

Quasi-antiphase Diel Patterns of Abundance and Cell Size/Biomass of Picophytoplankton in the Oligotrophic Ocean

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keypoints

Picophytoplanktonic quasi-antiphase diel cycles in abundance and cell size/biomass are likely a general feature of the oligotrophic ocean

Grazing pressure on *Prochlorococcus* and *Synechococcus* is as high during the day as during the night

Abstract

Picophytoplankton are the smallest, most abundant photosynthetic organisms in the ocean. Knowledge of the diel variability of these tiny microbes has important implications for the structure of microbial food webs and key biogeochemical processes. However, insight into the underlying mechanisms of picophytoplanktonic diel dynamics is limited. By combining a field survey with a published dataset, we found that cell numbers and cell sizes/biomasses

of picophytoplankton were tightly synchronized to the day-night cycle, but they were in a quasi-antiphase relationship to each other. This pattern is a confirmation and extension of previous studies. Mortality rates showed that *Prochlorococcus* and *Synechococcus* were subject to considerable grazing pressure throughout the day and night. The quasi-antiphase diel cycles in abundance and cell size/biomass are likely determined by the light-dependent diel behavior of cell growth and division and continuous losses to grazing. This work significantly improves our understanding of autotrophic picoplankton in the oligotrophic ocean.

Plain Language Summary

Picophytoplankton are tiny, single-celled photosynthetic organisms that contribute to almost all primary production in the vast euphotic zones of the oligotrophic ocean. Understanding their roles in that environment is critical but challenging, mainly because of their minuscule size and the complexity of microbial processes and interactions. Time-series observations based on flow cytometry, a powerful technique that provides information about the numbers and sizes of picophytoplankton cells, have elucidated many ecological and biogeochemical processes associated with picophytoplankton, but some questions remain. A field survey in the northern South China Sea combined with a published dataset revealed that picophytoplankton cell size and biomass tended to decrease (increase) during the night (day) when cell numbers were increasing (decreasing). Such quasi-antiphase cycles are likely a general feature of near-steady-state oligotrophic ecosystems and reflect the cycles of carbon fixation, energy storage, and cell growth during the daytime and cell division and energy depletion during the night. Mortality rates estimated via modified dilution experiments showed that *Prochlorococcus* and *Synechococcus* were subject to considerable grazing pressure throughout the day and night. This work significantly improves our understanding of these microorganisms and may have implications for the carbon cycle in oligotrophic marine ecosystems.

Introduction

The oligotrophic ocean accounts for approximately half of Earth's surface and is the habitat of the greatest number of phototrophs in the world. In that nutrient-deficient environment, the food web is dominated by the microbial loop, and picophytoplankton (*Prochlorococcus*, *Synechococcus*, and picoeukaryotes) account for most of the photosynthetic biomass and primary production (Fenchel, 2008). Picophytoplankton cells are quite small ($< 2 \mu\text{m}$ in diameter), but they are numerically the dominant photosynthetic organisms in the ocean and play pivotal roles in shaping planktonic community structure and regulating the carbon cycle in oligotrophic marine ecosystems. With an expansion of oligotrophic regions caused by climate change, these smallest of marine phytoplankton will increase in both numerical abundance and biomass in the future ocean (Flombaum et al., 2013, 2020).

Food webs in the oligotrophic ocean are highly dynamic. Diel cycles of cell division, cell numbers, and cell size of picophytoplankton are significant and well documented in tropical and subtropical near-surface oceanic waters (Vaulot et al., 1995; Vaulot and Marie, 1999; Binder and DuRand, 2002). An increase of cell numbers at night is driven primarily by concurrent cell division, and the decline of cell numbers during the day reflects an imbalance between gains from cell division and losses to grazing and viral lysis. However, it seems paradoxical that the decline of cell abundance is synchronized with photosynthetic production. Recent research performed in the North Pacific Subtropical Gyre (NPSG) has shown that picophytoplankton biomass exhibits marked diel oscillations—a diurnal increase and nocturnal decrease—that are synchronized with the concentrations of particulate organic carbon inferred from optical measurements (Henderikx-Freitas et al., 2020; Boysen et al., 2021). Although the diel periodicity of photosynthesis and cell division undoubtedly contribute to this cycle, mortality associated with grazing and viral infection are equally important determinants of phytoplankton abundance (Binder and DuRand, 2002). The diel pattern of cell numbers and biomass of picophytoplankton combined with estimates of their growth rates as well as losses to grazing and viral lysis allow consideration of both bottom-up and top-down control.

