Case Studies on Practical Use of Stable Isotope Probing

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providing management and consultancy services to the built and natural environment

15,000 people 300 offices 35 countries





bioremediation

mechanisms of degradation are unseen and not as well understood as horsepower based remediation



























agenda

start with overview of SIP





agenda start with overview of SIP see MI (microbe.com) can't avoid the underlying science but will try to minimize recorded webinars Mike Hyman Frank Loeffler Kirsti Ritalahti Matt Burns WSP

agenda

start with overview of SIP

can't avoid the underlying science but will try to minimize see MI (microbe.com) recorded webinars by:

Mike Hyman Frank Loeffler Kirsti Ritalahti Matt Burns

Pennsylvania case study



agenda start with overview of SIP see MI (microbe.com) can't avoid the underlying science but will try to minimize Mike Hyman Frank Loeffler Kirsti Ritalahti

recorded webinars

Matt Burns

Pennsylvania case study **Michigan case study**



agenda start with overview of SIP see MI (microbe.com) can't avoid the underlying science but will try to minimize by:

recorded webinars Mike Hyman Frank Loeffler

Kirsti Ritalahti Matt Burns

Pennsylvania case study **Michigan case study** another Michigan case study















































¹³C Stable Isotope Natural Abundance ~ 1%




















tear gas release site (CN)

Stable Isotope Probing (SIP) Study

port	MW-20	MW-32
P) Study		
PLFA		
total biomass:	2.29E+04 cells/bead	2.43E+06 cells/bead
¹³ C enriched biomass:	1.73E+01 cells/bead	2.68E+04 cells/bead
average PLFA del ¹³ C:	4 ‰	138‰
maximum PLFA del ¹³ C:	4‰	675‰

SITE LOGIC Repo	ort tudy	MW-20	MW-32
PLFA	total biomass: ¹³ C enriched biomass: average PLFA del ¹³ C:	2.29E+04 cells/bead 1.73E+01 cells/bead 4‰	2.43E+06 cells/bead 2.68E+04 cells/bead 138 ‰
DIC	DIC del ¹³ C: DIC del ¹³ C: percent ¹³ C DIC:	4‰ -20‰ 1.08%	675 ‰ 10,921 ‰ 11.76 %
			WSP

SITE LOGIC RE	port) Study	MW-20	MW-32
13C Phospholipid fatty acid	PLFA total biomass: ¹³ C enriched biomass: average PLFA del ¹³ C: maximum PLFA del ¹³ C:	2.29E+04 cells/bead 1.73E+01 cells/bead 4 ‰ 4 ‰	2.43E+06 cells/bead 2.68E+04 cells/bead 138 ‰ 675 ‰
double bond (or other	DIC DIC del ¹³ C: percent ¹³ C DIC:	-20 ‰ 1.08 %	10,921‰ 11.76%
structural variant)			

SITE LOGIC Re	eport) Study	MW-20	MW-32
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double bond (or other structural variant)	DIC DIC del ¹³ C: percent ¹³ C DIC:	-20‰ 1.08%	10,921‰ 11.76%
fatty acid structural groups provide finger print of the types of microbial groups	percent total PLFA) firmicutes (TerBrSats): proteobacteria (Monos): anaerobic metal reducers (brMonos): actinomycetes (MidBrSat): general (Nsat): eukaryotes (Polynoics):	0.0 22.9 0.0 0.0 77.1 0.0	6.2 59.2 1.1 0.8 30.3 2.5

Stable Isotope Probing (SIP) Study

definitively identified biodegradation as mechanism of benzene attenuation at the site (downgradient area)

Stable Isotope Probing (SIP) Study

		•	•
PLFA	total biomass:	9.91E+05 cells/bead	3.83E+06 cells/bead
	¹³ C enriched biomass:	4.57E+04 cells/bead	2.37E+05 cells/bead
	average PLFA del ¹³ C:	864‰	4202‰
	maximum PLFA del ¹³ C:	4469‰	6971 ‰
DIC	DIC del ¹³ C:	2227‰	3,235‰
	percent ¹³ C DIC:	3.44%	4.47%

MW-1

MW-2

Stable Isotope Probing (SIP) Study

		•	•
PLFA	total biomass:	9.91E+05 cells/bead	3.83E+06 cells/bead
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DIC	DIC del ¹³ C:	2227‰	3,235‰
	percent ¹³ C DIC:	3.44%	4.47%
Community Structure	firmicutes (TerBrSats):	4.0	5.5
(percent total PLFA)	proteobacteria (Monos):	76.4	73
	anaerobic metal reducers (brMonos):	0.5	4.1
	actinomycetes (MidBrSat):	0.0	0.05
	general (Nsat):	16.4	15.3
	eukaryotes (Polynoics):	2.7	1.6

