

# Description of Gloeomargarita lithophora gen. nov., sp. nov., a thylakoid-bearing, basal-branching cyanobacterium with intracellular carbonates, and proposal for Gloeomargaritales ord. nov.

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### International Journal of Systematic and Evolutionary Microbiology Description of Gloeomargarita lithophora gen. nov., sp. nov., a thylakoid-bearing basalbranching cyanobacterium with intracellular carbonates, and proposal for Gloeomargaritales ord. nov. --Manuscript Draft--

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Abstract:	A unicellular cyanobacterium, strain Alchichica-D10, was isolated from microbialites of the alkaline Lake Alchichica, Mexico. The cells were short rods (3.9 ± 0.6 µm in length and 1.1 ± 0.1 µm in width) forming biofilms of intense emerald green color. They exhibited red autofluorescence under UV light excitation. UV-visible absorption spectra revealed that they contain chlorophyll a and phycocyanin, and electron microscopy showed the presence of thylakoids. The strain grew within a temperature range of 15-30 °C. Genomic DNA G+C content was 52.2 mol%. The most remarkable feature of this species was its granular cytoplasm, due to the presence of numerous intracellular spherical granules (16-26 per cell) with an average diameter of 270 nm. These granules, easily visible under scanning electron microscopy, were composed of amorphous carbonate containing Ca, Mg, Ba, and Sr. A multi-gene phylogeny based on the analysis of 59 conserved protein markers supported robustly that this strain occupies a deep position in the cyanobacterial tree. Based on its phenotypic characters and phylogenetic position, strain Alchichica-D10 is considered to represent a new genus and novel species of cyanobacteria for which the name Gloeomargarita lithophora gen. nov., sp. nov. is proposed. The type strain is Alchichica-D10 (Culture Collection of Algae and Protozoa CCAP strain 1437/1; Collections de Cyanobactéries et Microalgues Vivantes of the Museum National d'Histoire Naturelle in Paris strain PMC 919.15). Furthermore, a new family, Gloeomargaritaceae, and a new order, Gloeoemargaritales, are proposed to accommodate this species under the International Code of Nomenclature for algae. fungi, and plants.

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- 29 The GenBank accession number for the sequences of 59 conserved genes of strain Alchichica-
- 30 D10 is CP017675.
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A unicellular cyanobacterium, strain Alchichica-D10, was isolated from microbialites of the 32 alkaline Lake Alchichica, Mexico. The cells were short rods  $(3.9 \pm 0.6 \,\mu\text{m}$  in length and  $1.1 \pm$ 33 0.1 µm in width) forming biofilms of intense emerald green color. They exhibited red 34 autofluorescence under UV light excitation. UV-visible absorption spectra revealed that they 35 contain chlorophyll a and phycocyanin, and electron microscopy showed the presence of 36 thylakoids. The strain grew within a temperature range of 15-30 °C. Genomic DNA G+C 37 content was 52.2 mol%. The most remarkable feature of this species was its granular 38 cytoplasm, due to the presence of numerous intracellular spherical granules (16-26 per cell) 39 40 with an average diameter of 270 nm. These granules, easily visible under scanning electron microscopy, were composed of amorphous carbonate containing Ca, Mg, Ba, and Sr. A multi-41 42 gene phylogeny based on the analysis of 59 conserved protein markers supported robustly that this strain occupies a deep position in the cyanobacterial tree. Based on its phenotypic 43 44 characters and phylogenetic position, strain Alchichica-D10 is considered to represent a new genus and novel species of cyanobacteria for which the name Gloeomargarita lithophora gen. 45 46 nov., sp. nov. is proposed. The type strain is Alchichica-D10 (Culture Collection of Algae and Protozoa CCAP strain 1437/1; Collections de Cyanobactéries et Microalgues Vivantes of the 47 Museum National d'Histoire Naturelle in Paris strain PMC 919.15). Furthermore, a new 48 family, Gloeomargaritaceae, and a new order, Gloeoemargaritales, are proposed to 49 accommodate this species under the International Code of Nomenclature for algae, fungi, and 50 51 plants.

