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## Flow cytometry as a rapid analytical tool to determine physiological responses to changing O<sub>2</sub> and iron concentration by Magnetospirillum gryphiswaldense strain MSR-1

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1 Flow cytometry as a rapid analytical tool to determine physiological responses to

2 changing O<sub>2</sub> and iron concentration by *Magnetospirillum gryphiswaldense* strain

3 MSR-1

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#### 12 Abstract.

13 Magnetotactic bacteria (MTB) are a diverse group of bacteria that synthesise 14 magnetosomes, magnetic membrane-bound nanoparticles that have a variety of diagnostic, 15 clinical and biotechnological applications. We present the development of rapid methods 16 using flow cytometry to characterize several aspects of the physiology of the commonly-used 17 MTB Magnetospirillum gryphiswaldense MSR-1. Flow cytometry is an optical technique that 18 rapidly measures characteristics of individual bacteria within a culture, thereby allowing 19 determination of population heterogeneity and also permitting direct analysis of bacteria. 20 Scatter measurements were used to measure and compare bacterial size, shape and 21 morphology. Membrane permeability and polarization were measured using the dyes 22 propidium iodide and bis-(1,3-dibutylbarbituric acid) trimethine oxonol to determine the 23 viability and 'health' of bacteria. Dyes were also used to determine changes in concentration 24 of intracellular free iron and polyhydroxylakanoate (PHA), a bacterial energy storage 25 polymer. These tools were then used to characterize the responses of MTB to different  $O_2$ 26 concentrations and iron-sufficient or iron-limited growth. Rapid analysis of MTB physiology 27 will allow development of bioprocesses for the production of magnetosomes, and will 28 increase understanding of this fascinating and useful group of bacteria.

29

INTRODUCTION. Magnetic nanomaterials are increasingly important products with myriad
 applications in diverse settings including but not limited to environmental pollution control,
 information and energy storage (1), catalysis (2), biotechnological (1, 3, 4) and especially

33 biomedical research (1, 5-7). While most are produced by chemical means there is growing 34 interest in harnessing the cellular machinery of certain naturally occurring bacteria (8-12) to 35 generate useful magnetic, and other metallic nanoparticle materials, biologically. In this 36 context, 'magnetosomes', magnetic nanoparticle based organelles naturally contained within 37 magnetotactic bacteria (MTB), are particularly important (8, 13-17). In most MTB, 38 magnetosomes are arranged in one or more highly ordered 'compass needle-like' chains of 39 single-domain permanently ferrimagnetic magnetite ( $Fe_3O_4$ ) or greigite ( $Fe_3S_4$ ) crystals (35-40 120 nm diameter) each wrapped in a phospholipid bilayer membrane containing a unique set 41 of magnetosome specific proteins, i.e. distinct from those of cytoplasmic and outer 42 membranes (8, 13, 15). These internal structures within MTB function as navigational 43 devices essential for magnetotaxis (18). Unique properties of magnetosomes, not normally 44 associated with chemically synthesized magnetic nanoparticles, of narrow size distribution, uniform morphology, high crystal purity, permanent magnetic character, high heating 45 46 capacity, low aggregation tendency, ready dispersion in aqueous solution, facile 47 functionalization, high biocompatibility, low toxicity and high specific absorption rates (10, 12) 48 make them especially attractive prospects for biotech and healthcare applications, i.e. in 49 immunoassays (19), magnetic affinity cell sorting (20), magnetic resonance imaging (21), 50 drug and gene delivery (22) and cancer therapy (12, 23).

51 It is recognised that future widespread application of magnetosomes will, to a large extent, 52 depend on the development of intensified high yielding manufacturing platforms for 53 magnetosomes (10, 12, 16). Fundamental to this are appropriate means for analysing MTB 54 growth, viability, physiology and biomineralization of magnetic iron minerals, in order to 55 understand and optimise magnetosome formation at any scale, from initial small (millilitre) 56 studies on strain isolation and cultivability in the laboratory, and pilot scale manufacture (10 57 - 100 L), to fully fledged commercial production in cubic metre scale bioreactors. Qualitative 58 evidence of magnetosome production within MTB can be obtained by observing a 59 shimmering effect in cell suspensions mounted on magnetic stirrer plates, and black 60 coloration of cell suspensions and/or colonies on agar plates, while magneto-61 spectrophotometric assay of cellular magnetism (Cmaa) of suspended cells provides a rapid 62 indirect measure of cellular magnetosome content (24, 25). Quantitative determination of 63 magnetosome content in cells and during subsequent recovery and purification operations 64 usually involves measurement of iron content by means of atomic absorption spectrometry 65 (dependent on species and cultivation conditions magnetosomes account for 80 to 99.5 % of 66 the total cell-bound iron in magnetic cells (18, 26, 27)), combined with imaging of 67 magnetosomes by transmission electron microscopy. Recent work indicates the importance 68 of monitoring physiological stress indicators to identify optimal conditions for magnetosome

69 formation (28), and the utility of transcriptome analysis for comparing magnetosome forming 70 and non-forming conditions in MTB (29). Other analytical methods especially pertinent to 71 pilot- and large-scale magnetosome production and downstream processing from high 72 biomass MTB fermentations include the tracking of polyhydroxyalkanoate (PHA) granules. 73 Here, PHA formation diverts cellular resources from growth, lowering yields, and high levels 74 of PHA would be likely to be a troublesome contaminant of magnetosome preparations. 75 Current procedures for the determination of PHA content employ lengthy procedures 76 involving solvent extraction, derivatization and gas chromatography (29).

77 With the exception of at line optical density and C<sub>mag</sub> measurements all of the 78 aforementioned techniques are labour intensive and/or time consuming. The development of 79 analytical methods is essential for the development of robust production processes, itself a 80 requirement for industrialisation implementation. It is desirable that such methods will be 81 rapid, requiring small volumes of samples and provide data of cellular parameters without 82 the need of further growth, thus giving a 'snapshot' of the current physiological state of the 83 cells. The flow cytometry (FCM) methods applied in this study fulfil these requirements. FCM 84 has previously been applied previously for rapid analysis of microbial physiology (30) and 85 expression of auto-fluorescent proteins (31), monitoring recombinant protein production (32) 86 and for investigating population heterogeneities in cultures. In FCM, multiple physical 87 characteristics of single particles suspended in a fluid can be measured concurrently as they 88 flow through a beam of light. FCM is a fast single-cell analysis technique well suited to 89 collection of large datasets (tens of thousands of cells can be analysed) and allows 90 determination of light scatter (relative size and granularity/internal complexity) and 91 fluorescence properties of individual cells and thus determination of population 92 heterogeneity. An important advantage of FCM is that it does not rely upon microbial growth 93 for analysis of cell viability. 'Viable but non-culturable' (VBNC) cells exist within most 94 microbial cell populations (33), but growth-based methods for determining viable cell 95 numbers (total viable counts generating colony forming unit, CFU data) will not detect the 96 VBNC phenotype, thus total viable cell concentrations are underestimated. FCM does not 97 share this limitation. MTB grow very slowly on agar plates, for example, M. gryphiswaldense 98 MSR-1 forms colonies after 7 – 10 days (26). Regardless of cell type FCM analysis can be 99 performed in a matter of seconds. Moreover, when combined with carefully selected 100 mixtures of fluorescent probes FCM can be employed to determine the physiological state of 101 single cells.

