



## Review article

Photosynthetic adaptation and multicellularity in the Antarctic psychrophile, *Chlamydomonas priscuii*

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## ABSTRACT

Acclimation, adaptation and survival in persistent cold polar environments are complex phenomena associated with myriad molecular, biochemical and physiological mechanisms. The psychrophile, *Chlamydomonas priscuii*, is endemic to Lake Bonney, Antarctica. Adaptation to its extreme polar environment includes homeoviscous adaptation of membranes, maintenance of energy balance through photostasis and surface area to volume ratio. In addition to these mechanisms, this psychrophile can exist in culture as motile, single cells or as immobile, multicellular palmelloids. Comparative biochemical, physiological, microscopic and spectroscopic analyses of purified single cells and palmelloids indicate that the conversion of single cells to multicellular palmelloids alters the composition and organization of the photosynthetic apparatus. This enhances photoprotection of the photosynthetic apparatus from light and low temperature stress by minimizing potential cellular energy imbalances and safely dissipating excessive excitation energy by nonphotochemical quenching mechanisms. In addition to decreased susceptibility to predation, enhanced photoprotection from photoinhibition associated with palmelloid formation may be a complementary, selective, evolutionary advantage for the induction of multicellularity in green algae.

## 1. Introduction

It has been estimated that the North Atlantic, above 30°N, and the Southern Ocean, below 50°S, account for 32 % of the global anthropogenic CO<sub>2</sub> inventory which makes these polar aquatic basins major sinks for the sequestration of atmospheric CO<sub>2</sub> [1]. Consequently, they are important components that govern global carbon balance and moderate the impact of global climate change on the biosphere.

These vast cold, Arctic and Antarctic ecosystems sequester at least 32 % of the total global CO<sub>2</sub> annually in the form of “blue carbon” [2,3]. As illustrated in Fig. 1, the aquatic photosynthetic productivity, estimated by chlorophyll *a* (Chla) concentration, is highest globally in the North Pacific, the North Atlantic and the Southern Ocean and is an order of magnitude more productive than the tropical regions of the Pacific and the Atlantic Oceans (Fig. 1). It is thought that the immense areas of these polar aquatic ecosystems more than compensates for the low species diversity with respect to their contributions as sinks for global

photosynthetic CO<sub>2</sub> assimilation compared to the tropical terrestrial ecosystems [2–5]. In addition to the polar Atlantic and Southern Oceans, permanently ice-covered lakes are characteristic of continental Antarctica [3–6]. Notwithstanding their important contributions to global CO<sub>2</sub> sequestration, these micro-organisms that dominate the polar oceans and permanently ice-covered lakes present in continental Antarctic represent some of the least studied organisms on our planet [6–10].

The ecosystem services provided by the polar photoautotrophs are a consequence of the light- and temperature-dependent process of photosynthesis which is essential for the maintenance and regulation of energy flow for the whole planet. This is achieved through the transformation of energy in the form of sunlight into biomass energy. This photosynthetic ecosystem service is reflected in the distribution of biomass on Earth. Plants and micro-organisms represent about 82 % and 17 % respectively of Earth's total biomass [11]. The latter includes the photosynthetic micro-organisms such as algae and cyanobacteria that

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dominate our vast oceans, lakes, rivers and streams. Energy from the Sun through the process of photosynthesis generated the 550 gigatons (Gt) of carbon biomass present on Earth today over millions of years of evolution.

Despite climate change and global warming, our planet remains a rather cold place [12]. Oceans that cover approximately 70 % of the Earth's surface have characteristic temperatures of  $\leq 5^{\circ}\text{C}$ . The polar Arctic and Antarctic regions constitute approximately 20 % of the surface area. Finally, the mountainous, alpine regions of the world constitute about another 5 % of the Earth's surface [12]. However, the study of non-model terrestrial and aquatic life endemic to polar regions provides an exceptional opportunity to understand the basis of novel adaptations to these extreme environments, which are not necessarily accessible through research focussed on model systems [14–31].

## 2. Photosynthetic adaptation to extreme polar environments

### 2.1. Cell membranes and cold adaptation

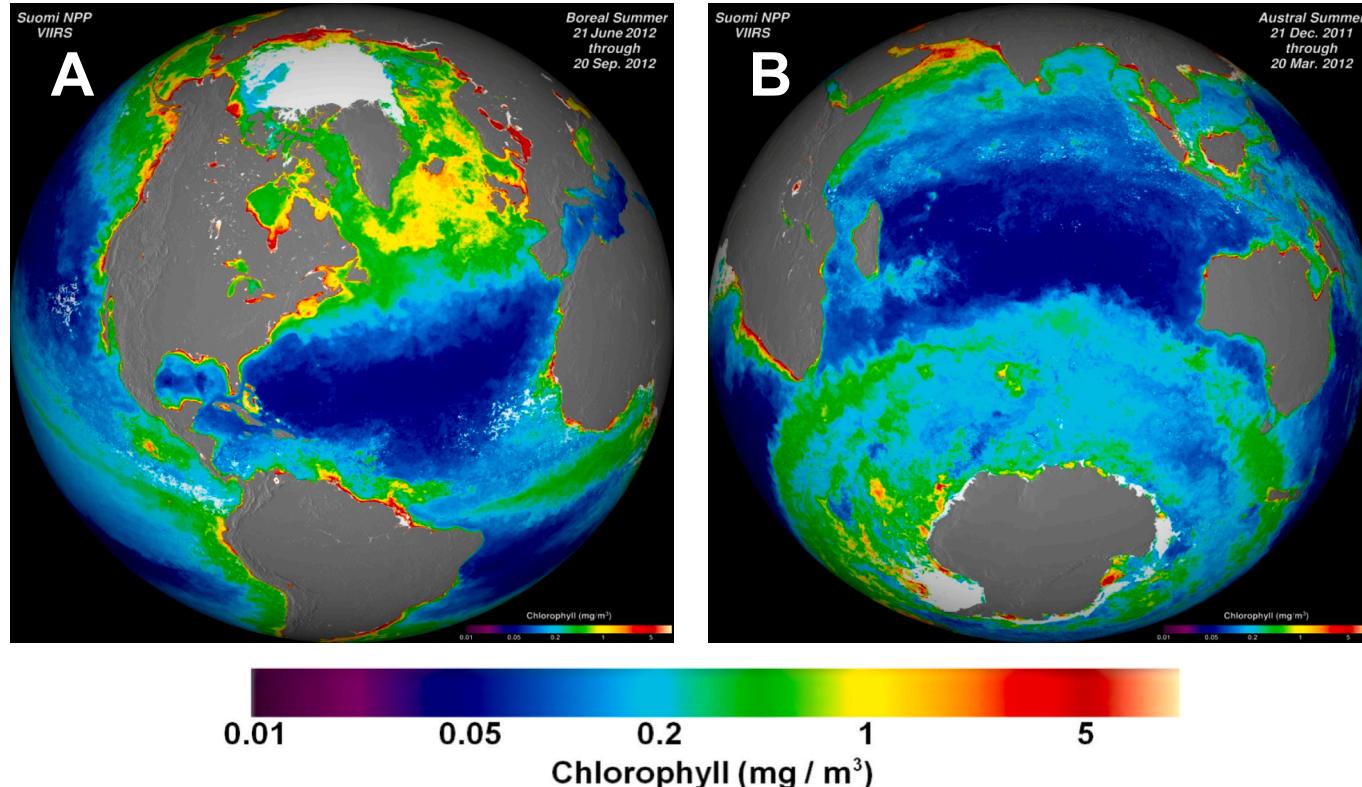
Poikilotherms, including photoautotrophs (plants, green algae, cyanobacteria) as well as heterotrophs (e.g. bacteria, fungi, reptiles, and fish) are unable to control their cellular temperature which is strictly dependent upon the vagaries of the ambient environment. Cellular function and compartmentation are strictly dependent upon the presence of cell membranes. However, changes in temperature can severely impact membrane stability and function [5,29–32]. To ensure the stability and function of these cellular membranes in response to cold temperature, poikilothermic organisms must adjust membrane lipid and fatty acid composition to maintain membrane fluidity in the cold. This response is called homeoviscous adaptation which, in plants, algae and cyanobacteria, generally has been focussed on modulation of the membrane lipid and the acyl chain composition in response to cold stress

and cold acclimation [32–44,46,47].

The Antarctic psychrophile, *C. priscuii*, cultured at its optimum low growth temperature ( $8^{\circ}\text{C}$ ) exhibits homeoviscous adaptation when compared to the model mesophile cultured at its optimum growth temperature ( $29^{\circ}\text{C}$ ); indeed, there is a 1.44 higher unsaturation index for *C. priscuii* (2.74) compared to *C. reinhardtii* (1.90) [48]. This probably contributes to a greater membrane fluidity at low temperature in the psychrophile than in the mesophile. Thylakoid membrane stability as a function of temperature can be assessed by monitoring the change in background Chla fluorescence yield ( $F_0$ ) [49–52]. The critical temperature for a change in membrane stability in *C. priscuii* ( $40^{\circ}\text{C}$ ), was 10° lower than that for *C. reinhardtii* ( $50^{\circ}\text{C}$ ) which is consistent with the observed differences in unsaturation index [48].

### 2.2. Photostasis and cold adaptation

Non-model organisms from the extreme polar habitats represent successful adaptation to life at the edge [5,28,52]. With respect to temperature adaptation, organisms are typically classified as either eurythermic or stenothermic. The former reflects organisms that can tolerate a wide temperature range whereas the latter can tolerate a much narrower range [53]. The Antarctic green alga, *C. priscuii*, is unique compared to the model green alga, *C. reinhardtii*, because the former is an extremophile that is adapted to persistent and consistent low temperature, high salt and low daily light levels, combined with extreme annual variation in photoperiod [5,28]. In addition, *C. priscuii* is a photopsychrophile (photosynthetic psychrophile) [53] that is obligately adapted to low temperature ( $0\text{--}12^{\circ}\text{C}$ ) and will die at temperatures  $\geq 20^{\circ}\text{C}$  [5,54]. Photopsychrophiles are photoautotrophs which distinguishes them from all other heterotrophic psychrophiles including fungi [55,56] and bacteria [30,57–59]. However, not all photosynthetic polar micro-organisms are stenothermic psychrophiles. Many polar microbes



**Fig. 1.** Global aquatic chlorophyll concentrations. A - Northern hemisphere; B - Southern hemisphere.  
[https://www.nasa.gov/mission\\_pages/NPP/multimedia/gallery/V20121732012264-NPP.html#.ZHYfiHbML3g](https://www.nasa.gov/mission_pages/NPP/multimedia/gallery/V20121732012264-NPP.html#.ZHYfiHbML3g).  
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are psychrotolerant eurytherms and, thus, abide a much wider range of temperatures than stenothermic psychrophiles [61,62].

For both psychrophilic or psychrotolerant organisms, the very rapid, temperature-insensitive photobiophysical processes of light absorption and energy transfer associated with the light harvesting complexes coupled to the photochemistry associated with PSI and PSII reaction centres are integrated with the much slower and highly temperature-dependent, redox reactions associated with photosynthetic electron transport and the soluble stromal enzymes involved in CO<sub>2</sub> assimilation through the Calvin-Benson-Bassham Cycle (CBB). The CBB consumes ATP and photosynthetic reductants to reduce C, N and S required for cellular maintenance, growth, development and reproduction. A major challenge for all polar photoautotrophs is to balance the flow of energy from the source, in the form of absorbed photons involved in photosynthetic photochemistry, to the metabolic sinks. These sinks consume the energy in the biochemical, enzyme-catalyzed reactions generating metabolites needed for growth, development, and reproduction in a thermodynamically challenging habitat [63–69]. The maintenance of cellular energy balance in photosynthetic organisms is referred to as photostasis [64,65,70,71] and is required to minimize the production of reactive oxygen species (ROS) and reduce the potential for oxidative damage. However, ROS are two-faced Janus. Not only does their accumulation during stress result in cellular damage, but ROS are also important stress-related signalling molecules [72–74].

