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Pelagic and ice-associated microalgae under elevated light and pCO₂: Contrasting physiological strategies in two Arctic diatoms

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Abstract

Sea ice retreat, changing stratification, and ocean acidification are fundamentally changing the light availability and physico-chemical conditions for primary producers in the Arctic Ocean. However, detailed studies on ecophysiological strategies and performance of key species in the pelagic and ice-associated habitat remain scarce. Therefore, we investigated the acclimated responses of the diatoms Thalassiosira hyalina and Melosira arctica toward elevated irradiance and CO_2 partial pressures (pCO₂). Next to growth, elemental composition, and biomass production, we assessed detailed photophysiological responses through fluorometry and gas-flux measurements, including respiration and carbon acquisition. In the pelagic T. hyalina, growth rates remained high in all treatments and biomass production increased strongly with light. Even under low irradiances cells maintained a high-light acclimated state, allowing them to opportunistically utilize high irradiances by means of a highly plastic photosynthetic machinery and carbon uptake. The ice-associated *M. arctica* proved to be less plastic and more specialized on low-light. Its acclimation to high irradiances was characterized by minimizing photon harvest and photosynthetic efficiency, which led to lowered growth. Comparably low growth rates and strong silification advocate a strategy of persistence rather than of fast proliferation, which is also in line with the observed formation of resting stages under low-light conditions. In both species, responses to elevated pCO₂ were comparably minor. Although both diatom species persisted under the applied conditions, their competitive abilities and strategies differ strongly. With the anticipated extension of Arctic pelagic habitats, flexible high-light specialists like *T. hyalina* seem to face a brighter future.

In the Arctic, rapid increases in air and sea surface temperature (Timmermans et al. 2015; Maturilli et al. 2019) accelerate the thinning and retreat of sea ice (Wadhams 2012; IPCC 2021). In currently still ice-covered regions, a transition trend from multi- to first-year-ice can be observed, with overall earlier ice breakup, melting of the light-reflective snow cover, as well as formation of melt ponds. These phenomena profoundly increase the irradiance intensity and variability, which is one of the most important drivers of productivity, not only in the upper ocean water column but also in underice habitats (Hill et al. 2018; Lannuzel et al. 2020; Nicolaus et al. 2012). Furthermore, rising atmospheric CO_2 partial pressure (pCO₂) causes ocean acidification, which is especially pronounced in the Arctic because of low temperatures and lower buffering capacities of the freshening sea water (AMAP 2018; Zhang et al. 2020).

Since food webs depend critically on photosynthetic primary producers, the described environmental transitions will likely affect many other organisms in the trophic network, such as invertebrates, fish larvae and ultimately also top predators (Søreide et al. 2010). Diatoms are the dominant primary producers in the Arctic (Armbrust 2009) and occupy diverse niches, including the open ocean as well as sea-ice. In line with these niches, recent studies found diatoms to follow different strategies to cope with changing environmental drivers, like light and pCO₂ (Croteau et al. 2021; Joli et al. 2021; Kvernvik et al. 2021).

A prime challenge for photosynthetic cells is to maximize photon (i.e., energy) harvest, while simultaneously minimizing damage to the photosynthetic apparatus caused by excess light energy. To achieve this, photoacclimation processes take place

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Additional Supporting Information may be found in the online version of this article.

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on time scales from seconds to days (Moejes et al. 2017). For example, the potential for photon harvest can be adjusted by changing pigment composition, controlling antenna size, and connectivity, as well as by dissipation of excess energy as heat, so-called non-photochemical quenching (NPQ). In the next step, the "harvested" energy is converted to reductive energy, as electrons are stripped from water and forwarded into the photosynthetic electron transport chain. Here. an "overreduction" can occur, and potentially harmful excess electrons need to be dissipated through a number of biochemical processes. For example, electrons can be fed back to oxygen to form either water (via the so-called Mehler reaction) or reactive oxygen species for which a number of detoxification systems are available in the cell (Foyer et al. 2009).

To achieve balanced growth, not only energy capture, but also downstream cellular processes, e.g., nutrient assimilation, carbon fixation, and macromolecule synthesis need to be adjusted and tuned to maximize the efficiency of the photosynthetic light reactions (Huner et al. 1998). Biochemical mismatches or imbalances can cause cellular "stress" and are costly to deal with. They can arise, for instance, from the kinetics of RubisCO, the key enzyme for carbon fixation, which suffers from slow turnover rates, especially at cold temperatures (Young and Schmidt 2020). Psychrophilic diatoms can compensate for this by expressing higher amounts of this enzyme (Gerecht et al. 2019), but protein biosynthesis is expensive in terms of energy as well as nitrogen resources, inducing a tradeoff between resource expenditure and the maximization of C-fixation (Young et al. 2015). The ability of microalgae to cope with different kinds of photosynthetic stress has been shaped over evolutionary timescales by the selective pressures experienced in the habitats they occupy. Accordingly, ice-associated algae typically struggle to cope with the high light (HL) intensities and the variability that occurs in open ocean waters (e.g., Kvernvik et al. 2020, 2021; Croteau et al. 2021). Pelagic species, on the other hand, have developed plastic photosynthesis apparatuses and effective protective mechanisms to thrive under such light regimes (Lacour et al. 2018; Kvernvik et al. 2020; White et al. 2020).

Effects of pCO_2 in diatoms are known to be small compared to other taxa, highly species-specific (Gao and Campbell 2014), and often interact with other drivers, especially with light (Li and Campbell 2013; Hoppe et al. 2015; Seifert et al. 2020). More specifically, elevated pCO_2 often stimulates photosynthesis and growth under low light (LL), while it can decrease it under HL (Gao et al. 2019). CO₂ effects are commonly explained with reallocation of energy among competing processes, which is also why they are modulated by the availability of energy, i.e., light. Under pre-industrial and contemporary aquatic CO₂ concentrations, CO₂ is potentially limiting the rate of carbon fixation by RubisCO, and most algae, therefore, employ so-called CO₂concentrating mechanisms (CCMs) that utilize the chemiosmotic gradient of photosynthesis to accumulate CO₂ in the vicinity of RubisCO (Rokitta et al. 2022). Since CCMs consume less chemiosmotic energy under the higher pCO_2 expected for the future, cells can profit from more energy available to photosynthesis (Rokitta et al. 2012; Van de Waal et al. 2019). At the same time, more energy may be required to maintain pH homeostasis within the cell under acidified conditions, together causing an optimum curve of CO₂ response (Bach et al. 2011).

