

ONE DEGREE RULE



- A small change can help correct our course and get us to our goal.
- Understanding the black box of microbiology allows us to correct quickly and cost effectively.
 - What are the dominant organisms present?
 - What is the genetic potential for complete reductive dechlorination?
 - What organisms are competing for available donors?
 - Are compounds biodegrading?
 - What is likely to happen in the future?
 - What is the rate of degradation?





In situ higramodiation of chlorinated solvents

Microbiology (1998), 144, 599-608

Printed in Great Britain

REVIEW ARTICLE

Bioremediation: towards a credible technology

Ian M. Head

of chlorinated solvents and chlorinated aliphatic hydrocarbon contaminants in groundwater may soon become a reality. Both anaerobic reductive dehalogenation on non-chlorinated products and complete aerobic oxidation offer greatly increased promise for potential applications.

Current Opinion in Biotechnology 1993, 4:323-330

contamination, and (o) an easily degraded, extracted, or

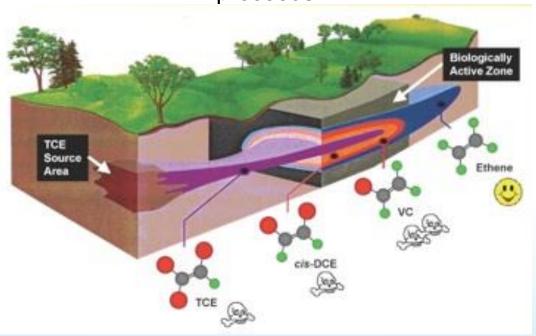
гентешация в опретити экс





REMEDIATION EFFORTS MUST LEAD TO COMPLETE DECHLORINATION

The toxicity of TCE transformation products increase as dechlorination proceeds.



This is problematic if the process stalls once cis-DCE and/or VC is produced.



DATA GAPS

- qPCR and QuantArray provide quantitative data for microorganisms and gene targets
 - What about activity?
- Next Generation Sequencing provides a bigger picture of the overall microbiome
 - What about the health of the microbiome?
- Compound Specific Isotope Analysis and Stable Isotope Probing provide proof of contaminant degradation
 - What about predicting future degradation trends?





ARTICLE

https://doi.org/10.1038/s41467-019-11311-9

OPEN

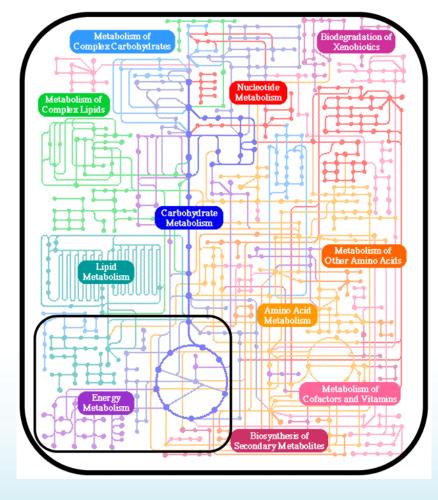
A metabolic profile of all-cause mortality risk identified in an observational study of 44,168 individuals

Joris Deelen et al.#





- Systematic study of the unique chemical fingerprints as the result specific cellular processes.
- Metabolome: the collection of all metabolites in a biological cell, tissue, organ or organism.
- Metabolic profiling: instantaneous snapshot of the cell physiology.



http://www.urmc.rochester.edu/labs/Munger-Lab/



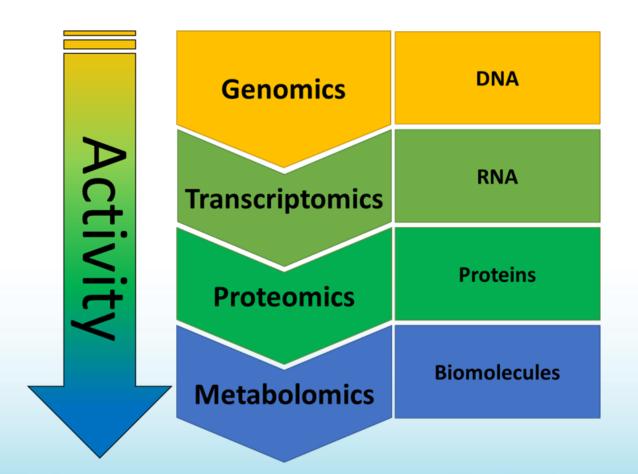
METABOLOMICS

- Analysis of all small molecules (MW<1100) within an environmental sample
- Identification of 80-100 known compounds
- Comparison of the overall metabolic profile
- Statistical Analysis and pattern recognition
 - Predictive capabilities
 - Activity of key degraders





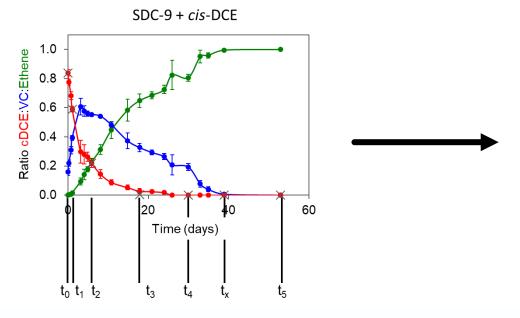
- Metabolomics analysis of thousands of small molecules from a biological sample
- Trends and patterns can be used to predict the health and activity of the entire microbiome
- Allows for a broad screening of reductive dechlorination potential



