

1           1           **Free ammonia inhibition in microalgae and cyanobacteria grown in**  
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3           2           **wastewaters: photo-respirometric evaluation and modelling**

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38           18           **Abstract:** The inhibitory effects of free ammonia (FA) on microalgae/cyanobacteria in  
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40           19           wastewater-treating photobioreactors (PBR) can strongly reduce their treatment efficiency,  
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42           20           increasing the operational costs and undermining the stability of the system. Although FA-  
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44           21           promoting conditions (high pH, temperature and ammoniacal nitrogen concentration) are  
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46           22           commonly met in outdoor PBRs, photosynthesis inhibition from FA has been scarcely explored  
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48           23           and is rarely considered in microalgae-bacteria growth models. Two pilot systems and a series  
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50           24           of lab-scale monocultures were tested using a photo-respirometry approach, to evaluate the  
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52           25           effects of FA (8.5 - 136 mg NH<sub>3</sub> L<sup>-1</sup>) on photosynthesis. Two mathematical inhibition models  
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54           26           were compared, with the aim of selecting best-fitting equations to describe photo-respirometric  
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56           27           experiments. A set of calibrated inhibition parameters was obtained for microalgae and  
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58           28           cyanobacteria, growing in monocultures or in mixed algae-bacteria consortia. Cyanobacteria

29 were more sensitive to FA than green microalgae and mixed phototrophs-bacteria consortia  
30 showed a higher resistance compared to monocultures. Estimated inhibition parameters were  
31 used to describe common operational/environmental conditions in algae-bacteria systems,  
32 demonstrating the potential drop in photosynthetic activity under those relevant operational  
33 conditions.

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35 **Keywords:** Free ammonia inhibition; Photosynthetic oxygenation modelling; Wastewater  
36 treatment; Microalgae and cyanobacteria; Algae/bacteria consortia

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38 **Highlights:**

- 39 • Free ammonia inhibition was studied in *chlorophyceae* and cyanobacteria
- 40 • Both monocultures and mixed phototrophs-bacteria consortia were investigated
- 41 • The EC<sub>50</sub> for free ammonia was higher for green microalgae than cyanobacteria
- 42 • The EC<sub>50</sub> was higher in mixed phototrophs-bacteria consortia than in monocultures
- 43 • Free ammonia inhibition is modelled under common weather/operational conditions

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## 45 1. INTRODUCTION

46 The design and operation of conventional biological wastewater treatment  
47 processes, such as activated sludge or anaerobic processes, rely on exhaustive  
48 knowledge, standardized methodologies and robust mathematical modelling,  
49 allowing to achieve high and stable removal efficiencies (van Loosdrecht et al.,  
50 2016). Aerobic processes are especially characterized by severe limitations: i)  
51 high operational costs and energy requirements for the aeration of mixed liquor  
52 and sludge dewatering (Plappally & Lienhard, 2012) and ii) high environmental

1 53 impacts for direct and indirect greenhouse gases (CO<sub>2</sub>, CH<sub>4</sub> and N<sub>2</sub>O)  
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3 54 emissions (Campos et al., 2016). In the last years there has been a renewed  
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5 55 interest in microalgae treatment systems, because it has been shown that  
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7 56 growing microalgae/cyanobacteria in consortia with wastewater-associated  
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9 57 bacteria, can lead to significant advantages, while guaranteeing effluent quality  
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11 58 (Muñoz et al., 2006; Campos et al., 2016). In fact, the algae-bacteria  
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13 59 metabolites exchange (CO<sub>2</sub> and O<sub>2</sub>), can significantly reduce costs for aeration.  
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15 60 Moreover, sludge disposal costs can be reduced by recovering nutrients  
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17 61 (biofertilizers/biostimulants) and/or energy (biogas) from the algal-bacterial  
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19 62 biomass (Moreno-García et al., 2017). Phototrophic microorganisms can adapt  
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21 63 and survive in extreme environmental conditions but some compounds, such as  
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23 64 antibiotics, herbicides/pesticides, heavy metals or unionized NH<sub>3</sub> (or free  
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25 65 ammonia, FA), can cause a severe inhibition of photosynthesis, hampering the  
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27 66 biological activity and therefore reducing the oxygen supply by phototrophs and  
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29 67 mining the efficiency of the system.  
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38 68 FA has been identified as an important short-term and species-specific inhibitor  
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40 69 of the photosynthetic process, with some species being particularly resistant to  
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42 70 elevated concentrations of ammonia and other being much more sensitive  
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44 71 (Collos & Harrison, 2014; Markou et al., 2016; Gutierrez et al., 2016). The  
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46 72 inhibition of photosynthesis can be associated to different mechanisms of  
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48 73 action: i) ammonia causes damages to the oxygen evolving complex (OEC) of  
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50 74 the photosystem II, acting as an uncoupler of the Mn cluster of the OEC and  
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52 75 displacing a water ligand (Drath et al., 2008; Markou and Muylaert, 2016;  
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54 76 Markou et al., 2016); ii) ammonia diffuses through membranes and  
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1 77 accumulates, acting as an uncoupler and disrupting the  $\Delta pH$  component of the  
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3 78 thylakoid proton gradient (Belkin and Bossiba, 1991, Markou and Muylaert,  
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5 79 2016; Gutierrez et al., 2016). Besides these effects, the activity of photosystem I  
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8 80 and the dark respiration rates are also negatively affected by FA and ammonia  
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10 81 toxicity also seems to be amplified at elevated light intensities, although the  
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12 82 mechanisms are not fully understood (Markou et al, 2016). Quantifying the  
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14 83 inhibition of photosynthetic activities is of particular interest in algae-based  
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16 84 wastewater treatment processes (Abeliovich & Azov, 1976; Kallqvist and  
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18 85 Svenson, 2003; Tan et al., 2016; Goto et al., 2019). Indeed, the equilibrium  
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20 86 reaction between FA and ammoniacal nitrogen ( $NH_4^+$ ) shifts toward FA under  
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22 87 the following conditions: i) high pH values associated to photosynthetic  
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24 88 processes, ii) high temperatures due to atmospheric conditions and iii) high total  
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26 89 ammoniacal nitrogen ( $TAN = NH_3 + NH_4^+$ ) concentrations (Anthonisen et al.,  
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28 90 1976; Collos and Harrison, 2014). The inhibition of photosynthesis due to the  
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30 91 presence of FA could be assessed in different ways: by carrying out batch  
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32 92 growth experiments (Källqvist and Svenson, 2003; Gutierrez et al., 2016; Bo et  
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34 93 al., 2016; Zhao et al., 2019); by measuring the uptake of nutrients or oxygen  
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36 94 (Abeliovich and Azov , 1976; Azov and Goldman, 1981; Boussiba et al., 1991;  
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38 95 Dai et al., 2008; Segura et al., 2017), or by coupling pulse-amplitude modulation  
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40 96 (PAM) with photo-respirometry or with some of the aforementioned techniques  
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42 97 (Drath et al., 2008; Markou et al., 2016; Markou and Muylaert, 2016; Li et al.,  
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44 98 2019; Wang et al., 2019). Despite the relevance of FA inhibition, and the  
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46 99 availability of mathematical models to describe this phenomenon (e.g.,  
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48 100 Andrews, 1968; Han and Levenspiel, 1987, among others), FA inhibition is  
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1 101 rarely considered when analysing and modelling the evolution of microalgae-  
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3 102 bacteria consortia (Shoener et al., 2019). In addition, FA inhibition assays on  
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5 103 microalgae-bacteria suspensions are not documented in literature, and almost  
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8 104 all experimental works are performed on axenic cultures using synthetic media,  
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11 105 apart from a few cases (Hernandèz et al., 2013; Beyl et al., 2019).  
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14 106 The objectives of this study were to quantify the inhibition of photosynthesis due  
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16 107 to FA in different phototrophic organisms, and to identify a mathematical photo-  
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18 108 oxygenation model, describing FA inhibition under typical operational conditions  
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21 109 in microalgae/cyanobacteria-based bioremediation systems. Photo-  
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23 110 respirometric tests (PRT) were carried out on cyanobacteria and green  
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26 111 microalgae monocultures and on mixed phototrophs-bacteria suspensions  
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28 112 dominated by microalgae and cyanobacteria, in order to analyse the inhibition of  
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31 113 different phototrophic microorganisms exploited in wastewater treatment. In  
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33 114 view of standardizing PRTs and comparing the results obtained with other  
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36 115 literature works, a detailed description of experimental procedures was also  
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38 116 included and discussed. The obtained dataset was used to compare two  
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41 117 different models (a simple non-competitive inhibition model and a logistic  
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43 118 sigmoidal function), and best-fit equations were used to describe common  
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46 119 operational/environmental conditions in algae-bacteria systems, demonstrating  
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48 120 the significance of the potential drop in photosynthetic oxygenation due to the  
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51 121 presence of FA. To the best knowledge of authors, this is the first study in which  
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53 122 the effects of FA on the photosynthetic activity of eukaryotic microalgae and  
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56 123 cyanobacteria were directly compared, using data from both monocultures and  
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58 124 mixed algae-bacteria consortia.  
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## 2. MATERIALS AND METHODS

### 2.1. Microorganisms and cultivation systems

#### 2.1.1. Pilot-scale phototrophs-bacteria cultivation systems

##### Microalgae-bacteria

The microalgae-bacteria consortium was grown in a pilot-scale high rate algal pond (HRAP) (volume: 1.2 m<sup>3</sup>, surface = 5.8 m<sup>2</sup>, liquid height = 0.2 m) located at the Bresso-Niguarda wastewater treatment plant (WWTP) (Milan, Italy). The liquid fraction of anaerobically digested sludge (LFAD) was separated by centrifugation and used to feed the HRAP, having demonstrated good characteristics as growth substrate. More information about the wastewater characterisation is reported in previous studies (Mantovani et al., 2019; Marazzi et al., 2019). The pilot scale HRAP was installed outdoor and covered with a polycarbonate roof to protect the pond from rain. The HRAP was inoculated with the microalgae-bacteria suspension with *Chlorella sp.* and *Scenedesmus sp.* as the dominant species and operated in continuous with undiluted LFAD as the feed at an average hydraulic retention time (HRT) of 10 d for seven months (April 2019 - October 2019).