102 Reports on the application of FCM to MTB are few in number (34, 35) and the full power of 103 the technique has not exploited in any case. FCM has been used for analysing gene expression in *M. gryphiswaldense* MSR-1 (34), and in the development of new expression systems for the same species (35). Green fluorescent protein (GFP) was used as a reporter in both studies, i.e. for magnetosomal localization and expression of GFP tagged magnetosome proteins under magnetite forming conditions (34); and for identification of promoters (fused to GFP) for efficient gene expression (35).

109 In this work, we present a battery of FCM methods tailored *a priori* to the study of *M*. 110 *gryphiswaldense* MSR-1 and other MTB, and applicable to cells grown in liquid cultures and 111 on agar plates. Specifically, we describe methods for determination of cellular concentration, 112 cell size distribution, single-cell physiology and relative changes over time of intracellular 113 contents of PHA and the chelatable iron pool.

#### 114 **RESULTS AND DISCUSSION**

#### 115 Morphological difference between cells grown on plates and in suspension.

116 FCM analysis was employed to monitor cell size and optical complexity of M. 117 gryphiswaldense MSR-1 by means of light scattering. In FCM, light scatter is collected at two 118 different angles: in the direction of the laser path (forward light scatter, FSC); and orthogonal 119 to it (side scatter, SSC). For spherical particles (e.g. of latex), FSC correlates with the 120 logarithm of particle diameter (36). For cells and other non-spherical particles, changes in 121 FSC are roughly indicative of changes in cell size. When applied to cells, SSC measures 122 'granularity' (37), a parameter that includes optical complexity caused by particulate material 123 contained within the cell. Figure 1 shows the results of comparative FCM scatter and light 124 microscopic analyses of MSR-1 cells cultured on ACA plates (resuspended in PBS) and in 125 the liquid medium FSM. Clear differences in the heterogeneity of cell populations cultured in 126 FSM (Fig. 1a) cf. those grown on ACA (Fig. 1b) can be discerned from the scatter patterns of 127 FSC vs. SSC dot plots (Figs 1a & 1b). Larger cells are represented by high FSC-A values (y-128 axis) whereas more granular cells are characterized by higher SSC-A (x-axis) values. Cells 129 grown in liquid FSM appear less heterogeneous, smaller and less granular than those grown 130 on ACA plates. Moreover, differences in particle size distribution and cell shape of 131 suspension and plate grown MSR-1 cells are respectively inferred from 'Count vs FSC-A' 132 histograms and light microscopy, with plate-grown cells appearing more polydisperse in size 133 (Fig 1c) and filamentous (Fig. 1e) compared to liquid-grown cells (Fig. 1d).

#### 134 **Determination of cell concentration by FCM.**

FCM analysis was also used as a rapid method for determining cell concentrations in shakeflask experiments. In auto-calibration mode and operating at a medium flow rate of 35

 $\mu$ L·min<sup>-1</sup> a strong correlation (R<sup>2</sup> > 0.95) between OD<sub>565</sub> and FCM event counts was 137 observed for MSR-1 cells (Supplemental Fig. S1a) with 1 OD<sub>565</sub> equivalent to 1.16 × 10<sup>9</sup> 138 139 cells mL<sup>-1</sup>. This relationship is strikingly different to Schultheiss and Schüler's (38) correlation 140 of OD<sub>565</sub> with CFU, i.e. 1 OD<sub>565</sub> equivalent to  $3.3 \times 10^8$  CFU mL<sup>-1</sup> and likely reflects an important advantage of FCM over CFU counting, namely its ability to detect viable but non-141 142 culturable cells (VBNC). We also used Syto<sup>®</sup>62, a permeant DNA dye, to stain MSR-1 cells 143 and so distinguish them from noise particles of similar size; with Syto®62-stained cells a similar correlation was found between  $OD_{565}$  and cell count ( $OD_{565} = 1.03 \times 10^9$  cells mL<sup>-1</sup>; 144 145 Supplemental Fig. S1b).

#### 146 Use of FCM to determine MSR-1 membrane polarization and cellular death.

147 Two fluorescent probes were used to monitor the respiratory potential and death of M. 148 gryphiswaldense MSR-1 cells using FCM. BOX (DiBac<sub>4</sub>(3); bis-(1,3-dibutylbarbituric acid) 149 trimethine oxonol) is a green lipophilic fluorescent probe that can only enter cells if their 150 membranes are depolarized (39). Healthy cells possess intact polarized cytoplasmic 151 membranes, which are impermeant to BOX (BOX<sup>-</sup>). In contrast, cells with depolarised 152 cytoplasmic membranes (whether injured, stressed or dead) permit BOX access and 153 fluoresce green (BOX<sup>+</sup>). PI (propidium iodide), a nucleic acid intercalator, is excluded by the 154 intact membrane of viable cells (PI<sup>-</sup>), but is taken up by dead cells which fluoresce red (PI<sup>+</sup>) 155 (39). Staining procedures were optimised using actively growing *M. gryphiswaldense* cells, 156 starving cells and dead cells killed with ethanol. Fig. 2 shows the two-colour fluorescence dot 157 plots of MSR-1 cells co-stained with BOX and PI (fluorescence being detected on FL1-A and 158 FL3-A channels, respectively). The fluorescence patterns from actively growing magnetic 159 cells (Fig. 2a) and non-magnetic cells (Fig. 2c) were strikingly alike; i.e.: 86-90 % of the cell 160 populations were 'healthy', staining negatively with both fluorescent markers (BOX<sup>-</sup> PI<sup>-</sup>, 161 Quadrant 1): 5-8 % were 'injured', staining positively with BOX, but negatively with PI (BOX<sup>+</sup> 162  $PI^{-}$ , Quadrant 2); and 3-4 % were dead (BOX<sup>+</sup>  $PI^{+}$ , Quadrant 3). The fluorescence patterns 163 from starving magnetic (Fig. 2b) and non-magnetic (Fig. 2d) MSR-1 cells were comparable 164 with one another, but indicated noticeably fewer healthy populations (31-39 % healthy, ~40% 165 injured, 19-26% dead) than those of actively growing cells (~90 % healthy, 5-8 % injured, <3 166 % dead). The low healthy population in starving cultures suggests the presence of large 167 numbers of VBNC cells. This confirms the observation that the correlation between OD<sub>565</sub> and cells·mL<sup>-1</sup> as measured by FCM is different to the correlation between OD<sub>565</sub> and CFU 168 169 mL<sup>-1</sup> (38) due to the presence of VBNC cells. Only 5 % of the positive control population, i.e. 170 cells killed with ethanol, were healthy MSR-1 cells (Fig. 2e).

