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1 [Manuscript for Applied and Environmental Microbiology]

2

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4 **Biomagnetic recovery of selenium: Bioaccumulating of selenium**
5 **granules in magnetotactic bacteria**

6

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19 **Running title:** Selenium granule formation in magnetotactic bacteria

20

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30 **Key words:** Magnetotactic bacteria, Magnetic recovery, Selenium, Magnetite

31 **Abbreviation used are:** MSGM, magnetic spirillum growth medium; MIC, minimum inhibitory
32 concentration; *M. magneticum* AMB-1, *Magnetospirillum magneticum* AMB-1; TEM,
33 transmission electron microscopy; EDX, energy dispersive X-ray spectrometry.

34

35 **ABSTRACT**

36 Using microorganisms to remove waste and/or neutralize pollutants from contaminated water is
37 attracting much attention due to the environmentally friendly nature of this methodology.
38 However, cell recovery remains a bottleneck and a considerable challenge for the development
39 of this process. Magnetotactic bacteria are a unique group of organisms that can be manipulated
40 by an external magnetic field due to the presence of biogenic magnetite crystals formed within
41 their cells. In this study, we demonstrated the first account of accumulation and precipitation of
42 amorphous elemental selenium nanoparticles within magnetotactic bacteria alongside and
43 independently to magnetite crystal biomineralisation when grown in a medium containing
44 selenium oxyanion (SeO_3^{2-}). Quantitative analysis shows that magnetotactic bacteria accumulate
45 the highest amount of target molecules (Se) per cell than any other previously reported of
46 non-ferrous metal/metalloid. For example, 2.4 and 174 times more Se is accumulated when
47 compared to Te uptaken into cells and Cd^{2+} adsorption onto the cell surface respectively.
48 Crucially, the bacteria with high levels of Se accumulation were successfully recovered with an
49 external magnetic field. This biomagnetic recovery and effective accumulation of target elements
50 demonstrate the potential for application in bioremediation of polluted water.

51

52 **IMPORTANCE**

53 The development of a technique for effective environmental water remediation is urgently
54 required across the globe. A biological remediation process of waste removal and/or
55 neutralization of pollutant from contaminated water using microorganism has great potential, but

56 cell recovery remains a bottleneck. Magnetotactic bacteria synthesize magnetic particles within
57 their cells, which can be recovered by a magnetic field. Herein, we report the first example of
58 accumulation and precipitation of amorphous elemental selenium nanoparticles within
59 magnetotactic bacteria independent of magnetic particle synthesis. The cells were able to
60 accumulate the highest amount of Se compared to other foreign elements. More importantly, the
61 Se accumulating bacteria were successfully recovered with an external magnetic field. We
62 believe magnetotactic bacteria confer unique advantages of biomagnetic cell recovery and of Se
63 accumulation, providing a new and effective methodology for bioremediation of polluted water.

64

65 INTRODUCTION

66 Environmental remediation, a technique of waste removal and/or neutralization of pollutant
67 from a contaminated site, is an attractive field because of the increasing difficulty and
68 importance of pure water acquisition in both developing and industrial countries. Among the
69 various technologies for environmental water remediation, biorecovery of waste using
70 microorganisms has great potential and is an environmentally friendly alternative to conventional
71 techniques such as reclamation treatment (1–3). Studies of the waste biosorption onto
72 microorganisms and uptake into cells have been well demonstrated, but cell recovery remains a
73 bottleneck in this approach because scale-up of collection methods such as centrifugation and
74 filtration provides a huge logistical and monetary challenge.

75 Magnetotactic bacteria are unique prokaryotes, recognized by their response to a magnetic field.
76 This is due to the presence of magnetic nanoparticles of Fe_3O_4 or Fe_3S_2 within the cells (4–6).

77 The particle formation occurs within an organelle, called a magnetosome, which is formed along
78 the intracellular filamentous structure (7–9). The magnetosomes confer a magnetic moment to
79 the cells, allowing them to migrate in aquatic environment under the influence of the Earth's
80 geomagnetic field. We have already investigated the use of magnetotactic bacteria for the
81 biomagnetic recovery of toxic and/or valuable metals and metalloid such as Cd (10, 11), Au (12),
82 and Te (13). In these studies, Cd^{2+} and AuCl_4^- were mainly adsorbed onto the cell surface (10,
83 12), while the Te oxyanion (TeO_3^{2-}) was reduced and biomineralized as discrete independent
84 elemental Te nano-crystals within the cells with no incorporation into the magnetite crystals (13).
85 The dual crystallization of tellurium and magnetite by magnetotactic bacteria enabled
86 approximately 70 times more bioaccumulation of the pollutant per cell than cell surface
87 adsorption. Therefore intracellular accumulation of target elements within magnetotactic bacteria
88 offers the most promising system for bioremediation due to the unique advantages of both
89 magnetic manipulation with external magnetic field and of effective target molecule
90 accumulation.

