RESEARCH ARTICLE

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Towards a quantitative assessment of inorganic carbon cycling in photosynthetic microorganisms

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Abstract

Photosynthetic organisms developed various strategies to mitigate high light stress. For instance, aquatic organisms are able to spend excessive energy by exchanging dissolved CO_2 (d CO_2) and bicarbonate (H CO_3^-) with the environment. Simultaneous uptake and excretion of the two carbon species is referred to as inorganic carbon cycling. Often, inorganic carbon cycling is indicated by displacements of the extracellular dCO₂ signal from the equilibrium value after changing the light conditions. In this work, we additionally use (i) the extracellular pH signal, which requires nonor weakly-buffered medium, and (ii) a dynamic model of carbonate chemistry in the aquatic environment to detect and quantitatively describe inorganic carbon cycling. Based on simulations and experiments in precisely controlled photobioreactors, we show that the magnitude of the observed dCO_2 displacement crucially depends on extracellular pH level and buffer concentration. Moreover, we find that the dCO₂ displacement can also be caused by simultaneous uptake of both dCO_2 and HCO_3^- (no inorganic carbon cycling). In a next step, the dynamic model of carbonate chemistry allows for a quantitative assessment of cellular dCO₂, HCO₃, and H⁺ exchange rates from the measured dCO_2 and pH signals. Limitations of the method are discussed.

KEYWORDS

carbonate chemistry, computational modeling, cyanobacteria, futile cycles, photosynthesis

1 | **INTRODUCTION**

Living organisms developed numerous adaptations to gain advantages in diverse environments. One branch of metabolic adjustments, widely spread throughout Bacteria, Plantae, Fungi, and Animalia kingdoms, is the evolution of so-called futile cycles. These cycles are part of regulation of the cellular energy status [1,2], reproduction processes [3], primary or secondary metabolism [4–7], redox components [8], signal transduction [9], and nutrient transport and assimilation pathways [10–14]. Despite their name, futile cycles have specifically addressed functions in cellular metabolism, and the term "futile" may be misleading. The common feature of these cycles, namely energy loss, is advantageous in many cases. In photosynthetic organisms, controlled energy dissipation is essential for cellular homeostasis. Without photoprotection

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Abbreviations: CA, carbonic anhydrase; dCO₂, dissolved carbon dioxide; DIC, dissolved inorganic carbon; dO₂, dissolved oxygen; ICC, inorganic carbon cycling; MIMS, membrane inlet mass spectrometry.

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mechanisms, sudden energy income can cause irreversible destruction of the photosynthetic apparatus. Photosynthetic organisms developed several strategies for the dissipation of excessive light energy such as adjustments of light-harvesting antennas, thermal energy dissipation, photo-protective adjustments of the electron transport chain, reactive oxygen species scavenging [15–20], and also light-dependent "futile" inorganic carbon cycling (ICC) between cells and their aquatic environment [2,21].

ICC, mediated through components of the carbon concentrating mechanism, can be defined as simultaneous uptake of dissolved carbon dioxide (dCO₂) and excretion of bicarbonate (HCO_{3}^{-}) , or vice versa simultaneous HCO_{3}^{-} uptake and dCO_2 efflux. In cyanobacteria, HCO_3^- can enter the cells via the transporters BCT1, SbtA and BicA, and dCO₂ by NDH- I_3 and NDH- I_4 [22–24]. In eukaryotic microalgae, dissolved inorganic carbon (DIC) transporters comprise HLA3, LCIA, possibly LCI1 and CCP1/2 (in Chlamydomonas reinhardtii), SLC4 and its homologues (in diatoms), and possibly also other transporters in other species [22]. Once inside the cell, dCO₂ is reduced to HCO₃ by carbonic anhydrase (CA) or CAlike enzymes, and HCO₃⁻ is transported for DIC assimilation by RuBisCo either to the carboxysome (in cyanobacteria), the pyrenoid (in some algae) or elsewhere in the chloroplast (in other algae). In algae, more pathways for DIC transport to chloroplast are known, e.g. the transport of intermediates of the C4-like metabolism [25]. During ICC, part of DIC also leaves the cell. The DIC efflux pathways are far less understood than DIC uptake pathways. CO_2 is known to diffuse out of the cell, either through membranes or pores, however, HCO_3^- excretion still remains an open question.

Although HCO₃ excretion transporters in cyanobacteria or algae have not yet been identified, it is assumed that HCO_3^- can be excreted through anion channels. HCO_3^- channels have been found in plants and have also been proposed for microalgae; for a recent review, see ref. [22]. Both constitutively transcribed transporters BTC1 and HLA3 require ATP (whereas the transporters SbtA and BicA, induced by low inorganic carbon, require only Na⁺ export, mediated by H⁺/Na⁺ antiporters such as NhaS3). Further, the conversion of dCO₂ to HCO₃, mediated by CA-like activity of NDH-I_{3/4} (where NDH-I₄ is also transcribed constitutively), requires reduction equivalents such as NADPH or ferredoxin [23]. Hence, a substantial amount of both energy and reductant equivalents formed under (excessive) light can be dissipated by ICC. A recent study also suggests that the CA-like activity of NDH-I (regulated through EcaB) prevents over-reduction of plastoquinone pool under high light [26].

ICC can be detected from the dCO_2 signal, measured by methods with high sensitivity and high temporal resolution, such as membrane inlet mass spectrometry (MIMS). A representative example of a Dark-Light-Dark experiment (to detect ICC) is shown in Figure 1; in the dark phases dCO_2 increases

PRACTICAL APPLICATION

Quantification of carbon fluxes in aquatic photoautotrophic communities is of significant importance for understanding and accurate assessment of cellular carbon partitioning and carbon economy. In this study, we present methods for a quantitative description of inorganic carbon cycling between photosynthetic microorganisms and their environment. The often overlooked phenomena of carbon cycling performed by some photosynthetic microorganisms can negatively influence the accuracy of the measurement of carbon-related properties of cells, e.g. their sequestration capacity, in dynamic culturing environments and can lead to incorrect bioenergetic calculations. Developed model and the technique for fast and accurate measurement of dissolved carbon dioxide provide a tool for identification and quantification of fast carbon fluxes that can be eventually used for smarter, model-based, design of biotechnological solutions.