#### 171 Accumulation of PHA aggregates in MSR-1.

172 It has been widely reported that limiting nitrogen and  $O_2$  availability under carbon excess 173 results in high-level accumulation of polyhydroxyalkanoates (PHA) in several species of 174 bacteria (40-44). Ban et al. (43) specifically examined the effect of hydrogen metabolism on 175 cell growth and magnetosome synthesis in *M. gryphiswaldense* MSR-1 concluding that in 176 MTBs PHA formation occurs under conditions of excess reducing power. Liu and co-workers 177 (45) succeeded in isolating an MSR-1 mutant capable of higher level magnetosome 178 production and lower PHA accumulation than the wild type, indicating a possible link 179 between the formation of PHA and magnetosomes in this bacterium. In more recent work, 180 genomic excision of the phbCAB operon in MSR-1 was shown to abolish PHA granule 181 formation albeit at the expense of much reduced growth (46). Collectively, the above findings 182 hint at the existence of an energy competition between the processes of PHA and 183 magnetosome formation. Here we have used FCM to investigate PHA accumulation within 184 individual bacteria of starved non-magnetic and magnetic MTB cultures. Cells were stained 185 with the lipophilic dye 1,3,5,7,8-pentamethylpyrromethene-difluoroborate complex 186 (pyrromethene-546 or Pyr-546) which on entering bacteria stains PHA green (47). Previous 187 studies have shown that Pyr-546 fluorescence correlates to intracellular PHA content (48) 188 and is superior to Nile red as a dye for PHA. After incubating for various times (10 – 300 s) 189 samples were immediately analysed by FCM. Fig. 3 shows that when used at a 190 concentration of 0.5 µg mL<sup>-1</sup> the timeframe for Pyr-546 penetration and near full staining of 191 intracellular PHA was the time taken to add the dye and analyse the sample (of the order of 192 10–15 s). No further enhancement in fluorescence occurred between 70 and 300 s exposure 193 to Pyr-546 for both non-magnetic (Fig. 3a) and magnetic (Fig. 3b) cells. This said, FCM 194 analysis reveals salient differences in the PHA content of magnetic and non-magnetic MSR-195 1, for example, revealing the presence of two discrete populations with low (Fig. 3b, labelled 196 '1') and high (Fig. 3b, labelled '2') PHA content in magnetic cells cf. just a single 'high PHA' 197 population in non-magnetic cells (Fig. 3a marked '2'). Fluorescence microscopy images of 198 cells containing different quantities of Pyr-546 stained PHA are shown in Fig. 3. Similarly to 199 our findings, recent studies performed with Cupriavidus necator observed sub-populations 200 with more and less PHA (47). Other works with Pseudomonas putida have recently observed 201 an asymmetric PHA distribution during cell division under carbon limiting conditions 202 suggesting that this could be explained by different cellular growth rates, distinct ability to 203 degrade PHA or uneven distribution of PHA granules to daughter cells (49).

#### 204 Measurement of intracellular chelatable iron.

The intracellular pool of chelatable iron is considered a critical component in the biomineralization of magnetosomes. Recent studies in *M. gryphiswaldense* MSR-1 suggest 207 that at least some of the iron transport for magnetite synthesis occurs through two copies of 208 the ferrous iron transporter FeoB which is common to most bacteria. Strains lacking feoB1 209 (50) and feoB2 (51) were found to have lower magnetite contents than the wild type. 210 Deletion of the iron response regulator, Fur, which activates feoB1 and feoB2 also resulted 211 in reduced magnetosome formation (52). All the above studies compare magnetosome 212 production of wild type and 'deficient' strains, but do not provide dynamic measurements of 213 iron transport in MSR-1. Moreover, it is well known that biomineralization depends not only 214 on iron, but also on  $O_2$  availability (26,53).

215 Typically, magnetosome production is quantified off-line (by measuring the iron content in 216 cells using atomic absorption spectroscopy), and is backed up by visualization of 217 magnetosomes under the transmission electron microscope. In both cases sample 218 preparation and analysis are laborious and time consuming. Therefore there is clearly need 219 of rapid new methods to interrogate and quantify magnetosome production and 220 biomineralization in MTBs, as well as inform the development and optimization of large-scale 221 magnetosome production strategies in bioreactors. It is this context that we developed a 222 FCM based method for detecting chelatable iron in *M. gryphiswaldense* MSR-1 cells using 223 phen green M SK (PG-SK), whose fluorescence is guenched by metal ions including  $Fe^{2+}$ 224 and  $Fe^{3+}$ . PG-SK has been previously used to study iron transporters (54) and efflux 225 systems in *E. coli* as well as applied to studies in human cell lines for iron, copper and silver 226 uptake (55,56).

227 Non-magnetic MSR-1 cells grown aerobically in FSM-Fe<sup>-</sup> (without iron) were stained with 228 PG-SK. The staining procedure was partially optimized with respect to staining time (600 – 229 900 s) and PG-SK concentration (0.2 – 10  $\mu$ M) at three different temperatures (22, 27 and 230 30 °C). The highest fluorescence was observed at a PG-SK concentration of 5 µM after 10 231 minutes of incubation (Fig. 4a). Doubling the PG-SK concentration did not enhance the 232 fluorescence intensity of stained MSR-1 cells further (Fig. 4a), and longer staining times 233 were not needed (Fig 4b). Peak fluorescence intensity was similar at all staining 234 temperatures employed (Fig. 4c).

