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Electrochemical Stimulation of Microbial Perchlorate Reduction

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As part of our studies into the diversity of dissimilatory perchlorate reducing bacteria (DPRB) we investigated the reduction of perchlorate in the cathodic chamber of a bioelectrical reactor (BER). Our results demonstrated that washed cells of *Dechloromonas* and *Azospira* species readily reduced 90 mg L⁻¹ perchlorate in the BER with 2,6-anthraquinone disulfonate (AQDS) as a mediator. No perchlorate was reduced in the absence of cells or AQDS, or in an open-circuit control. Similar results were observed when a natural microbial community was inoculated into a fed-batch BER. After 70 days of operation, a novel DPRB, strain VDY, was isolated which readily reduced perchlorate in a mediatorless BER. Continuous up-flow BERs (UFBERS) were seeded with active cultures of strain VDY, and perchlorate at a volumetric loading of 60 mg L⁻¹ day⁻¹ was successfully removed. Gas phase analysis indicated that low levels of H₂ produced at the cathode surface through electrolysis may mediate this metabolism. The results of these studies demonstrate that biological perchlorate remediation can be facilitated through the use of a cathode as the primary electron donor, and that continuous treatment in such a system approaches current industry standards. This has important implications for the continuous treatment of this critical contaminant in industrial waste streams and drinking water.

Introduction

Perchlorate (ClO₄⁻), a soluble anion, is known to affect mammalian thyroid hormone production potentially leading to neonatal neuropsychological development deficiencies (1–3). It is predominantly a synthetic compound with a broad assortment of industrial applications ranging from pyrotechnics to lubricating oils (4). Ammonium perchlorate represents 90% of all perchlorate salts manufactured and is used as an energetics booster or oxidant in solid rocket fuels and munitions (4). Its presence in the environment primarily results from legal historical discharge of unregulated manufacturing waste streams, disposal pond leachate, and the

periodic servicing of military inventories (5–7). Although a powerful oxidant, under most environmental conditions perchlorate is quite stable owing to the high energy of activation associated with its reduction (5, 6). Perchlorate salts readily dissociate in aqueous phases because of the large molecular volume and single anionic charge (5). Furthermore, perchlorate does not significantly sorb to soils or sediments, and in the absence of any biological interactions, its mobility and fate are largely influenced by the hydrology of the environment (8).

Remediation efforts for perchlorate contamination have focused primarily on microbial reduction (5, 9). Many recent studies have demonstrated that specialized microorganisms have evolved that can grow by the anaerobic reductive dissimilation of perchlorate into innocuous chloride (9, 10). More than 40 dissimilatory perchlorate-reducing bacteria (DPRB) are now in pure culture (9), and organisms capable of this metabolism are known to be ubiquitous in soil and sedimentary environments (11) making in-situ treatments relatively straightforward. Furthermore, several bioreactor designs are available for the ex-situ biological attenuation of perchlorate-contaminated waters. Recently, some of these reactor designs were conditionally approved by the California Department of Health Services for application in the treatment of perchlorate contaminated drinking water (URL <http://www.safedrinkingwater.com/archive/sdwn051502.htm>). However, these systems are still dependent on the continual addition of a chemical electron donor to sustain microbial activity and are always subject to biofouling issues (12, 13). In addition, a residual labile electron donor in the reactor effluent can stimulate microbial growth in water distribution systems and contribute to the formation of potentially toxic trihalomethanes during disinfection by chlorination (14, 15). To overcome these problems, chemolithotrophic perchlorate-reducing bioreactors utilizing H₂ as an electron donor have been proposed (16, 17). However, H₂ in bulk quantities is difficult to handle and is publicly perceived as representing a significant disaster threat due to its inherent explosive nature. Alternative inorganic electron donors including Fe(II), S⁰, or H₂S may offer a more practical approach; however, regular additions of these compounds to bioreactors are still required. Furthermore, H₂S is a malodorous toxic compound which can cause corrosion issues while the particulate ferric (hydr)oxides resulting from Fe(II) oxidation result in unpleasant taste and odor, clogged pump and treatment systems, and anodic corrosion of steel pipes and distribution lines (18–20).

Here we investigated the use of a negatively charged electrode (cathode) in the working chamber of a bioelectrical reactor (BER) as an electron donor for microbial perchlorate reduction. In this instance, the DPRB use the electrons on the electrode surface either directly or indirectly through an electron shuttle as a source of reducing equivalents for perchlorate reduction, while assimilating carbon from CO₂ or alternative available carbon sources. Such a process has the advantage of long-term, low-maintenance operation with ease of online monitoring and control while limiting the injection of additional chemicals into the water treatment process. This would negate downstream issues associated with corrosion and biofouling of distribution systems and the production of toxic disinfection byproducts.

