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**As release under the microbial sulfate reduction during redox
oscillations in the upper Mekong delta aquifers, Vietnam: a
mechanistic study**

1 **As release under the microbial sulfate reduction during redox oscillations**
2 **in the upper Mekong delta aquifers, Vietnam: a mechanistic study**

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31 **Abstract**

32 The impact of seasonal fluctuations linked to monsoon and irrigation generates redox oscillations in
33 the subsurface, influencing the release of arsenic (As) in aquifers. Here, the biogeochemical control
34 on As mobility was investigated in batch experiments using redox cycling bioreactors and As- and
35 SO_4^{2-} - amended sediment. Redox potential (E_h) oscillations between anoxic (-300 – 0 mV) and oxic
36 condition (0 – 500 mV) were implemented by automatically modulating an admixture of N_2/CO_2 or
37 compressed air. A carbon source (cellobiose, a monomer of cellulose) was added at the beginning of
38 each reducing cycle to stimulate the metabolism of the native microbial community. Results show
39 that successive redox cycles can decrease arsenic mobility by up to 92% during reducing conditions.
40 Anoxic conditions drive mainly the conversion of soluble As(V) to As(III) in contrast to oxic
41 conditions. Phylogenetic analyses of 16S rRNA amplified from the sediments revealed the presence
42 of sulfate and iron – reducing bacteria, confirming that sulfate and iron reduction are key factors for
43 As immobilization from the aqueous phase. As and S K-edge X-ray absorption spectroscopy
44 suggested the association of Fe-(oxyhydr)oxides and the importance of pyrite ($\text{FeS}_{2(s)}$), rather than
45 poorly ordered mackinawite ($\text{FeS}_{(s)}$), for As sequestration under oxidizing and reducing conditions,
46 respectively. Finally, these findings suggest a role for elemental sulfur in mediating aqueous
47 thioarsenates formation in As-contaminated groundwater of the Mekong delta.

48 **Keywords:** Redox oscillations, Arsenic release, Microbial sulfate reduction, Elemental sulfur

49

50 1. Introduction

51 Arsenic, a well-known environmental carcinogen, is subject to a variety of mobility-altering
52 processes (WHO IARC, 2004). The World Health Organization's provisional guideline value for
53 total As in drinking water is 10 µg/L. Concentrations of arsenic in groundwater higher than 50 µg/L
54 have been reported in several Southeast Asian countries including Bangladesh, India, Cambodia,
55 China, and Vietnam (e.g. Wang et al. 2018; Phan et al. 2017; He & Charlet, 2013; Fendorf, 2010;
56 Kocar et al., 2008; van Geen et al., 2008; Zheng et al., 2004; Buschmann et al., 2008). In particular,
57 the Mekong Delta floodplain, which stretches over 62100 km² in Southern Vietnam and South
58 Cambodia, harbors numerous wells with elevated As concentrations (Buschmann et al., 2008;
59 Nguyen et al., 2000). A national survey, conducted by the Department of Water Resources
60 Management monitored the As concentration in tube wells from 2002 to 2008, showed that large
61 areas of the Mekong Delta were contaminated by arsenic, in nearly 900 wells (Erban et al., 2013).
62 Areas with As concentrations in groundwater exceeding 1600 ppb are located along the main
63 Mekong river branches, from An Giang to Can Tho City (Wang et al., 2018; Phan et al., 2017; Hanh
64 et al., 2011; Hoang et al., 2010; Nguyen an Itoi, 2009). The origin of As in groundwater of the
65 Mekong Delta is naturally produced by microbial reduction of arsenic-bearing iron oxides in the
66 alluvial sediments (Polya et al., 2005; Rowland et al., 2007; Van Dongen et al., 2008; Kocar &
67 Fendorf, 2012; Polizzotto et al., 2008; Stuckey et al., 2015; Winkel et al., 2010). Following river
68 sediment transport, As bound to Fe-(oxyhydr)oxides is delivered to the delta from the Himalayan
69 mountain range. Arsenic is then released to pore water due to the establishment of reducing
70 conditions (Berg et al., 2007; Harvey et al., 2002; Polizzotto et al., 2005; Kocar et al., 2008; Nguyen
71 & Itoi, 2009; Stuckey et al., 2015). Natural organic matter (NOM) supplied from surface
72 sedimentary deposits feeds the dissimilatory reducing bacteria by serving as electron donors that
73 play the important role on As mobilization in this area (Kocar et al., 2008; Lawson et al., 2013;
74 Stuckey et al., 2015; Lawson et al., 2016; Magnone et al., 2017).

75 In nature, the seasonal reversal of flow direction - rivers to aquifers in the wet season and aquifers to
76 rivers in the dry season - influences the redox balance of the aquifers (Benner et al., 2008; Stuckey
77 et al., 2015). Redox oscillations occurring across the oxic-anoxic interfaces, which are frequent in
78 periodically inundated swamplands due to the changes of the water levels, may affect the release of
79 metal (loid)s (e.g., As, Fe) into the groundwater (Karimian et al., 2017). The behavior and transport
80 of As are closely related to the biogeochemical cycling of Fe, S and C (e.g. Bostick et al., 2003;
81 O'Day et al., 2004; Root et al., 2009). Microbial sulfate (SO₄²⁻) reduction, couples with the

82 oxidation of organic matter (OM) or H_2 , may also affect As behavior via the production of sulfide
83 (S(-II)) (Barton, 1995, Poulton et al., 2004; Rochette et al. 2000, Muyzer & Stams, 2008). Under
84 different conditions, HS^- can immobilize As by the precipitation of As sulfides (e.g., orpiment
85 ($As_2S_3(s)$) and realgar ($AsS(s)$) if the As concentration is high enough (O'Day et al., 2004; Root et
86 al., 2009), or form reduced inorganic sulfur such as $FeS_{2(s)}$ and $FeS_{m(s)}$ (Berner, 1970; Rickard, 2006;
87 Rickard & Iii, 2007) onto which As can adsorb (Wolthers et al., 2005, Couture et al., 2013b).
88 Arsenic can also co-precipitate with iron sulfide minerals to form arsenic-substituted pyrite, known
89 as “arsenian pyrite” (e.g. Rittle et al., 1995; Newman et al., 1998; Charlet et al., 2011; Langner et
90 al., 2012; Le Pape et al., 2017; Wang et al., 2018). Lastly, arsenite is known to bind strongly onto
91 sulfhydryl groups (thiols) of OM (ThomasArrigo et al., 2016; Langner et al., 2013; Couture et al.,
92 2013a; Langner et al., 2012). The dominance of one mechanism over the other is determined by
93 variety factors (e.g., aqueous As speciation, pH, Fe and S mineralogy) whose identification is
94 complicated by dynamic conditions.

95 There is significant interest in elucidating the processes driving As mobility in sediments under
96 sulfate reducing conditions and to understand hydrological transport controls on As fluxes from
97 sulfate-rich sediments in the Mekong Delta (Métral et al., 2008; Buschmann & Berg, 2009).
98 Nevertheless, experimental studies accurately isolating biogeochemical processes during the
99 oscillating redox conditions in Vietnamese paddy fields and shallow aquifers are still scarce. In
100 particular, the role of sulfate-reducing bacteria (SRB) and sulfate-oxidizing bacteria (SOB) in As
101 immobilization in the sediment during redox dynamic conditions remain poorly defined at such
102 sites. To determine the mechanisms controlling As mobility in sulfate-rich sediment of seasonally
103 flooded soils, we investigated the As, Fe, and S species generated by oscillating redox conditions in
104 laboratory experiments. We used automated bioreactors containing sediments with equal total As
105 concentration and variable sulfate contents. A 16S rRNA amplicon analysis was performed to
106 unravel the role of bacteria in the sediment through redox cycles and X-ray absorption near-edge
107 spectroscopy (XANES) was exploited to reveal solid-phase As species.

108 **2. Materials and methods**

109 **2.1 Field site characterization**

110 **2.1.1 The study area and sediment sampling**

111 The seasonally flooded sediment was collected at the Quoc Thai commune, An Giang province,
112 Vietnam, in a paddy soil near the Mekong River during the dry season, in 2014 (Fig. SI-1) where
113 was contained a high As concentration in groundwater (Wang et al., 2018, Phan et al., 2017). A

114 sediment core was drilled from a depth of 0 – 20 m and separated into sections as a function of
115 depth under an argon (Ar) flux and then immediately placed in the heat sealed Mylar[®] bags under N₂
116 atmosphere (Wang et al., 2018). The sediment samples were shipped in a cooler with ice back to
117 France and stored at +4°C until processing. Peat-rich sediment from 16m - deep layers was obtained
118 for the batch experiments. All preparation and samples conditioning were performed in a Jacomex[®]
119 under N₂ atmosphere (O₂ < 10 ppm) to ensure anoxic conditions.

120 2.1.2 Mineralogy and elemental composition analyses

121 The elemental composition was measured using the total acidic digestion (e.g. HNO₃ + H₂O₂ + HF,
122 H₃BO₃ + HF) followed by Inductively Coupled Plasma – Optical Emission Spectrometry (ICP-OES)
123 (Agilent 720-ES, Varian) (Cotten et al., 1995). X-Ray diffraction (XRD) measurements were
124 performed with CoK α radiation on a Panalytical[®] X'Pert Pro MPD diffractometer using a Debye–
125 Scherrer configuration using an elliptical mirror to obtain a high flux and parallel incident beam, and
126 an X'Celerator[®] detector to collect the diffracted beam. All sediment samples were transferred into
127 glass capillaries inside a glovebox Jacomex[®] (O₂ \leq 10 ppm) to avoid oxidation by air.

