


Article

Optimising Multispectral Active Fluorescence to Distinguish the Photosynthetic Variability of Cyanobacteria and Algae

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Abstract: This study assesses the ability of a new active fluorometer, the LabSTAF, to diagnostically assess the physiology of freshwater cyanobacteria in a reservoir exhibiting annual blooms. Specifically, we analyse the correlation of relative cyanobacteria abundance with photosynthetic parameters derived from fluorescence light curves (FLCs) obtained using several combinations of excitation wavebands, photosystem II (PSII) excitation spectra and the emission ratio of 730 over 685 nm ($F_o(730/685)$) using excitation protocols with varying degrees of sensitivity to cyanobacteria and algae. FLCs using blue excitation (B) and green–orange–red (GOR) excitation wavebands capture physiology parameters of algae and cyanobacteria, respectively. The green–orange (GO) protocol, expected to have the best diagnostic properties for cyanobacteria, did not guarantee PSII saturation. PSII excitation spectra showed distinct response from cyanobacteria and algae, depending on spectral optimisation of the light dose. $F_o(730/685)$, obtained using a combination of GOR excitation wavebands, $F_o(GOR, 730/685)$, showed a significant correlation with the relative abundance of cyanobacteria (linear regression, p -value < 0.01, adjusted $R^2 = 0.42$). We recommend using, in parallel, $F_o(GOR, 730/685)$, PSII excitation spectra (appropriately optimised for cyanobacteria versus algae), and physiological parameters derived from the FLCs obtained with GOR and B protocols to assess the physiology of cyanobacteria and to ultimately predict their growth. Higher intensity LEDs (G and O) should be considered to reach PSII saturation to further increase diagnostic sensitivity to the cyanobacteria component of the community.

Keywords: active fluorescence; multispectral; phytoplankton; cyanobacteria; algae; population dynamics; limnology



Citation: Courtecuisse, E.; Marchetti, E.; Oxborough, K.; Hunter, P.D.; Spyarakos, E.; Tilstone, G.H.; Simis, S.G.H. Optimising Multispectral Active Fluorescence to Distinguish the Photosynthetic Variability of Cyanobacteria and Algae. *Sensors* **2023**, *23*, 461. <https://doi.org/10.3390/s23010461>

Academic Editor: Marc Brecht

Received: 1 December 2022

Revised: 20 December 2022

Accepted: 24 December 2022

Published: 1 January 2023



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1. Introduction

Anthropogenic activities and climate change are primarily responsible for the increased biomass and occurrence of harmful algal blooms (HABs) [1–3]. In freshwater systems, eutrophic conditions promote cyanobacteria dominance [4–6], with the added risk of proliferation of toxin-producing species [7,8]. Although not every species of cyanobacteria that forms blooms is toxic, and not every strain of toxin-producing species will always produce toxins [9], regular monitoring for cyanobacteria blooms is critical to manage risk of toxin exposure to wildlife, recreational use or through water consumption, and to limit the economic cost of mitigation measures. Risk management may include regulation of nutrient loads into water bodies or access to them. Monitoring is required at the global scale and therefore requires cost-efficient methods. Regular monitoring should be able to assess the growth and physiology of cyanobacteria and to determine the status of the bloom or the potential for a bloom to form.

Current monitoring techniques include microscopy identification and quantification, toxin analysis [10–12], genetics and genomic detection [13,14], and methods based on

optical characteristics such as in vivo fluorometry [15], flow cytometry [16] and imaging flow cytometry [17]. Microscopic identification and quantification, toxicity analysis and genetic methods are highly diagnostic but can be time-consuming and costly. Optical methods, other than microscopy, include diagnostics based on size spectrum analysis, morphology, and absorption and fluorescence characteristics. With the possible exception of image-based classification, which requires training and supervision, these methods are not able to discriminate taxonomic detail beyond functional groups. Fluorometry-based techniques are also limited to distinguishing variations in photosynthetic pigment composition at group-level, but fluorescence can be used to determine photophysiological traits that can be related to population growth.

Several types of fluorometers exist to discriminate between phytoplankton groups, including cyanobacteria. The most affordable fluorometer designs (mostly handheld and sometimes submersible) measure fluorescence from single or multiple excitation wavebands and generally one emission waveband [18,19]. These instruments assist in obtaining phytoplankton biomass estimates but are not able to assess phytoplankton physiology [20]. In contrast, saturating flash fluorometers measure variable fluorescence from which photochemical efficiency can be determined [18]. The most elaborate fluorometers in this category saturate photosystem II (PSII) within a single turnover of all reaction centres, while fluorometers operating with a longer 'multiple-turnover' flash can be more affordable [20]. From the fluorescence emission during the saturating flash, the minimum quantum yield of fluorescence F_0 (in dark-adapted state, all reaction centres opened) and the maximum quantum yield of fluorescence F_m (all reaction centres closed) are obtained. From the determination of F_0 and F_m , the variable fluorescence F_v (from $F_m - F_0$) and the maximum charge separation at PSII (F_v / F_m) are calculated. By further modulating ambient light availability, daily photosynthetic rates can be modelled from the fluorescence response [21], which in turn can be combined with nutrient and light availability in the natural environment to model and predict the growth of phytoplankton in a specific sample. Few studies to date have attempted to separate the fluorescence response from phytoplankton associated with specific pigment groups from natural samples [22,23].

Due to overlapping photosynthetic absorption spectra, a combination of fluorescence markers is required to estimate the primary production and photochemistry efficiency of cyanobacteria. Several existing variable fluorescence instruments have multiple excitation wavebands to target different spectral groups [24–26]. To distinguish the fluorescence response of cyanobacteria in the community, the fluorometer requires at least an excitation waveband which excites the main light-harvesting pigments, the phycobilipigments allophycocyanin, phycoerythrin and phycocyanin [15,27–29]. Although rhodophytes and glaucophytes are also known to possess these pigments, their abundance is unlikely to be high in environments susceptible to cyanobacteria blooms [30–32]. To accurately interpret the fluorescence signal and assess the physiology of cyanobacteria, excitation protocols targeting cyanobacteria should be able to fully saturate PSII through these diagnostic pigments. Moreover, the choice of emission wavebands for fluorometers is important to consider. Fluorescence emission is recorded at PSII chlorophyll *a* (chl_a) emission around 685 nm in most fluorometers. In cyanobacteria, most of the chl_a is associated with photosystem I (PSI) rather than PSII [33] and therefore has non-variable fluorescence [34]. The interpretation of the fluorescence signal for the emission of PSII chl_a can be complex if the sample contains cyanobacteria. Consistently higher emission ratios of 730 over 685 nm and 660 over 685 nm in cyanobacteria cultures compared to algal cultures was shown by [35]. Consequently, they suggested implementation of an emission waveband centred on phycobilisomal emission around 660 nm and an emission waveband centred around PSI chl_a at 730 nm. Single-turnover fluorometers with emission at 650 nm have not been produced, possibly due to the challenge of cross-talk between excitation sources and emission filters centred on 650 nm.

A new fluorometer, the LabSTAF (Chelsea Technologies Group, UK), shows increased potential to target cyanobacteria in the natural environment. The LabSTAF is a portable

robust and compact instrument suited for regular on-site monitoring. The instrument is well suited for long-term continuous and autonomous measurements due to a number of software and hardware features. These include a mixing feature with a flow-through stirrer, temperature control, a sample exchange feature using a separate peristaltic pump controlled by the instrument, as well as an optional cleaning cycle using the pump and additional solenoid valve controller. The instrument has high signal sensitivity (suitable for oligotrophic conditions), which means that the full seasonal phytoplankton variability may be observed. Moreover, the instrument introduces several optical features which increase its potential to target distinct pigment groups, including those diagnostic of cyanobacteria. These include multiple excitation wavebands (see below), emission detection using 685 nm and 730 nm bandpass filters and the automated function to measure photosynthetic excitation spectra. The suitability of these specific features to target cyanobacterial photophysiology is discussed below.