91 Selenium (Se) is a rare element of high use in industry to produce various valuable materials
92 because of its unusual semiconducting and photo-optical physical properties (14). The increased
93 use of Se has led to its rising price and its increase in water contamination, which is in danger of
94 presenting both an ecological and human health risk (15, 16). Therefore, the growing demand for
95 Se in industrial technologies and the increased pollution effects of its byproducts into aquatic
96 environments is rendering the recovery and recycling of this valuable element a very attractive
97 global proposition. In aqueous environments, selenium is generally found as the toxic oxyanions

98 selenate (SeO_4^{2-} , +VI) and selenite (SeO_3^{2-} , +IV). The selenium oxide ions can adsorb
99 extracellularly to the cell surfaces of microorganisms (1, 17). In addition, some microorganisms
100 in the environment possess various strategies of detoxification such as methylation, assimilation
101 as selenoamino acid, and reduction that could provide the potential to effectively accumulate Se
102 within the cell (18, 19).

103 In this study we investigate the minimum inhibitory concentration (MIC) of selenium oxyanion
104 (SeO_3^{2-}) for the magnetotactic bacterium *M. magneticum* AMB-1; the effect of this anion on
105 magnetite crystal synthesis; and if uptaken, whether the Se dopes into the magnetite crystals
106 (similar to the Co and Mn previously reported) (20, 21) or forms discrete crystals/inclusions
107 within the cells (similar to the Te study) (13). Finally, the magnetic recovery of Se using
108 magnetotactic bacteria is investigated.

109

110 MATERIALS AND METHODS

111 **Determination of the minimum inhibitory concentration (MIC) of Selenite ion for *M.***
112 ***magneticum* AMB-1 growth.** *M. magneticum* AMB-1 (ATCC700264) (22) was
113 microaerobically cultured in magnetic spirillum growth medium (MSGM) at 28°C as previously
114 described (23). Microaerobic conditions were established by purging the cultures with argon gas.
115 The MIC of selenium for *M. magneticum* AMB-1 in MSGM was determined by growing the
116 cells in various initial concentrations of selenite salt (Na_2SeO_3): 0 (control), 5, 10, 20, 40, 60, 80,
117 100 and 250 μM . The cells were directly counted with a hemacytometer under an optical
118 microscope (Leica DML) after 7 days culture. Additionally the optical density (OD_{600}) was

119 recorded.

120 **Transmission electron microscopy (TEM) and energy dispersive X-ray (EDX) spectrometry**

121 **analyses of *M. magneticum* AMB-1 grown in the presence of SeO_3^{2-} .** Cultured bacterial cells

122 harvested from medium were washed with MilliQ three times and spotted onto 300-mesh

123 Formvar/Carbon coated copper grids (Agar Scientific Ltd). The samples were analyzed by TEM

124 operated at an accelerating voltage of 100 kV (Philips, CM10). High resolution TEM imaging

125 and analysis were conducted on a FEI CM200 field emission gun TEM running at 200 kV

126 equipped with an Oxford Instruments EDX spectrometer and a Gatan Imaging Filter. EDX

127 analysis was conducted for at least 6 crystals in different cells under the same experimental

128 conditions with representative spot data shown.

129 **Se accumulation in *M. magneticum* AMB-1.** To evaluate the amount of uptake and adsorbed

130 SeO_3^{2-} in/onto cells, an atomic absorption spectrophotometer (Shimadzu, AA-6600G) was used.

131 After the cells were collected by centrifugation (or in the case of the magnetic recovery assay,

132 collection by magnetic trap in a glass test tube), the precipitates were washed 3 times with

133 HEPES buffer (pH 7.4), dried and then dissolved with nitric acid solution (0.1N) with heating on

134 in oil bath. After discarding the supernatant, the cells were dissolved with same procedure as

135 described above. The dissolved solutions were quantitatively analyzed by atomic absorption

136 spectrophotometry, using a calibration curve derived from standard solutions. All assays were

137 performed three times.

