



Primary Productivity Dynamics in the Summer Arctic Ocean Confirms Broad Regulation of the Electron Requirement for Carbon Fixation by Light-Phytoplankton Community Interaction

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Specialty section:

This article was submitted to
Marine Biogeochemistry,
a section of the journal
Frontiers in Marine Science

Received: 30 November 2018

Accepted: 07 May 2019

Published: 29 May 2019

Citation:

Zhu Y, Suggett DJ, Liu C, He J, Lin L,
Le F, Ishizaka J, Goes J and Hao Q
(2019) Primary Productivity Dynamics
in the Summer Arctic Ocean Confirms
Broad Regulation of the Electron
Requirement for Carbon Fixation by
Light-Phytoplankton Community
Interaction. *Front. Mar. Sci.* 6:275.
doi: 10.3389/fmars.2019.00275

Predicting conversion of photosynthetic electron transport to inorganic carbon uptake rates (the so-called electron requirement for carbon fixation, K_C) is central to the broad scale deployment of Fast Repetition Rate fluorometry (FRRf) for primary productivity studies. However, reconciling variability of K_C over space and time to produce robust algorithms remains challenging, given the large number of factors that influence K_C . We have previously shown that light appears to be a proximal driver of K_C in several ocean regions and we therefore examined whether and how light similarly regulated K_C variability in the Arctic Ocean, during a summer cruise in 2016. Sampling transited ice-free and ice-covered waters, with temperature, salinity and Chl-a concentrations all higher for the ice-free than ice covered surface waters. Micro- and pico-phytoplankton generally dominated the ice-free and ice-covered waters, respectively. Values of K_C , determined from parallel measures of daily integrated electron transport rates and ^{14}C -uptake, were overall lower for the ice-covered vs. ice-free stations. As in our previous studies, K_C was strongly linearly correlated to daily PAR ($r = 0.68$, $n = 46$, $p < 0.001$) and this relationship could be further improved ($r = 0.84$, $n = 46$, $p < 0.001$) by separating samples into ice-free (micro-phytoplankton dominated) vs. ice-covered (Nano- and Pico-phytoplankton dominated water). We subsequently contrasted the PAR- K_C relationship from the Arctic waters with the previous relationships from the Ariake Bay and East China Sea and revealed that these various PAR- K_C relationships can be systematically explained across regions by phytoplankton community size structures. Specifically, the value of the linear slope describing PAR- K_C decreases as water bodies have an increasing fraction of larger phytoplankton. We propose that this synoptic trend reflects how phytoplankton

community structure integrates past and immediate environmental histories and hence may be a better broad-scale predictor of K_C than specific environmental factors such as temperature and nutrients. We provide a novel algorithm that may enable broad-scale retrieval of CO_2 uptake from FRRf with knowledge of light and phytoplankton community size information.

Keywords: primary productivity, Arctic Ocean, ice free/cover, electron requirement for carbon fixation, phytoplankton community, FRRf

INTRODUCTION

Accurate estimation of ocean primary productivity (PP) is central to understanding marine carbon geochemical cycling and the transfer of energy through food webs (Hancke et al., 2015). Compared to traditional incubation-based PP methods (Regaudie-de-Gioux et al., 2014), *in situ* bio-optical active chlorophylla fluorescence methods, notably Fast Repetition Rate fluorometry (FRRf, Kolber et al., 1998), afford unprecedented high spatial and temporal resolution needed to understand how the environment continually fine-tunes primary productivity (e.g., Behrenfeld et al., 2006; Suggett et al., 2009b). However, chlorophylla fluorescence measurements inherently quantify PP through the activity of photosystem II (PSII) light reactions, resulting in a “photosynthetic currency” of PSII photosynthetic electron transfer rate (ETR_{PSII} , $\text{mol e}^- \text{mol RCII s}^{-1}$, Suggett et al., 2009a; Hughes et al., 2018b). Additional knowledge of the electron requirement for C fixation (termed K_C , $\text{mol e}^- [\text{mol C}]^{-1}$, also termed $\Phi_{e,C}$, Lawrenz et al., 2013; Hughes et al., 2018a) is required to convert ETR_{PSII} to more conventional currencies of CO_2 uptake rates (Kromkamp et al., 2008). Studies have previously demonstrated that K_C appears highly variable in ocean and coastal waters, since numerous factors can cause cellular processes to consume ETR_{PSII} -derived energy and reductant that is otherwise used for CO_2 assimilation (e.g., Lawrenz et al., 2013; Halsey and Jones, 2015; Hughes et al., 2018b). Developing approaches to model K_C , and hence account for K_C variability, are therefore critical for efforts utilizing FRRf-based measurements for highly resolute CO_2 uptake rate retrieval.

Early efforts to reconcile conventional CO_2 uptake rates with FRRf-derived CO_2 uptake attributed FRRf “overestimates” to use of a constant K_C value (i.e., 4 or 5 $\text{mol e}^- (\text{mol C})^{-1}$) (Kromkamp et al., 2008; Mino et al., 2014), in particular under excess irradiance (Ralph et al., 2010). Subsequent studies demonstrated that miss-matches between CO_2 uptake rates and FRRf were largely due to K_C variability, which in turn could be explained (predicted) from co-variability with key environmental factors known to regulate PP, i.e., light, temperature and/or inorganic nutrient availability (Lawrenz et al., 2013; Hughes et al., 2018b; Ryan-Keogh et al., 2018). Schuback et al. (2015, 2016, 2017) and (Schuback and Tortell, 2019) further demonstrated that such K_C variability correlated with the extent of non-photochemical quenching (NPQ), interpreted as an indication of excess light energy, which leads to a decoupling of ETR_{PSII} and carbon uptake (see also Hughes et al., 2018b). We similarly recently reported that variance of K_C , derived from parallel

measures of FRRf-derived ETRs and C-uptake rates, could be generally explained by patterns in light availability (Zhu et al., 2016), but importantly was much improved when considering differences in phytoplankton size structures (Zhu et al., 2017). In fact, changes in predominant species within phytoplankton communities appears to be a factor increasingly important in explaining patterns of K_C (e.g., Kulk et al., 2018; Xie et al., 2018), likely reflecting selection of taxa by environmental factors that are not specifically measured as part of the assessment exercise of interest (see Hughes et al., 2018a). Patterns of K_C variability over space and time therefore remain problematic to fully resolve. For example, the strong linear relationship which we observed previously between light and K_C differed across disparate ocean ecosystem, Ariake Bay (Zhu et al., 2016), and East China Sea (ECS)/ Tsushima Strait (Zhu et al., 2017). Schuback et al. (2016) similarly reported differences in comparing K_C and NPQ derived from two locations in the subarctic Pacific where the regression slopes differed for two datasets, presumably as a result of alternate specific nutrient regimes and/or phytoplankton species. To therefore further examine, and reconcile cross-regional differences in the regulation of K_C variation by light and phytoplankton community structures, we applied our established daily integrated CO_2 -uptake and FRRf inter-comparison method (Zhu et al., 2016, 2017) to the Arctic Ocean, where the light environment and phytoplankton taxa are very different from those assessed previously.

The Arctic Ocean is recognized as a sentinel marine system that is currently proportionately affected by climate change compared to many other systems worldwide (Wassmann et al., 2011). Near-surface warming in the Arctic has been almost twice as large as the global average (Graversen et al., 2008), and consequently the extent of Arctic sea ice has decreased dramatically (Serreze and Stroeve, 2015) increasing sea ice meltwater to result in surface freshening (McPhee et al., 1998; Yamamoto-Kawai et al., 2009; Timmermans et al., 2011). Arctic ecosystems appear particularly sensitive to spatial and temporal variations in hydrographic properties. Ice cover loss leads to increased light availability and the phytoplankton growing season, thus primary productivity in the Arctic Ocean (Arrigo et al., 2008; Hill et al., 2017). However, ice melt also intensifies freshening and in turn stronger stratification that may act to suppress upward fluxes of nutrients leading to decreased primary productivity and an increase in the fraction of pico-phytoplankton, typical for low-nutrient environments (Li et al., 2009; Coupel et al., 2015; Mills et al., 2018) through selection for low nutrient pico-phytoplankton (see Li et al., 2009). The

non-uniform trends associated with changes in sea-ice and primary productivity (Brown and Arrigo, 2012; Fernández-Méndez et al., 2015) fundamentally require more data across a wide variety of assemblages that can more broadly resolve this issue. Notably, understanding how primary productivity is regulated in Arctic systems can be achieved by increasing the spatial and temporal coverage of *in situ* measurements via active chlorophyll fluorescence techniques (Hancke et al., 2015; Schuback et al., 2017).

In this study, we conducted parallel measures of ETR_{PSII} and ^{14}C carbon uptake rates for Arctic Ocean surface waters during a 2017 summer cruise. Variability of phytoplankton photophysiology and light-dependency of K_C was first examined to identify environmental drivers (e.g., light, nutrients, ice coverage, etc.) and/or phytoplankton community structure differences specific to this region. We then compared these observations with those from our previously reported Ariake Bay and East China Sea datasets to show for the first time that changes in dominated phytoplankton size structure may explain the variation amongst cross-regional light- K_C relationships. We discuss how this new insight can further help in applying FRRf based approaches for high resolution primary productivity retrieval.

MATERIALS AND METHODS

Sampling and Measurements of Physical and Biochemical Properties

Sampling was conducted on board the *R/V Chinare* (XueLong Arctic cruise), July 18th to September 10 th, 2016. A total of 50 stations were visited spanning three main regions, the Bering strait (12 stations), Chukchi shelf (15 stations) and Chukchi borderland (23 stations) (Figure 1, Supplementary Table 1). Water was sampled from depths of 5, 20, 30, 50, 75, 100, 125, 150, 175, and 200 m using a rosette equipped with 12 × 5L Niskin bottles (General Oceanics, USA) and a conductivity-temperature-depth profiler (CTD, 911+, SeaBird Electronics, USA). For safety reasons, the icebreaker cannot deploy CTD at locations where ice was heavily covered, but only in open water in the ice covered region (Figure 1). Data from the CTD was utilized for establishing hydrographic conditions of each station at the time of sampling. Mixed layer depth (MLD) was calculated using the density difference criterion ($\sigma = 0.1 \text{ kg m}^{-3}$) of Peralta-Ferriz and Woodgate (2015). Incident photosynthetically active radiation (PAR) at the sea surface ($PAR(0^+)$) was measured throughout the cruise period with a cosine irradiance sensor (AV-19Q, Avalon, USA) mounted on the ship (see Zhu et al., 2016).

