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1 **Coating Iron Oxide Nanoparticles with Mesoporous Silica Reduces their**
2 **Interaction and Impact on *S. oneidensis* MR-1**

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11 **Abstract**

12 Here, we investigate the impact of iron oxide nanoparticles (IONPs) and mesoporous silica-
13 coated iron oxide nanoparticles (msIONPs) on *S. oneidensis* in an aerobic environment, which
14 is likely the main environment where such nanoparticles will end up after use in consumer
15 products or biomedical applications. Monitoring the viability of *S. oneidensis*, a model
16 environmental organism, after exposure to the nanoparticles reveals that IONPs promote
17 bacterial survival, while msIONPs do not impact survival. These apparent impacts are correlated
18 with association of the nanoparticles with the bacterial membrane, as revealed via TEM and
19 ICP-MS studies, and upregulation of membrane-associated genes. However, similar survival in
20 bacteria was observed when exposed to equivalent concentrations of released ions from each
21 nanomaterial, indicating that aqueous nanoparticle transformations are responsible for the
22 observed changes in bacterial viability. Therefore, this work demonstrates that a simple
23 mesoporous silica coating can control the dissolution of the IONP core by greatly reducing the
24 amount of released iron ions, making msIONPs a more sustainable option to reduce
25 perturbations to the environment upon release of nanoparticles into the environment.

26 **Keywords**

27 Nanotoxicity, bacteria, iron oxide, silica, dissolution, gene expression

28 **1. Introduction**

29 The applications of magnetic iron oxide nanoparticles can range from data storage¹ to
30 biomedical imaging and various therapies.^{2,3} With such widespread use of these inexpensive
31 and easy-to-synthesize nanomaterials, it is inevitable that some of the nanomaterials will end up
32 in environment. Therefore, it is critical to understand the potential environmental impact of these
33 nanomaterials after their release, especially into aquatic environments.

34 Given that iron oxide nanoparticles (IONPs) are often susceptible to aggregation when in
35 suspension with complex biological milieu, a silica shell is commonly added to the iron oxide
36 nanoparticle core to improve their stability in complex matrices,^{4,5} expanding the range of their

37 utility for different applications. Since a mesoporous silica coating has pores in its structure, this
38 method of capping nanoparticles still allows access by water to the core material, an attribute
39 that is critical for performance in a variety of biomedical applications.⁶ For example, literature
40 has shown, when using iron oxide nanoparticles to image tumors in mice, that use of the
41 mesoporous silica shell is required to stabilize the iron oxide nanoparticles so their magnetic
42 properties facilitate the acquisition of images.⁷ With the relevance of this particular platform
43 (mesoporous silica-coated iron oxide nanoparticles) in mind, as well as the general concept of
44 pursuing nanoparticle design motifs that control a nanoparticle's effect on our ecosystem, this
45 work explores the impact of iron oxide and mesoporous silica-coated iron oxide nanoparticles
46 on a model bacterium, *Shewanella oneidensis* MR-1.

47 Bacteria are used as a surrogate for environmental health due to their important role as
48 decomposers to recycle nutrients for use by other organisms. Therefore, any impacts on
49 bacteria due to nanoparticle interaction could also ultimately affect other environmental
50 organisms. *S. oneidensis* MR-1, specifically, is commonly considered for bioremediation since it
51 is capable of respiring many different metals and is therefore an important part of the
52 geochemical nutrient cycle.⁸ *S. oneidensis* has three main mechanisms through which it
53 respires metals (like the constituents of the iron oxide nanoparticles under consideration here):
54 direct interactions with cytochromes on the bacterial surface, the secretion of flavins for
55 extracellular reduction, and growth of electrically conductive pili capable of metal reduction. *S.*
56 *oneidensis* synthesizes many different cytochromes, including cytochrome *c*,⁹ and these
57 cytochromes can be found associated with the outer membrane, periplasmic space, and inner
58 membrane. The system of cytochromes is capable of reducing metals when they come into
59 direct contact with the outer membrane-associated cytochromes.^{10,11} *S. oneidensis* also
60 secretes flavins, which are capable of extracellular electron transfer at distances greater than 50
61 μm from the bacterial cell surface.¹² The flavin system is intimately linked with the outer
62 membrane-associated cytochromes since they shuttle electrons from the cytochromes to the

63 extracellular metals.¹³ The third mechanism for *S. oneidensis* to reduce extracellular metals is
64 through direct contact with nanowires, which can grow to be longer than 10 μm .¹⁴ However, it
65 has been demonstrated that the nanowires are only present in significant quantities on *S.*
66 *oneidensis* when grown under O_2 -limited conditions,¹⁴ and therefore are not relevant in the
67 conditions considered herein.

68 Iron oxides are among the materials that *S. oneidensis* is capable of reducing.^{9,15,16} Often,
69 under anaerobic conditions, *S. oneidensis* reduces iron oxides to synthesize extracellular
70 magnetite.¹⁷⁻¹⁹ For example, when direct contact is made with hematite NPs, *S. oneidensis*
71 reduces them to form magnetite nanoparticles.^{20,21} *S. oneidensis* has also demonstrated an
72 ability to dissolve and reduce magnetite.^{22,23} While many of these studies are performed
73 anaerobically, a related species of *S. oneidensis*, *S. putrefaciens* 200R, has been shown to
74 attach to magnetite at similar levels under both aerobic and anaerobic conditions.²⁴ It has also
75 been observed that some reduction machinery is upregulated in *S. oneidensis* MR-1 under
76 aerobic conditions.²⁵ Furthermore, there is evidence that nitrate can be respiration in aerobic
77 conditions, as well as nitrite and fumarate, which can also be respiration, but only after bacteria
78 have entered the stationary phase; this is true for several electron acceptors that have been
79 studied.^{26,27} While there are many studies exploring the relationships between *S. oneidensis* and
80 bulk iron oxides, there is a dearth of literature looking at the interactions of nanoscale magnetite
81 with *S. oneidensis*. Given that nanoparticles have a much higher surface area to volume ratio
82 than their bulk counterparts, which imparts unique physicochemical properties to particles at the
83 nanoscale, simply extrapolating the interactions of *S. oneidensis* with bulk iron oxide to
84 nanoscale iron oxide may not wholly account for these increased complexities.

85 Herein, we report the interactions between *S. oneidensis* and iron oxide nanoparticles, and
86 then detail our use of a mesoporous silica coating around the IONPs to mitigate their impact.
87 The studies presented here investigate the impact of nanoscale magnetite under aerobic
88 conditions, as those are likely to be the prevailing conditions in aquatic environments where

89 nanoparticles are released. Colony counting was utilized to investigate the impact on bacterial
90 survival after nanoparticle exposure. To understand the differential survival, the binding of the
91 nanoparticles was investigated with TEM and ICP-MS. The contribution of ion release by the
92 nanoparticles was assessed, and any impact on the production of riboflavin was investigated
93 with HPLC. To probe these differences even further, changes in gene expression in *S.*
94 *oneidensis* was analyzed using quantitative reverse transcription polymerase chain reaction
95 (RT-qPCR). These studies showed that IONPs promote bacterial survival through the release of
96 iron ions, and that, by reducing ion dissolution, the addition of a mesoporous silica coating
97 mitigates that impact.

