

1 **New insight into colonies of *Microcystis* (Cyanobacteria) as multi-specific floating**  
2 **biofilms**

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28 **ABSTRACT**

29 The ability to form biofilms is a functional trait shared by many bacterial species. Biofilms provide  
30 bacteria a sheltered environment where the nutrients and oxygen gradients create a heterogeneous matrix  
31 and promote cells to differentiate their metabolism and functions according to the position they occupy  
32 inside the matrix. Species of the *Microcystis* genus are among the most common bloom-forming  
33 cyanobacteria. They are unicellular microorganisms able to form colonies and to reach high biomass  
34 during blooms in lakes, reservoirs and estuaries worldwide. Colonial lifestyle provides several  
35 advantages under stressing conditions, including adaptation to different light intensities, protection from  
36 toxic substances and grazing, while allowing them to grow when the nutrient supply is low. Although  
37 the biology, ecology and colony formation have been extensively recognized in *Microcystis* spp., the  
38 analysis of the progression from unicellular to multicellular phases in this cyanobacterium have been  
39 always addressed as individual phenotypic plasticity and rarely as a multi-specific community of  
40 interrelated microorganisms. Here, we re-interpreted the evidence coming from different studies about  
41 the *Microcystis* lifestyle and propose a new way to analyze the available information about this  
42 cyanobacterial group. We specifically address the characteristics shared by bacterial biofilms and  
43 *Microcystis* colonies and suggest that the morphological changes from single cells to colonies are due  
44 to a cascade of events leading to the formation of a multi-specific biofilm. Studying the formation of  
45 colonies using this framework would help to better understand the life cycle of *Microcystis*, its  
46 functional relationship with the associated microbiome and the factors triggering microcystin  
47 production, helping to design strategies for prevention and control of the blooms caused by these  
48 organisms. Taking into account the biology and the ecological strategies of *Microcystis*, a conceptual  
49 model of emergence and decay of these floating multi-specific biofilms is proposed.

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51 **Keywords:** multi-specific, biofilm, *Microcystis*, colonies, mucilage, EPS, holobiont

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## 53 INTRODUCTION

54 Bacterial biofilms are defined as aggregates of microbial cells surrounded by a self-produced  
55 polymer matrix that can be composed by a single (mono-specific) or several species (multi-specific)  
56 living in a collaborative way (Flemming et al., 2016). Biofilm growth of microorganisms was first  
57 defined in medical microbiology, when it was also demonstrated that biofilm-embedded organisms have  
58 an increased antimicrobial resistance compared to those growing as planktonic bacteria (Nickel et al.,  
59 1985). The classic conceptual model of biofilm formation involves motile planktonic cells that become  
60 attached to a surface in response to a variety of environmental signals, such as exposure to subinhibitory  
61 concentrations of antibiotics as demonstrated in *P. aeruginosa* and *Escherichia coli* (Hoffman et al.,  
62 2005). Attached cells produce a hydrated matrix of extracellular polysaccharides (EPS), extracellular  
63 DNA, proteins and lipids (Flemming and Wingender, 2010), changing their structure and functional  
64 relationships. There are several key features that distinguish the cells in a biofilm from the planktonic  
65 lifestyle. Although biofilm cells encounter stronger gradients of nutrients and waste products than during  
66 planktonic life (Stewart and Franklin, 2008), they are embedded in a more controllable environment.

67 In aquatic cyanobacteria, despite the increasing amount of information regarding their ecology, the  
68 biofilm concept is generally associated with benthic species, which form cyanobacterial mats in several  
69 aquatic ecosystems (Stal, 2012). Among the planktonic groups we will focus on *Microcystis* spp., a  
70 complex of cyanobacteria from the Chroococcales order that live in freshwater and brackish waters.  
71 They are Gram-negative bacteria that can be found as single cells or in colonies that float near the  
72 surface, reaching colony sizes that can be detected by naked eye. *Microcystis* blooms in eutrophic  
73 ecosystems and generally a size spectrum can be found, ranging from ca. 4  $\mu\text{m}$  (single cells) to hundreds  
74 of microns (large colonies) (Figure 1) (Reynolds et al., 1981). It has been described that *Microcystis*  
75 blooms are composed by populations able to produce secondary metabolites called microcystins, which  
76 are toxic to animals and humans (toxins), and by non-toxic populations (Vezie C, et al., 1998).  
77 Interestingly, some studies have shown that high temperature (between 25 and 30  $^{\circ}\text{C}$ ) promotes the  
78 growth of toxic *Microcystis* (Davis et al., 2009), while non-toxic populations seem to have less tolerance  
79 to extreme environmental conditions such as nutrient depletion or low light (Van de Waal et al., 2011).  
80 Therefore, it is very likely that under current climate warming and worldwide eutrophication scenarios,

81 a dominance of cyanobacterial blooms containing a higher percentage of toxic *Microcystis* cells will  
82 occur. This makes it very relevant to understand the biology and ecology of the whole *Microcystis*  
83 community (toxic and non-toxic), with special attention to the toxic component.

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98 **Figure 1. *Microcystis* colonies.** Large *Microcystis* colonies from Uruguay river (Uruguay-South America)  
99 observable at naked eye. A concentrated sample is shown at left.