In this study, we conducted time-series observations and parallel incubation experiments in the northern South China Sea (SCS), which is a typical oligotrophic marginal sea (Wong et al., 2007). We combined flow cytometric (FCM) analysis with an empirical laboratory calibration for cell size determination to assess the diel patterns of three picophytoplankton groups and found that there was a quasi-antiphase relationship between cell numbers and cell size/biomass of *Prochlorococcus* and *Synechococcus*. We complemented this study with a compiled dataset to ascertain the prevalence of this highly synchronized, quasi-antiphase diel cycles of cell numbers and cell size/biomass of picophytoplankton. Furthermore, the incubation experiment results and a simple model were helpful to understand the underlying mechanism of the quasi-antiphase diel patterns.

Materials and Methods

Sampling and Environmental Variables

The MARCO summer cruise (KK1904), a survey of the northern SCS, was conducted from 17 June to 04 July 2019 on board the R/V Tan Kah Kee (Figure S1). Three time-series stations, two on the slope (M4, K11) and one in the basin (SEATS), were occupied during the cruise. We conducted time-series surveys at K11, while Lagrangian observations following drifting sediment traps were carried out at SEATS and M4. During 24-h or 48-h sampling periods, surface water samples (~5 m) in triplicate were collected for FCM analysis every ~1.5 h using a CTD rosette sampler or a plexiglass water sampler. On each observation day, seawater samples were collected twice, at approximately 06:00 and 18:00 (local time), for modified dilution experiments.

Flow Cytometric (FCM) Analysis and Population-specific Carbon Biomass Calculation

Cell numbers and light scatter of three picophytoplankton populations (*Prochlorococcus*, *Synechococcus*, and picoeukaryotes) were determined using a BC CytoFLEX flow cytometer following procedures described previously (Marie et al., 1999). In accord with the commonly used flow cytometer-specific calibrations (DuRand et al., 2001; Jacquet et al., 2001; Worden

et al., 2004), forward light scatter (FSC) was converted to cell size (expressed as equivalent spherical diameter, ESD) using an empirical relationship ($\log ESD = 0.2504 \log FSC + 0.1351$, Figure S2) between FSC measured by this instrument and cell size determined by epifluorescence microscopy (Zeiss Imager.A2) for different exponentially growing phytoplankton cultures (eight marine picophytoplankton cultures at three different times of day). Additional details regarding FCM analysis and the empirical size-FSC calibration are described in the supporting information. The resulting cell size was used to calculate biovolume assuming spherical shape. Cell biovolume was then converted to carbon with a conversion factor of $280 \text{ fg C } \mu\text{m}^{-3}$, which was derived from an equatorial Pacific *Prochlorococcus* strain (Heldal et al., 2003). Carbon biomass of each group was estimated by multiplying per-cell carbon by cell numbers.

Growth and Loss Rate Estimates

To identify the biological factors responsible for the diel variations of picophytoplankton, modified dilution experiments were performed following the protocol of Kimmance and Brussaard (2010) in parallel to the time series. At about sunrise or sunset, natural surface seawater, gently passed through a 20- μm nylon net filter to remove microzooplankton, was combined with grazer-free filtrate ($< 0.1 \mu\text{m}$) or virus-and-grazer-free filtrate ($< 30 \text{ kDa}$) in proportions of 27%, 55%, 82%, and 100%. All mixtures in quadruplicate were incubated for 9 h in an on-deck Plexiglas incubator, which was screened with neutral density filters (LEE 298) to simulate the photosynthetically active radiation (PAR) intensity at $\sim 5 \text{ m}$. Incubation temperature was controlled by continuously flowing surface seawater. FCM samples were taken at the beginning and end of the incubation as described above. Mortality and intrinsic growth rates were calculated from linear regressions of apparent growth rate versus dilution factor (see more details in Text S2). The low abundance of picoeukaryotes made the rate estimates unreliable, and the calculated grazing rates were generally negative. In the following analysis, we therefore concentrated on *Prochlorococcus* and *Synechococcus*.

For a qualitative comparison, we calculated diurnal/nocturnal net growth rates from time-series observations and incubation experiments. The former ones were estimated from the changes between the two time points closest to sunrise and sunset, i.e., $\ln(P_2/P_1)/(t_2 - t_1)$, where P is the corresponding abundance or biomass and t represents time. The latter ones were determined from the difference between the intrinsic growth rates and the loss rates due to grazing and viral lysis. Sometimes the loss rates were negative, especially those associated with virus-induced mortality rates. The subtraction calculations were performed in two different ways: one was to include the negative loss rates in the calculations, and a second was to correct the negative loss rates to 0. All the calculations were based on both abundance and biomass.

