

1 Iron and Harmful Algae Blooms: Potential Algal-
2 Bacterial Mutualism between *Lingulodinium polyedrum*
3 and *Marinobacter algicola*

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22 **Abstract**

23 Phytoplankton blooms can cause acute effects on marine ecosystems either due to their
24 production of endogenous toxins or due to their enormous biomass leading to major impacts on
25 local economies and public health. Despite years of effort, the causes of harmful algal blooms
26 (HAB) are still not fully understood. Our hypothesis is that bacteria that produce photoactive
27 siderophores may provide a bioavailable form of iron to commensally associated phytoplankton,
28 which could in turn affect algal growth and bloom dynamics. Here we report a laboratory-based
29 study of binary cultures of the dinoflagellate *Lingulodinium polyedrum*, a major HAB species,
30 with *Marinobacter algicola* DG893, a phytoplankton-associated bacterium that produces the
31 photoactive siderophore vibrioferin. Comparing binary cultures of *L. polyedrum* with both the
32 wild type and the vibrioferin minus mutant of *M. algicola* shows that bacteria are necessary to
33 promote dinoflagellate growth and that this growth promotion effect is at least partially related to
34 the ability of the bacterium to supply bioavailable iron via the siderophore vibrioferin. These
35 results support the notion of a carbon for iron mutualism in some bacterial-algal interactions.

36

37 **Introduction**

38 Phytoplankton blooms are a frequent phenomenon in the coastal regions of every
39 continent in the world. Certain phytoplankton algae occurring in mass proliferations produce
40 toxins. Such harmful algal blooms (HABs) can, directly or indirectly, impact marine ecosystems
41 with repercussions on local economies and public health (Lewitus et al., 2012). Direct effects on
42 human health are often related to the consumption of shellfish that have ingested toxic
43 phytoplankton, resulting in the accumulation of toxins and, consequently upon human
44 consumption, paralytic, diarrheic, and neurotoxic poisoning which can sometimes be fatal
45 (Anderson 1994: Honner et al., 2012). HABs also have indirect impacts such as impairment of
46 water quality leading to losses in the tourism and recreational sectors. Even phytoplankton
47 blooms that do not produce toxins can be detrimental and lead to ecological impacts such as the
48 displacement of indigenous species, habitat alteration, or oxygen depletion (Glibert et al., 2005).
49 The economic effects caused by HABs in the U.S. alone were estimated at \$82 million per year
50 in 2006 (Hoagland and Scatasta, 2006). Current strategies to reduce health-related impacts due to
51 HABs are based on frequent coastal monitoring and early detection of HAB species and toxin
52 levels. Nevertheless, the factors leading to the origins of HABs are still not well understood.
53 Consequently, there is no strategy for their ultimate prevention.

54 There is an ongoing debate whether the factors leading to HABs are natural or
55 anthropogenic. Oceanographic factors and global climate change including and impacting
56 upwelling, reversal and relaxation of winds may be the major drivers initiating phytoplankton
57 blooms (Roegner et al., 2002: Tweddle et al., 2010). HABs have also been proposed to be the
58 consequence of human activities including eutrophication, changes in land use and agriculture,
59 overfishing, and ballast water discharge (Gilbert 2005). Others consider that better analytics

60 results in increased detection of HAB species (Burkholder, 1998). Nevertheless, HABs are a
61 global threat to human health, fisheries and aquaculture resources. While numerous studies
62 concentrate on the causes of HABs, few have considered that bacterial species coexisting with
63 microalgae could contribute to their development. However since Bell and Mitchell reported that
64 specific bacterial communities occur and microbial activity was altered in the so-called
65 phycosphere (Bell and Mitchell, 1972; Bell and Lang, 1974), more and more studies suggest that
66 such algal-bacterial interactions are very specific and important (Azam and Malfatti, 2007; Amin
67 et al., 2015; Bertrand et al., 2015; Ramanan et al., 2016; Seymour et al. 2017). A plausible
68 hypothesis suggests that the mutualistic association of some phytoplankton and bacteria is based
69 on nutrient exchange. Nitrogen and phosphorus are the best studied in this regard, although the
70 molecular nature of interactions involving these nutrients is not well understood (Hallegraeff and
71 Gollasch, 2006). Alternatively specific bacteria may affect algal growth and bloom dynamics by
72 their control of iron, a trace element which is often growth limiting to phytoplankton in the
73 marine environment (Miethke and Marahiel, 2007; Maldonado et al., 2005; Croot and Heller,
74 2012).

75 Iron is an essential element for all living organisms including phytoplankton and bacteria
76 due to its involvement in photosynthesis and respiration. Despite iron being the fourth most
77 abundant element on the Earth, its bioavailability in the marine environment is extremely low
78 due to its poor solubility under the mildly alkaline aerobic conditions present in the ocean
79 (Martin and Fitzwater, 1988; Wu and Luther III, 1994). To overcome this low bioavailability,
80 bacteria and fungi have evolved sophisticated systems to produce high-affinity iron-chelating
81 compounds called siderophores to acquire, transport, and process this essential metal ion (Sandy
82 and Butler, 2009). The major role of siderophores is to bind mineral phases of iron and to deliver