The instrument is based on the concept of a Single Turnover Active Fluorometer (STAF) and uses seven excitation wavebands centered at 416, 452, 473, 495, 534, 594 and 622 nm, which can be combined to target a range of photosynthetic pigments based on their spectral characteristics. Wavebands at 534 (green) and 594 nm (orange) are particularly useful, as these target different forms of phycoerythrin that are common in both oceanic and freshwater cyanobacteria, as well as the short-wavelength tail of phycocyanin absorption [36]. The LabSTAF is equipped with a broad-spectrum actinic light source to create Fluorescence Light Curves (FLCs) that are synonymous with photosynthesis–irradiance curves used to estimate and model primary production. The single turnover variable fluorescence is recorded at the waveband 685 nm to target PSII chl_a. In addition it obtains F_o , F_m and F_v at 730 nm at the start of a measurement sequence, using a linear actuator to switch between 685 nm and 730 nm bandpass filters (± 10 nm FWHM). This Dual Waveband Measurement (DWM) configuration was initially designed to correct for the pigment packaging effect [37,38]. We propose that this configuration may also be sensitive to the presence of cyanobacteria due to allocation of chl_a to PSI rather than PSII, resulting in a higher 730 over 685 nm emission ratio when chl_a light absorption is targeted. The LabSTAF is further preconfigured to measure the photosynthetic excitation spectrum across the seven excitation wavebands. This is done using one of two distinct protocols, differing in the light dose used at the spectral excitation maximum of algal and cyanobacterial pigment groups, respectively.

In this study, we assess the ability of the LabSTAF fluorometer to target physiological properties of cyanobacteria in a freshwater reservoir where cyanobacteria form annual blooms. Various combinations of fluorescence markers are tested and compared against relative phytoplankton abundance and nutrient availability. Specifically, we analyse the correlation of relative cyanobacteria abundance with photosynthetic parameters derived from FLCs obtained using combinations of excitation wavebands, the interpretation of PSII excitation spectra (both prokaryotic cyanobacteria and eukaryotic algae-optimised protocols), and the emission ratio of 730 over 685 nm obtained with excitation protocols targeting cyanobacteria versus algae. Appropriate predictors of cyanobacterial photophysiology within the phytoplankton community are expected to reflect a positive response of cyanobacteria to conditions favouring their growth, such as reduced N:P ratios and following depletion of silicate in the natural succession from spring (diatoms abundant) to summer (cyanobacteria dominant) community composition. Moreover, we expect fluorescence dynamics to reflect the phytoplankton group abundance estimates from microscopy counts. Comparing the various options for light-excitation protocols, we hypothesise that the photophysiological characterisation of algae and cyanobacteria is best achieved by interpreting single-turnover fluorescence from blue and green–orange light to target diagnostic pigment groups. Saturation and specificity of cyanobacteria fluorescence from green–orange excitation is the most critical test of the system, due to lower excitation energy achieved with light emitting diodes and the overlap with absorption by phycobilipigments and short-wavelength tails of both chlorophyll *c* and chl_a in this range of the spectrum.

2. Materials and Methods

2.1. Study Site

Roadford Lake is the largest drinking water reservoir in Devon County in the United Kingdom (Figure 1), part of the catchment of the river Tamar. It has a net storage of 34,500 million litres and a surface area of 295 hectares. Water is pumped from a draw-off tower situated in the south part of the reservoir, near the dam where there is also an overflow tower [39]. The lake is fed by the river Wolf in the northeast and from Westweek inlet, a stream to the northwest.

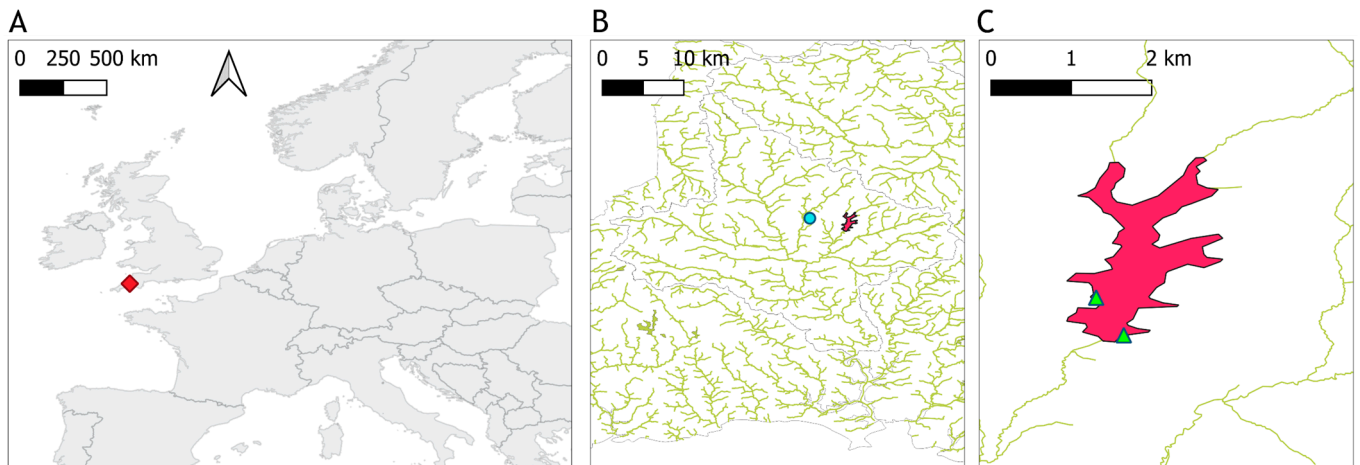


Figure 1. Location of Roadford Lake. (A) The location of the reservoir indicated with a red diamond marker. (B) The meteorological station “Virginstow, Beaworthy” represented by a turquoise circle to the west of the reservoir in magenta color. (C) Green triangles represent the sampling points on the reservoir.

High abundances of cyanobacteria have been recorded in samples analysed by the water utility company using light microscopy, since at least spring 2002, and often year-round (Figure 2, see Section 2.3 for methods). Water samples for this study, 23 in total, were collected between 12 April and the 16 October 2020. Samples were taken every two weeks during the sampling period, and weekly during the summer, when high cyanobacteria biomass was expected. Surface water was sampled from a jetty near the western shore with a bucket in the first 0.5 m from the surface in approximately 3 m-deep water (depending on reservoir volume), except for the first seven samples which were taken directly from the east shore whilst access to the jetty was restricted (Figure 1C). During calm weather, any accumulation of cyanobacteria visible at the surface was mixed with the surface water to ensure that the sample was homogeneously representative of conditions in the top layer. Water temperature was recorded at the time of sampling, and approximately 2 litres of water were brought to the laboratory in an insulated container, for further analysis on the same day. Total daily rainfall was retrieved from the National Meteorological Library and Archive of the UK Met Office for station “Virginstow, Beaworthy” ($50^{\circ}42'36.0''$ N $4^{\circ}17'45.6''$ W) which is located around 4 km west of Roadford Lake (Figure 1).

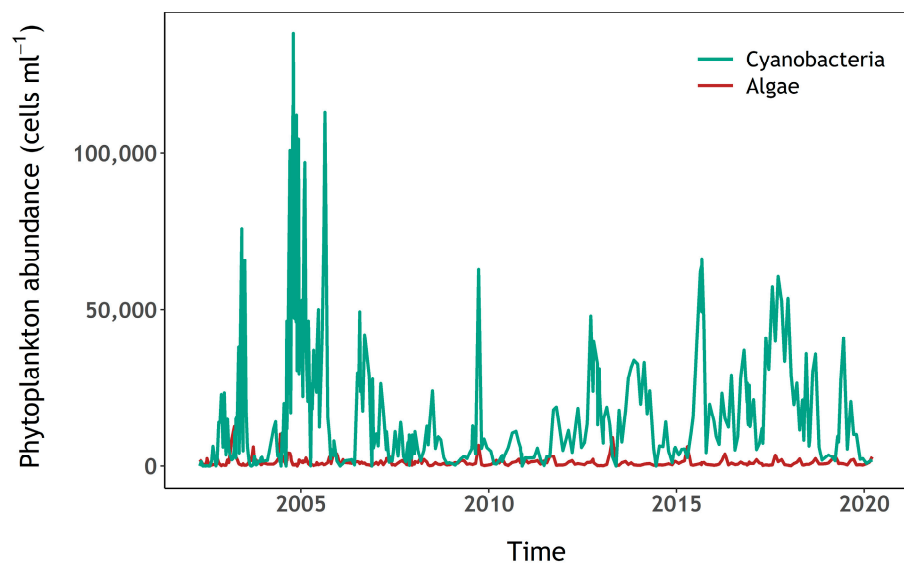


Figure 2. Algae and cyanobacteria abundance (cells mL⁻¹) at Roadford Lake.