Joint Analysis With SeaFlow Dataset

To demonstrate the generalization ability of the local time-series study, we extended a published dataset (SeaFlow data v1.3, Ribalet et al., 2020) that consisted of high-resolution, underway FCM observations in surface waters of the North Pacific and South Atlantic. The dataset included cell sizes and total carbon biomasses for picophytoplankton populations estimated using methodologies similar to ours. After data cleaning, data from 38 cruises (including our SCS cruise) in oligotrophic conditions remained for the subsequent analysis (Figure S3, Table S2). Details of data cleaning and other data processing procedure are presented in the supporting information.

In order to make a direct comparison of the data from different geographical regions and dates with differing daylight hours, all sampling times were adjusted to a “standard day” with sunrise at 06:00 and sunset at 18:00. Moreover, we standardized the measured values by dividing by the mesor (midline-estimating statistic of rhythm, a rhythm-adjusted mean calculated by the cosinor method) value in a 24-h rolling window. Because the time intervals between the data points in the datasets were not all the same and because many data gaps existed, the mesor could provide a better estimate of the mean level (the average value around which the variable oscillated) than the arithmetic mean (Refinetti, 2016). Normalization

resulted in a mean level of 1. In addition, the normalization led to the elimination of long-range trends, which could influence the assessment of 24-h periodicity (Leise, 2017). After identification and replacement of outliers and rolling smoothing, the final data were binned to half-hour intervals. To further explore diel variations of picophytoplankton, the diel periodicity analyses were conducted using the cosinor method by fitting a cosine curve with a 24-h period (Refinetti, 2016). A day with no more than 20 (41.7% of 48 half-hour intervals) missing values was considered a valid day. Altogether, 200 valid days from 250 days were included in the analysis. Because the reliability of the periodicity analyses and the accuracy of phase estimation based on a single-cycle's data were low (Leise, 2017), especially when there were many missing values, we conducted cosinor analyses using a rolling window of 3 days. Diel periodicity was statistically validated with $R^2 \geq 0.36$, and the corresponding clocktimes of acrophase were analyzed using circular boxplots (Buttarazzi et al., 2018).

