



Aerobic Biodegradation of Aromatic, Chlorinated Aliphatic, and Ether Contaminants by *Pseudonocardia sp.* strain ENV478 and Native Populations

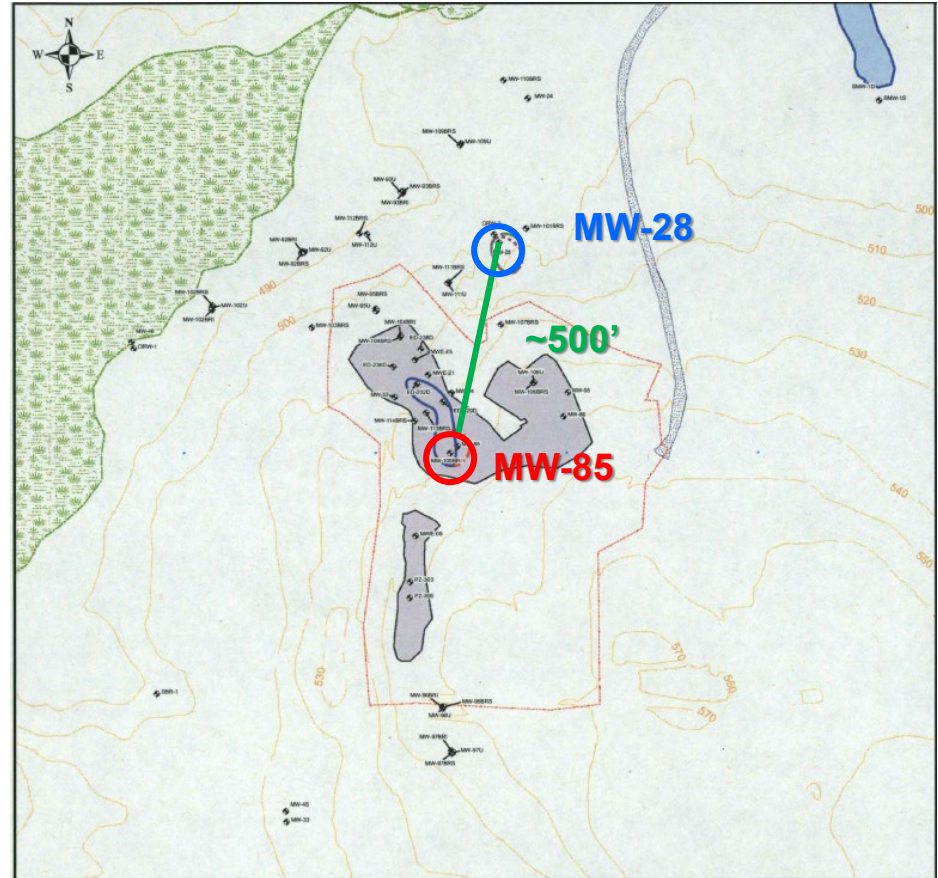