#### 235 Physiological changes of cells cultured with limited O<sub>2</sub> availability.

The effect of  $O_2$  limitation on growth and cellular magnetism of MSR-1 cells was investigated indirectly by varying the volume of headspace provided (i.e. 20 %, 40 %, 60 %, and 80 %) in tightly sealed 50 mL Falcon tubes. In all experiments the initial  $OD_{565}$  was 0.086 ± 0.004. After 48 h in culture,  $OD_{565}$  and  $C_{mag}$  values were recorded (Fig. 5a). Two clear and opposite trends were observed; whereas biomass production paralleled the increase in headspace 241 volume, and therefore O<sub>2</sub> availability, conversely, the magnetism of *M. gryphiswaldense* 242 MSR-1 cells dropped dramatically from strongly magnetic (C<sub>mag</sub> = 2) at 20 % (v/v) headspace 243 to very weakly magnetic  $C_{mag}$  = 1.1 at 60 % (v/v) headspace. These results are in agreement 244 with findings from previous studies (26). Fig. 5b shows corresponding FCM analyses for 245 relative quantification of intracellular iron and PHA content as a function of headspace 246 volume. The highest intracellular PHA accumulation was observed in cells cultured in tubes 247 with the lowest O<sub>2</sub> availability (i.e. lowest headspace volume of 20 %). Increased PHA formation during O<sub>2</sub> limitation has previously been reported (reviewed by 41). Conversely, 248 249 cells cultured at high O<sub>2</sub> concentrations (80% headspace volume) had the lowest PG-SK 250 fluorescence among the tested conditions and thus the highest free iron concentration. 251 Microarray data showed that iron transporter feoB1 is upregulated aerobically (29), 252 suggesting that iron transport into cells is highest aerobically.

Staining cells grown with different headspace volumes with PI and BOX (Fig. 5c) revealed that overall cell health was highest with 20 % headspace and lowest with 80% headspace volume. PI was also used for analysis of physiology of MSR-1 cells from ACA agar plates; this indicated that 15 - 20 % of cells on plates are dead (PI<sup>+</sup>). This highlights the difficulty in transferring MSR-1 cultures from single colonies to liquid cultures and emphasises the need to use a large amount of cells for setting up liquid cultures.

# 259 Physiological characterization of MSR-1 in shake flask experiments with free air260 exchange.

261 To gain new insight into *M. gryphiswaldense* MSR-1 physiology during the shift from O<sub>2</sub>-262 limited to aerobic conditions, we transferred cells grown under O<sub>2</sub>-limited conditions to O<sub>2</sub>-263 rich conditions with or without the supplementation of iron. Magnetic cells grown in FSM 264 batch medium and using a pH-stat feeding strategy in an O<sub>2</sub>-limited bioreactor were 265 aseptically transferred to non-baffled shake flasks containing fresh media, either FSM or 266 FSM without Fe (FSM-Fe), and grown in  $O_2$  rich conditions (free air exchange) at 30 °C on 267 an orbital shaker (150 rpm).  $OD_{565}$  and  $C_{mag}$  were monitored immediately before (t = 0 h) and 268 24 or 48 h after transfer and intracellular free iron and cellular PHA content were measured 269 and compared to the pre-transfer culture using FCM (Fig. 6). After O<sub>2</sub>-limited growth in the 270 bioreactor, and at the point of transfer to shake flasks (t = 0 h), MSR-1 cells were moderately 271 magnetic ( $C_{mag}$  = 1.71). Strong Pyr-546 fluorescence (Fig. 6b, 0 h) and electron and 272 fluorescence microscopy (Supplemental Fig. S2) confirmed that cells contained large 273 quantities of PHA. After transfer to aerobic conditions, cells grew better in the presence of 274 iron (FSM) compared to the absence (FSM-Fe<sup>-</sup>). After 24 h, C<sub>mag</sub> rose slightly from 1.71 to 275 1.84 (although variation was high at 24 h), but then dropped to 1.46 at 48 h. Cultures grown

in FSM had > 3 fold decreased Pyr-546 fluorescence at 24 h and 48 h, indicating a decrease
in PHA content, suggesting that growth utilised PHA as a substrate; FSC and SSC values
also dropped (Fig. 6c), indicative of decreasing cell size and potentially corroborating loss of
PHA granules. The impact of PHA utilisation on cell morphology has previously been
reported in *C. necator* (57) and *Pseudomonas oleovorans* (58). In addition, in a parallel
experiment, TEM analysis and fluorescence microscopy with Pyr-546 stained cells allow
comparison of PHA detection methods (Supplemental Fig. S3).

283 After transfer of magnetic cells to culture medium lacking iron citrate (FSM-Fe<sup>-</sup>), very little

growth ensued ( $OD_{565} \sim 0.3$ ; Fig. 6a).  $C_{mag}$  fell steadily (reaching 1.2 after 48 h; Fig. 6a). Pyr-546 fluorescence dropped to 65 % of its pre-transfer value (Fig. 6b), reflecting low PHA utilisation, whereas FSC and SSC increased, indicating an increase in cell size and granularity (Fig. 6c).

PG-SK fluorescence dropped over time for cultures grown in FSM but not in the absence of iron; as expected, this reflects an increase in chelatable iron concentration in cells grown in the presence of iron, but not in the absence. Headspace volume experiments revealed an increase in intracellular iron concentration under more aerobic conditions in the presence of iron (Fig. 4). As with the regulation of intracellular iron concentrations in response to  $O_2$ , high extracellular iron concentrations were shown to increase the expression of the *feo* iron transporters (59).

In summary, FCM analysis of viability, intracellular chelatable iron and PHA, employing PI / BOX, PG-SK and Pyr-546 dyes respectively provides valuable insight on the effects of  $O_2$ and iron levels on the growth, magnetosome and PHA production of MTBs. The data are rapidly obtained, does not require growth of MTBs on agar plates, and when used together with similarly fast measurements of optical density and C<sub>mag</sub> can be useful in the design of growth strategies for production of magnetosome rich cells.

301

#### 302 METHODS

#### 303 Strains, growth media and culture conditions.

304 *Magnetospirillum gryphiswaldense* MSR-1 was obtained from Leibniz-DSMZ (Deutsche 305 Sammlung van Mikroorganismen und Zellkulturen GmbH) and used in all experiments. 306 Unless otherwise indicated all chemicals were from Sigma-Aldrich Chemical Company Ltd 307 (Gillingham, Dorset, UK). *M. gryphiswaldense* MSR-1 cells were grown on solid activated

