



FEATURE ARTICLE

Interpretation of fast repetition rate (FRR) fluorescence: signatures of phytoplankton community structure versus physiological state

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ABSTRACT: Introduction of active chlorophyll *a* fluorescence protocols, in particular fast repetition rate (FRR) fluorometry, to oceanography and limnology 15 yr ago has enabled rapid assessment of photosynthetic physiology *in situ*. The FRR protocol generates simultaneous measurements of Photosystem II (PSII) effective absorption cross sections (termed σ_{PSII}) and photochemical efficiency (termed F_v/F_m). Both F_v/F_m and σ_{PSII} measurements have been utilised to examine the effects of physiological stress on the photosynthetic apparatus of phytoplankton in an ever growing number of fluorescence-based studies. However, it is now becoming clearer that *in situ* values of F_v/F_m and σ_{PSII} also contain taxonomic information. Here, we present a synthesis of previously unpublished and published data, which show that F_v/F_m and σ_{PSII} vary principally with broad-scale changes in community structure. These patterns observed *in situ* conform to trends observed in laboratory-grown cultures of a range of phytoplankton taxa. The magnitudes of variability in F_v/F_m and σ_{PSII} driven by changes in phytoplankton community structure often exceed that induced by nutrient limitation (as determined from controlled nutrient addition experiments). An exception to this general trend occurs in high-nutrient, low-chlorophyll *a* (HNLC) regions, where strong phenotypic changes in F_v/F_m and σ_{PSII} have been repeatedly demonstrated on relief of iron limitation. Overall, FRR fluorescence measurements of both F_v/F_m and σ_{PSII} in natural populations represent a combination of the taxonomic 'signature' (values of F_v/F_m and σ_{PSII} determined by the taxa present) within the phytoplankton community that is further modified according to the (photo-) physiological status. As such, fluorescence-based investigations of mixed populations must account for potential variations in phytoplankton community structure before interpretations of physiological status are made.



Fast Repetition Rate fluorometer (at left) attached to a CTD rosette sampler being deployed in the Celtic Sea.

Photo: Young-Nam Kim

KEY WORDS: Fast repetition rate · Fluorescence · PSII · Effective absorption cross section · Photochemical efficiency · Phytoplankton · Nutrient stress

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INTRODUCTION

An important milestone for aquatic research was the introduction of active chlorophyll *a* fluorometry to oceanography and limnology 15 yr ago (Falkowski & Raven 2007, Prášil et al. 2008). Technological and commercial development has since packaged various active fluorescence protocols into a number of plat-

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forms, from submersible profilers to bench-top imagers to aircraft remote-sensing light detection and ranging (LIDAR) platforms enabling its application across almost every field, from coral biology to ocean biogeochemistry (Prášil et al. 2008). As a direct result, active fluorometry has rapidly become an established tool by which scientists and managers evaluate the response of aquatic systems to environmental change.

Initial *in situ* studies immediately demonstrated the capabilities of active fluorescence for rapidly assessing the photosynthetic physiology of phytoplankton as they existed in nature (Kolber et al. 1990, Geider et al. 1993a, Greene et al. 1994). Conceptually, active chlorophyll *a* fluorescence measurements evaluate the efficiency by which absorbed light is utilised by photosynthesis: chlorophyll *a* fluorescence itself being a biophysical byproduct reflecting absorbed light not utilised for photochemistry by the oxygen evolving Photosystem II complex (PSII). Such measurements are strongly dependent on prior light exposure, and thus the PSII photochemical efficiency is at its highest (maximum) following a period of dark acclimation, i.e. when the primary acceptor molecules of PSII (quinones) are fully oxidised and ready to receive electrons. This maximum PSII photochemical efficiency typically decreases when autotrophs experience 'stressful' growth conditions, such as nutrient starvation or excessive light (Kolber et al. 1988, 1994, Geider et al. 1993b). It is this concept that has led to use of chlorophyll *a* fluorescence measurements for ascertaining the photosynthetic 'condition' of entire phytoplankton communities across large temporal (Suggett et al. 2006a) and spatial (Behrenfeld et al. 1996, Suzuki et al. 2002, Moore et al. 2005, 2006a,b, Suggett et al. 2006b) scales approaching those of oceanic ecosystems (Behrenfeld & Kolber 1999, Behrenfeld et al. 2006, Suggett et al. 2006b).

Of key interest to oceanographers has been a specific form of active fluorometry termed fast repetition rate (FRR; Kolber et al. 1998) and analogous techniques, including fluorescence induction and relaxation (FIRE; Bibby et al. 2008, Suggett et al. 2008). These fluorometers enable the high fluorescence sensitivity required for low chlorophyll *a* oligotrophic waters. Moreover these protocols result in the majority of PSII reaction centres being closed within the timescale of a single photochemical turnover (ST) (Kolber et al. 1998), enabling simultaneous measurement of the maximum PSII effective absorption cross section, termed σ_{PSII} , and the maximum PSII photochemical efficiency, termed F_v/F_m . The PSII effective absorption, also referred to as the functional absorption cross section, describes the functional 'target area' of the light-harvesting antenna that is energetically coupled to the O_2 -evolving reaction centres (RCIIs) (σ_{PSII} ; Mauzerall & Greenbaum 1989, Falkowski & Raven 2007).

Achieving simultaneous measurements of F_v/F_m and σ_{PSII} has proved to be an important step in active fluorometry by enabling autonomous estimates of the photosynthetic electron transfer (PET) rate (Kolber & Falkowski 1993). Under actinic light, both F_v/F_m and σ_{PSII} are modified according to how the absorbed light is utilised by PSII. Until the development of FRR, active fluorometry typically only measured F_v/F_m and so had to rely on independent measurements of light absorption to calculate PET rates in absolute terms (Kromkamp & Forster 2003, Johnsen & Sakshaug 2007). The measurements of σ_{PSII} also proved to be an additional advancement for active fluorometry, since this variable also appears to characterise an (eco-) physiological response to changes in environmental conditions (Greene et al. 1994, Kolber et al. 1994, Behrenfeld et al. 1996, Behrenfeld & Kolber 1999, Sosik & Olson 2002, Moore et al. 2006a,b, 2007, Suggett et al. 2006b, Six et al. 2008).

Substantial FRR fluorescence data sets now exist for phytoplankton from both laboratory-grown monocultures and natural communities. These data indicate that the biophysical characteristics of PSII, expressed as F_v/F_m and σ_{PSII} , contain a taxonomic 'signature', i.e. a unique identity, upon which is superimposed physiological variability (Suggett et al. 2004, Moore et al. 2005, 2006b, 2008, Fishwick et al. 2006). As such, FRR fluorescence measurements from natural communities cannot be interpreted on the basis of physiological stress alone (Prášil et al. 2008). Such a realisation is perhaps not surprising since light absorption is one of the key factors operated on by selective pressure during the evolution of phytoplankton taxa (Wood 1985, Glover et al. 1987, Stomp et al. 2007). As a result, the compliment and arrangement of pigments housed within the light-harvesting antennae are highly variable amongst phytoplankton groups (Lutz et al. 2001, MacIntyre et al. 2002, Johnsen & Sakshaug 2007). Such fundamental differences in antenna structure between phytoplankton groups are expected to result in mechanistic differences in both PSII light-harvesting potential and the efficiency with which this absorbed light is utilised (see Laverigne & Joliot 2000, Lutz et al. 2001).

Here, we present a synthesis of unpublished and previously published FRR fluorescence data collected from numerous laboratory-grown marine phytoplankton to demonstrate that FRR fluorescence measurements represent (eco-) physiological variability influenced by both phytoplankton community structure and physiological 'stress'. We also examine both unpublished and published FRR fluorescence data sets from (1) broad-scale environmental surveys and (2) nutrient-addition bioassay experiments to demonstrate conditions where and when FRR fluorescence measurements include important taxon-specific ecophysio-

logical traits of *in situ* phytoplankton populations. Consequently, we illustrate the relative magnitude of variability in F_v/F_m and σ_{PSII} within and between phytoplankton taxa, highlighting that care must be taken when using FRR fluorescence to assess physiological stress (principally nutrient stress) of natural phytoplankton populations. With these findings in mind, we briefly discuss current directions for FRR fluorometry.

MATERIALS AND METHODS

Common methodologies apply to both the unpublished and previously published data sets analysed here. We will briefly describe these methodologies; however, readers are referred to the original sources for full details.

FRR fluorescence. All FRR measurements were made using ‘standard’ protocols as described by Suggett et al. (2003, 2004) and Moore et al. (2005, 2006b). A Chelsea Instruments Fast^{track} FRR fluorometer was programmed to deliver ST saturation of PSII from 100 flashlets of 1.1 μs at 1 μs intervals. Induction curves were averaged over 160 to 320 individual sequences to minimize error. Fluorescence transients generated by the FRR were then fitted to the Kolber-Prasil-Falkowski (KPF; Kolber et al. 1998) model to yield values of F_v/F_m and σ_{PSII} (see Moore et al. 2006b). Importantly, the measured value of σ_{PSII} will be weighted to the spectral quality of light used to induce the fluorescence transient. The Fast^{track} has LEDs with peak excitation at 478 nm and a relatively narrow band width (25 nm); therefore, all values of σ_{PSII} reported by this method are weighted to the blue region of the photosynthetically active radiation (PAR) spectrum and should strictly be quoted as σ_{PSII} (~478 nm) (Moore et al. 2006b). Connectivity (ρ) was not fixed at the time of fitting the KPF model (see Suggett et al. 2004). However, values for ρ were always between ca. 0.2 and 0.4 (dimensionless), and extensive tests with subsets of the data indicated no sensitivity of our conclusions to assumptions about connectivity (not shown). Instrument non-linearity was corrected for at the time of curve fitting from fluorescence transients of chlorophyll *a* extracts following Laney & Letelier (2008). Blanks were also collected from fluorescence induction recordings of filtrates for each sample as described in detail by Moore et al. (2006b) and Suggett et al. (2006a).

Additional measurements. Additional information was required to interpret either field and/or laboratory FRR data. Methods for obtaining the PSII photosynthetic unit size (mol chl *a*:mol RCII), pigment concentrations, and excitation spectra for PSII fluorescence have been described previously (Suggett et al. 2004,

Moore et al. 2006b). Briefly, PSII photosynthetic unit sizes were determined from O_2 -flash yield measurements using a custom-built LED flash system (Suggett et al. 2003). Pigment concentrations were measured by high-performance liquid chromatography (HPLC) via identification and co-elution with pigment standards (see Leonardos & Geider 2004, Moore et al. 2006b). Finally, PSII fluorescence excitation spectra (400 to 715 nm) were measured with the monochromator on the detector set to 730 nm emission (F_{730}) upon samples pre-treated with DCMU (Lutz et al. 2001, Suggett et al. 2004). Values for F_{730} were normalized to values for σ_{PSII} at 478 nm, since this wavelength corresponded with the wavelength of peak excitation from the FRR fluorometer. As such, excitation spectra for PSII can be used to extrapolate measurements of the effective absorption measured by the FRR (σ_{PSII} , 487 nm) to other wavelengths (see Suggett et al. 2007).

