

***Magnetospirillum magneticum* as a living iron chelator induces transferrin receptor 1 upregulation in cancer cells**

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Abstract

Interest has grown in harnessing biological agents for cancer treatment as dynamic and responsive vectors with enhanced tumor targeting. While bacterial traits such as proliferation in tumors, modulation of an immune response, and local secretion of toxins have been well studied, less is known about bacteria as competitors for nutrients. Here, we investigated the use of a bacterial strain as a living iron chelator, competing for this nutrient vital to tumor growth and progression. We established an *in vitro* co-culture system consisting of the magnetotactic strain *Magnetospirillum magneticum* AMB-1 incubated under hypoxic conditions with human melanoma cells. Siderophores produced by 10^8 AMB-1/mL in human transferrin (Tf)-supplemented media were found to alter the Tf structure with an effect equivalent to $3.78 \mu\text{M} \pm 0.117 \mu\text{M}$ deferoxamine, a potent drug used in iron chelation therapy. Our experiments revealed an increased expression of transferrin receptor 1 (TfR1), indicating the bacteria's ability to influence iron homeostasis in human melanoma cells. Our results show the potential of a bacterial strain as a self-replicating iron-chelating agent, which could serve as an additional mechanism reinforcing current bacterial cancer therapies.

52 1. Introduction

53

54 Due to limited selectivity in systemically delivered cancer therapeutics, interest has grown in
55 harnessing bacteria as living, tumor-targeting anticancer agents. The therapeutic potential of
56 facultative anaerobic bacteria has been supported by studies demonstrating the delivery of non-
57 pathogenic strains of *Escherichia coli* to solid flank tumors with associated tumor regression.
58 Additionally, safe administration of *Salmonella typhimurium* (VPN20009) has been shown for
59 animal models and patients with metastatic melanoma (Chowdhury et al., 2019, Clairmont et
60 al., 2000, Toso et al., 2002). Bacteria can act therapeutically by secreting innate or engineered
61 toxins in situ (e.g. hemolysin E), transporting attached nanodrug formulations, or stimulating
62 an immune response (Duong et al., 2019, Sedighi et al., 2019, Din et al., 2016, Harimoto et al.,
63 2019). Colonizing bacteria can also engage in nutrient competition within the tumor
64 microenvironment (Forbes, 2010, Sznol et al., 2000, Song et al., 2018). While the starvation of
65 glucose as a crucial energy source to all cells has been studied intensively (Grasmann et al.,
66 2019, Vander Heiden et al., 2009, Pavlova and Thompson, 2016), other nutrients that are in
67 specifically high demand by cancer cells might serve as more specific, vulnerable targets for
68 deprivation.

69

70 For example, iron metabolism is significantly altered in mammalian tumor cells and recognized
71 as a metabolic hallmark of cancer (Wang et al., 2018, Torti and Torti, 2013). The main iron
72 uptake mechanism adopted by most cells utilizes the internalization of transferrin receptor 1
73 (TfR1) upon binding of iron-bound transferrin (Tf). TfR1 expression positively correlates with
74 cellular iron starvation and is upregulated in cancer cells, since malignant cells generally
75 require a nutrient surplus (Torti and Torti, 2013, Lane et al., 2015, Steegmann-Olmedillas,
76 2011). Accordingly, several types of iron scavenging molecules have been utilized to compete
77 with malignant cells for available iron sources and have demonstrated significant anti-
78 neoplastic activity both, *in vitro* and *in vivo* (Bedford et al., 2013, Ford et al., 2013, Lui et al.,
79 2013) Promising bacteria-derived iron-chelating siderophores, such as deferoxamine (DFO),
80 as well as synthetic iron chelators have been developed for therapeutic purposes (Hatcher et
81 al., 2009). However, non-negligible side effects, including systemic toxicity and low efficacy,
82 have hampered their translation into clinical trials as therapeutic agents for cancer treatment
83 (Richardson, 2002, Yu et al., 2012b, Saha et al., 2019).