As summarized in Table 1, myriad mechanisms have evolved in terrestrial plants and aquatic algae in response to cold adaptation and the attainment of photostasis. These include regulation of the absorption of light through modulation of the *physical size* of the light harvesting complexes associated with the photosystems via redox regulation of transcription of nuclear *Lhcb* genes [27,75–79], the transition between typical linear electron flow (LEF) through PSII and PSI versus PSI cyclic electron flow (CEF) as well as the reduction of O<sub>2</sub> via the plastid terminal oxidase (PTOX) in the process of chlororespiration [21,80–95]. In addition, state transitions, governed by redox regulated phosphorylation-dephosphorylation of PSII light harvesting polypeptides, control the energy distribution between PSII and PSI to fine tune photosynthesis and are considered essential in photoprotection of the photosynthetic apparatus from excessive excitation energy (EEE) [89] in plants and green algae [90–95].

Additional photoprotective mechanisms have evolved that govern the partitioning of energy between photochemical versus non-photochemical quenching (NPQ) to dissipate EEE safely as heat via photosystem II (PSII) light harvesting antennae and the xanthophyll cycle [96–103]. The induction of xanthophyll cycle activity affects the *functional size* of the photosystem light harvesting complexes through the conversion of the light harvesting carotenoid violaxanthin to the energy dissipating carotenoids antheraxanthin and zeaxanthin which reduces the photosynthetic efficiency of light absorption and ensures the safe dissipation of EEE as heat through NPQ [65,104–106,108–110]. Although the cost of these photoprotective mechanisms is a general decrease in photosynthetic light absorption efficiency, the benefits include enhanced probability of survival associated with short-term acclimation and long-term adaptation of photosynthetic organisms to EEE to reduce sensitivity to photoinhibition induced upon exposure to various stress conditions [111–117].

The regulation of photosynthetic electron flow, state transitions and energy partitioning contribute to the attainment of photostasis by protecting the photosynthetic apparatus from potential photodamage due to exposure to EEE in extreme environments. This includes restructuring and reorganizing the photosynthetic apparatus which is governed by redox retrograde signalling to regulate the expression of photosynthetic nuclear genes in plants and algae [118–125]. This usually affects changes in phenotype as reflected in major changes in pigmentation in algae such as *C. vulgaris* [25,26], *C. reinhardtii* [84,110], *D. tertiolecta* [116–118], *D. salina* [27,119,120] as well as the filamentous cyanobacterium, *Plectonema boryanum* [121,122]. The phenotypic changes

**Table 1**

Proposed cellular/molecular mechanisms of cold adaptation in psychrophilic (cold-adapted) microorganisms.

	Proposed cold-adaptation functions	References
Proteins/enzymes	Maintenance of adequate metabolic flux	Szyszka-Mroz et al. [129] Feller et al. [185] Devos et al. [186] Miyazaki et al. [187] Feller and Gerday [188] Casanueva et al. [183] Paredes et al. [189] Siddiqui et al. [29]
Lipids, membrane fatty acids, unsaturated fatty acids long-chain polyunsaturated fatty acids (LC-PUFA)	Maintenance of membrane fluidity	Russell and Fukunaga [191] Morgan-Kiss et al. [48] Mock and Kroon [192] Králová [198] Hassan et al. [194] Szyszka-Mroz et al. [128] Cvetkovska et al. [158] Morgan et al. [48] Mock and Valentín [196] Mock and Hock [195] Morgan-Kiss et al. [5] Zhang et al. [194] Chattopadhyay et al. [200] Jagannadham et al. [198] Leya et al. [199] Dieser et al. [201] Shen et al. [202] Ghobakhloo et al. [203] Fonseca et al. [204] Goordial et al. [205] Raymond et al. [206] Gilbert et al. [207] Muryoi et al. [208] Raymond et al. [209] Bar Dolev et al. [210] Tosco et al. [211] Yoshimune et al. [212] Romero-Romero et al. [213] Collins and Gerday [214] Cvetkovska et al. [54]
Photosynthetic apparatus	Maintenance of light harvesting capacity and photosynthetic electron flow	
Pigments: carotenoids	Modulation of membrane fluidity Photoprotection Antioxidants	
Compatible solutes	Osmoprotection against freezing induced osmotic stress, desiccation protection against freezing induced desiccation, cryopreservation	
Antifreeze proteins Ice-nucleating proteins	Ice growth and recrystallisation inhibition Extracellular ice crystal nucleation	
Chaperones	Maintenance of protein folding and stability Destabilisation of RNA/DNA secondary structures	

reflect photosynthetic acclimation characterized by decreases in total Chl, increases in the ratio of Chla/b and lower levels of light harvesting or phycobilisome polypeptides resulting in a decrease in the efficiency of photosynthetic light absorption.

A novel feature of the Antarctic psychrophile, *Chlamydomonas prisculii*, is that it exhibits minimal phenotypic changes in pigmentation in response changes in light levels or exposure to low temperature [5,11,123] compared to model plants such as *Arabidopsis thaliana* [124] and mesophilic green algae such as *C. reinhardtii* [84,110], *C. vulgaris* [25,26], *D. tertiolecta* [124–126] and *D. salina* [27,127,128]. Furthermore, *C. prisculii* is deficient in the capacity to perform state transitions [5,11,125,126]. This deficiency may be a consequence of an altered thylakoid protein kinase required for thylakoid protein phosphorylation [127–130] which preferentially phosphorylates a subunit of a PSI-Cytb6/f supercomplex which governs the high rates of PSI cyclic electron flow in *C. prisculii*. This is not observed in other model plant or green algal systems [135–137]; however, a similar deficiency in state transitions was observed for the Arctic psychrophile, *Chlamydomonas malina*. Notably, the Antarctic psychrophile, *Chlamydomonas* sp. ICE-MVD, was able to perform classic state transitions like the mesophile *C. reinhardtii* [11]. Clearly, the inability to perform state transitions is not a general characteristic of photopsychrophily in polar green algae. In addition to these novel characteristics, *C. prisculii* also exhibits a weak phototactic response compared to either the mesophile, *C. reinhardtii*, and the Antarctic psychrophile, ICE-MVD [130]. This weak phototactic response in *C. prisculii* is associated with a reduction in the repertoire of genes encoding photoreceptors important in phototaxis as well as its unusually small eyespot [130]. ICE-MVD exhibited a much stronger phototactic response than *C. prisculii*, the authors concluded that a weak phototactic response is not a characteristic of psychrophily, but rather, probably reflects shade-adaptation of *C. prisculii* to its natural, Lake Bonney habitat [130].

We suggest that homeoviscous adaptation and the maintenance of photostasis is essential for polar algae to survive life at the edge.

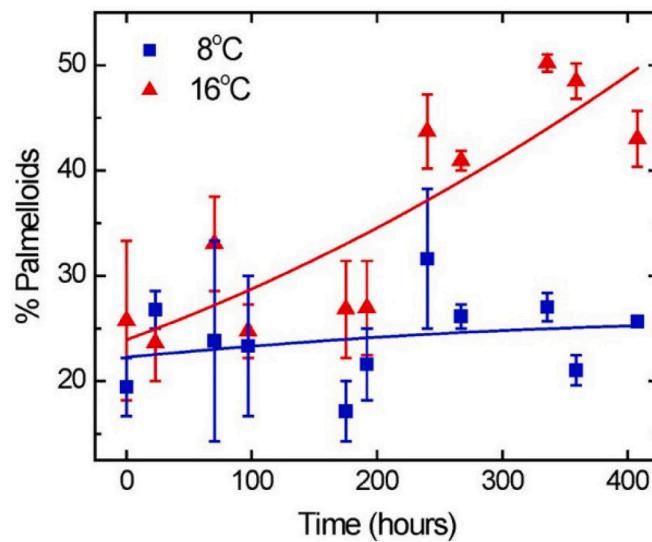
### 3. Multicellularity

Myriad acclimation strategies have evolved to enhance survival to environmental stress (Table 1). However, the intensity combined with the time-period over which an algal culture is exposed to a particular environmental stress is also a crucial factor that will ultimately determine survival. Are there alternative mechanisms to homeoviscous adaptation, phototaxis and photostasis that algae exploit to acclimate and adapt to conditions of prolonged stress? When the physiological and molecular mechanisms are no longer sufficient to protect against a particular stress, some algae have evolved higher-order, developmental strategies for survival through the conversion of single cells into either multicellular aggregates and/or multicellular structures called palmelloids [131–137]. The conversion of a motile, single cell to either an immobile, cellular aggregate consisting of hundreds of cells or a multicellular palmelloid consisting of 2 to 16 individual cells has been interpreted to represent stages in the onset of multicellularity in the evolution of eukaryotes [136–138]. In bacteria, cyclic di-GMP has been shown to regulate the transition from a motile lifestyle to a non-motile, sessile lifestyle important in the formation of complex biofilms [139–143]. Recently, the advent of complex cell architecture has also been reported in the Asgard archaea, *Candidatus Lokiarchaeum ossiferum*, from which it is suggested that multicellular eukaryotes may have emerged [144].

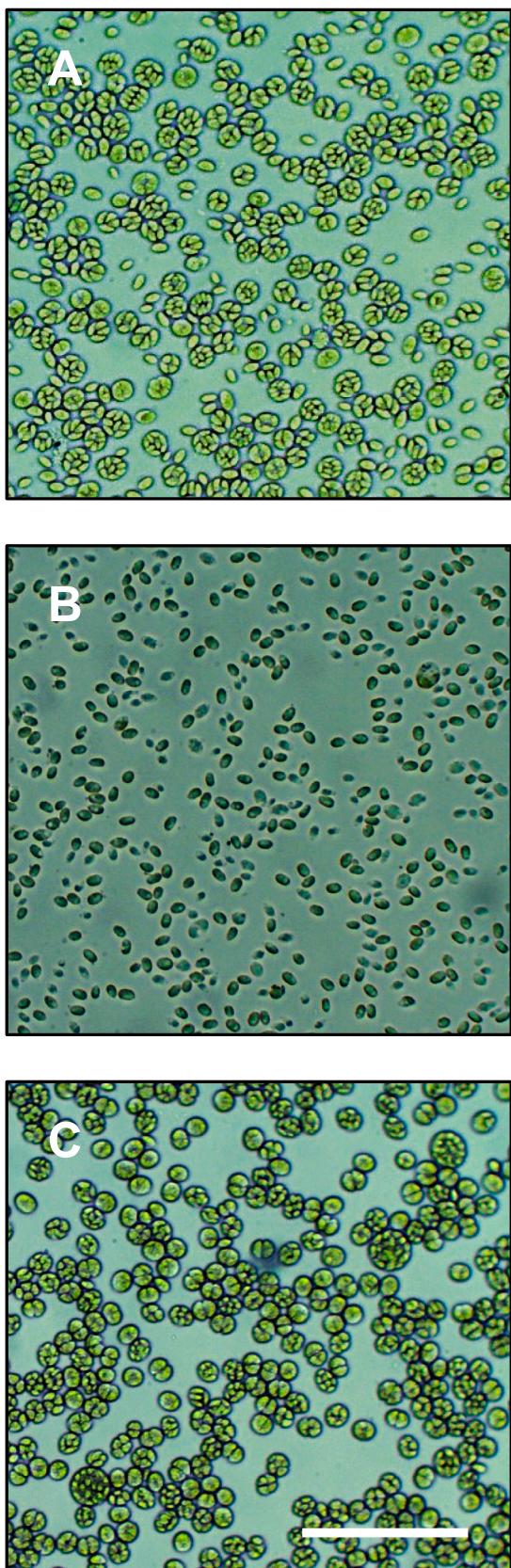
The transition from a single cell to a colonial palmelloid is the result of abnormal cell division and flagellar structure [145,146], which can be induced by various stress conditions including pollutants, low pH, temperature and oxidative stress [147–155]. Chloroplatinic acid induces palmelloid formation in *Chlamydomonas eugametos* by inhibiting the release of cells from the mother cell after cell division [152]. Interestingly, the cold-adapted Arctic strain of *Haematococcus pluvialis* exists

predominantly as an organized palmelloid [156]. It appears that any abrupt deviation from the optimal growth conditions potentially induces a transition from single cells to palmelloids and colonial aggregates in green algae. For example, exposure of the mesophile, *C. reinhardtii*, to a lower than the optimal growth temperature induces palmelloid formation whereas shifting the psychrophile, *C. prisculii*, to supra-optimal temperatures induces palmelloid formation (Figs. 2, 3) [157]. Most recently, it was reported that *C. reinhardtii* also forms palmelloids in response to salt stress [159,160]. Thus, we suggest that, in general, any prolonged environmental perturbation may impose a selective pressure for the transition to multicellularity in green algae.