Data sets 1: algal cultures. The phytoplankton data set includes 38 species spanning 10 classes (Table 1). Most species were grown at either the University of Essex (UK) or Horn Point Laboratory (Maryland, USA), data is also included from algal cultures grown at Bar-Ilan University (Israel, Ilana Berman-Frank) and the Finnish Institute of Marine Research (Mika Raateoja) (Table 1). Cultures were grown under 1 or more growth photon flux density (PFD) and were maintained in batch or semi-batch mode in nutrient replete media, as described previously (Raateoja et al. 2004, Suggett et al. 2004, Berman-Frank et al. 2007). Growth was performed under optimum temperatures and salinity, except where stated (Table 1). Cultures were maintained in exponential growth phase, as monitored using daily FRR fluorescence or haemocytometer cell counts, and sampled in triplicate from sequential cell generations.

In addition to our culture data, we have also included published FRR data for F_v/F_m and σ_{PSII} digitised from *Prochlorococcus* experiments of Steglich et al. (2001) and Bruyant et al. (2005) (Table 1). The digitised FRR measurements do not appear to account for cases of instrument non-linearity or sample blanks; however, the latter are probably unimportant where chlorophyll *a* concentrations exceed 1 to 10 mg m^{-3} (Suggett et al. 2006a), i.e. concentrations substantially below those for the *Prochlorococcus* cultures.

Data sets 2: field sampling. Primary field data sources were from selected cruises spanning a range of hydrographic regimes including seasonally stratified shelf sea (JR98; Moore et al. 2006b), Atlantic open ocean (AMT 15 and 17; Hickman & Suggett unpubl. data; Meteor 60, Moore et al. 2006a, 2008) and southern Indian ocean (CROZEX; Moore et al. 2007). The field data thus traversed large environmental and taxonomic gradients and yet are known to have consistently applied the same FRR techniques and data analysis throughout.

Table 1. Summary information for cultured marine phytoplankton examined using fast repetition rate (FRR) fluorescence. Culture reference banks are CCMP (Provasoli-Guillard National Center for Culture of Marine Phytoplankton Boothbay Harbour, Maine, USA), CCAP (Culture Collection of Algae and Protozoa, Dunstaffnage Marine Laboratory, Scotland, UK), PCC (Plymouth Culture Collection of Marine Algae, Marine Biological Association, Plymouth, UK), Choptank Isolates (Choptank River isolated by Alan Lewitus and maintained at Horn Point Laboratory, MD, USA) and Woods Hole (WH) isolate (Woods Hole Oceanographic Institute, MA, USA, by J. Waterbury). Cyanobacterial strains KAC and HEM are Baltic Sea isolates. All algae were grown under steady-state, nutrient-replete conditions as described by Suggett et al. (2003, 2004). All species were grown under 1 or more irradiances and at optimum growth temperatures (15 to 20°C) and salinity (30 to 35 PSU), except where strains were grown at 5°C and 15 to 25 PSU (*). All cultures were maintained in media at an N:P of 15 to 16, except where strains were grown at an N:P of 5, 15 or 45 (**). Steady-state FRR data provided by I. Berman-Frank for *Trichodesmium* spp., *Cyanothece* spp. and *Anabaena* spp. were from various iron concentrations, temperatures and salinities (refer to Berman-Frank et al. 2007)

Algal class	Species	Culture ID	Methods for cell growth and/or FRR data source
Diatom	<i>Chaetoceros muelleri</i>	CCAP1010/3	Leonardos & Geider (2004)**
	<i>Coscinodiscus wallensi</i>	PCC621	Estevez-Blanco & Suggett unpubl. data
	<i>Skeletonema costatum</i>	CCMP1332	Le Floc'H & Suggett unpubl. data*
	<i>Phaeodactylum cornutum</i>	CCMP 1327	Estevez-Blanco & Suggett unpubl. data
	<i>Thalassiosira weissflogii</i>	CCMP1047	Suggett et al. (2003, 2004)
	<i>Thalassiosira weissflogii</i>	CCMP1336	Estevez-Blanco & Suggett unpubl. data
	<i>Thalassiosira pseudonana</i>	CCMP1335	Estevez-Blanco & Suggett unpubl. data
	<i>Thalassiosira punctigera</i>	PCC623	Estevez-Blanco & Suggett unpubl. data
	<i>Thalassiosira minima</i>	CCAP1085/8	Estevez-Blanco & Suggett unpubl. data
	<i>Thalassiosira eccentrica</i>	CCAP1085/6	Estevez-Blanco & Suggett unpubl. data
Dinoflagellate	<i>Prorocentrum minimum</i>	CCAP1136/14	Le Floc'H & Suggett unpubl. data*
	<i>Prorocentrum minimum</i>	Choptank	Suggett et al. (2004)
	<i>Symbiodinium microadriaticum</i>	CCMP1633	Hennige & Suggett unpubl. data
Prymnesiophyte	<i>Coccolithus pelagicus</i>	PCC182	Estevez-Blanco & Suggett unpubl. data
	<i>Emiliania huxleyi</i>	PCC/B92	Suggett et al. (2007)
	<i>Emiliania huxleyi</i>	PCC/B11	Suggett et al. (2007)
	<i>Isochrysis galbana</i>	CCMP1323	Estevez-Blanco & Suggett unpubl. data
	<i>Prymnesium parvum</i>	PCC94	Estevez-Blanco & Suggett unpubl. data
Chlorophyte	<i>Dunaliella tertiolecta</i>	CCAP 19/6B	Suggett et al. (2003)
	<i>Dunaliella tertiolecta</i>	CCMP 1320	Suggett et al. (2003, 2004)
Cryptophyte	<i>Nannochloris atomus</i>	CCMP509	Suggett et al. (2003)
	<i>Rhodomonas salina</i>	CCAP 978/27	Le Floc'H & Suggett unpubl. data*
	<i>Rhodomonas salina</i>	CCMP1319	Estevez-Blanco & Suggett unpubl. data
Cyanophyte	<i>Storeatula major</i>	Choptank	Suggett et al. (2004)
	<i>Aphanizomenon</i> sp.	KAC15	Raateoja et al. (2004)
	<i>Nodularia spumigena</i>	HEM	Raateoja et al. (2004)
	<i>Synechococcus</i> sp.	CCMP837	Suggett unpubl. data*
	<i>Synechococcus</i> sp.	WH 7803	Suggett et al. (2004)
	<i>Trichodesmium</i> sp.	IMS101	Berman-Frank et al. (2007)
	<i>Katagnymene</i> sp.	WH isolate	Berman-Frank unpubl. data
	<i>Cyanothece</i> sp.	WH 8904	Berman-Frank et al. (2007)
Prochlorophyte	<i>Anabaena flos aqua</i>	UTEX 2557	Berman-Frank et al. (2007)
	<i>Prochlorococcus marinus</i>	PCC 9511	Bruyant et al (2005)
	<i>Prochlorococcus marinus</i>	CCMP 1375	Steglich et al. (2001)
Pelagophyte	<i>Aureococcus anophagefferens</i>	CCMP1790	Suggett et al. (2004)
	<i>Pelagococcus subviridis</i>	CCMP1429	Estevez-Blanco & Suggett unpubl. data
Prasinophyte	<i>Pycnococcus provasolii</i>	CCMP 1203	Suggett et al. (2004)
Rhodophyte	<i>Porphyridium cruentum</i>	Choptank	Suggett et al. (2004)

Full details of all cruise data synthesised are provided in Table 2. FRR fluorescence measurements were performed on discrete water samples drawn from Niskin bottles at various depths within the upper euphotic zone or from the ships' underway surface seawater supply. Water samples were collected pre-dawn (AMT 15, 17, JR98, Meteor 60) or were dark adapted for 45 to 60 min to ensure full dark adaptation. Nutrient perturbation experiments were carried out during CROZEX,

Meteor 60 and AMT 17. FRR protocols for AMT 15 and 17 followed those of Moore et al. (2006b), whilst nutrient perturbation protocols for AMT 17 followed those of Moore et al. (2006a, 2008).

Several additional field data sets were digitised from published sources for comparative purposes (Table 2). Values were extracted from continuous data sets corresponding to night-time or a PAR of zero to minimize the influence of reversible fluorescence quenching.

Table 2. Summary of re-examined field data sets where FRR fluorescence was used to yield values of PSII photochemical efficiency (F_v/F_m , dimensionless) and effective absorption cross section (σ_{PSII} , $\text{\AA}^2 \text{ quanta}^{-1}$). Previously unpublished data provided by Ondrej Prášil from nighttime vertical FRR casts was binned into 10 depths in the upper 200 m. Similarly, data from Anna Hickman and David Suggett were from dark-adapted measurements upon discrete water samples collected pre-dawn using Niskin bottles (see Suggett et al. 2006b)

Cruise	Geographical region	Location	Comments	Data set
IronEx II	Eastern Equatorial Pacific	3.5° S, 105° W	Surface water only	Behrenfeld et al. (1996)
SOIREE	Indian sector; south of the Antarctic Polar Front (APF)	61° S, 141° E	Depth integrated; binned to F_v/F_m intervals	Boyd & Abraham (2001)
RR09	Polar Front–Ross Sea (N to S)	50–78° S, 165° E–165° W	Mean in upper 40 m	Sosik & Olson (2002)
KH-99-3	Trans-Pacific (E to W, sub-Arctic)	45–55° N, 155° E–135° W	Surface water only	Suzuki et al. (2002)
	Mertz Polynya region of East Antarctica	64–67° S, 140–150° E	10–20 m and 45–55 m depth bins	Vaillancourt et al. (2003)
FISHES	N. Atlantic (Iceland Basin–Rockall Trough)	56–66° N, 0–30° W	Mean value in surface mixed layer	Moore et al. (2005)
	Ría de Vigo, NW Spain	42° 20' N, 8° 70' W	Two depths; seasonal	Cermeño et al. (2005)
	SW coastal Africa (Cape Town to Orange River)	33–29° S, 15–18° W	Depth of maximum in the enclosed chamber	Fishwick et al. (2006)
JR98	N. Atlantic (UK–Ireland shelf sea)	48–53° N, 5–10° W	Two depths, surface and thermocline	Moore et al. (2006b)
CROZEX	S. Atlantic (Polar-Frontal zone of Southern Ocean)	44–49° S, 49–56° E	Two depths, surface and thermocline	Moore et al. (2007)
Meteor 60	Trans-Atlantic (W to E, subtropical)	18–36° N, 66–24° W	Surface water only	Moore et al. (2008)
AMT 15	Trans-Atlantic (N to S)	48° N–40° S, 25° W–10° E	10 depths, upper 200 m	Hickman unpubl. data
BIOCOPE	Subtropical Pacific	31–34° S, 91–74° W	10 depths, upper 200 m	Prášil unpubl. data
AMT 17	Trans-Atlantic (N to S)	48° N–35° S, 38° W–13° E	10 depths, upper 200 m	Suggett unpubl. data

Again, it should be noted that these FRR data sets do not always appear to account for cases of non-linearity or sample blanks. The majority of these data sets employed a Chelsea Instruments FRR Fast^{tracka}, with excitation set to 478 nm; consequently, σ_{PSII} values reported here also strictly correspond to this wavelength, i.e. σ_{PSII} (478 nm). However, we also include data collected using an earlier ‘prototype’ variable fluorometer with an alternative excitation source (Kolber et al. 1988, 1994, Behrenfeld et al. 1996). Specifically, in this case, cut-off filters were used to isolate excitation by a xenon lamp between ca. 490 and 550 nm (Kolber & Falkowski 1993), and so peak excitation is broader and slightly more shifted to ‘green’ than that employed by the Fast^{tracka}. We consider how these alternative wavebands of excitation potentially affect the interpretation of σ_{PSII} and F_v/F_m values in response to changes in taxa and growth environment in later sections. Hereafter, when considering direct measurements with a 478 nm excitation source, we use the notation σ_{PSII} (478) to indicate the corresponding functional absorption cross section (Moore et al. 2006b). In contrast, the more general discussions of the ecological/evolutionary context behind functional antenna size will contain the implicit assumption that phytoplankton absorption must be considered in the context of the wavelength regime of interest (i.e. the *in situ* spectral growth irradiance).

