DYNAMIC INTERACTIONS IN AN ARTIFICIAL PHOTOTROPHIC BIOFILM FOR BIOTECHNOLOGICAL APPLICATIONS

Inaugural - Dissertation

zur

Erlangung des Doktorgrades

der Mathematisch-Naturwissenschaftlichen Fakultät

der Universität zu Köln

vorgelegt von

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Köln

2016

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Tag der letzten mündlichen Prüfung:

29.1.2016

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Acknowledgements

I would like first to thank my parents, who supported me in every aspect of my study. Thank you for being proud of me and for your trust in me! Thanks to my girlfriend for her support, for her understanding! Thanks to professor Michael Melkonian, for giving me the chance to do my Ph.D. study, for giving me great advices during the last 4 years and for being a great supervisor. Thanks to Dr. Björn Podola, who always gives me good advice and supported me with the production of all my publications. Thanks to Bastian Piltz for giving me advice for the manuscripts and for all his help in finishing this thesis. Thanks to Dr. Dirk de Beer and his workgroup for their advice and support in the microsensor measurements. And thanks to all the members of AG Melkonian for being the perfect colleges!

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1. Introduction

1.1. Microalgae and their biotechnological potential

"Algae" are defined as a polyphyletic group of oxygenic phototrophs that encompasses both prokaryotes and eukaryotes. The endosymbiotic theory hypothesizes that eukaryotic algae have originated from the endocytobiosis of cyanobacteria inside eukaryotic heterotrophic cells. Through several secondary endocytobiosis events, a variety of algal species have since evolved (Douglas 1998; Moreira and Philippe 2001; Whatley 1993). Algae can be divided into 10 monophyletic groups. Until now, there are 40,000 known algal species. However, it is estimated that the actual number is more than 10 million, and among these microalgae are the most numerous (Norton et al. 1996). This biodiversity of microalgae is the foundation of their biotechnological potentials. E.g., through various metabolic pathways, microalga can produce a wide range of substances, some of which can be synthesized chemically at high costs, whereas others cannot be synthesized at all (Cardozo et al. 2007; Shimizu 1996).

The utilization of microalga by human is not a novel concept: Arthrospira platensis (Spirulina platensis) has been used since centuries as food by people living in the tropical regions (Ciferri 1983). However, the controlled mass cultivation of microalgae was only introduced in the middle of the last century, and so far, of the known forty thousand microalgal species, only a fraction is used for biotechnological applications (Becker 1994; Moore 2001). Nevertheless, microalgae or substances derived from microalgae can be found today in a variety of different products, e.g. as human food and food supplements, as feed for aquaculture, as raw material for high value chemical compound production, or as additive in cosmetic and pharmaceutical products (Borowitzka et al. 1991; Muller-Feuga 2000). Also, microalga have been successfully applied in wastewater treatment processes (Abdel-Raouf et al. 2012; Davis et al. 2003; de la Noüe et al. 1992).

1.2. Microalgal cultivation system for biotechnological applications

The design of cultivation system is a major factor in the biotechnological application of microalga. Compared to heterotrophic biological systems for biotechnological applications (e.g. biomass and/or metabolites production, waste water treatment relying on bacteria or fungi), another factor has to be taken into consideration when designing bioreactors for microalgae: microalgal cells require light for photosynthesis. Hence, such bioreactors were named "photobioreactors". Photobioreactors can be divided into two major categories: suspension-based and biofilm-based systems (Gross et al. 2015; Olivieri et al. 2014).

In application, suspension system can be subdivided into open and closed system. In an open system, the suspension is directly exposed to the environment; typically, the suspension is kept in a pond-like structure (e.g. race way pond) and to achieve better light utilization and mass transfer efficiency, energy has to be put into the system to ensure proper light distribution and mass transfer (Olivieri et al. 2014). In a closed system, the suspension is isolated from the environment, thus minimizing the risk of contamination. However, high construction costs are typically associated with this type of photobioreactors, in order to achieve a sufficiently large illuminated surface area for optimal growth (Olivieri et al. 2014). Currently, commercial production of microalgal biomass relies heavily on suspension cultivation systems: E.g. open ponds for large scale microalgal biomass production and closed tubular (or of other forms) photobioreactors for more valuable products (Olivieri et al. 2014). Also, open 'high rate algal ponds' have been successfully applied to treat wastewater (Park et al. 2011). However, the cell densities in such suspended systems are typically low (e.g. 0.5 g L⁻¹ and 2 – 6 g L⁻¹ dry weight in open ponds and in photobioreactors, respectively; Davis et al. 2011), which results in low productivity (or efficiency in wastewater treatment) per volume and high cost of harvesting (i.e. separation of cells from culture medium).

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Type of	Suspens			
reactor	Open	Closed	- Biofilm-based	
Typical system setup	Suspension culture kept in a container open to the environment, stirring and aeration provided by mechanical mixing.	Suspension culture kept in an enclosed transparent container, stirring achieved by pumping the suspension through the system and/or aeration with gas bubbles.	Most of the algal cells are immobilized inside the biofilm, only the culture medium flows in the system. Mass transfer achieved mostly through diffusion.	
Prominent advantages	Simple construction (i.e. low construction cost).	Minimized risk of contamination. Controlled culture environment.	harvesting). Low operational cost (i.e. less energy required for pumping).	
Major drawbacks	Easily contaminated. High harvesting cost (i.e. dewatering).	High construction cost. High operational cost. High harvesting cost (i.e. dewatering).	Easily contaminated. Biofilm detachment.	

*Table I: Typical system design, major advantage and disadvantages of suspension-based and biofilm-based photobioreactors.

*a: Refer to the citations in section

A more recent development in photobioreactor design are the biofilm-based photobioreactors: typically, the algal cells are immobilized onto a substrate to form a biofilm, and the culture medium flows above the biofilm. Compared to suspension-based systems, such a system offers a much higher biomass density (10% - 20%) of the harvested biofilm is dry biomass, similar to post-centrifuge biomass from suspension cultures), and thus can reduce the harvesting cost (e.g. biomass dewatering) considerably. In addition, light transfer and mass transfer (especially for gases like CO₂) are more efficient in biofilm-based photobioreactors (Gross et al. 2015). As a result, various biofilm-based photobioreactors have been developed, and both bench-scale and pilot-scale experiments have already been carried out for biofilm-based systems both for biomass production and for wastewater treatment (Abdel-Raouf et al. 2012; Berner et al. 2015; de la Noüe et al. 1992; Olivieri et al. 2014).

Compared to suspension-based systems, biofilm-based photobioreactors offers several advantages. However, in a 'traditional' biofilm-based photobioreactor setup as described above, the biofilm remains completely or partially (spatially and/or temporally) submerged in culture suspension. Thus, such biofilms are subjected to shear forces due to the movement of the liquid phase above, which may cause detachment of the biofilm. Also, the liquid phase above the biofilm can lead to a fast spreading of contaminants, once a contaminating organism enters the system (Berner et al. 2015). The typical system design, advantage and disadvantages of suspension-based and biofilm-based photobioreactors as mentioned above are summarized in Tab. 1 (summarized from references cited previously).

1.3. Porous substrate biofilm photobioreactor (PSBR)

To overcome the problems in the existing photobioreactors, a novel biofilm-based photobioreactor was developed. The new system, developed first by Podola and Melkonian (Twin-Layer system; Podola and Melkonian 2003), and now known best as Porous Substrate Biofilm Photobioreactors (PSBRs), as defined by Murphy et al. (Murphy and Berberoglu 2014), eliminates the problems encountered in "traditional" biofilm-based systems by separating the bulk culture medium from the microalgal biomass. To achieve this, as inspired by membrane based bioreactors, in a PSBR, algal cells are immobilized on one side of a micro-porous membrane (contrary to a solid substrate in a "traditional" biofilm-based photobioreactor), and the culture medium is supplied from the other side of the micro-porous membrane by means of a medium-saturated macro-porous material or by simply applying the medium between two inoculated micro-porous membranes (Shi et al. 2007).





The schematic setup of a typical PSBR is illustrated in Fig. 1. The separation of algal cells and culture medium does not only eliminate the problem of biofilm detachment due to shear force, but also maximizes the gas transfer efficiency by directly exposing the surface of the biofilm to the ambient gas phase. In a PSBR, the nutrients are supplied to the biomass by diffusion through the micro-porous membrane (with pore size small enough to prevent algal cells from penetrating). The elimination of a flowing liquid phase around the algal cells also minimizes the spread of contamination, and the removal of a liquid phase (which could contain suspended solids such as detached algal cells) above the algal biofilm maximizes the light transfer efficiency. Furthermore, this absence of liquid phase above the biofilm further increases the dry biomass concentration in the algal biofilm (up to 40%, *Symbiodinium sp.* cultivated on Twin-Layer PSBR, unpublished data). To illustrate this, a photograph of a PSBR



Figure 2: 100x magnified Halochlorella rubescens PSBR biofilm cultivated on polycarbonate filter disk.

Previous studies have shown that PSBRs can have very high footprint productivities (productivity per ground area) compared to other cultivation systems under comparable conditions, as presented in Tab. II (Berner et al. 2015; Liu et al. 2013; Olivieri et al. 2014; Schultze et al. 2015). Also, studies carried out at both bench- and pilot-scales have proven the potential application of such PSBR in wastewater treatments (Li et al. 2015a; Shi et al. 2007; Shi et al. 2014). The data presented in these studies demonstrated clearly the potential of a viable large-scale PSBR for biomass production and/or for other applications (e.g. wastewater treatment). However, to transform a bench-scale system into a commercially viable system, several important advancements have to be made. Among these, an automated control system for growth optimization and/or an optimal growth strategy (e.g. light, CO₂ and nutrient supply, harvesting circle etc.) have to be developed. To achieve this, a comprehensive understanding of the dynamic processes inside such PSBR is essential.

Algal strain	Target pollutant	Operational mode	Removal efficiency	Source
Halochlorella rubescens isolated from wastewater	Nitrate Phosphate Ammonia	Batch	 b>85% per day b>70% per day b>70% per day 	Shi et al., 2014
Stichococcus bacillaris isolated from Zn contaminated	Zn	Batch	95% in 2 days	Li et al., 2015
stream		Continuous	35%	<i>,</i>

*a: Efficiency calculated as one minus concentration of targeted pollutant after treatment divided by that in the original untreated sample. b: In their study, different batches of wastewater samples were used.

Type of photobioreactor		Algal strain	Light intensity (µmol photons m ⁻² s ⁻¹) and ^b L/D	Growth medium	CO ₂ supple -ment (v/v)	"Productivity (g m ⁻² d ⁻¹)	Source	
Porous	Twin-Layer PSBR	Halochlorella rubescens	ver Halochlorella rubescens	1000, 14:10	Synthetic medium	/	51	Schultze et al., 2015
substrate biofilm photobioreact or (PSBR)	Attached cultivation PSBR	Scenedesmus obliquus	100, continuous	Synthetic medium	1%	80	Liu et al., 2013	
	Evaporation driven PSBR	Anabaena variabilis	110, continuous	Synthetic medium	/	°32	Murphy et al., 2014	
Submerged biofilm photobioreact or	Biofilm flow cell	Wastewater consortium	230, continuous	Synthetic wastewat er	/	°25	Boelee et al., 2011	
	Algal turf scrubber	Natural consortium	40-140, 16:8	Synthetic	/	42	Mulbry et al., 2008	
Closed suspension	Tubular	Spirulina sp.	80, continuous	Synthetic medium	/	°8.4	Converti et al., 2006	
	Flat-plate photobioreactor	Chlorella sp.	Natural sunlight	Synthetic medium	/	25	Carkozzi et al., 2000	
Open suspension	Open pond	Scenedesmus sp.	Natural sunlight	Synthetic medium	/	۶۹	Ketheesan and Nirmalakhandan, 2012	
	Raceway pond	Nannochloro psis salina	Natural sunlight	Synthetic medium	/	3.5	Crowe et al., 2012	

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*a: Productivity per footprint area (i.e. ground area); b: Light dark circle; c: Calculated from original data by the author; d: For further data on productivity of different photobioreactors, refer to citations in section 1.3.

1.4. Investigating dynamic processes inside PSBR biofilms

Unlike for bacterial biofilms or submerged phototrophic biofilms, due to its novelty, the processes inside PSBR biofilms have not yet been studied extensively. There are, until now, only few studies investigating processes inside PSBR biofilms: Murphy and Bergeroglu (2014) developed a model for a PSBR biofilm. However, the model focused more on the gradients (e.g. nutrients, dissolved inorganic carbon and growth etc.) along the medium flow direction rather than the gradients perpendicular to the medium flow (i.e. along the biofilm depth gradient) in an evaporation driven (instead of gravity) PSBR. Furthermore, an over-simplified model (i.e. effect of scattering was not considered in their model) was used to predict light distribution (discussed in the 3rd manuscript, Li et al. 2015b, presented in this thesis in detail), even though previous studies suggested otherwise (Berberoglu and Pilon 2007; Berberoglu et al. 2007; Jøsrgensen 1969; Richardson et al. 1983). In addition, no extensive experimental measurements were carried out to verify their proposed model. In another study, Wang et al. (2015) used a rather primitive method to investigate the effect of light on biomass productivity in PSBR biofilms with different thickness. They speculated that the respiration in the 'non-illuminated' zones in the biofilm caused the decrease of total biomass productivity in prolonged cultivation of PSBR biofilms. However, in their study, instead of a grown biofilm, cells were filtered onto micro-porous membranes to simulate PSBR biofilms grown to different thicknesses. Thus, the effects of adaptation of the biomass due to immobilization (e.g. cellular pigment content), as suggested by previous observations from other researchers (Jøsrgensen 1969; Richardson et al. 1983) and from observations made by the author (Fig. 3), were excluded in their study.

Schultz et al. (2015) observed an increased biomass productivity in PSBR biofilm with increasing surface light intensity up to a photosynthetically active radiation (PAR) intensity of about 1000 umol photon $m^{-2} s^{-1}$. This light intensity was well above the saturation PAR intensity expected in suspension cultures. This phenomenon has also been observed in other studies (Liu et al. 2013), and this is suspected to be caused by the following hypothesis: Higher light intensity on the biofilm surface led to deeper light penetration into the PSBR biofilm, thus increasing the total biomass productivity, even though the growth of the surface cells were inhibited by the strong PAR. However, until now, no experimental studies have been conducted to verify this hypothesis.

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Figure 3: Photo from a 200x magnified sectioned Halochlorella rubescens PSBR biofilm cultivated on polycarbonate filter disk (embedded in Epon and sectioned to 10 µm thick section along the direction of the depth gradient). The decoloration observed in the photo indicate the cells closer to the biofilm surface contain less piements.

In their study, Wang et al. (2015) have also observed a higher light dilution rate (light availability per cell in total biofilm/suspension volume) in a PSBR biofilm compared to a suspension with the same amount of biomass density per unit area. This can be translated to a more effective light transfer in the PSBR biofilm compared to that in suspensions. It has been observed in previous investigations, that in a very dense suspension culture, the forward-scattering of light can significantly increase the efficiency of light transfer (i.e. light dilution rate). Models have already been developed to describe this phenomenon, both in suspension and in phototrophic biofilms (Berberoglu et al. 2007; Murphy and Berberoglu 2014). However, until now, no experimental study has been carried out to verify the light distribution in PSBR biofilms as predicted by the model.

Moreover, as in a PSBR the nutrients are supplied from the opposite direction as light (and, inorganic carbon, if it is not supplied through bubbling the medium with CO₂, or by adding bicarbonates into the medium, Fig. 2). It could thus be imagined, that the nutrients might not be sufficiently supplied to the 'illuminated' zone in a PSBR biofilm. Also, submerged phototrophic biofilms are known to accumulated exceptionally high dissolved oxygen (DO) concentrations (Pringault and Garcia-Pichel 2000). This phenomenon, as summarized by Raven et al. (2008), can have negative effects on photosynthetic productivity. In a PSBR biofilm, due to its direct exposure to the ambient gas phase, a better gas exchange is expected, thus, a lower [DO] compared to submerged systems should be present. The pH is another important factor, since it affects photosynthetic productivity by changing the speciation of dissolved inorganic carbons (DICs) and/or due to other mechanisms (Danilov and Ekelund 2001; Heber et al. 1976). However, pH gradients in a PSBR biofilm along the biofilm depth gradient have not been investigated yet. Since most of the culture media used for microalgal cultivation are buffered, it is expected, if fresh medium is sufficiently supplied, that the pH gradient along the depth gradient of the PSBR biofilm would be rather stable (Murphy and Berberoglu 2014). All the above-mentioned hypotheses and/or speculations have not been verified by experimental studies or otherwise (e.g. using a well verified model) until now.

1.5. Framework of this study

The goal of this study is to create a solid basis (e.g. experimental methods and analytical approaches) in order to achieve a more comprehensive understanding of the dynamic processes (including: light transfer, mass transfer, growth dynamics and chemical equilibriums) in PSBR biofilms in future studies, and, the verification of previously established hypothesis. To realize this, experimental studies and a modeling study were carried out. The developed model was then applied to find explanations for the observations made in the experimental studies (e.g. by comparing results from simulated biofilms and the results from real experimental observations under identical culture conditions). To eliminate uncertainties originating from the behavior of different strains cultivated under the same conditions, a single algal strain that has been studied intensively in previous studies (green alga *Halochlorella rubescens*, CCAC 0126; Culture Collection of Algae at the University of Cologne; www.ccac.uni-koeln.de) was used throughout the present study.

To determine the distribution of different parameters (light intensity, DO, pH, and photosynthetic productivity) in the *H. rubescens* Twin-Layer PSBR (TL-PSBR) biofilm experimentally, microsensor measurements (in collaboration with Max-Planck Institute for Marine Microbiology; Bremen, Germany) were carried out. In this study, microsensors were used for the first time to measure parameters in a non-submerged phototrophic biofilm. In

consequence, several changes were made in the microsensor experimental setup for TL-PSBR biofilm measurement compared to microsensor setup used for measurements on submerged biofilms. Also, a new mathematical method was developed in order to evaluate the data (for photosynthetic productivity measurement) acquired from the microsensor measurements more accurately. The first manuscript presented in this thesis (Li et al. 2015c): A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to evaporation, was dedicated to these topics.

After the establishment of the experimental methods, a systematic microsensor measurement was carried out: PAR, DO, pH and photosynthetic productivity distributions were measured in *H. rubescens* PSBR biofilms cultivated at different surface PAR intensity and/or exposed to different CO₂ concentrations in the gas phase. The measurement data from this part of the study is presented in the second manuscript included in this thesis (Li et al. 2015b): *Microscale profiling of photosynthesis-related parameters in a highly productive biofilm photobioreactor*.

Finally, a model was developed for the *H. rubescens* TL-PSBR biofilm. To make the model more realistic, experiments were carried out in the framework of this study to determine the parameters required by the model. The model is described in the last manuscript presented in this thesis (Li et al., 2015d, manuscript under review): *Investigating dynamic processes in a porous substrate biofilm photobioreactor* – *A modeling approach*. Also, in this manuscript, the experimental results both from this study and from a previous study (Schultze et al. 2015) were compared with that from the model simulation.

2. Publications included in this thesis

1St manuscript: A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to evaporation (Published in 'Journal of Microbiological Methods', 2015, 117:100-107).

2nd manuscript: *Microscale profiling of photosynthesis-related variables in a highly productive biofilm photobioreactor* (Accepted in Oct. 2015 by 'Biotechnology and Bioengineering').

3rd manuscript: *Investigating dynamic processes in a porous substrate biofilm photobioreactor – A modeling approach* (Submitted to the journal 'Algal research', under review).

2.1. A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to

evaporation

Published in 'Journal of Microbiological Methods', 2015, 117:100-107.

DOI: 10.1016/j.mimet.2015.07.022

Presented here in the original published version, reformatted for the presentation in this thesis (with section

numbers removed and corresponding changes).

- Publications included in the thesis -

A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to evaporation

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Abstract

Phototrophic biofilms are widely distributed in nature and their ecological importance is well recognized. More recently, there has been a growing interest in using artificial phototrophic biofilms in innovative photobioreactors for production of microalgal biomass in biotechnological applications. To study physiological processes within these biofilms, microsensors have been applied in several studies. Here, the 'light-dark shift method' relies on measurement of photosynthetic activity in terms of light-induced oxygen production. However, when applied to non-submerged biofilms that can be found in numerous locations in nature, as well as in some types of photobioreactors, limitations of this approach are obvious due to rapid removal of gaseous species at the biofilm surface. Here, we introduce a mathematical correction to recover the distribution of the actual photosynthetic activity along the depth gradient in the biofilm, based on a numerical solution of the inversed diffusion equation of oxygen. This method considers changes in mass transport during the measurement period as can found on biofilms possessing a thin flow/mass transfer boundary layer (e. g. non-submerged biofilms). Using both simulated and real microsensor data, the proposed method was shown to be much more accurate than the classical method, which leads to underestimations of rates near the biofilm surface. All test profiles could be recovered with a high fit. According to our simulated microsensor measurements, a depth resolution of $\leq 20 \ \mu m$ is recommended near the surface. We conclude that our method strongly improves the quality of data acquired from light-dark measurements of photosynthetic activity in biofilms.

Keywords

Phototrophic biofilm, Microsensor, Gross photosynthetic activity, electron transfer rate, Microalgae, Tikhonov regularization.

Introduction

Phototrophic biofilms are attached microbial communities driven by photosynthesis, and their importance in nature has long been recognized (Roeselers et al., 2008; Seckbach and Oren, 2010). In recent years, phototrophic biofilms have been applied to biotechnology and environmental biotechnology of microalgae, as well. Here, most applications rely on submerged biofilms, where the light exposed biofilm surface is covered by a layer of water (Berner et al., 2014; Roeselers et al., 2008). Some more recent developments make use of the advantages of biofilms exposed to ambient air (Berner et al., 2014; Schultze et al., 2015; Shi et al., 2007), referred to as Porous Substrate Bioreactors (PSBRs) (Murphy et al., 2014). In view of the technical development of photobioreactors, detailed micro-scale (microsensor) analyses of parameters along the axis (gradient) perpendicular to the biofilm surface have been proposed (Roeselers et al., 2008; Schultze et al., 2015). Such studies might also be of interest to analyze non-submerged (aerial) phototrophic biofilms in the natural environment (Berner et al., 2014).

A number of approaches have been developed to investigate structures and processes inside phototrophic biofilms at the micro-scale level, such as confocal microscopy, microsensor studies, fluorometry, and mathematical modelling (Bachar et al., 2008; Beutler et al., 2009; Kühl and Polerecky, 2008; Ramanan et al., 2013; Revsbech et al., 1981; Wolf et al., 2007). Among these methods, the microsensor-based "light-dark shift method" is a well-established technique to estimate gross photosynthetic activity in phototrophic biofilms since its introduction in the 1980s (Revsbech and Jorgensen, 1983). The method determines the rate of decrease in the dissolved oxygen concentration by an oxygen microsensor, which is induced by applying a short period of darkness to a steady state phototrophic biofilm (Revsbech and Jorgensen, 1983). However, since the diffusion rate of oxygen changes during this dark period, the gross productivity estimated using the acquired data directly might be inaccurate. This limitation was soon recognized and solutions were suggested in serval previous studies, which established a solid basis for evaluating data acquired by light-dark shift method. However, these solutions require either a hypothetical distribution of photosynthetic activity or rely solely on the regression of measured data (Glud et al., 1992; Lassen et al., 1998; Revsbech and Jorgensen, 1983; Revsbech et al., 1986). For non-submerged biofilms, besides the limitation referred to above, other processes have to be taken into account: the dynamics of oxygen in a non-submerged biofilm can be described by Eq.1, assuming that the biofilm is homogenous in planes parallel to its surface (Bergman et al., 2011).

$$\frac{\partial c}{\partial t} = P_g - R_l + D_e \frac{\partial^2 c}{\partial x^2} + u \frac{\partial c}{\partial x} - R_s$$
(Eq.1),

The terms on the right hand side represent gross photosynthesis P_g , respiration during light R_l , diffusion and advection caused by evaporation on the surface, and the removal of dissolved gaseous species due to air flow above the surface R_s . D_e is the effective diffusion coefficient, C is the concentration of dissolved oxygen, u is the convective flow rate inside the biofilm,

$$u = q_e / (A_e \cdot \theta) \tag{Eq.2},$$

and can be calculated with the rate of liquid volume lost q_e , the exposed surface area of the biofilm A_e and the areal porosity of the biofilm θ (Eq.2). *t* is time and *x* is the depth. Without light, no oxygen evolution can be expected (Pessarakli, 2005). Thus after darkening oxygen decreases with a rate:

$$\frac{\partial c}{\partial t} = 0 - R_l + D_e \frac{\partial^2 c}{\partial x^2} + u \frac{\partial c}{\partial x} + R_s$$
(Eq.3).

Values measured by the light-dark shift method are the rate of change for the term on the left hand side of Eq.3 during the measurement period. Immediately after the removal of light (at 0 s of the measurement), $\partial C/\partial t$ does represent accurately the gross photosynthetic productivity. However, its value changes gradually with time, as the removal of light leads to change of the values on the right hand side (Glud et al., 1992; Revsbech, 1986; Revsbech and Jorgensen, 1983). In consequence, an accurate representation of the photosynthetic rate is provided only if all terms on the right hand side of Eq.3 remain constant with time (as the measurement time has to be > 0 s), which cannot be assumed: Previous work (Glud et al., 1992) concluded that the respiration rate can be considered stable during the measurement period, but the change in rate of diffusion can have a significant effect on the measured results (Glud et al., 1992; Revsbech and Jorgensen, 1983; Revsbech et al., 1986).

When applying the light-dark shift method for measuring photosynthetic activity to a non-submerged biofilm, two additional factors attributed to the absence of a thick liquid phase on the biofilm surface have to be considered (Bergman et al., 2011; Schlichting and Gersten, 2000):

- 1) For a non-submerged system directly exchanging dissolved gases with the gas phase above the biofilm surface, the flow/mass transfer boundary layer (compared to submerged biofilms) can be considered as non-existent in case of a considerable air flow. With the gas phase above the biofilm being in effect a sink for the oxygen produced inside the biofilm, during the light-dark cycle, the rate of change of oxygen concentration at the biofilm surface will always be 0 (or almost 0), and the change rate near the surface an underestimation of the photosynthetic activity. This effect also occurs in submerged system at the interface between the boundary layer and the bulk liquid, however, the effect in the biofilm itself is dampened due to the relatively thicker flow/mass transfer boundary layer.
- 2) The evaporation of water at the biofilm surface exposed to air induces a convective flow through the biofilm in the direction perpendicular to the biofilm surface, which in effect adds a mass flow towards the surface.

These two processes have a strong effect throughout a non-submerged biofilm, and, therefore, need to be considered for the interpretation of acquired experimental data to obtain a profile representing a valid photosynthetic activity.

In summary, direct results from the light-dark shift measurement do not accurately represent the actual photosynthetic activity in biofilms, especially for measurements performed in non-submerged biofilm systems. Thus, in the present contribution we propose a mathematical treatment to achieve an adequate recovery of the distribution of actual photosynthetic activity from measured microsensor data.

Materials and Methods

Suspension and biofilm cultivation of microalgae

In this work, a Twin-Layer algal biofilm, which is an immobilized biofilm system that separates the bulk medium and biomass by means of a micro-porous membrane (Nowack et al., 2005), was used as a model for a non-submerged biofilm. Stock cultures of the green algae *Halochlorella rubescens* (strain CCAC 0126; Culture Collection of Algae at the University of Cologne; www.ccac.uni-koeln.de) in suspension was kept at 16 °C at a light intensity of about 20 µmol photon $m^{-2} s^{-1}$ photosynthetically active radiation (PAR). To obtain sufficient material for biofilm inoculation, stock cultures were transferred into 2 L Erlenmeyer flask filled with 1.3 L of Bold's basal medium (BBM) (Bischoff and Bold, 1963), and were cultivated at a light intensity of about 50 µmol $m^{-2} s^{-1}$ PAR at 23 °C with 0.5 L min⁻¹ compressed air with 1% (v/v) supplementary CO₂. Biomass was harvested after 3 weeks of growth, and was inoculated onto polycarbonate filters (PC40, 0.4 µm pore size, 25 mm diameter, Whatman, Dassel, Germany) with a circular inoculation area of 2.55 cm² as described by Naumann et al. (2013). Inoculated filters were then transferred into a bench-scale Twin-Layer system for cultivation according to Schultze et al. (2015). Tubes were aerated with a flow rate of 0.75 L min⁻¹ with compressed air. Sodium discharge lamps (SON-T AGRO 400W, Philips, Hamburg, Germany) were used as light source with a 14-10 hour light-dark cycle at an intensity of 1,000 µmol m⁻² s⁻¹ PAR. Temperature in the tubes was 25±2 °C. 1 L BBM culture medium was used at a medium flow rate of 3 mL min⁻¹ per tube and the culture medium was replaced every 2 days.

Microsensor measurements

A schematic representation of the experimental setup for microsensor measurements based on previous works by Gieseke and de Beer (2004) is given in Figure 1: the biofilm immobilized on a polycarbonate membrane was placed inside the biofilm measurement cell under a controlled atmosphere of compressed air at a flow rate of 1 L min⁻¹ (Figure 1A). The non-inoculated area of the polycarbonate filter and glass fiber were covered with a black plastic foil to exclude light and gas exchange in this area. 50 mL of BBM culture medium were circulated though the measurement cell with a flow rate of 3 mL min⁻¹ by means of a peristaltic pump and were exchanged every 1 hour during the measurement. The evaporation rate was monitored by recording the change of the volume in the medium container. Biofilm temperature during the experimental period was 23 ± 0.5 °C. Light was supplied at an intensity of 1,000 μ mol m⁻² s⁻¹ (on biofilm surface) from the front side by a halogen lamp (KL-1500, Schott, Mainz, Germany) equipped with a 3-fold splitter to ensure even distribution. The movement of the sensor was enabled by the computer-controlled micromanipulator (Pollux Drive, PI miCos, Eschbach, Germany). Data were recorded with a depth resolution of 20 µm until 500 µm depth (at which the switch from light to dark has no significant effect on the oxygen concentration measured). Microsensor measurements were carried out on biofilms cultivated for 30 days. Microsensor signals were amplified and converted into digital data (DAQpad 6015 and 6009, National Instruments, Munich, Germany) prior to their further processing on a computer (Figure 1B). Software used for system control and data acquisition were custom made (can be acquired upon request from Dr. Lubos Polerecky, Utrecht University, Netherlands, l.polerecky@uu.nl). A picoampere meter and Clark-type oxygen microelectrodes with tip diameters $\leq 10 \ \mu m$, and a 95 % response time < 0.5 s were used for the light-dark shift measurement

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(Revsbech, 1989). Linear calibration of oxygen microsensor was carried out as described by Revsbech (1983) using BBM saturated with nitrogen gas or compressed air. A shutter connected to a timer was used to control light-dark cycling: the total dark period was 3 s, and a linear regression slope of oxygen concentration change between 0.7 and 2.1 s of the dark period was determined as the measured value. For each measurement depth, three light-dark cycles were performed after steady state was reached (this could be as long as 10 min), and these 3 values acquired were considered as triplicates. The PAR light intensity profile was acquired using a fiber optic light microsensor with a spherical tip (80 µm diameter) and a portable spectrometer (USB 4000, Ocean optics, Dunedin, USA) (Kühl et al., 1994a, 1994b). At each measurement depth, the average value of a 3 s measurement period was taken as the PAR intensity.



Figure 1: Experimental setup for microsensor studies in non-submerged microalgal biofilms. A: Schematic drawing of the vertical cross section of the measurement in artificial biofilms. Solid arrows in the glass fiber tissue indicate flow direction of the culture medium. B: Schematic drawing of the complete experimental setup, dotted lines with arrow indicate signal flow, whereas solid arrows represent medium and gas flows in the system. *The pH reference electrode and CO₂ input were installed but not used for measurements shown in the present paper.

Electron transfer rate (ETR) measurement

For measuring the photosynthetic performance at different depths in the algal biofilm by PAM (pulse amplitude modulated) chlorophyll fluorescence analysis, the biofilm was sectioned stepwise in parallel to its surface plane with a resolution of 50 µm by means of a custom-made hand microtome (Figure 2). The biofilm immobilized on a polycarbonate membrane was placed onto a height-adjustable stage on a thin layer of culture medium, whereas the stage allows levelling the biofilm sample into the sectioning plane by means of a micrometer screw. The sectioning plane was defined by a solid brass support over which a rigid microtome blade was passed manually to remove the overlying algal biofilm (Figure 2).