138 **Magnetic recovery assay of magnetotactic bacteria grown in the presence of selenite ions.**

139 To verify the ability of biomagnetic recovery of *M. magneticum* AMB-1 in the presence of

140 SeO_3^{2-} using magnetic force, a magnetic cell recovery assay was conducted. The *M. magneticum*
141 AMB-1 wild type strain was harvested at the late logarithmic phase of growth, cells were
142 counted and adjusted to 1.0×10^8 cells/ml of MSGM in the presence of the SeO_3^{2-} at different
143 concentrations (0, 25, 50 and 100 μM). Three milliliters of each sample was then transferred to
144 separate glass test tubes (Diameter: 7 mm, Height: 7.5 cm), each of which was sealed with a
145 rubber cork. Cylindrical neodymium-boron magnets (Diameter: 15 mm, Height: 1 cm) were
146 placed on the exterior of the horizontal centre of each test tube to allow cell recovery to take
147 place. At the designated times (1, 2, 4, 6, 8, 10, 15 and 20h), culture medium was collected by
148 inserting a syringe through the rubber cork and extracting culture medium (20 μl) from around
149 the water surface. A cell count was performed against the extracted culture medium samples.
150 After the magnetic separation, the amount of uptake and adsorbed SeO_3^{2-} in/onto magnetically
151 manipulated cells was evaluated using an atomic absorption spectrophotometer (Shimadzu,
152 AA-6600G). In addition, the magnetically collected cells and Se concentration were measured at
153 the endpoint for further verification.

154

155 RESULTS AND DISCUSSION

156 Effect of SeO_3^{2-} on cell growth and on magnetite biomineralisation in *M. magneticum*

157 **AMB-1.** The effect of selenium oxyanion (SeO_3^{2-}) on the growth of *M. magneticum* AMB-1 was
158 investigated at various concentrations (Fig. 1). Cells cultured in MSGM containing 0 and 5 μM
159 SeO_3^{2-} showed similar growth rates, with stationary-phase cell densities of approximately $2.2 \times$
160 10^8 cells/ml. Cell growth was negatively affected by the increase of SeO_3^{2-} concentration and no

161 cell growth was found at $\geq 250 \mu\text{M}$. The MIC of selenium oxyanion for *M. magneticum* AMB-1
162 was determined to be $250 \mu\text{M}$ under these experimental conditions. The result indicated that
163 SeO_3^{2-} is mildly toxic to this bacteria compared with the other chalcogen, tellurium oxyanion
164 (e.g. MIC = $60 \mu\text{M}$) (13). As *E. coli* has a MIC of 400 mM (SeO_3^{2-}), *M. magneticum* AMB-1 is
165 less resistant to this element. Similar observations have been previously found with respect to
166 Co^{2+} , Ni^{2+} , and Cu^{2+} showing approximately 90% less resistance than *E. coli* (20). It is of note
167 that light-orange colors developed during the cell growth in the presence of SeO_3^{2-} . Similar
168 observations were reported in various selenite-reducing bacteria (25, 26). The effect of the
169 chalcogen on magnetite crystal formation in magnetotactic bacteria was also investigated (Fig. 1).
170 The result showed a gradual decrease of magnetosomes with the increase of SeO_3^{2-} concentration
171 but magnetite formation was observed even in the presence of high concentrations ($100 \mu\text{M}$) of
172 SeO_3^{2-} . In addition, optical microscopy showed that approximately 100% and 70% of cells
173 grown in the presence of $25 \mu\text{M}$ and $100 \mu\text{M}$ of SeO_3^{2-} respectively responded to the external
174 magnetic field.

175 **Observation of discrete formation of magnetite crystals and Se granule in *M. magneticum***
176 **AMB-1 grown in the presence of SeO_3^{2-} .** Figure 2a shows representative transmission electron
177 microscope (TEM) images of *M. magneticum* AMB-1 grown in the presence ($100 \mu\text{M}$) and
178 absence of SeO_3^{2-} in the MSGM medium. Approximately 10 independent spherical granules
179 ($30\text{--}300 \text{ nm}$ diameter) were observed in the cell grown in the presence of SeO_3^{2-} (Fig. 2a), while
180 all cells revealed the presence of the magnetite crystals in a chain structure. The number and size
181 of Se inclusions within the cell increased with increasing initial concentration of SeO_3^{2-} in the