##### Cyanobacteria-bacteria

The cyanobacteria-bacteria consortia were grown in a set of three demonstrative-scale tubular semi-closed photobioreactors (PBR) (volume = 11.7 m<sup>3</sup> each), located in the Agròpolis experimental campus of Universitat

1 148 Politècnica de Catalunya (UPC) (Barcelona, Spain). Detailed information about  
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3 149 PBRs design, start-up and operations characterisation is available in García et  
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6 150 al. (2018). In brief, each PBR consisted of 2 lateral open tanks connected  
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8 151 through 16 transparent tubes (diameter = 125 mm, length = 47 m). Each tank  
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10 152 was equipped with a paddle wheel ensuring proper mixing and circulation of the  
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12 153 suspension through the tubes and the reduction of excess dissolved oxygen  
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14 154 (DO). The three PBRs were connected in series to promote the selection of  
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16 155 cyanobacteria and the production and the accumulation of biopolymers, using  
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18 156 agricultural runoff as feedstock medium. Nutrient concentrations in the first PBR  
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20 157 were adapted to reach the optimum ratio favouring the growth of cyanobacteria  
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22 158 over green microalgae, by adding an external source of NO<sub>3</sub> (potassium nitrate  
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24 159 inorganic fertilizer, NK13-46, 13% N-NO<sub>3</sub>). The culture was mainly dominated  
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26 160 by cyanobacteria of the species *Synechococcus* sp. and *Synechocystis* sp. In  
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28 161 the second PBR, a feast and famine regime was applied by adding an external  
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30 162 inorganic carbon source during 6 h d<sup>-1</sup>, in order to enhance the cyanobacterial  
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32 163 carbon uptake efficiency and subsequent biopolymers accumulation. In the third  
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34 164 PBR, the inorganic carbon was continuously provided to increase the  
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36 165 accumulation of biopolymers after the feast and famine phase. CO<sub>2</sub> and sodium  
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38 166 bicarbonate (NaHCO<sub>3</sub>) were used as external inorganic carbon source in both  
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40 167 the second and third PBRs. CO<sub>2</sub> was injected by means of diffusers in the  
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42 168 lateral open tanks of the PBRs and regulated by a pH-control system. NaHCO<sub>3</sub>  
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44 169 was added by a daily dose of a concentrated solution of NaHCO<sub>3</sub>. Detailed  
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46 170 information about the operational strategies adopted, wastewater characteristics  
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1 171 and biopolymers production can be found elsewhere (Díez-Montero et al., 2019,  
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3 172 Rueda et al., submitted).  
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## 10 174 **2.1.2. Cultivation of green microalgae and cyanobacteria**

### 11 12 175 **monocultures**

#### 13 14 15 176 Green microalgae

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18 177 Four different species of green microalgae were selected after microscope  
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20 178 observations in the HRAP and cultured in the laboratories of the *Istituto*  
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22 179 *Sperimentale Italiano Lazzaro Spallanzani* (Rivolta d'Adda, IT). In particular, two  
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24 180 strains of *Chlorella spp.* (*Chlorella vulgaris*, SAG211-11j and *Chlorella*  
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26 181 *sorokiniana*, SAG211-8k) and one strain of *Scenedesmus* (*Scenedesmus*  
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28 182 *quadricauda*, or *Desmodesmus armatus*, SAG276-4d) were acquired from the  
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31 183 Culture Collection of Algae at the University of Göttingen (SAG, Germany),  
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33 184 while one strain of *Scenedesmus spp.* (identified as *Scenedesmus obliquus*)  
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35 185 was isolated from an outdoor pond. All strains were cultured in 500 mL glass  
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37 186 Erlenmeyer flasks, using commercially available Modified Bold Basal Medium  
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39 187 (MBBM, Sigma-Aldrich), at room temperature (20 - 25 °C) and under controlled  
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41 188 irradiance (cool white fluorescent lamps, Philips F58W/33-640 58W) and 12  
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43 189 h/12 h light/dark (L/D) cycles. Sterile air (0.2 µm cutoff) was bubbled in the  
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45 190 PBRs to provide carbon dioxide and mixing. The cultivation of green microalgae  
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47 191 was achieved without pH-control.  
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#### 58 193 Cyanobacteria

1 194 Three different species of cyanobacteria were identified and isolated from the  
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3 195 semi-closed PBRs fed with agricultural runoff and cultured under controlled  
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5 196 conditions: *Synechococcus* sp., *Synechocystis* sp., and *Leptolyngbya* sp. The  
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8 197 strains were sampled from the first semi-closed PBR and inoculated in plates  
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10 198 prepared with 1% bacteriological agar and commercially available BG11  
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12 199 medium (Sigma-Aldrich, St. Louis, US), by direct streaking or after serial  
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14 200 dilutions in saline media, as explained in Rueda et al. (2020). Once  
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16 201 cyanobacteria colonies were obtained, they were transferred into 2 mL of  
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18 202 medium contained in 15 mL test tubes and scaled-up (scaling ratio = 1:5), until  
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20 203 1 L cultures were obtained. Finally, they were kept in Erlenmeyer flasks at room  
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22 204 temperature ( $30 \pm 2^\circ\text{C}$ ), under controlled irradiance (approximately  $36.2 \mu\text{E m}^{-2}$   
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24 205  $\text{s}^{-1}$  using 14W cool-white LED lights) under 15 h/9 h L/D cycles. Sterile air (0.2  
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26 206  $\mu\text{m}$  cutoff) was bubbled to provide mixing and  $\text{CO}_2$  and to remove accumulated  
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28 207 DO. The cultivation was achieved without pH-control. More information about  
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30 208 the cultivation of cyanobacteria monocultures can be found elsewhere Rueda et  
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32 209 al. (2020).  
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## 44 211 **2.2. Respirometric unit**

46 212 The activity of phototrophic organisms was evaluated using a fully equipped  
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48 213 photo-respirometric/titrimetric unit, provided with different options for DO- and  
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50 214 pH-control. The photo-respirometer (IDEA Bioprocess Technologies s.r.l.)  
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52 215 includes: a closed bioreactor (0.5 L glass bottle, DURAN protect, GLS80  
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54 216 headplate), a gas injection system (an airpump and a gas cylinder connected to  
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1 217 a set of electro-valves), a signal/communication and mixing (0 - 300 RPM) unit  
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3 218 and an acid/base dosage system (two 0 - 12 RPM peristaltic pumps). The DO-  
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5 219 control system was made of a DO probe (Hamilton VisiFerm, DO Arc 120) and  
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8 220 a DO-stat system by O<sub>2</sub>/N<sub>2</sub> bubbling, while the pH-control system was made of  
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11 221 a pH probe (Hamilton Polylite Plus, H Arc 120) and a pH-control system by CO<sub>2</sub>  
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13 222 bubbling or acid/base dosage. The entire system was controlled with an  
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15 223 industrial grade PC running a LabView-based control software (Figure 1). The  
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18 224 sampling interval for temperature, pH and DO data was set to 3 s. During  
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21 225 cyanobacteria and cyanobacteria-bacteria tests, the light source was an  
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23 226 incandescent light bulb (30 W), providing a photosynthetically active radiation  
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25 227 (PAR) level of about  $116 \pm 23 \mu\text{E m}^{-2} \text{s}^{-1}$ . During microalgae and microalgae-  
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28 228 bacteria tests, irradiance and temperature were controlled by placing the  
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31 229 vessels into a thermostatic chamber provided with irradiance and air  
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33 230 temperature regulation (F.lli Della Marca s.r.l., TS series). In this case, four  
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35 231 internal fluorescent elements (OSRAM L36W/965 - Deluxe cool daylight) were  
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38 232 switched on/off, reaching a similar light intensity as the one obtained for tests  
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40 233 performed on cyanobacteria ( $108 \pm 16 \mu\text{E m}^{-2} \text{s}^{-1}$ ). The incident PAR radiation  
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43 234 at 400 - 700 nm was measured along the internal surface of the glass bottle by  
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45 235 using a quantum sensor (Apogee Instruments, MQ-500).

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### 51 237 **2.3. Experimental procedures, test conditions and set of experiments**

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54 238 The effects of FA concentrations were assessed under standardized condition  
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57 239 with respect to: incident irradiance ( $110 \mu\text{E m}^{-2} \text{s}^{-1}$ ), pH (8.5) and temperature  
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1 240 (20 °C). The DO concentration was maintained in the range 100 - 130% of the  
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3 241 DO saturation at the test temperature by air bubbling. For cyanobacteria, FA  
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5 242 was tested at 8.5, 17, 34 and 68 mg NH<sub>3</sub> L<sup>-1</sup>, while microalgae were exposed to  
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8 243 higher concentrations, i.e. 17, 34, 68 and 136 and mg NH<sub>3</sub> L<sup>-1</sup>, due to the higher  
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10 244 tolerance to ammonium/ammonia reported in literature (Collos & Harrison,  
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12 245 2014). The selected range of FA concentrations was typically used in previous  
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14 246 FA inhibition assays, adequately covering the values expected during  
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16 247 microalgae-/cyanobacteria-based bioremediation and including quite high FA  
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18 248 concentrations to represent the case of high strength wastewaters with high  
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20 249 TAN concentrations (e.g. anaerobic digestates, landfill leachates). A total of  
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22 250 seven pure microalgae/cyanobacteria samples and four mixed microalgae-  
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24 251 /cyanobacteria-bacteria consortia were used for activity assessments at  
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26 252 different FA concentrations. A summary of performed PRTs and test conditions  
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28 253 is reported in Table 1.

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30 254 The test protocol was defined by slightly modifying a standardized protocol,  
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32 255 adopted for calibrating microalgae-bacteria models and evaluating best  
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34 256 equations to describe the effects of environmental conditions (Rossi et al.,  
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36 257 2018; Rossi et al., 2020a; Rossi et al., 2020b). The protocol included the  
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38 258 following steps: 1) sampling/transportation, 2) pre-treatments, 3) sample  
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40 259 characterisation, 4) addition of nitrifying inhibitors (except for monocultures), 5)  
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42 260 addition of nutrient solutions (except for monocultures), 6) acclimation to test  
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44 261 conditions (1.5 h), 7) excess DO removal, 8) alternation of L/D phases with  
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46 262 (inhibited reactor) and without (control reactor) NH<sub>3</sub> additions, 9) data  
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48 263 processing/modelling. Some of these aspects are discussed below.

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**2.3.1. Pre-treatments**

The samples were taken from the lab-scale cultivation systems and diluted with fresh media (MBBM and BG11 for microalgae and cyanobacteria monocultures, respectively), for reaching the desired optical density at 680 nm ( $OD_{680}$ ) and ensuring nutrient availability. After preliminary evaluations of the volumetric oxygen production and uptake rates (OPR and OUR, respectively) at different light intensities (data not shown), the initial  $OD_{680}$  was set to 0.2 to conduct PRTs under optimal (i.e. non-limiting and non-inhibiting) conditions of light availability, while avoiding a too fast DO accumulation in the vessel. The microalgae-bacteria suspension sampled from the HRAP were first screened with a 300  $\mu$ m mesh to remove detached biofilms and inert particles. The suspension was concentrated by centrifugation (5000 RPM, 10 min) and an appropriate amount of biomass was resuspended into a nutrient-free mineral medium up to  $OD_{680} = 0.2$ . The mineral medium used for resuspension was designed to mimic the LFAD (Rossi et al., 2018), according to the average concentrations of metals (Na, Mg, Ca, K, Al, Mn, Fe, Co, Ni, Cu and Zn) in the algae-bacteria suspension. Regarding cyanobacteria-bacteria consortia, the biomass developed in the semi-closed PBRs contained filamentous cyanobacteria flocs, making it difficult to determine the  $OD_{680}$ . Therefore, total suspended solids (TSS) concentrations were used to evaluate the amount of biomass to be diluted in the effluent of each PBR. Dilution with the effluent was preferred to dilution with synthetic media, since it was not possible to characterize the ionic composition of semi-closed PBRs suspensions.