Experimental evidence indicates that one evolutionary advantage of either cell aggregation or palmelloid formation is decreased susceptibility to predation by rotifers and other natural predators compared to the single cell state [136,137,148,149,153,154]. Exciting new evidence has been uncovered by de Carpentier et al. [136,137] elucidating the underlying molecular mechanism of cell aggregation in *C. reinhardtii*. The response of this model mesophile, to abiotic stress is to induce a collective behaviour between individual single cells to form large, random cell aggregates that protect the cells from heat shock. De Carpentier et al. [136,137] generated a family of socializer (*saz*) mutants of *C. reinhardtii* that aggregate spontaneously. Using a multi-omics approach, they showed that the observed cell aggregation is a result of genetic reprogramming and substantial modification of the secretome rather than passive agglutination. Exposure of WT cells to the media of *saz1* cells after those cells had been removed induced aggregation in WT cells. Analyses of the media of *saz1* cells indicated the presence of pherophorins, matrix metalloproteinases, serine-proline-rich proteins and lysis oxidases. Three pherophorin mutants (*phc30*, *phc41* and *phc50*) failed to aggregate and it was concluded that specific pherophorins are involved in the aggregation process in *C. reinhardtii*. Furthermore, the two pherophorin mutants (*phc28* and *phc35*) aggregated spontaneously [137], indicating that the products of these genes inhibit aggregation. These results provide key insights into the mechanisms and genes involved in the origins of multicellularity in green algae [138–141]. However, organization of single cells into a multicellular palmelloid must reflect a developmental response that is distinct from the apparent random aggregation reported by de Carpentier et al. [136,137]. Recent results show that, although the psychrophilic *C. prisculii* forms



**Fig. 2.** The effect of temperature on palmelloid formation in *C. prisculii*. *C. prisculii* was cultured at either 8 °C (blue, controls) or 16 °C (red, temperature stress) as described in detail in [157]. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)



**Fig. 3.** Light microscope images of *C. priscuii* control cells (A), isolated single cells (B) and isolated palmelloids (C). Magnification 40 $\times$ , scale bars 100  $\mu\text{m}$  (A, B and C).

palmelloids, it does not aggregate upon exposure to salt stress in contrast to *C. reinhardtii* (Fig. 4). However, *C. priscuii* does have a vegetative lytic enzyme (VLE) gene, homologous to that of *C. reinhardtii* and this gene appears to be complete and is present in three distinct copies in the *C. priscuii* genome [12a]. Thus, it would appear that *C. priscuii* has the potential to aggregate similar to *C. reinhardtii* but is inhibited from doing so. Further research is required to elucidate the regulation and molecular basis of palmelloid formation versus cellular aggregation in *C. priscuii*.

### 3.1. Multicellularity and the surface area to volume ratio

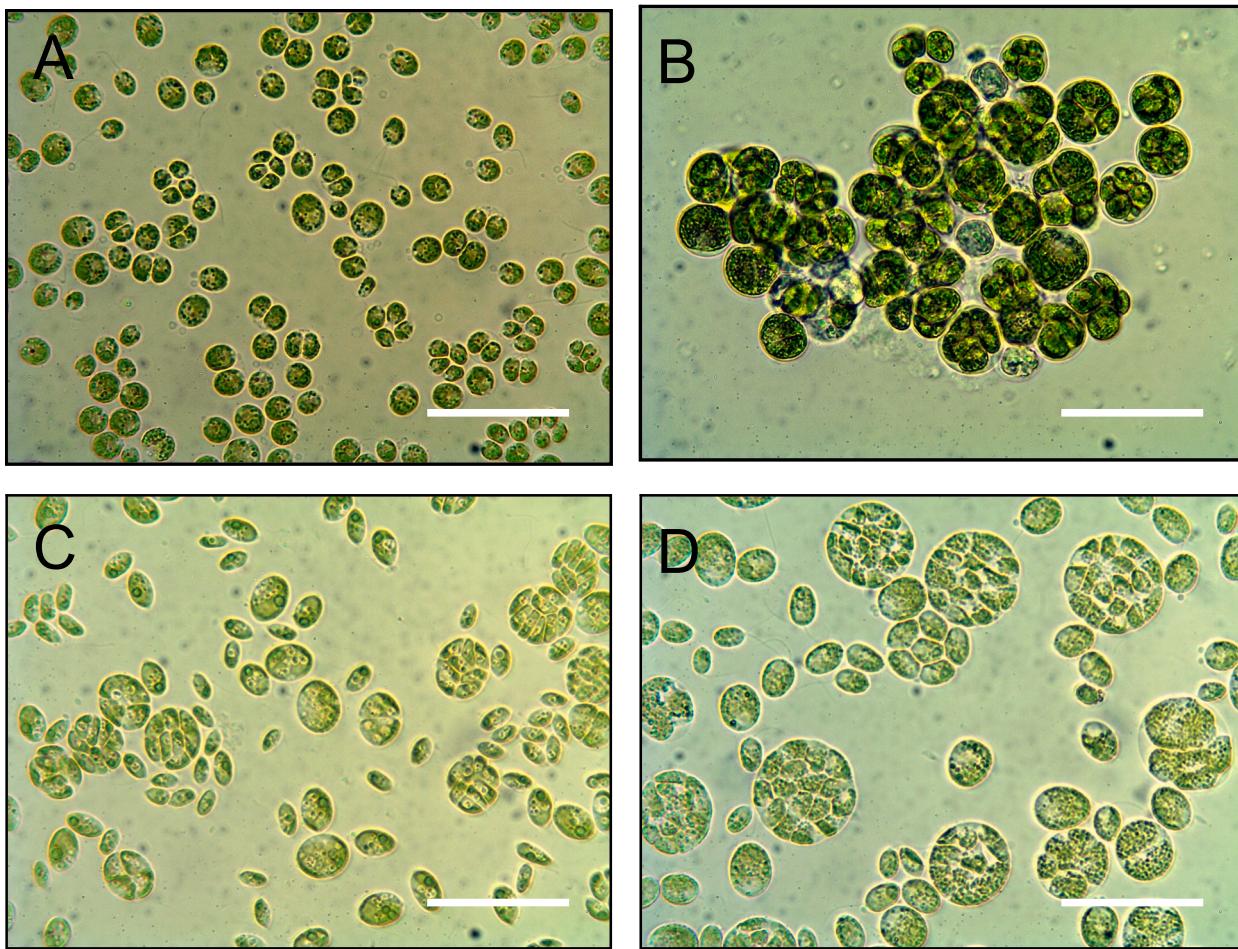
The growth rates of unicellular algal species are generally believed to depend on physiological, metabolic and environmental factors (temperature, salinity, etc.). In addition, several studies have shown that cell size is also a major factor for growth by its dependence on the surface area to volume (S/V) ratio and hence the rate of nutrient uptake [161,162]. Furthermore, it was hypothesised that the balance of selective pressures affecting S/V ratio is critical in aqueous environments, where autotrophic algal species compete for nutrients absorbed through their surface membranes [161–163]. Indeed, a positive correlation between growth rates, nutrient uptake, temperature, carbon assimilation, the maximum photosynthetic rate and S/V ratio has been reported in various algal species [163–171].

The origin and evolution of organismal complexity (i.e., the transition from unicellular to multicellular) have been studied extensively and require several selective and mechanistic prerequisites (see [176] for a review). Several previous studies [171–181], have shown that single-cell organisms can readily evolve multicellularity when subjected to the right environmental selective pressures [182]. It has been reported that the appearance of a multicellular structures derived from a unicellular one was accompanied by a remarkable change in *Chlamydomonas*-specific and volvocine-specific cell cycle and reproductive gene expression patterns [179].

The transition from motile unicellular organisms to colonial and/or larger multicellular aggregates is accompanied by a variety of requirements and ultimately has benefits and costs. Commonly, various environmental selective pressures are assumed to be the driving force for unicellular organisms to increase in size. The general constraint limiting the increase of cell size is the decrease in the surface to volume ratio (S/V ratio), which reduces the exchange of nutrients and thus the growth rates of the cells [181–184]. To avoid these constraints, several unicellular organisms including various algal species developed the strategy of holding the mitotic products (daughter cells) together by extracellular material, thus increasing in size by forming aggregates/palmelloids without significant alterations in S/V ratio of the individual cells [180–188]. However, forming a group (aggregates/palmelloids) would decrease the S/V ratio of the newly formed multicellular structure which was referred to as the “transport limitation” and was viewed as a general aspect of evolutionary transitions by Solari et al. [184]. Thus, the metabolic and viability constraints in larger structures (colonies/palmelloids) might provide a strong enough selective pressure to push the organismal design to germ-soma differentiation and higher complexity [29,184,185,187–189,191–193].

### 3.2. Multicellularity and photoprotection

Axenic cultures of *C. priscuii* propagated at low temperatures (4–10 °C) under continuous light represent a typical mixture of 20 % nonmotile palmelloids and 80 % motile single cells (Fig. 3) [157]. Compared to single cells, palmelloids of *C. priscuii* show decreased levels of the PSII light-harvesting complex II (LHCII) proteins consistent with a 2-fold higher Chl a/b ratio during growth at 16 °C than growth at 8 °C [157] which reduces light absorption efficiency. This is reflected in comparative difference 77 K fluorescence emission spectroscopy of *C. priscuii* single cells compared to palmelloids [157]. This spectroscopic



**Fig. 4.** Light microscope images of *Chlamydomonas reinhardtii* cultures grown in Bold's Basal medium (BBM), supplemented with 25 mM NaCl (A) and 200 mM NaCl (B). *Chlamydomonas priscuii* cultures grown in BBM, supplemented with 25 mM NaCl (C) and 700 mM NaCl (D). Magnification 40×. Scale bars, 20  $\mu$ m.

