

1 **Aspects of resilience of polar sea ice algae to changes in their environment**

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8

9 **Abstract**

10 Sea ice algae are the primary producers of the ice-covered oceans in both polar regions.  
11 Changes in sea ice distribution are potentially altering exposure to photosynthetically active  
12 (PAR) and ultra-violet (UV-B) wavelengths of light. Incubations using monospecific cultures  
13 of common species from the Ross Sea, Antarctic Peninsula and Arctic Ocean were carried  
14 out at ecologically relevant light levels over periods of 7 d to examine their tolerance to  
15 conditions likely to be faced during sea ice thinning and melt. Algal responses were assessed  
16 using chlorophyll fluorescence techniques and superoxide dismutase activity. Quantum yields  
17 of cultures incubated in the dark and at ambient light did not differ. At higher light levels, the  
18 Ross Sea and Arctic cultures showed no significant change in photosynthetic health, while  
19 those from the Antarctic Peninsula showed a significant decrease. Antarctic cultures showed  
20 no detectable changes in superoxide dismutase activity, while the Arctic culture showed  
21 dynamic changes, initially increasing, then decreasing to the end of the study. The general  
22 lack of significant changes signals the need for further parameters to be assessed during such

23 experiments. The coupling between the measured parameters appeared to protect  
24 photosynthetic health, even though significant effects have been detected in other studies  
25 when subjected to PAR or UV-B alone.

26

27 **Key words:** Sea ice algae, *Thalassiosira antarctica*, *Chaetoceros socialis*, photoprotection,  
28 stress, ultra-violet B, Ross Sea, Antarctic Peninsula, Arctic Ocean

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30

## 31 **Introduction**

32 Polar sea ice forms one of the largest ecosystems on Earth. In the Antarctic it covers an area  
33 of  $19 \times 10^6$  km<sup>2</sup> at its maximum extent in winter, diminishing to  $3 \times 10^6$  km<sup>2</sup> during the  
34 summer (Arrigo, 2014). In the Arctic it oscillates between 15 and  $5 \times 10^6$  km<sup>2</sup>. The long-term  
35 rate of change is currently greater in the Arctic, where summer ice extent has reduced by 45%  
36 in only the past three decades and, in areas of rapid sea ice decrease, the spring retreat now  
37 initiates two months earlier and autumn advance one month later as compared to 1979/1980,  
38 resulting in the ice-free summer season lengthening by three months (Maksym et al., 2012;  
39 Stammerjohn et al., 2012). In the Antarctic overall winter sea ice extent has slightly increased  
40 (Turner et al., 2013; <http://www.ipcc.ch/report/ar5/wg1/#.UuhKY2SBoYI>) but its distribution  
41 has changed, with increases in the Ross Sea region and decreases in the Weddell Sea (Arrigo,  
42 2014; Turner et al., 2013). In the Antarctic Peninsula and Bellinghousen Sea region, changes  
43 in sea ice retreat and advance timing have again led to a three month longer ice-free period  
44 (Stammerjohn et al., 2012). In contrast, in the Ross Sea region, sea ice retreat is one month  
45 later and advance is one month earlier, leading to a shortening of the summer ice-free season  
46 by almost two months. Such changes in sea ice duration and distribution can cause severe  
47 ecological disruptions, with potentially negative consequences for the whole ecosystem  
48 (Arrigo, 2014; Nicol et al., 2008).

49 Sea ice algal communities reach their peak standing biomass in spring, leading to a  
50 significant increase in pigment concentration in the ice (Arrigo, 2014). This reduces the  
51 amount of light available to the upper water column, affecting algal productivity in the water  
52 below the ice (Arrigo et al., 1991; SooHoo et al., 1987). Phytoplankton blooms in the water  
53 column are delayed until after the ice algal bloom, whose timing therefore controls that of the  
54 subsequent phytoplankton bloom (Arrigo et al., 2012). When the sea ice melts, some of the  
55 algae released can provide seed stock for blooms at the ice edge (Cunningham & Leventer,

56 1998; Mangoni et al., 2009), as well as in the benthic and epiphytic habitats beneath  
57 (Majewska et al. 2013). However, this depends on their surviving the melting transition and  
58 acclimating very rapidly to the higher photosynthetically active radiation (PAR) and ultra-  
59 violet-B (UV-B) conditions that they are then exposed to (Mundy et al., 2011), and not being  
60 lost through sedimentation (Riebesell et al., 1991).

61 Sea ice can form annually, or survive to become thicker multi-year ice (Arrigo & Thomas,  
62 2004) through which light transmission is reduced. Snow cover, and its changing patterns,  
63 can further influence the under-ice environment (Arrigo, 2014). An emigration of diatoms  
64 from Arctic sea ice has been reported under thickness reducing conditions that could lead to  
65 reductions in productivity from the ice algal bloom, thereby impacting overall productivity  
66 (Lund-Hansen et al., 2013).