83 the iron siderophore complex to specific outer membrane receptors on microbial cells. Several
84 hundred siderophores have been isolated and extensively studied with respect to their synthesis,
85 structures and transport mechanisms over the last three decades (Yamamoto et al., 1994; Challis,
86 2005; Sandy and Butler, 2009; Raymond et al., 2015). While much research has been done on
87 terrestrial siderophores, the study of marine siderophores is less extensive and only a relatively
88 few have been fully elucidated (Vraspir and Butler, 2009). Nevertheless one of the two major
89 attributes that seem to distinguish marine siderophores from those of terrestrial origin is the
90 tendency of the former to contain a α - and β -hydroxy acid group in their iron-binding domain
91 (Barbeau et al., 2001, 2002; Küpper et al., 2006). The significance of the presence of these
92 functional groups in siderophores is in their ability to make the iron-siderophore complex
93 photoreactive. The chelated Fe(III) in such siderophores is reduced in the presence of sunlight
94 via an internal redox process to release Fe(II), a more soluble form of iron (Amin et al., 2009). It
95 has been proposed that sunlight-driven reduction of the Fe(III) would transiently produce Fe(II)
96 which might be utilized not only by the siderophore-producing bacteria themselves but also by
97 non-siderophore producing bacteria and other organisms such as phytoplankton (Maldonado et
98 al., 2005; Naito et al., 2008; Amin et al., 2009, 2012).

99 While iron acquisition by bacteria is well understood, that of phytoplankton remains less
100 so. There are data to support the possibility of a variety of iron uptake mechanisms being
101 operative simultaneously depending on the species involved. These include an iron-reductive
102 route via a cell surface reductase, a direct xenosiderophore-mediated mechanism and cell
103 surface-enhanced processes (Sutak et al., 2012; McQuaid et al., 2018). However as of yet there
104 are no well documented examples, with some exceptions among the cyanobacteria, where
105 phytoplankton have been shown to produce their own siderophores for iron acquisition (Raven,

106 2013). Thus how phytoplankton effectively acquire this metal from the low concentration iron
107 environment is still unclear.

108 Our hypothesis is that algal growth and bloom dynamics may be affected by the increased
109 bioavailability of iron engendered by the presence of photoreactive siderophores produced by
110 mutualistic bacteria. Previously, we studied a bloom of the dinoflagellate, *Lingulodinium*
111 *polyedrum*, at the Scripps Pier (San Diego, CA, USA) in 2011. *L. polyedrum* is one of the HAB
112 species known to produce Yessotoxins, a group of polyether toxins which can accumulate in
113 shellfish and show high toxicity to mice via intraperitoneal injection (Tubaro et al., 2004). As the
114 first step to search for an association of phytoplankton with specific bacterial photoreactive
115 siderophore producers (producing petrobactin, aerobactin, and vibrio ferrin siderophores), we
116 monitored both the population of *L. polyedrum* and the bacterial siderophore producers before,
117 during, and after the 2011 bloom. The results showed that both *L. polyedrum* and the bacterial
118 siderophore producers simultaneously increased and decreased during the bloom period
119 (Yarimizu et al., 2014). At the *L. polyedrum* bloom maximum, the total number of bacterial
120 producers of photoreactive siderophore reached their maximum accounting for roughly 9% of the
121 total bacterial population suggesting that such high abundance of photoreactive siderophore
122 producers could potentially provide bioavailable iron to the phytoplankton in this environment.
123 Furthermore, when the PCR-derived amplicons were sequenced, a phylogenetic tree constructed
124 from the sequencing results showed that the community of siderophore producers in pre-bloom
125 was statistically significantly different than those found during and after the bloom by UniFrac
126 analysis suggesting that this particular bacterial community could be involved in bloom
127 initiation.

128 As a proof of concept, the present study was performed using laboratory culture data to
129 correlate the association of the vibrioferin siderophore producing bacterium, *Marinobacter*
130 *algicola* DG893 and the dinoflagellate *L. polyedrum* with iron as a nutrient. It is hoped that
131 finding an algal-bacterial mutualism based on iron availability may be useful for a more
132 thorough understanding of the mechanisms of HAB formation.

133

134 **Materials and Methods**

135 Trace iron cleaning procedures: The container cleaning technique was adopted from Bruland and
136 Franks (1979). All plastic and glass containers were washed with pure water and soaked in 3N
137 hydrochloric acid for at least two weeks at ambient temperature. The acid-washed containers
138 were rinsed with Milli-Q water and dried in a laminar-flow air bench. For those experiments
139 requiring sterile conditions, acid-washed containers were placed in pouches and autoclaved at
140 121°C for 30 minutes followed by a 45 minute dry cycle. Alternatively, purchased sterile
141 containers Nunc™ Cell Culture Treated Flasks with Filter Caps were used.

142 Growth media: Of eight sources of natural waters screened, Pacific Ocean oligotrophic water
143 was initially selected for our use due to its extremely low dissolved iron concentration (pM).
144 This was replaced later by Scripps Pier seawater because of its easier accessibility and
145 comparable *L. polyedrum* growth patterns observed in the two sources of seawater. Seawater
146 from Scripps Pier (32.153°N, 117.115°W, San Diego, CA) was collected (pH 8.2-8.4, 980
147 mOsm/Kg H₂O, total dissolved iron 3-4 nM) and filtered immediately through 0.22 mm pore size
148 membrane. The filtered seawater was mixed with approximately 0.005% metal free hydrochloric
149 acid, autoclaved for 30 minutes, cooled at ambient temperature for a day, and stored in 5°C until
150 use. The pH of the autoclaved seawater was ensured to be between 8.0 and 8.2 at ambient
151 temperature. L1 nutrient was added to sterile seawater per manufacturer's instruction (L1
152 medium). For those cultures requiring controlled iron concentrations in media, L1 trace element
153 solution without FeCl₃ and Na₂EDTA was prepared in house (Guillard and Hargraves, 1993) and
154 added along with other L1 nutrients (nitrate, phosphate, silicate, vitamins) to sterile seawater
155 (L1-Fe medium). Serial dilutions were made on L1 medium with L1-Fe medium to prepare
156 growth medium with defined total iron concentrations, [Fe]_T.