Materials and Methods

BER Construction. The BER was constructed with two 50 mm diameter glass chambers with three sample ports, connected with a 30 mm pinch clamp assembly (Laboratory

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Glass Apparatus, Berkeley, CA). Chambers were separated with a cation exchange membrane (Nafion 117). Sample ports were sealed with thick butyl rubber stoppers and aluminum crimp seals. The electrodes were $2.5 \times 1.25 \times 7.5 \text{ cm}^3$ unpolished graphite (G-10 Graphite Engineering and Sales, Greenville, MI) connected with watertight threaded fittings (Impulse, San Diego, CA) to wires and sealed with conductive silver epoxy (Epoxy Technology, Billerica, MA). Wires were fed through stoppers in the top of each chamber. Prior to initial use and before each use thereafter, the electrodes were washed repeatedly in 1 N HCl to remove residual minerals and any biomass buildup and rinsed in sterile deionized water. The silver (Ag/AgCl) reference electrode (Warner Instruments, Hamden, CT) was threaded through a butyl stopper after sterilization with ethanol. Prior to inoculation, the chambers and all stoppers were autoclaved to ensure sterility and subsequently flushed with filter sterilized ($0.22 \mu\text{m}$ pore size) $\text{N}_2\text{--CO}_2$ (80:20; v/v) gas. Throughout the study, the anodic chamber was filled with presterilized, anaerobic 30 mM (1.83 g L^{-1}) bicarbonate buffer (pH 6.8) while the cathodic chamber was filled with either bicarbonate buffer (cell suspensions) or basal growth medium (enrichment/growth studies) outlined below and amended with perchlorate as the sole electron acceptor. Unless otherwise noted, 500 μM AQDS (183 mg L^{-1}) was added as an appropriate electron shuttle. The growth medium was further amended with acetate (5.9 mg L^{-1}) as a suitable carbon source. Both chambers were bubbled continuously with filtered ($0.22 \mu\text{m}$ pore size) $\text{N}_2\text{--CO}_2$ (80:20; v/v) gas to ensure anaerobic operation and maintenance of pH throughout operation. A custom-built potentiostat (Berkeley BioScientific; Dublin, Ireland) was connected to a digital multimeter (Thurlby Thandar Instruments, Cambridgeshire, U.K.) for control of the electrochemical potential.

When run in continuous-flow experiments, 1/16 in. i.d. Norprene tubing (Saint Gobain Performance Plastics, Paris, France) was threaded through 1 mL syringe housings embedded in butyl stoppers in the ports of the cathodic chamber. Flow was established using a peristaltic pump (Rainin, Oakland, CA). The cathodic chamber was filled with powdered graphite that was held in place by a thin (0.5 cm) layer of acid washed sand. A single unpolished graphite (G-10 Graphite Engineering and Sales, Greenville, MI) electrode connected with watertight threaded fittings (Impulse, San Diego, CA) to wires and sealed with conductive silver epoxy (Epoxy Technology, Billerica, MA) was plunged into the powdered graphite bed to ensure electrical conductivity. The influent was pumped in an up-flow mode through the graphite bed, which was maintained at a potential difference of 500 mV relative to the reference electrode. The reactor was inoculated (10% by volume) with an active perchlorate-reducing culture of strain VDY which was spun down and resuspended in an equal volume of anoxic sterile growth media prior to injection into the graphite bed through the influent port.

Medium and Culturing Conditions. All pure cultures were obtained from lab stocks except strain VDY which was isolated in this study. Standard anaerobic culturing techniques were used throughout (21–23). The medium was boiled and cooled under $\text{N}_2\text{--CO}_2$ (80:20) to remove dissolved O_2 and then dispensed into anaerobic pressure tubes or serum bottles under $\text{N}_2\text{--CO}_2$, capped with thick butyl rubber stoppers, and sterilized by autoclaving. The basal medium was the bicarbonate-buffered freshwater medium that had previously been used for culturing perchlorate-reducing bacteria (24). Unless otherwise noted, sodium salts of acetate and perchlorate at 10 mM each (590 and 990 mg L^{-1} , respectively) were added from sterile anoxic aqueous stocks as the electron donor and acceptor, respectively.

Alternative electron donors and acceptors were added from sterile anoxic aqueous stocks prepared from the sodium salts to give final concentrations of 10 mM in the culture medium. All culture incubations were performed at room temperature without agitation in the dark.

Washed Cell Suspensions. Cells were pregrown anaerobically in 1 L volumes with acetate (590 mg L^{-1}) as the electron donor and perchlorate (990 mg L^{-1}) as the sole electron acceptor. After 18 h incubation, the cells were harvested by centrifugation ($3000 \times g$) and washed with anoxic bicarbonate buffer (2.5 g L^{-1}) under a $\text{N}_2\text{--CO}_2$ (80:20; v/v) headspace. Washed cells were resuspended in anoxic bicarbonate buffer (1 mL) and sealed in a 10 mL serum vial with a butyl stopper under the same headspace.