67 Microalgae are the primary producers found in the complex sea ice ecosystem, with diatoms  
68 (Bacillariophyceae) dominating (Arrigo, 2014) and reaching large stocks in the Antarctic due  
69 to the nutrient-rich waters of the Southern Ocean. The potential for nutrient limitation in the  
70 Antarctic is considered to be low, but local depletion can occur if there are other growth-  
71 limiting factors (Harrison & Cota, 1991) such as vertical stratification in the water column  
72 and depletion due to the sea ice algal bloom in spring. In both the Antarctic and Arctic factors  
73 other than nutrients, such as light and salinity, can also be limiting (Harrison & Cota, 1991).  
74 Algal cell concentrations in sea ice can vary by up to six orders of magnitude ( $<10^4$  to  $>10^9$   
75 cells  $l^{-1}$ ; Arrigo et al., 2010), a range that covers both typical global oceanic values and some  
76 of the highest recorded in any aquatic environment. Chlorophyll *a* (Chl *a*) biomass in sea ice  
77 varies by region, ice type and season. In the Arctic, volumetric Chl *a* concentrations range  
78 from 3 to 800 mg  $m^{-3}$ , and they can reach 10,100 mg  $m^{-3}$  in the Antarctic (Arrigo et al.,  
79 2010). The higher values in the Antarctic are again related to greater nutrient availability, and  
80 to lower annual light levels at the highest latitudes in the Arctic.

81 The community composition of the sea ice and associated ecosystems is determined by  
82 physical factors including light, salinity, nutrients and temperature (Arrigo & Thomas, 2004;  
83 Arrigo et al., 2010; Arrigo, 2014). Specific communities can include 30-170 diatom species.  
84 Commonly encountered genera in land-fast ice include *Nitzschia*, *Thalassiosira*,  
85 *Fragilariopsis* and *Navicula*. Microalgal biomass varies through the sea ice profile, with the  
86 highest levels found in the bottom 20 cm due to the stable light and temperature conditions  
87 present in this part of the sea ice (Arrigo & Thomas, 2004). There is also a constant supply of  
88 nutrients to this bottom layer through its interface with the underlying seawater (Arrigo,  
89 2014). Microalgal blooms in the sea ice are short-lived and are limited by low light and low  
90 temperatures (Ratkova et al., 2004). The distribution of algal biomass can also be patchy and  
91 show large variability (Rysgaard et al., 2001) due to local conditions such as surface snow  
92 cover and strong sub-ice water currents (Lund-Hansen et al., 2013).

93 Various studies have examined the effects of particular environmental parameters on  
94 monospecific cultures (e.g. Davidson et al., 1994; Hannach & Sigleo, 1998; Leu et al., 2010;  
95 Martin et al., 2012; Ryan et al., 2012). However, very few have examined the effect of  
96 multiple stressors (Halac et al., 2010; Hancke et al., 2008; Petrou et al., 2011; Salleh &  
97 McMinn, 2011) or compared common algal species from different parts of the polar regions  
98 (Petrou & Ralph, 2011). Brief periods of exposure to high light levels caused significant  
99 changes to the photosynthetic activity and composition (affecting its quality as a dietary item)  
100 of a common polar microalga, *Thalassiosira antarctica* var. *borealis* (Leu et al., 2006).

101 Exposure to higher PAR and UVR caused a decline in the quantum yield of photosystem II  
102 (PSII) and led to a significant reductions in C:P and N:P ratios. There was also a difference in  
103 the effects of PAR and UVR, with the algae affected strongly by increase in PAR but not  
104 UVR.

105 Ultra-violet radiation (UVR) reduces photo-protective capacity in diatoms such as  
106 *Phaedactylum tricorutum* (Halac et al., 2009), affecting the xanthophyll cycle and causing a  
107 decrease in photosynthetic health when exposed to saturating PAR. However, Halac et al.  
108 (2010) also demonstrated how increased temperature could counteract the negative effects of  
109 UVR, as well as variation in response due to length of exposure and size of cells, highlighting  
110 the importance of studying species-specific responses. High PAR and UVR stress can also  
111 lead to detrimental photoreactions (Janknegt et al., 2007), interrupting important metabolic  
112 pathways and causing an over-reduction of the electron transport chain (ETC). When this  
113 occurs, electrons from the ETC combine with molecular O<sub>2</sub> leading to the formation of  
114 reactive oxygen species (ROS). ROS cause damage to photosystem II reaction center  
115 proteins, thereby reducing the photosynthetic rate (Van De Poll et al., 2005). In response to  
116 this damage, cells produce antioxidant enzymes that actively scavenge ROS intermediates.  
117 The first ROS produced is a superoxide anion (O<sub>2</sub><sup>•-</sup>), which can be converted into the highly  
118 active hydroxyl radical (HO<sup>•</sup>) through a series of reductions. To avoid this, O<sub>2</sub><sup>•-</sup> is converted  
119 into hydrogen peroxide by the enzyme superoxide dismutase (SOD) (Gregory & Fridovich,  
120 1973). SOD is key to the cell's antioxidant response (Janknegt et al., 2007). Studies that  
121 describe SOD responses in marine polar microalgae are rare due to difficulties such as  
122 obtaining sufficient biomass for reliable measurements (Janknegt et al., 2009; Katayama &  
123 Taguchi, 2013; Van de Poll et al., 2009).

