1	Supplementary information for					
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3	Magnetotactic bacteria accumulate a large pool of iron distinct					
4	from their magnetite crystals					
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11 Supplementary text

12 Incubation of a mutant AMB-1 strain unable to form magnetosomes with FIP-1

13 An AMB-1 mutant unable to form magnetosomes from which the entire MAI has been deleted (Δ MAI strain) was used as a negative control for FIP-1 experiments. Δ MAI cells were 14 15 cultivated as described above in the absence of an iron source. To ensure that no iron 16 contamination from the glassware occurs, tubes containing Δ MAI cells were soaked with an oxalic acid solution (concentration of 4.25 g per liter) overnight. After incubation with FIP-1, 17 bacteria were centrifuged and suspended in fresh PBS to remove FIP-1 probes in the external 18 19 solution. Wild-type and Δ MAI AMB-1 were incubated with FIP-1 for 90 min, and observed by Structured Illumination Microscopy with a Carl Zeiss Elyra PS.1 Super Resolution 20 21 fluorescence microscope, as described in the main text. Results are shown in Figs. S4 and S5.

22

23 Magnetite size and shape factor

The length of magnetite crystals produced by wild-type AMB-1 was measured from electron microscopy images using the ImageJ software. Results are given in Fig. S2, and show longer crystals when AMB-1 is cultivated under higher iron conditions. The nanoparticle shape anisotropy was then quantified by measuring the shape factor (s) defined as:

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29
$$s = \frac{width}{length}$$
 (Eq. S1)

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Assuming a consistent shape factor, a linear relationship exists between the width and the length. Results are given in Fig. S3, and show a rough linear relationship between magnetite width and length, with correlation coefficients of 0.86 and 0.89 for iron concentration 34 conditions of 30 and 150 μ M, respectively. Shape factors showed similar values (0.80 and 35 0.82) almost similar to what has been measured in AMB-1 before (1).

36

37 Saturation magnetization in mutant strains

The saturation magnetization measured in $\Delta mamT$ samples showed a 100-fold decrease 38 39 compared to the wild-type strain (Table 5). From measurement of (i) the crystal size, (ii) the 40 number of crystals per cell, and (*iii*) the shape factor (width/length ratio) in $\Delta mamT$ and wild-41 type strains (Table S2), only a ~30-fold decrease of magnetite volume and associated saturation magnetization can be expected. The detection limit of the VSM we used for the 42 43 magnetic characterizations is 300 times lower than the magnetization in $\Delta mamT$ samples. Even though 2D projections of 3D objects can generate biases, the lower saturation 44 45 magnetization than expected in $\Delta mamT$ AMB-1 could be explained by crystal phases distinct 46 from magnetite and showing lower saturation magnetization. One example of such phases include hematite, which has a saturation magnetization of 0.3 emu/g(2). From single-crystal 47 48 characterization, Jones and co-workers identified only magnetite crystals in $\Delta mamT$ AMB-1 (3). Therefore, it is possible that some other crystal phases were missed. Dedicated 49 experimental work and bulk characterizations of iron phases in $\Delta mamT$ AMB-1 will be 50 51 needed to address this question.

52 Crystal phases in $\Delta mamP$ AMB-1 were characterized by Jones and co-workers following the 53 same single-crystal approach (3). Therefore, additional crystal phases than magnetite might as 54 well be contained in this mutant strain. However, the decrease in saturation magnetization of 55 $\Delta mamP$ compared to the wild-type strain expected from electron microscopy (~10-fold, see 56 Table S2) showed a much lower discrepancy than the one we measured (~20-fold, see Table 57 5). In this case, the discrepancy between expected and measured saturation magnetization 58 could also be explained from biases generated by 2D projection of 3D objects. Moreover,

- 59 magnetite porosity in MTB has never been characterized. The density of magnetite could also
- 60 show variations between the several AMB-1 strains we tested.

61 Supplementary references

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73 Supplementary figures



FIG S1 Hysteresis loops acquired from whole bacterial cultures, corresponding to wild-type AMB-1 cultivated under low- and high-iron conditions, and the $\Delta mamP$ and $\Delta mamT$ strains. Note the different y-axes.



FIG S2 Distribution of magnetite length produced in wild-type AMB-1 cultivated for three





FIG S3 Distribution of magnetite length produced in $\Delta mamP$ (black bars) and 150 $\Delta mamT$

^{85 (}white bars) AMB-1.



FIG S4 Magnetite width represented as a function of magnetite length in AMB-1 cultivated
with iron at 30 (left) or 150 (right) μM. Assuming a constant shape factor (s) (Eq. S1), a linear
relationship exists between width and length. Data are represented with the best linear fit, as
well as two linear fits for a given shape factor of 1 and 0.5.





FIG S5 Red (left panels), green (center panels), and merged (right panels) fluorescence images of wild-type AMB-1 incubated with FIP-1 for 90 min. Scale bars = 1 μ m.





FIG S6 Red (left panels), green (center panels), and merged (right panels) fluorescence images of Δ MAI AMB-1 incubated with FIP-1 for 90 min. Scale bars = 1 μ m.



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100 **FIG S7** Red (left panels), green (center panels) and merged (right panels) fluorescence 101 images of dividing wild-type cells incubated with FIP-1 for 90 min. Arrows point at higher 102 green fluorescence signal in the cell that may be located at the septum location during cell 103 division. Scale bars = 1 μ m.



106 FIG S8 Red (left panels), green (center panels) and merged (right panels) fluorescence

- 107 images of wild-type AMB-1 incubated with FIP-1 for 180 min. Scale bars = 1 μ m.
- 108



110 FIG S9 Red (left panels), green (center panels) and merged (right panels) fluorescence

¹¹¹ images of $\Delta mamP$ AMB-1 incubated with FIP-1 for 180 min. Scale bars = 2 μ m.



114 FIG S10 Red (left panels), green (center panels) and merged (right panels) fluorescence

¹¹⁵ images of $\Delta mamT$ AMB-1 incubated with FIP-1 for 180 min. Scale bars = 2 μ m.

117 Supplementary tables

Mass of iron in the initial medium (mg)	Mass of iron in the final medium (mg)	Mass of iron in bacteria (mg)	Mass of iron leaked outside of bacteria (mg)	Iron recovery (%)
2.31	2.20	0.06	0.28 10-3	96
2.31	2.15	0.07	0.17 10-3	96
2.37	2.31	0.05	0.27 10-3	100
2.27	2.11	0.08	0.23 10-3	97

Table S1 Iron mass balance in AMB-1 wild-type cultures.

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- 121 Table S2 Magnetite crystal parameters measured from electron microscopy observations in
- 122 wild-type (cultivated at either 30 or 150 μ M of iron), $\Delta mamP$ and $\Delta mamT$ AMB-1 strains.
- 123 Errors indicate 1 SD.

	Wild-type (30 µM of iron)	Wild-type (150 µM of iron)	∆ <i>mamP</i>	∆ <i>mamT</i>
Mean magnetite length (nm)	32.03	38.46	24.13	20.2
Width / length ratio	0.81	0.83	0.72	0.68
Number of crystals per cell	17.20 ± 4	20.75 ± 6	9.00 ± 3.57	21.00 ± 6.53
Number of crystals analyzed	344	351	205	274

126 **References**

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