FIGURE 1 Representative example of a Dark-Light-Dark experiment to detect inorganic carbon cycling from the dCO₂ signal, measured by a membrane inlet mass spectrometer. In the dark phases, dCO₂ fluxes are dominated by cellular respiration which causes a net increase of dCO₂, whereas in the light phase, dCO₂ fluxes are dominated by carbon fixation which results in net dCO₂ decrease. The abrupt dCO₂ displacement at the beginning and at the end of the light phase is typically explained as the effect of ICC (in this case as simultaneous dCO₂ uptake and HCO₃⁻ excretion), however, as discussed in the text, such a displacement can be observed even in the absence of ICC. The flat dCO₂ slopes represent the net effects of dCO₂ and HCO₃⁻ fluxes (between cells and the aquatic environment) close to equilibrium. Figure adapted from Tchernov et al., 2003 [2]

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mainly as a consequence of cellular respiration (although also carbon fixing processes can take place in the dark), whereas in the light phase dCO_2 decreases mainly as a consequence of DIC assimilation (although CO_2 producing processes such as respiration typically also take place at light). Most notably, the presence of ICC (in the case of Figure 1, simultaneous dCO_2 uptake and HCO_3^- excretion) causes an abrupt displacement of dCO_2 (a dynamics far from chemical equilibrium).

In this work, we show that in addition to the dCO_2 signal, also the pH signal can be used for ICC detection. For quantitative description of DIC exchange rates between photosynthetic cells and the aquatic environment, we developed a dynamic carbonate chemistry model. The model allows for detection of ICC activity and quantification of DIC reaction rates, and it extends previously published model [27] by considering dynamics far from equilibrium. The results show that the presence of ICC is not trivial to detect, in particular, that the dCO₂ displacement at the transitions between dark and light phases need not be caused by ICC. In fact, under high buffer concentrations such a displacement can be observed also in the absence of ICC. On the other hand, for low buffer concentrations and low pH (<7) it is not possible to detect ICC solely from the dCO_2 signal, but only in combination with the pH signal. We validated our model predictions experimentally, using MIMS in combination with well-controlled photobioreactors. For the assessment of ICC, we therefore recommend to measure dCO2 and pH simultaneously (with high accuracy and temporal resolution) and hence to use weak buffer concentrations in the cultivation media.

The model can be further used to quantitatively determine dCO_2 and HCO_3^- exchange rates between photoautotrophic cells and the aquatic environment from experimental data. Our unique combination of a highly controlled experimental setup with a mathematical model can be used as a tool for detection and quantification of ICC in photoautotrophic microorganisms.

2 | MATERIALS AND METHODS

2.1 | A dynamic model of carbonate chemistry in the aquatic environment

For the purpose of (i) a quantitative description of dCO_2 , HCO_3^- , and H^+ exchange rates between photoautotrophic cells and the aquatic environment and (ii) a quantitative assessment of ICC in photosynthetic microorganisms, we developed a dynamic model of carbonate chemistry and related processes. The model comprises the following variables (and constants):

[CO₂]: concentration of dCO₂ [HCO₃]: concentration of bicarbonate ions in Life Sciences



FIGURE 2 Reactions of carbonate species (together with water self-dissociation and buffer) in the aquatic environment and dCO₂, HCO₃⁻, and H⁺ exchange rates (called qCO₂, qHCO₃⁻, and qH⁺) between photoautotrophic cells and aquatic environment. The overall hydration rate constant k_1 is determined by the rate constants of the forward reactions 1', 1⁺, and 1⁻; see Equation 1. Reactions 1'', 2⁺, 2⁻, W, and B are considered as fast and equilibrium is assumed; see Table 1 for the equilibrium constants K_1 , K_2 , K_W , and K_B . Figure adapted from Nedbal et al., 2010 [27]

[H⁺]: concentration of protons

- CP: carbonate pool; see Equation 3d
- TA: total alkalinity; see Equation 3e
- *q*CO₂: CO₂ exchange rate between cells and the aquatic environment
- $qHCO_3^-$: HCO_3^- exchange rate between cells and the aquatic environment
- qH^+ : H⁺ exchange rate between cells and the aquatic environment
- v: net hydration rate of dCO_2 ; see Equation (2)
- $c_{\rm B}^{tot}$: concentration of buffer

The carbonate chemistry reactions are summarized in Figure 2, the corresponding rate and equilibrium constants are summarized in Table 1 and the resulting dynamic model is presented in Equations 3a–e. The model derivation and analysis is provided in the Supporting Information.

Reactions 1", 2⁺, 2⁻, W, and B of carbonate species, as shown in Figure 2, are considered as fast, and chemical equilibrium is assumed (as determined by the equilibrium constants K_1 , K_2 , K_W , and K_B as summarized in Table 1). Reactions 1', 1⁺, and 1⁻ are considered slow,

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TABLE 1 Rate and equilibrium constants, as measured in the cultivation medium BG-11, together with reference constants for sea water

		Value			Reference	
Constant	Comment	BG-11	Sea water	Unit	BG-11	Sea water
k_1	dCO ₂ hydration rate constant	0.04	0.04	[s ⁻¹]	Measured	[42]
K_1	Equilibrium constant of reaction $dCO_2 + H_2O \rightleftharpoons HCO_3^- + H^+$	$10^{-6.30}$	$10^{-5.86}$	$[mol L^{-1}]$	Measured	[42]
K_2	Equilibrium constant of reaction $HCO_3^- \Rightarrow CO_3^{2-} + H^+$	$10^{-9.00}$	$10^{-8.92}$	$[mol L^{-1}]$	Measured	[42]
$K_{\rm W}$	Water self-dissociation constant	$10^{-14.00}$	10-13.22	$[mol L^{-1}]$	[42]	[42]
$K_{\rm B}$	Equilibrium constant of HEPES buffer	$10^{-7.55}$		$[mol L^{-1}]$		

and the resulting overall hydration rate constant k_1 is given by:

$$k_{1} = k_{1'}^{-} + k_{1^{+}}^{-} + k_{1^{-}}^{-} \frac{K_{W}}{[H^{+}]}$$
(1)