124 The primary aims of the current study were (1) to examine the photosynthetic (measured  
125 using chlorophyll fluorescence techniques) and antioxidative (measured using the RF/NBT  
126 assay) responses of sea ice algae in laboratory treatments combining ecologically relevant  
127 PAR and UV-B exposures, in order to study the capacity of common species to tolerate  
128 increased radiation levels such as might be experienced during ice melt and the thinning of  
129 sea ice, and (2) to compare acclimatory abilities across species. The study used monospecific

130 cultures of common algal species obtained from three regions with different patterns of  
131 environmental variation, namely the Ross Sea, the Antarctic Peninsula and the Arctic Ocean.  
132 Assessing the capacity of these algae for photo-protection will provide a better foundation for  
133 predictions relating to their response in context of climate change.

134

## 135 **Methods**

### 136 Culture methodology

137 Monospecific stock cultures were maintained in Victoria University of Wellington at a PAR  
138 level of  $50 \mu\text{mol m}^{-2} \text{s}^{-1}$  and  $4^\circ\text{C}$  in f/2 medium (Ausaqua Pty Ltd., Australia) for 5-6 months  
139 prior to these experiments. The Ross Sea species, *Thalassiosira antarctica*, used in the  
140 experiments described below were isolated from sea ice at Cape Evans in the summer of  
141 2010/2011. A stock culture of *Chaetoceros socialis* from the Antarctic Peninsula was  
142 obtained from Dr. Claire Hughes (University of East Anglia) as part of studies under the  
143 Rothera Time Series maintained by the British Antarctic Survey near its Rothera Research  
144 Station in Ryder Bay (Marguerite Bay,  $67^\circ 34' \text{S}$ ,  $68^\circ 9' \text{W}$ ) during the austral summer of  
145 2008/2009 (<http://www.antarctica.ac.uk/staff-profiles/webspace/mmm/RaTS/RaTS.html>). A  
146 stock culture of the same nominate species from the Arctic was obtained from Assoc. Prof.  
147 Else Hegseth (University of Tromso), originally collected from Billefjorden, Svalbard, in  
148 summer 2011.

### 149 Light treatments

150 The light treatments were based on measurements taken in the field in the Ross Sea through  
151 the sea ice profile (MAR pers. obs.; see also Rajanahally et al., 2014) and are representative  
152 of the PAR and UV- B levels that sea ice algae would be exposed to from the middle to the  
153 bottom of the sea ice profile (Eicken, 1992; Petrou & Ralph, 2011; Rajanahally et al., 2014;

154 Ryan et al., 2011). Sub-samples (400 ml) of monospecific cultures (n=3 at each level) were  
155 incubated under each of four light levels (L0=0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ , L1=1  $\mu\text{mol m}^{-2} \text{s}^{-1}$  of PAR,  
156 L2=45  $\mu\text{mol m}^{-2} \text{s}^{-1}$  of PAR and 50  $\text{mW m}^{-2}$  of UV-B, L3=100  $\mu\text{mol m}^{-2} \text{s}^{-1}$  of PAR and 50  
157  $\text{mW m}^{-2}$  of UV-B) at 4°C for 168 h in a temperature controlled water bath (Haake, Cleveland,  
158 OH, USA). These light levels are intended to be representative of those found in the water  
159 column below the sea ice (L0), at the bottom of the sea ice (L1), a quarter of the way above  
160 the water-ice interface (L2), or at the middle of the sea ice profile (L3). The separate PAR  
161 and UV-B levels were combined to reflect that the increase in PAR is synchronous with an  
162 increase in UV-B through the sea ice profile.

163 The 400 ml samples were incubated following standard approaches described by Ryan et al.  
164 (2012) in 500 ml opaque containers which were placed under 5W LED lights (Greenlights,  
165 Taiwan) and a Philips Ultraviolet-B TL 12/40W UV-B tube. Different sections of the water  
166 bath were separated from each other with opaque black plastic sheets and the various levels  
167 of light and UV-B were obtained with appropriate shade cloth. The UV-B tube was also  
168 covered with a polyvinyl chloride (PVC) sheet to filter out harmful UV-C radiation. PAR  
169 levels were measured using a SpectroSense 2 meter with an SKR 1850 radiometer (bandwidth  
170 400 – 700 nm) and UV-B with an SKU 430 broad band UV-B radiometer (bandwidth 280 -  
171 315 nm) (Skye Instruments, UK).

172 Samples of 1 ml for cell counts were taken at 0, 48 and 168 h from all replicates in each  
173 incubation, preserved using 2% Lugol's Iodine and stored at 4°C. Fifty ml of sample was  
174 used for the measurement of MAA content following the protocol described by Ryan et al.  
175 (2002) and Rajanahally et al. (2014). However, as no detectable amounts of MAAs were  
176 produced, no further consideration of MAAs is included here.

177 Chlorophyll *a* content

178 Twenty-five ml of sample were used for the measurement of chlorophyll *a* content. This was  
179 filtered onto GF/F filters which were then placed in containers with 5 ml 100% methanol and  
180 extracted in the dark at 4°C for 24 h. The extract was then measured on a digital fluorometer  
181 (Turner 10AU, Australia) following the acidification method of Evans et al. (1987).