To investigate the responses of pelagic and ice-associated microalgae to the anticipated environmental changes, we chose two ecologically important species: Thalassiosira hyalina, a pelagic diatom that frequently dominates Arctic spring blooms, including ice-edge blooms (von Quillfeldt 2000; Hegseth and Sundfjord 2008), and has demonstrated high physiological plasticity toward changes in temperature, pCO₂, and light (Kvernvik et al. 2018; Wolf et al. 2018, 2019). Melosira arctica is known for forming long strands under multi-year sea-ice (Gran 1904; Syvertsen 1991; Melnikov et al. 2002) and can contribute disproportionally to carbon export to the deep ocean (Boetius et al. 2013), especially during ice-melt and breakup. To the best of our knowledge, no detailed ecophysiological investigations on M. arctica have been published to date, which is surprising considering its large contribution to under-ice biomass and carbon export, and thus its relevance for modeling efforts of ice-algal biomass production (e.g., Castellani et al. 2017; van Leeuwe et al. 2018).

In this study, physiological strategies of these two species were investigated through their acclimated responses to a matrix of two different light (LL = 50 and HL = $300 \,\mu mol$ photons $m^{-2} s^{-1}$) and pCO₂ regimes (380 and 1000 μ atm pCO₂). The LL level resembles an average irradiance in a wind-mixed pelagic habitat, or at the bottom of first-year ice and snow cover in late spring; the HL level was chosen to resemble a situation in a more stratified surface open ocean, or that during ice break-up and melt pond formation in summer (Nicolaus et al. 2012; Hill et al. 2018). The pCO₂ treatments resemble current and end-of-the-century pCO₂ conditions under a high-emission scenario. Based on their ecological niches, we expected T. hyalina to profit from HL and to experience only minor CO₂ effects, as observed in preceding studies (Wolf et al. 2018). M. arctica was expected to photosynthesize more efficiently under LL and to flexibly respond to changing carbonate chemistry, which tends to vary more strongly within dense aggregates under the ice compared to pelagic realm (McMinn et al. 2014). In addition to conventional analyses of growth, elemental composition and Chlorophyll a (Chl a) quota, we applied fastrepetition rate fluorometry (FRRF) and performed gas-flux measurements via membrane-inlet mass spectrometry (MIMS) to unravel photophysiological adjustments on a qualitative and quantitative level. The physiological strategies investigated here elucidate physiological mechanisms through which these species are adjusting to their environment, and thus allow a glimpse at how well each of them

may cope with the conditions to be expected in a future Arctic ocean.

Materials and methods

Culture conditions and experimental setup

T. hyalina and M. arctica cultures were obtained as singlecell isolates from the Kongsfjord, Svalbard (78°55'N, 11°56'E; isolated in 2014, strain WFB1) and from the central Arctic ocean (79° 33.23'N, 4° 50.29'W; isolated in 2015, strain PS93.1_030), respectively. Experiments were conducted in 2015/2016 as semi-continuous batch cultures in sterile 1-L glass bottles. In order to minimize changes in carbonate chemistry, to ensure nutrient-replete conditions throughout the experiments, and to remain close to natural population densities, cultures were diluted every 2-6 d, depending on cell density (T. hyalina max. 10,000 cells mL⁻¹, M. arctica max. 4000 cells mL⁻¹, i.e., up to 45 μ g Chl *a* L⁻¹). Sampling and dilution was conducted under sterile conditions, using a laminar flow hood. Cells were cultivated in 0.2-µm sterile-filtered Arctic seawater (salinity 33.5) enriched with macronutrients $(100 \ \mu \text{mol } \text{L}^{-1} \ \text{NO}_3^{-7}, \ 6.2 \ \ \mu \text{mol } \text{L}^{-1} \ \ \text{HPO}_4^{\ 2-}, \ 100 \ \ \mu \text{mol } \text{L}^{-1}$ SiOH₄), vitamins, and trace metals according to f/2 media (Guillard & Ryther, 1962). Experiments were performed in a temperature-controlled (3°C) room, with bottles immersed in water-filled aquaria for additional temperature stability. Continuous surveillance with a temperature logger (Almemo 2890, Ahlborn) ensured temperature stability at $3.4 \pm 0.15^{\circ}$ C.

Cells were grown under continuous light using daylight lamps (Biolux T8, 6500K, Osram). Irradiance levels were adjusted with a black mesh fabric and measured using a 4π sensor and data logger (ULM-500, Walz). Species were grown in a matrix of low and high light (LL vs. HL, 50 ± 1 vs. $300 \pm 4 \mu$ mol photons m⁻² s⁻¹) and low and high pCO₂ (380 vs. 1000 μ atm). The four resulting treatments (HL380, HL1000, LL380, LL1000) were tested with n = 4 biological replicates of each strain (except for *T. hyalina* LL where n = 3). In a side experiment with a reduced set of measured parameters, *M. arctica* was also cultured under the same conditions but at intermediate light (110 μ mol photons m⁻² s⁻¹). Prior to the experimental phase, all cultures were acclimated to treatment conditions for at least 6 d until growth rates stabilized (> 7 generations).

Carbonate chemistry

Cultures were continuously aerated (~170 mL min⁻¹) with humidified air of target pCO₂ delivered through sterile 0.2- μ m air-filters (Midisart 2000, Sartorius Stedim), provided by a custom-built gas mixing system (*see* Hoppe et al. 2015). Additionally, the growth medium was equilibrated (\geq 24 h) to the target pCO₂ at the respective temperature prior to inoculation. Stability of carbonate chemistry throughout the experiment was monitored by regular measurements of pH using a threepoint calibrated electrode and logger system (Aquatrode plus

Pt1000/pH 826 logger, Metrohm). The full cabonate system (Supplementay Table S1) was calculated for 3°C on the basis of pH and DIC values using the program CO2sys (Pierrot et al. 2006) with dissociation constants of carbonic acid by Mehrbach et al. (1973), refitted by Dickson and Millero (1987), as well as dissociation constants for sulfuric acid obtained by Dickson (1990). Samples of dissolved inorganic carbon (DIC) were taken on the day of final sampling, gently filtered through $0.2 - \mu m$ cellulose-acetate syringe-filters (Sartorius Stedim), and stored head-space free in 5-mL gastight borosilicate bottles in darkness at 3°C. DIC was measured colorimetrically in duplicates, according to the method of Stoll et al. (2001) with a OuaAAtro autoanalyzer (Seal Analytical), and corrected using Certified Reference Materials (A. Dickson, Scripps Institution of Oceanography). Cell-free control bottles with sterile medium were incubated alongside culture bottles, and provided baseline values of pH and DIC. Deviations in DIC from the incubations compared to these abiotic control bottles were typically $\leq 2\%$.