2.2. Single-Turnover Fluorescence

A dedicated LabSTAF instrument (serial number 19-0105-006) was used to characterise the photophysiology of phytoplankton at Roadford Lake. FLCs were measured on samples taken from 29 May to 16 October 2020, using the range of fluorescence excitation protocols described in Table 1. These protocols were selected to target semi-isolated pigment groups and the associated phytoplankton taxa through combinations of excitation wavebands expected to lead to full saturation of PSII, based on the work of [40,41]. Protocols differed in saturation pulse length (100 or 200 μ s) to accommodate saturation, and the combination of excitation wavebands denoted B(lue) = 452 nm, G(reen) = 534 nm, O(range) = 594 nm, and R(ed) = 622 nm (Table 1). The Light-Emitting Diodes (LEDs) used to provide the excitation energy were always set to their maximum intensities ($G = 10,500$, $O = 2896$ and $R = 7594 \mu\text{mol photons m}^{-2} \text{s}^{-1}$) except for the B LED which adapted automatically to adjust to the phytoplankton composition and abundance for the B protocol (2 are implemented in the instrument, B1 ranged between 14,108 and 16,819 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ and B2 between 14,068 and 16,719 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$) (Table 1). Optimal B intensity values within the GORB and GOB protocols were determined prior to any measurements, from exposure using only the B LEDs with a pulse length of 200 μ s. These values for the B LED were then adopted in the GORB and GOB protocols with the aim to see if results from separate protocols were additive (Table 1). Each FLC included 12 actinic light steps from 0 to 1200 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$. The actinic light spectra had a peak at 455 nm and shoulder from 480 nm to 655 nm. Samples were dark adapted for at least 20 min before each analysis to allow all reaction centres to open to correctly measure F_0 .

Photosynthetic parameters are expressed here as a function of excitation and emission waveband (λ_{ex} , λ_{em}). FLCs parameters and excitation spectrum were reported only for the emission band (λ_{em}) 685 nm, but single turnover curves were also recorded at $\lambda_{\text{em}} = 730$ nm. Where multiple excitation wavebands are considered, the excitation waveband is referred to by the abbreviations given in Table 1 (e.g., B, GOR). Several photosynthetic parameters were determined from each FLC: $\alpha_{\text{P}_{\text{II}}}$, $J_{\text{P}_{\text{II}}}$, P_{max} , $\sigma_{\text{P}_{\text{II}}}$, F_v/F_m and F_0 . The following definitions are repeated from the instrument manual [37] and the terminology of these photosynthetic parameters is derived from [26,37]. $\alpha_{\text{P}_{\text{II}}}$ defines the initial rate at which photons are used to drive PSII photochemistry during a single turnover (ST) pulse. This parameter is a proxy of saturation of PSII photochemistry. A value of $\alpha_{\text{P}_{\text{II}}}$ between 0.042 and 0.064 is considered optimal and reflects a good ST curve fit. $J_{\text{P}_{\text{II}}}$ measures the photon flux through the absorption cross section for PSII photochemistry provided by a single PSII complex. $J_{\text{P}_{\text{II}}}$ is interpreted as electron transport rate (ETR) on the assumption that

each photon used to drive PSII photochemistry results in the transfer of an electron out of PSII. ETR, in turn, determines the rate of electron transport which controls phytoplankton carbon fixation and growth [42]. ETR can express the ability of the phytoplankton to achieve metabolic processes. P_{\max} is the maximum specific photosynthetic rate which expresses the phytoplankton photosynthetic capacity in optimal ambient light conditions. σ_{PII} is the absorption cross section of PSII, which is the product of the absorption cross section for PSII light-harvesting and the probability that an absorbed photon will be used to drive PSII photochemistry. F_v/F_m measures the photochemistry efficiency, with F_v the variable part of fluorescence obtained from the difference between F_o and F_m , the minimum and maximum fluorescence, respectively. Values of F_o , F_v/F_m , α_{PII} and σ_{PII} tend to decrease along the light steps of each FLC. The largest value of these parameters may be observed at or shortly after the first light step. Consequently, reported values of F_o , F_v/F_m , α_{PII} and σ_{PII} were averaged over the first five light steps to characterise each FLC. The maximum value in each FLC was selected to describe the ETR parameter. P_{\max} is provided once per FLC.

Table 1. Excitation protocols tested to target pigments associated with specific phytoplankton groups, as described by [40,41]. Phycobilipigments include phycoerythrin, allophycocyanin and phycocyanin.

Protocol	Excitation Wavebands (nm at Centre)	Intensity Range ($\mu\text{mol Photons m}^{-2} \text{s}^{-1}$)	Pulse Length (μs)	Photosynthetic Pigment Groups Targeted	Phytoplankton Group Targeted
B	452, 452	28,175–33,538	100	Chlorophylls <i>a/b/c</i> , carotenoids	Algae, weak signal from cyanobacteria possible
GOR	534, 594, 622	20,990	200	Phycobilipigments Chlorophylls <i>a/b/c</i> and carotenoids	Cyanobacteria with weaker signal from algae likely
GORB	452, 452, 534, 594, 622	49,513–54,528	200	Chlorophylls <i>a/b/c</i> , carotenoids and phycobilipigments	Whole community
GOB	452, 452, 534, 594	41,919–46,933	200	Chlorophylls <i>a/b/c</i> , carotenoids and phycobilipigments	Whole community with a slower saturation response compared to GORB
GO	534, 594	13,395	200	Phycobilipigments, Chlorophyll <i>c</i> , carotenoids	Cyanobacteria, cryptophytes and rhodophytes

ST curves were recorded at both emission wavebands (685 and 730 nm) and with the B and GOR protocols, prior to running each FLC. Excitation spectra of $F_v(\lambda_{\text{ex}}, 685)$ and $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ were also collected before running FLCs. Depending on the $F_o(452, 685)$ response as a proxy for the presence of chlorophylls *b* and *c*, the LabSTAF optimises and scales the excitation spectrum to the response from either blue or orange excitation [37], here referred to as algae and cyanobacteria-optimised protocols, respectively.

2.3. Microscopy

Microscopy counts were collected from samples fixed in Lugol's medium. These samples were placed in a 1 mL Sedgewick Rafter Counting Chamber etched with a 20 row \times 50 column grid. Samples were examined with a LEICA DM IRB inverted microscope. 100 units of 1 μL each were randomly selected and counted to phytoplankton group level to quantify the abundance of diatoms, cyanophytes, chrysophytes, chlorophytes, euglenoids, dinoflagellates and unspecified single-celled eukaryotes. Counts were then multiplied by 10 to yield the abundance per mL.

Microscopy counts and identification to species level were also available from the water utility company (South West Water), from April 2002 to present, generally at monthly intervals. These long-term monitoring data were only used in Figure 2 to inform planning of the sampling campaign. Identification was made to species level, whereas the results reported here are aggregated to either “eukaryotic algae” or “prokaryotic cyanobacteria”. The samples were collected from water pumped at the draw-off tower from approximately 5 or 10 m depth, depending on surface water level.

2.4. Nutrients

Dissolved nutrient concentrations including nitrate + nitrite, nitrite, phosphate, ammonium and silicate were determined from 0.2- μm filtrate (Nalgene™ Sterile Syringe Filters 0.2 μm surfactant-free cellulose acetate membrane) using a 5-channel segmented flow colorimetric SEAL Analytical AAIII autoanalyser. The analytical methods were as described in [43].

3. Results

3.1. Phytoplankton and Nutrient Dynamics

Algae were dominant on the first sampling day followed by cyanobacterial dominance during the rest of the campaign (Figure 3A). Diatoms and chlorophytes dominated the algal community, with diatoms varying over a wide range between samples (Figure 3B,C). A succession of phytoplankton groups was observed through the sampling period and followed variability in nutrient concentrations (Figures 3 and 4). The beginning of the sampling period was marked by an initial increase in diatoms, cyanobacteria and chlorophytes until late April, during which silicate and phosphate concentrations decreased whilst a slight increase of available ammonium was observed. Cyanobacteria and diatoms abundance then decreased sharply until late May, while chlorophytes continued to increase, nitrate decreased, and nitrite availability slowly increased. From mid-May to mid-August, cyanobacteria increased while algae abundance remained relatively low. During this period, a decrease in nitrate and increase in nitrite was observed, while silicate and ammonium availability peaked. From mid-August until the end of the sampling period, algae abundance (notably diatoms and chlorophytes) increased while cyanobacteria showed an overall increasing trend with high variability between samples. Meanwhile, a slow increase of silicate and a decrease of nitrate and nitrite was observed.

