). At higher temperature,
380 cells thus have the potential for higher carboxylation rates provided sufficient CO₂ is available
381 (Kranz et al., 2015). Under elevated pCO₂ levels, diffusive CO₂ supply increases and/or costs
382 for active carbon acquisition decrease. Consequently, the positive effect of increasing
383 temperature on the carbon fixation rate can develop its full potential under OA.

384 In conclusion, elevated catabolic activity under warmer conditions can explain the
385 observed upward shift in the CO₂-optimum of growth with increasing temperature (Figure 1),
386 as the corresponding higher carbon demand causes CO₂ fixation to saturate under higher pCO₂
387 levels. In combination with a faster and more efficient machinery for pH homeostasis at
388 elevated temperatures (Morgan-Kiss et al., 2006), this could explain why declining growth rates
389 were only observed at relatively higher pCO₂ levels compared to those under low temperature
390 conditions (Figure 2).



391 **4.4 Implications for the current and future Arctic pelagic ecosystem**

392 Picoeukaryotes such as *M. pusilla* are considered to be potential winners of climate change:
393 They are not only thriving in warmer, more stratified environments, which are predicted to
394 further expand in the future, but also seem to benefit from OA (Li et al., 2009; Schulz et al.,
395 2017). Our results for *M. pusilla* confirm beneficial effects of warming and OA on growth and
396 biomass production under nutrient-replete conditions (Figure 1, Table 2). Hence, for current
397 summer and autumn situations where *M. pusilla* dominates Arctic phytoplankton assemblages
398 today, but also for a warmer spring situation of the future, this species may experience growth
399 stimulation under OA. Hence, this species can be expected to thrive well under conditions
400 expected for the end of this century (Stocker, 2014), potentially increasing its ecological
401 relevance even further. Regarding the importance of the nutrient availability, laboratory
402 experiments found beneficial OA effects on *M. pusilla* primary production to persist also under
403 P limitation (Maat et al., 2014), while in a mesocosm community, OA-dependent increases in
404 *M. pusilla* abundances disappeared when the system ran into P and N co-limitation (Engel et
405 al., 2008). Thus, it remains to be seen how the combined effects of warming and OA manifest
406 under low nutrient conditions.

407 A species' success in the environment does not only depend on individual performance, but
408 also on how it compares to that of competing species. When we compare our results with the
409 responses of the Arctic diatom *Thalassiosira hyalina*, isolated from the same location and
410 exposed to the same experimental conditions (Wolf et al., 2017), the diatom had higher growth
411 rates than the picoeukaryote under most treatment conditions, as can be expected for nutrient-
412 replete conditions (Sarthou et al., 2005). The relative increase in growth rates from ambient (2
413 or 3°C and 380 $\mu\text{atm pCO}_2$) to future conditions (6°C and 1000 $\mu\text{atm pCO}_2$) was, however,
414 much higher for *M. pusilla* than for *T. hyalina*. This indicates a stronger increase in fitness
415 (Collins et al., 2014) and could mean that *M. pusilla* gains another competitive advantage in the
416 future, in addition to the predicted benefit arising from changes in stratification (Li et al., 2009).
417 Thus, our findings suggest higher picoplankton contribution to future Arctic phytoplankton
418 assemblages.

419 Even though picoeukaryotes seem to contribute more to the downward export of organic
420 matter than previously assumed (Waite et al., 2000; Richardson and Jackson, 2007), in
421 comparison to e.g. diatoms, they are less efficient vectors for carbon export to depth and have
422 a lower energy transfer along trophic levels (Sherr et al., 2003). Consequently, Arctic food webs
423 dominated by picoeukaryotes would look very different from those fuelled by diatom
424 production (Sherr et al., 2003; Paulsen et al., 2015). Due to its motility and capability to grow



425 mixotrophically, *M. pusilla* is characterized by an exceptionally high cellular C:N ratio
426 compared to other Arctic phytoplankton (Table 2; Halsey et al., 2014; McKie-Krisberg and
427 Sanders, 2014). An increased importance of this species would thus not only affect the food
428 web due to its small size and concurrent grazer preferences, but also in terms of food quality
429 (van de Waal and Boersma, 2012). This could indicate that higher growth rates and thus
430 abundances of this species may strengthen the Arctic microbial food web. Together with a
431 concurrent weakening of the classical diatom-fuelled food web, this could have severe
432 implications for the flow of energy and nutrients through future marine Arctic ecosystems (Post,
433 2016).