charcoal agar (ACA) plates and in liquid media. ACA plates contained 3 g L<sup>-1</sup> activated 308 charcoal. 15 g L<sup>-1</sup> agar (Formedium, Hunstanton, Norfolk, UK), 0.1 g L<sup>-1</sup> yeast extract, 3 g L<sup>-1</sup> 309 310 soybean peptone, 3 g L<sup>-1</sup> sodium pyruvate, 0.34 g L<sup>-1</sup> NaNO<sub>3</sub>, 0.1 g L<sup>-1</sup> KH<sub>2</sub>PO<sub>4</sub>, 0.15 g L<sup>-1</sup> 311 MgSO<sub>4</sub>·7H<sub>2</sub>O and 2.38 g L<sup>-1</sup> 4-(2-Hydroxyethyl)piperazine-1-ethanesulfonic acid (HEPES) 312 buffer in MiliQ water. The pH was adjusted to 7.0 with sodium hydroxide (Heyen and 313 Schüler, 2003) before autoclaving. After autoclaving iron(III) citrate (final concentration of 314 500 µM) and 1.4-dithiothreitol (DTT; final concentration of 1 mM) were aseptically added to 315 the plate mix before pouring (38). Set ACA plates were incubated at 30°C in 12-plates 316 anaerobic jars with one Anaerocult<sup>®</sup>C sachet (Merck Chemicals Ltd, Beeston Notts, UK) to 317 achieve microaerobic conditions. Liquid cultures of M. gryphiswaldense MSR-1 were 318 routinely grown in a shaking incubator (30° C, 150 rpm) using a flask standard medium 319 (FSM) composed of 0.1 g L<sup>-1</sup> yeast extract, 3 g L<sup>-1</sup> soybean peptone, 3.5 g L<sup>-1</sup> potassium Llactate, 100  $\mu$ M iron(III) citrate, 0.34 g L<sup>-1</sup> NaNO<sub>3</sub>, 0.1 g L<sup>-1</sup> KH<sub>2</sub>PO<sub>4</sub>, 0.15 g L<sup>-1</sup> MgSO<sub>4</sub>·7H<sub>2</sub>O, 320 321 5 mL L<sup>-1</sup> of EDTA-chelated trace element mixture (60) and 2.38 g L<sup>-1</sup> HEPES buffer in 322 deionized water; the whole adjusted to pH 7.0 prior to sterilization in an autoclave. Cells 323 were grown at 30°C in a shaking incubator at 150 rpm. O<sub>2</sub>-limiting cultures were grown in 324 tightly sealed screw cap 50 mL Falcon tubes with variable headspace volumes (10 - 40 mL), 325 whereas aerobic cultivations were performed with 50 mL of media in 250 mL shake flasks 326 allowing free air exchange. Non-magnetic cells were cultured in FSM without iron (FSM-Fe<sup>-</sup>) 327 for a minimum of three sequential sub-cultures in an attempt to eliminate all trace of the 328 metal. Magnetic cells were obtained from cultures grown in bioreactor experiments under 329 controlled conditions using a growth strategy adapted from previous works (26, 28, 61).

#### 330 Flow cytometry (FCM).

331 Bacterial samples taken directly from agar plates or liquid cultures were resuspended in 332 phosphate-buffered saline (PBS) and then analysed directly in a BD Accuri C6 flow 333 cytometer (Becton, Dickinson and Company, Oxford, UK) for cell size and granularity, or 334 after staining with various fluorescent dyes (see Supplemental Table 1) using protocols 335 developed and detailed in the Results and Discussion. During FCM on fluorescently labelled 336 cells, samples were excited using a 488 nm solid-state laser and fluorescence was detected 337 using three different filters, i.e.: a 533/30 BP filter (FL1-A) for bis (1.3-dibutybarbituric acid) 338 trimethine oxonol (referred to here as bis-oxonol or BOX), pyrromethene-546 (Pyr-546) and 339 phen green<sup>™</sup> SK (PG-SK); a 670 LP filter (FL3-A) for propidium iodide (PI); and a 675/25 340 BP filter (FL4-A) for Svto<sup>®</sup>62.

#### 341 Analytical methods.

Culture optical densities were recorded at a wavelength of 565 nm (OD<sub>565</sub>) in an Evolution
 300 UV-Vis spectrophotometer (Thermo Fisher Scientific, Hemel Hempstead, Herts, UK)
 controlled by Thermo Scientific<sup>™</sup> VISION*pro*<sup>™</sup> software.

345 Cellular magnetic response (C<sub>mag</sub>) values of cultures were determined immediately after 346 OD<sub>565</sub> measurements using a purpose-built magnetic measurement system mounted within 347 the spectrophotometer. In common with previous magneto-spectrophotometry apparatus 348 (24, 25) our system features two pairs of Helmholtz coils arranged around the cuvette 349 holder, one pair perpendicular to the light beam and the other in line with it. Each pair of coils 350 is energized in turn, and the OD<sub>565</sub> is measured in each condition. Magnetic cells will align 351 with the magnetic field and thus orient in line with or perpendicular to the light beam; the 352 optical density will therefore change between the two conditions. Non-magnetic cells do not 353 align with the magnetic field, thus the optical density does not change on switching the field 354 orientation. C<sub>mag</sub> values for culture samples are calculated by dividing the OD<sub>565</sub> values of 355 suspensions of cells aligned parallel to the light beam by those obtained when the same 356 cells are perpendicularly aligned. C<sub>mag</sub> values greater than unity reflect the presence of 357 magnetic cells.

Cellular morphology was routinely examined by light microscopy using an Olympus BX50 optical microscope (Olympus Corporation, Tokyo, Japan). Images were captured using a MotiCam 1 (800×600 pixel) camera (Microscope Systems Limited, Glasgow, UK) and processed with Motic Images Plus 2.0 software (Motic Europe S.L.U., Barcelona, Spain).

Cells stained with fluorescent probes were observed and imaged using a Zeiss Axiolab microscope (Carl Zeiss Ltd., Cambridge, UK) fitted with a Zeiss AxioCam ICm1 camera, and the images were processed in auto-exposure mode with the aid of Zeiss ZEN Lite 2012 software. Samples were excited with a Zeiss VHW 50f-2b ultraviolet light source and a 520 LP filter was employed for detection of fluorescence from Syto<sup>®</sup> 9 and pyrromethene-546 (Pyr-546).

#### 368 **Data availability.**

The datasets generated during and analysed during the current study are available from the corresponding author on reasonable request.

371 **REFERENCES** 

- 1. Frey, N. A., Peng, S., Cheng, K. & Sun, S. Magnetic nanoparticles: synthesis,
- functionalization, and applications in bioimaging and magnetic energy storage. *Chem. Soc. Rev.* 38, 2532–2542 (2009).

Schätz, A., Reiser, O. & Stark, W. J. Nanoparticles as semi-heterogeneous catalyst
 supports. *Chem. Eur. J.* 16, 8950–8967 (2010).