There was evidence of phosphate limitation during the whole sampling period (Figure 4B). Phosphate concentration was slightly above the detection limit on the first sampling date and below the limit of detection (0.02 μM) during the remaining period (Figure 4B). Ammonium concentration (Figure 4B) peaked in mid-July, following a period of high precipitation during that period (Figure 5) which likely replenished both nitrite and ammonium from surrounding agricultural sources (Figure 4A,B). Silicate availability is a common driver of early-season phytoplankton succession. Silicate concentration decreased at the beginning of the sampling period when diatoms were relatively abundant, then increased up to the end of the sampling period while cyanobacteria were dominant and available ammonium appeared to be rapidly taken up (Figures 3 and 4A).

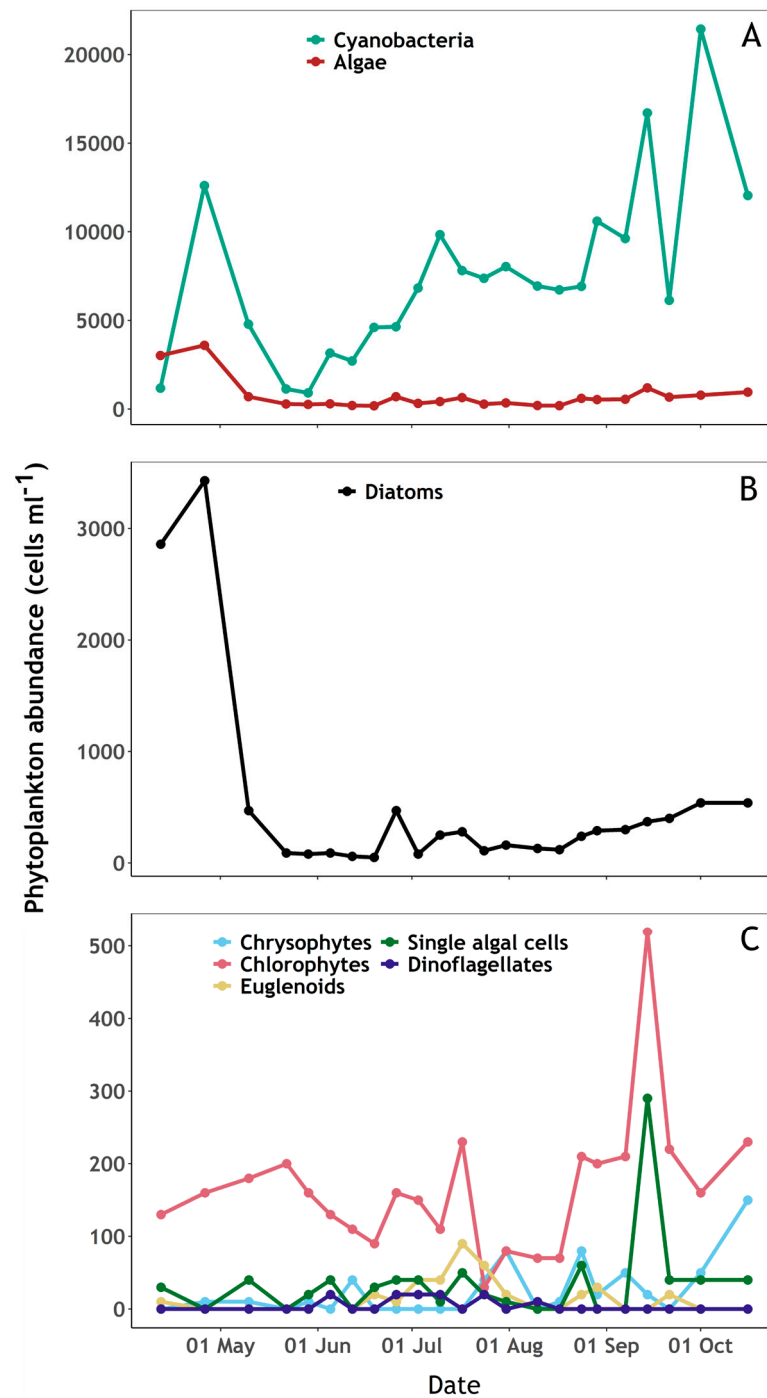


Figure 3. Phytoplankton abundance (cells mL⁻¹) during the sampling period at Roadford Lake. (A): Cyanobacteria and algae. (B): Diatoms. (C): Chrysophytes, Chlorophytes, Euglenoids, Dinoflagellates and single algal cells.

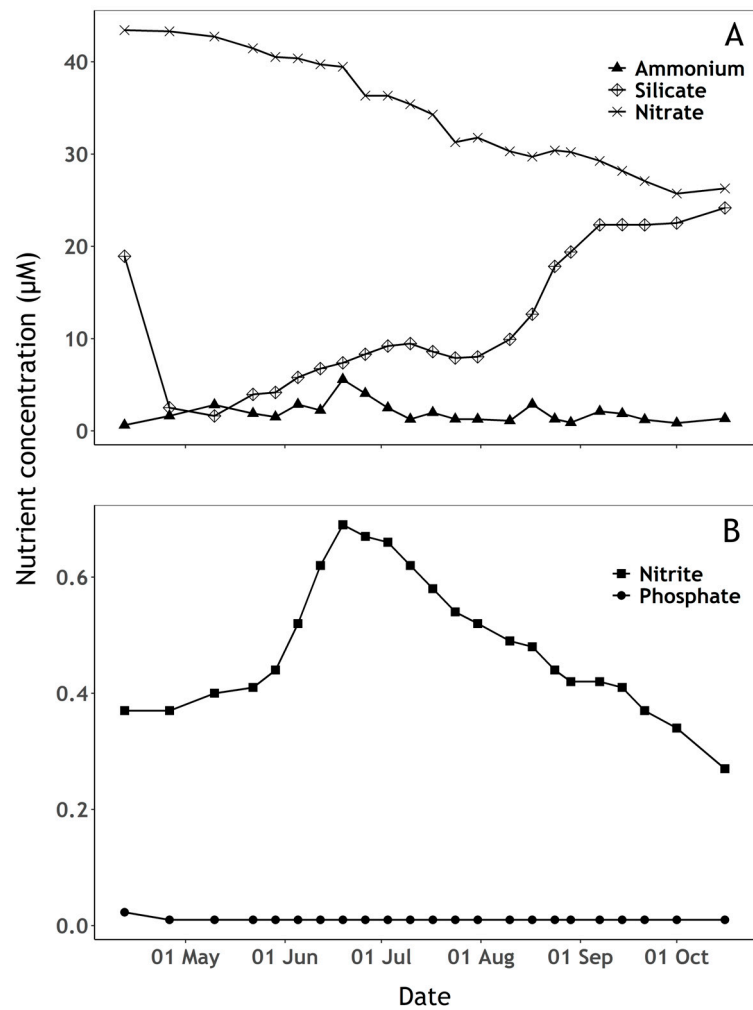


Figure 4. Nutrient concentrations (μM) over the sampling period. (A) Ammonium, silicate and nitrate concentration, (B) Nitrite and phosphate concentration. When the detection limit was not passed, phosphate concentrations below the limit of detection ($0.02 \mu\text{M}$) are displayed as $0.01 \mu\text{M}$.

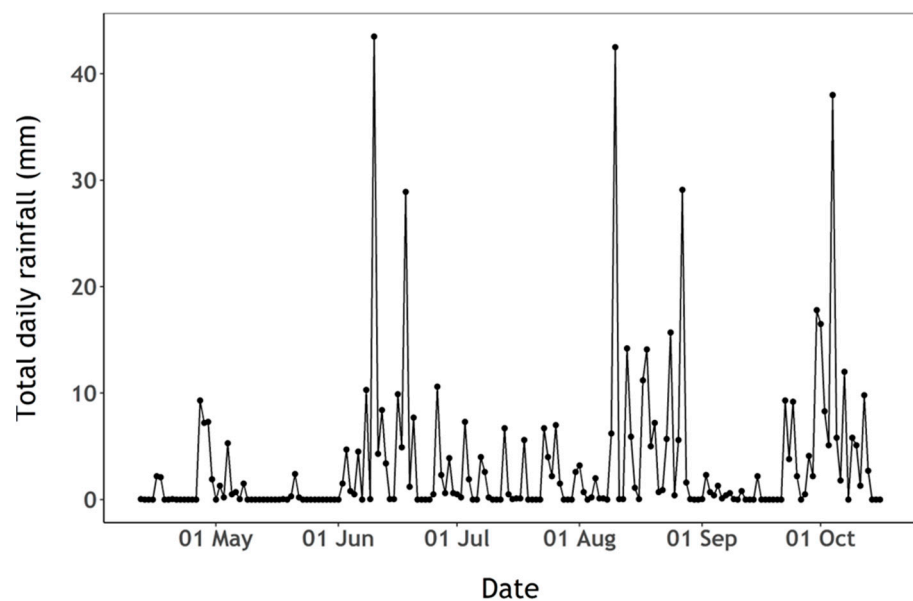


Figure 5. Total daily rainfall (mm) observed at the Virginstow, Beaworthy weather station.