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101 Current vision of organism's evolution is increasingly incorporating the concept of holobiont, which  
102 recognizes the widespread occurrence of host-associated microbiomes and makes emphasis in the  
103 multispecies nature of host-microbiome assemblage (Bordenstein and Theis, 2015). In *Microcystis*, the  
104 presence of an external layer of mucilage constituted by extracellular polysaccharides that is heavily  
105 populated by other bacterial species has been early discovered and is increasingly studied. The complex  
106 microbial structure generated in the mucilaginous colonies and its role in the survival and fitness of the  
107 cyanobacterium has started to be studied (Pérez-Carrascal et al., 2021; Schmidt et al., 2020) and points  
108 towards a holobiont lifestyle. In this context, studying the role of the heterotrophic counterpart of the

109 holobiont in *Microcystis* fitness and adaptation to different environmental conditions could provide the  
110 key for their global success.

111 The presence of an extracellular matrix with a microbiome, the evidence for a quorum sensing  
112 signaling system, the different metabolic capacities exhibited by single cells vs. colonies and field  
113 observations about *Microcystis* life cycle prompted us to propose that the colonies of these organisms  
114 are floating, multispecific biofilms. Here, we present a review of the literature focused on the main  
115 characteristics of *Microcystis* colonies resembling those of bacterial biofilms.

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### 117 **MAIN CHARACTERISTICS OF BACTERIAL BIOFILMS**

118 In a biofilm, the cells become embedded within a slimy extracellular matrix that is mainly composed  
119 of extracellular polymeric substances and is produced by the cells within the biofilm. This polymeric  
120 matrix is composed of EPS, proteins, lipids and DNA (Costerton et al., 1987; Stoodley et al., 2002).  
121 When growing as biofilms, bacteria are usually attached to surfaces in high cell density accumulation,  
122 where diffusion and physicochemical heterogeneity are linked to large physiological fluctuations  
123 (Stewart and Franklin, 2008). Bacterial biofilms can also be considered as hydrogels, which are complex  
124 polymers containing many times their dry weight in water.

125 Biofilms are not just bacterial slime layers but biological ecosystems that work as a functionally  
126 coordinated community. The complex network of interactions within the biofilm influences the growth  
127 rate and metabolic activity of the participating bacteria, which are affected by the differences in nutrients  
128 and oxygen availability within biofilms. The formation and dispersal of biofilms are tightly regulated at  
129 the genetic level and triggered by environmental signals (Fazli et al., 2014).

130 The mechanism involved in biofilm formation and regulation is known to be the quorum sensing  
131 (QS), a chemical language used for intercellular communication, which is based on small, self-generated  
132 signal molecules called autoinducers that are produced in low concentration by the cells. When enough  
133 bacteria are present the concentration of autoinducers reaches a threshold level and bacteria are able to  
134 sense their critical mass, repressing or activating target genes (De Kievit and Iglewski, 2000). It has  
135 been reported that the genes that are controlled by QS can build up to 10 % of the bacterial genome  
136 (Wagner et al., 2003). The functional differences between free-living cells and biofilms have been

137 extensively demonstrated in pathogenic bacteria, such as *Pseudomonas aeruginosa*, where QS regulates  
138 the expression of genes involved in lectins, EPS and exotoxin, among others (Kariminik et al., 2017).

139 Bacterial biofilms can also exist in the air-liquid interface, forming floating biofilms or pellicles.  
140 This interface provides access to oxygen and other gases from the air and nutrients from the liquid phase  
141 through opposing gradients (Armitano et al., 2014). In pellicle-forming bacteria, such as *B. subtilis* and  
142 *P. aeruginosa*, flagellar motility is required for wild-type pellicle maturation dynamics and is an  
143 important trait influencing whether or not cells can form pellicles (Hölscher et al., 2015). This is relevant  
144 in the case of heterotrophic bacteria lacking gas vesicles for flotation and need to reach the surface.  
145 When growing in culture under static conditions, the cells tend to accumulate at the bottom of the flask,  
146 meaning that floating at the surface would be only possible through the generation of positive buoyancy.  
147 Some taxa, such as *Gluconacetobacter* spp. are buoyant because they trap CO<sub>2</sub> bubbles generated during  
148 the respiration process. Other bacteria can secrete surface-active agents (e.g., surfactants) or synthesize  
149 a polysaccharides-rich matrix that avoid the mixing with the liquid medium (Angelini et al., 2009;  
150 Koizumi et al., 2008). In the case of *Microcystis*, the position relative to the surface can be achieved  
151 thanks to the presence of gas vesicles aggregations or aerotopes in the cytoplasm (Šmarda and Maršálek,  
152 2008), which allow them to regulate their vertical position in the water column and to form the colony  
153 in a suitable position receiving the right amount of light, oxygen and CO<sub>2</sub>. The EPS matrix of bacteria  
154 that are able to form pellicles are usually composed by glucose, galactose, rhamnose, mannose or  
155 cellulose (for a review see Armitano et al. 2014), which are also typical components of the extracellular  
156 matrix of *Microcystis* (Lei et al., 2007; Li et al., 2009). Thus, *Microcystis* colonies have more than one  
157 trait contributing to float near the surface.

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## 159 **CHARACTERISTICS OF *MICROCYSTIS* COLONIES RESEMBLING THOSE OF** 160 **BACTERIAL BIOFILMS**

161 Table 1 summarizes some of the main biofilm-like characteristics exhibited by *Microcystis* colonies.

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**Table 1.** Traits of bacterial biofilms exhibited by colonies of *Microcystis* spp.