- 377 3. Franzreb, M., Siemann-Herzberg, M., Hobley, T.J. & Thomas, O. R. T. Protein
- purification using magnetic adsorbent particles. *Appl. Microb. Biotechnol.* **70**, 505–516
  (2006).
- 380 4. Fischer, I., Hsu, C-C., Gärtner, M., Müller, C., Overton T. W., et al. Continuous
- 381 protein purification using functionalized magnetic nanoparticles in aqueous micellar two-
- 382 phase systems. J. Chromatogr. A. **1305**, 7–16 (2013).
- 383 5. Pankhurst, Q. A., Thanh, N. T. K., Jones, S. K. & Dobson, J. Progress in applications
  384 of magnetic nanoparticles in biomedicine. *J. Phys. D: Appl. Phys.* 42, 224001 (2009).
- Hao, R., Xing, R., Xu, Z., Hou, Y., Gao, S. & Sun, S. Synthesis, functionalization, and
  biomedical applications of multifunctional magnetic nanoparticles. *Adv. Mater.* 22, 2729–
  2742 (2010).
- Wilhelm, S., Tavares, A. J., Dai, Q., Ohta, S., Audet, J. *et al.* Analysis of nanoparticle
  delivery to tumours. *Nat. Rev. Mat.* 1, 16014 (2016).
- 390 8. Lang, C. & Schüler, D. Biogenic nanoparticles: production, characterization, and
- 391 application of bacterial magnetosome. J. Phys. Condens. Matter 18, S2815–S2828 (2006).
- 392 9. Arakaki, A., Nakazawa, H., Nemoto, M., Mori, T. & Matsunaga, T. Formation of
- 393 magnetite by bacteria and its application. J. R. Soc. Interface 5:977–999 (2008).
- 10. Yan, L., Zhang, S., Chen, P., Liu, H., Yin, H., Li, H. Magnetotactic bacteria,
- 395 magnetosomes and their application. *Microb. Res.* **167**, 507–519 (2012).
- Alphandéry, E. Applications of magnetosomes synthesized by magnetotactic bacteria
   in medicine. *Front. Bioeng. Biotechnol.* 2, 5 (2014).
- Jacob, J. J. & Suthindhiran, K. Magnetotactic bacteria and magnetosomes Scope
  and challenges. *Mater. Sci. Eng. C.* 68, 919–928 (2016).
- 400 13. Bazylinski, D.A. & Frankel, R.B. Magnetosome formation in prokaryotes. *Nat. Rev.*401 *Microbiol.* 2, 217–230 (2004).
- 402 14. Pollithy, A., Romer, T., Lang, C., Müller, F. D., Helma, J. et al. Magnetosome
- 403 expression of functional camelid antibody fragments (nanobodies) in Magnetospirillum
- 404 gryphiswaldense. Appl. Environ. Microbiol. 77, 6165–6171 (2011).
- 405 15. Póstfai, M., Lefèvre, C. T., Truitsyn, D., Bazylinski, D. A. & Frankel, R.B.
- 406 Phylogenetic significance of composition and crystal morphology of magnetosome minerals.
- 407 Front. Microbiol. 4, 344 (2013).

- 408 16. Uebe, R. & Schüler, D. Magnetosome biogenesis in magnetotactic bacteria. *Nat.*
- 409 Rev. Microbiol. 14, 621–637 (2016).
- 410 17. Barber-Zucker, S. & Zarivach, R. A look into the biochemistry of magnetosome
- 411 biosynthesis in magnetotactic bacteria. ACS Chem. Biol. **12**, 3–22 (2017).
- 412 18. Frankel, R.B. Magnetic guidance of organisms. *Annu. Rev. Biophys. Bioeng.* 13:85–
  413 103 (1984).
- 414 19. Wacker, R., Ceyhan, B., Alhorn, P., Schueler, D., Lang, C. & Niemeyer, C. M.
- 415 Magneto immuno-PCR: a novel immunoassay based on biogenic magnetosome
- 416 nanoparticles. *Biochem. Biophys. Res. Commun.* **357**, 391–396 (2007).
- 417 20. Yoshino, T., Hirabe, H., Takahashi, M., Kuhara, M., Takeyama, H. & Matsunaga, T.
- 418 Magnetic cell separation using nano-sized bacterial magnetic particles with reconstructed
   419 magnetosome membrane. *Biotechnol. Bioeng.* **101**, 470–477 (2008).
- 420 21. Boucher, M., Geffroy, F., Prévéral, S., Bellanger, L., Selingue, E. *et al.* Genetically
- 421 tailored magnetosomes used as MRI probe for molecular imaging of brain tumor.
- 422 Biomaterials **121**, 167–178 (2017).
- 423 22. Tang, Y.S., Wang, D., Zhou, C., Ma, W., Zhang, Y.Q. *et al.* Bacterial magnetic
- 424 particles as a novel and efficient gene vaccine delivery system. *Gene ther.* 19(12):1187–
  425 1195 (2012).
- 426 23. Felfoul, O., Mohammadi, M., Taherkhani, S., De Lanauze, D., Xu, Y. Z. *et al.*
- 427 Magneto-aerotactic bacteria deliver drug-containing nanoliposomes to tumour hypoxic
- 428 regions. Nat. Nanotechnol. **11**, 941–947 (2016).
- 429 24. Zhao, L., Wu, D., Wu, L. F. & Song, T. A simple and accurate method for
- 430 quantification of magnetosomes in magnetotactic bacteria by common spectrophotometer. J.
- 431 Biochem. Biophys. Methods **70**, 377–383 (2007).
- 432 25. Lefèvre, C. T., Song, T., Yonnet, J-P. & Wu, L-F. Characterization of bacterial
- 433 magnetotactic behaviors by using a magnetospectrophotometry assay. *Appl. Environ.*
- 434 *Microbiol.* **75**:3835–3841 (2009).
- 435 26. Heyen, U. & Schüler, D. Growth and magnetosome formation by microaerophilic
- 436 Magnetospirillum strains in an oxygen-controlled fermentor. Appl. Microbiol. Biotechnol. 61,
  437 536–544 (2003).
- 438 27. Grünberg, K., Müller, E. C., Otto, A., Reszka, R., Linder, D. *et al.* Biochemical and
- 439 Proteomic Analysis of the Magnetosome Membrane in *Magnetospirillum gryphiswaldense*.
- 440 Appl. Environ. Microbiol. **70**, 1040–1050 (2004).
- 441 28. Yang, J., Li, S., Huang, X., Tang, T., Jiang, W., Zhang, T. & Li, Y. A key time point for
- 442 cell growth and magnetosome synthesis of *Magnetospirillum gryphiswaldense* based on
- real-time analysis of physiological factors. *Front. Microbiol.* **4**, 1–7 (2013).