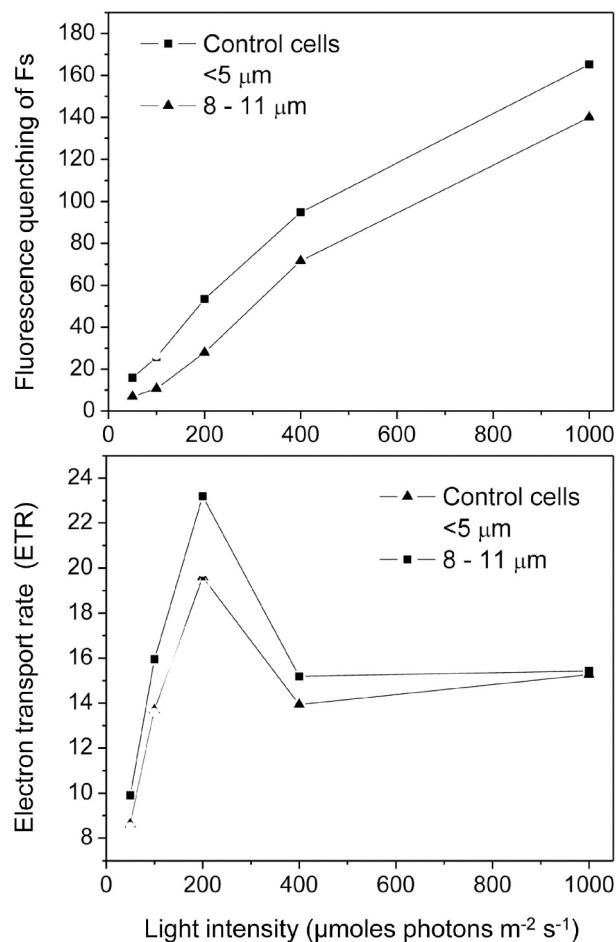
technique is very sensitive to changes in the pigment and polypeptide compositions as well as the organization of PSII and PSI and their functional interactions [50–52]. Thus, surprisingly, the transition from single cells to multicellular palmelloids appears to result in detectable alterations in the organization and composition of the PSI and PSII pigment-protein complexes [157].

Comparative room temperature Chla fluorescence induction combined with 77 K Chla fluorescence emission spectroscopic analyses of isolated cells and palmelloids indicated that, although the maximum PSII photochemical efficiency, measured as Fv/Fm, was comparable for isolated single cells and isolated palmelloids, excitation pressure, measured as 1-QL, was lower in palmelloids than single cells [157] consistent with an increased capacity to dissipate EEE in palmelloids compared to single cells. This differential level of excitation pressure was associated with 50 % higher light saturated rates of ETR in palmelloids relative to single cells exposed to high light at low temperature. The decreased sensitivity to high light was by enhanced non-regulated dissipation of EEE ( $\Phi_{NO}$ ) presumably through reaction centre quenching rather than normal antenna quenching via the xanthophyll cycle [63,64]. Functionally, the temperature-induced morphological transition of *C. priscuii* from a single to a multicellular state alters the composition and organization of the photosynthetic apparatus which predisposes it to dissipate EEE via non-regulated dissipation of excess absorbed energy [157]. Thus, single cells are morphologically and photosynthetically distinct from the single cells enclosed within a palmelloid. We suggest that in addition to decreased susceptibility to predation, palmelloid formation results in photoprotection from photoinhibition under EEE conditions. Thus, palmelloid formation may

be an additional selective criterion for multicellularity in some green algae which has not been considered previously, to our knowledge. However, the molecular mechanism underlying the observed alterations in the organization of the photosynthetic apparatus in response to the transition of single cells to multicellular palmelloids remains to be elucidated (Fig. 5).

#### 4. Conclusions

Cold adaptation of *C. priscuii* is dependent upon the combination of homeoviscous adaptation combined with the maintenance of photostasis and optimum S/V ratio. The exposure of *C. priscuii*, to increased growth temperatures close to non-permissive conditions (18–20 °C) is stressful for this stenothermic psychrophile. In response to this temperature stress, *C. priscuii*, shifts its morphology from motile, single cells to immobile, multicellular palmelloids which protects its photosynthetic apparatus from EEE by reducing light absorption efficiency by decreasing the abundance of the PSII light-harvesting complexes coupled with enhanced energy dissipation of excess absorbed light via NO quenching mechanisms. Thus, an advantage of palmelloid formation is decreased susceptibility to stress-induced photoinhibition [157] while maintaining an optimum S/V ratio. We suggest that in addition to decreased susceptibility to predation [144,145,148,149,153,154], enhanced photoprotection from photoinhibition associated with palmelloid formation may contribute an evolutionary advantage for the induction of multicellularity in some green algae.



**Fig. 5.** Palmelloids of *C. priscii* exhibit a decreased sensitivity to light. Isolated single cells (control cells, <5 μm) and isolated palmelloids (8–11 μm) were purified from *C. priscii* cultures exposed to increasing light intensity. Photosynthetic performance was assessed by quenching of room temperature Chla fluorescence (F<sub>s</sub>) and light saturated electron transport rates (ETR). Data are from [157].

#### CRediT authorship contribution statement

Conceptualization of the research by NPAH, AG and BS-M. Low temperature (77 K) chlorophyll fluorescence spectral analyses and room temperature chlorophyll fluorescence induction performed and analyzed by AG. Culturing of algae by BS-M, AG, VK and HL. Separation of single cells and palmelloids by BS-M, VK. Light microscopy and analyses by BS-M, VK. DRS performed the comparative genomic analyses. NPAH wrote the initial and the final drafts which were edited by all co-authors prior to submission.

#### Declaration of competing interest

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: Norman Hüner reports financial support was provided by Natural Sciences and Engineering Research Council of Canada.

#### Data availability

Data will be made available on request.

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#### References

- [1] C.L. Sabine, R.A. Feely, N. Gruber, et al., The oceanic sink for anthropogenic CO<sub>2</sub>, *Science* 305 (2004) 367–371.
- [2] N. Bax, C.J. Sands, B. Gogarty, R.V. Downey, C.V.E. Moreau, B. Moreno, C. Held, M. Paulsen, J. McGee, M. Haward, D.K.A. Barnes, Perspective: increasing blue carbon around Antarctica is an ecosystem service of considerable societal and economic value worth protecting, *Glob. Chang. Biol.* 27 (2020) 5–12.
- [3] C.W. Armstrong, N.S. Foley, D. Slagstad, M. Chierici, I. Ellingsen, M. Reigstad, Valuing blue carbon changes in the Arctic Ocean, *Front. Mar. Sci.* 6 (2019), 331.
- [4] S. Bielewicz, E. Bell, W. Kong, I. Friedberg, J.C. Priscu, R.M. Morgan-Kiss, Protist diversity in a permanently ice-covered Antarctic Lake during the polar night transition, *ISME J.* 5 (2011) 1559–1564.
- [5] R.M. Morgan-Kiss, J.C. Priscu, T. Pocock, L. Gudynaite-Savitch, N.P.A. Hüner, Adaptation and acclimation of photosynthetic microorganisms to permanently cold environments, *Microbiol. Mol. Biol. Rev.* 222–252 (2006).
- [6] J.C. Priscu, C.H. Fritsen, E.E. Adams, J. Giovannoni, H.W. Paerl, C.P. McKay, P.T. Doran, D.A. Gordon, B.D. Lanoil, J.L. Pinckney, Perennial Antarctic lake ice: an oasis for life in a polar desert, *Science* 280 (1998) 2095–2098.
- [7] M.C. Kennicutt, S.L. Chown, J.J. Cassano, D. Liggett, R. Massom, L.S. Peck, S. Rintoul, J.W.V. Storey, D.G. Vaughan, T.J. Wilson, W.J. Sutherland, Polar research: six priorities for Antarctic science, *Nature* 512 (2014) 23–25.
- [8] M.C. Kennicutt, S.L. Chown, J.J. Cassano, et al., A roadmap for Antarctic and Southern Ocean science for the next two decades and beyond, *Antarct. Sci.* 27 (2015) 3–18.
- [9] J.C. Xavier, A. Brandt, Y. Ropert-Coudert, et al., Future challenges in Southern Ocean ecology research, *Front. Mar. Sci.* 3 (2016) 94, <https://doi.org/10.3389/fmars.2016.00094>.
- [10] A.D. Rogers, Evolution and biodiversity of Antarctic organisms: a molecular perspective, *Phil. Trans. R. Soc. B* 362 (2007) 2191–2214.
- [11] Y.M. Bar-On, R. Phillips, R. Milo, The biomass distribution on Earth, *Proc. Natl. Acad. Sci. U. S. A.* 115 (2018) 6506–6511.
- [12] N.P.A. Hüner, D.R. Smith, M. Cvetkovska, X. Zhang, A.G. Ivanov, B. Szyszka-Mroz, I. Kalra, R.M. Morgan-Kiss, Photosynthetic adaptation to polar life: energy balance, photoprotection and genetic redundancy, *J. Plant Physiol.* 268 (2022), 153557. [a]X. Zhang, M. Cvetkovska, R.M. Morgan-Kiss, N.P.A. Hüner, D.R. Smith. Draft genome sequence of the Antarctic green alga *Chlamydomonas* sp. UW0241. *iScience* 24: 102084.
- [13] A. Casanueva, M. Tuffin, C. Cary, D.A. Cowan, Molecular adaptations to psychrophily: the impact of “omic” technologies, *Trends Microbiol.* 18 (2010) 374–381.
- [14] A. Banerjee, H. Wan-Shabir, A. Roychoudhury, Epigenetic control of plant cold responses, *Front. Plant Sci.* 8 (2017) 1643, <https://doi.org/10.3389/fpls.2017.01643>.
- [15] C. Crosatti, F. Rizza, F.W. Badeck, E. Mazzucotelli, L. Cattivelli, Harden the chloroplast to protect the plant, *Physiol. Plant.* 147 (2013) 55–63.
- [16] M. Griffith, M. Timonin, A.C.E. Wong, G.R. Gray, S.R. Akhter, M. Saldanha, M.A. Rogers, E.A. Weretilnyk, B. Moffatt, *Thellungiella*: an *Arabidopsis*-related model plant adapted to cold temperatures, *Plant Cell Environ.* 30 (2007) 529–538.
- [17] C.L. Guy, Cold acclimation and freezing tolerance: role of protein metabolism, *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 41 (1990) 187–223.
- [18] C.L. Guy, R. Porat, V. Hurry, Plant cold and abiotic stress gets hot, *Physiol. Plant.* 126 (2006) 1–4.
- [19] Y. Kazachkova, G. Eshel, P. Pantha, J.M. Cheeseman, M. Dassanayake, S., Barak Halophytism: what have we learnt from *Arabidopsis thaliana*: relative model systems? *Plant Physiol.* 178 (2018) 972.
- [20] C.-H. Lee, M.F. Thomashow, Photoperiodic regulation of the C-repeat binding factor (CBF) cold acclimation pathway and freezing tolerance in *Arabidopsis thaliana*, *Proc. Natl. Acad. Sci. U. S. A.* 109 (2012) 15054–15059.
- [21] P. Stepien, G.N. Johnson, Contrasting responses of photosynthesis to salt stress in the glycophyte *Arabidopsis* and the halophyte *Thellungiella*: role of the plastid terminal oxidase as an alternative electron sink, *Plant Physiol.* 149 (2009) 1154–1165.
- [22] M. Stitt, V. Hurry, A plant for all seasons: alterations in photosynthetic carbon metabolism during cold acclimation in *Arabidopsis*, *Curr. Opin. Plant Biol.* 5 (2002) 199–206.
- [23] M.F. Thomashow, Molecular basis of plant cold acclimation: insights gained from studying the CBF cold response pathway, *Plant Physiol.* 154 (2010) 571–577.
- [24] E. Ermilova, Cold stress response: an overview in *Chlamydomonas*, *Front. Plant Sci.* 11 (2020), 569437, <https://doi.org/10.3389/fpls.2020.569437>.
- [25] D.P. Maxwell, S. Falk, C.G. Trick, N.P.A. Hüner, Growth at low temperature mimics high-light acclimation in *Chlorella vulgaris*, *Plant Physiol.* 105 (1994) 535–543.

