# UC Berkeley UC Berkeley Previously Published Works

# Title

Bicarbonate impact on U(VI) bioreduction in a shallow alluvial aquifer

# Permalink

https://escholarship.org/uc/item/1nh6g1zn

# **Authors**

Long, Philip E Williams, Kenneth H Davis, James A <u>et al.</u>

# **Publication Date**

2015-02-01

# DOI

10.1016/j.gca.2014.11.013

Peer reviewed

# Bicarbonate impact on U(VI) bioreduction in a shallow alluvial aquifer

Philip E.Long<sup>a</sup>, Kenneth H.Williams<sup>a</sup>, James A.Davis<sup>a</sup>, Patricia M.Fox<sup>a</sup>Michael J.Wilkins<sup>b,a</sup>, Steven B.Yabusaki<sup>a</sup>, Yilin Fang<sup>b</sup>, Scott R.Waichler<sup>a</sup>, Elena S.F.Berman<sup>c</sup>, ManishGupta<sup>c</sup>, Darrell P.Chandler<sup>d</sup>, ChrisMurray<sup>b</sup>, Aaron D.Peacock<sup>a</sup>, Ludovic Giloteaux<sup>1</sup>, Kim M.Handley<sup>a<sup>2</sup></sup>, Derek R.Lovley<sup>a</sup>, Jillian F.Banfield<sup>a</sup> <sup>a</sup> Lawrence Berkeley National Laboratory, Berkeley, CA 94720, USA <sup>b</sup> Pacific Northwest National Laboratory, Richland, WA 99352, USA <sup>c</sup> Los Gatos Research, Mountain View, CA 94041-1529, USA <sup>d</sup> Akonni Biosystems, Inc., 400 Sagner Avenue, Suite 300, Frederick, MD 21701, USA <sup>e</sup> Haley & Aldrich, Inc., 501 River St. #100, Greenville, SC 29601, USA <sup>l</sup> Department of Molecular Biology and Genetics, 321 Biotechnology Building, Cornell University, Ithaca, NY 14853, USA <sup>e</sup> Earth and Planetary Science and Environmental Science, Policy, and Management, University of California-Berkeley, Berkeley, CA 94720, USA <sup>h</sup> Department of Microbiology, University of Massachusetts, Amherst, MA 01003, USA Available online 27 November 2014.

### https://doi.org/10.1016/j.gca.2014.11.013G

# Abstract

Field-scale biostimulation and desorption tracer experiments conducted in a uranium (U) contaminated, shallow alluvial <u>aquifer</u> have provided insight into the coupling of microbiology, biogeochemistry, and hydrogeology that control U mobility in the subsurface. Initial experiments successfully tested the concept that Fe-reducing bacteria such as Geobacter sp. could enzymatically reduce soluble U(VI) to insoluble U(IV) during in situ electron donor amendment (Anderson et al., 2003; Williams et al., 2011). In parallel, in situdesorption tracer tests using bicarbonate amendment demonstrated rate-limited U(VI) desorption (Fox et al., 2012). These results and prior laboratory studies underscored the importance of enzymatic U(VI)-reduction and suggested the ability to combine desorption and bioreduction of U(VI). Here we report the results of a new field experiment in which bicarbonate-promoted uranium desorption and acetate amendment were combined and compared to an acetate amendment-only experiment in the same experimental plot. Results confirm that bicarbonate amendment to alluvial aguifer sediments desorbs U(VI) and increases the abundance of Ca-uranyl-carbonato complexes. At the same time, the rate of acetatepromoted enzymatic U(VI) reduction was greater in the presence of added bicarbonate in spite of the increased dominance of Ca-uranyl-carbonato aqueous complexes. A model-simulated peak rate of U(VI) reduction was ~3.8 times higher during acetate-bicarbonate treatment than under acetate-only conditions. Lack of consistent differences in microbial community structure between acetatebicarbonate and acetate-only treatments suggest that a significantly higher rate of U(VI) reduction in the bicarbonate-impacted sediment may be due to a higher intrinsic rate of microbial reduction induced by elevated concentrations of the bicarbonate oxyanion. The findings indicate that

bicarbonate amendment may be useful in improving the engineered <u>bioremediation</u> of uranium in aquifers.

# 1. Introduction

The importance of direct enzymatic or microbially-mediated reduction of soluble U(VI) has been established by extensive laboratory and field experiments (Lovley et al., 1991, Gorby and Lovley, 1992, Suzuki et al., 2005, Wu et al., 2006, Wu et al., 2010, Kelly et al., 2008, Hwang et al., 2009, Kostka et al., 2009, Xu et al., 2010, Tang et al., 2013, Newsome et al., 2014), including those conducted in, or with material from, an alluvial <u>aquifer</u> at Rifle, CO, USA (Anderson et al., 2003, Holmes et al., 2007b, Komlos et al., 2008, Fang et al., 2009a, Bopp et al., 2010, Chandler et al., 2010, Williams et al., 2011, Druhan et al., 2014a, Druhan et al., 2014b). Other studies have likewise established the importance of bicarbonate concentrations on the desorption of U(VI) from subsurface sediments (Curtis et al., 2004, Davis et al., 2004, Hyun et al., 2009), and the slowing of bioreduction when ternary Ca-uranyl-carbonato aqueous species dominate U(VI) aqueous speciation (Dong and Brooks, 2006, Stewart et al., 2007). The idea of coupling U(VI) desorption using bicarbonate and microbial reduction of U(VI) was first proposed by Phillips et al. (1995) based on laboratory desorption of U(VI) from a variety of sediment types followed by bioreduction using Desulfovibrio desulfuricans. Subsequent field research at Rifle suggested the likely efficacy of such an approach under in situ aquifer conditions (Ortiz-Bernad et al., 2004a). An in situ bicarbonateonly desorption experiment at the Rifle site (referred to as "Little Rusty") confirmed that dissolved U(VI) concentrations increased significantly (1.2-2.6-fold above background levels), resulting from increases in bicarbonate alkalinity from injectate solution and Ca concentrations due to cation exchange (Fox et al., 2012). Earlier field experiments also conducted at the Rifle site (especially one referred to as "Big Rusty") had indicated that the predominance of Ca-uranyl-carbonato species did not prevent significant U(VI) bioreduction, which bolstered the idea that bicarbonate desorption could be combined with down-gradient, acetate-induced biostimulation to increase the mass of U(VI) reduced in a the subsurface aguifer (Williams et al., 2011). The field experimental results from Big Rusty were also the focus of an extensive reactive transport modeling effort (Yabusaki et al., 2011) that provided the underlying biogeochemical reaction network used in modeling for this paper. The objectives of this study (dubbed "Super 8") were to: (1) confirm and extend previous <u>uranium</u> desorption results using bicarbonate (Fox et al., 2012); (2) quantify uranium mobility during acetate amendment under conditions of varying alkalinity and where microbial iron reduction is the dominant metabolic process; (3) assess the extent to which higher alkalinity values would impact U(VI) speciation and the rate of enzymatic uranium reduction; and (4) compare the post-amendment rebound of U(VI) with and without desorption of U(VI) by bicarbonate amendment. Here we emphasize the conservative tracer, biogeochemistry, microbiology, and U(VI) reduction results from the bicarbonate-acetate and acetate-only parts of the Super 8 field experiment. Other studies conducted as part of the same experiment are reported elsewhere (Chandler et al., 2013, Shiel et al., 2013, Holmes et al., 2013a, Holmes et al., 2013b, Alessi et al., 2014, Bao et al., 2014, Handley et al., 2014). These studies investigated selected aspects of the experiment, including reactive transport

modeling, microbiology of selected wells, <u>gene expression</u>, U isotopic shifts during bioreduction, and predation by <u>protists</u> on bacteria during biostimulation. Here, we combine biogeochemistry, microbiology, conservative tracer, and modeling of U(VI) reduction from the bicarbonate-acetate and acetate-only treatments, to demonstrate that bicarbonate both increases the pool of uranium available for acetate-induced U(VI) reduction and also impacts the <u>microbial community</u> in a way that enhances the intrinsic rate of U(VI) bioreduction.

# 2. Materials and methods

Site location, site geohydrology, <u>analytical methods</u>, and materials are described elsewhere (<u>Williams</u> <u>et al., 2011</u>, <u>Fox et al., 2012</u> and references therein). For this experiment, we installed a new set of 34 wells in 2010 (Plot C) in a previously unstimulated portion of the Rifle site (Fig. 1) using the rotary sonic drilling method. The well layout was designed to combine <u>bicarbonate</u> and <u>acetate</u> amendment on one side of Plot C with the other side reserved solely for acetate addition. The timing of amendments was designed to span the period when microbial Fe reduction was the dominant terminal electron accepting process. However, the target initial bicarbonate concentration ( $\sim$ 30 mM) in the <u>aquifer</u> was chosen to mimic that produced during biostimulation when sulfate reduction is at a maximum. In <u>field experiments</u> at the Rifle site, acetate amendment initially stimulates Fe-reducing microbial populations that transitions to dominance of <u>sulfate reducers</u> after about 3 weeks (<u>Williams</u> <u>et al., 2011</u>), producing greater bicarbonate concentrations. Stoichiometrically, the background concentration of  $\sim$ 10 mM. However, Fe-reducers continue to reduce metals even during dominance of sulfate reduce metals even during dominance of sulfate reduce populations that populations.



Fig. 1. Well layout for Plot C, site of the Super 8 field experiment. The Rifle field site location is provided in Williams et al., (2011). Inset shows the two tanks containing the separate injectates used in the experiment.

Wells were emplaced to a depth corresponding to the contact between the aquifer alluvium and the largely impermeable Wasatch formation ( $\sim$ 6–7 m below ground surface, see <u>DOE (1999)</u> for additional information on geologic setting). Wells were typically screened over the entire saturated thickness of the aquifer ( $\sim$ 3–7 m below ground surface). Selected wells were designed to collect samples at 3 discrete depths.

Deuterium (<sup>2</sup>H or D) was used as a conservative tracer for the bicarbonate injectate and quantified (as water with one hydrogen (H), one D and one oxygen (O) or HDO) with a liquid <u>water</u> <u>isotope</u> instrument (Los Gatos Research Inc.) as described in <u>Berman et al. (2009)</u> except for differences detailed in <u>Appendix A. Supplementary Data</u>. NaBr was used as a conservative tracer for the acetate (CH<sub>3</sub>COONa·3H<sub>2</sub>O) amendment and analyzed by <u>ion chromatography</u> as described elsewhere (<u>Williams et al., 2011</u>).

# 2.1. Experimental design and timeline

The experimental objectives required that groundwater amendments (acetate and sodium bicarbonate) be introduced at spatially distinct locations and timed such that acetate followed bicarbonate <u>desorption</u> of U(VI) on the bicarbonate-acetate side of the experiment. To achieve these objectives, groundwater from a location with no previous biostimulation (well #655) was pumped into two separate tanks, (1) a 6000 L high-density <u>polyethylene</u>(HDPE) tank for bicarbonate plus

deuterium as a tracer and (2) a 2120 L stainless steel tank for acetate and NaBr as a tracer. NaHCO<sub>3</sub> (Fisher Scientific) and 70% D<sub>2</sub>O (Cambridge Isotope Laboratories, Inc.) were added to the HDPE tank to achieve a  $\delta D$  of tank water of ~380‰ and a bicarbonate concentration of 50 mM. The HDPE tank was sparged with CO<sub>2</sub> to prevent extensive oxygenation of stored groundwater, and achieve a pH of ~7. Tank contents were circulated for 4 days (first tank) and 2 days (second tank) to enhance mixing and dissolution. The HDPE tank remained open to atmosphere over the injection period (Table 1), with a 30–45 min daily CO<sub>2</sub>sparging to maintain pH at 7.1 ± 0.2. The stainless steel tank (2120 L) was amended with 50 mM CH<sub>3</sub>COONa·3H<sub>2</sub>O (Sigma Aldrich) and 20 mM NaBr (Sigma Aldrich) while sparging with N<sub>2</sub>. The tank remained sealed under N<sub>2</sub>headspace during the injection period.

Table 1. Summary of injection activities, wells utilized, duration and dates of injection, injected volume, and concentration of injectates for the "Super 8" field experiment.