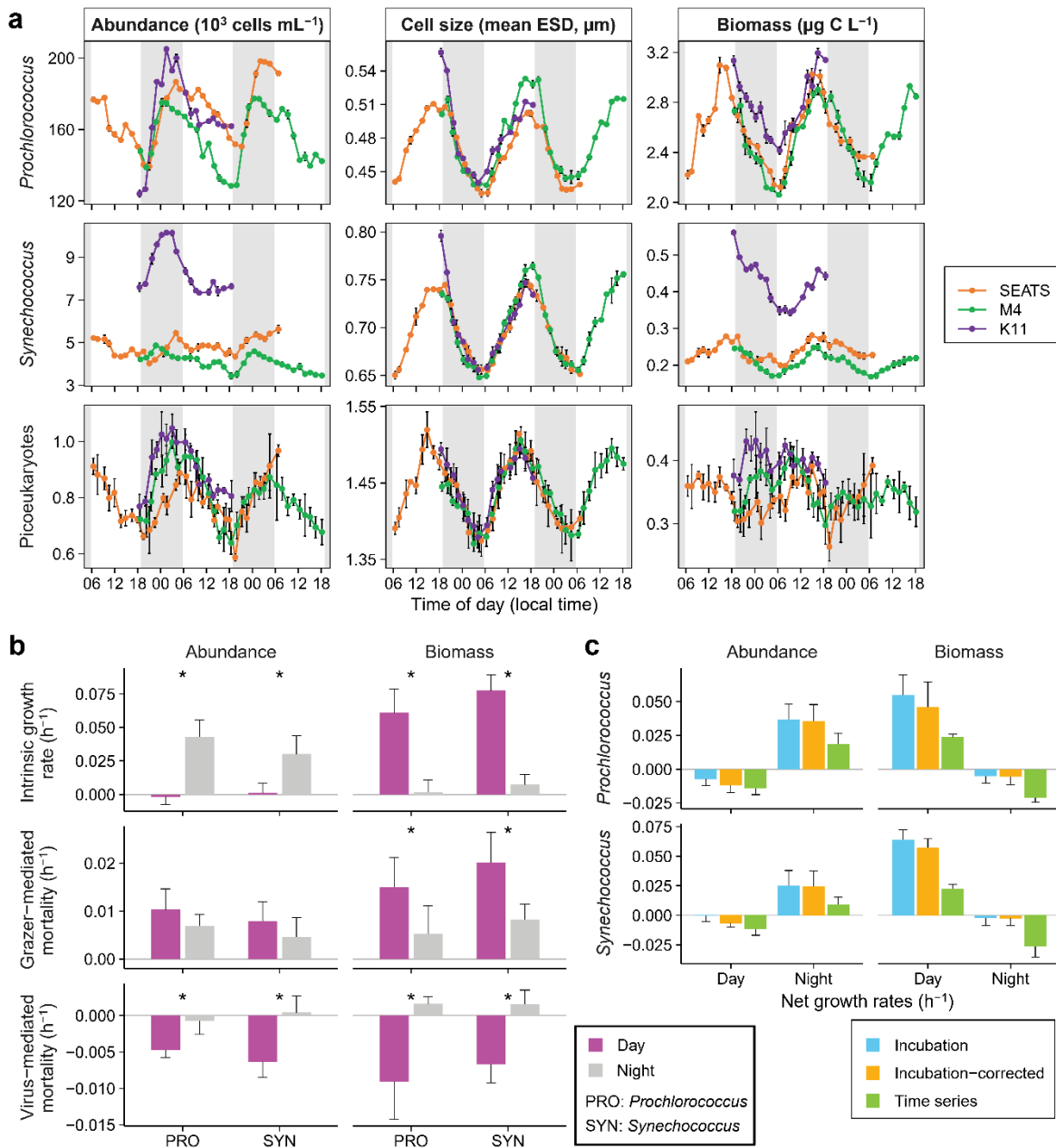


Figure 1. The field study in the northern SCS. (a) Time series of picophytoplankton (*Prochlorococcus*, *Synechococcus*, and picoeukaryotes) cell numbers, cell size, and biomass in surface waters at three stations during the MARCO cruise (June–July 2019). Night-time periods are shaded grey. (b) Incubation-based day versus night comparisons of picophytoplankton intrinsic growth, virus- and nanoflagellate-mediated mortality based on

cell numbers and biomass. Significant differences between day and night are indicated by asterisks ($p < 0.05$, Student's t -test). (c) Day versus night comparisons of picophytoplankton abundance- and biomass-based net growth rates estimated from incubation experiments and time series of in situ changes. Negative incubation-based loss rates are included or corrected to 0. Error bars denote standard deviations ($n = 3$ in (a); $n = 5$ in (b, c)) and are smaller than the data points when not apparent.

Results

Diel Variations of the Picophytoplankton Community in the Northern SCS

Survey results at three stations along the continental slope (K11 and M4) and in the basin (SEATS) indicated that environmental conditions in the northern SCS during the summer were relatively stable. Fluctuations of temperature and salinity were small, and PAR was more-or-less constant (Figure S1, Table S1). The surface Chl a concentrations at the three stations were quite low ($0.11\text{--}0.13 \mu\text{g L}^{-1}$), and biological variables at these three stations were strikingly similar (Table S1). Picophytoplankton community composition was dominated by *Prochlorococcus* (average abundance reached $156\text{--}170 \times 10^3 \text{ cells mL}^{-1}$); the cell numbers of *Synechococcus* and picoeukaryotes were 2–3 orders of magnitude lower, respectively. These characteristics indicated that the three stations could be considered typical of oligotrophic environments.

During 24-h or 48-h time series, the picophytoplankton in surface waters exhibited clear diel periodicity in cell numbers, cell size, and biomass (Figure 1a). Compared to *Prochlorococcus*, the oscillations were noisier for the cell numbers and biomasses of *Synechococcus* and picoeukaryotes, the abundances of which were relatively low. Generally, the cell numbers of the three picophytoplankton groups increased at night and decreased during the day, whereas cell sizes increased during the day and decreased at night. Analysis by the cosinor method revealed that the average daily percent increases (from trough to peak) in cell numbers of *Prochlorococcus*, *Synechococcus*, and picoeukaryotes were 27.7%, 25.9%, and 30.9%, respectively. The corresponding increases of cell sizes were 18.0%,

14.8%, and 7.5%, respectively. The biomasses of *Prochlorococcus* and *Synechococcus* displayed a pattern similar to those of cell sizes but almost antiphase with those of cell numbers; the biomasses and cell sizes increased during the day and decreased at night. The percent increases of the biomasses of *Prochlorococcus* and *Synechococcus* over the diel cycle were 29.2% and 34.6%, respectively. However, no diel periodicity was detected for the biomass of picoeukaryotes, and there were no diel patterns in the biomasses of heterotrophic bacteria and viruses (Figure S4).