DPRB Isolation. Perchlorate-reducing enrichments were established by transferring 1 g of electrode surface scrapings into 9 mL of prepared anoxic medium under a gas stream of $\text{N}_2\text{--CO}_2$ (80:20; v/v). Acetate (590 mg L^{-1}) was the electron donor, and perchlorate (990 mg L^{-1}) was the electron acceptor. Incubations were done at 30°C in the dark. Positive enrichments were identified by visual increase in optical density and by microscopic examination. Once a positive enrichment was established, the perchlorate-reducing culture was transferred (10% inoculum) into 9 mL of fresh anoxic medium. Isolated colonies were obtained from transfers of positive enrichments by the standard agar shake-tube technique outlined previously (24) with acetate as the sole electron donor and perchlorate (590 and 990 mg L^{-1} , respectively) as the sole electron acceptor.

16s rRNA Gene Sequencing and Analysis. Cells from 10-mL cultures of the isolated perchlorate reducing bacteria were harvested by centrifugation, resuspended in $40 \mu\text{L}$ sterile water, and lysed by the addition of $5 \mu\text{L}$ chloroform with a 10 min incubation at 95°C . Primers specific to bacterial 16S rDNA (8F: $5'\text{-AGAGTTTGATCCTGGCTCAG-3'}$; 1525R: $5'\text{-AAGGAGGTGATCCAGCC-3'}$) were used in a polymerase chain reaction (PCR) that consisted of 10 mM Tris-HCl (pH 9.0), 50 mM KCl, 0.1% Triton X-100, 1.2 mM MgCl_2 , 0.2 mM each dNTP, 75 ng of each primer, $0.5 \mu\text{L}$ Taq polymerase (Gibco/BRL), and $1 \mu\text{L}$ of lysed cells in a $50 \mu\text{L}$ reaction. Amplifications were performed at these parameters: 94°C for 3 min, followed by 30 cycles of 94°C for 1 min, 55°C for 1 min, and 72°C for 2 min with a final incubation of 10 min at 72°C . The amplification products were gel-purified (GeneClean II, BIO101) and cycle sequenced (ThermoSequenase, Amersham) using internal primers. Sequence entry and manipulation was performed with the MacVector 8.1 sequence analysis software program for the Macintosh (Oxford Molecular). Sequences of select 16S rRNAs were downloaded from the Ribosomal Database Project (25) and Genbank (26) into the computer program SeqApp (27). The partial 16S rDNA sequence (636 bp) from strain VDY was manually added to the alignment using secondary structure information for proper alignment. Distance analysis of the aligned sequences was performed using PAUP* 4.0d65 (28).

Analytical Methods. All experimental analyses were performed in triplicate to ensure reproducibility, and the results are expressed as the mean of these determinations. The concentration of perchlorate in cultures was determined using ion chromatography as previously described (29). Cell growth in active cultures was monitored by optical density at 600 nm (OD600), and by total direct cell count using phase-contrast microscopy as previously described (11).

Results

Bioelectrical Reduction of Perchlorate by Pure Cultures of DPRB. Active washed cell suspensions of *Dechloromonas agitata*, *D. aromatica*, and *Azospira suillum*, three representatives of the environmentally dominant perchlorate reducing bacterial genera, readily reduced 99 mg L^{-1} per-

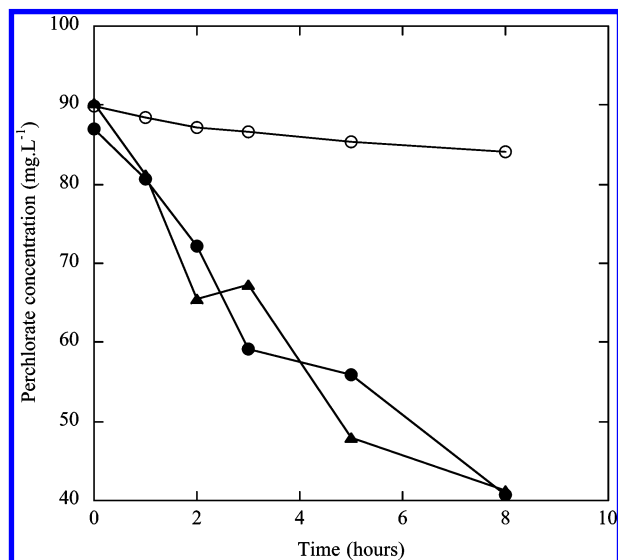


FIGURE 1. Perchlorate reduction by washed whole cell suspensions of *Azospira suillum* strain PS using acetate or the BER cathode as the primary electron donor. Closed circles depict perchlorate reduction in the BER, closed triangles depict perchlorate reduction with acetate (50 mg L^{-1}), and open circles depict perchlorate reduction in the open-circuit control.