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### **2.3.2. Addition of bacterial inhibitors and nutrient solutions**

During PRTs performed on mixed cultures of phototrophs/bacteria, Allylthiourea (ATU, 10 mg L<sup>-1</sup>) was added to inhibit nitrifying activity while the photosynthetic activity remained unaffected (Rossi et al., 2018; Rada-Ariza, 2018). Inorganic nutrients were also supplied in a control reactor, to determine the maximum photosynthetic activity, by injecting concentrated solutions of NH<sub>4</sub>Cl, K<sub>2</sub>HPO<sub>4</sub> and NaHCO<sub>3</sub> (15 mg N-NH<sub>4</sub><sup>+</sup> L<sup>-1</sup>, 5 mg P-PO<sub>4</sub><sup>3-</sup> L<sup>-1</sup>, 150 mg C-HCO<sub>3</sub><sup>-</sup> L<sup>-1</sup>). The background FA concentration corresponding to the NH<sub>4</sub>Cl injection was approximately 1.7 mg NH<sub>3</sub> L<sup>-1</sup>. The total volume of injected solutions (nutrients, inhibitors and acid/base titrants) did not substantially modify the biomass concentration and thus of the light extinction in the respirometer (maximum volume increase: 4%). Nutrients and bacterial inhibitors were not added during PRTs performed on microalgae/cyanobacteria monocultures, as the nutrient availability was guaranteed by the resuspension in the mineral medium.

### **2.3.3. Acclimation to test conditions and L/D phases**

Before starting each PRT, the biomass was incubated under the test conditions (1.5 h), for an equilibration of metabolic activities to the new light, temperature and pH. During PRTs, the biomass was exposed to light for 10 min, and then kept in the dark for 20 min. In the control reactor, this L/D cycle was repeated from three to six times for improving statistical significance and identifying stable initial conditions in terms of volumetric OPRs and OURs. In the inhibited

311 reactor, concentrated ammonia solutions were dosed at the beginning of each  
 312 light phase.

313

## 314 2.4. Data processing and calculations

### 315 2.4.1. Calculation of oxygen production rates

316 The DO dynamics was modelled by considering the concomitant occurrence of  
 317 (i) either a constant net photosynthetic oxygen production rate (during L  
 318 phases) or a respiratory oxygen uptake rate (during D phases), and (ii) the  
 319 oxygen mass transfer rate at the liquid-gas interface (OTR). The resulting  
 320 dynamic mass balance for the DO in the photo-respirometer is therefore  
 321 (Equations 1-3):

$$\begin{cases} \frac{d(\text{DO})}{dt} = \text{OPR}_{\text{NET},i} + \text{OTR} & \text{(Light phases, } L_i, i=1, \dots, 5) \\ \frac{d(\text{DO})}{dt} = \text{OUR}_{\text{RESP},i} + \text{OTR} & \text{(Dark phases, } D_i, i=1, \dots, 5) \end{cases} \quad \#(1)$$

$$\text{OTR} = \theta^{(T-293.15)} \cdot k_L a_{20} \cdot (\text{DO}_{\text{SAT}} - \text{DO}) \quad \#(2)$$

$$\text{DO}_{\text{SAT}} = p\text{O}_2 \cdot K_{\text{H},\text{O}_2}(T) = p\text{O}_2 \cdot K_{\text{H},\text{O}_2, \text{REF}} \cdot \exp\left(-\frac{\Delta_{\text{SOL}}H}{R} \cdot \left(\frac{1}{T} - \frac{1}{T_{\text{REF}}}\right)\right) \quad \#(3)$$

322 Where: DO [mg O<sub>2</sub> L<sup>-1</sup>] is the DO concentration at the time t [h], OPR<sub>NET,i</sub> [mg  
 323 O<sub>2</sub> L<sup>-1</sup> h<sup>-1</sup>] is the average net OPR during the i<sup>th</sup> light phases L<sub>i</sub>, OUR<sub>RESP,i</sub> [mg  
 324 O<sub>2</sub> L<sup>-1</sup> h<sup>-1</sup>] is the average respiratory OUR during the i<sup>th</sup> dark phases D<sub>i</sub>, OTR is  
 325 the oxygen mass transfer rate [mg O<sub>2</sub> L<sup>-1</sup> h<sup>-1</sup>], DO<sub>SAT</sub> [mg O<sub>2</sub> L<sup>-1</sup>] is the DO  
 326 saturation concentration at the temperature T [K], pO<sub>2</sub> = 0.21 [Atm] is the partial  
 327 pressure of oxygen in atmosphere, T<sub>REF</sub> = 298.15 [K] is the reference  
 328 temperature, K<sub>H,O<sub>2</sub></sub>(T) [mg O<sub>2</sub> L<sup>-1</sup> Atm<sup>-3</sup>] is the value of Henry's law solubility  
 329 constant for oxygen at the temperature T, K<sub>H,O<sub>2</sub>,REF</sub> = 40.5 [mg O<sub>2</sub> L<sup>-1</sup> Atm<sup>-1</sup>] is

330 the value of Henry's law solubility constant at  $T_{REF}$  (Sander, 2015),  $k_{La20} = 0.17$   
 331  $\pm 0.07$  [ $h^{-1}$ ] is the volumetric oxygen mass-transfer coefficient evaluated at 20  
 332 °C during abiotic tests, that was previously assessed for the photo-respirometer,  
 333 according to the nonlinear regression method (ASCE, 1993).

334 To compute the average  $OPR_{NET,i}$  and  $OUR_{RESP,i}$  and  $k_{La}$ , nonlinear least  
 335 square regression was performed using the *lsqcurvefit* function with the  
 336 software MATLAB R2019b (Optimisation Toolbox™, The MathWorks, Inc.,  
 337 USA). Raw DO data were fitted to estimate  $OPR_{NET}$  and  $OUR_{RESP}$ . According to  
 338 the methodology adopted in previous studies (Choi et al., 2010; Tang et al.,  
 339 2014; Najm et al., 2017; Rossi et al., 2018), the gross OPR ( $OPR_{GROSS}$ ) was  
 340 then calculated for each  $L_i/D_i$  determination, by subtracting the estimated  
 341  $OUR_{RESP}$  to the  $OPR_{NET}$ , and the result was divided by the TSS of the sample,  
 342 measured according to Standard Methods (APHA, 2017), to obtain specific  
 343 OPRs and OURs ( $sOPR_{GROSS}$  and  $sOUR_{RESP}$ , [ $mg\ O_2\ g\ TSS^{-1}\ h^{-1}$ ]) (Equations  
 344 5-7):

$$OPR_{GROSS,i} = OPR_{NET,i} - OUR_{RESP,i} \text{ (Phases 1, 2, 3) \#(4)}$$

$$sOPR_{GROSS,i} = \frac{OPR_{GROSS,i}}{TSS} \text{ (Light phases, } L_i \text{ } i=1, 2, 3) \#(5)$$

$$sOUR_{RESP,i} = \frac{OUR_{RESP,i}}{TSS} \text{ (Dark phases, } D_i \text{ } i=1, 2, 3) \#(6)$$

345 All models were fitted against  $sOPR_{GROSS}$ .

346

#### 347 **2.4.2. Inhibition models definition and selection**

1 348 The concentration of FA was calculated as a function of temperature, pH and of  
 2  
 3 349 total ammoniacal nitrogen (TAN) concentration (Anthonisen et al., 1976)  
 4  
 5  
 6 350 (Equation 7):  
 7

$$\text{NH}_3 = \text{TAN} \cdot \frac{\text{MW}_{\text{NH}_3}}{\text{AW}_\text{N}} \cdot \frac{10^{\text{pH}}}{\exp\left(\frac{6344}{T}\right) + 10^{\text{pH}}} \quad \#(7)$$

8  
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 11  
 12 351 Where: TAN = NH<sub>3</sub> + NH<sub>4</sub><sup>+</sup> [mg N L<sup>-1</sup>] is the total ammoniacal nitrogen, MW<sub>NH<sub>3</sub></sub>,  
 13  
 14 352 is the molecular weight of ammonia [g NH<sub>3</sub> mol NH<sub>3</sub><sup>-1</sup>], AW<sub>N</sub> is the atomic  
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 17 353 weight of nitrogen [g N mol N<sup>-1</sup>], pH is the pH of the suspension [-], T is the  
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 20 354 temperature of the suspension [K].  
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22 355 Two different inhibition models were chosen to describe the effect of FA on the  
 23  
 24 356 photosynthesis and respiration: the non-competitive inhibition model (Equation  
 25  
 26  
 27 357 8) used to evaluate FA inhibition in anaerobic digestion models (Angelidaki et  
 28  
 29  
 30 358 al., 1993) and a sigmoidal logistic curve, or Hill-type model (Equation 9), used to  
 31  
 32 359 describe dose-response curves (Prinz et al., 2010).  
 33

$$f_{\text{NH}_3} = \frac{\text{sOPR}_{\text{NH}_3}}{\text{sOPR}_{\text{CONTROL}}} = \frac{1}{1 + \frac{\text{NH}_3}{\text{EC}_{50,\text{NH}_3}}} \quad \#(8)$$

$$f_{\text{NH}_3} = \frac{\text{sOPR}_{\text{NH}_3}}{\text{sOPR}_{\text{CONTROL}}} = 1 - \frac{1}{1 + \left(\frac{\text{EC}_{50,\text{NH}_3}}{\text{NH}_3}\right)^N} \quad \#(9)$$

34  
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 41 360 Where: sOPR<sub>NH<sub>3</sub></sub> is the gross sOPR calculated in the reactor subject to FA  
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 43 361 inhibition [mg O<sub>2</sub> g TSS<sup>-1</sup> h<sup>-1</sup>], sOPR<sub>CONTROL</sub> is the gross sOPR calculated in the  
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 46 362 control reactor [mg O<sub>2</sub> g TSS<sup>-1</sup> h<sup>-1</sup>], NH<sub>3</sub> is the calculated initial FA concentration  
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 49 363 [mg NH<sub>3</sub> L<sup>-1</sup>], EC<sub>50,NH<sub>3</sub></sub> is the inhibition parameter of the non-competitive  
 50  
 51 364 inhibition model and the Hill model, representing the FA concentration causing a  
 52  
 53 365 50% inhibition of the photosynthetic activity [mg NH<sub>3</sub> L<sup>-1</sup>], N is the dimensionless  
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 56 366 shape parameter of the Hill model [-].  
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1 367 The models were applied to the dataset and different information criteria were  
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3 368 calculated, to select for the most appropriate model. In particular, the adjusted  
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5 369 R-squared ( $R_{ADJ}^2$ , equation 10) was calculated to evaluate the goodness of fit,  
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8 370 and the model resulting in a lower value of the Akaike Information Criterion  
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11 371 corrected for small samples (cAIC, equation 11, Hurvich and Tsai, 1991), was  
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13 372 considered as the most suitable to represent experimental results:

$$R_{ADJ}^2 = 1 - \left( \frac{n-1}{n-p} \right) * \frac{SSE}{SST} \quad \#(10)$$

$$cAIC = \frac{SSE}{n} * (1+2*p) + 2*p * \left( \frac{p+1}{n-p-1} \right) \quad \#(11)$$

21 373 Where: n is the number of experimental observations, p is the number of model  
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23 374 parameters, SSE is the sum of squared error and SST is the sum of squared  
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26 375 difference between each datum and the mean value of all data.