Daytime Versus Night-time Rate Estimates From FCM Abundances or Biomasses

For *Prochlorococcus* and *Synechococcus*, diel patterns of intrinsic growth rates based on cell numbers and biomass were completely opposite (Figure 1b, Table S3). The estimated intrinsic growth rate based on the rate of change of cell numbers was significantly lower during the day than at night, whereas the biomass-based intrinsic growth rate was obviously higher during the day than at night. Furthermore, net growth rates based on cell numbers were negative during the photoperiod and positive at night, whereas biomass-based estimates were just the opposite, no matter whether the calculation was based on incubation experiments or time-series data (Figure 1c). In contrast, nanoflagellate grazing rates derived from FCM abundances were not significantly different between the day and night, whereas biomass-based grazing rates were significantly higher during the daytime than at night (Figure 1b). We cannot speculate about the reason for these differences, and the later discussion will be based on the commonly used abundance parameters. Both abundance- and biomass-based virus-induced mortality rates were lower than nanoflagellate grazing rates, and most of them were even negative (Figure 1b). Yet it is unclear whether the negative viral mortality is biologically meaningful (i.e., viral effects are stimulatory) or just a methodological artifact (Pasulka et al., 2015).

The Prevalence of the Diel Patterns of Picophytoplankton Community

As expected, the results of the joint analysis were consistent with the local time series. When all the normalized data were simply aggregated, all three picophytoplankton populations

showed pronounced diel patterns, especially in terms of cell size and biomass (Figure 2a). Abundance data displayed great variability, and their diel patterns were less dramatic, even slightly different from those of the local study, that is, the abundances of *Prochlorococcus* and picoeukaryotes rose in the afternoon. This pattern suggested that the diel variation in abundance was less significant, or that the rhythmic pattern was less consistent.

Further analysis at the daily level confirmed this suggestion (Figure 2b, Figures S8–S10). Cosinor analysis showed that diel patterns of cell numbers of *Prochlorococcus*, *Synechococcus*, and picoeukaryotes were detectable in 66.0% ($n = 132$), 39.5% ($n = 79$), and 53.5% ($n = 107$), respectively, of the 200 valid days. For all three populations, cell numbers exhibited a diel peak mainly between midnight and sunrise, in which *Prochlorococcus* (01:59 [00:32–03:30], median and interquartile range) peaked earliest, followed by picoeukaryotes (03:52 [03:10–04:41]) and then *Synechococcus* (05:35 [04:18–06:42]). In the case of cell size, all three picophytoplankton populations almost always exhibited a clear diel variability (200 [100%], 190 [95.0%], and 188 [94.0%] for *Prochlorococcus*, *Synechococcus*, and picoeukaryotes, respectively). Diel oscillations of cell size were highly synchronized and phased almost identically for all three populations (*Prochlorococcus*, 17:11 [16:45–17:32]; *Synechococcus*, 17:24 [16:55–17:56]; and picoeukaryotes, 17:10 [16:47–17:32]) during nearly all valid days. Of all 200 valid days, 195 (97.5%), 97 (48.5%), and 169 (84.5%) were characterized by diel patterns of the biomasses of *Prochlorococcus*, *Synechococcus*, and picoeukaryotes, respectively. The corresponding acrophases occurred around sunset (*Prochlorococcus*, 17:35 [17:08–18:04]; *Synechococcus*, 16:51 [15:49–18:08]; and picoeukaryotes, 17:27 [16:59–17:56]) at times similar to the peaks of cell size.