RESULTS AND DISCUSSION

FRR fluorescence under unbalanced versus balanced growth

In order to examine taxonomic effects we will first consider how environment, in terms of growth rate, determines FRR-based measurements. Microalgae grown under batch and continuous culture approaches demonstrate the effect of unbalanced versus balanced growth upon PSII fluorescence characteristics (e.g. Fig. 1). Typically, in batch cultures, cells exhibit a constant value of F_v/F_m and σ_{PSII} (478) during logarithmic growth. However, a gradual decrease of F_v/F_m and increase of σ_{PSII} (478) is observed once nutrient availability decreases relative to cellular demand for photosynthesis and so the growth rate inevitably declines (summarized by Parkhill et al. 2001). Growth rates and cellular physiology are thus not in balance with the nutrient supply. Inverse relations between F_v/F_m and σ_{PSII} are observed when PSII function becomes compromised by nutrient starvation (Kolber et al. 1988, Geider et al. 1993b, Steglich et al. 2001) or photoinhibition by visible or UV radiation (Vassiliev et al. 1995, Ragni et al. 2008). Such a decline of F_v/F_m in response to stress can be interpreted to indicate a decline in the proportion of RCIIIs that are functional: when the pool of functional and damaged RCIIIs shares a common

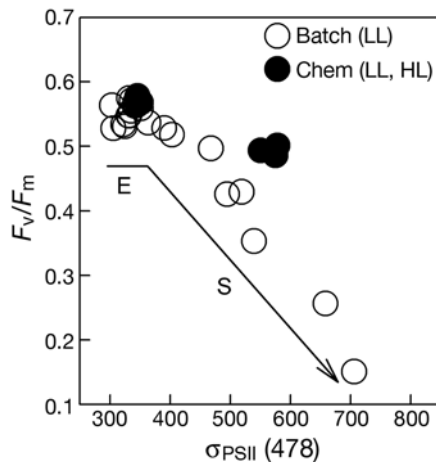


Fig. 1. Fast repetition rate (FRR) measurements of the maximum Photosystem II (PSII) photochemical efficiency (F_v/F_m , dimensionless) and effective absorption cross section (σ_{PSII} (478), $\text{\AA}^2 \text{ quanta}^{-1}$) for the diatom *Chaetoceros muelleri* when grown in either batch- or chemostat-culture modes (see Table 1). In batch mode (O) nutrients were not replenished post-inoculation and values of F_v/F_m and σ_{PSII} were recorded for cells growing at a single light intensity ($50 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$) through exponential (E) and stationary (S) phases. Each data point represents measurements made on consecutive days; E and S phases were identified by simultaneous daily quantification of cell number (Leonardos & Geider 2004). Values of F_v/F_m and σ_{PSII} shown are the means from triplicate cultures. In chemostat mode (●) nutrients were replenished continuously and values of F_v/F_m and σ_{PSII} were also recorded daily. Cultures were maintained under low-light (LL, $50 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$, growth rate [μ] = 0.3 d^{-1}) and high-light (HL, $700 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$, μ = 0.6 d^{-1}) conditions. Both LL and HL data clusters consist of 3 data points, each representing a chemostat delivered with a different N:P within the growth media (5, 15, 45 mol:mol; Table 1). Here, values of F_v/F_m and σ_{PSII} are the means from 15 to 22 consecutive days in steady-state growth

light-harvesting antenna, a decrease in the proportion of RCIIIs that are functional (i.e. a decrease in F_v/F_m) is accompanied by an increase in the effective cross section, σ_{PSII} (Greene et al. 1992, Vassiliev et al. 1995).

Cells in stationary-phase batch cultures, in which there is an abrupt exhaustion of nutrients, experience nutrient starvation. As such, batch cultures may provide a good model for the termination of a bloom by nutrient limitation in nature. However, under conditions of chronic nutrient limitation, regeneration of nutrients by heterotrophs can provide a continuous supply of nutrients. In the subtropical gyres, for example, there may be a high turnover of very small pools of dissolved nutrients due to a near balance between uptake of nutrients by phytoplankton and regeneration of nutrients by heterotrophs. In contrast to the batch approach, chemostat cultures continuously supply nutrients and the growth rate is strictly determined by the dilution rate, i.e. the rate of nutrient supply. Some chemostat studies have observed decreased val-

ues of F_v/F_m and increased values of σ_{PSII} as the growth rate is lowered (Kolber et al. 1988). However, such a trend is not the rule (Cullen et al. 1992, Parkhill et al. 2001). A continuous supply of nutrients eventually appears to enable cells to acclimate to chronic nutrient limitation. As a result, acclimation of the photosynthetic apparatus may alleviate or reduce the effects of nutrient limitation on F_v/F_m and σ_{PSII} . For example, although measured using an alternative active fluorescence technique, continuous culture studies have demonstrated little sensitivity of F_v/F_m to the nutrient-limited, steady-state growth rate (Parkhill et al. 2001, Price 2005).

Insensitivity of F_v/F_m and σ_{PSII} (478) to nutrient-limited growth conditions is also observed in FRR data collected during chemostat experiments of the diatom *Chaetoceros muelleri* (Fig. 1). Here, high light grown cells were allowed to grow at a higher dilution (growth) rate than were low light grown cells (Leonardos & Geider 2004). Lower values of F_v/F_m were not observed at the lower growth rates, but for cells grown under higher growth PFDs (at the higher growth rate). Therefore, these data indicate that the change in light availability outweighed any potential influence of nutrient limitation upon PSII functioning. In this experiment, cultures were also grown over a range of N:P concentrations for these same high-light, high-growth rate and low-light, low-growth rate conditions. However, values of F_v/F_m and σ_{PSII} (478) did not vary between N:P treatments (Fig. 1, see legend). In other words, FRR fluorescence measurements were also insensitive to adjustments to the nutrient ratio supplied (N:P) as long as the dilution rate was held constant. Thus, it appears that the overall functional capacity of the fully acclimated PSII pool is insensitive to the N:P supply ratio, despite the latter controlling the biomass that can be achieved (Geider et al. 1993b).

Taxonomic-dependence of F_v/F_m and σ_{PSII}

Under nutrient-replete growth conditions values of F_v/F_m are known to vary systematically amongst algal taxa (Koblížek et al. 2001, Suggett et al. 2003, Yentsch et al. 2004). The highest recorded FRR-based ST values of F_v/F_m are between ca. 0.65 and 0.70 and were measured for chlorophytes and diatoms (e.g. *Chlamydomonas reinhardtii* and *Scenedesmus quadricauda* [Koblížek et al. 2001] and *Thalassiosira weissflogii* [Fig. 2A, see also Yentsch et al. 2004]). However, values for F_v/F_m can be as low as 0.3 to 0.4 for the smallest (pico-) eukaryotes, such as the pelagophyte *Aureococcus anophagefferens*, the prasinophyte *Pycnococcus provasolii*, and the rhodophyte *Porphyridium cruentum* (Fig. 2A). Values of F_v/F_m for cyanobacteria are typi-

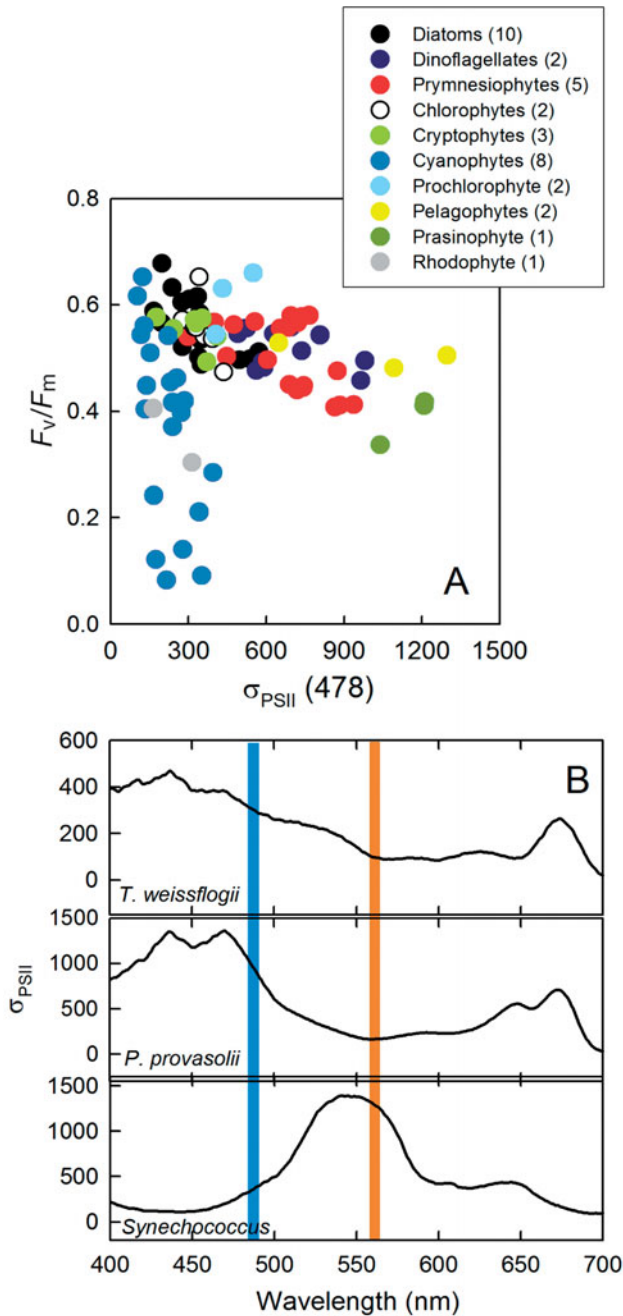


Fig. 2. (A) Values of the maximum PSII photochemical efficiency (F_v/F_m , dimensionless) and effective absorption cross section (σ_{PSII} (478), $\text{\AA}^2 \text{ quanta}^{-1}$) measured from FRR fluorometry for various species of prokaryotic and eukaryotic marine phytoplankton (see Table 1 for a full list). Regressions for the phycobilisome and chlorophyll-xanthophyll light-harvesting complex (LHC) 'trajectories' are $F_v/F_m = -0.0011 \times \sigma_{PSII}$ (478) + 0.621 ($r^2 = 0.322$, $n = 23$) and $F_v/F_m = -0.0002 \times \sigma_{PSII}$ (478) + 0.618 ($r^2 = 0.405$, $n = 79$), respectively. Species data have been pooled into phytoplankton classes indicated by different symbols. (B) Examples of σ_{PSII} -excitation spectra recorded for a diatom (*Thalassiosira weissflogii* CCMP1047), prasinophyte (*Pycnococcus provasolii* CCMP 1203) and cyanophyte (*Synechococcus* sp. WH7803). Solid lines indicate values of σ_{PSII} expected when exciting PSII with blue light (478 nm) and orange light (560 nm)

cally between 0.1 and 0.4 (Koblížek et al. 2001, see also Campbell et al. 1998), but values in excess of 0.60 to 0.65 have been measured for some species of nitrogen-fixing cyanobacteria (e.g. in the genera *Cyanothea* and *Anabaena*; Berman-Frank et al. 2007; see Fig. 2). Thus, despite early suggestions (Kolber et al. 1988, Geider et al. 1993a, Kolber & Falkowski 1993), a common maximum value for F_v/F_m cannot be assigned to all algae (see discussion by Suggett et al. 2004).