128 2.2 Experimental design and redox oscillation set-up

129 Two batch bioreactors, R1 and R2, were prepared with 1L of sediment suspension containing 100g
130 of dry sediment (<1 mm fraction), arsenite (As(III)), and two different concentrations of sulfate
131 (SO₄²⁻). The contaminants were prepared by dissolving of sodium (meta)arsenite (NaAsO₂, \geq 98%,
132 Sigma-Aldrich) and sodium sulfate (Na₂SO₄(s), \geq 99%, anhydrous, granular, Sigma-Aldrich), to
133 obtain a solution with final concentration of 50 μ M of As(III), 0.1 and 1 mM of SO₄²⁻ for R1 and
134 R2, respectively. The suspensions were pre-equilibrated inside the glove box for 2 weeks before
135 doing the redox experiments. More details of the reactor system were described in the previous
136 studies (e.g. Parsons et al., 2013, Markelova et al., 2018, Phan et al., 2018). At the start of the redox
137 experiment (t = 0), two mixture solutions containing the same concentration of As(III) and two
138 different concentrations of SO₄²⁻ were added in the two reactors (R1 and R2). Cellobiose (C₁₂H₂₂O₁₁
139 – Sigma-Aldrich), a byproduct of microbial hydrolysis of cellulose (Lynd et al., 2002;
140 Schellenberger et al., 2011), was manually replenished at the start of each anoxic half-cycles to
141 obtain the final concentration of DOC equal to 8.33 mM including the DOC in sediments. All input
142 solutions were adjusted for ionic strength (I = 30 mM) using sodium chloride (NaCl - Sigma-
143 Aldrich). Reactors were covered with aluminum foil to avoid the photo-degradation of organic
144 matter and contaminants.

145 Reduced – oxidized oscillations were carried out by automatically modulating the influx gas
146 between the mixture of N₂ and 300 ppm CO₂ in anoxic half-cycle (7 days) except for the first one
147 with 5 days, and compressed air of 1% CO₂, 79% N₂ and 20% O₂ in oxic half-cycle (7 days) using
148 an Agilent switching unit and a system of solenoid valves. Redox cycles were monitored for
149 approximately 40 days. Gas flow rate and temperature were kept constant at 30 mL/min and 30°C.
150 Suspension samples were collected five days/week of each half-cycle for aqueous, solid phase and
151 microbial analyses through a connection on the top of the reactors. Except for the days collecting the
152 plus 5 mL slurry for microbial analysis, a 15 mL of suspension was collected in the other days.
153 These suspensions were centrifuged at 14000 rpm for 20 min (Sigma 3-30 KS) to separate the solid
154 phase (e.g., XRD and XANES analysis), while the supernatant was then filtered through cellulose
155 hydrophilic 0.22µm membrane (Chromafil RC, Roth). A part of supernatant was acidified with
156 HNO₃ (2%, Sigma-Aldrich) or HCl (0.1 M, Roth) and preserved at 4°C for total element
157 concentration and DOC analysis, respectively. A 2 mL of supernatant was stored at 4°C without
158 acidification for anion measurement. The rest of supernatant, slurry samples were flash frozen using
159 liquid nitrogen and kept at – 80°C for another aqueous and microbial analysis.

160 **2.3 Aqueous phase characterization**

161 All standards and reagents, were prepared with ultra-pure water, and were of analytical grade from
162 Fluka, Sigma-Aldrich or Merck. The glass and plastic parts were firstly washed with 5% HNO₃ then
163 with ultra-pure water (18 MΩ.cm⁻¹). The E_h and pH data were recorded automatically every 10
164 seconds using an Aligent 34970A BenchLink Data Logger through the E_h and pH electrodes
165 (Mettler-Toledo Xerolyt Solid). Measured E_h value was corrected for the reference electrode's
166 voltage (+207 mV) relative to the standard hydrogen electrode (SHE), and corresponded to the
167 Ag/AgCl (3 M KCl) reference electrode. The pH electrodes were calibrated using the measured data
168 of three-point using pH buffers of 4, 7 and 10 at 25°C at the start and the end of experiment
169 displaying that the electrode response were not shifted significantly. The real pH values were then
170 calculated based on the calibration curves.

171 Analysis of total cations, including trace and major elements, were performed with ICP-OES
172 (Agilent 720-ES, Varian) after dilution and acidification by ultrapure HNO₃ (2%, Sigma-Aldrich)
173 with a detection limit of 0.02 and 0.1 mg. L⁻¹, respectively and a precision better than 5%. DOC was
174 determined using a Shimadzu VCSN analyzer (TOC-5000, Shimadzu) with a detection limit of 0.3
175 mg/L and a precision better than 2.5% for most samples. Nevertheless, it is important to note that for
176 concentrations close to the detection limit, the precision can be significantly lower (50-100%). DOC

177 tubes were heated at 400°C for 3 hours avoiding organic carbon contamination. Non-acidified
178 samples were used to analyze major anions (e.g. F⁻, Cl⁻, NO₃⁻, NO₂⁻, Br⁻, SO₄²⁻, PO₄³⁻) and acetate by
179 ion chromatography using a Metrohm 761 compact ion chromatography with a detection limit of 0.1
180 mg. L⁻¹ and a precision better than 5%. [Fe²⁺] and [S(-II)] were determined photometrically on the
181 sample filtered using the Ferrozine method (Lovley & Phillips, 1986; Viollier et al., 2000, Stookey,
182 1970) and Cline method (Cline, 1969), respectively. Standard stock solutions were prepared using
183 Na₂S*9H₂O for the calibration curve of S(-II) and FeCl₂*4H₂O for the calibration curve of Fe²⁺. The
184 standards and samples were then measured immediately using UV-Vis spectroscopy (Lambda 35,
185 Perkin Elmer) at 562 nm and 664 nm absorbance for Fe²⁺ and S(-II), respectively.

186 Aqueous As species (e.g. As(III), As(V), MMA and DMA) were analyzed at Plateforme AETE –
187 HydroScience/OSU OREME, Montpellier, France, using the coupling LC-ICP-MS system (Bohari
188 et al., 2001). The detail condition was described in Phan et al. (2018). The detection limit was 0.09
189 µg/L for As(III), 0.06 µg/L for DMA, 0.04 µg/L MMA and 0.41 µg/L for As(V), with a precision
190 better than 5%.

191 **2.4 Microbial community analysis**

192 The 16S rRNA gene diversity was accessed through paired-end Illumina MiSeq at Research and
193 Testing Laboratory (Texas, USA) following the method of Zhang et al. (2014). Slurry samples were
194 collected at the end of the reduced cycle and in the middle of the oxidized cycle in both reactors
195 (Fig. 1a-b). The overall approach entails 16S rRNA gene amplicon sequencing as a function of time
196 and geochemical conditions. A 0.25 g suspension sample was used for total genomic DNA
197 (gDNA) extraction using the PowerSoil[®] DNA Isolation kit (MO BIO Laboratories, Carlsbad,
198 USA). The DNA quantification was determined using a Nanodrop 1000 Spectrophotometer
199 (Thermo Fisher Scientific, USA). After the DNA quantification, the amplification of the 16S rRNA
200 gene was followed by the polymerase chain reaction (PCRs) using the forward (28F,
201 GAGTTTGATCNTGGCTCAG) and reverse (519R, GTNTTACNGCGGCKGCTG) primers that
202 amplify the V1-V3 region (Fan et al., 2012) (Biometra T3 Thermocycler, Germany). The PCR
203 reaction mix for amplification contained 5 µL 10×PCR buffer, 1 µL dNTPs (10 mM), 2 µL each of
204 the forward and reverse primer (5 µM) and 0.8 µL Taq polymerase and balance water with the total
205 volume of 50 µL. The thermal program, set up of a 5 min at 95°C, 1 min at 50°C, 1 min at 68°C, 34
206 cycles at 95°C for 1 min/cycle, 10 min at 68°C, was performed with the Lightcycler[®] 480 (Roche).
207 PCR products were conceived on an agarose gel, and the 16S rRNA band excised and purified using
208 the Wizard[®] SV Gel and PCR Clean-Up System (Promega, USA). The sequencing process of PCR

209 16S rRNA amplicons were merged, denosed, quality trimmed and demultiplexed using USEARCH
210 (Edgar, 2013). These sequences were then clustered into operational taxonomic units (OTUs) using
211 the UPARSE algorithm with a similarities higher than 97% (Edgar, 2013), and then checked for
212 Chimera detection using UCHIME (Edgar, 2011). A summarized database of high quality sequences
213 derived from the NCBI database was described in the OTUs table (Table 1), and the full data shown
214 in Table S-3 (Supporting Information).

215 2.5 Solid-phase characterization

216 2.5.1 X-ray absorption near edge structure (XANES)

217 Solid-phase samples were collected from the two reactors at the end of the redox cycles indicated in
218 Fig. 1a-b. X-ray absorption near edge structure (XANES) was used to characterize both As and S
219 speciation. S K-edge XANES data were collected at the XAFS beamline of the Elettra synchrotron
220 (Trieste, Italy), whereas As K-edge XANES data were done at the bending magnet beamline BM08-
221 LISA of the European Synchrotron Radiation Facility (ESRF, Grenoble, France).

222 Sulfur K-edge (2472 eV) measurements were performed using a Si(111) monochromator calibrated
223 relative to the white line of a $\text{Na}_2\text{S}_2\text{O}_3(\text{s})$ standard (Sigma-Aldrich) at 2471.64 eV. Spectra were
224 collected in fluorescence mode using a solid-state Si detector. A commercial elemental sulfur
225 reference was measured in fluorescence mode before each scan for accurate energy calibration. For
226 the linear combination fitting (LCF) of XANES spectra at the S K-edge, the following suite of sulfur
227 references was used: (i) FeS (mackinawite), synthesized under strict anoxic conditions with
228 solutions obtained after dissolution of $\text{FeCl}_2 \cdot 6\text{H}_2\text{O}$ and $\text{Na}_2\text{S} \cdot 9\text{H}_2\text{O}$ salts, (ii) pure FeS_2 (pyrite) was
229 synthesized according to the protocol described in Le Pape et al. (2017), (iii) S_8 (elemental sulfur,
230 Sigma – Aldrich), and (iv) commercial K_2SO_4 (sulfate). Some of the reference compounds spectra
231 (e.g. FeS, FeS_2 and K_2SO_4) were acquired on the 4-3 beamline at SSRL in the fluorescence mode.