182 medium. In a previous study, we have observed the doping of some metals (Cu, Mn, and Co)
183 into bacterial magnetite crystal under laboratory-controlled conditions (20). However, in this
184 study the elemental mapping showed no signal from Se in magnetite crystals (Fig. 2b). To verify
185 the elemental components in these Se particles, STEM-EDX spot spectra were recorded and
186 showed Se was the only element present (the Cu was from the TEM grid) (Fig. 2b and c). No
187 oxygen was detected, inferring the inclusions are composed of pure elemental Se (0), which
188 seems to be reduced and precipitated from SeO_3^{2-} in the cell. Selenium is a group 16 non-metal
189 (chalcogens), neighbored by sulfur and the metalloid tellurium. Thiosulfate ($\text{S}_2\text{O}_3^{2-}$), tellurite
190 (TeO_3^{2-}), and selenite (SeO_3^{2-}) are proposed to be taken up by bacteria and reduced to elemental
191 S, Te, and Se, respectively (25, 27, 28). This is supported by the fact that S-globules are present
192 in many microbes, including magnetotactic bacteria (29, 30), and we have also reported the
193 formation of Te nano-crystals in magnetotactic bacteria independent from the magnetosome (13).
194 Here we show for the first time that magnetotactic bacteria uptake, reduce and intracellularly
195 form discrete Se granules independent to magnetosomes, similar to Te crystal precipitation in the
196 same organism (13). The granules were examined by high-resolution TEM with selected area
197 electron diffraction which showed a diffuse pattern, revealing the amorphous Se structure.

198 **Time course measurements of Se accumulation in *M. magneticum* AMB-1.** The time course
199 of Se accumulation in magnetotactic bacteria was measured (Fig. 3). The cell growth and Se
200 accumulation were saturated within 7 days and the Se uptake in cells mainly occurred in the
201 stationary phase (for cells grown in 100 μM of SeO_3^{2-}). Under this condition, 68.1% of the initial
202 Se (100 μM) was accumulated by the cells, which accounts to 6.6×10^8 Se atoms per cell. In the

203 case of Te accumulation found in the previous study, the most effective condition revealed that
204 2.7×10^8 Te atoms were accumulated per cell, which indicates that 2.4 times more Se is
205 accumulated than Te. Furthermore, surface hexa-histidine expressing modified AMB-1 cells
206 have previously been shown to adsorb Cd^{2+} onto these sites on the cell surface, showing the
207 adsorption of 3.8×10^6 metal ions. Therefore, 2.4 and 174 times more Se was accumulated when
208 compared to Te in cell and Cd^{2+} adsorption onto cell surface. These results highlight the greater
209 loading of elemental Se into AMB-1 cells than any other metalloid or non-ferrous metal.

210 **Biomagnetic recovery of SeO_3^{2-} using *M. magneticum* AMB-1.** Magnetotactic bacteria
211 harboring our target element (Se) for recovery can be manipulated and isolated by an external
212 magnetic field, significantly magnifying the bioremediation potential of these cells for targeted
213 recovery from polluted water environments. Herein, biomagnetic recovery of magnetotactic
214 bacteria grown in the presence of SeO_3^{2-} was conducted. The result shown in Fig. 4 revealed that
215 almost all cells grown in $25 \mu\text{M}$ SeO_3^{2-} were successfully recovered within 8 hours. The time for
216 magnetic recovery of cells gradually increased with increasing concentration of SeO_3^{2-} . This
217 seems to be the result of the decreasing quantities of magnetite under higher Se concentration
218 conditions (Fig. 1). However, even in the presence of $100 \mu\text{M}$ SeO_3^{2-} , approximately 80% of
219 magnetotactic bacteria were magnetically recovered within 20 hours. To confirm the
220 biomagnetic recovery of Se, the amount of Se from magnetically recovered harvested cells was
221 measured and revealed 3.6×10^8 Se atoms per cell recovery. Though some Se was lost during the
222 recovery process (3.0×10^8 Se atoms after recovery), the result clearly shows that magnetotactic
223 bacteria could be applied in biomagnetic recovery of Se from SeO_3^{2-} containing water. We note

224 that a more effective recovery could be established by process optimization (e.g. cell number,
225 vessel size and magnetic force enhancement).

226 Current genetic and environmental microbiological research shows that magnetic particle
227 production within bacteria occurs across a diverse group of bacterial species. In fact, the genetic
228 region corresponding to magnetosome formation, called magnetosome island (MAI), is found
229 within microbes spread across the phylogenetic tree. As *M. magneticum* AMB-1 does not show
230 strong resistance to SeO_3^{2-} (Fig. 1), a magnetotactic bacterial species with higher tolerance and
231 effective accumulation of target molecule could be found and used to improve the biomagnetic
232 recovery; identified either from environments local to the bioremediation site or through
233 evolving conditions to those similar to the polluted environment for a range of candidate
234 magnetotactic bacteria. In addition, recently, magnetosome formation was enabled in another
235 bacterial species by artificially transferring key genetic regions of the MAI into the host
236 organism (31). Therefore, the induction of magnetosome formation within known bacteria
237 showing high resistance to target element is another promising approach to improve the
238 biomagnetic recovery efficiency.