#### 182 Superoxide dismutase activity

183 Twenty to 80 ml of the sample was stored in a container at -20°C for measurement of  
184 superoxide dismutase (SOD) activity following Janknegt et al. (2007). The variation in  
185 volumes collected for determination of SOD activity was due to varying growth rates  
186 between species. As SOD activity had not been previously determined for these cultures, it  
187 was not possible to identify the minimum volume required *a priori*. Therefore, a minimum of  
188 20 ml and, in some cases, a larger volume was collected as a contingency to ensure the level  
189 of SOD activity was assayable. These samples were centrifuged (2000 x g, 5 min, 4°C) in 1  
190 ml Eppendorf tubes and the pellets were stored at -80°C to preserve enzyme activity.

#### 191 *Cell lysis*

192 For enzyme measurements, cell pellets were thawed on ice, centrifuged (2000 x g, 5 min,  
193 4°C) and resuspended in 1 ml cold lysis buffer mixture of 300 µl potassium phosphate buffer  
194 (50 mM, pH 7.8) containing DTPA (0.2 mM), riboflavin (1.3 µM), L-methionine (10 mM),  
195 NBT (57 µM), and Triton X-100 (0.025% (v/v)). The suspension was then centrifuged (2000  
196 x g, 5 min, 4°C) again before the cells were resuspended in a final volume of 500 µl of lysis  
197 buffer and sonicated (Ultrasonic liquid processor W-380, Heat Systems-Ultrasonics, Inc.,  
198 New York) on ice for three 15 s pulses with a time interval of 10 s between each pulse. All  
199 lysates were centrifuged (16000 x g, 5 min, 4°C) and supernatants aliquoted and frozen at -  
200 80°C until further analysis. Total aqueous soluble protein content was determined using the  
201 improved Bradford assay with BSA as standard (Zor & Selinger, 1996).

202 *Superoxide dismutase (SOD)*

203 SOD assays were performed using the riboflavin/nitroblue tetrazolium (RF/NBT) assay in a  
204 microtiter plate format (Beauchamp & Fridovich, 1971; Fryer et al., 1998). Samples of 20  $\mu$ l  
205 of lysate or SOD standard (0.5-500 U ml<sup>-1</sup>) were added into a final reaction mixture of 300  $\mu$ l  
206 potassium phosphate buffer (50 mM, pH 7.8) containing EDTA (0.1 mM), riboflavin (1.3  
207  $\mu$ M), L-methionine (10 mM), NBT (57  $\mu$ M), and Triton X-100 (0.025% (v/v)). Absorbance  
208 was read at 560 nm both immediately and after 10 min incubation under a homogenous light  
209 field (130  $\mu$ mol m<sup>-2</sup> s<sup>-1</sup>) at 25°C. Standards and samples were measured using the same  
210 reaction mixture and a sigmoidal 5-parameter semi logarithmic standard curve (24 standard  
211 levels) used to determine SOD activity of samples. One unit of SOD activity was defined as  
212 the amount of enzyme that inhibited the NBT reduction by 50% (Beyer & Fridovich, 1987).

213 Sub-sampling, PAM fluorometry, and statistical analyses

214 PAM fluorometry is a widely used tool for studying the photosynthetic health of microalgae  
215 (Hancke et al., 2008; Katayama & Taguchi, 2013; McMinn & Hegseth, 2004; Parkhill et al.,  
216 2001; Ryan et al., 2009) as it provides a non-invasive method to study smaller cultures. A  
217 Water PAM fluorometer (Walz, Effeltrich, Germany) was used to measure effective quantum  
218 yield for photosystem II ( $\phi_{PSII}$ ). Before commencing observations, the cells were dark  
219 acclimated. Thereafter, all PAM measurements were made on cells exposed to light as the  
220 cells were subjected to actinic light used to make the PAM measurements. The Water PAM  
221 was also used to generate rapid light curves (RLC). An RLC describes the effective quantum  
222 yield as a function of irradiance (Ralph & Gademann, 2005). Each sample was treated with a  
223 series of eight increasing actinic light treatments (0, 8, 20, 35, 55, 75, 96, 114, 150  $\mu$ mol m<sup>-2</sup>  
224 s<sup>-1</sup>), after which a strong saturating pulse was applied and  $\phi_{PSII}$  was recorded. The RLC took  
225 90 s to generate. The electron transport rate (ETR) values were calculated by multiplying the  
226  $\phi_{PSII}$  value by the irradiance just applied. As the  $\phi_{PSII}$  value is a ratio and ETR is derived from

227 this parameter, it is termed relative ETR (rETR). An RLC permits derivation of different  
228 parameters that can be used to describe the photosynthetic properties of an algal sample,  
229 including rETR<sub>max</sub> (the maximum value for rETR), photosynthetic efficiency ( $\alpha$ ) and  
230 saturation irradiance ( $E_k$ ) (Ryan et al., 2009). To determine these parameters, the rETR data  
231 were imported into Microsoft Excel v 10.0 (Microsoft, USA) and the curve was fitted with a  
232 “waiting-in-line” function as described by Ritchie (2008). An RLC was generated for each  
233 sample at times 0, 2, 4, 6, 12, 24, 48, 72, 96, 120, 144 and 168 h.