232 For LCF of XANES spectra at the As K-edge, the following set of arsenic references compounds
233 was considered: (i) NaAsO_2 and NaHAsO_4 (Sigma-Aldrich), MTA(V) (monothioarsenate), DTA(V)
234 (dithioarsenate), and TeTA(V) (tetrathioarsenate), prepared using the method of Schwedt &
235 Rieckhoff (1996) and optimized by Suess et al. (2009); (ii) As-bearing minerals from natural
236 deposits were considered such as orpiment (As_2S_3), realgar (AsS), arsenian pyrite (As-substituted
237 FeS_2), and arsenopyrite (FeAsS) (Le Pape et al., 2017), (iii) As(III)- and As(V)-sorbed ferrihydrite
238 were used as proxies for As(III)-O and As(V)-O local molecular environments (Hohmann et al.,
239 2011; Ona-Nguema et al., 2005) and (iv) glutamyl-cysteinyl-glyciny-thiolarsenite (As(III)-Glu)

240 synthesized using the method of [Miot et al., \(2008\)](#) was used as representative of thiol bound As(III)
241 organic matter (OM) species (see in Table S-4 in Supporting information for more details on the
242 references used). Some of the reference compounds spectra (e.g. arsenian pyrite and thiol-bound
243 As(III)) were collected at 15K in fluorescence mode at the SAMBA beamline at SOLEIL
244 synchrotron ([Phan et al., 2018](#))

245 Both of As and S XANES data analysis were performed using the Athena software ([Ravel &](#)
246 [Newville, 2005](#)) and the relative contributions of the different compounds to the sample spectra
247 were achieved by linear combination fitting (LCF). A homemade program based on a Levenberg-
248 Marquardt algorithm was used to improve the fit quality following the method of Resongles et al.
249 (2016), which was described in detail in Phan et al. (2018).

250 **2.5.2 Acid-volatile sulfide and elemental sulfur analysis**

251 To complement the S K-edge XANES results, we measured acid-volatile sulfide (AVS) ([Hsieh et](#)
252 [al., 2002](#); [Burton et al., 2008](#); [Couture et al., 2016](#)). AVS extraction was based on a purge and trap
253 method ([Allen et al., 1993](#)), which relies on the conversion of sulfur compounds within sediments
254 after centrifuging and keep dry in the glovebox that are first purged with HCl to generate volatile
255 H₂S, and secondly trapped with NaOH. Finally, a complexation is done by adding a diamine reagent
256 to the trapped solution to form a methylene blue molecule ([Cline, 1969](#)), which is quantified by UV
257 spectroscopy at 670 nm. The calibration curve was performed by the same method using Na₂S
258 solution at 10mM. AVS purge and trap measurements were carried out within airtight Teflon
259 reactors under continuous flow of nitrogen to avoid oxidation of sulfide. A quantity of 0.18 to 0.50 g
260 of previously freeze-dried sediment was used for AVS extractions and quantification.

261 The amount of sulfur that remained in the solid phase at the surface in the form of elemental sulfur
262 (S₈) was analyzed using a perchloroethylene extraction combined with High Performance Liquid
263 Chromatography (HPLC) following the method described in [McGuire & Hamers \(2000\)](#).
264 Perchloroethylene-extractable sulfur was obtained after pretreatment of unfiltered samples with 250
265 μL of ZnAc (5%) to precipitate free sulfide ([Wan et al., 2014](#)). Sulfide fixation allows ZnAc to react
266 with S(-II) as well as with S_n²⁻ (n ≥ 2, polysulfides) leading to the precipitation of ZnS. After 10
267 minutes, 4 mL of perchloroethylene was injected into 0.5 mL of suspensions. The samples were
268 shaken for 3 hours and filtered through 0.22μm pore size membranes. The supernatants were then
269 analyzed by HPCL (PerkinElmer 2000 pump and auto-sampler, UV-vision detectors and software

270 AZUR V6.0 software) using a C18 column (Nucleosil 100-5PAH) and isocratic elution in methanol
271 95% at a flow rate of 0.4 mL/min. The detection was performed at a wavelength of 265 nm.

272 2.5.3 Quantitative total As concentration

273 To determine the bulk As concentration, sediment samples, collecting in the end of each anoxic an
274 oxic half-cycles, were dissolved completely using a hot plate digestion (Cotten et al., 1995). About
275 50 mg of dried and homogenized sample were first added with 0.84 mL of distilled HNO₃ (14 N)
276 and 0.2 mL of H₂O₂ to dissolve organic matter, then followed by mixing of 0.4 mL of distilled HF
277 (48%) under a fume hood, and put on a hot plate at 80°C for 3 days. This solution was let cool down
278 before adding 20 mL of H₃BO₃ (4.5%). The final extract was diluted with MilliQ water to 250 mL
279 and analysed for total As concentration using ICP-AES.

280 2.6 Thermodynamic modeling

281 Thermodynamic calculations were performed using the PHREEQC program version 3.3.10
282 (Parkhurst & Appelo, 2013). The following calculations were performed: (i) thermodynamic
283 aqueous species distribution during the anoxic and oxic half cycles, and (ii) the saturation indices
284 (SI = log IAP/K_{sp}) at each sampling point with respect to solid arsenic, sulfur and iron phases. The
285 WATEQ4F database for the As species was updated according to the data reported in the literature
286 (Helz & Tossell, 2008). The added species included As(III) species (H_nAsO₃ⁿ⁻³ (As(III) – arsenite),
287 H_nAsO₄ⁿ⁻³, H_nAsSO₂ⁿ⁻³ (MTA(III) – monothioarsenite), H_nAsS₂Oⁿ⁻³ (DMA(III) – dithioarsenite),
288 H_nAsS₃ (TTA(III) – trithioarsenite), and As(V) species (H_nAsO₄ⁿ⁻³ (As(V) – arsenate), H_nAsSO₃ⁿ⁻³
289 (MTA(V) – monothioarsenate), H_nAsS₂O₂ⁿ⁻³ (DTAs(V) – dithioarsenate), H_nAsS₃Oⁿ⁻³ (TTA(V) –
290 trithioarsenate) and H_nAsS₄ⁿ⁻³ (TeTA(V) – tetrathioarsenate)). Their reactions and equilibrium
291 constants are summarized in Table S-1 in Supporting Information.

292 3. Results

293 3.1 Aqueous chemistry

294 The aqueous phase results of the redox experiments carried out in reactors R1 and R2, containing
295 respectively with 0.1 mM and 1 mM of sulfate, are summarized in Fig. 1.

296 **E_h and pH cycling.** The reducing and oxidizing steps can be clearly identified monitoring the E_h
297 and pH values as a function of time (Fig. 1a-b). E_h ranges from -300 mV, in anoxic half-cycles, to
298 +500 mV, in oxic half-cycles, and pH varied in the range between 5.2 and 7.8 for both reactors. E_h
299 and pH underwent repeated cycling in the low sulfate reactor (R1). However, in the high sulfate
300 reactor (R2), oxidizing conditions were not fully attained in all cycles. Reducing half cycles were

301 characterized by a decrease in E_h and an increase in both pH and in the Fe^{2+} concentration, while the
302 oxidizing half cycles showed opposite trends (Fig. 1a-b). Intra-cycle E_h changes were similar to the
303 E_h -monitoring studies of flooded soils and previous redox oscillation experiments (Couture et al.,
304 2015; Parsons et al., 2013; Thompson et al., 2006). In the reductive processes, the decrease in E_h
305 may be driven by the consumption of successive terminal electron acceptors, such as Fe(III) and
306 SO_4^{2-} coupled to DOC oxidation by the microbial community (Essington, 2004), which resulted in
307 hydroxyl (OH^-) production and pH increase.

308 **DOC cycling.** A 8.33 mM of DOC was manually replenished after adding cellobiose at the start
309 every anoxic half-cycle (Fig. 1c-d). During the oxic half-cycle, in both reactors, DOC decreased
310 significantly together with the low concentration of acetate during each oxic half-cycles probably
311 due to respiration oxidation of Fe^{2-} , S(-II) and As(III). During the anoxic half-cycle, in R1 DOC also
312 decreased slightly, while in R2 with higher SO_4^{2-} , DOC increased significantly together with an
313 increase production of acetate. This is probably due to the release of DOC from sediment was
314 greater in amount of DOC consumption by heterotrophic bacteria (Parsons et al., 2013).

315 **Total Fe and Fe(II) cycling.** In both reactors, $[Fe]$ and $[Fe^{2+}]$ increased in the reducing cycles,
316 consistently with the reduction reaction of labile Fe-oxides (Essington, 2004; Thompson et al.,
317 2006). During the oxic cycles, $[Fe^{2+}]$ dropped when E_h values increased indicating that Fe^{2+}
318 originating from initial pyrite oxidation and Fe-(oxyhydr)oxides was oxidized to Fe(III), which
319 precipitated as poorly soluble phases, e.g., ferrihydrite and goethite. Furthermore, thermodynamic
320 predictions suggest that in anoxic sulfidic conditions, mackinawite ($FeS_{(s)}$) is the first iron sulfide to
321 precipitate, and it constitutes a major component of the empirically defined “acid volatile sulfides”
322 (AVS) (Rickard & Morse, 2005). The saturation index of FeS ($SI > 0$) was calculated using
323 PHREEQC in both reactors and is reported in Fig. 1e-f.