98 **2. Experimental**

99 **2.1 Materials**

100 Chloro-trimethyl silane was obtained from Fluka and 2-[methoxy(polyethyleneoxy)₉₋₁₂propyl]-
101 trimethoxysilane, tech 90 (PEG-silane, molecular weight 591-723 g/mol, 9-12 EO) was
102 purchased from Gelest, Inc. (Morrisville, PA). EMG 308 Ferrofluid was acquired from Ferrotec
103 (Santa Clara, CA). RNAzol® RT was acquired from Molecular Research Center, Inc. (Cincinnati,
104 OH) and iTaq™ Universal SYBR® Green Supermix was obtained from Bio-Rad Laboratories,
105 Inc. (Hercules, CA). Deoxyribonucleotides, random primers, SuperScript III reverse
106 transcriptase, and RNaseOUT™ recombinant ribonuclease inhibitors were acquired from
107 Invitrogen (Carlsbad, CA). Primers for specific genes for *S. oneidensis* were purchased from
108 Integrated DNA Technologies (Skokie, IL). *Shewanella oneidensis* MR-1 was graciously
109 provided by Dr. Jeffrey Gralnick (Dept of Microbiology, University of Minnesota). All other
110 materials used are listed in the Supplementary Information.

111 **2.2 Synthesis of Mesoporous Silica and Mesoporous Silica-coated Iron Oxide**

112 **Nanoparticles**

113 To synthesize MSNs, a previously published protocol was adapted and used.²⁸ Briefly,
114 cetyltrimethylammonium bromide (0.29 g) was mixed with 150 mL of 0.36 M NH₄OH [caution:

115 concentrated NH_4OH is both toxic and corrosive!] and stirred (1 h, 300 rpm, 50 °C). Then, 2.5
116 mL of 0.88 M tetraethylorthosilicate in ethanol was added dropwise and stirred (1 h, 600 rpm, 50
117 °C), followed by slow addition of 450 μL of 2-[methoxy(polyethyleneoxy)₉₋₁₂propyl]-
118 trimethoxysilane and stirring (30 min, 600 rpm, 50 °C). Finally, 68 μL of chlorotrimethylsilane
119 were added and stirred (30 min, 600 rpm, 50 °C) with the beaker covered.

120 A previously published method was used to prepare msIONPs.⁷ Since the purchased IONPs
121 have an anionic, proprietary ligand on their surface, they were first overcoated with PVP-10, a
122 hydrophobic coating that will allow for favorable interactions with the hydrophobic tails of CTAB,
123 used later in the synthesis. To make the EMG 308/surfactant suspension, the following
124 materials were added sequentially into 5 mL of ultrapure water while sonicating, with 1 h
125 sonication steps between each addition: 0.6 g PVP-10, 400 μL EMG 308 suspension, and 0.29
126 g CTAB. To an Erlenmeyer flask, 145 mL of ultrapure water was added. While sonicating, the
127 EMG 308/surfactant suspension was added dropwise; sonication continued for another hour to
128 ensure dispersity. The temperature was increased to 50 °C, and the suspension was stirred (15
129 min, 300 rpm). Then, 2.5 mL of 28% NH_4OH was added and stirred (15 min, 300 rpm, 50 °C),
130 followed by dropwise addition of 2.5 mL of 0.88 M ethanolic tetraethylorthosilicate with stirring (1
131 h, 700 rpm, 50 °C). Slowly, 450 μL of 2-[methoxy(polyethyleneoxy)₉₋₁₂propyl]-trimethoxysilane
132 was added and then stirred (30 min, 700 rpm, 50 °C) and lastly, 68 μL of chlorotrimethylsilane
133 was added and stirred (30 min, 700 rpm, 50 °C) with the flask covered by a glass Petri dish. The
134 suspension was then transferred to a clean 250 mL Erlenmeyer flask without a stir bar.

135 For gentle evaporation, both MSNs and msIONPs were then aged at 50 °C for ~20 h (care
136 was taken so that not all solvent evaporated), followed by hydrothermal treatment at 90 °C for
137 24 h to improve particle stability. Oxygen was removed from msIONPs prior to hydrothermal
138 treatment by purging the reaction vessel and suspension with nitrogen gas to reduce oxidation
139 of the cores at the higher temperatures of hydrothermal treatment. The NPs were purified by
140 ultracentrifugation (30 min, 61,579 $\times g$) and resuspension in 50 mL 6 g/L NH_4NO_3 for reflux (1 h,

141 300 rpm, 60 °C). The suspension was then ultracentrifuged (all subsequent ultracentrifugation
142 steps were done for 20 min at 61,579 $\times g$) and resuspended in 95% ethanol. This was
143 ultracentrifuged and resuspended in 6 g/L NH₄NO₃ to reflux again (1 h, 300 rpm, 60 °C). The
144 suspension was ultracentrifuged three more times and resuspended in the following order: 95%
145 ethanol, 99% ethanol, 99% ethanol. The final suspension was filtered through a 0.2 μ m GHP
146 syringe filter.

147 **2.3 Transmission Electron Microscopy**

148 To prepare the nanoparticles for imaging by transmission electron microscopy, they were first
149 diluted to a suspension of approximately 0.5 mg/mL (IONPs were used at ~2 mg/mL) and
150 sonicated for 10 min to ensure dispersal. Then, for MSNs and msIONPs, a 200 mesh copper
151 grid with Formvar and carbon supports (Ted Pella, Inc., Redding, CA) was briefly dipped into the
152 suspension. For IONPs, a 3 μ L drop of the suspension was placed onto the grid surface. All
153 grids were dried near an open 65 °C oven prior to imaging with a FEI Tecnai T12 transmission
154 electron microscope. To acquire the images, the microscope was used at an operating voltage
155 of 120 kV. Size analysis was performed on the images using ImageJ,²⁹ with size determined by
156 measuring the diameter of at least 500 randomly chosen nanoparticles (it is assumed that all
157 nanoparticles are spherical) using built-in functions of ImageJ.

158 **2.4 Hydrodynamic Diameter and Zeta Potential Measurements**

159 To determine the hydrodynamic diameter and zeta potential of the nanoparticles used in this
160 study, the nanoparticles were first suspended in water at a concentration of 0.5 mg/mL. The
161 hydrodynamic diameters and ζ -potentials were then analyzed using a Brookhaven ZetaPALS
162 instrument. The stability of the nanoparticles in the exposure medium (HEPES buffer) was also
163 assessed by suspending the nanoparticles (300 μ g/mL) in HEPES buffer and checking their
164 hydrodynamic diameter after 30, 60, and 120 min.