3.2. Fluorescence Dynamics

The succession towards increasing cyanobacteria abundance and dominance was clearly reflected in $F_V(\lambda_{exc}, 685)$ and $\sigma_{PII}(\lambda_{exc}, 685)$ (Figure 6). $F_V(\lambda_{exc}, 685)$ showed similar dynamics between the algae and cyanobacteria-optimised protocols (Figures 6A and 6B, respectively). $F_V(594, 685)$ and $F_V(622, 685)$ increased throughout the sampling period, matching changes in cyanobacteria abundance (see Figure 3A) while higher values of $F_V(416, 685)$ and $F_V(452, 685)$ corresponded to higher abundance of algae at the beginning of the sampling period (Figure 6A,B). It should be noted that the algae-optimised protocol showed higher values of F_V than the cyanobacteria-optimised protocol (Figure 6A,B). $\sigma_{PII}(\lambda_{exc}, 685)$ showed distinct responses from different parts of the community between the two protocols (Figure 6C,D). The blue part of the spectrum showed a gentle decrease in σ_{PII} under the algae-optimised protocol while under the cyanobacteria-optimised protocol, σ_{PII} was higher in early April with strong short-term variations until Mid-July. These variations showed contrasting behaviour between blue to orange–red excitation wavebands for σ_{PII} in the cyanobacteria-optimised protocol (Figure 6D), largely absent in the algae-optimised result (Figure 6C). Differences in σ_{PII} between the algae and cyanobacteria-optimised protocols were most significant in the red–orange excitation wavebands. $\sigma_{PII}(622, 685)$ increased markedly in the middle of the sampling period using the cyanobacteria-optimised protocol (Figure 6D) while $\sigma_{PII}(534, 685)$, $\sigma_{PII}(594, 685)$ and $\sigma_{PII}(622, 685)$ showed lower values under the algae-optimised protocol.

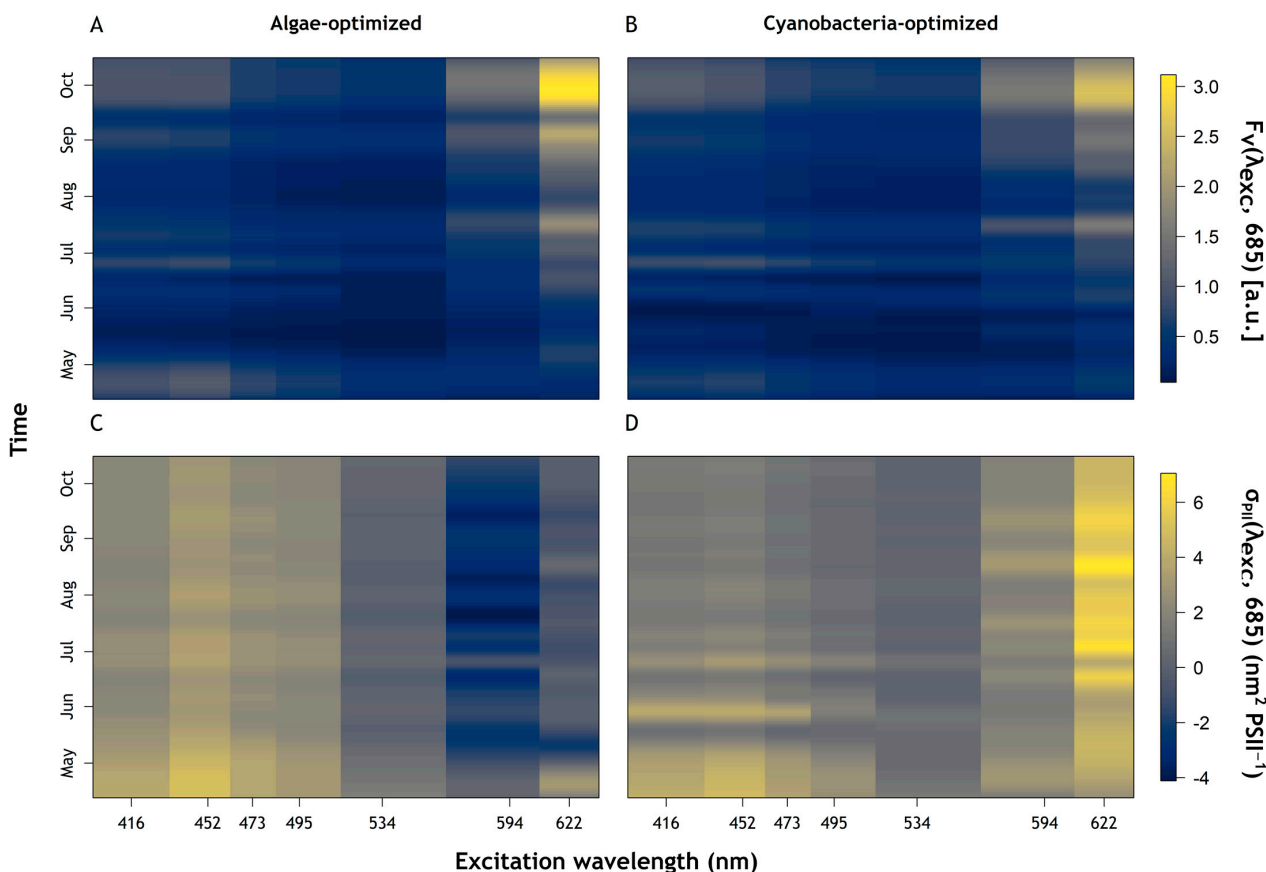


Figure 6. Photosynthetic Excitation spectrum results. (A) $F_V(\lambda_{exc}, 685)$ [a.u.], algae-optimised. (B) $F_V(\lambda_{exc}, 685)$ [a.u.], cyanobacteria-optimised. (C) $\sigma_{PII}(\lambda_{exc}, 685)$ ($\text{nm}^2 \text{PSII}^{-1}$), algae-optimised. (D) $\sigma_{PII}(\lambda_{exc}, 685)$ ($\text{nm}^2 \text{PSII}^{-1}$), cyanobacteria-optimised.

3.3. Relative Cyanobacteria Abundance and F_o Dynamics

$F_o(\text{GOR})$ trends followed variations in cyanobacteria abundance over time (Figure 7B, and see Figure 3A). $F_o(\text{GO})$ showed the same trend but with much lower values compared

to $F_o(\text{GOR})$, indicating that $F_o(\text{GO})$ was sensitive to cyanobacteria abundance despite lower excitation energy. Abundances of algae and cyanobacteria were significantly correlated with F_o obtained from the five excitation protocols. The abundance of cyanobacteria was more significantly correlated with $F_o(\text{GOR})$ (linear regression, p -value < 0.0001, adjusted $R^2 = 0.66$) than with other protocols (linear regression, p -values between p -value < 0.01 (GOB) and p -value < 0.0001 (GORB), adjusted R^2 ranged between 0.30 (GOB) and 0.58 (GORB)). Reciprocally, the abundance of algae was most strongly correlated with $F_o(\text{GORB})$ (linear regression, p -value < 0.0001, adjusted $R^2 = 0.57$) than other protocols (linear regression, p -value between p -value < 0.001 (GOR) and p -value < 0.001 (B), adjusted R^2 ranged between 0.46 (GOR) and 0.57 (B)).