	Injection activity	Injection well numbers	Date <sup>[time]</sup>	Duration [days]	Injected volume [L]	Reagent concentration¤[mM]	Isotope enrichment*[‰]
Precursory tracers							
	NaBr	CA02	10-Aug- 10 <sup>[1603]</sup>	0.58	400	5.8 [Br-]	-
	$H_{2}^{18}O$	CG04	15-Aug- 10 <sup>[1750]</sup>	0.53	350	-	23 [ <sup>18</sup> O]
Posterior tracers							
	$NaBr + D_2O$	CA02	20-Nov- 10 <sup>[1000]</sup>	0.58	400	5.8 [Br-]	500 [D]
Bicarbonate–deuterium							
	#1 start	CA01- CA03	16-Aug- 10 <sup>[0630]</sup>	11	6000	50 [HCO <sub>3</sub> -]	500 [D]
	#1 end	""	27-Aug- 10 <sup>[1400]</sup>				
	#2 start	""	29-Aug- 10 <sup>[1200]</sup>	10	6000	50 [HCO <sub>3</sub> -]	500 [D]
	#2 end	""	7-Sept- 10 <sup>[1912]</sup>				
Acetate-bromide							
	#1 start	CG01- CG10	23-Aug- 10 <sup>[1256]</sup>	14	2200	50 [CH <sub>3</sub> COO <sup>-</sup> ]	_
	#1 end	« »	7-Sept- 10 <sup>[0700]</sup>			20 [Br-]	
	#2 start <sup></sup>	« »	13- Sept- 10 <sup>[1845]</sup>	9	1500	50 [CH <sub>3</sub> COO <sup>-</sup> ]	-
	#2 end	""	22- Sept- 10 <sup>[1718]</sup>			20 [Br-]	

\* Concentration/enrichment within the injection tank.

\*\* Tank #2 injection was initially started on 9-Sept-10; however, a closed injection valve prevented flow from the tank; injection was re-started on 13-Sept-10, as indicated.

Both injectates were introduced to the aquifer using separate sets of <u>boreholes</u> oriented approximately orthogonal to <u>groundwater flow</u> direction and spaced at 1.5 m intervals (Fig. 1). Tank contents were introduced to each injection well at 3.5 and 5.5 m below ground surface at rates of 180 L well<sup>-1</sup> day<sup>-1</sup> (bicarbonate tank; 3 wells, CA01, CA02, CA03) and 16 L well<sup>-1</sup> day<sup>-1</sup> (acetate tank; 10 wells, CG01-CG10). Injection of acetate/bromide was interrupted for ca. 5 days between the first and second filling of the acetate tank (<u>Table 1</u>). Cross-well mixing was used to disperse the injectates across the zone of injection, with peristaltic pumps circulating fluids (0.7 L min<sup>-1</sup>) between adjacent wells through HDPE tubes (<u>Williams et al., 2011</u>). For both injectates, fluids from two wells (e.g. CA01 and CA03) were withdrawn from a depth of 6 m and injected simultaneously into an adjacent well (e.g. CA02) at a depth of 4 m to create head differences between adjacent wells. Water level elevations were monitored at 15-min intervals in all injection wells using pressure transducers. Pump directions were changed daily such that individual wells served alternately as extraction and injection wells for cross-well mixing.

NaHCO<sub>3</sub> (~50 mM) was delivered to the aquifer through wells CA01-CA03, such that NaHCO<sub>3</sub> entered the system upgradient of subsequent acetate amendment (Fig. 1). After injection,  $HCO_3^-$  was advected along the primary groundwater flow direction and intersected the region of acetate amendment in wells CG06-CG09. Wells CG01-CG05 were located in a region unimpacted by  $HCO_3^-$ . All of the CG wells (CG01-CG10) were used for injection of acetate and NaBr.

# 2.2. Aqueous U(VI) speciation calculations

Uranium(VI) speciation calculations were performed using the <u>thermodynamic</u> data published in <u>Hyun</u> <u>et al. (2009)</u> and the computer program FITEQL4 set to equilibrium mode (<u>Herbelin and Westall,</u> <u>1999</u>). Most of the uranium thermodynamic data are consistent with the NEA database (<u>Guillaumont</u> <u>et al., 2003</u>) with the exception of Ca-uranyl-carbonato complexes and Mg-uranyl-carbonato complex (<u>Dong and Brooks, 2006</u>, <u>Dong and Brooks, 2008</u>). Ionic strength corrections in FITEQL4 were made with the Davies equation.

# 2.3. Microbiological analyses

Bicarbonate-acetate and acetate-only treatments were compared microbiologically by 16S rRNA gene analysis of the groundwater. A 16S rRNA-targeted gel element 'amplification' microarray (combining amplification, labeling and <u>hybridization</u> in a single closed microfluidic chamber) method was used, which targets 24 genera of dissimilatory metal-, sulfate- and nitrate-reducers (<u>Chandler et al., 2013</u>). The link between <u>phylogeny</u> and function in the microarray targets has been established via culture-based methods (see <u>Andersen et al., 2010</u> and references therein). Total <u>nucleic acid</u> from 1 to 2 L groundwater was extracted from filter cartridges with a MoBio PowerSoil <u>DNA</u> Isolation Kit (Carlsbad, CA) following manufacturer's instructions. Three microliters of purified nucleic acid representing 30–

60 mL equivalent of groundwater was processed in duplicate as described in detail elsewhere (<u>Chandler et al., 2010</u>).

Microbial groundwater communities from a select time point during peak Fe(III) reduction were further analyzed using clone library analysis. DNA was extracted from groundwater collected on 9/14/2010 from acetate-only and bicarbonate-acetate treatment wells CD04 and CD14, respectively, as previously described (Giloteaux et al., 2013). Briefly, 16S rRNA sequences were amplified using primers 338F (Lane et al., 1985) and 907R (Amann et al., 1990) and clone libraries were constructed with a TOPO TA cloning kit (Invitrogen, Carlsbad, CA) according to the manufacturer's instructions. One hundred and fifty plasmid inserts per clone library were sequenced with the universal M13F/M13R primer set.

DNA from sediment-attached microbial populations was also extracted from sediment collected during drilling of well CD04 as part of a different study (Handley et al., 2014). In short, freshly obtained Rifle aquifer sediment was first packed into flow through columns, and incubated within wells during acetate-amendment. DNA was extracted from homogenized sediment using PowerMax Soil DNA Isolation Kits (MoBio Laboratories, Inc., Carlsbad, CA, USA). 16S rRNA genes were amplified using the universal bacterial primers 27F and 1492R, and gradient PCR (11 annealing temperature increments, 48–58 °C). Amplicons were fragmented to ~300 bp and paired-end libraries were constructed prior to sequencing on the HiSeq2000 platform (Illumina® Inc., San Diego, CA, USA). Quality trimmed reads (Q > 3) were reconstructed into full-length sequences using the EMIRGE method (Miller et al., 2011, Miller et al., 2013).

# 2.4. Reactive transport modeling

Modeling of the Super 8 experiment is based on a simulation capability developed for prior Rifle field biostimulation experiments (Yabusaki et al., 2007, Li et al., 2009, Li et al., 2010, Fang et al., 2009b, Yabusaki et al., 2011), and relies heavily on model parameters developed for the 2008 Big Rusty acetate biostimulation field experiment (Williams et al., 2011). eSTOMP, a massively parallel processing, multifluid flow and multicomponent reactive transport subsurface simulator provided the framework and the high performance computing (HPC) infrastructure to address high spatial resolution and the complex biogeochemical processes involved in the field experiment as described below. Simulations covered a 70-day period to capture the time period that includes the focus of this paper, which is comparing reduction rates under acetate-only and bicarbonate-acetate conditions. The 70-day cutoff was thus chosen to capture the behaviors of interest while conserving HPC resources.

A large biogeochemical reaction network is required to model Rifle field experiments in part because of the broad range of uranium mobility that is sensitive to pH, <u>alkalinity</u>, major ion chemistry, redox state and the surface reactivity of the subsurface sediments (<u>Morrison et al., 1995</u>, <u>Davis et al.,</u> 2002, <u>Fox et al., 2012</u>). The modeling challenge is that products of the biostimulation (e.g., bicarbonate, Fe(II), U(IV), sulfide) can alter the geochemical conditions controlling uranium mobility (<u>Fang et al., 2009b</u>, <u>Zhao et al., 2013</u>). Using the <u>Yabusaki et al. (2011</u>) biogeochemical reaction network (i.e., <u>stoichiometry</u>, thermodynamics, and rate laws), we simulated the 2010 Super 8 field

biostimulation experiment with 102 chemical species and 7 minerals addressing 3 microbiallymediated terminal electron accepting processes (i.e., Fe(III), U(VI), sulfate), and Fe(III)- and sulfatereducing microorganisms and their biomass. Major ion chemistry, mineral reactions (i.e., <u>calcite</u>, <u>siderite</u>, <u>goethite</u>, <u>magnetite</u>, FeS, elemental <u>sulfur</u>, uraninite), surface <u>complexation</u> (i.e., H<sup>+</sup>, Fe(II), U(VI)), and ion exchange processes (i.e., Ca<sup>++</sup>, Mg<sup>++</sup>, Na<sup>+</sup>, K<sup>+</sup>) were included. Ion exchange capacity and surface sites for U(VI) <u>sorption</u> were modified from the <u>Yabusaki et al. (2011)</u> model to provide an improved fit to the observed desorbed U(VI) and major <u>cation</u>concentrations, especially Ca<sup>++</sup> (see <u>SI for additional details</u>).

Two key assumptions in the model are (1) U(VI) aqueous species re-equilibrate, diffuse, or advect rapidly such that after formation or consumption of one species the equilibrium proportions for the local geochemical conditions are re-established rapidly, and (2) <u>uraninite</u> (UO<sub>2</sub>) is the product of bioreduction of U(VI). Uranyl <u>ion concentration</u> is used in the bioreduction reaction and rate law. Since the uranyl ion concentration is small relative to the total aqueous U(VI) concentration, its inclusion instead of total U(VI) does limit the conversion rate. However, the kinetic reductive removal of uranyl ion from solution is rapidly restored via the equilibrium speciation of the remaining U(VI) in solution. While monomeric (or non-crystalline) U(IV) is an important component of total U(IV) in bioreduced Rifle sediments (Bargar et al., 2013, Cerrato et al., 2013, Alessi et al., 2014), and Bargar et al. (2013) point out the need for reactive transport models that include monomeric U(VI) as bioreduction products, the lack of stability constants and uncertainty regarding biogeochemical controls for formation of monomeric U(IV) limit our ability to incorporate non-uraninite forms of U(IV) into the model. Uraninite is also assumed to be stable under subsurface conditions at Rifle that include low <u>dissolved oxygen</u> (typically <0.2 mg/L, Yabusaki et al., 2007) and based on *in situ* experiments at Rifle that show slow dissolution rates for uraninite (Campbell et al., 2011).

The geology in the well <u>field (Plot</u> C) is predominantly the sandy gravel <u>lithofacies</u> defined in <u>Yabusaki</u> <u>et al. (2011)</u> for an earlier field experiment at the site, the Big Rusty acetate biostimulation experiment conducted in the Summer and Fall of 2008. The lithofacies-based properties of the sandy gravel classification (30 m/d <u>hydraulic conductivity</u>, 0.230 <u>porosity</u>, and 3.25 m<sup>2</sup>/g specific surface area) were therefore assumed. As in the simulation for the Big Rusty experiment, longitudinal and transverse <u>dispersivity</u> were 0.4 and 0.04 m, respectively (based on analysis of bromide tracer data, see <u>Yabusaki et al. 2011</u>).

The modeled subsurface domain is approximately 14 m downgradient by 18 m transverse by 6.5 m deep, and is represented by 209,664 grid cells with nominal grid spacing of 0.25 m laterally and 0.10 m vertically. The <u>flow field</u> and saturated thickness are driven by time-dependent water levels on the domain boundary, and bicarbonate and acetate solutions introduced at injection wells. The model boundary conditions are based on water level measurements over the 70-day simulation period. In all cases, it was assumed that the total injectate volume was distributed equally over the specified set of injection wells for the duration of the release, a reasonable assumption for a newly established well field in the Rifle alluvial aquifer (<u>Yabusaki et al., 2007</u>, <u>Williams et al., 2011</u>).