Growth, cellular composition, and production

Growth rates of *T. hyalina* were determined by monitoring cell densities using a Coulter Multisizer III cell counter (Beckman-Coulter). Since *M. arctica* cell-chains tend to entangle, this species' growth rates were based on measurements of Chl *a* concentration in cultures over time (details see ahead). In both cases, specific growth rates μ (d⁻¹) were calculated by the formula $\mu = (lnN_t - lnN_0)/\Delta t$, where N_t refers to cell density or Chl *a* concentration, and Δt to the passed time (in days) since the start of the incubation. For each culture, we used the mean value of exponential fits of all cell densities or Chl *a* concentrations of two or more consecutive dilution cycles (each spanning 3–6 d, example see Supporting Information Fig. S2).

Cell counts of *M. arctica* were performed for filtration timepoints on an inverse light microscope (Axio Observer, Zeiss) in 10-mL Uthermöhl-chambers. We hereby also distinguished between regular cells and cells of divergent morphology. The latter exhibited < 50% of regular size, had a more oval cell shape, and were often brighter in color (Supporting Information Fig. S1a), resembling descriptions of resting stages in Kaczmarska and Jahn (2006). Since scanning electron micrographs of the silica frustules supported this interpretation (Supporting Information Fig. S1b), they are in the following referred to as resting stages.

For growth rate estimation and at the end of each incubation, Chl *a* samples were filtered on glass fiber filters (GF/F, 0.7μ m nominal pore size; Whatman), shock-frozen in liquid nitrogen, and stored at -80°C. For analysis, filters were shredded in acetone (90%) with glass beads (0.5–1 mm diameter) in a homogenizer (Precellys Evolution, Bertin Technologies). After overnight extraction at 4°C, Chl *a* was measured fluorometrically (TD-700, Turner Designs), including an acidification step (1 M HCl) (Knap et al. 1996). For particulate organic carbon (POC) and nitrogen (PON), cells were filtered onto precombusted (15 h, 500°C) GF/F filters, stored at -20°C, and then dried at 60°C. POC analysis was performed using a gas chromatograph CHNS-O elemental analyzer (Euro EA 3000, HEKAtech).

POC production rates per cell were obtained by multiplication of the respective elemental quota with corresponding growth rates μ [pg POC cell⁻¹ d⁻¹]. In addition, Chl *a*-specific POC production was calculated for better comparison with O₂ and inorganic carbon fluxes by normalizing cellular POC production with the Chl *a* quota of the respective replicate, yielding [mg C (mg Chl *a*)⁻¹ h⁻¹].

Variable Chl a fluorescence

Chl a variable fluorescence and derived photosynthesis vs. irradiance (PI) curves of relative PSII electron transfer rates were measured using a Fast Ocean FRR fluorometer (FRRF, FastOcean PTX) combined with a FastAct system (Chelsea Technologies). Prior to the assays, samples of T. hyalina and M. arctica were darkacclimated inside the FastAct chamber for at least 10 min. The excitation wavelength of the fluorometer was set to 450 nm. The FRRF was used in single turnover mode, with a saturation phase comprising 100 flashlets of 2-µs pitch and a relaxation phase comprising 40 flashlets of a $50-\mu s$ pitch. The minimum (F0) and maximum chlorophyll fluorescence (Fm) was derived from iterative fitting of the induction phase (Kolber et al. 1998) and the relaxation phase (Oxborough et al. 2012). The maximum darkacclimated quantum yield was calculated after Krause and Weis (1991) as Fv/Fm = (Fm - F0)/Fm. Relative electron transfer rates through PSII in the light were calculated accordingly as rETR = (F'm - F'0)/F'm *PAR. The nonregulated Y(NO) and regulated Y(NPQ) non-photochemical quenching was determined after Klughammer and Schreiber (2008) at 400 μ mol photons $m^{-2} s^{-1}$ (except for *M. arctica* at LL, where it was measured at 570 μ mol photons m⁻² s⁻¹), which were saturating light levels for cultures in all treatments. Additional parameters like the reopening rate (τ), absorption cross section (i.e., antenna size, σ), and connectivity between reaction centers (ρ) were estimated in the dark-acclimated state following Kolber et al. (1998) and Oxborough et al. (2012).

Gas-exchange measurements

To assess the microalgae's cellular exchange of O_2 and inorganic carbon (C_i) in PI curves, MIMS was applied following the approach of Badger et al. (1994) with modifications described in Kottmeier et al. (2016*a*). This approach is based on continuous measurements of O_2 and CO_2 during consecutive light–dark intervals, and makes use of the chemical disequilibrium during light-dependent C_i uptake to differentiate between CO_2 and HCO_3^- fluxes across the plasmalemma. The custom-built MIMS consists of a lightadjustable and temperature-controlled 8-mL cuvette, which is coupled to a sector-field multicollector mass spectrometer (Isoprime, GV Instruments) via a gas-permeable 0.01-mm polytetrafluorethylene-membrane (Reichelt Chemietechnik). After passing through the membrane, O_2 and CO_2 are conducted into the mass spectrometer so that the photosynthetic and respiratory fluxes of these gases can be monitored during light and dark phases, respectively.

Algal cells were concentrated by gentle filtration using polycarbonate filters (3-µm pore-size; Merck Millipore) and washed two times with 10 mL DIC-free F/2 R medium (buffered to the respective culturing pH using 50 mM HEPES) to exchange the media. Afterward, the concentrated cell culture was transferred to the pre-cooled cuvette. NaHCO3solution was added to final concentrations of 2.1 mmol L^{-1} to establish normal DIC backgrounds, while dextran-bound sulfonamide was added to final concentrations of ~ 25 μ mol L⁻¹ to suppress potential activities of extracellular carbonic anhydrase, which ensures that a chemical disequilibrium can develop during illumination. After a preacclimation step (3-6 min of LL followed by 6 min of darkness), PI-curves were assessed in 6-min light : dark cycles with increasing light intensities. During the transfer into the assay buffer and cuvette, cells were kept at the incubation temperature $\pm 1^{\circ}$ C.

