

PAR acclimation and UVBR-induced DNA damage in Antarctic marine microalgae

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ABSTRACT: Whether or not it is related to stratospheric ozone depletion, Antarctic microalgae experience ultraviolet-B radiation (UVBR) stress *in situ*, leading to decreased photosynthetic performance, DNA damage, and/or altered community composition. UVBR vulnerability is known to be species specific, but may also be affected by a range of environmental growth conditions, including the light history of the cells. This study investigates the influence of acclimation to photosynthetically active radiation (PAR) on the vulnerability to UVBR-induced DNA damage in Antarctic microalgae. *Chaetoceros dichaeta*, *Pyramimonas gelidicola*, *Phaeocystis antarctica* and *Polarella glacialis* were acclimated to 5 PAR levels, after which growth rate, pigment composition, malondialdehyde (MDA, a general indicator of oxidative stress) and UV-absorbing compounds were measured. Photoacclimated cultures were then exposed to a single UVBR treatment and the accumulation of UVBR-induced DNA damage was determined by the number of cyclobutane pyrimidine dimers (CPDs). Acclimation to increasing irradiance enhanced both the xanthophyll to chlorophyll *a* ratio and xanthophyll de-epoxidation in all species. Increased cellular MDA levels were found at the highest irradiance in all species except *P. gelidicola*. Yet, growth rates were only reduced at the 2 lowest PAR levels. *P. antarctica* and *P. glacialis* showed a strong linear induction of UV-absorbing compounds at increasing PAR intensities, whereas *P. gelidicola* showed no induction of these compounds. The UVBR treatment induced CPDs in *P. gelidicola* only, and CPD levels were elevated at the highest PAR acclimation intensities. Thus, sensitivity to UVB-induced CPD accumulation was species specific, and, counterintuitively, acclimation to high PAR increased the sensitivity of *P. gelidicola* to UVB-induced DNA damage.

KEY WORDS: DNA damage · Xanthophyll pigments · Antarctic microalgae · Malondialdehyde · MDA · Photoacclimation · UVBR · CPDs

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INTRODUCTION

Marine microalgae, including those inhabiting high latitude regions, experience ultraviolet-B radiation (UVBR: 280 to 315 nm wavelength) stress *in situ*. Natural UVBR levels are well known to structure Antarctic marine phytoplankton communities and affect overall productivity. There is a wealth of information on decreased photosynthetic rate (for reviews see Vernet 2000, Villafane et al. 2003) as well as on UVBR-induced DNA damage, expressed as cyclobutane

pyrimidine dimers (CPDs) (for review see Buma et al. 2003). At the same time, species-specific differences in UVBR vulnerability (Karentz et al. 1991, Davidson et al. 1994) may, under prolonged (elevated) UVBR exposure, alter phytoplankton community composition, as demonstrated for a number of field locations including Antarctica (Bothwell et al. 1993, Villafane et al. 1995, Mostajir et al. 1999, Mousseau et al. 2000, Davidson & Belbin 2002). In addition to the UVBR dose and dose rate, the extent to which microalgal communities suffer from *in situ* UVBR stress depends on the interaction

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with other environmental factors such as the rate and depth of vertical mixing (Helbling et al. 1994, Neale et al. 1998a, 2003) and nutrient concentrations (Behrenfeld et al. 1994, Lesser et al. 1994, Litchman et al. 2002, Shelley et al. 2002). In polar regions, UVBR levels are enhanced during spring time as a result of stratospheric ozone depletion, which may cause an integrated loss of carbon fixation in Antarctic waters between <0.15 and 12% (Smith et al. 1992, Holm-Hansen et al. 1993, Helbling et al. 1994).

The light history of cells can positively or negatively influence the UVBR tolerance. For example, acclimation to high irradiances of photosynthetically active radiation (PAR: 400 to 700 nm wavelength) or to ultraviolet-A radiation (UVAR: 315 to 400 nm wavelength) may induce production of mycosporine-like amino acids (MAAs), which protect the cells against incoming short wavelengths (Neale et al. 1998b, Banaszak 2003). Also, repair mechanisms, such as the enzymes that correct UVBR-induced DNA damage, may be more active when cells are acclimated to high *in situ* irradiances, as is found in higher plants (Takahashi et al. 2002, Waterworth et al. 2002). Conversely, high PAR and UVAR may cause oxidative stress, when excitation energy dissipates via pathways, leading to dangerous reactive intermediates such as singlet oxygen or other oxyradicals. In the latter case, radicals may damage proteins of the photosystem II (PSII) reaction centre and destroy membrane integrity, causing a decrease in photosynthetic performance or even viability loss (Poll et al. 2005). One recent study showed increased PAR- and UVR-induced viability loss, when cells were precultured at low irradiance levels, possibly related to increased pigmentation of the cells (Poll et al. 2005).

How photooxidative and antioxidative mechanisms affect UVBR sensitivity is largely unknown. Microalgae, as most organisms, develop or induce mechanisms that offer protection against excess solar irradiance (for review Banaszak 2003). Direct protection is provided by xanthophyll cycle activity (Demers et al. 1991, Olaizola et al. 1994), whereby excess excitation energy is dissipated as heat (Demers et al. 1991, Demming-Adams & Adams 1992) in the cyclic de-epoxidation/epoxidation of diadinoxanthin/diatoxanthin (chromophytes including diatoms, prymnesiophytes and dinoflagellates) or of violaxanthin/zeaxanthin/antheraxanthin (in chlorophytes and prasinophytes). The epoxidation state [EPS: epoxidated/(epoxidated + de-epoxidated pigments)] can, therefore, be considered an indicator of xanthophyll cycle activity under excess irradiance conditions. In addition, while acclimating to high irradiance, the pool of xanthophyll pigments may increase, whereas cellular light-harvesting pigments decrease (Demers et al. 1991).