165 **2.5 Bacterial Culture Conditions**

166 *S. oneidensis* MR-1 was stored at -80 °C until ready for use, when it was then plated on a
167 sterilized Luria-Bertani (LB) agar plate and incubated overnight at 30 °C. From the plate, 2
168 colonies were inoculated in 10 mL of LB broth overnight to reach the late log phase ($OD_{600} = 0.6$ -
169 1.0). The bacteria were washed by centrifuging (10 min, 750×g), resuspended in D-PBS, and
170 incubated at room temperature on a nutating mixer for 10 min. The bacteria were centrifuged
171 again (10 min, 750×g), and resuspended in a HEPES buffer (2 mM HEPES, 25 mM NaCl, pH
172 7.4) to the appropriate OD_{600} .

173 **2.6 Nanoparticle Dissolution in Bacterial Media**

174 To measure how much iron dissolves in the HEPES buffer after one hour, suspensions in
175 HEPES were made by mixing 450 μ L of nanoparticle stock with 5.55 mL of HEPES buffer to a
176 final concentration of 300 μ g/mL of iron oxide. These were left at room temperature for one hour
177 before being centrifuged at 4700×g for 30 min, followed by centrifuging the supernatant twice at
178 286,000×g for two hours. Removal of nanoparticles was confirmed with DLS, and the iron
179 content of the supernatants was measured with the Thermo Scientific XSERIES 2 ICP-MS.

180 **2.7 Drop plate Colony Counting Assays for Viability**

181 To assess the viability of *S. oneidensis* MR-1 after exposure to the NPs, the bacterial OD_{600} in
182 HEPES buffer was adjusted to 0.2 (which corresponds to $\sim 2 \times 10^8$ bacterial cells/mL) and then
183 diluted 1000-fold. To a suspension of bacteria (925 μ L), NP treatments were added (75 μ L), and
184 exposures lasted for 1 h. Both iron-containing nanoparticles were used at iron concentrations of
185 300 μ g/mL (as a control nanoparticle, the MSNs were used so that the silica mass matched that
186 of the silica in msIONPs (4.7 mg/mL)). Six 10 μ L drops of each suspension were dropped onto
187 dried, UV-sterilized LB agar plates and incubated for ~ 17 h at 30 °C. In separate experiments
188 assessing the viability of bacteria after exposure to iron ions, iron (III) chloride hexahydrate was
189 used for exposure concentrations of 7.6 and 1.0 ppb, to recreate the iron ion concentration
190 determined to be released from IONPs and msIONPs, respectively. The number of colonies that

191 grew in each treatment were counted and recorded. These colony counts were normalized by
192 dividing by the number of colonies that grow in the negative control samples to facilitate
193 comparison between trials.

194 **2.8 Nanoparticle Association with TEM**

195 To visualize the binding of nanoparticles to *S. oneidensis*, the bacterial OD₆₀₀ was adjusted to
196 0.8 (which corresponds to $\sim 8 \times 10^8$ bacterial cells/mL) before mixing the bacterial suspension
197 (925 μ L) with nanoparticles (75 μ L). The nanoparticles were used at the same concentrations as
198 for colony counting experiments. After a one-hour exposure, the samples were centrifuged at
199 800 $\times g$ for 5 min and the supernatant discarded.

200 The bacterial samples were prepared for TEM by adapting previously reported methods.^{30,31}
201 The samples were washed thrice without resuspension using 0.1 M cacodylate buffer,
202 centrifuging at 500 $\times g$ for 2 min between each wash step. To fix the sample, the pellet was
203 resuspended in 2.5% glutaraldehyde in 0.1 M cacodylate buffer for 50 min, followed by
204 centrifuging at 800 $\times g$ for 5 min. The pellet was again washed three times with 0.1 M cacodylate
205 buffer without resuspension. To dehydrate the samples, a series of ethanol washes was done at
206 increasing ethanol concentrations in water, using each concentration twice (30%, 50%, 70%,
207 80%, 95%, and 100% ethanol). The samples were then washed three times with propylene
208 oxide prior to using a 2:1 propylene oxide:resin mix for two hours, uncovered. This 2:1 mixture
209 was replaced with 1:1 propylene oxide:resin to soak overnight, after which it was replaced with
210 fresh 1:1 propylene oxide:resin for four hours. The samples were then incubated in pure resin
211 overnight, which was replaced with fresh resin the next day. To cure the resin, the samples
212 were put in a 40 °C oven for 24 h and a 60 °C oven for 48 h. The samples were sliced into ~ 70
213 nm thick sections using a LEICA EM UC6 ultramicrotome, which were stained with uranyl
214 acetate and lead citrate. The slices were placed on 200 mesh copper grids with carbon and
215 Formvar supports (Ted Pella, Inc., Redding, CA). Images of the samples were acquired using a
216 FEI T12 transmission electron microscope at an operating voltage of 120 kV.

217 **2.9 Nanoparticle Association using ICP-MS**

218 To measure the amount of iron oxide nanoparticles and mesoporous silica-coated iron oxide
219 nanoparticles that bind to *S. oneidensis*, the OD₆₀₀ of bacteria was adjusted to be 0.2 (~2×10⁸
220 bacterial cells/mL) and then exposed to nanoparticles by mixing 925 µL of bacterial suspension
221 with 75 µL of nanoparticles to achieve final concentrations of 300 µg/mL of iron oxide. Control
222 experiments were set up simultaneously that mixed HEPES buffer with nanoparticles. After the
223 one-hour exposure, the samples were centrifuged twice at 800×g, discarding all but 50 µL of the
224 supernatant since the pellet is easily disturbed. After the first centrifugation, the pellet was
225 resuspended carefully in 1 mL HEPES buffer, and after the second centrifugation, the pellet was
226 resuspended in 150 µL of HEPES buffer. The resuspended pellet (125 µL) was diluted with 375
227 µL of concentrated nitric acid (*Caution: use care when handling corrosive acids*) and the sample
228 was digested at 60 °C overnight. The samples were then diluted 14-fold and centrifuged at
229 17,000×g for 20 min to remove bacterial cell debris. The samples were analyzed using the
230 Thermo Scientific XSERIES 2 ICP-MS. To determine the amount of iron associated with the
231 bacteria, the iron concentration detected in the respective control and the iron contribution by
232 the bacteria were subtracted from the iron concentration in the experimental/bacterial samples,
233 and then the dilutions were taken into account.