- [26] D.P. Maxwell, S. Falk, N.P.A. Hüner, Photosystem II excitation pressure and development of resistance to photoinhibition I. LHCl abundance and zeaxanthin content in *Chlorella vulgaris*, *Plant Physiol.* 107 (1995) 687–694.
- [27] D.P. Maxwell, D.E. Laudenbach, N.P.A. Hüner, Redox regulation of light-harvesting complex II and *cab* mRNA abundance in *Dunaliella salina*, *Plant Physiol.* 109 (1995) 787–795.
- [28] R.M. Morgan-Kiss, Photosynthesis on the edge: phytoplankton life in the ice-covered lakes of the McMurdo Dry Valleys, *Can. Antarct. Res. Netw.* 21 (2006) 17–18.
- [29] K.S. Siddiqui, T.J. Williams, D. Wilkins, S. Yau, M.A. Allen, M.V. Brown, F. M. Lauro, R. Cavicchioli, Psychrophiles, *Annu. Rev. Earth Planet. Sci.* 41 (2013) 87–115.
- [30] T. Collins, R. Margesin, Psychophilic lifestyles: mechanisms of adaptation and biotechnological tools, *Appl. Microbiol. Biotechnol.* 103: 2857–2871.
- [31] Z. Zhang, C. Qu, R. Yao, Y. Nie, C. Xu, J. Miao, B. Zhong, The parallel molecular adaptations to the Antarctic cold environment in two psychophilic green algae, *Genome Biol. Evol.* 11 (2019) 1897–1908.
- [32] R. Ernst, C.S. Ejsing, B. Antony, Homeoviscous adaptation and the regulation of membrane lipids, *J. Mol. Biol.* 428 (2016) 4776–4791.
- [33] J. Browse, C. Somerville, Glycerolipid synthesis: biochemistry and regulation, *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 42 (1991) 467–506.
- [34] J.B. Ohlrogge, J. Browse, C.R. Somerville, The genetics of plant lipids, *Biochim. Biophys. Acta* 1082 (1991) 1–26.
- [35] B.Y. Moon, S.-I. Higashi, Z. Gombos, N. Murata, Unsaturation of the membrane lipids of chloroplasts stabilizes the photosynthetic machinery against low-temperature photoinhibition in transgenic tobacco plants, *Proc. Natl. Acad. Sci. U. S. A.* 92 (1995) 6219–6223.
- [36] I. Nishida, Chilling sensitivity in plants and cyanobacteria: the crucial contribution of membrane lipids, *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 47 (1996) 541–568.
- [37] N. Murata, Membrane fluidity and temperature perception, *Plant Physiol.* 115 (1997) 875–879.
- [38] J.L. Harwood, Involvement of chloroplast lipids in the reaction of plants submitted to stress, in: P.-A. Siegenthaler (Ed.), *Advances in Photosynthesis and Respiration, Lipids in Photosynthesis: Structure, Function and Genetics*, vol. 6, Kluwer Academic Publishers, Dordrecht, 1998, pp. 287–302.
- [39] P. Dormann, C. Benning, Galactolipids rule in seed plants, *Trends Plant Sci.* 7 (2002) 112–118.
- [40] C. Xu, H. Hartel, H. Wada, M. Hagio, B. Yu, C. Eakin, C. Benning, The *pgp1* mutant locus of *Arabidopsis* encodes a phosphatidylglycerolphosphate synthase with impaired activity, *Plant Physiol.* 129 (2002) 594–604.
- [41] D.A. Los, N. Murata, Sensing and responses to low temperature in cyanobacteria, in: K.B. Storey, J.M. Storey (Eds.), *Sensing, Signalling and Cell Adaptation*, Elsevier Science BV, Amsterdam, 2002, pp. 139–153.
- [42] A.C. Barnes, C. Benning, R.L. Roston, Chloroplast membrane remodeling during freezing stress is accompanied by cytoplasmic acidification activating *SENSITIVE TO FREEZING2*, *Plant Physiol.* 171 (2016) 2140–2149.
- [43] A.G. Ivanov, M. Krol, L.V. Savitch, B. Szyszka-Mroz, J. Roche, D.P. Spratt, E. Selstam, K.E. Wilson, R. Gardiner, G. Öquist, V.M. Hurry, N.P.A. Hüner, The decreased PG content of *pgp1* inhibits PSI photochemistry and limits reaction center and light-harvesting polypeptide accumulation in response to cold acclimation, *Planta* 255 (2022) 36, <https://doi.org/10.1007/s00425-022-03819-0>.
- [44] N. Ferrer-Ledo, L. Stegemüller, M. Janssen, R.H. Wijffels, M.J. Barbosa, Growth and fatty acid distribution over lipid classes in *Nannochloropsis oceanica* acclimated to different temperatures, *Front. Plant Sci.* 14 (2023), 1078998, <https://doi.org/10.3389/fpls.2023.1078998>.
- [45] N.J. Russell, N. Kukunaga, A comparison of thermal adaption of membrane lipids in psychophilic and thermophilic bacteria, *FEMS Microbiol. Rev.* 75 (1990) 171–182.
- [46] S. Kralova, Role of fatty acids in cold adaptation of Antarctic psychophilic *Flavobacterium* spp., *Syst. Appl. Microbiol.* 40: 329–333.
- [47] R. Morgan-Kiss, A.G. Ivanov, J.P. Williams, M. Khan, N.P.A. Hüner, Differential thermal effects on the energy distribution between photosystem II and photosystem I in thylakoid membranes of a psychophilic and a mesophilic alga, *Biochim. Biophys. Acta* 1561 (2002) 251–265.
- [48] J.A. Berry, O. Björkman, Photosynthetic response and adaptation to temperature in higher plants, *Annu. Rev. Plant Physiol.* 31 (1980) 491–543.
- [49] G.H. Krause, E. Weis, Chlorophyll fluorescence and photosynthesis: the basics, *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 42 (1991) 313–349.
- [50] N.R. Baker, Chlorophyll fluorescence: a probe of photosynthesis in vivo, *Annu. Rev. Plant Biol.* 59 (2008) 89–113.
- [51] T. Swoczyńska, H.M. Kalaji, F. Bussotti, J. Mojski, M. Pollastrini, Environmental stress- what can we learn from chlorophyll a fluorescence analysis in woody plants? A review, *Front. Plant Sci.* 13 (2022), 104852.
- [52] R.Y. Morita, Psychophilic bacteria, *Bacteriol. Rev.* 39 (1975) 144–167.
- [53] M. Cvetkovska, X. Zhang, G. Vakulenko, S. Benzaquen, B. Szyszka-Mroz, N. Malczewski, D.R. Smith, N.P.A. Hüner, A constitutive stress response is a result of low temperature growth in the Antarctic green alga *Chlamydomonas* sp. UWO241, *Plant Cell Environ.* 45 (2022) 156–177.
- [54] W.J. Newsted, N.P.A. Hüner, Major polypeptides associated with differentiation in psychophilic fungi, *Can. J. Bot.* 65 (1987) 1755–1761.
- [55] W.J. Newsted, N.P.A. Hüner, Major sclerotial polypeptides of psychophilic pathogenic fungi: intracellular localization and antigenic relatedness, *Protoplasma* 147 (1988) 162–169.
- [56] S. D'Amico, Fundamentals of cold-adapted enzymes, in: T. Collins, F. Roullier, F. Piette, J.-C. Marx, G. Feller, C. Gerday, R. Margesin, F. Schinner, J.-C. Marx, C. Gerday (Eds.), *Psychrophiles. From Biodiversity to Biotechnology*, Springer-Verlag, Berlin, 2008, pp. 211–227.
- [57] R. Margesin, F. Schinner, J.-C. Marx, C. Gerday, *Psychrophiles: From Biodiversity to Biotechnology*, Springer-Verlag, Berlin, 2008.
- [58] F. Zakhia, A.-D. Jungblut, A. Taton, W. Vincent, A. Wilmotte, Cyanobacteria in cold ecosystems, in: R. Margesin, F. Schinner, J.-C. Marx, C. Gerday (Eds.), *Psychrophiles. From Biodiversity to Biotechnology*, Springer-Verlag, Berlin, 2008, pp. 121–135.
- [59] E.P.Y. Tang, R. Tremblay, W.F. Vincent, Cyanobacterial dominance of polar freshwater ecosystems - high-latitude mat-formers adapted to low temperature, *J. Phycol.* 33 (1997) 171–181.
- [60] E.P.Y. Tang, W.F. Vincent, Strategies of thermal adaptation by high-latitude cyanobacteria, *New Phytol.* 142 (1999) 315–323.
- [61] N.P.A. Hüner, G. Öquist, F. Sarhan, Energy balance and acclimation to light and cold, *Trends Plant Sci.* 3 (1998) 224–230.
- [62] N.P.A. Hüner, G. Öquist, A. Melis, Photostasis in plants, green algae and cyanobacteria: the role of light harvesting antenna complexes, in: B.R. Green, W.W. Parson (Eds.), *Advances in Photosynthesis and Respiration, Light Harvesting Antennas in Photosynthesis*, vol. 13, Kluwer Academic Publishers, Dordrecht, 2003, pp. 401–421.
- [63] I. Ensminger, F. Busch, N.P.A. Hüner, Photostasis and cold acclimation: sensing low temperature through photosynthesis, *Physiol. Plant.* 126 (2006) 28–44.
- [64] K. Petrou, R. Hill, C.M. Brown, D.A. Campbell, M.A. Doblin, P.J. Ralph, Rapid photoprotection in sea-ice diatoms from the East Antarctic pack ice, *Limnol. Oceanogr.* 55 (2010) 1400–1407.
- [65] K. Petrou, R. Hill, M.A. Doblin, A. McMinn, R. Johnson, S.W. Wright, P.J. Ralph, Photoprotection of sea-ice microalgal communities from the east Antarctic pack ice, *J. Phycol.* 47 (2011) 77–86.
- [66] K. Petrou, S.A. Kranz, S. Trimborn, C.S. Hassler, S.B. Ameijeiras, O. Sackett, P.J. Ralph, A.T. Davidson, Southern Ocean phytoplankton physiology in a changing climate, *J. Plant Physiol.* 203 (2016) 135–150, <https://doi.org/10.1016/j.jplph.2016.05.004>.
- [67] D.M. Kramer, J.R. Evans, The importance of energy balance in improving photosynthetic productivity, *Plant Physiol.* 155 (2011) 70–78.
- [68] A. Melis, Photostasis in plants, in: Williams, Thistle (Eds.), *Photostasis and Related Phenomena*, Plenum Press, New York, 1998, pp. 207–220.
- [69] L. Hollis, N.P.A. Hüner, Retrograde operational sensing and signalling pathways maintain photostasis in green algae, cyanobacteria and terrestrial plants, *Trends Photochem. Photobiol.* 16 (2014) 47–61.
- [70] R. Mittler, S. Vanderauwera, N. Suzuki, G. Miller, V.B. Tognetti, K. Vandepoele, M. Gollery, V. Shulaev, F. van Breusegem, ROS signalling: the new wave? *Trends Plant Sci.* 16 (2011) 300–309.
- [71] K.-J. Dietz, R. Mittler, Recent progress in understanding the role of reactive oxygen species in plant cell signalling, *Plant Physiol.* 171 (2016) 1535–1539.
- [72] C.H. Foyer, G. Noctor, Redox homeostasis and signaling in a higher-CO<sub>2</sub> world, *Annu. Rev. Plant Biol.* 71 (2020) 157–182.
- [73] J.-M. Escoubas, M. Lomas, J. LaRoche, P.G. Falkowski, Light intensity regulates *cab* gene transcription via the redox state of the plastoquinone pool in the green alga, *Dunaliella tertiolecta*, *Proc. Natl. Acad. Sci. U. S. A.* 92 (1995) 10237–10241.
- [74] P.G. Falkowski, Y.-B. Chen, Photoacclimation of light harvesting systems in eukaryotic algae, in: B.R. Green, W.W. Parson (Eds.), *Advances in Photosynthesis and Respiration, Light Harvesting Systems in Photosynthesis*, vol. 13, Kluwer Academic Publishers, Dordrecht, 2003, pp. 423–447.
- [75] T. Masuda, A. Tanaka, A. Melis, Chlorophyll antenna size adjustments by irradiance in *Dunaliella salina* involve coordinate regulation of chlorophyll *a* oxygenase (CAO) and *Lhcb* gene expression, *Plant Mol. Biol.* 51 (2003) 757–771.
- [76] Y.-B. Chen, D.G. Durnford, M. Kobliezak, P.G. Falkowski, Plastid regulation of *Lhcb1* transcription in the chlorophyte alga, *Dunaliella tertiolecta*, *Plant Physiol.* 136 (2004) 3737–3750.
- [77] P. Cardol, B. Bailleul, F. Rappaport, E. Derelle, D. Bacal, C. Breyton, S. Bailey, F. A. Wollman, A. Grossman, H. Moreau, G. Finazzi, An original adaptation of photosynthesis in the marine green alga *Ostreococcus*, *Proc. Natl. Acad. Sci. U. S. A.* 105 (2008) 7881–7886.
- [78] C.H. Foyer, J. Neukermans, G. Queval, G. Noctor, J. Harbinson, The mechanisms contributing to photosynthetic control of electron transport by carbon assimilation in leaves, *Photosynth. Res.* 25 (1990) 83–100.
- [79] C.H. Foyer, J. Neukermans, G. Queval, G. Noctor, J. Harbinson, Photosynthetic control of electron transport and the regulation of gene expression, *J. Exp. Bot.* 63 (2012) 1637–1661.
- [80] A.E. McDonald, A.G. Ivanov, R. Bode, D.P. Maxwell, S.R. Rodermel, N.P.A. Hüner, Flexibility in photosynthetic electron transport: the physiological role of plastoquinol terminal oxidase (PTOX), *Biochim. Biophys. Acta Bioenerg.* 1807 (2011) 954–967.
- [81] J.-D. Rochaix, Regulation of photosynthetic electron transport, *Biochim. Biophys. Acta Bioenerg.* 1807 (2011) 375–383.
- [82] F. Chaux, G. Peltier, X. Johnson, A security network in PSI photoprotection: regulation of photosynthetic control, NPQ and O<sub>2</sub> photoreduction by cyclic electron flow, *Front. Plant Sci.* 6 (2015) 875, <https://doi.org/10.3389/fpls.2015.00875>.
- [83] G. Peltier, E.-M. Aro, T. Shikanai, NDH-1 and NDH-2 plastoquinone reductases in oxygenic photosynthesis, *Annu. Rev. Plant Biol.* 67 (2016) 55–80.
- [84] M.A.E. Miller, R. O'Cualain, J. Selley, D. Knight, M.F. Karim, S.J. Hubbard, G. N. Johnson, Dynamic acclimation to high light in *Arabidopsis thaliana* involves