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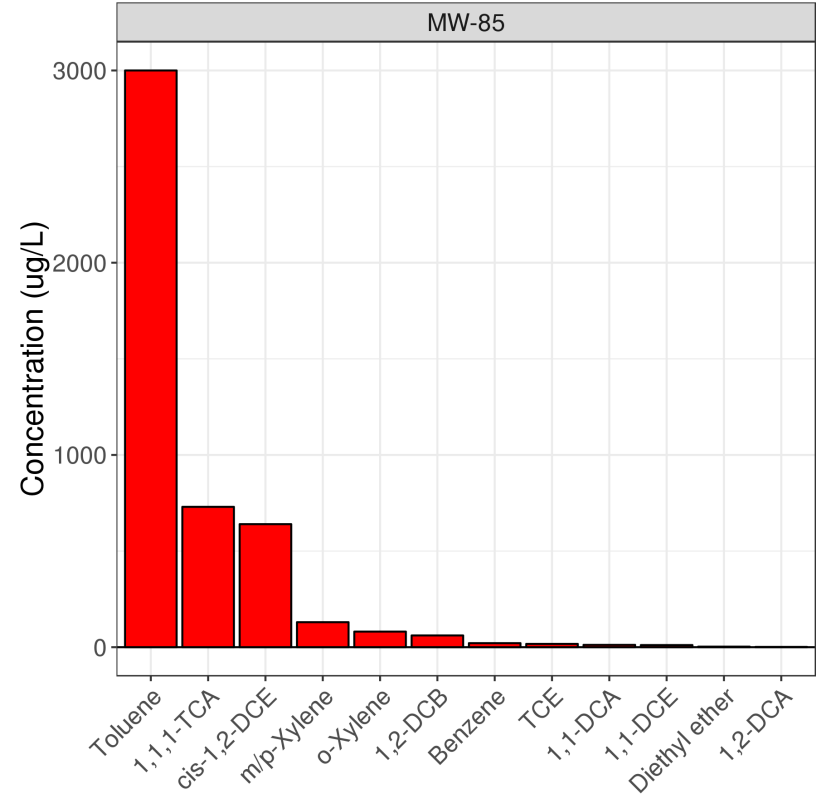
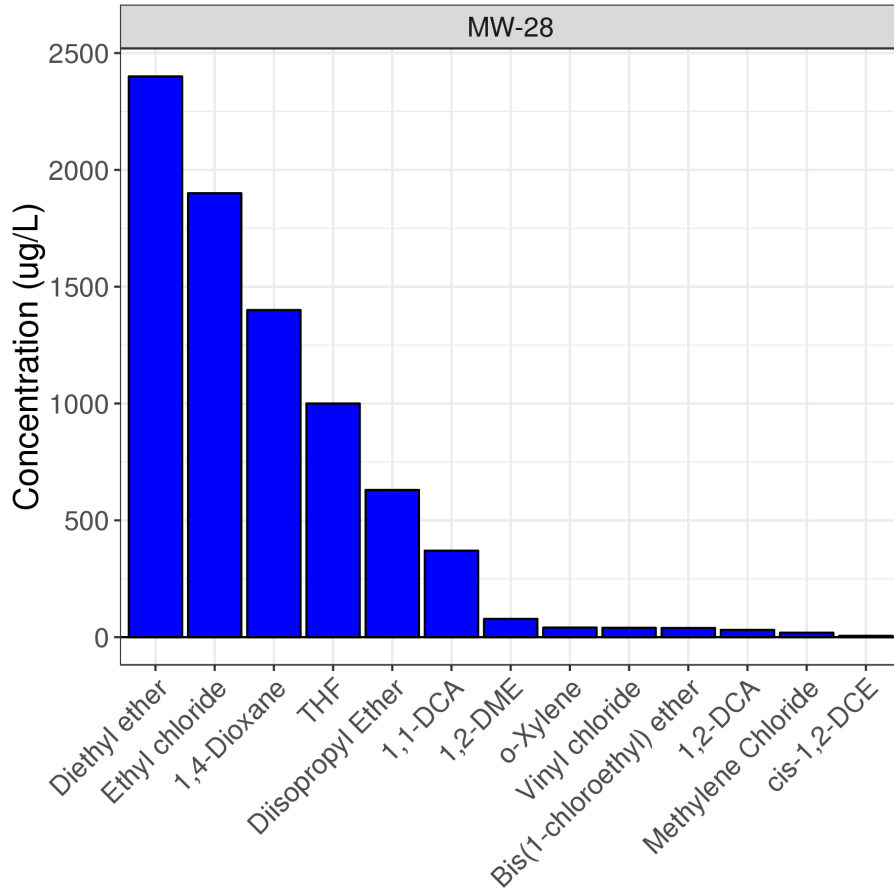
Paul Hatzinger and Simon Vainberg (APTIM)