- 444 29. Wang, X., Wang, Q., Zhang, Y., Wang, Y., Zhou, Y., et al. Transcriptome analysis 445 reveals physiological characteristics required for magnetosome formation in 446 Magnetospirillum gryphiswaldense MSR-1. Environ. Microbiol. Rep. 8, 371–381 (2016). 447 30. Geng, J., Beloin, C., Ghigo, J-M., Henry, N., Stewart, P., et al. Bacteria hold their 448 breath upon surface contact as shown in a strain of Escherichia coli, using dispersed 449 surfaces and flow cytometry analysis. PLoS One 9,e102049 (2014). 450 31. Lagendijk, E.L., Validov, S., Lamers, G. E. M., De Weert, S. & Bloemberg, G. V. 451 Genetic tools for tagging Gram-negative bacteria with mCherry for visualization in vitro and 452 in natural habitats, biofilm and pathogenicity studies. FEMS Microbiol. Lett. 305, 81–90 453 (2010). 454 32. Sevastsyanovich, Y., Alfasi, S., Overton, T., Hall. R., Jones., J. et al. Exploitation of 455 GFP fusion proteins and stress avoidance as a generic strategy for the production of high-456 quality recombinant proteins. FEMS Microbiol. Lett. 299, 86-94 (2009). 457 33. Khan, M. M. T., Pyle, B. H. & Camper, A.K. Specific and rapid enumeration of viable 458 but nonculturable and viable-culturable Gram-negative bacteria by using flow cytometry. 459 Appl. Environ. Microbiol. 76, 5088–5096 (2010). 460 34. Lang, C. & Schuler, D. Expression of green fluorescent protein fused to 461 magnetosome proteins in microaerophilic magnetotactic bacteria. Appl. Environ. Microbiol. 462 74, 4944–4953 (2008). 463 35. Lang, C., Pollithy, A. & Schüler, D. Identification of promoters for efficient gene 464 expression in Magnetospirillum gryphiswaldense. Appl. Environ. Microbiol. 75, 4206–4210 465 (2009). 466 36. Day, J.B., Kell, D.B. & Griffith, G.W. Differentiation of Phytophthora infestans 467 sporangia from other airborne biological particles by flow cytometry. Appl. Environ. Microbiol. 468 **68,** 37–45 (2002). 469 37. Lee, Y-H., Chen, S-Y., Wiesner, R.J., Huang, Y-F. Simple flow cytometric method 470 used to assess lipid accumulation in fat cells. J. Lipid Res. 45, 1162-7 (2004). 471 38. Schultheiss, D. & Schüler, D. Development of a genetic system for Magnetospirillum 472 gryphiswaldense. Arch. Microbiol. 179, 89–94 (2003). 473 39. Nebe-von-Caron, G., Stephens, P., Hewitt, C., Powell, J. & Badley, R. Analysis of 474 bacterial function by multi-colour fluorescence flow cytometry and single cell sorting. J. 475 Microbiol. Methods 42, 97-114 (2000). 476 40. Ward, A.C., Rowley, B.I. & Dawes, E.A. Effect of Oxygen and Nitrogen Limitation on 477 Poly-hydroxybutyrate Biosynthesis in Ammonium-grown Azotobacter beijerinckii. J. Gen. 478 Microbiol. 102, 61-68 (1977).
- 479 41. Kessler, B. & Witholt, B. Factors involved in the regulatory network of
- 480 polyhydroxyalkanoate metabolism. J. Biotechnol. 86, 97–104 (2001).

- 481 42. García-Torreiro, M., Lu-Chau, T. A. & Lema, J. M. Effect of nitrogen and/or oxygen
- 482 concentration on poly(3-hydroxybutyrate) accumulation by Halomonas boliviensis.
- 483 Bioprocess Biosyst. Eng. **39**, 1365–1374 (2016).
- 484 43. Ban, J., Jiang, W., Li, Y., Zhang, Y. P. & Li, J. L. Functional analysis of hydrogenases
- 485 and their effects on cell growth and magnetosome synthesis in *Magnetospirillum*
- 486 gryphiswaldense. Chinese Sci. Bull. 55:1271–1277 (2010).
- 487 44. Schultheiss, D., Handrick, R., Jendrossek, D., Hanzlik, M. & Schüler, D. The
- 488 presumptive magnetosome protein Mms16 is a poly(3-hydroxybutyrate) granule-bound
- 489 protein (phasin) in *Magnetospirillum gryphiswaldense*. J. Bacteriol. **187**, 2416–2425 (2005).
- 490 45. Liu, J., Ding, Y., Jiang, W., Tian, J., Li, Y. & Li, J. A mutation upstream of an ATPase
- 491 gene significantly increases magnetosome production in *Magnetospirillum gryphiswaldense*.
  492 *Appl. Microbiol. Biotechnol.* 81, 551–558 (2008).
- 493 46. Raschdorf, O., Plitzko, J. M., Schuler, D. & Muller, F. D. A tailored *galK*
- 494 counterselection system for efficient markerless gene deletion and chromosomal tagging in
- 495 Magnetospirillum gryphiswaldense. Appl. Environ. Microbiol. 80, 4323–4330 (2014).
- 496 47. Vizcaino-Caston, I., Kelly, C. A., Fitzgerald, A. V. L., Leeke, G. A., Jenkins, M.,
- 497 Overton, T. W. Development of a rapid method to isolate polyhydroxyalkanoates from
- 498 bacteria for screening studies J. Biosci. Bioeng. **121**,101–104 (2016).
- 499 48. Kacmar, J., Carlson, R., Balogh, S.J., Srienc, F. Staining and quantification of poly-3-
- 500 hydroxybutyrate in Saccharomyces cerevisiae and Cupriavidus necator cell populations
- using automated flow cytometry. *Cytometry A.* **69**, 27-35 (2006).
- 49. Karmann, S.; Panke, S.; Zinn, M. The Bistable Behaviour of *Pseudomonas putida*
- 503 KT2440 during PHA Depolymerization under Carbon Limitation. *Bioengineering.* **4**, 58 (2017).
- 505 50. Rong, C., Huang, Y., Zhang, W., Jiang, W., Li, Y. & Li, J. Ferrous iron transport
- 506 protein B gene (feoB1) plays an accessory role in magnetosome formation in
- 507 Magnetospirillum gryphiswaldense strain MSR-1. Res. Microbiol. 159, 530–536 (2008).
- 508 51. Rong, C., Zhang, C., Zhang, Y., Qi, L., Yang, J., et al. FeoB2 Functions in
- 509 magnetosome formation and oxidative stress protection in *Magnetospirillum*
- 510 gryphiswaldense strain MSR-1. J. Bacteriol. **194**, 3972–3976 (2012).
- 511 52. Uebe, R., Voigt, B., Schweder, T., Albrecht, D., Katzmann, E., et al. Deletion of a fur-
- 512 like gene affects iron homeostasis and magnetosome formation in *Magnetospirillum*
- 513 gryphiswaldense. J. Bacteriol. **192**, 4192–4204 (2010).
- 514 53. Schüler, D. & Baeuerlein, E. Dynamics of iron uptake and Fe<sub>3</sub>O<sub>4</sub> biomineralization
- 515 during aerobic and microaerobic growth of Magnetospirillum gryphiswaldense. J. Bacteriol.
- 516 **180**, 159–162 (1998).