Oxidative stress occurs when the rate of xanthophyll cycling or antioxidant enzyme action is exceeded by the oxiradical accumulation rate. One of the major consequences of oxidative stress is the oxidation of membrane lipids that may cause loss of cell permeability and, eventually, cell viability. During the process of membrane oxidation, harmful phospholipid residues may be formed, including malondialdehyde (MDA). MDA is considered a general indicator of oxidative stress in animal, human, as well as in micro- and macroalgal cells (Rijstenbil 2002, Bischof et al. 2003).

Considering the above, synergistic or antagonistic effects of PAR, UVAR and UVBR exposure are to be expected. Yet, the nature and extent to which this occurs in Antarctic microalgae needed to be further elucidated. One study with an Antarctic cyanobacterium showed UVBR vulnerability to be strongly dependent on the UVBR to UVAR ratio, with a lower ratio being more beneficial for the organisms (Quesada et al. 1995). This was suggested to be related to the induction of UVB protective stress proteins at higher UVAR levels. Here, we examined the effect of the PAR acclimation of 4 common Antarctic marine microalgae on their sensitivity to UVBR-induced DNA damage.

MATERIALS AND METHODS

Experimental set-up. Unialgal, non-axenic Antarctic microalgae were acclimated to PAR and exposed to UVBR in 500 ml polystyrene flasks (Techno Plastic Products TPP 90151) in GP5 medium (Loeblich & Smith 1968, and modified after Loeblich 1975) based on filter-sterilised natural seawater, in a temperature-controlled ($1.0 \pm 0.5^\circ\text{C}$) culture cabinet. The investigated species were: flagellates of the haptophyte *Phaeocystis antarctica* Karsten; the diatom *Chaetoceros dictyota* Ehrenb.; the dinoflagellate *Polarella glacialis* Montresor, Proccacani & Stoecker; and the prasinophyte *Pyramimonas gelidicola* McFadden, Moestrup & Wetherbee (CS-139). All cultures were obtained from the Australian Antarctic Division culture collection. *P. antarctica* remained in the flagellated stage for the duration of the experimental period.

PAR levels were applied by varying the distance between the culture bottles and 20 watt (W) cool white fluorescent tubes (Osram, Germany, see Fig. 1 for spectral composition), thereby giving 5 irradiance levels between 18 and 220 $\mu\text{mol m}^{-2} \text{s}^{-1}$ in a 16 h light:8 h dark cycle. PAR acclimation was achieved by keeping the cultures at their respective irradiances for a minimum of 2 wk, after which time the cultures were assumed to be acclimated. Cultures were regularly diluted with fresh medium to avoid nutrient depletion during the experiments.

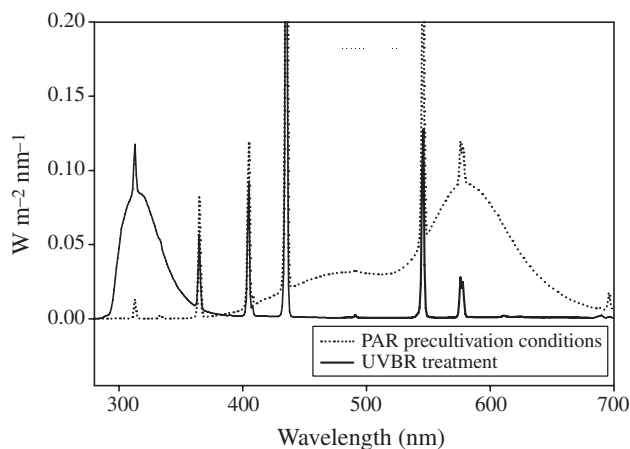


Fig. 1. Spectral irradiance conditions used for photosynthetically active radiation (PAR) acclimation (dotted curve) and the single ultraviolet-B radiation (UVBR) treatments (continuous curve). Spectra were corrected for spectral characteristics of the polystyrene flasks used in experiments. For irradiance levels, see 'Materials and methods'. Mean values of 2 replicates are presented

After acclimation, growth rates were measured over a 4 d period, and on Day 4, just prior to the UVBR treatment, samples were taken to measure pigments, UVR-absorbing compounds and MDA accumulation. Then, the flasks with the remaining culture suspensions were exposed to 1 single UVBR treatment for 2 h around noon (Philips TL 12, 0.1 W m^{-2} , for spectral composition see Fig. 1) at a temperature of 1.0°C ($\pm 0.5^\circ\text{C}$). The applied weighted dose (Minimal Erythral Dose [MED] 3.6), roughly corresponds with a daily erythral UVBR dose at approximately 4 m water depth at Davis Station (Antarctica) during a clear summer day, when assuming a UV attenuation constant in low-turbidity Antarctic marine coastal waters of 0.4 (Davidson & van der Hiejden 2000). Cells were harvested for CPD analysis after exposure to UVBR. The entire experiment was repeated after several weeks, giving 2 independent replicates.

