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Sulfur and oxygen stable isotope tracing in Zero valent iron based Insitu remediation system for metal contaminants

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Abstract:

In the present study, controlled laboratory column experiments were conducted to understand the biogeochemical changes during the microbial sulfate reduction. Sulfur and oxygen isotopes of sulfate were followed during sulfate reduction in zero valent iron incubated flow through columns at a constant temperature of 20 ± 1°C for 90 days. Sulfur isotope signatures show considerable variation during biological sulfate reduction in our columns in comparison to abiotic columns where no changes were observed. The magnitude of the enrichment in $\delta^{34}$S values (‰) ranged from +9.4 to +10.3‰ compared to initial value of 2.3‰, having total fractionation $\Delta S$ between biotic and abiotic columns as much as 6.1‰. Sulfur isotope fractionation was directly proportional to the sulfate reduction rates in the columns. Oxygen isotopes in this experiment seem less sensitive to microbial activities and more likely to be influenced by isotopic exchange with ambient water. A linear relationship is observed between $\delta^{34}$S and $\delta^{18}$O in biotic conditions and we also highlight a good relationship between $\delta^{34}$S and sulfate reduction rate in biotic columns.

Keywords: Sulfur isotopes, oxygen isotopes, zero valent iron, in-situ remediation
1. Introduction

In-situ groundwater remediation of metal contaminants has emerged as a sustainable option in recent years for various reasons, e.g. economic feasibility and energy consumption in pump and treats methods, site accessibility etc… (Farhadian et al., 2008). For in-situ remediation, various possibilities have also been explored for instance by enhancing natural attenuation, providing electron donors (Satyawali et al., 2010) or by using reactive barrier materials (Benner et al., 1999; Waybrant et al., 2002). Zero valent iron (Fe°) is getting large attention lately as a reactive material for in-situ applications (Dries et al., 2005; Burghardt and Kassahun, 2005; Wilkin and McNeil, 2003). The highly reducing nature and relatively larger available specific surface area make Fe° a suitable medium for groundwater contaminant removal. There have been already many successful installation of Fe° based remediation system in last decade (Phillips et al., 2010; Liyuan Liang, 2003; Gu et al., 1998; Rowland, 2002). Fe° has been successfully used in lab scale as well as field scale applications, dealing with wide range of groundwater contaminants e.g. chlorinated compounds and metals (Dries et al., 2005; Gandhi et al., 2002; Rangsivek and Jekel, 2005), radioactive material (Burghardt and Kassahun, 2005), arsenic (Su and Puls, 2003; Nikolaos et al., 2003) etc... However, the biogeochemical dynamics and contaminants behavior in subsurface environments are still poorly understood, as real field sites often encounter problems due to little or no control over fluxes and changes in subsurface processes with time or seasons. Regular chemical analysis and monitoring may not be a practical or economic option in many isolated sites. These complications often make it difficult to understand the actual processes (biotic/abiotic) contributing in contaminant removal, needless to say that this distinction is rather very important in designing realistic remediation strategy for any particular site.

Stable isotopes have emerged as a potential tool in understanding the dynamics of pollutants in water systems (Fritz et al., 1994; Houhou et al., 2010; Krouse, 2000; Mast et al., 2001; Wu et al., 2011). Relatively easy analysis and bulk information makes isotope study a practical and viable option. The characterization and quantification of electron-accepting processes, like nitrate, iron and sulfate reduction are extremely valuable in estimating the sustainability and longevity of degradation process in any subsurface system (Knöller et al., 2006). This characterization is of primary importance in case of in-situ treatment process, where there is little control over changing processes in subsurface environments. Biological sulfate reduction has been observed
multiple times in similar kind Fe° based treatments systems (Van Nooten et al., 2007; Gu et al., 2002).

Quantification of sulfate reduction by following dissolved concentrations of sulfate and co-existing sulfide in groundwater is often a challenge due to possible dilution, mixing or mineral precipitation processes. Precipitation of dissolved sulfide is particularly important in case of Fe° based systems where abundant Fe\(^{2+}\) is available in groundwater for possible FeS precipitation. It has been shown that sulfate reducing bacteria preferentially removes lighter isotopes of sulfur and oxygen from sulfate molecule resulting into isotopic enrichments of both heavier isotopes i.e. \(^{18}\)O and \(^{34}\)S in residual sulfate (Fritz et al., 1989).