Biofilm trait	In <i>Microcystis</i> colonies	References
Presence and composition of EPS matrix	Yes (glucose, xylose, galactose, fucose, rhamnose, arabinose)	Otsuka <i>et al.</i> , 2000; Lei <i>et al.</i> , 2007; Li <i>et al.</i> , 2009; Bi <i>et al.</i> , 2013; Li <i>et al.</i> , 2013b; Wang <i>et al.</i> , 2013; Liu, Huang, and Qin, 2018
Heterogeneity of the matrix	Yes	Sampognaro <i>et al.</i> , 2020
Buoyancy by EPS formation	Yes	Wang <i>et al.</i> , 2011; Xiao <i>et al.</i> , 2018; Chen <i>et al.</i> , 2019
Buoyancy by other mechanisms (gas vesicles)	Yes	Thomas and Walsby, 1985; Deacon and Walsby, 1990; Mlouka <i>et al.</i> , 2004
Physiological changes between single cells and multicellular stage	Yes	Gan <i>et al.</i> , 2012; Deus <i>et al.</i> , 2020; Harke and Gobler, 2013
Presence of a quorum sensing mechanism	Yes	Zhai <i>et al.</i> , 2012
Increased resistance to antimicrobial or toxic compounds	Yes	Bi <i>et al.</i> , 2013; Tan <i>et al.</i> , 2018

185 **Presence and composition of extracellular polysaccharides in colonies: its effect on *Microcystis***  
186 **morphology and ecology**

187 Cyanobacteria of the *Microcystis* genus exhibit high phenotypic plasticity. They can exist as single  
188 or paired cells when growing fast under axenic conditions in laboratory cultures, but in nature are usually  
189 found as colonies (Xiao *et al.*, 2017) displaying different morphologies, including irregular, sponge-  
190 like, spherical and elongated (Komárek and Komárková, 2002). These colonies consist of groups of  
191 cells stick to each other by a mucus envelope mainly composed of EPS (Hall-Stoodley *et al.*, 2004)

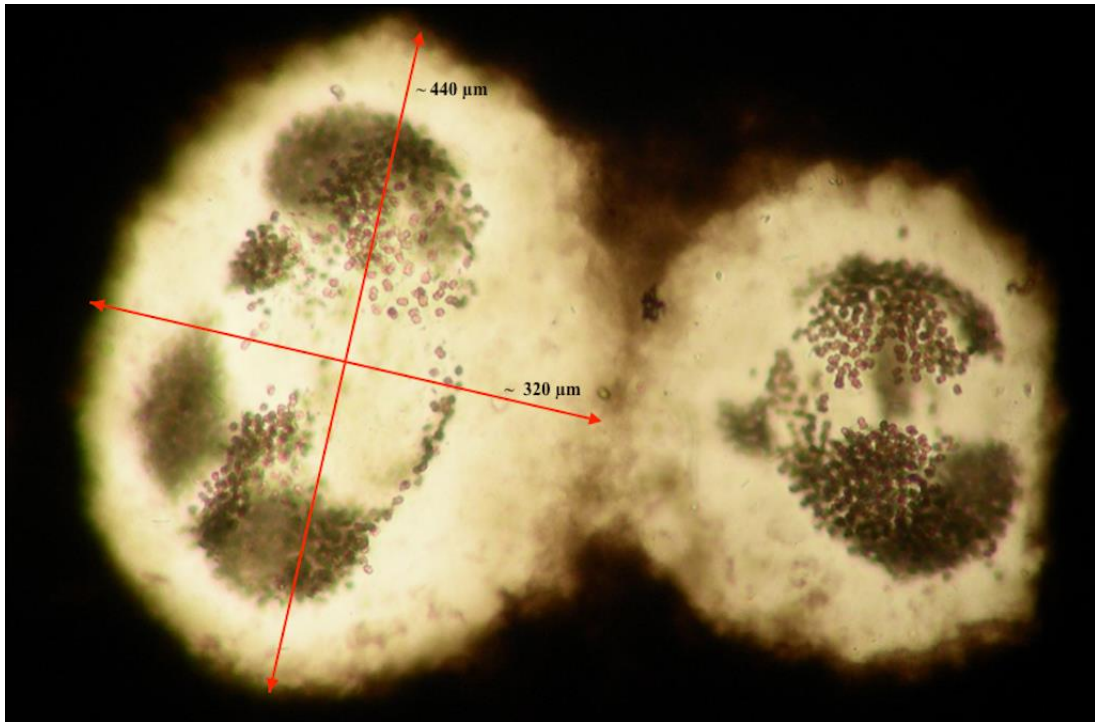
192 known as mucilage. This region that surrounds the cells creates a microenvironment known as the  
193 phycosphere (Bell and Mitchell, 1972), where complex ecological interactions between phytoplankton  
194 and bacteria occur (Jiang et al., 2007; Seymour et al., 2017).

195 Colony formation in *Microcystis* has been attributed to i) cell division: after binary fission cells  
196 remain attached and daughter cells become enveloped in a layer of mucilage that prevents their  
197 separation; or ii) cell adhesion: single cells aggregate via the secretion of sticky mucilage (Kessel and  
198 Eloff, 1975; Yang et al., 2012). In any case, colonies are extremely buoyant, commonly forming wind-  
199 blown scums (Znachor et al., 2006). As mentioned above, their buoyancy is achieved mainly by the  
200 presence of aerotopes (Šmarda and Maršálek, 2008) and helped by the presence of the thick matrix of  
201 EPS produced by the cells and typically composed by glucose, xylose, galactose, fucose, arabinose and  
202 rhamnose (Lei et al., 2007; Li et al., 2009) (Table 1) (Figure 2). It has been demonstrated that the  
203 morphology of *Microcystis* colonies in culture could change and the solubilization of the mucilage could  
204 induce changes in colonial morphology (Li et al., 2009; Otsuka et al., 2000; Wang et al., 2013).