157 Artificial Sea Water (ASW): The component solutions were prepared in three separate
158 containers, one with 15 g NaCl, 0.75 g KCl, 1 g NH₄Cl in approximately 500 mL water, one with
159 12.4 g MgSO₄·7H₂O in small volume of water, and one with 3.0 g of CaCl₂·2H₂O in a small
160 volume of water. The three solutions were mixed and 0.1 g disodium β-glycerol phosphate was
161 added. The pH was adjusted to 8.0-8.2 and volume was adjusted to 1L. The solution was
162 autoclaved at 121°C for 30 minutes.

163 Algal maintenance and growth monitoring: The *L. polyedrum* strain used in this study was
164 isolated from Venice Beach, California, and kindly provided to us by Avery Tatters (University
165 of Southern California). *L. polyedrum* cells were maintained in sterile one liter Erlenmeyer flasks
166 containing L1 medium and capped with a ventilation sponge. Cultures were exposed to 100 µmol
167 photons m⁻² s⁻¹ on a 12-hour alternating light and dark cycle at a temperature of 20±2 °C
168 (standard growth condition). The upper portion of a culture containing healthy cells was diluted
169 to 1/5 with L1 medium every three weeks. Algal growth was monitored by direct cell count
170 using an inverted microscope (Nikon Eclipse TE2000-U, 4x objective). To prepare cells in
171 controlled iron medium, cultures containing 10⁴-10⁵ cells per mL was placed in a 15-mL Falcon
172 tubes and centrifuged for a few seconds to pellet cells which were gently washed three times and
173 re-suspended to an appropriate volume with controlled iron medium.

174 “Axenic” *L. polyedrum* preparation: Non-axenic *L. polyedrum* cells were grown to 10⁵ cells/mL
175 in L1 medium. Antibiotics 0.1% (v/v) (Penicillin 5units/mL, Streptomycin 5 µg/mL, Neomycin
176 10 µg/mL) were added to a non-axenic culture and incubated for 24 hours under the standard
177 growth condition. Antibiotic-treated cultures were placed in 15 mL Falcon tubes and gently
178 centrifuged for a few seconds. The pelleted cells were washed three times and re-suspended with
179 L1 medium containing 0.05% (v/v) antibiotics in a sterile culture flask to appropriate volume.

180 Antibiotics 0.05% (v/v) was added every 3 days to maintain the culture “axenic”. The absence of
181 cultureable bacteria in the antibiotic treated *L. polyedrum* was tested by spreading a 10 μ L
182 sample of the culture on a marine broth plate (5 g/L peptone, 1 g/L yeast extract, 15 g/L agar in
183 75% seawater) followed by incubation at 25°C for 2-3 days. The absence of any visible bacterial
184 colonies was considered indicative of an “axenic” culture of *L. polyedrum*.

185 Bacteria strains and growth monitoring: The model bacterium used was *Marinobacter algicola*
186 DG893 (GenBank code NZ_ABCP0000000.1) since it has been isolated by Green et al. (2004)
187 and well characterized as a producer of the photoactive siderophore vibrio ferrin (Amin et al.,
188 2007). The mutant *ΔpvsAB*-DG893 was made as described by Amin et al. (2012) by 872 base
189 pair deletion from the wild type to knock out two siderophore biosynthesis genes, *pvsA* and *pvsB*.
190 To monitor bacterial growth, two methods were used: Optical density at 600 nm was applied to
191 apparently turbid samples which generally contained $\geq 10^7$ /mL of bacterial cells. Enumeration of
192 bacteria by serial dilution was applied for samples containing lower numbers of bacteria cells
193 ($<10^7$ /mL). For enumeration techniques, serial dilution was made on those samples with sterile
194 seawater, each diluted sample was spread on marine broth plates, and the number of colony
195 forming units (CFU) recorded from the lowest diluted sample plate. CFU in original sample per
196 mL was calculated by (CFU in diluted sample) / (volume plated in mL) x dilution factor.

197 Stock bacteria solution: One liter of marine broth was prepared (5 g/L peptone, 1 g/L yeast
198 extract, 15 g/L agar in 75% seawater, pH 8.2, sterile) and dispensed into sterile tubes with a 5
199 mL volume. A few colonies of bacteria picked from stock plates were transferred into the marine
200 broth tubes and shaken at 25°C for 1-2 days until they reached log phase. The cells were
201 collected by centrifugation, washed three times and re-suspended in sterile medium to an optical
202 density (OD600) of 0.3 (stock bacteria solution).

203 CAS-dye assay: The method was adopted from Alexander and Zuberer (1991) to detect
204 siderophore production by DG893. Using acid washed glassware, the following stock solutions
205 were prepared: 10 mM HDTMA in water, 2 mM CAS in water, 1 mM $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ in water, 50
206 mM piperazine anhydrous buffer (pH5.6), and 0.2 M 5-sulfosalicylic acid in water. CAS solution
207 was prepared by pouring a mixture of 15 mL of CAS and 3 mL of $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ slowly into 12 mL
208 of HDTMA followed by addition of 50 mL of piperazine. The final volume was adjusted to 200
209 mL with water. CAS shuttle solution was prepared by adding 20 μL of 5-sulfosalicylic acid to
210 every mL of CAS solution (used within one day of preparation). Samples to be tested were
211 centrifuged to remove particles and 0.5 mL supernatant was mixed with 0.5 mL of CAS shuttle
212 solution followed by incubation in the dark at ambient temperature for 10 minutes. A color
213 change from dark purple to clear red indicated siderophore production.