234 **2.10 Riboflavin Secretion Measurements**

235 A previous method was adapted to determine the amount of riboflavin secreted by bacteria after
236 nanoparticle exposure.³² To begin, the OD₆₀₀ in HEPES buffer was adjusted to 0.2. The bacteria
237 were then exposed to NPs for 1 h and then centrifuged at 17,000×g for 20 min to pellet both the
238 bacteria and NPs. The supernatant was collected, and 200 µL was transferred to an HPLC vial
239 with 250 µL glass insert. The supernatant was analyzed using an Agilent 1200 HPLC fitted with
240 a Zorbax Eclipse XDB-C₁₈ analytical column (4.6 x 150 mm, 5 µm packing material) and an
241 Eclipse XDB-C₁₈ analytical guard column (4.6 x 12.5 mm, 5 µm packing material) ahead of the
242 fluorescence detector. Isocratic HPLC was performed with a mobile phase composed of 70/30

243 20 mM citric acid buffer (pH 3.3)/methanol with an injection volume of 30 μ L, flow rate of 1
244 mL/min, and run time of 8 min. Excitation and emissions wavelengths of 450 and 530 nm,
245 respectively, were used to detect riboflavin.

246 **2.11 Extracting RNA from *S. oneidensis* after Nanoparticle Exposure**

247 The RNA extraction was performed using a Zymo Research Direct-zolTM RNA MiniPrep Plus kit.
248 To do the one-hour nanoparticle exposure prior to RNA extraction, the bacterial OD₆₀₀ in
249 HEPES buffer was adjusted to 1.0 (corresponding to $\sim 1 \times 10^9$ bacterial cells/mL), and 1.85 mL of
250 bacterial suspension was mixed with 150 μ L of nanoparticle suspension (or water as a negative
251 control), using nanoparticle concentrations that matched the colony counting experiments.
252 Afterward, the suspensions were centrifuged at 2000 $\times g$ for 10 min and the pellet was
253 resuspended in 200 μ L of RNAzol RT. Centrifugation at 16,000 $\times g$ for 1 min removed particulate
254 debris, and 200 μ L of 200 proof molecular biology grade ethanol was added to the supernatant
255 and mixed. This mixture was added to a Zymo-Spin IIC column in a collection tube and
256 centrifuged at 16,000 $\times g$ for 30 sec, discarding the flow through. To the column, 400 μ L of RNA
257 wash buffer was added and centrifuged at 16,000 $\times g$ for 30 sec. To do a DNase I treatment, 5
258 μ L of DNase I and 75 μ L of DNA digestion buffer were mixed separately and then transferred to
259 the column surface, incubating at room temperature for 15 min to digest DNA. Then, the column
260 was washed twice by sequential additions of 400 μ L of Direct-zol RNA PreWash and
261 centrifuging at 16,000 $\times g$ for 30 sec, discarding the flow through. 700 μ L of RNA wash buffer
262 was added to the column and centrifuged at 16,000 $\times g$ for 2 min. The column was transferred to
263 a clean, RNase-free Eppendorf tube where 80 μ L of DNase/RNase-free water was added and
264 centrifuged at 16,000 $\times g$ for 30 sec to collect the RNA product. The concentration and quality of
265 the RNA was measured using a Thermo Scientific NanoDropTM One^C.

266 **2.12 Monitoring Gene Expression Changes in *S. oneidensis* after Nanoparticle Exposure**

267 Total purified RNA was reverse transcribed into complementary deoxyribonucleic acid
268 (cDNA) as previously described.³³ cDNA was synthesized using 100 ng of total RNA template

269 and was incubated with an aliquot of deoxyribonucleotides and random primers at 65 °C for 5
270 min followed by chilling on ice for 1 min. SuperScript III reverse transcriptase, dithiothreitol, and
271 RNaseOUT™ recombinant ribonuclease inhibitors were added into the mixture after the
272 incubation following the temperature program of 25 °C for 5 min, 50 °C for 60 min, and 70 °C for
273 15 min for primer extension. Synthesized cDNA was then stored at -20 °C.

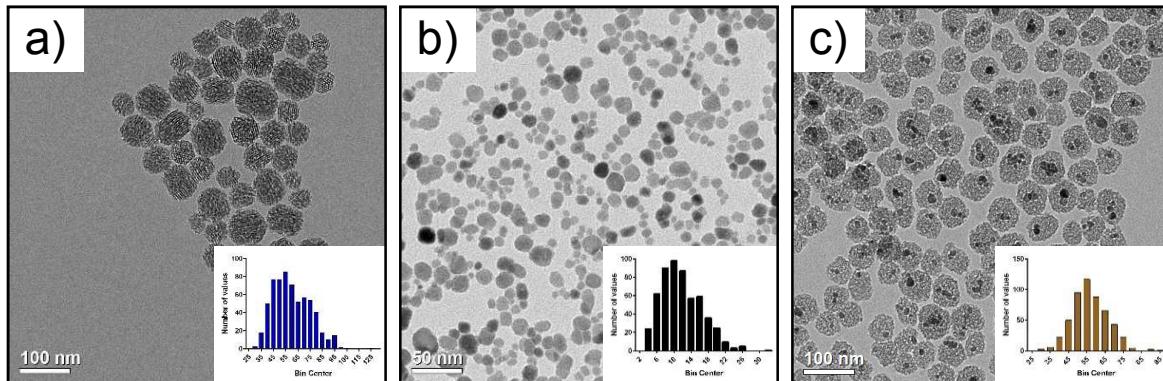
274 Quantitative reverse transcription PCR was run to determine the expression levels of several
275 genes involved in iron acquisition, storage, and utilization for *S. oneidensis*, with *radA* and *gyrB*
276 genes serving as housekeeping genes for analysis. The list of genes used in this study can be
277 found in the Supplementary Information (Table S1); these genes were selected to assess
278 different aspects of the interaction between *S. oneidensis* and iron. An iQ5 real-time PCR
279 detection system (Bio-Rad Laboratories) using SYBR Green for the fluorescent intercalating dye
280 (iTaq™ Universal SYBR® Green Supermix) was used for this process. Each of the qPCR
281 reactions containing cDNA (1 µL) mixed with primers (1 µL) with fluorescent dye (10 µL) and
282 nuclease-free water (8 µL) was carried out in 96-well PCR plates, centrifuging the plates at
283 1000×g for 10 min at 4 °C prior to running the qPCR. The polymerase chain reactions started at
284 95 °C for 10 min for DNA denaturing, then underwent 40 real-time PCR amplification cycles (15
285 s at 95 °C, followed by 30 s at 60 °C). Fluorescence of the SYBR Green was then detected at
286 the end of each PCR cycle. All samples were analyzed with technical duplicates.

287 **3. Results and Discussion**

288 **3.1 Characterization of MSNs, IONPs, and msIONPs**

289 The nanomaterials used in this study were characterized by a variety of methods. Using TEM,
290 the morphology and size distribution of the materials could be determined (Figure 1). The
291 mesoporous silica nanoparticles were found to have a diameter of 60 ± 15 nm (n=632), similar
292 to the diameter of 57 ± 10 nm (n=521) found for mesoporous silica-coated iron oxide

293 nanoparticles. The purchased iron oxide nanoparticles were smaller, with a diameter of 12 ± 5
 294 nm (n=557).