In this study, for phytoplankton size structure measurement, we separated each sample to Micro-, Nano- and Pico in these three groups based on size-fractionated chlorophyll a (Chl-a) measurements, according to standardized methods used in marine phytoplankton size research (Cormeño et al., 2006). Size-fractionated Chl-a concentration was determined on a 250 ml volume sample, which was filtered sequentially through 20 μm nylon membrane, 2 μm polycarbonate, and 0.7 μm pore size GF/F filters. Filtering through 20 and 2- μm was done under

gravity and 0.7- μm under low vacuum pressure ($<0.02 \text{ MPa}$), and cells retained by the 20 μm filters belong to the micro-phytoplankton whereas those retained by the 2 and 0.7 μm pore size constitute, respectively, nano-phytoplankton and pico-phytoplankton. Filters were extracted in 8 mL 90% acetone for 24 h in darkness under -20°C and the Chl-a concentration determined fluorometrically using a pre-calibrated fluorometer (Turner Trilogy, USA) following Holm-Hansen et al. (1965). A second aliquot of 5 mL for nitrate + nitrite (NO_x^-), phosphate (PO_4^{3-}), and silicate (DSi) was measured using a continuous flow-analyzer (Skalar SA-400, Skalar, Breda, The Netherlands) and concentrations determined following Grasshoff et al. (1999). Detection limits based on this approach were 0.1, 0.03 and 0.1 μM for NO_x^- , PO_4^{3-} and DSi, respectively.

Primary Productivity

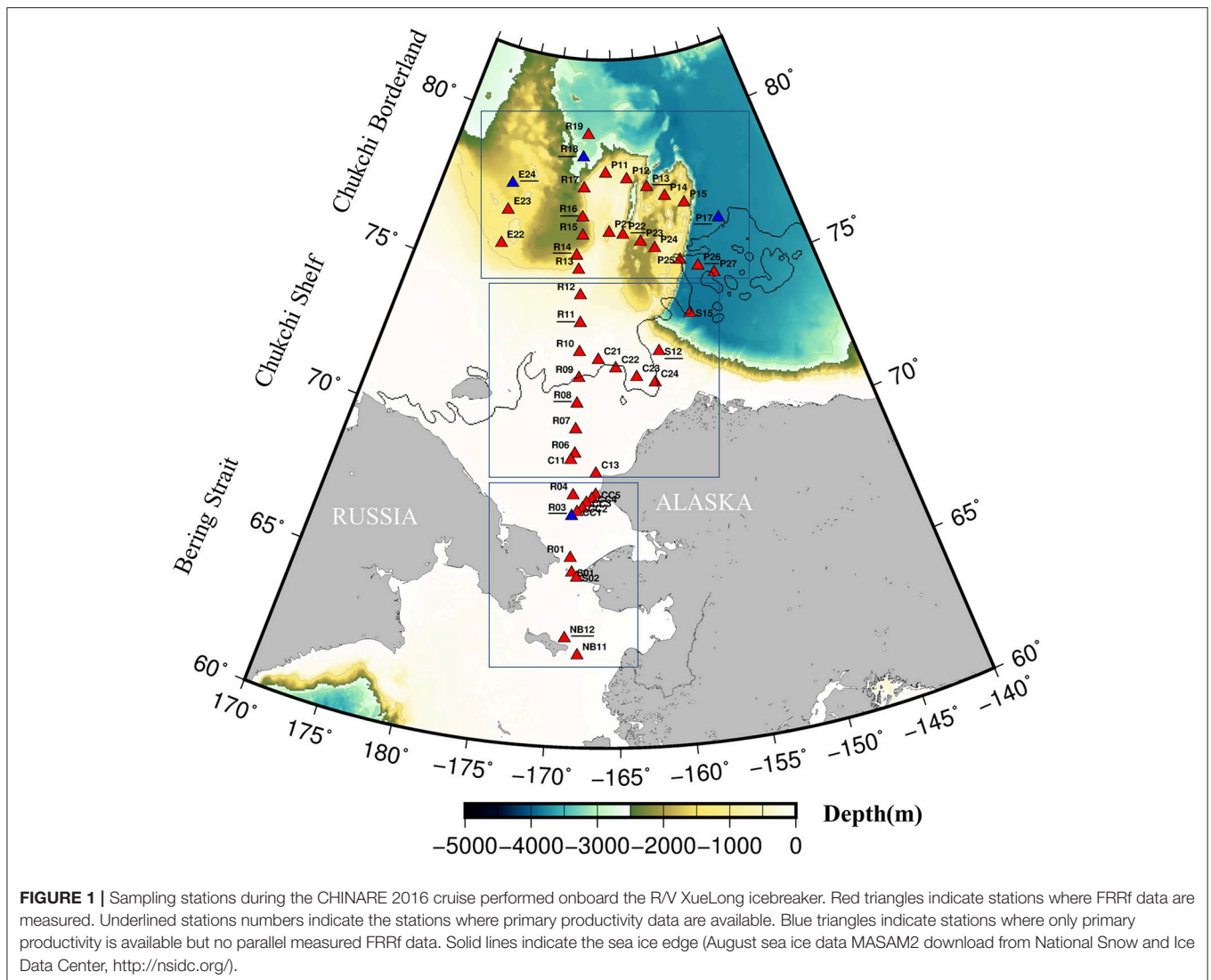
^{14}C uptake experiments for surface water samples ($\sim 5 \text{ m}$) only, were conducted using 6 h (local time 8:00 to 14:00) on-deck incubations as per Steemann Nielsen (1952) [modified by Evans et al. (1987) and (Ning and Liu, 1988)]. Briefly, water samples were prescreened through 200 μm mesh and then filled into two light and one dark acid-cleaned polycarbonate bottles of 250 mL 10 μCi of $\text{NaH}^{14}\text{CO}_3$ was added to each bottle, which was then placed in an on-deck incubator. When incubating samples from within the upper mixed layer, light intensity in the bottles was adjusted to its mean value within the mixed layer by covering neutral density screening. However, the effects of ultraviolet radiation were not considered in this study and we will discuss this later. A seawater circulation system-maintained incubators at the *in-situ* temperature. After incubation, water samples were filtered through GF/F filters under a low vacuum ($<0.02 \text{ MPa}$). The filters were then fumed by concentrated HCl overnight to remove inorganic carbon on the filters and preserved within scintillation vials -20°C for later measurement (Ning and Liu, 1988). Upon returning to the laboratory, 10 mL of scintillation counting cocktail (PerkinElmer, US) was added to each vial. Radioactivity was measured with a liquid scintillation counter (Tri-carb Packard 2050, US) that used an internal standard for a quenching correction. Carbon uptake rates (P^C , $\text{mgC m}^{-3} \text{ h}^{-1}$) were determined as daily rates ($\text{mgC m}^{-3} \text{ d}^{-1}$) as,

$$\text{daily } P^C = P^C \times \text{day length (h[d]}^{-1}) \quad (1)$$

where day length was calculated as per Kirk (1994). Daily Chl-a specific primary productivity (P_B^C , $\text{mgC mgChl-a}^{-1} \text{ d}^{-1}$) was then calculated as *daily* $P^C(z)$ divided by total Chl-a concentration.

Fast Repetition Rate Fluorometry (FRRf)

At each station, surface water was also taken for FRRf measurements (Act2-Based Laboratory system, Chelsea Technologies, West Molesey, UK) using the ship's seawater intake system (from a depth of ca. 5 m, similar with ^{14}C water sampling depth). All samples were acclimated to dark for $\sim 10 \text{ min}$ before measurement. The FRRf was programmed to deliver a single-turnover protocol with a saturation phase



comprising 100 blue flashlets (450 nm LED) on a 2 μs pitch and a relaxation phase comprising 40 flashlets on a 50 μs pitch (as per Hoppe et al., 2015; Hughes et al., 2018b). Measurements recorded were from the average of 58 consecutive acquisitions at 100-ms intervals. Each FRRf acquisition was subsequently fitted to the KPF model (Kolber et al., 1998) using FastPRO software (Chelsea Technologies Group) to yield the minimum fluorescence yield, maximum fluorescence yield, effective absorption, and photochemical efficiency of photosystem II (PSII) for darkness (F_o , F_m , σ_{PSII}^{450} and $F_v (= F_m - F_o)/F_m$) and for each actinic light level (F' , F'_m , σ_{PSII}^{450} and $F'_q (= F'_m - F')/F'_m$). The normalized Stern-Volmer quenching coefficient (NPQ_{NSV} , McKew et al., 2013) was calculated from these parameters as F'_o/F'_v where F'_o was estimated as $F'_o = F_o/(F_v/F_m + F_o/F'_m)$ (Oxborough and Baker, 1997) and $F'_v = (F'_m - F'_o)/F'_m$.

FRRf measurements were made continuously throughout light response protocols following an initial dark step. The sample was sequentially exposed to 10 actinic light levels (0, 48, 106, 175,

258, 358, 477, 621, 793, 1,000 $\mu\text{mol quanta m}^{-2} \text{s}^{-1}$, white LEDs) within the FastAct2, to retrieve a fluorescence-light response curve, whereby each light step was delivered for 40 s.

The instantaneous PSII reaction center normalized electron transport rate (ETR_{PSII} , $\text{mol e}^- [\text{mol PSII}]^{-1} \text{s}^{-1}$) for each light level was calculated as per Kolber and Falkowski (1993),

$$ETR_{PSII} = \text{PAR} \times \sigma_{PSII}^{450} \times q_p \times \Phi_{RC} \times 6.022 \times 10^{-3} \quad (2)$$

where PAR is in units of $\mu\text{mol quanta m}^{-2} \text{s}^{-1}$ and σ_{PSII}^{450} is the effective absorption cross section of PSII specific to light of 450 nm wavelengths. Φ_{RC} (electrons quanta^{-1}) accounts for the assumption that one electron is produced from each RCII charge separation (see Kolber and Falkowski (1993)), and the constant value 6.022×10^{-3} that converts $\mu\text{mol quanta}$ to quanta, PSII to mol PSII and A^2 to m^2 (e.g., Suggett et al., 2001). The term q_p (dimensionless) is the PSII operating efficiency and accounts for the extent of photochemical energy conversion

by PSII, determined as $(F'_m - F')/(F'_m - F'_0)$. To account for a lack of phytoplankton absorption and *in situ* light spectral measurements, an empirical relationship between “correction factor” (f , dimensionless) and optical depth was applied to spectrally correct values of σ_{PSII} (see Suggett et al., 2006b for detail). Specifically, for our surface data, an f value of 1.6 was used as optical depth was 0. As such, the spectrally corrected ETR_{PSII} is equal to ETR_{PSII} / f .