84

85 Here we investigate the potential of a specific bacterial strain with high demand for iron to
86 serve as local, self-replicating iron chelator that could thereby reduce systemic effects.
87 Magnetotactic bacteria, like other bacteria, possess the ability to secrete iron-scavenging
88 siderophores. Unlike other bacteria, their demand for iron is particularly high, since this
89 mineral is crucial both for their survival and synthesis of unique intracellular organelles called
90 magnetosomes. (Mirabello et al., 2016, Faivre and Schüler, 2008). These biomineralized
91 magnetic nanocrystals are arranged in chains enclosed in a lipid bilayer and enable the bacteria
92 to align with magnetic fields (Bazylinski and Williams, 1970, Yan et al., 2012, González et al.,
93 2015). Furthermore, MTB are aerotactic, possessing an oxygen-sensing system that regulates
94 motility in an oxygen gradient (Lefèvre et al., 2014). These features have previously been
95 leveraged to magnetically guide MTB to the hypoxic core of solid tumors, yielding
96 significantly higher tumor accumulation and penetration compared to their administration in
97 the absence of external magnetic fields (Felfoul et al., 2016). Once on site, nutrients from the
98 tumor microenvironment are sourced to maintain proliferation and growth, and we hypothesize
99 that MTB could induce iron deprivation of cancer cells.

100

101 To study this, we employed *Magnetospirillum magneticum* strain AMB-1 and first quantified
102 the production of siderophores, benchmarked with molar concentrations of DFO. We then
103 investigated the influence of AMB-1 on cell surface TfR1 expression using human melanoma
104 cells and demonstrated the ability of AMB-1 to affect iron homeostasis. The iron scavenging
105 capabilities of bacterial strains with naturally high or enhanced siderophore production may
106 act as an additional mechanism for bacterial cancer therapy, complementing or augmenting
107 established bacterial anticancer mechanisms.

108

109 **2. Materials and Methods**

110

111 **2.1. Bacterial strain and culture condition**

112

113 *Magnetospirillum magneticum* AMB-1, a strain of magnetotactic bacteria, was purchased from
114 ATCC (ATCC, Manassas, Virginia, USA). AMB-1 bacteria were grown anaerobically at 30°C,
115 passaged weekly and cultured in liquid growth medium (ATCC medium: 1653 Revised
116 Magnetic Spirillum Growth Medium). *Magnetospirillum magneticum* Growth Media (MSGM)
117 contained the following per liter: 5.0 mL Wolfe's mineral solution (ATCC, Manassas, Virginia,
118 USA), 0.45 mL Resazurin, 0.68 g of monopotassium phosphate, 0.12 g of sodium nitrate, 0.035
119 g of ascorbic acid, 0.37 g of tartaric acid, 0.37 g of succinic acid and 0.05 sodium acetate. The
120 pH of the media was adjusted to 6.75 with sodium hydroxide (NaOH) and then sterilized by
121 autoclaving at 121°C. 10 mM ferric quinate (200x) Wolfe's Vitamin Solution (100x) (ATCC,
122 Manassas, Virginia, USA) were added to the culture media shortly before use. The
123 concentration of AMB-1 in solution was determined by optical density measurement (Spark,
124 Tecan, Männedorf, Switzerland) and the approximate number of bacteria was extrapolated
125 from a standard curve.

126

127 **2.2. CAS assay to assess siderophore quantification**

128

129 *Magnetospirillum magneticum* AMB-1 were cultured in 1.7 mL phenol red-free DMEM
130 (11054020, Invitrogen, Carlsbad, California, USA) supplemented with GlutaMAX (35050061,
131 Invitrogen, Carlsbad, California, USA) in a sealed 1.5 mL Eppendorf tube at 37°C for 24 h.
132 FBS was excluded from the media and replaced with a known concentration of iron source; 25
133 µM holo-transferrin (T0665, Sigma-Aldrich, St. Louis, Missouri, USA). Quantification of
134 siderophores produced by AMB-1 was performed using the Chrome Azurol S (CAS) assay
135 (199532, Sigma-Aldrich, St. Louis, Missouri, USA) (Schwyn and Neilands, 1987). 100 µL of
136 each sample's supernatant was collected and mixed with 100 µL CAS assay solution on a
137 transparent 96-well plate. The assay was then incubated in the dark at room temperature for 1
138 h before the absorbance was measured at 630 nm on a multimode microplate reader (Spark,
139 Tecan, Männedorf, Switzerland). The measurement was expressed in siderophore production
140 unit (s.p.u.), which was calculated as follows:

$$141 \text{ Siderophore production unit (s. p. u.)} = \frac{OD_{630,ref} - OD_{630}}{OD_{630,ref}}$$

142 DMEM supplemented with different concentrations of deferoxamine mesylate salt (DFO,
143 D9533, Sigma-Aldrich, St. Louis, Missouri, USA) was prepared by serial dilution and used to
144 generate a calibration curve (Supplementary 1).

145

146 **2.3. Analysis of human transferrin using SDS-PAGE Electrophoresis**

147

148 AMB-1 bacteria (1×10^8 cells/mL) were cultured in 1.7 mL phenol red-free DMEM
149 (11054020, Invitrogen, Carlsbad, California, USA) in a sealed 1.5 mL Eppendorf tube at 30°C

150 for 48 h. Excess volume was used to ensure no or minimal air was trapped in the tubes. 25 μ M
151 holo-transferrin (holo-Tf, T4132, Sigma-Aldrich, St. Louis, Missouri, USA), or 25 μ M apo-
152 transferrin (apo-Tf, T2036, Sigma-Aldrich, St. Louis, Missouri, USA) respectively were added
153 to the mammalian cell culture media. Changes in transferrin molecular mass during the growth
154 of AMB-1 were evaluated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis
155 (SDS-PAGE) analysis of culture supernatant. Electrophoresis was conducted using the
156 protocol described by Laemmli (Laemmli, 1970) and protein loading of each sample was
157 normalized to 2 μ g. Proteins were visualized using SYPRO ruby protein stain (1703126, Bio-
158 rad, Hercules, California, USA). The electrophoresis chamber and the reagents were purchased
159 from Bio-rad. Stained gels were imaged using a fluorescent scanner (Sapphire Biomolecular
160 Imager, Azure Biosystems, Dublin, California, USA) at 488 nm excitation and 658 nm
161 emission.

162

163 **2.4. Mammalian cell culture**

164

165 Human melanoma MDA-MB-435S cells (ATCC, Manassas, Virginia, USA) were cultured in
166 high glucose Dulbecco's Modified Eagle's Medium (DMEM, Invitrogen, Carlsbad, California,
167 USA) supplemented with 10% fetal bovine serum (FBS, Biowest, Nuaille, France) and 1%
168 penicillin-streptomycin (CellGro, Corning, New York, USA). All cells were incubated at 37°C
169 in a humidified atmosphere with 5% CO₂ at 37°C.

170

171 **2.5. Co-culture of mammalian cancer cells with magnetotactic bacteria**

172

173 Human melanoma MDA-MB-435 cells (1×10^5 cells) were cultured on 12-well plates and
174 incubated in a 5% CO₂ incubator at 37°C for 24 h. For microscopic analysis at high
175 magnification (> 40x), a circular cover slip was placed in each well prior to cell seeding.
176 Following incubation, *Magnetospirillum magneticum* AMB-1 ($1 \times 10^6 - 1 \times 10^8$ cells) were
177 introduced into the wells. The well plate was stored in a sealable bag and the bag was flushed
178 with nitrogen for 15 min in order to produce hypoxic conditions. The setup with the 12-well
179 plate was then incubated at 37°C for 48 h. To serve as negative and positive controls, 0, 10 μ M
180 and 25 μ M of the iron-chelating agent deferoxamine mesylate (D9533, Sigma-Aldrich, St.
181 Louis, Missouri, USA) was added to the MDA-MB-435S cell culture in place of AMB-1
182 bacteria.