239 In conclusion, in this study we showed the first account of amorphous elemental Se particle
240 formation from the reduction of SeO_3^{2-} within the magnetotactic bacterial cell, completely
241 independent of the crystallization of magnetite within the cells' magnetosomes. The cells were
242 accumulated the highest amount of Se compared to any other foreign elements. For example, 2.4
243 and 174 times more Se was accumulated as compared to Te in cells and Cd^{2+} adsorption onto cell
244 surfaces. Importantly, the Se accumulating bacteria were successfully recovered with an external

245 magnetic field. Therefore, we believe magnetotactic bacteria have the unique advantage of
246 biomagnetic cell recovery, providing a new effective methodology for bioremediation of polluted
247 water and additional potential to utilize the pollutant product for further material applications.

248

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257 N. H., and S. S. performed the experiments. All authors analyzed the data. M. T., A. A., S. S., and
258 T. M. wrote the paper. All authors have no conflict of interest directly relevant to the content of
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260

261 REFERENCES

- 262 1. **Adams GO, Fufeyin PT, Okoro SE, Ehinomen I.** 2015. Bioremediation, biostimulation
263 and bioaugmentation: A Review. *Int J Environ Bioremediation Biodegrad* **3**:28–39.
- 264 2. **Tordoff G., Baker AJ, Willis A.** 2000. Current approaches to the revegetation and
265 reclamation of metalliferous mine wastes. *Chemosphere* **41**:219–28.
- 266 3. **Kleindienst S, Paul JH, Joye SB.** 2015. Using dispersants after oil spills: impacts on the
267 composition and activity of microbial communities. *Nat Rev Microbiol* **13**:388–96.
- 268 4. **Jogler C, Schüler D.** 2009. Genomics, genetics, and cell biology of magnetosome
269 formation. *Annu Rev Microbiol* **63**:501–21.
- 270 5. **Matsunaga T, Suzuki T, Tanaka M, Arakaki A.** 2007. Molecular analysis of
271 magnetotactic bacteria and development of functional bacterial magnetic particles for
272 nano-biotechnology. *Trends Biotechnol* **25**:182–8.
- 273 6. **Rahn-Lee L, Komeili A.** 2013. The magnetosome model: insights into the mechanisms of
274 bacterial biomineralization. *Front Microbiol* **4**:352.
- 275 7. **Komeili A, Li Z, Newman DK, Jensen GJ.** 2006. Magnetosomes are cell membrane
276 invaginations organized by the actin-like protein MamK. *Science* **311**:242–5.
- 277 8. **Scheffel A, Schüler D.** 2007. The acidic repetitive domain of the *Magnetospirillum*
278 *gryphiswaldense* MamJ protein displays hypervariability but is not required for
279 magnetosome chain assembly. *J Bacteriol* **189**:6437–46.
- 280 9. **Blakemore RP.** 1975. Magnetotactic bacteria. *Science* **19**:377–9.
- 281 10. **Arakaki A, Takeyama H, Tanaka T, Matsunaga T.** 2002. Cadmium recovery by a
282 sulfate-reducing magnetotactic bacterium, *Desulfovibrio magneticus* RS-1, using magnetic
283 separation. *Appl Biochem Biotechnol* **98-100**:833–40.
- 284 11. **Tanaka M, Nakata Y, Mori T, Okamura Y, Miyasaka H, Takeyama H, Matsunaga T.**
285 2008. Development of a cell surface display system in a magnetotactic bacterium,
286 “*Magnetospirillum magneticum*” AMB-1. *Appl Environ Microbiol* **74**:3342–8.
- 287 12. **Tanaka M, Kawase M, Tanaka T, Matsunaga T.** 2011. Gold biorecovery from plating
288 waste by magnetotactic bacterium, *Magnetospirillum magneticum* AMB-1. *MRS Proc*
289 **1169**:1169–Q03–12.
- 290 13. **Tanaka M, Arakaki A, Staniland SS, Matsunaga T.** 2010. Simultaneously discrete
291 biomineralization of magnetite and tellurium nanocrystals in magnetotactic bacteria. *Appl*
292 *Environ Microbiol* **76**:5526–32.
- 293 14. **Yang C-P, Yin Y-X, Guo Y-G.** 2015. Elemental selenium for electrochemical energy
294 storage. *J Phys Chem Lett* **6**:256–66.
- 295 15. **Lemly AD.** 2002. Symptoms and implications of selenium toxicity in fish: the Belews Lake
296 case example. *Aquat Toxicol* **57**:39–49.
- 297 16. **Srikanth Lavu R V., Van De Wiele T, Pratti VL, Tack F, Du Laing G.** 2015. Selenium
298 bioaccessibility in stomach, small intestine and colon: comparison between pure Se
299 compounds, Se-enriched food crops and food supplements. *Food Chem* **197**:382–87.
- 300 17. **Johansson CL, Paul NA, de Nys R, Roberts DA.** 2016. Simultaneous biosorption of
301 selenium, arsenic and molybdenum with modified algal-based biochars. *J Environ Manage*