Background

- Superfund site in NE US
- Decades of remedial efforts
 - ppt – low ppm **aromatics**, cVOCs, **ethers** remaining
- **Geology**
 - MW-28 –competent bedrock (32-42 ft bgs)
 - MW-85 - overburden (25-29 ft bgs), weathered bedrock (29-30 ft bgs)
 - Wells are not hydraulically connected
- **Geochemistry**
 - pH: 5.8 – 6
 - DO: 0.24 – 0.5 mg/L
 - ORP: -36.6m to -78 mV



COCs of interest



COCs differ between MW-28 and MW-85

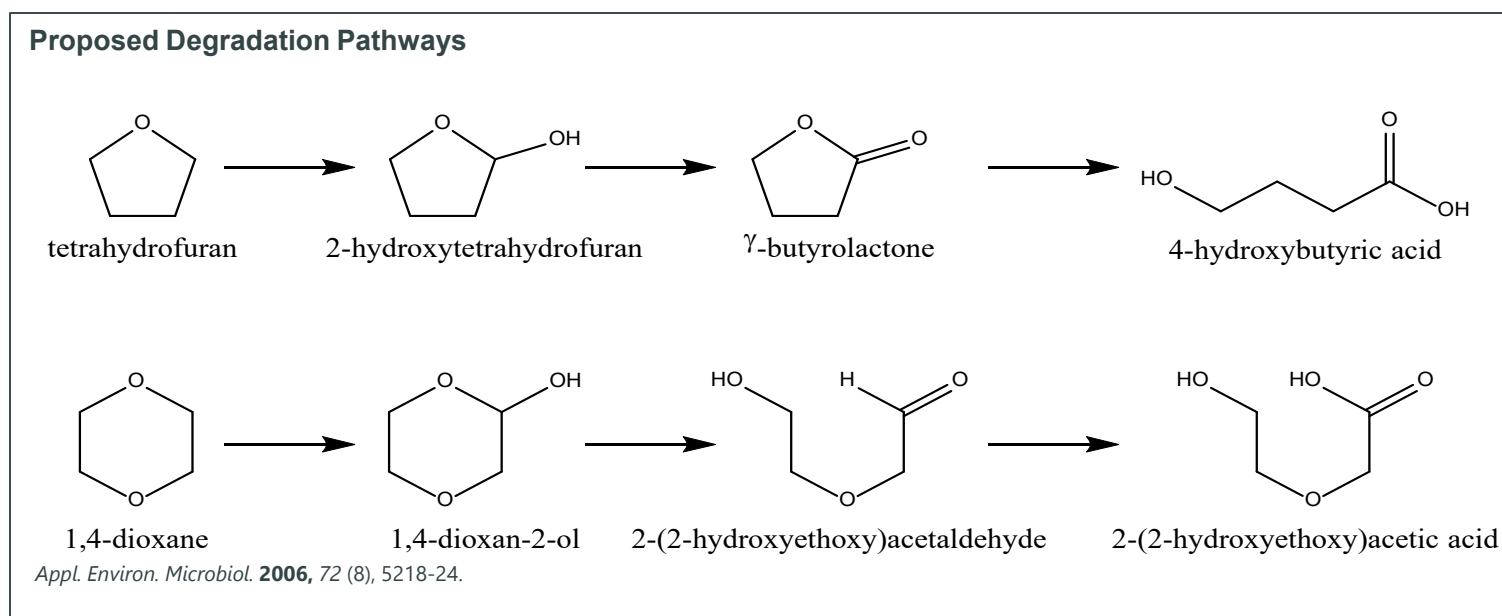
Objectives

Evaluate potential bioremediation options

- Aerobic biostimulation & bioaugmentation
- **THF** present as growth substrate for 1,4-DX degradation

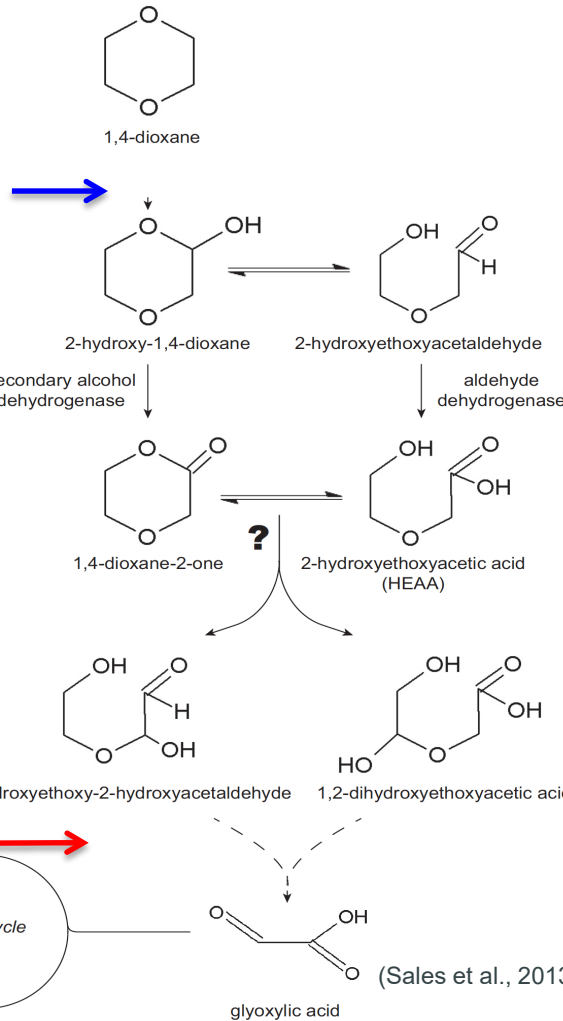
Evaluate biodegradation of site COCs using *Pseudonocardia* sp. strain **ENV478**

- **Co-metabolic** ether degrader (1,4-DX, BCEE, MTBE)
- Degradation of **aromatics and cVOCs not previously tested**



Pathways

MO catalyzed reaction



Potential ALDH catalyzed reaction

Potential ALDH catalyzed reaction

(Sales et al., 2013; Grostern et al., 2012; Mahendra et al., 2007)

CENSUS (qPCR) Data

MW-28

Functional Gene	Abbreviation	cells/mL	Percentile (%)
Total Eubacteria	EBAC	1.83 x 10⁶	70
<i>Dioxane/THF Monooxygenase</i>	<i>DXMO</i>	<i>6.00 x 10⁻¹ (J)</i>	---
Aldehyde Dehydrogenase	ALDH	<5.00	---
Propane Monooxygenase	PPO	7.17 x 10 ¹	---
Soluble Methane Monooxygenase	SMMO	9.86 x 10²	38
Particulate Methane Monooxygenase	PMMO	<5.00	---
Phenol Hydroxylase	PHE	3.71 x 10³	72
Toluene-2-Monooxygenase	RDEG	4.93 x 10³	82
Toluene-3,4-Monooxygenase	RMO	2.27 x 10³	75