In this study, we followed the evolution of \(\delta^{34}\)S and \(\delta^{18}\)O from dissolved sulfate in groundwater within lab scale Fe° based flow through columns, which were primarily designed for heavy metal remediation from groundwater. Aim of this study was to characterize the isotope fractionation and changes during the long-term remediation processes and also to determine the impact of microbiology on isotopic signature in subsurface redox environment. We summarize here the results of sulfur and oxygen isotope fractionation in Fe° based lab scale in-situ remediation systems over 90 days of experiment.

2. Material and Methods

2.1. Column design

Four double jacketed flow through glass columns (30 cm length x 4.5 cm I.D, total liquid volume \(\approx 480 \text{ mL}\)) were setup in laboratory for \(\sim 30\) weeks at controlled temperature \((20 \pm 1{\degree}\text{C})\), with the primary aim of groundwater contaminant (metals) removal using Fe° as a reactive material. Columns were filled with sediment obtained from a heavy metal contaminated site in Belgium from a depth of \(\approx 32\) m, more description about this site is given elsewhere (Vanbroekhoven et al., 2008). Efforts were made to design a lab scale concept of in-situ reactive barrier using two types of Fe° differing in particles size and source, i.e. granular zero valent iron (gFe°, Gotthard Maier, Germany) and micro zero valent iron (mFe°, Högenäs, Sweden) with an average particle size of 0.25-2 mm and 20-40 µm respectively. For each column, the first bottom half (\(\sim 240 \text{ mL}\)) was filled with an aquifer/ Fe° mixture with ratios of 80:20 and 98:2 v/v for gFe° and mFe°
respectively. The second (upper) half of all columns was filled only with aquifer (Figure 1).

Filling of columns was performed under nitrogen atmosphere in a glove box. Simulated groundwater, which was prepared in lab corresponding to the site characteristics (Table 1), was injected in parallel through the columns using a peristaltic pump at a constant flow rate of 1 ± 0.2 mL/h. A slight over pressure (0.01 Bar) of N\textsubscript{2} was maintained in the feeding bottles to avoid air contact during column feeding. All tubes and fittings used in the experiment were acid washed and flushed with nitrogen before use.

For each Fe\textsuperscript{0} type, two columns were set-up, of which one was fed with a small dose of glycerol (0.1% v/v of inlet water) to enhance indigenous microbial activity, and the column was exposed to gamma radiations (IONISOS, Dagneux, France), with minimum absorbed radiation dose of 25 kGy, before injecting groundwater to restrict all microbial activities.

### 2.2. Analytical methods

For observation and counting of bacterial cells, defined volumes of samples (from 50 to 1000 µL) were extracted from the columns using a nitrogen filled plastic syringe, by injecting the nitrogen and extracting equal amount of liquid from column. Samples were immediately diluted in deionized water and filtered onto a black polycarbonate filter, 0.22 µm (Nuclepore, Whatman). The filter was incubated 15 min in the dark with 1 mL filtered deionized water mixed with 1 µL DAPI (4',6-diamidino-2-phenylindole) solution (1 mg/mL, SIGMA). This mixture was removed by filtration, and the filter was rinsed 2 times with 1 mL filtered deionized water. The filter was then mounted on a glass slide with Citifluor (Biovailey), and observed with an optical microscope (Zeiss Axio Imager Z1) equipped with Filter Set 49 for DAPI, UV HBO lamp and a digital camera. Bacteria were enumerated on 10 independent fields (each of 5800 µm\textsuperscript{2}). Cell counts were calculated considering the volume of the sample used and filter surface area calculations on an average basis. Sulfate concentration was analyzed with a spectrophotometer operating at \( \lambda \) 540° using specific analysis kits (Merck Spectroquant\textsuperscript{®} kit 1.14548.001, Germany).

Samples for sulfur and oxygen isotope analysis were collected at the outlet of columns using 250 mL pre acid washed plastic perplex bottles. Cd-Acetate was already added in the bottles (5% v/v) prior to sample collection, to fix sulfur as CdS, and then an aliquot was filtered through a 0.2 µm-nitrocellulose filter before chemical determination of residual sulfate. The amount of sample
collected varied at different time points during the experiment as the sulfate concentration in the outlet solution changed over the time. However, in any case, a minimum of 5 mg of SO$_4$ was collected for every sampling point. The analysis was performed as described by (Fritz et al., 1989).