3. Results

Samples from ~33 wells were analyzed during the course of the <u>field experiment</u>; here we plot <u>geochemistry</u> results from three wells that show the results from different parts of the experiment: bicarbonate-only (well CU03), acetate-only (well CD01) and bicarbonate-acetate (well CD14). Other wells exhibit similar trends depending on location in the experimental plot (experimental data for all wells are available at doi: 10.1594/PANGAEA.830272). Observations are well matched by the <u>reactive transport</u> model (e.g., Fig. S-3). Bicarbonate-only well CU03 was located ~1 m downgradient from the <u>bicarbonate</u> injection wells and 1 m upgradient from the <u>acetate</u> injection wells, such that it was exposed to the NaHCO<sub>3</sub>/deuterium injection but minimally impacted by the CH<sub>3</sub>COONa/NaBr injection. Sporadic, low concentrations of acetate and bromide were occasionally detected during sampling of well CU03, in part due to cycling of cross-well mixing within ~1 m of the CG01-CG10 injection gallery. Wells CD01 and CD14 were both impacted by the acetate injection, with both wells located ~2.5 m downgradient from the region of acetate injection (Fig. 1).

# 3.1. Groundwater quality parameters and conservative tracer breakthrough

The effects of bicarbonate and/or acetate on pH, specific/electrical conductance, Fe(II) concentrations, and  $\delta D$  values are illustrated in Fig. S-1. As expected, breakthrough of <u>deuterium</u> was primarily observed downgradient from NaHCO<sub>3</sub> injection wells CA01-CA03, with increased dilution observed between CU03 and CD14 (peak  $\delta D \approx 340\%$  at CU03 versus  $\delta D \approx 175\%$  at CD14). A +5‰ increase over baseline in  $\delta D$  was observed at CD01 (Fig. S-1; inset), something we attribute to the cross-well mixing process. A significant increase in HDO ( $\delta D$  of >35‰) was observed in wells CU04, CD11-CD17 (as intended), as well as CD06, CD08, CD09, and CD10 (not shown). Such breakthrough is generally consistent with the predicted flow direction (175–180° <u>azimuth</u> from north), but the channel feature along the top of the Wasatch Formation may have impacted flow of the relatively dense deuterium-bearing injectate (see Fig. 1 for isopleths of the contact between the Wasatch and the alluvium). Overall the conservative tracers (D and NaBr) demonstrate that the injectates were distributed over the experimental domain as intended.

Irrespective of the injectate (acetate, bicarbonate, or both), pH values generally remained within the 7–7.5 range; the abrupt excursion (pH > 7.5) observed in CU03 (bicarbonate only) accompanied a rapid rise in pH within the NaHCO<sub>3</sub>/D<sub>2</sub>O injection tank, which was eliminated by CO<sub>2</sub> sparging. Changes in groundwater <u>electrical conductivity</u> (EC) tracked the delivery of both injectates (50 mM NaHCO<sub>3</sub>, and 50 mM NaCH<sub>3</sub>COO + 20 mM NaBr, <u>Table 1</u>), with the greatest increases over baseline values observed in wells that were impacted by the bicarbonate injection. EC increases observed at CD01 and other wells solely impacted by the CH<sub>3</sub>COONa/NaBr injectate increased by <40%. Although the tank concentrations of the CH<sub>3</sub>COONa/NaBr injectate were similar in ionic strength that of the bicarbonate tank, injection flow rates were different for the two tanks such that a greater in-well dilution occurred for the acetate-only injection. As a consequence, the peak total <u>inorganic</u> carbon (TIC) in CU03 (bicarbonate-only) is ~50 mM (10–13 mM of which is background TIC), whereas peak acetate plus the NaBr tracer in CD01 (acetate-only) is ~9 mM. For bicarbonate-acetate (CD14) peak TIC + NaBr + acetate is ~40 mM, including 10–13 mM TIC background.

Considering the background TIC, the ionic strength from amendments in the acetate-only part of the experiment is  $\sim$ 50% of that of bicarbonate-acetate.

Increases in ferrous iron concentration were observed downgradient from the region of acetate addition, reflecting the reduction of Fe(III) by acetate-stimulated, <u>iron-reducing bacteria</u>, consistent with all previous acetate injections at the Rifle site. Some increase in Fe(II) was observed at CU03, which was likely caused by cation exchange (50 mM Na<sup>+</sup>) releasing Fe(II) sorbed to surface exchange sites (<u>Fox et al., 2013</u>) and/or movement of stimulated groundwater from the vicinity of the acetate injection wells toward CU03 (<1 m).

# 3.2. Geochemical changes accompanying bicarbonate injection

Expanding upon previous NaHCO<sub>3</sub> injection experiments at Rifle (Fox et al., 2012), geochemical changes in well CU03 were used to evaluate the impact of sustained (>20-day) levels of elevated HCO<sub>3</sub><sup>-</sup> on <u>uranium</u> mobility, metals (V, Fe, Mn), as well as its impact on other exchangeable <u>cations</u> (Ca, Mg, K, Na, Fig. 2). The results are consistent with an ion exchange model (Ca, Mg, K) driven by the ~70 mM Na in the amendments and enhanced <u>desorption</u> of uranium driven by the formation of stable aqueous species (e.g.,  $Ca_2UO_2(CO_3)_3^\circ$ , the predominant U(VI) species under the conditions of the experiment). Abrupt and prolonged decreases in total Mn concentration are inferred to result from the low solubility of <u>manganese</u> carbonates.



Fig. 2. Concentration of selected geochemical constituents in well CU03 plotted versus time in days after 8/16/2010 (bicarbonate-only conditions). NaHCO<sub>3</sub> injection period delineated by vertical bars. Relevant, maximum contaminant levels (MCL) and UMTRA standard concentrations are denoted by horizontal dashed lines with <u>concentration values</u> as shown.

Concentrations of multiple cations (Ca, Mg, and K) fell below their pre-injection values once NaHCO<sub>3</sub> injection ceased likely due to re-exchange of these cations with Na following the NaHCO<sub>3</sub> flush. This effect was previously observed in the Little Rusty field experiment (Fox et al., 2012, Fox et al., 2013) and was simulated in an updated ion exchange model. Following injection, influx of groundwater from regions upgradient of the zone of NaHCO<sub>3</sub> injection leads to removal of U(VI) from solution, as <u>sorption</u> sites for U(VI) species are re-filled. Once such sites are largely occupied, groundwater U(VI) concentrations rebound to pre-injection levels. In comparison with Little Rusty (Fox et al., 2012), both cation and U(VI) concentrations were generally slower to rebound following the Super 8 injection, likely due to the longer duration of the Super 8 injection (21 versus 0.18 d). In contrast, total Fe (Fe<sub>TTL</sub> = Fe<sup>III</sup> + Fe<sup>III</sup> ionic species) remained elevated following NaHCO<sub>3</sub> injection, potentially the result of enhanced release of Fe(III) <u>colloids</u> (<45 µm particle size) during injection (note that Fe(II) levels determined colorimetrically also fall to pre-injection values; Fig. S-1).

# 3.3. Geochemical changes accompanying acetate injection

The acetate-only portion of Plot C provided a direct comparison to the NaHCO<sub>3</sub>-impacted portion of the flow cell. Geochemical changes in well CD01 are shown in Fig. 3. Desorption and re-sorption of exchangeable cations was also observed during and after CH<sub>3</sub>COONa/NaBr injection, respectively, although the magnitude of the effect was less than that observed at CU03. The much smaller increase in ionic strength (and hence less injected Na<sup>+</sup>) accounts for the observed difference between the two sections of Plot C. The presence of acetate, absence of amended bicarbonate, and resulting stimulation of metal-reducing bacteria also led to markedly different changes in concentrations of redox-sensitive metals in CD01 relative to CU03. The contrast was especially strong for U(VI) and V, both of which can be reductively immobilized via enzymatic reduction by microorganisms (Lovley et al., 1991, Ortiz-Bernad et al., 2004b, Yelton et al., 2013). For ~5–7 days there was a slight increase in U(VI) and total V groundwater concentrations upon delivery of acetate to the wellbore region, followed by a rapid decrease in both U(VI) and total V in groundwater via reductive immobilization. A decrease in total V concentration in groundwater occurs immediately following the slight increase, similar to the observations made during the first Rifle field experiment (Ortiz-Bernad et al., 2004b), whereas removal of U(VI) was delayed by approximately 5 days. Groundwater U(VI) concentrations fell below the EPA's maximum contaminant level (MCL =  $0.126 \mu$ M) so long as acetate remained at detectable levels, as previously observed at the site (Williams et al., 2011). Once acetate injection ceased, U(VI) concentrations steadily rebounded due to the influx of U(VI)-bearing groundwaters from upgradient locations and a slowing of microbial reduction in the absence of acetate addition. In contrast, total V concentrations in groundwater remained below the cleanup target concentration (0.67 µM) far beyond the period where acetate fell below detectable levels (Ortiz-Bernad et al., 2004b, Yelton et al., 2013).



Fig. 3. Concentration of selected geochemical constituents in well CD01 plotted versus time in days after 8/16/2010 (acetate-only conditions). <u>Acetate</u> injection period delineated by vertical grey bars. MCL's and UMTRA standard are as shown in <u>Fig. 2</u>.

# 3.4. Combined Impact of bicarbonate and acetate injection

As anticipated, the temporal behavior of exchangeable cations, metals, and uranium at CD14 was a combination of the processes observed at CU03 and CD01 (Fig. 4, Fig. 5). Following an initial increase in U(VI) and V (as also observed in CU03), both metals were rapidly removed to very low concentrations. In contrast to the CU-03 and CD01 locations, U(VI) concentrations remained below the MCL for ~55 days after acetate fell below detection. Total V concentration remained below the cleanup target at the point of exposure (DOE, 2001) of 6.48 µM for more than 55 days after acetate levels fell below detection. <u>Vanadium</u> showed no indication of rebound, similar to the results obtained in well CD01 and observed in earlier experiments (Ortiz-Bernad et al., 2004b). Fe(II) reached greater maximum concentrations at CD14 than at CD01 (150 µM at CD01 versus 225 µM at CD14, Fig. 3, Fig. 4). While the impact of elevated HCO<sub>3</sub>- concentration on Fe(III) reduction is still uncertain, the significant increases in Fe(II) at CD14 relative to CD01 may simply be the result of a heterogeneous distribution of bioavailable Fe(III) in subsurface sediments or a combination of Fe(III) reduction and Fe(II) release through ion exchange (as observed in CU03). Uranium(VI) concentration increased twofold at CU03 (ca. 0.65–1.3 µM), whereas the increase at CD14 prior to acetate-induced U(VI) bioreduction was fourfold (ca.  $0.5-2 \mu$ M). The likely explanation is a longer transport distance (i.e., a larger volume of extracted sediments) over which bicarbonate is able to desorb U(VI). Specifically, the distance between the NaHCO<sub>3</sub>injection wells and monitoring well CU03 is ~1 m versus well CD14 which is  $\sim$ 3.5 m from the NaHCO<sub>3</sub> injection wells (cf. <u>Fox et al., 2012</u>).



Fig. 4. Concentration selected geochemical constituents in well CD14 plotted versus time in days after 8/16/2010 (bicarbonate-acetate conditions). <u>Acetate</u> injection period delineated by vertical bars. MCL's and UMTRA standard are as shown in <u>Fig. 2</u>.



Fig. 5. Comparison of the behavior of aqueous U(VI), D and bromide tracers, <u>acetate</u>, and total <u>inorganic carbon</u> (TIC) in CU03 (A), CD01 (B) and CD14 (C). TIC concentrations result from the NaHCO<sub>3</sub> injection (start and stops represented by dashed vertical lines). Starts and stops for acetate injection denoted by vertical grey bars (see <u>Table 1</u>). The inset in (A) shows acetate with an expanded scale where acetate is above detection. MCL is as shown in <u>Fig. 2</u>.

There was a large difference in total inorganic carbon (TIC) concentration associated with NaHCO<sub>3</sub> addition that resulted in a 6-fold difference in U(VI) desorption from sediments between

CD01 and CD14 (Fig. 5). Regardless of TIC concentration (and by extension [HCO<sub>3</sub>-]), the *apparent* rates of U(VI) decrease ( $\Delta$ U(VI)<sub>obs conc</sub>/t) associated with stimulated <u>microbial activity</u> were largely indistinguishable under high and low <u>alkalinity</u> and both wells reached relatively low values of U(VI) concentration (CD01: 0.03 µM on day 42 and CD14: 0.06 µM on day 53). The apparent rates are significant from the perspective of applied <u>bioremediation</u>, but understanding the underlying mechanisms and associated rates requires separating competing processes of aqueous speciation changes, desorption, bioreduction, and precipitation of new solid phases. We address this need by reactive transport modeling, establishing estimates of the *actual* rate of bioreduction of U(VI), as described in Section <u>3.8</u>.