Net photosynthesis was determined from the steady-state O_2 fluxes at the end of the light phase, while respiratory O_2 uptake was measured during the subsequent dark phase. Photosynthetic and respiratory O₂ fluxes were converted to C_i fluxes by applying a photosynthetic quotient of 1.3 (Spilling et al. 2015) and a respiratory quotient of 1.0, respectively (Burkhardt et al. 2001; Rost et al. 2007). Cellular CO₂ uptake rates were derived from the rate of CO₂ depletion during steady-state photosynthesis at saturating light levels, which were further corrected for the simultaneous interconversion between CO₂ and HCO₃⁻ under the respective chemical disequilibrium. Cellular HCO₃⁻ uptake was calculated from a mass balance equation, i.e., the difference between net photosynthetic C_i uptake and net CO₂ uptake. All rates were normalized to Chl a concentration in the cuvette. For further details on the calculations of the photosynthetic fluxes, please refer to Badger et al. (1994) and Kottmeier et al. (2016a).

Estimation of photosynthetic parameters (PI-curves)

Both FRRF and MIMS assess photosynthesis, but while FRRF describes the relative electron transport from photosystem II, approximating gross photosynthesis, the MIMS measures O_2 fluxes, including the mitochondrial respiratory activity, hence net photosynthesis. To accommodate for the different measures of those methods and still use the same fit function for both, we modified a function from Platt et al. (1980) to allow for respiration (i.e., "negative" photosynthesis rates at light intensities below the compensation point) as well as an inhibition term to depict photophysiological stress at high irradiances:

$$V = \left(V_{\text{max;theor}} \times \left(1 - e^{\left(\frac{-\alpha \times I}{V_{\text{max;theor}}} \right)} \right) \times e^{\left(\frac{-\beta \times I}{V_{\text{max;theor}}} \right)} \right) + \text{Resp} \quad (1)$$

$$V_{\max} = V_{\max; \text{theor}} \times \left(\frac{\alpha}{\alpha + \beta}\right) \times \left(\frac{\beta}{\alpha + \beta}\right)^{\binom{\beta}{\alpha}}$$
(2)

$$I_k = \frac{V_{\max}}{\alpha} \tag{3}$$

$$I_{Inh} = \frac{V_{\text{max; theor}}}{\alpha} \times ln\left(\frac{\alpha+\beta}{\beta}\right) \tag{4}$$

In this photosynthesis function, I is the incident irradiance, while $V_{\text{max:theor}}$ represents a theoretical value of maximum photosynthesis, which would be approached in absence of light inhibition. V_{max} represents the apparent maximum photosynthesis rate, I_k the irradiance where the curve transitions from light limitation to light saturation, and α is the initial light-limited slope, also known as the maximum light-use efficiency. The parameter β represents the inhibition coefficient, I_{Inh} the irradiance where photoinhibition begins, and Resp represents the respiration, i.e., "negative" O₂ evolution in darkness (I = 0). After obtaining the parameters of the PI-curves, the equation can be solved for V = 0 in order to obtain the compensation point c, i.e., the irradiance where photosynthesis exactly balances respiration. Values for the photosynthesis at acclimation irradiance (Vaccli) were obtained by using the curve parameters and entering the irradiance values during acclimation in the LL and HL treatments, respectively. In order to differentiate between the photosynthetic parameters based on electron transport (FRRF) and O₂ evolution (MIMS), the respective parameters are distinguished by a subscript (i.e., α_e , $V_{max;e}$, $I_{k;e}$ and α_{O2} , $V_{max;O2}$, $I_{k;O2}$).

Data handling and statistics

All parameters were statistically tested using two-way-ANOVA (R Core Team, 2017, version 4.0.2) to identify the effects of light and CO₂ treatments, as well as their interaction (significance level ≤ 0.05). Data were tested for normality (Levene's test > 0.05) and homogeneity (Shapiro-Wilk test > 0.05), using log transformation of data where assumptions were violated. Only few parameters, derived from FRRF measurements in *M. arctica*, were not normally distributed (Ik: e, V_{max;e}, α_e, Fv/Fm, ρ). An overview of all statistical results is provided in Supplementary Table S3. Correlations between ratios of resting stages and growth or stoichiometric parameters of M. arctica were based on data in the LL and HL treatment, as well as the side experiment at an intermediate light level, and tested using simple linear models in the R-software. One biological replicate (B) of M. arctica in the HL-380 treatment was strongly dominated by resting stages (72%), while the other replicates had very low resting stage abundance (3%–9%). We removed this outlier from further analysis of cell properties to prevent a large distortion in this treatment.

Results and discussion

T. hyalina: opportunistic all-rounder with high plasticity *Physiological stability under both light regimes*

In T. hyalina, specific growth rates (μ) remained stable between 0.8 and 0.9 d^{-1} under all tested conditions (Fig. 1a), suggesting that fast growth has a high priority, which may be an advantageous competitive strategy of this species. Cellular POC quotas, however, were significantly higher under HL conditions (+ 65%, p = 0.002, Supplementary Table S2), and thus cellular POC production increased (+ 60%, p < 0.001, Fig. 1b). Slightly increased C : N ratios (from 5.1 to 5.2, p = 0.04, Supplementary Table S2) and stable BSi : POC ratios of 0.3 (p = 0.12; Fig. 1f) suggest that both, N-acquisition and silification, were upregulated alongside with C-fixation under HL to maintain biomass stoichiometry. This hints toward a controlled investment of additional energy into the buildup of cellular biomass under high irradiance, which has previously been observed in T. hyalina (Kvernvik et al. 2020) as well as in other pelagic diatom species (Thompson et al. 1991).

Chl *a* : POC ratios were reduced under HL (p < 0.001, Fig. 1e), which is a typical response to elevated light (Kvernvik et al. 2020; Lewis et al. 2018; MacIntyre et al. 2002). As the Chl a quota remained unaltered (p = 0.11, Fig. 1d), however, this photoacclimation was driven by changes in POC quotas. Increased light thus fueled higher biomass production per cell in T. hyalina, while most other cellular parameters remained stable despite a six-fold increase in irradiance. Not only Chl a quotas but also other fluorescence-derived parameters of rETR at PSII (i.e., Fv/Fm, $V_{max;e}$, $I_{k;e}$, τ , σ , ρ ; see Fig. 2 and Supplementary Table S2) remained remarkably unaltered in our treatments. Hence, acclimation did not occur on the level of light harvesting, nor did we observe signs of substantial light stress, as values of maximum quantum yield (Fv/Fm, Fig. 2a) remained unaffected. Similarly, non-photochemical quenching (Y(NPQ), Fig. 2b) remained unchanged, without upregulation of photoprotective energy dissipation in the antennae (i.e., the xanthophyll cycle).