324 **S cycling.** $[SO_4^{2-}]$ decreased during the anoxic half-cycles and increased during the oxic half-cycles
325 and reached about 1.1 mM in R1 and 2 mM in R2, respectively, both were > 1 mM, compared to the
326 added SO_4^{2-} of 0.1 mM at R1 and 1 mM at R2. This difference is probably due to the oxidation of
327 pyrite from the sediment (Fig. 1g-h). Additionally, complete $[S(-II)]$ depletion was observed during
328 the oxidation cycles, while in reduction cycles, $[S(-II)]$ varied in the range between 0.8 - 8.7 μM in
329 R1, and 1.4 - 4.8 μM in R2 (Fig. 1g-h). Low $[S^0]$ was detected in supernatants, which is consistent
330 with the fact that elemental sulfur is typically associated with the solid phase (S_8) (McGuire &
331 Hamers, 2000; Wan et al., 2014). $[S_8]$ was found in the suspension at concentration ranges between

332 0.06 – 8.7 μM and 0.1 – 2.7 μM in R1 and R2, respectively. This finding suggests that sulfate
333 reduction were followed by sulfide oxidation to sulfur and sulfate.

334 **As cycling.** Intra-cycle mobilization of aqueous As was observed during the anoxic half cycles.
335 Particularly, As was released in the initial anoxic half-cycle, reaching up to 32 μM and 27 μM in R1
336 and R2, respectively, and then sequestered in the solid phase in subsequent anoxic cycles.
337 Conversely, oxidizing conditions re-immobilized As, returning to a base-level of approximately
338 1 μM in both R1 and R2. The As decreasing trend and the concomitant increasing in E_h were
339 observed (Fig. 1k-l). When the oxidizing condition prevail, the most thermodynamically favorable
340 As species was As(V) (H_2AsO_4^- and HAsO_4^{2-}). The reducing condition was established during the
341 anoxic half-cycles, which favors the formation of arsenite (H_3AsO_3). Successive redox cycles
342 resulted in 92% and 83% removal of total dissolved As in R1 and R2, respectively.

343 **3.2 Microbial community analysis**

344 Operational taxonomic unit (OTU) libraries were generated for the samples from day 19 (anoxic,
345 R1- or R2-19) and 22 (oxic, R1- and R2-22), as shown by the red arrows in the Fig. 1a-b. Numerous
346 bacterial phyla were identified in the microbial community during the reducing and oxidizing phases
347 mainly consist of Proteobacteria including Alpha-, Beta-, Gamma-, Delta- and Epsilon-
348 proteobacteria classes, and the phyla Bacteroidetes and Firmicutes. The full data are presented in
349 Table S-3, Supporting Information.

350 In reactor R1, there were only three phylotypes that were more abundant under anoxic conditions than
351 under oxic conditions and present at greater than 1% of OTUs: *Arcobacter* sp., represented the
352 largest group in R1, with more than 60% of the OTUs, *Klebsiella* p., representing ~6% of the OTUs,
353 and *Acidovorax* sp., representing 2% of the OTUs in R1. The first two organisms were also more
354 abundant under anoxic conditions in R2 (Table 1).

355 In contrast, and somewhat surprisingly, taxa related to SRB and iron-reducing bacteria were
356 detected under both oxic and anoxic conditions. *Geobacter* species (*Geobacter* sp. and *Geobacter*
357 *bremensis*), known iron and sulfur reducers, were enriched in the two experiments (R1 and R2)
358 during both the oxic and anoxic cycles, representing ~2.7 and ~6.7% (anoxic and oxic, respectively)
359 of the OTUs in R1 and ~6 and ~25% of the OTUs in R2 (anoxic and oxic, respectively). SRB,
360 including *Desulfobulbus* sp., *Desulfomicrobium* sp., and *Desulfovibrio* sp., were detected in both
361 reactors with higher contributions in reactor R2 than in reactor R1 (Table 1).

362 In the oxic half-cycles, taxa including iron- and sulfur-oxidizing bacteria such as *Thiobacillus* sp.
363 were detected suggesting the microbial oxidation of Fe^{2+} , S^{2-} or sulfur (Hedrich & Johnson, 2011).
364 This was particularly true for R1 (*Thiobacillus* sp. represented 3.3% of OTUs) and less so for R2
365 (0.4% of OTUs). Iron oxidizers, e.g., *Rhodobacter* sp., potentially able to use reduced sulfur
366 compounds (S^0 , HS^- , $\text{S}_2\text{O}_3^{2-}$), H_2 , or organic compounds as electron donors, were also identified,
367 particularly in R2 (17% and 14.3% of OTUs in the anoxic and oxic half-cycle, respectively).
368 Additionally, the genus *Hydrogenophaga* sp., are facultatively autotrophic aerobic hydrogen-
369 oxidizing bacteria (Willems et al., 1989) and OTUs related to this genus were predominantly
370 identified in the oxic phase in R1 (11.8% of OTUs in oxic and 3.6% in anoxic) and in the anoxic
371 phase in R2 (7.2% of OTUs in oxic and 9.3% in anoxic).

372 Finally, the presence of many heterotrophic bacteria, capable of operating under both aerobic and
373 anaerobic conditions, such as *Rhizobium* sp. and *Klebsiella* sp., may account for the oxidation of
374 OM during both phases (Eller & Frenzel, 2001; Franciscon et al., 2009). Other heterotrophic
375 bacteria, OM respiring bacteria, take a prominent part of the overall community in both reactors
376 representing from 17 – 45% of total OTUs (Table 1).

377 3.3 Solid sulfur and arsenic dynamics

378 The initial anoxic sediments contained 32.1 g/kg dry sediment organic C, 0.7 g/kg total S, 15.6 g/kg
379 total Fe (Table S-1). Smectite, quartz, muscovite, chlorite, albite, and pyrite were identified in the
380 input sediment (Table S-1). The total As content was lower than detection limit as determined by
381 total digestion and ICP-MS, but As K-edge XANES indicated that As was initially present in the O-
382 bound As(V) and As(III) oxidation states, S bound As(III), and arsenian pyrite.

383 3.3.1 Solid S speciation

384 LCF of the XANES spectra at the S K-edge revealed that the majority of S accumulated in R1 and
385 R2 as $\text{FeS}_{2(s)}$, sulfate (SO_4) and S_8 , (Fig. 2 and Table 2). Most S was in the form of $\text{FeS}_{2(s)}$ during
386 the last anoxic cycles (e.g. samples R1 33, R2 33). While S_8 mostly deposited in R1 during the
387 anoxic half-cycles, about 11% of S_8 rested in the second oxic half-cycle, and increased to 22% in the
388 last anoxic half-cycle (Table 2). The AVS values indicated of 1.8 and 0.17 $\mu\text{M}\cdot\text{g}^{-1}$ dry sediment
389 during the second anoxic cycles (e.g., samples R1 19, R2 19), respectively (Table 2), while S spectra
390 show that the fitting procedure with $\text{FeS}_{(s)}$ was not obtained a good result.

391 3.3.2 Solid As speciation

392 To determine the effect of redox cycling on solid arsenic speciation, X-ray absorption spectra were
393 recorded at the As K-edge and the bulk As concentration was analyzed using total digestion on
394 sediments sampled at the end of each half-cycle for both reactors, and on the initial sediment. It is
395 worth noting that Linear Combination Fitting XANES analysis is able to detect only species that are
396 present in amounts higher than 5 – 10%, and to distinguish only species with sufficiently different
397 spectral features. Arsenic K-edge XANES data suggested the presence of four distinct As species
398 (Fig. 3, Fig. 4). The initial sediment included O-bound As(V), O-bound As(III), S-bound As(III),
399 and arsenian pyrite (i.e arsenic-substituted pyrite) (Fig. 3). The bulk As concentrations in the same
400 samples were analyzed using total digestion and ICP. The total concentration of As during the oxic
401 conditions were higher than during the anoxic conditions in both reactors (Fig. 4). An increasing As
402 trend in the sediment during the anoxic half-cycles were also observed. The presence of an
403 admixture of O-bound As(III) and O-bound As(V) species in all samples, with an increase in the
404 proportion of As(V) over As(III) during the oxidizing cycles (Fig. 4). In the last oxic cycle, the
405 difference between As(V) and As(III) is quite small: 57% (or 53%) of As(V) and 44% (or 47%) of
406 As(III) (Table 3, Fig. 4). In R1, in the anoxic half cycles, and partly also in oxic half cycles, 8 – 15%
407 of the total As solid phase was attributed to thiol-bound As(III) (Fig. 3, Fig. 4a). In R2, LCF
408 suggested that S-bound As compounds contributed to 8 – 16% of the total As, but they were not
409 found to contribute in the last cycles, both R1 and R2 (Fig. 3; Fig. 4b).

410 **4. Discussion**

411 **4.1 Microbial sulfate reduction and sulfide oxidation**

412 The analytical results of pore water and microbial analysis in both reactors suggest the occurrence of
413 Fe and SO_4^{2-} reduction during the anoxic half-cycles. Sulfate-reducing microorganisms (e.g.,
414 *Desulfobulbus* sp., *Desulfomicrobium* sp. and *Desulfovibrio* sp.) and iron-reducing ones (e.g.,
415 *Geobacter* sp., *Dechloromonas* sp.) were detected, supporting the occurrence of microbial SO_4^{2-} and
416 Fe reduction processes are ongoing. The detected species claimed to be involved in arsenate-
417 respiration in the mobilization of As in Mekong Delta region (Héry et al., 2008, 2015; Lear, Song et
418 al., 2007; Rizoulis et al., 2014). It was surprising that *Geobacter* species were more abundant during
419 oxic half-cycles than anoxic half-cycles. *Geobacter* species were previously considered strict
420 anaerobes (Caccavo et al., 1994; Lin, et al., 2004). However, evidence of the use of oxygen as a
421 terminal electron acceptor by some *Geobacter* species have emerged as has their ability to reduce O_2
422 under microaerophilic conditions (Lin et al., 2004; Parsons et al., 2013). Sulfate-reducing bacteria
423 (SRB), such as *Desulfovibrio* sp., were also observed in oxic half-cycles. *Geobacter* species, can

424 survive oxygen exposure and also contain enzymes to reduce oxygen (Cypionka, 2001).
425 Nonetheless, these findings suggest the persistence of anaerobic micro-environments in which these
426 organisms are able to grow during the nominally oxic phase. It can be hypothesized that the duration
427 of the half-cycles was sufficient to lead to bulk geochemical parameters, indicating oxic conditions
428 but probably not to fully oxidize microenvironments in the reactor.