# 3.5. Uranium rebound

One of the most pronounced differences between the bicarbonate-acetate versus the acetate-only parts of the experiment is the extent to which uranium rebounds after acetate injection ends (Fig. 5). Irrespective of the large increase in U(VI) concentrations associated with HCO<sub>3</sub>--mediated desorption, uranium was not only removed rapidly from groundwater at the wells impacted by both bicarbonate-acetate, but it also remained at low levels (below the MCL) long after acetate injection ended (well CD14, Fig. 5C). The same was typically not true for locations exposed solely to acetate. There, uranium concentrations rebounded immediately or soon after acetate concentrations fell to levels below detection (well CD01, Fig. 5B). The most likely explanation for this phenomenon involves the extent to which re-sorption of advecting U(VI) occurs in areas where desorption was greatly enhanced by NaHCO<sub>3</sub> injection. As noted above, movement of upgradient groundwater with U(VI) into the part of the bicarbonate-acetate part of the experiment is impacted by sorption onto newly vacant sorption sites; the same effect does not occur in wells close to the injection point (CD03, CD02, CD01, Fig. S-B) and amended with acetate-only because U(VI) desorption was limited in the acetate-only part of the experiment (an increase in U(VI) of ca. 0.7–0.9  $\mu$ M, much less than for either CU03 or CD14).

Fox et al. (2012) demonstrated that there is a kinetic limitation to U(VI) desorption in Rifle sediments, likely due to a mass transfer limitation related to intragranular porosity of the sediments, a pore-size constraint on movement of U(VI) off mineral surfaces into groundwater. Kinetic limitation to U(VI) desorption likely caused a limited availability of sorbed inventories of U(VI) to microbial reduction (Ortiz-Bernad et al., 2004a). Were the sorbed pool of U(VI) not kinetically limited, a similar refilling of vacant sorption sites with U(VI) in upgradient groundwater (and the suppression of U(VI) concentrations) would have been expected under acetate-only conditions (e.g. at CD01). This is because removal of U(VI) from groundwater by bioreduction of U(VI), without the kinetic limitation, would result in additional desorption of U(VI). The rate of U(VI) desorption is apparently slow enough in the acetate-only region that the sorbed pool of U(VI) is not significantly reduced. Microbial reduction of the sorbed pool of U(VI) which is rapidly replenished by U(VI) desorption from sediments). Apparently neither of these mechanisms is significant under the *in situ* conditions at the Rifle site, otherwise longer-term suppression of U(VI) concentrations would be expected under

acetate-only conditions. An important assumption here is that U(VI) is not reduced abiotically or biotically during the post-acetate amendment time period, or otherwise scavenged by incorporation into a newly formed phase such as <u>calcite</u>. While such mechanisms have been proposed (<u>N'Guessan et al., 2008</u>, <u>Hyun et al., 2012</u>), the process of filling of vacated sorption sites is consistent with reactive transport modeling described below, and provides the simplest explanation of the differences in U(VI) rebound under bicarbonate-acetate and acetate-only conditions.

3.6. Speciation of aqueous U

**Fig. 6** compares the calculated U(VI) speciation in groundwater over the duration of the experiment in wells CD01 (acetate-only) and CD14 (bicarbonate-acetate). The dissolved U(VI) speciation was dominated by the same five complexes in both cases, with uncharged calcium-uranyl tricarbonato species  $(Ca_2)UO_2(CO_3)_3^{\circ}$  making up ~97% of the dissolved U, followed by  $CaUO_2(CO_3)_3^{2-}$ , MgUO<sub>2</sub>(CO<sub>3</sub>)<sub>3</sub><sup>2-</sup>, and UO<sub>2</sub>(CO<sub>3</sub>)<sub>3</sub><sup>4-</sup>. Note that these species all contain 3 carbonate ligands, so we refer to the sum of these four species as the uranyl-tricarbonato species, and the group of all other dissolved U(VI) species as the non-tricarbonato species. The UO<sub>2</sub>(CO<sub>3</sub>)<sub>2</sub><sup>2-</sup>species has the fifth highest concentrations of dissolved U(VI) in both wells; its concentration was more than 3 orders of magnitude lower than the uncharged, calcium-uranyl tricarbonato species, but was about an order of magnitude higher under acetate-only conditions (CD01) than under bicarbonate-acetate conditions (CD14).



Fig. 6. Plots of the total concentration of dissolved U and calculated U(VI) aqueous species concentrations as a function of time in wells CD01 (A) and CD14 (B). Vertical lines are the same as in Fig. 5. See "Section 2" for details on calculations.

The concentrations of dissolved U(VI) in wells CD01 and CD14 were slightly different at the beginning and varied over the duration of the experiment. In well CD01 (acetate-only), the initial concentration was 0.75  $\mu$ M, and 0.051% was calculated to be present as non-tricarbonato species. For CD14 (acetate-bicarbonate), the values were 0.66  $\mu$ M and 0.064%. Fig. 7 shows the ratio of the relative fraction of non-tricarbonato species in well CD14 (bicarbonate-acetate) to that in well CD01 (acetateonly) over the course of the experiment. The non-tricarbonato ratio is slightly greater than 1 at the beginning of the experiment, but then the ratio decreases rapidly to 0.25 during the bicarbonate and acetate additions to the groundwater (Fig. 7), as the increased bicarbonate concentration causes a greater proportion of dissolved U(VI) to be present as uranyl-tricarbonato species in well CD14. Later in the experiment (>40 days), the water chemistry at CD14 changes significantly as the bicarbonate plume is transported away and the non-tricarbonato ratio for CD14:CD01 is again greater than 1 for a period of time.



Fig. 7. Plot of the ratio of the calculated fractional contribution of non-tricarbonato species to total dissolved U in well CD14 versus well CD01 as a function of time. Vertical lines are the same as in Fig. 5.

As shown by <u>Ulrich et al. (2011)</u>, the kinetics of uranium bioreduction are highly sensitive to U(VI) aqueous speciation, and thus depend significantly on pH and calcium and bicarbonate concentrations. The greater proportion of uranyl-tricarbonato species present under bicarbonate-amended conditions is therefore expected. However,  $\Delta U(VI)_{obs conc}/t$  under bicarbonate-acetate and acetate-only conditions were still indistinguishable in spite of the laboratory observation that Ca-uranyl-tricarbonato species in particular are bioreduced at slower rates than non-tricarbonato species (Dong and Brooks, 2006, Stewart et al., 2007).

# 3.7. Microbiology

Relative changes in Plot C microbiology with time are shown in Fig. 8, and provide more extensive data than the amplification microarray data reported previously for this experiment (Chandler et al. 2013). Relative to the CU01 background (i.e., unimpacted by acetate or bicarbonate) signatures, the bloom of metal-reducing bacteria in all wells was dominated by *Geobacter* and *Pelobacter* 16S rRNA gene signatures, as expected. The transition from iron to sulfate reduction was accompanied by a decline in planktonic metal-reducers and an increase in *Desulfotomaculum* and *Clostridium* (data not shown, see doi: 10.1594/PANGAEA.830272). The relative increase in *Desulfotomaculum* and *Clostridium* signatures is interesting in light of recent data indicating that some *Desulfotomaculum* and *Clostridium* species are capable of U(VI) reduction under fermentative conditions, even as spores(Junier et al., 2009, Bernier-Latmani et al., 2010, Junier et al., 2011). Discernable differences among the composition of microbial communities were not evident between acetate-only and bicarbonate-acetate amendments.



Fig. 8. <u>Spatial distribution</u> and time variation of metal reducers, <u>sulfate reducers</u> and fermenters in selected wells based on microarray data. The *Y*-axis scale on graphs is the <u>signal to noise ratio</u>, SNR, *X*-axis scale is days since the start of the experiment. All graphs have the same scales (*X*-axis 0–70 days, *Y*-axis 0–500 SNR). Charts show stacked results for the three functional groups of bacteria. Wells CD11, CD14 and CD16 represent bicarbonate-acetate impacted wells as compared to CD02, CD05, and CD07 as acetate-only wells. Well CD-17 was missed by <u>bicarbonate</u> but was impacted by <u>acetate</u>. Day zero was 16 August 2010.

16S rRNA clone libraries constructed from groundwater samples collected from wells CD04 and CD14 on 9/16/2010 (Day 31) were collected as part of another microbiological study of the Super 8 experiment but provide results that are consistent with the <u>hybridization</u> microarray results (Fig. 9, upper panel). As expected for the acetate-only well CD04 (~1.5 m downgradient from CD01), a large fraction (~75%) of the community belongs to the Deltaproteobacteria (*Geobacter*, sp. mainly, some *Desulfobacteraceae*). CD14 is also enriched in Betaproteobacteria, Gammaproteobacteria, and, notably, fermenters, including microorganisms from candidate division OD1 (Wrighton et al., 2012). The variability between the proportion of Deltaproteobacteria between the two wells is within the range of that observed previously (e.g., <u>Anderson et al., 2003</u>).





Fig. 9. 16S rRNA gene clone library results (topmost chart) summarized to the class level for groundwater samples from wells CD-04 (acetate-only) and CD14 (bicarbonate-acetate) on 16 September 2010. For comparison, reconstructed 16S rRNA gene results are given below for unamended sediment and sediment from 4 replicate flow-through columns collected from acetate-only wells on 12th and 19th of September 2010 (CD03, CD01 and CD05). Reconstructed sequences used were present at  $\geq 1\%$  relative abundance.

Microbial community data from a more extensive time series (13 samples from *in situ* sediment columns) representing the acetate-only treatment have been analyzed in a manner similar to data reported on earlier Rifle experiments (Handley et al., 2012, Handley et al., 2013, Handley et al., 2014, Wrighton et al., 2012). In brief, results indicate a community structure that is analogous to the groundwater communities in CD14 and CD04 at a similar time point in the experiment (Fig. 9, lower panel), with the exception that bacteria belonging to the Bacteroidetes phylum are more prevalent in the sediment (namely unclassified Sphingobacteria, Flavobacteria and Bacteroidetes). As in the groundwater acetate-bicarbonate CD14 community, Deltaproteobacteria and Betaproteobacteria dominate in near equal proportions within the sediment and are clearly enriched relative to unamended background sediment and groundwater. Overall results indicate a similar microbiological response was achieved during both acetate and acetate-bicarbonate experiments that also reflects

results obtained during previous experiments (<u>Anderson et al., 2003</u>, <u>Handley et al., 2012</u>). Recognizing that 16S rRNA data on groundwater and sediments may not be exactly comparable to the hybridization microarray data, the 16S rRNA data indicate that the microbial community during biostimulation was typical for Rifle *in situ* acetate amendment experiments. They are broadly consistent with the hybridization microarray results and thus provide independent corroboration of the microarray data.

# 3.8. Reactive transport modeling and U(VI) reduction rates

Observed and simulated aqueous U(VI) concentrations were well correlated for most of the 27 monitoring locations over a 70-day simulation period (Fig. S-3). Simulated aqueous U(VI) concentrations under bicarbonate-only conditions (CU03 and CU04) rapidly increased from 0.5 to  $\sim$ 1.5 µM before a continuous decline. In the three rows of monitoring wells directly downgradient from the bicarbonate injection (CD11/12, CD13/14/15, and CD16/17), the peak concentration of the simulated aqueous U(VI) pulse progressively increased to  $\sim$ 2.0 µM. This peak reflects the accumulation of desorbed U(VI) as more of the aquifer sediments were impacted by the bicarbonate front moving through the aquifer. The occurrence of the simulated peak before the observed peaks in U(VI) concentration are related to the use of an equilibrium model, and the attenuation could be better described with a kinetic model for U(VI) desorption (Fox et al., 2012). The duration of this elevated U(VI) pulse is limited by both depletion of adsorbed U(VI) and U(VI) bioreduction that begins with injection of acetate on day 8.

Fig. 10 shows simulated U(VI) reduction rates at all monitoring locations over the 70-day simulation period. These rates are normalized to U(VI) concentration to account for the expected increase in U(VI) reduction rate with increasing U(VI) concentration based on Monod kinetics or the *in silico*model for iron-reducing bacteria (FeRB) used for this plot (Fang et al., 2012). Figs. S-4, S-6, S-7, and S-8 show absolute rates and rates normalized to FeRB biomass using the method of Bao et al. (2014). Per the modeling schema, the simulated rates are exclusively the result of U(VI) bioreduction and are derived from the time-dependent change in amount of U(VI) reduced (i.e., slope of graphs in Fig. S-2), providing the local reaction rate for U(VI) reduction.