183

184 **2.6. Immunofluorescence labelling of MDA-MB-435S cells**

185

186 After the co-culture, cells were washed with ice cold 1X Dulbecco's Phosphate-Buffered
187 Saline solution (DPBS, Gibco, Carlsbad, California, USA) and then blocked with a 1% Bovine
188 Serum Albumin (BSA, Sigma-Aldrich, St. Louis, Missouri, USA) solution diluted in 1X
189 DPBS. The cells were then incubated with 10 μ g/mL primary anti-TfR1 antibody (ab84036,
190 Abcam, Cambridge, UK) on ice in dark for one h. Subsequently, the cells were washed with
191 ice-cold DPBS and incubated with 20 μ g/mL secondary goat anti-rabbit antibody (ab150077,
192 Abcam, Cambridge, UK) and 25 μ g/mL Hoechst 33342 (H3570, Thermo Fisher Scientific,
193 Waltham, Massachusetts, USA) on ice in dark for another hour. Next, the cells were washed
194 with ice-cold 1X PBS twice and fixed with a 2% paraformaldehyde (PFA) solution. Fixed cells
195 were washed three times with 1X DPBS and the cover slips were mounted on glass slides and
196 stored overnight in dark at 4°C. A Nikon Eclipse Ti2 microscope equipped with a Yokogawa
197 CSU-W1 Confocal Scanner Unit and Hamamatsu C13440-20CU ORCA Flash 4.0 V3 Digital
198 CMOS camera were used for visualization. Microscope operation and image acquisition was

199 performed using Nikon NIS-Elements Advanced Research 5.02 (Build 1266) software. ImageJ
200 v2.0 (NIH) was used to process the obtained images.

201

202 **2.7. Evaluation of fluorescently labelled MDA-MB-435S cells by flow cytometry**

203 Flow cytometry was used to measure the expression of fluorescently labelled TfR1 on the
204 surface of MDA-MB-435S cells. Cells were harvested at different timepoints during co-culture
205 (0h, 6h, 12h, 24h) and washed in cold 1X DPBS (Gibco Carlsbad, California, USA). Harvested
206 cells were stained with primary anti-TfR1 antibody (ab84036, Abcam, Cambridge, UK) at a
207 concentration of 10 $\mu\text{g}/\text{mL}$. After 1 h of incubation on ice, cells were washed twice with 1X
208 DPBS and then stained with 20 $\mu\text{g}/\text{mL}$ secondary goat anti-rabbit antibody (ab150077, Abcam,
209 Cambridge, UK). Finally, cells were washed twice with 1X DPBS and analyzed by flow
210 cytometry with BD LSRFortessa (BD Biosciences, San Jose, California, USA) using a 488nm
211 excitation laser and 530/30 and 690/50 band pass emission filters for detection. FlowJoTM (Tree
212 Star) software was used to evaluate the data.

213 Flow cytometry was used to assess the cell membrane integrity of MDA-MB-435S cells. Cells
214 were harvested at different timepoints during co-culture (0h, 6h, 24h) and washed in cold 1X
215 DPBS. Collected cells were stained with 1 $\mu\text{g}/\text{mL}$ Propidium Iodide (V13242, Thermo Fisher
216 Scientific, Waltham, Massachusetts, USA) and incubated for 30 in a humidified atmosphere
217 with 5% CO₂ at 37°C. Finally, cells were washed twice with 1X DPBS and analyzed by flow
218 cytometry with BD LSRFortessa (BD Biosciences, San Jose, California, USA) using a 488nm
219 excitation laser and 610/10 bandpass emission filters for detection. FlowJoTM (Tree Star)
220 software was used to evaluate data and graphs were plotted using Prism 8.0 (GraphPad).

221

222 **2.8. Statistics and data analysis**

223

224 All graphs and statistical analyses were generated using Prism 8.0 (GraphPad). Statistical
225 significance and number of replicates of the experiments are described in each figure and figure
226 legend. Error bars, where present, indicate the standard error of the mean (SD). P values are
227 categorized as * P<0.05, ** P<0.01, and *** P<0.001.