chlorate over a 24 h trial when incubated in the cathodic chamber of the electrochemical cell at a poised potential of -450 mV (relative to the Ag reference electrode) containing 183 mg L^{-1} ($500 \mu\text{M}$) 2,6-anthraquinone disulfonate (AQDS) (Figure 1). In all cases, the rate and extent of perchlorate reduction were almost identical to those observed in the positive control to which acetate (59 mg L^{-1}) was added as the sole electron donor (Figure 1). In contrast, no significant perchlorate reduction was observed in identical incubations in which the electrical circuit was incomplete (open-circuit control) (Figure 1), suggesting that perchlorate reduction was an enzymatically mediated reaction using electrons derived from the surface of the cathode. Although perchlorate was continuously removed, none of the tested DPRB (*D. agitata*, *D. aromatica*, or *A. suillum*) grew in the BER (as determined by microscopic observation and change in optical density at 600 nm), when similar experiments were performed in the BER using basal growth medium amended with 5.9 mg L^{-1} acetate as a suitable carbon source (data not shown).

In all incubations, no significant perchlorate reduction occurred relative to the open-circuit control if the AQDS was omitted from the medium even after an extended incubation period of 72 h (data not shown). In order to ensure that the observed perchlorate reduction was not due to the biological catabolism of the added AQDS, perchlorate concentrations were monitored in active anaerobic cultures of *D. agitata* incubated with AQDS as the sole electron donor in sealed serum vials. Again, no perchlorate reduction was observed unless acetate was added as an additional electron donor (data not shown). In contrast, if AH_2DS (the reduced hydroquinone form of AQDS) was provided, *D. agitata* readily reduced the perchlorate with the concomitant oxidation of the AH_2DS to AQDS (Figure 2). In the absence of either perchlorate or cells, no AH_2DS oxidation was apparent, and in the absence of AH_2DS , no perchlorate reduction was observed (data not shown). Analysis of the concentration of AQDS and AH_2DS in the medium during incubation revealed that the total anthraquinone concentration remained constant during incubation and was not being biodegraded as a carbon source by the organisms (data not shown). Furthermore, the oxidation of 1.54 g L^{-1} (4.2 mM) AH_2DS resulted in the reduction of 98 mg L^{-1} (0.99 mM) of

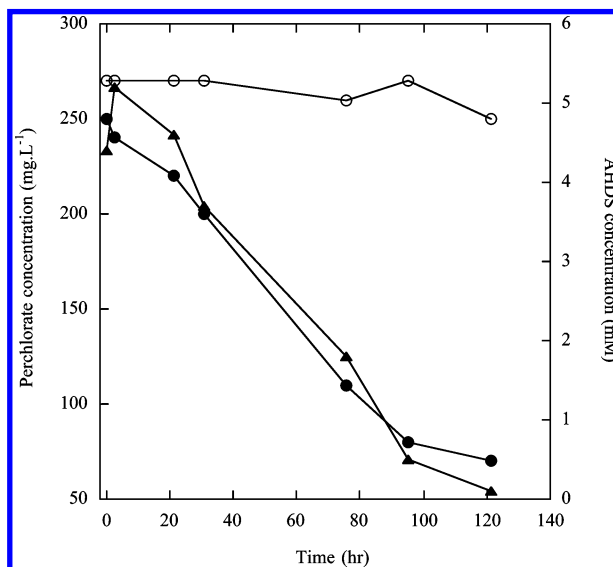
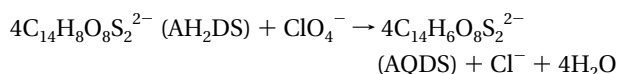


FIGURE 2. Perchlorate reduction coupled to oxidation of the reduced hydroquinone form of the quinone electron shuttle AH_2DS in an anaerobic batch culture by *Dechloromonas agitata* strain CKB. Closed circles depict perchlorate reduction in the presence of 5 mM AH_2DS , closed triangles depict AH_2DS oxidation in the presence of 2.5 mM perchlorate, and open circles depict perchlorate reduction in the presence of 5 mM AQDS.

perchlorate giving a stoichiometry of 4.24, which is 106% of the theoretical value according to



These results indicate that perchlorate reduction in the BER was microbially mediated by electrons being shuttled from the cathode surface to the bacterial cells by the AQDS.

Bioelectrical Reduction of Perchlorate by Natural DPRB Populations. Significant perchlorate removal was observed under growth conditions when the cathodic chamber of the BER was filled with sterile anoxic basal media amended with perchlorate, 183 mg L^{-1} ($500 \mu\text{M}$) AQDS, poised at -500 mV (vs Ag/AgCl), and inoculated with 10% v/v water from Strawberry Creek on the UC Berkeley campus that had been filtered with $11 \mu\text{m}$ filter paper. Over the first three weeks, almost 200 mg L^{-1} perchlorate was removed in the enrichment compared to an open-circuit control (Figure 3). At this time, the medium in both the open- and closed-circuit BERs was amended with 0.1 mg L^{-1} yeast extract for continued cell maintenance. As expected, this resulted in a sharp increase in the amount of perchlorate consumed in both systems, but thereafter perchlorate levels stabilized in the open-circuit control while the closed-circuit BER continued to remove perchlorate. An amendment of perchlorate and an additional 0.1 mg L^{-1} yeast extract at day 30 showed similar results, and after 70 days of operation, the closed-circuit enrichment had removed 1000 mg L^{-1} more perchlorate than the open-circuit control.