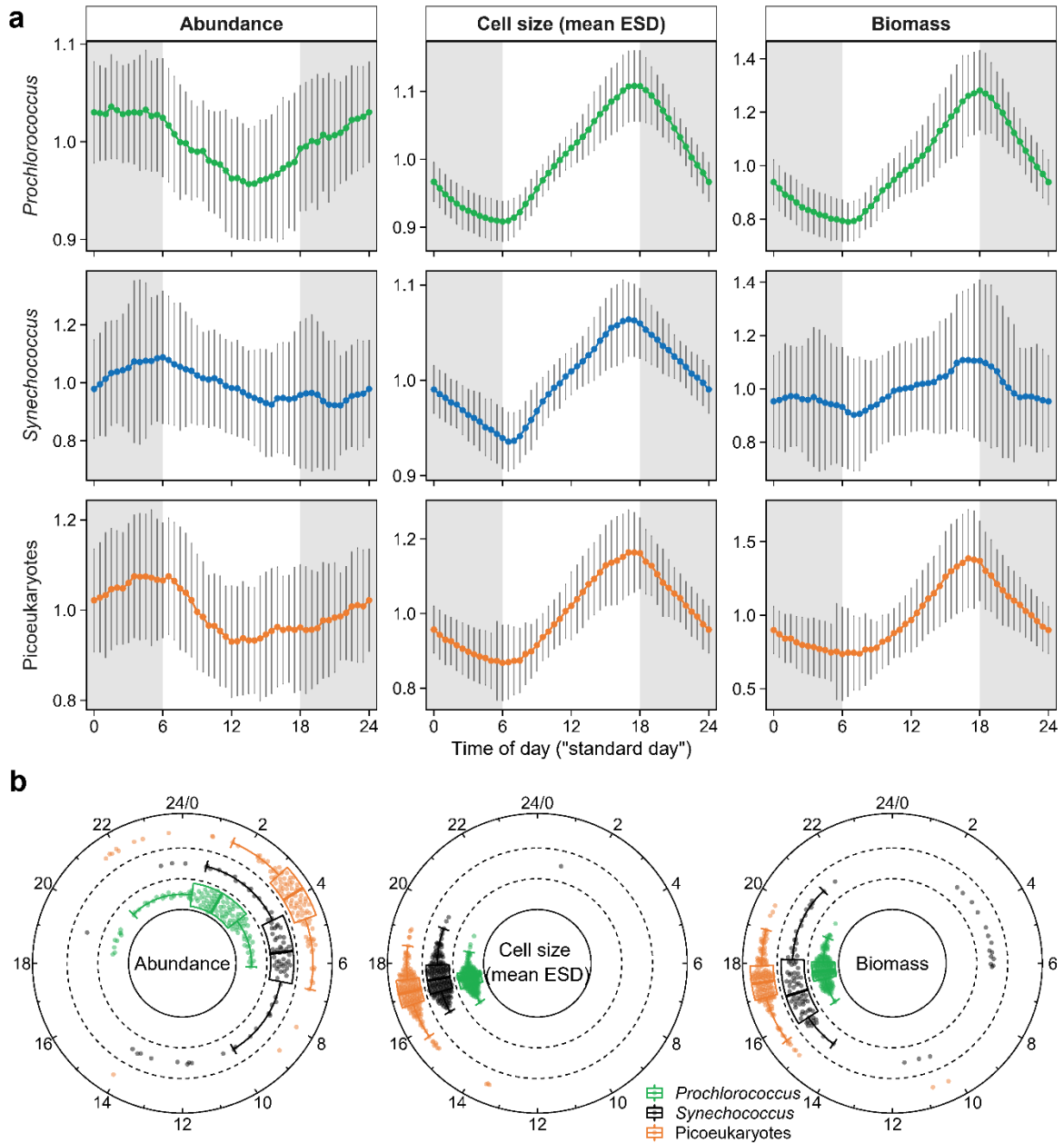


Figure 2. Joint analysis of the SCS observations and SeaFlow dataset. (a) Average half-hourly values of cell numbers, cell sizes, and biomasses of three picophytoplankton groups across all 38 cruises. Note that because the values have been normalized, so they are unitless and fluctuate about 1. Error bars denote standard deviations (n ranges from 181 to 218). (b) Circular boxplots overlaid with beeswarm plots showing the distribution of the acrophases

(peak times) of the diel periodicity in cell numbers, cell sizes, and biomasses of three picophytoplankton groups. Each dot represents a valid day with a statistically significant 24-h periodicity. From left to right, $n = 132, 200, 195$ for *Prochlorococcus*; $n = 79, 190, 97$ for *Synechococcus*; and $n = 107, 188, 169$ for picoeukaryotes. The boxes contain the central 50% of the data surrounding the circular median; whiskers correspond to 1.5 times the interquartile range.

Discussion

For obvious reasons, the physiological processes of phytoplankton are closely correlated with the light-dark cycle. In this study, we found marked differences in picophytoplanktonic diel cycles between cell numbers and cell size/biomass in the northern SCS (Figure 1a). The differences were consistent with the net growth rates estimated during the day/night with our modified dilution experiments (Figure 1c). The cycles of abundance and cell size in our study were similar to those observed in the equatorial Pacific, although phased slightly differently (Vaulot and Marie, 1999). In the case of *Prochlorococcus* and *Synechococcus*, the cycles of biomass were consistent with the observations in studies from the NPSG, in which biomass has exhibited marked diel oscillations and coherent synchronization with bulk optical properties (Henderikx-Freitas et al., 2020; Boysen et al., 2021). We did not find a similar diel pattern in picoeukaryotic biomass, possibly because of their low abundance and the fact that the upper threshold (2 μm) that we used may have excluded some cells that became larger with growth (Figure S11). Direct measure of group-specific picophytoplankton biomass under field conditions has not been possible, although such measurements are simple in laboratory cultures. Field studies normally make use of some indirect method, such as photosynthetic pigments (Mackey et al., 1996) and FCM-derived estimates (Ribalet et al., 2019; Boysen et al., 2021). Estimates of carbon per cell based on FCM analyses are confounded by uncertainties associated with the conversions of forward scatter to size, size to biovolume (assuming spherical shape), and biovolume to carbon. However, we found that the daily percent increases in biovolume based on the FCM-derived sizes of picophytoplankton were similar in the field and laboratory (Figure S11). Furthermore, when

four different empirical functions (Worden et al., 2004; Ribalet et al., 2019) were used to convert cell biovolume to carbon, the cellular quotas of carbon differed, but the broad trends of biomass were unchanged (Figure S12). We therefore felt that the patterns in the diel oscillations of picophytoplanktonic biomass were robust.