205 The presence of a mucilaginous envelope also provides *Microcystis* with an advantage to withstand  
206 alterations of the physical environment, such as osmotic stress (Kehr and Dittmann, 2015). Moreover,  
207 it has been shown that after strong mixing events large mucilaginous colonies have a faster recovery of  
208 the near to surface position than single cells and small-sized colonies (Kruk et al., 2017). Floating rates  
209 of medium size *Microcystis* colonies (ca. 100  $\mu\text{m}$ ) rarely exceeded  $\pm 30 \mu\text{m s}^{-1}$ , whereas colonies  
210 considerably larger than this are reported to achieve flotation rates of 300  $\mu\text{m s}^{-1}$  (Ganf, 1974;  
211 Humphries and Imberger, 1983; Reynolds, 2007). This buoyancy would favor the access to sunlight by  
212 the cells located at the center of the colony. In addition, it has been shown that estuarine to marine  
213 salinity levels promoted an increase in the thickness of the mucilage and a decrease of cell-free space  
214 resulting in higher cell density, which serve as a defense mechanism to cope with salinity stress  
215 (Sampognaro et al., 2020). Since *Microcystis* colonies contain many times its dry weight in water they  
216 would match the criteria for being classified as hydrogels.



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229 **Figure 2. *Microcystis* mucilage.** Chinese ink staining of *Microcystis* spp. colonies reveals the thick layer of  
230 mucilage. Brightfield, 40x magnification.

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Colony formation in *Microcystis* can be induced by abiotic factors, such as low temperature (15 °C) and low light intensity ( $10 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ ), which enable them to develop colonies up to 100  $\mu\text{m}$  diameter (Li et al., 2013; Xu et al., 2016; Yang et al., 2012). On the opposite hand, an increased growth rate is observed at higher light intensities, with a concomitant increase of intracellular polysaccharides consumption that provokes a decreased propensity to form colonies (Xiao et al., 2018). It has been described that in presence of high concentration of calcium (Sato et al., 2017; Wang et al., 2011) and lead (Bi et al., 2013), the formation of colonies reaching more than 100  $\mu\text{m}$  diameter can be induced. In addition, the exposure to metals also showed to induce the secretion of EPS that helps to precipitate the metal ions, acting as a mechanism to avoid metal poisoning (Bi et al., 2013). The ability to form colonies was also linked to antibiotic resistance. In this sense, low concentrations of aminoglycoside antibiotics induced the aggregation of *Microcystis* cells, suggesting a protective role for the EPS (Zhang et al., 2018).

244 Nutrients can also affect colony formation, as previously demonstrated by Ma et al. (2014), who  
245 found that addition of nitrogen and phosphorus provoked disaggregation of colonies in culture; while  
246 (Zhu et al., 2016) found a general decrease in colony size at increasing nutrient concentrations in the  
247 field, potentially resulting from increased growth rate. Interestingly, the EPS amounts produced by  
248 single-celled laboratory strains have shown to be intensely reduced compared to freshly isolated  
249 colonies, which can have up to 10-fold higher quantities of EPS (Wang et al., 2011). Thus, the  
250 production of an EPS-rich mucilaginous envelope seems to be a response to adverse environmental  
251 conditions.

### 252 **Evidence for a quorum sensing (QS) mechanism in *Microcystis***

253 Early studies suggested that microcystins could act as a signaling or QS molecule (Dittmann et al.,  
254 2001). The MrpA protein (a microcystin-related protein) was found to be strongly expressed in wild-  
255 type *Microcystis* PCC 7806, but became undetectable in a mutant lacking a gene involved in microcystin  
256 synthesis, *mcyB*. This protein showed similarity to the RhiA protein from *Rhizobium leguminosarum*,  
257 which is encoded by the *rhiABC* operon and it is controlled by quorum-sensing mediators  
258 (Supplementary Table 1). This finding led the authors to suggest a QS role of microcystins (Dittmann  
259 et al., 2001).

260 Zhai et al. (2012) reported evidence indicating the presence of QS-related signal molecules, the  
261 acylated homoserine lactones (AHLs) in cultures of *M. aeruginosa* PCC-7820. Electron microscope  
262 photographs of *M. aeruginosa* supplemented with AHLs showed a shift from single free-living cells to  
263 a biofilm-like membrane, which led to a stronger aggregation of the cells compared to controls without  
264 AHLs. This suggests that QS might play an important role in the environmentally-driven morphological  
265 changes of *M. aeruginosa*, providing strong evidence that it regulates colony formation (Zhai *et al.*,  
266 2012) through a coordinated multicellular behavior, as described for biofilms. More recently, Herrera  
267 and Echeverry (2021) applied several AHLs known to be involved in QS in Gram negative bacteria to  
268 cultures of *Microcystis* and found a correlation with colony-forming activity for most of them. This  
269 finding is very interesting, since it also points to a QS-based mechanism associated with the growth of  
270 the colonies. Moreover, using ELISA assays, they found increased microcystin levels with some AHLs.  
271 They propose that the source of the AHLs could be *Microcystis* or members of the microbiome present

272 in the phycosphere, meaning that the QS would be an ability conferred by the cyanobacterium and its  
273 microbiome, acting cooperatively as a holobiont. Further studies analyzing the presence of the genes  
274 encoding for the AHLs synthesis should be performed to confirm this.