Analyses. For growth rate measurements, samples were taken on 4 consecutive days, fixed with Lugol's iodine (0.1% final concentration), diluted with filter-sterilised seawater when necessary and settled in 5 ml counting chambers; the entire chamber (1200 to 2500 cells) was screened using a Zeiss Axiovert inverted microscope under Nomarski optics at $100\times$ or $400\times$ magnification. Biovolumes of 200 cells of each species and acclimation irradiance were measured using stereometric formulas for flagellates and diatoms, as given by Hillebrand et al. (1999).

For pigment analyses with HPLC, between 50 and 80 ml of culture was filtered onto Whatman GF/F filters under dim light and stored immediately in liquid nitrogen. To avoid de-epoxidation of xanthophyll cycle pigments, the time in between sampling, filtration and

storage was typically <2 min. Samples were extracted in 3 ml of 100% MeOH under dim light, using *apo-8'*-carotenal as internal standard. Pigments were analysed using the procedure described by Zapata et al. (2000). Pigments were detected using a Waters 996 photodiode array connected to Hitachi fluorescence detectors. Standards were isolated by semi-preparative HPLC, using procedures described by Repeta & Bjørnland (1997), from well-characterised species including *Amphidinium carterae*, *Dunaliella tertiolecta*, *Emiliania huxleyi*, *Micromonas pusilla*, *Pavlova lutheri* and *Pelagococcus subviridis*, as recommended by Jeffrey & Wright (1997).

For UVB-absorbing compounds, between 50 and 80 ml of culture was filtered onto GF/F filters and stored at -135°C until analysis, which was done within a few weeks. UV-absorbing compounds were measured spectrophotometrically following Tartarotti & Sommaruga (2002), using 3 ml of aqueous 25% methanol extraction at 45°C for 2 h. Extracts were filtered through GF/F filters, and the absorption of the filtrate was recorded between 250 and 700 nm on a Hitachi spectrophotometer, revealing at least 1 UV absorbance peak around 323 nm. Baseline correction of all UV absorbance peaks was done using 25% MeOH as a blank, and further background correction (non-specific turbidity) was done by subtracting extrapolated background signals between 250 and 700 nm, using Hitachi software. UV-absorbing compounds were expressed as absorption units per cell or biovolume.

MDA formation was assessed using the procedure of Heath & Parker (1968). This spectrophotometric method measures total oxidated phospholipid content, with MDA as the major constituent. Therefore, the observed colorimetric response is generally referred to as TBAR (thiobarbituric acid reactive), rather than MDA. For TBAR measurements, between 70 and 100 ml of culture (18 and $220 \mu\text{mol m}^{-2} \text{ s}^{-1}$ PAR-acclimated cultures only) was filtered onto GF/F filters and stored at -135°C until further analysis. Extraction was done for 30 min at 98°C in 3 ml of solvent consisting of 1.5 ml MilliQ and 1.5 ml of trichloroacetic acid (TCA, 20% in MilliQ) with thiobarbituric acid (TBA, 0.5% [w/v], Sigma) and then put on ice immediately. After cooling, the extract was filtered to remove cell debris, after which absorption spectra between 450 and 600 nm were recorded on a Hitachi spectrophotometer. The extraction solvent (TCA-TBA/MilliQ, 50:50 [v/v]) was used as a blank. To correct for non-specific turbidity, average background absorption was measured over the whole range of the spectrum and subtracted from the TBAR peak at 532 nm using Hitachi software. MDA was calculated assuming a molar extinction coefficient of 155.

For CPD analysis, 80 to 100 ml of culture was harvested onto Osmonics polycarbonate membrane filters (47 mm, 0.2 μm pore size) immediately after the UVBR treatment, and the filters were frozen at -135°C until analysed. In short, DNA was extracted from the filters using the same procedure as described in Buma et al. (2001), which is a method modified from Doyle & Doyle (1991). To remove RNA, the extracts were incubated for 1 h with 75 mg ml^{-1} RNase (Boehringer Mannheim) at room temperature. DNA concentrations of the extracts were determined fluorometrically using Pico-green dsDNA quantitation reagent (dilution 1:400, Molecular Probes) on a 1420 Victor multilabel counter (EG&G Wallac, excitation 485 nm, emission 535 nm). The amount of CPDs was determined using the immuno-dotblot procedure described in Boelen et al. (2001), employing a primary antibody (H3, Affitech) directed mainly to thymine dimers. Each blot contained 2 dilution series of standard DNA with known amounts of CPDs. Significance of all results was tested using paired *t*-tests at $p < 0.05$.