- High levels of bacteria and functional gene expression measured
- Estimable amounts of DXMO suggest some dioxane/THF degradation may be ongoing

CENSUS (qPCR) Data

MW-85

Analyte	Abbreviation	cells/mL	Percentile (%)
Total Eubacteria	EBAC	1.32 x 10⁶	65
Dioxane/THF Monooxygenase	DXMO	<4.70	---
Aldehyde Dehydrogenase	ALDH	<4.70	---
Propane Monooxygenase	PPO	9.40	---
Soluble Methane Monooxygenase	SMMO	9.86 x 10²	22
Particulate Methane Monooxygenase	PMMO	<5.00	---
Phenol Hydroxylase	PHE	3.71 x 10³	65
Toluene-2-Monooxygenase	RDEG	4.93 x 10³	62
Toluene-3,4-Monooxygenase	RMO	2.27 x 10³	66

- High levels of bacteria and functional gene expression were measured

Treatments

Treatment	Amendment	Amendment Amount	pH
Biostimulation			
Abiotic Control	HgCl ₂	250 mg/L	Native
Live Control	---	---	Native, 7
DAP	Diammonium phosphate*	9.6 mg/L	Native, 7
DAP + propane	Diammonium phosphate* Propane	9.6 mg/L 3 mL @ 1 atm	Native, 7
Bioaugmentation			
Groundwater	ENV478	5 mL	Native
Basal Salt Medium (BSM)	ENV478 + COCs	5 mL	7

Microcosms prepared in triplicate

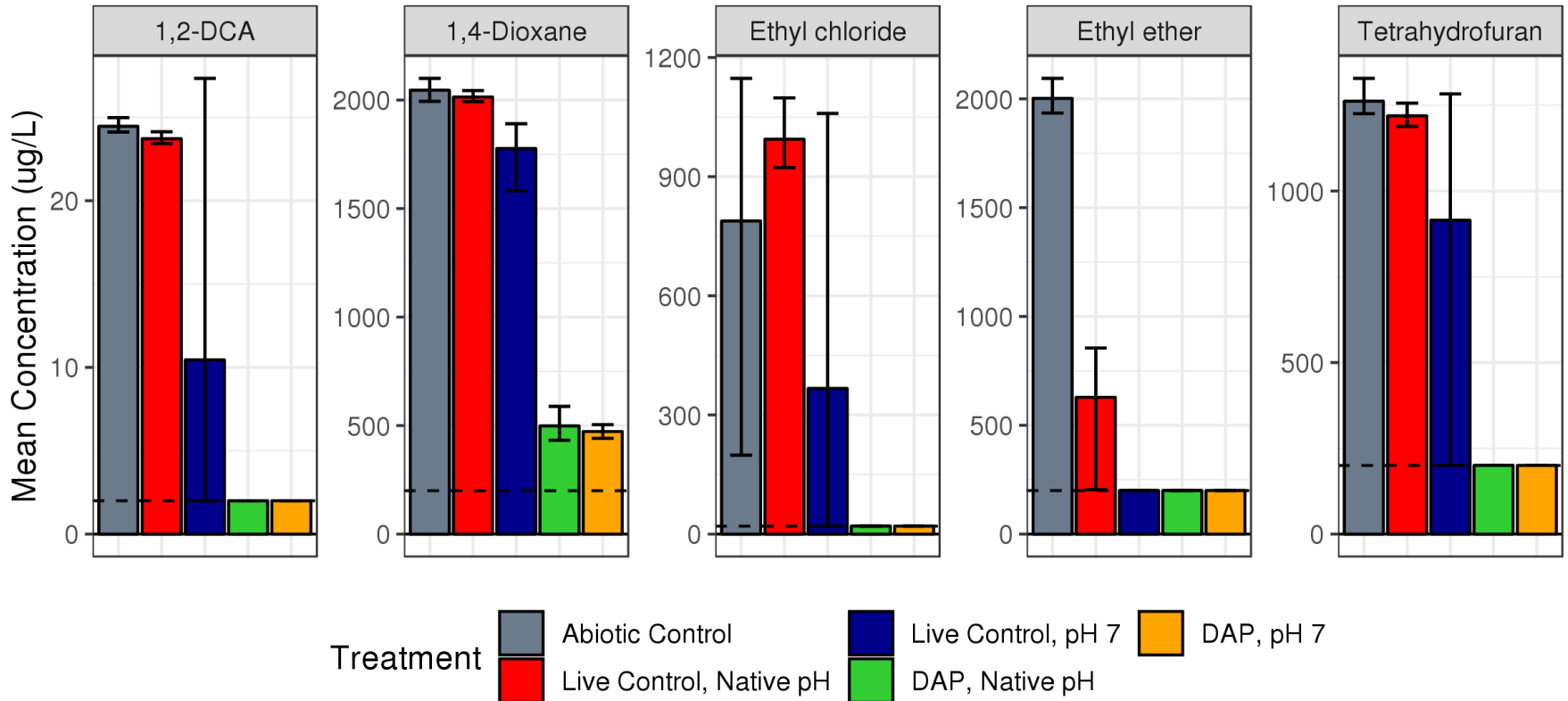
All treatments except BSM contain 125 mL groundwater from MW-28 or MW-85