Figure 2: Schematic drawing of the experimental setup used for pulse amplitude modulated fluorometry and biofilm sectioning. The sample holder allows moving the biofilm in vertical direction by means of a micrometer screw (indicated by the thin solid arrows). The biomass above the sectioning plane was removed with a solid microtome blade prior to the fluorescence measurement in the respective layer. The cutting direction is indicated by the thick arrow. Measurements of chlorophyll fluorescence were carried out from surface to a depth of 500 um without removing the biofilm from the stage. Using a PAM-2000 Chlorophyll Fluorometer (WALZ, Effeltrich, Germany), PAM chlorophyll a fluorescence of photosynthetically active cells of the exposed biofilm surface was measured by application of the saturation pulse method: The fiber-optic cable of the chlorophyll fluorometer was centered in a 20 mm distance perpendicular to the biofilm surface (Figure 2). A weak beam of modulated (20 kHz) red light (0.03 µmol m⁻² s⁻¹ PAR with a maximum intensity at 650 nm and a maximum cutoff wavelength at 670 nm) was used to excite chlorophyll fluorescence (F_{θ} or F), which was then detected by a photodiode with a minimum cutoff wavelength of 700 nm. Maximal fluorescence F_m or F'_m was induced by applying a saturation pulse of white light of about 2.000 mol m^{-2} s⁻¹ and a duration of 600 ms. Fluorescence data were recorded during rapid light curves with an ascending light intensity applied to each section: directly after each sectioning procedure, the biofilm was dark-adapted for 10 min and F_0 and F_m were recorded at the end of the dark period to determine the maximal quantum efficiency F_{v}/F_{m} . Subsequently, a rapid light curve was recorded using actinic white light applied stepwise at 11 consecutive intensities (7, 12, 18, 38, 66, 122, 209, 373, 642, 1099, and 1762 mol m⁻² s⁻¹ PAR) for 60 s (Herlory et al., 2007), followed by the determination of F and F'm to calculate effective quantum yield of electron transport $\Delta F/F'_m$ according to Genty et al. (Genty et al., 1989) for each of the light intensities. During the measurement, the samples were kept water-saturated by regularly adding a few drops of medium to the edges of the membrane filter (resulting in a total volume of 50 - 100 µL culture medium for the whole experimental period). ETR measurements were carried out on biofilms cultivated for 97 days.

The electron transfer rate (ETR) can be used to estimate the primary productivity of photosynthetic biofilms and is related to (gross) rates of oxygen production. The ETR was calculated as (Hofstraat et al., 1994; Ralph et al., 1999):

$$ETR = 0.5 \times I_{PAR} \times \Delta F / F'_m \times Abs$$
 (Eq.4)

 I_{PAR} is the PAR intensity and *Abs* represents the PAR absorptivity of the sample material. *Abs* was determined by an IMAGING-PAM Chlorophyll Fluorometer (equipped with Maxi-Head; WALZ, Effeltrich, Germany) for all measured depths. Measurements were performed in triplicates (three independent biofilm samples). From rapid light curve data for every section, light intensity was plotted against the ETR and was mathematically fitted using the Photosynthesis-Irradiance (P-I) model of Platt et al. (Platt et al., 1980) by means of GraphPad Prism v5.02 statistical software (GraphPad Software Inc., La Jolla, USA). Based on microsensor measurements of the light intensity profiles, available light intensities for all of the measured depths can be determined by using the "Point to point" function in GraphPad Prism. These data were used to derive the potential ETR for the sectioned depths to obtain an overall ETR depth profile.

Mathematical treatment

The following mathematical treatment was proposed for correcting data acquired from the light-dark shift method: Take the derivative of time on both sides of Eq.3, and assume the time t and spatial variables x are not dependent (as during the measurement period, the thickness of the biofilm hardly changes):

$$\partial (\frac{\partial c}{\partial t})/\partial t = 0/\partial t - \partial R_l/\partial t + D_e \frac{\partial^2 (\frac{\partial c}{\partial t})}{\partial x^2} + u \frac{\partial (\frac{\partial c}{\partial t})}{\partial x} - \partial R_s/\partial t$$
(Eq.5).

The term $\partial R_s / \partial t$ represents the rate change of the removal of dissolved gaseous species at the submerged biofilm surface due to the flow of bulk liquid, or, in the case of an aerial biofilm (or PSBR biofilm), due to air above the biofilm:

$$\partial R_s / \partial t = (D_e \frac{(C - C_s)}{x^2} + u \frac{(C - C_s)}{x}) / \partial t$$
$$= (\frac{D_e}{x^2} + \frac{u}{x}) \cdot \partial C / \partial t$$
(Eq.6).

 C_s represents the oxygen concentration on the very surface of the aerial biofilm, or at the interface between the boundary layer and the bulk liquid in the case of a submerged biofilm, which is in equilibrium with the phase above and is considered to be a constant. Considering that the rate of respiration remains stable during the measurement period (Glud et al., 1992), and take $\partial C/\partial t = P(x, t)$, as $\partial C/\partial t$ is a function of depth *x* and time *t*. Substitute Eq.6 into $\partial R_s/\partial t$ and $\partial C/\partial t$ to P(x, t), Eq.5 becomes:

$$\frac{\partial P(x,t)}{\partial t} = 0 - 0 + D_e \frac{\partial^2 \left(P(x,t) \right)}{\partial x^2} + u \frac{\partial \left(P(x,t) \right)}{\partial x} - \left(\frac{D_e}{x^2} + \frac{u}{x} \right) \cdot P(x,t)$$
(Eq.7).

Eq.7 is a nonhomogeneous diffusion equation with a source term (here a negative source, or a sink) and can be solved by using any of the methods summarized by Crank or Polyanin (Crank, 1975; Polyanin, 2002). Using initial condition P(x, t = 0) = P(x, 0) and boundary condition $\partial P(x = 0, t)/\partial t = 0$. The exact solution of Eq.7 at $t = \tau$ can be expressed as:

$$P(x,\tau) = \int_0^{\delta} P(y,0) \cdot \left[e^{\left(\frac{-(y-x+u\cdot\tau)}{4\cdot D_e \cdot \tau}\right)} - e^{\left(\frac{-(y+u\cdot\tau)}{4\cdot D_e \cdot \tau}\right)} \cdot erfc(\frac{x}{\sqrt{4\cdot D_e \cdot \tau}}) \right] \cdot dy$$
(Eq.8),

 δ is the total depth of the biofilm, or, alternatively, the depth until which the shift from light to dark has an effect on the dissolved oxygen concentration. $P(x, \tau)$ is the measured value from the light-dark shift method with measurement time τ at measurement depth x, and P(y, 0) is the actual photosynthetic activity at position y (the integrating variable), which is the desired term. For measurements taken at all depths, the measured value $P(x, \tau)$ at a depth x depends on the P(y, 0) across the complete photosynthetic active region of the biofilm (surface to depth δ). The effect of diffusion and advection inside the biofilm is reflected by the first term in the square brackets (inner term). The effect of surface flow removal is reflected by the second term in the same brackets (surface term, and *erfc* represents the complement error function). In this form, the terms representing the inner and surface effects are separated.

Eq.8 is a Fredholm integral equation of the first kind (Kress, 2014), and can be solved by using numerical methods (Baker et al., 1964; Neumaier, 1998). The numerical approach used in this paper is: Discrete the system perpendicular to the biofilm surface using a simple 1D upwind scheme by setting the dissolved oxygen concentration in a discretion element equal to the value at its knot closer to the biofilm surface (Courant et al., 1952); then calculate the approximated solution of Eq.8 using Tikhonov regularization (Tikhonov et al., 1995) with a non-negative constrain applying the active-set algorithm purposed by Gill et al. (1981). Discrete δ by *n*, and let $P(x_j, \tau) = g_j$, $P(y_h, 0) = h_i$; (i, j = 1, 2, ..., n) and write Eq.8 in the discrete operator notion:

$$g_j = \sum_i^n k_{i,j} \cdot h_i \tag{Eq.9},$$

 $k_{i,j}$ is the operator term which contains both the term that reflects the effect of diffusion and advection of dissolved oxygen inside the biofilm (inner term) and the term that reflects the effect of dissolved oxygen removal due to surface flow (surface term). In matrix notion, let *G* be a vector containing g_j , *H* a vector containing $h_{i,j}$ and *K* a matrix containing the operator terms $k_{i,j}$:

$$G = K \cdot H \tag{Eq. 10}.$$

Since a profile with only positive values is expected, and the diffusion is expected to be the dominant process affecting the measured value, generalized Tikhonov regularization with a second derivative operator is used (Hansen, 1994; Varah, 1983). Apply the regularization and non-negative constrain to Eq.10, and let L be the second derivative operator, an approximation of H can be acquired numerically by solving the minimization problem:

$$\min_{H>0} \{ \|G - K \cdot H\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \}$$
(Eq. 11).

The problem represented by Eq.11 can easily be solved by using solvers designed for this kind of problems as explained by Stewart (1973). The regularization parameter λ can be found by using several methods (Calvetti et al., 2000; Hansen, 1994; Hansen and O'Leary, 1993), in this paper, the L-curve method purposed by Hansen & O'Leary (1993) was used to find the best λ .

The solution of H for problems represented by Eq.11 is straightforward. However, as the operator K contains terms that represent both the effect of diffusion and advection inside the biofilm and removal due to surface flow, a direct solution might be inaccurate near to the surface if the noise level in the input data is high. Since the term representing surface flow removal tends to smoothen the profiles near to the surface (Hansen, 1992), a peak in the true profile near the surface can be smoothened out in the solution. This, however, can be resolved by mathematical manipulation of the operator K. Rewrite Eq.10 as:

$$(G_{in} + G_{surf}) = (K_{in} + K_{surf}) \cdot H$$
(Eq.12),

The subscripts *in* and *surf* represent the effect from the inner term and the surface term in Eq.8 respectively, operators K_{in} , K_{surf} can be calculated using Eq.8, as well as g_{in} and G_{surf} , if an *H* is given. The problem in Eq.11 thus became:

$$\min_{F \ge 0} \left\{ \left\| \left(G_{in} + G_{surf} \right) - \left(K_{in} + K_{surf} \right) \cdot H \right\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \right\}$$

or,

$$\min_{F \ge 0} \left\{ \left\| (G_{in} - K_{in} \cdot H) + (G_{surf} - K_{surf} \cdot H) \right\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \right\}$$
(Eq.13).

Observing the surface term in Eq.8, notice the maximum value of the surface term is controlled by the position of the actual activity *y*, but a complement error function dependent solely on the position of the measurement taken (*x*) controls the final value. Furthermore, for a given *G*, under the same measurement conditions, G_{surf} always has the same set of values. Thus, for a given *G*, the minimization of the term $||G_{surf} - K_{surf} \cdot H||^2$ will always yield the same solution. The problem in Eq. 13 can be simplified to:

$$\min_{f \ge 0} \{ \|G_{in} - K_{in} \cdot H\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \}$$
(Eq. 14).

Eq.14 does not contain the surface term, but the solution of the problem still leads to H, which is the desired original profile.

The complete procedure of the calculation is given below:

- 1) Discrete the system and calculated L as the discrete second derivative operator.
- Calculate operator K_{in}, K_{surf} and K using Eq.7. No knowledge other than diffusion coefficient D_e, flow rate u (calculated from Eq.2) inside the biofilm and measurement time τ is required. A λ was calculated using L and the operator K.
- A preliminary H (here denoted as H₀) was calculated by solving Eq.11 using the measured data as G, the operator L, K and the λ calculated from the previous step.
- G_{in} and G_{surf} were calculated by using H₀, K_{in}, K_{surf} and Eq.12. And a new λ was calculated using operator L and K_{in}.
- 5) Solve Eq.14 for H using the calculated G_{in} , K_{in} , L and the λ calculated from the previous step.

 H_0 does not necessarily give a faithful representation of the true profile, but gives the same measured distribution G, and thus G_{in} and G_{surf} (as G_{surf} has a fixed value for the same measurement condition) when used in Eq.10 and Eq.12 respectively. In total, 5 inputs are needed: measured data points from the light dark shift method, effective diffusion coefficient in the biofilm, evaporation rate, porosity of the biofilm and measurement time.

The treatment procedure was coded in MATLAB (version 2013a, MathWorks, Ismaning, Germany), and the inbuilt *quadprog* was used for solving the minimization problems. *L-curve* function from the regularization tools (Hansen, 1994) was used to find the λ value. The mean values of triplicates acquired from the microsensor measurement were randomly added or subtracted with a random value in range of the calculated standard deviation (SD) at the same position. These values were used as input for the calculation. The calculation procedure was repeated 3 times, and the mean value of the 3 results was taken as the final result (a SD can also be calculated). Data interpolation, if needed, was done by simply connecting two measured data points with a straight line (linear interpolation, using MATLAB inbuilt function *interp1*).

Test problems setup

To test the proposed treatment, first, simulated measurement data points were generated using known actual profiles, in total, 5 different test profiles adopted from previously reported possible photosynthetic activity distribution in phototrophic biofilms were used (Glud et al., 1992; Lassen et al., 1998; Revsbech and Jorgensen, 1983; Revsbech et al., 1986). These known profiles, together with assumed diffusion coefficient, evaporation rate, porosity and measurement time (which were similar or identical to the values encountered in actual biofilm measurements) were put into Eq.8 to generate simulated measurement data points. The acquired profiles (simulated measurement data points), were then sampled every 5, 10, 20, 40 or 80 μ m to simulate the different depth resolutions used. Then a random error $\leq 20\%$ of the maximum value in the simulated data was added to simulate

noise in the measurement. The simulated profiles together with other parameters were then used for recovering the original distribution.

The test was also carried out using data acquired from light-dark shift measurements on a *H. rubescens* Twin-Layer biofilm: The porosity of the biofilm was estimated from the volume of cells (biovolume) in the biofilm divided by the total biofilm volume. The biovolume was calculated as described by Hillebrand et al. (1999) from suspended biofilm samples with a 100 μ m resolution to a depth of 500 μ m using the sectioning technique described above (n = 3 filters). The determination of porosity showed no significant differences in depth sections of 100 μ m each (one-way ANOVA test using GraphPad Prism v5.02, as described above), and the mean value of 0.76 was taken as volumetric porosity for the calculation. Using this value, the diffusion coefficient was estimated using the relationship suggested by Weissberg (1963), and effective diffusion coefficients were calculated to be 0.68 of that in pure water. Evaporation rate during the measurement was 0.68 mL min⁻¹, which corresponded to an advective flow rate of 5.4 μ m s⁻¹ inside the biofilm. 2.0 × 10⁻⁹ m² s⁻¹ was taken as diffusion coefficient of dissolve oxygen in water (Bergman, 2011) for all calculations. The recovered photosynthetic profile was than compared with the measured ETR profile. The n values used for discretion for both simulated and real profiles were 10, 25, 50, 100, 200 and 500.

Results

To assess the performance of the proposed mathematical method, a comparison between the input test distribution used for generating the simulated measurement data points and the recovered distribution using the proposed method with different simulated depth resolutions was performed (Figure 3). Only recovered distributions using n= 50 were shown, since further increase in n did not lead to significant improvement in the results and a decrease in n, in some cases, tends to produce physically impossible results (in combination with a low depth resolution). The maximum computation time (n = 500) using a DuoCore 2.8 GHz CPU (Intel, Santa Clara, USA) was less than 1s. The following test profiles were used (Figure. 3A-E): A test distribution following the shape of measured light distribution in the biofilm (3A); a biofilm that has lower activity at the surface as proposed by Glud et al. (1992) (3B); distributions following the shape proposed by Revsbech et al. (1983) and Revsbech & Jorgensen (1986) (3C+D); a profile with a double peak as described by Lassen et al. (1998) (3E). All profiles tested were recovered with relatively high fidelity by the proposed method. However, a more accurate representation of the actual profiles was acquired with higher depth resolutions in the simulated measurement. For all test profiles, resolution of 20 µm produced profiles that are not different from the original distributions, whereas, at 40 µm and 80 µm resolution, this was observed only for some profiles (Figure 3A-D and 3B, respectively). A further increase in resolution to < 20µm produced better results in only one of the test profiles (Figure. 3E). Even though all sharp edges in the original distribution in the biofilm were smoothened, the discontinuities on the surface were successfully recovered in all cases.



Figure 3: Comparison between the simulated, actual, and recovered distribution of photosynthetic activity (oxygen production in terms of μ mol O₂ L⁺ s⁻¹) for five different test profiles (panel A to E): Black dashed lines show the original/actual distributions, whereas the simulated profiles are given in empty circles (with 5 μ m resolution); solid lines represent the recovered distributions using the proposed method, and the simulated measurement depth resolutions used for recovery decreases with increasing brightness.

Results for the test conducted using real measurement data from the *H. rubescens* biofilm are presented in Figure 4. The data points from direct measurement showed reduced activity at the biofilm surface. In comparison, the recovery profile follows the trend of light profile recorded in the biofilm, displaying a monotonical decrease of photosynthetic activity with increasing depth. At a depth of about 450 μ m, the recovered photosynthetic activity dropped to 0 μ mol O₂ L⁻¹ s⁻¹ in congruence with the measured PAR light intensity (Figure 4A). Furthermore, the recovered distribution of photosynthetic activity from microsensor measurements and the photosynthetic electron transfer rates (ETRs) measured by means of chlorophyll fluorescence showed an almost identical distribution of photosynthetic activity (Figure 4B) in the *H. rubescens* biofilm. This indicates that the corrected profile is a better representation of the actual photosynthetic activity distribution.



Figure 4: Comparison between measured photosynthetic activity (oxygen production in µmol O2 L-1 s-1and ETR in µmol electrons m⁻² s⁻¹), and the calculated distribution of photosynthetic activity in terms of oxygen production in biofilm of the microalgae Halochlorella rubescens. A: Measured distribution of photosynthetic activities (empty squares, error bar represent standard deviation of triplicates) and the recovered actual distribution (solid black line). Grey solid line gives the values of the photosynthetic activity that should have been measured if the solid black line was the actual photosynthetic activity profile. B: ETR (electron transfer rate, empty circles are mean values from triplicates, solid line represents the smoothed curve) of the microalgal biofilm. Distribution of photosynthetic active radiation (PAR) intensity were given in both panels as dotted lines (in µmol photon m⁻² s⁻¹).

Discussion

Generally, using data acquired directly from the light-dark shift method for estimating the photosynthetic activity in phototrophic biofilms can be inaccurate due to the change in mass transfer rates during the dark period and a loss of dissolved species because of surface removal of dissolved oxygen due to the advection in the phase above the biofilm (as described in introduction). The latter effect becomes even more pronounced when measurements are to be carried out on non-submerged biofilms. For submerged biofilms, to mitigate this inaccuracy, an agar layer has been applied onto the biofilm surface to stabilize the flow boundary layer (Glud et al., 1992). However, this procedure could change the structure or the mass transfer at the biofilm surface. For non-submerged biofilms, this agar treatment is inappropriate, since the biofilm will no longer be in direct contact with the gas phase above its surface. Light-dark shift measurements have been carried out previously on non-submerged biofilms, but no additional treatment of data was used (Bernstein et al., 2014). Such results, as shown by the comparison of original photosynthetic profiles and the simulated measured data points in Figure 3 (the effect of the surface removal), can be very misleading, especially for biofilms with both phototrophic and heterotrophic components (Glud et al., 1992; Lassen et al., 1998; Revsbech and Jorgensen, 1983; Revsbech et al., 1986).

To a certain degree, a better experimental set up (e.g. a shorter measurement time), can reduce the inaccuracy mentioned above. However, a mathematical treatment is required to improve the results further. And this requires the exact solution of the diffusion equation (Eq.7) (Glud et al., 1992; Revsbech et al., 1986). This can be derived using various methods (Crank, 1975; Polyanin, 2002). In the proposed mathematical treatment, the expression of Eq.8 was chosen so that the terms representing the effect of the process in the biofilm and the effect from the surface are separated (inner and surface terms in the square brackets). Eq.8 is a Fredholm integral equation of the first kind. Its solution requires the solution of an inversed diffusion equation, which has been discussed in detail since the 1960s and numerous solution methods have been developed (Hansen, 1992, 1994; Tikhonov et al., 1995). As described above, a generalized Tikhonov regularization with an approximated second derivative operator was chosen and the L-curve method was used for finding the best regularization parameter, which defines the balance point between the regularization error and the residue norm (Hansen and O'Leary, 1993). The regularization method, gives a smoothened approximated solution, and thus the recovery of the exact values should never be expected (Hansen, 1992). As a result, none of the sharp edges in the original distribution inside the biofilm was recovered (Figure. 3). Another problem caused by the smoothening is, as mentioned previously, a peak in the actual distribution near to the surface of a biofilm might be simply smoothened out if no precaution is taken to avoid this (e.g. direct solution of Eq.11). However since the Eq.8 separates the inner and surface terms, the minimization of Eq.11 can be transformed into the minimization of Eq.14. This yields a solution more accurately representing the original distribution, and the activity at the surface is always recovered when appropriate depth resolutions and nvalues were used (Figure 3). Following this approach, in the complete mathematical treatment procedure, the final approximated solution was solved using a previously calculated approximated solution (Step 3 and 4 in the last section of material and method). This did increase the numerical error compared to procedures that contain only a single approximated solution (Hansen, 1992; Tikhonov et al., 1995). However, in comparison to the solutions

acquired by directly solving Eq.10 (which could not recover the peaks near the biofilm surface in some cases, even though 5 μ m depth resolution and n = 500 were used, results not shown), this increment in numerical error is acceptable.

For the simulated test problems using different measurement depth resolutions, accuracy of the recovered profile increases with a higher depth resolution, which, for a real measurement, would increase time consumed by experimental work. In all test problems presented here, a 20 µm depth resolution gave satisfactory results, and is considered as a reasonable resolution to perform experimental measurements.

The number of the discretion elements, the value *n*, is generally a very important parameter affecting the accuracy of a numerical method. With smaller *n* values, the solutions acquired generally become less accurate or even unstable, whereas a larger *n* value leads to higher computational cost (Hutton, 2003). In the simulated test problems, with n < 50, oscillating results were sometimes generated when low depth resolution was used. However, with n = 50, no oscillating solutions were generated, and an increase of *n* to values higher than 50 produced no significant improvement in the results acquired (not shown). Interpolation of data points is needed if an *n* larger than the number of measurement positions is used (as for a depth resolution of 20 µm and n = 50 for a 500 µm thick biofilm). This, as described above, was carried out by simply assuming that the value changes linearly between two adjacent points. Several curve fitting methods can be used to improve the quality of the interpolation, however, if this is to be used with lower measurement resolutions, caution must be taken, since the fitted curve might reflect the wrong trend (Guest, 1961; Motulsky and Ransnas, 1987; Wahba, 1990).

Microsensor data collected from a highly active non-submerged phototrophic biofilm of Halochlorella rubescens cultivated using a Twin-Layer photobioreactor system (Schultze et al., 2015) was chosen in this study to test the proposed mathematical treatment: The correction of direct microsensor data recovered the real photosynthesis profile in terms of ETR favourably compared to the uncorrected profile (directly measured). In the uncorrected profiles, a decrease of activity near the biofilm surface was observed, but this was not reflected by the ETR measurement. The ETR was used in several studies to estimate the gross photosynthetic activity in phototrophic biofilms (Banares-Espana et al., 2013; Barranguet and Kromkamp, 2000; Flameling and Kromkamp, 1998) and ETR values reported here are in the same ranges as observed e.g. by Kromkamp et al. (2001) in a biofilm dominated by a cyanobacterium. A direct comparison between the ETR and the recovered distribution from the light-dark shift method, however, has to be treated with caution: the microsensor method is generally recognized as a non-destructive and in-situ method (Nivens et al., 1995). In contrast, the ETR determination in combination with biofilm sectioning as applied in this work is neither in-situ nor non-destructive. Sectioning places cells at the biofilm surface, which previously have been located deeper inside the biofilm. This changes the concentrations of oxygen and CO₂ that the cells were originally exposed to. Nevertheless, a general comparison of the trends of the recovered photosynthesis and measured ETR profiles can be made. In this sense, the recovered profile reflects the photosynthetic activity distribution better than the uncorrected direct measurement data. The reason for this discrepancy between recovered and direct profiles was very likely, the removal of dissolved oxygen at the biofilm surface.

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The accuracy of the proposed method depends strongly on the accurate determination of the effective diffusion coefficient and the advection rate in the biofilm. The effective diffusion coefficient in a biofilm depends on porosity and tortuosity, and the molecular size of the diffusing species as well as its affinity to biomass (IWA Task Group on Biofilm Modelling, 2006; Melo, 2005). Several methods have been developed for measuring or estimating the effective diffusion coefficient of dissolved oxygen in biofilms, however, all of these methods show individual weaknesses (Stewart, 1998; Wood et al., 2002). Considering a biofilm with a strongly depth-dependent structural heterogeneity, the situation becomes even more complex (Ramanan et al., 2013). In the simulated test problems shown here, the coefficient used to generate the simulated data points was the same as that used for recovery, thus no error is expected to originate from the effective diffusion coefficient has been estimated from porosity in the real biofilm test problems in this study, as described in section 2.5, since the observation during the sectioning suggested the cells in the biofilm behaved similarly to a densely packed porous medium), an inaccurate effective diffusion coefficient can lead to false estimations. On the other hand, the determination of the advection rate is relatively straightforward (using Eq.2), and, therefore, is no critical factor for the accuracy of the method.

We conclude that the proposed mathematical procedure can be applied with no further modification to microsensor data acquired by the light-dark shift method from biofilms with a very thin flow boundary layer as can be found in aerial biofilms or submerged biofilms subjected to a high flow rate in the bulk liquid. Furthermore, with minor modifications of the boundary conditions used for discretion, the method can also be used to recover the actual distribution of photosynthesis from measurements done on biofilms with a thicker boundary layer to overcome limitations of the currently used methods.

Acknowledgements

This study was supported by the University of Cologne (KST 158901001).

The authors thank Dr. Anthony Dron (Max-Planck-Institute für Marine Mikrobiologie, Bremen, Germany) for support during the installation of the experimental microsensor setup, Dr. Arjun Chennu (Max-Planck-Institute für Marine Mikrobiologie, Bremen, Germany) and Bastian Piltz (Universität zu Köln, Köln, Germany) for their valuable opinions. - Publications included in the thesis

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2.2. Microscale profiling of photosynthesis-related variables in a highly productive biofilm photobioreactor.

Accepted in Oct. 2015 by 'Biotechnology and Bioengineering'.

DOI: 10.1002/bit.25867

Presented here in the accepted version, reformatted for the presentation in this thesis.

Publications included in the thesis

Microscale profiling of photosynthesis-related variables in a highly productive biofilm photobioreactor

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Running head: Profiling biofilm in photobioreactor

Abstract

In the present study depth profiles of light, oxygen, pH and photosynthetic performance in an artificial biofilm of the green alga *Halochlorella rubescens* in a porous substrate photobioreactor were recorded with microsensors. Biofilms were exposed to different light intensities $(50 - 1,000 \ \mu mol photons \ m^{-2} \ s^{-1})$ and CO₂ levels $(0.04 - 5 \ \% \ v/v$ in air). The distribution of photosynthetically active radiation showed almost identical trends for different surface irradiances, namely: a relatively fast drop to a depth of about 250 μ m, (to 5% of the incident), followed by a slower decrease. Light penetrated into the biofilm deeper than the Lambert-Beer Law predicted, which may be attributed to forward scattering of light, thus improving the overall light availability. Oxygen concentration profiles showed maxima at a depth between 50 and 150 μ m, depending on the incident light intensity. A very fast gas exchange was observed at the biofilm surface. The highest oxygen concentration of 3.2 mM was measured with 1,000 μ mol photons m⁻² s⁻¹ and 5% supplementary CO₂. Photosynthetic productivity increased with light intensity and/or CO₂ concentration and was always highest at the biofilm surface; the stimulating effect of elevated CO₂ concentration in the gas phase on photosynthesis was enhanced by higher light intensities. The dissolved inorganic carbon concentration profiles suggest that the availability of the dissolved free CO₂ has the strongest impact on photosynthetic productivity. The results suggest that dark respiration could explain previously observed decrease in growth rate over cultivation time in this type of PSBR.

Our results represent a basis for understanding the complex dynamics of environmental variables and metabolic processes in artificial phototrophic biofilms exposed to a gas phase and can be used to improve the design and operational parameters of PSBRs.

Key words

Phototrophic biofilm, Microalgae, Photobioreactor, Microsensor, Photosynthesis parameters

Introduction

Recent advances in the design of bioreactors for the cultivation of microalgae have focussed on immobilization of these organisms as a phototrophic biofilm (reviewed by Berner et al. 2014; Olivieri et al. 2014; Gross et al. 2015). An efficient type of biofilm bioreactor features a non-submerged biofilm exposed directly to light and a gas phase, and separated from the bulk of culture medium by a fine-porous sheet-like material (e.g., a microporous membrane). The principle of this biofilm bioreactor was introduced a decade ago by Podola and Melkonian (2003) as the 'Twin-Layer' (Nowack et al. 2005), now referred to in a more general term as 'porous substrate bioreactors' (PSBRs) (Murphy and Berberoglu 2014). Compared to other photobioreactors (e.g. suspension systems), PSBRs display fundamental advantages with respect to: biomass harvesting, water/energy consumption, the spectrum of applications and microalgal species that can be cultivated (e.g. Shi et al., 2007; Naumann et al., 2013; Benstein et al. 2014). Recent studies using high light and CO₂ levels and/or modifications of reactor design have documented high biomass productivities in PSBRs with respect to reactor surface (> 30 g *dw* m⁻² d⁻¹; Schultze et al., 2015) as well as ground area (> 60 g *dw* m⁻² d⁻¹; Liu et al., 2013; Zhang et al., 2015). Also, research performed at pilot-scale (Naumann et al., 2013; Shi et al., 2014; Zhang et al., 2015) demonstrated the proof of principle and a significant potential of PSBRs for commercial-scale microalgal cultivation.

Consequently, as a next step, more basic research is required to understand the dynamics of light, mass transfer, productivity and their interactions in PSBR biofilms, since this knowledge is essential for a systematic and comprehensive approach aimed at process optimization. For example, in several studies continuously decreasing growth rates were reported during prolonged cultivation of microalgae in PSBRs (Cheng et al., 2013; Ji et al., 2014; Schultze et al., 2015), possibly caused by respiration in the increasingly thicker non-illuminated zones of a growing biofilm. Moreover, oxygen generated by photosynthesis can accumulate to high concentrations in photosynthetic biofilms (Pringault and Garcia-Pichel, 2000). In suspension cultures oxygen accumulation is known to inhibit net productivity by photorespiration, or by formation of free oxygen radicals.

Several high resolution methods (Revsbech and Jorgensen, 1983; Wolf et al., 2007; Kühl and Polerecky, 2008; Ramanan et al., 2013) have been used on natural biofilms. Among others, microsensors offer direct, non-destructive and in-situ measurements and have been successfully applied to submerged phototrophic biofilms since the 1980s (Revsbech, 1983; Revsbech and Jorgensen, 1983). The data from these previous studies, however, does not apply to non-submerged artificial biofilms due to significant differences in the system setup (e.g. direct exposure to the atmosphere) and species composition of the biofilm (Tampion and Tampion, 1987; Podola and Melkonian, 2003; Seckbach and Oren, 2010).

In this study, we used microsensors to acquire depth profiles of light, oxygen, pH, and photosynthetic activity inside a highly active biofilm of the green alga *Halochlorella rubescens*, cultivated in a Twin-Layer PSBR at different light intensities and exposed to various CO_2 levels. The data obtained were used to describe the dynamics in a PSBR for the first time in detail. Based on the results obtained, we discuss the specific strengths of PSBRs and identify parameters for further optimization in these novel photobioreactors.

Materials and Methods

Microalgal cultivation

A suspension culture of the green alga *Halochlorella rubescens* (CCAC 0126; Culture Collection of Algae at the University of Cologne; www.ccac.uni-koeln.de) was used for biofilm inoculation and cultivation according to Schultze et al. (2015). The same strain has been used in previous studies, and is known to form a fast-growing phototrophic biofilm (Schultze et al., 2015). After inoculation, biofilms were pre-cultivated with a PAR light intensity of 300 µmol photons $m^{-2} s^{-1}$ at the biofilm surface. After 14 days, filters were subjected to different PAR light intensities of 50, 300, and 1,000 µmol photons $m^{-2} s^{-1}$ for 10 days. Microsensor measurements were then also conducted at these three light intensities. For cultivation and during measurements the temperature was kept at 23 ± 1 °C, and Bold's Basal Medium (BBM; Bischoff and Bold, 1963) was used.

