1	Title: An engineered community approach for industrial cultivation of microalgae			
2	Authors: Elena Kazamia ^{1*} , Anthony S. Riseley ² , Christopher J. Howe ² and Alison G. Smith ¹			
3	Affiliations:			
4 5	 Department of Plant Sciences, University of Cambridge, Downing Street, Cambridge, CB2 3EA, United Kingdom. 			
6 7	 Department of Biochemistry, University of Cambridge, Tennis Court Road, Cambridge, CB2 1QW, United Kingdom. 			
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9	Running title: Community approaches for robust algal cultures			
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11	*Corresponding author information:			
12	Dr Elena Kazamia			
13	Ek288@cam.ac.uk			
14	+441223330219			
15				
16	Plant Metabolism Group			
17	Department of Plant Sciences			
18	University of Cambridge			
19	Downing Street			
20	Cambridge			
21	CB2 3EA			
22	United Kingdom			
23				

25 Abstract

26 In Nature, no species live in isolation. Traditionally, efforts to grow organisms for use in 27 biotechnology have focused on a single-species approach, particularly where a high-value 28 product is required in pure form. In such scenarios, preventing the establishment of 29 contaminants requires considerable effort that is justified economically. However, for algal 30 biotechnology, in particular where the focus is on fuel production, axenic culture is not 31 necessary, provided yields of the desired strain are not hampered by unwanted 32 contaminants. In the following article we review what is known about inter-specific 33 interactions of natural algal communities, the dynamics of which are likely to parallel 34 contamination in industrial systems. Furthermore, we discuss the opportunities to improve 35 both yields and the stability of cultures by growing algae in multi-species consortia.

36

37 **1. Background**

38 Microalgae (eukaryotic photosynthetic microbes) and cyanobacteria (oxygenic 39 photosynthetic bacteria), are a highly diverse collection of micro-organisms. They live in a 40 range of environments, including all aquatic ecosystems, both fresh-water and marine, and 41 species are also found in terrestrial habitats including on hard surfaces and snow. Many taxa 42 are capable of growing heterotrophically as well as phototrophically, and some obligate 43 heterotrophs also exist that, although ancestrally photosynthetic, have lost the ability to 44 photosynthesise. These include the dinoflagellate Crypthecodinium cohnii, which is of commercial importance as a source of docosahexaenoic acid (DHA).¹ Algae are currently 45 46 cultivated on a relatively small scale for high value products such as the carotenoid

astaxanthin from *Haematococcus pluvialis* and the phycobiliprotein phycocyanin from the
cyanobacterium *Aphanizomenon flos-aquae*.² Certain strains are marketed as dietary
supplements, such as the cyanobacterium *Spirulina* sp. (*Arthrospira platensis*) and *Chlorella vulgaris*.

51 Bulk growth of algae for products of lower value to displace commodities traditionally made from fossil oil has received a lot of research attention.³ However, the scale-up required to 52 53 achieve this poses a wide range of problems, ranging from the energy costs of maintaining 54 large-scale photobioreactors, lower yields in large-scale cultures arising from factors such as 55 poor light penetration, mass transfer (where exogenous carbon dioxide is supplied, or 56 oxygen needs to be removed) and biological contamination, as well as the energy costs of downstream processing and product.⁴⁻⁶ We concentrate here on consideration of how 57 58 understanding the ecology of the organisms under cultivation, that is their interaction with 59 others in the environment, can be harnessed to enhance productivity and thus increase financial and environmental benefits achieved by cultivating algae. 60