Dissolved sulfate was precipitated as BaSO$_4$ at pH $< 4$ (in order to remove HCO$_3^-$ and CO$_3^{2-}$ species) by adding a BaCl$_2$ solution. The isotopic analyses on BaSO$_4$ were carried out using a Delta+XP mass spectrometer coupled in continuous-flow mode to a Thermo Elemental Analyzer. Sulfate-isotope compositions are reported in the usual δ-scale in ‰ with reference to V-CDT (Vienna Canyon Diablo Troilite) and V-SMOW (Vienna Standard Mean Ocean Water) according to $\delta_{\text{sample}}$ (‰) = {(R$_{\text{sample}}$/R$_{\text{standard}}$) - 1} x 1000, where R is the $^{34}\text{S}/^{32}\text{S}$ and $^{18}\text{O}/^{16}\text{O}$ the atomic ratios. Sulfate-isotope compositions ($\delta^{34}\text{S} (\text{SO}_4)$ and $\delta^{18}\text{O} (\text{SO}_4)$) were measured with a precision of ± 0.3‰ vs. CDT for $\delta^{34}\text{S} (\text{SO}_4)$ and ±0.8 ‰ vs. VSMOW for $\delta^{18}\text{O} (\text{SO}_4)$ respectively.

### 3. Results and Discussion

#### 3.1. Sulfate

Sulfate reduction is a common phenomenon observed in Fe$^0$ based PRB’s due to favorable growth environment i.e. close to neutral pH and a very low ORP (Oxidation Reduction Potential) conditions (Gu et al., 2002). The efficiency of Fe$^0$ in increasing pH and decreasing ORP by releasing OH$^-$ ions and scavenging O$_2$ (see reaction 1) and production of water born H$_2$, is well known (Johnson et al., 2008). This water born H$_2$ can also be a potential electron donor for sulfate reducing bacteria (Karri et al., 2005). In the present study, dissolved sulfate concentration in the column outlet solution was analyzed regularly to follow behavior and activity of sulfate reducing bacteria.

$$\text{Fe}^0 + 2 \text{H}_2\text{O} \rightarrow \text{Fe}^{2+} + \text{H}_2 + 2 \text{OH}^- \quad (1)$$

Dissolved sulfate concentrations in the column outlet solution decreased from initial inlet concentrations of 3.95 mmol/L to 0.03 and 0.09 mmol/L in gFe$^0$ and mFe$^0$ biotic columns respectively (Table 2, Figure 2). Appearance of black patches (believed to be FeS precipitation)
in both biotic columns were in agreement with microbial sulfate reduction. Providing glycerol probably also stimulated bacterial growth by acting as carbon source along with availability of H\textsubscript{2}. In this experiment no extra microbial culture was added, so only natural population of sulfate reducers were expected to grow.

However, in the abiotic columns, sulfate concentrations were unaffected throughout the experiment (Figure 2), which was expected due to absence of viable microbial cells after gamma radiation exposure to sediment and the small dose of formaldehyde (100µL/L) that was added with inlet water to avoid any possibility of microbial growth.

Potential sulfate reduction rates (SRR, nmol.cm\textsuperscript{-3}.h\textsuperscript{-1}) were calculated using equation (2) described by Stam et al., 2011.

\[
SRR = \Delta C \times Q / V \tag{2}
\]

Where Q represents the volumetric flow rate of solution through the reactor, \(\Delta C\) is the difference between inflow (\(C_0\)) and outflow (C) sulfate concentration and V is the volume of the sediment contained in the reactor.

Sulfate reduction rate (SRR) values in this experiment are consistent with the sulfate reduction as the maximum rate was achieved after 30 days of incubation, after which similar values of SRR were obtained (value of 4 - 4.1 nmol.cm\textsuperscript{-3}.h\textsuperscript{-1}, Figure 3a) for both granular and micro biotic columns. Equal and steady values of SRR throughout the experiment are consistent with the equal and limited supply of sulfate in the system. In general the rate of sulfate reduction is also believed to influence the isotope fractionation, and in this study sulfate reduction rate was directly related to the sulfur isotope fractionation (Figure 3b). It is also argued in literature that substrate type may also influence the fractionation, considering H\textsubscript{2} gas producing lower fractionation than organic substrate (Kemp and Thode, 1968; Rittenberg., 1964). As in the present study H\textsubscript{2} as well as glycerol were available as electron donor, individual contribution cannot be established in this experimental setup.

### 3.2. Cell counting and microscopic observations

Bacterial cell counting was performed using DAPI imaging. Changes in cell numbers during the experiment can give general trends of microbial activity; increases or decreases in bacterial cell
count can be related to the overall sulfate reduction rate. Normally, if the sulfate is not limiting and energy source is available, the cell count could increase with time which would further enhance the sulfate reduction.