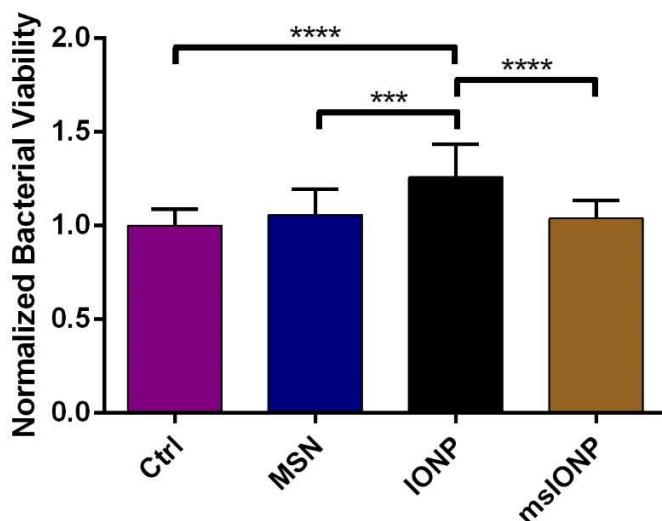
296 **Figure 1.** Transmission microscope images reveal the size and dispersity of a) mesoporous silica
 297 nanoparticles, b) iron oxide nanoparticles, and c) mesoporous silica-coated iron oxide nanoparticles. An
 298 inset in each image shows the size histogram determined for each nanoparticle type. Note that the scale bar
 299 for image (b) is half the size of that for the other two images.

300 The hydrodynamic diameters of the material were also determined for MSNs (86 ± 21 nm),
 301 msIONPs (90 ± 6 nm), and IONPs (55 ± 6 nm), reflecting the trend seen by TEM size analysis.
 302 The ζ -potential for msIONPs (-39 ± 12 mV) was more negative than that for MSNs (-18 ± 9 mV)
 303 or IONPs (-23 ± 15 mV); these values suggest that all three nanoparticle formulations should be
 304 relatively colloidally stable. In fact, throughout the course of the exposure experiment, all three
 305 nanoparticles were stable, as indicated by their unchanging hydrodynamic diameter in HEPES
 306 buffer over 2 h (Figure S1).

307 **3.2 Impact of Nanoparticles on *S. oneidensis***

308 After a one-hour exposure to the different nanoparticles, the impact to *S. oneidensis* was
 309 assessed using a colony counting assay (Figure 2). The results demonstrate that while
 310 mesoporous silica-coated iron oxide nanoparticles and mesoporous silica nanoparticles have no
 311 impact on bacterial survival compared to the control, the iron oxide nanoparticles enhance the
 312 growth of the bacteria in HEPES buffer and therefore have an enhanced survival rate. Iron is an
 313 essential nutrient for bacteria,³⁴ thus it is reasonable that the presence of exogenous iron could
 314 enhance their growth. Specifically for *S. oneidensis* MR-1, iron oxide is a metal that they are

315 capable of respiring anaerobically as a means of obtaining energy;⁹ alternately, dissolution
 316 product iron ions may contribute to bacterial sustenance. To probe the mechanism by which iron
 317 helps sustain bacteria in HEPES buffer, the contributions of the various means by which *S.*
 318 *oneidensis* can respire iron oxide were determined, as well as any impact of iron dissolution
 319 from the iron oxide nanoparticles. To evaluate how the mesoporous silica coating mitigates the
 320 impact by IONPs, effects on these endpoints by IONPs and msIONPs were compared.



321
 322 **Figure 2.** The colony counts determined from each nanoparticle type after being normalized to the control
 323 indicates that mesoporous silica and mesoporous silica-coated iron oxide nanoparticles have no impact on
 324 survival while the iron oxide nanoparticles enhance bacterial survival compared to a negative control. The
 325 error bars represent standard deviations from sixteen replicates. Statistical significance was evaluated using
 326 a one-way ANOVA with Tukey's multiple comparisons post-hoc test. ***p<0.001, ****p<0.0001

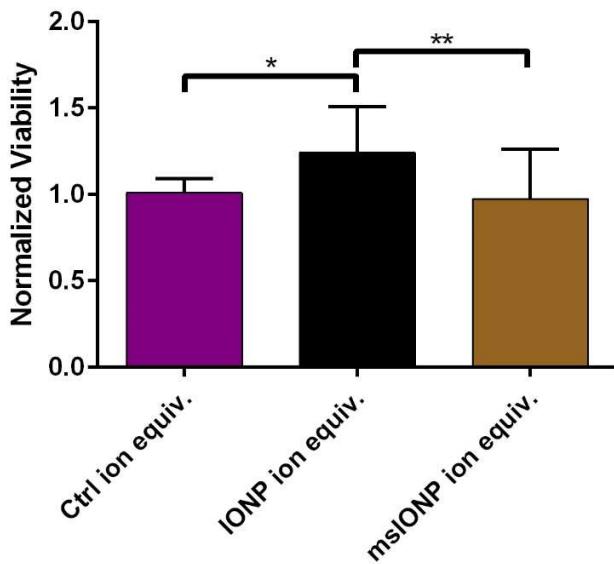
327 **3.3 Differential Ion Release is Observed for Nanoparticles**

328 To determine the concentrations of iron that were released from IONPs and msIONPs during
 329 their one-hour exposure to bacteria in HEPES buffer, ICP-MS was used to measure iron
 330 concentrations from an abiotic supernatant after removing the NPs from suspension. Results
 331 showed that IONPs released significantly more iron (7.6 ± 0.2 ppb) compared to msIONPs ($1.0 \pm$
 332 0.4 ppb) after an hour in HEPES buffer ($p<0.0001$). This observation suggests that a silica
 333 coating mitigates dissolution of the nanoparticle core.³⁵ While these results indicate the iron
 334 concentrations released from the nanoparticles, they do not specify the oxidation states of the

335 dissolved iron. MINTEQ modeling was used to show that at equilibrium, Fe^{3+} is the dominant
336 oxidation state for the dissolved iron (Table S2), prompting the use of iron (III) chloride
337 hexahydrate for the ion control studies performed.