As with F_v/F_m , FRR-based values of σ_{PSII} appear to vary across algal taxa (Koblížek et al. 2001, Suggett et al. 2004). In fact, plotting corresponding values of F_v/F_m and σ_{PSII} (478) reveals that the 2 variables negatively co-vary across a wide range of algal classes (Fig. 2A). These data sets include species that have been grown at >1 PFD and in both chemostat and batch culture modes (Table 1). We have already shown that F_v/F_m and σ_{PSII} (478) can be modified in a manner similar to that in Fig. 2A when *Chaetoceros muelleri* was grown under 2 PFDs (Fig. 1); therefore, some of this co-variability may arguably be the result of photo-acclimation. However, the over-riding trend across the range of values for both F_v/F_m and σ_{PSII} (478) from the cultures is clearly driven by taxonomic clusters of data (Fig. 2A). Few data sets are available to investigate whether this observation is also maintained within an algal class. However, our data for diatoms under nutrient-replete growth indicated a significant trend between F_v/F_m and σ_{PSII} (478) (10 species, $r^2 = 0.426$) for the substantial range of species tested (Fig. 3A). Recent studies have also demonstrated that some degree of co-variability between F_v/F_m and σ_{PSII} (478) extends to the intra-clonal level of a single eukaryotic species (e.g. *Emilinia huxleyi* and *Symbiodinium* sp.; Suggett et al. 2007, 2008). However, the degree of variability is considerably smaller than that observed between species within an algal class (Fig. 3A), which is, in turn, small when compared to differences between eukaryotic groups (Fig. 2).

For the data examined here, the algae appear to group along 2 distinct 'trajectories'. The slope (or 'trajectory') of the F_v/F_m versus σ_{PSII} (478) relationship is much steeper for cyanobacteria than it is for eukaryotes (Fig. 2A). The eukaryotic algae cluster (from high- F_v/F_m , low- σ_{PSII} (478) to low- F_v/F_m , high- σ_{PSII} (478)) in the following general order: diatoms, chlorophytes and cryptophytes, prymnesiophytes and dinoflagellates, nanoflagellates (pelagophytes and prasinophytes) along a common trajectory. For the taxa measured, this ordering broadly corresponds to a decreasing trend in cell size. A similar relationship with cell size also appears within the diatom class (Fig. 3B).

The notable exceptions are for the prokaryotic prochlorophytes and the eukaryote rhodophyte *Porphyridium cruentum*, for which phenotypic co-variability

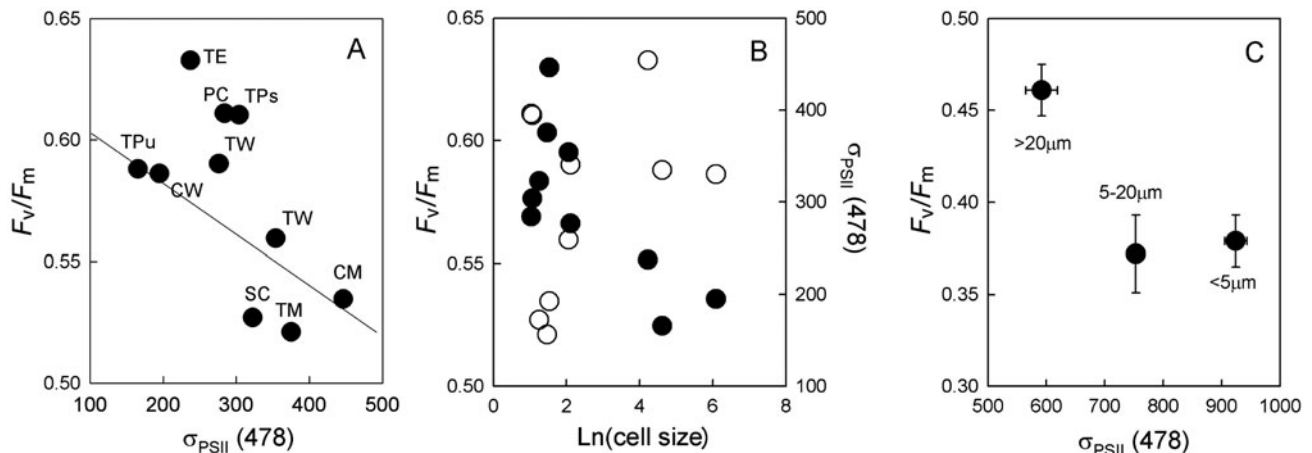


Fig. 3. Comparison of: (A) maximum PSII photochemical efficiency (F_v/F_m , dimensionless) versus effective absorption cross section ($\sigma_{PSII} (478)$, $\text{\AA}^2 \text{ quanta}^{-1}$) for all diatom species: CM (*Chaetoceros muelleri*), CW (*Coscinodiscus wallensi*), SC (*Skeletonema costatum*), PC (*Phaeodactylum cornutum*), TE (*Thalassiosira eccentrica*), TM (*Thalassiosira minima*), TP_s (*Thalassiosira pseudonana*), TP_u (*Thalassiosira punctigera*), TW (*Thalassiosira weissflogii*) (see Table 1). Note that each data point represents the mean value for all growth environments (where multiple data sets were available) to attempt to remove potential additional influences of growth light environment upon the relationship. Regression is $F_v/F_m = -0.0003 \times \sigma_{PSII} (478) + 0.665$ ($r^2 = 0.426$, $n = 10$). The solid line indicates the chlorophyll-xanthophyll LHC 'trajectory' determined for regressions applied to laboratory culture data from Fig. 2. (B) Comparison of F_v/F_m (O) and $\sigma_{PSII} (478)$ (●) versus $\ln(\text{cell size}, \mu m)$. Only the relationship between $\sigma_{PSII} (478)$ and cell size was significant: $\sigma_{PSII} (478) = -0.360 \times \ln(\text{cell size}) + 319$ ($r^2 = 0.372$, $n = 10$). (C) F_v/F_m versus $\sigma_{PSII} (478)$ measured for 3 size classes isolated from natural phytoplankton communities from the Ría de Vigo, Spain (Cermeño et al. 2005) (Table 2). Data represent the means (\pm SE) of all size-fractionation samples throughout the experimental period. Additional data to those already published were supplied by Patricia Estevez-Blanco and Pedro Cermeño

in F_v/F_m and σ_{PSII} followed a trajectory similar to that for the eukaryotes and cyanobacteria, respectively. Rhodophytes and cyanobacteria are related by the characteristic that both employ phycobilisomes as light-harvesting antennae for PSII. Relatively low values of both F_v/F_m and $\sigma_{PSII} (478)$ have been documented previously for cyanobacteria, particularly those containing substantial concentrations of phycocyanin (Campbell et al. 1998, Raateoja et al. 2004). The fluorescence emission band from phycocyanin overlaps with that of chlorophyll *a* from PSII and, hence, can 'contaminate' the minimum fluorescence yield and so lead to lower values of F_v/F_m (Campbell et al. 1998). This same feature likely explains the presence of the rhodophyte *P. cruentum* sp. within the 'cyanobacteria' rather than the 'eukaryotic' trajectory. However, some cyanobacteria with relatively low concentrations of phycocyanin within their phycobilipigment complement, e.g. *Synechococcus* sp. WH7803, still have relatively low values of both F_v/F_m and $\sigma_{PSII} (478)$. A substantial proportion of chlorophyll *a* within cyanobacteria is associated with PSI; therefore, it is possible that additional 'contamination' of the minimum yield (F_0) may originate from PSI chlorophyll *a* fluorescence (Campbell et al. 1998).

Low values of $\sigma_{PSII} (478)$ for cyanobacteria almost certainly result from the blue excitation waveband employed by the Fast^{track} FRR fluorometer, which is an inefficient excitation source for these organisms since

PSII absorption is dominated by phycobilisomes (Raateoja et al. 2004, Suggett et al. 2004, Johnsen & Sakshaug 2007). Fluorescence excitation spectra demonstrate that the PSII action spectrum for cyanobacteria is typically much lower in blue than in orange light (Fig. 2B); for example, *Synechococcus* sp. WH7803 exhibits a value of $\sigma_{PSII} (478)$ that is only 20% of that when excited at 550 nm, i.e. $\sigma_{PSII} (550)$ will be a factor of 5 greater than $\sigma_{PSII} (478)$. For the blue excitation wavelengths currently employed by the FRR, a clear difference in trajectory is thus evident between phytoplankton with a phycobilisome- (cyanobacteria and rhodophytes) versus a chlorophyll-xanthophyll- (chl *a/b*, chl *a/c* or chl *a*/peridinin) based light-harvesting complex (LHC) (Fig. 2). Consideration of the excitation wavelength is thus crucial when interpreting *in situ* data. For example, rigorous interpretation of observed gradients in values of $\sigma_{PSII} (478)$ in an ecological context strictly requires normalization to the *in situ* irradiance spectrum, i.e. $\sigma_{PSII} (in situ)$ (Moore et al. 2006b, Suggett et al. 2006b). However, we note that $\sigma_{PSII} (478)$ is likely to correlate well with $\sigma_{PSII} (in situ)$ for many natural waters, due to the dominance of blue wavebands, particularly where dominance of light absorption is by chlorophyll-xanthophyll LHCs.