429 Typically, carbon compounds are incompletely oxidized to acetate by SRB, which may explain the
430 accumulation of acetate under anoxic conditions in R2 and, to lesser extent in R1. Coincidentally,
431 Fe(II) and S(-II) were released repeatedly during three anoxic half-cycles for both high and low
432 sulfate contents. These results seem to indicate an increasing trend in iron and sulfate reduction
433 activities with the increase in sulfate concentration.

434 *Arcobacter* sp. represents 60% of the OTUs in the anoxic half-cycle in R1. There are several
435 possible interpretations for the predominance of this genus. One is that the *Arcobacter* species is a
436 microaerophilic sulfide oxidizer and has grown in the reactor during the anoxic phase, as dissolved
437 oxygen concentrations were decreasing. The time point at day 19, simply might represent the anoxic
438 end point of prolonged microaerobic conditions. Another explanation may be that this particular
439 species is capable of reducing sulfur or other intermediate valence sulfur compounds. The
440 prevalence of *Hydrogenophaga* sp., which are typically aerobic hydrogen oxidizers, during the
441 anoxic half-cycle of R1 reactor would support the first interpretation.

442 Typical aerobic sulfide oxidizing bacteria, such as *Thiobacillus* sp., *Rhodobacter* sp., were also
443 relatively abundant in oxic cycles, particularly in R2. These organisms are typically phototrophic,
444 anaerobic iron oxidizers. However, since the reactors were wrapped in aluminum foil, phototrophy
445 seems unlikely. They are also chemotrophs and it is conceivable that they could oxidize sulfide
446 anaerobically with nitrate as an electron acceptor (Cytryn et al., 2005).

447 **4.2 Mechanism of As mobility during redox oscillation**

448 *4.2.1 Control on As release under oxic condition*

449 During the oxic half-cycles, 92% and 83% of dissolved As was removed in R1 and R2, respectively.
450 Moreover, As K-edge XANES spectra show that O-bound As(V) and As(III), which can sorb onto
451 Fe-(oxyhydr)oxides were the main As solid phase species in R1 and R2 (Fig. 4; Table 2). Other
452 forms of As species, present in the initial sediment (e.g. arsenian pyrite and thiol bound As(III)),
453 were absent in the oxic half-cycles, probably due to their oxidation. The ratio of As(V)/As(III)
454 increased in the first two oxic half-cycles, and remained constant in the last oxic half-cycle. The

455 precipitation of arsenic pentoxide (As_2O_5) may occur during the oxidation process. Although,
456 calculations performed at pH values of 5.5 – 5.7 (SI of arsenic pentoxide (As_2O_5) \geq 4.5), indicated
457 that As_2O_5 may form in both R1 and R2 during the oxic half-cycles, As_2O_5 is relatively easy to
458 dissolve to As(V) in the aqueous phase. Accordingly, As_2O_5 was not detected by As K-edge
459 XANES analysis. Adsorption onto freshly formed Fe-(oxyhydr)oxides is the most likely mechanism
460 to explain its removal from the aqueous phase in nature or through the redox cycling batch
461 experiments (Nickson et al., 2000; van Geen et al., 2003; Root et al., 2007; Couture et al., 2015;
462 Parsons et al., 2013). Altogether, these results suggest that As immobilization during the oxidizing
463 conditions is controlled by As adsorption on Fe-(oxyhydr)oxides.

464 4.2.2 Control on As release under anoxic condition

465 Under anaerobic conditions, microbially driven oxidation of organic matter coupled to the
466 dissimilatory reductive dissolution of As-bearing Fe-(oxyhydr)oxides is commonly accepted to
467 cause the transfer of As from the solid to the aqueous phase (Anawar et al., 2003; Hoque et al.,
468 2009; Horneman et al., 2004; Nickson et al., 2000). As observed in a previous study performed on
469 the same core of sediment collected in An Giang (Vietnam), under the effect of pyrite oxidation,
470 which inhibits the sulfate-reducing bacteria, dissolved As was backfilled cycle after cycle (Phan et
471 al., 2018). In contrast, this study shows that the successive anoxic half-cycles exhibit a decreased
472 concentration of dissolved As (e.g. 92% in R1 and 83% in R2), and that As does not desorb at the
473 end of the experiment, at both different sulfate concentrations (Fig. 1). This demonstrates that
474 microbial SO_4^{2-} reduction led to the sequestration of a portion of aqueous As.

475 The formation of As sulfide precipitations (e.g. orpiment ($\text{As}_2\text{S}_3(\text{s})$) and realgar ($\text{AsS}(\text{s})$)) is one of the
476 possible mechanisms to remove As from the aqueous phase by direct reaction with dissolved S(-II)
477 (Couture et al. 2013a; O'Day et al. 2004). In this study, thermodynamic calculations indicate that
478 $\text{AsS}(\text{s})$, $\text{As}_2\text{S}_3(\text{s})$, and amorphous As sulfide ($\text{As}_2\text{S}_3(\text{s})$) were slightly oversaturated by SI of 0.7, where
479 HS^- is estimated at the maximum aqueous sulfide concentration (e.g. 8 and 5 μM in R1 and R2,
480 respectively), and at slightly acidic to quasi-neutral pH (Fig. 5). In contrast, As K-edge data suggest
481 no evidence for the formation of $\text{AsS}(\text{s})$ and $\text{As}_2\text{S}_3(\text{s})$ during each 7-day anoxic half-cycles. This is
482 probably due to low sulfide concentration and slow kinetics (Burton et al., 2013).

483 Another mechanism is linked to thiols formed during OM sulfurization (Couture et al., 2013a).
484 Recently, Wang et al. (2018) have reported the accumulation of thiol-bound As(III) in an organic-
485 rich sediment of the Mekong Delta in Vietnam. Our finding of S-bound As(III) species in the initial

486 samples are consistent with this. However, over successive redox cycles, As K-edge XANES data
487 suggest that S-bound As(III) gradually decreased and remained negligible in the last anoxic half-
488 cycle in both reactor (Fig. 4). Since the high porewater Fe(II) concentration favorably buffers
489 aqueous S(-II) to very low concentration, it subsequently limits the sulfurization of organic matter
490 (Burton et al., 2014). Consequently, in the successive cycles, the thiol bound As(III) complex
491 formation is unlikely to be a dominant mechanism of aqueous As immobilization.

492 Arsenic may also sorb onto mackinawite ($\text{FeS}_{(s)}$) under the effect of microbial SO_4^{2-} reduction
493 (Burton et al., 2014). The thermodynamic calculation also indicates that $\text{FeS}_{(s)}$ was oversaturated
494 during anoxic half-cycles (Fig. 1e-f). However, the S K-edge XANES data indicated the presence of
495 S(-II) and S^0 , which suggests that the experimental conditions might be favorable for pyrite
496 formation rather than FeS precipitation (Berner, 1970; Raiswell & Berner, 1985; Hurtgen et al.,
497 1999). In agreement, As K-edge results showed no evidence for As(V) or As(III) adsorption onto Fe
498 sulfides during redox cycling experiments. Hence, the As sequestration on $\text{FeS}_{(s)}$ may be marginal.

499 The actual As-sequestration mechanism could be the sorption of As on $\text{FeS}_{2(s)}$, a dominant Fe sulfide
500 detected by S K-edge XANES investigation. The As(III) and As(V) species often sorb more strongly
501 than the thioarsenates onto Fe-(oxyhydr)oxides and $\text{FeS}_{(s)}$, whereas MTA(V) sorbs most strongly on
502 FeS_2 and leads to O-bound As(III), because thiorarsenics donate S atoms to FeS_2 (Couture et al.,
503 2013b). Based on the thermodynamic modeling constants, As species are expected to be dominated
504 by As(III), MTA(V), TTA(V), and TeTA(V) both at low (R1) and high (R2) sulfate contents (Fig.
505 5). Bath experiments revealed a mix of O-bound As(III) and As(V) species, with a ratio of
506 As(III)/As(V) considerably increasing during the reducing half-cycles, and higher O-bound As(III)
507 content than O-bound As(V) one, was observed in the last anoxic half-cycles, probably due to the
508 enrichment of thioarsenate on $\text{FeS}_{2(s)}$ (Fig. 4, Table 3). However, the mechanism of thioarsenate
509 sequestration in the Fe-As-S system requires further investigation to be confirmed, e.g., by
510 performing EXAFS analysis to reveal the coordination of O-bound As(III) on Fe sulfides (e.g.,
511 $\text{FeS}_{(s)}$, $\text{FeS}_{2(s)}$) at low aqueous As concentrations. The fact that during anoxic half-cycles, the
512 proportion of O-bound As(III) increases, and that aqueous As is removed during anoxic cycles could
513 be due to the formation of aqueous thioarsenate sorbed on FeS_2 . This would explain the observed
514 increment of the O-bound As(III) species during the last anoxic half-cycles.