As a consequence, the net hydration rate of dCO_2 (determined by the slow reactions 1', 1⁺, and 1⁻) can be written as:

$$v = k_1 \left([CO_2] - [HCO_3^-] \frac{[H^+]}{K_1} \right)$$
 (2)

After introducing the carbonate pool *CP* and total alkalinity *TA*, we can present our dynamic model for the five variables [CO₂], [HCO₃⁻], [H⁺], CP, TA (for given exchange rates qCO₂, qHCO₃⁻, and qH⁺). The differential algebraic equation system contains three ordinary differential equations and two algebraic equations:

$$\frac{d[CO_2]}{dt} = -v + qCO_2 \tag{3a}$$

$$\frac{\mathrm{dCP}}{\mathrm{d}t} = +v + q\mathrm{HCO}_3^- \tag{3b}$$

$$\frac{\mathrm{d}TA}{\mathrm{d}t} = +q\mathrm{HCO}_3^- - q\mathrm{H}^+ \tag{3c}$$

$$CP = [\text{HCO}_3^-] \left(1 + \frac{K_2}{[\text{H}^+]}\right)$$
(3d)

$$TA = [\text{HCO}_{3}^{-}] \left(1 + \frac{2K_{2}}{[\text{H}^{+}]} \right) - [\text{H}^{+}] + \frac{K_{\text{W}}}{\text{H}^{+}} + c_{\text{B}}^{tot} \frac{1}{1 + \frac{[\text{H}^{+}]}{K_{B}}}$$
(3e)

Further details including model derivation as well as model analysis are provided in the Supporting Information. For numerical solutions of the differential algebraic equation system (3), we used *Wolfram Mathematica 10* (Wolfram Research, Champaign, IL, USA) in particular, the function *NDSolve* with default options.

2.2 | Experimental setup (for validating the model predictions)

2.2.1 | Inoculum culture conditions

Model predictions were validated experimentally, using a cyanobacterium *Synechocystis* sp. PCC 6803. The strain was kindly provided by Dr. Martin Hagemann. The inoculum cultures were pre-cultivated in 250 mL Erlenmeyer flasks on air on a standard orbital shaker at 31°C, under 110 µmol(photons) m⁻² s⁻¹ of warm white LED light and in cultivation medium BG-11 [28] supplemented with 17 mM HEPES (Sigma-Aldrich, St. Louis, MO, USA). Prior to the experiments, cells from Erlenmeyer flask (c. 100 mL) were centrifuged (2000 *x g*, 10 min), supernatant was discarded, and pellet was washed two times with HEPES-free BG-11 medium, to ensure minimal buffer concentration in the culture media. The culture was then inoculated into the photobioreactor.

2.2.2 | Photobioreactor

For the validating experiments, we used a flat-panel photobioreactor described in detail in [29]. The bioreactor illumination was secured by red and blue LED panels with chess board LEDs configuration (red: $\lambda_{max} \approx 633$ nm, $\Delta \lambda_{1/2} \approx 20$ nm, Luxeon LXHL-PD09; blue: $\lambda_{max} \approx 445$ nm, $\Delta \lambda 1/2 \approx 20$ enm, Luxeon LXHL-PR09; both manufactured by Future Lighting Solutions, Montreal, QC, CA). The culture pH was continuously monitored by InPro3253 electrode (Mettler-Toledo, 1900 Polaris Parkway, Columbus, OH 43240, USA), the owndeveloped scripts allowed measurement of pH in temporal resolution of 1 s. Culture temperature was monitored by the same electrode and controlled by a Peltier cell incorporated in the instrument base. All other photobioreactor accessories and properties were such as described in Sinetova et al., 2012 [30].

2.2.3 | Gas exchange rate measurement

Dynamics in dCO_2 and dissolved oxygen (dO_2) exchange between cells and cultivation media was measured by MIMS, described in detail in Zavřel et al., 2016 [31]. Briefly, MIMS consisted of three main parts: (i) the mass spectrometer (PrismaPlusTM QMG 220 M1 with open ion source, Pfeiffer Vacuum, Asslar, Germany), (ii) silicon membrane at the system inlet (SILASTIC 508-006, Dow Corning, Midland, MI, USA), and (iii) tubing between the membrane inlet and the analyzer; stainless steel tubing connected with 90° or 180° stainless steel fittings (all manufactured by Swagelok, Solon, Ohio, USA) with U-shape water trap constantly tempered to -100 to -80°C for capturing water molecules crossing the silicone membrane, according to Tu et al., 1987 [32]. The MIMS was configured to detect m/z 32 (O₂), 44 (CO₂), and 40 (Ar).

2.2.4 | Carbonate system parametrization experiments

Equilibrium constants K_1 and K_2 of BG-11 cultivation medium at 30°C were determined using high-sensitivity Total Organic Carbon Analyzer (TOC-V_{CSH}, Shimadzu Scientific Instruments, Kyoto, Japan). All experiments for carbonate system parametrization were performed in a highly controlled environment of the photobioreactor to secure accurate and stable conditions, and the carbon analyzer was calibrated using certified premium range standards (TOC/TIC Standards, Reagecon Diagnostics, Shannon, Ireland).

First, to determine concentration of the dCO₂ in BG-11 medium, we adjusted pH of BG-11 medium to pH 4.0 by addition of 1 M hydrochloric acid and aerated the medium with 15 000 ppm CO₂ + air mixture (calibration cylinder 15 000 \pm 75 ppm of CO₂ in dry air, SIAD Czech, Rajhradice, Czechia) long enough to eliminate HCO₃⁻; the bicarbonate elimination was validated by sequential sampling for total inorganic carbon quantification in the time interval of several hours.