Microsensor measurements setup

The experimental setup and calibration of the microsensor system is based on the work of Gieseke and de Beer (2004) and the experimental setup is described in detail in the supplementary materials (Li et al., 2015). Oxygen microsensors with tip diameters $< 10 \ \mu m$ and a response time ≤ 0.5 s were build and used as described previously (accurate to µM range, and with a stirring effect of less than 4%; Revsbech, 1989). Liquid ion exchange (LIX) type pH microsensors with accuracy of more than 0.01 pH unit and measurement range from pH 3 to 11 were used in the present study to measure pH profiles (Lee and de Beer, 1995). pH microsensors were calibrated according to de Beer et al. (2008). Irradiance profiles were recorded for all three incident light intensities with spherical tip irradiance microsensor with a tip diameter of 80 µm (Lassen et al, 1992), which was connected to a spectrophotometer (USB 4000, Ocean optics, Dunedin, USA). The measured irradiance data was integrated over wavelength bands to give estimations of the total, red, green and blue PAR intensities (400 - 700, 400 - 500, 500 -600 and 600 – 700 nm, respectively). For each incident irradiance, the concentration of dissolved oxygen ([DO]), the pH and the photosynthetic activity profiles were recorded for different CO₂ levels in aeration applied onto the biofilm surface: 0.039 % (dry ambient air), and 1%, 3%, and 5 % CO₂ (dry ambient air with supplementary CO₂ v/v). When CO₂ levels were changed, biofilms in the measurement cell were adapted for 15 min prior to the microsensor measurements (which was sufficient for the biofilm to reach steady state, as measurements taken between 5 min to 1 hour after measurement conditions had been changed have produced profiles with no significant difference). Profiles were also recorded in biofilms subjected to darkness and aerated with dry ambient air (dark measurement). Prior to dark measurements, biofilms were incubated in the measurement cell for 3 hours without illumination.

All microsensor measurements were carried out with a 20 μ m depth resolution. [DO] and pH profiles were acquired from 3 positions on the same biofilm sample (triplicates). Light and gross photosynthetic productivity profiles were acquired by using the method described in the supplementary materials (Li et al., 2015). Gross productivity measurements were carried out until a depth at which the shift from light to dark did not have significant effects on the [DO] within the measurement period (200, 400 and 500 μ m for biofilms exposed to 50, 300 and 1,000 μ mol m⁻² s^{-1} , respectively). Attenuation coefficients (extinction per unit depth, in mm⁻¹) of the PAR in the biofilms were calculated as described by Lassen et al. (1992).

Calculation for net productivity and dissolved inorganic carbon profiles

Net photosynthetic productivity was calculated from the steady state [DO] profiles using the mass balance equation (Bergman et al., 2011, Revsbech and Jorgensen, 1983):

$$P_n = -D_{e,O_2} \frac{\partial^2 [\text{DO}]}{\partial x^2} - u \frac{\partial [\text{DO}]}{\partial x}$$
(Eq. 1)

 P_n is the net productivity, D_{e,o^2} is the effective diffusion coefficient of oxygen in biofilm, x is the depth and u is the convective flow rate in the biofilm perpendicular to its surface (due to surface evaporation). D_{e,o^2} and u were estimated from the measured biofilm porosity and the measured evaporation rate using the technique as described in supplementary materials (Li et al., 2015). The measurement showed no significant difference between biofilms exposed to different PAR light intensities and at different depths (Two-way ANOVA test). Porosity was measured to be 0.76 in *the H. rubescens* Twin-Layer biofilm, and D_{e,o^2} was calculated to be 0.68 of that in pure water, u was calculated to be 5.4 µm s⁻¹ (Weissberg, 1963). To acquire the derivatives needed in Eq.1, 6th order polynomial curves were fitted to the steady state [DO] profiles (R² values ≥ 0.98 ; *curve fitting toolbox*, MATLAB, version 2013a, MathWorks, Ismaning, Germany).The derivatives were then calculated from the polynomial curves (de Beer, 2001). The thickness of the biofilms was estimated using a hand microtome as described in supplementary materials (Li et al., 2015). All statistical analysis were performed using GraphPad Prism v5.02 (GraphPad Software Inc., La Jolla, USA). For calculation, 1.99×10^{-9} m² s⁻¹ was taken as diffusion coefficient of oxygen in water (Han and Bartels, 1996).

An estimation of the distribution of dissolved inorganic carbon concentrations ([DIC]) can be calculated using the DIC consumption rate distribution together with the CO_2 concentration in the gas phase and the measured pH profiles: Assuming DIC consumption to oxygen production has a 1:1 ratio (mole to mole), the photosynthetic productivity profiles can be used to estimate the consumption rate distribution of the [DIC]. Using the equation provided by Morel and Hering (1993), the total [DIC] at the biofilm surface can be calculated as (considering that the hydration of CO_2 at the biofilm surface is instantaneous, which is the ideal case):

$$DIC_{surface} = H_{CO_2} \cdot p_{CO_2} \cdot \left(1 + \frac{\kappa_{a_1}}{10^{-pH_{surface}}} + \frac{\kappa_{a_1} \kappa_{a_2}}{\left(10^{-pH_{surface}}\right)^2} \right)$$
(Eq.2),

 H_{CO2} is the Henry's constant for CO₂, and has a value of 0.304 mM kpa⁻¹; p_{CO2} is the CO₂ partial pressure in the gas phase, K_{a1} and K_{a2} are the dissociation coefficients of carbonic acid and bicarbonate ion, with value of 10^{-6.3} and 10^{-10.3}, respectively; $pH_{surface}$ is the measured pH value at the biofilm surface (Murphy and Berberoglu, 2014; Wolf et al., 2007). Distribution of the total [DIC] at depth x can be acquired from a transformation of mass balance equation:

$$[DIC]_x = [DIC]_{surface} - \frac{\int_0^x P_n dx}{\left(\frac{D_{e,DIC}}{x}\right)}$$
(Eq.3),
$D_{e,DIC}$ is the effective diffusion coefficient of DICs in biofilm, and was assumed to equal the value of bicarbonate ions, which has a value of 1.18×10^{-9} m² s⁻¹ in pure water (Murphy and Berberoglu, 2014). DIC speciation at depth *x* was calculated from the measured pH profiles:

$$\begin{cases} [CO2]_{x} = [DIC]_{x} \cdot \frac{1}{1 + \frac{K_{a1}}{10^{-PH_{x}}} + \frac{K_{a1}K_{a2}}{(10^{-PH_{x}})^{2}}} \\ [HCO3^{-}]_{x} = [DIC]_{x} \cdot \frac{K_{a1}/10^{-PH_{x}}}{1 + \frac{K_{a1}}{10^{-PH_{x}}} + \frac{K_{a1}K_{a2}}{(10^{-PH_{x}})^{2}}} \\ [CO3^{2^{-}}]_{x} = [DIC]_{x} \cdot \frac{K_{a1}K_{a2}/(10^{-PH_{x}})^{2}}{1 + \frac{K_{a1}K_{a2}}{10^{-PH_{x}}} + \frac{K_{a1}K_{a2}}{(10^{-PH_{x}})^{2}}} \end{cases}$$
(Eq.4).

Results

Biofilm thickness and appearance

The thickness of the biofilms grown at different light intensities was significantly different: around 320, 700 and 800 μ m for biofilms exposed to 50, 300 and 1000 μ mol photons m⁻² s⁻¹, respectively. Visually, no color difference between the biofilms exposed to different light intensities could be observed.

Light

PAR profiles in *H. rubescens* biofilms exposed to different light intensities as the percentage of the total PAR measured at the biofilm surface and the calculated attenuation coefficients are presented in Fig. 1. Since spherical irradiance sensors with tip diameters of 80 μ m were used, the irradiances of the first 80 μ m were overestimated (Fig. 1), as the tip was not fully immersed in the biofilm. The PAR distributions from biofilms exposed to 50, 300 and 1000 μ mol photons m⁻² s⁻¹ exhibit nearly identical trends: A relatively sharp decrease from the surface to about 250 μ m depth, followed by a smoother reduction to less than 1% of the surface irradiance at a depth of around 350 μ m. Irradiance in the blue, green and red ranges of the PAR also behaved similarly in biofilms exposed to different PAR intensities at the surface. In all cases, the intensity of blue light at the biofilm surface was the lowest and the first to be absorbed completely by the biomass. Attenuation coefficients showed similar tendencies for biofilms exposed to different light intensities: An increase from the biofilm surface to a maximal value (at around 200 μ m), followed by a decrease. However, biofilms exposed to lower irradiance were observed to have higher attenuation coefficients (both the measured maximum and the average). Also, in Fig. 1, a 'dark zone' (non-illuminated region) can be observed in thicker biofilms (biofilm cultivated at 300 or 1000 μ m0 m⁻² s⁻¹).



Figure 1: Intensity and attenuation coefficient of photosynthetically active radiation (PAR) in *Halochlorella rubescens* biofilms exposed to different light intensities. A: Biomass exposed to 50 μ mol photons $m^2 s^{-1}$; B: Biomass exposed to 300 μ mol photons $m^2 s^{-1}$; C: Biomass exposed to 1000 μ mol photons $m^2 s^{-1}$. Intensities are presented in percentage of the total irradiance measured at the biofilm surface. Total PAR intensity, blue (400 to 500 nm), green (500 to 600 nm) and red (600 to 700 nm) are represented by different line styles; the lines represent average values measured in a 3 second period. Bars represent the attenuation coefficient of the complete PAR range. Lengths of x-axis indicates the thickness of the biofilm. The thin solid grey lines show the light distribution profile predicted by Lambert-Beer law with a fixed attenuation coefficient (denoted as LB-curve, and the absorption coefficient used for the calculation of the LB-curve was selected so, that the LB-curve have the same penetration depth as the measured profile). The grey areas indicate the depth in which the tip of the sensor was not fully immersed in the biofilm.

Oxygen and pH

[DO] profiles measured under illumination displayed similar shapes for all incident PAR and CO₂ levels (Fig. 2): Increasing from air saturation (0.274 mM) at the biofilm surface to a maximum at a depth between 50 and 150 μ m, depending on light intensity and CO₂ levels. The maximum was followed by a decrease to values lower than the [DO] at the biofilm surface (from 180, 460 and 630 μ m inwards for biofilms exposed to 50, 300 and 1000 μ mol photons m⁻² s⁻¹, respectively). The [DO] and its maximal value increased with light intensities and/or with an increase of CO₂ in the gas phase. The effect of CO₂ was more prominent when combined with higher PAR light intensities. The highest [DO] of 3.2 mM was measured in biofilms exposed to 1000 μ mol photons m⁻² s⁻¹ aerated with 5% supplementary CO₂ (Fig. 2).



Figure 2: Concentration of dissolved oxygen ([DO]) profiles in *Halochlorella rubescens* biofilms exposed to different light intensities and areated with different carbon dioxide concentrations A: Biomass exposed to 50 µmol photons $m^2 s^4$; B: Biomass exposed to 300 µmol photons $m^2 s^4$; C: Biomass exposed to 1000 photons µmol $m^2 s^4$. The marks and the error bars represent the means and standard deviations of triplicates; dotted line represents [DO] in BBM (Bold's Basal Medium) saturated with air. Lengths of x-axis indicates the thickness of the biofilm. Note the differences in oxygen concentration scales (y-axis) for different light intensities.

Fig. 3 shows pH profiles measured in illuminated biofilms. With 50 μ mol photons m⁻² s⁻¹ PAR light intensity, the pH profiles displayed a maximum at the biofilm surface and decreased with depth. In biofilms exposed to 300 and 1,000 μ mol photons m⁻² s⁻¹, however, the pH first increased with depth, resulting in a maximum between 100 and 200 μ m. Stronger PAR light intensities lead to higher pH values, whereas higher CO₂ concentrations reduced the

pH (Fig. 3). Furthermore, high CO₂ concentrations smoothened the peaks of the pH profiles in biofilms exposed to 300 and 1000 μ mol photons m⁻² s⁻¹. The maximal pH value measured was more than 10 (at 1,000 μ mol photons m⁻² s⁻¹ with no additional CO₂), and was much higher than the pH value measured at the interface between biomass and support membrane (i.e. at the bottom) of the same biofilm.



Figure 3: pH profiles in *Halochlorella rubescens* biofilms exposed to different light intensities and aerated with different carbon dioxide concentrations. A: Biomass exposed to $50 \ \mu mol m^2 s^1$, B: Biomass exposed to $300 \ \mu mol photons m^2 s^1$; C: Biomass exposed to $1000 \ photons \ \mu mol m^2 s^1$. Biomass exposed to $1000 \ photons$ multiplicates; dotted line represents pH value of freshly prepared BbM (Bold's Basal Medium). Lengths of x-axis indicates the thickness of the biofilm.

Fig. 4 presents O_2 and pH profiles in biofilms in the absence of light and supplementary CO_2 . Without illumination, the [DO] decreased with depth (Fig. 4 A). In thicker biofilms (biofilm pre-cultivated at 300 and 1000 µmol photons $m^{-2} s^{-1}$) this decrease reached values of nitrogen gas saturated medium (0 mM) at a depth of around 330 µm and was followed by an increase in deeper layers from 400 µm on. The pH values in the biofilm in the dark did not change with depth and were around 6.8 for all measured depths (Fig. 4 B).



Figure 4: A: Dark oxygen profiles in *Halochlorella rubescens* biofilms exposed to different light intensities during cultivation and aerated with dry ambient air. Dotted line represents [DO] in BBM (Bold's Basal Medium) saturated with air. B: Dark plf profiles in *H. rubescens* biofilms exposed to different light intensities during cultivation and aerated with dry ambient air. Dashed line represents the pH value of freshly prepared BBM. The marks and the error bars represent the means and standard deviations of triplicates.

Photosynthetic activity

Fig. 5 describes the gross and the net productivity in the Twin-Layer biofilm. The trends of gross and net photosynthetic productivities were similar for all light intensities and CO_2 levels: A productivity maximum was found at the biofilm surface, which then decreased with depth. The net productivity dropped below 0 μ mol oxygen m⁻³ s⁻¹ in the deeper parts of the biofilms, where respiration was higher than gross photosynthesis. Increase in light intensity always led to an increase in both the maximal productivity and the total productivity (estimated from the total area under the productivity curve, Fig. 5). In contrast, an increase in CO₂ concentration alone did not necessarily lead to an increase in productivity. E.g. for biofilm exposed to 300 μ mol m⁻² s⁻¹ an increase from 3% to

5% CO₂ did not lead to an increase in productivity. Generally, the positive effects of CO₂ concentrations on both the gross and net photosynthetic productivities were more pronounced at higher light intensities.

Dissolved inorganic carbon

The calculated [DIC] distributions are given in Fig. 6. Both light intensity and CO₂ in the gas phase influenced the total [DIC] and DIC speciation in the *H. rubescens* biofilms. Total [DIC] in biofilms aerated with additional CO₂ (> 1% v/v) remained almost constant along the depth gradient. Bicarbonate was the dominant DIC species in all measured biofilms, whereas carbonate was only significantly preset in biofilms subjected to high light intensity and low gas phase CO₂ levels (e.g. 1000 µmol photons $m^{-2} s^{-1}$ and normal air). For all measured biofilms subjected to 300 µmol photons $m^{-2} s^{-1}$ and aerated with normal air, the total [DIC] dropped to almost 0 mM at around 200 µm depth. Generally, a higher gas phase CO₂ concentration led to higher dissolved free CO₂ levels in the biofilms, but not necessarily to a higher total [DIC] (3% and 5% CO₂ with 1000 µmol photons $m^{-2} s^{-1}$).



Figure 5: Photosynthetic productivity (measured as rates of change of [DO], for details, refer to the method for measuring gross productivity in the supplementary materials) profiles in *Halochlorella rubescens* biofilms exposed to different light intensities and aerated with different carbon dioxide concentrations. Solid line represents the corrected gross productivity; Dotted line gives the calculated net productivity (for graph clarity, SDs were calculated but not shown). The difference between gross productivity and net productivity gives light respiration. Lengths of x-axis indicates the thickness of the biofilm. Light intensities and aeration CO₂ concentrations are marked at the top of each panel (µmol photons $m^2 s^4 / % v/v$). Note the differences in photosynthetic productivity scales (y-axis) for different light intensities.



Figure 6: Calculated dissolved inorganic carbon concentration ([DIC]) distribution profiles in *Halochlorella rubescens* biofilms exposed to different light intensities and aerated with different carbon dioxide concentrations. Solid line represents the total [DIC] concentration; the speciation of DIC is represented by the bands under the solid line (Dark grey, light grey and white for free CO₂, bicarbonate and carbonate, respectively). Lengths of x-axis indicates the thickness of the biofilm. Light intensities and aeration CO2 concentrations are marked at the top of each panel (µmol photons $m^2 s^{-1} / \%$ v(v). The arrows give the position and value of the lowest free CO₂ concentration in biofilm. Note the differences in concentrations scales (y-axis) for different panels; carbonate bands (white) are invisible in most of the panels due to very low concentrations.

Discussion

Light distribution as the driving factor for photosynthetic productivity

In the *H. rubescens* biofilms, light profiles did not follow the log distribution of the Lambert-Beer absorption curve (Fig. 1). In addition, no significant difference in the distribution of biomass density along the depth profile was found (see Materials and Methods). Wang et al. (2015) observed that the mass extinction (extinction per unit biomass) was lower in an algal biofilm than in suspension culture. These observations indicate that forward scattering of light by cells and/or adaptation of cellular pigment content plays an important role in light transfer in phototrophic biofilms, reducing the attenuation of light with depth (Kühl et al., 1994; Pringault and Garcia-Pichel, 2000; Berberoglu et al., 2007). As a result, light penetrates deeper into the biofilm, leading to a higher light dilution rate (photons available per cell) in the biofilm, which, in general, is favorable for photobioreactors (e.g. Olivieri et al., 2014). Our measured irradiance profiles showed similar trends for biofilms exposed to different light intensities, however, differences can be observed in the attenuation coefficients (Fig 1): The maximal attenuation coefficients decreased with increasing light intensity, which indicates that a more efficient light uptake was achieved by biofilms exposed to lower light intensities. This suggests an adaptation of the biomass to the light intensity it was exposed to, as it is well known that algal cells can adapt to different light intensities by regulating their pigment content (e.g. Jørgensen, 1969; Richardson et al., 1983). The observed increase of the attenuation coefficient with depth within about 200 µm from the biofilm surface can be a combined effect of this adaptation and the forward

scattering of light. Low irradiances in the deeper parts of the biofilm may lead to pigment degradation, and thus to lower attenuation coefficients. Attenuation distributions with similar trends have been observed by Pringault and Garcia-Pichel (2000) in a submerged artificial cyanobaterial biofilm.

Interactions between photosynthesis and chemical parameters

Since the results showed that the [DO] at the surface of all the measured biofilms were not significantly different from air saturated medium, it was assumed that [DO] at the surface of all biofilms was in equilibrium with the ambient gas phase. This supports the assumption of the existence of only a very thin mass transfer boundary layer in the gas phase adjacent to the biofilm surface that should only offer little resistance to gas transfer (Schlichting and Gersten, 2000). [DO] in the *H. rubescens* biofilm was controlled by photosynthesis, respiration, diffusion, advection inside the biofilm and the removal of dissolved gaseous species at the biofilm surface. Near the surface, [DO] increased with depth. This is due to an excess in production compared to consumption (see Eq. 1). At depths below the [DO] maximum, consumption of oxygen due to respiration and a reduced production (less light and/or CO₂) in the deeper parts of the biofilm led to decreasing [DO] with depth. Similar trends have been observed in submerged phototrophic biofilms (Glud et al., 1992; Pringault and Garcia-Pichel, 2000; Bernstein et al., 2014).

Despite the extremely thin boundary layer, and thus an effective exchange between the gas phase and the biofilm, high oxygen concentrations of up to 3.2 mM were observed in the biofilm (Fig. 2). This is more than 11 times higher than the [DO] in medium (BBM) saturated with air, and is equivalent to a partial pressure of oxygen of 2.4 atm. [DO] reaching a 6-fold concentration of air-saturated water has been measured by Pringault and Garcia-Pichel (2000) in a submerged artificial cyanobacterial biofilm exposed to similar light intensities. How biofilms and mats can establish this oversaturation, and why no development of gas bubbles was observed in the biofilm used in this study needs to be investigated further. Such high oxygen levels should inhibit net photosynthetic productivity, as they favor photorespiration (Spreitzer and Salvucci, 2002). This detrimental effect could be counteracted by carbon concentrating mechanisms (CCM), which elevate the CO_2/O_2 ratio around the RuBisCO (review by Raven et al., 2008).

In biofilms subjected to darkness, oxygen concentrations inside the biofilm were lower than at the surface of the biofilm, presumably due to respiration (Fig. 4). The slight increase of [DO] near the interface between the biofilm and substrate layer observed in thicker biofilms (biofilms grown at 300 and 1000 μ mol photons m⁻² s⁻¹, respectively) can be attributed to the dissolved oxygen supplied from culture medium through the membrane. This was presumably also responsible for the lower [DO] gradients observed in biofilms grown at 50 μ mol photons m⁻² s⁻¹ during cultivation. During dark periods, biomass was consumed rather than produced. The loss of productivity due to respiration in the dark period is a known and undesired feature of microalgal mass cultivation (e. g. Grobbelaar and Soeder, 1985), and is particularly pronounced when large suspension volumes are used (e. g. in a raceway pond): the suspension stores the heat absorbed during the day, and due to the large liquid volume, the stored heat is released only relatively slowly (i.e. the temperature remains high for a relatively longer period after the onset of darkness), and thus resulting in higher respiration rates during the night. Since the lowest temperatures in open pond cultivation occur during the morning hours when light is already available for photosynthesis,

suboptimal photosynthetic productivity characterizes the performance of such systems during the first half of the day. Here, a biofilm-based bioreactor might be advantageous by offering a 100 to 1,000-fold lower water content compared to a suspension culture, leading to a significantly faster adoption of ambient temperature by the biomass. Another approach to increase productivity could be the application of continuous illumination. However, artificial illumination for biomass production is clearly uneconomical at a technical scale (e.g., Blanken et al., 2013).

The pH is an important variable regulating biomass productivity in phototrophic biofilms as it controls inorganic carbon speciation (free CO₂, bicarbonate and carbonate; Fig. 6). The pH, has been shown to influence the affinity of algal cells to free CO₂ (Azov, 1982). A high pH value represents a low dissolved free CO₂ concentration, as dissolved free CO₂ can only exist in large quantities in solutions with a pH value lower than 7.5. A further increase in pH in combination with high net oxygen production thus indicates uptake of bicarbonate ions for photosynthesis: at elevated pH, as CO₂ level decreases, it is more likely that organisms rely on uptake of bicarbonate (Fig. 6). Thus the role of carbonic anhydrase, that catalyzes the rapid interconversion of carbon dioxide to carbonic acid, is presumably important for DIC uptake in such biofilms. The uptake of nitrate and phosphate during growth can also elevate the pH. However, due to a slower uptake, the effect can be considered insignificant when compared to the effect of inorganic carbon uptake on pH.

For light intensities of 50 and 300 µmol photons m⁻² s⁻¹, photosynthesis did not increase further at higher CO₂ levels (e.g. from 3% to 5% CO₂ at 300 µmol m^{-2} s⁻¹; Fig. 2), indicating that light rather than CO₂ was limiting photosynthesis. At 1000 µmol m⁻² s⁻¹, increasing CO₂ from 3% to 5% did indeed lead to a further increase in photosynthetic activity, showing that the CO₂ supply should be tuned to light intensity to optimize the harvesting of light energy into biomass. In the dark, pH values were stable across the whole biofilm for biofilms exposed to all three light intensities during their cultivation period. This indicates that CO₂ production through respiration during the dark period was not strong enough to influence the local pH, and the phosphate buffer supplied in the medium was sufficient to stabilize the pH. In the 'dark zone' of a light exposed biofilm however, an elevation in pH above the value of the medium was observed. This, instead of being caused by direct CO₂ and/or HCO₃⁻ consumption by the biomass, may be caused by the shifting of the chemical equilibrium due to transport of CO_2 and/or HCO_3^- to the phototrophically active region, connected with the transport of protons in the opposite direction. With additional CO₂ in the gas phase, the total [DIC] stayed stable along the depth gradient of the biofilms. This indicates that with high CO₂ concentration in the gas phase (> 1% v/v), diffusion was not limiting the supply of DIC to the deeper part of the biofilm. However, the dissolved free CO₂ concentration changed significantly along the depth gradient and with the gas phase CO₂ levels. A higher dissolved free CO₂ resulted in higher productivity, even though a lower total [DIC] was observed (Fig. 5 and 6; biofilms exposed to1000 µmol photons m⁻² s⁻¹, compare 3% and 5% CO₂). This shows that although the algal cells can utilize bicarbonate for photosynthesis, this is not as efficient as CO₂ assimilation (reviewed by Singh, 2014). By elevating the CO_2 level in the gas phase, the dissolved free CO_2 in the biofilm is increased through an increase in total [DIC] in the biofilm and/or by decreasing the pH in the biofilm.

Biological efficiency and photosynthetic productivity

Gross photosynthetic productivity decreased continuously with depth. Higher light intensities increased the maximum gross productivity, which was found at the surface of all biofilms (Fig. 5). Total gross photosynthetic productivity also increased with higher light intensity due to the increase in the maximum gross productivity as well as the increase in gross productivity in the deeper layers of the biofilm. The latter was caused by deeper light penetration and/or a higher light intensity at a given depth (Fig. 1). Gross photosynthetic productivity is a useful term for evaluating the functional status of a phototrophic biofilm (Gieseke and de Beer, 2004). However, it does not provide information on biomass net productivity, which is an important parameter for evaluating the overall performance of a phototrophic biofilm. Net photosynthetic productivity can be used for such estimations (Gieseke and de Beer, 2004). In a highly active biofilm grown in a porous substrate bioreactor under high light and CO₂ regimes, photorespiration may consume significant amounts of the oxygen as well as biomass produced by photosynthesis considering the very high [DO] (Fig. 2) (Pope, 1975; Birmingham et al., 1982; Larkum et al., 2003). As a result, the net photosynthetic productivities in such biofilms were lower compared to gross photosynthetic activities measured (Fig. 5). Our results show that both the net- and gross productivity integrated over the whole photosynthetic biofilm increased with light intensity, especially when supplementary CO₂ was available. However, when considering a realistic cultivation scenario, the effect of high light intensity and high CO₂ concentration on total biomass productivity would be much more prominent at the start of biofilm cultivation: The increase of the 'dark zone' with time (with biofilm thickness increase) would reduce the net productivity of the biofilm as a whole. Net productivity in these 'dark zones' was reduced to negative values, likely due to respiration. Moreover, for all biofilms, the net productivity dropped to negative values whenever the PAR intensity was below 25 µmol m-2 s-1 ('low light zones', Fig 1 and 5). In these 'low light zones', oxygen produced by photosynthesis was less than the consumption by respiration. This leads to biomass loss in both the 'low light zones' and the 'dark zones', and as a result, a reduction of the net productivity of the biofilm as a whole. This information can be used to determine the time point of biomass harvest, to maintain optimal productivity of the photobioreactor.

Interpretation of the calculated gross productivity data must be made with caution, because the calculation relies on an estimated effective diffusion coefficient of oxygen in the biofilm, which can be a source of error. For the calculation of the net productivity, in addition to the diffusion coefficients, a 6th order polynomial fit can be another source of inaccuracy, even though the R^2 value is high. In the studied biofilm system, the maximum net photosynthetic productivities were always found on the biofilm surface. In submerged biofilms, however, maxima usually occur deeper within the biofilms (Glud et al., 1992;Lassen et al., 1998; Pringault and Garcia-Pichel, 2000). This may be explained by the direct exposure of a non-submerged biofilm to the gas phase with a relatively high gas flow rate. Compared to a submerged surface, a non-submerged biofilm allows much faster removal rates of dissolved oxygen (produced in the biofilm) from its surface due to the very thin mass transfer/flow boundary layer (Schlichting and Gersten, 2000; Bergman et al., 2011). Compared to the deeper parts, cells near the biofilm surface of a non-submerged system had higher gross productivity, but were exposed to lower [DO], which leads to lower photorespiration rates (Larkum et al., 2003). Also, a higher dissolved free CO₂ was observed near the biofilm surface (Fig. 6). This resulted in a higher net photosynthetic productivity near the biofilm surface. The increase in the net productivity (from negative values towards zero) near the bottom of the biofilm, as observed in thicker biofilms could be explained by limiting dark respiration due to very low oxygen concentrations in this part of the biofilm (Fig. 5).

Conclusions

The present study, using a highly productive *Halochlorella rubescens* biofilm grown in a Twin-Layer porous substrate photobioreactor (TL-PSBR), provides the first in-depth analysis of dynamic processes within such biofilms employing microsensors. The knowledge gained on the distribution of light, oxygen, pH and photosynthetic productivity within the Twin-Layer biofilms provides a sound basis to improve the design of this PSBR towards optimizing microalgal biomass productivity. The present study confirmed the existence of a 'dark zone' in PSBRs subjected to prolonged cultivation, and that its presence can reduce the productivity of PSBRs. However, a dedicated harvesting scheme can be used to minimize this problem (i.e. harvest before the biofilm reaches a thickness to develop 'dark zones'). Furthermore, the data presented could facilitate the modeling of growth in PSBR biofilms in general.

Acknowledgements

This study was supported by the University of Cologne (KST 158901001). The authors thank Meike Hahn for her assistance in experimental work.

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Supplementary materials

The method descrided here are described in detail by Li et al. (2015), for more detailed information, please refer to their publication.

Estimation of evaporation rate and effective diffusion coefficient in biofilm

The porosity of the biofilm (θ) was estimated from the volume of cells (biovolume) in the biofilm divided by the total biofilm volume. The biovolume was calculated as described by Hillebrand et al. (1999) from suspended biofilm samples with a 50 µm resolution by means of a custom-made hand microtome (Fig. S1). The biofilm immobilized on a polycarbonate membrane was placed onto a height-adjustable stage on a thin layer of culture medium. The stage allows levelling the biofilm sample into the sectioning plane by means of a micrometer screw. The sectioning plane was defined by a solid brass support over which a rigid microtome blade was passed manually to remove the overlying algal biofilm (Fig. S1).

Evaporation rate during the measurement was 0.68 mL min⁻¹ (monitored by recording the change of the volume in the medium container, Fig S2), which corresponded to a convective flow rate of 5.4 μ m s⁻¹ inside the biofilm, which was calculated as:

$$u = q_e / (A_e \cdot \theta) \tag{Eq.S1},$$

 q_e is the rate of liquid volume lost, A_e is the exposed surface area of the biofilm.

Microsensor setup

A schematic representation of the experimental setup for microsensor measurements based on previous works by Gieseke and de Beer (2004) is given in Fig. S2: the biofilm immobilized on a polycarbonate membrane was placed onto a moist glass fiber inside the biofilm measurement cell under a controlled atmosphere of compressed air at a flow rate of 1 L min⁻¹ (Fig. S2A). The non-inoculated area of the polycarbonate filter and glass fiber were covered with a black plastic foil to exclude light and gas exchange in this area. 50 mL of BBM culture medium were circulated though the measurement cell with a flow rate of 3 mL min⁻¹ by means of a peristaltic pump and were exchanged every 1 hour during the measurement. Light was supplied from the front side by a halogen lamp (KL-1500, Schott, Mainz, Germany) equipped with a 3-fold splitter to ensure even distribution. The movement of the sensor was enabled by a computer-controlled micromanipulator (Pollux Drive, PI miCos, Eschbach, Germany).





Microsensor signals were amplified and converted into digital data (DAQpad 6015 and 6009, National Instruments,

Munich, Germany) prior to their further processing on a computer (Fig. S2B). Software used for system control and data acquisition was custom made (can be acquired upon request from Dr. Lubos Polerecky, Utrecht University, Netherlands, l.polerecky@uu.nl). Light-dark shift measurements (Revsbech, 1989) were carried out, and the data acquired was processed as described in the following section. Linear calibration of oxygen microsensor was carried out as described by Revsbech (1983) using BBM medium saturated with nitrogen gas or compressed air. A shutter connected to a timer was used to control light-dark cycling: the total dark period was 3 s, and a linear regression slope of oxygen concentration change between 0.7 and 2.1 s of the dark period was determined as the measured value. For each measurement depth, three light-dark cycles were performed after steady state was reached (this could be as long as 10 min), and these 3 values acquired were considered as triplicates. The PAR light intensity of a fixed depth was determined as the average value of a 3 s measurement period.