Additional consideration of the expected relationship between F_v/F_m and σ_{PSII} for the prochlorophytes is more problematic. Few FRR data currently exist for

prochlorophytes under controlled growth conditions. Published data by Steglich et al. (2001) and Bruyant et al. (2005) both demonstrate that *Prochlorococcus marinus* is characterized by relatively high values of F_v/F_m (ca. 0.55 to 0.65) and low values of $\sigma_{\text{PSII}}(478)$ (ca. 300 to 600), suggesting that they fall along the eukaryote rather than the cyanobacteria trajectory (Fig. 2). Such a suggestion would seem logical since prochlorophyte pigments are dominated by divinyl chlorophylls with only relatively low concentrations of phycoerythrin (Penno et al. 2000). Furthermore, *Prochlorococcus* sp. appear to have values for RCI:RCII that are lower than for *Synechococcus* sp. by a factor of 3 (Partensky et al. 1997), and so are more similar to eukaryotes in this regard (see Suggett et al. 2007). Consequently, prochlorophytes likely maintain a relatively high PSII absorption cross section and potentially exhibit little 'contamination' of PSII fluorescence by phycobilipigment or PSI fluorescence that would act to lower F_v/F_m . However, given the apparent wide diversity of the prochlorophytes, we cannot consider this limited data set to be indicative of the physiology of the group as a whole.

Understanding the mechanisms that drive the co-variability between F_v/F_m and σ_{PSII} observed across eukaryotic algal classes requires us to examine the underlying nature of the functional absorption cross section. Several studies have previously demonstrated that variability of σ_{PSII} can be accounted for by the corresponding changes to the concentration, type and arrangement of pigmentation within the thylakoid membranes in addition to the associated RCII concentration (Suggett et al. 2004, Moore et al. 2006b). Conse-

quently, a tight relationship exists between measurements of $\sigma_{\text{PSII}}(478)$ and the ratio of photosynthetic carotenoid (PSC) to RCII concentration for those taxa whose absorption at 478 nm is dominated by these carotenoids (Fig. 4).

The clear inverse correlation between F_v/F_m and $\sigma_{\text{PSII}}(478)$, which additionally appears to correlate with cell size for eukaryotes, may be driven by a number of factors related to the energetics of light absorption and excitation energy transfer to reaction centres. These factors have likely acted as selective pressures during phytoplankton evolution. Overall an inverse relationship between the photochemical efficiency of PSII (F_v/F_m) and the amount of pigment within the antenna serving the reaction centre (which will largely dictate σ_{PSII}) is expected in energetic terms. Increased numbers of pigment molecules within an antenna will increase the lifetime of an exciton within the antenna and, hence, the probability of thermal dissipation (Lavergne & Joliot 2000). As such, photochemical efficiencies are predicted to decrease with increasing antenna size (Lavergne & Joliot 2000), as was observed (Fig. 2).

Many physiological traits have a strong dependence on cell size (Geider et al. 1986, Finkel 2001, Irwin et al. 2006), which consequently appears to have experienced strong evolutionary selection pressure (Finkel et al. 2007). Such evolutionary pressure on cell size has presumably also driven a diversification of light-harvesting strategies, resulting in the association of higher σ_{PSII} (and, hence, lower F_v/F_m) with smaller cells (Figs. 2A & 3). Phytoplankton frequently inhabit nutrient-limited environments (e.g. Cullen et al. 1992, Kolber et al. 1994, Graziano et al. 1996, Moore et al. 2006a, Behrenfeld et al. 2006); consequently, selection pressures for these photoautotrophs would be broadly expected to maximise light harvesting according to a given amount of biomass (carbon) or resource availability (nutrient). A number of hypotheses can be formulated from these basic premises.

Firstly, physical constraints associated with pigment and RCII packaging are well known to alter the light-harvesting capacity of eukaryotic algae (Duyens 1956, Morel & Bricaud 1981, Falkowski & Raven 2007). As pigment concentrations increase within a cell, self shading within the thylakoid membranes effectively reduces the efficiency of light capture by individual pigment molecules; this is known as the package effect (Morel & Bricaud 1981, Berner et al. 1989). Given a constant cellular pigment concentration, the absorption per unit pigment is predicted to decrease as cell size increases due to the susceptibility of larger cells to the package effect (Morel & Bricaud 1981). Consequently a decreased σ_{PSII} might be expected to result from a higher package effect within large cells (Fig. 3;

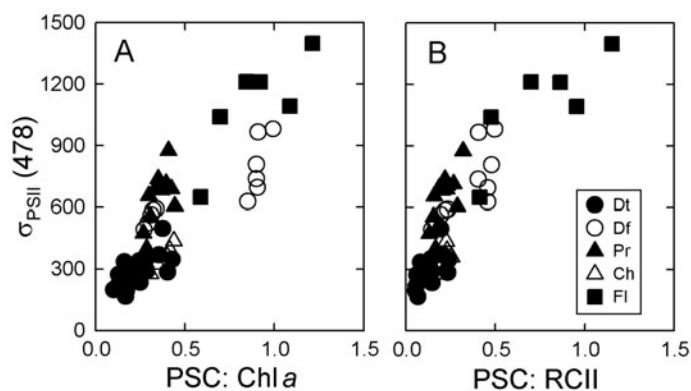


Fig. 4. Relationship between FRR measures of the effective absorption cross section ($\sigma_{\text{PSII}}(478)$, $\text{\AA}^2 \text{ quanta}^{-1}$) and the ratio of: (A) photosynthetic carotenoids (PSC) to chlorophyll a (chl a) concentration (g g^{-1}) and (B) PSC to PSII reaction centre (RCII) concentration ($\text{g } \mu\text{mol}^{-1}$). Data are included only for species of diatoms (Dt), dinoflagellates (Df), prymnesiophytes (Pr), chlorophytes (Ch) and the flagellated pelagophytes and prasinophytes (Fl) from Fig. 2, since species in all other groups lack PSC

Moore et al. 2005). However, packaging cannot explain all of the noted trends. In particular the strong correlation of higher σ_{PSII} (478) with PSC:chl *a* and PSC:RCII (Fig. 4) along with decreased F_v/F_m (Fig. 2A) suggest that the antenna size also varies between groups and, in particular, is larger for the smaller taxa.

A second consideration is the area of photosynthetic membrane space available per unit cell volume, which will decrease as cell size decreases (Raven 1986, 1999). Photosynthetic membranes contain light-harvesting complexes and reaction centre complexes, both of which contain pigmented and non-pigmented complexes, including the cytochrome b6/f complex and the chloroplast ATP synthase. Assuming that the membrane is densely packed (Kirchhoff 2008), competition for space between non-pigmented and pigmented complexes will result in the light absorbed per unit membrane surface being higher when the ratios of light-harvesting complexes relative to reaction centre and non-pigmented complexes are higher (Laverne & Joliot 2000). Larger antenna sizes will thus increase the light absorbed per unit membrane surface (Laverne & Joliot 2000) and, consequently, may offset the decreased membrane space per unit cell volume that accompanies smaller cell size (Raven 1986, 1999). Small eukaryotes may therefore have evolved higher σ_{PSII} (i.e. larger LHCs or a higher LHC:RCII ratio) in order to maintain biomass-specific light-harvesting rates.

Thirdly, the cellular resources required for the production of reaction centres are greater than those required for pigment-protein antenna complexes (this is particularly the case for iron). Consequently, selective pressure for larger σ_{PSII} (and σ_{PSI}) has been hypothesised to occur in resource-limited environments (Bibby et al. 2001, Strzepek & Harrison 2004, Moore et al. 2006b), which will also favour small cells (Irwin et al. 2006).

Finally, for prokaryotes, large membrane intrinsic antennae have been hypothesised to be a limiting factor in PSII repair, again imposing selective pressure on σ_{PSII} (Six et al. 2008). In the case of eukaryotes a small cellular pool of RCII and a higher LHC:PSII ratio (and thus larger σ_{PSII}) may be adequate in the stable light regimes frequently dominated by small cells. In contrast, the resource-rich but highly variable light environments, typically inhabited by larger diatoms, for example (Falkowski & Oliver, 2007), may favour a smaller antenna (i.e. lower σ_{PSII}), which can be readily disassembled to enable rapid replacement of the core D1 protein of PSII. Consideration of membrane diffusion rates is likely required in this context (Kirchhoff 2008). Overall each of the hypotheses proposed to explain higher σ_{PSII} at smaller cell sizes warrant further investigation in future studies.

Mixed community signals of F_v/F_m versus σ_{PSII}

Many common features have been observed in FRR fluorescence data from field studies. In particular, values of F_v/F_m are highest and σ_{PSII} (478) are lowest for coastal well-mixed waters dominated by large fast-growing diatom cells. Values of F_v/F_m decrease and values of σ_{PSII} (478) increase with enhanced water column stability and dominance by small flagellated eukaryotes (Aiken et al. 2004, Moore et al. 2005, 2006b, Fishwick et al. 2006). This same trend is observed when different size fractions of mixed community samples are individually measured (Cormeño et al. 2005; Fig. 3C), specifically, that cells within the largest size fraction exhibit the highest values of F_v/F_m and the lowest values of σ_{PSII} (478). The cell size or algal group within the mixed community that contributes the greatest proportion of the total chlorophyll *a* fluorescence appears to dominate the mixed community value measured for F_v/F_m and σ_{PSII} (478) (Suggett et al. 2004).

Field data demonstrate that the inverse relationship between F_v/F_m and σ_{PSII} (478) observed for laboratory-grown monocultures is conserved for mixed phytoplankton communities examined across relatively large spatial scales (Fig. 5). However, the trajectories are highly variable between data sets (Fig. 5). Considering the laboratory FRR fluorometry, the variability in trajectories could be interpreted as different mixtures of *Synechococcus* sp. and eukaryotic populations within the community's chlorophyll *a* fluorescence yield. In the case of the polar studies (Mertz-Polynya; Vaillancourt et al. 2003; CROZEX; Moore et al. 2007), *Synechococcus* sp. were rare along these sampling transects and, consequently, the 2 data sets overlap along the same trajectory. Unfortunately, it is impossible to rule out the possibility of instrument and sampling artefacts when comparing these various data sets, specifically, calibration of σ_{PSII} (478) and not accounting for background fluorescence (the effect of the blank; Cullen & Davis 2003). Generating absolute values of σ_{PSII} (478) requires accurate calibration of the intensity of the LEDs used to induce the fluorescence yields. Whilst such calibrations are performed at the time of instrument service, it is impossible to gauge the amount of subsequent drift without regular internal standardisation by the operator. Accurate values of σ_{PSII} (478) are also dependent upon whether instrument bias has been accounted for at the time of fitting the KPF model to the fluorescence transients (Laney & Letelier 2008). Further, the consideration of an additional sample blank is important for maximizing the accuracy of values of F_v/F_m , but not σ_{PSII} (Cullen & Davis 2003, Laney & Letelier 2008). Background fluorescence will lead to significant reductions to F_v/F_m for chlorophyll *a* concentrations less than ca.