- widespread reengineering of the leaf proteome, *Front. Plant Sci.* 8 (2017) 1239, <https://doi.org/10.3389/fpls.2017.01239>.
- [87] W.J. Nawrocki, N.J. Tourasse, A. Taly, F. Rappaport, F.-A. Wollman, The plastid terminal Oxidase: its elusive function points to multiple contributions top plastid physiology, *Annu. Rev. Plant Biol.* 2015 (66) (2015) 49–74.
- [88] W.J. Nawrocki, B. Bailleul, D. Picot, P. Cardol, F. Rappaport, F.-A. Wollman, P. Joliot, The mechanism of cyclic electron flow, *Biochim. Biophys. Acta Bioenerg.* 1860 (2019) 433–438.
- [89] S. Karpinski, H. Reynolds, B. Karpinska, G. Wingsle, G. Creissen, P. Mullineaux, Systemic signaling and acclimation in response to excess excitation energy in *Arabidopsis*, *Science* 284 (1999) 654–657.
- [90] F.-A. Wollman, State transitions reveal the dynamics and flexibility of the photosynthetic apparatus, *EMBO J.* 20 (2001) 3623–3630.
- [91] J.-D. Rochaix, Fine-tuning photosynthesis, *Science* 342 (2013) 50–51.
- [92] J.-D. Rochaix, Regulation and dynamics of the light-harvesting system, *Annu. Rev. Plant Biol.* 65 (2014) 287–309.
- [93] S. Eberhard, G. Finazzi, F.-A. Wollman, The dynamics of photosynthesis, *Annu. Rev. Genet.* 42 (2008) 463–515.
- [94] J. Minagawa, State transitions—the molecular remodeling of photosynthetic supercomplexes that controls energy flow in the chloroplast, *Biochim. Biophys. Acta Bioenerg.* 1807 (2011) 897–905.
- [95] J. Minagawa, R. Tokutsu, Dynamic regulation of photosynthesis in *Chlamydomonas reinhardtii*, *Plant J.* 82 (2015) 413–428.
- [96] P. Horton, A. Ruban, The role of light-harvesting complex II in energy quenching, in: N.R. Baker, J.R. Bowyer (Eds.), *Photoinhibition of Photosynthesis*. From Molecular Mechanisms to the Field, Bios Scientific, Oxford, UK, 1994, pp. 111–128.
- [97] A.A. Pascal, Z. Liu, K. Broess, B. van Oort, H. van Amerongen, C. Wang, P. Horton, B. Robert, W. Chang, A. Ruban, Molecular basis of photoprotection and control of photosynthetic light-harvesting, *Nature* 436 (2005) 134–137.
- [98] B. Demmig-Adams, W.W. Adams III, The role of xanthophyll cycle carotenoids in the protection of photosynthesis, *Trends Plant Sci.* 1 (1996) 21–26.
- [99] Ecophysiology of the xanthophyll cycle, in: B. Demmig-Adams, W.W. Adams, V. Ebbert, B.A. Logan, H.A. Frank, A.J. Young, G. Britton, R.J. Cogdell (Eds.), *Advances in Photosynthesis*, The Photochemistry of Carotenoids, vol. 8, Kluwer Academic, Dordrecht, 1999, pp. 245–269.
- [100] B. Demmig-Adams, J.J. Stewart, T.A. Burch, W.W. Adams, Insights from placing photosynthetic light harvesting into context, *J. Phys. Chem. Lett.* 5 (2014) 2880–2889.
- [101] A.S. Verhoeven, Sustained energy dissipation in winter evergreens, *New Phytol.* 201 (2014) 57–65.
- [102] E. Erickson, S. Wakao, K.K. Niyogi, Light stress and photoprotection in *Chlamydomonas reinhardtii*, *Plant J.* 82 (2015) 449–465.
- [103] G. Öquist, Effects of low temperature on photosynthesis: a review, *Plant Cell Environ.* 6 (1983) 281–301.
- [104] G. Öquist, W.S. Chow, J.M. Anderson, Photoinhibition of photosynthesis represents a mechanism for long-term regulation of photosystem II, *Planta* 186 (1992) 450–460.
- [105] S.P. Long, S. Humphries, P.G. Falkowski, Photoinhibition of photosynthesis in nature, *Annu. Rev. Plant Physiol. Plant Mol. Biol.* 45 (1994) 633–662.
- [106] N.P.A. Hüner, G. Öquist, V.M. Hurry, M. Krol, S. Falk, M. Griffith, Photosynthesis, photoinhibition and low temperature acclimation in cold tolerant plants, *Photosynth. Res.* 37: 19–39.
- [107] G.Y. Nie, S.P. Long, N.R. Baker, The effects of development at sub-optimal growth temperatures on photosynthetic capacity and susceptibility to chilling-dependent photoinhibition in *Zea mays*, *Physiol. Plant.* 85 (1992) 554–560.
- [108] D.B. Hayden, N.R. Baker, M.P. Percival, P.B. Beckwith, Modification of the photosystem II light-harvesting chlorophyll a/b protein complex in maize during chill-induced photoinhibition, *Biochim. Biophys. Acta* 851 (1986) 86–92.
- [109] T. Pfannschmidt, Chloroplast redox signals: how photosynthesis controls its own genes, *Trends Plant Sci.* 8 (2003) 33–41, [https://doi.org/10.1016/s1360-1385\(02\)00005-5](https://doi.org/10.1016/s1360-1385(02)00005-5).
- [110] T. Pfannschmidt, C. Yang, N. Peter, K. Chong, The hidden function of photosynthesis: a sensing system for environmental conditions that regulates plant acclimation responses, *Protoplasma* 249 (2012) 125–136.
- [111] A. Nott, H.-S. Hou-Sung, S. Koussevitzky, J. Chory, Plastid-to-nucleus retrograde signalling, *Annu. Rev. Plant Biol.* 57 (2006) 739–759.
- [112] J.D. Woodson, J. Chory, Coordination of gene expression between organellar and nuclear genomes, *Nat. Rev. Genet.* 9 (2008) 383–395.
- [113] H.-S. Jung, J. Chory, Signaling between chloroplasts and the nucleus: can a systems biology approach bring clarity to a complex and highly regulated pathway? *Plant Physiol.* 152 (2010) 453–459.
- [114] M. Jokal, X. Johnson, G. Peltier, E.-M. Aro, Y. Allahverdiyeva, Hunting the main player enabling *Chlamydomonas reinhardtii* growth under fluctuating light, *Plant J.* 94 (2018) 822–836.
- [115] D.G. Durmford, P.G. Falkowski, Chloroplast redox regulation of nuclear gene transcription during photoacclimation, *Photosynth. Res.* 53 (1997) 229–241.
- [116] A. Sukenik, J. Bennett, P.G. Falkowski, Changes in the abundance of individual apoproteins of light-harvesting chlorophyll a/b complexes of photosystem I and II with growth irradiance in the marine chlorophyte *Dunaliella tertiolecta*, *Biochim. Biophys. Acta* 932 (1988) 2016–2215.
- [117] J. Laroche, A. Mortain-Bertrand, P.G. Falkowski, Light intensity-induced changes in cab mRNA and light harvesting complex II apoprotein levels in the unicellular chlorophyte *Dunaliella tertiolecta*, *Plant Physiol.* 97 (1991) 147–153.
- [118] M.R. Webb, A. Melis, Chloroplast response in *Dunaliella salina* to irradiance stress, *Plant Physiol.* 107 (1995) 885–893.
- [119] M. Krol, D.P. Maxwell, N.P.A. Hüner, Exposure of *Dunaliella salina* to low temperature mimics the high light-induced accumulation of carotenoids and the carotenoid binding protein (Cbr), *Plant Cell Physiol.* 38 (1997) 213–216.
- [120] E. Miskiewicz, A.G. Ivanov, J.P. Williams, M.U. Khan, S. Falk, N.P.A. Hüner, Photosynthetic acclimation of the filamentous cyanobacterium, *Plectonema boryanum* UTEX 485, to temperature and light, *Plant Cell Physiol.* 41 (2000) 767–775.
- [121] E. Miskiewicz, A.G. Ivanov, N.P.A. Hüner, Stoichiometry of the photosynthetic apparatus and phycobilisome structure of the cyanobacterium, *Plectonema boryanum* UTEX 485 are regulated by both light and temperature, *Plant Physiol.* 130 (2002) 1414–1425.
- [122] T. Pocock, A. Vetterli, S. Falk, Evidence for phenotypic plasticity in the Antarctic extremophile *Chlamydomonas raudensis* Ettr. UWO 241, *J. Exp. Bot.* 62 (2011) 1169–1177.
- [123] D. Rosso, R. Bode, W. Li, M. Krol, D. Saccon, S. Wang, L.A. Schillaci, S. R. Rodermel, D.P. Maxwell, N.P.A. Hüner, Photosynthetic redox imbalance governs leaf sectoring in the *Arabidopsis thaliana* variegation mutants *immutans*, *spotty*, *var1*, and *var2*, *Plant Cell* 21 (2009) 3473–3492.
- [124] R. Morgan-Kiss, A.G. Ivanov, N.P.A. Hüner, The Antarctic psychrophile, *Chlamydomonas subcaudata*, is deficient in state I-state II transitions, *Planta* 214 (2001) 435–445.
- [125] S. Stahl-Rommel, I. Kalra, S. D'Silva, M.M. Hahn, D. Popson, M. Cvetkovska, R. Morgan-Kiss, Cyclic electron flow (CEF) and ascorbate pathway activity provide constitutive photoprotection for the photopsychrophile, *Chlamydomonas* sp. UWO 241 (renamed *Chlamydomonas priscii*), *Photosynth. Res.* 151 (2022) 235–250.
- [126] B. Szyszka, A.G. Ivanov, N.P.A. Hüner, Psychrophily is associated with differential energy partitioning, photosystem stoichiometry and polypeptide phosphorylation in *Chlamydomonas raudensis*, *Biochim. Biophys. Acta Bioenerg.* 1767 (2007) 789–800.
- [127] B. Szyszka-Mroz, P. Pittock, A.G. Ivanov, G. Lajoie, N.P.A. Hüner, The Antarctic psychrophile *Chlamydomonas* sp. UWO 241 preferentially phosphorylates a photosystem I-cytochrome b6/f supercomplex, *Plant Physiol.* 169 (2015) 717–736.
- [128] B. Szyszka-Mroz, M. Cvetkovska, A.G. Ivanov, D.R. Smith, M. Possmayer, D. P. Maxwell, N.P.A. Hüner, Cold-adapted protein kinases and thylakoid remodeling impact energy distribution in an Antarctic psychrophile, *Plant Physiol.* 180 (2019) 1291–1309.
- [129] M. Poirier, P. Osmers, K. Wilkins, R.M. Morgan-Kiss, M. Cvetkovska, Aberrant light sensing and motility in the green alga *Chlamydomonas priscii*, *BioRxiv* (2023), <https://doi.org/10.1101/2023.02.02.526531>.
- [130] K. Iwasa, S. Murakami, Palmelloid formation of *Chlamydomonas*, *Physiol. Plant.* 21 (1968) 1224–1233.
- [131] K. Iwasa, S. Murakami, Palmelloid formation of *Chlamydomonas* II. Mechanism of palmelloid formation by organic acids, *Physiol. Plant.* 22 (1969) 43–50.
- [132] E. Harris, Introduction to *Chlamydomonas* and its laboratory use, in: *The Chlamydomonas Source Book*, 2nd edition vol. 1, 2009 (932p).
- [133] S. Sathe, P.M. Durand, Cellular aggregation in *Chlamydomonas* (Chlorophyceae) is chimaeric and depends on traits like cell size and motility, *Eur. J. Phycol.* 51 (2015) 129–138.
- [134] D.K. Khona, S.M. Shirolkar, K.K. Gawde, E. Hom, J.S. D'Souza, Characterization of saltstress-induced palmelloids in the green alga, *Chlamydomonas reinhardtii*, *Algal Res.* (2016) 16434–16448.
- [135] F. de Carpenter, S.D. Lemaire, A. Danon, When unity is strength: the strategies used by *Chlamydomonas* to survive environmental stresses, *Cells* 8 (2019) 1307, <https://doi.org/10.3390/cells8111307>.
- [136] F. de Carpenter, A. Mae, C.H. Marchand, C. Chung, C. Durand, P. Crozet, S. D. Lemaire, A. Danon, How abiotic stress-induced socialization leads to the formation of massive aggregates in *Chlamydomonas*, *Plant Physiol.* 190 (2022) 1927–1940.
- [137] A. Rokas, Origins of multicellularity and the early genetic toolkit for animal development, *Annu. Rev. Genet.* 42 (2008) 235–251.
- [138] S.E. Prochnik, J. Umen, A.M. Nedelcu, A. Hallmann, S.M. Miller, I. Nishii, P. Ferris, A. Kuo, T. Mitros, L.K. Fritz-Laylin, U. Hellsten, J. Chapman, O. Simakov, S.A. Rensing, A. Terry, J. Pangilinan, V. Kapitonov, J. Jurka, A. Salamov, H. Shapiro, J. Schmutz, J. Grimwood, E. Lindquist, S. Lucas, I. V. Grigoriev, R. Schmitt, D. Kirk, D.S. Rohksar, Genomic analysis of organismal complexity in the multicellular green alga *Volvox carteri*, *Science* 329 (2010) 223–226.
- [139] B.J.S.C. Olson, From brief encounters to life-long unions, *eLife* 2 (2013), e01893.
- [140] B. Calla, Stronger together: how unicellular algae respond to stress by socialization, *Plant Physiol.* 190 (2022) 1554–1555.
- [141] U. Römling, M. Galperin, M. Gomelsky, Cyclic di-GMP: the first 25 years of a universal bacterial second messenger, *Microbiol. Mol. Biol. Rev.* 77 (2013) 1–52.
- [142] U. Romling, A. Moglich, Seeing the light brings more food in the deep sea, *EMBO J.* 42 (2023), e114091.
- [143] T. Rodrigues-Oliveira, F. Wollweber, R.I. Ponce-Toledo, J. Xu, S.K.M.R. Rittmann, A. Klingl, M. Pilhofer, C. Schleper, Actin cytoskeleton and complex cell architecture in an Asgard archaeon, *Nature* 613 (2023) 332–339, <https://doi.org/10.1038/s41586-022-05550-y>.
- [144] M.E. Boraas ME, D.B. Seale, J.E. Boxhorn, Phagotrophy by a flagellate selects for colonial prey: a possible origin of multicellularity, *Evol. Ecol.* 12: 153–164.
- [145] M.D. Herron, J.M. Borin, J.C. Boswell, J. Walker, I.C.K. Chen, C.A. Knox, M. Boyd, F. Rosenzweig, W.C. Ratcliff, *De novo* origins of multicellularity in response to predation, *Sci. Rep.* 9 (2019) 2328, <https://doi.org/10.1038/s41598-019-39558-8>.