517	54.	Große, C., Scherer, J., Koch, D., Otto, M., Taudte, N. & Grass, G. A new ferrous iron-
518	uptake transporter, EfeU (YcdN), from Escherichia coli. Mol. Microbiol. 62, 120–131 (2006).	
519	55.	Xu, G., Ahn, J., Chang, S., Eguchi, M., Ogier, A., et al. Lipocalin-2 induces
520	cardiomyocyte apoptosis by increasing intracellular iron accumulation. J. Biol. Chem. 287,	
521	4808–4817 (2012).	
522	56.	Zhao, G., Di, D., Wang, B., Zhang, P., Xu, Y. Iron regulates the expression of
523	ferropo	ortin 1 in the cultured hFOB 1.19 osteoblast cell line. Exp. Ther. Med. 8, 826–830
524	(2014).	
525	57.	Tian, J., Sinskey, A. J. & Stubbe, J. Kinetic studies of polyhydroxybutyrate granule
526	formation in Wautersia eutropha H16 by transmission electron microscopy. J. Bacteriol. 187,	
527	3814–3824 (2005).	
528	58.	Ruiz, J. A., López, N. I., Fernández, R. O. & Méndez, B. S. Polyhydroxyalkanoate
529	degradation is associated with nucleotide accumulation and enhances stress resistance and	
530	survival of Pseudomonas oleovorans in natural water microcosms. Appl. Environ. Microbiol.	
531	<b>67</b> , 225–230 (2001).	
532	59.	Wang, Q., Liu, J. X., Zhang, W. J., Zhang, T. W., Yang, J. & Li, Y. Expression
533	patterns of key iron and oxygen metabolism genes during magnetosome formation in	
534	Magnetospirillum gryphiswaldense MSR-1. FEMS Microbiol. Lett. 347,163-72 (2013).	
535	60.	Widdel, F. & Bak, F. Gram-negative mesophilic sulfate-reducing bacteria in The
536	Prokaryotes (eds. Balows, A., Troper, H., Dworkin, M., Harder, W. & Schleifer, K.) 3352-	
537	3378 (Springer, 1992)	
538	61.	Sun, J-B., Zhao, F., Tang, T., Jiang, W., Tian, J., <i>et al</i> . High-yield growth and
539	magne	tosome formation by Magnetospirillum gryphiswaldense MSR-1 in an oxygen-
540	controlled fermentor supplied solely with air. Appl. Microbiol. Biotechnol. 79, 389–397	
541	(2008)	
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### 546 Author Contributions

- 547 AFC and HL completed the practical work. ORTT and TWO supervised the work. AFC,
- 548 ORTT and TWO wrote the manuscript. All authors reviewed the manuscript.
- 549 Additional Information

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- 551

#### 552 Figure legends

- 553 Figure 1. Analysis of *M. gryphiswaldense* MSR-1 using FSM and light microscopy.
- 554 Scatter plots (Forward scatter, FSC-A vs. Side scatter, SSC-A) of cells cultured (a) in liquid
- 555 FSM and (b) on ACA plates; (c) comparison of individual particle count vs FSC-A plots for
- liquid (red trace) and plate (blue trace) grown cells; light microscope images of (d) liquid and(e) plate grown cells.
- **Figure 2. Viability analysis of MSR-1 cells using FCM.** MSR-1 cells were co-stained with BOX (fluorescence measured on FL1-A channel, y axis) and PI (fluorescence measured on FL3-A channel, x axis). Key: (a) actively growing magnetic cells; (b) starving magnetic cells; (c) actively growing non-magnetic cells; (d) starving non-magnetic cells; and (e) cells incubated with absolute ethanol for 10 minutes, centrifuged and then resuspended in phosphate buffered saline. The numbers of cells in each of the four quadrants of all plots are indicated in red font and are expressed as percentages of the total population.
- **Figure 3. Analysis of PHA content using FCM.** Fluorescence intensity histograms of starved (a) non-magnetic and (b) magnetic cells after staining with Pyr-546 ( $0.5 \ \mu g \cdot mL^{-1}$ ) for various times. The numbers '1' & '2' marked on the inserts and the fluorescence micrographs correspond to non-magnetic and magnetic cells, respectively to identify those with low and high PHA content. The scale bars indicate a length of 5  $\mu m$ .
- 570 **Figure 4. Analysis of intracellular iron by FCM.** Fluorescence intensity histograms of non-571 magnetic MSR-1 cells growth in FSM-Fe<sup>-</sup> after staining: (a) with various concentrations of 572 PG-SK for 600 s at 30°C; (b) for various times with 10  $\mu$ M PG-SK at 30°C; and (c) at various 573 temperatures using 10  $\mu$ M PG-SK for 600 s.
- Figure 5. Effect of  $O_2$  limitation on physiology. MSR-1 cultures were grown in tubes with different headspace volumes for 48 h. (a)  $OD_{565}$  (pale grey bars) and cellular magnetism ( $C_{mag}$ ; black bars). Error bars are standard deviation. (b) Fluorescence of cells stained with PG-SK (dark grey bars) and Pyr-546 (white bars) as measured using FCM. Error bars are covariance. (c) Viability as determined using FCM and staining with PI and BOX; percentage of healthy (black bars), injured (pale grey bars) and dead (dark grey bars) cells are shown. Experiments were performed in triplicate.

581 Figure 6. Physiology of magnetic cells during shift to aerobic conditions. MSR-1 cells 582 grown under  $O_2$ -limited conditions in a bioreactor were transferred to  $O_2$ -sufficient conditions 583 with either iron-containing (FSM) or iron-lacking (FSM-Fe<sup>-</sup>) media. (a) OD<sub>565</sub> (pale grey bars) 584 and C<sub>mag</sub> value (black bars). Error bars are standard deviation; cells were taken from a single 585 bioreactor into three replicate flasks for each condition. (b) Mean fluorescence intensity of 586 cells stained with 0.5 µg·mL<sup>-1</sup> pyrromethene-546 (Pyr546) (white bars) or 5 µM phen green<sup>™</sup> 587 SK (PG-SK) (dark grey bars). Error bars are covariance. (c) Forward scatter (FSC, white 588 bars) and side scatter (SSC, grey bars) of cells as determined by FCM. Error bars are 589 covariance. 25 000 events were analysed per sample by FCM. 590





FL3 (PI)-A

7.2









Headspace volume (%)





Time (h)