### 275 **Phenotypic and functional differences between single cells and colonies**

276 The cells growing in a *Microcystis* colony are physiologically distinct from single cells (Table 1),  
277 for example in terms of toxin production. In this sense, there is a growing body of evidence relating the  
278 colony size to toxicity and identifying that colonies in the size range from 60 to 150  $\mu\text{m}$  (diameter or  
279 maximum linear dimension) are those producing higher amount of microcystins compared to single cells  
280 or colonies smaller than 20  $\mu\text{m}$  diameter (Deus Álvarez et al., 2020; Gan et al., 2012). Cultures of *M.*  
281 *wesenbergii* DC-M1, *M. ichthyoblabe* TH-M1 and *Microcystis* sp. FACHB1027 treated with  
282 microcystin-RR developed colonies significantly larger than the control and provoked the upregulation  
283 of genes related to the synthesis of polysaccharides: *capD*, *csaB*, *tagH* and *epsL*, resulting in a significant  
284 increase of EPS (Supplementary Table 1). This is especially relevant in the case of the non-toxic species  
285 *M. wesenbergii*, since the interaction with exogenous microcystin affected growth and colony size and  
286 suggests that during a bloom the toxin produced by toxic species would also promote the growth of non-  
287 toxic ones. On the other hand, depletion of extracellular microcystin concentrations caused a decrease  
288 in colony size, indicating that released microcystins may be involved in maintaining the colony size of  
289 *Microcystis* (Gan et al., 2012), regardless of their toxin-production ability.

290 It has been also shown that when subjected to intense grazing *Microcystis* cells increase the  
291 abundances of transcripts encoding extracellular polysaccharides and gas vesicles (Harke and Gobler,  
292 2013). This is in agreement with early findings by Reynolds et al. (1981), who reported that *Daphnia*  
293 grazing pressure is stronger on small colonies having less amount of mucilage, indicating that the  
294 increase of EPS production could be a mechanism to avoid predation (Reynolds et al., 1981).

295 Interestingly, genes encoding for type IV pili (e.g., *pilT*) have been found in *Microcystis aeruginosa*  
296 PCC 7806 (Nakasugi and Neilan, 2005) (Supplementary Table 1). These pili are present in many gram-  
297 negative bacteria systems and are involved in several functions such as cell adhesion, twitching motility,  
298 and natural transformation (Mattick, 2002). In several bacterial species, they are also related to biofilm  
299 formation via bacterial migration (Barken et al., 2008). For example, in the case of *P. aeruginosa* are

300 necessary for cap formation of the mushroom-shaped structured biofilm (Klausen et al., 2003), while in  
301 *Clostridium difficile* promote early biofilm formation (Maldarelli et al., 2016). We hypothesize that the  
302 presence of *pilT* in *Microcystis* cells (see Nakasugi and Neilan, 2005 for images reference) reveals their  
303 ability to move and could have a role during the initial arrangement of the cells inside the growing  
304 colony. When individual cells initiate a biofilm, they are surrounded by small amounts of EPS and are  
305 probably capable of independent movement by means of twitching motility mediated by these pili. As  
306 the colony grows and the biofilm starts to mature, water channels develop and a differentiation in  
307 physiological processes among cells start to establish in response to conditions in their particular  
308 environments (Stoodley et al., 2002).

309 In a previous study addressing the individual activity of cells in the colony using the redox dye 5-  
310 cyano-2,3-ditolyl tetrazolium chloride (CTC), we found that cells located at the inner part of the colonies  
311 exhibited a lower respiration activity than those in the peripheric, suggesting a less active metabolic  
312 state (Kruk et al., 2017). This kind of differential activity can be also found in mature biofilms of several  
313 bacterial species. In *P. putida* and *E. coli* the cellular activity in the center of cell clusters diminished as  
314 the clusters grew larger. This activity can be restored after carbon sources addition, suggesting that cell  
315 activity in the inner part of the clusters would be controlled by resources availability (Sternberg et al.,  
316 1999).

317 When growing under laboratory conditions, *M. aeruginosa* colonies disaggregate and resulting  
318 single cells have significantly lower chlorophyll a, phycocyanin and total carbohydrate than cells in  
319 colonies (Reynolds et al., 1981; Wang et al., 2015). This induction of a unicellular lifestyle has not been  
320 explored enough and might be the consequence of dilution and selection in a favorable milieu. This  
321 change from colonies to single cells that occurs in cultures might reflect the different ecological  
322 requirements between both morphological states and call for caution when analyzing the ecology and  
323 environmental preferences of these organisms by using isolates from the laboratory.

#### 324 **Potential role of the *Microcystis* microbiome in colony formation**

325 As it can be seen in Figure 3, the mucilage of *Microcystis* is populated by a high number of  
326 heterotrophic bacteria growing at expense of the EPS carbon, as well as other cyanobacteria and  
327 sometimes even protists. The interactions occurring between individual organisms within this

328 phycosphere have an ecosystem-level effect on several processes, e.g., nutrient cycling, toxin  
329 biosynthesis, etc. (Bell and Mitchell, 1972; Seymour et al., 2017). The incorporation of bacteria into the  
330 phycosphere likely occurs through chemotaxis, random contacts and vertical transmission (Seymour et  
331 al., 2017).

332 In the case of *Microcystis*, it has been shown that heterotrophic bacteria stimulated cyanobacterial  
333 growth and induced the production of EPS (Wang et al., 2016). In fact, their presence was early  
334 documented by Reynolds et al. (1981), who found that *Microcystis* mucilage was crowded with rod-  
335 shaped bacteria that were frequently observed on the periphery of planktonic colonies in their stationary  
336 or decline phases (Table 2).