434

435 **4.5 Conclusions**

436 This study is the first to show synergistic effects of warming and OA on *M. pusilla*, one of the
437 most abundant species of the worlds' oceans. Individually, both warming and OA cause more
438 efficient biomass build-up. Beneficial effects manifest, however, even more strongly in
439 combination, when facilitated carbon acquisition (e.g. due to higher diffusive CO₂ supply) co-
440 occurs with higher fixation rates (e.g. due to higher turnover-rates of RuBisCO). Our results
441 provide an explanation for the observations of previous mesocosm studies, which indicated
442 beneficial effects of OA and warming on *M. pusilla* and other picoeukaryotes. This study is the
443 first attempt to characterise the responses of this Arctic key species to warming and OA, which
444 will help us to develop mechanistic phytoplankton functional types and more realistic model
445 representation of phytoplankton assemblages as well as their responses to multiple drivers.
446 Future studies are needed to elucidate further multifactorial environmental changes, addressing
447 both abiotic (e.g. changes in light and nutrients) as well as biotic (e.g. heterotrophy,
448 competition, grazers, viruses) interactions.



449 **Author Contributions**

450 C.J.M.H. and B.R. designed the study. C.J.M.H. and C.F. conducted the experiment. C.J.M.H.

451 analysed the data and prepared the manuscript with contributions from B.R. and C.F.

452

453 The authors declare that they have no conflict of interest.

454

455

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Table 1: Seawater carbonate chemistry at the end of the experiments ($n=3$; mean ± 1 s.d.). CO_2 partial pressure (pCO_2) and Dissolved CO_2 concentrations were calculated from total alkalinity (A_T) and pH_{total} at 2 or 6°C, a salinity of 32.7 using CO_2SYS (Pierrot et al., 2006), and phosphate and silicate concentrations of 10 and 100 $\mu\text{mol kg}^{-1}$, respectively. n.a. indicates that values are not available for this specific treatment.

Temperature [°C]	pCO_2 level [μatm]	pH total scale	A_T [$\mu\text{mol kg}^{-1}$]	C_T [$\mu\text{mol kg}^{-1}$]	dissolved CO_2 [$\mu\text{mol kg}^{-1}$]	pCO_2 [μatm]
2	180	8.3 ± 0.01	2264 ± 9	2024 ± 6	11.6 ± 0.2	197 ± 3
	380	8.11 ± 0.01	2244 ± 30	2124 ± 11	19.0 ± 0.7	323 ± 12
	1000	7.68 ± 0.01	2255 ± 45	2215 ± 23	56.4 ± 1.3	959 ± 22
	1400	7.52 ± 0.02	2243 ± 5	n.a.	81.1 ± 3.1	1380 ± 53
6	180	8.3 ± 0.01	2243 ± 28	1969 ± 10	10.0 ± 0.3	198 ± 6
	380	8.04 ± 0.01	2256 ± 21	2058 ± 7	20.0 ± 0.5	394 ± 10
	1000	7.65 ± 0.01	2262 ± 22	2178 ± 14	52.6 ± 1.6	1036 ± 31
	1400	7.52 ± 0.01	2265 ± 5	n.a.	73.6 ± 0.9	1449 ± 18



Table 2: Growth rate constants, POC production rates and cellular quota of Chl *a*, POC and PON as well as their ratios of *M. pusilla* at the end of the experiment under the different treatment conditions (n=3; mean ± 1 s.d.). Results from statistical analysis can be found in Table SI2.