61

2. Applying community ecology to algal cultivation

62 Most studies that target increasing yields in industrial cultures are aimed at an individual 63 species level, which assumes that cultivation will be in monoculture. However, because 64 contamination is inevitable without stringent sterile practice, which is neither cost-effective nor likely to be achievable at industrial scale,⁷ understanding the growth dynamics of an 65 algal population growing in reactors is fundamentally an ecological problem.⁸ Moreover, 66 67 monocultures are by their nature unstable and prone to perturbation. Their genetic uniformity encourages quick proliferation of pathogens and invaders, a common problem for 68 traditional single-crop agriculture (reviewed in Smith et al., 2014)⁹. Monocultures are 69

predicted to be unstable by classical theories of community ecology, which describe natural systems as increasing in complexity over time (e.g. Elton, 1958)¹⁰. Given the chance, for a given habitable environment, multiple species with diverse niche specificities will coexist alongside each other, maximising the use of the available resources. "Invasions" by organisms from neighbouring environments will continue until a "climax" stable state is assembled, which is predicted to be resilient to change provided abiotic conditions remain constant (May, 1977)¹¹.

Therefore, a new and emerging approach is to consider community approaches to cultivation. The reasoning is that by starting with what would be an "end-point" consortium in a natural system, it may be possible to avoid the development of unwanted alternatives. In the following section we review the advantages of growing algae in consortia of species, rather than as monocultures. The principles that we draw on are from aquatic community ecology, and key concepts are summarised in Table 1.

83

2.1 Maximising productivity

84 One of the tenets of community ecology is that productivity is enhanced when diverse 85 organisms are grown together. This has been illustrated for a range of habitats, and 86 famously in a long-term experiment on grasslands. For a period of seven years, it was 87 demonstrated that 16-species grassland plots attained 2.7 times more biomass than the respective monocultures.⁹ An aquatic experiment showed that diverse algal communities 88 89 (grown in biofilms) increased the uptake and storage of nitrate from streams, and significantly increased in biomass content compared to monocultures.¹⁰ Overyielding is said 90 91 to occur when the biomass production of a consortium of species is greater than that of the average monoculture of the species contained in the mixture.¹¹ *Transgressive* overyielding is 92

93 said to occur when the mixture outperforms even the most productive of the monocultures of the constituent species.¹² There is evidence that, when functionally diverse algae with 94 complementary light requirements are grown together, the resulting communities are more 95 96 productive than monocultures of individual species. Behl et al. analysed the rate of carbon 97 uptake and productivity for 85 assembled communities composed of species from four functional groups: chlorophytes, diatoms, cyanobacteria and chrysophytes.¹³ The 98 99 researchers found that all algal communities consisting of species from two, three or four 100 different functional groups showed overyielding compared with their respective 101 monocultures, with transgressive overyielding in more than half of the assemblages studied. 102 This is interesting as it suggests that positive interactions beyond resource use 103 complementarity occurred between species. A possible way this could occur is through 104 mutualistic interactions, reviewed in Section 2.3 below.

105 An important explanation for increased productivity in diverse cultures is through resource 106 use complementarity. When species that have different growth requirements are grown 107 together, competition between members of the community is reduced compared with that 108 experienced by individuals in dense monocultures. This allows more individuals to cohabit, 109 increasing the net biomass of the culture. One of the traits that distinguishes algal species is 110 the portfolio of pigments they use to absorb light. Although oxygenic photosynthetic 111 organisms use chlorophyll a as the major pigment in the photosystems, the accessory light 112 harvesting pigments differ (Figure 1). In cyanobacteria grown under iron replete conditions, 113 phycobilisomes on the surface of the thylakoid membranes contain the phycobilin pigments 114 phycocyanin and phycoerythrin. These pigments are also found in red algae, whereas green 115 algae (chlorophytes) contain chlorophyll b, as do all land plants. Chlorophyll c is the major

accessory pigment in the Chromalveolata. A possible explanation for overyielding of diverse
algae grown in cocultures as observed by Behl et al. therefore could be due to maximised
use of available light resource.

119 2.2 *Crop protection*

Contaminating organisms that invade algal cultures can reduce yields in different ways: predators and pathogens are able to do so directly by killing the algae in culture, whilst competing microalgae can take over as the dominant strain. The latter is a problem when a specific algal strain is required, such as an oil producer or a strain with useful pigments. In principle it could be possible to address all of these challenges by growing algae in culture with carefully selected cohabiting species.