ETR_{PSII} and PAR data from the FRRf-light response curves were then fit to the photosynthesis-light dependency model of Jassby and Platt (1976), Equation (3).

$$ETR_{PSII} = ETR_{PSII}^{\max} \times \tanh\left(\frac{\alpha PAR}{ETR_{PSII}^{\max}}\right) \quad (3)$$

Using knowledge of α and ETR_{PSII}^{\max} , we were then able to retrieve the surface ETR_{RCII} for any given value of surface PAR at any given time. As such, the daily integrated ETR_{PSII} ($\text{mol e}^- [\text{mol PSII}]^{-1} \text{d}^{-1}$) at the surface was finally determined as:

$$\text{daily } ETR_{PSII} = \int_{t1}^{t2} ETR_{PSII} dt \quad (4)$$

where the period between $t1$ and $t2$ is daylength (h).

In order to convert ETR normalized to PSII content (ETR_{PSII}), normalized to Chl-*a* content, and hence ETRs that could be directly compared with parallel measures of carbon uptake to retrieve K_C (Lawrenz et al., 2013), knowledge of the RCII per Chl-*a* (i.e., n_{PSII} , $\text{mol RCII} [\text{mol Chl-}a]^{-1}$) is required. Direct measurement of n_{PSII} under natural conditions is extremely challenging (Moore et al., 2006; Suggett et al., 2006a) often requiring that the RCII concentration be determined indirectly (see Suggett et al., 2010). Since there are almost no cyanobacteria that exist in the Arctic region (Pedrós-Alió et al., 2015), here we assumed a “standard” n_{PSII} value of $0.002 \text{ mol RCII} [\text{mol chl}]^{-1}$ for eukaryotes (Kolber and Falkowski, 1993) in our ETR calculation.

Thus, a daily Chl-*a* specific ETR ($\text{mmol e}^- \text{ mg Chl-}a^{-1} \text{ d}^{-1}$) was calculated as follows:

$$\text{daily } ETR = \text{daily } ETR_{PSII} \times 0.002 \times 893^{-1} \quad (5)$$

the constant factor 893 converts $\text{mol Chl-}a$ to $\text{mg Chl-}a$ and mol e^- to mmol e^- .

Finally, K_C ($\text{mol e}^- (\text{mol C})^{-1}$) was defined to be the ratio of the two independently determined variables, ETR and P_B^C as per Zhu et al. (2016, 2017):

$$K_C = \frac{\text{daily } ETR}{P_B^C} \times 12 \quad (6)$$

where P_B^C is the daily-integrated carbon assimilation per unit Chl-*a* ($\text{mgC mg Chl-}a^{-1} \text{ d}^{-1}$), and the factor 12 converts g C to mol C .

Sea Ice Concentration

In this study the sea ice information was derived from MASAM2 sea ice concentration product (3 July 2012 to present; Fetterer et al., 2015) with 4 km resolution, which is bundled into monthly NetCDF data (<https://nsidc.org/data/g10005#>). We applied August 2016 data to obtain the ice concentration in the Arctic Ocean (60°N- 90°N) (<ftp://sidacs.colorado.edu/pub/DATASETS/NOAA/G10005/>). Sea ice concentration values >70% was defined as an ice cover area, for plotting the boundary of the ice free/ice cover region (Figure 1).

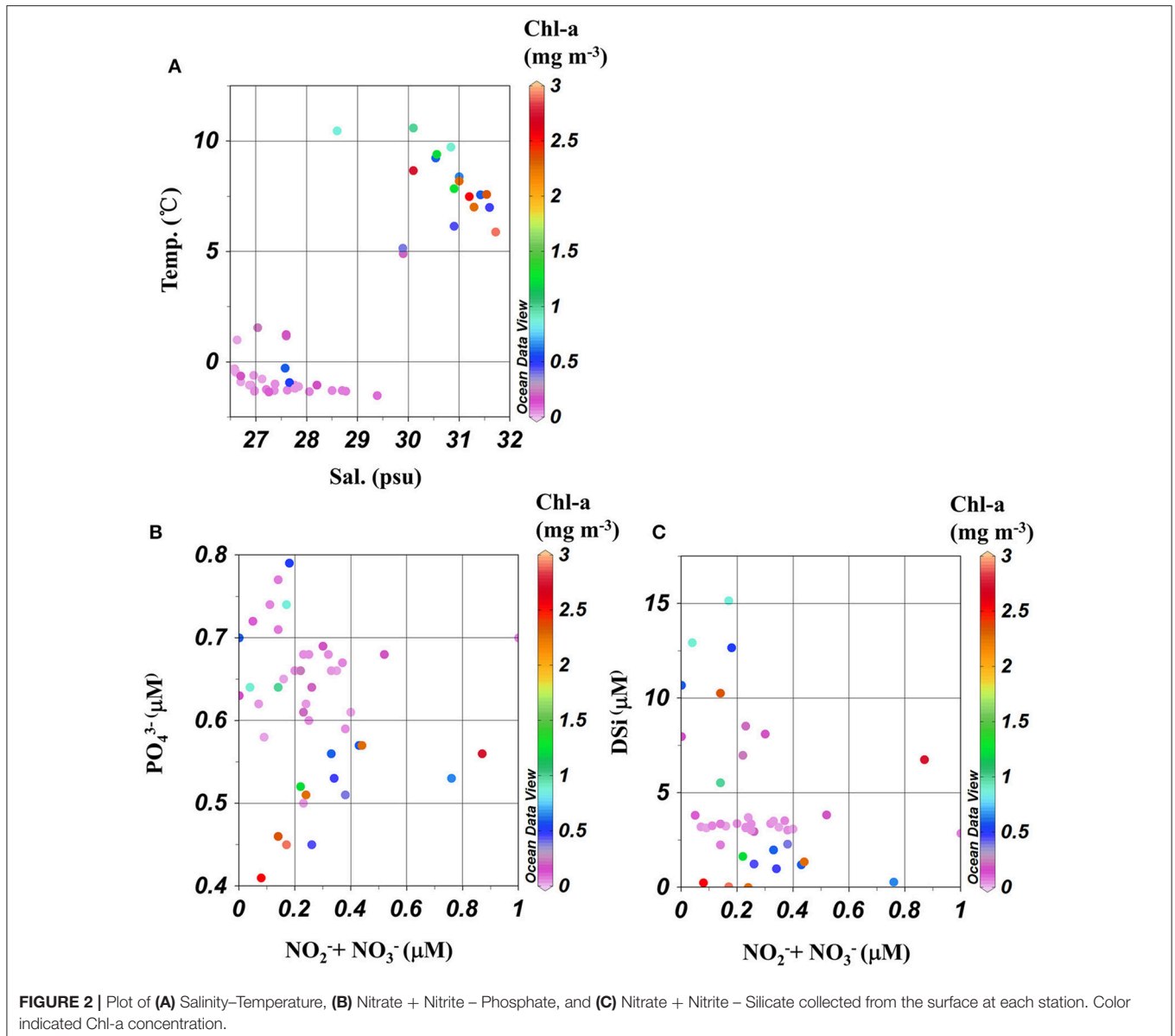
Statistical Analyses

Spearman and Pearson rank correlation analysis was utilized to examine the covariance of K_C with physico-chemical (or taxonomic) parameters. Kolmogorov Smirnov tests were applied to examine for data normality. Welch *t*-test, ANOVA and ANCOVA were then applied for testing the significant differences between clusters or groups of data and the linear regression models (as per Zhu et al., 2017). All statistical analyses and curve fittings were performed using open source statistical software R version 3.2.3 (R Core Team, 2014). Figures were plotted by GMT 4, Ocean Data View 5 (Schlitzer, 2018) and R software.

RESULTS

Physical and Nutrient Properties in the Study Area

Transiting across the Bering strait, Chukchi shelf and Chukchi borderland captured a gradient of physico-chemical conditions in the Arctic surface waters (Figure 1), whereby sea surface temperature (SST) decreased progressively from lowest (63°N) to highest (79.7°N) latitude, (mean \pm standard error, SE) of $8.8 \pm 0.3^\circ\text{C}$ in the Bering strait, $2.6 \pm 0.8^\circ\text{C}$ in the Chukchi shelf and $-1.1 \pm 0.1^\circ\text{C}$ in the Chukchi borderland (Supplementary Figure 1A, Figure 2A). Surface salinity exhibited a similar trend than SST, with the highest value of 31.54 at station NB12 in the Bering Strait, and decreasing northward to the lowest salinity (26.5 PSU) at station P26 in the Chukchi borderland (Supplementary Figure 1B, Figure 2A). Parallel changes of SST and salinity were observed between stations R8 and R9, when crossing the boundary of ice free and ice cover regions (Figure 1, Supplementary Figure 1), decreasing from 5°C to $<2^\circ\text{C}$, and 30 to 27 PSU, respectively (Supplementary Figures 1A,B). As such, this area delineated two water mass assemblies in the surface layer of cold, low-salinity water predominantly in the ice-free region and warm high-salinity water in the ice-cover area (T-S plot, Figure 2A). Mean (\pm SE) SST for the water mass in the ice-free area (stations included from NB11 to R8, $n = 17$) was $8.0 \pm 0.4^\circ\text{C}$, compared to $-0.56 \pm 0.2^\circ\text{C}$ in the ice cover region (stations located north than R8, $n = 32$). Salinity was 30.7 ± 0.2 vs. 27.5 ± 0.1 for the ice-free and ice-cover region, respectively. MLDs generally remained constant across the cruise transect, at ca. $14 \pm 2 \text{ m}$ (Table 1).