214 Materials and reagents: The following materials were purchased from Sigma-Aldrich: Chrome
215 azurol S (CAS, C-1018, MW605.28), antibiotics (Penicillin/Streptomycin/Neomycin, P4083-100
216 mL), peptone (P-1265), iron(III) chloride hexahydrate ($\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$, 10025-77-1, MW270.30),
217 manganese(II) chloride tetrahydrate ($\text{MnCl}_2 \cdot 4\text{H}_2\text{O}$, 13446-34-9, MW197.91),
218 hexadecyltrimethylammonium bromide (HDTMA, H6269-100G, MW364.45), 1,4-
219 diazacyclohexane, diethylenediamine (Piperazine, Sigma-Aldrich, P45907-100G, MW 86.14), 5-
220 sulfosalicylic acid (390275, MW218. 18), zinc sulfate heptahydrate ($\text{ZnSO}_4 \cdot 7\text{H}_2\text{O}$, 1088830500,
221 MW287.54), cobalt(II) chloride hexahydrate ($\text{CoCl}_2 \cdot 6\text{H}_2\text{O}$, 8025400010, MW129.83), copper(II)
222 sulfate pentahydrate ($\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$, 209198-5G, MW249.69), sodium molybdate dehydrate
223 ($\text{Na}_2\text{MoO}_4 \cdot 2\text{H}_2\text{O}$, 331058-5G, MW241.95), selenous acid (H_2SeO_3 , 211176-10G, MW128.97),
224 nickel(II) sulfate hexahydrate ($\text{NiSO}_4 \cdot 6\text{H}_2\text{O}$, 227676-100G, MW262.85), sodium orthovanadate
225 (Na_3VO_4 , 450243-10G, MW183.91), potassium chromate (K_2CrO_4 , 216615-100G, MW194.19),

226 salicylaldoxime (SA, 84172-100G). The following materials were purchased from Fisher
227 Scientific: trace metal-free hydrochloric acid (Optima, A466-250), yeast extract (BP9727-500),
228 agar (BP1423-500), Casamino Acids (BP1424-500), NuncTM Cell Culture Treated Flasks with
229 Filter Caps (12-565-57), ethylenediaminetetraaceticacid, tetrasodium salt dihydrate
230 (Na₄EDTA·2H₂O, BP121-500, MW416.2), boric acid (AAJ67202A1). Other materials were
231 purchased as follows: 0.22 μ m filter membrane (Millipore, MillexGV, SLGV033RS), L1
232 medium kit (Bigelow, MKL150L), pure water (Barnstead water system, 18.2 m Ω). Vibrioferin
233 (VF, MW434.35) was isolated and purified in house according to Amin et al. (2007).

234

235 **Results**

236 Bacterial growth: The growth of DG893 and mutant *ΔpvsAB*-DG893 in L1 medium containing
237 different total iron concentrations $[Fe]_T$ (1,000, 100, 10, and 0 nM) was monitored. All media
238 were divided in two parts, one maintaining [EDTA] at 10,000 nM (L1 nutrient level) with
239 variable $[Fe]_T$, and the other reducing [EDTA] along with $[Fe]_T$ to maintain molar ratio of Fe:
240 EDTA of 1:1. All media were prepared with and without sterile Casamino Acids. In each
241 medium, ca. 10^6 cells/mL of DG893 or mutant *ΔpvsAB*-DG893 were added and shaken in the
242 dark at 30°C. The bacterial growth in each medium was monitored for 5 days by optical density
243 at 600 nm. Neither WT DG893 nor the mutant grew in simple seawater or L1-Fe medium
244 without a carbon source (Figure 1A). Addition of iron to seawater or L1-Fe medium also did not
245 maintain growth (Figure 1B). However upon addition of Casamino Acids, both DG893 and the
246 mutant grew and their growth increased with increasing $[Fe]_T$ in media (Figure 1C). There was
247 no remarkable difference in growth pattern between DG893 and the mutant, consistent with the
248 report by Amin et al. (2012). In summary, both DG893 and the mutant require a carbon source to
249 grow to a detectable level in seawater medium.

250 Siderophore production and total iron concentration $[Fe]_T$: The ability to produce the siderophore
251 vibrioferrin (VF) by DG893 but not by the mutant was confirmed by the CAS-dye assay. Since
252 siderophore biosynthesis is generally turned on under low iron concentrations and turned off
253 under high concentrations (Braun, 1995 and 1998; Miethke and Marahiel, 2007), WT was grown
254 in L1 medium containing various $[Fe]_T$ with 0.3% Casamino Acids at 30°C in dark for 24 hours
255 to find the approximate $[Fe]_T$ under which DG893 produces VF. The WT did not produce VF in
256 the media with $[Fe]_T \geq 10 \mu M$ but did so with a $[Fe]_T$ of 1 μM or less. The mutant *ΔpvsAB*-
257 DG893 did not show signs of siderophore production under any of the conditions tested. The

258 results suggest that under normal seawater conditions, where $[Fe]_T$ is estimated to be nM to pM,
259 that siderophore biosynthesis in DG893 is likely turned on.

