Seasonal cycles in phytoplankton mycosporine-like amino acids and the attenuation of ultraviolet radiation in coastal surface waters in Sagami Bay

SATORU TAGUCHI1*, KEN-ICHI YAMAO1,2, YOSHINORI YAMADA1,2, JUN-ICHI HAGIMOTO1,2, AKIRA TAKEUCHI1,2, KIONOBU CHIBA1,2, TOMOYO KATAYAMA1, SHOZO MOTOKAWA1, AI MURATA1 & HITOMI Taira1

1 Laboratory of Biological Oceanography, Soka University, 1–236 Tangi-Cho, Hachioji, Tokyo 192–8577, Japan
2 Mailing address, 1238–1–306 Ishikawa-Cho, Hachioji, Tokyo 192–0032, Japan

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Abstract: The concentration of mycosporine-like amino acids (MAAs) of phytoplankton and the attenuation coefficients of ultraviolet radiation (UVR) were determined in coastal surface waters in Sagami Bay during the period from 2004 to 2012 to examine the seasonal cycle. Four major MAAs, including mycosporine-glycine, palythine, shinorine, and porphyra-334, were detected throughout the year. Mycosporine-glycine was the most dominant type, reaching more than 100 nM, whereas porphyra-334 was the least dominant type, barely reaching 20 nM. The total concentration of the four MAAs was high in the high water temperature (WT) months (February–July) and low in the low WT months (August–January). The seasonal cycle was repeatedly sustained for six years. The attenuation of ultraviolet radiation at two wavelengths, 320 and 340 nm, also followed a similar seasonal cycle: high in the warm period and low in the cool period. The attenuation of UVR at two wavelengths ($k_{320}$ and $k_{340}$) was linearly correlated with the concentration of MAAs ($p<0.01$). Solar energy was the environmental driving force responsible for the seasonal cycles of MAAs, which were further enhanced by increased biomass and nutrient availability. Individual groups in the phytoplankton community might share roles to protect the entire community from UVR in the marine ecosystem by producing various MAAs.

Key words: Chl a, mycosporine-glycine, palythine, PAR, porphyra-334, shinorine, UVR attenuation

Introduction

High levels of ultraviolet radiation (UVR) are environmentally stressful to phytoplankton (e.g. Helbling & Zagarese 2003). An increase in solar UVR (from 280 to 400 nm) and a high UVR proportion compared to total solar photosynthetically active radiation (PAR from 400 to 700 nm) in addition to UVR is expected in coastal water ecosystems at mid-latitudes (e.g. Sugawara et al. 2003). Solar UVR is considered to be a major controlling factor for the abundance of mycosporine-like amino acids (MAAs) throughout the year (Llewellyn & Harbour 2003). The downwelling solar irradiance of UVR ranges from 45 µW cm⁻² in the summer and 25 µW cm⁻² in the winter in the western mid-North Pacific (Kuwahara et al. 2015). However, both the quantity and quality of UVR are altered when UVR penetrates into surface seawater (Kirk 1994, Hargreaves 2003). Consequently, the attenuation of UVR at a given wavelength should be partly affected by the concentrations of MAAs in the water column. In the water column, the attenuation of UVR is primarily regulated by dissolved organic matter (DOM) and particulate matter (PM). Dissolved MAAs are included in DOM, and the cellular MAAs of phytoplankton are included in PM. The attenuation of PAR is primarily regulated by the cellular pigment contents of phytoplankton in the surface mixed layer (Kirk 1994). Dissolved MAAs are an exudate of DOM released from phytoplankton (Vernet & Whitehead 1996). The cellular MAAs are primarily synthesized de novo via the shikimate pathway (Shick & Dunlap 2002). Phytoplankton intercellular MAAs are also partially responsible for regulating the attenuation of UVR in the surface mixed layers. The origin of MAAs in coastal waters is primarily from cyanobacteria (Sinha et al. 2001), micro-
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algae (Jeffrey et al. 1999), and red algae (e.g. Takano et al. 1979, Tsujino et al. 1980). The accumulation of MAAs in algal cells might enhance UVR attenuation within the cell, primarily between 310 and 340 nm, thus providing a UVR shield for DNA (Buma et al. 1995). Although the short wavelengths of UVR, particularly 280–315 nm (UVB), are more responsible than the long wavelengths of UVR, such as 315–400 nm (UVA), for DNA damage in cells (Helbling et al. 2001), UVB is attenuated more rapidly than UVA in surface mixed layers in coastal waters at temperate latitudes (Kuwahara et al. 2015). Different effects of UVR can be expected in the surface mixed layer depending on the wavelength because the shorter wavelengths cause severe detrimental effects.

