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APPLICATION OF MICROBIAL BIOMASS AND ACTIVITY MEASURES TO ASSESS IN SITU BIOREMEDIATION OF CHLORINATED SOLVENTS

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Key words: bioremediation, activity, monitoring, in situ, biotechnology, microbiology

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INTRODUCTION

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Evaluating the effectiveness of chlorinated solvent remediation in the subsurface can be a significant problem given uncertainties in estimating the total mass of contaminants present. If the remediation technique is a biological activity, information on the progress and success of the remediation may be gained by monitoring changes in the mass and activities of microbial populations.

The in situ bioremediation demonstration at the U.S. Department of Energy (DOE) Savannah River Site (SRS) is designed to test the effectiveness of methane injection for the stimulation of trichloroethylene (TCE) degradation in sediments. Past studies have shown the potential for TCE degradation by native microbial populations (Fliermans et al. 1988). The design and implementation of the SRS Integrated Demonstration is described in this volume (Hazen et al. 1993). A control phase without treatment was followed by a phase withdrawing air. The next phase included vacuum extraction plus air injection into the lower horizontal well located below the water table. The next period included the injection of 1% methane in air followed by injection of 4% methane in air. Based on the literature (Little et al. 1988, Wilson and Wilson 1985), it was hypothesized that the injection of methane would stimulate methanotrophic populations and thus accelerate biological degradation of TCE. Measuring the success of bioremediation is a complex effort that includes monitoring of changes in microbial populations associated with TCE degradation. These monitoring efforts are described in this paper and in related papers (Brockman et al. 1993, etc) in this volume.

METHODS

The primary source of samples for monitoring bioremediation was groundwater obtained from a series of 12 wells (Figure 1, Hazen et al. 1993) located at and around the site at Savannah River.

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Wells were sampled twice monthly and experiments were initiated onsite or samples were stored on ice until processed at the University of Tennessee (UT) or Oak Ridge National Laboratory (ORNL).

Microbial biomass was evaluated using turbidimetric most probable number (MPN) techniques for methanotrophic and methylotrophic populations. A phosphate- and bicarbonate-buffered mineral salts medium, supplemented with 5% methane (vol/vol, headspace), was used to enumerate methanotrophs exhibiting turbidity. The methylotrophic medium was supplemented with 3 mg L⁻¹ of yeast extract, 5 mM methanol, and 3% headspace propane. Microbial activity was assessed by measuring TCE mineralization in time course experiments and enrichments as described previously (Phelps et al. 1989). Assays used 10 mL groundwater, 1.0 mL sterile water and 0.5 μ Ci of carrier-free [1,2-¹⁴C]TCE and were incubated 30 days. Methane and CO₂ were assayed using a Shimadzu GC-8A gas chromatograph (GC) with a thermal conductivity detector (TCD). Radioactive CO₂, VC, TCE, and PCE were quantified using a gas proportional counter (GPC) analyzing the effluents from the gas chromatograph (GC-GPC). Groundwater samples (1 L) were filtered through 0.2 μ m inorganic filters and analyzed for biomass and community structure. Microbial phospholipids were extracted (Phelps et al. 1989) and methyl esters of the phospholipid fatty acids (PLFAs) were analyzed by GC.

Specific nitrogen transformations were evaluated by measuring urease activity (adding urea and measuring ammonia production) and ammonia uptake (by measurement of the loss of ammonia after addition) in well water. Ammonia concentrations were measured using a Technicon autoanalyzer.

RESULTS AND DISCUSSION

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The numbers of methanotrophs and methylotrophs increased significantly during the treatments (Table 1). During the control period, 19% of the analyses indicated presence of methanotrophs in

10mL samples. None of the analyses from the control, vacuum extraction or air injection periods indicated > 10³ methanotrophs/mL. Methanotrophic populations increased dramatically with the initiation of methane injection. Wells exhibiting > 10¹ methanotrophs/ml increased from < 10% to 47% with 1% methane additions and to 85% during the 4% methane injections. During the final 2 months of 1% methane injection, 53% of the samples indicated > 10³ methanotrophs/mL. During the control period, 61% of the samples indicated the presence of > 10¹ methylotrophs/mL. Between the control and air injection periods, the percentage of wells with > 10³ methylotrophs increased from 11% to 81%. These results indicated that methanotrophic and methylotrophic populations increased > 100-fold during the treatments. Increased methanotrophic densities within several individual wells were 4 to 5 orders of magnitude (data not shown). Interestingly, both methanotrophic and methylotrophic densities decreased in many wells during the 4% methane injection phase, suggesting that nutrients or other constraints may have become limiting.

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Further evidence of increased methanotrophic densities was shown by the PLFA analyses (i.e. well MHT-2C, Figure 1). Well MHT-2C showed an increase in lipid biomass from < 5 pmol PLFA/mL to values often > 30 pmol/mL shortly after the onset of methane injection. Increased methanotrophic populations were detected by signature PLFAs of type II methanotrophs resembling *Methylosinus trichosporium* OB3b. When total lipid biomass was compared to mole percent PLFA for type II methanotrophs, the increases in methanotrophic PLFA corresponded to the increases in lipid biomass. By the end of the 1% methane injection, lipids typical of type II methanotrophs, including 18:1w8c, 18:1w7c, 18:2a, and 18:2b, represented more than half of the total PLFA present.

Enrichment experiments examining ¹⁴CO₂ production from [1,2- ¹⁴C]TCE showed potential for TCE degradation throughout the operations at the SRS. For example, during the 1% methane injection, the amount of TCE degradation in the enrichments typically was <10% (Figure 2) but there was substantially more TCE degradation (>20%) in some of the analyses (6C, 7C, 8C).

Maximum TCE degradation in the enrichments was 56% after 30 days. If bioremediation were optimized, most analyses should reveal mineralization within 10 days without the need for enriching nutrients, a criterion that fewer than 5% of the analyses met.

SUMMARY

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There have been substantial changes in biological activity and biomass with the increasingly aggressive measures to promote TCE degradation. The data indicate the success in stimulating TCE-degrading populations. Other nutrients may be limiting, and a further phase (methane injection with nutrient addition) is currently being implemented to further increase critical population levels and degradative activity.

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Treatment	> 0.1/mL Presumptive		> 10 ¹ /mL Presumptive		> 10 ³ /mL Presumptive	
	Methy.	Methane	Methy.	Methane	Methy.	Methane
Control	100	19	61	3	11	0
Vacuum Extraction	100	13	100	8	42	0
Air Injection	100	67	100	8	83	0
1% Methane	100	82	100	47	79	32
4% Methane	100	92	100	85	81	38

Table 1. Average percent of wells with presumptive methanotrophs (methane) and methylotrophs (methy.) at greater than 0.1/mL, 10¹/mL and 10³/mL.





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LIST OF FIGURE AND TABLE CAPTIONS

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- TABLE 1.Average percent of wells with presumptive methanotrophs (methane.) and methylo-
trophs (methy.) at greater than 0.1/mL, 101/mL and 103/mL.
- FIGURE 1. Methanotrophic PLFA and total biomass over time. Mole percent of methanotrophic PLFA (squares) was compared to biomass (diamonds) in well MHT-2C. Type II methanotrophic PLFA included 18:1w8c, 18:1w7c, 18:2a, and 18:2b.
- FIGURE 2. Mineralization of $[1,2^{-14}C]TCE$ to ${}^{14}CO_2$ by microorganisms enriched from groundwater. ${}^{14}CO_2$ in blanks and control tubes after 30-d experiments were $1.4 \pm 1.0\%$.

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