U(IV) bioreduction rate normalized to aqueous U(VI) concentration



Fig. 10. Plots of the simulated U(VI) reduction rate normalized to U(VI) concentration as a function of time across the experimental domain. *Y*-axis values are mol  $L^{-1} d^{-1} mol_{U(VI)}^{-1} L^{-1}$  or  $d^{-1}$ .Well numbers with a digit after a decimal point (e.g., CD-13.5) indicate wells with discrete sampling ports ranging from 1 to 3 m above the base of the modeling domain (elevation 1612.85 m above MSL). The ground surface is 6.5 m above the base of the modeling domain.

The highest simulated U(VI)-bioreduction rates (normalized to U(VI) concentration, Fig. 10) occur in the bicarbonate-acetate part of the experiment at day 37 to day 55, depending on well location. At wells CD10, CD11, CD12, CD13, CD14, and CD16, peak simulated reduction rates range from 0.30 to 0.35 mol L<sup>-1</sup> d<sup>-1</sup> mol<sub>u(VI)</sub><sup>-1</sup> L<sup>-1</sup> (or d<sup>-1</sup>), with a mean value of 0.33 (1 $\sigma$  = 0.014, *n* = 6). Locations unimpacted by the bicarbonate injection (e.g., CD01) have simulated rates of ~0.06–

0.1 mol L<sup>-1</sup> d<sup>-1</sup> mol<sub>u(V)</sub><sup>-1</sup> L<sup>-1</sup> at the times of peak reduction rate in the wells above with a mean value = 0.087 (1 $\sigma$  = 0.015, *n* = 6). Thus peak simulated rates in bicarbonate-acetate wells are ~3.8 times higher than in the acetate-only wells.

The model output estimates the U(VI) bioreduction rate during the experiment for either bicarbonateacetate or acetate-only wells. Results clearly show that the estimated bioreduction rates for the acetate-bicarbonate treatment (well CD11) are higher than those for the acetate-only treatment (well CD01) 7 to ~25 days from the start of acetate amendment for both the un-normalized U(VI) bioreduction rate and for the U(VI) reduction rate normalized to biomass (Fig. S-6). For the U(VI) bioreduction rate normalized to U(VI) concentration (Fig. 10 and Fig. S-5), the estimated rate for the bicarbonate-acetate treatment exceeds rates for the acetate-only treatment from ~7 to ~62 days from the start of the experiment. The rates diverge from ~13 to ~35 days, with increasingly greater rates for the bicarbonate-acetate treatment while rates in CD01 (for example) remain nearly constant. This is the appropriate comparative time period during which acetate levels are similar on both sides of the experiment. As expected, the time of peak estimated U(VI) reduction rates shift to later times in further down-gradient wells.

The model does show significant simulated reduction rates for wells CU03 and CU04 that are  $\sim 1$  m upgradient from the line of acetate injection wells. Non-zero U(VI) reduction rates are expected in these wells due to transport of acetate away from injection wells during cycling of cross-well mixing. Peak rates of 1/3 or 1/2 of the peak rates in nearby down-gradient wells are likely overestimated given the low measured acetate concentrations in these wells.

# 4. Discussion

# 4.1. U(VI) desorption and changes in U aqueous complexes

The impact of U(VI) <u>desorption</u> on U(VI) aqueous concentrations is clearly demonstrated by this experiment. Results from monitoring wells downgradient from the <u>bicarbonate</u> injection (e.g., CU03) suggest that U(VI) desorption began to deplete the available pool of sorbed U(VI) based on the declining U(VI) concentrations, while bicarbonate concentrations remained at or near their peak concentrations. The implication of this observation is that near-complete desorption of labile U(VI) is feasible over reasonably short time frames (days to weeks) and that it might be practical to use desorption as a remedial technology either as a stand-alone U(VI) flushing process or in combination with bioreduction to enhance <u>bioavailability</u> of U(VI). Results from the experiment confirm earlier work (<u>Phillips et al., 1995</u>, <u>Williams et al., 2011</u>) that shows no apparent inhibition of the reduction of U(VI) even when calcium-uranyl tricarbonato complexes dominate aqueous species.

However, the net effect of bicarbonate addition to the system is complex, increasing total dissolved U(VI) but also decreasing the proportion of more readily bioavailable non-tricarbonato species. Note that the decrease in U(VI) concentration begins even when DIC and  $\delta D$  are still rising, indicating that microbial reduction is outpacing U(VI) desorption on the bicarbonate-acetate side of the experiment (e.g. well CD14). The net effect on enzymatic U(VI) reduction is thus also complex, with the simulated *absolute* rate of reduction faster in CD14 than at CD01 (for days 7–23, Fig. S-4), partly

related to the greater total dissolved U(VI) concentration. The simulated rate of reduction per mole of U(VI) is faster on the bicarbonate-acetate side for a longer duration (days 12–62). The proportion of uranyl-tricarbonato species during days 10–40 on the bicarbonate side of the experiment likely plays a limited role in the U(VI) reduction rates (Fig. 7 cf. Fig. 10 CD01 and CD14), but the mechanistic interaction between U speciation and microbial reduction of U(VI) is still poorly understood. Given the increase abundance of uranyl-tricarbonato species under conditions of bicarbonate-acetate (Ulrich et al., 2011 and Fig. 7), it would be expected that there would be lower  $\Delta U(VI)_{obs core}/t$  on the acetate-bicarbonate amended side of the experiment. This difference might be expected to be significant enough to show up directly in U(VI) concentrations as function of time. As this is not the case, it suggests that (1) the model assumption that U(VI) aqueous species re-equilibrate, diffuse, or advect rapidly such that there is no decrease in the  $\Delta U(VI)_{obs core}/t$  with bicarbonate amendment is correct, or (2) under Rifle field conditions the *in situ* biostimulated microbial community is not sufficiently sensitive to U(VI) aqueous speciation to affect the rates of bioreduction (Ortiz-Bernad et al., 2004a).

# 4.2. Impacts of bicarbonate on the microbial community and U(VI) bioreduction rates

Univariate and multivariate statistical analysis of the microarray data show no consistent differences in the microbial community structure between the bicarbonate-acetate and the acetate-only treatments (Fig. S-9). The analysis indicates that total biomass is not correlated with TIC, acetate concentration and metal reducers but positively correlated with sulfate reducers and fermenters, consistent with the microbial succession during the *field experiments* at Rifle (i.e., biomass increases with time or the degree of stimulation). This analysis is consistent with the temporal differences in microbial biomass (total signal to noise ratio, SNR) in bicarbonate-impacted wells in Fig. 8. In addition, simulated U(VI) reduction rates normalized to U(VI) concentration are higher for the bicarbonate-acetate treatment than for the acetate-only treatment. These results suggest that although the acetate-bicarbonate treated microbial community is compositionally indistinguishable from the acetate treated community, the former is more effective at U(VI) reduction, irrespective of total planktonic biomass (Fig. S-6), or concurrent reduction of bioavailable oxidized iron (Bao et al., 2014). Heterogeneity in bioavailable Fe(III) could partially account for differences in Fe-reducers across the experimental domain, but these difference do not translate into modeled differences in U(VI) reduction rate. Since there are no consistent differences in community structure, or biomass between the bicarbonate-acetate and acetate-only treatments, the difference in U(VI) reduction rate must be related to the difference in bicarbonate and its impact on the intrinsic U(VI) reduction rate of that part of the microbial community responsible for U(VI) reduction, most likely the metal reducers (Fig. 8).

<u>Chandler et al. (2013)</u> suggested that dissolved CO<sub>2</sub> may enhance the reverse tricarboxylic acid cycle of U-respiring organisms, leading to increased rates of U(VI) reduction per cell. Oxyanions, sulfate, <u>silicate</u>, and phosphate have been shown to favor microbial precipitation of monomeric U(IV) by increasing bacterial extracellular polymeric substances (EPS) and bacterial viability (<u>Stylo et al.,</u> 2013), providing further evidence that the oxyanion bicarbonate may also impact U(IV) species and

reduction rates. On the other hand, the oxyanion bicarbonate favors formation of <u>uraninite</u> (UO<sub>2</sub>) and suppresses the production of monomeric U(IV) (<u>Bernier-Latmani et al., 2010</u>). The general importance of bicarbonate to microbial <u>redox processes</u> is also supported by its effect on microbial products of Fe(II) oxidation, in which the formation of <u>goethite</u> ( $\alpha$ -FeOOH) at the expense <u>lepidocrocite</u> ( $\gamma$ -FeOOH) was favored under conditions of high increased bicarbonate and <u>humic acid(Larese-Casanova et al., 2010</u>). However, bicarbonate does not seem to have a strong impact without the presence of microbes. For example, abiotic reduction of U(VI) by FeS was not impacted by the presence or absence of bicarbonate (<u>Gallegos et al., 2013</u>), again supporting the concept that bicarbonate increases U(VI) reduction rates by impacting the metabolism of the subsurface microbial community.

The attached microbial community may also play a role in governing U(VI) reduction rates under Fereducing conditions (Bargar et al., 2013), which would not be fully accounted for by sampling the planktonic community in groundwater in this study. However, attached and planktonic *Geobacter* sp. are similar during acetate biostimulation (Holmes et al., 2007a), although other important differences in community membership may exist (e.g. Fig. 9). Based on currently available data, it seems unlikely that the un-sampled, attached population of Fe-reducers accounts for the difference in U(VI) reduction rate on the two sides the Super 8 experimental plot. Since the Holmes et al. (2007a) data were collected only under conditions of acetate stimulation, the possibility that attached versus unattached Fe-reducer populations are different under conditions of bicarbonate-acetate cannot be entirely ruled out. In addition, changes in DNA signatures over time do not necessarily correlate with metalreducing activity or rates in situ. For example, analysis of samples collected from CD04 during the Super 8 experiment in conjunction with ex situ column experiments demonstrate that rpsC gene expression correlates better with the rate of *Geobacter* sp. growth and metabolism than changes in cell abundance (Holmes et al., 2013a). U(VI) is a trace electron acceptor at Rifle compared to the major electron acceptors that are available for anaerobic respiration (e.g. Fe(III)), and as noted above, a wide range of organisms can reduce U(VI), so relatively low-abundance of any U(VI)reducing cells could account for the difference in U(VI) reduction rate.

Putative fermenters are present in most wells (Fig. 8), commonly increasing as the experiment progressed. Part of the taxonomically inferred fermentative community includes organisms from the OD1 candidate division (Wrighton et al., 2012). Recent research indicates that the fermentative community is important in carbon, <u>sulfur</u>, and nitrogen cycling under biostimulated conditions with implication for utilization of natural dissolved and detrital <u>organic carbon</u> in sediments (Wrighton et al., 2014). Some of the organisms inferred to be obligate fermenters described in these studies also cycle hydrogen which may drive U(VI) reduction and could account for some of the U(VI) reduction in the later stages of the experiment (Bernier-Latmani et al., 2010, Bargar et al., 2013). In addition, abiotic reduction of U(VI) by sorbed Fe(II) or by FeS could make a small contribution to the total observed U(VI) reduction under these experimental conditions (Hyun et al., 2012, Fox et al., 2013).

4.3. Complexity of processes governing U(VI) concentrations during bioreduction

In the case of both iron and <u>uranium</u>, it is particularly difficult to quantitatively discern the individual contributions of processes governing their concentrations from measured concentrations alone. This is because both elements have complex aqueous and solid species that are directly involved in aqueous and surface <u>complexation</u>, microbially mediated terminal electron accepting processes (TEAPs), and mineral reactions, which are sensitive to geochemical parameters such as pH, Eh, Fe(III), and <u>alkalinity</u> which are in turn controlled by reactions involving other elements. For example, minor increases in bicarbonate as a byproduct of microbial respiration during electron donor field experiments (Williams et al., 2011) not only promotes U(VI) desorption, but also may exert a <u>positive</u> feedback for enhanced microbial reduction of U(VI), albeit at a lower rate than when high concentrations of bicarbonate are amended to the subsurface.