A high biomass of microalgae is typically observed during the warm season at temperate latitudes. During the period of high biomass, the concentration of MAAs in surface waters is also expected to increase, reflecting the high abundance of dinoflagellates and diatoms (Llewellyn & Harbour 2003). Dinoflagellates and diatoms are the most active taxonomic groups for the synthesis of MAAs in the surface mixed layers of coastal waters (Jeffrey et al. 1999). UVR-protective MAAs have been observed in twenty forms in marine environments (Llewellyn & Airs 2010). Although species-specific MAAs can be expected in the sea, mycosporine-glycine, shinorine, palythine, porphyra-334, palythenic acid, and palythene are particularly considered to be the dominant MAAs (Llewellyn & Airs 2010). The first four MAAs are also common in seven toxic dinoflagellates (Carreto et al. 2001). The effects of UVR on the synthesis of primary MAAs, such as mycosporine-glycine, shinorine, and porphyra-334 are positive, but not necessarily positive on the synthesis of secondary MAAs, such as palythine (Shick 2004). MAAs are synthesized under environmentally stressful conditions, such as high UVR. Not only is an increase in solar UVR expected during the summer in coastal waters, but a high UVR component of total radiation (PAR+UVR) is also expected (Ichikawa & Taguchi 2001, Sugawara et al. 2003). The period of high UVR levels typically coincides with high temperature and low nutrient availability in temperate coastal waters. The concentration of MAAs might increase when elevated temperatures are combined with increased UV exposure (Shick & Dunlop 2002). Low nitrogen availability under exposure to UBV has been reported to depress the synthesis of cellular MAAs in diatoms (Lesser et al. 1994), dinoflagellates (Litchman et al. 2002) and macroalgae (Korbee et al. 2005).

MAA concentrations are expected to be high during warm periods and low during cool periods, as indicated by the absorption of phytoplankton MAAs in the UV range in the ocean (Morrison & Nelson 2004). This prediction likely reflects the linear relationship between the Chl a biomass, an indicator of the phytoplankton biomass, and the concentration of MAAs in natural assemblages of phytoplankton (Carreto & Carignan 2011). The objectives of the present study were two-fold: to determine the seasonal cycle of phytoplankton MAAs in temperate coastal waters with respect to the Chl a biomass and the stability-reproducibility of the seasonal cycle of phytoplankton MAAs over six years in the western mid-North Pacific, and to determine UVR regulation by phytoplankton MAAs in the same region.

Materials and Methods

Sampling

Surface water was collected using a bucket at St. M, located at 35°09′N and 139°10′E off Manazuru Harbour, Sagami Bay, in central Japan (Fig. 1). Monthly cruises were conducted onboard the R.V. Tachibana of Yokohama National University during the periods from May 2004 to January 2007 and May 2011 to May 2012. The surface water was pre-screened through a 183-µm mesh plankton net cloth and stored in carboys in a dark cold room at 4°C. As soon as the samples were returned to the laboratory, the subsamples were filtered onto 25-mm glass fiber filters (GF/F, Whatman, Dallas, USA) to analyze the Chl a concentrations and 47-mm Whatman glass fiber filters (GF/F) for the analysis of MAAs. The analyses of Chl a and MAAs concentrations were conducted within three months after sampling. The filtrate was passed through Millipore membrane filters (Milex HA, 0.2-µm pore size, Whatman, Dallas, USA) for the analysis of nutrients. All filtrate samples were stored in vials below −20°C under dark conditions.

Optical measurements

Solar irradiation was measured using a PUV-500 submersible radiometer and a PUV-510 surface radiometer ( Biospherical Instruments Inc., USA), which provided measurements of cosine-corrected downwelling irradiance at St. M (Fig. 1). The instruments were designed to measure absolute UVR irradiance values with 10-nm bands centered at 320 and 340 nm. The mean surface irradiance just under the surface (∼0) was determined from the data collected during the underwater profile measurements. Vertical profiles of temperature and salinity were determined using a factory-calibrated conductivity-tempera-
ture-depth sensor (JFE Advantech, ASTDI100, Tokyo, Japan). Water density was estimated as sigma-t ($\sigma_t$, kg m$^{-3}$), and the surface mixed layer depth was determined based on a threshold value (0.15 kg m$^{-3}$) larger than the average of the first 5 m.