515 **5. Conclusion**

516 This study was aimed to better understand whether the high As concentrations detected in the
517 aquifer in An Giang (Vietnam) can be attributed to redox dynamics. It has been shown that S and Fe
518 biogeochemistry play a key role in determining the fate of As in deltaic sediments during redox
519 oscillations. In particular, the results of this study indicate that: (i) As is first released during Fe
520 reducing conditions but is then sequestered during SO_4^{2-} reduction; (ii) As adsorbs/desorbs on Fe-
521 (oxyhydr)oxides, (iii) aqueous thioarsenic can form and be adsorbed on Fe sulfide minerals (e.g.,
522 FeS_2); and (iv) the reactions leading to the decrease of aqueous As concentrations over subsequent
523 cycles are driven by microbial activity, which induces fermentation during anoxic periods, and
524 respiratory consumption during oxic ones. While previous studies have shown that As mobility is
525 controlled by reductive dissolution of Fe-(oxyhydr)oxides in the reducing cycles, the results of the
526 present work suggests that As sequestration in seasonally saturated sediments in the Mekong Delta
527 Vietnam is controlled by the combined impact of surface hydrology, and by the reaction of Fe
528 sulfide minerals and soluble As compounds (e.g. thioarsenic). This mechanism requires further
529 investigation, e.g. by using techniques able to reveal the coordination of O-bound As(III) species on
530 FeS_2 , such as As K-edge EXAFS analysis.

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542 **Supplementary data**

543 Supplementary data related to this article can be found at Supporting Information part.

544

545

546 **Reference**

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869 **Table Caption**

870 **Table 1.** Analysis of taxa (contribute $\geq 1\%$ of OTUs) corresponding to potential metabolism and
871 oxygen tolerance of selected samples at R1 and R2

872 **Table 2.** Proportion of the different S solid phases in the sediment samples by applying a linear
873 combination fitting procedure (LCF) on XANES spectra at the S K-edge (Fig. 3). The acid
874 volatile sulfide (AVS, $\mu\text{mol/g}$ of dry soil) was calculated from sequential extractions. Goodness
875 of fit is estimated by both the classical R-factor (R_f) and the reduced chi-square ($\text{Red}\chi^2$)
876 indicators.

877 **Table 3.** Proportion of the different As solid phases in the initial sediment sample (R0) and in
878 sediments extracted from bioreactors (R1 and R2) as revealed by applying a linear combination
879 fitting (LCF) performed on the corresponding XANES spectra. The numbers in parenthesis
880 give the error on percentages for each fitting component. Goodness of fit is estimated by both
881 the classical R-factor (R_f) and the reduced chi-square ($\text{Red}\chi^2$) indicators.

882

883 **Figure Caption**

884 **Figure 1.** Aqueous chemistry measured E_h (SHE) and pH (a-b), DOC and acetate (c-d), Fe(tot) and
885 Fe^{2+} (e-f), S(tot) and SO_4^{2-} , HS^- and S_8 in suspension (g-h), As(tot), As(V) and As(III) (i-j)
886 data with time in reactor R1 (0.1 mM SO_4^{2-}) (left) and R2 (1.0 mM SO_4^{2-}) (right). Blue and
887 white shaded areas indicate the anoxic and oxic half-cycles, respectively. Sampling points for S
888 K-edge XANES and microbial community analysis are shown on the E_h curve (As and S K-
889 edge XANES = open purple circles, 16S rRNA = red arrows), cellobiose (OC) adding points =
890 green arrows, and saturation index of porewaters (e-f) with respect to FeS.

891 **Figure 2.** Normalized S K-edge XANES spectra of sediment samples collected from reactors R1
892 (0.1 mM SO_4^{2-}) and R2 (1 mM SO_4^{2-}) at the end of each anoxic or oxic half-cycles, as shown in
893 Fig. 1. The spectra of the samples (open circles) are reported together with the LCF curves
894 superimposed (solid lines). The spectra of the reference compounds used for LCF (K_2SO_4 , S_8
895 and FeS_2) are also shown. All spectra are vertically shifted for clarity.

896 **Figure 3.** The spectra of the reference compounds used for LCF (As(III/V) goethite, $FeAsS$, As_2S_3 ,
897 thiol-bound As(III)) are shown. Normalized As K-edge XANES spectra of sediment samples
898 collected from reactors R1 (0.1 mM SO_4^{2-}) and R2 (1 mM SO_4^{2-}) at the end of each anoxic or
899 oxic half-cycle, as shown in Fig. 1. The spectra of the samples (open circles) are reported
900 together with the LCF curves superimposed (solid lines). All spectra are vertically shifted for
901 clarity.

902 **Figure 4.** Arsenic quantitative speciation in reactors (a) R1 (0.1 mM SO_4^{2-}) and (b) R2 (1 mM SO_4^{2-})
903 at the end of each anoxic or oxic half-cycles. The As speciation is estimated by multiplying
904 total As concentration sediments samples analyzed using the total digestion (the fraction of As
905 as each of four species was obtained by LCF of As K-edge XANES data)

906 **Figure 5.** Thermodynamically aqueous (white region) and solid (brown region) As speciation with
907 R1 in the anoxic (a) and oxic cycle (b) ($\Sigma As = 50 \times 10^{-6}$ and $\Sigma S = 100 \times 10^{-6}$ (M)) and R2 in the
908 anoxic (c) and oxic cycle (d) ($\Sigma As = 50 \times 10^{-6}$ and $\Sigma S = 1000 \times 10^{-6}$ (M)) using PHREEQC
909 code. The blue vertical areas represent pH values of the cycling experiments.

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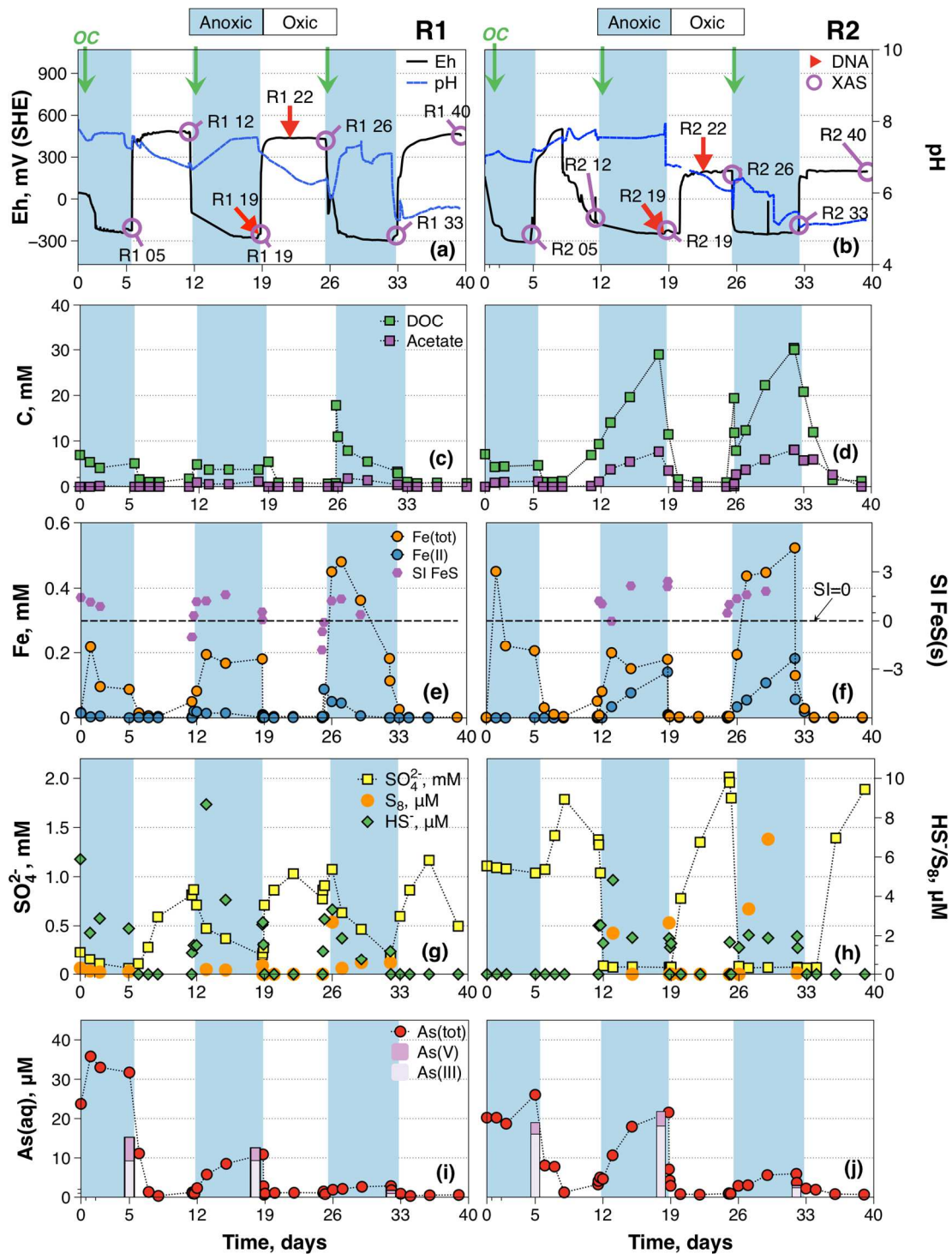


Figure 1. Aqueous chemistry measured E_h (SHE) and pH (a-b), DOC and acetate (c-d), Fe(tot) and Fe²⁺ (e-f), S(tot) and SO₄²⁻, HS⁻ and S₈ in suspension (g-h), As(tot), As(V) and As(III) (i-j) data with time in reactor R1 (0.1 mM SO₄²⁻) (left) and R2 (1.0 mM SO₄²⁻) (right). Blue and white shaded areas indicate the anoxic and oxic half-cycles, respectively. Sampling points for S K-edge XANES and microbial community analysis are shown on the E_h curve (As and S K-edge XANES = open purple circles, 16S rRNA = red arrows), cellobiose (OC) adding points = green arrows, and saturation index of porewaters (e-f) with respect to FeS.