228

229

230 **3. Results**

231

232 **3.1. AMB-1 produced siderophores affect human transferrin structure in mammalian** 233 **cell culture medium**

234

235 First, we sought to determine to what extent AMB-1 would produce siderophores in Dulbecco's
236 Modified Eagle's medium (DMEM). Using the Chrome Azurol S (CAS) assay (Supplementary
237 Fig. 1), 10⁸ AMB-1 bacteria were found to produce 0.10 ± 0.005 siderophore units in DMEM
238 supplemented with 25 μM holo-transferrin (holo-Tf), while siderophore production in
239 transferrin-free DMEM was negligible (Fig. 1A). AMB-1 siderophore production was
240 compared to the widely used iron chelator deferoxamine. It was found that the siderophores
241 produced by 10⁸ AMB-1 in Tf-supplemented media was equivalent to $3.78 \mu\text{M} \pm 0.117 \mu\text{M}$
242 deferoxamine (Fig. 1B).

243

244 Having established the ability of AMB-1 to produce siderophores in Tf-supplemented media,
245 we next determined whether AMB-1 would have an effect on human transferrin structure. SDS-
246 PAGE analysis was used to compare DMEM supplemented with either iron-containing holo-
247 Tf or iron-depleted apo-Tf. The apo-Tf appeared as a broader band on the SDS-gel compared
248 to holo-Tf (Fig. 1C). Furthermore, we ascertained that holo-Tf structure was not affected by a

249 48 h incubation period at 30°C. To test whether the bacteria induced changes in Tf, AMB-1
250 were inoculated in DMEM supplemented with holo-Tf. This approach revealed that holo-Tf
251 formed a broader band very similar to that seen for the apo-Tf band incubated in DMEM (Fig.
252 1C, lane 6). These experiments demonstrate that AMB-1 produced a quantifiable amount of
253 siderophore when holo-Tf was supplemented to the mammalian cell culture media and that the
254 structure of holo-Tf was altered by the bacteria.

255

256 **3.2. AMB-1 upregulates TfR1 expression in human melanoma cells**

257

258 To test whether AMB-1 can affect the iron uptake machinery in mammalian tumor cells we
259 co-cultured AMB-1 with the human melanoma cell line MDA-MB-435S and monitored TfR1
260 expression using immunofluorescence. The surface expression of TfR1 increased 2.7-fold on
261 cancer cells co-cultured with live bacteria at AMB-1:MDA-MB-435S ratios as low as 10:1 (10⁶
262 AMB-1). The TfR1 upregulation was shown to increase with increasing bacteria ratios (Fig.
263 2A, B). Deferoxamine was used here to create iron-deficient cell culture conditions as a
264 positive control. MDA-MB-435S cells showed a significant and increasing upregulation of
265 TfR1 surface expression up to 5.6-fold. To ensure that the upregulation of TfR1 expression
266 was on the cell surface and not cytoplasmic, cell membrane integrity in the cultures was
267 monitored. Less than 5% of cells were stained by the cell-impermeant DNA stain propidium
268 iodide (PI), indicating cell membrane preservation over time (Fig. 2C).

269

270 To gain insights on the TfR1 expression kinetics of the cell population, AMB-1-induced
271 increase of cell surface TfR1 expression was analyzed over time. The effect, at an AMB-
272 1:MDA-MB-435S ratio of 1000:1 was already apparent after 6 h of co-culture (Fig. 1D). The
273 fluorescent intensity after 24 h of co-culture was 1.8 times higher than the initial value, while
274 the change reached 95% of the final value after 12 h (Fig. 1E). Untreated cancer cells did not
275 display any increase in fluorescence (Fig. 1F). Altogether, these findings show an upregulation
276 of TfR1 on the cell surface of human melanoma cancer cells in the presence of AMB-1, thereby
277 suggesting a direct link between AMB-1 induced disruption of iron uptake and TfR1
278 expression.