- 302 **165**:117–23.
- 303 18. **Silva IR, Serrão VHB, Manzine LR, Faim LM, da Silva MTA, Makki R, Saidemberg**
304 **DM, Cornélio ML, Palma MS, Thiemann OH.** 2015. Formation of a ternary complex for
305 selenocysteine biosynthesis in bacteria. *J Biol Chem* **290**:29178–88.
- 306 19. **Nancharaiah Y V., Lens PNL.** 2015. Ecology and biotechnology of selenium-respiring
307 bacteria. *Microbiol Mol Biol Rev* **79**:61–80.
- 308 20. **Tanaka M, Brown R, Hondow N, Arakaki A, Matsunaga T, Staniland S.** 2012. Highest
309 levels of Cu, Mn and Co doped into nanomagnetic magnetosomes through optimized
310 biomineralisation. *J Mater Chem* **22**:11919.
- 311 21. **Staniland S, Williams W, Telling N, Van Der Laan G, Harrison A, Ward B.** 2008.
312 Controlled cobalt doping of magnetosomes *in vivo*. *Nat Nanotechnol* **3**:158–62.
- 313 22. **Matsunaga T, Sakaguchi T, Tadakoro F.** 1991. Magnetite formation by a magnetic
314 bacterium capable of growing aerobically. *Appl Microbiol Biotechnol* **35**:651-5
- 315 23. **Tanaka M, Okamura Y, Arakaki A, Tanaka T, Takeyama H, Matsunaga T.** 2006.
316 Origin of magnetosome membrane: proteomic analysis of magnetosome membrane and
317 comparison with cytoplasmic membrane. *Proteomics* **6**:5234–47.
- 318 24. **Nies DH.** 1999. Microbial heavy-metal resistance. *Appl Microbiol Biotechnol* **51**:730–50.
- 319 25. **Klonowska A, Heulin T, Vermeglio A.** 2005. Selenite and tellurite reduction by
320 *Shewanella oneidensis*. *Appl Environ Microbiol* **71**:5607–9.
- 321 26. **Ruta L, Paraschivescu C, Matache M, Avramescu S, Farcasanu IC.** 2010. Removing
322 heavy metals from synthetic effluents using “kamikaze” *Saccharomyces cerevisiae* cells.
323 *Appl Microbiol Biotechnol* **85**:763–71.
- 324 27. **Baesman SM, Bullen TD, Dewald J, Zhang D, Curran S, Islam FS, Beveridge TJ,**
325 **Oremland RS.** 2007. Formation of tellurium nanocrystals during anaerobic growth of
326 bacteria that use Te oxyanions as respiratory electron acceptors. *Appl Environ Microbiol*
327 **73**:2135–43.
- 328 28. **Rathgeber C, Yurkova N, Stackebrandt E, Beatty JT, Yurkov V.** 2002. Isolation of
329 tellurite- and selenite-resistant bacteria from hydrothermal vents of the Juan de Fuca Ridge
330 in the Pacific Ocean. *Appl Environ Microbiol* **68**:4613–22.
- 331 29. **Bazyliniski DA, Dean AJ, Williams TJ, Long LK, Middleton SL, Dubbels BL.** 2004.
332 Chemolithoautotrophy in the marine, magnetotactic bacterial strains MV-1 and MV-2.
333 *Arch Microbiol* **182**:373–87.
- 334 30. **Spring S, Amann R, Ludwig W, Schleifer KH, van Gemerden H, Petersen N.** 1993.
335 Dominating role of an unusual magnetotactic bacterium in the microaerobic zone of a
336 freshwater sediment. *Appl Environ Microbiol* **59**:2397–403.
- 337 31. **Kolinko I, Lohße A, Borg S, Raschdorf O, Jogler C, Tu Q, Pósfai M, Tompa E, Plitzko**
338 **JM, Brachmann A, Wanner G, Müller R, Zhang Y, Schüler D.** 2014. Biosynthesis of
339 magnetic nanostructures in a foreign organism by transfer of bacterial magnetosome gene
340 clusters. *Nat Nanotechnol* **9**:193–7.
- 341
- 342

343 **FIGURE LEGENDS**

344 **Fig. 1. Tolerance of *M. magneticum* AMB-1 to SeO_3^{2-} and magnetite nano-particle**
345 **synthesis.**

346 The number of cells (\square) and magnetite crystals (\bullet) grown in different concentrations (0, 5, 10,
347 20, 40, 60, 80, 100, and 250 μM) of SeO_3^{2-} were directly counted. To evaluate the number of
348 magnetite within the cells, over 50 cells randomly selected were manually counted. Error bars
349 show SDs.