Photographs of DAPI-stained samples (Figure 4) show the presence of dominant rod shaped bacterial cells with an average length of 5 µm. All cells were visually similar in shape and sizes, suggesting the growth of a dominant species. No efforts were made at this step of the experiment to identify the bacterial species.

Bacterial cells were counted at day 30 of column test, when dissolved sulfate was completely reduced in biotic columns. An average cell count of 1.9x10^7 and 1.2x10^7 cells/mL were obtained in gFe° biotic and mFe° biotic column, respectively. At the end of the experiment (day 90), the cell concentrations were 1.7x10^7 and 6.5x10^7 cell/mL in gFe° and mFe° biotic column, respectively. These observations suggest that the cell concentration during the experimental period did not change significantly, which is consistent with limited and uniform supply of sulfate and almost constant SRR in the columns. These observations also suggest that due to the limited sulfate doses, the sulfate-reducing bacterial population would continue to consume sulfate in the columns, so the cell concentrations are more likely to be related to the dissolved sulfate supply in this case.

3.3. δ³⁴S variations and origin of sulfur isotope fractionation

Although isotopes of the same element behave the same physically and chemically, reaction rates differ due to the mass difference between the isotopes. This mass difference causes a preferential partitioning, namely isotope fractionation, that results in varying isotopic compositions during reaction. On the one hand, in the biotic columns δ³⁴S enrichment was evident (Figure 5), explicable by a preferred use of lighter sulfur element of SO₄ (³²S) by sulfate reducers, which results into abundance of heavier element ³⁴S in the remaining sulfate molecules. On the other hand, no sulfate reducers were active in the abiotic columns, so δ³⁴S values remain almost the same as the initial value (Figure 5). There is no mechanism reported till date for abiotic sulfate reduction.

A large range of variation in δ³⁴S (from 4 up to 46‰) has been observed in pure bacterial cultures where sulfate was available abundantly (Kaplan and Rittenberg, 1964; Detmers
In the present study, the maximum values of $\delta^{34}S$ observed equal +10.3‰ in the mFe° biotic columns, with final enrichment values of +9.4‰. A total $\Delta S$ was as much as +6.1‰, while compared between biotic and abiotic processes, where $\Delta S$ is the difference of $\delta^{34}S$ (‰) between biotic and abiotic columns. The isotope measurement were followed until 90 days of column operation, after this period SRR was constant so no further fractionation was expected; also sample collection was not possible as the dissolved sulfate concentration was nearly zero.

Sulfide produced by sulfate reducing bacteria was not considered to be associated with isotope fractionation in this study as sulfide was very likely to be precipitated with Fe(II) available in the plume to form insoluble iron monosulfide and this phenomenon is known to not be associated with sulfur isotope fractionation effect (Canfield et al., 1992).

Figure 6 shows that the dissolved sulfate concentration and $\delta^{34}S$ (‰) follow an inverse trend in both biotic columns, where dissolved sulfate concentration decreases as microbial activity increases, on the other hand $\delta^{34}S$ (‰) value increases through time. This observation confirms that the sulfur isotope fractionation only originates from microbial sulfate reduction avoiding any contribution of sulfate from sediment.

3.4. $\delta^{18}O_{(SO_4)}$ variations and origin of the oxygen isotope fractionation

Although significant enrichment of $^{34}S$ was evident in residual sulfate, a corresponding enrichment of $^{18}O$ was not observed in this study. Similar observations were previously reported by other researchers (Spence et al., 2005). Sulfur and oxygen isotope followed a different isotopic pattern during sulfate reduction probably because of fundamental differences in the enrichment mechanism. Sulfur isotope fractionation is a purely kinetic effect, whereas oxygen isotope fractionation is influenced by catalysis of isotopic exchange between water and sulfate during sulfate reduction (Fritz et al., 1989., Freney and Ivanov, 1983).

AN identical pattern shift was observed in $\delta^{18}O$ behavior in both biotic and abiotic processes (Figure 7), so it is very likely that the relatively small variation observed in $\delta^{18}O$ could be due to shift in isotopic equilibrium with ambient water. This leads to the widely believed assumption that the oxygen isotope exchange dominated over kinetic isotope fractionation. $\delta^{18}O$ of ambient water was not analyzed in this experiment, so we were not able to confirm this hypothesis. Direct chemical or microbial oxidation of H₂S to sulfur is accompanied by much smaller oxygen isotope effect. The $\delta^{18}O$ fractionation is normally controlled by ambient surface water. This
hypothesis is considered in many studies recently, however no established explanation of this process has been reported yet in literature (Knöller et al., 2006).