279

280

281 **4. Discussion**

282

283 Magnetotactic bacteria acquire iron through siderophore-mediated uptake, as ferric and ferrous
284 ions cannot directly enter bacteria cells. We quantified the number of siderophores produced
285 by strain *Magnetospirillum magneticum* AMB-1 and investigated the bacteria's effect on
286 human transferrin's structure in mammalian cell culture medium. The addition of AMB-1 to
287 the media led to a broader holo-Tf band, similar to that of apo-Tf in DMEM, suggesting that
288 the bacteria induced changes in Tf's molecular mass and structure (Fig. 1C, lane 6). Comparing
289 our findings to studies involving the proteolytic cleavage of transferrin by *Prevotella*
290 *nigrescens* we can deduce that specific cleavage of the protein did not occur, as sub-products
291 with lower molecular mass were not detected on the gel (Duchesne et al., 1999). Therefore, our
292 results suggest a loss of iron ions by holo-Tf, which is consistent with bacteria-produced
293 siderophores having a higher affinity for Fe ions compared to human transferrin (Wilson et al.,
294 2016, Holden and Bachman, 2015). This higher affinity could be exploited by AMB-1 to
295 efficiently compete for ferric ions with the host cells, resulting in iron starvation for the latter.

296

297 We then showed that AMB-1 inoculation with human melanoma cell cultures affects iron
298 homeostasis of the cancer cells. Iron starvation is mainly characterized by alterations in the

299 iron import machinery, specifically by an upregulation of the transferrin receptor 1 on the cell
300 surface. Increased TfR1 expression found on MDA-MB-435S melanoma cancer cells
301 correlates with increasing bacteria ratios. This finding suggests that AMB-1 effectively
302 competes for free iron ions and therefore limits the mineral's availability to MDA-MB-435S
303 cells (Fig. 2A, B). Moreover, a significant increase of TfR1 expression could already be
304 detected 6 h after inoculation (Fig. 2D-F). Similarly, the cancer cells showed a significant
305 upregulation of TfR1 surface expression after incubation with deferoxamine (10 μ M and 25
306 μ M), in line with previous reports on cellular iron deficiency (Torti and Torti, 2013, Lane et
307 al., 2015, Bajbouj et al., 2018). These observations demonstrated that AMB-1 affect the iron
308 import mechanisms of human melanoma cells, acting as an effective competitor for iron when
309 in co-culture with MDA-MB-435S cells.

310
311 Our data support the idea that AMB-1 have the ability to act as living iron chelators by secreting
312 a quantifiable amount of siderophores. We showed that 10^8 AMB-1/mL can produce high-
313 affinity iron scavenging molecules equivalent to 3.78 μ M deferoxamine over 24 h (Fig. 1B).
314 Previous studies demonstrated that the treatment of different cell lines with 10 μ M - 30 μ M
315 deferoxamine significantly reduced cell viability *in vitro* (Ford et al., 2013, Bajbouj et al.,
316 2018). Moreover, a significant diminution of cell viability was even detected at the lower
317 deferoxamine concentration of 2.5 μ M when combined with the chemotherapeutic drug
318 cisplatin (Ford et al., 2013). Nonetheless, the implementation of molecular iron scavenging
319 molecules in translational medicine is hampered by elevated systemic toxicity, as well as
320 limited tumor selectivity. These challenges might be overcome by implementing bacteria as
321 direct competitors for nutrients at the tumor site. The ability of AMB-1 to self-replicate and
322 secrete comparable, sustained doses of siderophores qualifies them as promising candidates for
323 further studies.

324
325 The intrinsic magneto-aerotactic capability of MTB allows them to regulate their motility
326 towards environments with low oxygen concentration as well as react to externally applied
327 magnetic fields (Lefèvre et al., 2014, González et al., 2015). Aerotaxis and hypoxic traits have
328 also been leveraged in *Salmonella* strains, enabling them to act as bacterial anti-cancer agents
329 that target necrotic tumor microenvironments with poor oxygen supply (Mengesha et al., 2006,
330 Yu et al., 2012a, Kasinskas and Forbes, 2007). In addition to enhanced tumor accumulation,
331 native bacterial cytotoxicity, expression of anticancer agents, and immunomodulation have
332 been exploited for their use in clinical cancer therapy (Din et al., 2016, Chowdhury et al., 2019,
333 Forbes, 2010, Duong et al., 2019). By combining iron chelation with these additional benefits
334 of bacterial cancer therapy, we envision that MTB could become a valid therapeutic agent to
335 implement against cancer.