126 The effect of predators can be decreased through biomanipulation of the food web, 127 whereby an ecosystem is deliberately altered by adding or removing species. This is common 128 practice in the freshwater management industry, where the goal is to minimise algal production.¹⁴ In the context of algal cultivation, which is the reverse scenario, if production 129 130 were to be hampered by invading zooplankton, the addition of zooplanktivores (such as small fish) to the reactors might increase yields.^{8,15} However, this is unlikely to be possible 131 132 for closed photobioreactors, but may also not be practical in open ponds because most 133 reactors are very shallow, and would not be suitable for fish. An alternative solution is that 134 of crop protection through "interference". By introducing multiple inedible algal species to 135 grow alongside the desired strain, the foraging efficiency of invading zooplankton may be decreased due to the increased energetic costs of finding their desired prey.¹⁶ This technique 136 137 of pest control was recently investigated by Shurin et al. in a set of laboratory experiments, the results of which are summarised in Figure 2.¹⁷ Communities containing 1, 2, 5 and 10 138

species of algae in various combinations were subjected to grazing by *Daphnia pulex*. Although the total biomass of algal food resources increased with diversity, survival of introduced *Daphnia* grazers declined markedly when 5 or 10 species of algae were grown together.

143 However, there may be a cost to co-cultivation of a range of algal species when only a single 144 species is of commercial interest. It is possible to imagine a scenario where the growth of a 145 desired strain is decreased in a dense polyculture due to increased shading by co-cultured 146 strains. Where stability of a monoculture against invasions is the primary concern, this may 147 be enhanced by manipulating the abiotic environment to make the establishment of 148 competitors less likely. This is why extremophiles have been preferred in commercial 149 cultures, such as Spirulina sp., which is grown in highly alkaline conditions, or Dunaliella 150 salina, which is cultured in highly saline medium. A community solution to the problem of 151 competitors may be engineered through co-culturing with partners that produce allelopathic 152 chemicals. Chemical interactions are an important part of phytoplankton competition and are particularly functionally important with dinoflagellates and cyanobacteria.¹⁸ These 153 154 organisms are able to produce chemicals that are toxic to most other algae in the 155 environment, allowing the former to bloom under the right conditions for growth, often causing what are known as Harmful Algal Blooms, HABs.¹⁹ However, some species have 156 157 evolved to withstand the toxins produced during HABs and are able to cohabit with the toxin 158 producing strains. If either HAB-forming or HAB-tolerant species were identified as 159 interesting candidates for biofuel production, growth in consortia with toxin producing 160 strains could be a possible solution to competitive invasion. A similar approach is taken in 161 water treatment, where often barley straw is used to control populations of unwanted

algae. Toxins produced from the straw liquor are known to inhibit the growth of some algae
but not others.²⁰

164 Finally, bacterial contaminants often invade cultures of algae, as they are able to scavenge 165 algal exudates, which provide a source of carbon. If the bacteria compete with algae for 166 other nutrients, they often overtake the growth of the microalgae and can lead to the 167 establishment of anoxic conditions (REFS by Val). Bacterial fouling (surface growth) is very 168 severe in closed bioreactors, requiring these systems to be shut down and fully flushed 169 before operation can resume. This leads to yield losses and has an associated financial 170 burden. We have previously suggested that bacterial contamination may be decreased through co-culturing algae with symbiotic (probiotic) bacteria that enhance algal growth.²¹ 171 172 When bacteria are present in the culture medium, invading bacteria are less likely to 173 establish as the bacterial niche is already occupied. There is some empirical evidence from 174 fish aquaculture that supports this theory. For example, Sharifah and Eguchi report that 175 Roseobacter clade bacteria that are symbiotic with Nannochloropsis oculata (grown 176 commercially for fish food) successfully inhibited the growth of the fish pathogen Vibrio anguillarum.²² 177