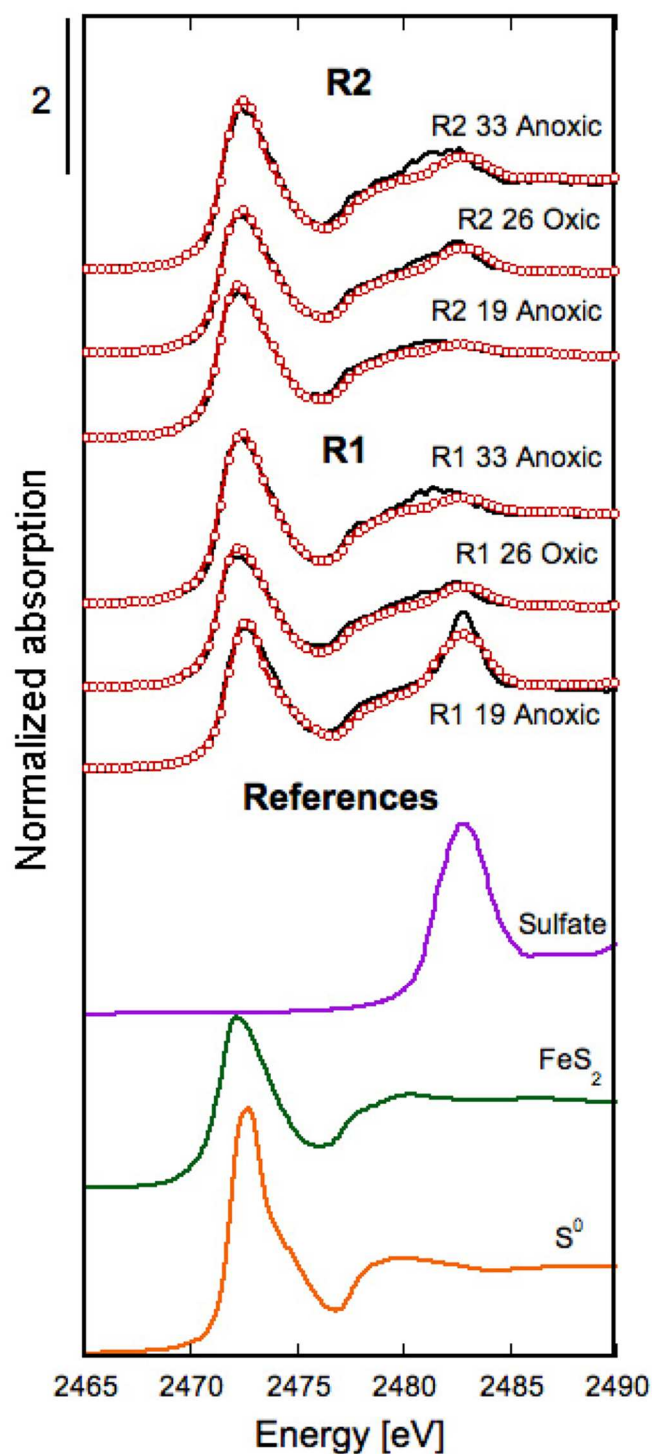


Figure 2. Normalized S K-edge XANES spectra of sediment samples collected from reactors R1 (0.1 mM SO_4^{2-}) and R2 (1 mM SO_4^{2-}) at the end of each anoxic or oxic half-cycles, as shown in Fig. 1. The spectra of the samples (open circles) are reported together with the LCF curves superimposed (solid lines). The spectra of the reference compounds used for LCF (SO_4 , S_8 and FeS_2) are also shown. All spectra are vertically shifted for clarity.

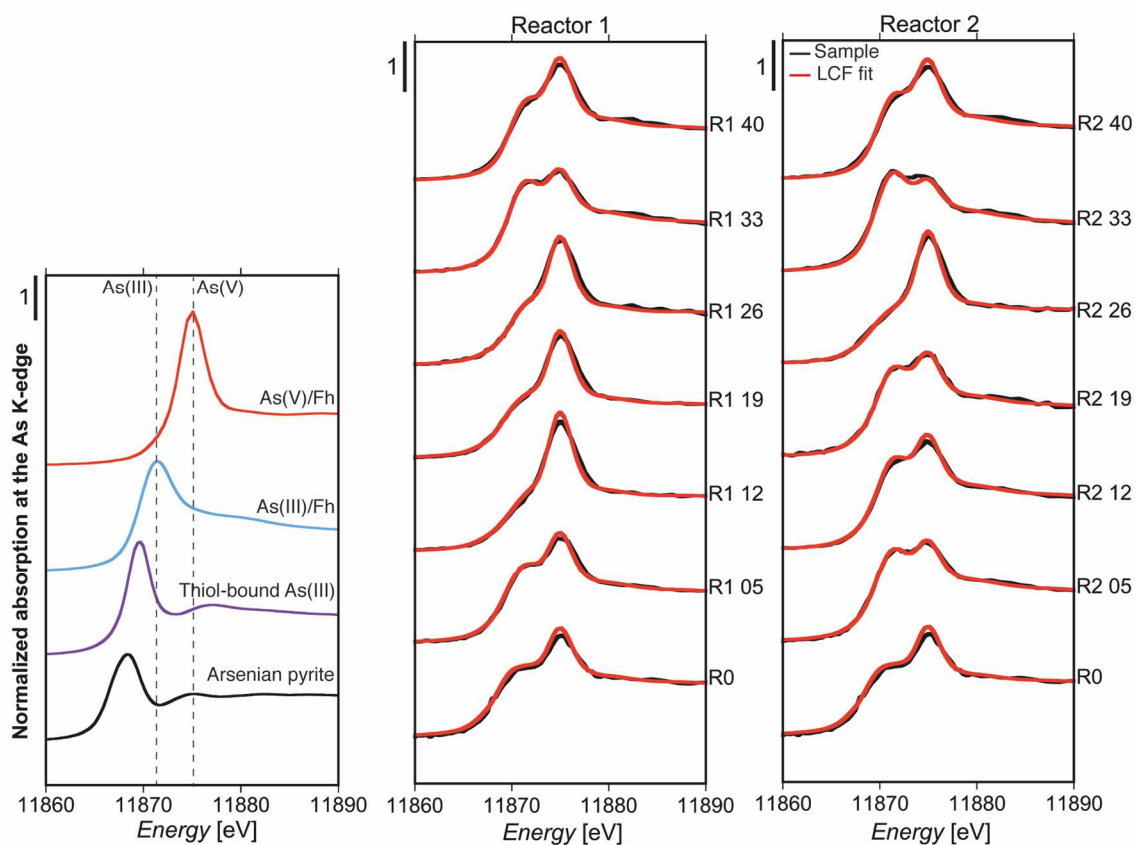


Figure 3. The spectra of the reference compounds used for LCF (As(III/V) goethite, FeAsS, As₂S₃, thiol-bound As(III)) are shown in the left panel. Center and right panels: Normalized As K-edge XANES spectra of sediment samples collected from reactors R1 (0.1 mM SO₄²⁻) and R2 (1 mM SO₄²⁻) at the end of each anoxic or oxic half cycle, as shown in Fig. 1. The spectra of the samples (open circles) are reported together with the LCF curves superimposed (solid lines). All spectra are vertically shifted for clarity.

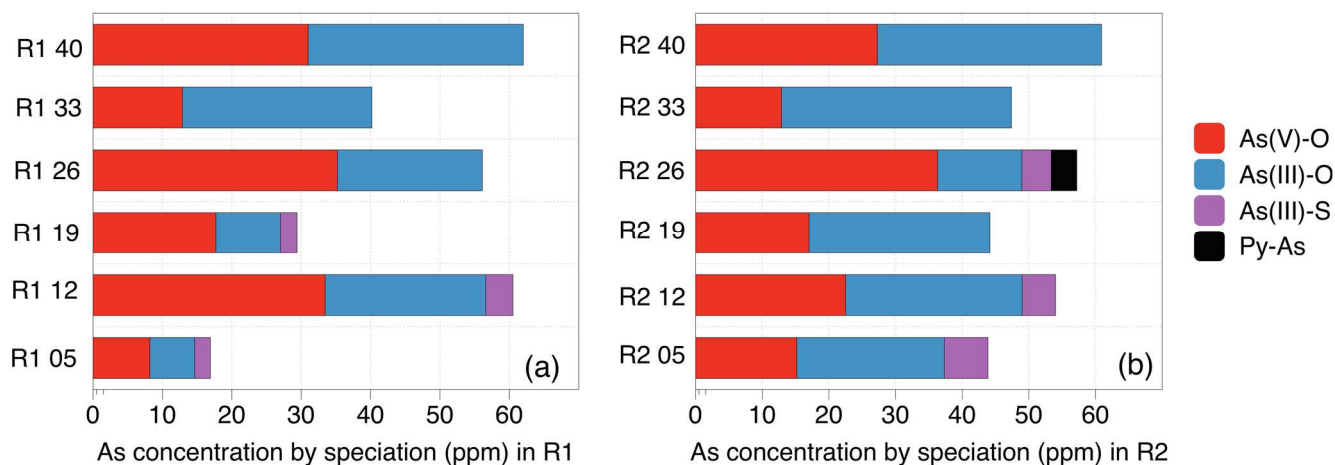


Figure 4. Solid arsenic quantitative speciation in reactors (a) R1 (0.1 mM SO_4^{2-}) and (b) R2 (1 mM SO_4^{2-}) at the end of each anoxic or oxic half-cycles. The solid As speciation is estimated by multiplying total As concentration sediments samples analyzed using the total digestion (the fraction of As as each of four species was obtained by LCF of As K-edge XANES data)