Isolation and Characterization of Strain VDY. To identify some of the DPRB present in the BER inoculated with Strawberry Creek groundwater, samples (1 g) were scraped from the surface of the cathode after the 70-day incubation and transferred into fresh basal medium with acetate as the electron donor and perchlorate as the sole electron acceptor. After two weeks incubation, growth was visually apparent in the primary enrichments of these samples. These enrichments were transferred into fresh basal medium (10% inoculum) where good growth was again observed after 24 h as determined by increase in optical density and micro-

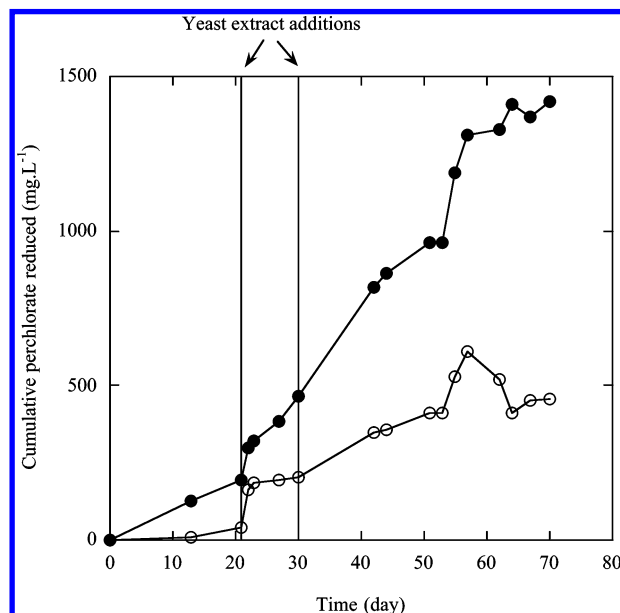


FIGURE 3. Cumulative perchlorate reduction in the bioelectrical reactor and open-circuit control by the indigenous dissimilatory perchlorate-reducing population in Strawberry Creek waters collected from UC Berkeley campus. Closed circles depict cumulative perchlorate removal in the BER, and open circles depict cumulative perchlorate removal in the open-circuit control.

scopic examination. Highly enriched, perchlorate-reducing cultures were obtained by sequential transfer over the following week prior to serial dilution into agar tubes. Small (1–2 mm diameter) pink colonies of consistent morphology were apparent after two weeks of incubation, and a dissimilatory perchlorate-reducing isolate, strain VDY, was selected for further characterization.

Strain VDY was a gram-negative, facultative anaerobe. Cells, 0.2 μm diameter by 7 μm length, showed a consistent spirillum morphology. Strain VDY completely oxidized organic electron donors to CO_2 in the presence of a suitable electron acceptor. Alternatively, strain VDY grew fermentatively in anoxic basal medium amended with glucose (1.80 g L^{-1}), yeast extract (0.1 g L^{-1}), and casamino acids (0.1 g L^{-1}). Spores were not visible in wet-mounts by phase contrast microscopy, and no growth was observed in fresh acetate–perchlorate medium after pasteurization at 80 $^\circ\text{C}$ for 3 min. In addition to acetate, strain VDY used lactate, AH_2DS , ethanol, and H_2 as electron donors and perchlorate, chlorate, nitrate, or O_2 as electron acceptors. Analyses of the partial 16S rDNA sequence (636 bp) indicated that strain VDY was closely related (>99% 16S rDNA sequence identity) to the known DPRB *Dechlorospirillum anomalous* strain WD in the alpha subclass of the proteobacteria (Figure 4).

Pure Culture BER Studies with Strain VDY. As with the other DPRB tested (*D. agitata*, *D. aromatica*, and *A. suillum*), perchlorate was rapidly removed in the cathodic chamber of a BER poised at -500 mV when inoculated with strain VDY (Figure 5). Although no significant growth was observed, the cell density in the BER remained constant throughout the incubation, while that of the open-circuit control rapidly declined (data not shown). No perchlorate removal was observed in the open-circuit control, however, in contrast to the results obtained with the other DPRB, strain VDY was capable of reducing perchlorate in the BER in the absence of the mediator AQDS (Figure 5), although this removal was significantly slower than that in the BER amended with AQDS. Analysis of H_2 production in the BER indicated that under operational conditions $0.78\text{ }\mu\text{g min}^{-1}$ was produced through electrolysis of water at the surface of the cathode which is