*Diammonium phosphate (DAP) based on C estimate from COCs

BSM treatments to differentiate degradation by ENV478 from native organisms

Biostimulation: MW-28

Native pH = 6.0

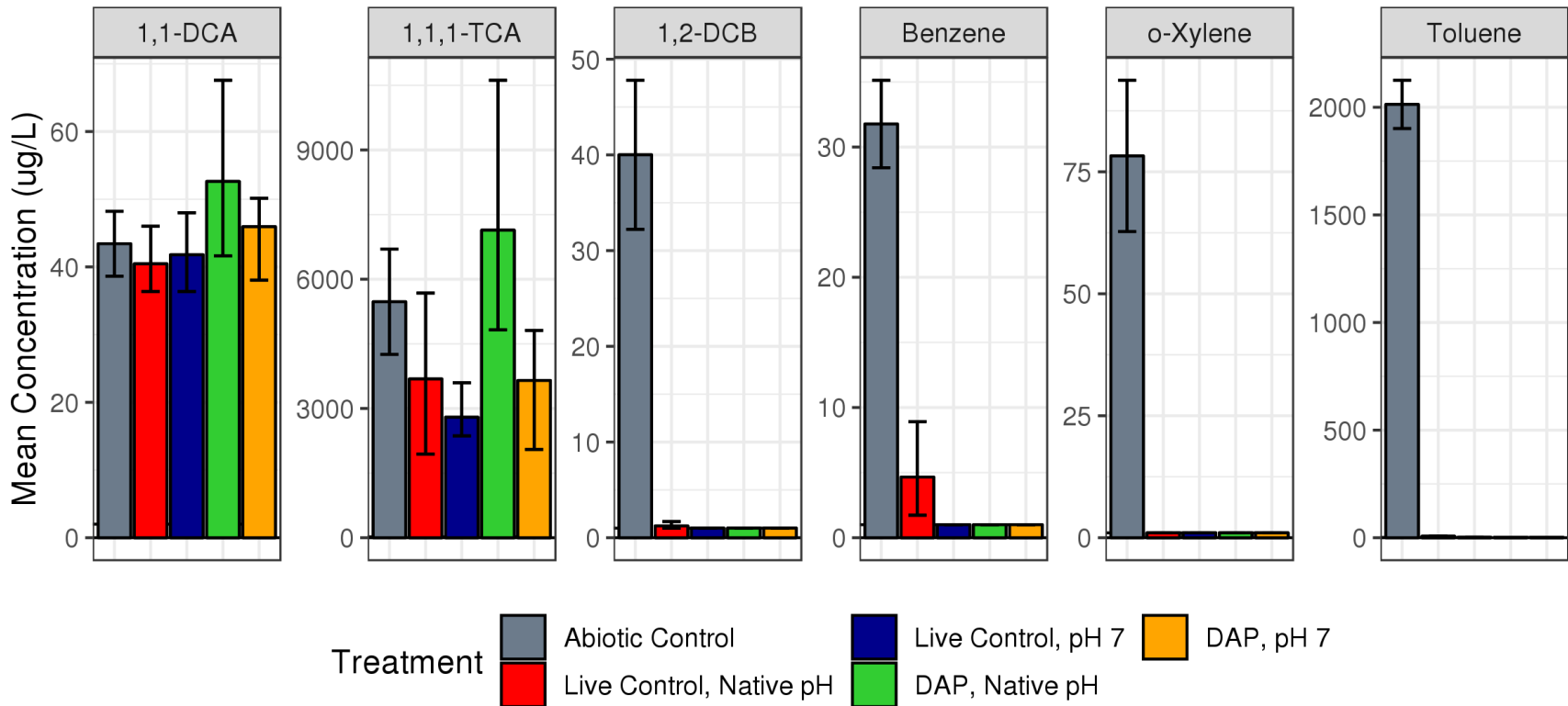


Bars represent the range of replicate data. Data collected at 180 d.