Relatively higher $\text{NO}_2^- + \text{NO}_3^-$ surface concentrations were observed for water masses in the ice-free than for the ice-cover waters, with mean (\pm SE) values of $0.31 \pm 0.05 \mu\text{M}$ and $0.25 \pm 0.03 \mu\text{M}$ (Table 1, Supplementary Figure 1C). In contrast, higher PO_4^{3--} ($0.66 \pm 0.01 \mu\text{M}$) and DSi ($4.3 \pm 0.4 \mu\text{M}$) concentrations were found for the ice cover region than for the ice free waters (PO_4^{3--} : $0.54 \pm 0.02 \mu\text{M}$; DSi: $3.8 \pm 0.4 \mu\text{M}$, Table 1, Supplementary Figures 1D,E). Even so, similarly low ratios of $\text{NO}_3^- + \text{NO}_2^-$ and PO_4^{3--} concentrations were found for both sub-regions (N:P of ca. 0.5), indicating nitrate availability was the limiting nutrient throughout the entire observation area (Tremblay and Gagnon, 2009). Nitrogen to silicate ratios ($\text{NO}_3^- + \text{NO}_2^-$ and DSi; N:Si) were ca. 10 times higher for the ice free (0.71 ± 0.36), than for the ice cover (0.07 ± 0.01) samples (Table 1, Figures 2B,C).

Phytoplankton Chl-a, Community Size Distribution and Photo-Physiology Variability

Chl-a varied by 2 orders of magnitude across the study area, from 0.01 to 2.73 mg m^{-3} (with mean \pm standard error (SE) of $0.52 \pm 0.11 \text{ mg m}^{-3}$ for all samples). As observed for SST and salinity, Chl-a concentration and phytoplankton size structure differed for water masses across the ice-free and ice-cover boundary. Chl-a decreased from 0.4 mg m^{-3} at R8 to 0.2 mg m^{-3} of R9, accompanied by a relative decrease in micro- and increase in pico-phytoplankton fractions (32 to 4% and 27 to 64% from R8 to R9, respectively). Higher Chl-a concentrations that were also comprised of more micro-phytoplankton were characteristic of the ice-free area (mean \pm SE of $1.4 \pm 0.2 \text{ mg m}^{-3}$ and $44 \pm 3\%$) compared to the

TABLE 1 | Mean \pm standard error (SE) of physico-chemical and biological parameters within ice free/ice cover regions.

Sampling parameters	Ice free	Ice cover	p-value
Temp.	8.0 \pm 0.4	-0.56 \pm 0.21	<0.01
Sal.	30.7 \pm 0.2	27.5 \pm 0.1	<0.01
MLD	16.5 \pm 1.8	12.2 \pm 0.7	0.01
NO _x ⁻	0.31 \pm 0.05	0.25 \pm 0.03	0.3
PO ₄ ³⁻	0.54 \pm 0.02	0.66 \pm 0.01	<0.01
DSi	3.9 \pm 1.1	4.3 \pm 0.4	0.6
N:P	0.57 \pm 0.09	0.38 \pm 0.04	0.04
N:Si	0.71 \pm 0.36	0.07 \pm 0.01	0.02
Chl-a	1.36 \pm 0.21	0.09 \pm 0.02	<0.01
Micro%	44% \pm 3%	6% \pm 1%	<0.01
Nano%	35% \pm 2%	36% \pm 5%	0.8
Pico%	21% \pm 2%	58% \pm 5%	<0.01
σ_{PSII}^{450}	490 \pm 2	474 \pm 12	0.3
F_v/F_m	0.52 \pm 0.01	0.42 \pm 0.02	<0.01
α_{PSII}	1.45 \pm 0.01	1.43 \pm 0.04	0.6
ETR_{PSII}^{max}	670 \pm 24	465 \pm 29	<0.01

Welch t-test results are shown comparing the difference between the groups. Value in bold indicate significant difference where $p < 0.05$.

ice-cover area (0.09 ± 0.01 mg m⁻³ and $6 \pm 1\%$) (Table 1, Supplementary Figures 2A,B). In contrast, pico-phytoplankton dominated the ice-covered samples, with mean values of $58 \pm 5\%$ (Supplementary Figure 2D). The nano-phytoplankton had similar fraction in two regions ($34 \pm 2\%$ and $35 \pm 4\%$ for ice-free and ice-cover area, Supplementary Figure 2C). Overall, Chl-a concentration and the proportion of biomass in the micro-phytoplankton fraction was strongly positively correlated throughout this cruise transect ($r = 0.8$, $n = 50$, $p < 0.001$).

Spatial distribution for photo-physiological parameters was highly variable throughout the region, whereby changes in values likely reflect a changing phytoplankton community structure (Chl-a biomass and size fractionation), (Supplementary Figure 3), as reported previously (Moore et al., 2005; Suggett et al., 2009b). Values of σ_{PSII}^{450} and F_v/F_m ranged from 370 to 654 Å² quanta⁻¹ and 0.10 to 0.58 respectively across the study area. However, F_v/F_m was higher (mean \pm SE) for samples in the ice free waters (0.52 ± 0.01) than for ice covered waters (0.42 ± 0.02) (Table 1, Supplementary Figure 3E) whereas values of σ_{PSII}^{450} were generally similar for both (490 ± 2 vs. 474 ± 12 Å² quanta⁻¹, respectively (Table 1, Supplementary Figure 3D). Values of α_{PSII} ranged from 1.2 to 2.4 mol e⁻ mol PSII⁻¹ s⁻¹ (μ mol quanta m⁻² s⁻¹)⁻¹, varying little throughout the cruise (mean \pm SE of 1.44 ± 0.18 mol e⁻ mol RCII⁻¹ s⁻¹ (μ mol quanta m⁻² s⁻¹)⁻¹) (Supplementary Figure 3A). In contrast, ETR_{PSII}^{max} ranged from 119 to 831 mol e⁻ mol PSII⁻¹ s⁻¹ with higher values in the ice free waters than in the ice covered waters (670 ± 24 vs. 465 ± 29 mol e⁻ mol PSII⁻¹ s⁻¹, Table 1, Supplementary Figure 3B). As such, the light saturation of PSII charge separation, E_K ($= ETR_{PSII}^{max}/\alpha_{PSII}$) were also higher for samples taken from ice free waters (460 ± 17 μ mol quanta m⁻² s⁻¹) compared to ice covered waters (335 ± 22 μ mol quanta m⁻² s⁻¹

(Supplementary Figure 3C). The extent of non-photochemical quenching (dark acclimated NPQ_{NSV} at light intensity of 48 μ mol quanta m⁻² s⁻¹) were all within the range of 0.66–2.07 (1.13 ± 0.06), except three relatively high values (ca. 3, 5, and 10) for stations R15, R16, and R17 (Supplementary Figure 3F).

To further evaluate the potential influence of phytoplankton taxa on photo-physiology and photosynthetic parameters, all data were subsequently binned according to the dominant phytoplankton fraction as per Zhu et al. (2017); specifically selecting samples based on their larger proportion of cells within micro-, nano-, or pico- fraction to yield three effective phytoplankton size dominated groups (f_{micro} -, f_{nano} -, and f_{pico} -dominated consisting of nine, 15 and 26 data points, respectively). Hydrographic properties generally differed for samples binned according to these three size groups (ANOVA test, $p < 0.05$, $df = 2$; Table 2). High temperature, salinity and Chl-a concentration was a consistent feature for samples characterized by the dominance of the largest size phytoplankton dominated group (*micro fraction* > 50%, Table 2, Supplementary Figure 4), and largely for stations from the Bering strait. For f_{nano} and f_{pico} dominated samples, Chl-a, temperature and salinity were always lower. Whilst PO₄³⁻ concentrations exhibited relatively little variation across all 3 size-class bins, higher NO₂⁻ + NO₃⁻ associated with lower DSi concentration were a common feature of f_{micro} dominated samples (Table 2, Supplementary Figure 4). ETR_{PSII}^{max} and F_v/F_m significantly increased for samples dominated by increasingly larger fractions (ANOVA test, $p < 0.01$, $df = 2$, Table 3, Figure 3), whereas α_{PSII} and σ_{PSII}^{450} did not vary amongst the size groups (ANOVA test, $p = 0.5$ and 0.6 , $df = 2$, Table 3, Figure 3); However, ANOVA test analysis further demonstrated that the ETR_{PSII}^{max} and F_v/F_m for f_{pico} were significantly different than for f_{nano} and f_{micro} ($df = 1$, $p < 0.001$) but not for f_{nano} compared to f_{micro} ($df = 1$, $p = 0.3$).

Variability of ETR, Carbon Uptake Rates, and K_C

Daily integrated ETR ranged from 12.7 to 54.9 mmol e⁻ (mg Chl-a)⁻¹ d⁻¹, with a mean (\pm SE) of 33.6 ± 1.4 mmol e⁻ (mg Chl-a)⁻¹ d⁻¹ from all samples. Daily ETR was significantly linearly correlated with daily integrated PAR (with range of 6.0 to 21.1 mol quanta m⁻² d⁻¹) ($R^2 = 0.86$, $n = 46$, $p < 0.001$; Supplementary Figure 5A), revealing that variability of daily ETR was highly dependent on light availability throughout this study region, and as previously identified from other regions (Zhu et al., 2016, 2017).

Whilst relatively few ¹⁴C uptake rate data points ($n = 13$) were ultimately obtained from this cruise, values of surface P^C were highly correlated with corresponding values of Chl-a, and with a variability best described by a second-order polynomial function ($R^2 = 0.99$, $n = 13$; Supplementary Figure 5B). We therefore applied this empirical model to derive the carbon uptake rate for all 46 stations where both surface Chl-a and FRRf data were available. From this, surface P^C varied from 0.18 mgC m⁻³ d⁻¹ at the Chukchi Borderland (P21-P26) to 102.5 mgC m⁻³ d⁻¹ at Bering Strait (S01), with average values of 1.7 ± 0.4 and 37.9 ± 8.2 mgC m⁻³ d⁻¹ for the ice covered and ice free water masses,

TABLE 2 | Mean (\pm SE, standard error) of phytoplankton size fractions (Micro%, Nano%, Pico%), biomass (Chl-a, mg m^{-3}) and environmental parameters [Temp., Temperature ($^{\circ}\text{C}$); Sal, Salinity; MLD, Mixed Layer Depth (m); $\text{NO}_x^- = \text{NO}_2^- + \text{NO}_3^-$ (μM), PO_4^{3-} (μM), $\text{DSi}(\mu\text{M})$] within three dominated size classes of phytoplankton.