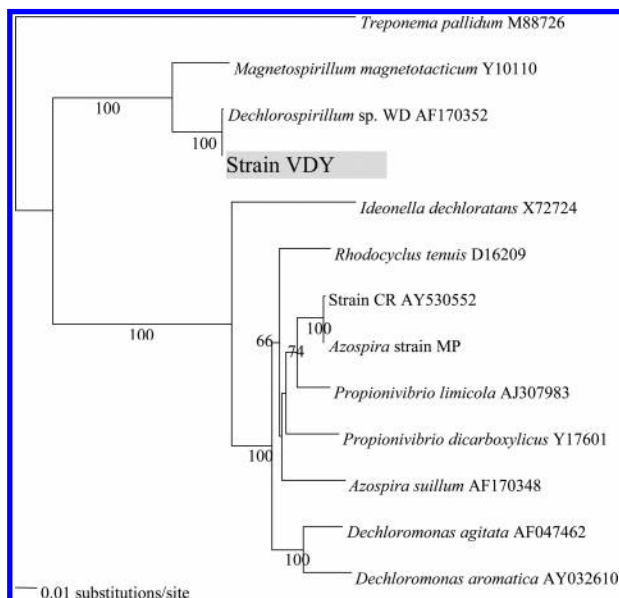


FIGURE 4. Phylogenetic tree resulting from heuristic analysis of a 16S rRNA dataset consisting of 1411 characters using the Kimura 2-parameter distance setting. Bootstrap values of >50% from 500 replicates are indicated near the relevant nodes.

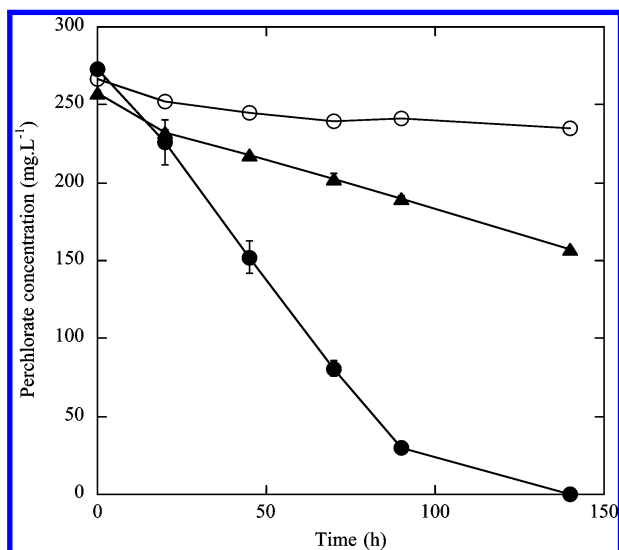
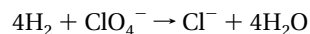


FIGURE 5. Perchlorate removal by strain VDY in a batch culture in the bioelectrical reactor at -500 mV with and without the electron shuttle AQDS. Closed circles depict perchlorate reduction in the BER in the presence of AQDS, closed triangles depict perchlorate reduction in the BER in the absence of AQDS, and open circles depict perchlorate reduction in the open-circuit control.

more than enough reducing equivalents to account for the observed reduction of the perchlorate ($2\text{ }\mu\text{g min}^{-1}$) in the mediator-less BER throughout the incubation assuming a theoretical stoichiometry of



Continuous Treatment of Perchlorate in an UFBER. In order to determine the applicability of the BER system combined with strain VDY to the continuous treatment of perchlorate, bench-scale up-flow bioelectrical reactors were constructed in triplicate in the same BER architecture used for the batch experiments. The active chamber contained a packed bed of graphite powder charged by a graphite electrode inserted into the bed to maximize the surface area available for electron donation by the electrode. The reactors

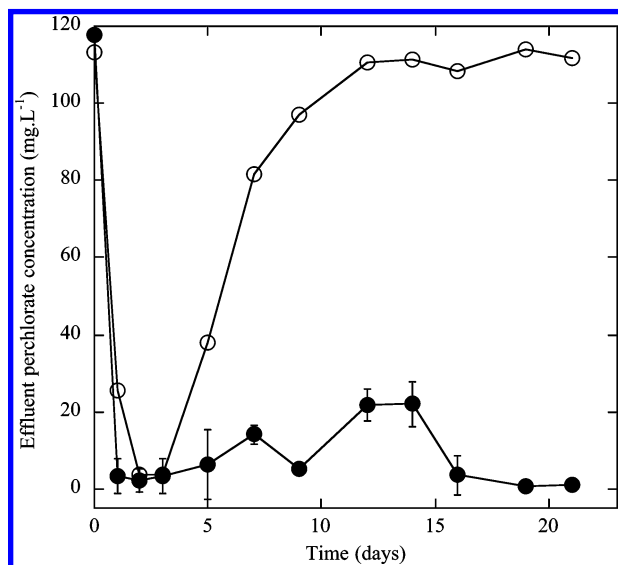


FIGURE 6. Continuous treatment of a perchlorate-containing simulated wastestream amended with 500 μM AQDS as an electron shuttle in the upflow bioelectrical reactor inoculated with strain VDY. The results depicted are the mean of triplicate determinations. Closed circles depict effluent perchlorate concentrations in the UFBER, and open circles depict effluent perchlorate concentrations in the open-circuit control.