- Oxygenation of groundwater is not sufficient for degradation
- Addition of N&P stimulates biodegradation
- Increasing pH helped accelerate removal
- 1,4-DX degradation stalled

Biostimulation: MW-85

Native pH = 5.8

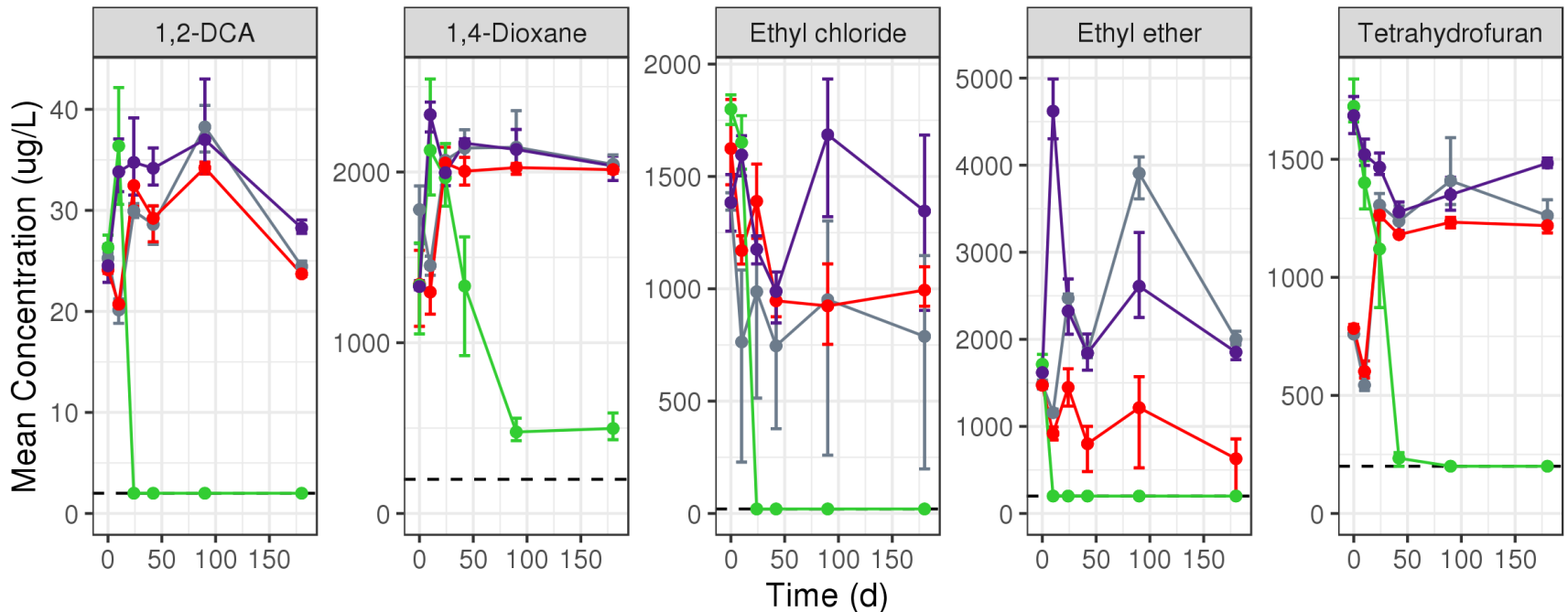


Bars represent the range of replicate data. Data collected at 180 d.

- Oxygenation of groundwater was sufficient for degradation of aromatics
- Amendment with nutrients and pH adjustment accelerated degradation
- **No significant degradation of 1,1-DCA and 1,1,1-TCA**

Biostimulation: Effect of Propane

Native pH = 6.0
MW-28



Treatment — Abiotic Control — Live Control, Native pH — DAP, Native pH — Propane + DAP, Native pH

Bars represent the range of replicate data.