Figure S2: Experimental setup for microsensor studies in non-submerged microalgal biofilms. A: Schematic drawing of the vertical cross section of the measurement in artificial biofilms. Solid arrows in the glass fiber tissue indicate flow direction of the culture medium. B: Schematic drawing of the complete experimental setup, dotted lines with arrow indicate signal flow, whereas solid arrows represent medium and gas flows in the system.

Gross photosynthetic productivity calculation

Take the derivative of time on both sides of the dissolved oxygen mass transfer equation, and assume the time t and spatial variables x are not dependent (as during the measurement period, the thickness of the biofilm hardly changes) (Bergman et al., 2011, Revsbech and Jorgensen, 1983):

$$\partial (\frac{\partial C}{\partial t})/\partial t = 0/\partial t - \partial R_l/\partial t + D_e \frac{\partial^2 (\frac{\partial C}{\partial t})}{\partial x^2} + u \frac{\partial (\frac{\partial C}{\partial t})}{\partial x} - \partial R_s/\partial t$$
(Eq.S2).

The term $\partial R_s/\partial t$ represents the rate change of the removal of dissolved gaseous species at the submerged biofilm surface due to the flow of bulk liquid, or, in this case, of the gas phase above the biofilm:

$$\partial R_s / \partial t = \left(D_e \frac{(C - C_s)}{x^2} + u \frac{(C - C_s)}{x} \right) / \partial t$$
$$= \left(\frac{D_e}{x^2} + \frac{u}{x} \right) \cdot \partial C / \partial t$$
(Eq.S3).

 C_s represents the oxygen concentration on the very surface of the aerial biofilm, which is in equilibrium with the phase above and is considered to be a constant. Considering that the rate of respiration remains stable during the measurement period (Glud et al., 1992), and take $\partial C/\partial t = P(x, t)$, as $\partial C/\partial t$ is a function of depth x and time t. Substitute Eq.S3 into $\partial R_s/\partial t$ and $\partial C/\partial t$ to P(x, t), Eq.S2 becomes:

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$$\frac{\partial P(x,t)}{\partial t} = 0 - 0 + D_e \frac{\partial^2 (P(x,t))}{\partial x^2} + u \frac{\partial (P(x,t))}{\partial x} - (\frac{D_e}{x^2} + \frac{u}{x}) \cdot P(x,t)$$
(Eq.S4)

Eq.S4 is a nonhomogeneous diffusion equation with a sink and can be solved by using any of the methods summarized by Crank or Polyanin (Crank, 1975; Polyanin, 2002). Using an initial condition P(x, t = 0) = P(x, 0) and a boundary condition $\partial P(x = 0, t)/\partial t = 0$, the exact solution of Eq.S4 at $t = \tau$ can be expressed as:

$$P(x,\tau) = \int_0^{\delta} P(y,0) \cdot \left[e^{\left(\frac{-(y-x+u\tau)}{4\cdot D_e \cdot \tau}\right)} - e^{\left(\frac{-(y+u\tau)}{4\cdot D_e \cdot \tau}\right)} \cdot erfc(\frac{x}{\sqrt{4\cdot D_e \cdot \tau}}) \right] \cdot dy$$
(Eq.S5),

 δ is the total depth of the biofilm, or, alternatively, the depth until which the shift from light to dark has an effect on the dissolved oxygen concentration. $P(x, \tau)$ is the measured value from the light-dark shift method with measurement time τ at measurement depth x, and P(y, 0) is the actual photosynthetic activity at position y (the integrating variable), which is the desired term. For measurements taken at all depths, the measured value $P(x, \tau)$ at a depth x depends on the P(y, 0) across the complete photosynthetically active region of the biofilm (surface to depth δ).

Eq.S5 is a Fredholm integral equation of the first kind (Kress, 2014), and its approximated solution can be calculated using Tikhonov regularization (Tikhonov et al., 1995) with a non-negative constrain applying the active-set algorithm purposed by Gill et al. (1981): Discrete δ by n, and let $P(x_j, \tau) = g_j$, $P(y_i, 0) = h_i$; (i, j = 1, 2, ..., n) and write Eq.S5 in the discrete operator notion.

$$g_j = \sum_i^n k_{i,j} \cdot h_i \tag{Eq.S6},$$

 $k_{i,j}$ is the operator term which contains both the term that reflects the effect of diffusion and advection of dissolved oxygen inside the biofilm (inner term, first term in square bracket in Eq.S5) and the term that reflects the effect of dissolved oxygen removal due to surface flow (surface term, second term in square bracket in Eq.S5). In matrix notion, let *G* be a vector containing g_{i} , *H* a vector containing h_{i} , and *K* a matrix containing the operator terms $k_{i,j}$:

$$G = K \cdot H$$
 (Eq. S7).

Rewrite Eq.S7 as:

$$(G_{in} + G_{surf}) = (K_{in} + K_{surf}) \cdot H$$
(Eq.S9),

The subscripts *in* and *surf* represent the effect from the inner term and the surface term in Eq.S5 respectively, operators K_{in} , K_{surf} can be calculated using Eq.S5, as well as g_{in} and G_{surf} , if an *H* is given. Apply the Tikhonov regularization, the problem in Eq.11 thus becomes:

$$\min_{F\geq 0}\left\{\left\|\left(G_{in}+G_{surf}\right)-\left(K_{in}+K_{surf}\right)\cdot H\right\|^{2}+\lambda^{2}\cdot\|L\cdot H\|^{2}\right\}$$

or,

$$\min_{F \ge 0} \left\{ \left\| \left(G_{in} - K_{in} \cdot H \right) + \left(G_{surf} - K_{surf} \cdot H \right) \right\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \right\}$$

(Eq.S10).

Observing the surface term in Eq.S5, notice the maximum value of the surface term is controlled by the position of the actual activity *y*, but a complement error function dependent solely on the position of the measurement taken (*x*) controls the final value. Furthermore, for a given *G*, under the same measurement conditions, G_{surf} always has the same set of values. Thus, for a given *G*, the minimization of the term $||G_{surf} - K_{surf} \cdot H||^2$ will always yield the same solution. As a result, the problem in Eq. S10 can be simplified to:

$$\min_{t \ge 0} \{ \|G_{in} - K_{in} \cdot H\|^2 + \lambda^2 \cdot \|L \cdot H\|^2 \}$$
(Eq.S11).

Eq.S11 does not contain the surface term, but the solution of the problem still leads to H, which is the desired original profile. The L-curve method purposed by Hansen & O'Leary (1993) was used to find the best regularization parameter λ .

The treatment procedure was coded in MATLAB (version 2013a, MathWorks, Ismaning, Germany), and the inbuilt *quadprog* was used for solving the minimization problems. *L-curve* function from the regularization tools (Hansen, 1994) was used to find the λ value. The mean values of triplicates acquired from the microsensor measurement were randomly added or subtracted with a random value in range of the calculated standard deviation (SD) at the same position. These values were used as input for the calculation. The calculation procedure was repeated 3 times, and the mean value of the 3 results was taken as the final result (a SD can also be calculated). Data interpolation, if needed, was done by simply connecting two measured data points with a straight line (linear interpolation, using MATLAB inbuilt function *interp1*).

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2.3. Investigating dynamic processes in a porous substrate biofilm photobioreactor – A modeling approach. Published in 'Algal Research', 2016(13):30-40.

DOI: 10.1016/j.algal.2015.11.006

Presented here in the original published version, reformatted for the presentation in this thesis (with section numbers removed and corresponding changes).

Investigating dynamic processes in a porous substrate biofilm photobioreactor $-\mathbf{A}$ modeling approach

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Running head: Modeling of non-submerged algal biofilm

Abstract

In the present study, a one dimensional kinetic model was developed for a porous substrate biofilm photobioreator (PSBR), a biofilm bioreactor for the production of microalgae and other biotechnological applications. Light transfer was modeled with a radiative transfer equation (RTE), considering absorption, scattering and biomass pigment adaptation. Dissolved chemical species were modeled using mass balance equations (partial differential equations), with terms describing diffusion, convection, biomass growth, biomass consumption and chemical conversions. pH was modeled as a state variable, and a novel approach of modeling biomass increase in a phototrophic biofilm was introduced. The model was colved using numerical methods and model parameters were acquired either from literature or from experimental work carried out in the framework of this study. The proposed mode was applied to simulated gradients of light, dissolved oxygen concentration, pH etc. in a PSBR biofilm. The simulated results were compared with experimental data acquired in previous studies.

Our results show that the proposed model can accurately predict light intensity distribution in the modeled PSBR biofilm, provided the optical properties of the biomass were measured experimentally. The prediction of the concentration of dissolved oxygen and pH profiles also reflected experimental data with high fidelity. The results also strongly suggest that facilitated CO_2 transfer due to the presence of extracellular carbonic anhydrase in the biofilm matrix plays an important role in DIC transport in the modeled PSBR biofilm at low gas phase CO_2 concentration; and the pH gradient along the depth gradient of the modeled biofilm was mainly caused by the uptake of dissolved inorganic carbon. In addition, the simulation can predict biomass growth in the modeled PSBR with minimal error, thus, with minor modifications, the model can also be applied to predict biomass growth in larger-scale PSBRs.

Keywords

Phototrophic biofilm, Biofilm photobioreactor, dynamic modeling

Introduction

Porous substrate photobioreactors (PSBRs) have been described in previous studies as highly productive systems for the cultivation of various microalgal species for biotechnological applications (e. g. Liu et al., 2013; Murphy et al., 2014; Schultze et al., 2015). One variety of PSBR, the Twin-Layer PSBR (TL-PSBR), has been shown in recent studies to have very high biomass productivity and a pilot-scale study showed its potential in large scale commercial operations (Naumann et al. 2013; Schultze et al. 2015; Shi et al. 2014). Until now, the dynamics behind these promising observations are still not well understood, however, such knowledge should be important for further optimization and up-scaling of the system.

In general, a model of such a biological system can help to improve its understanding, and may be particularly useful when designing, optimizing and controlling bioreactors based on a biological system in a prototype or larger scale (Dunn et al., 2003). Several models have already been developed for submerged phototrophic biofilms or other photobioreactors (e.g.,Berberoglu et al., 2007; Ogbonna et al., 1995; Olivieri et al., 2014; Wolf et al., 2007). However, only one model so far was developed specifically for PSBR biofilms: Murphy and Berberoglu (2014) described a two dimensional kinetic model, giving a first insight into the kinetic interactions inside the biofilm. However, this work focused more on the gradients in the direction of medium flow (i.e. gradient parallel to biofilm surface) rather than the gradients perpendicular to the medium flow (i.e. along the biofilm depth gradient), as pH was considered invariant along the depth gradient. This however, is not the case, as shown recently by microsensor measurements performed on a TL-PSBR biofilm (Li et al., 2015b).

In this study, a comprehensive kinetic model with respect to light and mass transfer, chemical equilibrium, biological processes and biomass production of a PSBR biofilm focusing on the gradients perpendicular to the biofilm surface (along a depth gradient) was developed. Several experiments were carried out in the framework of this study to determine the required parameters. The model was implemented using a numerical method. Simulated growth experiments were carried out in silico using the developed model with several growth condition scenarios, and the simulated results were compared with experimental data measured with microsensors from a previous study.

Materials and Methods

General assumptions and model setup

In the present study, gradients are assumed to exist in the direction perpendicular to the biofilm surface (along the depth gradient) only, thus are one dimensional (1D). Phototrophic growth is assumed to depend solely on irradiance, dissolved oxygen concentration, inorganic carbon availability and the availability of macro-nutrients (N and P). Trace elements are considered to be sufficiently supplied, and, therefore, their concentrations are supposed to have no effect on growth (e.g. Mg, Fe, Zn etc.). Biomass in the biofilm is assumed to be homogeneous: Dry biomass concentration (X_F) is considered as a constant; in consequence, porosity of the biofilm (θ) is also a constant (as was observed experimentally, see supplementary materials). The intracellular composition of the biomass is assumed to be constant over time and the depth gradient (i.e. biomass has a fixed C:N:P ratio, as it was observed experimentally, see supplementary materials).



Figure 1: One-dimensional (1D) compartmentalization of a non-submerged algal biofilm grown in a porous substrate bioreactor as implemented in the proposed model. Compartments and boundaries are marked at the top of the figure. Direction of the depth gradient, Z, are given at the bottom. Growth related processes boldy occur in the biofilm compartment. Arrow directions indicate the directions of the processes. L_f and L_{mathware} are thicknesses of the biofilm and microporous substrate membrane, respectively. C1 to C4 represent: bulk gas compartment, biofilm compartment, boundary layer (BL) compartment and bulk liquid compartment, respectively. B1 and B2 represent the surface and bottom boundaries of the biofilm compartment.

Table I: "Nomenclature of symbols presented in Fig. 2.

Model parameters			State variables		
Symbol	Definition	Symbol	Definition		
$D_{e,x}$	Effective diffusion coefficient of dissolved species x.	S_x	Concentration of dissolved species x in biofilm compartment.		
u _z	Convective flow rate in biofilm due to evaporation.	Ι	Irradiance in the biofilm compartment.		
p_x	Partial pressure of gas x in bulk gas.	μ	Growth rate of the biomass.		
Iin	Incident irradiance.	L_F	Biofilm thickness.		
S _{x,surface}	Concentration of dissolved species x at biofilm surface.	pH ^b	pH in biofilm.		
$S_{x,medium}$	Concentration of dissolved species x in bulk liquid.				
pH _{medium}	pH in bulk liquid.				

^aThe subscript x representing species x are given in the subscripts in Fig. 2. ^bpH itself is not a state variable, and is calculated from state variable S_{H} :

Since the proposed model focuses on modeling the kinetics of processes inside the biofilm rather than fluid dynamics at the interfaces, in the proposed model, the biofilm was simplified into 4 compartments without any flow boundary layer (Fig. 1): Bulk gas compartment (the ambient gas phase); biofilm compartment (the biomass, and the only compartment containing state variables); mass transfer boundary layer (BL) compartment (the micro-porous membrane) and bulk liquid compartment (the liquid phase in a macro porous material). Considering the fast mass transfer in bulk gas and bulk liquid compartments (displaying high convective flow rates), both compartments are assumed to be well-mixed (homogeneous). The micro-porous membrane is considered as a mass transfer boundary layer for dissolved species and is assumed to have a flow only in the direction parallel to the depth gradient. Growth related processes (e.g. consumption of nutrients, production of O2 etc.) take place only in the biofilm compartment. Evaporation at the biofilm surface creates a convective flow in both the biofilm compartment and the boundary layer compartment. However, the gas flow boundary layer at the biofilm surface is neglected (Bergman et al., 2011; Schlichting and Gersten, 2000). Concentration gradients exist in both the biofilm compartment and the boundary layer compartment, and lead to diffusive fluxes in these compartments. Exchange of all dissolved species between biomass and bulk medium occurs through the BL compartment. In addition, direct exchange of dissolved gaseous specious (dissolution and/or escape of O2 and CO2) occurs at the biofilm surface (interface between bulk

gas and biofilm). Illumination at the biofilm surface is assumed to be emitted from a diffuse source (equal intensity in all directions). State variables only exist in the biofilm compartment, which has a surface/top boundary (interface between biomass and gas phase) and a bottom boundary (interface between biomass and the porous membrane). A schematic representation of the parameters, state variables and their interactions are given in Fig. 2 and Table I (Nomenclature of the symbols in Fig. 2 can be found in Tab. I). Processes included in the proposed model are:

 Transport processes: Light transfer (absorption and scattering), mass transport process (diffusion of dissolved species and convection due to evaporation).

- Biological processes: Photosynthesis (growth), respiration (both dark and photo-respiration), biomass pigment
 adaptation due to exposure to different light intensities and increase in biofilm thickness due to growth.
- Chemical processes: Acid-base equilibrium (speciation of inorganic carbon and inorganic phosphate), and dissolution/release of gases (O₂ and CO₂) at biofilm surface.



Figure 2: Schematic representation of the variables and parameters (text boxes) and their interactions (processes, indicated by arrows) considered in the p roposed model. Grey box includes variables, parameters and interactions inside the biofilm compartment. The dash line box includes variables affected by the mass transfer processes: D_{ex} presents diffusion, and u, represents advection caused by surface evaporation. Solid arrows represent growth-related interactions. Thick grey arrows show interactions that cross the boundaries (interaction between compartments). Notice interactions can be one-way or bi-directional (indicated by the arrow/arrows). Nomenclature of the symbols can be found in Tab. 1. *Subscript totalP denote total inorzanic phosphate.

Light transfer and phototrophic growth

The proposed model uses the radiative transfer equation (RTE) to model light transfer in the biofilm, as previous observations showed that the effect of scattering in densely packed suspension (here, a biofilm) cannot be neglected (Berberoglu et al., 2007). The 1D form of RTE is (Murphy and Berberoglu, 2014):

$$\frac{\partial I_{\lambda}(z,\hat{s}_{a})}{\partial z} = -\left(\kappa_{\lambda,F,z} + \sigma_{\lambda,F}, z\right) \cdot I_{\lambda}(z,\hat{s}_{a}) + \frac{\sigma_{\lambda,F,z}}{4\pi} \int_{0}^{4\pi} I_{\lambda}(z,\hat{s}_{b}) \cdot \Phi(\hat{s}_{b},\hat{s}_{a}) \, d\Omega_{b} \tag{Eq.1},$$

here, z represents the depth, $I_{\lambda}(z, \hat{s}_a)$ and $I_{\lambda}(z, \hat{s}_b)$ are the spectral radiant intensities of wavelength λ at depth z, travelling in the direction \hat{s}_a and \hat{s}_b , respectively. $\kappa_{\lambda,F,z}$ and $\sigma_{\lambda,F,z}$ are the effective spectral absorption coefficient and the effective spectral scattering coefficient of the biofilm at depth z. $\Phi(\hat{s}_b, \hat{s}_a)$ is the scattering phase function from direction \hat{s}_i to \hat{s}_i , and is estimated from Henry-Greenstein approximation as described by (Berberoglu et al., 2007); the scattering of algal cells was considered to be strongly forward, and an asymmetry factor of 0.98 was taken for calculation (Berberoglu et al., 2007). Ω_b is the solid angel surrounding direction \hat{s}_b . To simulate the effect of adaptation of the biomass by adjusting their cellular pigment content (verified experimentally, see supplementary materials), a simple linear function dependent on the position of the cell in the biofilm for all photosynthetically active radiation (PAR) wavelengths:

$$\kappa_{\lambda,F,z} = \frac{z}{L_{F,light}} \cdot \kappa_{\lambda,F} \qquad (z \le L_{F,light})$$
(Eq.2),

 $L_{F,light}$ is the thickness of the illuminated part of the biofilm. $\kappa_{\lambda,F}$, which is the maximal absorption coefficient of the biomass, was determined experimentally together with the maximal scattering coefficient of the biomass $\sigma_{\lambda,F}$ (see supplementary materials; $\sigma_{\lambda,F}$, was considered to remain constant throughout the biofilm, i.e. $\sigma_{\lambda,F,z} = \sigma_{\lambda,F}$).



Figure 3: Schematic representation of the radiation transfer equation (RTE) boundary conditions considered in the proposed model (as described by Eq. 3 and 4). Thick solid line arrows: Total irradiance at the biofilm surface and the biofilm-membrane interface. Thick dotted line arrows: Reflected irradiance at the biofilm surface and the hiofilm-membrane Thin interface arrows: Directional irradiation inside the biofilm due to direct incident irradiance and/or reflected irradiance by the membrane surface

At the biofilm surface (top boundary), light can be reflected before entering the biofilm. Assuming a diffuse light source, the boundary condition at the top boundary (z = 0) according to (Murphy and Berberoglu, 2014), and considering the vertical placement of the modeled biofilm (perpendicular to the ground as described by Shi et al., 2007) is:

$$I_{\lambda}(z=0,\omega_n) = (1-r_{bs}) \cdot \frac{I_{in\lambda}}{\pi} \qquad \left(-\frac{\pi}{2} < \omega_n < \frac{\pi}{2}\right)$$
(Eq.3).

Similarly, light can be reflected by the microporous membrane before leaving the biofilm at the biofilm–membrane interface (bottom boundary). Thus, the boundary condition at the bottom boundary ($z = L_F$) following Murphy and Berberoglu (2014) is written as:

$$I_{\lambda}(z = L_F, \omega_n) = r_{ms} \cdot \frac{I_{L_F\lambda}}{\pi} \qquad \left(-\pi < \omega_n < -\frac{\pi}{2} \text{ and } \frac{\pi}{2} < \omega_n < \pi\right) \qquad (\text{Eq.4}).$$

In Eq.3 and 4, ω_n is the angle relative to the normal of the biofilm surface; L_f is biofilm thickness; r_{bs} and r_{ms} are the reflectance of the biofilm and membrane surface, respectively (Fig. 3).

The proposed model does not differentiate the efficiency of PARs with different wavelengths in photosynthesis. In consequence, at any depth z, the calculated intensities in all directions and all bands in PAR range are integrated into total PAR (I_z):

$$I_z = \int_{400nm}^{700nm} \int_0^{4\pi} I_\lambda(z, \hat{s}_a) \, d\Omega_a \, d\lambda \tag{Eq.5}.$$

Irradiance for one discrete element was calculated as the average value of its left and right nodes, and this was fed into an equation (P-I curve) describing the kinetics of phototrophic growth as described by (Platt et al. 1980):

$$f_I = \left(1 - e^{\frac{-\alpha i_Z}{ETR_{max}}}\right) \cdot e^{\frac{-\beta i_Z}{ETR_{max}}}$$
(Eq.6),

whereas f_I represents the effect of irradiance on photosynthetic growth. ETR_{max} is the maximal electron transfer rate (ETR), α is the initial slope of the ETR-irradiance curve and β is a term representing the degree of photo-inhibition (In the present study, ETR_{max} , α and β were determined experimentally, see supplementary materials).

The effect of oxygen on phototrophic growth was expressed as an inhibition term dependent on the concentration of dissolved oxygen ([DO], denoted as $S_{O:}$), similar to the approach used by Murphy and Berberoglu (2014) for inorganic carbon and phosphate but without the saturation term (i.e. only inhibition is considered, as observed experimentally by Pope, 1975 and by Grötzschel and de Beer, 2002):

$$f_{O_2} = \frac{S_{O_2}}{S_{O_2} + S_{O_2}^{-2}/K_{O_2,In}}$$
(Eq. 7).

 f_{O_2} represents the effect of [DO] on phototrophic growth, and $K_{o_2,In}$ is the inhibition constant for [DO] on photosynthesis. In the present study, this inhibition was considered as an effect of photorespiration. $K_{O_2,In}$ is a dependent variable controlled by the CO₂ to O₂ ratio, R_{CO_2/O_2} , and follow a Monod type kinetic, as suggested by the data presented by Ku and Edwards (1978):

$$K_{O_2,In} = K_{O_2,In,max} \frac{R_{CO_2/O_2}}{R_{CO_2/O_2} + K_{R_{CO_2/O_2,S}}}$$
(Eq.8).

 $K_{O_2,In,max}$ is the assumed maximal inhibition (i.e. photorespiration) coefficient of [DO] on phototrophic growth, and $K_{R_{CO_2/O_2,S}}$ is the assumed half saturation constant of the CO₂ to O₂ ratio on photorespiration. The effects of dissolved macro-nutrients (N as nitrate; S_{NOi} ; P as total phosphate: sum of monohydrogenphosphate, S_{HPOi} and dihydrogenphosphate, S_{HPOi} and phosphate ion S_{POi}) on growth were modeled using Monod type kinetics (IWA Task Group on Biofilm Modelling 2006), the effect of inhibition by high CO2 concentration and high nutrients concentration are not considered in the present study, as suggested by the data presented by Schultze et al. (2015) and in the supplementary material (Fig. S9):

$$f_x = \frac{S_x}{K_{x,S} + S_x} \tag{Eq. 9}.$$

 f_x represents the effect of S_{NOI} (f_{NOI}) or $S_{POI,Iotal}$ (f_{POI}) on phototrophic growth, and their saturation constants are represented by $K_{x,S}$. However, only monohydrogenphosphate and dihydrogenphosphate ions were modeled as diffusive species in the present study, as significant concentrations of phosphate ions were not expected in such biofilms as suggested by previous investigations (Li et al., 2015b).

In the present study, carbon concentrating mechanisms (CCMs) were taken into account by assuming that both CO_2 and HCO_3 (direct uptake of bicarbonate and conversion of bicarbonate to CO_2 at the cell surface before uptake were not distinguished in the present study, and are considered to have the same effect) are able to support phototrophic growth (Raven et al. 2008), and an approach similar to that described by Wolf et al. (2007) is used: Monod type kinetics are applied to describe phototrophic growth on CO_2 and HCO_3 expressed as

$$f_{CO_2} = \frac{S_{CO_2}}{K_{CO_2,S} + S_{CO_2}}$$
(Eq.10),

and

$$f_{HCO_3} = \frac{S_{HCO_3}}{\kappa_{HCO_3,s} + S_{HCO_3}} \cdot \frac{S_{CO_2}}{S_{CO_2/KCO_2,in}^2 + S_{CO_2}}$$
(Eq.11),

respectively. S_{CO_2} and S_{HCO_3} represent concentrations of dissolved CO₂ and HCO₃⁻. $K_{CO_2,S}$ and $K_{HCO_3,S}$ are the saturation constants for CO₂ and HCO₃⁻ growth, respectively. $K_{CO_2,in}$ is the assumed inhibition constant of CO₂ on growth using bicarbonate as carbon source.

Finally, the phototrophic growth rate was calculated using a similar method as described by Wolf et al. (2007), however, irradiance and [DO] are considered to have an influence on the growth rate (μ) independently of the nutrient concentrations:

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$$= \mu_{max} \cdot f_I \cdot f_{O_2} \cdot \left[\min(f_{NO_3}, f_{PO_4}, f_{CO_2}) + \min(f_{NO_3}, f_{PO_4}, f_{HCO_3}) \right]$$
(Eq.12).

The maximal growth rate of the biomass is represented by μ_{max} (estimated experimentally, see supplementary materials). Another effect of the CCM is the production of extracellular carbonic anhydrase (CA) at low dissolved free CO₂ concentrations (Raven et al., 2008). This results in an increased interconversion rate of CO₂ – bicarbonate (increased to identical value as dehydration of carbonic acid). Another effect of the extracellular CA is a facilitated CO₂ transport in the biofilm, which leads to significantly accelerated CO₂ transport (Bao and Trachtenberg, 2006; Gros et al., 1976; Trachtenberg et al., 1999). In the present study, this effect was modeled by multiplying the effective diffusion coefficient of CO₂ with a simple exponential function dependent on local dissolved free CO₂ concentration, as suggested by the results from Gros et al. (1976):

$$D_{e,CO_2}^* = D_{e,CO_2} \cdot (f_a \cdot e^{f_b \cdot S_{CO_2}} + 1)$$
 (Eq. 13).

 f_a and f_b are two pre-determined factors, and D_{e,Co_2} is the calculated effective diffusion coefficient of CO₂ in the biofilm (see section below). Eq. 13 describes a function influencing the effective diffusion coefficient of CO₂ in biofilm, at low CO₂ concentrations the effective diffusion coefficient of CO₂ can be increased significantly (several fold), whereas at high CO₂ concentrations the effective diffusion coefficient of CO₂ is hardly affected, as suggested by the data presented in previous studies (Bao and Trachtenberg, 2006; Gros et al., 1976; Trachtenberg et al., 1999).

Dark respiration rate was considered to be [DO] dependent and follows the Monod kinetic (Wolf et al., 2007):

$$R_d = R_{d,max} \cdot \frac{S_{O_2}}{S_{O_2} + K_{O_2,Sa}}$$
(Eq. 14),

in which $K_{O_2,Sa}$ is the saturation constant, and $R_{d,max}$ is the maximal dark respiration rate (determined experimentally, see supplementary materials). The biofilm thickness (L_F) is a pseudo-state variable in the proposed mode, and the rate of biofilm thickness change is calculated as (Wolf et al., 2007; the approach for modeling change in biofilm thickness is described in detail below):

$$u_{L_F} = \frac{dL_F}{dt} = \int_0^{L_F} (\mu - R_d) \, dz \tag{Eq.15}.$$

It is possible, that besides aerobic respiration, the algal cells respire anaerobically (Atteia et al., 2013), however, due to a lack of knowledge on this topic, this process was not included in the present study.

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The consumption/production of dissolved species by biological processes are calculated based on biomass growth, as described in detail by (IWA Task Group on Biofilm Modelling 2006), and generally:

$$\gamma_{x,F} = (\mu - R_d) \cdot X_F \cdot \frac{1}{Y_x}$$
(Eq.16).

 $\gamma_{x,F}$ is the consumption/production rate of the dissolved species x due to growth, X_F is the biomass concentration of the biofilm (dry weight per volume). Y_x is the yield of species x with respect to biomass production or consumption. For inorganic carbon, nitrogen and phosphorus, the yields are calculated as the reciprocal of their contents in biomass (IWA Task Group on Biofilm Modelling, 2006). In the present study, it is assumed that photosynthesis utilizing CO₂ and bicarbonate have identical yields on oxygen (O₂:C = 1.3:1; Atkinson and Smith, 1983; Raven et al., 2014) and biomass.

Mass balance and chemical equilibrium

Mass transfer in the proposed model is described by the 1D mass balance equation (Bergman et al., 2011):

$$\frac{dS_x}{dt} = D_{e,x}\frac{d^2S_x}{dz^2} + u_z\frac{dS_x}{dz} + \gamma_{x,F} + \gamma_{x,Che}$$
(Eq.17),

 S_x is the concentration of dissolve species x, $D_{e,x}$ is the effective diffusion coefficient of species x in the biofilm and u_z is the convective flow rate in the biofilm caused by evaporation at the biofilm surface. $\gamma_{x,F}$ and $\gamma_{x,Che}$ are production/consumption due to biomass growth/respiration and chemical conversion, respectively. Chemical conversions in the present study are controlled by the rate constant (k) and the equilibrium constant (K) of the reaction, e.g., for reaction A \leftrightarrow B+C, the conversion rate of A is expressed as:

$$\gamma_{A,Che} = k_{A/BC} \cdot \left(S_A - \frac{S_B \cdot S_C}{\kappa_{A/BC}} \right)$$
(Eq.18)

Values of rate and equilibrium constants can be found in Tab. III.

The boundary conditions required to solve Eq.17 are given below (IWA Task Group on Biofilm Modelling, 2006):

1) At the bottom boundary ($z = L_F$, biofilm – membrane interface), for all dissolved species, the flux of the species *x* across the BL compartment equals the flux at the bottom of the biofilm compartment ($j_{m,x}$):

$$i_{m,x} = D_{membrane,x} \cdot \frac{s_{x,medium} - s_{x,L_F}}{L_m} + u_z \cdot s_{x,medium}$$
(Eq.19).

 $D_{membrane,x}$ is the effective diffusion coefficient of x in the porous membrane, $S_{x,medium}$ is the concentration of species x in the liquid phase (bulk liquid compartment), and L_m is the thickness of the membrane (BL compartment).

 At the surface boundary (z = 0, biofilm surface), for gaseous species, the concentrations of O₂ and CO₂ are assumed to be constant, and are governed by Henry's law. For oxygen:

$$S_{O_2,z=0} = p_{O_2} \cdot K_{Hen,O_2}$$
 (Eq.20),

and for dissolved inorganic carbons (DICs):

$$S_{DIC,z=0} = p_{CO_2} \cdot K_{Hen,CO_2} \cdot \left(1 + \frac{K_{a_1,H_2CO_3}}{S_{H,z=0}} + \frac{K_{a_1,H_2CO_3} \cdot K_{a_2,H_2CO_3}}{S_{H,z=0}^2}\right)$$
(Eq.21),

in which $S_{0_2,z=0}$ and $S_{DIC,z=0}$ ($S_{x,surface}$ in Fig. 2) are the concentration of dissolved O₂ and DIC at the biofilm surface, p_x is the partial pressure of x in the gas phase, and $K_{Hen,x}$ is the Henry's constant for species x (For CO₂, chemical equilibria of the dissolved inorganic carbons are also taken into consideration. As described by Murphy and Berberoglu (2014), K_{a1,H_2CO_3} , K_{a2,H_2CO_3} and $S_{H,z=0}$ are the dissociation constants for H₂CO₃ and the proton concentration at biofilm surface, respectively.). For non-gaseous dissolved species however, the surface boundary is considered as a no flux boundary ($J_x = 0$). This simulates the increase in concentration of a dissolved species as a result of evaporation at the biofilm surface.

For initial conditions (at t = 0), an initial biofilm thickness is given as $L_{F,t=0}$; the concentrations of non-gaseous dissolved species were set to identical values as found in freshly prepared medium; for dissolved gaseous species, the concentrations throughout the biofilm at t = 0 were assumed to be identical to their concentration at the biofilm

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surface. In the present study, effective diffusion coefficients were determined as described by Li et al. (2015a; see also supplementary materials).