178 **2.3**

Capitalising on mutualisms

There is a range of ways in which it is possible to capitalise on mutualisms in industrial biotechnology of microalgae. Mutualistic exchange of metabolites can replace external inputs of scarce or expensive resources. For example, half of all algae are known to require vitamin B₁₂ (cobalamin) for growth, while no eukaryotic microalgae are able to synthesise it. Model laboratory consortia have been described in which vitamin B₁₂ dependent algae can obtain cobalamin from vitamin B₁₂-synthesising bacteria, in exchange for a source of fixed 185 carbon,^{21,23} and indeed in the case of the *Dinoroseobacter shibae* partnership with its 186 dinoflagellate host, vitamin B_1 is also exchanged (Figure 3A). If this system were to be 187 employed industrially, the bacteria could replace exogenous addition of vitamins into the 188 medium, reducing material and energy inputs into the system. Other described mutualisms 189 include the provision of iron via siderophores from bacteria to algae in exchange for fixed 190 carbon.²⁴

191 It is possible to envisage a system where the mutualism between algae and bacteria 192 depends on provision of nitrogen by the bacteria, a macronutrient that is acknowledged as one of the key drivers of microalgal productivity in natural systems.^{25,26} Modelling the 193 194 potential for algal biodiesel production in the USA indicated that the availability of nitrogen and phosphorus fertilisers were the major limiting factors to large scale cultivation.²⁷ In a 195 196 recent study, Azotobacter vinelandii, a nitrogen-fixing bacterium, was genetically engineered to excrete ammonium into the surrounding medium.²⁸ When the strain was co-cultured in 197 198 medium that did not contain exogenous carbon or nitrogen with oil producing microalgae 199 including Chlorella sorokiniana, Pseudokirchineriella sp. and Scenedesmus obliquus, the algae 200 were able to grow and accumulated lipid of up to 30% of their dry weight (Figure 3B). This 201 shows the potential for growing algae industrially in the absence of nitrogenous fertiliser 202 input by co-culturing with appropriate bacteria. As nitrogenous fertiliser is made through the 203 energy-intensive Haber-Bosch process that has been estimated to contribute up to 40% of all energy inputs into microalgae biofuel systems,²⁹ provision of nitrogen *via* a symbiont could 204 205 significantly reduce the lifecycle energy and carbon footprint of the resulting fuel. It must be 206 noted that a sustainable alternative could be to grow algae on waste water that is rich in nitrogen and phosphorus, thus recycling nutrients from domestic and agricultural effluent.³⁰ 207

208 It is likely that the range of options for co-culturing algae with bacteria will increase as our 209 understanding of inter-specific interactions between these organisms improves. Evidence 210 suggests that microalgal interactions with bacteria are ubiquitous, although the physiological 211 basis for these is often not known. For example, Park et al. describe that 6 out of the 8 212 contaminants isolated from a Chlorella elipsoidea culture enhanced algal growth when coinoculated with the species in a controlled co-culture.³¹ Similarly, Do Nascimento et al. 213 214 described that the inoculation of Rhizobium strain 10II into cultures of oleaginous 215 microalgae Ankistrodesmus sp. strain SP2-15, resulted in up to 30% increased accumulation 216 of chlorophyll, biomass and lipids compared with axenic monocultures of the alga.³² The 217 bacteria influenced the metabolism of the microalgae, redirecting it towards lipid 218 accumulation.