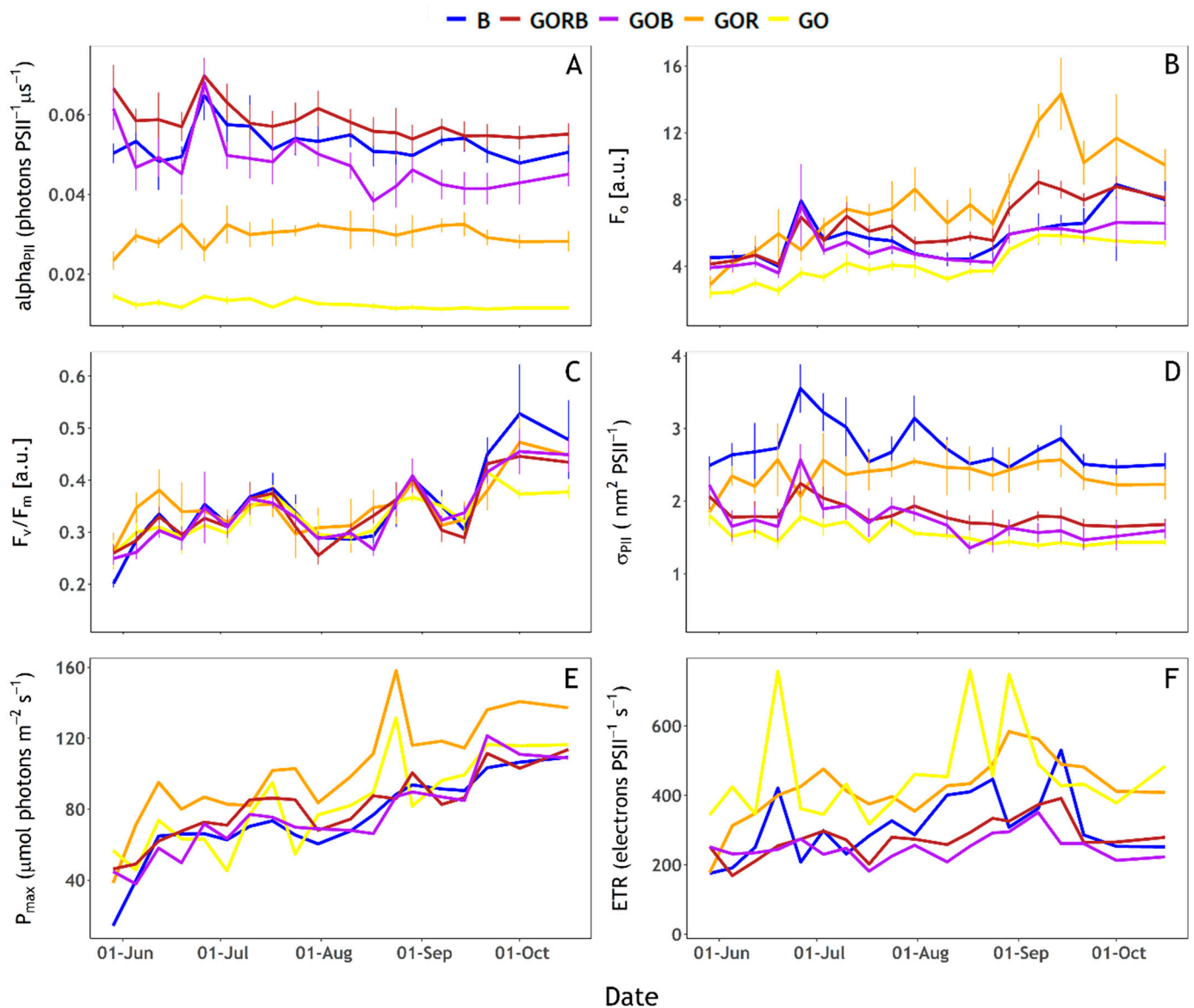


Figure 7. Fluorescence Light Curves (FLCs) photosynthetic parameters obtained with the five excitation protocols (B), GORB, GOB, GOR and GO, see Table 1). (A) $\alpha_{\text{P}_{\text{II}}}$ (photons $\text{PSII}^{-1} \mu\text{s}^{-1}$), (B) F_o [a.u.], (C) F_v/F_m [a.u.], (D) $\sigma_{\text{P}_{\text{II}}}$ ($\text{nm}^2 \text{PSII}^{-1}$), (E) P_{max} ($\mu\text{mol photons m}^{-2} \text{s}^{-1}$) and (F) ETR (electrons $\text{PSII}^{-1} \text{s}^{-1}$).

3.4. Photophysiological Characterisation

When targeting cyanobacteria, full saturation of PSII was not reached with the GOR and GO protocols. $\alpha_{\text{P}_{\text{II}}}(\text{GO})$ was < 0.02 and $\alpha_{\text{P}_{\text{II}}}(\text{GOR})$ around 0.03 (Figure 7A). The

other protocols (B, GORB and GOB) did achieve saturation of the community with $\alpha_{P_{II}}$ near or above 0.04. The fitting of ST curves to obtain photophysiological parameters does not necessarily require full saturation as long as the initial fluorescence rise and asymptote (F_m) can be extrapolated from the curve. Nevertheless, GO protocol results should be interpreted with caution.

Photophysiological parameters derived from the FLCs showed multiple trends. P_{max} , F_v/F_m and F_o all increased over time (Figure 7B,C,E). F_v/F_m increased from 0.2–0.3 at the end of May to approximately 0.4–0.5 at the end of the sampling period with minor variations between excitation protocols, detailed below (Figure 7C). By contrast, $\sigma_{P_{II}}$ showed more short-term variations and a generally decreasing trend (Figure 7D). ETR generally increased up to mid-September before it decreased again, while ETR(GO) results were highly variable (Figure 7F). Assessing specific differences between excitation protocols, P_{max} (GOR) and ETR(GO) showed higher values than other protocols (Figure 7 E,F). P_{max} (GOR) and P_{max} (GO), as well as ETR(GOR) and ETR(GO), followed similar rising trends. When combined with B excitation (GOB and GORB protocols), these trends remained similar, whilst P_{max} (B) lacked some of the short-term variability over the duration of the rising trend. ETR(B) showed a contrasting trend from ETR(GORB) and ETR(GOB). While ETR(GO) and ETR(GOR) trends seem mostly driven by cyanobacteria abundance, ETR(B) trends seem to follow algae abundance (Figure 7F). Moreover, changes in diatoms, chlorophytes and other single-celled algae abundance followed short-term variations of ETR(B) even when algal abundance was low (Figure 7F). $\sigma_{P_{II}}$ (B) and $\sigma_{P_{II}}$ (GOR) were both higher than the other excitation light compositions (Figure 7D). $\sigma_{P_{II}}$ (B) showed three peaks in the time series during late June, late July and mid-September. Only the steady decrease following the September peak was also visible in $\sigma_{P_{II}}$ (GOR). Short-term variations of $\sigma_{P_{II}}$ (B) and $\sigma_{P_{II}}$ (GOR) seemed to follow short-term variations of, respectively algae and cyanobacteria abundance (Figure 7D). The other protocols showed similar trends between them with values decreasing slowly from the end of May up to the end of the sampling period, and a $\sigma_{P_{II}}$ peak observed at the end of June with GORB, GOB, and GO protocols.

Photochemistry efficiency (F_v/F_m) was similar between excitation light protocols except marked differences at the start of June (higher F_v/F_m (GOR)) and at the end of September (higher F_v/F_m (B) and lower F_v/F_m (GO), Figure 7C). F_o (GOR) was higher than F_o obtained with the other protocols (Figure 7B). F_o (GO) was consistently lower compared to other combinations of excitation wavebands. F_o (GORB) showed higher values than F_o (B), F_o (GOB) and F_o (GO) protocols from July until the end of the sampling period. F_o (GORB) was lower than F_o (GOR), contrary to expectations considering the GORB protocol operated the GOR LEDs at the same intensities between protocols and received additional energy from the B LEDs.

3.5. Emission Ratio of 730 nm over 685 nm

The emission ratio of 730 over 685 nm, or F_o (GOR, 730/685), corresponded to cyanobacteria biomass as seen in Figure 8. F_o (GOR, 730/685) was positively and significantly correlated with the relative abundance of cyanobacteria of the phytoplankton community (linear regression, p -value < 0.01, adjusted $R^2 = 0.42$). F_o (B, 730/685), in contrast, was not significantly correlated with the relative abundance of either group (linear regression, p -value: 0.076, adjusted $R^2 = 0.12$), although some correspondence of F_o (B, 730/685) with cyanobacteria biomass could be observed.