RESULTS

PAR acclimation affected the biovolumes of *Polarella glacialis* and *Phaeocystis antarctica*. For *P. glacialis*, the biovolume increased from $176 \pm 51.5 \mu\text{m}^3$ at a PAR irradiance of $18 \mu\text{mol m}^{-2} \text{s}^{-1}$ to $535 \pm 350 \mu\text{m}^3$ at $220 \mu\text{mol m}^{-2} \text{s}^{-1}$ PAR. For *P. antarctica* flagellates, this was $13.1 \pm 4.7 \mu\text{m}^3$ at $18 \mu\text{mol m}^{-2} \text{s}^{-1}$ and $19.8 \pm 11.8 \mu\text{m}^3$ at $220 \mu\text{mol m}^{-2} \text{s}^{-1}$, whereas the biovolumes of *Chaetoceros dichchaeta* (mean biovolume of $2319 \pm 106 \mu\text{m}^3$) and *Pyramimonas gelidicola* ($202 \pm 28 \mu\text{m}^3$) were not affected by irradiance conditions. PAR acclimation strongly affected the pigment content and/or ratio in all species. In addition to chlorophyll *a* (chl *a*) and β,β -carotene, which occurred in all species, the significant pigments detected were: *C. dichchaeta*—chl c_2 , chl c_3 , fucoxanthin, diadinoxanthin, diatoxanthin; *P. antarctica*—chl c_2 , chl c_3 , 19'-butanoyloxyfucoxanthin, 19'-hexanoyloxyfucoxanthin, 4-keto-19'-hexanoyloxyfucoxanthin, diadinoxanthin, diatoxanthin; *P. glacialis*—chl c_2 , chl c_3 , peridinin, diadinoxanthin,

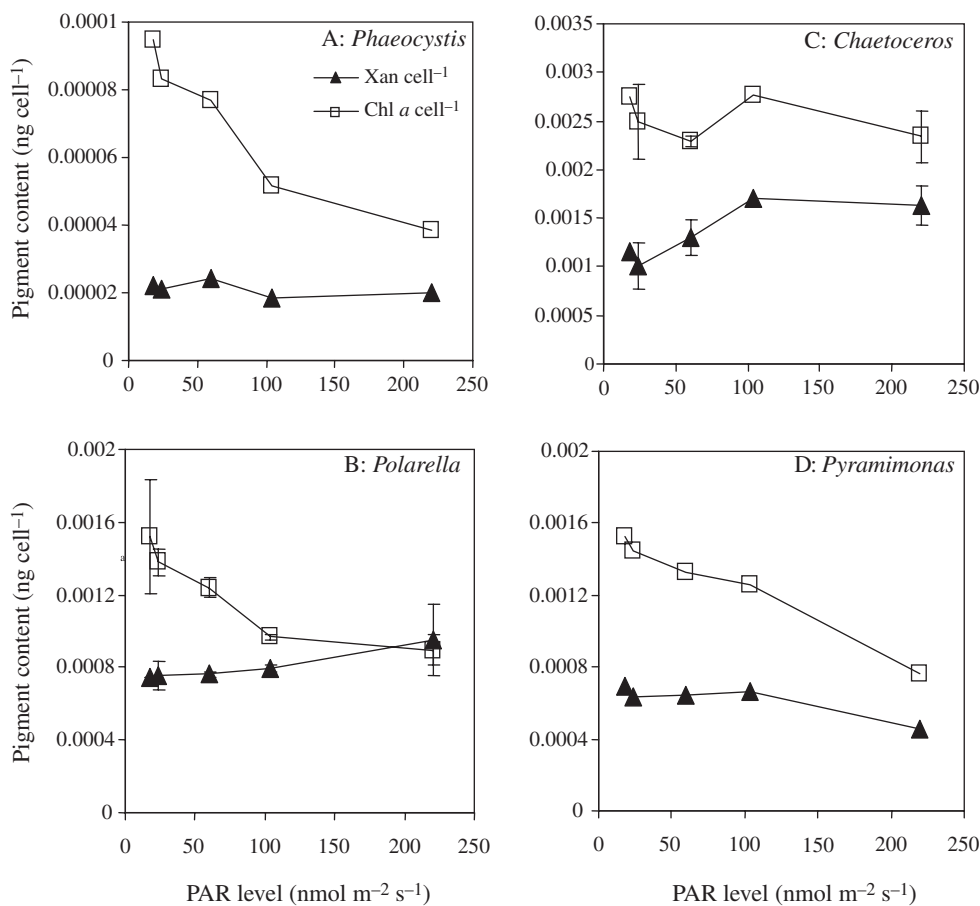


Fig. 2. Effects of PAR acclimation on pigment content in 4 Antarctic microalgae: (A) *Phaeocystis antarctica*, (B) *Polarella glacialis*, (C) *Chaetoceros dichchaeta* and (D) *Pyramimonas gelidicola* (\blacktriangle : total xanthophyll [Xan] pigments cell^{-1} ; \square : chlorophyll *a* [chl *a*] cell^{-1} ; error bars: $\pm\text{SE}$). Mean values of 2 replicate experiments are presented

diatoxanthin, dinoxanthin; and *P. gelidicola*—chl *b*, neoxanthin, violaxanthin, antheraxanthin (minor), zeaxanthin (minor), lutein, β,ϵ -carotene. With the exception of *C. dictyota*, cellular chl *a* content decreased sharply upon acclimation to the higher PAR intensities (Fig. 2). *C. dictyota* and *P. glacialis* showed increased cellular contents of the xanthophyll cycle pigments (diadinoxanthin and diatoxanthin) between the lowest and highest PAR intensities, with concentrations rising from 0.00075 to 0.00095 ng cell⁻¹ in *P. glacialis* and from 0.0011 to 0.0016 ng cell⁻¹ in *C. dictyota* (Fig. 2B,C). Acclimation to the highest irradiance level in *P. glacialis* resulted in a xanthophyll/chl *a* ratio >1 (Fig. 3B) due to a simultaneous decrease in chl *a* content and increase in cellular xanthophyll pigment concentrations. For the other 3 species, PAR acclimation

caused a significant increase in this ratio as well; however, these changes were not as pronounced as for *P. glacialis* (Fig. 3). Xanthophyll cycle activity was measured as the EPS (Fig. 3), which significantly increased upon acclimation to higher PAR levels. EPS values were lowest for *P. glacialis* (Fig. 3B).