Another example of the complexity of U biogeochemistry is the structure and bonding of U(IV) products of bioreduction. The simulated U(VI) bioreduction rates are the most readily available estimate because of the practical difficulty of obtaining subsurface sediment samples and the challenge of simultaneously measuring sorbed U(IV) and the concentration of monomeric U(IV) and uraninite in bioreduced sediments (Alessi et al., 2012, Stoliker et al., 2013). Moreover, X-ray spectroscopy has been very successful at identifying U redox status and bonding of U phases, but has not yet been able to quantify U(IV) at typical concentrations in a sediment matrix (total U in Rifle sediments is 0.4-0.9 mg/kg, see Bargar et al., 2013Table 1). In-well columns in which the concentration of U(VI) was increased by amendment with 20 µM uranyl do enable both reliable bicarbonate extractions and synchrotron X-ray spectroscopy (Alessi et al., 2012, Alessi et al., 2014, Bargar et al., 2013, Cerrato et al., 2013). In these studies, limited sampling (e.g., 3 time points over 102 days, Alessi et al., 2014) precludes calculation of rates comparable to the simulated rates reported here. Sediments from the in-well columns do show the prevalence of U(IV) (90%), two thirds of which is non-crystalline U(IV) (Alessi et al., 2014). Broadly the thermodynamics of uraninite and non-crystalline U(VI) formation from soluble U(VI) are likely similar since both involve the same electron transitions, supporting the model use of uraninite formation constants for this study. However, the reactivity of non-crystalline U(VI) is clearly greater than that of uraninite (<u>Cerrato et al., 2013</u>). Future reactive transport models will be more useful for assessing the post-bioreduction behavior of U(IV) if they incorporate the differences in reactivity between uraninite and non-crystalline U(IV). Fe(III) concentrations in <u>aquifer</u> sediments provide another example of complex relationships among geochemical reactions. In the reactive transport model described in this paper, neither physical nor geochemical heterogeneity is required in the model for it to match observations, suggesting that the biogeochemical process behaviors dominate any impacts of heterogeneity in this experiment. Modeling of the same experimental data by <u>Bao et al. (2014)</u> is based on a fundamentally different view of the role of geochemical subsurface heterogeneity in Fe(III). In the <u>Bao et al. (2014)</u>model, an inverse relationship between permeability and the concentration of bioavailable oxidized Fe is posited for the subsurface aquifer hosting the Super 8 experiment. The model simulation of Bao et al. did show higher bioreduction rates in the bicarbonate injection zone, similar to the simulation results presented here. The Bao et al. rates are, on average, similar, but their simulations show significant heterogeneity in bioreduction rates, which derives from the variable content of Fe(III)-containing

minerals in the model domain inferred from the correlation between Fe(III) <u>mineral, clay</u> content, and the associated <u>hydraulic conductivity</u> based on the <u>borehole flow meter</u>measurement. Another key difference between the two models is that the fully 3-D model described in this paper accounts for changes in the saturated thickness and gradient magnitude, whereas the 2-D Bao et al. model uses a constant <u>pressure gradient</u> and saturated thickness. The differences in the two models offers opportunity for detailed comparison in future research that would make it possible to determine if biogeochemical heterogeneity, including reduced phases (<u>Borch et al., 2012</u>, <u>Bargar et al.,</u> <u>2013</u>, <u>Zhao et al., 2013</u>), or the fundamental capabilities of the microbial community are more important in determining outcomes of U(VI) bioreduction at Rifle. Results of such a determination would have broad implications for subsurface biogeochemical processes globally.

# 5. Conclusions

The <u>field experiment</u> described here compares, for the first time, bicarbonatepromoted <u>uranium desorption</u> and <u>acetate</u> amendment within the same subsurface experimental plot. Enzymatic U(VI)-reduction was not inhibited by the addition of <u>bicarbonate</u> and the associated increase in the predominance of Ca-uranyl-carbonato aqueous complexes. Instead, the simulated peak rate of U(VI)-reduction during acetate-bicarbonate amendment was ~3.8 times higher than under acetate-only conditions. Lack of statistically significant differences between the microbial populations under conditions of elevated versus background concentrations of bicarbonate suggest that metabolic processes associated with increased bicarbonate concentrations may be responsible for the increased rate of bioreduction of U(VI); however, additional research is needed to elucidate the mechanisms involved. Overall, the experiment indicates that, under the conditions studied, the efficacy of <u>bioremediation</u> of uranium in <u>aquifers</u> can be enhanced by bicarbonate amendment, and that natural cycling of the redox status of uranium in the subsurface may be more sensitive to bicarbonate concentrations than previously thought.

# Acknowledgements

This work was supported by the Director, Office of Science, Biological and Environmental Research, Subsurface <u>Biogeochemistry</u> Program of the U.S. Department of Energy under Contract No. <u>DE-AC02-05CH11231</u>. We thank the entire Rifle, Colorado Integrated Field Challenge (IFRC) team for their contributions to the Super 8 Experiment, including the U.S. Department of Energy, Grand Junction Office for their excellent field support. We also thank Jason Greenwood for assistance with Fig. 1, Roelof Versteeg for help with uploading data to <u>Pangaea</u>, and anonymous Reviewer #2 for extensive, detailed and very helpful comments that significantly improved the manuscript.

Appendix A. Supplementary data

Download Word document (1MB)

# Help with docx files

Supplementary data 1. This document file contains Supplementary Materials.

# Research data for this article

#### PANGAEA

Data Publisher for Earth & Environmental Science

Data associated with the article:

(Tab 1) Biogeochemistry of groundwater during acetate and bicarbonate amendment to an alluvial aquifer along the Colorado River in Rifle, Colorado, USA

(Tab 3) Signal to noise ratio (SNR) by genus for microarray probes in an alluvial aquifer along the Colorado River in Rifle, Colorado, USA

<u>Biogeochemistry and microbiology of groundwater during acetate and bicarbonate amendment to an alluvial aquifer,</u> <u>supplement to: Long, Philip E; Williams, Kenneth H; Davis, James A; Fox, Patricia M; Wilkins, Michael J; Yabusaki, Steven</u> <u>B; Fang, Yilin; Waichler, Scott R; Berman,...</u>

(Tab 2) Average signal to noise ratio (SNR) for individual microarray probes in an alluvial aquifer along the Colorado River in Rifle, Colorado, USA

# References

### Alessi et al., 2012

D.S. Alessi, B. Uster, H. Veeramani, E.I. Suvorova, J.S. Lezama-Pacheco, J.E. Stubbs, J.R. Bargar, R. Bernier-LatmaniQuantitative separation of monomeric U(IV) from UO<sub>2</sub> in products of U(VI) reduction

Environ. Sci. Technol., 46 (2012), pp. 6150-6157, <u>10.1021/Es204123z</u>

CrossRefView Record in Scopus

# Alessi et al., 2014

D.S. Alessi, J.S.L. Pacheco, N. Janot, E.I. Suvorova, J.M. Cerrato, D.E.Giammar, J.A. Davis, P.M. Fox, K.H. Williams, P.E. Long, K.M. Handley, R. Bernier-Latmani, J. Bargar**Speciation and reactivity of uranium products formed during in** situ bioremediation in a shallow alluvial aquifer

Environ. Sci. Technol., 48 (2014), pp. 12842-12850, 10.1021/es502701u

CrossRefView Record in Scopus

Amann et al., 1990

R.I. Amann, B.J. Binder, R.J. Olson, S.W. Chisholm, R. Devereux, D.A.Stahl**Combination of 16s ribosomal-RNA**targeted oligonucleotide probes with flow-cytometry for analyzing mixed microbial-populations

Appl. Environ. Microb., 56 (1990), pp. 1919-1925

View Record in Scopus

Andersen et al., 2010

G.L. Andersen, Z. He, T.Z. DeSantis, E.L. Brodie, J. Zhou**The use of microarrays in microbial ecology** Environ. Mol. Microbiol. (2010), pp. 87-109

View Record in Scopus

Anderson et al., 2003

R.T. Anderson, H.A. Vrionis, I. Ortiz-Bernad, C.T. Resch, P.E. Long, R.Dayvault, K. Karp, S. Marutzky, D.R. Metzler, A. Pe acock, D.C. White, M. Lowe, D.R.Lovley**Stimulating the in situ activity of** *Geobacter* species to remove uranium from the groundwater of a uranium-contaminated aquifer

Appl. Environ. Microb., 69 (2003), pp. 5884-5891, 10.1128/Aem.69.10.5884-5891.2003

CrossRefView Record in Scopus

#### Bao et al., 2014

C. Bao, H. Wu, L. Li, K.H. Williams, P. Long, D. Newcomer, C. Steefel**Uranium bioreduction rates across scales: biogeochemical "Hot Moments" and "Hot Spots" during a field biostimulation experiment at Rifle, Colorado** Environ. Sci. Technol. Rev., 48 (2014), pp. 10116-10127, <u>10.1021/es501060d</u>

# CrossRefView Record in Scopus

#### Bargar et al., 2013

J.R. Bargar, K.H. Williams, K.M. Campbell, P.E. Long, J.E. Stubbs, E.I.Suvorova, J.S. Lezama-Pacheco, D.S. Alessi, M. S tylo, S.M. Webb, J.A. Davis, D.E.Giammar, L.Y. Blue, R. Bernier-Latmani**Uranium redox transition pathways in** 

### acetate-amended sediments

Proc. Natl. Acad. Sci. U.S.A., 110 (2013), pp. 4506-4511, 10.1073/Pnas.1219198110

### CrossRefView Record in Scopus

#### Berman et al., 2009

E.S.F. Berman, M. Gupta, C. Gabrielli, T. Garland, J.J. McDonnell**High-frequency field-deployable isotope analyzer for hydrological applications** 

Water Resour. Res., 45 (2009), p. W10201, 10.1029/2009wr008265

#### Bernier-Latmani et al., 2010

R. Bernier-Latmani, H. Veeramani, E.D. Vecchia, P. Junier, J.S.Lezama-Pacheco, E.I. Suvorova, J.O. Sharp, N.S. Wiggint on, J.R. Bargar**Non-uraninite products of microbial U(VI) reduction** 

Environ. Sci. Technol., 44 (2010), pp. 9456-9462, 10.1021/Es101675a

#### CrossRefView Record in Scopus

# <u>Bopp et al., 2010</u>

C.J.I. Bopp, C.C. Lundstrom, T.M. Johnson, R.A. Sanford, P.E. Long, K.H.Williams**Uranium 238U/235U isotope ratios as** indicators of reduction: results from an in situ biostimulation experiment at Rifle, Colorado, U.S.A

Environ. Sci. Technol., 44 (2010), pp. 5927-5933, <u>10.1021/es100643v</u>

# CrossRefView Record in Scopus

# Borch et al., 2012

T. Borch, N. Roche, T.E. Johnson**Determination of contaminant levels and remediation efficacy in groundwater at a** former in situ recovery uranium mine

J. Environ. Monit., 14 (2012), pp. 1814-1823, <u>10.1039/c2em30077j</u>

### CrossRefView Record in Scopus

### Campbell et al., 2011

K.M. Campbell, J. Davis, J. Bargar, D. Giammar, R. Bernier-Latmani, R.Kukkadapu, K. Williams, H. Veramani, K.-U. Ulrich , J. Stubbs**Composition, stability, and measurement of reduced uranium phases for groundwater bioremediation** 

### at Old Rifle, CO

Appl. Geochem., 26 (2011), pp. S167-S169

ArticleDownload PDFView Record in Scopus

Cerrato et al., 2013

J.M. Cerrato, M.N. Ashner, D.S. Alessi, J.S. Lezama-Pacheco, R. Bernier-Latmani, J.R. Bargar, D.E. Giammar**Relative** reactivity of biogenic and chemogenic uraninite and biogenic noncrystalline U(IV)

Environ. Sci. Technol., 47 (2013), pp. 9756-9763

### CrossRefView Record in Scopus

### Chandler et al., 2010

D.P. Chandler, A. Kukhtin, R. Mokhiber, C. Knickerbocker, D. Ogles, G.Rudy, J. Golova, P.E. Long, A. Peacock**Monitoring** microbial community structure and dynamics during in situ U(VI) bioremediation with a field-portable microarray analysis system

Environ. Sci. Technol., 44 (2010), pp. 5516-5522, 10.1021/es1006498

#### CrossRefView Record in Scopus

#### Chandler et al., 2013

D.P. Chandler, C. Knickerbocker, L. Bryant, J. Golova, C. Wiles, K.H.Williams, A.D. Peacock, P.E. Long**Profiling in situ** microbial community structure with an amplification microarray

Appl. Environ. Microb., 79 (2013), pp. 799-807

CrossRefView Record in Scopus

#### Curtis et al., 2004

G.P. Curtis, P. Fox, M. Kohler, J.A. DavisComparison of in situ uranium *K*<sub>b</sub> values with a laboratory determined surface complexation model

Appl. Geochem., 19 (2004), pp. 1643-1653

#### ArticleDownload PDFView Record in Scopus

#### Davis et al., 2002

Davis, J. A., Payne, T. E., and Waite, T. D. (2002) Simulating the pH and pCO<sub>2</sub> dependence of uranium(VI) adsorption by a weathered schist with surface complexation models. In *Geochemistry of Soil Radionuclides* (eds. P.-C. Zhang and P. V. Brady). SSSA Special Publication Number 59, Soil Science Society of America, Madison, WI. pp. 61–86, pp. 61–86.