260 Axenic culture preparation: The difficulty in preparing and maintaining axenic cultures of many
261 marine algae has been reported in the past (Green et al., 2004; Jauzein et al., 2015; Liu et al.,
262 2017). In fact, Lupette et al. (2016) noted that completely axenic cultures of a model green algal
263 species could not be maintained despite the use of antibiotic treatment protocols. We were
264 partially successful in our efforts to prepare axenic *L. polyedrum*. Initially, non-axenic *L.*
265 *polyedrum* cultures containing 10^5 cells/mL were treated with antibiotics (1% v/v Penicillin 50
266 units/mL, Streptomycin 50 μ g/mL, Neomycin 100 μ g/mL) for 24 hours under the normal growth
267 conditions. The treated cells were washed and re-suspended with sterile L1 media and while the
268 “axenicity” of the culture at this point was confirmed, bacterial colonies reappeared after only a
269 few days. An attempt was made to keep 1% antibiotics in the culture longer than 24 hours,
270 however, dinoflagellate cells began to disintegrate after 2 days suggesting that 1% antibiotics
271 must be removed from the culture within 24 hours. The fact that the negative control and sterile
272 L1 medium without phytoplankton cells remained bacteria-free for weeks ruled out a possibility
273 of culture contamination. We believe that the antibiotics are most likely eliminating only free-
274 living bacteria leaving attached bacteria unaffected. The next effort of preparing axenic cultures
275 was made by immediate dilution of antibiotic-treated cultures. The rationale of this attempt was
276 to minimize attached bacteria on *L. polyedrum* cells by dilution. The antibiotic treated *L.*
277 *polyedrum* culture was diluted 200-fold with sterile L1 medium and incubated under standard
278 growth conditions. The diluted culture remained bacteria-free for the first 10 days, however
279 microbial growth was observed on the 12th day as the number of *L. polyedrum* cells also
280 increased. Although this dilution method suggested that the culture stayed bacteria free for up to

281 10 days, the *L. polyedrum* cell count was too dilute to be useful for further study. The next
282 attempt at axenic culture preparation was made with three sequential antibiotic treatments. The
283 non-axenic culture was treated with 1% antibiotics for 24 hours, washed three times, and re-
284 suspended with sterile L1 medium. The cells were allowed to recover under normal growth
285 conditions for a day and then again treated with 1% antibiotics. This procedure was repeated
286 three times. This procedure also failed as the dinoflagellate cells began to disintegrate. Finally,
287 using a similar procedure but reducing the concentration of antibiotics to 0.1% for the first
288 treatment and adding 0.05% subsequently every 2-3 days resulted in *L. polyedrum* cultures
289 without visible cell damage that appeared to be free of free-living bacteria for more than a
290 month.

291 *L. polyedrum* growth screening: A matrix containing thirty cultures of *L. polyedrum* in L1
292 medium containing different numbers of DG893 cells in the initial inoculum (10^7 , 10^6 , 10^5 , 10^4 ,
293 10^3 , 0 CFU/mL) and different $[Fe]_T$ (10^4 , 10^3 , 10^2 , 10^1 , and 0 nM) was set up. The same matrix
294 was set up with the mutant $\Delta pvsAB$ -DG893. Bacterial growth was monitored by enumeration of
295 bacteria by serial dilution and phytoplankton growth was monitored by direct cell counts under
296 the microscope. Regardless of the starting bacterial cell number, both DG893 wild type and
297 mutant grew in all the binary culture matrices over 28 days to eventually reach a bacterial
298 population of ca. 10^5 CFU/mL. The group of media containing 1,000 nM $[Fe]_T$ showed the
299 greatest growth of *L. polyedrum* regardless of the starting bacterial inoculum (Figure 2A). The
300 group of media containing ≤ 10 nM $[Fe]_T$ showed very slow *L. polyedrum* growth regardless of
301 starting bacterial inoculum. No significant differences in *L. polyedrum* growth co-cultured with
302 DG893 or the mutant were observed, which left a question as to whether VF secreted by DG893
303 would have a significant effect on *L. polyedrum* growth. It was speculated that under these

304 growth conditions, the amount of VF being produced by DG893 was insufficient to see
305 detectable changes in *L. polyedrum* growth. This was later confirmed (*vide infra*) when excess
306 VF artificially added to the culture produced pronounced *L. polyedrum* growth. Finally, the
307 growth pattern of axenic *L. polyedrum* was similar to that of a non-axenic culture. This
308 observation left open another question as to whether DG893 and *L. polyedrum* were
309 commensally associated. However this also was confirmed by experiments described below
310 where after subsequent culturing the “axenic” *L. polyedrum* ceased to grow while binary *L.*
311 *polyedrum* DG893 kept growing exponentially (Figure 3).

312 Subsequent batch culture growth of *L. polyedrum* in DG893 binary culture: *L. polyedrum* grown
313 in L1 media containing various [Fe]_T and DG893 (Figure 2A) for 28 days were diluted with
314 appropriate media to adjust *L. polyedrum* to 10² cells/mL in the 2nd subsequent batch culture. The
315 *L. polyedrum* growth in this 2nd subsequent culture was monitored for another 28 days under
316 each growth condition. The result showed that the 2nd subsequent batch culture *L. polyedrum* in
317 the higher two [Fe]_T (10,000 and 1,000 nM) maintained their exponential growth while those in
318 the lower three [Fe]_T did not grow (Figure 2B). Although the cells under the latter conditions did
319 not completely die out, the few cells that survived had their swimming activity under the
320 microscope clearly reduced.