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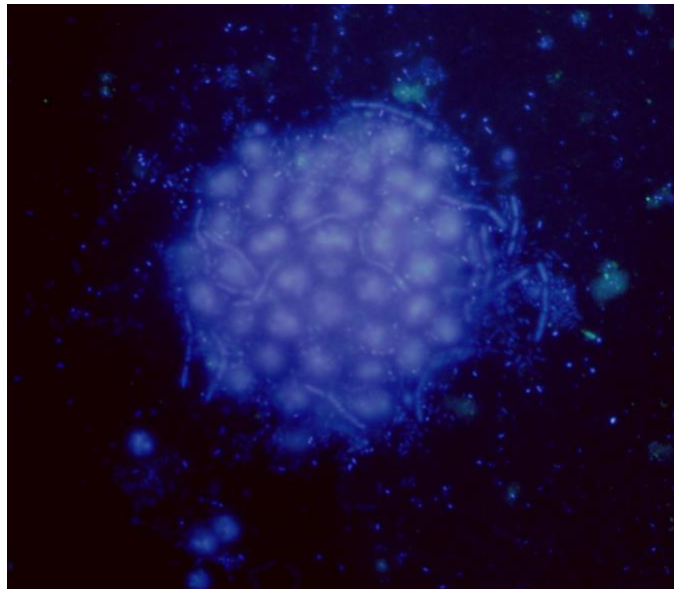
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347 **Figure 3. *Microcystis* microbiome.** DAPI-stained *Microcystis* spp. colony showing the bacteria attached to the  
348 mucilage. 1000x magnification.

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354 **Table 2.** Bacterial groups that have been described as associated to *Microcystis* colonies through  
 355 different methodological approaches (bacterial cultures, 16S rDNA and shotgun metagenomics,  
 356 metagenome-assembled genomes).

Phylum	Class	Order/Family/Genus	Found in colony or free-living	References
Proteobacteria	Alphaproteobacteria	Rhizobiales, Sphingomonadales, Rhodospirillales, Caulobacterales	Colony-attached	Li et al., 2018; Jankowiak and Gobler, 2020; Cook et al., 2019; Wu et al., 2019; Pérez-Carrascal et al., 2021; Jackrel et al., 2019; Li et al., 2018;
		Rhodocyclaceae	Colony-attached	
		<i>Pleomorphomonas</i>	Colony-attached	
		<i>Candidatus Phycosocius bacilliformis</i>	<i>M. aeruginosa</i> phycosphere	
		<i>Rhodobacter</i>	Colony-attached	
		<i>Methylobacterium</i>	Colony-attached	
		<i>Roseomonas</i>	Colony-attached	
		<i>Pseudanabaena</i>	Colony-attached	
	Betaproteobacteria	<i>Burkholderia</i>	Colony-attached	
		<i>Alcaligenaceae</i>		
	Gammaproteobacteria		Colony-attached	
	Epsilonproteobacteria		Colony-attached	
Bacteroidetes	Cytophagia, Sphingobacteriia, Chitinophagia	Chitinophagaceae, Cytophagales	Found in free-living and colony-attached fractions	Wu et al., 2019; Pérez-Carrascal et al., 2021
Firmicutes				Wang et al., 2016
Gemmatimonadetes		<i>Gemmatimonas</i>	Colony-attached	Yang et al., 2017; Pérez-Carrascal et al., 2021
Verrucomicrobia			Colony-attached	Cai et al., 2014

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358 The bacterial community inhabiting the mucilage has shown to differ markedly from that present  
 359 in free-living *Microcystis* (Wu et al., 2019). This highly diverse microbiome surrounding *Microcystis*  
 360 colonies and living in close cooperative way allows the cyanobacteria the access to specific compounds,  
 361 such as vitamins and some components of the outer membrane lipopolysaccharide, while providing  
 362 bacteria with highly bioavailable carbon. The bacterial community in the phycosphere has shown to be

363 highly structured and related to the size of the colonies, suggesting highly specific conditions within the  
364 *Microcystis* (Cai et al., 2014). Interestingly, it has been shown that although *Microcystis* microbiomes  
365 diverged in taxonomy along a phosphorus concentration gradient, they converged in function, indicating  
366 a metabolic interdependence between the host and its microbiome (Jackrel et al., 2019). Cook et al.  
367 (2020) found that *Microcystis* microbiome is highly similar across global blooms regardless of the  
368 environmental differences, pointing out the existence of a stable associated microbiome that might be  
369 involved in colony formation (Cook et al., 2020). Moreover, it has been recently found by single-colony  
370 metagenomic sequencing that *Microcystis* microbiome is genotype-specific, and that closely related  
371 genotypes have similar microbiomes (Pérez-Carrascal et al., 2021). Similarly, Tu et al. (2019) found  
372 that colonies of the same *Microcystis* species have very similar community composition. Indeed, the  
373 non-toxic *M. wesenbergii* microbiome harbored a very different composition compared to that from  
374 toxic *M. aeruginosa* and *M. panniformis*, which implies that microcystin could play a structuring role  
375 in the community.