350

351 **Fig. 2. Transmission electron micrographs, and STEM-EDX analyses for magnetite and**
352 **Se within magnetotactic bacteria.**

353 (a) TEM micrographs of magnetotactic bacteria grown i) in the presence of SeO_3^{2-} (100 μM) and
354 ii) in its absence. Characteristic intracellular granules were indicated with arrows. Scale bar
355 indicates 100 nm. (b) TEM image and STEM-EDX maps of Se, Fe, and O taken using a probe
356 size of approximately 5 nm. (c) Spot EDX spectra of *i and *ii in b) as a representation of Se
357 and magnetite. The Cu signal is from copper TEM grid.

358

359 **Fig. 3. SeO_3^{2-} removal during magnetotactic bacterial cell growth.**

360 SeO_3^{2-} removal using magnetotactic bacteria (\circ) and cell growth (\blacklozenge) was evaluated in the
361 presence of 100 μM SeO_3^{2-} for 7 days. The average values from three independent experiments
362 were obtained. Error bars show standard deviations.

363

364 **Fig. 4. Magnetic recovery assay of Se granule-containing *M. magneticum* AMB-1.** The
365 percentage of recovered cells is calculated from the initial cell numbers (1.0×10^8 /ml) by
366 counting the number of dispersed cells left within the culture medium. In addition, the number of
367 cells recovered by magnetic force was also verified by counting the cells recovered at the end
368 points. *M. magneticum* AMB-1 was cultured and assayed with the respective concentrations of
369 SeO_3^{2-} (SeO_3^{2-} concentration = 0 μM (control) (■), 25 μM (○), 50 μM (▲), and 100 μM (◇)).
370 The average values from three independent experiments were obtained. Error bars show standard
371 deviations.