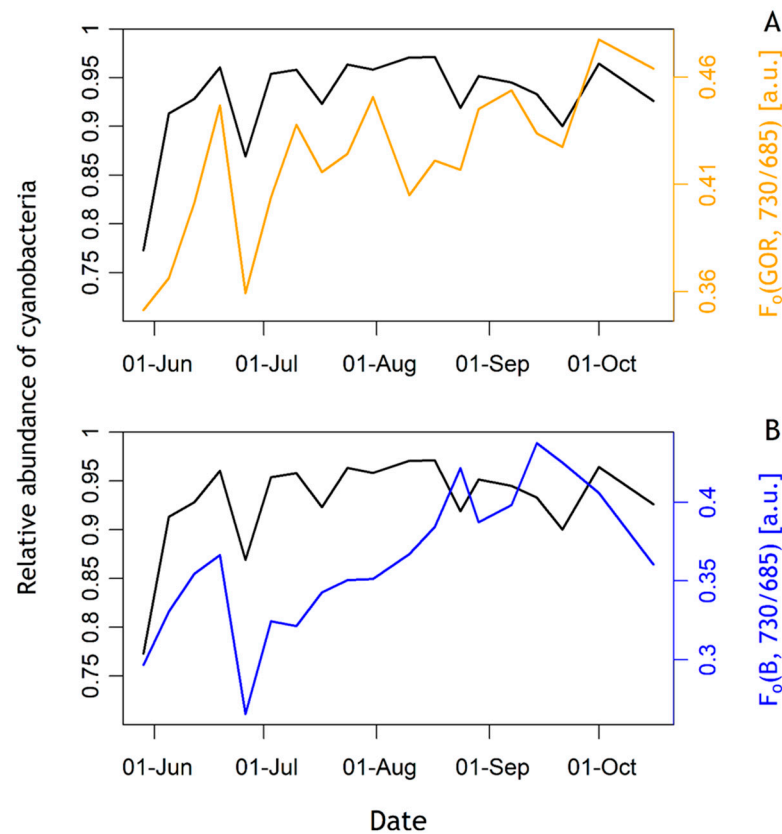


Figure 8. F_o emission ratio using B and GOR excitation protocols compared to the relative abundance of cyanobacteria (black drawn line). (A) $F_o(\text{GOR}, 730/685)$ [a.u.], (B) $F_o(\text{B}, 730/685)$ [a.u.].

4. Discussion

4.1. Phytoplankton, Nutrient and Fluorescence Dynamics

The succession of phytoplankton observed from early spring to late summer is typical of the seasonal succession of phytoplankton in temperate lakes [44,45]. The increase of cyanobacteria may be explained by a combination of several factors: temperature increase, replenishment of ammonium, and the ability of cyanobacteria to store phosphorus [46].

The use of fluorescence excitation spectra to distinguish pigments groups in phytoplankton forms the basis of all available multispectral phytoplankton fluorescence excitation instruments, and has been described in great detail e.g., [15,47]. In this study, the shift observed from algae to cyanobacteria dominance was clearly visible in $F_o(\text{GOR})$, and $F_o(\text{GO})$ which tracked variations of cyanobacteria abundance over time. In addition, spectra of $F_v(\lambda_{\text{ex}}, 685)$ and $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ provide insight into the photosynthetic light uptake potential of the two major phytoplankton groups. In spectra of $F_v(\lambda_{\text{ex}}, 685)$, a shift occurred from shorter to longer excitation wavebands as the relative abundance of cyanobacteria increased. Similarly, spectra of $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ showed distinct responses from algae and cyanobacteria. This was observed with $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ obtained with the algae-optimised protocol at the blue part of the spectrum, following algae abundance. On the other hand, at red–orange wavebands, higher $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ values obtained with the cyanobacteria-optimised protocol tracked the abundance of cyanobacteria. These results illustrate the use of PSII excitation spectra to determine the photosynthetic response by either phytoplankton group during natural succession. The fact that the algae-optimised protocol showed low $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ in the red–orange part of the spectrum when algae were most abundant, while the cyanobacteria-optimised protocol induced higher $\sigma_{\text{PII}}(\lambda_{\text{ex}}, 685)$ at blue wavebands illustrates that both protocols induced signal in the other phytoplankton group. The former is explained by the presence of red light at 622 nm in the cyanobacteria-optimised protocol which will induce some chl_a absorption in both groups, while the latter points to blue

light absorption by chl_a in algae and cyanobacterial PSII chl_a. The parallel use of the two protocols ensures that there is sufficient signal to extrapolate the photosynthetic parameters, and that differences are associated with the phytoplankton group for which each protocol is optimised. During periods where neither group dominates, no major differences should be expected between the protocols.

4.2. Emission Ratio of 730 nm over 685 nm

The F_o emission ratio $F_o(\text{GOR}, 730/685)$ was positively and significantly correlated with the relative abundance of cyanobacteria in the community. $F_o(\text{B}, 730/685)$ was not significantly correlated but followed the overall trend and short-term variations of the relative abundance of cyanobacteria. As discussed above, the B and GOR excitation signals are not exclusively diagnostic to either group. It is, therefore, interesting to observe that the emission ratio is sensitive to cyanobacteria in either excitation protocol. These results confirm the potential of interpreting the ratio of 730 nm over 685 nm emission to identify cyanobacteria in the community suggested by [35], as well as the importance of the choice of the excitation wavebands. LEDs targeting the orange-to-red spectrum will favour PSII emission from cyanobacteria compared to algae, whilst avoiding the excitation of chl_a associated with PSI further enhances the fluorescence emission ratio, as indeed observed here. Nevertheless, we did not observe major variability in time in the relative abundance of cyanobacteria in the community, so these results should be interpreted with some caution. To further confirm the utility of $F_o(\text{GOR}, 730/685)$ to identify cyanobacteria, measurements should be repeated alongside group-specific biomass determinations at higher frequency. In this context, the light-acclimation and nutrient history of the samples would have to be considered as these can be expected to influence the ratio of PSI over PSII emission through the expression of accessory pigment as well as photophysiological efficiency. It is known that state transitions induce further variations of the PSII to PSI ratio [48]. In this study, samples were dark-adapted before measuring F_o at both emission wavebands so that state transitions should not influence the observed variability in $F_o(\text{GOR}, 730/685)$. Moreover, as described by [49], light conditions in which the cyanobacterium *Synechocystis sp.* PCC 6803 were grown as well as the excitation wavebands have an influence on the PSII:PSI ratio. Ref. [49] showed that the cyanobacterium grown under blue light had a lower PSI:PSII ratio than under orange or red light. Under blue excitation (440 nm), PSI:PSII ratios were higher than under orange excitation (590 nm) [49]. This phenomenon is explained by the fact that the phycobilisomes in cyanobacteria do not efficiently transfer energy from blue light absorption to PSII.

4.3. Photophysiological Characterisation

Differentiated trends between the photosynthetic parameters highlight changes in the physiology of phytoplankton in the lake. The photochemistry efficiency expressed by F_v/F_m more than doubled over the observed period, likely corresponding with improved nutrient availability to the dominant group. F_v/F_m did not vary between the protocols, suggesting that algae and cyanobacteria populations adjusted similarly to environmental conditions (as also observed by [22]). In the case of the algal population, this included replacement of diatoms by chlorophytes during a period of silicate depletion.

P_{max} and F_o expectedly increased with overall biomass. σ_{PII} showed a decreasing trend over the sampling period which could be associated with seasonally increasing light availability. Ref. [50] also showed an inverse correlation between $F_v/F_m(478, \lambda_{\text{em}})$ and $\sigma_{\text{PII}}(478, \lambda_{\text{em}})$, which can be explained by energy requirements: the increasing numbers of pigment molecules in the light-harvesting antenna induce a higher probability of thermal dissipation and a decrease of photochemistry efficiency [50,51]. Moreover, the excitation spectra of σ_{PII} showed higher values of $\sigma_{\text{PII}}(560, \lambda_{\text{em}})$ than $\sigma_{\text{PII}}(478, \lambda_{\text{em}})$ in cyanobacteria while the opposite was observed in algae [50]. In our study, algae abundances were lower than cyanobacteria except at the beginning of the sampling period. We would, thus, expect to see higher $\sigma_{\text{PII}}(\text{GOR})$ than $\sigma_{\text{PII}}(\text{B})$ during periods of cyanobacteria dominance. Neverthe-

less, $\sigma_{\text{P}_{\text{II}}}(\text{B})$ and $\sigma_{\text{P}_{\text{II}}}(\text{GOR})$ followed the short-term variations in algae and cyanobacteria abundance, respectively. The fact that $\sigma_{\text{P}_{\text{II}}}(\text{B})$ was higher than $\sigma_{\text{P}_{\text{II}}}(\text{GOR})$ can be explained, either by the fact that $\sigma_{\text{P}_{\text{II}}}(\text{B})$ also targets cyanobacteria or that the ratio of chl_a pigments over phycobilipigments was low. This corroborates with the fact that phytoplankton growth was not light-limited and that cyanobacteria regulate their pigment expression [9]. Moreover, $\sigma_{\text{P}_{\text{II}}}$ is calculated as the product of $\alpha_{\text{P}_{\text{II}}}$ and EST which is the photon irradiance provided to the sample by the LEDs during a ST pulse ($\sigma_{\text{P}_{\text{II}}} = \alpha_{\text{P}_{\text{II}}} \times 100 \text{ EST}$, see [37]). Our results suggest lower $\alpha_{\text{P}_{\text{II}}}$ and EST values induced by the GOR protocol compared to the B protocol, which can explain the lower values of $\sigma_{\text{P}_{\text{II}}}(\text{GOR})$ compared to $\sigma_{\text{P}_{\text{II}}}(\text{B})$. The interpretation of $\sigma_{\text{P}_{\text{II}}}$ obtained with the GOR protocol and the B protocol is, therefore, not straightforward here.