376 It has been reported that high temperatures (32 °C) provoke changes in the composition of  
377 heterotrophic bacterial communities embedded into the mucilage of *Microcystis* (Dziallas and Grossart,  
378 2011). These associated heterotrophic bacteria can stimulate cyanobacterial growth and induce the  
379 production of EPS that is relevant for the optimum development of *Microcystis* (Reynolds, 2007).  
380 Besides, interspecies interactions could promote EPS production (Yang et al., 2008), and co-cultivation  
381 of axenic, single-celled cultures of *Microcystis* with heterotrophic bacteria isolated from *Microcystis*  
382 colonies stimulated the production of EPS, allowing to reconstitute colony formation (Shen et al., 2011).  
383 Moreover, removing the EPS had a detrimental effect on the auto-aggregation abilities of heterotrophic  
384 bacteria isolated from *Microcystis* colonies, which suggest that EPS plays a relevant role in the  
385 recruitment of bacteria by promoting their attachment (Zhang et al., 2018). More recently, Schmidt et  
386 al. (2020) have found that during invasion experiments the ability of *M. aeruginosa* to compete with  
387 other phytoplankton species is not determined by the ability to produce the toxin, but by genes from its  
388 microbiome. This points to an important role of host-associated bacteria in mediating phytoplankton  
389 interspecies interactions.

390 The resulting cooperative microbial network, which strongly agrees with the holobiont concept as  
391 the ecological and evolutionary unit, might be the key for *Microcystis* success under changing  
392 environments. The role of the *Microcystis* microbiome is just starting to be discovered and further  
393 research is needed to shed light on the role of specific heterotrophic organisms in colony formation and  
394 in the persistence of *Microcystis*.

#### 395 **Further evidence on the similarity between the colonial lifestyle and biofilms**

396 Sigee et al. (2007) reported the presence of programmed cell death (PCD) during a late summer  
397 bloom of *Microcystis*. More recently, Hu and Rzymiski (2019) proposed that under certain stressing  
398 conditions (excessive salt concentration, exogenous oxidants, ultraviolet radiation, herbicides, among  
399 others) a PCD-like mechanism would cause apoptosis and significant release of microcystin in  
400 *Microcystis*. The released microcystin would have an extracellular function, not yet described, which  
401 benefits the rest of the population inside the colony. These authors also speculate on the similarities  
402 between *Microcystis* colonies and bacterial biofilms, emphasizing the central role of the PCD and  
403 microcystin release in bloom development. They also propose that non-microcystin producing  
404 *Microcystis* would benefit from the microcystin released by the toxic populations (Hu and Rzymiski,  
405 2019).

406 These antecedents constitute additional evidence that point towards a kind of multicellular behavior  
407 within the colonies, such as that existing in the bacterial biofilms. In multicellular organisms, PCD  
408 removes cells having molecular errors in order to keep homeostasis and a healthy organism  
409 development, prioritizing the benefit of the organism over the survival of the cell. In the case of bacteria,  
410 this behavior would not provide any benefit to an individual cell but would confer an advantage to the  
411 remaining cells (Allocati et al., 2015; Bayles, 2014; Lewis, 2000). In *Microcystis*, colony formation  
412 provides a highly efficient survival strategy under adverse environmental conditions (low nutrient  
413 levels, high grazing pressure, high ultraviolet radiation).

414

#### 415 **PROPOSAL OF A CONCEPTUAL MODEL FOR BIOFILM FORMATION IN *MICROCYSTIS***

416 **SPP.**



417 The colonies of *Microcystis* spp. share several characteristics with bacterial biofilms. They can  
 418 switch from single planktonic cells to aggregates (colonies) organized into a coordinated functional  
 419 community that is embedded in an EPS matrix, which composition highly resembles that found in  
 420 bacterial biofilms and that is teemed with a diversity of heterotrophic bacteria living in a cooperative  
 421 manner with the cyanobacterial cells (Figure 4). The change from single or few cells to multicellular  
 422 organization would be triggered by autoinducers molecules, such as AHLs, which could be synthesized  
 423 either by the cyanobacterium or by the microbiome, and that build up during exponential growth under  
 424 resource-rich conditions. As the population grows, the resources become less available and the AHLs  
 425 upregulate a number of functional genes (e.g., the microcystin biosynthesis cluster *mcy*) that allow the  
 426 organisms to thrive under conditions that would not be favorable, such as nutrients and light shortage.  
 427 This kind of multi-specific biofilm is not built from the attachment of the cells to an abiotic or biotic  
 428 surface, but on the attachment of cells to each other to form a floating biofilm and allowing *Microcystis*  
 429 spp. to thrive in a highly diverse array of environmental conditions.

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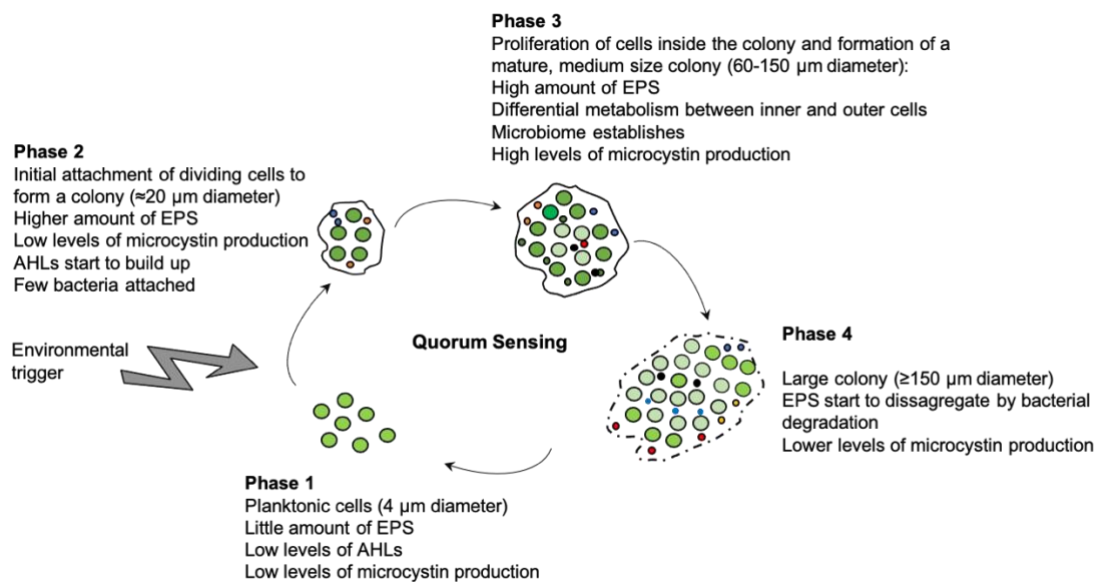
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441 **Figure 4. Proposal for the floating biofilm formation in colonies of *Microcystis*.** Four phases can be