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33 377 **2.4.3. Statistical analysis**

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36 378 An unbalanced one-way analysis of variance (ANOVA) was applied to the  
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38 379 datasets of experiments performed on microalgae and cyanobacteria, to  
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41 380 evaluate statistically significant differences ( $\alpha = 0.05$ ) between calculated values  
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43 381 of inhibition parameters for monocultures and mixed groups. The software  
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45 382 MATLAB R2019b was used for the analysis (Statistics and Machine Learning  
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47 383 Toolbox™, function *anova1*).

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54 385 **2.5. Definition of free ammonia inhibition scenarios under typical**  
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57 386 **operational conditions**

1 387 To evaluate the need for considering FA inhibition in algae/bacteria modelling,  
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3 388 several scenarios were analysed by calculating theoretical FA concentration  
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5 389 profiles during typical operational days in the pilot plants. Scenarios were  
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8 390 defined by varying: i) the season (spring, summer and autumn), ii) the setpoint  
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10 391 of the pH-control system (7, 8 and 9 for microalgae and 8.5, 9.5 and 10.5 for  
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12 392 cyanobacteria) and iii) the initial TAN concentration in the suspension (5, 10 and  
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14 393 20 mg N-TAN L<sup>-1</sup> for cyanobacteria and 35, 70 and 140 mg N-TAN L<sup>-1</sup> for  
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16 394 microalgae). Typical daily patterns were defined for incident PAR and water  
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18 395 temperature by averaging hourly data collected over a long-term period  
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20 396 (January 2017 - November 2019) (Figure 3). Irradiance data were collected  
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22 397 from the closest weather stations located near each pilot-plant site, and water  
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24 398 temperature was logged by temperature probes in pilot reactors. Daily average  
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26 399 trends for each season are shown in Figure 3. The pH setpoints and TAN  
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28 400 concentrations were chosen according to values measured in pilot plants during  
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30 401 the photo-respirometric campaigns, in order to reflect relevant conditions that  
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32 402 are commonly met in wastewater-treating outdoor PBRs. The measured pH  
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34 403 value in the microalgae-bacteria system was on average 7.0 (maximum pH:  
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36 404 8.5), as a consequence of the high nitrification rates reported (Mantovani et al.,  
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38 405 2019) and compared with the higher values measured in the cyanobacteria-  
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40 406 bacteria systems (average pH = 8.4, maximum pH = 9.5). Likewise, the  
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42 407 measured TAN concentration in the microalgae-bacteria system was on  
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44 408 average = 34 mg N-TAN L<sup>-1</sup> (maximum TAN = 71 mg N-TAN L<sup>-1</sup>). This condition  
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46 409 reflects the high concentration of NH<sub>4</sub><sup>+</sup> in the LFAD (240 ± 55 mg N-NH<sub>4</sub><sup>+</sup> L<sup>-1</sup>, on  
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48 410 average) and the presence of residual NH<sub>4</sub><sup>+</sup> concentration in the suspension,  
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possibly due to low transitory algal activities. On the contrary, cyanobacteria scenarios were characterized by lower TAN concentrations, as the influent was a low strength wastewater stream (agricultural runoff). The biomass was subject to starvation to promote the accumulation of biopolymers and fed with nitrate as nitrogen source, with the ammoniacal nitrogen being almost absent during the entire experimentation (average TAN = 0.3 mg N-TAN L<sup>-1</sup>, maximum TAN = 2.9 mg N-TAN L<sup>-1</sup>).

In addition to modelling FA inhibition, switch functions for light ( $f_I$ , Equation 12, Bernard and Rémond, 2012), temperature ( $f_T$ , Equation 13, Bernard and Rémond, 2012) and pH dependence ( $f_{pH}$ , Equation 14, Ippoliti et al., 2016) were evaluated to describe a more realistic photosynthetic sOPR trend, during typical days. The model describing the overall trend of oxygen production,  $f_{TOT}$  (equation 15), is the product of all the switch functions (Equation 8, and 12 - 14).  $f_{pH}$  is constant, as an ideal pH-control is implemented. No light/solute-gradients were included (0-D model), therefore average irradiance and perfect mixing are considered.

$$f_I = sOPR_{MAX} * \frac{I}{I + \frac{sOPR_{MAX}}{\alpha} * \left(\frac{I}{I_{OPT}} - 1\right)^2} \quad \#(12)$$

$$f_T = \frac{(T - T_{MAX}) * (T - T_{MIN})^2}{(T_{OPT} - T_{MIN}) * ((T_{OPT} - T_{MIN}) * (T - T_{OPT}) - (T_{OPT} - T_{MAX}) * (T_{OPT} + T_{MIN} - 2 * T))} \quad \#(13)$$

$$f_{pH} = \frac{(pH - pH_{MAX}) * (pH - pH_{MIN})}{(pH_{OPT} - pH_{MIN}) * ((pH_{OPT} - pH_{MIN}) * (pH - pH_{OPT}) - (pH_{OPT} - pH_{MAX}) * (pH_{OPT} + pH_{MIN} - 2 * pH))} \quad \#(14)$$

$$f_{TOT} = f_I * f_T * f_{pH} * f_{NH_3} \quad \#(15)$$

Where:  $sOPR_{MAX}$  is the maximum sOPR obtained at the optimal light intensity [mg O<sub>2</sub> g TSS<sup>-1</sup> h<sup>-1</sup>],  $I$  is the actual irradiance [ $\mu E m^{-2} s^{-1}$ ],  $\alpha = 0.45$  is a dimensionless shape parameter [-],  $I_{OPT} = 313 [\mu E m^{-2} s^{-1}]$  is the optimal light

1 430 intensity corresponding to the maximum photosynthetic activity,  $T$  is the actual  
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3 431 temperature of the suspension [ $^{\circ}\text{C}$ ],  $T_{\text{MAX}} = 41.7$  [ $^{\circ}\text{C}$ ] is the maximum  
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6 432 temperature above which the photosynthetic activity stops,  $T_{\text{OPT}} = 28.1$  [ $^{\circ}\text{C}$ ] is  
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8 433 the optimal temperature corresponding to the maximum photosynthetic activity,  
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10 434  $T_{\text{MIN}} = 0.1$  [ $^{\circ}\text{C}$ ] is the minimum temperature below which the activity stops,  $\text{pH}$  is  
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13 435 the actual  $\text{pH}$  value [-],  $\text{pH}_{\text{MAX}} = 11.1$  [-] is the maximum  $\text{pH}$  above which the  
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15 436 activity stops,  $\text{pH}_{\text{OPT}} = 7.5$  [-] is the optimal  $\text{pH}$  corresponding to the maximum  
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18 437 photosynthetic activity,  $\text{pH}_{\text{MIN}} = 0.2$  [-] is the minimum  $\text{pH}$  below which the  
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20 438 activity stops.

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23 439 The parameters characterizing the described switch functions were calibrated  
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25 440 on the microalgae-bacteria consortium during a previous PRT campaign, in  
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27 441 which the effects of a wide range of irradiance, temperature and  $\text{pH}$  values were  
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29 442 assessed (Rossi et al., submitted). Since no experimental data were available  
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31 443 to describe the effects of environmental conditions on cyanobacteria, and the  
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33 444 optimal conditions for cyanobacteria can significantly differ from microalgal  
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35 445 optima (Giannuzzi et al., 2019), the evolution of switch functions was only  
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37 446 modelled for the HRAP case study.

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### 45 46 47 448 **3. RESULTS AND DISCUSSION**

#### 48 49 449 **3.1. Model selection**

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52 450 The experimental  $\text{sOPR}_{\text{GROSS}}$  values were calculated as described in section  
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54 451 2.4 and the dataset obtained from PRTs was used to fit to the non-competitive  
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56 452 inhibition model (equation 9) and to the sigmoidal logistic curve (equation 10).  
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1 453 Model information criteria ( $R_{ADJ}^2$  and cAIC) are shown in Table 2. Both models  
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3 454 can describe the entire photo-respirometric dataset and show a high value of  
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6 455  $R_{ADJ}^2$  (all higher than 0.98). Regarding cAICs values, low differences among  
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8 456 models are observed, however the non-competitive model was preferred  
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11 457 because a similar predicting ability was obtained with one parameter less, thus  
12  
13 458 reducing computational costs and facilitating parameter estimation. The  
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16 459 predicting ability of this model was able to correctly describe the inhibition  
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18 460 process for both mono and mixed cultures. However, the variability of some  
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21 461 estimated parameter was quite high, as proven by the large extension of 95%  
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23 462 confidence bounds. Predictions for cyanobacterial monocultures were more  
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25 463 accurate than for microalgae. The highest variabilities were found for  
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28 464 *Scenedesmus quadricauda* and *Chlorella sorokiniana*. With respect to this  
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30 465 variability, decreasing the measurement noise (e.g. by reducing/eliminating gas-  
31  
32 466 liquid transfer) and/or adjusting the experimental protocol (e.g. by increasing the  
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35 467 number of replications to obtain more robust inhibition data) could be desirable  
36  
37 468 improvements to the proposed methodology. Moreover, the variability of  
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40 469 estimated parameters was generally higher in PRTs performed on mixed  
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42 470 consortia, compared to monocultures (especially for the sample from the  
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45 471 microalgae-bacteria system). This variability can be due to the presence of  
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47 472 other microorganisms in the suspension, possibly contributing to the DO mass  
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49  
50 473 balance (protozoa, heterotrophic bacteria) and constituting an additional  
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52 474 biological noise. In order to improve data reliability, further research is  
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55 475 suggested regarding the pre-treatment of the sample and the possibility of using