234

### 235 Statistical analyses

236 First, a repeated measures ANOVA was applied to test specifically for changes over time  
237 within each treatment. The assumptions of normality and equal variance were satisfied. As  
238 the data from successive time points are likely not to be independent the covariance matrix -  
239 the matrix of all covariances between time points - was examined. If that matrix was  
240 spherical (as confirmed by Mauchly’s test of sphericity), then a standard repeated measures  
241 ANOVA calculation was carried out. If, however, the sphericity assumption did not hold,  
242 then the Greenhouse Geisser adjustment was applied (Greenhouse & Geisser, 1959).  
243 Repeated measures ANOVA was used to analyse effective quantum yield and rETR<sub>max</sub>, with  
244 ‘treatment’ being the irradiance treatments and the different time points. The significance of  
245 differences between pairs of time points was adjusted using Bonferroni’s correction. Finally,  
246 one-way ANOVA was used to compare the different treatment groups at specific time points.  
247 *Post hoc* tests using Bonferroni’s correction were used for subsequent pair-wise comparisons,  
248 e.g. between different pairs of treatment groups at a time point, or between different time  
249 points within a treatment group. All differences referred to are statistically significant at  $p <$   
250 0.05.

251

252 **Results**

253 Effect of different light levels on the photosynthetic performance of *Thalassiosira antarctica*  
254 (Ross Sea), *Chaetoceros socialis* (Antarctic Peninsula) and *C. socialis* (Arctic)

255 Initially a two way repeated measures ANOVA was conducted for all parameters with time as  
256 the ‘within’ factor and treatment and species as ‘between’ factors (Table 1). This  
257 demonstrated that both time alone and all the interaction combinations (excepting the three-  
258 way time-treatment-species interaction on  $E_k$ ) had a highly significant influence on each  
259 parameter. Since the significant 3-way interactions make it impossible to interpret the  
260 treatment effects, repeated measures analyses were carried out for each species separately to  
261 test for the effect of all treatments and time.

262 *Quantum Yield ( $\phi_{PSII}$ )*

263 A repeated measures ANOVA was carried out for each species separately to test for effect of  
264 all treatments and time (Table 2). Although the general response, except for *Chaetoceros*  
265 *socialis* (Antarctic Peninsula) at L2, appeared to show consistent patterns over time, the  
266 repeated measures ANOVA for algae incubated at all light treatments was not significant for  
267  $\phi_{PSII}$  over 168 h.

268 All the algal cultures showed no significant difference in quantum yield between those  
269 incubated in the dark (L0) or at ambient low light (L1) levels. For both the Antarctic species,  
270 quantum yield was lowest for algae incubated at L2 and highest for cultures incubated in the  
271 dark or at ambient light. For both species, quantum yield decreased significantly over the first  
272 two hours. However, quantum yield for *T. antarctica* at L2 did not show a significant change  
273 over 168 h while *C. socialis* showed an overall decrease. In the Arctic *C. socialis*, there was  
274 no overall change in quantum yield over 168 h. However, at the end of the incubation, the  
275 quantum yield for the algae at ambient low light level was the highest and that of those in the

276 dark and at the highest light exposure were the lowest. The general response for  
277 *Thalassiosira antarctica* (Ross Sea) (Figure 1(a)) at L2 and L3 was an initial decrease  
278 followed by a recovery in  $\phi_{PSII}$ , then maintenance of a stable value over the rest of the  
279 incubation. Algae at L0 and L1 showed very little change over 168 h. The general response  
280 for *C. socialis* (Antarctic Peninsula) (Figure 1(b)) at L0 and L1 was a slight increase followed  
281 by  $\phi_{PSII}$  remaining stable over 168 h. For algae incubated at L2 and L3, there was a sharp  
282 decrease over the first 2 h, followed by a recovery in  $\phi_{PSII}$  between 2 and 4 h. However, algae  
283 at the higher light treatments of L2 and L3 then showed a gradual decrease in  $\phi_{PSII}$  to reach  
284 levels similar to those at 2 h. Although the repeated measures ANOVA showed no significant  
285 change in  $\phi_{PSII}$  over 168 h for this species at L0, L1 and L3, those at L2 showed a significant  
286 decrease over the full 168 h period. *C. socialis* (Arctic Ocean) (Figure 1(c)) showed a gradual  
287 decrease in  $\phi_{PSII}$  for algae incubated at L0 and a gradual increase for algae incubated at L1.  
288 Algae incubated at L2 showed an initial decrease in  $\phi_{PSII}$  followed by a recovery to reach the  
289 same levels as at the start of the incubation. Algae incubated at L3 also showed an initial  
290 decrease, but this was followed by a recovery to reach a  $\phi_{PSII}$  lower than that at time 0 h.

291 *rETR<sub>max</sub>*

292 A repeated measures ANOVA was carried out for each species separately to test for effect of  
293 all treatments and time (Table 3). Although the general response showed changes over time,  
294 the repeated measures ANOVA showed no significant change in  $rETR_{max}$  over 168 h for  
295 algae at all treatments, except for both *Chaetoceros* cultures at L2.