3.5. Isotope variation: $\delta^{18}$O vs. $\delta^{34}$S

In Figure 8, linear relationships were observed between $\delta^{34}$S and $\delta^{18}$O in the biotic experiments irrespective of the type of Fe° (either granular or micro Fe), although the slopes are slightly different for each in Figure 8. These results suggest that the particle size may affect the overall sulfate reduction process in some cases.

It is also argued in literature that there is no simple relationship between SRR and isotope fractionation (Detmers et al., 2001), however in the present study we observed a linear relationship (Fig 8b). Brunner et al. (2005) proposed a combined investigation of the influence of sulfate reducing bacteria on the sulfur and oxygen isotopic composition of residual sulfate could be the key to a better understanding of sulfate reduction rate. Böttcher et al. (1998) hypothesized that $\delta^{34}$S$_{SO4}$ vs $\delta^{18}$O$_{SO4}$ relationships reflect sulfate reduction rates in marine sediments; the steeper the slope the slower the sulfate reduction rates. In the present study, we observed not so steep slope for $\delta^{34}$S$_{SO4}$ vs $\delta^{18}$O$_{SO4}$ relationship (Fig 8), which is consistent with the experimental condition where SRR was stable after initial increase.

It is also noteworthy that the $\delta^{34}$S$_{SO4}$ vs $\delta^{18}$O$_{SO4}$ relationship does not reflect the bulk SRR but rather cell-specific SRR. So basically, a large number of bacteria with slow cell specific SRR or a small number of bacteria with high cell specific SRR, both can achieve a high bulk SRR (Brunner et al., 2005). But in present study, the number of bacterial cell in both the biotic columns did not increase significantly during the experimental period.

4. Conclusions

In the present study, we report results from a long-term experiment, designed for groundwater treatment using real site sediment. Sulfur isotope analysis is a good and practically viable option to characterize sulfate reduction activities and sulfate reduction rate in any subsurface system without going for microbial analysis and characterization. Oxygen isotope analysis is also
important but needs to be considered in light of ambient water oxygen isotope exchange, as it is more likely to be controlled by ambient water. Sulfur and oxygen isotopes both provide important information and supports the actual practical data obtained during the experiment. However, it is important to see the long term effect of isotope behavior as very less knowledge is available in literature for these kinds of system for long time operations, which is very typical for in-situ remediation treatment systems.

Isotope geochemistry can also be used for precise identification of pollution sources, effectiveness of remediation process, and can provide crucial insight into contaminants fate and transport (examples for CAH removal are). Considering the simple methods for isotopic analysis and less analytical cost, these standard analytical tools can easily be incorporated into typical field sampling. As the standard geochemical analysis always retain certain level of uncertainty at times important question might left unanswered. In combination with other chemical and biogeochemical techniques, isotopic analysis can be used for better understanding the processes.
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References


Table and Figure Caption

Table 1
Groundwater characteristics.

Table 2
Dissolved sulfate (nmol/L), sulfur ($\delta^{34}S$, ‰) and oxygen isotope ($\delta^{18}O$, ‰) values in the columns with time (days).

Figure 1
Column setup (left schematic diagram and right a picture of column setup in the laboratory).

Figure 2
Sulfate concentration (mmol/L) as a function of time (days) in column outlets.

Figure 3
(a) Sulfate Reduction Rate (SRR, nmol.cm$^{-3}$.h$^{-1}$) in biotic columns as a function of time (days).
(b) sulfur isotope values ($\delta^{34}S$, ‰) as a function of Sulfate Reduction Rate (SRR, nmol.cm$^{-3}$.h$^{-1}$)
in biotic columns.

Figure 4
Photographs of DAPI staining (a) gFe$^\circ$ biotic column and (b) mFe$^\circ$ biotic column.

Figure 5
Sulfur isotope variation ($\delta^{34}S$, ‰) as a function of time (days) in biotic (red) and abiotic columns (blue).

**Figure 6**

Comparison of dissolved sulfate (mmol/L) and $\delta^{34}S$(‰) as a function of time for both biotic columns (micro mFe° and granular gFe°).

**Figure 7**

Oxygen isotope ($\delta^{18}O$, ‰) values in columns as a function of time (days).

**Figure 8**

$\delta^{34}S$ (‰) vs. $\delta^{18}O$ (‰) in sulfate for biotic columns.