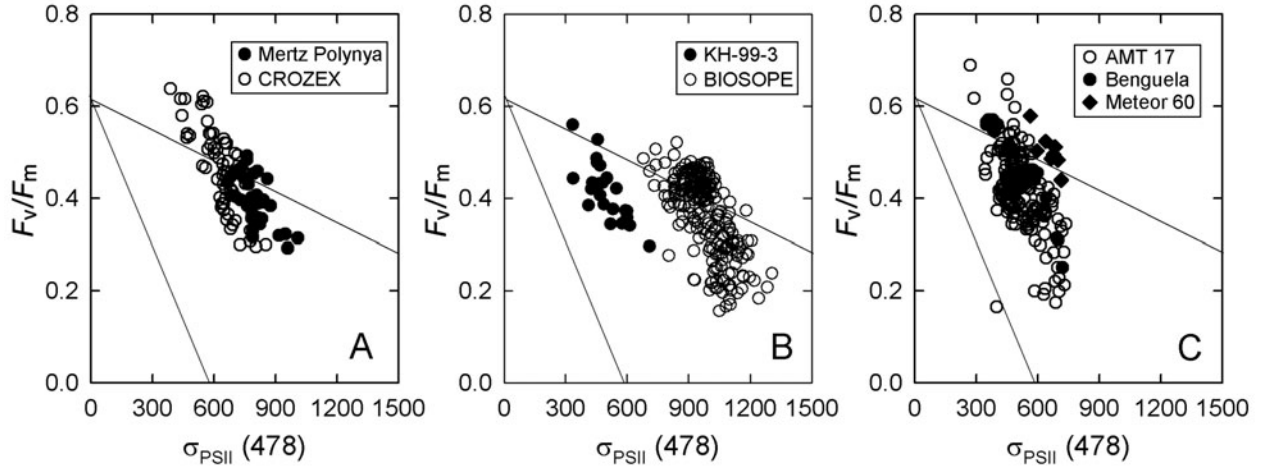


Fig. 5. Values of maximum PSII photochemical efficiency (F_v/F_m , dimensionless), and effective absorption cross section ($\sigma_{\text{PSII}}(478)$, $\text{\AA}^2 \text{ quanta}^{-1}$), measured by FRR fluorometry from: (A) Mertz-Polynya (Vaillancourt et al. 2003) and CROZEX (Moore et al. 2007) polar waters, (B) KH-99-3 (Suzuki et al. 2002) and BIOSOPE (Prášil unpubl. data) and (C) AMT 17 and Meteor 60 (Moore unpubl. data) and Benguela upwelling (Fishwick et al. 2006) and in the Atlantic. Details of the original sources of these data sets are given in Table 2. Phycobilisome and chlorophyll-xanthophyll LHC ‘trajectories’ determined for regressions applied to laboratory culture data from Fig. 2 are overlaid

1 to 10 mg m^{-3} (Suggett et al. 2006a), i.e. concentrations observed across the majority of trans-oceanic transects for the FRR data in Fig. 5. To eliminate such uncertainty, we next focus on data collected with the same inter-calibrated instruments and protocols (see Moore et al. 2005, 2006a).

Sampling campaigns that cover substantial changes in phytoplankton community structure typically display large differences in FRR fluorescence signatures (Fig. 6). Cruises JR98 and FISHES encountered phytoplankton communi-

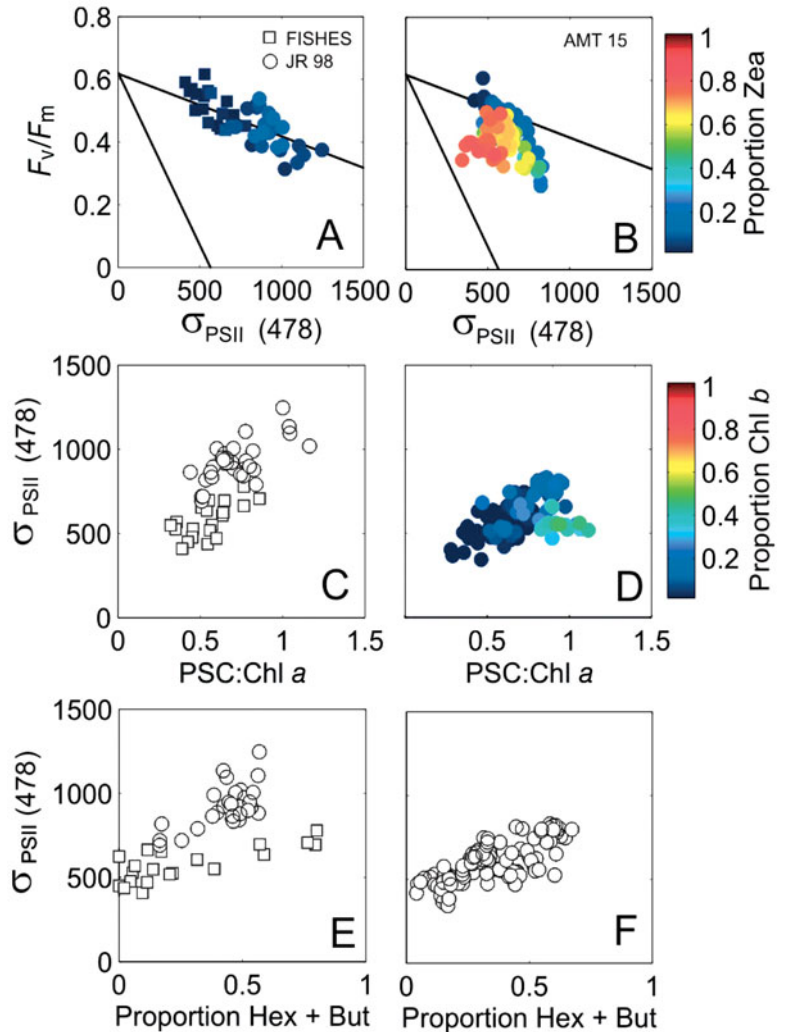


Fig. 6. Values of maximum PSII photochemical efficiency (F_v/F_m , dimensionless) and effective absorption cross section ($\sigma_{\text{PSII}}(478)$, $\text{\AA}^2 \text{ quanta}^{-1}$) measured by FRR fluorometry during: (A) FISHES (Moore et al. 2005) and JR98 (Moore et al. 2006b) and (B) during AMT 15 (Hickman unpubl. data). Phycobilisome and chlorophyll-xanthophyll LHC ‘trajectories’ determined for regressions applied to laboratory culture data from Fig. 2 are indicated. Colours indicate the relative proportion of the cyanobacterial diagnostic pigment (DP) zeaxanthin ($\text{g zeax:g } \Sigma\text{DP}^{-1}$). ΣDP refers to the sum of fucoxanthin, peridinin, alloxanthin, 19'-butanoyloxy-fucoxanthin (but), 19'-hexanoyloxyfucoxanthin (hex), chlorophyll *b* and zeaxanthin (see ‘Results and discussion: Mixed community...’). (C, D) Values of $\sigma_{\text{PSII}}(478)$ compared against the ratio of PSC to chl *a* concentration (g g^{-1}) for: (C) FISHES and JR98 and (D) AMT 15. Colours indicate the relative proportion of the prochlorophyte DP chl *b* ($\text{g chl } b:\text{g } \Sigma\text{DP}^{-1}$). (E, F) Values of $\sigma_{\text{PSII}}(478)$ compared against the proportion of total community DP contributed by both 19'-hex and 19'-but (i.e. $\text{Hex}/\Sigma\text{DP} + \text{But}/\Sigma\text{DP}$) for: (E) FISHES and JR98 and (F) AMT 15

ties that shifted from dominance by diatoms to nano-flagellates (Moore et al. 2006b) and by large diatoms to small diatoms and flagellates (Moore et al. 2005), respectively. Trends of F_v/F_m and σ_{PSII} (478) collapse clearly along a single trajectory (Fig. 6A). Although *Synechococcus* sp. was encountered during JR98, the diagnostic marker pigment for cyanobacteria, zeaxanthin, remained relatively low throughout that cruise. *Synechococcus* sp. therefore remained a minor component of the phytoplankton community and did not significantly affect the physiological indices. As demonstrated for the laboratory-grown monocultures, the values of σ_{PSII} (478) closely correspond with the PSC to chlorophyll *a* ratio (Fig. 6C).

In contrast to the study regions for JR98 and FISHES, basin-scale open-ocean cruises encounter varying proportions of prochlorophytes, *Synechococcus* sp. and picoeukaryotes and yield more complex relationships between F_v/F_m and σ_{PSII} (478) (Figs. 5 & 6). For example, for the AMT 15 data (Fig. 6B), populations with low cyanobacterial dominance (i.e. low zeaxanthin:chlorophyll ratios) display a similar inverse correlation between σ_{PSII} (478) and F_v/F_m as that found in eukaryotic laboratory cultures (Fig. 2). However, regions with relatively high zeaxanthin concentrations and, hence, increased dominance of the community biomass by *Synechococcus* sp. were associated with a cluster of low- σ_{PSII} (478), low- F_v/F_m values (Fig. 6B). It follows that the AMT 15 data are consistent with a mixture of the 2 trajectories as would be expected given the community structure. Considerable co-variability was also observed between σ_{PSII} (478) and the PSC:chl *a* ratio in the AMT 15 data (Fig. 6C). Regions with significant relative increases of chlorophyll *b*, a diagnostic marker pigment indicating prochlorophytes and/or chlorophytes, exhibited relatively low σ_{PSII} (478) (Fig. 6D); this is also generally consistent with the low values of σ_{PSII} observed for these phytoplankton taxa in culture experiments (Fig. 2).

Analysis of data from the Benguela upwelling off the south-west coast of South Africa has also demonstrated that cross-shelf trends of F_v/F_m and σ_{PSII} (478) can be broadly correlated with community structure (Fishwick et al. 2006). Once again, values of F_v/F_m increased and σ_{PSII} (478) decreased as the community changed from diatoms to flagellated (pico-) eukaryotes. Diagnostic pigments (DP) for diatoms (fucoxanthin), dinoflagellates (peridinin), picoeukaryotes (alloxanthin, 19'-butanoyloxyfucoxanthin, 19'-hexanoyloxyfucoxanthin, chlorophyll *b*) and prokaryotes (zeaxanthin) were used to partition the phytoplankton community into the percentage contributions of various taxonomic groups (Fishwick et al. 2006). This approach demonstrated that 87 % of the variance in σ_{PSII} (478) and 71 % of the variance in F_v/F_m could be explained by variability

in the percent of picoeukaryotes. In contrast, only 32 % of the variance in σ_{PSII} and 27 % of the variance in F_v/F_m could be explained by variability in the percent diatoms. The cause of correlation coefficients observed between FRR variables and the %DP that were lower for diatoms than for picoeukaryotes is not clear, but may reflect limitations inherent to DP-type reconstructions. For example, in the current context, certain groupings used by Fishwick et al. (2006) will contain organisms with very different PSII antenna structures. For example, the picoeukaryote grouping will contain both pelagophytes (19'-butanoyloxyfucoxanthin) and chlorophytes (chl *b*), which have high and low values of σ_{PSII} (478), respectively (Fig. 2).