The equilibrium constant K_1 (relating dCO₂ and HCO₃⁻) was determined from a series of measurements in BG-11 medium for slightly acidic to neutral pH levels, adjusted by addition of 1 M hydrochloric acid, following similar approach as for determination of dCO₂; at each pH level the medium was aerated with 15 000 ppm CO₂ + air mixture and the total inorganic carbon content was measured when the carbonate system equilibrated.

Finally, the equilibrium constant K_2 (relating HCO₃⁻ and CO₃²⁻) was determined from a series of measurements in BG-11 medium under slightly alkaline pH levels, adjusted by addition of 1 M sodium hydroxide. Calculated equilibrium constants according to the carbonate system description (Figure 2 and Supporting Information Section 1.2) are summarized in Table 1.

2.2.5 | Validating experiments

The experiments validating the model predictions were performed in a photobioreactor cuvette. Cells in nonbuffered BG-11 cultivation media were inoculated into the cuvette, and kept at pH 8.5 and modest irradiance of 10 μ mol(photons) m⁻² s⁻¹ of both red and blue light for 24 h for acclimation. The initial cell density was approximately 3×10^7 cells mL⁻¹. Temperature was stabilized at 30°C during all experiments. in Life Sciences

The dynamics in dO₂, dCO₂, and pH during Dark-Light-Dark experiments was measured under pH levels 6.5, 7.3, 7.7, and 8.1. After 24 h of acclimation period, pH in the photobioreactor was decreased to 6.5 by addition of 1 M hydrochloric acid. dCO₂ formed from HCO_3^- pool was bubbled out by air until the dCO₂ concentration in the culture reached approximately 30 μ M. Then, simultaneous measurement of gas exchange and pH was initiated. Each measurement consisted of three steps: (i) initial measurement in dark, (ii) measurement of photosynthetic activity at light, and (iii) final measurement in dark. Each step took 4 min.

Between the measurements, culture homogenization was secured by air bubbling through the photobioreactor cuvette (flow rate 200 mL min⁻¹). During the measurements, bubbling was turned off and culture homogenization was secured by rotation of magnetic stirrer bar ($\phi 5 \times 35$ mm) in a vertical plane (500 rpm). To secure light saturated conditions for the cells [33], light intensity during each measurement was set to 150 μ mol(photons) m⁻² s⁻¹ of red light and 900 μ mol(photons) m⁻².s⁻¹ of blue light. Between the measurements, the intensity of both lights was reduced to $10 \,\mu\text{mol}(\text{photons}) \,\text{m}^{-2} \,\text{s}^{-1}$. Once each measurement was completed, pH of the culture was increased by addition of 1 M sodium hydroxide to the next pH level (6.5-7.3-7.7-8.1). After pH increase, culture was bubbled with air until dCO_2 level reached approximately 30 µM. Stabilization of pH took no longer than 45 min after each induced pH increase.

3 | RESULTS

3.1 | Model predictions

3.1.1 | Magnitude of the dCO₂ displacement during inorganic carbon cycling strongly depends on buffer concentration and pH level

Using the dynamic model of carbonate chemistry, we can simulate dCO_2 and pH signals dynamics under diverse conditions. By specifying the exchange rates between cells and their aquatic environment (qCO_2 , $qHCO_3^-$, and qH^+), both dark respiration and light-dependent carbon fixation (as a sum of all dCO_2 , HCO_3^- , and H^+ fluxes between cells and their aquatic environment) can be simulated together to abrupt dCO_2 displacements at the beginning and at the end of the light phase as caused by inorganic carbon cycling (Figure 1).

First, we studied the visibility of both dCO₂ and pH displacements under standard laboratory conditions. In previous publications related to ICC, strongly buffered cultivation media with slightly alkaline pH was used [2,34]. In our simulations, we considered two scenarios: a highly buffered system ($c_{\rm B}^{tot} = 17 \text{ mM}$ [21,33,35]), but also a weakly buffered system ($c_{\rm B}^{tot} = 1 \mu$ M). In dark, we assumed excretion of both dCO₂ and HCO₃⁻ from the cells to the cultivation media due

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to dark respiration. Under light, we assumed carbon fixation together with ICC, where dCO_2 is assimilated and HCO_3^- is excreted from the cells to the cultivation medium. To preserve charge balance, we assumed $qH^+ = qHCO_3^-$, which results in constant total alkalinity *TA*, see Equation 3c.

More specifically, we simulated dynamics in dCO_2 and pH signals for pH levels of 7, 7.5, and 8 and under low and high buffer concentrations. Under light, the exchange rate qCO₂ was set to -150 nmol L⁻¹ s⁻¹ (CO₂ uptake), whereas $qHCO_3^-$ was set to +100 nmol L⁻¹ s⁻¹ (HCO₃⁻ excretion). The assumed magnitudes of the exchange rates qCO_2 and $qHCO_{2}^{-}$ are based on the previously published experimental data [36–38]. Results of the simulations show that with low buffer concentration ($c_{\rm B}^{tot} = 1 \ \mu M$), the magnitude of the dCO₂ displacement decreases with decreasing pH (from 8 to 7), although identical exchange rates qCO_2 and $qHCO_2^-$ are assumed (Figure 3A-C). Under low buffer, the simulations also predict an increasing pH displacement with decreasing pH level, as a consequence of decreasing HCO_{2}^{-} concentration in the aquatic environment under decreasing pH (Figure 3A-C). On the other hand, under high buffer ($c_{\rm B}^{tot} = 17$ mM), the magnitude of the dCO₂ displacement remains almost constant (in the pH range 7-8), and the pH signal is shifted only negligibly (Figure 3D-F).