321 Axenic *L. polyedrum* culture and rescue: The growth of “axenic” and binary cultures of *L.*
322 *polyedrum* with DG893 in L1 was compared over several subsequent batch cultures. No
323 significant difference in growth was observed in the 1st culture, however, a clear difference was
324 observed in the 2nd subsequent batch culture of *L. polyedrum* with and without DG893. While *L.*
325 *polyedrum* with DG893 grew fully in both the 1st and 2nd subsequent batch cultures, “axenic” *L.*
326 *polyedrum* ceased to grow (Figure 3). When DG893 was added to the 2nd subsequent “axenic”

327 batch culture, *L. polyedrum* cells began to regrow exponentially demonstrating that DG893 could
328 rescue “axenic” *L. polyedrum*.

329 Non-axenic *L. polyedrum* growth and rescue: The non-axenic *L. polyedrum* was maintained in
330 L1 medium and assumed to contain not only siderophore-producing bacteria but also other
331 unknown bacteria. The non-axenic culture was divided in two parts, one diluted to 1/5 with L1
332 medium and other diluted to 1/5 with L1-Fe medium (batch subculture 1). When the cell count
333 reached approximately 10^4 cells/mL, both cultures were further diluted with appropriate medium.
334 Therefore, one culture maintained $[Fe]_T$ at L1 level throughout the subsequent batch cultures
335 while the other contained successively reduced $[Fe]_T$. The *L. polyedrum* growth was monitored
336 for 28 days for each subculture. After the 6th subculture where $[Fe]_T$ in the medium was
337 estimated to be 2 nM, *L. polyedrum* stopped growing completely (Figure 4). On 29th day of the
338 6th subculture, an iron supplement to give a $[Fe]_T$ of 5850 nM was added to the culture and
339 growth was continued to be monitored for additional 28 days. The results indicate that the *L.*
340 *polyedrum* culture which stopped growing under 2 nM $[Fe]_T$, could be rescued by addition of
341 iron after 28 days (Figure 5).

342 *L. polyedrum* growth in DG893 supernatant: Earlier it was stated that when *L. polyedrum* was
343 co-cultured with either DG893 or the mutant $\Delta pvsAB$ -DG893, little difference was observed in
344 its growth pattern. We speculated that this may have been due to the small amount of VF
345 produced by the equilibrium population of DG893 in the binary culture. To test this idea,
346 samples of the supernatant of the media from the WT DG893 grown under ideal conditions and
347 containing detectable amounts of VF by CAS-dye assay, as well as the mutant $\Delta pvsAB$ -DG893
348 which did not, respectively, were added to *L. polyedrum* cultures and growth under these
349 different conditions was compared. It should be noted that while we traditionally grow DG893

350 and the mutant in marine broth containing Casamino Acids, these were found to be toxic to *L.*
351 *polyedrum* cells. In this experiment, we instead used ASW with succinic acid (at from 0.001% to
352 0.1%) as a carbon source (pH 8.2) to grow DG893 and the mutants prior to their supernatant
353 being added to *L. polyedrum* cultures. *L. polyedrum* growth was enhanced the most by the
354 addition of the DG893 supernatant. The mutant supernatant also showed a slightly positive effect
355 on *L. polyedrum* growth compared to media without bacterial supernatant, however the degree of
356 growth was not as large as that induced by the DG893 supernatant (Figure 6). These results
357 suggest that VF has potential influence on *L. polyedrum* growth, but that other extracts from
358 bacteria (vitamin B₁₂?) may also have a potential effect on their growth. Further experiments
359 were performed using purified VF itself. When purified VF at 57 μ M was directly added to a *L.*
360 *polyedrum* culture in L1 media containing 100 nM [Fe]_T, *L. polyedrum* reached cell counts
361 approximately 1.5 times as large as those in media without VF.

362 DG893 growth in *L. polyedrum* supernatant: “Axenic” *L. polyedrum* cells were resuspended in
363 L1 media containing 100 nM, 1,000 nM, 10,000 nM [Fe]_T and incubated under the standard
364 growth conditions for two days to collect their organic extracts. The *L. polyedrum* cultures were
365 divided in two parts, one of which was filtered through a 0.22 μ m membrane to collect only
366 organics and the other was kept without filtration to keep phytoplankton cell debris. The *L.*
367 *polyedrum* supernatant with and without cells was added to the stock DG893 containing 0.3%
368 Casamino Acids, and the mixture shaken at 30°C for several days. The control was prepared in
369 the same manner without *L. polyedrum* supernatant and cells. In the media containing 1,000 nM
370 and 10,000 nM [Fe]_T, DG893 growth was enhanced by the presence of *L. polyedrum* supernatant
371 and was even more pronounced with the supernatant containing cell debris (Figure 7). In media

372 with 100 nM $[Fe]_T$, only a subtle difference was observed in DG893 growth with and without *L.*
373 *polyedrum* supernatant.

374 **Discussion**

375 Before proceeding to a discussion of the results presented here it is important to
376 operationally define what we mean by $[Fe]_T$ and “axenic” cultures. Here we used the term total
377 iron concentrations $[Fe]_T$ to mean the total analytical amount of iron added to culture which
378 includes all phases of iron present in the solution. This is different from the total dissolved iron
379 concentration which we routinely measure for environmental seawater samples (and which is
380 approximately 2-6 nM for Scripps Pier water for example) as determined by cathodic stripping
381 voltammetry (CSV). Total dissolved iron in such environmental samples is processed by
382 filtration through a 0.22 μm membrane followed by acidification. As a result, total dissolved iron
383 in a sample is much lower than the total analytical iron concentration. We use this operational
384 definition as it is not possible to measure total dissolved iron in culture media using the CSV
385 method because equilibrium between the various iron oxo/hydroxo species potentially present in
386 seawater at pH 8.2 is reached only extremely slowly and thus concentrations of total dissolved
387 iron are constantly changing over the time course utilized in these experiments. Using a straight
388 forward expression of total analytical iron concentration is thus a potentially more reproducible
389 approach with which to compare the effects of iron on bacterial and algal growth in a laboratory
390 setting.