#### Davis et al., 2004

J.A. Davis, D.E. Meece, M. Kohler, G.P. CurtisApproaches to surface complexation modeling of uranium (VI) adsorption on aquifer sediments

Geochim. Cosmochim. Acta, 68 (2004), pp. 3621-3641

ArticleDownload PDFView Record in Scopus

<u>DOE, 1999</u>

DOE. (1999) Final site observational work plan for the UMTRA project old rifle site. GJO-99-88-TAR Rev 1. U.S.

Department of Energy, Grand Junction Office, Grand Junction, Colorado.

DOE, 2001

DOE. (2001) Ground water compliance actin plan for the Old Rifle, Colorado, UMTRA project site. In *Energy* (ed. U.S.D.o.). U.S. Government, Grand Junction, Colorado. p. 18.

### Dong and Brooks, 2006

W. Dong, S.C. BrooksDetermination of the formation constants of ternary complexes of uranyl and carbonate with alkaline earth metals (Mg<sup>2+</sup>, Ca<sup>2+</sup>, Sr<sup>2+</sup>, and Ba<sup>2+</sup>) using anion exchange method

Environ. Sci. Technol., 40 (2006), pp. 4689-4695

# CrossRefView Record in Scopus

### Dong and Brooks, 2008

W.M. Dong, S.C. BrooksFormation of aqueous  $MgUO_2(CO_3)(3)(2-)$  complex and uranium anion exchange mechanism onto an exchange resin

Environ. Sci. Technol., 42 (2008), pp. 1979-1983, 10.1021/Es0711563

#### CrossRefView Record in Scopus

### Druhan et al., 2014a

J.L. Druhan, M. Bill, H. Lim, C. Wu, M.E. Conrad, K.H. Williams, D.J.DePaolo, E.L. Brodie**A large column analog** experiment of stable isotope variations during reactive transport: II. Carbon mass balance, microbial community structure and predation

Geochim. Cosmochim. Acta, 124 (2014), pp. 394-409

ArticleDownload PDFView Record in Scopus

# Druhan et al., 2014b

J.L. Druhan, C.I. Steefel, M.E. Conrad, D.J. DePaoloA large column analog experiment of stable isotope variations during reactive transport: I. A comprehensive model of sulfur cycling and  $\delta_{34}$  S fractionation

Geochim. Cosmochim. Acta, 124 (2014), pp. 366-393

ArticleDownload PDFView Record in Scopus

#### Fang et al., 2009a

Y. Fang, S. Yabusaki, S. Morrison, J.P. Amonette, P. Long**Multicomponent reactive transport modeling of uranium** bioremediation field experiments

Geochim. Cosmochim. Acta, 73 (2009), pp. 6029-6051, 10.1016/j.gca.2009.07.019

ArticleDownload PDFView Record in Scopus

#### Fang et al., 2009b

Y. Fang, S.B. Yabusaki, S.J. Morrison, J.P. Amonette, P.E. LongMulticomponent reactive transport modeling of

uranium bioremediation field experiments

Geochim. Cosmochim. Acta, 73 (2009), pp. 6029-6051

# ArticleDownload PDFView Record in Scopus

#### Fang et al., 2012

Y.L. Fang, M.J. Wilkins, S.B. Yabusaki, M.S. Lipton, P.E. Long**Evaluation of a genome-scale in silico metabolic model** for geobacter metallireducens by using proteomic data from a field biostimulation experiment

Appl. Environ. Microb., 78 (2012), pp. 8735-8742, 10.1128/Aem.01795-12

# CrossRefView Record in Scopus

#### Fox et al., 2012

P.M. Fox, J.A. Davis, M.B. Hay, M.E. Conrad, K.M. Campbell, K.H. Williams, P.E. Long**Rate-limited U(VI) desorption** during a small-scale tracer test in a heterogeneous uranium-contaminated aquifer

Water Resour. Res., 48 (2012), p. W05512, 10.1029/2011WR011472

### Fox et al., 2013

P.M. Fox, J.A. Davis, R. Kukkadapu, D.M. Singer, J. Bargar, K.H. WilliamsAbiotic U (VI) reduction by sorbed Fe(II) on natural sediments

Geochim. Cosmochim. Acta, 117 (2013), pp. 266-282

http://dx.doi.org/10.1016/j.gca.2013.05.003

# ArticleDownload PDFView Record in Scopus

### Gallegos et al., 2013

T.J. Gallegos, C.C. Fuller, S.M. Webb, W. Betterton Uranium (VI) interactions with mackinawite in the presence and

#### absence of bicarbonate and oxygen

Environ. Sci. Technol., 47 (2013), pp. 7357-7364, 10.1021/Es400450z

CrossRefView Record in Scopus

#### Giloteaux et al., 2013

L. Giloteaux, D.E. Holmes, K.H. Williams, K.C. Wrighton, M.J. Wilkins, A.P. Montgomery, J.A. Smith, R. Orellana, C.A. Tho mpson, T.J. Roper, P.E. Long, D.R.Lovley**Characterization and transcription of arsenic respiration and resistance genes during in situ uranium bioremediation** 

ISME J., 7 (2013), pp. 370-383, 10.1038/Ismej.2012.109

CrossRefView Record in Scopus

#### Gorby and Lovley, 1992

Y.A. Gorby, D.R. Lovley Enzymatic uranium precipitation

Environ. Sci. Technol., 26 (1992), pp. 205-207, 10.1021/Es00025a026

#### CrossRefView Record in Scopus

# Guillaumont et al., 2003

R. Guillaumont, T. Fanghänel, J. Fuger, I. Grenthe, V. Neck, D.A.Palme, M.H. Rand**Update on the Chemical Thermodynamics of Uranium, Neptunium, Plutonium, Americium and Technetium** 

Elsevier, Amsterdam (2003)

#### Handley et al., 2012

K.M. Handley, K.C. Wrighton, Y.M. Piceno, G.L. Andersen, T.Z. DeSantis, K.H. Williams, M.J. Wilkins Mj, A. Lucie N'Guessan, A. Peacock, J. Bargar, P.E. Long, J.F. Banfield**High-density PhyloChip profiling of stimulated aquifer microbial communities reveals a complex response to acetate amendment** 

FEMS Microbiol. Ecol., 81 (2012), pp. 188-204, <u>10.1111/j.1574-6941.2012.01363.x</u>

# CrossRefView Record in Scopus

#### Handley et al., 2013

K.M. Handley, N.C. VerBerkmoes, C.I. Steefel, K.H. Williams, I. Sharon, C.S. Miller, K.R. Frischkorn, K. Chourey, B.C. Tho mas, M.B. Shah, P.E. Long, R.L.Hettich, J.F. Banfield**Biostimulation induces syntrophic interactions that impact C, S** and N cycling in a sediment microbial community

ISME J., 7 (2013), pp. 800-816, 10.1038/Ismej.2012.148

# CrossRefView Record in Scopus

### Handley et al., 2014

K.M. Handley, K.C. Wrighton, C.S. Miller, M.J. Wilkins, R.S. Kantor, B.C.Thomas, K.H. Williams, J.A. Gilbert, P.E. Long, J.
F. BanfieldDisturbed subsurface microbial communities follow equivalent trajectories despite different structural starting points

Environ. Microbiol. (2014), 10.1111/1462-2920.12467

#### Herbelin and Westall, 1999

Herbelin, A. L., Westall, J. C. (1999) FITEQL 4.0: a computer program for determination of chemical equilibrium constants from experimental data. Oregon State University Report 99-01, Corvallis, Oregon.

#### Holmes et al., 2007a

D.E. Holmes, R.A. O'Neil, H.A. Vrionis, L.A. N'Guessan, I. Ortiz-Bernad, M.J. Larrahondo, L.A. Adams, J.A. Ward, J.S. Ni coll, K.P. Nevin, M.A. Chavan, J.P.Johnson, P.E. Long, D.R. Lovley**Subsurface clade of Geobacteraceae that** predominates in a diversity of Fe(III)-reducing subsurface environments

ISME J., 1 (2007), pp. 663-677, <u>10.1038/Ismej.2007.85</u>

CrossRefView Record in Scopus

Holmes et al., 2007b

D.E. Holmes, R.A. O'Neil, H.A. Vrionis, L.A. N'Guessan, I. Ortiz-Bernad, M.J. Larrahando, L.A. Adams, J.E. Ward, J.S. Ni coll, K.P. Nevin, M.A. Chavan, J.P.Johnson, P.E. Long, D.R. Lovley**Subsurface clade of Geobacteraceae that** 

predominates in a diversity of Fe(III)-reducing subsurface environments

ISME J., 1 (2007), pp. 663-677

CrossRefView Record in Scopus

#### Holmes et al., 2013a

D.E. Holmes, L. Giloteaux, M. Barlett, M.A. Chavan, J.A. Smith, K.H.Williams, M. Wilkins, P. Long, D.R. Lovley**Molecular** analysis of the in situ growth rates of subsurface *Geobacter* species

Appl. Environ. Microb., 79 (2013), pp. 1646-1653, 10.1128/aem.03263-12

#### CrossRefView Record in Scopus

#### Holmes et al., 2013b

D.E. Holmes, L. Giloteaux, K.H. Williams, K.C. Wrighton, M.J. Wilkins, C.A. Thompson, T.J. Roper, P.E. Long, D.R. Lovley Enrichment of specific protozoan populations during in situ bioremediation of uranium-contaminated groundwater

ISME J., 7 (2013), pp. 1286-1298, 10.1038/ismej.2013.20

#### CrossRefView Record in Scopus

#### Hwang et al., 2009

C.C. Hwang, W.M. Wu, T.J. Gentry, J. Carley, G.A. Corbin, S.L. Carroll, D.B. Watson, P.M. Jardine, J.Z. Zhou, C.S. Criddl e, M.W. Fields**Bacterial community succession during in situ uranium bioremediation: spatial similarities along** 

# controlled flow paths

ISME J., 3 (2009), pp. 47-64, <u>10.1038/Ismej.2008.77</u>

#### CrossRefView Record in Scopus

#### Hyun et al., 2009

S.P. Hyun, P.M. Fox, J.A. Davis, K.M. Campbell, K.F. Hayes, P.E. Long**Surface complexation modeling of U(VI)** adsorption by aquifer sediments from a former mill tailings site at Rifle, Colorado

Environ. Sci. Technol., 43 (2009), pp. 9368-9373

### CrossRefView Record in Scopus

#### Hyun et al., 2012

S.P. Hyun, J.A. Davis, K. Sun, K.F. Hayes**Uranium (VI) reduction by iron (II) monosulfide mackinawite** Environ. Sci. Technol., 46 (2012), pp. 3369-3376

CrossRefView Record in Scopus

### Junier et al., 2009

P. Junier, M. Frutschi, N.S. Wigginton, E.J. Schofield, J.R. Bargar, R.Bernier-Latmani**Metal reduction by spores of** desulfotomaculum reducens

desunotomaculum reducens

Environ. Microbiol., 11 (2009), pp. 3007-3017, 10.1111/J.1462-2920.2009.02003.X

# CrossRefView Record in Scopus

# Junier et al., 2011

P. Junier, E. Dalla Vecchia, R. Bernier-LatmaniThe response of desulfotomaculum reducens MI-1 to U(VI) exposure:

a transcriptomic study

Geomicrobiol. J., 28 (2011), pp. 483-496

10(1080/01490451), 2010, 512031

# CrossRefView Record in Scopus

Kelly et al., 2008

S.D. Kelly, K.M. Kemner, E.J. O'Loughlin, W.M. Wu, C. Criddle, T.L. Marsh**Monitoring uranium transformations** Geochim. Cosmochim. Acta, 72 (2008) A462–A462

#### Komlos et al., 2008

J. Komlos, A. Peacock, R.K. Kukkadapu, P.R. JafféLong-term dynamics of uranium reduction/reoxidation under low sulfate conditions

Geochim. Cosmochim. Acta, 72 (2008), pp. 3603-3615

ArticleDownload PDFView Record in Scopus

# Kostka et al., 2009

J.E. Kostka, S. Green, W. Overholt, W.M. Wu, C. Criddle, D.B. Watson, P.Jardine**Subsurface microbial community** structure correlates with uranium redox phases during in situ field manipulation in a contaminated aquifer Geochim. Cosmochim. Acta, 73 (2009)