Temperature [°C]	pCO ₂ [µatm]	Growth rate constant [d ⁻¹]	POC production [fmol cell ⁻¹ d ⁻¹]	POC quota [fmol cell ⁻¹]	PON quota [fmol cell ⁻¹]	Chl <i>a</i> quota [fg cell ⁻¹]	C:N [mol mol ⁻¹]	C:Chl <i>a</i> [g g ⁻¹]
2	180	0.75 ± 0.04	178 ± 8	239 ± 20	28.7 ± 2.6	28.6 ± 2.1	8.3 ± 0.1	100 ± 7
	380	0.85 ± 0.03	202 ± 11	237 ± 22	30.9 ± 1.8	24.7 ± 1.4	7.7 ± 0.3	115 ± 10
	1000	0.79 ± 0.05	156 ± 16	196 ± 25	24.9 ± 4.9	20.0 ± 2.9	8.0 ± 0.6	118 ± 5
	1400	0.82 ± 0.05	164 ± 9	199 ± 16	23.9 ± 2.3	22.0 ± 1.5	8.4 ± 0.1	109 ± 4
6	180	1.06 ± 0.03	261 ± 10	245 ± 3	26.9 ± 0.6	21.1 ± 0.8	9.1 ± 0.1	140 ± 7
	380	1.05 ± 0.03	237 ± 27	226 ± 26	25.9 ± 2.5	22.1 ± 2.6	8.7 ± 0.3	123 ± 11
	1000	1.25 ± 0.05	344 ± 8	275 ± 6	33.7 ± 2.0	27.2 ± 0.9	8.2 ± 0.6	122 ± 6
	1400	1.05 ± 0.04	277 ± 9	263 ± 7	31.1 ± 1.7	28.6 ± 3.2	8.5 ± 0.3	111 ± 16



Table 3.: FRR-fluorometrical PSII photochemistry measurements – PSII quantum yield efficiency Fv/Fm, functional absorption cross section (σ_{PSII}), rate of PSII re-opening (τ_{ES}), maximum non-photochemical quenching at 672 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ (NPQ_{max}), maximal absolute electron transfer rates through PSII (ETR_{max}), maximum light-use efficiency (initial slope α) and the light saturation index (E_K) under the different temperature and pCO_2 treatments ($n=3$; mean ± 1 s.d.). [ds.] indicates dimensionless units. Results from statistical analysis can be found in Table SI2.

Temp [°C]	pCO_2 [μatm]	Fv/Fm [ds.]	σ_{PSII} [$\text{nm}^{-2} \text{PSII}^{-1}$]	τ_{ES} [ms]	NPQ_{max} [ds.]	α [ds.]	ETR_{max} [ds.]	E_K [$\mu\text{mol photons m}^{-2} \text{s}^{-1}$]
2	180	0.50 \pm 0.01	8.66 \pm 0.35	439 \pm 8	2.26 \pm 0.18	0.42 \pm 0.05	33 \pm 2	81 \pm 13
	380	0.43 \pm 0.09	8.93 \pm 0.26	425 \pm 4	3.51 \pm 0.55	0.32 \pm 0.15	25 \pm 5	91 \pm 44
	1000	0.45 \pm 0.08	8.55 \pm 0.07	448 \pm 1	3.96 \pm 0.71	0.42 \pm 0.03	31 \pm 2	75 \pm 10
	1400	0.47 \pm 0.10	9.06 \pm 0.05	422 \pm 14	2.45 \pm 0.44	0.43 \pm 0.08	31 \pm 7	75 \pm 31
6	180	0.49 \pm 0.01	9.22 \pm 0.22	412 \pm 6	2.51 \pm 0.37	0.49 \pm 0.08	28 \pm 10	59 \pm 28
	380	0.43 \pm 0.12	8.83 \pm 0.17	427 \pm 6	2.83 \pm 0.59	0.38 \pm 0.09	35 \pm 14	90 \pm 17
	1000	0.41 \pm 0.07	8.91 \pm 0.22	422 \pm 11	4.94 \pm 1.46	0.33 \pm 0.09	32 \pm 5	100 \pm 21
	1400	0.45 \pm 0.04	8.71 \pm 0.50	428 \pm 19	2.93 \pm 0.50	0.38 \pm 0.04	40 \pm 5	104 \pm 6



Figure 1: Growth rate constants (A) and POC production (B) of *M. pusilla* under low (open symbols) and high temperatures (filled symbols) as a function of $p\text{CO}_2$ ($n=3$; mean ± 1 s.d.). Results from statistical analysis can be found in Table S12.