Consequently, we can repeat the approach used by Fishwick et al. but considering instead a more detailed DP classification to account for each of the (pico)-eukaryotic groups. Thus, we assume that 19'-butanoyloxyfucoxanthin/ Σ (fucoxanthin, peridinin, alloxanthin, 19'-butanoyloxyfucoxanthin, 19'-hexanoyloxyfucoxanthin, chlorophyll *b* and zeaxanthin) (hereafter But/ Σ DP) provides some measure of relative dominance by pelagophytes; similarly, Hex/ Σ DP is the proportion of prymnesiophytes; Fuco/ Σ DP, diatoms; etc. For the JR98 and FISHES data sets that were dominated by eukaryotes, high positive co-variability was observed between σ_{PSII} (478) (Fig. 6E) or F_v/F_m (not shown) and Hex/ Σ DP + But/ Σ DP (Moore et al. 2005, 2006b). This is fully consistent with the laboratory data that indicated that pelagophytes and prymnesiophytes have a high σ_{PSII} (478) relative to other eukaryotes (Fig. 2). Similarly, despite the complicated community structure encountered during AMT 15 (Fig. 6D), >50 % of the variance in σ_{PSII} (478) across the Atlantic Ocean from ~50°N to 40°S could be explained by Hex/ Σ DP + But/ Σ DP (Fig. 6F). However, the combined dominance of these groups was a poor predictor of F_v/F_m ($r^2 < 0.1$, $n = 118$) in this case, due to the influence of the *Synechococcus* sp. populations in parts of the transect (Fig. 6B). In other words, a more sophisticated multivariate approach is required to account for natural communities composed of mixtures of different populations with a range of PSII characteristics. Finally, stepwise linear regression for each of the DP/ Σ DP against FRR measures of F_v/F_m or σ_{PSII} (478) for the AMT 15 transect results in the following relationships:

$$F_v/F_m = 0.395 \times \text{But}/\Sigma\text{DP} - 0.492 \times \text{Hex}/\Sigma\text{DP} - 0.095 \times \text{Zea}/\Sigma\text{DP}$$

$$\sigma_{PSII} (478) = 941 \times \text{Hex}/\Sigma\text{DP} - 128 \times \text{Chl } b/\Sigma\text{DP} + 92.1 - \text{Fuco}/\Sigma\text{DP}$$

where these are linear combinations and can explain the majority of the variability in F_v/F_m ($r^2 = 0.61$, $n = 118$, $p < 0.001$) and σ_{PSII} ($r^2 = 0.71$, $n = 118$, $p < 0.001$). Data sets collected throughout the Atlantic Ocean thus demon-

strate the inherent relationship between phytoplankton community structure and the fundamental characteristics of PSII as measured using FRR fluorescence.

Physiological signatures of nutrient limitation

Selective pressure from prevailing environmental conditions between water bodies simultaneously drives both the taxonomic composition of the phytoplankton community and the physiological status of the resident taxa. It is well established that well-stratified, low-nutrient oceanic waters become dominated by small-celled communities composed of *Prochlorococcus* sp., *Synechococcus* sp. and picoeukaryotes (e.g. Irwin et al. 2006). As described in the preceding section, changes in the taxonomic composition of the community (e.g. as inferred from diagnostic pigments) will impart a taxonomic signature on F_v/F_m and σ_{PSII} (478). However, F_v/F_m and σ_{PSII} may also be modified in response to physiological stress (Kolber et al. 1988, Herzig & Falkowski 1989, Geider et al. 1993b). Significantly, within a taxon, F_v/F_m decreases and σ_{PSII} (478) increases in response to nutrient starvation and other environmental stresses, e.g. photoinhibition and/or UV radiation (Vassiliev et al. 1994, Ragni et al. 2008). This pattern of change essentially reflects that observed in the taxonomic shift from diatoms to picoplankton (compare Fig. 1 with Fig. 2A).

Growth rates appear to be lower than would be predicted for picoplankton-dominated, low-nutrient oceanic waters. Such low growth rates would suggest nutrient limitation (e.g. Marañón 2005), despite the documented potential for balanced growth in these regions (e.g. Goldman et al. 1979, Falkowski et al. 1992). As discussed above, the low values of F_v/F_m and high values of σ_{PSII} (478) frequently observed in these regions cannot only be attributed to environmental stress (e.g. nutrient starvation). Furthermore, values of F_v/F_m can be maintained at the maximal values for a given taxon under nutrient-limited balanced growth (Parkhill et al. 2001). These considerations confound unambiguous interpretation of F_v/F_m and σ_{PSII} (478) for natural populations, primarily because the observed magnitudes likely represent a composite of both the phytoplankton taxonomy and nutritional status, but secondarily because taxonomic shifts and stress could both result from environmental change. Consequently, FRR measurements will be indicative of both long- and short-term environmental control upon marine primary productivity. Therefore, a key question to ask is just how much of the variability of FRR fluorescence signatures is driven by nutrient availability acting in a physiological (stress) sense? We address this question in the following section.

Nutrient limitation of natural communities

One of the fundamental applications of FRR fluorometry has been to examine the nutrient status of phytoplankton communities (Kolber et al. 1994, Behrenfeld et al. 1996, 2006, Olson et al. 2000, Boyd & Abraham 2001, Sosik & Olson 2002, Moore et al. 2006a, 2008). The majority of these studies have focused on low-Fe, high-nutrient, low-chlorophyll *a* (HNLC) ocean regions. Oceanic transects have demonstrated the effect of HNLC conditions on phytoplankton physiology. For example, FRR fluorescence measured on transects by Sosik & Olson (2002) demonstrated that decreases in F_v/F_m coincided with an increase in cell size, as the community changed from cyanobacteria- to diatom-dominated, and as waters became more Fe-depleted (Fig. 7). From the laboratory data, we would expect the larger diatoms to exhibit higher values of F_v/F_m if growth were nutrient replete. However, the most direct evidence of the effect of nutrient availability upon FRR fluorescence signatures has come from experimental nutrient addition experiments upon natural populations.

IronEx I was the first experiment on an unenclosed natural community to demonstrate the effect of nutrient enrichment upon FRR-based measures of phytoplankton physiology; specifically, a rapid (24 to 48 h) and substantial (70 to 100 %) increase of F_v/F_m and simultaneous decrease (20 to 50 %) of σ_{PSII} followed fertilisation of HNLC waters with iron (Kolber et al. 1994). Such a response reflects the relief from nutrient starvation, as discussed previously. Changes in F_v/F_m during IronEx I were observed consistently across all size classes, indicating a general physiological response by all components of the community. Other *in situ* iron release experiments have similarly shown a clear physiological response within 24 h of the experiment initiation (Boyd & Abraham 2001, Gervais et al. 2002). For example, Boyd & Abraham's (2001) FRR fluorometry from the SOIREE iron release programme detected a 35 % increase in F_v/F_m within 24 h of Fe-addition, but a much larger response of both F_v/F_m and σ_{PSII} (478) over the subsequent 6 d. It should also be noted that community changes from picoeukaryote- to diatom-dominated were simultaneously occurring over this period (Fig. 7). Similar changes on the order of 10 to 30 % have been observed in a number of such iron-release experiments.

Bottle experiments can also demonstrate the short timescale physiological response to nutrient addition. Such experiments can be conducted over relatively short periods (48 h) so as to largely preclude a change in the taxonomic composition of the community and therefore allow the physiological response to be assessed (Moore et al. 2008). For example, in the

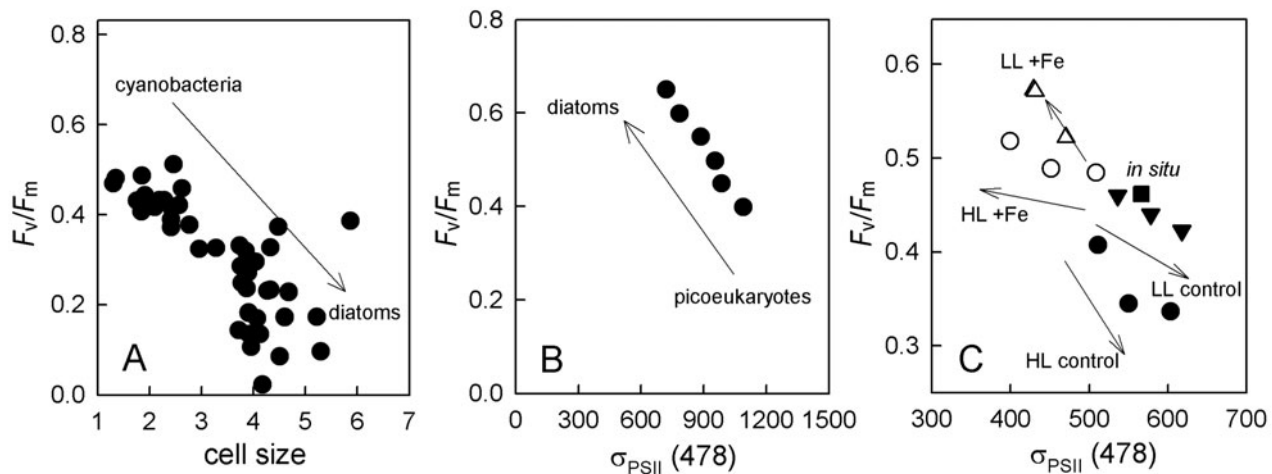


Fig. 7. (A) Relationship of maximum PSII photochemical efficiency (F_v/F_m , dimensionless) measured by FRR fluorometry to mean cell diameter (μm) from a transect across the Antarctic Polar Frontal Zone (Sosik & Olson 2002) and (B) relationship of F_v/F_m to σ_{PSII} (478) ($\text{\AA}^2 \text{ quanta}^{-1}$) within the Polar Front Antarctic Circumpolar Current during the SOIREE iron fertilisation experiment (Boyd & Abraham 2001). Values of σ_{PSII} (478) are from all depths, but are plotted as the means from bins of F_v/F_m separated by 0.05 units. (C) Relationship between FRR fluorescence measurements of F_v/F_m and σ_{PSII} (478) from discrete bioassay incubations of surface water during CROZEX (Moore et al. 2007; our Table 2). Incubations were performed in the presence (+Fe, open symbols) and absence (control, filled symbols) of iron under low-light (LL, triangles) and high-light (HL, circles) photon flux densities. Also shown is the initial value (filled square). FRR measurements were made on Days 1, 3 and 5 of the incubation. Each data point is the mean of triplicate incubations for each treatment. Arrows indicate the increase in time of incubation and the corresponding direction of change for F_v/F_m and σ_{PSII} for the 4 treatments (with respect to the corresponding *in situ* mean value recorded at the time of sample collection). Note the different axis scales between Panels B and C

case of combined light and Fe-manipulations during CROZEX (Moore et al. 2007; our Fig. 7C), the physiological response to increased Fe-supply was observed but modulated according to the simultaneous photoinhibitory damage caused by increased irradiance. Smaller scale physiological changes have also been observed for bottle experiments conducted during the Atlantic Ocean spring bloom in response to iron addition (an increase of F_v/F_m and decrease of σ_{PSII} (478) by 10 to 15%; Moore et al. 2006a; see Fig. 8).