296 The general response of *Thalassiosira antarctica* (Ross Sea) (Figure 2(a)) at L2 and L3 was  
297 an initial increase followed by  $rETR_{max}$  varying little throughout the incubation and not  
298 showing any change from time 0 h. Algae at L0 showed a slight decrease in  $rETR_{max}$  before it  
299 stabilized for the rest of the incubation. Algae incubated at L1 showed an increase in  $rETR_{max}$   
300 over the 168 h. The general response of *Chaetoceros socialis* (Antarctic Peninsula) (Figure

301 2(b)) for all light treatments other than at L0 was a slight increase in rETR<sub>max</sub> followed by a  
302 decrease to reach a level higher than at time 0 h. Algae at L0 did not show any change over  
303 time. Those incubated at L2 showed a significant increase in rETR<sub>max</sub> between 144 and 168  
304 h. However, there was no overall significant increase. The general response of *C. socialis*  
305 (Arctic Ocean) (Figure 2(c)) for all light treatments other than at L0 was a slight increase in  
306 rETR<sub>max</sub> followed by a decrease to reach a level similar to that at time 0 h. Algae at L0 did  
307 not show any change over time. Algae incubated at L2 showed an overall significant increase  
308 in rETR<sub>max</sub>.

### 309 Chlorophyll *a*

310 Repeated measures ANOVA indicated a significant effect of time ( $p < 0.005$ ) and the  
311 interaction between time and species ( $p < 0.005$ ), but no significant interaction between time  
312 and light. However, Levene's test for equality of variances was significant at each time point,  
313 indicating that the data were highly variable. The chlorophyll data were, thus, too variable to  
314 permit conclusions to be drawn about the responses of the algae.

### 315 Superoxide dismutase

316 Two way repeated measures ANOVA conducted with time as the 'within' factor and  
317 treatment and species as 'between' factors identified significant 3-way interactions. Repeated  
318 measures analyses were, therefore, carried out for each species separately to test for the effect  
319 of all treatments and time (Table 4).

320 Although the general responses appeared to show small and consistent patterns of change  
321 over time, the repeated measures ANOVA showed no significant change in SOD activity  
322 over 168 h for algae at all light treatments, except for *Chaetoceros socialis* (Antarctic  
323 Peninsula) which had a significant increase in the first 48 h for all light treatments followed  
324 by a significant decrease by 168 h, resulting in no overall change over 168 h.

325 The general response of *Thalassiosira antarctica* (Ross Sea) (Figure 3(a)) in the dark and at  
326 low light levels was an increase in SOD activity. At the two higher light levels, SOD activity  
327 increased over the first 48 h, followed by a decrease by the end of the incubation at 168h. The  
328 general response of *Chaetoceros socialis* (Antarctic Peninsula) (Figure 3(b)) at all light levels  
329 was an increase in SOD activity over the first 48 h, followed by a decrease to reach a similar  
330 level at 168 h to that at the start of the incubation. That of *C. socialis* (Arctic Ocean) (Figure  
331 3(c)) for all light treatments was a large decrease in SOD activity over the first 48 h, followed  
332 by a slight recovery by 168 h.

333

### 334 **Discussion**

335 Each of the species studied here is a common member of ice algal communities from each of  
336 three regions with different patterns of environmental variation, namely the Ross Sea, the  
337 Antarctic Peninsula and the Arctic Ocean. In the western Antarctic Peninsula (WAP) region,  
338 diatoms respond positively to increase in irradiance, with the effect being modulated by cell  
339 volume (Vernet et al., 2008). Although there has been a decrease in summer sea ice in this  
340 region leading to a decrease in phytoplankton blooms containing large, chain-forming  
341 diatoms (Garibotti et al., 2005), flagellate blooms are still able to maintain daily productivity  
342 (Vernet et al., 2008). These species-specific tolerances emphasise the variability that could be  
343 observed in environmentally similar regions with varying community composition.

344 In terms of the response of quantum yield to exposure to different light levels, differences  
345 between the same species from different regions could indicate specialization according to  
346 the niche they occupy. Species-specific sensitivities to environmental change have been  
347 observed elsewhere, suggesting that there is a link between photosynthetic capacity and  
348 ecological niche occupancy (Petrou & Ralph, 2011).

349 The combinations of PAR and UV-B exposures used, which mimic those that might  
350 reasonably be expected in the natural environment, mean that it is difficult to differentiate  
351 why a lack of change in quantum yield was observed at the higher light exposures. These  
352 exposures combine two parameters that are complementary to each other in the natural  
353 ecosystem, giving a better indication of how algae may tolerate increased radiation levels  
354 during summer and/or due to thinner sea ice conditions. The failure to find detectable levels  
355 of MAA production may indicate that MAAs are not amongst the mechanisms used by these  
356 algal species for photo-protection, unlike Rajanahally et al.'s (2014) clear demonstration of  
357 MAA production in experimental exposures of mixed multi-species algal communities soon  
358 after collection in the field in the same region of the Ross Sea used to source one of the  
359 cultures used here.

360 When the three species were compared at each light level, there was no significant change in  
361 quantum yield for algae incubated in the dark or at ambient low light, while those incubated  
362 at higher light exposures showed an overall significant decrease. At both higher light levels,  
363 the Arctic culture of *C. socialis* had the highest quantum yield at 168 h, perhaps indicating a  
364 better ability to tolerate higher light exposure. During ice melt, algae are obviously exposed  
365 to both varying and/or increasing levels of light and other parameters which are more  
366 complex than those used in the current study. Van de Poll et al. (2009), studying the Antarctic  
367 marine diatom *Chaetoceros brevis*, identified no difference in growth rates of algae exposed  
368 to constant or dynamic irradiance regimes in iron-limited or replete conditions. Iron  
369 limitation and wind-driven vertical mixing that resulted in a dynamic irradiance regime were  
370 frequently co-occurring conditions that affected photosynthetic health of algae. However,  
371 their study identified differences in pigment composition, quantum yield and antioxidant  
372 capacity between algae exposed to the two irradiance regimes. Cellular pigment  
373 concentrations increased three-fold under dynamic as compared to constant irradiance, under

374 iron-replete conditions. In iron-limited conditions smaller differences were detected in  
375 cellular pigment concentrations between the two regimes, suggesting reduced acclimation  
376 potential.