219

2.4 Improving the persistence of a desired strain

220 A similar degree of regulation has been observed in the specific mutualism between the 221 vitamin B₁₂-dependent green alga Lobomonas rostrata and the soil bacterium 222 Mesorhizobium loti, where the ratio of algal to bacterial numbers equilibrated to around 1:30 in semi-continuous co-culture.²¹ Regulation can be defined in accordance with Smith 223 and Douglas (1987) whereby a state of balance and stability between two organisms' growth 224 and population numbers is reached as a result of their symbiosis (living together).³³ 225 226 Mathematical modelling of the dynamics of the two species in coculture revealed that the 227 population growth of one organism could be predicted entirely based on the expected 228 carrying capacity of the cocultured symbionts Grant et al. (2014) . Although the mechanism 229 remains unknown, the biological implication is that the symbionts are controlling the 230 amount of each other's growth when in coculture. Understanding regulatory mechanisms in

symbioses can benefit biotechnology by providing a mechanism for maintaining the long
term maintenance of a culture and its fidelity. If the growth of a desired algal is regulated by
a bacterium (or vice-versa) yields can be maintained despite a changing environment.

234 Environmental fluctuation, such as temperature and irradiance changes, is inevitable in all 235 large scale production systems, and could exert a selection pressure for a community of 236 algae to change from what is optimal for production (for example away from producing high 237 yields of lipids). Furthermore, if genetically modified organisms are considered, a changing 238 environment may exert pressure for the transformed strains to revert back to their original 239 form (the wild type) or drift randomly to an alternative genetic composition. For example, a 240 recent large scale effort to re-sequence strains of wild type Synechocystis sp. PCC6803 241 (originally from Berkeley as described by Stanier et al., 1971) maintained in various culture 242 collections around the world revealed that strains that had been presumed identical had in fact accumulated mutations that are likely to have effects on glucose tolerance, metabolism, 243 motility, phage resistance and stress responses.^{34,35} 244

245 Culturing organisms that have been genetically engineered to be interdependent might 246 provide a selection pressure to prevent reversion, which would decrease the fitness of both 247 partners in the consortium. In fact, it was recently shown that engineered co-dependence is stable even against the evolution of "cheaters" within the system,³⁶ although modelling 248 249 studies suggest that when the cost of cooperation is very high revertants will dominate.^{37,38} 250 Nonetheless it has been argued that co-dependence is so valuable to production it should be genetically engineered.³⁹ Hosoda et al. engineered a syntrophic (cross-feeding) community 251 252 of Escherichia coli, where 2 strains co-habited: one auxotrophic for isoleucine and the other for leucine.⁴⁰ Neither strain was able to survive on its own, but growth was possible in 253

synergistic co-culture. Kerner et al. engineered a similar system, where *E. coli* were either tyrosine or tryptophan auxotrophs, but improved on the previous attempts by introducing an element of control to the system.⁴¹ By tuning the metabolic exchange via gene expression or chemical inducer they were able to regulate the growth rates and strain ratios. Finally, more recently engineered inter-species associations have been demonstrated successfully. An *E. coli* strain auxotrophic for glutamine was engineered to provide lipoic acid to *Dictyostelium discoideum,* an amoeba, in exchange for the amino acid.⁴²

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3. Towards designing algal communities

262 There is increasing awareness amongst the scientific community that microorganisms are 263 very social. Evidence is continuously emerging to demonstrate that microorganisms rely on 264 interactions with other species for a range of functions and communicate and cooperate to 265 perform activities such as dispersal, foraging, construction of biofilms, reproduction, chemical warfare, and signalling.⁴³ Interactions range from necessary or advantageous to 266 267 growth, to competitive or even fatal. Ignoring the importance of interspecific interactions in 268 biotechnology dismisses the problems associated with contamination and misses the 269 opportunity to capitalise on the beneficial associations that can be harnessed to maximise 270 productivity.

We have identified four main advantages for using community approaches for the cultivation of microalgae. It is possible to increase productivity of microalgal cultures (by cultivating consortia of species that have complementary functional traits and therefore overyield) or to decrease loss of productivity, by cultivating microalgae with species from other life domains (such as non-photosynthetic bacteria and zooplanktivores), which can increase resistance to predators and contaminants. We have highlighted the importance of

engineering co-dependence amongst introduced members to the consortium via mutualisms with the benefit of reducing energy and material inputs. Finally, in agreement with Brenner et al.³⁹ we believe that for a stable and robust culture, whenever a new organism is introduced into a consortium, it should be contributing something useful to the culture 'economy' alongside receiving something in return for example through the division of labour or specialisation. In that way interacting organisms rely on each other through trading to establish a stable and long-lasting culture.