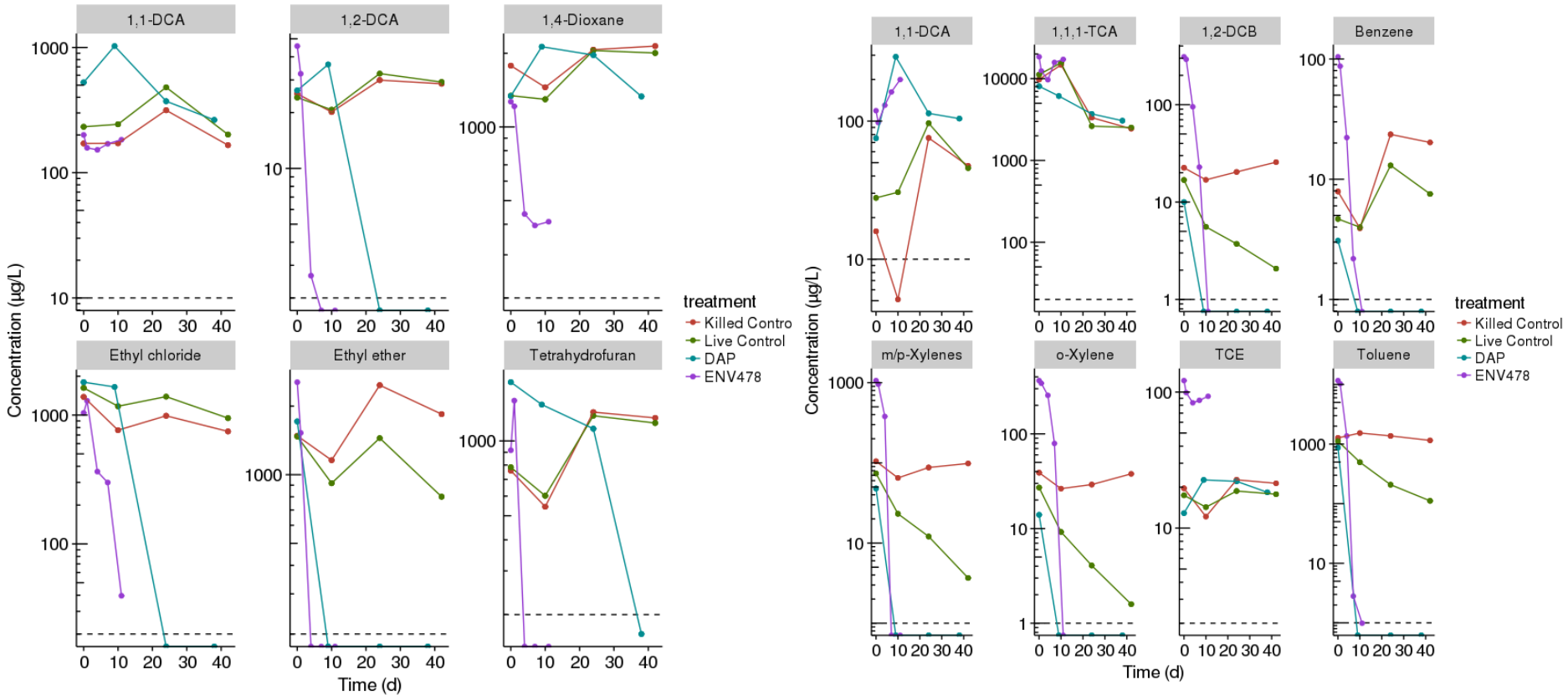
**Propane amendment inhibited degradation of COCs
→ preferential C source?**

Bioaugmentation Experiments

MW-28

Native pH

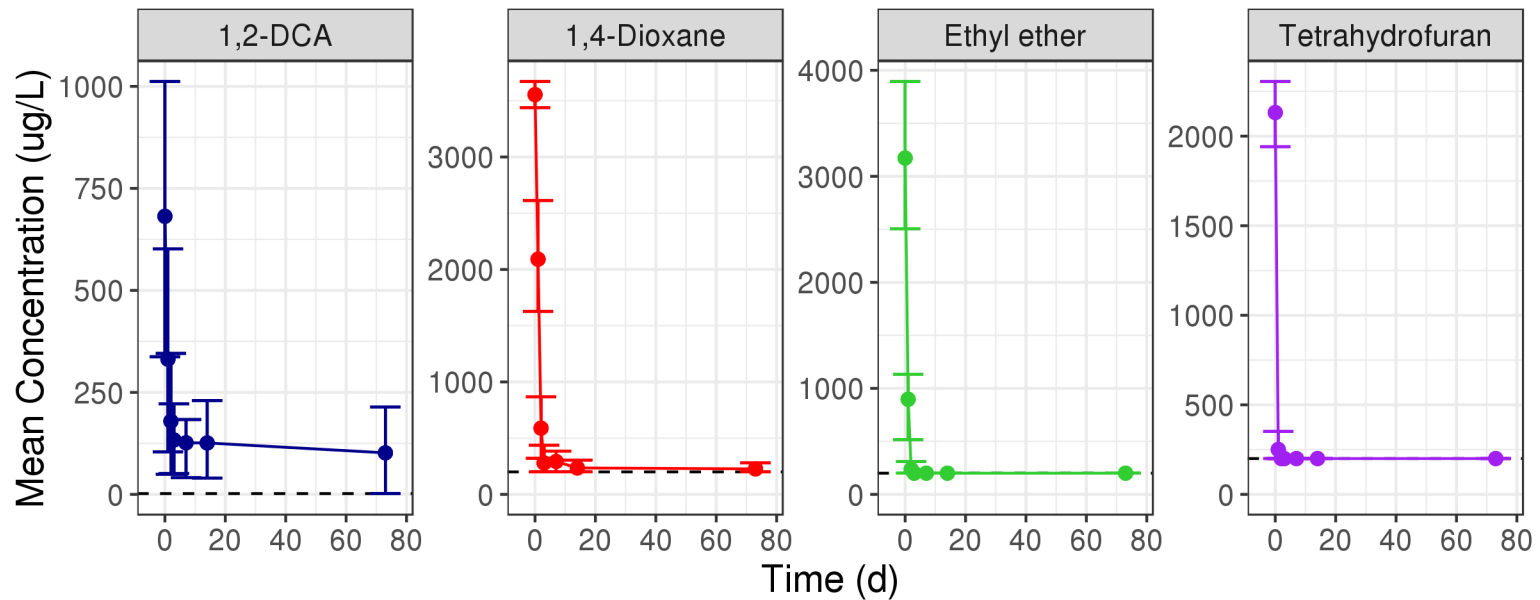
MW-85



- Addition of **ENV478** resulted in faster degradation of ethers, ethyl chloride & 1,2-DCA than DAP; aromatics showed similar removal to DAP
- **No significant degradation of 1,1-DCA & 1,1,1-TCA observed**