1 476 wide spectrum inhibitors/antibiotics to suppress undesired in biological  
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3 477 activities.  
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6 478 **3.2. Effects of free ammonia on pure microalgae and mixed microalgae-**  
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9 479 **bacteria consortia**

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12 480 **Microalgae and cyanobacteria monocultures**

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15 481 Experimental  $sOPR_{NH_3}/sOPR_{CONTROL}$  values quantifying the reduction of  
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17 482 photosynthetic activity due to the exposure to FA for microalgae and  
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19 483 cyanobacteria monocultures are shown in Figure 2A and Figure 2B,  
20  
21 484 respectively, together with the fit of the non-competitive inhibition model. The  
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23 485 photosynthetic activities of all monocultures decreased at increasing FA, as  
24  
25 486 expected due to the inhibitory effects on photosynthesis, and no stimulatory  
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27 487 effects due to ammonia assimilation were observed. FA only affected the  
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32 488  $sOPR_{NET}$ , and the observed  $sOUR_{RESP}$  did not vary significantly regardless of  
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34 489 the FA concentration applied (data not shown), coherently with the findings of  
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37 490 Abeliovich & Azov (1976). However, this behaviour might be species-specific  
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39 491 and should not be generalized. For instance, Markou et al. (2016) measured  
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41 492 decreasing respiration rates at increasing FA in *Arthrospira platensis sp.* and  
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44 493 *Chlorella vulgaris sp.*

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47 494 Regarding results on microalgae monocultures (Table 2), the value of  $EC_{50,NH_3}$   
48  
49 495 for *Chlorella vulgaris* ( $60.9 \text{ mg NH}_3 \text{ L}^{-1}$ ) was close to  $54 \text{ mg NH}_3 \text{ L}^{-1}$  obtained by  
50  
51 496 Markou et al. (2016) under similar conditions. An  $EC_{50,NH_3} = 96.3 \text{ mg NH}_3 \text{ L}^{-1}$   
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53 497 was obtained for *Chlorella Sorokiniana*, coherently with the absence of inhibition  
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55 498 reported by Gutierrez et al. (2016), and with the higher concentrations obtained  
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1 499 for different *Chlorella Sorokiniana* strains by Wang et al. (2019). The inhibition  
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3 500 parameter for *Scenedesmus quadricauda* was 77.7 mg NH<sub>3</sub> L<sup>-1</sup>, but literature  
4  
5 501 values are not available for this species and a direct comparison is not possible.  
6  
7  
8 502 The value obtained for *Scenedesmus obliquus* (52.6 mg NH<sub>3</sub> L<sup>-1</sup>) is instead very  
9  
10 503 similar to that obtained at the same pH by Abeliovich & Azov (1976) (51 mg NH<sub>3</sub>  
11  
12 504 L<sup>-1</sup>). EC<sub>50,NH3</sub> for *Scenedesmus obliquus* (60.9 mg NH<sub>3</sub> L<sup>-1</sup>) are slightly lower,  
13  
14 505 but in the same order of magnitude, than what reported by Azov & Goldman  
15  
16 506 (1982) and Collos & Harrison (2014).  
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21 507 Regarding cyanobacterial monocultures, estimated inhibition parameters and  
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23 508 model fits are reported in Figure 2B. As mentioned, contrarily to other findings  
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25 509 (Collos & Harrison, 2014), the adverse effect of FA on photosynthesis was more  
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27 510 pronounced in cyanobacteria than in microalgae. This is confirmed by  
28  
29 511 comparing the average EC<sub>50,NH3</sub> for the two types of organisms (Table 2). For  
30  
31 512 cyanobacterial monocultures, the average EC<sub>50,NH3</sub> was 14.1 mg NH<sub>3</sub> L<sup>-1</sup>, with  
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33 513 similar values among the different strains adopted. Unluckily, only a few authors  
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35 514 reported inhibition parameters for cyanobacteria, and most available data are  
36  
37 515 for the strain *Arthrospira platensis* sp., typically characterized by a high  
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39 516 resistance to FA (Markou et al., 2014; 2016). However, all the values obtained  
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41 517 for cyanobacterial monocultures and mixed populations fall in the range  
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43 518 indicated by Collos & Harrison (2014) (4.3 - 34.8 mg NH<sub>3</sub> L<sup>-1</sup>).  
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#### 519 Mixed phototrophs-bacteria consortia

520 The value of EC<sub>50,NH3</sub> for the mixed microalgae-bacteria consortium was among  
521 the highest and also the coefficients determined for cyanobacteria mixed

1 522 cultures were higher than those obtained from monocultures data. Due to the  
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3 523 increase in the  $EC_{50,NH_3}$  for both microalgae and cyanobacteria monocultures to  
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5 524 mixed cultures (Table 2), a first interpretation of results would suggest that the  
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8 525 environmental conditions in which the mixed cultures are grown selected  
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10 526 phototrophic strains that are more robust and tolerant to adverse conditions,  
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13 527 including inhibitory compounds. Unravelling this aspect would contribute to a  
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15 528 better understanding of the interactions between microorganisms in wastewater  
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18 529 treatment processes with microalgae-bacteria (e.g., optimizing influent TAN  
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20 530 loading rates, or adopting dynamic pH setpoints based on TAN).  
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23 531 However, the difference in the effect of FA on microalgae mixed and  
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25 532 monocultures could not be explained by ANOVA ( $p$ -value = 0.501), therefore  
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27 533 microalgae monocultures and mixed consortia could be described by an  
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29 534 average value of  $EC_{50,NH_3} = 75 \text{ mg NH}_3 \text{ L}^{-1}$ . On the contrary, cyanobacteria  
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31 535 growing in monocultures and mixed cultures were characterized by statistically  
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33 536 different values of the inhibition parameter. In this case, an average value of  
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36 537  $EC_{50,NH_3} = 14 \text{ mg NH}_3 \text{ L}^{-1}$  was estimated for monocultures, which is significantly  
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39 538 different from the  $EC_{50,NH_3}$  of  $26 \text{ mg NH}_3 \text{ L}^{-1}$  obtained for mixed cultures ( $p$ -value  
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41 539 = 0.029).

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49 541 **3.3. Free ammonia inhibition scenarios in microalgae-based wastewater**  
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51 542 **treatment**

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55 543 The utilisation of parameter estimates for the obtained FA inhibition model can  
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57 544 be particularly useful to evaluate the extent of the inhibition due to FA in several  
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1 545 common operational conditions of the phototrophs-bacteria cultivation  
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3 546 processes. For example, rising TAN concentrations can result from limited  
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6 547 removal rates during start-up periods or due to adverse atmospheric conditions.  
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8 548 Similarly, the pH value can rise during the day, as a result of the photosynthetic  
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11 549 activity. As an example, as explained in paragraph 2.5, the photosynthesis  
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13 550 inhibition model was run using the time-series describing the daily evolution of  
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15 551 FA during typical days in each scenario (i.e. by varying the season, the average  
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18 552 pH and total TAN). To predict the FA response, eq. 9 was used with the  
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20 553 estimated values of inhibition parameters (i.e.,  $EC_{50,NH_3} = 88.4 \text{ mg NH}_3 \text{ L}^{-1}$  for  
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22 554 microalgae-bacteria and  $EC_{50,NH_3} = 26.2 \text{ mg NH}_3 \text{ L}^{-1}$  for cyanobacteria-bacteria).  
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25 555 The evolution of the inhibition function ( $f_{NH_3}$ ) under the identified  
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28 556 environmental/operational conditions is depicted in Figure 4 for microalgae and  
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30 557 cyanobacteria. Although the value of the microalgae-bacteria inhibition  
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32 558 parameter is high, which means a high resistance to FA, severe inhibition levels  
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35 559 can be reached under the worst conditions. In particular, the values of  $f_{NH_3}$   
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37 560 during autumn indicate a photosynthesis inhibition of 30%, while the inhibition  
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40 561 can reach values higher than 40%, during summer times. Temperature  
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42 562 variations seem to have a lower influence on FA production, compared to the  
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44 563 other parameters. When comparing summer and autumn, maximum  $f_{NH_3}$   
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46 564 variations fall within 30% due to temperature variations, while larger effects are  
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49 565 associated to the variation of other parameters (TAN and pH). At low and  
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52 566 average pH,  $f_{NH_3}$  is close to one, regardless of the TAN concentration or the  
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54 567 seasonal condition imposed. A drastic drop in photosynthetic sOPRs occurs  
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57 568 with higher pH values. Similarly, in the cyanobacterial mixed culture, the  
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1 569 inhibition function can result in a limited photosynthetic oxygen evolution during  
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3 570 the day, due to the high pH values and temperatures.  $f_{\text{NH}_3}$  can reach very low  
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5 571 values (up to 75% inhibition), thus depicting severe inhibition, even if the  
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8 572 considered TAN concentrations are seven times lower than those expected in  
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10 573 HRAP scenarios. This clearly indicates the high influence of pH on FA  
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12 574 generation, confirming that pH should be strictly controlled in wastewater  
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14 575 treatment PBRs, to prevent reductions in the photosynthetic oxygenation by  
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16 576 phototrophs.  
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21 577 For the microalgae-bacteria consortium case study, the trends for  $f_{\text{TOT}}$  and for  
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23 578 the functions expressing the dependence of photosynthesis on FA and  
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25 579 environmental conditions were also constructed. Switch functions are shown in  
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27 580 Figure 5, for a TAN concentration of  $60 \text{ mg N-TAN L}^{-1}$  and for different seasons  
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29 581 and pH conditions. Among the studied variables, temperature is the one most  
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31 582 directly affecting photosynthetic sOPR: the value assumed by the  $f_{\text{T}}$  switch  
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33 583 function is always the lowest (excluding the irradiance switch function, which is  
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35 584 obviously zero during the night). This is particularly evident during autumn  
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37 585 (Figure 5B and Figure 5D), when temperature is lower than the optimum. During  
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39 586 summer, temperature approaches the optimal value, resulting in  $f_{\text{T}}$  close to one  
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41 587 for almost all the daytime. Although summer temperatures are close to optimal  
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43 588 values resulting in higher  $f_{\text{T}}$  values, the increase in temperature also favours the  
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45 589 FA formation, what inhibits photosynthesis (Figure 5 A and Figure 5B). The  
46  
47 590 combined effects of temperature and pH are evident when the pH is 9 (Figure  
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49 591 5C and Figure 5D):  $f_{\text{TOT}}$  reaches approximately 0.6-0.65 during summer and is  
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51 592 reduced to approximately 0.5 during autumn. Regarding  $f_{\text{NH}_3}$ , it has comparable  
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1 593 or lower values than the  $f_T$  during summer, indicating that under these  
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3 594 conditions the inhibition due to FA is the most relevant limitation occurring in the  
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5 595 reactor. It is also important to notice that the pH value can be responsible for a  
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7 596 reduction of the photosynthetic activity in itself. This reduction is negligible at pH  
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9 597 7, but at a pH of 9 causes a reduction of approximately 15% of the sOPR.  
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## 16 599 **CONCLUSIONS**

20 600 Microalgae and cyanobacteria were differently inhibited by FA, with microalgae  
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22 601 showing higher resistance than cyanobacteria. The simulation of different  
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24 602 weather/operational conditions showed that FA can drastically impact photo-  
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26 603 oxygenation processes in algae-bacteria wastewater treatment systems. The  
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28 604 results suggest that considering FA inhibition in existing mathematical models  
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30 605 describing microalgae/cyanobacteria-based wastewater treatment processes  
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32 606 could lead to more reliable predictions and to a more rational design of  
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34 607 treatment units. In addition, the proposed procedure can be used to generate a  
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36 608 dataset of  $EC_{50}$  values and theoretical dose-response curves, for different  
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38 609 chemicals known to be inhibitory for the algal-bacterial biomass (e.g.  
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40 610 herbicides/pesticides or organic compounds).  
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1 769 **Figure captions**

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3 770 **Figure 1.** Scheme of the experimental setup for photo-respirometric test. Legend: 1 = Mixing  
4 771 and signal communication unit, 2 = Glass bottles, 3 = DO, temperature and pH probes, 4 =  
5 772 Acid/base dosage pumps, 5 = Air/gas pumps, 6 = Normally closed electro-valves, 7 = Gas  
6 773 cylinder (CO<sub>2</sub>/O<sub>2</sub>/N<sub>2</sub>), 8 = Acid/base solutions, 9 = Industrial grade PC, 10 = light source.