Overall, our data for *T. hyalina* implies that larger energy input under HL are directly invested into C-fixation and biomass buildup without prominent signs of stress, which is in line with the previous finding for this species by Kvernvik et al. (2020). The stability in properties other than C-fixation, however, also means that photosynthetic efficiency was not further increased under LL. Such a "non-responsive" lightharvesting system will generate electrons according to incident light and thus yields opportunistically high energy input when the environment allows, without investing resources into costly photophysiological adjustments.

HL acclimation as a ground state, modulated by respiration

Along with Chl *a* quota, also antenna size (approximated as σ , Fig. 2c) remained stable across treatments, suggesting also constant light-harvesting efficiency of reaction centers and



Fig. 1. Stoichiometric and growth responses of *T. hyalina* in the four treatments (LL/HL = low/high light, $380/1000 = \mu \text{atm pCO}_2$). Points around boxplots show biological replicates. Significant differences between light or CO₂ treatments are marked in the title as follows: *** < 0.001; ** < 0.01; * < 0.05. Abbreviations: BSi: biogenic silica; Chl *a*: chlorophyll *a*; POC: particulate organic carbon.

thus making trends in Chl a-specific MIMS data comparable to RCII-specific FRRF data (despite not knowing specific RCII to Chl *a* ratios). As expected, net photosynthesis under acclimation irradiance, as measured by MIMS, was significantly higher under HL ($V_{accli;O2} + 80\%$; Fig. 1m, p < 0.001), and was remarkably well aligned with the observed increase in Chl a-specific POC production (+ 86%, Fig. 2g, Supplementary Table S2). This roughly doubled net photosynthesis under HL, however, does not match the almost tripled electron transport rate (Vaccli;e, Fig. 2d). Furthermore, while the maximal light-use efficiency of photochemical energy $(\alpha_e, Fig. 2e)$ remained stable across light treatments, the maximum light-use efficiency of net photosynthesis (α_{O2}) was significantly lower in HL compared to LL (p < 0.001, Fig. 2h). While the former observation could potentially also suggest a change in the ratio of PSII per Chl a, both of these apparent discrepancies between electron transport and production rates could be explained by the strongly stimulated dark respiration (+ 260%, p < 0.001; Fig. 2k), which likely consumed larger parts of the photosynthate generated in HL-acclimated cells. Accordingly, photosynthesis seems to become less efficient under HL conditions (trends in Chl a-specific POC production and Vaccli;O2 compared to trends in V_{accli;e}, Fig. 2d vs. 2g) and the initial light-limited slope of O_2 production is lowered (α_{O2} , Fig. 2h), since the relative amount of photosynthetic energy consumed by respiration is larger at the lower photosynthetic rates under low incident light. The strongly elevated dark respiration suggests that HL-acclimated cells had an increased metabolic demand, even independently of acute incident light. This may reflect higher maintenance costs of the higher biomass per cell itself (POC quota +-65%, Supplementary Table S2), higher energy-consuming detoxification mechanisms, or enhanced photosystem turnover in the HL treatment (Apel and Hirt 2004; Campbell and Serôdio 2020).

Under saturating light, maximum net O_2 production $(V_{max;O2})$ and C_i uptake $(V_{max;Ci})$ was similar in all treatments (~ 120–150 μ mol C (mg Chl *a*)⁻¹ h⁻¹, *p* = 0.07, Supplementary Table S2). While HL-acclimated cells nearly reached these maximum rates under acclimation conditions, LL-acclimated cells could only build up biomass at a lower C_i uptake rate. Accordingly, HL-acclimated cells exhibited light acclimation values below the acclimation irradiance ($I_{k;e} \sim 135 \,\mu$ mol photons m⁻² s⁻¹, $I_{k;O2} \sim 150-180 \,\mu$ mol photons m⁻² s⁻¹, Fig. 2f,i), i.e., experienced light saturation at growth conditions. I_k values of LL-acclimated cells, on the other hand, were far above acclimation irradiance ($I_{k;e} \sim 135 \,\mu$ mol photons m⁻² s⁻¹, $I_{k;O2}$ in LL ~ 95–130 μ mol photons m⁻² s⁻¹), i.e., they had grown in the light-limited part of the PI-curve. This explains why biomass production under HL was elevated (POC production,



Fig. 2. Responses in rate measurements of *T. hyalina* in the four treatments (LL/HL = low/high light, 380/1000 = μ atm pCO₂). Points around boxplots show replicate measurements. Significant differences between light or CO₂ treatments are marked in the title as follows: *** < 0.001; *< 0.05. Abbreviations: dark resp: dark respiration; Vaccli: e-transport rate/ O₂ production at culturing conditions; Y(NPQ): regulated photochemical quenching. Y(NPQ) was measured at 400 μ mol photons m⁻² s⁻², the unit for alpha_e is rETR (μ mol photons m⁻² s⁻¹)⁻¹, for alpha_{O2} μ mol O₂ h⁻¹ mg Chl a⁻¹ (μ mol photons m⁻² s⁻¹)⁻¹.

Fig. 1b), and underlines once more that even LL-acclimated cells are ready to use sudden increases in incident light for C-fixation.

All this advocates that HL conditions are rather the "comfort zone" of *T. hyalina*, and that this species' light-harvesting systems are maintained ready for high irradiances at all times. They thus avoid the danger of high-light stress, while accepting light limitation under LL conditions, under which cells coped well by lowering respiration as well as their carbon quota.

Flexible carbon acquisition helps minimizing CO₂ effects

Maximum net photosynthesis (V_{max;O2}) and total C_i uptake (V_{max;Ci}) remained similar in both pCO₂ treatments (Supplementary Table S2), but T. hyalina flexibly adjusted their relative uptake rates of CO₂ vs. HCO₃⁻. Under HL, where cells were acclimated to a higher carbon demand, HCO3⁻ uptake was significantly upregulated (i.e., relative CO₂ uptake decreased by one third, p < 0.001, Fig. 21), while under high CO₂ conditions, a significantly larger percentage of inorganic carbon was taken up as CO_2 (LL: from 60% to > 80%; HL: from 30% to > 60%; p < 0.001, Supplementary Table S2). CCM regulation was thus affected by both, light as well as pCO₂, which is in agreement with findings that CO₂ and HCO₃⁻ uptake systems are regulated as a function of carbon demand and supply (Rost et al. 2006b; Kottmeier et al. 2016a). In the coccolithophore Emiliania huxleyi, for instance, increased carbon demand under HL was met by stronger CCM activity based on HCO₃⁻ uptake, while acidified conditions inhibit HCO_3^- transport into the cell (Kottmeier et al. 2016b).

