## 1 SUPPLEMENTAL MATERIAL





FIG S1 Morphological identification of strain XQGS-1 cells with a coupled FISH-SEM
approach. (A) Fluorescence micrograph of bacteria hybridized *in situ* with the 5'-FAMlabeled universal bacterial probe EUB338. (B) Fluorescence micrograph of bacteria
hybridized *in situ* with the 5'-Cy3-labeled XQGS-1-specific probe XQGS1-945. (C)

- Overlay of images (A) and (B). (D) Coordinated SEM image of the same field of view
  as in image (C). (E), (F), (H), (I), (J), (K), and (L) High-magnification SEM image of
  the boxed areas in (D). (G) Partial enlargement of the boxed area in (F).
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12 FIG S2 Morphological identification of strain XQGS-1 cells with a coupled FISH-SEM

approach. (A) Fluorescence micrograph of bacteria hybridized *in situ* with the 5'-FAMlabeled universal bacterial probe EUB338. (B) Fluorescence micrograph of bacteria
hybridized *in situ* with the 5'-Cy3-labeled XQGS-1-specific probe XQGS1-945. (C)
Overlay of images (A) and (B). (D) Coordinated SEM image of the same field of view
as in image (C). (E) - (M) High-magnification SEM image of the boxed areas in (C).



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FIG S3 EDXS elemental mappings of strains AMB-1 cells with polyhydroxybutyrates
(PHB) and magnetosome particles and XQGS-1 cells with magnetosome particles used
for coupled FISH-SEM experiments. (A) and (D) SEM images according with FIG S11
and FIG 1G. (B) and (E) EDXS elemental maps of C (C Kα). (C) and (F) EDXS
elemental maps of Fe (Fe Kα).

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FIG. S4 EDXS elemental mapping of five XQGS-1 cells in the STEM-HAADF mode.
(A) STEM-HAADF image with positions of five type-III (calcium carbonate granule)
inclusions (yellow hollow circles). (B) - (F) EDXS elemental maps of (B) C (C Kα),
(C) O (O Kα), (D) Fe (Fe Kα), (E) P (P Kα), and (F) Ca (Ca Kα). (G) RGB map with P
(cyan), Ca (green), and Fe (red). (H) EDXS spectra for selected (~50 nm) regions of
five type-III (calcium carbonate granule) inclusions in (A).





FIG. S5 EDXS elemental mapping of five XQGS-1 cells in the STEM-HAADF mode. 35 (A) STEM-HAADF image with positions of ten type-II (calcium phosphate granule) 36 inclusions (red hollow circles) and five type-III (calcium carbonate granule) inclusions 37 (yellow hollow circles). (B) - (F) EDXS elemental maps of (B) C (C Ka), (C) O (O Ka), 38 (D) Fe (Fe Ka), (E) P (P Ka), and (F) Ca (Ca Ka). (G) RGB map with P (cyan), Ca 39 (green), and Fe (red). (H) EDXS spectra for selected (~50 nm) regions of ten type-II 40 (calcium phosphate granule) inclusions and five type-III (calcium carbonate granule) 41 inclusions in (A). 42

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## 44 SUPPLEMENTAL TABLE

45 Table S1. 16S rRNA sequences retrieved from Xingqinggong Lake

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	Strain	No. of clones	Percentage of clones	Most similar strain	Accession	Identity	Reference
	XQGS-1	22	73.3%	Caenispirillum salinarum	NR_117049	92.6%	(1)
	Magnetospirillum sp.	6	20%	<i>Magnetospirillum</i> sp. clone WYH-24	JX537774	99.8%	(2)
	Methylobacillus sp.	2	6.7%	<i>Methylobacillus</i> sp. clone CH1-43	JX079398	97.1%	Unpublished

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## 48 **REFERENCES**

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