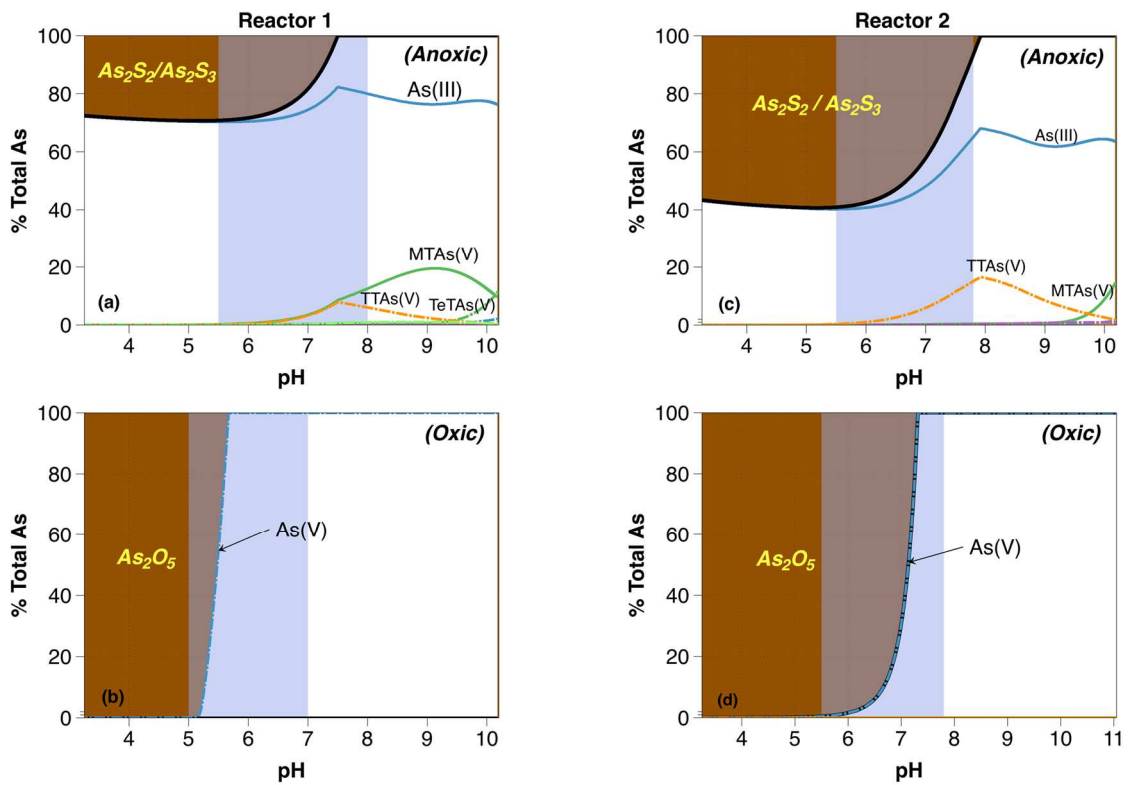


Figure 5. Thermodynamically aqueous (white region) and solid (brown region) As speciation with R1 in the anoxic (a) and oxic cycle (b) ($\Sigma\text{As} = 50 \times 10^{-6}$ and $\Sigma\text{S} = 100 \times 10^{-6}$ (M) and R2 in the anoxic (c) and oxic cycle (d) ($\Sigma\text{As} = 50 \times 10^{-6}$ and $\Sigma\text{S} = 1000 \times 10^{-6}$ (M) using PHREEQC code. The blue vertical areas represent pH values of anoxic/oxic half-cycles.

Table 1. Analysis of taxa (contribute $\geq 1\%$ of OTUs) corresponding to potential metabolism and oxygen tolerance of selected samples at R1 and R2

Species	% OTUs				Potential metabolism	O2 tolerance	Reference
	R1-19 Anoxic	R2-19 Anoxic	R1-22 Oxic	R2-22 Oxic			
<i>Arcobacter sp.</i>	60.487	2.931	0.862	1.077	Sulfide oxidizing	Microaerophilic, autotrophic	Wirsen et al., 2002
<i>Klebsiella sp.</i>	6.381	10.511	3.704	2.735	Fermentation	Microaerophilic–aerobic	Franciscon et al., 2009
<i>Rhizobium sp.</i>	2.804	3.328	2.725	1.668	Nitrogen fixing	Microaerophilic	Schmeisser et al., 2009
<i>Geobacter bremensis</i>	2.490	5.109	2.690	5.674	Iron and sulfur reduction	Aerotolerant	Snoeyenbos-West et al., 2000
<i>Clostridium sp.</i>	2.158	2.435	3.253	3.295	Iron and sulfur reduction	Microaerophilic	Dobbin et al., 1999; Li et al., 2011; Thabet et al., 2004
<i>Acidovorax delafieldii</i>	2.147	0.405	1.735	0.507	Definication	Aerobic	Willems et al., 2015
<i>Hydrogenophaga taeniospiralis</i>	2.020	3.095	7.484	5.679	Hydrogen oxidizing	Microaerophilic–aerobic	Willems et al., 1989
<i>Acinetobacter sp.</i>	1.879	0.124	2.637	0.847	Methane oxidation	Aerobic	Wang et al., 2012
<i>Hydrogenophaga pseudoflava</i>	1.624	8.728	1.582	1.255	Hydrogen oxidation, thiosulfate oxidation	Microaerophilic–aerobic	Willems et al., 1989; Graff et al., 2003
<i>Thiobacillus thioparus</i>	0.466	2.388	0.211	0.251	Iron and sulfide oxidation	Microaerophilic–aerobic	Hedrich & Johnson, 2011
<i>Candidatus Accumulibacter sp.</i>	0.788	1.690	4.689	2.986	Iron oxidizing	microaerophilic–aerobic	Hedrich et al., 2011
<i>Geobacter sp.</i>	0.237	1.628	3.311	19.114	Iron and sulfur reduction	Microaerophilic	Snoeyenbos-West et al., 2000, Lin et al., 2004
<i>Rhodobacter sp.</i>	0.095	1.447	1.371	3.263	Iron oxidation	Microaerophilic–aerobic	Poulain and Newman, 2009
<i>Thiobacillus sp.</i>	0.120	1.083	0.064	0.178	Iron and sulfide oxidation	Microaerophilic–aerobic	Poulain and Newman, 2009
<i>Gemmobacter sp.</i>	0.343	1.076	0.955	0.847	Carbon and nitrogen degradation	Facultative anaerobic	Kumaresan et al., 2015
<i>Flavobacterium sp.</i>	0.011	1.026	0.000	0.000	Decompose several	Aerobic	Schellenberger et al., 2011

					polysaccharides		
<i>Rhodobacter capsulatus</i>	0.198	0.745	15.619	10.987	Fe(II) oxidation	Microaerophilic–aerobic	Hedrich & Johnson, 2012
<i>Dechloromonas hortensis</i>	0.304	0.793	2.837	1.072	Organic matter degradation	Aerobic	Wolterink et al., 2005
<i>Vogesella indigofera</i>	0.035	0.064	1.993	1.229	Organic matter degradation	Aerobic	Luan & Medzhitov, 2016
<i>Sulfurospirillum sp.</i>	0.735	0.050	1.178	1.914	Sulfur reduction	Microaerophilic	Tang et al., 2009
<i>Desulfobulbus sp.</i>	0.286	0.600	0.615	1.537	Sulfate reducing	Anaerobic	Muyzer & Stams, 2008; Pokorna & Zabranska, 2015
<i>Desulfomicrobium sp.</i>	0.102	0.231	0.820	1.517	Sulfate reducing	Anaerobic	Muyzer & Stams, 2008; Pokorna & Zabranska, 2015
<i>Prolixibacter sp.</i>	0.512	0.969	0.399	1.412	Fermentation	Microaerophilic	Holmes et al., 2007
<i>Novosphingobium sp.</i>	0.441	0.857	0.586	1.124	Organic matter degradation	Microaerophilic	Nguyen et al., 2014

Table 2. Solid-phase Sulfur speciation. Proportion of the different S solid phases in the sediment samples by applying a linear combination fitting procedure (LCF) on XANES spectra at the S K-edge (Fig. 2). The error on percentages is estimated to be in the range between 5 and 15%. The acid volatile sulfide (AVS, $\mu\text{mol/g}$ of dry soil) was calculated from sequential extractions. Goodness of fits is estimated by the reduced chi-square ($\text{Red}\chi^2$) values.

Sample	AVS ($\mu\text{mol/g}$)	LCF (%)			Red- χ^2 ($\times 10^2$)
		K ₂ SO ₄	S ₈	FeS ₂	
R1 19	1.817	35	38	27	3.9
R1 26	-	15	-	85	1.8
R1 33	-	10	13	77	2.7
R2 19	0.167	8	-	92	1.2
R2 26	-	19	11	70	1.5
R2 33	-	16	22	63	3.9

	Anoxic
	Oxic

Table 3. Solid phase Arsenic speciation. Proportion of the different As solid phases in the initial sediment sample (R0) and in sediments extracted from bioreactors (R1 and R2) as revealed by applying a linear combination fitting (LCF) performed on the corresponding XANES spectra. The error on percentages is estimated to be in the range between 5 and 10%. Goodness of fits can be estimated by the reduced chi-square ($\text{Red}\chi^2$) values.

Sample	LCF (%)				Red- χ^2 (x 10 ³)
	As(V)-O	As(III)-O	As(III)-S	Py-As	
R0	40	34	15	14	1.5
R1 05	43	47	12	-	1.0
R1 12	68	27	8	-	1.8
R1 19	59	31	8	6	1.1
R1 26	64	38	-	-	1.9
R1 33	33	70	-	-	0.7
R1 40	50	53	-	-	1.1
R2 05	37	54	16	-	0.7
R2 12	45	53	10	-	1.0
R2 19	37	59	-	-	0.9
R2 26	66	23	8	7	0.5
R2 33	28	75	-	-	0.7
R2 40	47	58	-	-	0.8

	Anoxic
	Oxic