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Cyanobacteria thrive in a variety of aquatic and terrestrial habitats, where their ability, unique 53 54 among bacteria, to carry out oxygenic photosynthesis makes them ecologically significant (Castenholz, 2001). They are also important from an evolutionary point of view since they 55 56 were responsible for the early oxygenation of the Earth's atmosphere (Buick, 2008) and an ancestral cyanobacterium was the endosymbiont that gave rise to the chloroplasts now found 57 58 in eukaryotic algae and plants (Gray & Doolittle, 1982). In addition, cyanobacteria have a rich 59 fossil record. The massive fossil stromatolites dating back to at least 2.7 billion years ago are considered to have been built by microbial communities dominated by cyanobacteria 60 (Altermann et al., 2006) and unequivocal calcified cyanobacterial fossils are common since 61 62 the base of the early Cambrian, with Girvanella as the first undisputed occurrence at 700 million years ago (Riding, 2006). The capacity of several cyanobacteria to precipitate calcium 63 carbonate may have enhanced their preservation and explain, at least partly, their extensive 64

fossil record. In fact, several species induce calcium carbonate precipitation by their 65 photosynthetic activity (Arp et al., 2001; Kamennaya et al., 2012), which increases locally the 66 concentration of  $CO_3^{2-}$  by the disproportionation of  $HCO_3^{-}$  to  $CO_3^{2-}$  and  $CO_2$ , the latter being 67 fixed by the enzyme ribulose-1,5-bisphosphate carboxylase-oxygenase (RuBisCO). The 68 export of alkalinity from the intracellular to the extracellular medium, by a mechanism that 69 remains poorly known, raises the saturation index for carbonate minerals in the immediate 70 cell environment and thus leads to mineral precipitation if free cations (e.g., Ca2+) and 71 nucleation sites are present. It has also been proposed that the cell surface, in particular the 72 73 exopolysaccharidic matrix, may serve as a nucleation site for carbonates (Obst et al., 2009). In all cases, the precipitation of carbonates by cyanobacteria has been regarded as an 74 75 extracellular uncontrolled process.

76 However, we recently reported the discovery of a cyanobacterial species that contained intracellular carbonate inclusions (Couradeau et al., 2012). It was the first time that this 77 capacity was found in cyanobacteria. More recently, this capability has also been found in a 78 79 small number of other cyanobacterial taxa (Benzerara et al., 2014; Li et al., 2016). Intracellular carbonates appear to be generally rare in bacteria since, up to the recent 80 discovery in cyanobacteria, their occurrence had been described only in a single species, the 81 proteobacterium Achromatium oxaliferum (Head et al., 1996). The new cyanobacterial strain 82 83 Alchichica-D10, which we provisionally named Candidatus Gloeomargarita lithophora 84 (Couradeau et al., 2012), was isolated from microbialite samples collected in the alkaline (~43 mM HCO3<sup>-</sup>, pH ~8.9) Lake Alchichica (Mexico) in 2007 and maintained alive in 85 laboratory aquaria since then. Here, we describe formally this new species and its 86 87 phylogenetic position within the Cyanobacteria.

88 Lake Alchichica microbialites are mostly composed of hydromagnesite [Mg<sub>5</sub>(CO<sub>3</sub>)<sub>4</sub>(OH)<sub>2</sub>•4(H<sub>2</sub>O)] and aragonite (CaCO<sub>3</sub>), and the microbial community inhabiting 89 them is largely dominated by very diverse cyanobacteria (Couradeau et al., 2011; Saghai et 90 91 al., 2015). After several years of growth in laboratory aquaria, the microbialites collected in 92 2007 were still inhabited by a large diversity of cyanobacteria similar to that found in the Lake, which suggests that this community is highly resilient (Couradeau et al., 2011). 93 Microbialites and the aquaria walls were covered by extensive biofilms. These biofilms 94 contained different cyanobacterial morphotypes, with a particularly abundant one consisting 95 of small rod-shaped cells with granular cytoplasm, noticeable under optical microscopy (Fig. 96 1). To enrich this cyanobacterial species, we disrupted a biofilm sample and filtered the 97