Model predictions of dCO_2 signal dynamics under exchange rates qCO_2 and $qHCO_3^-$ ranging between -150 to +100 nmol L⁻¹ s⁻¹ (simulating uptake or excretion of both dCO_2 and HCO_3^-) are summarized in Supporting Information Figure S1.

3.1.2 | dCO₂ displacement need not be caused by inorganic carbon cycling in strongly buffered systems

Figure 3 demonstrates the effect of pH and buffer on the visibility of dCO₂ displacement in the presence of ICC. Figure 4 shows that with high buffer and initial pH 8.0 a visible dCO_2 displacement can be observed even in the absence of ICC — in the case of sole carbon fixation where both dCO_2 and $HCO_3^$ are taken up by the cells. This effect can be explained by analyzing the dynamics of dCO_2 hydration which brings the system back (close) to equilibrium. As the model analysis shows (see Supporting Information Section 1.4), the net dCO₂ hydration rate v (Equation 2) does not directly follow the hydration rate constant k_1 (Equation 1), but the "apparent" dCO₂ hydration rate defined as $k_1(1+\alpha)$; see Supporting Information Equation 4. The parameter $\alpha \ge 0$ depends on buffer concentration and pH in a complicated way (Supporting Information Section 1.4 and Figure S2). The highest value of α (with $\alpha \gg 1$) and thus the fastest hydration dynamics is achieved at zero buffer concentration and pH 6.3 (equal to pK_1). The lowest value $\alpha \approx 0$ and thus the slowest dynamics occurs for high buffer concentration and alkaline pH. Only then, the net hydration rate *v* follows the hydration rate constant k_1 , and the slow dynamics of the system returning back (close) to equilibrium is visible as a dCO₂ displacement. The dynamics of the dCO₂ hydration rate *v* during Dark-Light-Dark experiments (caused by shifts of dCO₂, HCO₃⁻, and H⁺ exchange rates; see Supporting Information Equations 2 and 5 for further details) under pH 7.0 and 8.0 as well as under low and high buffer is summarized in Supporting Information Figure S3.

As a consequence, under high buffer, it can be hard or even impossible to distinguish between sole carbon fixation and carbon fixation with ICC, when only measuring dCO_2 . In Figure 4, we set qCO_2 and $qHCO_3^-$ such that the resulting DIC fixation rate was $-(qCO_2 + qHCO_3^-) =$ $-50 \text{ nmol } \text{L}^{-1} \text{ s}^{-1}$, both in absence and presence of ICC. In Figure 5, we show that by choosing different DIC fixation rates in absence and presence of ICC, almost identical dCO_2 dynamics can be obtained. Indeed, the particular choice of exchange rates qCO_2 and $qHCO_3^-$ as used in Figure 5 is only one possible combination out of many.

The model can also be used to simulate the effect of various buffers (with different pKa values) on the visibility of the dCO_2 displacement. In the simulations summarized in Supporting Information Figure S4, we considered buffers with pKa ranging from 5.59 (piperazine) to 10.51 (CAPS). In the presence of ICC, the dCO_2 displacement is almost identical for all buffers, whereas in the absence of ICC, the dCO_2 displacement decreases with an increasing difference between the pKa value of a particular buffer from the initial pH value of 8 considered in the simulation.

Based on the results of the simulations, we conclude that optimal conditions for the detection of ICC as well as for identifying the exchange rates qCO_2 and $qHCO_3^-$ from experimental data (see Section 3.5), weak pH buffer and simultaneous monitoring of the dCO_2 and pH signals dynamics is required. In the next section, we present a case study of such an experimental system where a precisely controlled photobioreactor is coupled with a membrane inlet mass spectrometer.

3.2 | Experimental validation of the model predictions

For the validation of our numerical simulations of the dCO_2 and pH displacements under varying pH levels and buffer concentrations, we performed a series of Dark-Light-Dark experiments. High magnitudes of the dCO_2 displacement under high buffer concentration are well documented in the literature [2,21,34,36,39,40]. Therefore, in this work, we focused on the experimental validation of ICC visibility in more natural, weakly buffered systems. To be consistent with the model predictions as summarized in Figure 3A–C, we performed a series of measurements under initial pH values of 6.5–8.1, and we monitored the dCO_2 and pH dynamics in a *Synechocystis* sp. PCC 6803 culture during Dark-Light-Dark experiments.

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FIGURE 3 Simulations of dCO₂ (blue lines) and pH (black lines) signals dynamics in the presence of inorganic carbon cycling, for initial pH 7.0 (left panels A, D), 7.5 (middle panels B, E), and 8.0 (right panels C, F), with buffer (HEPES) concentration of 1 µM (upper panels A, B, C) and 17 mM (lower panels D, E, F) during Dark-Light-Dark experiments. Dark phases (gray rectangles): exchange rates qCO_2 and $qHCO_3^-$ both set to +10 nmol L⁻¹ s⁻¹ (DIC excretion). Light phases: exchange rates set to $-150 \text{ nmol } L^{-1} \text{ s}^{-1}$ for qCO_2 (uptake) and $+100 \text{ nmol } L^{-1} \text{ s}^{-1}$ for $qHCO_3^{-1}$ (excretion). Initial dCO2 concentration set to 30 µM