377 In the current study, the Arctic cultures had the highest SOD activity and the two Antarctic  
378 cultures showed equal activity at the start of all incubations. At all light exposures, the SOD  
379 activity did not show a difference between treatments at 168 h. This lack of variation in SOD  
380 production at the different light treatments was unexpected (cf. Janknegt et al., 2009; Van de  
381 Poll et al., 2009), with algae incubated at the higher light levels predicted to express greater  
382 activity. It is possible that even the higher irradiance exposures used in this study were not  
383 high enough to cause stress in these algal cultures. Other studies that have observed oxidative  
384 stress have often used irradiances exceeding  $1000 \mu\text{mol m}^{-2} \text{s}^{-1}$  (Van de Poll & Buma, 2009).  
385 Janknegt et al. (2009), conversely, found that SOD activity in some microalgae decreased  
386 with exposure to high irradiance. An increase in the de-epoxidation of the xanthophyll  
387 pigments led to more efficient recovery of PS II and reduced the production of hydrogen  
388 peroxide, in turn reducing the need for production of SOD. The variation in SOD activity was  
389 correlated with the cell size of the species examined, with those that had a higher surface area  
390 to volume ratio showing an increase in SOD activity whereas those with lower ratio showed a  
391 decrease. In the species examined in the current study, *C. socialis* cells are known to be  
392 smaller than those of *T. antarctica*. SOD activity in *C. socialis* from the Arctic (but not the  
393 Antarctic Peninsula) was higher than that of *T. antarctica*. The difference in SOD activity  
394 between cultures of *C. socialis* obtained from the Antarctic Peninsula and the Arctic could be  
395 related to the degree of change in the two regions, although to confirm this would require  
396 detailed measuring and experimental studies of cultures obtained from multiple locations.

397 This study took place over a one week manipulation period in order to examine any effect of  
398 length of incubation on acclimatory abilities. For all three species, there were very few

399 significant changes in photosynthetic parameters between 2 and 7 d, which strongly supports  
400 the utility of shorter incubation periods in experimental studies of this type. In the natural  
401 environment, the time taken for sea ice to melt can vary considerably and is influenced by  
402 various factors such as snow cover and initial thickness (Thomas & Dieckmann, 2003).

403 In this study, the combination of exposure to PAR and UV-B led to a lack of differentiation  
404 in algal photosynthetic health, contrasting with the significant effects on photosynthetic  
405 parameters and MAA production that was previously observed in monospecific Ross Sea  
406 cultures when subjected to increases in PAR or UV-B separately (Rajanaahally, 2014). During  
407 and after sea ice melt, algae face increases in PAR, UV-B and temperature and decreases in  
408 salinity, and the consequences of these effects cannot be estimated simply in an additive  
409 fashion (Folt et al., 1999). These responses will be further complicated by responses to  
410 changing CO<sub>2</sub> concentrations and ocean acidification processes (e.g. Gao et al., 2012, Rost et  
411 al., 2006). It is therefore essential for multivariate studies to be conducted that combine as  
412 many of these factors as possible in order to give insight into how algae tolerate the entirety  
413 of the processes of ice melt and release into the pelagic environment. Studies including  
414 evaluation of the photo-protective responses of these algae, such as that described here, will  
415 help assess their ability to survive current and predicted magnitudes of climate change and,  
416 hence, assess risks to the productivity of these ice-covered regions of the global oceans.

417

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427

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Table 1. Summary of results of statistical analyses of changes in quantum yield of photosystem II,  $rETR_{max}$ , Alpha( $\alpha$ ) and  $E_k$  for *Thalassiosira antarctica* (Ross Sea), *Chaetoceros socialis* (Antarctic Peninsula) and *C. socialis* (Arctic) over 168 h experimental treatments at different combinations of PAR and UV-B levels (L0=0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L1=1  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L2=45  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 50  $\text{mWm}^{-2}$ ; L3=100  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 100  $\text{mWm}^{-2}$ ) at 4°C, analysed using repeated measures ANOVA. Detailed analyses are presented in Tables 2 and 3.

Source of variation	Time	Time x Treatments	Time x Species	Time x Treatments x Species
$\phi_{PSII}$	<0.001	<0.001	<0.001	<0.001
$rETR_{max}$	<0.001	<0.001	<0.001	<0.001
Alpha( $\alpha$ )	<0.001	<0.001	<0.001	<0.001
$E_k$	0.002	<0.001	0.001	0.064

Table 2. Changes in quantum yield of photosystem II over 168 h experimental treatments at different combinations of PAR and UV-B levels (L0=0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L1=1  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L2=45  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 50  $\text{mWm}^{-2}$ ; L3=100  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 100  $\text{mWm}^{-2}$ ) at 4°C, analysed using repeated measures ANOVA. Significant changes identified by *post hoc* pairwise analyses are also listed.