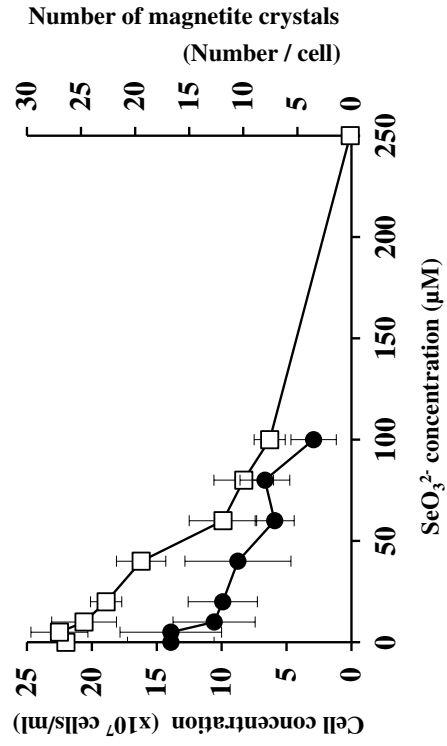


Figure 1

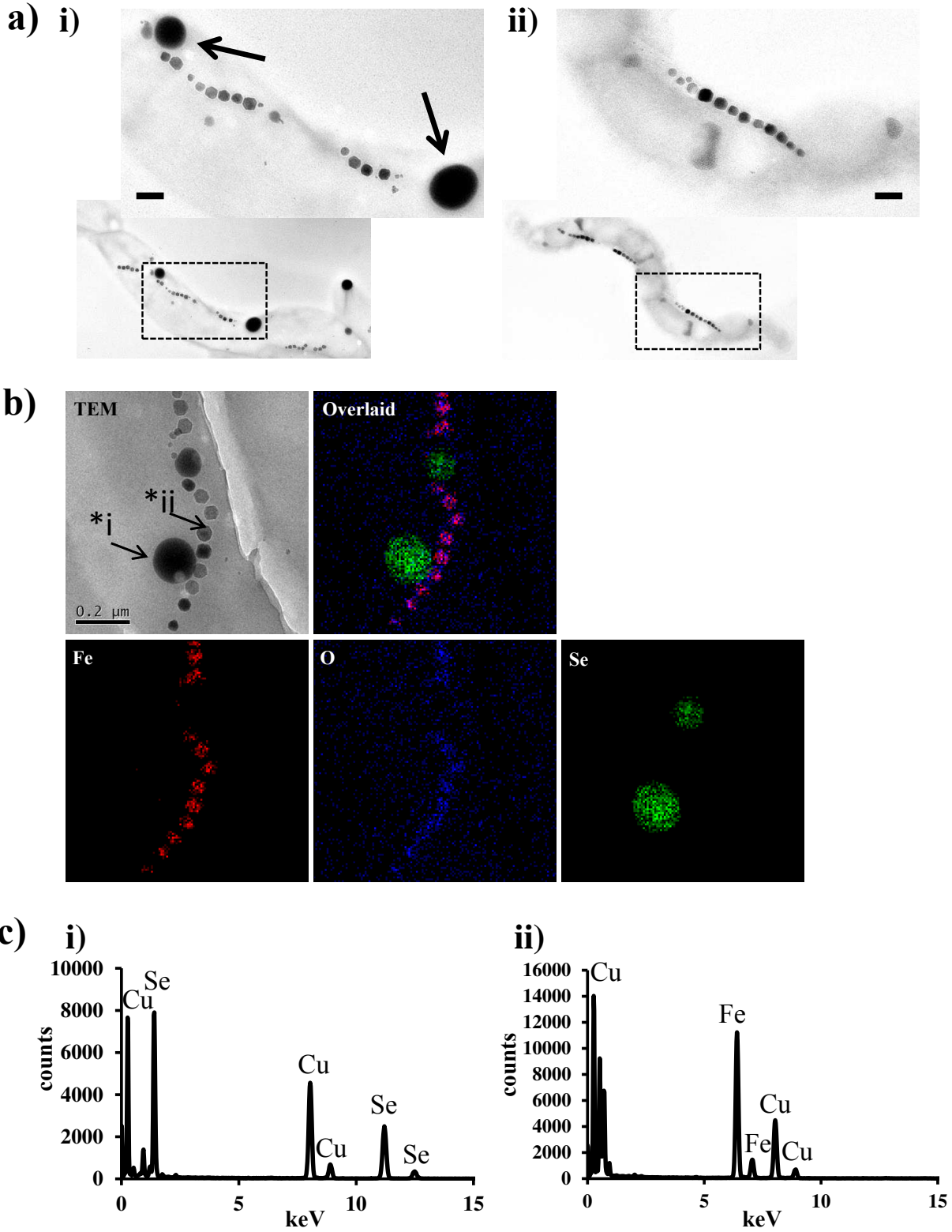


Figure 2

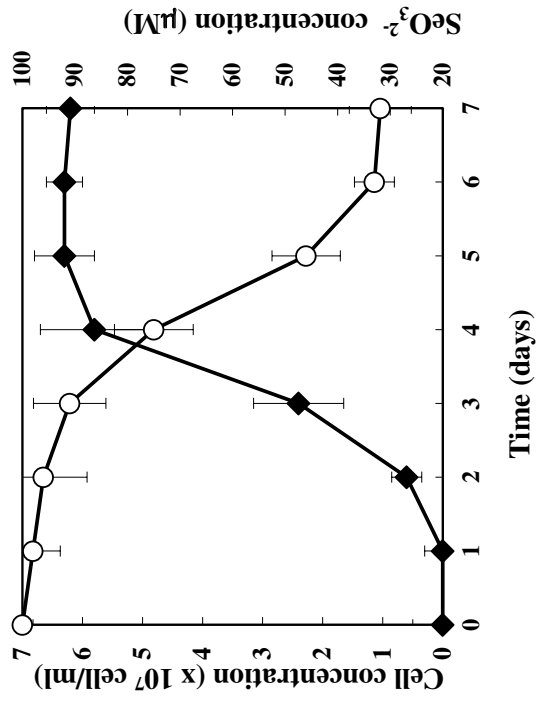


Figure 3

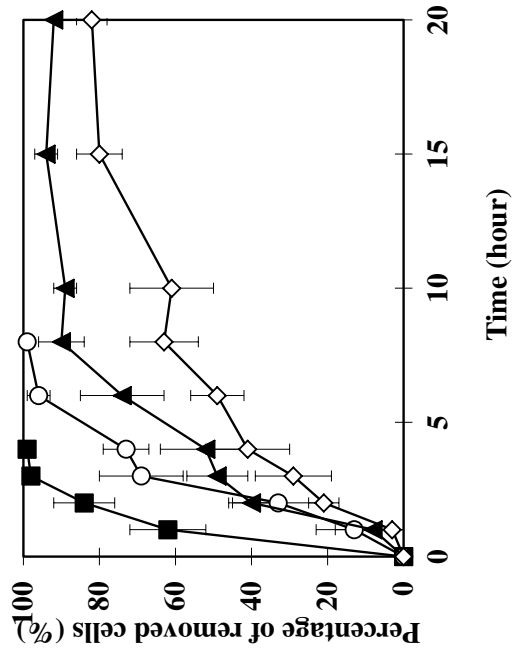


Figure 4