284 Of course the use of consortia of microbes in biotechnology is not novel; multi-species 285 systems are often employed to increase yields in microbial-based processes such as anaerobic digestion, fermentation and bioremediation (reviewed in Sabra et al.)⁴⁴. In these 286 287 traditional systems microbial communities are allowed to develop naturally; the most 288 efficient assemblages are chosen for application and subsequently carefully maintained. 289 Although this approach is not common in algal biotechnology, recently Mooij et al. 290 demonstrated that by providing a selection pressure for algae to accumulate storage 291 compounds linked directly to fitness, communities rich in starch and/or lipid assembled stochastically, and were able to outperform monocultures of known lipid producers.⁴⁵ 292

These directed selection approaches will prove very useful to understanding the complex and advantageous interactions of microorganisms. In parallel to these efforts, we proposed a *Synthetic Ecology* approach to consortium assembly of cultures aimed to be more productive and/or more resistant to contamination (15)(Kazamia et al., 2012a). Synthetic ecology differs from the selection approaches by introducing an element of design and using transferrable building blocks (namely specific species, engineered symbioses and growth conditions) to assemble a desired community of microorganisms. We believe that by

300 focusing on species specific interactions and engineered metabolic exchanges we can 301 advance the understanding of fundamental microbial physiology without compromising on 302 creative solutions for biotechnology. However, with all community approaches to 303 cultivation, their efficacy remains questionable until proven at scale. Stability of an 304 engineered consortium may face the same challenges as monocultures. A range of 305 unanswered questions remain: such as how much complexity within a consortium is 306 required before challenges faced by monocultures (instability, invisibility etc.) are 307 surpassed?

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318 **References**

Mendes A, Reis A, Vasconcelos R, et al. Crypthecodinium cohnii with emphasis on DHA
 production: a review. J Appl Phycol 2008:21(2):199–214.

321	2.	Benedetti S, Benvenuti F, Pagliarani S, et al. Antioxidant properties of a novel
322		phycocyanin extract from the blue-green alga Aphanizomenon flos-aquae. Life Sci
323		2004:75(19):2353–2362.
324	3.	Stephens E, Ross IL, King Z, et al. An economic and technical evaluation of microalgal
325		biofuels. Nat Biotechnol 2010:28(2):126–128.
326	4.	Scott SA, Davey MP, Dennis JS, et al. Biodiesel from algae: challenges and prospects.
327		Curr Opin Biotechnol 2010:21(3):277–286.
328	5.	Mata TM, Martins AA, Caetano NS. Microalgae for biodiesel production and other
329		applications: A review. <i>Renew Sustain Energy Rev</i> 2010:14(1):217–232.
330	6.	Singh A, Olsen SI. A critical review of biochemical conversion, sustainability and life
331		cycle assessment of algal biofuels. <i>Appl Energy</i> 2011: 88(10):3548–3555.
332	7.	Letcher PM, Lopez S, Schmieder R, et al. Characterization of Amoeboaphelidium
333		protococcarum, an algal parasite new to the cryptomycota isolated from an outdoor
334		algal pond used for the production of biofuel. <i>PLoS One</i> 2013 8(2):e56232.
335	8.	Smith VH, Sturm BSM, Denoyelles FJ, Billings SA. The ecology of algal biodiesel
336		production. <i>Trends Ecol Evol</i> 2010:25(5):301–9.
337	9.	Tilman D, Reich PB, Knops J, et al. Diversity and productivity in a long-term grassland
338		experiment. <i>Science</i> 2001:294(5543):843–845.