Dominated groups	Size composition (%)&Biomass				Hydrographic properties					
	Micro%	Nano%	Pico%	Chla	Temp.	Sal.	MLD	NO_x^-	PO_4^{3-}	DSi
Micro- ($n = 9$)	53.1 (3.9)	30.6 (3.4)	16.3 (2.3)	1.81 (0.28)	8.3 (0.3)	30.9 (0.2)	17.8 (2.2)	0.39 (0.09)	0.52 (0.02)	2.8 (1.2)
Nano- ($n = 15$)	18.2 (4.1)	54.3 (5.3)	27.5 (3.8)	0.40 (0.09)	3.3 (1.1)	29.0 (0.5)	13.8 (1.6)	0.24 (0.03)	0.61 (0.02)	4.9 (1.1)
Pico- ($n = 26$)	8.8 (2.4)	23.5 (3.2)	67.4 (4.6)	0.11 (0.03)	0.1 (0.6)	27.6 (0.2)	11.7 (0.9)	0.25 (0.04)	0.66 (0.01)	4.6 (0.6)
ANOVA test	$p < \mathbf{0.001}$	$p < \mathbf{0.01}$	$p < \mathbf{0.001}$	$p < \mathbf{0.001}$	$p < \mathbf{0.001}$	$p < \mathbf{0.001}$	$p = \mathbf{0.04}$	$p = 0.3$	$p < \mathbf{0.01}$	$p = 0.4$

ANOVA test results are shown comparing the difference between the groups. Values in bold indicate significant differences where $p < 0.05$.

TABLE 3 | Mean (\pm SE, standard error) of photo-physiology parameters α_{PSII} ($\text{mol e}^- \text{ mol RCII}^{-1} \text{ s}^{-1}$ ($\mu \text{ mol quanta m}^{-2} \text{ s}^{-1}$) $^{-1}$), ETR_{PSII}^{max} ($\text{mol e}^- \text{ mol RCII}^{-1} \text{ s}^{-1}$), σ_{PSII}^{450} ($A2 \text{ quanta}$), F_v/F_m within three dominated size classes of phytoplankton.

Dominated groups	α_{PSII}	ETR_{PSII}^{max}	σ_{PSII}^{450}	F_v/F_m
Micro- ($n = 8$)	1.48 (0.01)	675 (32)	490 (29)	0.51 (0.01)
Nano- ($n = 15$)	1.39 (0.02)	615 (39)	468 (13)	0.49 (0.02)
Pico- ($n = 23$)	1.45 (0.05)	436 (31)	483 (13)	0.4 (0.02)
ANOVA test	$p = 0.5$	$p < \mathbf{0.001}$	$p = 0.6$	$p = \mathbf{0.004}$

ANOVA test results are shown comparing the difference between the groups. Values in bold indicate significant differences where $p < 0.01$.

respectively. After normalizing to Chl-a, surface P_B^C appeared higher for stations in the ice free water (mean \pm SE: $25.9 \pm 1.6 \text{ mgC mg Chl-a}^{-1} \text{ d}^{-1}$) than in the ice covered area ($17.4 \pm 0.2 \text{ mgC mg Chl-a}^{-1} \text{ d}^{-1}$). Binning P_B^C data to the different dominated size phytoplankton groups (i.e., f_{micro} , f_{nano} and f_{pico}) demonstrated that P_B^C decreased with the phytoplankton size decrease, 30.2 ± 2.8 , 19.6 ± 0.6 and $17.5 \pm 0.3 \text{ mgC mg Chl-a}^{-1} \text{ d}^{-1}$ for f_{micro} , f_{nano} , and f_{pico} , respectively (Table 4).

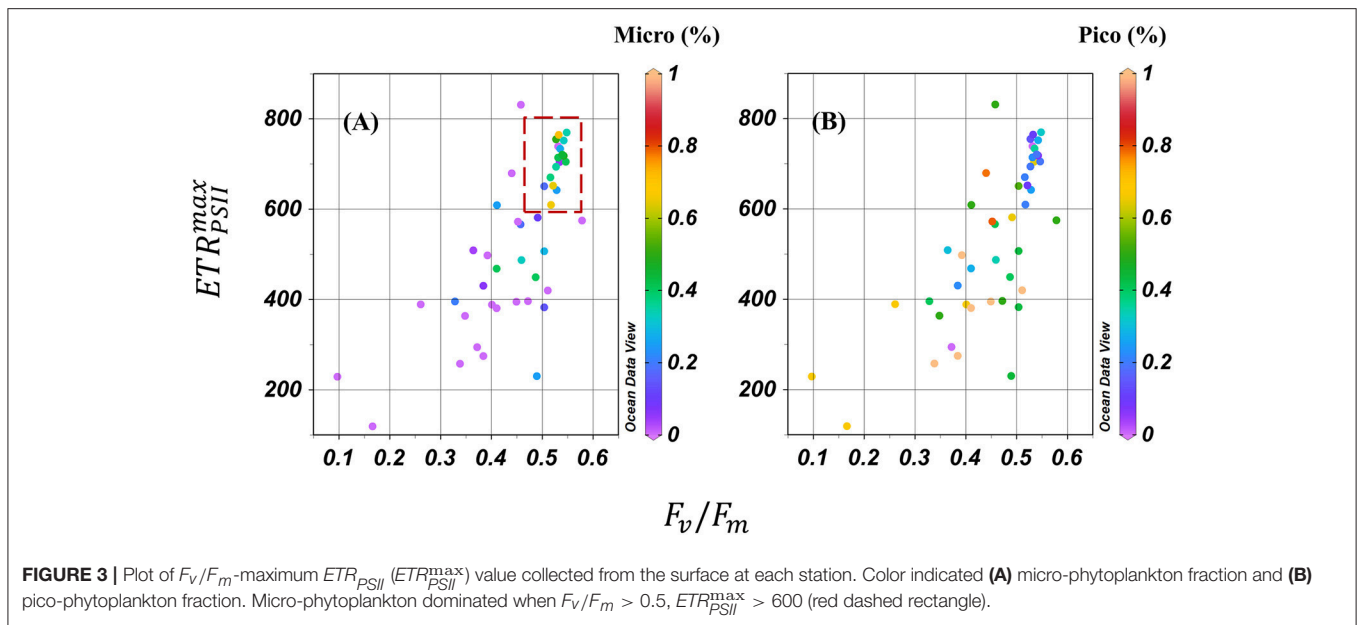
Values of K_C derived from corresponding daily ETR and C-uptake rates (Equation 6) are key in understanding phytoplankton electron usage efficiency for carbon fixation. K_C exhibited a range from 8.8 to $38.5 \text{ mol e}^- (\text{mol C})^{-1}$ with an average of $20.5 \pm 6.3 \text{ mol e}^- (\text{mol C})^{-1}$ during the cruise. Lower K_C values were observed for ice-free than for ice-covered samples (mean \pm SE, 19.5 ± 1.6 vs. $21.0 \pm 1.2 \text{ mol e}^- (\text{mol C})^{-1}$) or for samples dominated by f_{micro} than for f_{nano} and f_{pico} (16.8 ± 1.8 vs. 24.0 ± 1.2 and $19.4 \pm 1.0 \text{ mol e}^- (\text{mol C})^{-1}$). ANOVA demonstrated that K_C was different amongst the 3 phytoplankton size groups, with Welch t -tests identifying that K_C was the same for f_{micro} and f_{nano} ($df = 1$, $p = 0.02$), f_{nano} and f_{pico} ($df = 1$, $p = 0.05$), but not for f_{micro} and f_{pico} ($df = 1$, $p = 0.3$).

Relationships between PAR and K_C were further examined for data binned according to water mass (ice-free vs. ice-cover samples) and then according to the three dominated size groups. Variability of K_C for the entire dataset could be described by a simple PAR-dependent linear model ($K_C = 0.98\text{PAR} + 7.4$, $R^2 = 0.43$, $n = 46$, $p < 0.001$, Figure 4A) as we have demonstrated previously for other regions in the Ariake Bay and East China Sea (Zhu et al., 2016, 2017). However, linear regression models

describing the relationship between K_C and PAR were different for samples pooled for ice free ($K_C = 0.98\text{PAR} + 4.02$, $R^2 = 0.35$, $n = 16$, $p = 0.01$) vs. ice cover ($K_C = 1.42\text{PAR} + 3.93$, $R^2 = 0.77$, $n = 30$, $p < 0.001$) regions (Figure 4B), or f_{micro} ($K_C = 0.54\text{PAR} + 8.2$, $R^2 = 0.21$, $n = 8$, $p = 0.25$), f_{nano} ($K_C = 1.62\text{PAR} - 0.4$, $R^2 = 0.75$, $n = 15$, $p < 0.001$), and f_{pico} ($K_C = 1.2\text{PAR} + 5.7$, $R^2 = 0.69$, $n = 23$, $p < 0.001$) (Figure 4C). For the water mass bins, the linear regression slopes appeared lower for ice free (0.98) than for ice cover (1.42) samples, but were not significantly different (ANCOVA, $df = 1$, $p = 0.2$). In contrast, for the size group bins, ANCOVA analysis demonstrated that the regression slope describing the K_C vs. PAR relationships for f_{micro} was significantly lower than for f_{nano} and f_{pico} ($df = 1$, $p = 0.03$ and 0.05) but not for f_{nano} compared to f_{pico} ($df = 1$, $p = 0.2$). As such, data from f_{nano} and f_{pico} bins were subsequently pooled to generate the PAR- K_C linear relationship for samples dominated by small phytoplankton ($f_{\text{nano+pico}}$, $< 20 \mu\text{m}$) (Figure 4D). In this case, the regression slope was lower for samples dominated by micro-phytoplankton ($f_{\text{micro}} > 20 \mu\text{m}$) than those dominated by small phytoplankton ($f_{\text{nano+pico}}$, $< 20 \mu\text{m}$) (slope: 0.54 vs. 1.36, ANCOVA, $df = 1$, $p = 0.02$). The correlation coefficient between K_C and PAR was much improved for data binned by water mass for the ice free region ($R^2 = 0.77$, Figure 4B) or by taxa for the $f_{\text{nano+pico}}$ group ($R^2 = 0.74$, Figure 4D) compared to that previously where all data was pooled together ($R^2 = 0.43$, Figure 4A). Overall, the data pooled as two taxonomic groups (compared to two water masses) appeared best at explaining covariation between PAR and K_C .