A688–A688

#### Lane et al., 1985

D.J. Lane, B. Pace, G.J. Olsen, D.A. Stahl, M.L. Sogin, N.R. Pace**Rapid-determination of 16s ribosomal-RNA** sequences for phylogenetic analyses

Proc. Natl. Acad. Sci. U.S.A., 82 (1985), pp. 6955-6959, 10.1073/Pnas.82.20.6955

CrossRefView Record in Scopus

#### Larese-Casanova et al., 2010

P. Larese-Casanova, S.B. Haderlein, A. Kappler**Biomineralization of lepidocrocite and goethite by nitrate-reducing** Fe(II)-oxidizing bacteria: effect of pH, bicarbonate, phosphate, and humic acids

Geochim. Cosmochim. Acta, 74 (2010), pp. 3721-3734

ArticleDownload PDFView Record in Scopus

### Li et al., 2009

L. Li, C.I. Steefel, K.H. Williams, M.J. Wilkins, S.S. Hubbard**Mineral transformation and biomass accumulation** associated with uranium bioremediation at Rifle, Colorado

Environ. Sci. Technol., 43 (2009), pp. 5429-5435

# CrossRefView Record in Scopus

### Li et al., 2010

L. Li, C.I. Steefel, M.B. Kowalsky, A. Englert, S.S. HubbardEffects of physical and geochemical heterogeneities on mineral transformation and biomass accumulation during biostimulation experiments at Rifle, Colorado J. Contam. Hydrol., 112 (2010), pp. 45-63, 10.1016/j.jconhyd.2009.10.006

ArticleDownload PDFView Record in Scopus

### Lovley et al., 1991

D.R. Lovley, E.J.P. Phillips, Y.A. Gorby, E.R. Landa Microbial reduction of uranium

Nature, 350 (1991), pp. 413-416, <u>10.1038/350413a0</u>

# CrossRefView Record in Scopus

### Miller et al., 2011

C.S. Miller, B.J. Baker, B.C. Thomas, S.W. Singer, J.F. Banfield**EMIRGE: reconstruction of full-length ribosomal genes** from microbial community short read sequencing data

Genome Biol., 12 (2011), p. 12

CrossRefView Record in Scopus

Miller et al., 2013

C.S. Miller, K.M. Handley, K.C. Wrighton, K.R. Frischkorn, B.C. Thomas, J.F.Banfield**Short-read assembly of full-length 16S amplicons reveals bacterial diversity in subsurface sediments** 

PLoS One, 8 (2013), p. e56018, 10.1371/journal.pone.0056018

#### **CrossRef**

#### Morrison et al., 1995

S.J. Morrison, R.R. Spangler, V.S. TripathiAdsorption of uranium(VI) on amorphous ferric oxyhydroxide at highconcentrations of dissolved carbon(IV) and sulfur(Vi)

J. Contam. Hydrol., 17 (1995), pp. 333-346

ArticleDownload PDFView Record in Scopus

# N'Guessan et al., 2008

A.L. N'Guessan, H.A. Vrionis, C.T. Resch, P.E. Long, D.R. Lovley**Sustained removal of uranium from contaminated** groundwater following stimulation of dissimilatory metal reduction

Environ. Sci. Technol., 42 (2008), pp. 2999-3004

#### CrossRefView Record in Scopus

### Newsome et al., 2014

L. Newsome, K. Morris, J.R. Lloyd**The biogeochemistry and bioremediation of uranium and other priority** radionuclides

Chem. Geol., 363 (2014), pp. 164-184

ArticleDownload PDFView Record in Scopus

#### Ortiz-Bernad et al., 2004a

I. Ortiz-Bernad, R.T. Anderson, H.A. Vrionis, D.R. LovleyResistance of solid-phase U(VI) to microbial reduction

during in situ bioremediation of uranium-contaminated groundwater

Appl. Environ. Microb., 70 (2004), pp. 7558-7560, 10.1128/Aem.70.12.7558-7560.2004

### CrossRefView Record in Scopus

# Ortiz-Bernad et al., 2004b

I. Ortiz-Bernad, R.T. Anderson, H.A. Vrionis, D.R. LovleyVanadium respiration by Geobacter metallireducens: novel strategy for in situ removal of vanadium from groundwater

Appl. Environ. Microb., 70 (2004), pp. 3091-3095, 10.1128/Aem.70.5.3091-3095.2004

CrossRefView Record in Scopus

Phillips et al., 1995

E.J.P. Phillips, E.R. Landa, D.R. Lovley**Remediation of uranium contaminated soils with bicarbonate extraction and** microbial U(Vi) reduction

J. Ind. Microbiol., 14 (1995), pp. 203-207, <u>10.1007/Bf01569928</u>

### CrossRefView Record in Scopus

#### Shiel et al., 2013

A.E. Shiel, P.G. Laubach, T.M. Johnson, C.C. Lundstrom, P.E. Long, K.H.WilliamsNo measurable changes in 238U/235U due to desorption–adsorption of U (VI) from groundwater at the Rifle, Colorado, integrated field research challenge site

# Environ. Sci. Technol., 47 (2013), pp. 2535-2541

### CrossRefView Record in Scopus

### Stewart et al., 2007

B.D. Stewart, J. Neiss, S. FendorfQuantifying constraints imposed by calcium and iron on bacterial reduction of uranium(VI)

#### J. Environ. Qual., 36 (2007), pp. 363-372

#### CrossRefView Record in Scopus

#### Stoliker et al., 2013

D. Stoliker, K. Campbell, P.M. Fox, D. Singer, N. Kaviani, M. Carey, N.E.Peck, J. Bargar, D. Kent, J.A. Davis**Evaluating** chemical extraction techniques for the determination of uranium oxidation state in reduced aquifer sediments Environ. Sci. Technol., 47 (2013), pp. 9225-9232, <u>10.1021/es401450v</u>

CrossRefView Record in Scopus

#### Stylo et al., 2013

M. Stylo, D.S. Alessi, P.P. Shao, J.S. Lezama-Pacheco, J.R. Bargar, R.Bernier-LatmaniBiogeochemical controls on the product of microbial U(VI) reduction

Environ. Sci. Technol., 47 (2013), pp. 12351-12358

#### CrossRefView Record in Scopus

#### Suzuki et al., 2005

Y. Suzuki, S.D. Kelly, K.M. Kemner, J.F. Banfield**Direct microbial reduction and subsequent preservation of uranium** in natural near-surface sediment

Appl. Environ. Microb., 71 (2005), pp. 1790-1797

CrossRefView Record in Scopus

# Tang et al., 2013

G.P. Tang, W.M. Wu, D.B. Watson, J.C. Parker, C.W. Schadt, X.Q. Shi, S.C.Brooks**U(VI) bioreduction with emulsified vegetable oil as the electron donor – microcosm tests and model development** 

Environ. Sci. Technol., 47 (2013), pp. 3209-3217, 10.1021/Es304641b

#### CrossRefView Record in Scopus

#### Ulrich et al., 2011

K.-U. Ulrich, H. Veeramani, R. Bernier-Latmani, D.E. Giammar**Speciation-dependent kinetics of uranium(VI)** 

# bioreduction

Geomicrobiol. J., 28 (2011), pp. 396-409

# CrossRefView Record in Scopus

# Williams et al., 2011

K.H. Williams, P.E. Long, J.A. Davis, C.I. Steefel, M.J. Wilkins, A.L.N'Guessan, L. Yang, D. Newcomer, F.A. Spane, L.J. K erkhof, L. McGuinness, R.Dayvault, D.R. Lovely**Acetate availability and its influence on sustainable bioremediation** 

# of uranium-contaminated groundwater

Geomicrobiol. J., 28 (2011), pp. 519-539

10(1080/01490451), 2010, 520074

#### CrossRefView Record in Scopus

#### Wrighton et al., 2012

K.C. Wrighton, B.C. Thomas, I. Sharon, C.S. Miller, C.J. Castelle, N.C.VerBerkmoes, M.J. Wilkins, R.L. Hettich, M.S. Lipto n, K.H. Williams, P.E. LONG, J.F.Banfield**Fermentation, hydrogen and sulfur metabolism in multiple uncultivated** 

# bacterial phyla

Science, 337 (2012), pp. 1661-1665, <u>10.1126/science.1224041</u> <u>CrossRefView Record in Scopus</u> <u>Wrighton et al., 2014</u> K.C. Wrighton, C.J. Castelle, M.J. Wilkins, L.A. Hug, I. Sharon, B.C.Thomas, K.M. Handley, S.W. Mullin, C.D. Nicora, A. Si ngh, M.S. Lipton, P.E. Long, K.H. Williams, J.F. Banfield**Metabolic interdependencies between phylogenetically novel** *fermenters and respiratory organisms in an unconfined aquifer* 

ISME J. (2014), <u>10.1038/ismej.2013.249</u>

#### Wu et al., 2006

Q. Wu, R.A. Sanford, F.E. Loffler**Uranium(VI) reduction by** *Anaeromyxobacter dehalogenans* strain 2CP-C Appl. Environ. Microb., 72 (2006), pp. 3608-3614, <u>10.1128/Aem.72.5.3608-3614.2006</u>

# CrossRefView Record in Scopus

# Wu et al., 2010

W.M. Wu, J. Carley, S.J. Green, J. Luo, S.D. Kelly, J. Van

Nostrand, K. Lowe, T. Mehlhorn, S. Carroll, B. Boonchayanant, F.E. Lofller, D. Watson, K.M. Kemner, J.Zhou, P.K. Kitanidi s, J.E. Kostka, P.M. Jardine, C.S. Criddle**Effects of nitrate on the stability of uranium in a bioreduced region of the** subsurface

Environ. Sci. Technol., 44 (2010), pp. 5104-5111, 10.1021/es1000837

CrossRefView Record in Scopus

#### Xu et al., 2010

M. Xu, W.M. Wu, L. Wu, Z. He, J.D. Van Nostrand, Y. Deng, J. Luo, J. Carley, M. Ginder-

Vogel, T.J. Gentry, B. Gu, D. Watson, P.M. Jardine, T.L. Marsh, J.M. Tiedje, T. Hazen, C.S. Criddle, J. Zhou**Responses of** microbial community functional structures to pilot-scale uranium in situ bioremediation

ISME J., 4 (2010), pp. 1060-1070, <u>10.1038/ismej.2010.31</u>

# CrossRefView Record in Scopus

#### Yabusaki et al., 2007

S.B. Yabusaki, Y. Fang, P.E. Long, C.T. Resch, A.D. Peacock, J. Komlos, P.R. Jaffe, S.J. Morrison, R.D. Dayvault, D.C. W hite, R.T. Anderson**Uranium removal from groundwater via in situ biostimulation: field-scale modeling of transport and biological processes** 

J. Contam. Hydrol., 93 (2007), pp. 216-235, <u>10.1016/j.jconhyd.2007.02.005</u>

ArticleDownload PDFView Record in Scopus

#### Yabusaki et al., 2011

S.B. Yabusaki, Y. Fang, K.H. Williams, C.J. Murray, A.L. Ward, R.D.Dayvault, S.R. Waichler, D.R. Newcomer, F.A. Spane, P.E. Long**Variably saturated flow and multicomponent biogeochemical reactive transport modeling of a uranium bioremediation field experiment** 

J. Contam. Hydrol., 126 (2011), pp. 271-290, <u>10.1016/j.jconhyd.2011.09.002</u>

ArticleDownload PDFView Record in Scopus

### Yelton et al., 2013

A. Yelton, K.H. Williams, J. Fournelle, K.C. Wrighton, K.M. Handley, J.F.Banfield**Vanadate and acetate biostimulation of contaminated sediments decreases diversity, selects for specific taxa and decreases aqueous V5+ concentration** Environ. Sci. Technol., 47 (2013), pp. 6500-6509, <u>10.1021/es4006674</u>

### CrossRefView Record in Scopus

### Zhao et al., 2013

J. Zhao, T.D. Scheibe, R. MahadevanModel-based analysis of mixed uranium(VI) reduction by biotic and abiotic pathways during in situ bioremediation

Chem. Geol., 357 (2013), pp. 215-222, <u>10.1016/j.chemgeo.2013.08.037</u>

ArticleDownload PDFView Record in Scopus

<sup>1</sup>Current address: School of Earth Sciences and Department of Microbiology, The Ohio State University, Columbus, OH 43210, USA.

<sup>2</sup>Current address: Department of Ecology and Evolution, University of Chicago, Chicago, IL 60637, USA.