detached cells through an isopore filter of 3 µm pore size. We then inoculated a 96-well plate 98 containing BG11 medium with the filtered cells. After one month of incubation at 21 °C 99 applying a diel cycle, we observed growth of the targeted morphotype in 6 wells. Sequencing 100 of the 16S rRNA gene from these 6 cultures yielded identical sequences (Couradeau et al., 101 2012; accession number JQ733894). We further purified this cyanobacterium by growth on 102 BG11-agar plates and single colony isolation. This allowed us to obtain cultures with this 103 single cyanobacterial species. However, sequencing of 16S rRNA genes amplified with 104 universal bacterial primers revealed the presence of a contaminant alphaproteobacterium 105 106 closely related to several species of the genus Sandarakinorhabdus (with 97% 16S rRNA gene sequence similarity). We have been unable to eliminate this contaminant from our 107 108 cultures, partly due to the very slow growth rate of the cyanobacterium. Interestingly, some of these Sandarakinorhabdus species have also been reported in association with other 109 110 cyanobacteria, such as the strain Sandarakinorhabdus sp. A14 that is found in cultures of Microcystis aeruginosa (Shi et al., 2009). Nevertheless, observations using epifluorescent 111 optical microscopy and scanning electron microscopy (SEM) showed that the cultures were 112 largely dominated by the cyanobacterium and that the contaminant appeared to be rare. Thus, 113 even if non-axenic, the cultures were suitable for the description of the new cyanobacterial 114 species using a variety of techniques. 115

Cyanobacterial cells belonging to the new Alchichica-D10 strain grown in BG11 medium 116 measured  $3.9 \pm 0.6 \,\mu\text{m}$  in length and  $1.1 \pm 0.1 \,\mu\text{m}$  in width. The most conspicuous feature of 117 those cells observed under SEM (using secondary electron mode) was the presence of 118 numerous bright intracellular spherical granules (3-19 per cell) measuring between 60 and 119 120 380 nm in diameter (Fig. 2a). As previously determined by Benzerara et al. (2014) using energy-dispersive x-ray spectrometry (EDXS), these inclusions were composed of Ca 121 122 carbonate. In addition, cells grown in BG11 contained a relatively small number of larger and darker inclusions rich in P that corresponded to polyphosphate granules (Fig. 2b). When 123 124 grown in the aquarium water, highly alkaline and rich in Ca, Mg, and other cations but poor in 125 P, the cells contained more carbonate granules (16 to 26 per cell) with an average diameter of 270 nm. In contrast with cells grown in BG11, the chemical composition of these inclusions 126 contained Mg, Ba, and Sr as major elements in addition to Ca, and polyphosphate granules 127 128 were rare (Fig. S1, available in the online Supplementary Material). Interestingly, Ba/Ca and Sr/Ca atomic ratios in the inclusions were 1370 and 86 times higher, respectively, than those 129 measured in the aquaria solution (Couradeau et al., 2012), although Ca, Sr and Ba are usually 130

supposed to be incorporated relatively conservatively by carbonates. This suggested that thecells controlled the chemical composition of the inclusions.

The ultrastructure of the cells was studied by transmission electron microscopy (TEM). In 133 134 addition to the typical Gram negative double cell membrane, an important structural feature was the presence of thylakoids, clearly visible as several concentric layers parallel to the cell 135 136 periphery (Fig. 3). The occurrence of thylakoids clearly differentiates Gloeomargarita lithophora from the deep-branching genus Gloeobacter, which lacks these endomembrane 137 138 structures (Rippka et al., 1974). Intracellular structures were observed in the same cells with a contrast different from that of carbonate or polyphosphate inclusions (see arrows in Fig. 3) 139 140 but with morphological and contrast similarities with carboxysomes found in other cyanobacteria (e.g., Porta et al., 2000). 141

Cells observed by confocal laser scanning microscopy (CLSM) under UV light (405 nm) excitation showed intense red autofluorescence (Fig. S2, available in the online Supplementary Material). The absorption spectrum of pigments extracted with 90% acetone showed peaks at wavelengths of 620 and 664 nm, indicating the presence of phycocyanin and chlorophyll *a* (Fig. S3, available in the online Supplementary Material), which are typical pigments of cyanobacteria.