The findings of the local investigation were further confirmed and generalized in the joint analysis (Figure 2). Of the three picophytoplankton groups, cycles with a period of 24 h were more evident for *Prochlorococcus* than for *Synechococcus* and picoeukaryotes, which were present in relatively low abundance. The evidence of a diel pattern was strongest for cell size, weakest for cell numbers, and intermediate for biomass, which depended on the product of cell numbers and cell size. It should be noted that only a few underway observations were conducted in the same water mass, and the obtained percentages of significant diel variation were definitely lower than the percentages obtained using Lagrangian observations (similar to stations SEATS and M4). In addition, the acrophase distribution was more concentrated for cell size and biomass than for abundance. These differences reflected the fact that cell size was regulated by a combination of the cell division cycle and the 24-h photosynthesis/respiration cycle (Binder and DuRand, 2002), whereas cell numbers were regulated by a complex combination of biotic and abiotic factors. Moreover, the daily cycles of biomass were clearly driven largely by changes in cell size and less by the dynamics of abundance. Cell size is controlled mainly by the light-dark cycle, whereas grazing and physical processes have little effect on cell size, but they do affect cell numbers. Mixing of different water masses due to horizontal advection or vertical mixing could add or remove cells and dramatically perturb the diel patterns of cell numbers (André et al., 1999). Tsai et al. (2009) have reported that strong winds and heavy rains during the passage of a typhoon seriously disturbed the diel pattern of *Synechococcus* abundance but had little effect on the cell division cycle.

Furthermore, the joint analysis revealed that the almost antiphase cycles between abundance and cell size/biomass were evident in most cases (Figure 2b). In fact, the underlying mechanism of such quasi-antiphase relations is easy to understand. In the absence of any

physical disturbance, picophytoplankton diel dynamics are driven primarily by three processes: cell division, cell growth, and loss processes. For a photosynthetic organism, biomass in terms of carbon increases only during the day, decreases at night, and peaks at the end of the photoperiod. Although the timing varies among these three groups and between different times and locations, cell division generally occurs at night and/or in the late afternoon, whether in culture or in the field (Jacquet et al., 2001; Binder and DuRand, 2002). In addition, cell division during steady state growth occurs when the biomass per cell has doubled, and cellular biomass can increase only during the photoperiod. Carbon fixed by photosynthesis is therefore used mainly for cell growth during the daytime. Cell size tends to decrease at night because of cell division, respiration, and exudation. Thus, like biomass, cell size exhibits significant diel periodicity and peaks near sunset. In contrast, diel variations of abundance are determined by cell division and loss processes (e.g., grazing, viral lysis). The fact that these two processes can occur simultaneously and affect abundance in opposite directions confounds deconvolution of diel abundance cycles (Binder and DuRand, 2002) and explains, to some extent, the relatively heterogeneous behavior of those cycles (Figure 2b). However, decreases in cell number can result only from loss processes, whereas increases can result only from cell division. The cell numbers increase when the division rate exceeds the loss rate and peak when the former declines and/or the latter increases until they are equal. Figure 2b shows that peak times of abundance occurred mainly between midnight and sunrise. This pattern was consistent with the results of most previous studies (Binder and DuRand, 2002, and references therein), with the exception of Ribalet et al. (2015). The acrophases of *Prochlorococcus* abundance during the two cruises (CN111D and TN271) studied by Ribalet et al. (2015) are the outliers in Figure 2b. During those cruises, cell numbers and cell sizes of *Prochlorococcus* both increased during the day and decreased at night (Figures S8 and S9). This pattern differed from the pattern we observed in most cases and may have been exceptions to the more common diel cycle of *Prochlorococcus* cell numbers.

The idea that grazing generally accounts for the majority of phytoplankton mortality in the ocean (Calbet and Landry, 2004) is consistent with the observations in Figure 1b. However,