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8 774 **Figure 2.** Effects of free ammonia inhibition on microalgae and cyanobacteria: reduction of f<sub>NH<sub>3</sub></sub>,  
9 775 non-competitive inhibition model fit and estimated model parameters (A: Microalgae, B:  
10 776 Cyanobacteria). Samples abbreviations: M<sub>1</sub> = *Chlorella vulgaris* sp., M<sub>2</sub> = *Scenedesmus*  
11 777 *quadricauda* sp., M<sub>3</sub> = *Chlorella sorokiniana* sp., M<sub>4</sub> = *Scenedesmus obliquus* sp., MB = Sample  
12 778 from the HRAP; C1 = *Synechocystis* sp., C2 = *Synechococcus* sp., C3 = *Leptolyngbya* sp., CB<sub>1</sub>  
13 779 = Sample from the semi-closed PBR<sub>1</sub>, CB<sub>2</sub> = Sample from the semi-closed PBR<sub>2</sub>, CB<sub>3</sub> = Sample  
14 780 from the semi-closed PBR<sub>3</sub>. Shaded areas and error bars represent 95% confidence intervals.

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16 781 **Figure 3.** Typical daily variations of water temperature and irradiance for different seasons (A =  
17 782 water temperature in the HRAP, B = water temperature in semi-closed PBRs, C = irradiance  
18 783 data for the HRAP, dataset: January 2017 - November 2019).

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20 784 **Figure 4.** Simulated evolution of the inhibition function (f<sub>NH<sub>3</sub></sub>) under typical daily variations for  
21 785 different seasons, pH values and TAN concentrations. Inhibition functions are calculated  
22 786 considering microalgae (A - D) and cyanobacteria (E - H) as dominant species in mixed  
23 787 phototrophs-bacteria consortia.

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25 788 **Figure 5.** Simulated evolution of switch functions in the HRAP at 60 mg N-TAN L<sup>-1</sup> for different  
26 789 seasons and pH values (A = summer, low pH, B = autumn, low pH, C = summer, high pH, D =  
27 790 autumn, high pH).

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31 792 **Table captions**

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33 793 **Table 1.** Free ammonia inhibition tests performed and conditions applied. Temperature and  
34 794 irradiance data are reported as mean ± standard deviation.

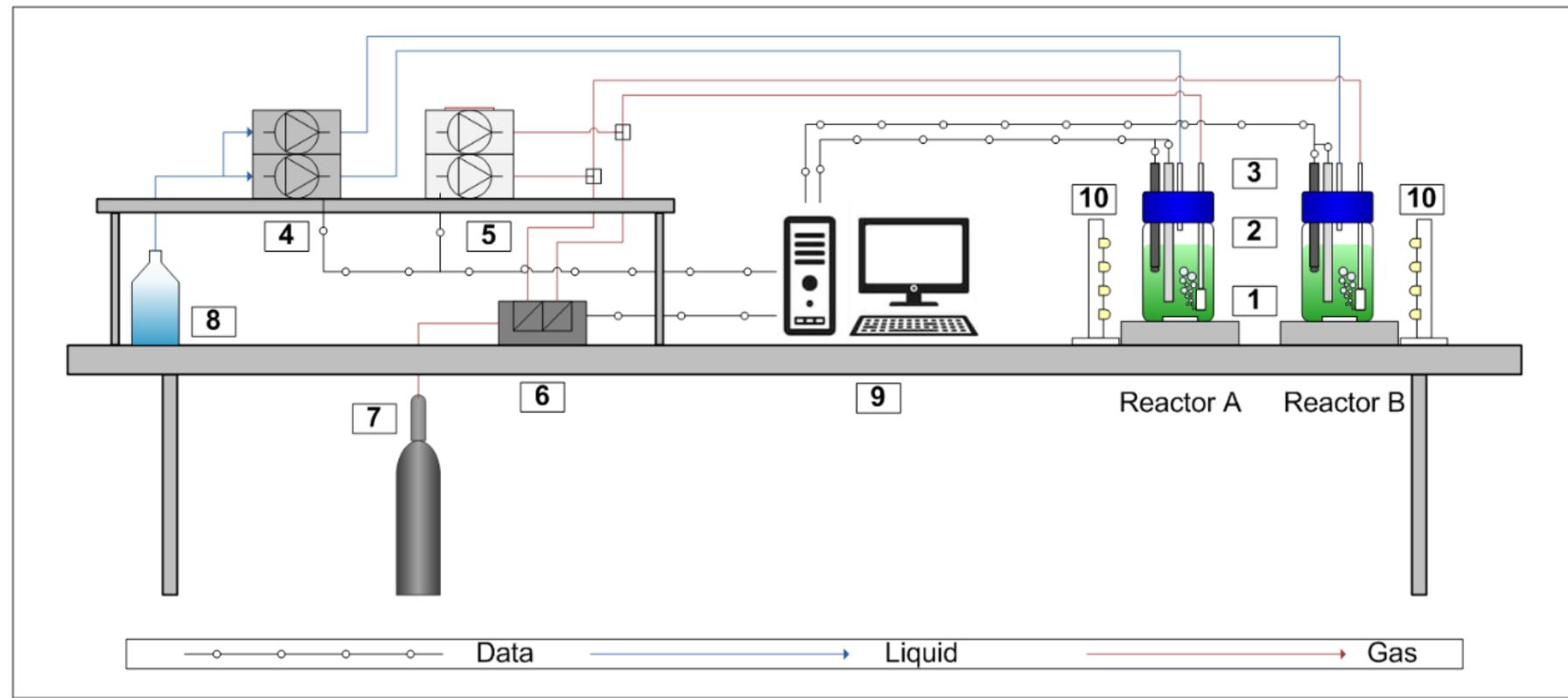
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36 795 **Table 2.** Selection criteria and estimated parameters for selected free ammonia inhibition  
37 796 models. 95% confidence intervals on estimated parameters are reported in square brackets.  
38 797 Averaged data are reported as mean ± standard deviation.

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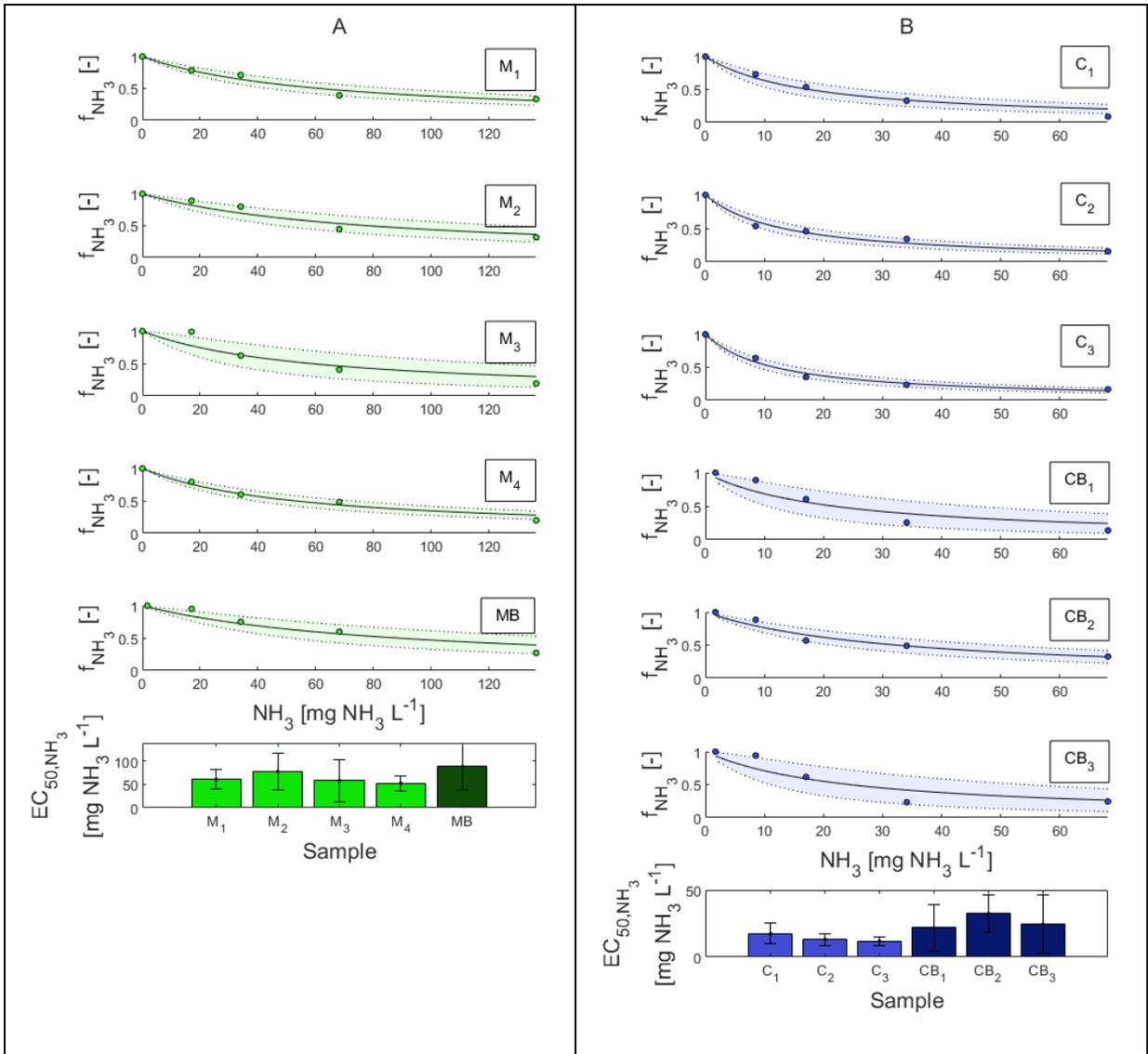
799 **Figure 6.**