A clear PAR-related induction of UV-absorbing compounds was found for *Phaeocystis antarctica* and *Polarella glacialis* (Fig. 4A,B), while *Chaetoceros dictyota* showed a maximum when cultivated at 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ (Fig. 4C). No UV-absorbing compounds were found in *Pyramimonas gelidicola* (not shown). Normalisation of UV-absorbing compounds/chl *a* content showed a significant increase for *P. glacialis*, increasing to 0.007 at the highest PAR level (Fig. 4B). *P. antarctica* also showed a clear increase;

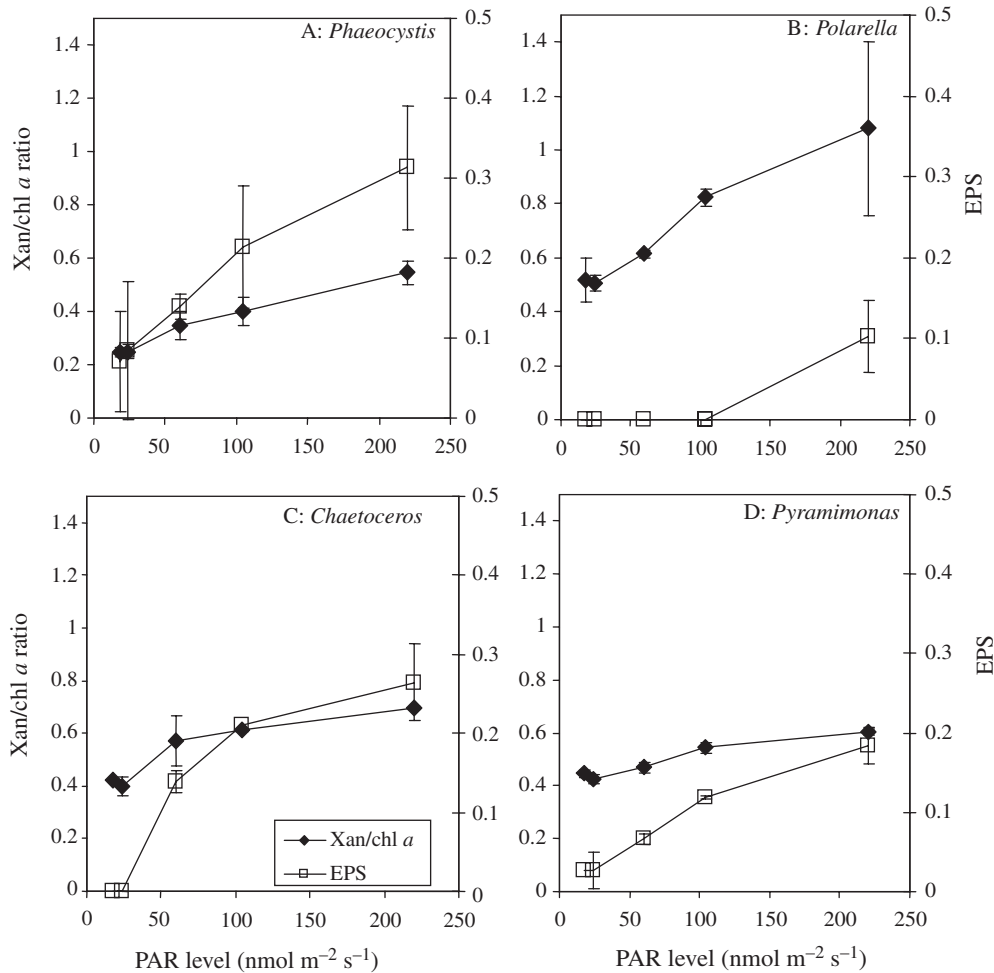


Fig. 3. Effects of PAR acclimation on xanthophyll pigments in 4 Antarctic microalgae: (A) *Phaeocystis antarctica*, (B) *Polarella glacialis*, (C) *Chaetoceros dictyota* and (D) *Pyramimonas gelidicola* (◆: ratio of total xanthophyll [Xan] pigments/chlorophyll *a* [chl *a*]; □: epoxidation state [EPS: epoxidated xanthophyll pigment relative to total xanthophyll cycle pool, i.e. violaxanthin/(violaxanthin + antheraxanthin + zeaxanthin) for *Pyramimonas* and diadinoxanthin/(diadinoxanthin + diatoxanthin) for other species tested]; error bars: \pm SE). Mean values of 2 replicate experiments are presented

however, the UV-absorbing compounds/chl *a* ratio remained much lower at the highest light intensity. Like the cell-specific UV-absorbing content, *C. dichæta* had a maximum UV-absorbing compounds/chl *a* ratio at around 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ (Fig. 4C). The strong UV absorption response of *P. glacialis* was further underlined when the UV-absorbing content was normalised to cell volume (Table 1).