Strain Alchichica-D10 colonies grew very slowly on agar plates. They exhibited an intense 148 149 emerald color and were surrounded by a thick mucilaginous cover (Fig. S4a, available in the online Supplementary Material). Individual cells appear to be able to glide on the plate 150 151 surface to initiate the growth of new peripheral colonies (Fig. S4b, available in the online Supplementary Material). This phenomenon can lead to the formation of migration fronts that 152 153 provide a stratified structure to the margins of mature colonies (Fig. S4a, available in the online Supplementary Material). To determine the optimal growth conditions in the 154 laboratory, we combined 3 different buffered pH values (8.0, 8.5, and 9.0), 5 temperatures 155 (15, 20, 25, 30, and 37 °C), and 3 light intensities (photon flux of 5, 10, and 41  $\mu$ moles m<sup>-2</sup>s<sup>-1</sup>) 156 in both liquid and solid BG11 media buffered with HEPES. Growth was extremely slow at 157 15° C and did not occur at 37 °C. Thus, we focused on the intermediate temperatures. In all 158 cases, growth was slow and took at least 6 weeks to become noticeable by the development of 159 160 visible colonies on solid medium or by the appearance of green color in the liquid cultures. In these liquid cultures, the highest cell densities were observed at pH 8.0 and 8.5 at 25 and 30 161 162 °C, whereas in solid medium the optimal condition appeared to be at a pH of 8.5 with low

light intensity (photon flux of 5-10  $\mu$ moles m<sup>-2</sup>s<sup>-1</sup>) and a temperature of 25-30 °C (Fig. S5, available in the online Supplementary Material).

Preliminary phylogenetic analyses based on 16S rRNA gene sequences suggested the 165 166 proximity of strain Alchichica-D10 to the basal order Gloeobacterales (Couradeau et al., 2012). However, this relationship was not strongly supported and was based on unrooted 167 168 phylogenetic trees. In fact, a 16S rRNA rooted tree published later showed that G. lithophora did not branch as sister of the Gloeobacter species but as the second branch to diverge within 169 170 the Cyanobacteria after *Gloeobacter*, though still with weak statistical support (Saw et al., 2013). To resolve this uncertainty, we carried out a multi-gene phylogenetic analysis. For this 171 172 purpose, we extracted G. lithophora genomic DNA that was sequenced using the Illumina Genome Analyzer II technology, which yielded 2.1 Gbp of DNA sequences (with a G + C173 174 content of 52.2). Among these sequences, we fetched 59 conserved genes involved in transcription and translation (Table S1, available in the online Supplementary Material). Their 175 translated protein sequences were aligned with the respective homologous sequences found in 176 all completely sequenced cyanobacterial genomes and several other bacteria included as 177 outgroups. Alignments were trimmed to eliminate ambiguously aligned regions and 178 concatenated to build a 7,220 amino acids-long concatenation, which was analyzed using 179 Bayesian inference to reconstruct a phylogenetic tree. The resulting tree was highly supported 180 and placed G. lithophora in an early-diverging position, as the third most basal cyanobacterial 181 182 branch, just after the two available *Gloeobacter* species and a group containing the strain Synechococcus sp. PCC 7336 and the two thermophilic strains Synechococcus sp. JA-2-183 3B'a(2-13) and JA-3-3Ab isolated from Yellowstone (Fig. 4 and Fig. S6, available in the 184 online Supplementary Material). Therefore, Gloeobacter and Gloeomargarita are not sister 185 groups, contradicting our previous single gene-based suggestion that strain Alchichica-D10 186 187 might be a divergent species belonging to the order Gloeobacterales. Indeed, the presence of thylakoids in strain Alchichica-D10 (see above) constituted a major difference with the 188 189 thylakoid-lacking Gloeobacter genus (Rippka et al., 1974), in agreement with their placement in independent branches in the multi-gene phylogenetic tree. 190

Although *G. lithophora* is the only species available in culture for this new genus, a large diversity of related environmental 16S rRNA gene sequences has been detected, indicating that it belongs to a diverse clade found in various environments, in particular freshwater microbialites and microbial mats (Ragon *et al.*, 2014). Interestingly, several sequences have been retrieved from microbial mats thriving in continental hot springs from various locations, such as Yellowstone, central Tibet, and Algeria (Amarouche-Yala *et al.*, 2014; Lau *et al.*,
2009; Turner *et al.*, 1999). Moreover, cells with morphological characteristics similar to those
of *G. lithophora*, including the presence of numerous carbonate inclusions in the cytoplasm,
were observed by electron microscopy in the Algerian hot spring samples (Ragon *et al.*,
2014). These results indicate that the different lineages related to *Gloeomargarita* have
adapted to a wide range of temperatures.