Source of variation	df	MS	F	p	<i>post hoc</i> comparisons
<b><i>Thalassiosira antarctica</i> (Ross Sea)</b>					
<b>Within subjects</b>					
Time	2.340	0.072	54.188	<0.001	No change over 168 h
Time x Treatments	7.021	0.027	20.116	<0.001	(see text for details)
Error	18.723	0.001			
<b>Between subjects</b>					
Treatments	3	0.479	91.685	<0.001	(L0=L1) > L3 > L2
Error	8	0.005			
<b><i>Chaetoceros socialis</i> (Antarctic Peninsula)</b>					
<b>Within subjects</b>					
Time	4.325	0.034	41.242	<0.001	Decrease over 168 h
Time x Treatments	12.974	0.029	34.796	<0.001	(see text for details)
Error	34.597	0.001			
<b>Between subjects</b>					
Treatments	3	0.687	143.726	<0.001	(L0=L1) > L3 > L2
Error	8	0.003			
<b><i>C. socialis</i> (Arctic Ocean)</b>					
<b>Within subjects</b>					
Time	3.249	0.074	67.017	<0.001	No change over 168 h
Time x Treatments	9.746	0.029	26.218	<0.001	(see text for details)
Error	25.991	0.001			
<b>Between subjects</b>					
Treatments	3	0.329	78.456	<0.001	L1 > (L0=L3) > L2
Error	8	0.004			

Table 3. Changes in  $rETR_{max}$  of photosystem II over 168 h experimental treatments at different combinations of PAR and UV-B levels (L0=0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L1=1  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L2=45  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 50  $\text{mWm}^{-2}$ ; L3=100  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 100  $\text{mWm}^{-2}$ ) at 4°C, analysed using repeated measures ANOVA. Significant changes identified by *post hoc* pairwise analyses are also listed.

Source of variation	df	MS	F	p	<i>post hoc</i> comparisons
<b><i>Thalassiosira antarctica</i> (Ross Sea)</b>					
<b>Within subjects</b>					
Time	1.755	707.828	43.814	<0.001	No change over 168 h
Time x Treatments	5.266	229.137	14.184	<0.001	(see text for details)
Error	14.043	16.155			
<b>Between subjects</b>					
Treatments	3	703.075	149.478	<0.001	L1 > L2 > L3 > L0
Error	8	4.704			
<b><i>Chaetoceros socialis</i> (Antarctic Peninsula)</b>					
<b>Within subjects</b>					
Time	2.824	157.000	38.798	<0.001	No change over 168 h
Time x Treatments	8.471	82.856	20.475	<0.001	(see text for details)
Error	22.590	4.047			
<b>Between subjects</b>					
Treatments	3	489.269	119.725	<0.001	L1 > L2 > L3 > L0
Error	8	4.087			
<b><i>C. socialis</i> (Arctic Ocean)</b>					
<b>Within subjects</b>					
Time	2.929	620.532	283.378	<0.001	Increase over 168 h
Time x Treatments	8.786	78.510	35.853	<0.001	(see text for details)
Error	23.429	2.190			
<b>Between subjects</b>					
Treatments	3	670.216	82.660	<0.001	L2 > (L1 = L3) > L0
Error	8	8.108			

Table 4. Changes in SOD activity over 168 h experimental treatments at different combinations of PAR and UV-B levels (L0=0  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L1=1  $\mu\text{mol m}^{-2} \text{s}^{-1}$ ; L2=45  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 50  $\text{mWm}^{-2}$ ; L3=100  $\mu\text{mol m}^{-2} \text{s}^{-1}$  and 100  $\text{mWm}^{-2}$ ) at 4°C, analysed using repeated measures ANOVA. Significant changes identified by *post hoc* pairwise analyses are also listed.

Source of variation	df	MS	F	p	<i>post hoc</i> comparisons
<b><i>Thalassiosira antarctica</i> (Ross Sea)</b>					
<b>Within subjects</b>					
Time	2	55299.731	3.817	0.044	No change over 168 h
Time x Treatments	6	23650.055	1.632	0.202	(see text for details)
Error	16	14488.0404			
<b>Between subjects</b>					
Treatments	3	35853.695	2.271	0.157	L0 = L1 = L2 = L3
Error	8	15790.600			
<b><i>Chaetoceros socialis</i> (Antarctic Peninsula)</b>					
<b>Within subjects</b>					
Time	2	326604.643	23.878	<0.001	No change over 168 h
Time x Treatments	6	18888.579	1.381	0.281	(see text for details)
Error	16	13677.826			
<b>Between subjects</b>					
Treatments	3	11406.406	0.588	0.640	L0 = L1 = L2 = L3
Error	8	19382.839			
<b><i>C. socialis</i> (Arctic Ocean)</b>					
<b>Within subjects</b>					
Time	1.255	1307937.01	47.731	<0.001	Decrease over 168 h
Time x Treatments	3.766	16582.805	0.605	0.659	(see text for details)
Error	10.042	27402.151			
<b>Between subjects</b>					
Treatments	3	12677.400	0.910	0.478	L0 = L1 = L2 = L3
Error	8	13924.998			