DISCUSSION

Summer has been a recent focus of study within the Arctic Ocean because of the rapid reduction of sea ice that appears to drive a series of dramatic changes in ocean biogeochemistry. Sea ice melting results in warming and freshening of surface waters (Screen and Simmonds, 2010), which subsequently enhances the stratification of the upper Arctic Ocean. Stratification suppresses the water exchange between the surface and deep layers resulting in nutrient deficiency in the upper Arctic Ocean waters after phytoplankton blooms (Li et al., 2009). As a result, phytoplankton must respond to evolving physico-chemical conditions through changes in community structure (Ardyna et al., 2011), growth period (Arrigo and van Dijken, 2004; Ardyna et al., 2014), and



biomass (Pabi et al., 2008; Wassmann et al., 2011), which together influence the entire food web (Palmer et al., 2013) and ecosystem dynamics (Smetacek and Nicol, 2005). We used parallel measures of FRRf and ^{14}C uptake to better understand phytoplankton responses to dynamically changing Arctic Ocean surface waters in the summer, encompassing both ice free and ice cover water masses. It needs to be noted that this study, concerning the surface (upper mixed layer) only, may not fully reflect photophysiology and primary productivity characteristics of the Arctic Ocean, especially in the lack of sub-surface Chla maximum (SCM), which has great importance in Arctic regions. Hill et al. (2013) recently suggested a constant PP underestimation of 75% throughout the summer for the entire Arctic Ocean when smoothing out the SCM. Moreover, the light- K_C relationship derived from the surface may be different, or not applicable for data from below the mixed layer (Schuback et al., 2017), especially for the SCM where the phytoplankton size structure differs with regards to the surface layer (Wang et al., 2014). We also acknowledge that taxa could change *in situ* with a similar cell size and only focusing on phytoplankton cell size structures may not be enough to fully explain our results (Suggett et al., 2004). Unfortunately, the data cannot address it currently, but we have noted that it may have contributed to the unexplained variance (**Figures 4C,D**). Arctic Ocean surface/upper waters were our focus because of the sensitivity to ocean warming, ice retreat etc., which further affects primary production and organic matter circulation (Benner et al., 2005; Arrigo et al., 2008; Steele et al., 2008; Brown and Arrigo, 2012). Thus, we are confident our work contributes to a better understanding of the relationship between phytoplankton size structure and photo-physiology of Arctic Ocean upper water. Our observations of distinct photophysiological and phytoplankton community signatures for both ice free and ice cover water masses in the summer Arctic Ocean, is consistent with previous investigation

in this area (2010–2011), where differences in light and nutrient availability associated with the seasonal sea ice zone resulted in strong patterns of (carbon-based) photophysiology (Palmer et al., 2013). However, we demonstrate that different water masses also display inherently distinct properties in how PSII electron transport (ETR_{PSII}) is invested into carbon uptake, and hence values of K_C . In the following sections, we consider how such differences in K_C may contribute to PP dynamics in the Arctic Ocean surface water, and how these observations compare to environmental (notably light) -dependency of K_C in other ocean systems.

Environmental and Phytoplankton Community Characteristics of Summer Waters in the Arctic Ocean

The Arctic Ocean has been warming over the past few decades (Zhang, 2005) whereby increasing surface atmospheric heat fluxes have effectively entered through the Bering Strait due to the absence of ice cover (Woodgate et al., 2006; Steele et al., 2008). At the same time, accelerating ice melt has been enhancing summer Arctic Sea freshening (Li et al., 2009; Cai et al., 2010). Significant regional variability for temperature and salinity from our study appears to be most associated with sea ice cover and crossing the ice cover edge (i.e., R8 to R9, **Figure 1, Supplementary Figure 1**). Lower salinity within the upper layer of ice melt is likely to result in stronger stratification (Frey et al., 2014). Weaker mixing will in turn reduce winter nutrient renewal to the euphotic layer and thus limit summer primary productivity (McLaughlin and Carmack, 2010). Indeed, whilst MLD remained constrained throughout the transect, MLD was generally deeper in ice-free waters (16.5 m) than for ice-covered water (12 m) (see **Table 1**). Less nutrient availability is expected for phytoplankton in the upper layer of ice covered waters since nutrient rich Pacific winter water is much deeper over the Chukchi Borderland (100 m) than

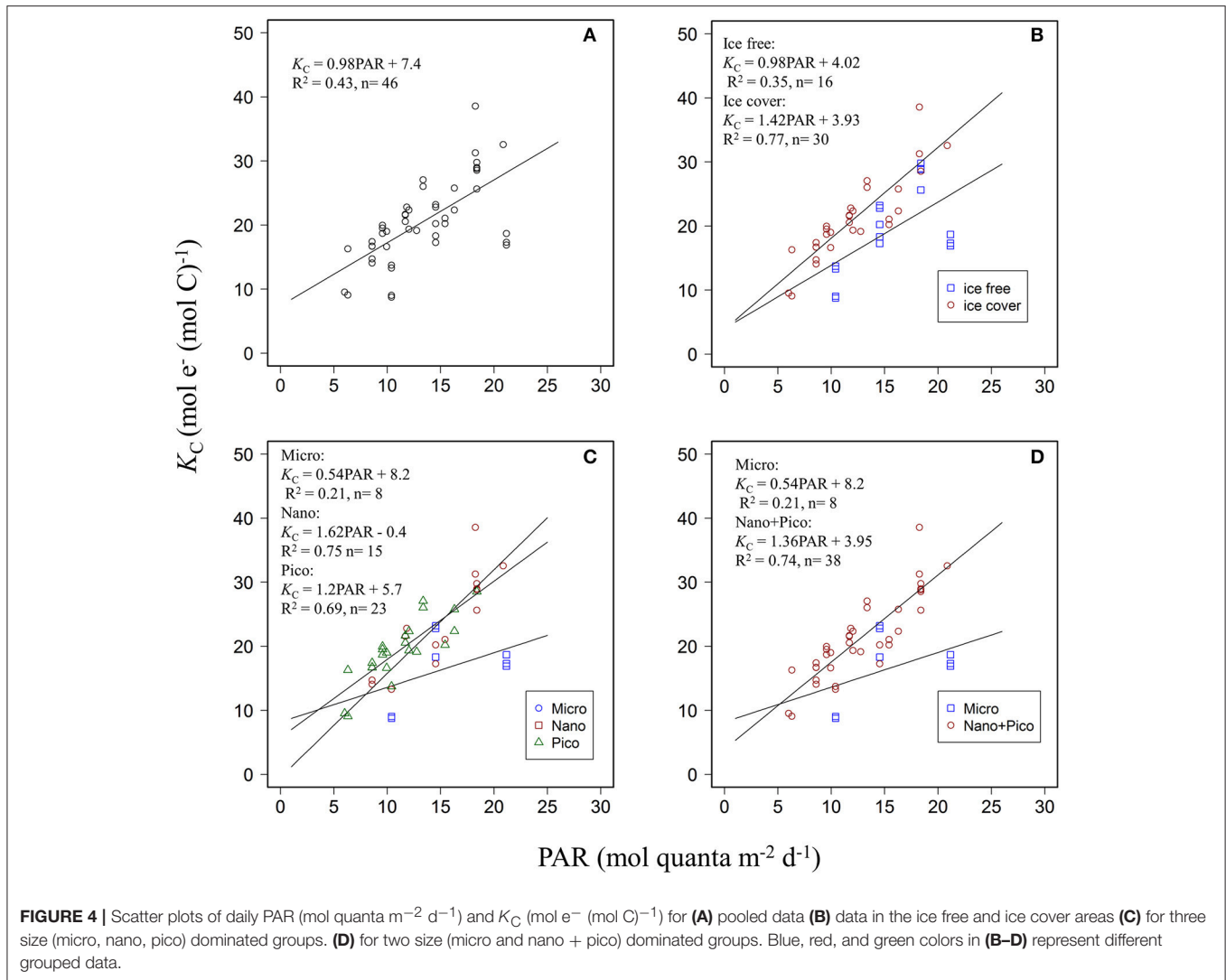


FIGURE 4 | Scatter plots of daily PAR (mol quanta m⁻² d⁻¹) and K_C (mol e⁻ (mol C)⁻¹) for (A) pooled data (B) data in the ice free and ice cover areas (C) for three size (micro, nano, pico) dominated groups. (D) for two size (micro and nano + pico) dominated groups. Blue, red, and green colors in (B–D) represent different grouped data.

the Chukchi Shelf (20–50 m) (Coupel et al., 2015). However, in our study, $\text{NO}_2^- + \text{NO}_3^-$ was only slightly higher for the ice free than for the ice covered waters (and DSi and PO_4^{3-} in fact lower) and similar results were also observed in Palmer et al. (2013). Whilst our data therefore appears at odds with the expected trends, it possibly reflects enhanced phytoplankton nutrient uptake in the ice free water mass (Cermeño et al., 2006). We return to this point below.

Phytoplankton biomass and size structure observed in our study agree well with values for summer Arctic phytoplankton communities previously reported in the same region (Coupel et al., 2012, 2015). Our range and spatial distribution for Chl-*a* is consistent with Coupel et al. (2015), who showed high mean concentrations ($\sim 1 \text{ mg m}^{-3}$) on the Chukchi shelf compared to $< 0.1 \text{ mg m}^{-3}$ further north. Large cells, such as diatoms *Chaetoceros* sp., *Fragilaria* sp., *Cylindrotheca* sp., and dinoflagellates dominate the Bering Strait and Chukchi shelf (Coupel et al., 2012), and thus presumably contribute to the high micro-phytoplankton fraction found throughout

this similar area (Supplementary Figure 2B). In contrast, prasinophytes such as *Micromonas* sp. ($< 2 \mu\text{m}$ diameter) mainly dominate surface waters with low Chl-*a* ($< 0.1 \text{ mg m}^{-3}$), i.e., values consistent with our pico-phytoplankton fraction (Supplementary Figure 2D). Nutrient availability is often the major factor influencing phytoplankton size structure (e.g., Marañón et al., 2015), and we observed larger cell fractions (and more Chl-*a* biomass) associated with slightly more NO_x^- (but less DSi and PO_4^{3-}) within the ice free waters. Decoupling between dissolved nutrient supply and apparent concentrations may reflect alternate utilization rates by prevailing phytoplankton communities (Cermeño et al., 2006), and thus it is plausible that nutrients may have been consumed and invested into biomass (higher Chl-*a*) in the ice-free waters immediately prior to the observation period. This notion is supported by the significant negative relationship between nutrient (PO_4^{3-} , DSi) and micro%, (PO_4^{3-} : $r = -0.42$, $p < 0.01$; DSi: $r = -0.3$, $p = 0.04$, spearman, Supplementary Figure 5), suggesting a large draw down by diatoms in the ice free region (see also Lewis et al.,

1996), and the increase in large phytoplankton in parallel with total biomass is in agreement with many studies indicating fast opportunistic growth by diatoms in response to nutrient and light availability (Cermeño et al., 2005a; Kameda and Ishizaka, 2005; Brewin et al., 2010).