336
337 Our work motivates the use of living AMB-1 as self-replicating iron scavenging organisms
338 actively competing for this vital nutrient, with the possibility of compromising the survival of
339 cancer cells. Further application could include the use of tumor-targeting organisms both as a
340 monotherapy and as a combination therapy with established anti-neoplastic drugs to obtain
341 optimal clinical outcomes. Moreover, the unique characteristics of magnetotactic bacteria
342 could be exploited to engineer iron-scavenging strains of surrogate commensal and attenuated
343 bacteria that have already been established as anti-cancer agents. This work lays the foundation
344 for future investigations which combine iron chelation with bacterial cancer therapy to enhance
345 existing therapeutic strategies and open new frontiers for combating cancer.

346
347

348 **5. Author contributions**

349
350 SS and PSH conceived and designed the experiments. PSH and SM collected and analyzed
351 data. SM wrote the manuscript. TG assisted the authors and contributed to the paper revisions.
352 SS supervised the study and helped writing the paper.

353
354

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356
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362
363

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369
370

371 **8. Conflict of interest**

372
373 The authors declare no conflict of interest.

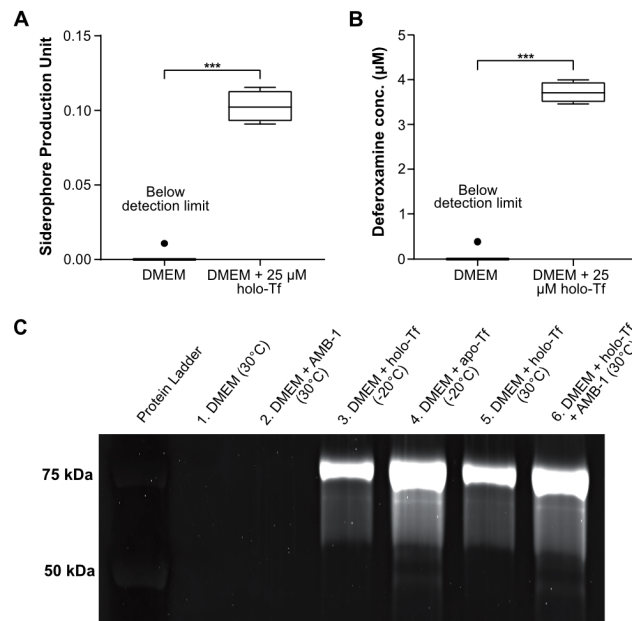
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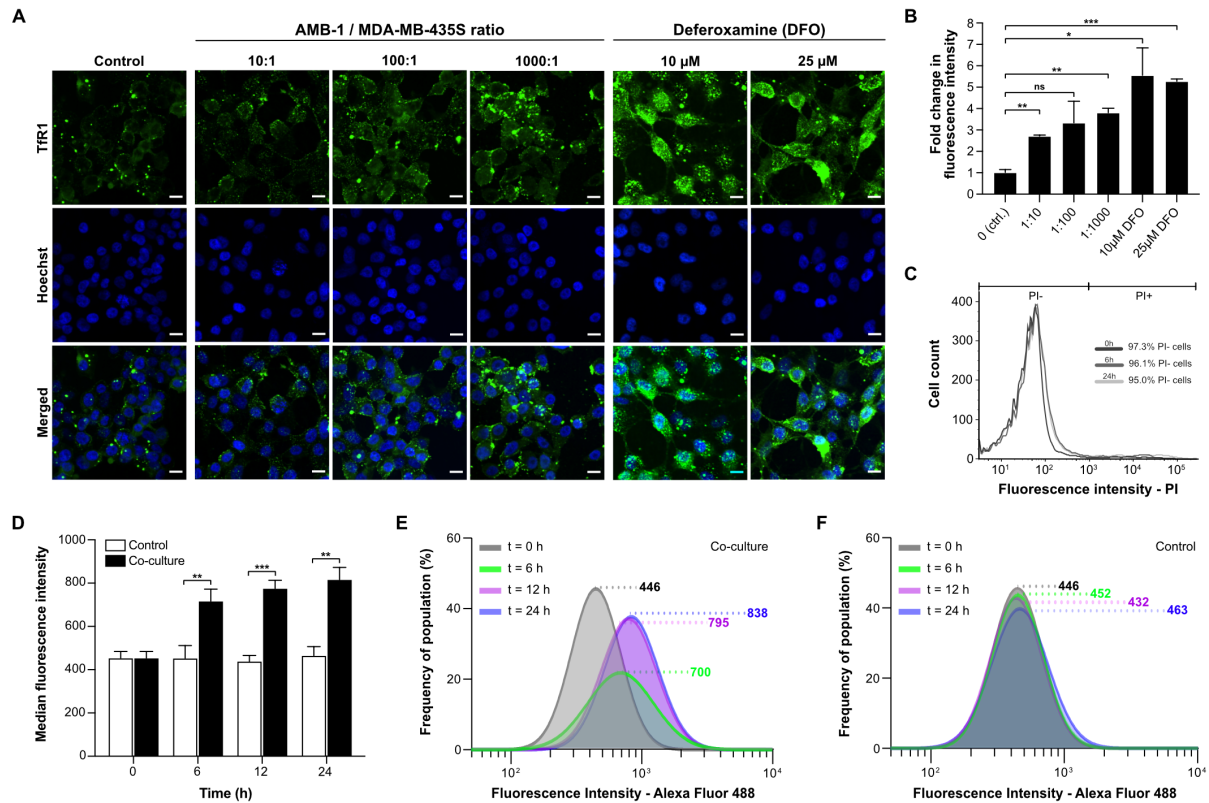
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504 **Figure 1:** Quantification of siderophores produced by *Magnetospirillum magneticum* AMB-1
505 and analysis of their interaction with human transferrin. **(A)** Siderophores produced by AMB-