Regarding traits other than those related to the CCM, CO₂ effects were minor or completely absent. Respiration rates, for instance, were slightly lower under elevated pCO₂ in both light levels (p = 0.03. Fig. 2k), which could be attributed to a general down-regulation of mitochondrial activity under low pH (Strobel et al. 2013; Melzner et al. 2020). Furthermore, slightly higher irradiances were necessary to saturate photosynthesis at high CO₂ ($I_{k;O2}$, p = 0.037, Fig. 2i): lower respiration and easier carbon acquisition under high pCO₂ may be responsible for the slightly higher maximum O₂ production (V_{max:O2}, Supplementary Table S2), but these energy demands were met only at irradiances higher than acclimation conditions. Otherwise, no significant interactions between light and pCO₂ treatments were observed here, which is unlike some other diatoms, where higher pCO₂ stimulated growth and primary production under LL conditions or reduced photoinactivation at HL levels (Gao et al. 2012 and references within; Li and Campbell 2013). In an experiment testing the effect of sudden exposure to HL treatments (20 vs. 380 µmol photons $m^{-2} s^{-1}$) under similar pCO₂ treatments, effects of elevated CO₂ were found to be much stronger (Kvernvik et al. 2020). Thus, some of those stress responses may be transient and disappear during the acclimation process, or may only become apparent under these slightly larger spreads in light levels.

Overall, *T. hyalina* showed only weak acclimation responses to LL conditions, and instead remained "photosynthetically fit" and ready for coping with and profiting from sudden increases in irradiances. Although this strategy may not be maximizing energy output under LL conditions, it had no impact on their growth rate due to the high plasticity in carbon quota. Plasticity toward changing pCO_2 levels was also large, and the flexible CCM regulation according to their own demand and environmental supply may aid this species in achieving this robust acclimation with correspondingly little changes in traits over a wide range of environmental conditions.

Melosira arctica: low-light specialist with insufficient photoprotection

Resting stage formation under LL

M. arctica showed high variability among replicates in most parameters, especially in LL-acclimated cells, which correlates with a significant fraction of resting stages (up to 30%, Supporting Information Fig. S1c). While HL cultures contained up to 6% resting stages, a side experiment under intermediate light $(110 \,\mu \text{mol photons m}^{-2} \text{ s}^{-1})$ contained an intermediate fraction (3%-13%, Supporting Information Fig. S1c), suggesting a light dependence of resting stage formation. One of the eight HL replicate bottles contained 70% of resting stages and was treated as an outlier. Notably, growth rate and the percentage of resting stages showed a weakly positive correlation (Supporting Information Fig. S1d; p = 0.05), contradicting the notion that these cells are mostly inactive (McQuoid and Hobson 1996). Variability in cell quotas (Chl a, POC, PON) were overall higher in treatments with high number of resting stage (i.e., the LL treatments), while variability for C : N and Chl a : POC ratios was consistently low in all treatments (Fig. 3, Supplementary Table S2). BSi : POC ratios showed a higher degree of silification in cultures with higher number of resting stages (Supporting Information Fig. S1e, p = 0.02), which is in line with resting spores often having thicker cell walls than vegetative cells (McQuoid and Hobson 1996; Ellegaard and Ribeiro 2018).

High trait variability and little gain from higher light

In contrast to the pelagic T. hyalina, the ice-associated M. arctica significantly decreased its growth rate under HL conditions from 0.6 to 0.4 d⁻¹ (p < 0.001, Fig. 3a), implying a reduced competitiveness in terms of proliferation. Average cell quotas and POC production rates showed larger variability between the LL replicates that contained a high fraction of resting stages, and therefore no significant light effects could be determined (Fig. 2b, Supplementary Table S2). This is in contrast to T. hyalina, which strongly increased its POC production under HL. Biomass stoichiometry of M. arctica was also more variable than in T. hyalina, e.g., C : N ratios were significantly elevated under HL (p < 0.001, Supplementary Table S2). This was due to a decrease in N quota rather than an increase in C quota (Supplementary Table S2), suggesting N-assimilation and related processes may have been hampered. M. arctica exhibited on average five-fold higher BSi : POC ratios than T. hyalina (1.8 vs. 0.4; Fig. 3f). Slower growth often coincides with stronger silification in diatoms



Fig. 3. Stochiometric and growth responses of *M. arctica* in the four treatments (LL/HL = low/high light, $380/1000 = \mu \text{atm pCO}_2$). Points around boxplots show individual replicates. Significant differences between light or CO₂ treatments are marked in the title as follows: *** < 0.001; ** < 0.01; * < 0.05. Abbreviations: BSi: biogenic silica; Chl *a*: chlorophyll *a*; POC: particulate organic carbon.

(Martin-Jézéquel et al. 2000), which could indicate that grazing defense is prioritized over fast proliferation in *M. arctica*. Furthermore, BSi : POC content was increased under LL (+ 33% BSi : POC, p = 0.007, Supplementary Table S2), which is likely attributable to the high amount of resting stages.

Chl *a* : POC ratios were less variable than respective quotas, and were significantly reduced by ~ 60% in HL-acclimated cells (p < 0.001, Fig. 3e). While a similar reduction in *T.hyalina* (~ 50% Chl *a* : POC, Fig. 1e) was driven by an increase of POC quota, in *M. arctica*, this was due to reduced pigment content rather than stimulation of biomass production (Fig. 3d, Supplementary Table S2), indicating the cells' attempt to reduce energy input. Such a typical photoacclimation response (Brunet et al. 2011) is well in line with other studies on ice algae (Hegseth 1992) and indicates that HL acclimation in *M. arctica* is also more focused on preventing photodamage than exploiting elevated light availability.