Belton, C.S. The Ecology of Invasions by Animals and Plants. 1958. University of Chicago
Press, Chicago, USA.

341	11.	May, R.M. Thresholds and breakpoints in ecosystems with a multiplicity of stable
342		states. Nature 1977:269, 471–477.
343	10.	Cardinale BJ. Biodiversity improves water quality through niche partitioning. Nature
344		2011: 472(7341):86–89.
345	11.	Hector A, Bazeley-White E, Loreau M, et al. Overyielding in grassland communities:
346		testing the sampling effect hypothesis with replicated biodiversity experiments. Ecol
347		Lett 2002:5(4):502–511.
348	12.	Schmid B, Hector A, Saha P, Loreau M. Biodiversity effects and transgressive
349		overyielding. J Plant Ecol 2008:1(2):95–102.
350	13.	Behl S, Donval A, Stibor H. The relative importance of species diversity and functional
351		group diversity on carbon uptake in phytoplankton communities. Limnol Oceanogr
352		2011: 56(2):683–694.
353	14.	Perrow MR, Meijer M-L, Dawidowicz P, Coops H. Biomanipulation in shallow lakes:
354		state of the art. Hydrobiologia 1997:342-343:355–365.
355	15.	Kazamia E, Aldridge DC, Smith AG. Synthetic ecology – A way forward for sustainable
356		algal biofuel production? <i>J Biotechnol</i> 2012:162(1):163–169.
357	16.	Duffy JE. Biodiversity and ecosystem function: the consumer connection. <i>Oikos</i>
358		2002:99(2):201–219.
359	17.	Shurin JB, Abbott RL, Deal MS, et al. Industrial-strength ecology: trade-offs and
360		opportunities in algal biofuel production. <i>Ecol Lett</i> 2013:16(11):1393–1404.

361	18.	Legrand C, Rengefors K, Fistarol GO, Granéli E. Allelopathy in phytoplankton -
362		biochemical, ecological and evolutionary aspects. <i>Phycologia</i> 2003:42(4):406–419.
363	19.	Heisler J, Glibert PM, Burkholder JM, et al. Eutrophication and harmful algal blooms: A
364		scientific consensus. <i>Harmful Algae</i> 2008:8(1):3–13.
365	20.	Ferrier MD, Butler BR, Terlizzi DE, Lacouture R V. The effects of barley straw (Hordeum
366		vulgare) on the growth of freshwater algae. <i>Bioresour Technol</i> 2005:96(16):1788–
367		1795.
368	21.	Kazamia E, Czesnick H, VTT Nguyen, et al. Mutualistic interactions between vitamin
369		B ₁₂ -dependent algae and heterotrophic bacteria exhibit regulation. <i>Environ Microbiol</i>
370		2012:14(6):1466–1476.
371	22.	Sharifah EN, Eguchi M. The phytoplankton Nannochloropsis oculata enhances the
372		ability of Roseobacter clade bacteria to inhibit the growth of fish pathogen Vibrio
373		anguillarum. PLoS One 2011:6(10):e26756.
374	23.	Wagner-Döbler I, Ballhausen B, Berger M, et al. The complete genome sequence of
375		the algal symbiont Dinoroseobacter shibae: a hitchhiker's guide to life in the sea. ISME
376		J 2010:4(1):61–77.
377	24.	Amin SA, Green DH, Hart MC, et al. Photolysis of iron-siderophore chelates promotes
378		bacterial-algal mutualism. PNAS 2009:106(40):17071–17076.