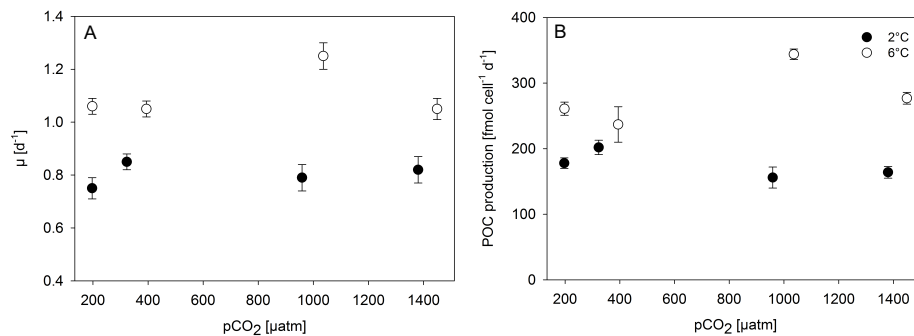




Figure 2: Cellular composition, i.e. POC (A) and Chl a quota (B) as well as as C:N (C) and C:Chl a ratios (D), of *M. pusilla* under low (open symbols) and high temperatures (filled symbols) as a function of $p\text{CO}_2$ ($n=3$; mean ± 1 s.d.). Results from statistical analysis can be found in Table SI2.

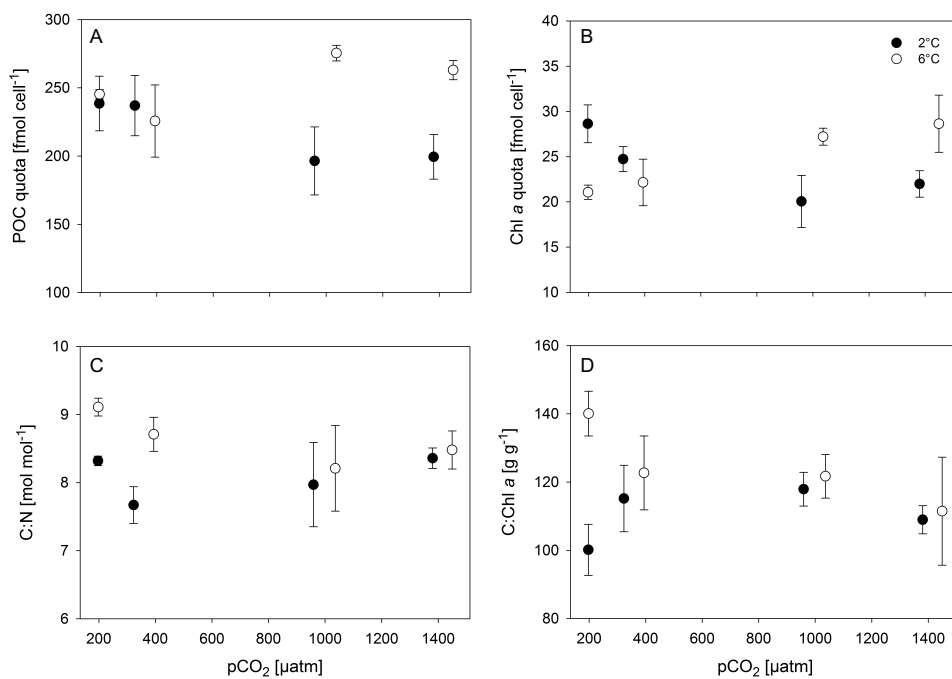




Figure 3: Schematic illustration of results for both temperatures over the entire range of pCO₂ levels as well as focusing on the responses between 380 and 1000 μatm (as the representation for commonly used OA treatments) and their modulation by temperature.

Parameter	Response curves	380 - 1000 μatm effect 2°C → 6°C
Growth		
POC quota		
PON quota		
Chla quota		
C:N		
C:Chla		