Electron transport rates under acclimation light intensity (V_{acclie}) approximately doubled in response to HL (rising from 20 to 50; p < 0.001, Fig. 4d). Notably, also Chl *a*-specific net photosynthesis ($V_{accliO2}$) increased to a similar degree (Fig. 4g; reaching rates close to those of *T. hyalina*, Fig. 2g), and also Chl *a*-specific POC production increased accordingly (Fig. 3c). The similar effect sizes in these three parameters are remarkable, since values originate from different measurement techniques, which assess vastly different timescales (seconds to

days). When normalizing these parameters to POC, however, i.e., taking the decrease in Chl *a* quota into account, these effects disappeared, showing that there was no effect of light on biomass production (POC production per cell and per POC, Supplementary Table S2). This is also in line with the variable but substantially increased dark respiration under HL (+ 72% p = 0.07, Fig. 4k). Hence, the HL acclimation of *M. arctica* involves throttling its photosynthetic performance via decreased Chl *a* content rather than exploiting increased energy availability.

High-light stress in spite of acclimation

In contrast to *T. hyalina*, and somewhat counterintuitively, maximum electron transport rates were significantly lower under HL than under LL ($V_{max;e} - 61\%$, p < 0.001, Supplementary Table S2). This was also mirrored in lower light acclimation indices under HL than under LL ($I_{k;e}$, p < 0.001, Fig. 4f), which suggests that HL-acclimated cells capped their light harvest far below treatment conditions, at around 100 μ mol photons m⁻² s⁻¹. LL-acclimated *M. arctica* cells thus continued to harvest incoming photons at irradiances far beyond their acclimation light, while in HL-acclimated cells, Chl *a* quota where drastically lowered in order to prevent excessive excitation pressure and oxidative stress. This is in line with the significantly downregulated absorption cross-section per RCII in the HL treatment (antenna size, p < 0.001, Fig. 4c).



Fig. 4. Responses in rate measurements of *M. arctica* in the four treatments (LL/HL = low/high light, 380/1000 = μ atm pCO₂). Points around boxplots show individual replicates. Significant differences between light or CO₂ treatments are marked in the title as follows: *** < 0.001; ** < 0.01; * < 0.05; ^ < 0.1. Abbreviations: dark resp.: dark respiration; Vaccli: e-transport rate/ O₂ production at culturing conditions; Y(NPQ): regulated photochemical quenching. Y(NPQ) was measured at 400- μ mol photons m⁻² s⁻² (in LL treatments at 570- μ mol photons m⁻² s⁻²), the unit for α_e is rETR (μ mol photons m⁻² s⁻¹)⁻¹.

Accordingly, the maximum net photosynthesis ($V_{max;O2}$, Supplementary Table S2) remained unchanged by light treatment, since this parameter was normalized to Chl *a*. Taking the

lower Chl *a* content into account by normalizing rates to cell or POC, also net photosynthesis followed this trend of lower maximal values in HL acclimation (data not shown). In other words, HL-acclimated *M. arctica* set an upper limit to its maximum energy input by employing smaller antenna complexes and lower Chl *a* content, while LL-acclimated cells maintained a highly efficient harvesting machinery, aiming at maximizing photosynthetic yield at the cost of a higher risk for photodamage.

Evidence of M. arctica being exposed to high-light stress and possible photodamage even in the HL-acclimated state can be found in the potential photochemical efficiency of dark-acclimated cells (Fv/Fm), which was significantly decreased in HL compared to LL conditions (p < 0.001, Fig. 4a). One protective way to channel surplus light energy away from photochemistry, and thus to avoid potential damage, is heat dissipation, or non-photochemical quenching. This can occur in a regulated way, e.g., via the xanthophyll cycle (Y(NPQ)) or in a non-regulated way, e.g., through passive dissipation, mainly due to closed PSII reaction centers (Y(NO), Klughammer and Schreiber 2008). For M. arctica, regulated NPQ was only slightly higher in HL conditions compared to LL (Y(NPQ), p = 0.02, Fig. 4b), but approximately twice as high as in *T. hyalina* (Fig. 2b). This suggests that regulated heat dissipation played a role in its acclimation response and thus in limiting the maximal electron transport during sudden light increases, as seen in other ice algae (Croteau et al. 2021; Kvernvik et al. 2020).

Non-regulated NPQ (Y(NO), Supplementary Table S2), on the other hand, remained stable, suggesting that sudden highlight exposure did not block reaction centers, and potential photodamage may instead occur further down in the electron transport chain. Concomitantly, turnover time of PSII (reopening rate τ) was increased in HL- compared to LLacclimated cells (p = 0.03, Fig. 4j). Thus, acclimation to HL did not vield a more efficient downstream machinery, as could be expected in response to increased energy input (Barlow et al. 1988), but rather became more inefficient, e.g., due to substrate limitations during C-fixation (see ahead). Insufficient capacities of the downstream processes have the potential for detrimental overreduction in the photosynthetic electron transport chain, e.g., resulting in cytotoxic ROS production (Apel and Hirt 2004). The lack of photophysiological protection against light variability under LL conditions, along with the apparent light stress in HL even after acclimation, leaves the impression of a species neither well adapted to fluctuating, nor to stable high-light regimes.

No pCO₂ effects in spite of inflexible carbon acquisition

Unlike other ice-algal species (Kvernvik et al. 2020), *M. arctica* showed no significant responses to ocean acidification, or any interactive effects of pCO_2 with light (Supplementary Table S3). This supports findings by Torstensson et al. (2021), who found assemblages of sea-ice algae dominated by *M. arctica* to be insensitive toward pCO_2 perturbations. In line with this, maximum carbon uptake rates per Chl *a* were stable in all treatments (~ 90 μ mol (mg Chl *a*)⁻¹ h⁻¹, Supplementary

Table S2) and *M. arctica* relied mainly on CO_2 uptake instead of adjusting its preferred carbon species (Fig. 41). Such an unaltered C_i uptake behavior may also limit the ability to match higher carbon demands under increasing light (Rost et al. 2006*a*; Hoppe et al. 2015). Thus, it may be the origin of a slower photosynthetic downstream machinery, i.e., the increased reopening rate (and decreased Fv/Fm) under HL. This was in contrast to *T. hyalina*, which changed its carbon source and exhibited higher C_i uptake rates, suggesting an overall less efficient and dynamic regulation of photosynthesis in *M. arctica*. Another reason for slow downstream machinery under HL might be a rate limitation imposed by slow Rubisco kinetics at low temperatures (*see* Young et al. 2015). This is unlikely, however, as the culturing temperature was comparably high for an ice-algae like *M. arctica*.