FRRf-based photophysiology parameters (i.e., F_v/F_m and σ_{PSII}) are expected to covary with phytoplankton taxa (Moore et al., 2005; Suggett et al., 2009b), where larger F_v/F_m and smaller σ_{PSII} are expected for larger taxa (cells). Our FRRf values of F_v/F_m were higher for larger size phytoplankton dominated samples (Table 3), which further corroborates these dynamics. That said, F_v/F_m was also significantly positively correlated with Chl a and micro% ($r = 0.5$ and 0.6 , respectively; $p < 0.01$, Supplementary Figure 4), but negatively correlated with PO_4^3 and DSi ($r = -0.5$ and -0.3 , respectively; $p < 0.05$, Supplementary Figure 4) again consistent with possible recent relief from nutrient limitation for large size phytoplankton. Whilst σ_{PSII} is also expected to increase as cell size declines (Suggett et al., 2009b), we observed no differences in σ_{PSII} amongst sizes bins ($p = 0.6$, Table 3) and the reason for this is unclear. However, examining σ_{PSII} for the pico-phytoplankton dominated dataset in more detail along the R section from R17 to R11 (Figure 1), in fact demonstrates that σ_{PSII}^{450} was relatively large (mean \pm SE, $553 \pm 151 \text{ \AA}^2 \text{ quanta}^{-1}$) compared to the P section (P27 to P21; $405 \pm 70 \text{ \AA}^2 \text{ quanta}^{-1}$, Supplementary Figure 3D) suggesting additional localized environmental regulation of σ_{PSII} within this region that presumably confounds the role of taxa alone in predominantly influencing σ_{PSII} .

Phytoplankton Size Related ETR, Production, and K_C Variation in the Summer Arctic Ocean

Whilst values of σ_{PSII}^{450} were unrelated to phytoplankton size (see above, Table 3), both ETR_{PSII}^{max} and F_v/F_m were higher for the larger size fractions (f_{micro} and f_{nano}) (Table 3). Higher values of both F_v/F_m and ETR_{PSII}^{max} are expected for larger size phytoplankton (Cermeño et al., 2005b; Giannini and Ciotti, 2016) and from laboratory cultures (Suggett et al., 2009a; Blache et al., 2011), suggesting larger cell phytoplankton are indeed growing under more favorable conditions (Cermeño et al., 2005a,b). Larger phytoplankton dominating in the ice free area were acclimated to higher light regimes as compared to those under ice cover (E_k : 455 ± 23 vs. $218 \pm 18 \mu\text{mol quanta m}^{-2} \text{ s}^{-1}$), which also explains higher ETR_{PSII}^{max} achieved for this group (Schuback et al., 2017, Table 4). Average surface instantaneous PAR was ca. $190 \mu\text{mol quanta m}^{-2} \text{ s}^{-1}$ ($=13 \text{ mol quanta m}^{-2} \text{ d}^{-1}$) and hence below values of E_k , suggest that phytoplankton in both ice free and ice cover waters were growing under light limitation, and as such one would expect PAR and ETR to highly co-vary (Supplementary Figure 5A, Table 4, Hancke et al., 2015).

Values of P_B^C were higher in samples where the large size fraction dominated (Table 4), indicating that larger phytoplankton have higher photosynthetic efficiency when growing under optimum environmental conditions, such as higher light, temperature and nutrients (Palmer et al., 2013). Higher F_v/F_m and enhanced photochemical efficiency of PSII for

TABLE 4 | Summary of mean value (standard errors) of daily PAR (mol quanta $\text{m}^{-2} \text{ d}^{-1}$), daily ETR (mmol $\text{e}^- [\text{mgChl-a}]^{-1} \text{ d}^{-1}$), P_B^C (mg C $[\text{mgChl-a}]^{-1} \text{ d}^{-1}$), and K_C (mol $\text{e}^- (\text{mol C})^{-1}$) at surface layer of dominated size classes of phytoplankton.

Dominated groups	PAR	daily ETR	P_B^C	K_C
Micro- ($n = 8$)	15.9 (1.5)	40.3 (3.3)	30.23 (2.28)	16.8 (1.8)
Nano- ($n = 15$)	15.1 (1.0)	38.2 (2.3)	19.58 (0.65)	24.0 (1.3)
Pico- ($n = 23$)	11.1 (0.7)	28.2 (1.4)	17.45 (0.25)	19.4 (1.0)
ANOVA test	$\rho = \mathbf{0.001}$	$\rho < \mathbf{0.001}$	$\rho < \mathbf{0.001}$	$\rho = \mathbf{0.02}$

ANOVA test results are shown comparing the difference between 3 groups. Values in bold indicate significant differences where $p < 0.05$.

larger phytoplankton dominated samples may provide a possible mechanism to account for the higher P_B^C (Cermeño et al., 2005b). Furthermore, phytoplankton acclimated to higher irradiance appeared to lower Chl-a concentration per cell, leading to an increase of Chl-a normalized productivity, which also explains our results that P_B^C value appeared larger in the ice free region than in the ice cover area.

Concomitant changes in ETRs and C-uptake rates resulted in lower values of quantum requirement for carbon fixation (i.e., K_C) for f_{micro} than for f_{nano} and f_{pico} (16.8 ± 1.8 vs. 24.0 ± 1.3 and $19.4 \pm 1.0 \text{ mol e}^- (\text{mol C})^{-1}$). This observation (lowest K_C was observed for f_{micro} , Table 4) is consistent with the observations of Suggett et al. (2009a) that K_C was smallest for diatoms among six different eukaryotic algal taxa grown under identical nutrient replete conditions in the laboratory but contrasts with our recent observations from the East China Sea where K_C was higher for the large phytoplankton dominated community (Zhu et al., 2017). Phytoplankton productivity is however expected to vary under different environmental conditions (e.g., temperature, light, nutrient etc.) even for the same size community (Shiomoto et al., 1997; Staehr et al., 2002; Halsey et al., 2011; Zhu et al., 2017), and confirms that both environment and taxonomy interact to regulate K_C variation (Lawrenz et al., 2013).

Light availability has previously been shown to strongly regulate K_C (e.g., Brading et al., 2013; Zhu et al., 2016), and we therefore examined the variation of PAR- K_C relationships as a result of phytoplankton community changes as per Zhu et al. (2016, 2017). Spearman correlation and nMDS analyses revealed that PAR indeed had the highest correlation coefficient with K_C (Supplementary Table 2; Supplementary Figure 6, $r = 0.66$, $p < 0.001$), which is consistent with our previous results (Zhu et al., 2016, 2017). Changes in light availability drive an increase in the need to dissipate absorbed excitation energy through non-photochemical dissipation (McKew et al., 2013) and hence a positive association between K_C and NPQ may also be expected, as broadly observed previously (Schuback et al., 2015, 2016, 2017; Hughes et al., 2018b; Ryan-Keogh et al., 2018). Since the K_C of this study was derived from ETR and primary productivity in daily scales, similar to the method applied for daily ETR calculation, we also integrated the NPQ_{NSV} values according to their light dependent functions (Supplementary Figure 7A). However, no positive relationship between NPQ_{NSV} and K_C was

evident for this data set (**Supplementary Figure 7B**). Lack of a relationship (or negative relationship) between NPQ or PAR with K_C was also reported before (Schuback et al., 2017; Ryan-Keogh et al., 2018). Whilst the reason is not well-known yet, it may reflect a decoupling of the NPQ_{NSV} (or PAR)- K_C relationship under relatively low light (Schuback et al., 2017) and further investigation is needed.

Lower vs. higher K_C observed for larger vs. smaller phytoplankton (i.e., f_{micro} and $f_{\text{pico+nano}}$) resulted in differences in PAR- K_C linear regressions (**Figure 4D**), confirming that knowledge of the dominant phytoplankton taxa (or at least size class) present is crucial for the improvement of light-dependent retrieval of K_C (Zhu et al., 2017). The smaller slope value of the PAR- K_C linear regression for large phytoplankton, presumably diatoms, suggests that under relatively high light intensities micro-phytoplankton have a higher efficiency of electron usage for carbon fixation. When grown at higher and/or fluctuant irradiances, warmer temperature and without nutrient limitation, diatoms tend to express higher F_v/F_m values (Cermeño et al., 2005b; Suggett et al., 2009b), which inherently increases the transfer of excitation energy to the PSII reaction center to sustain electron transport activity compared to other species (Owens, 1986; Ott et al., 1999; Nymark et al., 2009; Brunet and Lavaud, 2010). Higher F_v/F_m values lead to higher Chl-a normalized C-assimilation rates (Falkowski, 1980; Zhao et al., 2015). Such inherent physiology may explain the lower K_C observed for f_{micro} than for $f_{\text{pico+nano}}$, especially when daily PAR was relatively high (i.e., >20 mol quanta $\text{m}^{-2} \text{d}^{-1}$, **Figure 4D**).

Contrasting PAR- K_C Estimate Models Between Different Study Regions

A major factor in the power of FRRf-type sensors is the potential for describing primary productivity over extremely broad but resolute temporal and spatial scales (Fujiki et al., 2008; Suggett et al., 2009b; Cheah et al., 2011; Hughes et al., 2018a). To further improve the accuracy with which carbon uptake rates are estimated from FRRf, many recent studies have attempted to develop K_C prediction models from environmental variables to avoid errors introduced by assuming K_C is constant (Lawrenz et al., 2013; Schuback et al., 2015; Zhu et al., 2016, 2017; Hughes et al., 2018b). Light variability appears to be the primary driver of variability in K_C , which is reflective of ETRs decoupled from C-fixation under light stress. However, it is important to note that the tight correlation between PAR (or NPQ) and K_C may be due to the two parameters not being entirely independent of K_C , since PAR (or F_v) is used in the derivation of both ETR (or NPQ) and K_C , (Schuback et al., 2015; Zhu et al., 2017). Nevertheless, using NPQ or PAR as a predictor of K_C is highly desirable over regional scales (Schuback et al., 2015, 2016, 2017; Zhu et al., 2016, 2017; Hughes et al., 2018b; and Ryan-Keogh et al., 2018). While examining the empirical light- K_C regression with larger data sets, however, variations of this relationship (e.g., their regression slopes differ) have been observed within different water layers, water mass types (Schuback et al., 2017; Zhu et al., 2017; Ryan-Keogh et al., 2018), nutrient conditions (Schuback et al., 2016; Hughes et al.,

2018b) as well as phytoplankton taxa (Robinson et al., 2014; Zhu et al., 2017 and this study), which limits the application of this approach for wide use. Thus, further understanding variability of light- K_C relationships collected using similar approaches from datasets spanning multiple oceanic regions is highly desirable.