were inoculated with anaerobic active washed cells of strain VDY previously grown on acetate and perchlorate. To promote establishment of the DPRB culture in the bed and to provide a starting baseline for the closed- and open-circuit reactors, acetate (82 mg L^{-1}) was added with strain VDY. AQDS was added at 183 mg L^{-1} ($500 \mu\text{M}$) to these initial experiments, and the hydraulic loading rate of perchlorate in the influent was initially set at $60 \text{ mg L}^{-1} \text{ day}^{-1}$ for the first 12 days. As expected in response to the acetate injections, perchlorate levels in the effluent of all reactors were negligible for the first 3 days of operation (Figure 6). However, the perchlorate levels in the open-circuit control soon began to rebound to the influent concentration and stabilized at that level for the remainder of the incubation (Figure 6). In contrast, perchlorate was consistently removed throughout the incubation in the closed-circuit reactors, and at least 80% removal efficiency was continuously achieved in each of the reactors for the three weeks of operation (Figure 6). Variations in effluent perchlorate concentration throughout the incubation were due to adjustments of the reactor residence time, which was optimized to 3 days at day 16, corresponding to a volumetric loading rate of $40 \text{ mg L}^{-1} \text{ day}^{-1}$. After residence time optimization, effluent perchlorate concentrations were consistently below detection for the remainder of the incubation (Figure 6). Furthermore, one of the UFBER replicates was kept operational after the experiment was concluded at day 21, and while there was no concurrent negative control, this reactor successfully maintained nondetectable perchlorate concentrations in the effluent for a further 40 days without any fluctuations in efficiency (data not shown).

While successful for treating perchlorate, the presence of AQDS was not optimal for water treatment as it adds cost to the treatment and would require an additional step to remove it from the treated effluent. Furthermore, headspace gas analysis indicated that significantly more H_2 was produced ($3.72 \mu\text{g min}^{-1}$) in the UFBER configuration than in the batch BERs, probably because of the much larger cathodic surface area, which would theoretically provide enough reducing equivalents to reduce all of the perchlorate at a volumetric loading rate of $66 \text{ mg L}^{-1} \text{ day}^{-1}$. In order to determine the

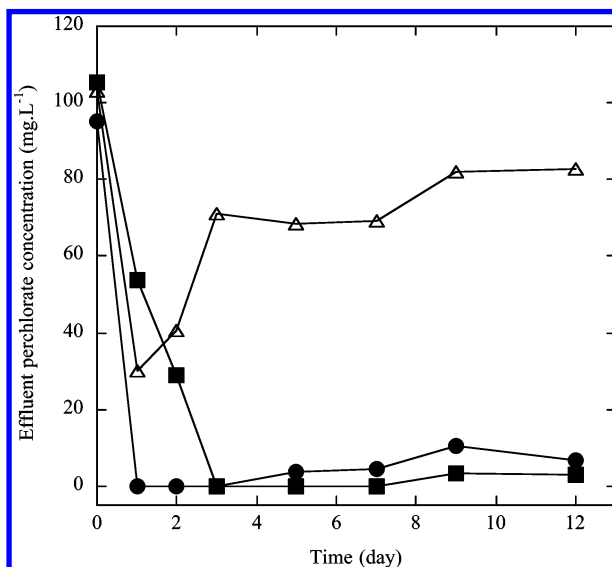


FIGURE 7. Comparison of the continuous-flow treatment of perchlorate in the upflow bioelectrical reactor inoculated with strain VDY in the presence and absence of the electron shuttle AQDS. Closed circles depict effluent perchlorate concentrations in the absence of AQDS, closed squares depict effluent perchlorate concentrations in the presence of $1 \mu\text{M}$ AQDS, and open triangles depict effluent perchlorate concentrations in the open-circuit control.

necessity of the mediator, several UFBERs were constructed, inoculated with an active perchlorate reducing culture of strain VDY, and amended with 59 mg L^{-1} acetate as before. An initial perchlorate volumetric loading rate of approximately $60 \text{ mg L}^{-1} \text{ day}^{-1}$ was set. Where necessary, AQDS was added to the influent to give a final concentration of 0.4 mg L^{-1} ($1 \mu\text{M}$). As predicted, immediate perchlorate removal was apparent in the presence and absence of the AQDS as well as in the open-circuit control because of the initial amendment with acetate. However, the effluent perchlorate content of the open-circuit control rapidly rebounded to levels approaching the influent concentration within 48 h (Figure 7). In contrast, perchlorate removal was consistent both in the presence and absence of AQDS in the closed-circuit UFBERs for the remainder of the operation. Removal efficiencies were essentially identical regardless of the presence or absence of AQDS, and both UFBERs continuously achieved removal efficiencies of greater than 95% (Figure 7), indicating that AQDS was not necessary to achieve effective treatment efficiencies at high loading rates in these reactors.