- [147] J.R. Warr, A. McVittie, J. Randall, J.M. Hopkins, Genetic control of flagellar structure in *Chlamydomonas reinhardtii*, *Genet. Res.* 7 (1966) 335–351.
- [148] J.R. Warr, A mutant of *Chlamydomonas reinhardtii* with abnormal cell division, *J. Gen. Microbiol.* 52 (1968) 243–251.
- [149] I. Visviki, D. Santikul, The pH tolerance of *Chlamydomonas appanata* (Volvocales, Chlorophyta), *Arch. Environ. Contam. Toxicol.* 38 (2000) 147–151.
- [150] M. Lürling, E. Van Donk, Morphological changes in *Scenedesmus* induced by infochemicals released in situ from zooplankton grazer, *Limnol. Oceanogr.* 42 (1997) 783–788.
- [151] M. Lürling, W. Beekman, Palmelloid formation in *Chlamydomonas reinhardtii*: defence against rotifer predators? *Int. J. Limnol.* 42 (2006) 65–72.
- [152] K. Nakamura, D.F. Bray, E.B. Wagenaar, Ultrastructure of *Chlamydomonas eugametos* palmelloids induced by chloroplatinic acid treatment, *J. Bacteriol.* 121 (1975) 338–343.
- [153] K. Nakamura, D.F. Bray, E.B. Wagenaar, Ultrastructure of a palmelloid-forming strain of *Chlamydomonas eugametos*, *Can. J. Bot.* 56 (1978) 2348–2356.
- [154] V. Zachleder, I. Ivanov, M. Vitova, K. Bisova, Cell cycle arrest by supraoptimal temperature in the alga *Chlamydomonas reinhardtii*, *Cells* 8 (2019) 1237, <https://doi.org/10.3390/cells8101237>.
- [155] K. Nakamura, M. Sakon, M.K. Hatanaka, Chemical factors affecting palmelloid-forming activity of chloroplatinic acid on *Chlamydomonas eugametos*, *Physiol. Plant.* 36 (1976) 293–296.
- [156] T. Klochkova, M.S. Kwak, J.W. Han, T. Motomura, Cold-tolerant strain of *Haematococcus pluvialis* (Haematococcaceae, Chlorophyta) from Blomstrandhalvoya (Svalbard), *Algae* 28 (2013) 185–192.
- [157] B. Szyszka-Mroz, A.G. Ivanov, C.G. Trick, N.P.A. Hüner, Palmelloid formation in the Antarctic psychrophile, *Chlamydomonas prisculii*, is photoprotective, *Front. Plant Sci.* 13 (2022), 911035, <https://doi.org/10.3389/fpls.2022.911035>.
- [158] M. Cvetkovska, B. Szyszka-Mroz, M. Possmayer, P. Pittock, G. Lajoie, D.R. Smith, N.P.A. Hüner, Characterization of photosynthetic ferredoxin from the Antarctic alga *Chlamydomonas sp.* UWO241 reveals novel features of cold adaptation, *New Phytol.* 219 (2018) 588–604.
- [159] E. Devadasu, S.D. Kanna, S. Neelam, R.M. Yadav, S. Nama, P. Akhtar, T.F. Polga, B. na Ughy, G. Garab, P.H. Lambrev, R. Subramanyam, Long-term and short-term acclimation of the photosynthetic apparatus to salinity in *Chlamydomonas reinhardtii*. The role of St7 protein kinase, *Front. Plant Sci.* 14 (2023), 1051711.
- [160] A. Farkas, B. Pap, O. Zsiros, R. Patai, P. Shetty, G. Garab, T. Biro, V. Ordog, G. Maroti, Salinity stress provokes physiological responses of eukaryotic unicellular microalgae, *Algal Res.* 73 (2023), 103155.
- [161] G.E. Fogg, Algal Cultures and Phytoplankton Ecology, 2nd edition, University of Wisconsin Press, Madison and Milwaukee, 1975.
- [162] W.M. Lewis, Surface/volume ratio: implications for phytoplankton morphology, *Science* 192 (1976) 885–887.
- [163] I.O.W. Findlay, Effects of external factors and cell size on the cell division rate of a marine diatom, *Coscinodiscus pavillardii* Forti, *Int. Revue ges Hydrobiol. Hydrogr.* 57 (1972) 523–533.
- [164] R.H. Foy, C.E. Gibson, R.V. Smith, The influence of daylength, light intensity and temperature on the growth rates of planktonic blue-green algae, *Br. Phycol. J.* 11 (1976) 151–163.
- [165] R.H. Foy, The influence of surface to volume ratio on the growth rates of planctonic blue-green algae, *Br. Phycol. J.* 15 (1980) 279–289.
- [166] E.S. Friebele, D.L. Correll, M.A. Faust, Relationship between phytoplankton cell size and the rate of orthophosphate uptake: *in situ* observations of an estuarine population, *Mar. Biol.* 45 (1978) 39–52.
- [167] I. Findenegg, Relationship between standing crop and primary productivity, *Memorie Ist. Ital Idrobiol.* 18 (Suppl) (1965) 271–289.
- [168] S. Taguchi, Relationship between photosynthesis and cell size of marine diatoms, *J. Phycol.* 12 (1976) 185–189.
- [169] R. Margalef, Modifications induced by different temperatures on the cells of *Scenedesmus obliquus* (Chlorophyceae), *Hydrobiologia* 6 (1954) 83–91.
- [170] K.J. Niklas, S.A. Newman, The many roads to and from multicellularity, *J. Exp. Bot.* 71 (2020) 3247–3253.
- [171] J.H. Koschwanez, K.R. Foster, A.W. Murray, Improved use of a public good selects for the evolution of undifferentiated multicellularity, *eLife* 2 (2013), e00367.
- [172] L. Becks, S.P. Ellner, L.E. Jones, N.G. Hairston, The functional genomics of an eco-evolutionary feedback loop: linking gene expression, trait evolution, and community dynamics, *Ecol. Lett.* 15 (2012) 492–501.
- [173] M.D. Herron, W.C. Ratcliff, J. Boswell, F. Rosenzweig, Genetics of a de novo origin of undifferentiated multicellularity, *R. Soc. Open Sci.* 5 (2018), 180912.
- [174] W.C. Ratcliff, R.F. Denison, M. Borrello, M. Travisano, Experimental evolution of multicellularity, *Proc. Natl. Acad. Sci. U. S. A.* 109 (2012) 1595–1600.
- [175] W.C. Ratcliff, M.D. Herron, K. Howell, J.T. Pentz, F. Rosenzweig, M. Travisano, Experimental evolution of an alternating uni- and multicellular life cycle in *Chlamydomonas reinhardtii*, *Nat. Commun.* 4 (2013) 2742.
- [176] R.K. Grosberg, R.R. Strathmann, The evolution of multicellularity: a minor major transition? *Annu. Rev. Ecol. Evol. Syst.* 38 (2007) 621–654.
- [177] L. Becks, S.P. Ellner, L.E. Jones, N.G. Hairston, Reduction of adaptive genetic diversity radically alters eco-evolutionary community dynamics, *Ecol. Lett.* 13 (2010) 989–997.
- [178] C.A. Solari, S. Ganguly, J.O. Kessler, R.E. Michod, R.E. Goldstein, Multicellularity and the functional interdependence of motility and molecular transport, *Proc. Natl. Acad. Sci. U. S. A.* 103 (2006) 1353–1358.
- [179] C.A. Solari, J.O. Kessler, R.E. Goldstein, A general allometric and life-history model for cellular differentiation in the transition to multicellularity, *Am. Nat.* (2013) 369–380.
- [180] K.J. Niklas, Plant Allometry: The Scaling of Form and Process, University of Chicago Press, Chicago, 1994.
- [181] K.J. Niklas, The evolution of plant body plans: a biomechanical perspective, *Ann. Bot.* 85 (2000) 411–438.
- [182] L.E. Graham, L.W. Wilcox, Algae, Prentice Hall, Englewood Cliffs, NJ, 2000.