The pH in the biofilm compartment is modeled using the charge balance equation by (Wolf et al., 2007), considering the concentration of protons (S_{H}) only depending on the charge balance (assumed to be a non-diffusive species). The speciation of inorganic carbon and inorganic phosphate are assumed to be pH dependent, only. The charge balance equation applied in the present study is:

 $S_{H^+} - S_{OH^-} = S_{HCO_3^-} + 2 \cdot S_{CO_3^{2-}} + S_{H_2PO_4^-} + 2 \cdot S_{HPO_4^{2-}} + 3 \cdot S_{PO_4^{3-}} - S_{cations}$ (Eq.22). S represents concentration of the different species, and K_w is the disassociation constant of water. In the present study, Na⁺ is considered to be the sole cation present. Define $Ch(pH) = S_H - S_{OH}$ (a pH dependent parameter), according to (Campos and Flotats, 2003), S_H can be calculated as:

$$S_{H} = \begin{cases} \frac{ch(\mathbf{pH}) + \sqrt{ch(\mathbf{pH})^{2} + 4 \cdot K_{w}}}{2}, \ ch(\mathbf{pH}) \ge 0\\ \frac{2 \cdot K_{w}}{-ch(\mathbf{pH}) + \sqrt{ch(\mathbf{pH})^{2} + 4 \cdot K_{w}}}, \ ch(\mathbf{pH}) < 0 \end{cases}$$
(Eq.23).

However, in the proposed model, instead of calculating S_H at the end of each numerical integration step, S_H is considered to be a state variable, and, therefore is calculated using the algorithms described by (Campos and Flotats, 2003):

$$\frac{dS_H}{dt} = \frac{dCh(\text{pH})}{dt} \cdot \left\{ \begin{array}{l} \frac{1}{2} \cdot \left(\frac{Ch(\text{pH})}{\sqrt{Ch(\text{pH})^2 + 4 \cdot K_w}} + 1 \right), \ ch(\text{pH}) \ge 0\\ - \frac{S_H}{2 \cdot K_w} \left(\frac{Ch(\text{pH})}{\sqrt{Ch(\text{pH})^2 + 4 \cdot K_w}} - 1 \right), \ ch(\text{pH}) < 0 \end{array} \right.$$
(Eq.24).

Model implementation and parameters

In the proposed model, the biofilm compartment is divided into discrete elements using a simple upwind scheme (Courant et al., 1952) into *n* elements (for the i^{th} element: I = 1, 2, ..., n), with the 1st element being the biofilm surface (top boundary) and the n^{th} element representing the biofilm–membrane interface (bottom boundary). In the present study, the model was coded using MATLAB (version 2013a, MathWorks, Ismaning, Germany; the code can be acquired from the first author upon request).

To implement the RTE into the proposed model, the discrete ordinate method (DOM) was applied to divide the solid angle (4π) into two hemispheres, each with 24 distinct directions (Berberoglu et al., 2007). This generated for each discrete element a system of ordinary differential equations (ODEs), which were then solved using the method provided by Colomer et al. (2000) with the MATLAB *linsolve* function. A grid sensitivity study was carried out and 1 discrete elements per 5 μ m simulated biofilm was determined to be numerically sufficient (< 1% change with a further increase in number of discrete elements; Berberoglu et al., 2007).

Partial differential equations (PDEs) describing the mass balances were transformed into systems of ODEs using the method of lines (MOL; Schiesser, 1991). The acquired ODE systems were solved using a variable order numerical differentiation formula (*ode15s*, MATLAB). The biofilm thickness (L_F) is a pseudo-state variable in the proposed model: All non-surface (non-top boundary) discrete elements are set to contain the same amount of biomass, and, hence, have the same thickness ($L_{F,i\neq I} = dL_F$) during the simulated growth period, and were kept constant. Increase in biofilm thickness is modeled as the addition of a new layer of biomass onto the surface of the existing biomass (into the surface discrete element, $L_{F,i=1}$, which has an initial thickness of 0). Addition of the new surface element is triggered when the increase in thickness reached dL_F (detected by the *event* function of the *ode15s* solver) (Fig. 4). Directly after addition of this new discrete element, the variables in new discrete element are assumed to have the same value as the element below ($S_{x,i=1} = S_{x,i=2}$) and the system was solved again using this as the new initial condition. A grid sensitivity study showed one discrete element per 10 µm of the simulated biofilm thickness was numerically sufficient (< 1% change with a further increase in number of discrete elements).



Figure 4: Schematic representation of the implementation of biofilm thickness (L_r) as a pseudo-state variable. dL_r is the pre-defined biofilm thickness for the non-surface discrete element (i = 2, ..., n). L_{Fr-1} is the biofilm thickness of the surface discrete element (i = 1). The sum of biofilm thickness increase from all discrete elements $(2_{T}^{T}L_{F_{T}})$ is added to $L_{F_{T-1}}$. Once $L_{F_{T-1}}$ reached dL_r , a new discrete element was added on top of the former $L_{F_{T-1}}$

In the present study, the parameters used in the model were adjusted to a *Halochlorella rubescens* (CCAC 0126; Culture Collection of Algae at the University of Cologne; www.ccac.uni-koeln.de) TL-PSBR biofilm cultivated with Bold's basal medium (BBM; Bischoff and Bold 1963), which served as a model system in previous experimental PSBR biofilm studies (Schultze et al., 2015; Li et al, 2015; Li et al 2015).

In the present study, in order to achieve a realistic representation of the investigated system, experiments were carried out in the framework of this study to acquire the parameters required in the model equations (coefficients and constants found in Eq. 1 to Eq. 24). However, it was not possible to acquire all the parameters experimentally, as a result, some of the parameters were taken from previously published studies, and others had to be assumed due to lack of knowledge (e.g. the biomass was assumed to be able to utilize bicarbonate directly as a carbon source). The assumed parameters were acquired by running the model with different sets of unknown parameters (within a theoretically reasonable range) and selecting the parameter set that produced results that fitted the experimental observations. Model parameters used in the present study, their value and their methods of determination or references can be found in Tab. II and III (for detailed experimental method see supplementary materials, as indicated in the text above).

Reaction	Equilibrium constant, K	Rate constant, k, in d-1	Source
$H_2O \leftrightarrow H^+ + OH^-$	10-14	Instantaneous	Morales-Rodriguez et al., 2011
$CO_2 + H_2O \leftrightarrow HCO_3 + H^+$	10.6.35	107.54	Gibbson and Edsall, 1963
$CO_2 + OH^- \leftrightarrow HCO_3^-$	10-6.35	10 ^{8.86}	Wolf et al., 2000
$HCO_3^- \leftrightarrow CO3^{2-} + H^+$	10-10.33	104.6	Morales-Rodriguez et al., 2011
$H_3PO_3 \leftrightarrow H_2PO_3^{2*} + H^+$	10 ^{-2.15}	Instantaneous	Morales-Rodriguez et al., 2011
$H_2PO_3^- \leftrightarrow HPO_3^{2-} + H^+$	10 ^{-7.2}	10 ^{9.74}	Morales-Rodriguez et al., 2011
$\text{HPO}_3^{2*} \leftrightarrow \text{PO3}^{3*} + \text{H}^+$	10-12.1	Instantaneous	Morales-Rodriguez et al., 2011

Table III: "Chemical reactions integrated into the proposed model and their rate and equilibrium constants.

^aAt 25°C.

Table II	: "Model	parameters	applied	in the	present	study
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Symbol	Definition	Value/Unit		Source	
ETR_{max}	Maximal electron transfer rate	49.6	mole m ⁻² s ⁻¹	b	
$D_{cations}$	Considered here as diffusion coefficient of Na ⁺ in water	$1.15\times10^{\text{-}4}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011	
D_{CO_2}	Diffusion coefficient of CO2 in water	$1.65\times10^{\text{-}4}$	$m^2 \; d^{\text{-}1}$	Morales-Rodriguez et al. 2011	
D_{CO_3}	Diffusion coefficient of CO32- in water	$1.02\times10^{\text{-}4}$	$m^2 d^{*1}$	Morales-Rodriguez et al. 2011	
D_{HCO_3}	Diffusion coefficient of HCO3 in water	$7.9\times10^{\cdot5}$	$m^2 d^{\cdot 1}$	Morales-Rodriguez et al. 2011	
D_{NO_3}	Diffusion coefficient of NO3 in water	$1.47\times10^{\text{-}4}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011	
$D_{O_{2}}$	Diffusion coefficient of O2 in water	$1.73\times 10^{\text{-4}}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011	

Table II (continued)						
Symbols	Definition	Value/Unit		Source		
$D_{H_2PO_4}$	Diffusion coefficient of H ₂ PO ₄ ⁻ in water	$6.2\times10^{\text{-5}}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011		
f_a	Factor determining facilitated CO2 transfer	100	/	с		
f_b	Factor determining facilitated CO2 transfer	10-4	/	с		
K_{Hen,CO_2}	Henry's constant of CO2 dissolution in water	$3.04\times 10^{\text{-7}}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011		
K_{Hen,O_2}	Henry's constant of oxygen dissolution in water	$1.3\times 10^{\text{-8}}$	$m^2 d^{-1}$	Morales-Rodriguez et al. 2011		
$K_{R_{CO_2/O_2,S}}$	Saturation coefficient representing effect of CO_2 to O_2 ration on O_2 inhibition	0.35	/	c		
$K_{O_2,In,max}$	Maximal inhibition coefficient of oxygen on net photosynthetic productivity	1×10^{-3}	М	c		
$K_{CO_2,S}$	Half saturation coefficient of CO_2 on photosynthetic productivity	$4\times 10^{\text{-4}}$	М	'Goldman et al. 1974		
$K_{CO_2,in}$	Inhibition coefficient of CO2 on photosynthetic productivity using bicarbonate	$5\times 10^{\text{-4}}$	М	с		
$K_{HCO_3,S}$	Half saturation coefficient of HCO ₃ ⁻ on photosynthetic productivity	$4\times 10^{\text{-5}}$	М	'Goldman et al. 1974		
$K_{NO_3,S}$	Half saturation coefficient of NO ₃ on photosynthetic productivity	$1\times 10^{\text{-4}}$	М	Wolf et al. 2000		
$K_{O_2,S}$	Half saturation coefficient of O2 on dark respiration	$1\times 10^{\text{-4}}$	М	Wolf et al. 2000		
$K_{PO_4,total,S}$	Half saturation coefficient of total phosphate on photosynthetic productivity	$1.7\times10^{\text{-}5}$	М	Wolf et al. 2000		
L_m	Substrate membrane thickness	$5 imes 10^{-5}$	m	c		
p_{o_2}	O2 partial pressure at the biofilm surface	21000	Ра	/		
$p_{CO_2,medium}$	Dissolved CO2 partial pressure in medium	41	Ра	/		
$p_{co_2,surface}$	Dissolved CO2 partial pressure at biofilm surface	41	Ра	/		
pH_{medium}	pH of the medium	6.8	/	Bischoff and Bold, 1963		
$R_{d,max}$	Maximal biomass dark respiration rate	0.133	d^{-1}	b		
r_{bs}	Surface reflectance of the biofilm	0.1	/	c		
r_{ms}	Surface reflectance of the membrane	0.5	/	с		
$S_{cations,medium}$	Sum of cations concentrations in medium, here considered as Na ⁺ (calculated as [NO ₃ ⁻] + 2·[HPO ₄ ²⁻] + [H ₂ PO ₄ ⁻] in fresh medium)	$5.1\times10^{\text{-4}}$	М	'Bischoff and Bold, 1963		
$S_{CO_2,medium}$	Dissolved CO2 concentration in medium	/	М	e		
$S_{CO_3,medium}$	Dissolved bicarbonate ion concentration in medium	/	М	e		
$S_{HCO_3,medium}$	Dissolved carbonate ion concentration in medium	/	М	e		
$S_{O_2,medium}$	Dissolved oxygen concentration in medium	0	М	с		
$S_{NO_3,medium}$	NO3 ⁻ concentration in medium	$2.94\times 10^{\cdot 3}$	М	Bischoff and Bold, 1963		
$S_{HPO_4,medium}$	HPO ₄ ³⁻ concentration in medium	$4.31\times 10^{\text{-4}}$	М	Bischoff and Bold, 1963		
$S_{H_2PO_4,medium}$	H ₂ PO ₄ ³ concentration in medium	$1.29\times 10^{\cdot 3}$	М	Bischoff and Bold, 1963		
uz	Convection rate due to evaporation on the	0.23	m d ⁻¹	Li et al., 2015		

"At 23°C. bDetermined experimentally, see supplementary materials. cAssumed. dProduct data sheet. °Calculated from chemical equilirium.

Results and Discussion

Light transfer in TL-PSBR

Simulated irradiance distributions in a 1,000 μ m thick TL-PSBR biofilm subjected to 300 μ mol photon m⁻² s⁻¹ are presented in Fig. 5: Comparison between irradiance profile modeled, considering neither forward scattering nor pigment adaptation, with forward scattering but no pigment adaptation and in-scattering plus pigment adaptation are given in Fig. 5A. Fig. 5B compares the modeled results and experimental data as measured by Li et al. (2015b). The results show that both forward scattering and pigment adaptation of the biomass have significant effects on the distribution of PAR in a PSBR biofilm. By adding the forward scattering term and/or the term describing the pigment adaptation of the biomass, the PAR penetrates deeper into the biofilm (Fig. 5A). Comparison between the simulated results with both forward scattering and biomass pigment adaptation taken into consideration showed that the proposed model can very accurately describe the PAR distribution in H. rubescens TL-PSBR (Fig. 5B). Based on the model developed by Murphy and Berberoglu (2014), it is concluded that the intensity distributions calculated from optical thickness with a fixed mass extinction coefficient are a sufficient modeling approximation. Our results, however, show that both forward scattering and adaption of the biomass (changing the absorption coefficient along the depth gradient, i.e. lowering biomass absorption coefficient in the part of the biofilm near the surface) increased the penetration depth of the PAR (as shown by the high fit of modeled data to the experimental data). Thus, light penetration depth and light dilution rate (i.e. available photons per cell) would be significantly underestimated if the in-scattering and/or biomass pigment adaptation were not taken into consideration in such biofilm models, as could be suspected from previous studies as reviewed by (Richardson et al., 1983).





Delivery of dissolved inorganic carbon

Figure 6 compares the simulated results (with and without taking facilitated CO_2 transport due to extracellular CA into account) of biofilms exposed to 300 µmol photon m⁻² s⁻¹ surface irradiance and different additional gas phase CO_2 concentrations (0% and 5% v/v). Compared to the modeled result without considering facilitated CO_2 transfer in the model, when no additional CO_2 was supplied in the gas phase, the simulated results with a term describing facilitated CO_2 transfer reflected the experimental data presented by Li et al. (2015b) more faithfully. In contrast to these findings, at high CO_2 concentration (5% supplementary CO_2 in the gas phase), no significant effect of the implementation of facilitated CO_2 transfer could be observed (Fig. 6B).

The phenomenon of facilitated CO_2 transport in solutions with CA is well documented, and known to accelerate intracellular CO_2 transfer and CO_2 transfer in liquid membranes up to 200-fold (Bao and Trachtenberg, 2006; Trachtenberg et al., 1999). Our results strongly suggest that this is also the case in *H. rubescens* biofilms: Without taking into account facilitated CO_2 transfer, the simulated results predict a much lower dissolved oxygen

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concentration ([DO]) compared to the experimental data, which is also valid for pH values. Thus, the results suggest that facilitated CO_2 transport plays an important role in supplying DIC into the biofilm at low gas phase CO_2 concentrations. In the present study, the facilitated CO_2 transport was modeled using an exponential function dependent on the local CO_2 concentration, However, previous studies suggested that this phenomenon also depends on the concentration of CA, the concentration of other dissolved species and/or their structures as well (Bao and Trachtenberg, 2006; Gros et al., 1976; Trachtenberg et al., 1999). In contrast, the approach used here appears to be crude. Nevertheless, under the assumption of all other factors being constant, a CO_2 concentration-dependent relationship as found here has been previously observed by other researchers (Gros et al., 1976). It can be expected, that this facilitated CO_2 transport exists also in submerged phototrophic biofilms. However, due to a significantly thicker boundary layer at the biofilm surface and the liquid phase instead of a gas phase covering the biofilm, the effect of the facilitated CO_2 transport would be much less pronounced in submerged biofilms due to limited transport at the biofilm surface compared to a PSBR biofilm.



Figure 6: Dissolved oxygen concentration ([DO]), pH and dissolved inorganic carbon (DIC) profiles in Halochlorella rubescens biofilms exposed to 300 µmol photon m⁻² s⁻ Panel A and B present data simulated/measured with 0% and 5% additional CO2 in the gas phase. In each panel: [DO] profiles are presented in top-left subplots, whereas bottom-left subplots give pH profiles (in both left subplots, solid lines: simulated distribution with facilitated CO2 transfer; dashed lines: simulated distribution without facilitated CO2 transfer; empty squares: experimental data measured by Li et al., 2015b). The two subplots on the right present simulated DIC distributions with profiles considering facilitated transfer on top (solid lines: total [DIC]: light grav area: bicarbonate; dark gray area: dissolved free CO2). Notice at 5% CO2, no significant difference can be observed between simulated distributions with and without considering facilitated CO2 transfer.

Nutrient supply in TL biofilms subjected to high irradiance

Gradients in simulated *H. rubescens* PSBR biofilms exposed to 1,000 μ mol photon m⁻² s⁻¹ and different CO₂ concentrations in the gas phase are presented in Fig. 7. The simulated [DO] and pH profiles are compared with the experimental data measured by Li et al. (2015b): The simulated profiles generally reflect the experimental data with a good fit. However, at 5% additional CO₂ in the gas phase, the simulated [DO] are about 20% lower than the measured values. The simulated results also show that the supply of N and P to the growing region of the biofilm (illuminated zone) is not limiting at irradiances and CO₂ concentrations used for the simulation.

The observed lower [DO] in the simulated results using 5% supplementary gas phase CO₂ compared to the experimental data might be explained by the determination of the model parameters from biomass grown at a lower light intensity (300 μ mol photon m⁻² s⁻¹), which may lead to an inaccurate set of input parameters (e.g. a higher *ETR_{max}* and/or different Monod constants) for this specific growth conditions.

Figure 7: Dissolved oxygen concentration ([DO]), pH, dissolved inorganic carbon (DIC) and macronutrients (N as nitrate and P as total inorganic phosphate) profiles in Halochlorella rubescens biofilms exposed to 1,000 µmol photon m⁻² s⁻¹, and considering facilitated CO₂ transfer. Panel A and B present data simulated/measured with 0% and 5% additional CO2 in the gas phase, respectively. In each panel: [DO] profiles are presented in top-left subplots and bottom-left subplots give the pH profiles (in both left subplots, solid lines: simulated distribution; empty squares: experimental data measured by Li et al. 2015b). The top-right subplots present simulated DIC profiles (solid lines: total [DIC]; light gray area: bicarbonate; dark gray area: dissolved free CO2). The show the bottom-right subplots simulated macronutrient profiles (dark gray solid lines: N; light gray solid lines: P).



At 0% CO₂, the concentrations of macro-nutrients N and P decreased only slightly at the biofilm surface compared to the biofilm-membrane interface, whereas the [DIC] changed dramatically along the depth gradient. At 5% CO₂, the concentrations of N and P, especially nitrate, changed significantly along the depth gradient, however, this was a minor change compared to the change in [DIC] (5-fold greater difference than NO₃). Thus, it is very likely that the change of N and P concentrations along the depth gradient in the modeled biofilm has only a minor effect on the pH gradient. The main contributor shaping the pH gradient is the consumption of DIC.

Our data suggest that supply of N and P to the growing region of the biofilm was also not limiting at high irradiances and high CO₂ concentrations, even when the 20% underestimation of [DO] in the modeled results was taken into account (assuming a lower [DO] directly translates into lower productivity), provided that the nutrient concentrations in the bulk liquid phase remain constant. Thus, a direct upscaling of the model to 2D should be avoided (as the nutrient concentrations change with vertical distance from the medium outlet). Nevertheless, a pseudo-2D model based on the proposed model can be easily developed (IWA Task Group on Biofilm Modelling 2006), that can be used to predict growth along a vertically oriented cultivation area.

Effect of buffering on growth and DIC distribution

Fig. 8 presents the simulated growth rates and [DIC] profiles in *H. rubescens* PSBR biofilms exposed to 300 μ mol photon m⁻² s⁻¹ and three supplementary CO₂ concentrations (0%, 1% and 5% v/v, respectively), as well as medium with different buffering capacities (BBM without additional buffer and BBM with excessive buffering capacity, the latter was achieved by modeling the pH as a constant with a value of 6.8). The results show that the medium buffering capacity can influence both total [DIC] and its speciation in the biofilm. At 0% supplementary CO₂ in the gas phase, the growth for both well buffered and normal medium is mostly due to bicarbonate assimilation. The additional buffering of the medium actually decreased the [DIC]. At 1% supplementary CO₂, the growth based on bicarbonate surpasses growth based on CO₂ slightly. The buffered medium causes a decrease in [DIC] near the biofilm surface, however, this has no significant impact on growth rate. At 5% CO₂, growth utilizing CO₂ exceeds bicarbonate.

Figure 8: Simulated growth and dissolved inorganic carbon (DIC) profiles in *Halochlorella rubescens* biofilms exposed to 300 µmol photon m⁵ s⁷, and considering facilitated CO₂ transfer. Panel **A**, **B** and **C** present data simulated/measured with 0%, 1% and 5% additional CO₂ in the gas phase, respectively. In each panel: The left subplots present simulated growth profiles (black solid lines: CO₂ growth with normal medium; gray solid lines: CO₂ growth with normal medium; gray solid lines: CO₂ growth with normal medium; gray solid lines: CO₂ growth with excessively buffered medium; gray dashed lines: bicarbonate growth with excessively buffered medium). DIC profiles are presented in the right subplots with the profiles simulated with normal medium on top (solid lines: total [DIC]; light gray area: bicarbonate; diask gray area: dissolved free CO₂).



The results suggest that at low gas phase CO_2 concentration (no additional CO_2 supplied in the gas phase), by buffering the medium excessively, the pH at the biofilm surface is lowered. Thus, less bicarbonate is formed at the biofilm surface, leading to less DIC entering the biofilm. Finally, this results in reduced growth rates in buffered medium compared to normal medium (photosynthesis using bicarbonate is reduced). At 1% supplementary CO_2 in the gas phase, the buffering of the medium produces no significant change in growth rate distribution, even though the DIC distribution was affected. This might be caused by the limitation of growth by other factors (such as the CO_2 : O_2 ratio) rather than by the limiting supply of DIC at the given CO_2 concentration in the gas phase. By increasing the ambient CO_2 concentration to 5%, a slight increase in CO_2 -mediated growth was observed, most likely induced by an increase in the CO_2 : O_2 ratio. However, this was countered by the inhibiting effect of a high CO_2 concentration on bicarbonate utilization. As a result, the total growth rate (sum of CO_2 and bicarbonate growth) was not affected.

In the present study, the *H. rubescens* strain used for simulation is assumed to be pH insensitive (i.e. pH has no direct influence on growth) and can utilize bicarbonate for photosynthesis with relatively high efficiency. Thus, the buffering of the medium generally does not have a large effect on growth. However, if a more pH-sensitive strain and/or a strain with less efficient bicarbonate uptake mechanisms would be considered, the effect of medium buffering would be much more pronounced.

Long term biomass growth and effect of dark respiration

Figure 9 presents biomass growth data using continuous illumination (as the model currently is numerically unstable when a light/dark circle is considered) from both experimental (Schultze et al., 2015) and simulated data. In this growth simulation, the effect of temperature is taken into consideration: Algal growth was simulated with both the dark respiration rate measured at 23° C (see supplementary materials) and an adjusted dark respiration rate increased by a factor of 1.5, as the temperature in the growth experiments was elevated by about 5°C (Schultze et al., 2015). Here, the elevated temperature was assumed to have the same impact on gross photosynthetic production and photorespiration, thus the net growth (i.e. gross photosynthetic production minus photorespiration rate is affected by the elevated temperature (i.e. increase with increasing temperature) in the proposed model. And the correction factor for the increase of dark respiration rate with elevated temperature was estimated based on the observation of Hancke and Glud (2004). The simulated results show that biomass growth of *H. rubescens* was significantly affected by the dark respiration rate (i.e. temperature). Considering a higher dark respiration rate adjusted to the modified experimental conditions, simulated values fit the experimental data with very high fidelity.



Figure 9: Experimental (Schultze et al., 2015) and simulated growth data of *Halochlorella rubescens* biomass in a Twin-Layer porous substrate bioreactor using continuous illumination. Panel A, B and C give growth data from biofilms cultivated at 600, 800 and 1000 µmol photon m² s¹ surface irradiance, respectively. In all panels: Grey squares represent previously measured experimental data; solid lines gives the simulated data using an adjusted dark respiration rate to match the experimental temperature of 28 °C. Dashed line show the simulated data using dark respiration rate measured at 23°C.

Another possible reason for the increased dark respiration rate could be: in their experimental setup, Schultze et al. did not exclude gas transfer between the macro-porous material and the ambient air (Schltze et al., 2015), thus oxygen was supplied from the bottom of the biofilm to the non-illuminated region of the biofilm, and this presumably led to an increase in dark respiration. However, this effect affects only the bottom part of the biofilm, and its influence was small compared to the overall effect of temperature (data not shown).

The results presented here simulate growth of continuously illuminated *H. rubescens* biofilms, and thus should not be directly applied to systems with a light/dark cycle. The results strongly suggest that the dark respiration rate, during the dark phase or in the non-illuminated region of the biofilm, has a strong impact on biomass productivity of the TL-PBSR. In practice, by reducing this dark respiration (e.g. reduced temperature during the dark period), a higher total biomass productivity can be achieved.

Conclusions

Our results show that the proposed model in the present study can accurately predict the distribution of light intensity in TL-PSBR biofilms, provided the optical properties of the biomass are measured experimentally. With other model parameters acquired both from the literature and from experiments, the prediction of [DO] and pH profiles also reflect experimental data with high fidelity. Also, the simulated biomass growth reflexes the experimental results with high accuracy.

In the present study, the model was used to simulate TL-PBSR subjected to several growth scenarios for one algal strain by applying different sets of model parameters. The model can be used to predict the distribution of gradients under any given culture condition and/or for different algal strains, provided experiments are carried out to determine different parameters. Improvement of the model can be made in several aspects, e.g. a better description for the facilitated CO₂ transport and/or the effect of temperature; or, a more accurate description of the pigment adaptation mechanism. However, such improvements require more experimental observations and will increase the complexity of the model considerably. By presenting this approach, we have proven that our model considering various complex dynamics in a biofilm does have the capability to predict total biofilm growth rates with considerable accuracy. Therefore, with some further modification, it may be a valuable tool in system optimization in PSBR and process design

Acknowledgements

This study was supported by the University of Cologne (KST 158901001). The authors thank Bastian Piltz (University of Cologne, Cologne, Germany) for his valuable opinions, Nico Helmsing (Netherlands Institute of Ecology, the Netherlands) for his assistance in determination of biomass compositions, Lisa Marie Blümel

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(University of Cologne, Cologne, Germany) and Kenneth Dumack (University of Cologne, Cologne, Germany) for their assistance in model parameter determinations.

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Supplementary materials

General information

In this study, a Twin-Layer biofilm of the green alga *Halochlorella rubescens* (CCAC 0126; Culture Collection of Algae at the University of Cologne; www.ccac.uni-koeln.de) was modeled. Some of the parameters used in the present study were acquired experimentally, as described below. All statistical analysis of the experimental results presented here were carried out using Prism (version 5.1, Graphpad, GraphPad Software Inc., La Jolla, USA).

Experimental studies

Biofilm cultivation

To investigate the growth of the biofilm and to characterize the biomass, H. rubescens biofilms with an inoculation biomass concentration of 5 g m-2 were produced and cultivated as described by Li et al. (2015) (batch setup, with circulating medium): BBM culture medium (Bischoff and Bold, 1963) and a light intensity of 300 μ mol photons m-2 s-1 with a 14/10 light/dark cycle were used for the whole cultivation period, and the biofilms were aerated with 0.75 L min-1 ambient air.

The freshly harvested biofilm samples were measured for fresh weight, and biofilm thickness was measured using a hand microtome as described by Li et al. (2015). The biofilm volume was estimated by multiplying biofilm area and thickness. The samples were then dried at 105 °C for 2 hours prior to dry weight (DW) determination.

Long term growth experiment

To investigate the growth of the biofilm and to characterize the biomass, H. rubescens biofilms were cultivated for 100 days, and triplicate samples were taken twice a week. Subsequently, biomass was digested, and its N and P contents were determined using the method by (Hu and Barker, 1999). Finally, to achieve a more accurate determination, the nitrogen and phosphorus contents (w/w) of the N and P contents of samples from 100 day were also determined using an elemental analyzer (Flash 1112, Interscience, Rockland, MA, USA). Biomass concentration (X_F) was calculated as dry biomass per fresh biofilm volume (in kg m⁻³). In the present study, the biomass concentration is considered to be constant, as: A linear regression of the data does yield a line with a slope significantly different from zero. However, after day 40, most of the dry biomass concentration data points were not significantly different from each other, as suggested by ANOVA test. Thus, we suspect that the significant different at the beginning of the cultivation is caused by error in biofilm thickness measurements, as the biofilm is too thin to be measured accurately. Even if the biomass concentration change observed was caused by reasons other than inaccurate biofilm thickness measurements, the exact reason and mechanism are unclear to us. From our point of view, and also for the simplicity of the model the assumption that the biomass concentration remains constant during the cultivation period was made. The results are present in Fig. S1, S2 and S3. Biomass yield on C, N and P are calculated as the reciprocal of their respective biomass contents. A C:N:P ratio of 40:7:1 was used for the present study (for calculated yields, refer to Tab. II).



'Figure SI: Biomass of Halochlorella rubescens biofilms during 100 days of cultivation. A: Biomass dry weight (DW), the dotted line is a one-phase association curve fitted to the data. B: Biomass water content in w/w (DW/fresh weight), dotted line represents the linear regression line of the data. C: Biofilm thickness, dotted line represents a linear regression. D: Biomass concentration (DW/fresh volume), dotted line represents a linear regression. In all panels, error bars represent the standard deviations of triplicates.



Figure S2: Biomass N and P contents of Halochlorella rubescens biofilms during 100 days of cultivation. A: Biomass N content (w/w), empty circles and error bars represent the means and standard deviations of triplicates; dotted line represents the linear regression line of the data. B: Biomass P content (w/w), empty squares and error bars represent the means and standard deviations of triplicates; dotted line represents the linear regression line of the data.

Figure S3: Carbon, nitrogen and phosphorus content of dry biomass of *Halochlorella rubescens* measured with an elemental analyzer. The grey bars and error bars represent means and standard deviations of 6 replicates. The horizontal dot-dash lines and the numbers next to the bars give the C, N and P contents of biomass used in the proposed model.



Experiments investigating effects of nutrient concentrations on biomass

To investigate the effects of different nitrogen and phosphorus concentrations in culture medium on biomass N and P content, a separate growth experiment with the same cultivation conditions except using a continuous system (medium was pump only once through the system, instead of circulating, as described by Li et al. (2015), was carried out. In this experiment, after inoculation, BBM medium with modified N or P concentrations was supplied to the Twin-Layer system. The biofilms were cultivated for 4 days, and 6 parallel samples were analyzed. The

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samples were first dried for DW determination and the N and P contents were then determined with digestion (as described above). The results are shown in Fig. S4. The biomass composition is considered stable during the simulated growth (i.e. the N, P concentrations do not have impact on biomass composition), as indicated by the experimental results.



Figure S4: N and P contents of *Halochlorella rubescens* biomass from biofilms cultivated using medium with modified N or P concentrations. A: N content (w/w) of biofilms cultivated using medium with modified N concentrations. B: P content (w/w) of biofilms cultivated using medium with modified P concentrations. C: N content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations. D: P content (w/w) of biofilms cultivated using medium with modified P concentrations of N or P in non-modified standard Bold's basal medium (BBM).