Discussion

Electrodes as Electron Donors for Microbial Metabolism.

These studies represent the first demonstration that electrodes can serve as a primary electron donor for microbial perchlorate reduction and have resulted in the isolation and characterization of a novel isolate, *Dechlorospirillum* strain VDY, that can exist stably in the BER for extended treatment periods. Previous studies have similarly demonstrated the use of an electrode as the primary electron donor for the dissimilatory reduction of nitrate by *Geobacter* species (30, 31), fumarate by both *Geobacter* and *Actinobacillus* species (30, 32), hexavalent uranium by *Geobacter* species (33), and CO_2 by an undefined enrichment (31). Furthermore, electrochemical reduction of soluble iron by a cathode has also been shown to support growth and CO_2 fixation by the iron-oxidizing *Acidithiobacillus* species (34, 35). In contrast to those studies, however, electrochemically generated H_2 through the electrolysis of water at the cathode surface is likely to play a role in the microbial reduction of perchlorate observed in our BER's amended with strain VDY in the

absence of AQDS. This may also explain the conflicting results obtained with strain VDY and the other DPRB tested in this study, none of which could reduce perchlorate in the BER in the absence of AQDS. This is because H₂ is not utilized as an electron donor by the *Dechloromonas* or *Azospira* species tested in this study (24, 36–38) while, in contrast, physiological characterization revealed that strain VDY could readily use H₂ as an electron donor for respiration.

However, other alternatives to H₂ mediation may also exist. The previous *Geobacter*-based studies showed members of this genus were able to directly access electrons off the electrode surface (30) without the need for electrolytically produced H₂. *Geobacter* species are similarly known to pass electrons directly onto an anodic surface in the absence of electron shuttling compounds when oxidizing organic electron donors (39). Whether or not these organisms use a reversal of a single pathway to mediate these contrasting electron transfers is unknown. Additionally, microbially produced electron shuttles have also been shown to play an important role in electron transfer in the anodic chamber of microbial fuel cells (40). While strain VDY has been shown to utilize hydrogen for the reduction of perchlorate, the possibility still exists that it may also utilize electrons directly from the electrode surface or produce an electron shuttling compound to further supplement its metabolism of perchlorate in the cathodic chamber, and this represents an important future direction for this research. Regardless of the mechanism of electron transfer off the electrode surface, the removal of AQDS or any other additional organic carbon source from the BER system is advantageous for treatment cost-effectiveness as well as effluent water quality.

Reactor Effectiveness. The results obtained in this study indicate that the stimulation of microbial perchlorate reduction in the UFBER is already competitive with existing state of the art bioreactors without any electrochemical optimization. Current high-throughput treatment of concentrated perchlorate wastestreams is over 400 mg L⁻¹ day⁻¹ (41); however, average volumetric loads are one tenth of that. The UFBER in this study showed consistent removal of 60 mg L⁻¹ reactor volume day⁻¹, without addition of a chemical electron donor. This is significant given the inherent inefficiencies of the H-style electrochemical cell employed for these studies. The limitations to microbial stimulation stem from high internal resistance that can be overcome easily with design modifications already in use for a variety of microbial fuel cell (MFC) technologies. These modifications included increased cation-specific membrane surface area or removal of the cation-specific membrane altogether, as well as decreased distances between the anode and cathode. Given the order of magnitude improvements in power production from these types of modifications in MFCs (42), it would be expected that similar modifications would improve UFBER performance for perchlorate treatment.

Significance. The results presented here indicate that microbial perchlorate reduction can be coupled to the removal of electrons off the surface of an electrode. This has important implications with regard to the continuous long-term treatment of perchlorate-contaminated waters and wastestreams. Although previous studies have resulted in the development of various alternative bioreactor designs (13), all of these are limited by the requirement for a continuous addition of a suitable chemical electron donor. Because microbial perchlorate reduction is generally inhibited by the presence of O₂ and to some extent nitrate (29, 43), excess electron donor must be added to biologically remove these components from reactor influents prior to the stimulation of perchlorate reduction. Such additions must be carefully monitored to prevent the presence of residual labile electron donor in the reactor effluent which may result in biofouling of distribution systems and the formation of

trihalomethanes (14, 15). This is especially true if the total electron accepting capacity of the perchlorate present in the contaminated stream is small relative to that of the nitrate and dissolved O₂ content which is the case for most contaminated waters (9). Bioelectrical reduction at the cathode surface overcomes many of these issues because no chemical electron donor is added to the bioreactor.

In conclusion, the results presented here demonstrate the exciting potential for the application of bioelectrical reduction for the treatment of perchlorate contamination without many of the limitations normally associated with bioreactor-based processes.

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