To be consistent with the simulations, the initial dCO_2 concentration during the experiments was set to 30 µM. Additionally, all simulations shown in Figures 3-5 were performed to predict dCO₂ and pH signals for "CO₂ users" — organisms taking up a higher fraction of dCO_2 compared to HCO_3^- . The measured dCO₂ dynamics in Synechocystis during our experiments also resembles dynamics typical for "CO₂ users" [2]. However, since we used low CO₂ concentration (air), it is likely that all DIC transporters were active [41] and therefore Synechocystis was most likely able to uptake both dCO_2 and HCO₃. The results of the experiments are summarized in Figure 6 (left panels), and the results agree with the model predictions; under decreasing pH, the dCO₂ displacement became less visible and, on the contrary, the pH displacement became more pronounced. In particular, the pH displacement was the most visible at pH 6.5 ($\Delta pH = 0.051$, corresponding to $\Delta H^+ = 40.9 \text{ nM}$), less visible at pH 7.3 ($\Delta pH = 0.042$, corresponding with $\Delta H^+ = 4.8$ nM), even less visible at pH 7.7 $(\Delta pH = 0.036, \text{ corresponding with } \Delta H^+ = 1.8 \text{ nM})$, and the least visible at pH 8.1 ($\Delta pH = 0.016$, corresponding with $\Delta H^+ = 0.3$ nM). It is important to note that this agreement between simulations and experiments was achieved although the rates qCO_2 and $qHCO_3^-$ as identified from the experiments (summarized in the right panels of Figure 6 and further described in the next section 3.3) differed from rates considered in the simulations. There, we considered qCO_2 and $qHCO_3^-$ rates as constant, whereas in the experiments both rates were identified as dynamic (Figure 6, right panels). The net dO₂ evolution (as a sum of all fluxes between Synechocystis cells and the aquatic environment under light) was almost



FIGURE 4 Simulations of dCO₂ (blue lines) and pH (black lines) signal dynamics during Dark-Light-Dark experiments in absence (left panels A, C) and presence (right panels B, D) of inorganic carbon cycling (ICC) at initial pH 8.0, with HEPES concentration of 1 μ M (dashed lines) and 17 mM (full lines). Dark phases (gray rectangles): exchange rates qCO_2 and $qHCO_3^-$ both set to +10 nmol L⁻¹ s⁻¹. Light phases (white rectangles): in absence of ICC, exchange rates set to -50 nmol L⁻¹ s⁻¹ for qCO_2 (uptake) and 0 nmol L⁻¹ s⁻¹ for $qHCO_3^-$; in presence of ICC, exchange rates set to -150 nmol L⁻¹ s⁻¹ for qCO_2 (uptake) and +100 nmol L⁻¹ s⁻¹ for $qHCO_3^-$ (excretion). Initial dCO₂ concentration set to 30 μ M

identical for all pH levels (6.5–8.1), which indicates a negligible effect of the pH level on the light-dependent photosynthetic reactions (see Supporting Information Figures S6 and S7).

3.3 | Identifying exchange rates *q*CO₂ and *q*HCO₃⁻ from experimental data

The mathematical model can also be used for identifying the exchange rates qCO_2 and $qHCO_3^-$ (and qH^+) from the measured dCO₂ and pH signals. Mathematically, the identification of three time series from two noisy time series is an ill-posed problem. The development of a numerically stable identification algorithm is beyond the scope of this work. As a first step, we assume equal HCO_3^- and H^+ exchange rates (and hence constant total alkalinity) and implement a simple algorithm that involves data differentiation; see Supporting Information Subsection 1.5. This approach works for sufficiently high S/N; in our case, for low pH values and low buffer concentrations. The measured data (Figure 6 left panels) were used for identifying the exchange rates qCO_2 and $qHCO_2^-$; the results are summarized in Figure 6, right panels. As it turned out, our simple algorithm is problematic only for a high pH of 8.1.

Inorganic carbon cycling (simultaneous dCO_2 uptake and HCO_3^- excretion) was identified to be both qualitatively and quantitatively different for initial pH levels of 6.5–8.1. For

initial pH 6.5 (Figure 6A–B), we identified ICC (with decreasing magnitude) throughout the whole light period and partially also during the dark period after light (Figure 6B). For initial pH 7.3 (Figure 6C–D), the exchange rate qCO_2 was comparable to pH 6.5, whereas a significant exchange rate $qHCO_3^-$ was present only for the first 90 s of the light phase (Figure 6D). For initial pH 7.7 (Figure 6E–F), massive ICC was identified for the first 20 s of and sole carbon fixation for the rest of the light phase. For initial pH 8.1 (Figure 6G–H), both qCO_2 and $qHCO_3^-$ were qualitatively identical as for pH 7.7, however, the $qHCO_3^-$ magnitude was higher than for pH 7.7. Our results suggest that the magnitude of ICC can vary significantly over time and in fact, ICC may be present only at the beginning of the light phase.

4 | **DISCUSSION**

4.1 | Detection of inorganic carbon cycling

In the present work, we developed methods for a quantitative description of inorganic carbon cycling (ICC) between photosynthetic microorganisms and their aquatic environment. In particular, (i) we developed a mathematical model that quantitatively describes dCO_2 and HCO_3^- fluxes (qCO_2 and $qHCO_3^-$) between cells and the environment, (ii) we predicted the visibility of ICC under various buffer concentrations and



FIGURE 5 Simulations of similar dCO₂ (blue lines) and pH (black lines) signals dynamics during Dark-Light-Dark experiments in absence of inorganic carbon cycling (ICC, dashed lines) and presence of ICC (full lines), with initial pH set to 8.1, initial dCO₂ concentration set to 30 μ M, and with buffer (HEPES) concentration set to 17 mM. Dark phases (gray rectangles): exchange rates set to +2 nmol L⁻¹ s⁻¹ for *q*CO₂ and +10 nmol L⁻¹ s⁻¹ for *q*HCO₃⁻. Light phases: in absence of ICC (fixation only), exchange rates set to -40 nmol L⁻¹ s⁻¹ for *q*CO₂ and -10 nmol L⁻¹ s⁻¹ for *q*HCO₃⁻ (uptake); in presence of ICC, exchange rates set to -45 nmol L⁻¹ s⁻¹ for *q*CO₂ (uptake) and +15 nmol L⁻¹ s⁻¹ for *q*HCO₃⁻ (excretion)

pH levels, and (iii) we experimentally validated the model predictions, using the combination of a precisely controlled photobioreactor and a high resolution membrane inlet mass spectrometer. In both simulations and experiments, we focused on slightly acidic to slightly alkaline pH range, since first, with pH increase, the pH displacement becomes hardly detectable even in non-buffered systems due to (self-) buffering capacity of the carbonate system, Figure 3A–C, Figure 6), and second, at pH < 6, dCO₂ uptake can become significantly reduced or even inhibited [34].