506 1 were quantified by a chrome azurol S (CAS) assay in DMEM and DMEM supplemented with
507 25 μM holo-transferrin (n=4 per group, statistical significance was assessed with an unpaired
508 two-tailed *t*-test). **(B)** Siderophore production units plotted in terms of the inferred equivalent
509 concentration of deferoxamine (n=4 per group, statistical significance was assessed with an
510 unpaired two-tailed *t*-test). **(C)** SDS-PAGE analysis displaying the effect of AMB-1 on the
511 structure of human transferrin. Tested conditions are indicated in the figure, with holo-Tf
512 corresponding to saturated transferrin and apo-Tf corresponding to non-saturated transferrin.

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536 **Figure 2:** Analysis of TfR1 upregulation and cell surface expression on MDA-MB-435S. (A)
 537 Representative immunofluorescence images of human melanoma cells co-cultured under
 538 hypoxic conditions for 48 h with different ratios of AMB-1 bacteria and different
 539 concentrations of deferoxamine as a positive control. Images show MDA-MB-435S cells
 540 marked by anti-TfR1 antibody (green) and Hoechst 33342 (blue), (scale bar: 10 μ M). (B)
 541 Quantification of the fold change in fluorescence intensity relative to the control condition,
 542 (n=2 per condition, statistical significance was assessed with an unpaired two-tailed *t*-test). (C)
 543 Membrane integrity was measured as a graphical representation of PI negative and PI positive
 544 cell populations after 0, 6 and 24 h. (D) TfR1 median fluorescence intensity measured over 24
 545 hours, (n=3 per timepoint, statistical significance was assessed with an unpaired two-tailed *t*-
 546 test). (E) Representative log-normal fitted fluorescence intensity histograms of cell surface
 547 TfR1 expression on MDA-MB-435S cells in co-culture model and (F) negative control, (n=3
 548 per timepoint).