Irradiance profiles in the surface mixed layer (down to 40 m depth at 0.5 m per second) were obtained to determine the attenuation of UVR. The data were collected at 1-sec intervals during the profiling. The measured irradiance profiles of the down-cast were recorded. All surface and profile casts were conducted within 3 h of solar noon, during which special attention was paid to the shadow of the ship. Dark corrections, according to Biospherical Instruments’ protocols, were conducted prior to every profile. The sub-surface radiometer was consistently lowered into the water prior to measurements to minimize temperature effects. The current drag and tilt of the submersible sensor was maintained at a minimum during the study. Special attention was also paid to the rate at which the radiometer was lowered for each profile.

Diffuse attenuation coefficients

The data were cleaned to remove data points either outside the detection limit of the radiometer sensors or near the surface. Diffusive attenuation coefficients ($K_d$) for the downwelling irradiance at each wavelength were subsequently determined from the slope of the linear regression of the natural logarithm of downwelling irradiance at each respective wavelength ($i$) against the depth, assuming that solar irradiance reduced exponentially,

$$E_d = E_d(-0)e^{-K_d Z}$$  \hspace{1cm} (1)

where $E_d$ is the downwelling irradiance at depth $Z$ and $E_d(-0)$ is the downwelling irradiance just below the surface.

Analysis

NUTRIENTS

Concentrations of nitrate, phosphate, and silicate were determined using a Bran+Luebbe Autoanalyzer model AACS II according to Parsons et al. (1984).

CHLOROPHYLL PIGMENTS

Cellular material on the 25-mm Whatman GF/F glass fiber filters was extracted in opaque vials using N,N-dimethylformamide for 24 h at 4°C (Suzuki & Ishimaru 1990). The chlorophyll pigments were analyzed using a Turner Design Fluorometer model 10-AU according to Holm-Hansen et al. (1965) for the samples collected during the period from May 2004 to January 2006. For the samples collected during other periods, the extracts were sonicated for 10 min in an ice bath (International Electric Elutekku, Sine Sonic 150, Japan) and filtered through a membrane filter with a 0.2-µm pore size (Millipore, Millex-GC, USA) for analysis on a high performance liquid chromatograph (HPLC). The filtrates were analyzed by HPLC (Beckman Coulter, 168 Diode Array Detector, USA) with a C18 reverse-phase Ultrasphere 3 mm column, using a solvent gradient based on the modified method of Wright et al. (1997). The two instruments were intercalibrated to confirm the significant relationship.

MICOSPORINE-LIKE AMINO ACIDS

Cellular material on the 25-mm Whatman GF/F glass fiber filters was extracted in 1.5 mL of 20% (v/v) methanol (HPLC grade, Wako, Japan) for at least 24 h at 37°C in the dark, after sonication of the harvested cell materials in an ice bath for 10 min at full strength. The methanolic extract was subsequently centrifuged at 10,000 rpm for 5 min. The supernatant (1 mL) was evaporated under reduced pressure and re-dissolved in 1 mL of 100% methanol. The extracts were vortexed and centrifuged, and the supernatant was carefully separated. The supernatant was evaporated until completely dry at 45°C. The residue was re-dissolved in 100 µL of MilliQ water and filtered through a 0.2-µm membrane. Further analysis was performed by HPLC using a CAPCELL PAK C18 UG120 5 µm (250 mm×4.6 mm I.D., Shiseido, Tokyo, Japan) and Shiseido SG 120 5-µm guard column (35 mm×4.6 mm I.D. Japan). During each run, the column was equilibrated with 100% solvent A [80% Milli-Q water and 20% (v/v) 0.5 M ammonium acetate (Wako, Japan)] at a flow rate of 1 mL min$^{-1}$, followed by 100% eluent B [75% Milli-Q water, 25% (v/v) methanol and 0.2% acetic acid (Wako, Japan)] for 20 min at a room temperature. Peak absorbance was detected at 310 and 334 nm. The MAA concentrations were calculated from the peak areas detected at 334 nm using the standard calibrated peak areas for individual MAAs. Common MAAs were identified based on the retention time, absorption spectra from 290 to 400 nm, and co-chromatography with authentic standards, shinorine, palythine, and porphyra-334 obtained from the red algae Chondrus yendoi, Chondrus vancouveriensis, and Porphyra yezoensis, respectively (Taira et al. 2004a). Mycosporine-glycine was obtained from cultures of the dinoflagellate Alexandrium tamarense (Taira et al. 2005b). The concentrations of MAAs were calculated using the following molar extinction coefficients: shinorine: $\varepsilon_{334}$=44,700 (Tsujino et al. 1980), palythine: $\varepsilon_{330}$=36,200 (Takano et al. 1978), porphyra-334: $\varepsilon_{334}$=42,300 (Takano et al. 1979), and mycosporine-glycine: $\varepsilon_{330}$=28,100 (Ito & Hirata 1977). The ratios of MAAs to Chl $a$ were calculated based on the total MAAs in nM and Chl $a$ in ng L$^{-1}$.