the diel pattern of protistan grazing on picophytoplankton has not been well defined, and some contradictory findings have been reported (Connell et al., 2020; Fowler et al., 2020). Some reports (Tsai et al., 2005; Ribalet et al., 2015; Connell et al., 2020) have indicated that protistan grazing occurs primarily at night. The discovery in our study of comparable grazing loss during the daytime as at night (Figure 1b) was similar to the results of some other studies (Dolan and Šimek, 1999; Ng and Liu, 2016; Fowler et al., 2020). This grazing coincided with a diurnal decrease in cell numbers (Figure 1a). As mentioned above, the *Prochlorococcus* abundance cycles observed by Ribalet et al. (2015) may have been exceptions to the general pattern, and if so, the diel patterns of *Prochlorococcus* mortality would have been exceptions as well. Tsai et al. (2005) have calculated diurnal/nocturnal growth rates of *Synechococcus* based on cell abundance and then multiplied those growth rates by in situ biomasses (derived from a constant carbon conversion factor) to obtain biomass-based production rates and grazing rates. We feel that their calculations were biased by the fact that they did not take into account diel patterns of cell size and biomass. Furthermore, the pattern of grazing can also be inferred from laboratory incubations, in which clear diel variations of picophytoplankton abundance have been observed when cells were grown under light-dark cycles. The growth curves in conventional batch cultures follow a step-like pattern with an ascending period during the night and a stable period during the day (e.g., Jacquet et al., 2001; Zinser et al., 2009; Waldbauer et al., 2012), whereas temporal periodicity of abundance similar to those of natural populations can be observed only under continuous culture conditions (Mori et al., 1996; Claustre et al., 2002). The key features that distinguish continuous culture systems from batch cultures are continuous removal of culture and continuous addition of fresh medium in the former. The implication is that loss processes under realistic field conditions may be continuous and may even occur at a more-or-less constant rate.

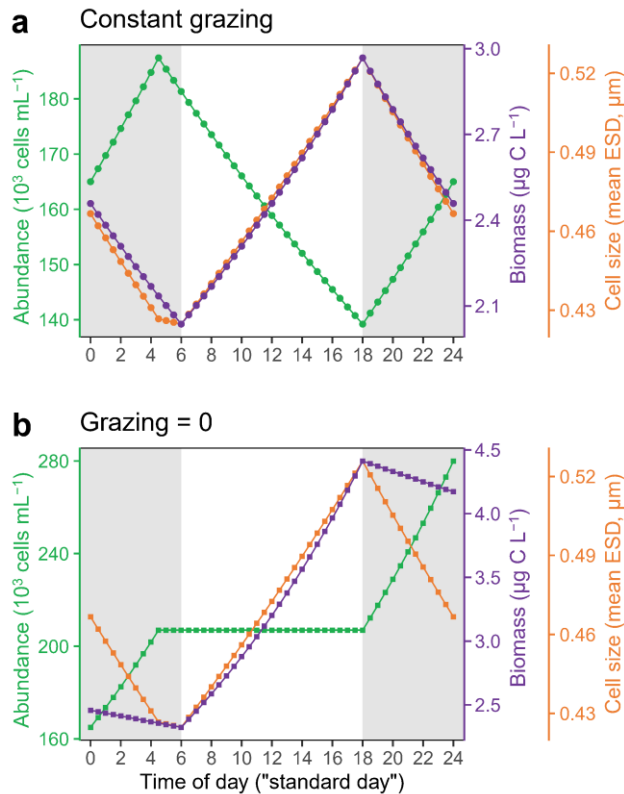


Figure 3. A simple model simulating the daily patterns of cell numbers, cell size, and biomass of *Prochlorococcus* when the grazing rate is constant (a) or equal to 0 (b). All vital rates are assumed to be constant, with net primary production during the day, respiration throughout the night, grazing throughout 24 hours, and cell division beginning at sunset and continuing through the first 10.5 hours of darkness (the average peak time was ~04:30 for *Prochlorococcus* in the SCS study). The model was optimized using the SCS *Prochlorococcus* data (Figure 1a) and adjusted so that the numbers were almost identical after 24 hours and at the beginning, i.e., the system was in steady state.

Based on the aforementioned results and discussion, we developed a simple mathematical model to simulate the behavior of cell numbers, cell size, and biomass of picophytoplankton over a period of 24 hours (Figure 3a, see more details in Text S4). In this simulation, there was clearly an almost antiphase relationship between cell numbers and cell size/biomass. If the grazing rate is adjusted to 0 (with other parameters unchanged), the diel patterns would

be very similar to that of laboratory batch cultures (Figure 3b). Although this model is very simplistic, it can explain the diel patterns of picophytoplankton in the field (Figure 1a) and in culture and supports the rationale of the above explanation.

Conclusions

By combining a field survey with a published dataset, this study confirmed and extended previous results of diel patterns of autotrophic picoplankton in the oligotrophic ocean. We suggest that the quasi-antiphase diel cycles in abundance and cell size/biomass of picophytoplankton are likely a general feature of near-steady-state oligotrophic ecosystems. Light is recognized as a key driver in the dynamics of microbial food webs. The quasi-antiphase diel patterns of cell numbers and cell size/biomass of these tiny phototrophs reflect the cycles of carbon fixation, energy storage, and cell growth during the daytime and cell division and energy depletion during the night. The loss processes through grazing seem to occur throughout the day and night, but many details remain to be discovered.

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