- Schindler DW, Hecky RE, Findlay DL et al. Eutrophication of lakes cannot be controlled
 by reducing nitrogen input: results of a 37-year whole-ecosystem experiment. *PNAS*2008:105(32):11254–11258.
- 382 26. Conley DJ, Paerl HW, Howarth RW, et al. Ecology. Controlling eutrophication: nitrogen
 383 and phosphorus. *Science* 2009:323(5917):1014–1015.
- Pate R, Klise G, Wu B. Resource demand implications for US algae biofuels production
 scale-up. *Appl Energy* 2011: 88(10):3377–3388.
- 386 28. Ortiz-Marquez JCF, Do Nascimento M, Dublan M de LA, Curatti L. Association with an
- 387 ammonium-excreting bacterium allows diazotrophic culture of oil-rich eukaryotic
- 388 microalgae. *Appl Environ Microbiol* 2012:78(7):2345–2352.
- 389 29. Clarens AF, Resurreccion EP, White MA, Colosi LM. Environmental life cycle
- 390 comparison of algae to other bioenergy feedstocks. *Environ Sci Technol*
- 391 2010:44(5):1813–1819.
- 392 30. Pittman JK, Dean AP, Osundeko O. The potential of sustainable algal biofuel
- 393 production using wastewater resources. *Bioresour Technol* 2011:102(1):17–25.
- 394 31. Park Y, Je K-W, Lee K, et al. Growth promotion of *Chlorella ellipsoidea* by co-
- inoculation with *Brevundimonas* sp. isolated from the microalga. *Hydrobiologia* 2007
 :598(1):219–228.
- 397 32. Do Nascimento M, Dublan MDLA, Ortiz-Marquez JCF, Curatti L. High lipid productivity
- 398 of an Ankistrodesmus-Rhizobium artificial consortium. Bioresour Technol 2013:
- 399 146:400–407.

400	33.	Smith D, Douglas A. The biology of symbiosis. London: Edward Arnold; 1987. p. 302.
401	34.	Kanesaki Y, Shiwa Y, Tajima N, et al. Identification of substrain-specific mutations by
402		massively parallel whole-genome resequencing of Synechocystis sp. PCC 6803. DNA
403		Res 2012: 19(1):67–79.
404	35.	Trautmann D, Voss B, Wilde A, et al. Microevolution in cyanobacteria: re-sequencing a
405		motile substrain of Synechocystis sp. PCC 6803. DNA Res 2012:19(6):435–448.
406	36.	Pande S, Merker H, Bohl K, et al. Fitness and stability of obligate cross-feeding
407		interactions that emerge upon gene loss in bacteria. <i>ISME J.</i> 2013:1–10.
408	37.	Gore J, Youk H, van Oudenaarden A. Snowdrift game dynamics and facultative
409		cheating in yeast. <i>Nature</i> 2009:459(7244):253–256.
410	38.	Nadell CD, Foster KR, Xavier JB. Emergence of spatial structure in cell groups and the
411		evolution of cooperation. <i>PLoS Comput Biol</i> 2010:6(3):e1000716.
412	39.	Brenner K, You L, Arnold FH. Engineering microbial consortia: a new frontier in
413		synthetic biology. Trends Biotechnol 2008:26(9):483–489.
414	40.	Hosoda K, Suzuki S, Yamauchi Y, et al. Cooperative adaptation to establishment of a
415		synthetic bacterial mutualism. PLoS One 2011:6(2):e17105.
416	41.	Kerner A, Park J, Williams A, Lin XN. A programmable Escherichia coli consortium via
417		tunable symbiosis. PLoS One 2012:7(3):e34032.
418	42.	Kubo I, Hosoda K, Suzuki S, et al. Construction of bacteria-eukaryote synthetic
419		mutualism. <i>Biosystems</i> 2013:113(2):66–71.

- 420 43. West SA, Diggle SP, Buckling A, et al. The Social Lives of Microbes. *Annu Rev Ecol Evol*421 *Syst* 2007:38(1):53–77.
- 422 44. Sabra W, Dietz D, Tjahjasari D, Zeng A-P. Biosystems analysis and engineering of
- 423 microbial consortia for industrial biotechnology. *Eng Life Sci* 2010:10(5):407–421.
- 424 45. Mooij PR, Stouten GR, Tamis J, et al. Survival of the fattest. *Energy Environ Sci*425 2013:6(12):3404.