338 **3.4 Iron Ion Exposure Recapitulates Effect Seen by Nanoparticles**

339 *S. oneidensis* were exposed to the concentrations of iron ions released during nanoparticle
340 exposure, and their viability was assessed using the colony counting assay. From this assay, it
341 was seen that the iron ion exposure recapitulates the results of nanoparticle exposure, with the
342 iron ion concentrations equivalent to the release from the control and msIONP trials both
343 showing a similar viability around 1.0, and the iron ion concentrations equivalent to release from
344 IONPs showing an increased bacterial survival (Figure 3). The bacterial viability after exposure
345 to the ion concentration representing IONP exposure was 1.24 ± 0.27 , which is not statistically
346 distinct from the 1.26 ± 0.18 viability seen after exposure to IONPs. These comparisons
347 demonstrate that the dissolved ion constituents of these nanoparticles are dominating the
348 observed effect of the nanoparticles on *S. oneidensis*. In addition, these results show that, since
349 the presence of a mesoporous silica coating reduces dissolution to levels that do not impact *S.*
350 *oneidensis*, this reduced dissolution is the major contributor to the mechanism by which the
351 mesoporous silica coating mitigates IONP impact.



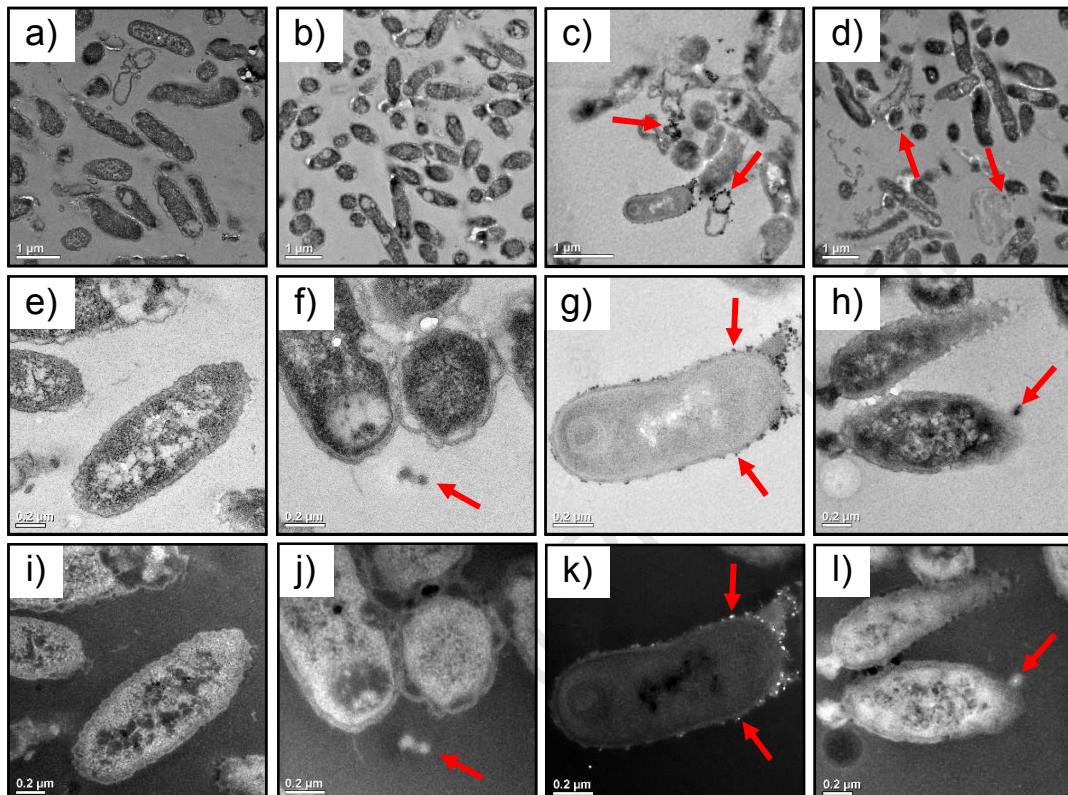
352

353 **Figure 3.** At Fe^{3+} ion concentrations that match those released during nanoparticle exposure, it can be seen
 354 that 7.6 ppb iron ions (equivalent to the released concentration from IONPs) are enhancing bacterial survival
 355 to the same extent as observed with IONP exposure, while at 1.0 ppb (equivalent to the released
 356 concentration from msIONPs), no impact on growth is observed. These data demonstrate that the impact
 357 from the iron ions is recapitulating the impact observed from the nanoparticle exposures, and therefore ion
 358 dissolution is the main contributor to the nanoparticles' effects. The error bars represent the standard
 359 deviations from fifteen replicate trials. One-way ANOVA with Tukey's multiple comparisons test was used to
 360 evaluate statistical significance. * $p < 0.05$, ** $p < 0.01$

361 **3.5 IONPs Display Significant Binding to the Bacterial Surface**

362 While the ion studies indicate that ion release is the major contributor to the impacts of the
 363 nanoparticles, the other mechanisms by which *S. oneidensis* can interact with iron oxide
 364 nanoparticles were assessed as well. Some of these mechanisms involve direct binding with the
 365 cell wall, thus, the interactions of the NPs with bacteria were visualized by first fixing the
 366 samples and embedding them in resin. Images of the resin-embedded samples were then
 367 acquired with TEM. For samples containing IONPs and msIONPs, the TEM was performed in
 368 dark-field mode to verify that nanoparticles were observed by taking advantage of the scattering
 369 efficiency of crystalline iron oxide nanoparticles.³⁶ These images clearly show that the IONPs
 370 are binding to *S. oneidensis* to a greater extent than MSNs or msIONPs (Figure 4). Given that
 371 all three nanoparticles used in this study have a negative surface charge, which often leads to
 372 electrostatic repulsion from the net-negative charge of the bacterial membrane, it is not

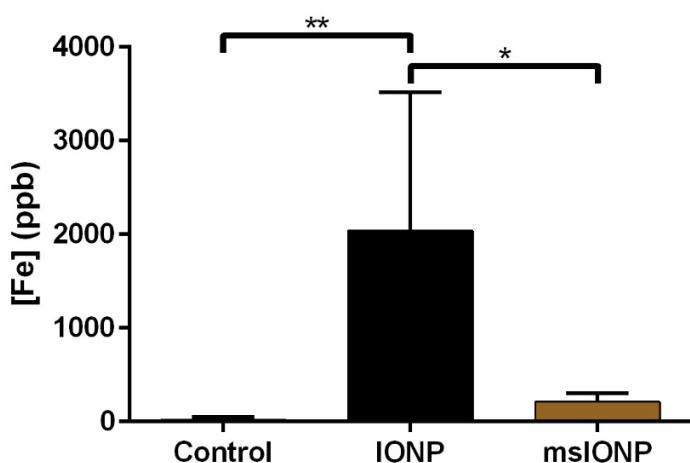
373 surprising that there is no interaction between the membrane and MSNs or msIONPs. However,
 374 it is interesting that there is such prevalent binding of IONPs, which implies specific interactions
 375 by the bacteria with this material.