Although ICC has been first described already in 1998 [21], available methods used for ICC detection still rely on experimental evaluations of the dCO_2 displacement at the beginning and at the end of the light phase (Figure 1). However, as we show in the simulations, in highly-buffered cultivation media, such dCO_2 displacement can be observed even in the absence of ICC (Figures 4 and 5). On the other

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hand, in weakly buffered systems, dCO_2 displacements are a clear sign of ICC. Quantitatively, we show that the magnitude of dCO_2 displacements induced by ICC decreases between pH 8 and 7 (Figure 3A–C, Figure 6); and at pH \leq 7, a dCO_2 displacement cannot be detected any more (Figure 3A–C).

The model simulations further show that, with low buffer concentrations, it is possible to detect ICC from pH displacements (Figure 3A–C). Similarly to dCO₂ displacements, also pH displacements result from abrupt changes of the exchange rates qCO_2 , $qHCO_3^-$, and qH^+ . Thereby, the displacements occur with a certain delay corresponding to the "apparent" hydration rate constant $k_1(1+\alpha)$; see Supporting Information Equation 4. For pH = 7, when the hydration rate v shows a very short delay compared to pH = 8 (Supporting Information Figure S3), simulations (Figure 3A) as well as experimental data (Figure 6) show only slightly delayed pH displacements. Since the "acceleration" parameter α decreases with increasing pH between 7-8 (Supporting Information Figure S3), the simulations predict that the corresponding delay of the pH displacement increases with increasing pH (in the range 7-8, Figure 3A-C). In the experiments, however, the exchange rates qCO_2 , $qHCO_3^-$ (and qH^+) identified from the experimental data (Figure 6) showed some delay (at the beginning of the light phase) under all tested pH levels, which indicates that the assumption of abrupt change used in the simulations may not be justified.

A further analysis showed that the magnitude of the pH displacement during ICC decreases with increasing pH in the range of 7–8 (Figure 3A–C, Figure 6), which is a consequence of the increased (self-) buffering capacity of the carbonate system as well as water self-dissociation; with pH increasing from 7 to 8, the concentrations of OH⁻ and HCO⁻₃ (in chemical equilibrium) increase 10 times, and protons that are excreted from the cells (pressumably together with HCO^-_3) react with the both anions to a higher extent which leads to a reduced amount of "free" protons available for pH displacement.

As discussed above, the dCO_2 displacement can be detected only when the dCO₂ hydration rate constant k_1 (part of the "apparent" hydration rate constant k_1 (1+ α)) is small enough — in particular, as shown in Supporting Information Figure S5, less than 1 s⁻¹. In this work, we experimentally determined the value of k_1 as 0.04 s⁻¹ (Table 1), which has been also reported for sea water [42]. Some photosynthetic microorganisms contain extracellular carbonic anhydrase (CA) [43], an enzyme that can increase k_1 significantly. The turnover number k_{cat} of CA is usually reported in the range of 10^4 – 10^6 s⁻¹, that is 10^6 – 10^8 times higher than k_1 . As shown in Supporting Information Figure S5, already a k_1 value of 0.08 s⁻¹ reduces the dCO₂ displacement by one half, and at $k_1 \ge 4$, the dCO₂ displacement cannot be detected anymore. For ICC detection in strains with external CA, it is therefore extremely important to ensure complete CA



FIGURE 6 Experimental data of *Synechocystis* sp. PCC 6803 cultivated in non-buffered BG-11 medium during Dark-Light-Dark cycles: measurements of dCO₂ (full blue lines) and pH signals (full black lines) signals (left panels) for initial pH 6.5 (panel A), 7.3 (panel C), 7.7 (panel E), and 8.1 (panel G). Four biological replicates with qualitatively identical results for each condition were collected; data from one representative experiment are shown. Identification of the exchange rates qCO_2 (dotted blue lines, right panels) and $qHCO_3^-$ (dotted black lines, right panels), identified from the measured data involving data differentiation and the assumption of $qH^+ = qHCO_3^-$ is represented in right panels for initial pH 6.5 (panel B), 7.3 (panel D), 7.7 (panel F), and 8.1 (panel H). Positive values in the right panels indicate dCO_2 or HCO_3^- excretion, negative values indicate dCO_2 or HCO_3^- uptake. For further experimental details, see Section 2

inhibition since even a minimal CA activity can prevent the ICC visibility.

4.2 | Assumptions and limits of the presented carbonate chemistry model

Our mathematical model is based on several assumptions regarding carbonate chemistry such as fast equilibrium for reactions W, B, 1', and $2^+/2^-$ (see Figure 2 and Supporting Information Section 1.3 for further details). To preserve constant total alkalinity and charge balance, we also assume co-transport of HCO₃⁻ and H⁺ (q HCO₃⁻ = qH⁺). The current understanding of proton efflux in cyanobacteria is fully

consistent with this assumption. Even though current carbon concentrating mechanism models consider two HCO₃⁻ transporters (SbtA and BicA) that require extracellular Na⁺ which is exchanged with H⁺ by the antiporter NhaS3, the photosynthetic and respiratory models consider at least three complexes in the cytoplasmic membrane that all work as proton (efflux) pumps: NDH_{1/2}, COX, and P-ATPase [44]. Indeed, proton efflux channels are present also in eukaryotic algae [24,45]. There is also direct evidence for HCO₃⁻ and H⁺ cotransport in some strains [46]. Clearly, the assumption of equal integrals of $qHCO_3^-$ and qH^+ over time is reasonable; otherwise the cells would become acidified or basified over time. However, eventual fast fluctuations in both $qHCO_3^-$ and qH^+ (on the time scale of seconds) can result in temporal $qHCO_3^-$ and qH^+ disbalance. This can lead to (temporally) incorrect qCO_2 and $qHCO_3^-$ rates identification from experimental data (see the following section for further discussion).