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802 **Figure 7.**

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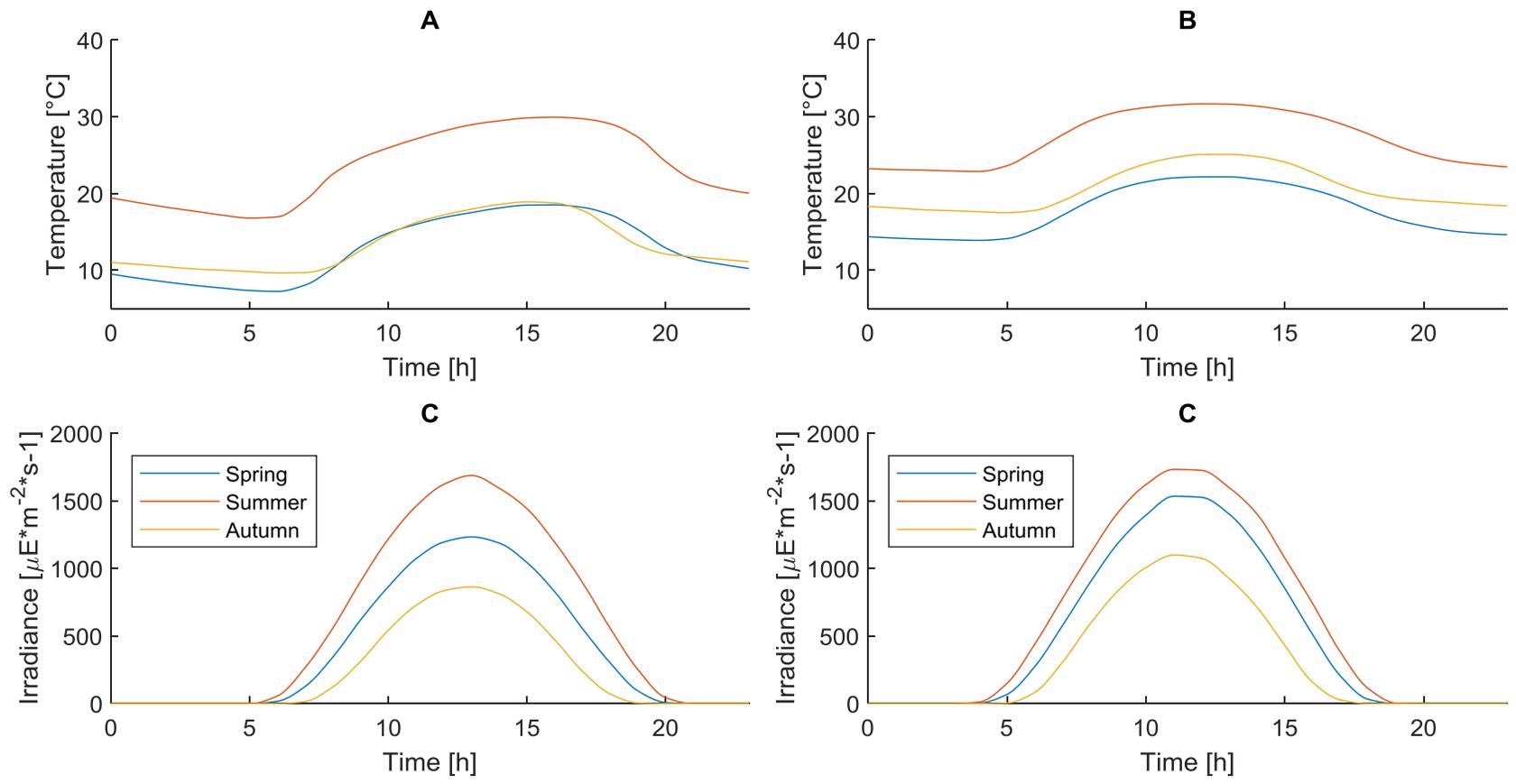


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805 **Figure 3**

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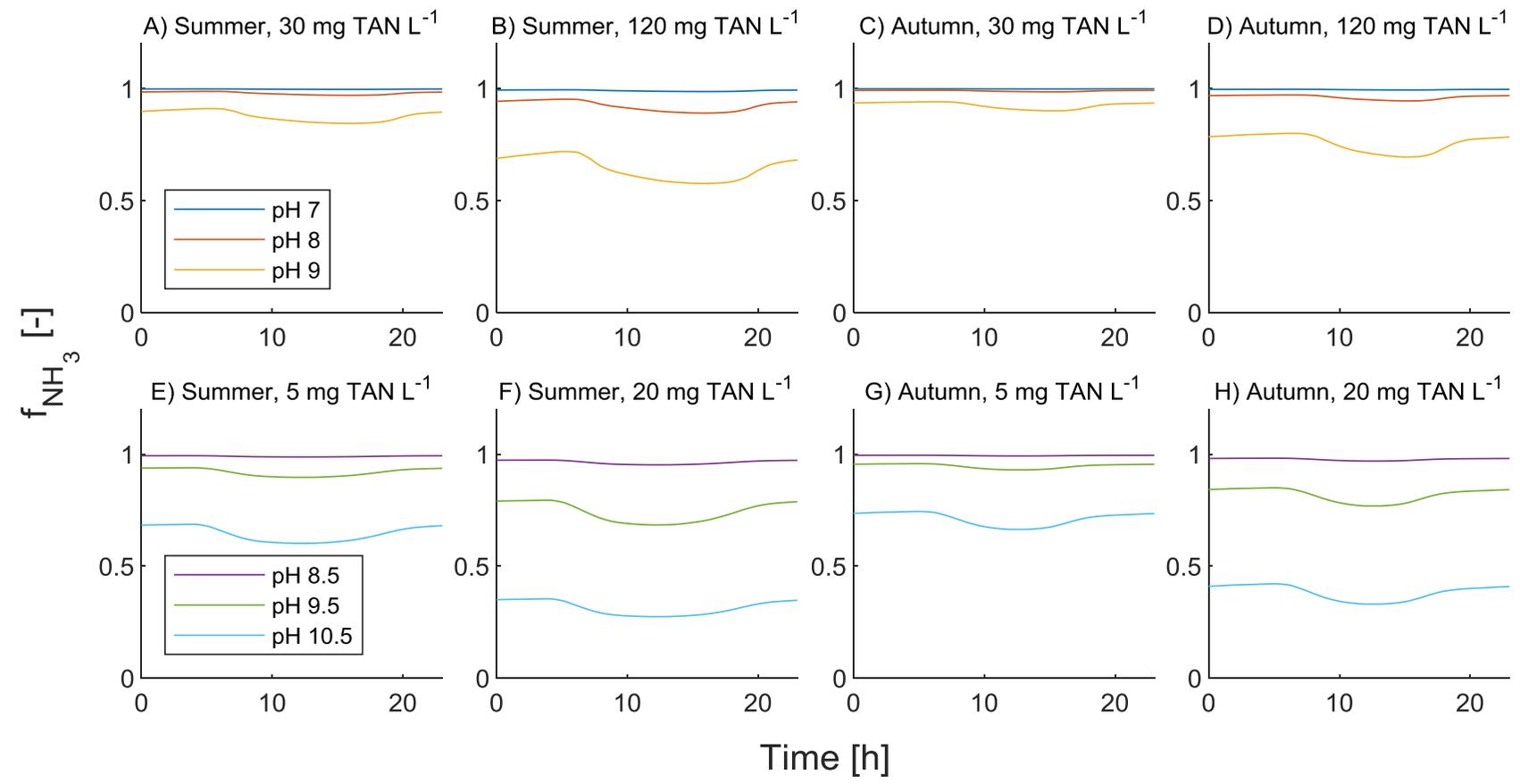


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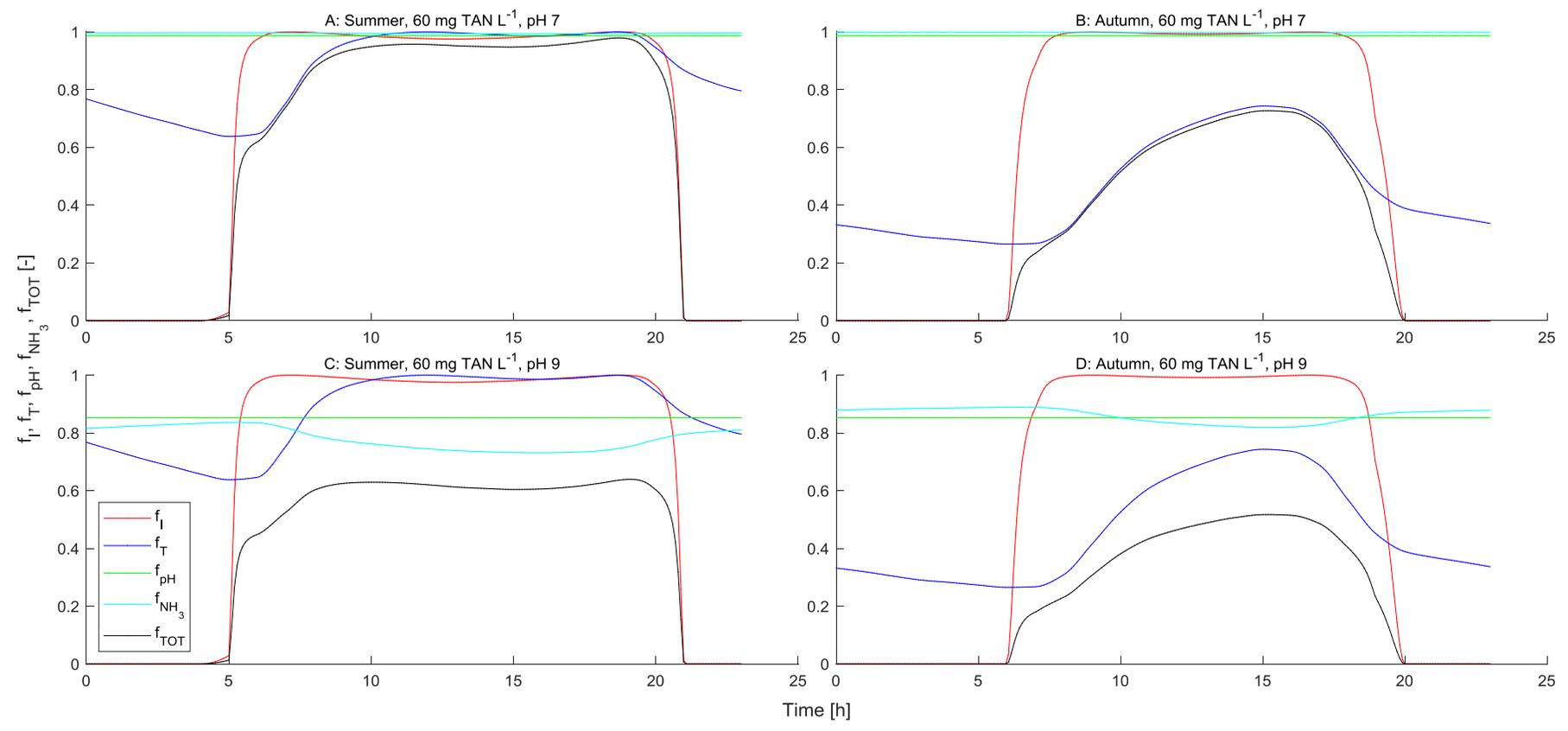