Bioaugmentation with ENV478

MW-28 Simulated Sample (BSM)



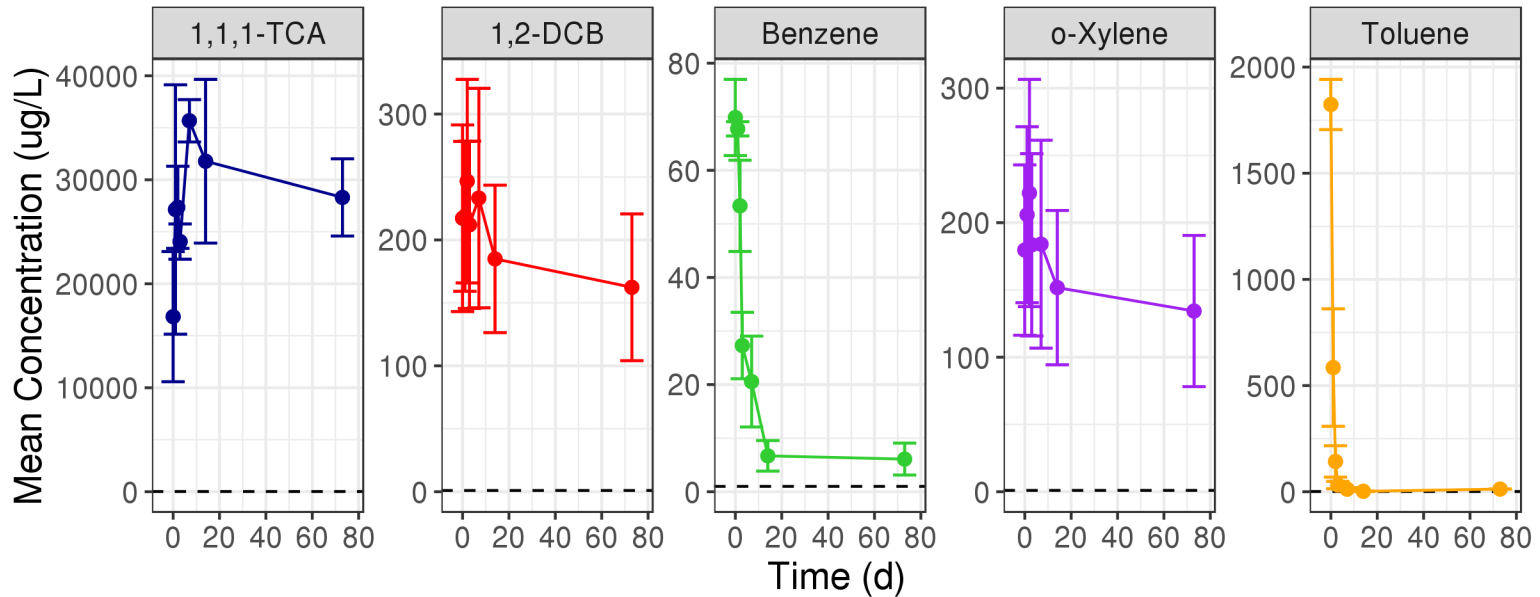
Treatment ● 1,2-DCA ● 1,4-Dioxane ● Ethyl ether ● Tetrahydrofuran

Bars represent the range of replicate data.

- **Rapid degradation of all ethers observed**
- **Initial rapid degradation of 1,2-DCA stalled after consumption of ethers → co-metabolic?**

Bioaugmentation with ENV478

MW-85 Simulated Sample (BSM)



Treatment —●— 1,1,1-TCA —●— 1,2-DCB —●— Benzene —●— o-Xylene —●— Toluene

Bars represent the range of replicate data.

- **ENV478 rapidly degraded benzene and toluene**
- **No significant degradation for other COCs**

Summary

- Lack of nutrients & low pH are limiting aerobic biodegradation
- >95% removal of THF, diethyl ether, ethyl chloride and 1,2-DCA after biostimulation in less than 50 days while 1,4-DX showed about 75% removal
- Similarly non-detectable levels of benzene, toluene, xylene, and 1,2-DCB were observed under biostimulating conditions
- Treatments with propane to induce native co-metabolic degradation were unsuccessful
- ENV478 rapidly degraded all ethers within 2 weeks
- ENV478 also degraded benzene, toluene to ND & 1,2 DCA by ~ 85%
- 1,1-DCA & 1,1,1-TCA were persistent under all conditions tested



— Thank You

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