The model further assumes that both kinetic and equilibrium constants (as summarized in Table 1) are known for the given physical and chemical properties of the aquatic environment such as temperature, salinity, total alkalinity, pH, or pressure. Finally, only three fluxes between cells and their aquatic environment are considered in the model (qCO_2 , $qHCO_3^-$, and qH^+ , Figure 2), and fluxes of all other ions that can influence total alkalinity (such as Na⁺, K⁺, NO₃⁻⁻, NH₄⁺ [47,48]) are neglected. Despite these limitations, the model provides a valuable contribution towards understanding and quantitative evaluation of inorganic carbon cycling.

4.3 | Towards a reliable method for the identification of qCO_2 and $qHCO_3^-$

In addition to predicting the visibility of ICC, the model was further used to identify the exchange rates $qHCO_2^{-1}$ and qCO_2 from the experimental data. We assessed ICC of various durations and magnitudes in the pH range 6.5-8.1 (Figure 6, right column). The qCO_2 and $qHCO_3^-$ identification algorithm involved data differentiation which turned out to be appropriate for data measured at pH < 7.7. For pH = 8.1, this method amplified data noise in the pH signal, caused by an increasing (self-)buffering capacity of carbonate chemistry and hence a higher S/N with increasing pH. Noise reduction in the identified exchange rate $qHCO_3^-$ at higher pH can be achieved by advanced data fitting methods involving regularization techniques. However, without direct measurement of the HCO_3^- dynamics, the identification of qCO_2 and $qHCO_3^-$ fluxes may still depend on model assumptions such as $q \text{ HCO}_3^- = q \text{H}^+$ (see the previous section for further discussion).

Being aware of the model limitations outlined above, the exchange rates qCO_2 and $qHCO_3^-$ identified from experimental data can be interpreted only as rough guides of ICC in general. The first analysis of our experimental data showed that ICC could take place throughout the whole light period (Figure 6B), only partially during the light period (Figure 6D,F,H), and also partially in dark (Figure 6B), suggesting highly dynamic ICC and/or $qHCO_3^-$ and qH^+ co-transport.

The amount of ATP "burned" during ICC by the cells can be significant since transport of HCO₃⁻ against the concentration gradient requires ATP [24,25]. Giving an oversimplified example based on our experimental data, qHCO₃⁻ as identified during the initial displacement at light was as high as 40 μ M s⁻¹ (excretion, Figure 6 right column) which corresponds with 20 amol cell⁻¹ s⁻¹. Assuming (for simplicity) that each HCO₃⁻ molecule requires 1/3 ATP to be excreted from the cell (only one out of three total HCO₃⁻ transporters in Life Science

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require ATP), such a rate would burn 6.7 amol cell⁻¹ s⁻¹ ATP, that is, 35% of the ATP generated by photosynthetic light reactions: assuming 3 ATP produced per each O₂ molecule, the dO₂ production rate of 6.4 amol cell⁻¹ s⁻¹ (as derived from the data presented in Supporting Information Figures S6 and S7) would result in an ATP production rate of 19.2 amol $cell^{-1} s^{-1}$ (and 6.7/19.2 = 35%). However, we note that a precise accounting of ATP (and reducing equivalents) production/consumption is tricky since the stoichiometry of both dO₂:ATP production and HCO₃⁻ excretion:ATP consumption can vary considerably over time. In addition, under high blue light as used in our experiments, one can expect a strong cyclic electron flow around photosystem I (PSI) that can shift the ratio of ATP:NADPH production significantly [49]. Considering higher ICC rates at the beginning of the light phase, compared to the rest of the light phase (Figure 6, right panels), our results suggest that ICC can serve as a temporal ATPburning mechanisms until other ATP-consuming processes such as inorganic carbon fixation become activated.

The photosynthetic quotient in our experiments (PQ, net O_2 release:CO₂ fixation) was ranging between 5.7–0.3 in the pH range 6.5–8.1. Such values represent a higher PQ variation than reported previously for *Synechocystis*. However, in previous works [50–52] light of lower intensity and different wavelength was used. As mentioned above, in *Synechocystis* blue light preferentially affects PSI, resulting in cyclic electron flow [49].

The model analysis also showed that the negative slope of the dCO_2 signal during the light phase does not reflect the true inorganic carbon fixation rate. In fact, it equals $\alpha/(1+\alpha)$ times the true inorganic carbon fixation rate $(qCO_2 + qHCO_2^-)$, see Supporting Information Equation 8. Thus, the dCO₂ signal slope depends on the pH level (Figure 3), which was also confirmed experimentally (Figure 6 left column, Supporting Information Figure S7). The "apparent" rate is closest to the true rate around pH = pK1 = 6.3 (for large α), and the difference between the "apparent" and true inorganic carbon fixation rates increases with increasing pH and buffer concentrations - following the dependence of the "acceleration" parameter α on both pH and buffer concentration (see Supporting Information Equation 4 and Figure S2). As shown in Supporting Information Figure S7, the "apparent" and true inorganic carbon fixation rate, as identified from our experimental data, can differ by a factor of 10 under the highest tested pH 8.1.

A reliable method for the dynamic identification of both exchange rates qCO_2 and $qHCO_3^-$, and thus for the assessment of ICC is still missing. This work summarizes the complexity and limitations of ICC detection related to the effects of carbonate chemistry and to the current state of the art of the experimental techniques. Until a method for the dynamic and quantitative measurement of HCO_3^- concentrations will be available, both qCO_2 and $qHCO_3^-$ quantification will depend

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on a mathematical analysis involving extra assumptions. In this work, we outlined future developments of both experimental and modeling methods as necessary for a reliable ICC detection.

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OPEN DATA

Source data for Figure 6 are available online here: https:// identifiers.org/ecyano.experiment:31. Mathematical model is available online here: https://identifiers.org/ecyano.model:46.

CONFLICT OF INTEREST

The authors have declared no conflict of interest.

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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