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812 **Figure 5.**



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814 **Table 3.**

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Test ID	Cultivation system (volume)	Type of culture and dominant phototrophs	Species	Growth substrate (N-source)	Free ammonia concentration	Temperature	Irradiance
-	-	-	-	-	mg NH <sub>3</sub> L <sup>-1</sup>	°C	μE m <sup>-2</sup> s <sup>-1</sup>
MB	HRAP (1.2 m <sup>3</sup> )	Phototrophs-bacteria	<i>Chlorella/Scenedesmus sp.</i>	Anaerobic digestate (NH <sub>4</sub> )	17, 34, 68, 134	20.3 ± 0.2	108 ± 16
CB <sub>1</sub>	Semi-closed PBRs (11.7 m <sup>3</sup> )	Phototrophs-bacteria	<i>Synechocystis sp.</i> , <i>Synechococcus sp.</i>	Agricultural runoff (NO <sub>3</sub> )	8.5, 17, 34, 68	21.5 ± 0.3	116 ± 23
CB <sub>2</sub>	Semi-closed PBRs (11.7 m <sup>3</sup> )	Phototrophs-bacteria	<i>Synechocystis sp.</i> , <i>Synechococcus sp.</i>	Agricultural runoff (NO <sub>3</sub> )	8.5, 17, 34, 68	21.6 ± 0.3	116 ± 23
CB <sub>3</sub>	Semi-closed PBRs (11.7 m <sup>3</sup> )	Phototrophs-bacteria	<i>Synechocystis sp.</i> , <i>Synechococcus sp.</i>	Agricultural runoff (NO <sub>3</sub> )	8.5, 17, 34, 68	21.8 ± 0.4	116 ± 23
M <sub>1</sub>	Lab-scale PBR (1 L)	Green algae monoculture	<i>Chlorella vulgaris</i>	MBBM (NO <sub>3</sub> )	17, 34, 68, 134	19.6 ± 0.1	108 ± 16
M <sub>2</sub>	Lab-scale PBR (1 L)	Green algae monoculture	<i>Scenedesmus quadricauda</i>	MBBM (NO <sub>3</sub> )	17, 34, 68, 134	19.7 ± 0.1	108 ± 16
M <sub>3</sub>	Lab-scale PBR (1 L)	Green algae monoculture	<i>Chlorella sorokiniana</i>	MBBM (NO <sub>3</sub> )	17, 34, 68, 134	19.9 ± 0.1	108 ± 16
M <sub>4</sub>	Lab-scale PBR (1 L)	Green algae monoculture	<i>Scenedesmus obliquus</i>	MBBM (NO <sub>3</sub> )	17, 34, 68, 134	20 ± 0.0	108 ± 16
C <sub>1</sub>	Lab-scale PBR (1 L)	Cyanobacteria monoculture	<i>Synechocystis sp.</i>	BG11 (NO <sub>3</sub> )	8.5, 17, 34, 68	20.0 ± 0.4	116 ± 23
C <sub>2</sub>	Lab-scale PBR (1 L)	Cyanobacteria monoculture	<i>Synechococcus sp.</i>	BG11 (NO <sub>3</sub> )	8.5, 17, 34, 68	20.6 ± 0.3	116 ± 23
C <sub>3</sub>	Lab-scale PBR (1 L)	Cyanobacteria monoculture	<i>Leptolyngbia sp.</i>	BG11 (NO <sub>3</sub> )	8.5, 17, 34, 68	20.8 ± 0.3	116 ± 23

816 **Table 4.**

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<b>Cyanobacteria</b>						
<b>Test ID</b>	<b>Model 1 (Non-competitive inhibition)</b>			<b>Model 2 (Sigmoidal logistic function)</b>		
	<b>cAIC</b>	<b>R<sub>ADJ</sub><sup>2</sup></b>	<b>Estimated parameters</b>	<b>cAIC</b>	<b>R<sub>ADJ</sub><sup>2</sup></b>	<b>Estimated parameters</b>
C <sub>1</sub>	-16.7	0.9478	EC <sub>50,NH3</sub> = 17.5 mg NH <sub>3</sub> L <sup>-1</sup> [9.9, 25.0]	-17.8	0.9820	EC <sub>50,NH3</sub> = 18.5 mg NH <sub>3</sub> L <sup>-1</sup> [14.7, 22.2], N = 1.43 [0.96, 1.89]
C <sub>2</sub>	-19.7	0.9990	EC <sub>50,NH3</sub> = 13.1 mg NH <sub>3</sub> L <sup>-1</sup> [8.8, 17.5]	-16.1	0.9993	EC <sub>50,NH3</sub> = 11.8 mg NH <sub>3</sub> L <sup>-1</sup> [6.1, 17.5], N = 0.78 [0.34, 1.21]
C <sub>3</sub>	-21.0	0.9982	EC <sub>50,NH3</sub> = 11.7 mg NH <sub>3</sub> L <sup>-1</sup> [8.2, 15.2]	-15.3	0.9981	EC <sub>50,NH3</sub> = 12.3 mg NH <sub>3</sub> L <sup>-1</sup> [7.9, 16.6], N = 1.15 [0.58, 1.71]
<i>Avg</i>	-	-	<i>EC<sub>50,NH3</sub> = 14.1 ± 3.0 mg NH<sub>3</sub> L<sup>-1</sup></i>	-	-	<i>EC<sub>50,NH3</sub> = 14.2 ± 3.7 mg NH<sub>3</sub> L<sup>-1</sup>, N = 1.12 ± 0.32</i>
CB <sub>1</sub>	-10.3	0.9972	EC <sub>50,NH3</sub> = 21.8 mg NH <sub>3</sub> L <sup>-1</sup> [4.4, 39.2]	-17.6	0.9998	EC <sub>50,NH3</sub> = 21.4 mg NH <sub>3</sub> L <sup>-1</sup> [17.8, 25.0], N = 2.01 [1.37, 2.67]
CB <sub>2</sub>	-16.6	0.9983	EC <sub>50,NH3</sub> = 32.4 mg NH <sub>3</sub> L <sup>-1</sup> [18.4, 46.5]	-10.4	0.9979	EC <sub>50,NH3</sub> = 31.6 mg NH <sub>3</sub> L <sup>-1</sup> [15.3, 47.9], N = 1.12 [0.33, 1.91]
CB <sub>3</sub>	-9.3	0.9832	EC <sub>50,NH3</sub> = 24.4 mg NH <sub>3</sub> L <sup>-1</sup> [2.8, 46.1]	-7.4	0.9903	EC <sub>50,NH3</sub> = 22.7 mg NH <sub>3</sub> L <sup>-1</sup> [11.8, 33.6], N = 1.92 [0.23, 3.61]
<i>Avg</i>	-	-	<i>EC<sub>50,NH3</sub> = 26.2 ± 5.5 mg NH<sub>3</sub> L<sup>-1</sup></i>	-	-	<i>EC<sub>50,NH3</sub> = 25.2 ± 5.6 mg NH<sub>3</sub> L<sup>-1</sup>, N = 1.68 ± 0.49</i>
<b>Microalgae</b>						
<b>Test ID</b>	<b>Model 1 (Non-competitive inhibition)</b>			<b>Model 2 (Sigmoidal logistic function)</b>		
	<b>cAIC</b>	<b>R<sub>ADJ</sub><sup>2</sup></b>	<b>Estimated parameters</b>	<b>cAIC</b>	<b>R<sub>ADJ</sub><sup>2</sup></b>	<b>Estimated Parameters</b>
M <sub>1</sub>	-18.8	0.9996	EC <sub>50,NH3</sub> = 60.9 mg NH <sub>3</sub> L <sup>-1</sup> [39.7, 82.1]	-12.4	0.9995	EC <sub>50,NH3</sub> = 60.3 mg NH <sub>3</sub> L <sup>-1</sup> [34.5, 86.1], N = 1.08 [0.42, 1.75]
M <sub>2</sub>	-15.1	0.9996	EC <sub>50,NH3</sub> = 77.7 mg NH <sub>3</sub> L <sup>-1</sup> [37.7, 117.7]	-13.6	0.9998	EC <sub>50,NH3</sub> = 71.1 mg NH <sub>3</sub> L <sup>-1</sup> [49.0, 93.1], N = 1.49 [0.73, 2.24]
M <sub>3</sub>	-12.7	0.9960	EC <sub>50,NH3</sub> = 96.3 mg NH <sub>3</sub> L <sup>-1</sup> [31.2, 161.3]	-9.5	0.9972	EC <sub>50,NH3</sub> = 54.2 mg NH <sub>3</sub> L <sup>-1</sup> [34.2, 74.2], N = 1.80 [0.62, 2.97]
M <sub>4</sub>	-19.6	0.9993	EC <sub>50,NH3</sub> = 52.6 mg NH <sub>3</sub> L <sup>-1</sup> [26.1, 66.4]	-15.3	0.9994	EC <sub>50,NH3</sub> = 52.4 mg NH <sub>3</sub> L <sup>-1</sup> [37.1, 67.7], N = 1.20 [0.68, 1.73]
MB	-14.2	0.9994	EC <sub>50,NH3</sub> = 88.4 mg NH <sub>3</sub> L <sup>-1</sup> [37.9, 138.9]	-16.6	0.9999	EC <sub>50,NH3</sub> = 78.7 mg NH <sub>3</sub> L <sup>-1</sup> [61.5, 95.9], N = 1.63 [1.01, 2.26]
<i>Avg</i>	-	-	<i>EC<sub>50,NH3</sub> = 75.2 ± 18.3 mg NH<sub>3</sub> L<sup>-1</sup></i>	-	-	<i>EC<sub>50,NH3</sub> = 63.3 ± 11.3 mg NH<sub>3</sub> L<sup>-1</sup>, N = 1.44 ± 0.30</i>

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