- [183] M.B. Short, C.A. Solari, S. Ganguly, T.R. Powers, J.O. Kessler, R.E. Goldstein, Flows driven by flagella of multicellular organisms enhance log-range molecular transport, *Proc. Natl. Acad. Sci. U. S. A.* 103 (2006) 8315–8319.
- [184] C.A. Solari, J.O. Kessler, R.E. Michod, A hydrodynamics approach to the evolution of multicellularity: flagellar motility and the evolution of germ-soma differentiation in volvocalean green algae, *Am. Nat.* 167 (2006) 537–554.
- [185] G. Feller, J.L. Arpigny, E. Nminx, Ch. Geday, Molecular adaptations of enzymes from psychrophilic organisms, *Camp. Biochem. Physiol.* 118A (1997) 495–499.
- [186] N. Devos, M. Ingouff, R. Loppes, R.F. Matagne, Rubisco adaptation to low temperatures: a comparative study in psychrophilic and mesophilic unicellular algae, *J. Phycol.* 34 (1998) 655–660.
- [187] K. Miyazaki, P.L. Wintrode, R.A. Grayling, D.N. Rubingh, F.H. Arnold, Directed evolution study of temperature adaptation in a psychrophilic enzyme, *J. Mol. Biol.* 297 (2000) 1015–1026.
- [188] G. Feller, C. Gerday, Psychrophilic enzymes: hot topics in cold adaptation, *Nat. Rev. Microbiol.* 1 (2003) 200–208.
- [189] D.I. Paredes, K. Watters, D.J. Pitman, C. Bystroff, J.S. Dordick, Comparative void-volume analysis of psychrophilic and mesophilic enzymes: structural bioinformatics of psychrophilic enzymes reveals sources of core flexibility, *BMC Struct. Biol.* 11 (2011) 42.
- [191] N.J. Russell, N. Fukunaga, A comparison of thermal adaptation of membrane lipids in psychrophilic and thermophilic bacteria, *FEMS Microbiol. Rev.* 75 (1990) 171–182.
- [192] T. Mock, M.B. Kroon, Photosynthetic energy conversion under extreme conditions. II: the significance of lipids under light limited growth in Antarctic Sea ice diatoms, *Phytochemistry* 61 (2002) 53–60.
- [193] S. Králová, Role of fatty acids in cold adaptation of Antarctic psychrophilic *Flavobacterium* spp., *Syst. Appl. Microbiol.* 40 (2017) 329–333.
- [194] N. Hassan, A.M. Anesio, M. Rafiq, J. Holtvoeth, I. Bull, A. Haleem, A.A. Shah, F. Hasan, Temperature driven membrane lipid adaptation in glacial psychrophilic bacteria, *Front. Microbiol.* 11 (2020) 824.
- [195] T. Mock, H. Hock, Long-term temperature acclimation of photosynthesis in steady-state cultures of the polar diatom *Fragilaropsis cylindrus*, *Photosynth. Res.* 85 (2005) 307–317.
- [196] T. Mock, K. Valentini, Photosynthesis and cold acclimation: molecular evidence from a polar diatom, *J. Phycol.* 40 (2004) 732–741.
- [198] M.V. Jagannadham, M.K. Chattopadhyay, C. Subbalakshmi, M. Vairamani, K. Narayanan, C.M. Rao, S. Shivaji, Carotenoids of an Antarctic psychrotolerant bacterium, *Sphingobacterium antarcticus*, and a mesophilic bacterium, *Sphingobacterium multivorum*, *Arch. Microbiol.* 173 (2000) 418–424.
- [199] T. Leya, A. Rahn, C. Lütz, D. Remias, Response of arctic snow and permafrost algae to high light and nitrogen stress by changes in pigment composition and applied aspects for biotechnology, *FEMS Microbiol. Ecol.* 67 (2009) 432–443.
- [200] M.K. Chattopadhyay, M.V. Jagannadham, M. Vairamani, S. Shivaji, Carotenoid pigments of an Antarctic psychrotrophic bacterium *Micrococcus roseus*: temperature dependent biosynthesis, structure, and interaction with synthetic membranes, *Biochem. Biophys. Res. Commun.* 239 (1997) 85–90.
- [201] M. Dieser, M. Greenwood, C.M. Foreman, Carotenoid pigmentation in Antarctic heterotrophic bacteria as a strategy to withstand environmental stresses, *Arct. Antarct. Alp. Res.* 42 (2010) 396–405.
- [202] L. Shen, Y. Liu, N. Wang, N. Jiao, B. Xu, X. Liu, Variation with depth of the abundance, diversity and pigmentation of culturable bacteria in a deep ice core from the Yuzhufeng Glacier, *Tibetan Plateau, Extremophiles* 22 (2018) 29–38.
- [203] A.-F. Ghobakhloo, A. Johnston, L. Harris, H. Antoun, S. Laberge, Microarray transcriptional profiling of Arctic *Mesorhizobium* strain N33 at low temperature provides insights into cold adaption strategies, *BMC Genomics* 16 (2015) 383.
- [204] F. Fonseca, J. Meneghel, S. Cenard, S. Passot, G.J. Morris, Determination of intracellular vitrification temperatures for unicellular microorganisms under conditions relevant for cryopreservation, *PLoS One* 11 (2016), e0152939.
- [205] J. Goordial, I. Raymond-Bouchard, Y. Zolotarov, L. de Bethencourt, J. Ronholm, N. Shapiro, T. Woyke, M. Stromvik, C.W. Greer, C. Bakermans, L. Whyte, Cold adaptive traits revealed by comparative genomic analysis of the eurypsychrophile *Rhodococcus* sp. JG3 isolated from high elevation McMurdo Dry Valley permafrost, Antarctica, *FEMS Microbiol. Ecol.* 92 (2) (2016), fiv154, <https://doi.org/10.1093/fems/fiv154>.
- [206] A. Raymond, R. Morgan-Kiss, S. Stahl-Rommel, Glycerol is an osmoprotectant in two Antarctic *Chlamydomonas* species from an ice-covered saline lake and is synthesized by an unusual bidomain enzyme, *Front. Plant Sci.* 11 (2020) 1259.
- [207] J.A. Gilbert, P.J. Hill, C.E. Dodd, J. Laybourn-Parry, Demonstration of antifreeze protein activity in Antarctic lake bacteria, *Microbiology* 150 (2004) 171–180.
- [208] N. Muryoji, M. Sato, S. Kaneko, H. Kawahara, H. Obata, M.W. Yaish, M. Griffith, B. R. Glick, Cloning and expression of *afpA*, a gene encoding an antifreeze protein from the arctic plant growth-promoting rhizobacterium *Pseudomonas putida* GR12-2, *J. Bacteriol.* 186 (2004) 5661–5671.
- [209] J.A. Raymond, B.C. Christner, S.C. Schuster, A bacterial ice-binding protein from the Vostok ice core, *Extremophiles* 12 (5) (2008) 713–717.
- [210] M. Bar Dolev, I. Braslavsky, P.L. Davies, Ice-binding proteins and their function, *Annu. Rev. Biochem.* 85 (2016) 515–542.
- [211] A. Tosco, L. Birolo, S. Madonna, G. Lolli, G. Sannia, G. Marino, GroEL from the psychrophilic bacterium *Pseudoalteromonas haloplanktis* TAC 125: molecular characterization and gene cloning, *Extremophiles* 7 (2003) 17–28.

- [212] K. Yoshimune, A. Galkin, L. Kulakova, T. Yoshimura, N. Esaki, Cold-active DnaK of an Antarctic psychrotroph *Shewanella* sp. Ac10 supporting the growth of dnaK-null mutant of *Escherichia coli* at cold temperatures, *Extremophiles* 9 (2005) 145–150.
- [213] M.L. Romero-Romero, A. Inglés-Prieto, B. Ibarra-Molero, J.M. Sanchez-Ruiz, Highly anomalous energetics of protein cold denaturation linked to folding-unfolding kinetics, *PLoS One* 6 (2011) e23050.
- [214] T. Collins, C. Gerday, Enzyme catalysis in psychrophiles, in: R. Margesin (Ed.), *Psychrophiles: From Biodiversity to Biotechnology*, 2nd edn, Springer, Cham, 2017, pp. 209–235.