As *G. lithophora* does not belong to the Gloeobacterales, the erection of a new genus, family and order to accommodate this new cyanobacterial species is required (see below). As far as we know, the genus name *Gloeomargarita* has never been used in botanical literature, so it can be validly published as a new cyanobacterial genus under the International Code of Nomenclature for algae, fungi and plants (McNeill *et al.*, 2012).

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#### 208 Description of *Gloeomargarita* gen. nov.

209 *Gloeomargarita* (Gloe.o.mar.ga.ri'ta. Gr. n. *gloios* glutinous substance; L. fem. n. *margarita*210 pearl; N.L. fem. n. *Gloeomargarita* glutinous cells containing pearls).

Unicellular rods with oxygenic photoautotrophic metabolism and gliding motility. Contain
chlorophyll *a* and phycocyanin and photosynthetic thylakoids located peripherally.
Reproduction by transverse binary fission in a single plane. Do not produce well-defined
sheath layers. Contain spherical inclusions of earth alkaline carbonates in the cytoplasm. The
type species is *Gloeomargarita lithophora* sp. nov.

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#### 217 Description of *Gloeomargarita lithophora* sp. nov.

218 *Gloeomargarita lithophora* (li.tho'pho.ra. Gr. masc. n. *lithos* stone; Gr. masc. n. *phoros*219 carrier; N.L. fem. n. *lithophora* carrier of stones).

Exhibits the following properties in addition to those given in the genus description. Cells are 1.1  $\mu$ m wide and 3.9  $\mu$ m long in average. Growth occurs at 15-30 °C (optimum 25 °C) in alkaline freshwater and BG11 medium. The G + C content of the genomic DNA of the type strain is 52.2 mol%. The type strain, Alchichica-D10 (=CCAP 1437/1, =PMC 919.15), was isolated from microbialites of the alkaline Lake Alchichica (Mexico) preserved in laboratory aquaria at Orsay (France). The 16S rRNA gene sequence of the type strain is available in GenBank under accession number JQ733894. 227 The holoptype of *G. lithophora* is the specimen PueAl-43a in the FCME Herbarium of the

Faculty of Sciences at the UNAM. Type locality: Alchichica Lake (Mexico). Living cultures
CCAP 1437/1 and PMC 919.15 are ex-holotypes.

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#### 231 Description of *Gloeomargaritaceae* fam. nov.

232 Gloeomargaritaceae (Gloe.o.mar.ga.ri.ta.ce'ae. N.L. fem. n. Gloeomargarita type genus of

the family; suff. -aceae ending to denote a family; N.L. fem. pl. n. Gloeomargaritaceae the

- family of the genus *Gloeomargarita*).
- 235 The description is the same as for the genus *Gloeomargarita*.
- 236 Type genus is *Gloeomargarita* gen. nov.
- 237

#### 238 Description of *Gloeomargaritales* ord. nov.

239 Gloeomargaritales (Gloe.o'mar.ga.ri.ta'les. N.L. fem. n. Gloeomargarita type genus of the

240 order; suff. -ales ending denoting an order; N.L. fem. pl. n. *Gloeomargaritales* the order of

- the genus *Gloeomargarita*).
- 242 The description is the same as for the genus *Gloeomargarita*.
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#### 332 Figure legends

Fig. 1. Differential interference contrast (DIC) image of several Alchichica-D10 cells
collected from an aquarium biofilm. Bar, 10 μm.

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Fig. 2. Electron microscopy analyses of Alchichica-D10 cells grown in BG11. (a) Scanningtransmission electron microscopy image in high angle annular dark field (STEM-HAADF)
mode: Ca-carbonates appear as brighter round-shaped inclusions, while polyphosphate
granules are darker, sometimes bigger globules. (b) Scanning-transmission electron
microscopy energy dispersive X-rays spectrometry (STEM-EDX) map of the same area:
Calcium is in red, phosphorus in green and carbon in blue; as a result, Ca-carbonates appear
in red and polyphosphate granules in green. Bars, 1 μm.

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Fig. 3. Bright-field TEM image of a thin section of Alchichica-D10 cells embedded in EPON
resin. Several concentric thylakoid membranes are visible under the cell membrane.
Arrowheads indicate structures that may correspond to carboxysomes. Bar, 1 µm.