442 distinguished during the development of a *Microcystis* community according to lifestyle (single celled vs attached),

443 EPS and microcystin production, presence of an established microbiome and autoinducers concentration (AHLs).

444 Phase 1 is composed of single cells (4 μm diameter, green circles) having little amount of EPS mucilage, low

445 levels of microcystin production and low levels of AHLs. Phase 2 starts with the initial attachment of dividing  
446 cells to each other to form a colony surrounded by a higher amount of EPS mucilage, cells probably mobilize  
447 inside the colony and they have low levels of microcystin synthesis while AHLs start to build up and other bacteria  
448 (smaller red, blue and black circles) start to attach to the EPS. In Phase 3, the proliferation of cells inside the colony  
449 allows the formation of a mature biofilm, with elevated amounts of EPS, high levels of microcystin production  
450 and clearly different metabolism between inner and outer cells. A microbiome is well established. The Phase 4 is  
451 characterized by large, amorphous colonies, low levels of microcystin production and disaggregation of the  
452 mucilage by bacterial degradation of the EPS (typically at the end of a bloom). We propose that the onset of a  
453 bloom will depend on abiotic and biotic conditions and on the phase of the *Microcystis* community, being more  
454 likely to develop a high biomass in a short time period during phase 3 (active cells, with high microcystin  
455 production rates).

456

457 It must be noticed that, as it has been described for temperate lakes, the sediment can be the source  
458 of *Microcystis* colonies that will trigger the water column colonization when environmental conditions  
459 are favorable, especially in temperate lakes where this kind of annual cycle has been described  
460 (Reynolds et al., 1981; Yang et al., 2020). The phase at which these colonies will be recruited after  
461 winter would depend on the lake temperature and the physiological state of the overwinter organisms.

462 More data and information gathered from studies specifically addressing the biofilm formation in  
463 *Microcystis* are needed in order to develop mathematical models describing the growth and dispersal of  
464 the colonies. This, together with morphological, gene expression and functional studies of different  
465 *Microcystis* species would bring new insight into the life cycle of this relevant group of organisms.

466 Understanding the mechanism underlying biofilm formation in *Microcystis* spp. and the role of  
467 heterotrophic bacterial community in toxin synthesis and environmental performance will improve the  
468 current models of growth, fitness and dispersal of these cyanobacteria.

## 469 CONCLUSIONS

470 1) The information gathered so far about colony formation in *Microcystis* spp. suggests that the  
471 mechanisms involved in this process are the same as those defined for biofilm formation in a number of  
472 bacterial species. Single-celled *Microcystis* are able to multiply while producing a mucilaginous

473 envelope that contributes to the differentiation into a colony. Colonies are described to form either by  
474 cell division or by cell aggregation; in any case, the presence of an extracellular matrix ensures the  
475 confinement of the cells into a tridimensional, secluded structure. The triggers to switch between both  
476 lifestyles would involve environmental cues that induce cellular stress such as salinity, oxidative damage  
477 due to ROS (inducing microcystin production and export), predation and low nutrient availability.

478 2) The colonial or biofilm stage, while provokes a reduced specific growth rate, provides *Microcystis*  
479 with a sheltered milieu. As a consequence, several gradients of resources (oxygen, light, nutrients) are  
480 generated, with the concomitant creation of different micro-environments inside the colony. Thus, cells  
481 located in these different micro-environments will have different metabolic rates (e.g., respiration,  
482 photosynthesis, microcystin production, etc.).

483 3) The main components of the biofilm mucilage are EPS, DNA from lysed cells, proteins, lipids and  
484 heterotrophic bacteria that live embedded in this mucus. The bacterial community associated with  
485 *Microcystis* colonies is starting to be explored and recent findings suggest that there is a quite constant  
486 microbiome, in which functional relationships with the cyanobacterium are closely intertwined. This  
487 relationship involves the trade of different goods that allows the survival and increases the fitness of the  
488 colony as a multispecies biofilm community.

489 4) The concentration of microcystin has been shown to be linked to the production of EPS by the cells,  
490 suggesting that the toxicity potential of *Microcystis* could be the consequence of cells adopting a biofilm  
491 lifestyle.

492 5) The presence of AHLs, the autoinducers molecules involved in bacterial QS behavior, has been  
493 detected in *M. aeruginosa* and experimental evidence suggests that the increase of AHLs provokes a  
494 shift from single-cell to a biofilm-like status. This strongly suggests that the QS system regulates the  
495 coordinated cellular behavior leading to biofilm formation.

496 6) As stated by Xiao et al. in 2018, we still do not have a clear explanation for the observed differences  
497 in colonial morphology among different *Microcystis* species. This gap in knowledge could be shortened  
498 if colonies start to be studied using the biofilm approach, since morphology differences (differences in  
499 biofilm architecture) could be attributed to different phases of biofilm maturation and would account  
500 for the environmental conditions to which the organisms were subjected.

501

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504

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