With the exception of *Chaetoceros dichæta*, which showed little change, cellular concentrations of MDA (expressed as TBARs) were higher in high-PAR-acclimated cultures than in those exposed to low PAR irradiances (Table 1). Concentrations of MDA in

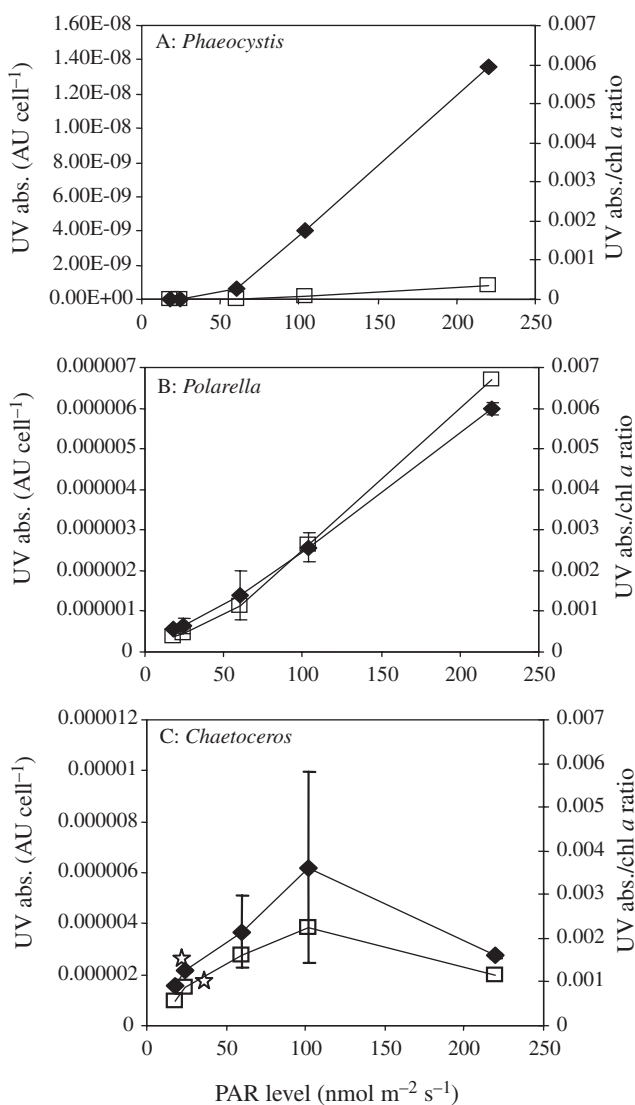


Fig. 4. Effect of PAR acclimation on UV-absorbing compounds (UV abs.): (A) *Phaeocystis antarctica*, (B) *Polarella glacialis* and (C) *Chaetoceros dichæta* (◆: UV abs. cell⁻¹; □: UV abs./chl *a* ratio; error bars: \pm SE). Mean values of 2 replicate experiments are presented (AU: arbitrary units)

Table 1. Biovolume-specific contents of UV-absorbing compounds (UV abs.), malondialdehyde (MDA) and chlorophyll *a* (chl *a*) in *Phaeocystis antarctica*, *Polarella glacialis*, *Chaetoceros dichæta* and *Pyramimonas gelidicola* under high and low PAR irradiance (HL: culture acclimated to high light [PAR irradiance 220 $\mu\text{mol m}^{-2} \text{s}^{-1}$]; LL: culture acclimated to low light [PAR irradiance 18 $\mu\text{mol m}^{-2} \text{s}^{-1}$])

Species	PAR condition	UV abs. (arbit. units)	MDA (mmol μm^{-3})	Chl <i>a</i> ($\mu\text{g } \mu\text{m}^{-3}$)
<i>C. dichæta</i>	HL	1.19×10^{-9}	2.00×10^{-11}	1.01×10^{-6}
<i>C. dichæta</i>	LL	6.6×10^{-10}	1.19×10^{-11}	1.19×10^{-6}
<i>P. gelidicola</i>	HL	0	4.44×10^{-13}	3.76×10^{-6}
<i>P. gelidicola</i>	LL	0	5.99×10^{-13}	7.55×10^{-6}
<i>P. glacialis</i>	HL	1.12×10^{-8}	1.94×10^{-12}	1.67×10^{-6}
<i>P. glacialis</i>	LL	4.19×10^{-10}	8.87×10^{-13}	2.84×10^{-6}
<i>P. antarctica</i>	HL	6.88×10^{-10}	6.51×10^{-10}	1.95×10^{-6}
<i>P. antarctica</i>	LL	0	2.44×10^{-10}	4.79×10^{-6}

Polarella glacialis, *C. dichæta* and *Phaeocystis antarctica* were around 10, 30 and 100 times higher, respectively, than in *P. gelidicola* (Table 1).

Increased EPS and MDA formation did not reduce growth of any of the species at the highest PAR level (220 $\mu\text{mol m}^{-2} \text{s}^{-1}$). However, light limitation reduced growth of all species at the lowest PAR levels (Table 2). UVBR exposure for 2 h caused significant CPD induction in *Pyramimonas gelidicola*, whereas no CPD induction was found for the other species. In *P. gelidicola*, CPD accumulation was observed for all PAR acclimation levels, whereas CPD accumulation increased >4-fold at higher PAR acclimation levels (Fig. 5).