Determination of dark respiration rate

For estimating the dark respiration rate, 24 samples were harvested on day 12 of the 100 day growth experiment (see section below). 12 samples were dried for DW determination, the other 12 were put into a bioreactor system under the same cultivation conditions but covered with aluminum foil to exclude light. After 24 hours, these 12 samples were removed from the system and their DW was determined. The dark respiration rate (in d^{-1}) was determined as:

$$R_d = \ln \left(\frac{DW_{t=1}}{DW_{t=0}}\right) \tag{Eq. S1},$$

 $DW_{t=0}$ and $DW_{t=1}$ are the mean DWs of samples dried before and after the dark incubation in darkness, respectively. An R_d of 0.13 d⁻¹ was calculated for the present study.

Measurement of the electron transfer rate

A depth profile of the electron transfer rate (ETR, in µmol electrons m⁻² s⁻¹) of the 100 days old *H. rubescens* biofilms (3 parallels, i.e. 3 independent samples) was acquired between 0-600 µm using the pulse amplitude modulation (PAM) method described by Li et al. (2015). The data acquired were fitted with a P-I model (Platt et al., 1980). Based in the fitted model the maximal ETR (ETR_{max}) and the initial ETR curve slope (α) were determined. The results of the ETR measurement are presented in Fig. S5 and S6. In the present study, ETR_{max} and α was set to be 49.6 mole m⁻² s⁻¹ and 0.24, respectively (the average value of the 'illuminated zone'; as suggested by data presented by Li et al., 2015, Fig S5). The maximal biomass growth rate (μ_{max}) was estimated from the ETR_{max}: Assuming 1 mole of electron transferred corresponds to 0.16 mole of 0₂ production (Morris and Kromkamp, 2003); and the ratio of O₂ production to inorganic carbon consumption is 1:1.3 (Atkinson and Smith, 1983). In the present study, the calculation yielded a maximal growth rate of 8 d⁻¹.



Figure S5: Electron transfer rate (ETR) measured at different depths in *Halochlorella rubescens* biofilms grown at a surface irradiance of 300 µmol photons m⁻² s⁻¹. A: ETR at 0 (biofilm surface, empty circles), 50 (empty squares), 100 (empty triangles) and 150 µm depth (empty diamonds). B: ETR at 200 (empty circles), 250 (empty squares), 300 (empty triangles) and 350 µm depth (empty diamonds). C: ETR at 400 (empty circles), 450 (empty squares) and 600 (empty triangles). In all panels, the symbols and error bars give the means and standard deviations of triplicates, and the dotted lines represent the fit curve of a P-1 model (see text).



Figure S6: Maximal electron transfer rate (ETR) and initial slope of the ETR curve plotted as a depth profile in the Halachiorelia rubescens biofilms. Values were calculated from the P-I model fitted to the ETR curve (see Fig. S3 and text). A: Maximal electron transfer rate (ETR_{max}). B: Initial slope of the ETR curve (α). C: Inhibition coefficient estimated from ETR curve (β). The biofilms were separated into two sections according to the results presented by Li et al. (2015): A well illuminated section subjected to low light (solid circles, < 5 µmol photons m² s⁻¹) and a section subjected to low light (solid circles, < 5 µmol photons m² s⁻¹). In all panels, the dot-dash lines and numbers represent the average values (values used in the present study as model parameters) in well illuminated sections.

Determination of optical properties of biomass

To measure the optical properties of the biomass, fresh biomass samples (triplicate, i.e. 3 samples) from a 3 days old Twin-Layer biofilm were suspended in BBM to obtain a biomass concentration of 0.05 g L⁻¹. Ultrasonication was carried out to separate the cells. The specific extinction, absorption and scattering coefficients (β_{λ} , κ_{λ} and σ_{λ} ; extinction = absorption + scattering) of the cell suspension were measured as described by (Berberoglu and Pilon, 2007): A photospectrometer (UV-2450, Shimadzu, Duisburg, Germany) equipped with an integrating sphere (ISR-240A, Shimadzu, Duisburg, Germany) were used; and a flow cell with 0.1 mm light path (FT04-06, Thermo, Duisburg, Germany) was employed to keep the cells in suspension during the measurements. The mass extinction, mass absorption ($M_{absorption}$ and $M_{scattering}$) and mass scattering were calculated according to (Berberoglu et al., 2007) with 1 nm wavelength resolution (coefficient divided by biomass concentration, m² kg⁻¹). The results of the measurements of the optical properties are given in Fig. S7. The absorption and scattering coefficient of the *H. rubescens* Twin-layer biomass in the present study ($\kappa_{\lambda,F}$ and $\sigma_{\lambda,F}$), are calculated from $M_{absorption}$, $M_{scattering}$ and X_{F} :

$$\kappa_{\lambda,F} = M_{absorption} \cdot X_F \tag{Eq. S2}$$

and,

$$\sigma_{\lambda,F} = M_{scattering} \cdot X_F \tag{Eq. S3}.$$



Figure S7: Optical properties of biomass from Halochlorella rubescens biofilms grown at a surface irradiance of 300 µmol photons m² s² A. Si Diomass mass extinction in the photosynthetically active radiation (PAR) region. B: Biomass mass absorption in the PAR region. C: Biomass mass scattering in the PAR region. In all panels, empty circles represent mean values of triplicates. Notice the differences in y-axis scales.

Determination of effective diffusion coefficients in biofilm

In present study, the X_F was set to 220 kg m⁻³ and assumed to be a constant both over time and biofilm depth (even though the linear regression showed a slope significantly different from 0, Fig. S1); Biomass C, N and P content are also considered to be stable over time and biofilm depth (as experimental results indicated little effect of both different medium nutrient concentrations and cultivation time on biomass composition). The effective diffusion coefficients (of a species *x*) in biofilm and membrane ($D_{e,x}$ and $D_{m,x}$) are estimated from biomass and membrane porosity ($\theta = 0.76$ and 0.35, respectively, Whatman PC-40, Whatman, Freiburg, Germany; Li et al., 2015) using the formula:

$$D_{x,e} = D_x \cdot \frac{\theta}{1+0.5 \cdot (1-\theta)} \tag{Eq. S4}$$

(Weissberg, 1963). In Eq. S4, D_x represents the diffusion coefficient of species x in water, whose values can be found in Tab. II.

Experimental verification of biomass pigment adaptation

In order to verify biomass pigment adaptation, an experiment was carried out to determine the effect of light intensity on biomass chlorophyll content. *H. rubescens* biofilms were cultivated using the batch setup with identical cultivation conditions as previously described (see 0). However, instead of using only one light intensity, several light intensities were used ($50 - 1000 \mu$ mol photons m² s⁻¹ surface irradiance). Biofilms were harvest after 3 days and the biomass resuspend in BBM prior to mass absorption determination. Chlorophyll a content of biomass was estimated with the mass absorption as 675 nm. The results are presented in Fig. S8, which shows a significant effect of irradiance on mass absorption at 675 nm (One-way ANOVA).





Effect of nutrients concentrations on biomass growth rate

Using a similar approach as described in supplementary material section 2.3, and with identical cultivation conditions, the growth rates of *H. rubescens* biofilms using BBM medium with different nitrate and phosphate concentrations were determined. The biomass growth was observed from day 2 to day 6 after inoculation, and the

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growth rates were taken as the slope of biomass increase between day 2 and day 6. The results are present in Fig. S9. The data suggest that at the concentrations encountered in normal cultivation conditions, N and P do not inhibit biomass growth.



Figure S9: Effect of BBM medium with modified nutrient concentrations on the growth of *Halochlorella rubescens* biofilms. A: Growth of *H. rubescens* biofilms using BBM medium with modified nitrate concentrations (as indicated in the x-axis label). B: Growth of *H. rubescens* biofilms using BBM medium with modified phosphate concentrations (as indicated in the x-axis label). Filled bars and error bars represent the mean values and standard deviations of 6 replicates.

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3. Discussion

As the first step in this study, a new experimental setup and new mathematical method were developed to analyze the data acquired during the experimental studies. Then, a comprehensive and systematic investigation was carried out to study the dynamic processes inside a PSBR. In this second part, both experimental investigations with microsensors and modeling of the PSBR phototrophic biofilm were carried out. The present study established a solid basis for future investigations of the PSBR biofilms. The results acquired by this study offer the first deep insight into the PSBR biofilm. From the results, several valuable conclusions on previously established hypotheses and new hypotheses have been made.

3.1. Improvement of data analysis through the new method

As discussed previously (see manuscript 1; Li et al. 2015c), the drawbacks of the gross photosynthetic productivity measurement with oxygen microsensors in phototrophic biofilms have been known since the early 1980s, and various methods have been developed to improve its accuracy (Glud et al. 1992; Lassen et al. 1998; Revsbech 1983). However, all the methods developed until now require an assumed 'true' distribution or modification of the biofilm surface (e.g. application of a thin agar layer onto the biofilm surface), and cannot be applied if the shape of the 'true' (i.e. original gross productivity distribution) is unknown. In general, the data treatment in such measurements tries to correct the inaccuracy caused by the loss of dissolved oxygen (DO) through the biofilm surface to the phase above during the measurement period (Revsbech 1983). In a biofilm whose surface is directly exposed to a gas phase, the inaccuracy caused by this phenomenon leads to a large error when the data was analyzed directly or when previously developed methods were applied to correct the raw data.

As a result, a new mathematical method was developed in the framework of this study to correct for this effect. The results (both numerical and experimental) show the effectiveness of the proposed method. The results show clearly the significant impact that an exposed biofilm surface has on gas exchange. An important point made in the present study is that, in order to effectively apply this new data treatment method, a fine measurement resolution ($\leq 20 \,\mu$ m) has to be used. The establishment of this method paved the way for the photosynthetic productivity measurements of the investigated PSBR biofilm using microsensors, and deepened the understanding of the transport processes (especially DO) in such biofilms. The proposed method was developed to analyze data acquired using the light-dark shift method for a non-submerged biofilm, however, with minor modifications of the boundary conditions applied (as discussed in manuscript 1), the method can also be used for data acquired using the same light-dark shift method from submerged biofilms.

3.2. Light transfer in phototrophic biofilms

As observed by Wang et al. (2015), the immobilization of algal cells in biofilms increases the light dilution rate of the total biovolume (i.e. photons available per cell) compared to that in a suspension culture. It has been long suspected, that in a phototrophic biofilm, besides adsorption, both forward-scattering of light and the adaptation of biomass (e.g. changes in pigment contents) can have significant impact on light transfer (Berberoglu and Pilon 2007; Jøsrgensen 1969). It has also been suspected, that these effects would result in a more efficient light utilization by cells immobilized in biofilms.

In this study, a direct verification of this hypothesis was carried out for the first time, by comparing modeled results using three different scenarios: 1) incorporating only extinction (no forward-scattering or biomass pigment adaptation); 2) with forward-scattering but no biomass pigment adaptation and 3) with both forward-scattering and pigment adaptation taken into account to experimental data measured with irradiance microsensors. The high accuracy of the model prediction when both forward-scattering and pigment adaptation proved clearly, that the above mentioned hypothesis is correct. Thus, in future studies investigating radiative transfer (i.e. light transfer) in phototrophic biofilms, the effects of both forward-scattering and biomass pigment adaptation have to be taken into account. However, to achieve an accurate evaluation of the effects of forward-scattering and cell pigment adaptation, experiments similar to that conducted in this study have to be

carried out in order to determine the algal strain and/or cultivation condition specific parameters (e.g. scattering coefficient).

An important point that has not been addressed in this study is the effect of photosynthetically active radiation (PAR) of different wavelengths. The solution of the radiative transfer equation (RTE) applied in the model as proposed in the present study does consider the effects of different wavelengths on light transfer. However, in this study, the PAR with different wavelengths is considered to have the same effect on photosynthetic productivity, hence the term 'total PAR' was used. However, it is general knowledge, that different PAR wavelengths have different effects on photosynthetic efficiency. Thus, in future studies, one should consider incorporating this effect, and experimental studies have to be carried out to determine strain and/or cultivation condition dependent cellular pigment composition. The results can be then used to evaluate the efficiency of PAR with different wavelengths.

3.3. Effects of surface evaporation of a non-submerged biofilm surface

The analysis reveals that the fundamental processes in a PSBR biofilm and a 'traditional' (i.e. submerged) phototrophic biofilm are very similar, as both are subjected to diffusion, convection and have identical biological processes (e.g. photosynthesis and respiration; Murphy and Berberoglu 2014; Wolf et al. 2007). However, due to their essential differences in system configuration, compared to a submerged biofilm, some processes in PSBR biofilms are enhanced (e.g. gas transfer at the biofilm surface), and others weakened (e.g. accumulation of DO inside the biofilm). A marked difference between a PSBR biofilm and a submerged biofilm is that the former is subjected to surface evaporation, as in a PSBR the biofilm surface is exposed directly to the ambient gas phase. This surface evaporation creates an outward flow of liquid in the PSBR biofilm perpendicular to the biofilm surface. This, in effect, adds a flux inside the PSBR biofilm towards the biofilm surface (Bergman et al. 2011). The effect of this additional flux is an enhanced mass transfer in the direction perpendicular to the biofilm surface towards the surface of the biofilm. This leads to an enhanced nutrients transfer from the bottom of the biofilm to its surface. Also, due to an extremely thin mass transfer boundary layer at the biofilm surface, an enhanced gas exchange between the biofilm and the phase above (a gas phase for a PSBR biofilm and a liquid phase for a submerged biofilms) is achieved (Bergman et al. 2011; Schlichting and Gersten 2000). The combined effect of the additional flux and the enhanced gas exchange is an increased removal rate of oxygen produced by photosynthesis from the inside of the biofilm. The effect on [DO] in the biofilm is guite significant, as has been concluded in the first manuscript (Li et al. 2015c) presented in this thesis. However, even with this enhanced gas transfer, exceptionally high [DO] has been measured in the studied biofilm (equals to 12 times of the normal oxygen partial pressure in air under normal condition). Results from the modeling study proved that it is theoretically possible to achieve such high DO level in the investigated biofilm, but only if no oxygen is released as gas from the liquid phase inside the biofilm. However, the mechanism behind this over-saturation of oxygen in the investigated biofilm without the formation of gas bubbles is still not completely clear. Also, it is not clear how the investigated algal cells still maintained relatively high photosynthetic activity at this high level of DO. This phenomenon should be investigated further in future studies. Nevertheless, this over-saturation of oxygen could prove useful in biotechnological applications (e.g. oxygenation and/or odor removal in wastewater treatment), if this over-saturation would be transferred to the bulk liquid phase (i.e. culture medium).

Similar to oxygen, the exchange of CO₂ between the biofilm and the phase above is enhanced in a PSBR biofilm. However, this enhanced CO₂ exchange alone is not sufficient to supplied enough dissolved inorganic carbon into the deeper part of the biofilm to support the high photosynthetic activity (i.e. fast growth) that has been observed in investigations conducted both in the framework of this study (e.g. comparing the modeled data with measured data) and elsewhere (Schultze et al. 2015). A new hypothesis has been developed based on the findings made in the present study, which is discussed in the following section.

3.4. Supply of inorganic carbons in phototrophic biofilms

In this study, the following is hypothesized: At low CO_2 concentrations (i.e. without additional CO_2 in the gas phase) the facilitated CO_2 transfer in the investigated biofilm significantly increased the DIC transport efficiency

inside the PSBR biofilms. The phenomenon of facilitated CO_2 in solutions when carbonic anhydrase (CA) is present have long been observed, and has already been applied to enhance CO_2 transfer in artificial membrane systems (Bao and Trachtenberg 2006; Gros et al. 1976; Trachtenberg et al. 1999). Thus, it can also be expected in the studied biofilm. It is further hypothesized: in phototrophic biofilm, both the liquid between the cells and the surface of the cells serves as 'carrier' for the facilitated CO_2 transport, thus leading to a very strong DIC transport enhancement at low CO_2 concentrations (Bao and Trachtenberg 2006; Gros et al. 1976).

Until now, facilitated CO_2 transfer has not been reported in phototrophic biofilms. This study presents the first evidence of the existence of facilitated CO_2 transfer, and its effect on dissolved inorganic carbon (DIC) transport in a highly active phototrophic biofilm. The lack of reports on observed facilitated CO_2 transport in such biofilms until now might be due to several reasons: Firstly, the most extensively studied phototrophic biofilms are natural phototrophic biofilms, and most of these biofilms are mixed biofilms (containing significant portion of both heterotrophic and phototrophic components) as defined by Wolf et al. (2007). As CO_2 is produced by heterotrophic components inside the biofilm, which is then provided to the phototrophic organisms (i.e. increases the inorganic carbon availability inside the biofilm), the effect of the facilitated CO_2 from biofilm surface into the inside of the biofilm becomes less significant. Secondly, it is very likely, as discussed in the 2nd manuscript (Li et al. 2015b) presented in this thesis, that the DIC supply in a submerged biofilm is limited due the liquid phase above the biofilm (i.e. compared to a direct exposure to the ambient gas phase) and not by the transport inside the biofilm itself. Nevertheless, this hypothesis has to be verified independently, and additional experimental studies have to be carried out (e.g. direct determination of DIC profiles in PSBR biofilms, or verification of extracellular CA in the biofilm).

3.5. Carbon availability and pH

As suggested by the results, the facilitated CO_2 transfer plays an important role in transporting DIC to the deeper part of the biofilm when CO_2 is supplied in very low concentration in the gas phase. However, as the algal cells can only take up DIC in free CO_2 , and some strains also in bicarbonate form (as assumed in the present study), the total [DIC] in the biofilm does not equal to the available DIC (e.g. when carbonate dominates, the biologically available carbon is much less than total [DIC]). As discussion previously (see manuscript 2 and 3, Li et al., 2015b and Li et al. Manuscript under review), the speciation of DIC is pH dependent, and their relationship can be estimated using a standard textbook DIC speciation chart: Generally, the higher the pH, the higher the proportion of carbonate in the total DIC, which is not available for uptake by the algal cells. However, in a phototrophic biofilm system, the interaction between pH and DIC speciation and consumption is a bidirectional process: DIC uptake (i.e. free dissolved CO₂ and bicarbonate) increases the local pH, this leads to less CO₂ and bicarbonate in the total DIC, which again influence the DIC uptake rate. In the present study, the proposed model can describe this process with relatively high accuracy, as indicated by the results.

Also, contrary to previous assumptions, the results show that uptake of macronutrients (i.e. N as nitrate and P as total phosphate) has no significant or only very small influence on local pH. Results from both experimental and modeling investigations show, that the pH in the investigated PSBR biofilm can reach high values even when a buffered medium (e.g. BBM, buffered with phosphate buffer in the present study) is used for cultivation. This indicates the existence of a cellular level pH regulation mechanism, at least in the case of *H. rubescens* used in the study, which sustained the photochemical activity of the cell at very high pH values. This pH regulation mechanism is possibly related to the carbon concentrating mechanism, as reviewed by Raven et al. (2008). And as has been shown in the present study, this mechanism plays a very important role in the investigated PSBR biofilms.

In the present study, the concentrations of solutes (e.g. macronutrients and DIC) in the medium (i.e. the bulk liquid phase, see manuscript 3) were considered to be time-independent. This was experimentally achieved by two different approaches: 1) By changing the medium very often and assumes the minor change in solute concentrations which occurs between each change had no significant influence on the results (as in all experimental investigations carried out in this study using a batch setup) or, 2) by keeping the concentrations in the medium constant (as in experiments using a continuous setup and in the model). In a real cultivation situation however,

Disccusion

when the medium is circulated in the system for a longer period of time, the cumulative effect of nutrients uptake, as observed by (Shi et al. 2007), and its effect on pH of the medium have to be taken into account.

3.6. Long term growth of PSBR biofilm

As discussed previously, due to the absence of a moving liquid layer above the PSBR biofilm, PSBR biofilms are not subjected to biofilm detachment. However, a reduction of growth rate with time in prolonged cultivation has been observed both in this study and in other studies (Liu et al. 2013; Li et al, 2015; Schultze et al. 2015). In the present study, the proposed model was used to predict the growth of the investigated biofilm in a 10 days period under continuous illumination. The results show that the shape of the growth curve was directly influenced by the dark respiration rate. The analysis of results indicates, it is very likely, that this reduction in growth observed during prolonged cultivations is due to the thickening of the 'dark region' (i.e. region of the biofilm that is not illuminated) as a result of the increase of the thickness of the biofilm. As in these dark regions, the biomass is consumed (i.e. respired) rather than produced. In the present study, growth simulation with a light/dark circle was not performed due to the instability of the model. When a light/dark circle is applied however, a thicker biofilm, thus resulting also in a reduced daily growth rate of the thicker biofilm compared to the thinner one under identical cultivation conditions.

3.7. Closing remarks

Through this study, important new insights into the dynamic processes in PSBR biofilms have been acquired (as discussed in the sections above). A solid basis for investigation together with powerful tools (microsensor measurements, the proposed model etc.) has been developed to help understand the processes inside such biofilms. By modifying the model parameters, the model can easily be applied to BSPR biofilms with other algal strains and/or different cultivation conditions (e.g. medium, temperature). However, experimental investigations similar to that performed in this study have to be carried out in order to acquire the parameters required by the model to achieve a more accurate prediction. With minor modifications, the model can also be applied for system optimization in large-scale operations. The present study provides several interesting and important topics that should be investigated in the future: E.g., the effect of irradiance spectrum on photosynthetic activity, investigations into the facilitated CO₂ transfer in the studied biofilm; the mechanism behind the exceptionally high [DO] measured in the biofilm and a more comprehensive modeling study aimed at system optimization.

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Summary

In the present study, a comprehensive investigation of the dynamic processes in an artificial algal biofilm immobilized on a porous substrate has been conducted. Experimental investigations including microsensor measurements were carried out. For this purpose, the microsensor setup used for profiling submerged biofilms was modified to enable measurement on the investigated biofilms. To achieve an accurate evaluation of the data acquired through microsensor measurements, a new mathematical method was developed, and a 20 µm depth resolution has been suggested for future photosynthetic activity measurements.

After the establishment of the microsensor methods, a systematic microsenser investigation was carried out: the distribution of dissolved oxygen, pH value and photosynthetic productivity profiles of algal biofilms in a porous substrate biofilm photobioreactor (Twin-Layer photobioreactor) exposed to different surface irradiance and/or exposed to different gas phase CO_2 concentrations were measured. The results acquired from these experiments offered important insights into the processes in such biofilms: E.g. light penetration depth, maximal dissolved oxygen concentration and pH distribution. The results show, as expected, photosynthesis in the biofilms occurs only near the biofilms surface (i.e. in the illuminated zones), and dark respiration in the inner part of the biofilm surface irradiance and/or gas phase CO_2 concentrations led to an increase in photosynthetic productivity of the investigated biofilm. No photoinhibition was observed in the studied biofilms, although exceptionally high dissolved oxygen concentrations (12 times of that in normal atmosphere) have been recorded.

The model (as described in the 3^{rd} manuscript, Li et al. 2015d) developed in the study has proven to be very effective in predicting experimental observations. The results show clearly the importance of taking into account not only adsorption and scattering, but also the adaptation of the pigment content of the biomass for investigating radiative transfer in PSBR biofilms. Also, through the development of the model, important insights into the dynamic processes in the investigated biofilm were acquired: E.g., it is very likely that the facilitated CO₂ transfer plays an important role in inorganic carbon transport in the studied biofilms when the CO₂ concentration supplied in the gas phase is low; macronutrients (N and P) do not limit growth even at high surface irradiance and high gas phase CO₂ concentrations as long as they are sufficiently supplied in the medium; and the buffering of the medium with a strong buffer will have significant effects on the inorganic carbon availability in the studied biofilm.

Through this study, a solid basis has been established for future investigation on PSBR biofilms. The methods and model developed in this study are established specifically for investigating biofilms grown in the Twin-Layer porous substrate biofilm photobioreactor. However, with minor modifications and/or additional experimental measurements, they can be easily applied to other phototrophic biofilm systems or for investigation and/or optimization of commercial scale systems.

Zusammenfassung

In der vorgelegten Studie wurden die dynamischen Prozesse in einem auf porösem Substrate künstlich gezüchteten Algenbiofilm umfassend untersucht. Experimentelle Untersuchungen, u.a. Mikrosensormessungen, wurden durchgeführt. Zu diesem Zweck wurde ein neuer Versuchsaufbau, sowie eine neue mathematische Berechnungsmethode zur Auswertung der durch Mikrosensormessungen aufgenommenen Daten entwickelt. Die Untersuchungen zeigten, dass um die Daten sinnvoll auswerten zu können, zukünftige Mikrosensormesungen am untersuchten Biofilm mit einer Tiefenauflösung von 20 µm durchgeführt werden sollten.

In nachfolgende systematische Mikrosensormessungen wurden die Profile von gelöstem Sauerstoff, pH und photosyntetischer Produktivität an Biofilmen gemessen. Die Messungen wurden an bei vieschiedenen Lichtintensitäten gezüchtet und/oder verschiedenen CO₂ Konzentrationen ausgesetzt Biofilmen durchgeführt. Die Ergebnisse ermöglichen wichtige Einsichten in den obengenannten Biofilm, wie: z.B. Lichtverfügbarkeit, maximale Konzentration von gelöstem Sauerstoff und pH Werte. Die Ergebnisse zeigen, dass die Photosynthese wie erwartet nur nahe der Biofilm-Oberfläche, beziehungsweise in der beleuchteten Region des Biofilms, stattfindet. Außerdem trägt die Dunkelresipration in der tieferen Schicht, beziehungsweise der nicht oder nur schwach beleuchteten Region des Biofilms zu der beobachteten Abnahme von Biomassewachstum bei längerer Kutivierungszeit bei. Die Daten zeigen ebenfalls, dass die photosynthetische Produktivität im untersuchten Biofilm trotz der gemessenen außgewöhnlich hohen gelöste Sauerstoffkonzentration (12 fach der Konzentration in normaler Luft) nicht beobachtet.

Die Ergebnisse beweisen, dass das in dieser Studie vorgelegte Model die experimentellen Beobachtungen mit angemessener Präzision vorhersagen kann. Um das Transferverhältnis des Lichts in dem untersuchten Biofilm präzise modellieren oder untersuchen zu können, sind nicht nur die Absorption des Lichts sondern auch dessen Streuung, sowie die Anpassung der Algenzellen an verschiedene Lichtstärken zu berücksichtigen. Das Model offenbart weiterhin wichtige Einsichten in die dynamischen Prozesse des untersuchten Biofilms: z.B. 1) Es wurde festgestellt, dass der sogenannte "Facilitated CO₂ Transfer" eine sehr wichtige Rolle für den Transport des gelösten anorganischen Kohlenstoffs in dem untersuchten Biofilm, insbesondere bei geringem CO₂ Gehalt in der Gasphase spielt; 2) Wenn ausreichend Makro-Nährstoffe (Stickstoff und Phosphor) im Kuturemedium vorhanden sind, stellt der Transport (z.B. Diffusion) der Makro-Nährstoffe kein Hindernis für das Wachstum dar, d.h. Makro-Nährstoffe Verfügbarkeit des gelösten anorganischen Kohlenstoff in deutlichem Maße beeinflussen.

Die vorgelegte Studie hat eine solide Basis für zukünftige Untersuchungen am PSBR Biofilm geschaffen. Die im Rahmen dieser Studie entwickelten Methoden eignen sich für die Untersuchungen des obengenannten Biofilms. Allerdings können mit geringfügigen Modifikationen und ggf. mit zusätzlichen experimentellen Untersuchungen die hier vorgelegten Methoden auch für andere phototrophische Biofilme und/oder für die Optimierung kommerzieller Systeme, die auf die untersuchten Biofilm basieren, verwendet werden.

Appendix I: MATLAB code for measured photosynthetic productivity data treatment

The MATLAB code presented here was written as described and applied in the 1st manuscript presented in this

thesis.

```
function [
xfinal,Measured,STD,calback]=
PIcorrectNN( testX, testLfcum
,realdata,nLf,dataSTD )
,realdata,nLf,dataSTD )
%input data:testX,testLfcum:measured
value in, thickness of in test problem.
realdata:Real data in matrix form, nLf:
thickness of the biofilm in discrete
form, dataSTD: STD of the measured data
if nargin<3 || realdata==0,
     realdata=0;
testXSTD=zeros(1,length(testLfcum));
    nLf=length(testX);
     originaldata=[];
end,
%% parameters input
DO2=1.97e-9; %diffusion coefficient of
02 in water in m^2/sec
Poro=0.32; %volumetric porosity
DO2bio=D02.*(ones(size(testLfcum)).*
Poro^1); %Diffsion coefficient
inside biofilm
DO2bio=DO2.*Poro^1;
uz=4.5E-06./(ones(size(testLfcum)).*
Poro^(1/3)); %advection rate caused by evaporation on the surface
uz=4.5E-06/Poro^(1/3);
t=1;
%% fit curve to the realdata
if realdata==1
     if nargin<4 && realdata==1,
error('For real data enter
number discretion points'),
     end.
if nargin<5 && realdata==1,
    warning('No STD input for re
data, set to 5% of the realdata'),
                                            for real
          dataSTD=testX.*0.01;
     end.
originaldata=[testLfcum;testX;dataST
D1:
     [~,~,dCdtcurvefit ,STDcurvefit] =
dCdtcurvetestSlm(
testX,testLfcum,dataSTD ,nLf); %use
Slm tool for data curve fitting
testLfcum=linspace(0,testLfcum(end),
nLf);
     testX=dCdtcurvefit.
     testXSTD=STDcurvefit;
end,
    calculation
[ xfinal,Measured,STD,calback] =
testGtikhonovopsNN( testX, testLfcum
,realdata,testXSTD,DO2bio,uz,t,origi
naldata);
end
function [
runction [
xfinal2,Btotal1,BSTD,Recoverdfx2] =
testGtikhonovopsNN(testX,testLfcum
,realdata,testXSTD,D02bio,uz,t,origi
naldata)
            effective diffusion
D02bio: effective diffusion
coefficient of D0, uz: evaporation
caused convection rate, t: measurement
%% calculation
[L,~] = get_l(length(testLfcum),2);
%data input, here a test problem or real
data if realdata==1
x2sum=zeros(length(testX),1);
Btotallsum=x2sum;
```

BSTDsum=x2sum; if realdata == 1. ncal=3; else ncal=1: end, for i=1:ncal, if realdata == 1, [Atotall, Ain1, ~, ~, ~, ~, ~] = testABgeneration (testX,testLfcum,DO2bio,uz,t); Btotal0=testX': % real data input [UUcal,smcal,XXcal] = cgsvd [xcal] =tikhonov(UUcal,smcal,XXcal,Btotal0, lambdacal); ~, ~, Btotal0, Bin0, ~] = [~,~,~, testABgeneration BSTD=((testXSTD)'-(-(testXSTD)')).*r and(size((testXSTD)'))+(-(testXSTD)' and(size((testXSTD)'))+(-(testX); %add artificial random error representing noising data Btotall=Btotal0+BSTD; Btotal1(Btotal1<0)=0; Bin1=Bin0+BSTD; Bin1(Bin1<0)=0; [~, Ain1, ~, Btotal0, Bin0, ~] estABgeneration (testX,testLfcum,DO2bio,uz,t); % error input BSTD=max(Btotal0)/50*randn (size(Btotal0));%.*(Btotal0./max(Bto tal0)); %add artificial random error representing noising data Btotal1=Btotal0+BSTD; Btotal1(Btotal1<0)=0; Bin1=Bin0+BSTD: Bin1(Bin1<0)=0; end [UU1,sm1,XX1] = cgsvd (Ain1,L); lambdal = 1_curvenoP (UU1, sml, Bin1); [x1] =tikhonov(UU1, sm1,XX1, Bin1, lambda1); x1(x1<0)=0; %projected
non-constrain tikhonov</pre> % solve for non-negative solution x0=x1:%zeros(size(testLfcum)); Lcal=[zeros(1,length(testLfcum));zer
os(1,length(testLfcum));L]; H=(Ain1'*Ain1+lambda1^2.*Lcal'*Lcal) f=(-Ainl'*Binl)'; opts3 optimoptions('quadprog','Algorithm', 'active-set','Display','iter'); tive-set','Display','ite
lb=ones(size(Bin1)).*0; ub=[]; x2 = quadprog(H,f,[],[],[],lb,ub,x0,op ts3); v2g11m=v2g11m+v2. Btotallsum=Btotallsum+Btotall; BSTDsum=BSTDsum+abs(BSTD); end, if realdata == 1, x2=x2sum./ncal;