STATISTICAL ANALYSIS

The means and standard deviations were calculated based on subsamples analyzed at least in triplicate. Significant differences between seasons were tested for using the Student’s t-test for unpaired comparisons and a paired t-test, the Wilcoxon signed-rank test for paired comparisons, and the Kruskal–Wallis one-way analysis of variance on ranks for paired multiple comparisons using the Sigma-Plot software program (System Software, version 11.0, San Francisco, USA).
Results

Attenuation of UVR

The intensity of UVR at 320 and 340 nm decreased rapidly with depth as it was rapidly attenuated, and an optical depth of 10% was typically shallower than the deepest extent of the surface mixed layer, ranging from approximately 12 m in the summer to 82 m in the winter. The attenuation of UVR, indexed based on the extinction coefficient of downwelling UVR ($K_d(\lambda)$), indicated a seasonal cycle with an increase in low water temperature months (February–July) and a decrease in high water temperature months (August–January), although the attenuation was variable among years (Fig. 2). The highest attenuation was approximately 0.98 m$^{-1}$ at 320 nm and 0.68 m$^{-1}$ at 340 nm, which was always observed in the high water temperature months. The lowest attenuation was approximately 0.16 m$^{-1}$ at 320 nm and 0.13 m$^{-1}$ at 340 nm, observed during the low water temperature months.

Temperature, salinity, and nutrients

The surface temperature rapidly increased from April to a maximum of 28°C in July and gradually decreased to a minimum of 14°C in February (Fig. 3A). The surface salinity was low (ca. 31.5) at the highest temperature and high (ca. 35.5) at the lowest temperature (Fig. 3B). The high (February–July) and low (August–January) water temperature period were almost equivalent in length. In contrast with the harmonic cyclic changes in the surface temperature, the seasonal changes in nutrients were highly variable between years (Fig. 4A, B, and C). The highest concentrations of nitrate, phosphate, and silicate were 7.01 µM
The concentrations of mycosporine-glycine, shinorine, palythine, and porphyra-334 indicated highly variable seasonal changes between years (Fig. 5). Mycosporine-glycine occurred at the highest (130 nM) and lowest concentrations (0.1 nM) in the summer and winter, respectively, but there was no stable seasonal trend, although concentrations of the other three MAAs exhibited a strong seasonal trend, increasing in high (February–July) and decreasing in low (August–January) water temperature periods. The concentrations of palythine ranged from 0.5 nM in the winter to 90 nM in the summer. The concentrations of shinorine ranged from 0.4 nM in the winter to 70 nM in the fall. The concentrations of porphyra-334 ranged from 0.1 nM in the winter to 27 nM in the summer. The concentrations of total MAAs were generally less than 20 nM during the period from November to March, and values larger than 10 nM were observed during the period from April to October (Fig. 6A). The Chl a concentrations in these three periods were 0.4–6, 0.4–8, and 0.3–9 mg Chl a m$^{-2}$, respectively (Fig. 6B). The total variation of phytoplankton MAAs was on the order of 10$^2$, and the total variation in the Chl a concentrations was also of the order of 10$^2$. The total concentrations of MAAs (nM) were significantly correlated with the concentrations of Chl a (mg m$^{-3}$) (MAAs=40 Chl a+13, r=0.39, p<0.05). The highest relative proportion of mycosporine-glycine versus total MAA concentrations reached 90%, while shinorine, palythine, and porphyra 334 reached 85%, 75%, and 37%, respectively. The respective relative abundances of MAAs were highly variable throughout the year (Fig. 7). A high relative abundance of mycosporine-glycine was observed during the warm period, while a high relative abundance of shinorine was observed during the cool period. All pairwise multiple comparison tests indicated significant relationships between MAAs/Chl a, $K_d$(320), $K_d$(340), nitrate, and phosphate ($p<0.05$).