391 Secondly we use the term “axenic” in quotation marks for describing cultures of *L.*
392 *polyedrum*, to mean the absence of any observable *cultureable* bacteria based on the absence of
393 visual bacterial colonies when cultures were placed on solid media and incubated for several
394 days. We were only partially successful making the culture “axenic” by treating the culture with

395 low concentrations of antibiotics and sequential addition of the antibiotics to the culture over
396 extended times. Using this method the culture appeared to remain bacteria free for at least a
397 month. However when the sequential addition of antibiotics to the culture was stopped, bacteria
398 growth was observed to return within a few days. Thus clearly the cultures were not truly axenic.
399 Nevertheless our main concern was that the dominant species of bacteria growing in our binary
400 culture should be DG893 and that “axenic” cultures should contain insignificant numbers of
401 unknown bacteria. We confirmed the former by qPCR and the latter was further supported by
402 presence of few if any bacteria seen via DAPI staining. To our knowledge, there is at present no
403 general protocol for preparing a completely axenic culture of *L. polyedrum* hence our use of the
404 operational definition of “axenic” described here.

405 We initially investigated conditions under which the bacterium *Marinobacter* DG893 and
406 the dinoflagellate *L. polyedrum* could grow independently. As expected, in the absence of an
407 added carbon source DG893 could not grow in natural unamended seawater or in L1
408 supplemented ASW irrespective of the iron concentration. In the presence of an adequate
409 artificial source of carbon however bacterial growth was controlled by the available
410 concentration of iron. Under idealized conditions, both DG893 and the VF null mutant reached a
411 final concentration of around 10^5 CFU/mL independent of the initial inoculum. However even
412 without any externally added carbon source DG893 grew well in the presence of *L. polyedrum*
413 indicating that the dinoflagellate leaked enough dissolved organic carbon (DOC) to fully support
414 the growth of the bacterium. The notion that DG893 could utilize *L. polyedrum* as a carbon
415 source was further supported by the observation that growth of DG893 was stimulated when it
416 was incubated in the media containing the supernatant from a filtered *L. polyedrum* culture
417 which contains non-specific organics secreted from *L. polyedrum*. These may possibly include

418 dimethylsulfoniopropionate (DMSP) a labile carbon and sulfur source known to “leaked” from
419 dinoflagellates (Caruana and Malin, 2014).

420 In the case of both “axenic” and nonaxenic cultures of the photosynthetic dinoflagellate
421 *L. polyedrum*, iron proved to be a growth limiting nutrient. Thus maximum growth occurred at
422 total iron concentrations of 100-1000 nM while higher concentrations proved toxic and lower
423 concentrations (0-10 nM) were growth limiting. In the first subculture, both “axenic” and non-
424 axenic cultures grew equally well. However, their growth showed a remarkable difference in the
425 second batch subculture. Here the non-axenic *L. polyedrum* culture continued to grow
426 exponentially, although now only at the higher $[Fe]_T$. When the iron concentrations were kept
427 high, *L. polyedrum* cultures continued to thrive up to at least 6 subcultures. However, when the
428 iron concentrations were allowed to be reduced by dilution in subsequent subculturing, the
429 growth gradually ceased. Poorly growing subcultures in low iron media could however be
430 rescued by the addition of additional iron to the culture. Remarkably “axenic” *L. polyedrum*
431 subcultures ceased to grow regardless of $[Fe]_T$ but could be rescued by the addition of DG893.
432 These results emphasize the importance of both bacteria and iron to *L. polyedrum* growth.

433 Our results further suggest that the *L. polyedrum* growth promoting effects of the
434 bacterium *Marinobacter* DG893 were in fact not unspecific but at least partially related to its
435 production of the photoactive siderophore vibrioferin. Thus only the supernatant of DG893 that
436 contained sufficient amounts of vibrioferin facilitated *L. polyedrum* growth. While the addition
437 of the supernatant from a culture of the VF null mutant of DG893 also improved growth of *L.*
438 *polyedrum* the effect was much less than that from the WT. This observation was further
439 confirmed by direct addition of purified VF into the *L. polyedrum* culture which growth
440 increased 1.5 times more than those without VF. These results imply that the presence of the

441 siderophore vibrioferin clearly influences *L. polyedrum* growth, but that other factors such as
442 vitamin B12 (Cruz-Lopez and Maske, 2015) provided by bacteria may also be important in
443 promoting this apparent bacterial-algal mutualism.