BSTD=BSTDsum./ncal: else BSTD=abs(BSTD); if max(x2)<=max(Btotall),</pre> warning('May yield false result, try change regularization parameters'), end if max(x2)>=le2*max(Btotall),
 warning('yielding false result, regularization parameters have to be nged'), end, %% calculate recovered f(x,0) xfinal2=x2': [~,~,~, Recoverdfx2,~,~]=testABgeneration (x2',testLfcum,DO2bio,uz,t); %% plotting figure('name','test problem, Tikhonov regulerization') hold on if realdata~=1. plot(testLfcum,testX,'black--','line width'.3); plot(testLfcum,Btotall(:,end),'bo'); 0100 errorbar(originaldata(1,:),originald ata(2,:), originaldata(3,:), 'blacks', linewidth',2); plot(testLfcum,xfinal2,'-b','linewid plot(testLfcum,Recoverdfx2,'-r','lin ewidth',3);
if realdata~=1, legend('Real f(x,0)'...
,'measured g(x,T)' ...
,'Calculated f(x,0) with non-negative' ...
,'recoverd g(x,T) with non-negative') else legend('Measured data points, mean & STD TD' ... ,'Calculated f(x,0) with non-negative non-negative'
, 'recoverd g(x,T) with
non-negative'); set(gca,'children',flipud(get(gca,'c hildren'))); end xlim([0 testIfcum(end)]); ylim([0 ceil(max(max(testX),max(xfinal2))*10 >/10]); set(gca,'XTick',0:testLfcum(end)/10: testLfcum(end), ... 'YTick',0:ceil(max(max(testX),max(xf inal2))*10)/10/5:ceil(max(max(testX)
,max(xfinal2))*10)/10) title('Mathematical treatment of the title('Mathematical creatment of c measured data'); xlabel('Depth into biofilm'); ylabel('Photosynthetic activity in mole/m^3/sec 02'); set(gca,'XGrid','on');

Btotall=Btotallsum./ncal;

end

Appendix _______ % is identified and the corresponding

Bin, Bsurf] = testABgeneration (testX,testLfcum,DO2bio,uz,t) &UNTITLED Summary of this function goes her Detailed explanation goes here 8 %% constructe the known kernel, matrix of Ai xcal=testLfcum; acal=(4*pi*DO2bio*t)^0.5; %a=(4*pi*D*t)^0.5 in meter bcal=acal.^2/pi; %b=4*D*t deltay=xcal(2)-xcal(1);
weighing factor \$doltay xcall=repmat(xcal,length(xcal),1); % start-repmat(xcal,tength(xcal),1);
% y or x0 in nLf discretion
xcal2=repmat(xcal',1,length(xcal))-0
.*deltay; % x in nLf discretion xend=xcal2./bcal.^0.5: erfx=erf(xend); Ain=(exp(-(xcal2-xcal1+uz.*t).^2./bc
al))./acal.*deltay; %y or x0 position dependent Ain(xcal2<0)=0: Asurf=(exp(-(0-xcall+uz*t).^2./bcal)
.*(1-erfx))./acal.*deltay; %y or x0
position dependent
heurg(uc)) Asurf(xcal2<0)=0: Atotal=Ain-Asurf: %% constructe the known lefthand side. matrix of B, measured dCdt using Splinefit Bin=sum(repmat(testX,length(testLfcu m),1).*(Ain),2);%y or x0 position Bsurf=sum(repmat(testX,length(testLf cum),1).*(Asurf),2); Btotal=sum(repmat(testX,length(testL fcum),1).*(Atotal),2); ond function [reg_corner,rho,eta,reg_param] = 1 curvenoP(U,sm,b,method,L,V) %L_CURVE Plot the L-curve and find its "corner". % [reg_corner,rho,eta,reg_param] = l curve(U,s,b,method) l_curve(U,sm,b,method) , sm =
[sigma,mu] l curve(U,s,b,method,L,V) Plots the L-shaped curve of eta, the solution norm || x || or % semi-norm || L x ||, as a function of rho, the residual norm
% || A x - b ||, for the following methods:
% method = 'Tikh' : Tikhonov
regularization (solid line) % method = 'tsvd' : truncated SVD or GSVD (o markers) % method = 'dsvd' : damped SVD or (dotted line)
 (dotted line)
 method = 'mtsvd' : modified TSVD GSVD (x markers) % The corresponding reg. parameters are returned in reg param. If no % method is specified then 'Tikh' is default. For other methods use plot lc. Note that 'Tikh', 'tsvd' and 'dsvd' require either U and s (standard-% form regularization computed by the function csvd, or U and sm (general-% form regularization) computed by the function cgsvd, while 'mtvsd' % requires U and s as well as L and V computed by the function csvd. % If any output arguments are specified. then the corner of the L-curve

function [Atotal, Ain, Asurf, Btotal,

reg. parameter reg_corner is % returned. Use routine 1 corner if an upper bound on eta is required. % Reference: P. C. Hansen & D. P. O'Leary, "The use of the L-curve in % the regularization of discrete ill-posed problems", SIAM J. Sci. % Comput. 14 (1993), pp. 1487-1503. % Modified from code by Per Christian Hansen, DTU Compute, October 27, 2010. & Set defaults if (nargin==3), method='Tikh'; end % if (nargin==), method into , end o Tikhonov reg. is default. npoints = 200; % Number of points on the L-curve for Tikh and dsvd. smin_ratio = 16*eps; % Smallest regularization parameter. % Initialization. [m,n] = size(U); [p,ps] = size(sm); if(nargout>0), locate = 1; else locate = 0; end beta = U'*b; beta2 = norm(b)^2 norm(beta)^2; if (ps==1) $s = sm \cdot beta = beta(l \cdot n) \cdot$ else s = sm(p:-1:1,1)./sm(p:-1:1,2); beta = beta(p:-1:1); end xi = beta(1:p)./s; xi(isinf(xi)) = 0; if (strncmp(method,'Tikh',4) |
strncmp(method,'tikh',4)) eta = zeros(npoints,1); rho = eta; req param = eta; s2 = s.^2; reg_param(npoints) =
max([s(p),s(1)*smin ratio]); ratio = (s(1)/reg param(npoints))^(1/(npoint s-1)); i=npoints-1:-1:1, reg_param(i) = ratio*reg_param(i+1); end for i=1:npoints f = s2./(s2 + reg_param(i)^2); eta(i) = norm(f.*xi); rho(i) = norm((1-f).*beta(1:p)); end end if (m > n & beta2 > 0), rho = sqrt(rho.^2 + beta2); end marker = '-'; txt = 'Tikh.'; elseif (strncmp(method, 'tsvd',4) strncmp(method, 'tgsv',4)) eta = zeros(p.1): rho = eta: eta = zeros(p,1); rho = eta; eta(1) = abs(xi(1))^2; for k=2:p, eta(k) = eta(k-1) + abs(xi(k))^2; end eta = sqrt(eta); if (m > n) (m > n) if (beta2 > 0), rho(p) = beta2; else rho(p) = eps^2; end rho(p) = eps^2; end for k=p-1:-1:1, rho(k) = rho(k+1) + abs(beta(k+1))^2; end rho = sqrt(rho); reg_param = (1:p)'; marker = 'o'; f (ps==1) U = U(:,] U(:,1:p); txt = 'TSVD'; else U = U(:,1:p); txt = 'TGSVD'; elseif (strncmp(method, 'dsvd', 4) | strncmp(method, 'dasy', 4)) eta = zeros(npoints,1); rho = eta; req param = eta; reg_param(npoints) = max([s(p),s(1)*smin ratio]); ratio (s(1)/reg_param(npoints))^(1/(npoint s-1)); r i=npoints-1:-1:1, reg_param(i) = ratio*reg_param(i+1); end for i=1:npoints f = s./(s + reg_param(i));

end if (m > n & beta2 > 0), rho = sqrt(rho.^2 + beta2); end
marker = ':'; if (ps==1), txt = 'DSVD'; else txt = DGSVI end elseif (strncmp(method, 'mtsv',4)) if (nargin~=6) error('The matrices L and V must also be specified') ond [p,n] = size(L); rho = zeros(p,1); eta rho: [Q,R] = qr(L*V(:,n:-1:n-p),0); [Q,R] = qr(L*V(:,n:-1:n-p), for i=1:p k = n-p+i; Lxk = L*V(:,1:k)*xi(1:k); zk = R(1:n-k,1:n-k)\(Q(:,1:n-k)'*Lxk); zk zk(n-k:-1:1); eta(i) = norm(Q(:,n-k+1:p)'*Lxk); if (i < p) rho(i) = norm(beta(k+1:n) + s(k+1:n).*zk); else rho(i) = eps; end end if (m > n & beta2 > 0), rho =
sqrt(rho.^2 + beta2); end reg_param = (n-p+1:n)'; txt = 'MTSVD'; U = U(:,reg_param); sm = sm(req param); marker = 'x'; ps = 2; % General form
reqularization. 0100 error('Illegal method') end % Locate the "corner" of the L-curve. if required. if (locate) [reg_corner,rho_c,eta_c] = corner(rho,eta,reg param,U,sm,b,me thod); end

eta(i) = norm(f.*xi); rho(i) = norm((1-f).*beta(1:p));

Appendix II: MATLAB code of the model presented in this study

The MATLAB code presented here was written as described and applied in the 3rd manuscript presented in this

thesis.

function [Tcomp,Datacomp,BM]=TL Model test(Iz inPer.Irrsurf,pC02,BMt0)
% input, Calculate irradiance from
function "RTE_TL" Irradiance in microE, pCO2 in Pa, initial biomass in kg % the whole spatial domain was discreted into n=n0+1 layers, 1 to n0 th % layer are the inside layers, and have % layer are the inside layers, and have initial biomass at t0, the nth % layer is the surface layer, which is assumed to have no biomass at t0 or % directly after a new discretion, all 1 to n layer contribute to growth, 8 but all the biomass increments are only inside the nth layer %% general discretion setups c=10; %number of variables calculated using ODEs implicitly tend=12; %time of simulation in days LDratio=14/10; %light dark cycle ratio %% Paramotors XF=220: %biomass concentration in BioPoro=0.76: %biofilm porosity. Li et SubPoro=0.35; %substrate layer porosity DCO2b,DHCO3b,DCO3b,DNO3b,DHPO4b,DH2P O4b,DO2b,DNab,DCO2s,DHCO3s ... ,DCO3s,DNO3s,DHPO4s,DH2PO4s,DO2s,DNa % according to porosity and diffusion coefficient in water in m^2/d^-1 DCO2b=DCO2b*1; SNO3medium=2.94e-3; %NO3 in bulk liquid SH2PO4medium=1.29e-3; %HPO4 in bulk liguid in M SHP04medium=0.431e-3: %H2P04 in medium Sionsmedium=SNO3medium+SH2PO4medium+ 2*SHPO4medium; %cations in bulk liquid, calculated as Na in M KsCO2=4e-4; %assumed half saturation constant for CO2 as carbon source in M, 4e-6 M from literature, KsHCO3=4e-5; %assumed half saturation constant for HCO3 as carbon source in M, 4e-6, literature states the value with pH KsNO3=1e-5; %assumed half saturation constant for N in M, 5e-4 KsPO4=1.7e-5; %assumed half saturation nstant for P in M. 1.7e-5 KiCO2toHCO3=5e-4; %assumned inhibition constant for CO2 on growth using HCO3, le-6 PHOBIA KiO2max=le-3; %assumned maxim inhibition constant for O2 in M, represent photorespiration, high = less photorespiration KsO2=1e-4; %assumed half saturation constant for O2 on dark respiration Lsub=1e-5; %membrane thickness in meter

uz=0.23; %convection velocity in m/day 0 23 RHCO3transCO2=1: %efficience of HCO3 assimilation compare to CO2 assimilation YxsCO2=30; %bimass yield on CO2 consumed in kg biomass/kmol C, from digestion meas romont YxsHCO3=YxsCO2.*RHCO3transCO2; %bimass vield on HCO3 consumed in kg_biomass/kmol C, from digestion asurement. RHCO3DvsC=1: %ratio of direct hco3 uptake vs extracellular conversion to YxsO2= YxsCO2/1.3;%biomass yield of O2 by respiration, assume biomass C:O ratio is 1:1.5 mole PCO2O2=1; % production ratio of O2 by Co2 assimilation mole to mole PHCO302=1.*RHCO3transCO2; % production ratio of 02 by HCO3 sassimilation, mole to mole YxsN=200; %bimass yield on DN consumed in kg biomass/kmol N in kg_Diomass/kmol N YksP=3100; %bimass yield on DP consumed in kg_biomass/kmol P KalC03=10^(-6.35); %equilibrium constant for C02+H20 to H+HC03 Ka2CO3=10^(-10.33); %equilibrium Ka2C03=10"(-10.33); sequilibrit constant for HCO3 to H+CO3 KaCaOH=10^(7.64); sequilibrium constant for CO2+OH to HCO3 HenCO2=3.04*10^(-7);%Henry's constant for CO2 at 25 degree in M/Pa. For co2 at 25 degree in M/Pa. Hen02=1.3*10^(-8);%Henry's constant for 02 at 25 degree, in M/Pa. Kw=10^(-14); %equilibrium constant of KalPO4=10^(-2.15); %equilibrium constant for H3P04 to H+H2P04 Ka2P04=10^(-7.2); %equilibrium constant for H2PO4 to H+HPO4 Ka3PO4=10^(-12.1); %equilibrium Kd3P04=10"(-122.1); %equilibrium constant for HPO4 to H+PO4 kC02aH2O=10^7.54; %rate constant of C02+H2O to H2C03 in d^-1, Gibbson and Edsall, 1963, 10^7.54 or from book below, kCO2aOH=10^8.86: %rate constant of CO2+OH to HCO3 in d^-1 kHCO3tCO3=10^4.6; %rate constant of HCO3 to H+CO3 in d^-1, rate constants from product and process modelling, a case study approach, 2011 kH2PO4tHPO4=10^9.74; %rate constant of H2PO4 to HPO4 in d^-1 pO2=21000; %partial pressure of O2 in Pa, 21000 in atm SO2medium=pO2*HenO2*0.01; %medium O2 concentration pO2*HenO2 or 0 SCO2medium=pCO2*HenCO2: %medium CO2 concentration pCO2*HenCO2; smella concentration pCO2*HenCO2 or 0 rs=0.05; %surface reflectance umax=8; % maximal growth, d^-1 BMc=0.6; % umax=1.5e-4; % maximal growth in biomass (here as thickness) in m s^-1 Ures=0.134; %respiration rate in day^-1 Lft0=BMt0/XF; %initial biofilm n0=fix(Lft0/10e-6): %set n0 for initial dicretion determination n=n0+1; %set spatial grid, n=n0+1 indicate the nth layer pHmedium=6.8, %pH in medium %% preset surface pH and bottom pH, from surement if needed pHSurfaceM=6.8; pHBottomM=6.8;

HSurfaceM-pHmedium,n); % a correction factor for initial pH guess span tL=LDratio ./ (1 + LDratio); %illuminated time in day %% Initial conditions c*n rows, last cth elements indicate values in nth layer ICS=repmat([10^(-pHmedium), SCO2medium, KalCO3/10^(-pHmedium)*SCO2medium, KalCO3*Ka2CO3/(10^(-pHmedium))^2*SCO 2medium,... SNO3medium, SHPO4medium, SH2PO4medium, Sionsmedium, pO2*HenO2, Lft0/n01.1.n); H (guess for initial), HCO3, • c02, CO3 >DN. DHPO4. DH2PO4. DO2. Sions. sLf. % nth layer biomass has to be se to 0 att0, to avoid singlarity, set as a very small value instead of 0 ICS(c*n)=ICS(c*n)/10^10; ICS(1:c:c*n)=ICS(1:c:c*n).*10.^(-CO2 cor); % TCS(c*(n-1)+1)=pC02*HenC02: . ICS(c*(n-1)+2)=KalCO3/10^(-pHmedium) *pCO2*HenCO2 -ICS(c*(n-1)+3)=KalCO3*Ka2CO3/(10^(-p Hmedium))^2*pCO2*HenCO2; ee timo ener tspan=[0 tend]; %% call ode solver tcal=0; %set starting point for dtermine is tend is reached Datacomp=[]; %set matrix for compiling pH data, each column is one spatial %discretion unit (total n columns), each row is one time step (solver defined) %pages represent variables,1 to 9 ---> H, CO2, HCO3, CO3, Dn. %DP, Sions, DO2,Lf (c pages). Tcomp=[]; %set matrix for compiling time data EventGC=[]; %set matrix for compiling event tim e data ncal=1: %the number of discretions used during 0 to tend ReG=0; %growth higher than respiration while tcal<tend, %initial adapting shing of the finite elements meshing of the finite elem
while ncal<120&&tcal<tend,</pre> %set absolute tolenrance for each component ATol=repmat([le-14, pCO2*HenCO2/le2, KalCO3/l0^(-pHmedium)*pCO2*HenCO2/le (KalCO3*Ka2CO3/(10^(-pHmedium))^2*pC 02*HenCO2)/1e2,... % SNO3medium/1e2, % SNOSMEdium/162, SPO4medium/162, (SNO3medium+(129/172.1)*SPO4medium+2 *(43.1/172.1)*SPO4medium)/le2, pO2*HenO2/le2, Lft0/n0/le2],1,n);

CO2cor=linspace(pHBottomM-pHmedium,p

Appendix

ATo1=10e-14; NonNeg=1:c*n:

options=odeset('Vectorized', 'on', 'Ab
sTol',ATol, 'RelTol', le-6, 'Events',...

@EventsLf, 'NonNegative', NonNeg); %set solver option

[T, calS, GridCT, ~, ~]=odel5s(@ODE_dS,t span, ICS, options);

%after dLf reached set value(dLf=Lf(t=0)/n), reset initial
conditions %first accumulate solution

len=size(Datacomp,1); %calculate size length of the previous data compling matrix lenT=size(T.1):

%compile data

Datacal=permute(reshape(calS',[cn lenT]),3:-1:1);

Datacomp((len+1):(len+lenT),:,:)=Dat acal(:,:,:): % compiling data in 3D matriv

%compiling time data if tcal==0,

Tcomp((len+1):(len+size(T,1)))=T(:);

Tcomp((len+1):(len+size(T,1)))=T(:)+ Tcomp(len); end

acompiling event data EventGC(ncal)=GridCT;

\$then test if tend is reached tcal=T(length(T));

%update storage space for solution Datacomp=[Datacomp zeros(size(Datacomp,1),1,c)];

calS(size(calS,1),c*n)<=Lft0/10^10, %dark, respiartion > growth, thickness
decrease

%set new initial conditions, consider as new layer added on top and hae

%the same concentration of everything as the layer below, except %biomass, which is initially 0 ICS=ICS(1:c*(n-1)); new grid

n=n+1:

warning('growth stoped'); break,

else

%set new initial conditions, consider as new layer added on top and

%the same concentration of everything as the layer below, except %biomass, which is initially

ICS=[calS(size(calS,1),:),calS(size(calS,1),(n-1)*c+1:n*c)];

%set new grid
n=n+1;

%reset nth layer thickness to 0 ICS(c*n)=ICS(c*n)/10^10; end.

%set new time span and start new iteration tspan=[0 (tend-tcal)];

ncal=ncal+1; end,

%% ploting plotting End;

function plotting End

for flipi=1:c; % flip the x-axis for better view

Datacomp(:.:.flipi)=fliplr(Datacomp(:,:,flipi)); end.

figure('Name',['TL-model with
',num2str(Irrsurf),' PAR irradiance

.num2str(round((pCO2-39)/1e5*100)). additional CO ٠ı.

subplot(3,3,2),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,2).*le3,'o-blue hold on:

grid on

subplot(3,3,2),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,3).*le3,'s-blue

subplot(3,3,2),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*1e6,Dataco mp(length(Tcomp),2:n,4).*1e3,'*-blue

subplot(3,3,2),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,(Datac omp(length(Tcomp),2:n,2)...

+Datacomp(length(Tcomp),2:n,3)+Datac omp(length(Tcomp),2:n,4)).*le3,'-bla ck');

title('DIC in mM vs. depth');

subplot(3,3,4),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,5).*le3,'o-blue);

hold on; arid on

subplot(3,3,4),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,6).*le3,'s-red'

subplot(3,3,4),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,7).*le3,'o-red'

subplot(3,3,4),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,(Datac omp(length(Tcomp),2:n,6)+...

Datacomp(length(Tcomp),2:n,7)).*le3, red'): %total

subplot(3,3,4),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*1e6,Dataco mp(length(Tcomp),2:n,8).*1e3,'*-gree n');

title('NO3, HPO4, H2PO4, total
P, Cations(Na) in mM vs depth');

subplot(3,3,3),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*le6,Dataco mp(length(Tcomp),2:n,9).*le3,'o-blac i); hold on;

arid on: title('02 in mM vs depth');

pHcal=-log10(Datacomp(:,:,1));

subplot(3,3,5),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10)).*1e6,pHcal(length(Tcomp),2:n),'o-black'); hold on; grid on;

title('pH vs depth');

%Plot Irr(explicit)

nt=length(Tcomp); Lfaccleft=zeros(nt,n);

Lfaccleft(:,1:n)=cumsum(Datacomp(:,: .10).2);

Irrcalleft(1:nt,1:n)=(Irrsurf.*(1-rs

Lfaccright=zeros(size(Lfaccleft)):

Lfaccright(:,1:size(Lfaccright,2)-1)
=Lfaccleft(:,2:size(Lfaccleft,2));

Irrcalright(1:nt,1:n)=(Irrsurf.*(1-r s)).*polyval(IzinPer,Lfaccright(1:nt ,1:n));

Irrcal=(Irrcalleft+Irrcalright),/2;

subplot(3,3,1),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10).*1e6),Irrcal (length(Tcomp),l:n-1), 'o-black');

hold on; grid on;

title('Irradiance in microE vs depth'):

%plot growth rate(explicit)

GCO2=zeros(nt,n-1);GHCO3=zeros(nt,n-1):Rd=zeros(nt.n-1);

GCO2(:,:)=umax.*PCO2O2.

.*Datacomp(:,2:n,9)./(Datacomp(:,2:n ,9)+Datacomp(:,2:n,9).^2./(KiO2cal(D atacomp(:,2:n,2),Datacomp(:,2:n,9)))

.*GroIrr TL(Irrcal(:,1:n-1))... light

.*min(Datacomp(:,2:n,2)./(KsCO2+Data comp(:,2:n,2)),...CO2

min(Datacomp(:,2:n,5)./(KsNO3+Dataco mp(:,2:n,5)),...NO3

(Datacomp(:,2:n,6)+Datacomp(:,2:n,7))./(KsP04+(Datacomp(:,2:n,6)+Datacom p(:,2:n,7)))); %P04

GHCO3(:,:)=umax.*PHCO3O2.*RHCO3DvsC.

.*Datacomp(:,2:n,9)./(Datacomp(:,2:n ,9)+Datacomp(:,2:n,9).^2./(KiO2cal(D atacomp(:,2:n,2),Datacomp(:,2:n,9)))) 02

.*GroIrr TL(Irrcal(:.1:n-1))...light

*min(Datacomp(:,2:n,3)./(KsHCO3+Dat acomp(:,2:n,3)),...HCO3

min(Datacomp(:,2:n,2)./((Datacomp(: 2:n,2)+Datacomp(:,2:n,2).^2./KiCO2to HCO3)),... CO2 inhibition on HCO3

min(Datacomp(:,2:n,5)./(KsNO3+Dataco mp(:,2:n,5)),... NO

(Datacomp(:,2:n,6)+Datacomp(:,2:n,7))./(KsPO4+(Datacomp(:,2:n,6)+Datacom p(:,2:n,7))))); %PO4

Rd(:,:)=-Ures.*Datacomp(:,2:n,9)./(K sO2+Datacomp(:,2:n,9)); %Respiration GrowthR(:,:)=GCO2+GHCO3+Rd;

subplot(3,3,6),plot(cumsum(Datacomp) length(Tcomp),1:n-1,10).*1e6),Growth
R(length(Tcomp),1:n-1),'o-black'); hold on; grid on;

subplot(3,3,6),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10).*1e6),GCO2(1
ength(Tcomp),1:n-1),'o-green');

subplot(3,3,6),plot(cumsum(Datacomp(length(Tcomp),1:n-1,10).*le6),GHCO3(length(Tcomp),1:n-1),'s-green');

subplot(3,3,6),plot(cumsum(Datacomp(length(Tcomp),l:n-1,10).*le6),Rd(len gth(Tcomp),l:n-1),'*-red'); title('Growth in dr-^1 vs depth');

Appendix

%Plot Biomass

Lfcal(1:nt)=sum(Datacomp(:,:,10),2); BM=Lfcal.*XF; subplot(3,3,[7 8 9]),plot(Tcomp,BM.*1e3);

hold on; arid on:

title('Biomass in g/m^2 vs. time');

ventGCcum=cumsum(EventGC); if length(EventGCcum)-=1 &&

idyBM= find(Tcomp>EventGCcum(j),1); %find

and mark the first time step after rediscretion subplot(3,3,[7 8

9]),plot(EventGCcum(j),BM(idxBM).*le 3,'or'); end.

end, if tcal~=tend&&ReG==1,

plot(Tcomp(length(Tcomp)), BM(length(Tcomp().*le3,'sr'); 0100

plot(Tcomp(length(Tcomp)), BM(length(Tcomp)).*1e3,'or'); end,

end % plot end condition

%% function for ODE systems

function dS = ODE dS(t, S)

%Function handle for diffusion-convection-reaction PDE %ODEs from MOL with upwind scheme,

original PDE is: %dDS./dt=Dbio.*d(dDS)./dz.^2 +

uz.*dDS./dz -Umax.*DS(i)./(KsDS+DS(i))./YxsS.*Xf for

%DS, and

dLf(i)./dt=Lf(i).*dzd.*Umax.*DS(i)./

(KsDS+DS(i)); %upwind scheme: d(dDS)./dz.^2= DSbio./.*(DS(i-1)-2.*DS(i)+DS(i+1)). /dz.^2, and

%dDS./dt=uz./Lf.*(DS(i)-DS(i-1))./dz St ODEs: dS=zeros(c*n,size(S,2));

%% ODEs describing pH (column i) dDCO2./dt (column i+1),ODEs describing dDHCO3./dt (column i+2), ODEs describing dDCO3./dt (column i+3)

---bottom s ____ boundary: 1 %where

Dbio.*(D(i+c)-D(i-c))./(2.*dz)=Dsub. /Lsub.*(SDm-D(i))

- $i = 1 \cdot$
- \$C02

dS(i+1,:)=DCO2bcal(S,i)./(Lft0./n0). ^2.*((S(i+1+c,:)+2.*(Lft0./n0).*DCO2 s./(Lsub.*DCO2bcal(S,i)).*(SCO2mediu m-S(i+1,:)))-2.*S(i+1,:)+S(i+1+c,:))diffusio

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DCO 2s./(Lsub.*DCO2bcal(S,i)).*(SCO2medi um-S(i+1,:)))...%convection

-kCO2aH2O.*(S(i+1,:)-(S(i+2,:).*S(i, /KalCO3))-kCO2aOH.*(S(i+1,:).*Kw. /S(i,:)-S(i+2,:)./KaCaOH) ... %reaction to HCO3

-UcalCO2(S,i)./YxsCO2.*XF ... consumption

(1-RHCO3DvsC).*UcalHCO3(S.i)./YxsHC 03.*XF ... from anhydrase +RdO2cal(S,i)./YxsO2.*XF; %prodction

SHCO3

dS(i+2,:)=DHCO3b./(Lft0./n0).^2.*((S (i+2+c,:)+2.*(Lft0./n0).*DHCO3s./(Ls ub.*DHCO3b).*(SCO2medium.*KalCO3./(1 0.^(-pHmedium))-S(i+2,:)))-2.*S(i+2,

0.^(-pHmedium))-S(1+2,:)))-2.*S(1+2, :)+S(1+2+c;))...diffusion +uz./(Lft0./n0) .*(2.*(Lft0./n0).*DHCO3S./(Lsub.*DHC O3b).*(SOOZmedium.*KalCO3./(10.^(-pH medium))-S(1+2,:))...%convection

+kCO2aH2O.*(S(i+1,:)-(S(i+2,:).*S(i, :)./KalCO3))... re CO2+820

+kCO2aOH.*(S(i+1.:).*Kw./S(i.:)-S(i+ 2,:)./KaCaOH)... reaction from CO2+OH

-kHCO3tCO3.*(S(i+2,:)-S(i+3,:).*S(i, :)./Ka2CO3)... reaction to CO3

-UcalHCO3(S,i)./YxsHCO3.*XF; \$consumption

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dS(i+3,:)=DCO3b./(Lft0./n0).^2.*((S(i+3+c,:)+2.*(Lft0./n0).*DCO3s./(Lsub .*DCO3b).*(SCO2medium.*KalCO3.*Ka2CO 3./(10.^(-pHmedium).^2-S(i+3,:))-2 .*S(i+3,:)+S(i+3+c,:))... diffusion

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DCO 3s./(Lsub.*DCO3b).*(SCO2medium.*KalC O3.*Ka2CO3./(10.^(-pHmedium)).^2-S(i +3.:))) ...%convection

+kHCO3tCO3.*(S(i+2,:)-S(i+3,:).*S(i, :)./Ka2CO3); %reaction from HCO3

\$DN

dS(i+4,:)=DNO3b./(Lft0./n0).^2.*((S(i+c+4,:)+2.*(Lft0./n0).*DNO3s./(Lsub .*DNO3b).*(SNO3medium-S(i+4,:)))-2.* S(i+4,:)+S(i+c+4,:)) ... diffusion

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DNO 3s./(Lsub.*DNO3b).*(SNO3medium-S(i+4 (:))) ...convection

-(UcalCO2(S,i)./YxsN+UcalHCO3(S,i)./ YxsN).*XF...consumption +RdO2cal(S,i)./YxsN.*XF; %prodction

dS(i+5.:)=DHPO4b./(Lf+0./n0).^2 .*((S(i+c+5,:)+2.*(Lft0./n0).*DHPO4s ./(Lsub.*DHPO4b).*(SHPO4medium-S(i+5 (i)))-2.*S(i+5,:)+S(i+c+5,:))...
diffusion

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DHP O4s./(Lsub.*DHPO4b).*(SHPO4medium-S(i+5,:))) ...convection

+kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i,:)./Ka2PO4)... reaction from H2PO4

-(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P).*XF...consumption

+(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction

%DH2PO4

dS(i+6,:)=DH2PO4b./(Lft0./n0).^2 .*((S(i+c+6,:)+2.*(Lft0./n0).*DH2PO4 s./(Lsub.*DH2PO4b).*(SH2PO4medium-S(i+6,:)))-2.*S(i+6,:)+S(i+c+6,:))... diffusion

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DH2 PO4s./(Lsub.*DH2PO4b).*(SH2PO4medium -S(i+6,:))) ...convection

-kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i,:)./Ka2PO4)... reaction to HB

-(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P).*XF...consumption

+(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction

*Deione

dS(i+7,:)=DNab./(Lft0./n0).^2.*((S(i +c+7,:)+2.*(Lft0./n0).*DNas./(Lsub.* DNab).*(Sionsmedium-S(i+7,:)))-2.*S(i+7,:)+S(i+c+7,:)) ... diffusion

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DNa s./(Lsub.*DNab).*(Sionsmedium-S(i+7,

dS(i+8.:)=D02b./(Lft0./n0).^2.*((S(i uls(i+0;.)=b02b./(lit0./n0). 2.*((S(i
+8+c,:)+2.*(Lft0./n0).*D02s./(Lsub.*
D02b).*(S02medium=S(i+8,:)))=2.*S(i+ 8,:)+S(i+c+8,:)) ... diff

+uz./(Lft0./n0).*(2.*(Lft0./n0).*DO2 s./(Lsub.*DO2b).*(SO2medium-S(i+8,:)))...%convection

+(UcalCO2(S,i)./YxsO2+UcalHCO3(S,i). /Yxs02.*RHC03DvsC).*XF ...productic -Rd02cal(S,i)./Yxs02.*XF;

&consumption

8-----boundary 2:n-1

i=(c+1):c:((n-1)*c):

8C02

dS(i+1,:)=DCO2bcal(S,i)./(Lft0./n0).