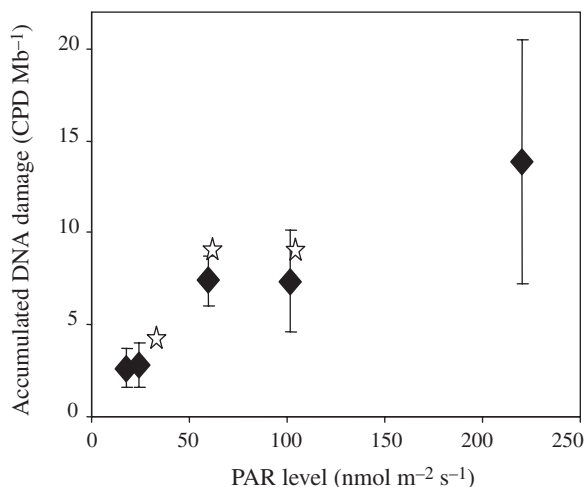


Fig. 5. *Pyramimonas gelidicola*. Effect of PAR acclimation on cyclobutane pyrimidine dimer (CPD per Mega base [Mb]) accumulation. Mean values of 2 replicates are presented (error bars: \pm SE; stars: significant differences compared with the highest irradiance condition [*t*-test, $p < 0.05$])

Table 2. Specific growth rates (d^{-1}) of the 4 Antarctic microalgal species after PAR acclimation

PAR level ($\mu\text{mol m}^{-2} \text{s}^{-1}$)	<i>Chaetoceros dichaeta</i>	<i>Polarella glacialis</i>	<i>Pyramimonas gelidicola</i>	<i>Phaeocystis antarctica</i>
220	0.295 \pm 0.006	0.229 \pm 0.031	0.225	0.512 \pm 0.024
104	0.259 \pm 0.024	0.161 \pm 0.030	0.182	0.464 \pm 0.069
60	0.273 \pm 0.001	0.225 \pm 0.068	0.170	0.337 \pm 0.015
24	0.267 \pm 0.010	0.132 \pm 0.040	0.110	0.237 \pm 0.129
18	0.199 \pm 0.063	0.111 \pm 0.114	0.075	0.292 \pm 0.013

DISCUSSION

The present study has shown inter- and intra-specific differences in vulnerability to accumulated UVB-induced DNA damage in Antarctic microalgae. Our experimental set-up excluding UVR and PAR from the UVBR treatments was chosen because active photoregulation, photoprotection, as well as DNA repair during UVBR exposure had to be avoided. Therefore, results cannot be extrapolated to the field; the experiment was merely designed to collect fundamental information on PAR acclimation and subsequent CPD induction vulnerability in relation to pigmentation and UV absorption characteristics.

No CPD accumulation was found for *Polarella glacialis*, *Chaetoceros dichaeta*, or *Phaeocystis antarctica* at any PAR acclimation level, whereas CPDs accumulated in *Pyramimonas gelidicola*, with the highest CPD accumulation at the highest PAR acclimation levels. Seemingly in contrast, other studies we recently performed showed increased UVBR tolerance (PSII efficiency and viability) in high-PAR-acclimated diatoms (Poll et al. 2005). Lower UVBR vulnerability was also found, when cells had been acclimated to high PAR first, compared with low-light-acclimated cultures (Neale et al. 1998b, Litchman & Neale 2005). In a natural Antarctic phytoplankton community, UVR effects were lower during noontime hours than during morning hours, suggesting that UVBR protection is induced during the course of the day (Figueroa et al. 1997). Other field observations revealed that low-light-acclimated Antarctic phytoplankton communities were more vulnerable to UVR photoinhibition than high-light-acclimated communities (Helbling et al. 1994, Neale et al. 1994, Villafane et al. 1995, Bracher & Wiencke 2000). It is clear that discrepancies between observed UVBR effects, whether or not in relation with the biotic or abiotic environment, are primarily determined by the parameter under study (growth, viability, photosynthetic rate or efficiency, DNA damage) (Buma et al. 2003, Villafane et al. 2003).

In the current experiments, all species acclimated to increasing PAR by decreasing their light-harvesting

pigment content and by increasing both xanthophyll/chl *a* ratios as well as EPS. Increases in xanthophyll/chl *a* ratios are commonly observed in microalgae and higher plants subjected to high irradiance, including UVR (Paerl et al. 1983, Demers et al. 1991, Demmig-Adams & Adams 1992, Buma et al. 2000). *Polarella glacialis*, in particular, was shown to be well equipped for high irradiance exposure, judging from its very high xanthophyll/chl *a* ratio, its low EPS value and very high content of UV-

absorbing compounds at the highest PAR acclimation levels. These characteristics clearly match the natural habitat of *P. glacialis*, since this species is found primarily in Antarctic sea ice, particularly in the brine of the upper fast ice (Montresor et al. 1999, Thomson et al. 2004). Here, irradiance conditions (including UVR) are consistently high in comparison with open, mixed waters or the under-ice environment. Furthermore, high UV absorbance is consistent with the observations made by Jeffrey et al. (1999) who demonstrated high structural MAA levels in a large number of dinoflagellate species. These and other observations (Vernet et al. 1989, Neale et al. 1998b) indicate that bloom-forming dinoflagellates and other microalgae are well equipped to grow at high natural irradiance levels, although a good correlation was not always found between MAA content and collection depth in dinoflagellates (Banaszak et al. 2000). In addition, the observed low intracellular MDA concentrations in *P. glacialis* further seem to indicate that intracellular metabolic activities or other antioxidant actions prevent rapid accumulation of harmful oxiradicals that could otherwise oxidise cellular membranes.