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**Fig. 4.** Bayesian phylogenetic tree based on the analysis of a concatenation of 59 conserved proteins (7220 amino acids) reconstructed using PhyloBayes MPI (Lartillot *et al.*, 2013) with the CAT GTR model. Numbers at branches are posterior probabilities (only those >0.50 are shown). For space constraints, the outgroup and the *Synechococcus/Prochlorococcus* group have been replaced by triangles (for the complete tree see Fig. S6, available in the online Supplementary Material). The scale bar indicates the number of substitutions per position.

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#### Supplementary Table S1, Proteins used for phylogenetic reconstruction

Protein	Protein name
EFG	elongation factor G
EFTs	elongation factor Ts
EFTu	elongation factor Tu
IF-1	initiation factor 1
IF-2B	initiation factor 2B
IF-3	initiation factor 3
rpL1	ribosomal protein L1
rpL10	ribosomal protein L10
rpL11	ribosomal protein L11
rpL13	ribosomal protein L13
rpL14	ribosomal protein L14
rpL15	ribosomal protein L15
rpL16	ribosomal protein L16
rpL17	ribosomal protein L17
rpL18	ribosomal protein L18
rol 19	ribosomal protein L 19
rp = 10	ribosomal protein L2
rpl 20	ribosomal protein L20
rol 21	ribosomal protein L20
rol 22	ribosomal protein L21
rol 23	ribosomal protein L22
rol 24	ribosomal protein L20
rol 27	ribosomal protein L24
$rp \perp 27$	ribosomal protein L27
rpL20	ribosomal protein L20
rpL23	ribosomal protein L2
rpL3	ribosomal protein L31P
rpL31F	ribosomal protein L32
rpL32	ribosomal protein L32
rpL33	ribosomal protein L33
rpL34	ribosomal protein L34
rpL35	ribosomal protein L35
rp∟4	ribosomal protein L4
rpL5	ribosomal protein L5
rpL6	ribosomal protein L6
rpL7-L12	ribosomal protein L7-L12
rpL9	ribosomal protein L9
rpPSRP3	ribosomal protein PSRP3
rpS10	ribosomal protein S10
rpS11	ribosomal protein S11
rpS12	ribosomal protein S12
rpS13	ribosomal protein S13
rpS14	ribosomal protein S14
rpS15	ribosomal protein S15
rpS16	ribosomal protein S16
rpS17P	ribosomal protein S17P
rpS18	ribosomal protein S18
rpS19	ribosomal protein S19
rpS1P	ribosomal protein S1P
rpS2	ribosomal protein S2

rpS20	ribosomal protein S20
rpS21	ribosomal protein S21
rpS3	ribosomal protein S3
rpS30EA	ribosomal protein S30EA
rpS4	ribosomal protein S4
rpS5	ribosomal protein S5
rpS6	ribosomal protein S6
rpS7	ribosomal protein S7
rpS8	ribosomal protein S8
rpS9	ribosomal protein S9



**Supplementary Fig. S1.** Electron microscopy analyses of Alchichica-D10 cells grown in aquarium water. (a) Scanning-transmission electron microscopy high angle annular dark field (STEM-HAADF) image: Ca-carbonates appear as brighter round-shaped inclusions. (b) Energy dispersive X-ray spectrometry (EDXS) map of the area outlined in (a): Calcium is in red, phosphorus in green and carbon in blue; as a result, Ca-carbonates appear in red and PolyP granules in green. (c) EDXS spectrum of a carbonate inclusion showing the presence of Ca, Sr, Mg and Ba.



**Supplementary Fig. S2.** Autofluorescence of Alchichica-D10 cells observed under confocal laser scanning microscopy with UV light illumination (405 nm). Bar,  $10 \mu m$ .



**Supplementary Fig. S3.** Absorption spectra of strain Alchichica-D10 pigments extracted with 90% acetone.



**Supplementary Fig. S4.** (a) Alchichica-D10 strain colony growing on a BG11-agar plate. (b) Magnification of the colony border. Bars, 1 mm (a) and 100  $\mu$ m (b).

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**Supplementary Fig. S5.** Growth of strain Alchichica-D10 in liquid and solid BG11 medium under different temperature, pH, and light intensity conditions.



**Supplementary Fig. S6.** Bayesian phylogenetic tree based on the analysis of a concatenation of 59 proteins (7220 amino acids) reconstructed using PhyloBayes MPI with the CAT GTR model. Numbers at branches are posterior probabilities (only those >0.50 are shown). The scale bar indicates the number of substitutions per position.

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