376
 377 **Figure 4.** A low magnification view of bacteria after exposure to a) nanoparticle-free suspension, b) MSNs, c)
 378 IONPs, and d) msIONPs as well as a high magnification view of e) NP-free suspension, as well as bacteria
 379 exposed to f) MSNs, g) IONPs, and h) msIONPs reveal that there is more binding with IONPs than the other
 380 two nanomaterials. No instance of direct binding by MSNs could be found, which indicates that if they do
 381 occur, they are a rare occurrence. Where present, instances of nanoparticles in each image are highlighted
 382 with arrows. Dark-field images of i) NP-free bacteria are included as a comparison to bacteria exposed to j)
 383 MSNs, k) IONPs, and l) msIONPs. The bright points are due to the high scattering efficiency of IONPs in both
 384 images and are not seen in the control image. Mesoporous silica nanoparticles do not exhibit the same
 385 scattering intensity in dark-field mode due to their noncrystalline nature.

386 Beyond the visual evidence from TEM, quantitative evidence of increased IONP binding was
 387 acquired by exposing the bacteria to nanoparticles, pelleting the bacteria, and digesting them to
 388 quantify the amount of iron material that was associated with the bacteria (likely the surface,
 389 based on TEM data). Since iron quantitation was the endpoint, this does not reveal anything
 390 about the binding of MSNs to the bacterial surface. Corroborating the TEM images, a significant
 391 increase in iron content from bacteria that were exposed to IONPs was found compared to

392 those that were exposed to msIONPs (Figure 5). In fact, msIONPs displayed bacterial
 393 association that was not statistically different from the negative control. These quantitative data
 394 demonstrate that *S. oneidensis* is bound to 0.9% of the available NPs when exposed to IONPs
 395 and to 0.01% of the available msIONPs in that exposure. This was calculated by dividing the
 396 bound iron mass by the total iron content present in each exposure.



397

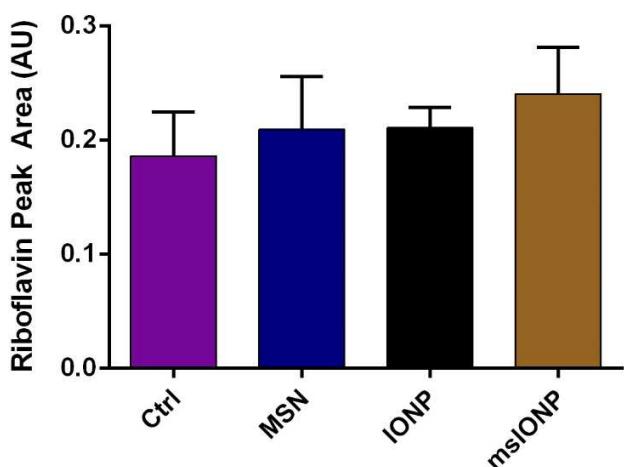
398 **Figure 5.** The iron content determined from bacteria that had been exposed to IONPs and msIONPs shows
 399 that there is significant iron bound to the surface of IONP-exposed bacteria. Statistical testing was performed
 400 with a one-way ANOVA of at least three replicates, using Tukey's multiple comparisons test to compare the
 401 effects of each treatment. * $p<0.05$, ** $p<0.01$

402 Taken together, these data show that there is significantly more binding of IONPs to the
 403 bacterial surface than MSNs or msIONPs. While this observation is important, it was shown in
 404 Section 3.4 that the bacterial viability after exposure to nanoparticles was recapitulated by the
 405 released iron ions, demonstrating that this increased association has negligible contributions to
 406 the effects of IONPs to bacterial survival. Therefore, while coating the IONPs with a
 407 mesoporous silica shell does reduce the association, this reduced physical association does not
 408 contribute to the mitigation of the impact of IONPs on bacterial viability.

409 **3.6 Riboflavin Production is Unchanged by Nanoparticle Exposure**

410 Riboflavin secretion is one method by which *S. oneidensis* is capable of respiring
 411 extracellular iron oxide. To assess the mechanism that extracellular electron shuttling via

412 riboflavin secretion is contributing to the beneficial impacts seen with IONP exposure, the
 413 production of riboflavin in bacteria was monitored via HPLC. Secreted riboflavin, which elutes
 414 around 6.3 min with the separation method used, was measured from all of the samples;
 415 however, it is noted that the secretion of riboflavin from *S. oneidensis* is not changed by the
 416 presence of any of the nanomaterials (Figure 6). Given that riboflavin is secreted by bacteria to
 417 perform extracellular electron transfers to respiration metals, these findings indicate that this
 418 riboflavin-mediated extracellular electron transport mechanism is not a major contributor to the
 419 beneficial impact seen with IONPs.



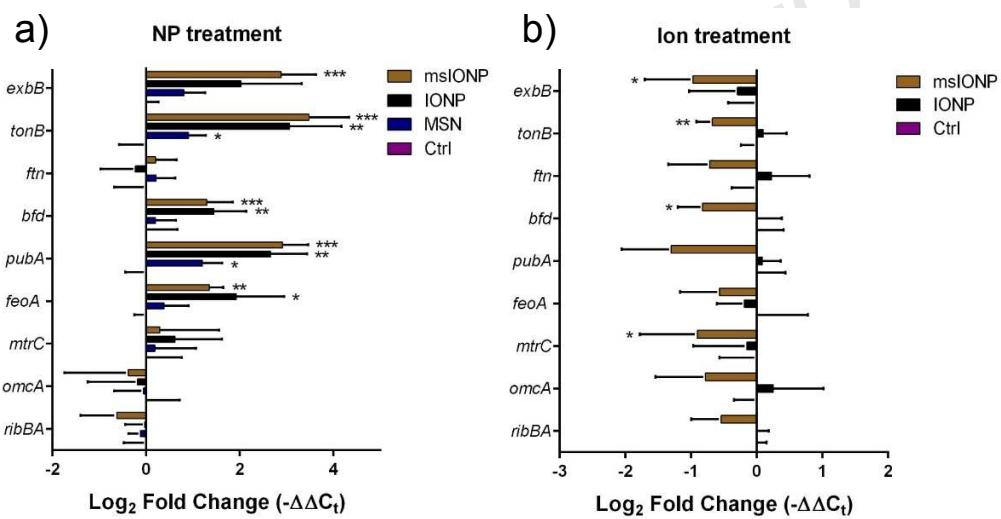
420
 421 **Figure 6.** The secretion of riboflavin by *S. oneidensis* is not impacted by exposure to any of the
 422 nanomaterials used in this study. The error bars represent the standard deviations from six replicates. One-
 423 way ANOVA with Tukey's multiple comparisons test was used to determine statistical significance between
 424 the treatments.
 425