Behrenfeld et al. (1996) demonstrated a large alteration in the physiology of prokaryotic-dominated communities within 24 h of Fe-addition during IronEx II (Fig. 8). Specifically, F_v/F_m and σ_{PSII} increased by a factor of 2 and 4, respectively (Fig. 8). Curiously, this physiological response is not what would be expected from preferential RCII re-activation/synthesis from laboratory observations, as noted previously. Also, this result does not appear to be consistent with other Fe-enrichment experiments. Recently Behrenfeld et al. (2006) interpreted these physiological changes as indicative of a re-coupling to PSII of detached iron-stress-induced antenna proteins, resulting in a simultaneous increase of σ_{PSII} and lower values of F_o and F_m , hence increasing F_v/F_m . Such a response may be specific to relieving Fe-limitation of prokaryotes.

Iron additions within HNLC regions provide key examples where FRR fluorescence indeed provides a

(relative) indicator of physiological state. In contrast, relief of nutrient limitation in nutrient-addition experiments conducted in low-nitrate, low-chlorophyll *a* regions typically produces no detectable difference (Graziano et al. 1996, Behrenfeld et al. 2006, Moore et al. 2008), or a relatively small change in F_v/F_m or σ_{PSII} (478) (Fig. 8). A number of bioassay experiments have now been performed upon Atlantic open ocean communities to examine co-limitation by multiple nutrients (N-P-Fe) (e.g. Moore et al. 2006a, 2008). In nitrogen-limited environments, only nutrient additions containing nitrogen consistently yield an increase in phytoplankton chlorophyll *a* biomass. A small physiological response can accompany this change in biomass (Fig. 8). However, the observed response is not the same as would be predicted on the basis of culture data, indicating an increase in F_v/F_m on relief of nitrogen starvation (Geider et al. 1993b). Typically, values of F_v/F_m decrease by ca. 10 to 15%, whilst values of σ_{PSII} (478) increase by ca. 15 to 20% (e.g. Fig. 8) following relief from N-limitation. One potential explanation may involve preferential synthesis of phycobilisomes leading to decreased F_v/F_m due to an artifactually high F_o as explained above, but only if cyanobacteria dominated the physiological response. Similar low physiological sensitivity (sometimes accompanied by a decrease in F_v/F_m of <50%; Behrenfeld et al. 2006) following release from N-limitation has also been ob-

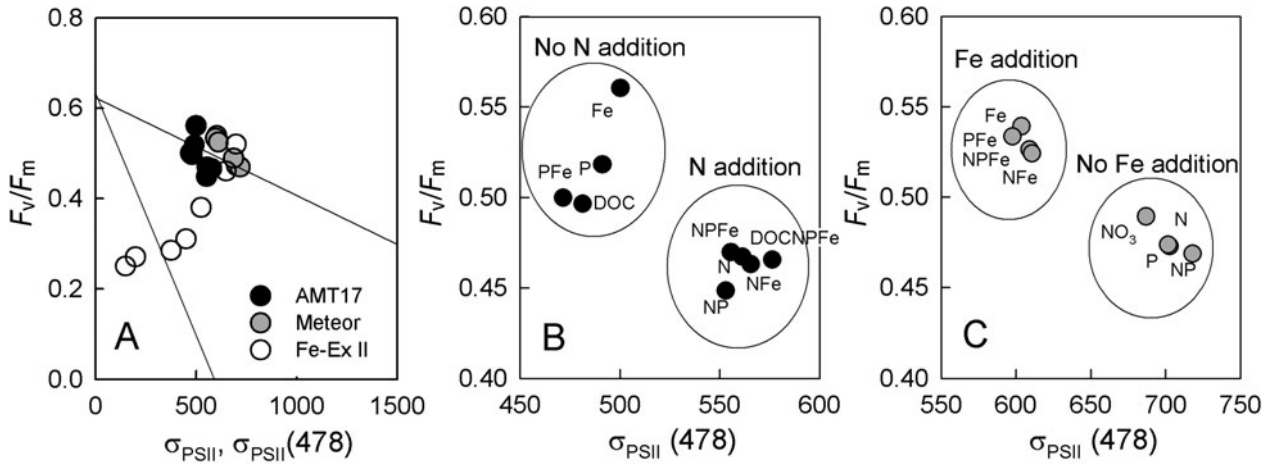


Fig. 8. Values of maximum PSII photochemical efficiency (F_v/F_m , dimensionless) and effective absorption cross section (σ_{PSII} , $\text{\AA}^2 \text{ quanta}^{-1}$) plotted for: (A) the IronEx II *in situ* fertilisation experiment (Behrenfeld et al. 1996; our Table 2), and F_v/F_m and σ_{PSII} (478) for on-deck nutrient addition incubation experiments during: (B) AMT 17, 23° S, 17° W, November 2005 (Suggett & Moore unpubl. data) and (C) METEOR 60, 42° N, 42° W, April 2004, (Moore et al. 2006a). Phycobilisome and chlorophyll-xanthophyll LHC 'trajectories' determined for regressions applied to laboratory culture data from Fig. 2 are overlaid; data from the nutrient-addition experiments have been rescaled (Panels B, C) to more clearly illustrate the small but significant effects of different nutrient treatments: +N, +P, +Fe, +NP, +NFe, +PFe, +NPFe, +NO₃, +dissolved organic carbon (DOC) combinations (note the initial and control have been removed for clarity). Each data point is the mean of triplicate incubations for each treatment. Two data clusters appear for both the AMT 17 and METEOR 60 experiments: these correspond to: (B) one for +N (low F_v/F_m , high σ_{PSII} (478)) and -N (high F_v/F_m , low σ_{PSII} (478)) for AMT17, and (C) +Fe (high F_v/F_m , low σ_{PSII} (478)) and -Fe (low F_v/F_m , high σ_{PSII} (478)) for METEOR 60

served in the sub-tropical Pacific (Behrenfeld et al. 2006) and in a high biomass subtropical estuary (Juhl & Murrell 2008).

Nutrient-limited communities thus appear to yield a FRR fluorescence signal that is indicative of a physiological response in many cases. However, the direction of change can vary between environments, and the extent of change of both F_v/F_m and σ_{PSII} in response to nutrient addition (Fig. 8) is frequently small compared to differences between taxa (Fig. 2A). Together, these data sets imply that across non-HNLC regions much of the variability in F_v/F_m and σ_{PSII} (478) can be accounted for by taxonomic gradients (Fig. 6). Lack of sensitivity of F_v/F_m to limiting macro-nutrient amendment (Graziano et al. 1996, Behrenfeld et al. 2006, Moore et al. 2008) also suggests that nutrient starvation may be rare in these systems. Much of the open ocean may thus potentially be in balanced nutrient-limited growth, with the resident phytoplankton communities physiologically acclimated to nutrient limitation and potentially experiencing environmental variability of either low frequency or magnitude (Moore et al. 2008). Under such a scenario, significant physiologically driven variability of F_v/F_m and σ_{PSII} might only be expected where steady-state conditions break down, for example, during transient nutrient-enrichment events (e.g. Falkowski et al. 1991), which cause a proportion of the community structure to experience rapid growth followed by nutrient starvation.

Interpreting future FRR measurements

FRR fluorometry has gained popularity in oceanographic studies as a means to provide a rapid assessment of phytoplankton physiology. Since its introduction, F_v/F_m (and to a lesser extent σ_{PSII}) has frequently been used as an unambiguous diagnostic of nutrient limitation, e.g. (Kolber et al. 1988, 1994, Geider et al. 1993a, Behrenfeld et al. 1996). However, as we have highlighted here, signals generated by FRR fluorescence are the product of a complex mix of acclimation, inhibition and adaptive processes. Based on the wide variability of FRR fluorescence characteristics observed for marine phytoplankton grown in culture, as well as the many consistent features observed in FRR fluorescence data sets collected across ocean regions, it is apparent that FRR fluorescence measurements *in situ* can exhibit a strong taxonomic (adaptive) component. Furthermore, the short-term stress components of fluorescence-based measurements of photosynthetic physiology appear to decrease as phytoplankton acclimate over longer periods to nutrient limitation in culture (Parkhill et al. 2001, Cullen et al. 1992, Price 2005). Such acclimation to nutrient limitation may be analogous to growth conditions in some low-nutrient, open-ocean regions (Marañón 2005, Moore et al. 2008). Together, these observations clearly demonstrate that using FRR fluorescence parameters to explicitly characterise (stress) physiology

comes with a caveat: observations of F_v/F_m and σ_{PSII} will inevitably consist of a physiological signal superimposed upon a taxonomic signal. In many cases, taxonomic gradients, as well as steady-state acclimation to long-term nutrient limitation, may thus override any effect of direct physiological stress.

Any interpretation of FRR fluorescence measurements must certainly account for changes in the community structure; in particular, if we are to improve the confidence with which FRR fluorometry accurately gauges the (eco-) physiological response of aquatic systems to environmental change. To date, a key use of FRR-based measurements of F_v/F_m and σ_{PSII} has been the estimation of PSII electron transport, and, hence, following a variety of assumptions (conversion factors), overall phytoplankton community photosynthesis *in situ* (Kolber & Falkowski 1993, Kromkamp & Forster 2003, Suggett et al. 2003, Smyth et al. 2004). Detailed discussion of the influence that the taxonomic variability described herein will have within these models is beyond the scope of the current manuscript and will be considered elsewhere. However, of key importance is that any algorithm that relies on a taxon-independent 'typical' or 'constant' value for either F_v/F_m or σ_{PSII} may ultimately be susceptible to errors.

Increasing availability of FRR data for key taxa under controlled growth environments will no doubt continue to aid the interpretation of field data; however, broad-scale environmental assessments should consider more substantial truthing of FRR 'signatures' for key species that are specific to the system under consideration. One suggestion for future FRR fluorometry would, therefore, be a movement towards multi-sensor bio-optical platforms that take advantage of the recent technological advances in instrumentation for examining microalgal community structure, for example, multi-spectral radiometers and/or submersible flow cytometers, alongside these physiological measurements.

Observations of taxonomic variability inherent to FRR-based measurements are derived from instruments tuned to detect fluorescence following excitation of a specific (narrow) waveband, hence, the difficulties in reconciling observations using a FastTracka versus a custom-built FRR fluorometer, for example. Taxonomic variability in fluorescence-based parameters simply reflects differences in antenna size, arrangement and functioning for this region of the photosynthetically active spectrum. Consequently, interpretation of observed patterns is likely to be improved by a move towards multi-spectral FRR fluorometers. The advantages are potentially 2-fold, with a means not only to examine changes in taxonomy according to bulk fluorescence yields but also to generate values of F_v/F_m and σ_{PSII} that are closer to

those that might be expected under 'natural' light fields. Either way, better resolution of the spectral dependence of the fluorescence-based description of phytoplankton physiology will generate much novel information on algal types and their response to environmental change.

Finally, we stress again that, in addition to short-term physiological stress responses, changes in bulk community ecophysiology resulting from taxonomic shifts also convey important information on system responses to environmental forcing. The taxonomic traits of different phytoplankton groups/sizes, including the variable fluorescence yield and the effective absorption cross section of PSII, will have experienced considerable evolutionary pressure. Consequently, the fundamental differences in PSII characteristics which we have discussed likely contribute significantly to phytoplankton biogeography in the modern ocean.

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