Except for *Pyramimonas gelidicola*, all species showed a clear induction of UVB-absorbing compounds with increasing irradiance of PAR acclimation, as described before for other Antarctic microalgae, such as *Phaeocystis* sp. (Riegger & Robinson 1997, Moisan & Mitchell 2001) or other prymnesiophytes and dinoflagellates (Carreto et al. 1990, Hannach & Sigleo 1998). Earlier work (Marchant et al. 1991) had shown undetectable amounts of MAAs in *Phaeocystis antarctica* flagellates when cultivated under low PAR irradiance, which was confirmed in our experiments. Davidson & Marchant (1992) and Peperzak et al. (2000) proposed that the flagellate stage in the life cycle of *Phaeocystis* spp. is an escape mechanism from the confines of colonial metabolism, the latter indicating that this life stage was initiated by low light. We found that photoacclimation of this alga to a range of PAR irradiances elicited the greatest percentage change in concentrations of intracellular UV absorbance and chl *a* and the greatest growth rate range. Thus, *P. antarctica* flagellates had a high capacity to acclimate to changes in the light climate. If indicative of

the photophysiology of *Phaeocystis* spp. flagellates in general, this plasticity may contribute to the reported persistence of this life stage throughout the year (Parke et al. 1971, Veldhuis et al. 1986, Verity et al. 1988) and the ubiquity of *Phaeocystis* spp. in temperate and polar waters (Davidson & Marchant 1992).

Although a PAR-related difference was found in MDA levels for all species except *Pyramimonas gelidicola* (Table 1), we question whether these increases indicate substantial oxidative stress under our experimental conditions. Other marine studies showing enhanced MDA concentrations at excess irradiances were performed under (UV) irradiance levels much higher than ours (Malanga et al. 1997, Butow et al. 1998, Rijstenbil 2001, 2002, Bischof et al. 2003). Since oxidative stress is thought to be an imbalance between oxiradical action and antioxidative responses, it would be expected that growth rates at the highest irradiance level were affected. However, this was not observed (Table 2). The differences in MDA concentration may merely reflect differences in photosynthetic activity, thereby generating higher intracellular concentrations of radicals without causing chronic stress. In support of this, *Phaeocystis antarctica* growth was very rapid at the highest light intensities, while MDA concentrations were up to 2 magnitudes higher than the levels found in other species (Table 1). Therefore, the observed MDA levels might result from active cell metabolism, without impairment of photosynthesis or growth.

For some MAAs, antioxidant properties have been suggested in addition to their UV-absorbing function (Dunlap & Yamamoto 1995). However, our analytical spectrophotometric procedure did not characterise the various UV-absorbing compounds. *Pyramimonas gelidicola* was the only species that accumulated CPDs, whereas, at the same time, it did not synthesise UV-absorbing compounds. It has to be stated that *P. gelidicola* might have induced UV-absorbing compounds under different spectral irradiance conditions (UVAR or a higher proportion of blue wavelengths). For this reason, *P. gelidicola* may show a different UVBR response under natural solar irradiance conditions. Our results with *P. gelidicola* could indicate a protective role of UV-absorbing compounds against CPD accumulation. On the other hand, no CPD accumulation was found in *Phaeocystis antarctica* cells that were acclimated to low PAR irradiance, which contained undetectable amounts of UV-absorbing compounds. Therefore, inter-specific vulnerability for UVBR-induced DNA damage may be due to other factors, such as structural differences in CPD repair capacity (Karentz et al. 1991, Poll et al. 2001) or other photoprotection mechanisms.

It has been postulated before (reviewed in Buma et al. 2003) that larger cells would be less vulnerable to

CPD induction than smaller cells, because of the differences in the intracellular light path (Karentz et al. 1991, Buma et al. 2001). However, the present study shows that other features must also be involved. We found CPD were not accumulated in the smallest species (*Phaeocystis antarctica*), but were accumulated in *Pyramimonas gelidicola* that had a 10 times larger biovolume (Fig. 5). Photorepair enzymes have been reported in marine red macroalgae inhabiting littoral areas exposed to high irradiance as opposed to sublittoral species. This hints at a strong habitat-related vulnerability to UVBR (Poll et al. 2001), as has also been suggested for microalgae (Davidson 2006). It is likely that organisms adapted to high mean irradiance habitats are well equipped to simultaneously overcome high PAR, UVAR and UVBR stress. For example, MAAs are found to be present in a range of microalgae, even when cultured under relatively low PAR levels (Jeffrey et al. 1999). This supports our observations of UV absorbance in *Chaetoceros dichæta* and *Polarella glacialis* under PAR conditions as low as $18 \mu\text{mol m}^{-2} \text{s}^{-1}$. Contrasting with other species within the genus, *P. gelidicola* appeared less equipped for high irradiance environments, as xanthophyll cycle pigments decreased at the highest irradiances and other protective mechanisms such as UV-absorbing compounds were not induced.

Finally, intracellular chromophores can absorb UVBR photons, thereby reducing DNA damage via reduced intracellular UVBR transmission. The strong decline in cellular chl *a* content in high-PAR-acclimated *Pyramimonas gelidicola*, combined with a low photoprotection capacity (including the absence of UV-absorbing compounds), may therefore have contributed to enhanced CPD formation. Also, the higher growth rate at high PAR irradiances could have resulted in a higher proportion of DNA being unfolded for DNA replication (DNA synthesis phase). The complexity of photobiological responses in Antarctic microalgae needs to be further elucidated, especially given the observed shifts in the light climate in the Southern Ocean.

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