*(S(i+1-c,:)-2.*S(i+1,:)+S(i+1+c,:)) diffusion

+uz./(Lft0./n0).*(S(i+1-c,:)-S(i+1+c

.:))...convection

-kCO2aH2O.*(S(i+1,:)-(S(i+2,:).*S(i, :)./Ka1CO3))-kCO2aOH.*(S(i+1,:).*Kw. /S(i,:)-S(i+2,:)./KaCaOH) ... %reaction to HCO3

-UcalCO2(S,i)./YxsCO2.*XF

....consumption

+(1-RHCO3DvsC).*UcalHCO3(S,i)./YxsHC 03.*XF ... from anhydrase +RdO2cal(S,i)./YxsO2.*XF;

%prodction

%HCO3

8CO3

.:))... convection

dS(i+2,:)=DHCO3b./(Lft0./n0).^2 .*(S(i+2-c,:)-2.*S(i+2,:)+S(i+2+c,:)

)...diffusion

+uz./(Lft0./n0).*(S(i+2-c,:)-S(i+2+c .:))... convection

+kCO2aH2O.*(S(i+1,:)-(S(i+2,:).*S(i,

-kHCO3tCO3.*(S(i+2,:)-S(i+3,:).*S(i,

dS(i+3,:)=DCO3b./(Lft0./n0).^2.*(S(i

+uz./(Lft0./n0).*(S(i+3-c,:)-S(i+3+c

+3-c,:)-2.*S(i+3,:)+S(i+3+c,:)) ...diffusion

:)./Ka2CO3)... reaction to CO

-UcalHCO3(S,i)./YxsHCO3.*XF;

:)./KalCO3))... reaction from CO2+H2O +kCO2aOH.*(S(i+1,:).*Kw./S(i,:)-S(i+ 2,:)./KaCaOH)... reaction from CO2+OH

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+kHCO3tCO3.*(S(i+2,:)-S(i+3,:).*S(i, :)./Ka2CO3); %reaction from HCO3

dS(i+4,:)=DNO3b./(Lft0./n0).^2.*(S(i +4-c,:)-2.*S(i+4,:)+S(i+4+c,:)) ... diffusion

+uz./(Lft0./n0) .*(S(i+4-c,:)-S(i+4+c,:))...convecti on

-(UcalCO2(S,i)./YxsN+UcalHCO3(S,i)./ YxsN).*XF...consumption +RdO2cal(S,i)./YxsN.*XF;

%DHPO4

dS(i+5,:)=DHPO4b./(Lft0./n0).^2.*(S(i+5-c,:)-2.*S(i+5,:)+S(i+5+c.:))

+kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i,:)./Ka2PO4)... reaction from H2PO4

-(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P).*XF...consumption

+(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction

dS(i+6.:)=DH2PO4b./(Lf+0./n0).^2.*(S (i+6-c,:)-2.*S(i+6,:)+S(i+6+c,:)) ...diffusion +uz./(Lft0./n0) .*(S(i+6-c,:)-S(i+6-c,:))...

nvection

-kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i,:)./Ka2PO4)... reaction to HPO4

-(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P).*XF...consumption

+(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction

%Dsions

dS(i+7,:)=DNab./(Lft0./n0).^2.*(S(i+

+uz./(Lft0./n0).*(S(i+7-c,:)-S(i+7+c .:)): %convection

\$D02

dS(i+8,:)=DO2b./(Lft0./n0).^2.*(S(i+ 8-c,:)-2.*S(i+8,:)+S(i+8+c,:)) ... diffusion

+uz./(Lft0./n0).*(S(i+8-c,:)-S(i+8+c (:)) ... convection in

+(UcalCO2(S i) /VxsO2+UcalHCO3(S i) /Yxs02.*RHC03DvsC).*XF ...productio -Rd02cal(S,i)./Yxs02.*XF; %consumption

۰. -----top boundary n

i=(c*n-(c-1));

\$C02

dS(i+1,:)=0;

%HCO3

dS(i+2,:)=DHCO3b./(Lft0./n0).^2 .*(S(i+2,:)-2.*S(i+2,:)+S(i-c+2,:)). .*(S(1+2,., ..diffusion +uz./(Lft0./n0) 2''+2.:))... .*(S(i+2-c,:)-S(i+2,:))..

%convection in

+kCO2aH2O.*(S(i+1,:)-(S(i+2,:).*S(i, :)./KalCO3))... r tion from CO2+H2O

+kCO2aOH.*(S(i+1,:).*Kw./S(i,:)-S(i+ 2,:)./KaCaOH)... reaction fi

-kHCO3tCO3.*(S(i+2.:)-S(i+3.:).*S(i. :)./Ka2CO3)... reaction to CO3

-UcalHCO3(S,i)./YxsHCO3.*XF; \$consumption

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dS(i+3,:)=DCO3b./(Lft0./n0).^2.*(S(i +3,:)-2.*S(i+3,:)+S(i-c+3,:)) ...diffusion

+uz./(Lft0./n0).*(S(i+3-c,:)-S(i+3,:)) ...%convection in

+kHCO3tCO3.*(S(i+2,:)-S(i+3,:).*S(i, :)./Ka2CO3); %reaction from HCO3

ds(i+4 ·)=DNO3b /(Lft0 /n0) ^2 *(S(i +4,:)-2.*S(i+4,:)+S(i-c+4,:)) ... diffusion

+uz./(Lft0./n0) .*(S(i+4-c,:)-S(i+4,:))...convection

-(UcalCO2(S,i),/YxsN+UcalHCO3(S,i),/ YxsN).*XF... consumption +RdO2cal(S,i)./YxsN.*XF;

%prodction

9DUD04

dS(i+5,:)=DHPO4b./(Lft0./n0).^2 .*(S(i+5,:)-2.*S(i+5,:)+S(i-c+5,:)). ..diffusion

+uz./(Lft0./n0).*(S(i+5-c,:)-S(i+5,:)) ... convection in

+kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i.:)./Ka2PO4)... reaction from H2PO4

-(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P) *XF ... consumption

+(S(i+5,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction \$DH2PO4

dS(i+6,:)=DH2PO4b./(Lft0./n0).^2 .*(S(i+6,:)-2.*S(i+6,:)+S(i-c+6,:)).

+uz./(Lft0./n0).*(S(i+6-c.:)-S(i+6.:)) ... convection in

-kH2PO4tHPO4.*(S(i+6,:)-S(i+5,:).*S(i,:)./Ka2PO4)... reaction to HF

-(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*(U calCO2(S,i)./YxsP+UcalHCO3(S,i)./Yxs P).*XF ... consumption

+(S(i+6,:)./(S(i+5,:)+S(i+6,:))).*Rd O2cal(S,i)./YxsP.*XF; %prodction

*Deione

dS(i+7.:)=DNab./(Lft0./n0).^2.*(S(i+ 7,:)-2.*S(i+7,:)+S(i-c+7,:)) ...diffusion (out)

+uz./(Lft0./n0).*(S(i+7-c,:)-S(i+7,:)); %convection in

%DO2

dS(i+8,:)=0;

8_____ - special case for pH, with have no boundary condition, and is discribed as a algebraic equations

%dH as state variables, CO3 and HCO3 calculated in their own ODEs, so here their concentrations are not dependent on pH

i=1:c:c*n:

dS(i,:)=(Chcal(S).*(-dS(i+7,:)+dS(i+ 4 ·) Cations NO3

+((S(i.:)).^2.*KalPO4./((S(i,:)).^3+ (S(i,:)).^2.*KalPO4+S(i,:).*KalPO4.* Ka2PO4+KalPO4.*Ka2PO4.*Ka3PO4).*(dS(i+5,:)+dS(i+6,:)))... H2PO4

+2.*(S(i,:).*Ka1PO4.*Ka2PO4./((S(i,:)).^3+(S(i,:)).^2.*KalPO4+S(i,:).*Ka 1PO4.*Ka2PO4+KalPO4.*Ka2PO4.*Ka3PO4) .*(dS(i+5.:)+dS(i+6.:)))... HPO4

+3.*(KalPO4.*Ka2PO4.*Ka3PO4./((S(i.)).^3+(S(i,:)).^2.*KalPO4+S(i,:).*Ka 1PO4.*Ka2PO4+KalPO4.*Ka2PO4.*Ka3PO4) .*(dS(i+5,:)+dS(i+6,:)))... PO4

+(((S(i,:)).*KalCO3)./((S(i,:)).^2+(S(i,:)).*KalCO3+KalCO3.*Ka2CO3).*(dS (i+1,:)+dS(i+2,:)+dS(i+3,:))) ... HCO3

+2.*((Ka1CO3.*Ka2CO3)./((S(i,:)).^2+ (S(i,:)).*Ka1CO3+Ka1CO3.*Ka2CO3).*(d S(i+1,:)+dS(i+2,:)+dS(i+3,:)))) ...

./(1-(Chcal(S).*(-0+0 ...Cations, NO:

+(1./(Ka2PO4.*Ka3PO4).*(-(S(i,:)).^4 ./(KalPO4.*Ka2PO4.*Ka3PO4)+(S(i,:)). ^2./Ka3PO4+2.*(S(i,:)))./((S(i,:)).^ 3./(KalPO4.*Ka2PO4.*Ka3PO4)+(S(i,:)) 04+1).^2).*(S(i+5,:)+S(i+6,:)) ...

+2.*(1./Ka3PO4.*(-2.*(S(i.:)).^3./(K +2.*(1./Ka3P04.*(-2.*(S(1;:)).^3./(K a1P04.*Ka2P04.*Ka3P04)-(S(i;:)).^2./ (Ka2P04.*Ka3P04)+1)./((S(i;:)).^3./(Ka1P04.*Ka2P04.*Ka3P04)+(S(i;:)).^Xa3P04+1 /(Ka2P04.*Ka3P04)+(S(i;:))./Ka3P04+1).^2).*(S(1+5,:)+S(1+6,:)) ... HPO4

+3.*((3.*(S(i,:)).^2./(KalPO4.*Ka2PO 4.*Ka3PO4)+2.*(S(i,:))./(Ka2PO4.*Ka3 PO4)+1./Ka3PO4)./((S(i,:)).^3./(KalP O4.*Ka2PO4.*Ka3PO4)+(S(i,:)).^2./(Ka 2P04.*Ka3P04)+(S(i,:))./Ka3P04+1).^2).*(S(1+5,:)+S(1+6,:)) ... PO4

+(1/Ka2CO3.*(-(S(i,:)).^2./(Ka1CO3.* Ka2CO3)+1)./((S(i,:)).^2./(Ka1CO3.*K a2CO3)+(S(i,:))./Ka2CO3+1).^2.*(S(i+ 1.:)+S(i+2.:)+S(i+3.:))) ... HCO3

+2.*(-1.*(2.*(S(i,:))./(KalCO3.*Ka2C O3)+1./Ka2CO3)./((S(i,:)).^2./(KalCO 3.*Ka2CO3)+(S(i,:))./Ka2CO3+1).^2.*(S(i+1,:)+S(i+2,:)+S(i+3,:))))); %CO3

cases for biofilm thickness. Irr in this case calculated explicitly,

Irr(i,:)=(((Irrsurf.*rs).*exp(-Mex.*
Xf.*sum(Lf(i:n))))
i=1:c:(c*n-(c-1)); %for 1st to

n0=(n-1)th layer, no thickness increase. dS(i+9,:)=0;

i=(c*n-(c-1)); %for nth layer (top most), the cummulative thickness increase from 1st to nth layers

dS(i+9,:)=(sum(S(c:c:c*n,:).*(UcalCO 2(S,1:c:c*n).*BMc))... CO2 growth

+sum(S(c:c:c*n,:).*(UcalHCO3(S,l:c:c *n).*BMc.*RHCO3DvsC))... HCO3 growth

-sum(S(c:c:c*n,:).*RdO2cal(S,l:c:c*n)./BMc));%decrease from respiration

%% Ch cal, determine which function should be used for the

Appendix

calculation direvitive, whether ph>or<7

function B=Chcal(Scal) Ch= -Scal(i+6,:)+Scal(i+4,:)...Cations.

+(Scal(i,:)).^2.*KalPO4./((Scal(i,:)).^3+(Scal(i,:)).^2.*KalPO4+Scal(i,:)).*KalPO4.*Ka2PO4+KalPO4.*Ka2PO4.*Ka 3P04).*(Scal(i+5,:)+Scal(i+6,:))... H2PO4

+2.*(Scal(i,:).*KalPO4.*Ka2PO4./((Sc al(i,:)).^3+(Scal(i,:)).^2.*KalPO4+S cal(i,:).*KalPO4.*Ka2PO4+KalPO4.*Ka2 P04.*Ka3P04).*(Scal(i+5.:)+Scal(i+6. :)))... HPO

+3.*(KalPO4.*Ka2PO4.*Ka3PO4./((Scal) i,:)).^3+(Scal(i,:)).^2.*KalPO4+Scal (i,:).*KalPO4.*Ka2PO4+KalPO4.*Ka2PO4 *Ka3PO4).*(Scal(i+5,:)+Scal(i+6,:)))... PO4

+(((Scal(i,:)).*KalCO3)./((Scal(i,:)).^2+(Scal(i,:)).*KalCO3+KalCO3.*Ka2 CO3).*(Scal(i+1,:)+Scal(i+2,:)+Scal(i+3,:))) ... HCO3

+2.*((KalCO3.*Ka2CO3)./((Scal(i,:)). ^2+(Scal(i,:)).*KalCO3+KalCO3.*Ka2CO 3).*(Scal(i+1,:)+Scal(i+2,:)+Scal(i+ 2).*(Scal(i+1,:)+Scal(i+2,:)+Scal(i+2)) 3):"(Scalle,,,, 3,:))); %CO3 %to maintain stablility

if Ch>=0,

B=0.5.*(Ch./(Ch.^2+4.*Kw).^(1/2)+1); else

B=-(Scal(i,:)).^2./(2.*Kw).*(Ch./(Ch .^2+4.*Kw).^(1/2)-1); end,

end

%% U, growth rate function for CO2 function U=UcalCO2(Scal,j) U=umax.*PCO2O2..

.*Scal(i+8,:)./((Scal(i+8,:)+Scal(i+ 8,:).^2./(KiO2cal(Scal(j+1,:),Scal(j
+8,:)))))... 02

.*UaveTrrLfcal(Scal.i)... light

.*min(Scal(j+1,:)./(KsCO2+Scal(j+1,:)),... CO2

min(Scal(j+4,:)./(KsN03+Scal(j+4,:))

(Scal(j+5,:)+Scal(j+6,:))./(KsPO4+(S cal(j+5,:)+Scal(j+6,:)))); % PO4 end

%% U, growth rate function for нсоз

function U=UcalHCO3(Scal,j)
U=umax.*PHCO3O2 ...

*Scal(j+8,:)./((Scal(j+8,:)+Scal(j+ 8,:),^2./(KiO2cal(Scal(j+1,:),Scal(j +8,:)))))... 02

.*UaveIrrLfcal(Scal.i)... light

.*min(Scal(j+2,:)./(KsHCO3+Scal(j+2, :)),... HCO3

min(Scal(j+1,:)./((Scal(j+1,:)+Scal(j+1,:).^2./KiCO2toHCO3)), ... CO2 inhibition on HCO3

min(Scal(j+4,:)./(KsNO3+Scal(j+4,:)) NOT

(Scal(i+5,:)+Scal(i+6,:))./(KsPO4+(S cal(j+5,:)+Scal(j+6,:)))))); %P04 end

%% Dark respiration function Rd=RdO2cal(Scal,j)

Rd=Ures.*(Scal(i+8.:)./(Ks02+Scal(i+ 8,:))); end

%% Calculated maximal growth rate per element when only light is limiting function

Uave=UaveIrrLfcal(Scal,j) % function for calculating light intensity in the biofilm

if t-fix(t)<=tL,</pre> determine if is light else %or dan. Irrcal=0; Irrcal=Irrsurf.

Lfcumleft=zeros(size(Scal)):

Lfcumleft(c*n:-c:c.:)=cumsum(Scal(c* n:-c:c.:),1);

IrrLfleft=(Irrcal.*(1-rs)).*polyval(IzinPer,Lfcumleft(j+9,:));

Lfcumright=zeros(size(Scal));

Lfcumright(c*n-c:-c:c,:)=Lfcumleft(c *n:-c:c+c,:);

IrrLfright=(Irrcal.*(1-rs)).*polyval (IzinPer,Lfcumright(j+9,:));

Uave=(GroIrr_TL(IrrLfright)...Right, to sufac

+GroIrr TL(IrrLfleft))... Left, Further away from surface /2; %take average value of left and right as growth in dLf end

ond

%% Event function, for determing whether dLf has reached Lf(t=0)/n %-----

function [value,isterminal,direction]=EventsL f(t,Scal)

8 the event is now when the nth % the event is now when the nth layer's thickness reaches the preset % Lft0/n0 value, which means the biomass thickness has increased one

whole % grid, and a new discretion is

needed

value=[Scal(c*n,1)-(Lft0./n0),Scal(c
*n,1)-(Lft0./n0)./10^10] ; %Lf(n,t)
reached Lf(t=0)/n0, or the growth is negative

isterminal=[1,1]; direction=[0,-1];

%% Estimate effect of CO2/O2 ration on photorespiration

photorespiration function KiO2=KiO2cal(DCO2,DO2) % function for for estimating the photorespiration, Monod type KRC02to02=0.35; %factors

kRO22002-0.53, should be determined by the control of the control photorespiration factor

Ki02=Ki02max.*(DC02./D02)./(KRC02to0
2+DC02./D02)+Ki02min;
end

%% Estimate the facilitated CO2 transport by EPS + CA (possiblz also cell surface) in biofilms function DCO2bfacilitated=DCO2bcal(Scal,j)

% use a simple log curve to estimat facilitated CO2 transport,local CO2 % concentration dependent fa=100.2;

fb=-le4; fc=1:

DCO2bfacilitated=DCO2b.*(fa.*exp(fb. *Scal(j+1,:))+fc);

end

% H as state variable

function [Ipar] = RTE_TL(XF)
%calculated PAR irradiance inside the
biofilm with Sthe RTE method using DOM method and finite volume with upwind scheme, for %DOM 2 hemisphere each with 24 discrete SDUM 2 nemisphere each with 24 discrete directions are used, weighting %factor are aquired from literature. This method also take into account the %case of optically thin and shallow biofilm, and the reflection by the %substrate layer membrane is taken into account.

LF=1000e-6; % assume a very thick biofilm, i.e. no light penetration. g=0.98; % assumed asymmetry factor g=0.98; % assumed asymmery factor Nnodes=200; %set the number of discrete elements in depth direction.

load('DOM.mat'): % Load discrete ordinate load('OptPro'); % Load biomass optical

propertities, mass absorption: Amass; % mass scttering: Smass; and spectral irradiance at biofilm

% surface (300 total): SpeIsur; reflectivity of biofilm SpeIsur; reflections; surface, rbios; % reflectivity of

[v,disdir]=size(DOM); %angel and ighting factors for DOM, 1st row index

%2nd row angle, 3rd angle in radius,4th row row cos(angle), %5th row weighing

factore

error('check discrete ordinate system setup'),

end.

if g>1||g<-1, fprintf('g value out of range, calculating as no inscattering'), q=2;

and

dz=LF/Nnodes:

dLF=0:dz:LF;

[a,m]=size(SpeGsur); %m:number of different wavelength, a:2(1st row wavelength, 2nd row intensity) [n]=length(dLF); %n:bumber of depth
steps(row is the depth of the nodes) [b,o]=size(Amass); %o:number of different wavelength,b:2(1st row wavelength,2nd row mass absorption) [c,p]=size(Smass); %p:number of different wavelength, b:2(lst row wavelength,2nd row mass scattering)

-=o||a~=2||b~=2||c~=2||o~=p||p~=m, error('ERROR: wavelegth steps are not compatible'), end, %check if the datais compatible

Cabs=zeros(2,m); %absorption coefficient of the biomass, 1st row wavelength, 2nd row coefficient Csca=zeros(2,m); %scattering coefficient of the biomass, 1 wavelength, 2nd row coefficie . lst row

for i=1:m, %calculate absorption and scattering coefficient Cabs(1,i)= Amass(1,i);

Cabs(1,1) - Amass(1,1); Cabs(2,i) = Amass(2,i)*XF; Csca(1,i) = Smass(1,i); Csca(2,i) = Smass(2,i)*XF;

- end,

- Appendix -

ncal=n+1;

SpeIsur=zeros(m,disdir);

for i=1:m,
 for j=1:disdir/2,

SpeIsur(i,j)=((1-rbios)*SpeGsur(2,i)
)/2/pi;
end,
end, %set the diffuse incident

end, %set the diffuse incident irradiance at different wavelength and discrete directions, 12 indicate only for 1/4 sphere

Normalizer=zeros(1,24); for j=1:24, for i=1:24, Normalizer(1,j)=Normalizer(1,j)+DOM(5,i)*(HenGrePhase(DOM(3,i),DOM(3,j), g)+HenGrePhase(DOM(3,i),DDM(3,j),g)

Normalizer(1,j)=1/Normalizer(1,j);
end,

end,

start iferative procedure Spelite=zeros(m,disdir,ncal); start iferative procedure Spelite=zeros(1,2); %set stopping criterion. factor-mese(ncal,m); %set adaptation factor. for k=1ncal-1; Afactor(k,;)=dLF(k)/250e-6; Afactor(kfactor)]=1;

end
while
SpeIbot(1,1)==SpeIbot(1,2)||abs(SpeI
bot(1,1)-SpeIbot(1,2))/SpeIbot(1,2)>
0.0,
ittinuity
ittinuity
ittinuity
ittinuity
ittinuity
ittinuity
iterative process
ite=ite+1;
for i=:m.

Spelite(i,j,1)=Spelsur(i,j); end, end, end,%apply new surface boundary

condition

Bi=zeros(m,disdir); ai=zeros(m,disdir); bi=zeros(m,disdir,disdir); ci=zeros(m,disdir,ncal); Ai=zeros(disdir,disdir); Ci=zeros(disdir,1);

for i=1:m,
 for k=2:ncal-1,
 for j=1:disdir ,

Bi(i,j)=Cabs(2,i)*Afactor(k,i)+Csca(2,i)-Csca(2,i)*DOM(5,j)*(HenGrePhase (DOM(3,j),-DOM(3,j),g)+HenGrePhase(D OM(3,j),DOM(3,j),g)*Normalizer(1,j)

if j>disdir/2,

ai(i,j)=Bi(i,j)*dz-DOM(4,j);

ci(i,j,k)=-DOM(4,j)*Spelite(i,j,(k+1
));
else

ai(i,j)=Bi(i,j)*dz+DOM(4,j);

```
ci(i,j,k)=DOM(4,j)*SpeIite(i,j,(k-1));
```

end, for j1=1:disdir, if j==j1,

j represent inclient discretedirection, j represents outgoing discrete direction end,

end, for p=1:disdir, for q=1:disdir, Ci(p,1)=ci(1,p,k); Ai(p,q)=bi(1,p,q); end, end, %establish matrix for linsolve function

Spelite(i,j,k)=Spelk(j,l); end,&integrate solution result end, Calculate local spectral irradiance for i=1:m,

for i=1:m,
 for j=1:disdir,
 if j>disdir/2

Spelite(i,j,ncal)=rmems*Spelite(i,di
sdir_j+1,ncal-1);
 end,
 end,

end, stransfer new boundary condition at the substrate biofilm inteface.

for i=1:m,
 for k=ncal-1:-1:2,
 for j=1:disdir ,
 if j>disdir/2,

ci(i,j,k)=-DOM(4,j)*SpeIite(i,j,(k+1
));
else

end, for p=1:disdir, for q=1:disdir, Ci(p,1)=ci(i,p,k); Ai(p,q)=bi(i,p,q); end,

end,%establish matrix for linsolve function

end, %Calculate local spectral irradiance

for k=1:ncal, for i=1:m, for j=1:disdir. SpeGzite(i,k)=SpeGzite(i,k)+2*pi*Spe Iite(i,j,k)*DOM(5,j); end, if Cabs(1.i)<=500. GpariteB(1,k)=GpariteB(1,k)+SpeGzite (i,k); elseif 500<Cabs(1,i)&&Cabs(1,i)<=600. GpariteG(1,k)=GpariteG(1,k)+SpeGzite (i,k); else GpariteR(1,k)=GpariteR(1,k)+SpeGzite (i,k); ond end end, Scalculate local spectral intensity and PAR irradiance at depth z, *2 indicate 2 heremisphere for total and RGB GpariteT=GpariteB+GpariteG+GpariteR; Local total PAR irradianc SpeIbot(1,1)=SpeIbot(1,2);
SpeIbot(1,2)=sum(GpariteT(1,:)); if g==2, break, end. if ite>300, warning('ite > 30, exit loop'); break. end, ***** %end of iterative process end, Ipar=zeros(5,n-1);
Ipar(1,:)=dLF(1:n-1); for k=2:n: Ipar(2,k-1)=GpariteT(1,k); %Total Ipar(3,k-1)=GpariteB(1,k); %Blue Ipar(4,k-1)=GpariteG(1,k); %Green Ipar(5,k-1)=GpariteR(1,k); %Red ond if g==0, gcha='g=0,isotropic scattering'; elseif (0<g)&&(g<1), gcha=['g=',num2str(g),',forward
cattering']; scattering']; elseif (-1<g)&&(g<0), gcha=['g=',num2str(g),',backward scattering']; else gcha='No inscattering'; ond figure('Name', 'Irradiance
distribution in Twin-layer biofilm'); hold on; plot (Ipar(1,:), plot (lpar(1;), Ipar(2;)./Ipar(2,2).*100, '-black'); plot (lpar(1;), Ipar(3;)./Ipar(2,2).*100, '-blue'); plot (Ipar(1,:), Ipar(4,:)./Ipar(2,2).*100, '-green');

SpeGzite=zeros(m.ncal); % Local

GpariteG=zeros(1,ncal); % Local

GpariteR=zeros(1.ncal); % Local red

spectral intensity
GpariteB=zeros(1,ncal); % Local

blue PAR irradiance

green PAR irradia

DAR irradiance

plot (Ipar(1,:), Ipar(5,:)./Ipar(2,2).*100, '-red'); xlabel('Depth into the biofilm in meter'); ylabel('Local irradiance in % of the total PAR at biofilm surface'); title('Irradiance distribution in Twin-layer biofilm'); text(Tpar(i,int8(n/2)),Tpar(2,int8(n /2)),[`\leftarrow' goha]; HorizontalAlignment','left',' FontSize',12);

bre

function [PhasefunHG] = HenGrePhase(Ain,Aout,g) %HenGreOhase calculate the phase function apprimation with the %Henyey-Greenstein method, PhasefunHG returns the calculated phase function.

This function calculate the phase % This function calculate the phase function for strongly forwarding % medium, and need to variables for the approximation, g is the % coeffcient, generally between 0.95

coerclent, generally between 0.55 to 0.99 for strongly forwarding % medium, and the scattering angle, which can be found in variable DOM.

u0=cos(Ain+Aout); PhasefunHG=(1-g^2)/(1+g^2-2*g*u0)^(3 /2)/2; %calculate the HG phase function

end

function [fIrr] = GroIrr_TL(Irr)
%Growth when irradiance is the only factor aI, bI, cI factors, refer to Platt and

Gallegos, 1980 %% factors for irradiance's influence

on growth on growth
ETRmax=49.6; % micromole electron per
m^2 per s, max ETR
alpha=0.24; % initial slope of ETR curve
beta= 0.0068; % factor representing
photo inhibition

PBs=ETRmax; a=alpha; b=beta;

fIrr=(1-exp(-a.*Irr./PBs)).*exp(-b.* Irr./PBs); %Growth rate when irrandiance is the only factor

ond

function [

DbioCO2, DbioHCO3, DbioCO3, DbioNO3, Dbi oHPO4, DbioH2PO4, DbioO2, DbioNa, DbioH,

DsubCO2, DsubHCO3, DsubCO3, DsubN, DsubH PO4, DsubH2PO4, DsubO2, DsubNa, DsubH 1....

= DiffCoCal_TL(BioPoro, SubPoro) %Calculated effective diffusion coefficient in TL biofilm

%%%%% diffusion coefficients in water \$%%% diffusion coefficients in water DO2=1.65=4; %diffusion coefficient of CO2 in water in m²/day^, 1.65=4 PRC03=1.02=4; %diffusion coefficient of HCO3 in water in m²/day DCO3=7.9=-5; %diffusion coefficient CO3 in water in m²/day DMO3=1.47=4; %diffusion coefficient of NO3 in water in m^2/day DHPO4=8.2e-5; %diffusion coefficient of HPO4 in water in m^2/day DH2PO4=6.5e-5; % diffusion coefficient of H2PO4 in warer in m^2/day DO2=1.73e-4; %diffusion coefficient of O2 in water in m^2/day $_{\rm V2}$ in water in m^2/day DNa=1.15e-4; %diffusion coefficient of Na in water in m^2/day DH=6.05e-3; %diffusion coefficient of H in water in m^2/day %%%%%

BioRDC=BioPoro/(1+0.5*(1-BioPoro)); % SubRDC=SubPoro/(1+0.5*(1-SubPoro));% refer to Weissberg, 1963

DbioCO2=BioRDC*DCO2; DbioHCO3=BioRDC*DHCO3: DbioCO3=BioRDC*DCO3; DbioNO3=BioRDC*DNO3; DbioHPO4=BioRDC*DHPO4 DbioH2PO4=BioRDC*DH2PO4; DbioH2PO4=BioRDC*DH2PO4; DbioO2=BioRDC*DO2; DbioNa=BioRDC*DNa; DbioH=BioRDC*DH;

DsubCO2=SubRDC*DCO2; DsubHCO3=SubRDC*DHCO3; DsubCO3=SubRDC*DCO3; DsubN=SubRDC*DNO3; DeubHPO4=SubPDC*DHPO4. DsubH2P04=SubRDC*DH2P04; DsubH2P04=SubRDC*DH2P04; DsubO2=SubRDC*DO2; DsubNa=SubRDC*DNa; DsubH=BioRDC*DH;

ond

Appendix III: Author's contribution to the publications included in this thesis

1St manuscript: A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to evaporation (Published in the journal 'Journal of Microbiological Methods').

Authors: Tong Li, Björn Podola, Dirk de Beer and Michael Melkonian

Authorship: First Author

- Contribution in experimental work (%): Sample preparation; Experimental measurement with the microsensors (50%).
- Contribution in theoretical part (%): Development of the mathematical concept; Implementation of the mathematical concept; Analysis of data (80%).

Contribution in writing (%): Preparation of the original manuscript including introduction, methods and materials, results and discussion (70%).

- 2nd manuscript: Microscale profiling of photosynthesis-related parameters in a highly productive biofilm photobioreactor (Submitted to the journal 'Biotechnology and Bioengineering', under review).
 - Authors: Tong Li, Bastian Piltz, Björn Podola, Anthony Dron, Dirk de Beer and Michael Melkonian
 - Authorship: First Author
 - Contribution in experimental work (%): Sample preparation; Experimental measurement with the microsensors (90%).
 - Contribution in theoretical part (%): Data analysis; Discussion of results (80%).
 - Contribution in writing (%): Preparation of the original manuscript including introduction, methods and materials, results, discussion and supplementary materials (75%).
- 3rd manuscript: Investigating dynamic processes in a porous substrate biofilm photobioreactor A modeling approach (finished manuscript).

Authors: Tong Li, Björn Podola and Michael Melkonian

- Authorship: First Author
- Contribution in experimental work (%): Sample preparation; Experimental work carried out in the framework of the study (90%).
- Contribution in theoretical part (%): Model development and implementation; Data analysis (90%).

Contribution in writing (%): Preparation of the original manuscript including introduction, methods and materials, results, discussion and supplementary materials (85%).

Erklärung

Ich versichere, dass ich die von mir vorgelegte Dissertation selbständig angefertigt, die benutzten Quellen und Hilfsmittel vollständig angegeben und die Stellen der Arbeit – einschließlich Tabellen, Karten und Abbildungen –, die anderen Werken im Wortlaut oder dem Sinn nach entnommen sind, in jedem Einzelfall als Entlehnung kenntlich gemacht habe; dass diese Dissertation noch keiner anderen Fakultät oder Universität zur Prüfung vorgelegen hat; dass sie – abgesehen von oben (Teil 2 und Appendix III) angegebenen Teilpublikationen – noch nicht veröffentlicht worden ist, sowie, dass ich eine solche Veröffentlichung vor Abschluss des Promotionsverfahrens nicht vornehmen werde. Die Bestimmungen der Promotionsordnung sind mir bekannt. Die von mir vorgelegte Dissertation ist von Prof. Dr. Michael Melkonian betreut worden.

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Lebenslauf

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Promotion in Prüfungsfach Botanik

Thema der Dissertation: 'Dynamic interactions in an artificial phototrophic biofilm for biotechnological applications'

Forschungsbereiche:

Algenbiotechnologie, Biofilm Modeling, Biofilm Microsensormessungen, Biofilm Imaging

Publikationen:

Continuous removal of zinc from wastewater and mine dump leachate by a microalgal biofilm PSBR. Tong Li, Gengyi Lin, Björn Podola, Michael Melkonian, Journal of Hazardous Materials 297 (2015) 112–118.

A method to determine photosynthetic activity from oxygen microsensor data in biofilms subjected to evaporation. Tong Li, Björn Podola, Dirk de Beer, and Michael Melkonian, Journal of Microbiological Methods 117 (2015) 100-107.

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