444 Here we have focused on a bacterium that produces the photoreactive siderophore,
445 vibrioferin. Vibrioferin was chosen for more detailed study for a number of reasons. First
446 vibrioferin has been isolated from several different bacteria which are known to be closely
447 associated with HAB species (Amin et al., 2007) and we found that bacteria that had the ability
448 to synthesize vibrioferin were by far the most numerous as compared to the producers of other
449 photoreactive siderophores which we followed during the bloom of *L. polyedrum* at Scripps Pier
450 in 2011. We initially supposed that vibrioferin was a good candidate for a siderophore that could
451 provide bioavailable iron for both the bacterial producers and their algal partners as it is the most
452 rapidly photolyzed of the photoactive siderophores tested and that photolyzed vibrioferin has no
453 further affinity for iron. This feature is unusual in that most other photoactive siderophores retain
454 the ability to strongly bind Fe(III) even after photolysis (Amin et al., 2009). However we have
455 recently showed that while vibrioferin was the best source of iron for *L. polyedrum* of the ones
456 we tested (Yarimizu et al., 2017), it was not due to its photolysis since iron uptake from VF was
457 the same in the dark as in the light. We therefore attribute its ability to provide bioavailable iron
458 to the dinoflagellate to its relatively weak iron binding properties (with respect to other more
459 traditional siderophores) as it lacks the sixth donor group required to complete the octahedral
460 coordination geometry preferred by Fe(III) (Amin et al., 2009). This in turn leads to a less
461 negative reduction potential (it falls within the biological range) so that the iron can readily be
462 released from it by the cell surface reductases known to be present in *L. polyedrum*.

463 As a final note, while this laboratory study strongly supports a carbon for iron mutualism
464 between the dinoflagellate *L. polyedrum* and *Marinobacter* DG893 as previously proposed
465 (Amin et al, 2009), it is unclear at present how, or indeed if, such a mutualism will operate in the
466 field. To this end we have collected data from several field studies involving both HAB and non-
467 HAB bloom events and are currently searching for relationships between phytoplankton,
468 vibioferrin producing bacteria, and available iron. We hope that both these laboratory and field
469 studies will complement each other and advance our knowledge of the potential role of bacterial-
470 algal interactions in understanding the mechanisms of HAB formation.

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599 **Figure 1. DG893 growth in media with and without Fe and carbon**

600 A) DG893 growth in media without added Fe. The bold lines represent media containing 10
601 mg/mL Casamino Acids. The dotted lines represent media without Casamino Acids. The grey
602 lines represent L1-Fe media. The black lines represent seawater. The VF null mutant of DG893
603 showed a similar pattern of growth (data not shown). B) DG893 incubated in L1 media
604 containing different $[Fe]_T$ but without an added carbon source. The black bold line, grey bold
605 line, grey wide dotted line, and grey narrow dotted line represent L1 media with 1,000 nM, 100
606 nM, 10 nM, 0 nM $[Fe]_T$, respectively. The mutant DG893 showed a similar pattern. C) DG893
607 incubated in L1 media containing 10 mg/mL Casamino Acids and different $[Fe]_T$. The black bold
608 line, grey bold line, grey wide dotted line, and grey narrow dotted line represent L1 media with
609 1,000 nM, 100 nM, 10 nM, 0 nM $[Fe]_T$, respectively.

610
611 **Figure 2. *L. polydruum* co-cultured with a starting inoculum of DG893 at 10^3 cells/mL in**

612 media with various $[Fe]_T$

613 A) 1st generation growth of *L. polydruum* in the presence of DG893. The starting inoculum of
614 DG893 was 10^3 cells/mL which reached 10^5 cells/mL after 28 days incubation under the standard
615 growth conditions. A similar pattern was observed in all other groups: i.e. media containing a
616 starting inoculum of DG893 of 10^7 , 10^6 , 10^5 , 10^4 , 10^3 , 0 cells/mL and $[Fe]_T$ of 10^4 , 10^3 , 10^2 , 10^1 ,
617 or 0 nM. B) 2nd generation growth of *L. polydruum* in the presence of DG893. After 28 days the
618 cultures in panel A were diluted to a *L. polydruum* concentration of 10^2 cells/mL with
619 appropriate media and incubated under the standard growth condition for an additional 28 days.

620
621 **Figure 3. Rescue of *L. polydruum* by addition of DG893**

622 A) 1st generation of *L. polydruum* growth maintained in L1 media either “axenic” or co-cultured
623 with DG893. B) Growth of *L. polydruum* in the 2nd subculture either “axenic” or co-cultured with
624 DG893. After the 2nd subculture of “axenic” *L. polydruum* was incubated under the standard
625 growth conditions for 21 days, an inoculum containing approximately 10^2 cells/mL of DG893
626 was added and incubation continued for an additional 14 days.

627
628 **Figure 4. Non-Axenic *L. polydruum* growth in L1 media and reduced $[Fe]_T$ over 6**
629 **subcultures**

630 Solid line) cultures maintained with a $[Fe]_T$ at the L1 level throughout the subculturing Dotted
631 line) cultures where the $[Fe]_T$ continually decreased over the subcultures due to dilution.

632
633 **Figure 5. Iron starved non-axenic *L. polydruum* growth rescued by iron addition**

634 The iron sufficient culture (solid black line) contained 11700 nM $[Fe]_T$ while the iron deficient
635 culture contained 2 nM (dotted black line). After 28 days 5850 nM $[Fe]_T$ was added to the iron
636 deficient culture which rescued *L. polydruum* growth.

637
638 **Figure 6. *L. polydruum* growth in media with DG893 supernatant, mutant supernatant,**
639 **and without bacterial supernatant**

640 The black lines, grey lines, and dotted lines represent *L. polyedrum* growth in media with added
641 DG893 supernatant, with added VF null mutant supernatant, and without any bacterial
642 supernatant, respectively.

643

644 **Figure 7. DG893 growth in media containing different $[Fe]_T$ with and without presence of**
645 ***L. polyedrum* supernatant and cell debris**

646 The solid black lines represent growth of DG893 in the presence of *L. polyedrum* supernatant
647 with their cell debris, the grey lines are DG893 growth in *L. polyedrum* supernatant without their
648 cell debris, and the dotted lines are DG893 without any *L. polyedrum* supernatant.

649