PAM (Pulse Amplitude Modulation) and FFR (Fast Repetition Rate) Fluorometers using several induction protocols have been used to estimate the physiology and the biomass of distinct phytoplanktonic groups, including cyanobacteria. Ref. [23] used the FastOcean (FRRf) and a combination of LEDs to accurately estimate ETR_{PSII} of freshwater cyanobacteria. Ref. [52] could discriminate cyanobacteria from other phytoplanktonic groups using a PAM fluorometer and blue, green and red LEDs. Ref. [53] warned that ST fluorometers equipped solely with blue LEDs lack sensitivity to cyanobacteria without phycoerythrin. Some improvements to group-specific and bulk phytoplankton sensitivity are therefore expected from modern instruments such as used here, equipped with multiple excitation channels and broad-spectrum actinic light. In this work, P_{max} increased over the sampling period and was higher when obtained with GOR and GO protocols, explained by the increased abundance of cyanobacteria over time. This confirms the importance of using protocols targeting cyanobacteria and especially the GOR protocol (or the GO protocol if saturation can be improved) when trying to predict group-specific population growth. Short-term variations of P_{max} obtained with GOR and GO protocols followed cyanobacteria abundance over time while short-term variations in P_{max} obtained using the other excitation protocols followed algae abundance. The general increasing trend of P_{max} between the excitation protocols suggest that protocols other than GOR and GO also induced a signal in cyanobacteria. ETR generally increased up to mid-September and subsequently decreased with $\text{ETR}(\text{GOR})$ and $\text{ETR}(\text{GO})$ showing higher values than other protocols. $\text{ETR}(\text{GOR})$ followed trends driven by cyanobacteria abundance while $\text{ETR}(\text{B})$ trends followed abundance of algae. Moreover, $\text{ETR}(\text{B})$ showed a different trend from P_{max} , especially at the end of the sampling period. Wide variability in the ‘electron-to-carbon exchange rate’ was described by [54], which highlights the difficulty in linking ETR to C fixation. This is confirmed by the fact that this study focuses on a single lake over a relatively short period of time while nutrient concentrations and light availability remained relatively stable. It was explained by [55] that alternative electron sinks can occur for other metabolic processes and physiological mechanisms. Moreover, light and nutrient history effects on ETR values cannot be determined well from this data set due to the relatively long sampling intervals.

4.4. Suitability of LabSTAF for In Situ Assessment of Cyanobacteria

The configuration of the LabSTAF used in this study could not guarantee PSII saturation in the protocols we expected to be most diagnostically sensitive to cyanobacteria, which should be improved using higher intensity LEDs. Ideally, the R LED would only be used comparatively (i.e., combined with B or GO), to determine how well the GO protocol describes the cyanobacteria component of the community. A complete assessment would consist of the spectral excitation measurements providing insights into the efficiency of light uptake through key pigments in the community (providing the first insight into cyanobacteria presence) followed by GO and B fluorescence–excitation curves to determine photophysiological growth parameters which would indicate whether any differences in photosynthetic efficiency are likely between the two major phytoplankton groups.

Due to the nature of overlapping photosynthetic pigment absorption profiles between the major phytoplankton groups, it is essential to consider several optical markers, as confirmed by results presented here. Notably, we recommend using in parallel the ratio of $F_o(\text{GOR}, 730/685)$ and the PSII excitation protocols obtained with cyanobacteria and algae-optimised protocol. Ultimately, parameters derived from the FLCs obtained with GOR and B protocols can then be used to gain insight into likely growth and succession in the phytoplankton community. The DWM emission feature is promising because replacement filters could, in theory, be considered to look at other diagnostic wavebands for targeting cyanobacteria using this approach. The 650 nm emission waveband has been recommended by [35] to be implemented in a fluorometer targeting cyanobacteria. Although emission recorded at 650 nm could yield weak fluorescence compared to emission recorded at 685 nm, $F_o(\text{GOR}, 650/685)$ would likely give additional valuable diagnostic information on the presence of phycobilisome pigments.

5. Conclusions

This study assessed the ability of a new active fluorometer, the LabSTAF, to diagnostically assess the physiology of freshwater cyanobacteria in a reservoir exhibiting annual blooms.

The reservoir had a typical seasonal succession of phytoplankton of a temperate lake with diatoms being more abundant in spring and cyanobacteria more abundant in summer. PSII excitation spectra optimised for algae or for cyanobacteria showed distinct responses, illustrating the suitability of the instrument to determine the photosynthetic response by either group during their natural succession. FLCs parameters obtained with GOR and B protocols captured physiology parameters of, respectively, cyanobacteria and algae. However, the GO protocol, expected to be most diagnostically sensitive of cyanobacteria, did not reach saturation. Moreover, $F_o(\text{GOR}, 730/685)$ was significantly correlated with the relative abundance of cyanobacteria, which shows the potential of the ratio of 730 nm over 685 nm emission to identify cyanobacteria in a mixed phytoplankton community.

Due to the nature of overlapping photosynthetic pigment absorption profiles between the major phytoplankton groups, it is essential to consider several optical markers. We recommend using in parallel excitation spectra obtained with both protocols and $F_o(\text{GOR}, 730/685)$ to detect different pigment groups and assess cyanobacteria presence, followed by FLCs obtained with GOR and B protocols to assess the physiology and potential to grow of cyanobacteria and algae. Increased intensity of GO LEDs should be achieved to correctly assess the physiology of cyanobacteria and the R LED should only be used comparatively. Moreover, a 650 nm bandpass filter could be implemented in the DWM feature of the LabSTAF to measure the ratio of 650 nm over 685 nm emission, $F_o(\text{GOR}, 650/685)$. The potential of $F_o(\text{GOR}, 650/685)$ to be diagnostic of cyanobacteria should also be verified. According to this study, the LabSTAF is a good candidate to assess the presence and physiology of cyanobacteria in the natural environment.

This instrument is well suited for long-term continuous measurements and its optical features provided suitable optical markers to target cyanobacteria. Improvement or addition of some features are nevertheless recommended to increase the potential of the LabSTAF to be diagnostic of cyanobacteria.

Author Contributions: Conceptualisation, E.C., S.G.H.S., G.H.T., E.S., P.D.H. and K.O.; methodology, E.C. and S.G.H.S.; formal analysis, E.C.; investigation, E.C. and E.M.; resources, K.O.; data curation, E.C.; writing—original draft preparation, E.C.; writing—review and editing, S.G.H.S., G.H.T., E.S., P.D.H., E.M. and K.O.; visualisation, E.C.; supervision, S.G.H.S., G.H.T., E.S., P.D.H. and K.O.; project administration, S.G.H.S.; funding acquisition, S.G.H.S. and K.O. All authors have read and agreed to the published version of the manuscript.

Funding: This research was funded by Natural Environment Research Council, grant number 1983680 to E.C. and by European Union’s Horizon 2020 research and innovation programme, grant number 776480 to S.G.H.S. GHT was funded by European Regional Development Fund through the INTER-REG VA France-Channel-England program called ‘Sentinel-3 products for detecting EUtROphication and Harmful Algal Bloom events’ (Contract No. 106 S3-EUROHAB). The APC was funded by NERC.

Institutional Review Board Statement: Not applicable.

Informed Consent Statement: Not applicable.

Data Availability Statement: The data underlying this paper are held in a public repository under digital object identifier 10.5281/zenodo.7469162.

Acknowledgments: We thank E. Malcolm S. Woodward (Plymouth Marine Laboratory) for nutrient analysis. We thank the team of the National Meteorological Library and Archive of the UK Met Office for the meteorological data. We South West Water for nutrient and microscopy data, and the South West Lakes Trust for facilitating access to Roadford Lake.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript; or in the decision to publish the results.

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