MW-1

MW-2

PLFAtotal biomass:9.91E+05 cells/bead3.83E+06 cells/bead1³C enriched biomass:4.57E+04 cells/bead2.37E+05 cells/beadaverage PLFA del 1³C:864 ‰4202 ‰maximum PLFA del 1³C:4469 ‰6971 ‰DICDIC del 1³C:2227 ‰3,235 ‰percent 1³C DIC:3.44 %4.47 %Community Structure (percent total PLFA)firmicutes (TerBrSats):4.05.5proteobacteria (Monos):76.473anaerobic metal reducers (brMonos):0.54.1actinomycetes (MidBrSat):0.00.05general (Nsat):16.415.3eukaryotes (Polynoics):2.71.6	Stable Isc	t udy	MW-1	MW-2
Index entriched biomass:4.57E+04 cells/bead2.37E+05 cells/beadaverage PLFA del 13C:864 %4202 %maximum PLFA del 13C:4469 %6971 %DICDIC del 13C:2227 %3,235 %percent 13C DIC:3.44 %4.47 %Community Structurefirmicutes (TerBrSats):4.05.5(percent total PLFA)proteobacteria (Monos):76.473anaerobic metal reducers (brMonos):0.54.1actinomycetes (MidBrSat):0.00.05general (Nsat):16.415.3eukaryotes (Polynoics):2.71.6	PLFA	total biomass:	9.91E+05 cells/bead	3.83E+06 cells/bead
average PLFA del ¹³ C: 864‰ 4202‰ maximum PLFA del ¹³ C: 4469‰ 6971‰ DIC DIC del ¹³ C: 2227‰ 3,235‰ percent ¹³ C DIC: 3.44% 4.47% Community Structure (percent total PLFA) firmicutes (TerBrSats): 4.0 5.5 anaerobic metal reducers (brMonos): 76.4 73 anaerobic metal reducers (brMonos): 0.5 4.1 actinomycetes (MidBrSat): 0.0 0.05 general (Nsat): 16.4 15.3 eukaryotes (Polynoics): 2.7 1.6		¹ ^o C enfiched biomass:	4.57 E+04 cells/bead	2.37 E+05 cells/bead
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DICDIC del 13C: percent 13C DIC:2227 ‰ 3.44 %3,235 ‰ 4.47 %Community Structure 		maximum PLFA del ¹³ C:	4469‰	6971 ‰
Die dei * O.ZZZ / MB5,200 / MBpercent 13C DIC:3.44 %4.47 %Community Structure (percent total PLFA)firmicutes (TerBrSats):4.05.5proteobacteria (Monos):76.473anaerobic metal reducers (brMonos):0.54.1actinomycetes (MidBrSat):0.00.05general (Nsat):16.415.3eukaryotes (Polynoics):2.71.6	DIC	DIC del ¹³ C·	2227%	3 235 %
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		004 /00	
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(porcent total PLEA)	ninicules (Terbi Sals).	4.0	0.0 73
(percent total i El A)	anaerobic metal reducers (brMonos):	0.5	4 1
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Contaminant Loss	¹³ C Benzene pre-deployment:	185±12 µg/bd	185±12µg/bd
	¹³ C Benzene post-deployment:	130±11 µg/bd	144±11 µg/bd

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		-	-

Stable Isotope Probing (SIP) Study

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Stable Isotope Probing (SIP) Study

quantified expected full-scale performance of two remedial options











Stable Isotope Probing (SIP) Study

Contaminant Loss	¹³ C Benzene pre-deployment:	202±12 µg/bd
	¹³ C Benzene post-deployment:	156±11 µg/bd
	first order rate:	0.0021 day-1
DIC	DIC del ¹³ C:	-35 ‰
	percent ¹³ C DIC:	1.06%



MW-102



13C DNA	Separate 13C from 12 C using centrifuge
	Standard WW0224 WW 10224 CT3 10224
Separate DNA and sequence to identify microbes	10 10 10 10 10 10 10 10 10 10

		MW-102	
Study		Ļ	
Contaminant Loss	 ¹³C Benzene pre-deployment: ¹³C Benzene post-deployment: first order rate: 	202±12 μg/bd 156±11 μg/bd 0.0021 day ⁻¹	
DIC	DIC del ¹³ C: percent ¹³ C DIC:	-35 ‰ 1.06 %	
			ſ





6 6 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	(i) 2 bands of enriched DNA! Acinetobacter spp. Trichococcus spp.	Co Di
1.10		

		MW-102
Study		Ļ
Contaminant Loss	¹³ C Benzene pre-deployment: ¹³ C Benzene post-deployment: first order rate:	202±12 μg/bd 156±11 μg/bd 0.0021 day ⁻¹
DIC	DIC del ¹³ C: percent ¹³ C DIC:	-35 ‰ 1.06 %
		_
		WSP



Stable Isotope Probing (SIP) Study

identified unexpected pathway that can be enhanced







definitively demonstrate biodegradation



evaluate inhibition

quantifiably compare remedial options

identify unexpected mechanisms



Case Studies on Practical Use of Stable Isotope Probing

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