We compared several PAR- K_C relationships developed from upper water samples from three different ocean regions, including a semi-closed bay (Ariake Bay, AB, Zhu et al., 2016), marginal sea (East China Sea, ECS, Zhu et al., 2017) and the Arctic Ocean (AO, this study) The highest regression slope value (0.98) was observed for the Arctic Ocean dataset, followed by the East China Sea (0.74), and the lowest value for Ariake Bay (0.51) (**Figure 5A**). ANCOVA analysis revealed that these slopes were only different for AO and AB datasets ($df = 1$, $p = 0.01$), and not for AO and ECS or ECS and AB ($df = 1$, $p = 0.3$). **Supplementary Table 3** summarizes the physical and biochemical parameters associated with these PAR- K_C regression slope values across the three regions. Subsequent correlation analysis of environmental variables vs. K_C -light regression slope values revealed that accounting for phytoplankton size structures yields better correlation with slope value (spearman, $r = 0.98$ for pico-phytoplankton fraction, $p = 0.09$, and $= -0.94$ for micro-phytoplankton fraction, $p = 0.2$) than other environmental parameters (**Table 5**, **Figures 5B,C**). As such, phytoplankton community composition appears to be a key descriptor of the overall phytoplankton physiological status that integrates immediate and historical environmental exposure (Boyd and Abraham, 2001; Koblizek et al., 2001; Cermeño et al., 2005a,b; Moore et al., 2005; Suggett et al., 2009a,b), better than the immediate environmental state, such as nutrient concentration (Behrenfeld et al., 2006; Moore et al., 2008) or phytoplankton species (Geider et al., 1986; Cermeño et al., 2005b) that may in fact be decoupled from the physiological status. We therefore observed that the phytoplankton community composition co-varies well with K_C as well as PAR- K_C regression slopes (**Figures 5B,C**).

Here we show a novel correlated factor of K_C in the complex summer waters of the Arctic Ocean. Our observations suggest that a large fraction of the variation in K_C can be best explained by PAR on a daily timescale. However, we acknowledge that in this study, the limitation of approaches, for example the correction method for σ_{PSII} , the assumption for n_{PSII} (0.002 mol RCII [mol Chla] $^{-1}$), ignoring the variability of ETR-I curve over the daytime and lack consideration of UV effects on primary production would raise uncertainty and weaken our results (Gao et al., 2007; Schuback et al., 2016). We thus applied an uncertainty assessment of the linear relationship between PAR and K_C , as per (Zhu et al., 2017), to examine its reliability. Specifically, $\pm 20\%$ and $\pm 50\%$ random error was added to ETR, respectively, to account for the overall uncertainty caused by the methods limitation. Determination of uncertainty totally repeated 50 times. The resultant K_C was calculated for each new ETR and P_B^C and compared against daily PAR, as per the original data. We then calculated the mean for each correlation coefficient (R^2), slope and intercept from the entire 50 simulations combined, which is $K_C = 1.03 \times \text{PAR} + 6.8$ with $R^2 = 0.40$ and $K_C = 1.2 \times \text{PAR}$

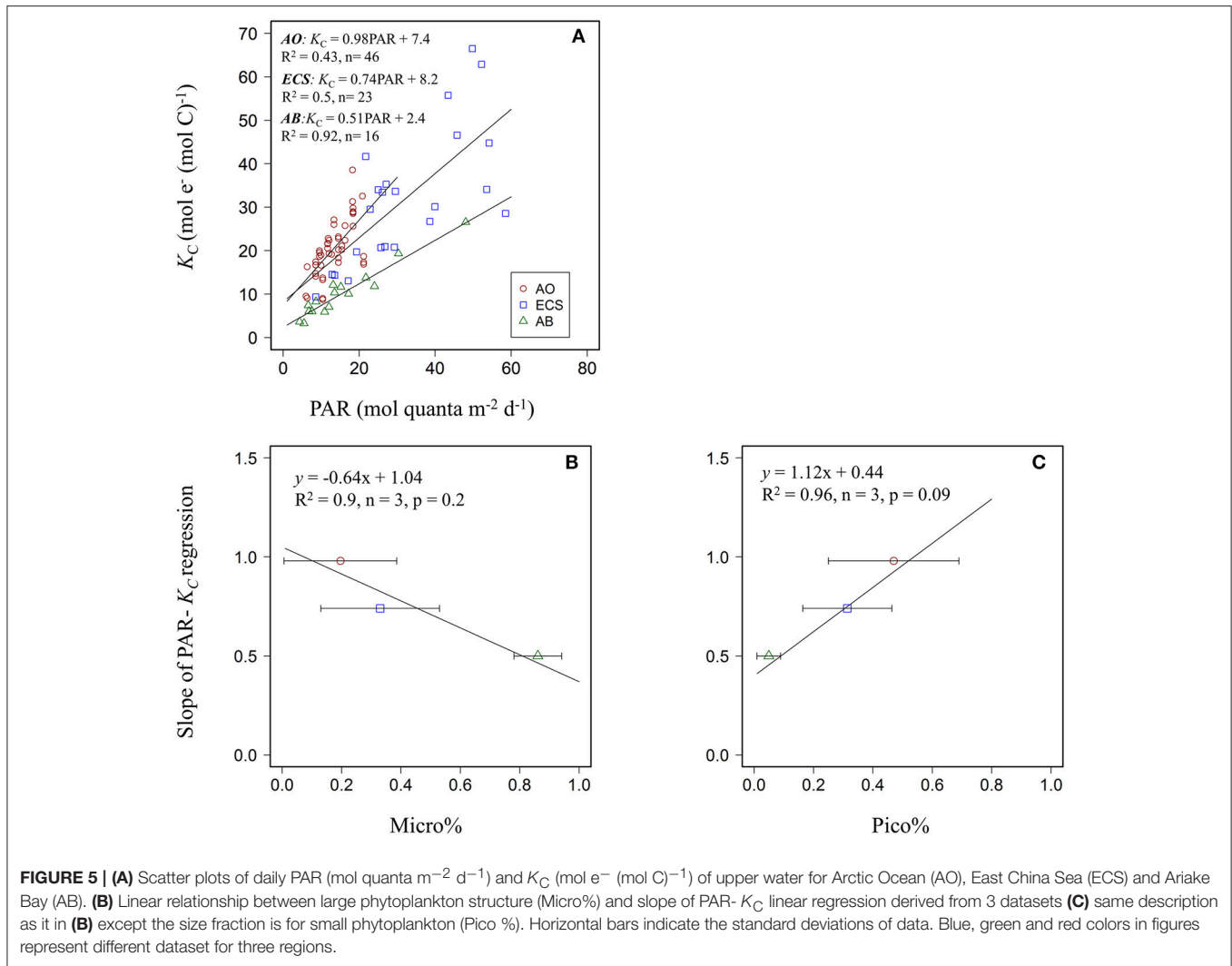


FIGURE 5 | (A) Scatter plots of daily PAR (mol quanta $m^{-2} d^{-1}$) and K_C (mol e^- (mol C) $^{-1}$) of upper water for Arctic Ocean (AO), East China Sea (ECS) and Ariake Bay (AB). **(B)** Linear relationship between large phytoplankton structure (Micro%) and slope of PAR- K_C linear regression derived from 3 datasets **(C)** same description as it in **(B)** except the size fraction is for small phytoplankton (Pico %). Horizontal bars indicate the standard deviations of data. Blue, green and red colors in figures represent different dataset for three regions.

TABLE 5 | Pearson correlation coefficients for correlations between slopes of Par- K_C regression derived from 3 datasets and environmental variables.

	Temp.	Sal.	$NO_3^- + NO_2^-$	PO_4^{3-}	DSi	Chla	Micro%	Nano%	Pico%
Regression slope	-0.7	-0.85	-0.85	0.76	-0.9	-0.91	-0.94	0.82	0.98
n	3	3	3	3	3	3	3	3	3
p	0.5	0.35	0.4	0.45	0.27	0.27	0.2	0.4	0.09

Values in bold indicate the two highest correlation coefficients.

+ 5.8 with $R^2 = 0.38$, for 20% and 50% error, respectively. Compared to the original data ($K_C = 0.98 \times PAR + 7.4$, $R^2 = 0.43$), although absolute terms changed and R^2 decreased with error adding, the extent of covariance explained by a linear model is broadly equivalent, which confirms that the linear relationship between light and K_C observed in the Arctic Ocean is robust.

Further comparing the present data from the Arctic with parallel data sets collected elsewhere, allows us to constrain factors regulating variance of K_C as well as PAR- K_C regressions. Our study thus shows, for the first time, that the variation of

the PAR- K_C relationship under different ocean regimes can be correlated by phytoplankton size structure, which partially overcomes previous FRRf application limitations and represents a step forward for the application of the FRRf technique to derive net primary productivity. Moreover, the cell size (or particle size), is actually a parameter quite amiable to monitoring by optical sensors (i.e., backscattering, which is also a satellite product, as per Dall’Olmo et al., 2009), and our study provides the opportunity for combing FRRf and other techniques to estimate primary productivity across broad regions.

AUTHOR CONTRIBUTIONS

We confirm that all individuals listed as authors have agreed to be listed as such and approve the submitted version of the manuscript. YZ analyzed the FRRf and ^{14}C data, developed the methods of this study and wrote the first draft of this manuscript. DS was responsible for the partial FRRf-ETR calculation and data analyses. He contributed to the interpretation of the results, manuscript writing and revision. CL and FL contributed to the FRRf and primary productivity in field measurements. JH and LL were responsible for nutrient and Chl-a data collection and analyzing. JI and JG specified the structure and the logic of this manuscript and contributed to the manuscript revision. QH developed the research design and interpreted the nutrient and phytoplankton community results. He also contributed to the writing and revisions of the manuscript.

FUNDING

This research was supported by the National Natural Science Foundation of China (Grant No: 41206181),

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ACKNOWLEDGMENTS

We would like to thank the captain, officers and crew of the *R/V Chinare* for their admirable assistance during onboard sampling and measurements. Many thanks to Dr. Yinning Zhang for the FRRf technique and data analysis support.

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2019.00275/full#supplementary-material>

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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