**Discussion**

**MAAs**

There are more than 20 characterized MAAs with maximum absorption peaks ranging from 309 to 362 nm (Carreto & Carignan 2011). In this study, four MAAs, specifically palythine, mycosporine-glycine, porphyra-334, and shinorine, were positively identified, although some unidentified peaks were also observed in the analyses. These peaks were not able to be assigned to specific MAAs, reflecting the uncertainty in peak positions and lack of standards available for laboratory studies. In coastal waters, two additional MAAs, palythene and palythenic acid, are commonly identified in addition to the four MAAs listed above (Llewellyn & Airs 2010). The inclusion of these two
MAAs in the present MAAs analysis would undoubtedly improve current understanding of the natural distribution and abundance of MAAs and UVR penetration in the sea, although the relative abundance of the two combined MAAs is expected to be 20% at most (Llewellyn & Airs 2010).

Seasonal cycle

Seasonal cycles are common in biological systems, with amplitude and phase varying according to latitude (Demarcq et al. 2012). In most boreal waters, the biomass of phytoplankton, measured through the proxy of Chl $a$ concentration, increases during warm periods and decreases during cool periods. However, during the present study the Chl $a$ concentration fluctuated remarkably (10-fold at maximum) during the year. The seasonal patterns of Chl $a$ obtained in the present study were similar to those that have been previously observed in the region (Kuwahara et al. 2015). When Chl $a$ was present in the surface mixed waters, the presence of phytoplankton MAAs was also confirmed throughout the year, not only in the western North Pacific Ocean but also in other mid-latitude coastal waters (e.g., Llewellyn & Mantoura 1997, Whitehead & Vernet 2000, Riemer et al. 2007).

The amplitude and phase of the seasonal cycle of the total concentrations of MAAs has been reported to be associated with the attenuation of solar UVR (Zepp 2003). In addition, Llewellyn & Harbour (2003) and Riemer et al. (2007) clearly confirmed a seasonal pattern, in that the maximum in total MAA concentrations occurred in summer in phytoplankton from the surface mixed layer of the English Channel and off Otago coast, New Zealand, respectively. The summer maximum in total MAA concentrations also coincided with the summer maximum in UV absorption by phytoplankton in a multiyear seasonal study by Morrison & Nelson in Bermuda (2004).

The ratio of total MAAs, represented as the four major MAAs investigated in the present study, (i.e. mycosporine-glycine, shinorine, palythine, and porphyra-334) to the Chl $a$ concentration (0.048±0.055 nM MAAs [ng Chl $a$] $^{-1}$) might provide some insight into the seasonal cycle of MAAs, as suggested by Carreto & Carignan (2011). These four MAAs were present throughout the years examined in the present study, although the relative abundances were highly variable, presumably as a result of the variable community structure of the phytoplankton. The weak relationship between the total concentration of MAAs and Chl $a$ might suggest variable photo-adaptation and photo-acclima-

![Seasonal variations in the concentrations of mycosporine-glycine (A), palythine (B), shinorine (C), and porphyra-334 (D). Symbols indicate the values in 2004 (solid circles), 2005 (open circles), 2006 (reversed solid triangles), 2007 (open triangles), 2011 (solid squares) and 2012 (open squares).](image-url)
Ratios of MAAs/Chla were significantly correlated with concentrations of phosphate and nitrate ($p < 0.05$) in the present study. This suggests nutrient-dependency of MAAs/Chla, as observed in diatoms (Lesser et al. 1994) and dinoflagellates (Litchman et al. 2002).

The ratios of MAAs to Chla ($0.048\pm0.055$ nM MAAs [ng Chla $^{-1}$]), which are analogous to the slope of the relationship as mentioned above, agreed well with those described in previous studies in natural assemblages of phytoplankton in the mixed layer. In natural assemblages of phytoplankton, the spring ratio of MAAs to Chla ranged from 0.00067 to 0.15 nM MAAs (ng Chla $^{-1}$) in the coastal waters off California (Whitehead & Vernet 2000), 0.0073 nM MAAs (ng Chla $^{-1}$) in the estuary of the Chesapeake Bay (Banaszak & Neal 2001), from 0.0007 to 0.0027 nM MAAs (ng Chla $^{-1}$) in the Southern Ocean (Moisan & Mitchell 2001), and from 0.00011 to 0.016 nM MAAs (ng Chla $^{-1}$) in the English Channel (Llewellyn & Harbour 2003). A high ratio of 0.92 nM MAAs (ng Chla $^{-1}$) was observed in the oligotrophic subtropical waters of the Brazil Current (Carreto & Carignan 2011). The variability in the ratios obtained from natural assemblages of phytoplankton reflects two types of information concerning the species composition responsible for total MAA and Chla concentrations as suggested by Bancroft et al. (2007). In diatom-dominated assemblages mycosporine-glycine has been reported to be consistently abundant, with porphyrins334 and shinorine occasionally being observed (Carreto & Carignan 2011). Diatoms such as Thalassiosira spp. and Coscinodiscus spp. were abundant in the summer in this region, in terms of relative biovolume versus total phytoplankton cells (Mori & Taguchi, unpublished data). Even in cultured populations of dinoflagellates, the ratio of MAAs to Chla can be highly variable, ranging from 0.019$\pm$0.001 pM MAAs (pg Chla $^{-1}$) for Scrippsiella sweeneyae (Taira et al. 2004b) to 0.12 pM MAAs (pg Chla $^{-1}$) for Alexandrium catenella (Carreto et al. 2001).