Overall, *M. arctica* was physiologically less performant under HL than under LL, exhibiting decreased growth and signs of photophysiological stress. In contrast to *T. hyalina*, considerable acclimation to the different light levels took place, mainly through restructuring of light-harvesting antenna to maximize the energy input under LL, while minimizing it under HL conditions.

Ecological implications

T. hyalina exhibited the typical physiology of a species relying on fast proliferation, and seemed especially well equipped for high and variable irradiances. These properties are characteristic for pelagic "boom and bust" species (Assmy and Smetacek 2009) and enable them to occupy their ecological niche: similar to other ecological r-strategists often found in unstable high-nutrient environments (Prosser et al. 2007; Southwood et al. 1974), *T. hyalina* prioritizes fast growth as soon as enough light is available in early spring and while nutrients are still plentiful to develop large bloom events (von Quillfeldt 2000).

Furthermore, acclimation to LL appeared minimal, which is a good strategy when thriving in the upper mixed water column, where light regimes are rapidly fluctuating, and damage by sudden increases in irradiance is likely. Accordingly, T. hyalina keeps its photosynthetic apparatus extremely stable under different conditions, but plastic enough to cope efficiently with fluctuations and to profit from favorable conditions. In line with this, (Kvernvik et al. 2020) could show that T. hyalina uses potent protective mechanisms (e.g., xanthophyll cycle and oxidative stress defense) to overcome sudden increases in irradiance within less than 48 h without experiencing disadvantages. Furthermore, T. hyalina's ability to adjust the fraction of HCO_3^- and CO_2 uptake (i.e., CCM regulation) allows for efficient C_i fixation, independently of CO₂ supply and its internal C-demands. This may also help diatoms like T. hyalina to thrive in dynamic mixed layers by fulfilling quickly changing carbon demands and thus preventing photodamage (Hoppe et al. 2015). Rising temperatures in

the water column may further fuel the fast growth rates of this species, as it has been shown to have a thermal optimum at temperatures above common Arctic temperatures (Wolf et al. 2018).

T. hyalina's strategy can be summarized as an opportunistic approach to the tradeoffs of growing in the sunlit mixed layer: rather than investing into fine-tuned photoacclimation to LL conditions, which entails the danger of photodamage during suddenly peaking irradiances, *T. hyalina* ensures stable growth, but otherwise endures phases of light limitation (e.g., by lowering its carbon quota) while waiting for better conditions, e. g., for a shoaling of the mixed layer.

M. arctica, on the other hand, showed rather opposite strategies and preferences. Growth rates were generally lower than those of T. hyalina and decreased even further in HL conditions. Acclimation to high irradiance involved mainly photoprotection rather than exploitation of the higher energy input, and nevertheless, cells showed signs of stress. In an LLacclimated state, M. arctica possessed an efficient lightharvesting system, but showed little tolerance toward light fluctuations. Slow C_i uptake and growth require less investment into CCM regulation, but this adaptation may limit capacities to balance high energy inputs. *M. arctica* is typically not exposed to high or strongly fluctuating light in its sub-ice habitat, where it forms long, mucilaginous threads that grow into the underlying water column (Melnikov et al. 2002). Therefore, its physiology is tuned for optimal light harvest under stable LL conditions, often even far below the here applied irradiance (Castellani et al. 2021). High or fluctuating light situations would mainly be found in a breakup or meltpond scenario, when the ice disintegrates and the heavy algae mats sink quickly, leading to massive, pulsed carbon export (Boetius et al. 2013).

These ecological considerations may also explain the conspicuous formation of resting stages. Traditionally, resting stage formation is thought to be initiated by nutrient-limitation, high cell densities or other kinds of non-favorable conditions (McQuoid and Hobson 1996; Pelusi et al. 2020). In the case of M. arctica, however, resting stage formation was found in their rather favored LL treatment, while under higher, more stressful light conditions they were nearly absent (Supporting Information Fig. S1c). As high irradiance in the sea-ice habitat is an indicator of ice-breakup, however, production of resting stages under LL conditions seems more reasonable: it allows their preservation in multi-year ice or ridges, where M. arctica is often found (Syvertsen 1991; Poulin et al. 2014). Another unexpected observation was that resting stages differed from vegetative cells mainly in their size (i.e., cell quota) and their BSi : POC ratio, but not in terms of other elemental ratios and even growth rates. Although SEM images showed typical morphology of resting stages (Kaczmarska and Jahn 2006; Supporting Information Fig. S1), this poses the question whether these cells were indeed resting, or whether the difference between a vegetative and a resting state may be rather gradual. According to our data, their main adaptation may be the stronger grazing protection, which would primarily reduce loss rates once the cells are in an environment not permitting growth (e.g., darkness). To which extend these cells actively contribute to overall growth rates remains yet to be understood.

Low growth rates, extremely strong silicification, and high percentages of resting stages under favorable conditions, all of this suggests that *M. arctica* follows a k-like strategy of slow, but optimized growth and persistence rather than one of fast, opportunistic proliferation. In comparison to pelagic species, ice-associated species have on the one hand increased encounter rates with grazers (Kohlbach et al. 2016), but on the other hand smaller loss terms due to sinking. Strong silicification can thus be used as a grazer defense without the tradeoff of fast sinking.

In conclusion, T. hyalina showed characteristics that explain its success as a dominant spring bloom species in northern waters. As they cope well with ocean acidification and strongly benefit from higher irradiance, this species is likely to profit from the larger areas of open, sunlit ocean as the sea ice retreats. Their final success, however, will also strongly depend on other variables such as nutrient conditions, temperature, and especially on the respective response of their direct competitors and grazers. In contrast, M. arctica seemed to require low irradiances and stable light conditions. Both, their photophysiology as well as their ecological strategy of forming thick, grazer-protected mats under the ice, seem disadvantageous in view of anticipated changes in the Arctic. Although M. arctica is traditionally described as a dweller in multi-year ice (Syvertsen 1991), there are also frequent observations in first-year and new ice, and even floating aggregates at the halocline and between ice floes (Gran 1904; Poulin et al. 2014). Nevertheless, in view of its specialized physiology, it seems likely that the relevance of this currently important Arctic biomass producer will decline alongside the retreating sea-ice, making it one of the "losers" of climate change.

Data availability statement

The data supporting the findings of this study are openly available in the Pangea Database https://www.pangaea.de/.

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Conflict of Interest

The autors declare no conflit of interest.

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