426 **3.7 Changes in Gene Expression are Nanoparticle-Specific**

427 Since it is clear that *S. oneidensis* is using the iron oxide nanoparticles, it was expected that
 428 there would be changes in gene expression of bacterial genes relating to the transport, storage,
 429 and utilization of iron after exposure to IONPs and their released iron equivalents. To probe this,
 430 changes in gene expression after exposure to nanoparticles or to the released iron ion
 431 equivalents were monitored using RT-qPCR (Figure 7). There was no noted change in the
 432 expression of *ribBA*, whose function is in the synthesis of riboflavin. This corroborates the lack

433 of increased production of riboflavin after NP exposure in the HPLC analysis shown in Figure 6,
434 and verifies that this mechanism is not contributing significantly to the impact of IONPs on *S.*
435 *oneidensis*. Most of the genes that encode for proteins that are associated with the membrane
436 of the bacteria, *exbB*, *tonB*, *feoA*, have been upregulated by IONPs and msIONPs. However,
437 *pubA*, which does not encode for a membrane-associated protein, was also upregulated after
438 nanoparticle treatment. Instead, *pubA*, encodes for a siderophore that complexes with ferric iron
439 and shuttles it back to the bacterial cell for storage via a TonB-dependent siderophore receptor,
440 which suggests that its function is still related to processes at the bacterial surface.³⁷ Given that
441 *exbB* and *tonB* both have functions related to the intracellular uptake of iron,³⁸ it is clear that *S.*
442 *oneidensis* is working to sequester at least some of the extra iron that it is encountering in the
443 presence of the IONPs or msIONPs; however, the increased bacterial association seen with
444 IONPs helps to facilitate this process better than with msIONPs. For the gene, *ftn*, which
445 corresponds with iron sequestration, there is no change in expression noted, which makes
446 sense given that in an iron-rich environment, the bacterium would not need to store it for later
447 use. Conversely, for *bfd*, a gene that encodes for a protein that initiates use of sequestered iron,
448 upregulation is observed. An interesting observation is that there is increased expression of
449 *feoA*, which is specific for ferrous ion transport into the bacterial cell.³⁹ Since our MINTEQ
450 calculations showed that the oxidation state of the majority of released iron would be Fe³⁺, the
451 increased *feoA* expression suggests that *S. oneidensis* may be processing the nanoparticles to
452 make ferrous ion, which is then transported into the bacterial cell for use. The fact that gene
453 expression changes after exposure to IONPs and msIONPs appear to be very similar, and yet
454 only IONP exposure is aiding bacterial survival, indicates that while genes for iron uptake and
455 usage are being upregulated in both exposures, there is more iron present in the IONP
456 exposures for the bacteria to actually use. This highlights the fact that gene expression changes
457 are more sensitive to environmental changes than more macro-level endpoints such as overall
458 bacterial survival.

459 These observed gene expression changes appear to be nanoparticle-specific, as treatment
 460 with equivalent doses of released ions did not induce the same changes. In fact, exposure to
 461 ions caused very few gene expression changes, with just a few genes being downregulated
 462 upon exposure to 1 ppb Fe^{3+} . We speculate that the observed association of the nanoparticles
 463 to the bacteria may be initiating these gene expression changes by dissolving to form a higher
 464 localized concentration of iron ions right at the bacterial surface, especially if the bacteria are
 465 also assisting in the dissolution by processing the nanoparticles to generate ferrous ion, as
 466 suggested by the upregulation of *feoA*.



467

468 **Figure 7. Gene expression changes in *S. oneidensis* after exposure to a) nanoparticles and b) equivalent**

469 released ion concentrations. The error bars represent standard deviations from five replicates. One-way

470 ANOVA with a Dunnett's multiple comparisons test was used to determine statistical significance between

471 the different treatments and the control. *p<0.05, **p<0.01, ***p<0.001

472 **4. Conclusions**

473 The work presented herein investigates the impact of IONPs and their mesoporous silica-coated
 474 counterparts on *S. oneidensis*. After exposure to the nanoparticles, colony counting reveals that
 475 IONPs promote bacterial survival. Exposing *S. oneidensis* to ferric ion at a dose equivalent to
 476 released iron from IONPs and msIONPs recapitulates the viability seen with nanoparticle
 477 exposure, indicating that ion release is the major contributor to the bacterial impact of IONPs.
 478 As expected, the presence of a mesoporous silica shell on IONPs reduced the iron dissolution

479 observed, which explains the mitigated impact on viability of msIONPs on the bacteria.
480 Association between the nanoparticles and *S. oneidensis* shows that there is the greatest
481 bacterial association with IONPs, with minimal association with msIONPs or MSNs.
482 Interestingly, while the ions account for the enhanced bacterial survival after exposure to the
483 nanoparticles, at the genetic level, a nanoparticle-specific effect is observed since the ion
484 controls did not induce the same gene expression changes. Given that the genes impacted
485 upon nanoparticle exposure mostly encode for membrane-associated proteins, we speculate
486 that the observed association of nanoparticles with the bacterial membrane may cause
487 upregulation of these genes. This could be due to a higher localized concentration of iron
488 released near the bacterial surface or simply due to the direct interaction between the
489 nanoparticles and bacterial membrane. Since perturbations to the environment, whether by
490 increasing or decreasing survival for select organisms, can be detrimental in some situations,
491 the mesoporous silica-coated iron oxide nanoparticles are deemed here to be a better option for
492 sustainability. Given that a mesoporous silica shell should reduce dissolution for other core
493 materials, this strategy could also be applied to more toxic NPs whose major toxicity mechanism
494 is related to dissolution of ions, where it would similarly be expected to reduce harmful impacts
495 to organisms.

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508 **6. Competing Interests**

509 The authors declare no competing financial interest.

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518 **7. Supplementary Information**

519 Materials used; list of genes analysed in gene expression studies; stability of nanoparticles in
520 exposure medium; MINTEQ evaluation of dissolved iron species; representative HPLC
521 chromatograms.

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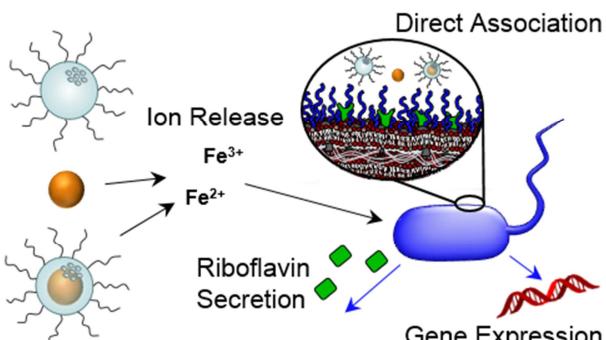
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647 **9. TOC Graphic**



648

Highlights

- Iron oxide nanoparticles enhance survival of the bacterium, *Shewanella oneidensis*.
- Dissolved iron released from IONPs recapitulates growth pattern observed with NP exposure.
- A mesoporous silica coating reduces IONP dissolution, thus mitigating their impact.
- Changes in gene expression demonstrate a nanoparticle-specific effect.