**Fig. 6.** Seasonal variations in the total MAA concentrations (A) and Chla concentrations (B). Symbols indicate the values in 2004 (solid circles), 2005 (open circles), 2006 (reversed solid triangles), 2007 (open triangles), 2011 (solid squares) and 2012 (open squares).
study, were within the range of the lowest values (0.16 m$^{-1}$ and 0.10 m$^{-1}$) yet reported (Boelen et al. 2001) to the highest reported values (obtained in tropical coastal water) of 1.45 m$^{-1}$ and 0.99 m$^{-1}$, respectively (Garza & Suttle 1998). The total attenuation of UVR depends on attenuation due to water, DOM, detritus, and phytoplankton, following the equation:

$$K_d(\text{UVR}) = K_d(\text{water}) + K_d(\text{DOM}) + K_d(\text{detritus}) + K_d(\text{phytoplankton})$$

(2)

where $K_d$ is downwelling attenuation (m$^{-1}$).

At the present sampling location, detritus and DOM from local river outflow immediately diffuse into the surrounding water upon entering Sagami Bay (Koyama et al. 2007), and these components have been found not to directly influence the seasonal attenuation of UVR or nutrient concentrations at St. M (Kuwahara et al. 2015). The attenuation of UVR is primarily controlled by DOM and MAA concentrations in ambient water and within phytoplankton cells (Moisan & Mitchell 2001). This supposition is widely subscribed to, although few in situ measurements of the attenuation of UVR in relation to phytoplankton MAA concentrations have, in fact, been simultaneously conducted. Although UV-absorption at 330 nm was only nominally correlated with the particulate concentration of MAAs off the southern California coast (Whitehead & Vernet 2000), the significant relationships between the total concentrations of MAAs and the attenuation coef-
coefficients of UVA and UVB ($p<0.01$, Fig. 8) suggest that phytoplankton MAAs absorbed UVA (ca. 340 nm) in the present study. The absorption peaks of the four MAAs determined in the present study ranged from 320 to 334 nm. The maximum absorption of UVR was reported to range from 313 to 335 nm for oceanic waters near Bermuda (Morrison & Nelson 2004), although dissolved MAAs could also influence attenuation (Whitehead & Vernet 2000).

**Ecological significance of MAAs**

The intracellular MAAs of microalgae contribute to the attenuation of UVR in the surface mixed layer year-round, in addition to the effects of dissolved MAAs. MAAs in microalgae are synthesized year-round, although the synthesis rate can be species-specific for any given MAA (Carreto & Carignan 2011). In natural assemblages of phytoplankton, several species synthesize several groups of MAAs, and the correlation between the concentration of MAAs and Chl $a$ is not necessarily significant because synthesis of MAAs is controlled by environmental conditions (Carreto & Carignan 2011). The quality and quantity of light and nutrients are critical environmental factors. The quality and quantity of UVR influences the synthesis of MAAs (Moisan & Mitchell 2001), and a low availability of nutrients can also depress the synthesis of MAAs (Korbée et al. 2005). Attenuation of UVR might be greater in warmer periods because of high phytoplankton biomass and higher concentrations of intercellular MAAs. Primary MAAs, such as mycosporine-glycine, might be synthesized during the warm period, and secondary MAAs might be synthesized in subsequent periods (Carreto & Carignan 2011). At shorter wavelengths of UVR, such as UV-B, increased attenuation of UVR occurs. In particular, the attenuation of UV-A (315–400 nm) by total MAAs might assist in recovery from photo-damage, as suggested by the high levels of accumulation of MAAs in natural assemblages of phytoplankton cells, and also by laboratory experiments (Carreto et al. 1990, Buma et al. 1995).

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