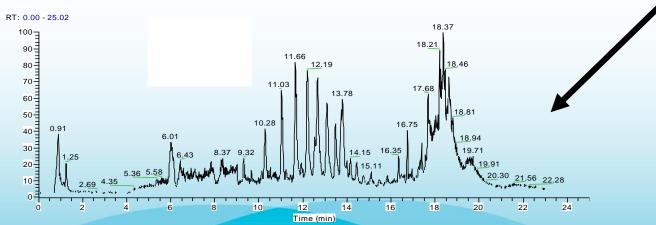




SBIR METABOLOMICS STUDY



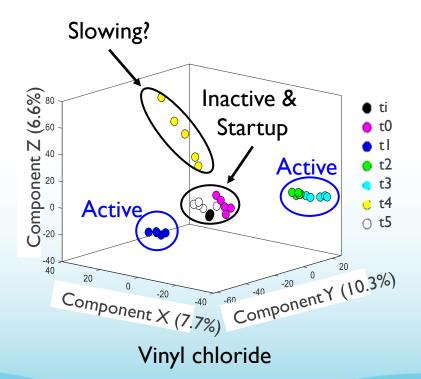


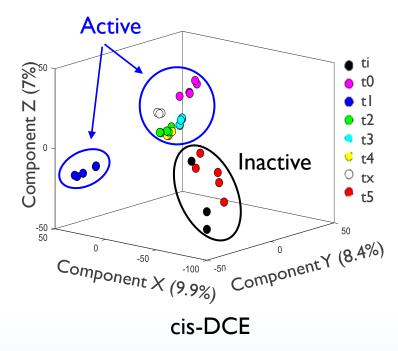






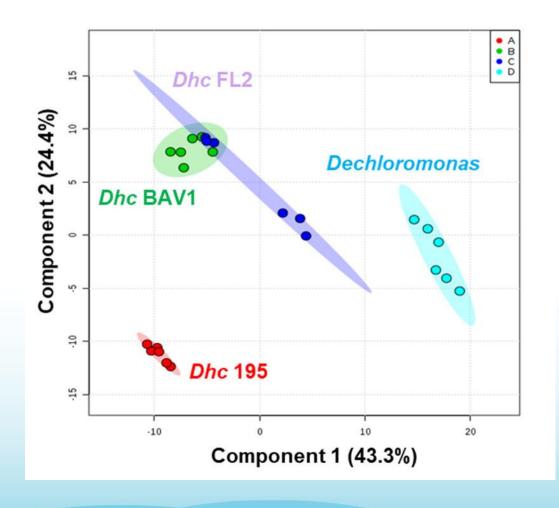
SBIR METABOLOMICS STUDY





- CommercialBioaugmentationconsortium (SDC-9)
- Sampling performed in intervals to capture stage of dechlorination
- Over 10,000 metabolites identified
- Both the known and unknown metabolites were analyzed using PLS-DA plots
- Differences in stages of dechlorination









SBIR METABOLOMICS STUDY

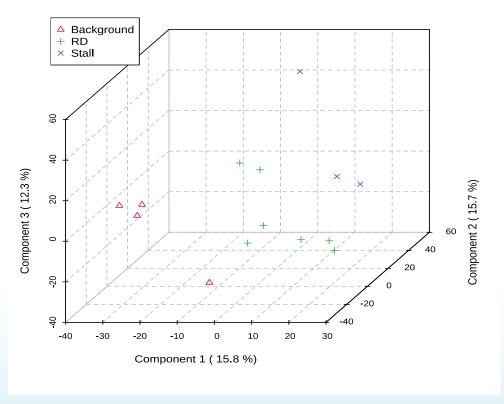


Figure 10: 3D scores plot between the selected PCs. The explained variances are shown in brackets.

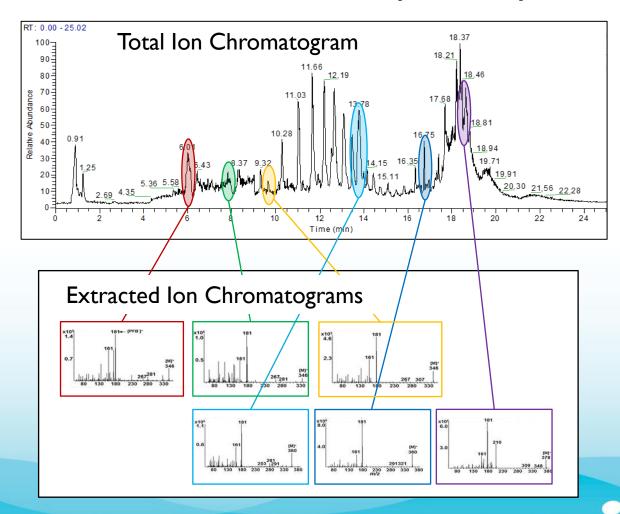
- Three chlorinated sites
 - Belleville Industrial (NJ)
 - Former Vickers Facility (MO)
 - Earnhart Site (TN)
- Different stages of Dechlorination
 - Active RD
 - Stall
 - Background
- Over 12,000 metabolites identified
- PLS-DA plot clearly distinguishes each stage
- VIP scoring identified 5 metabolites that could be sentinel for robust RD!





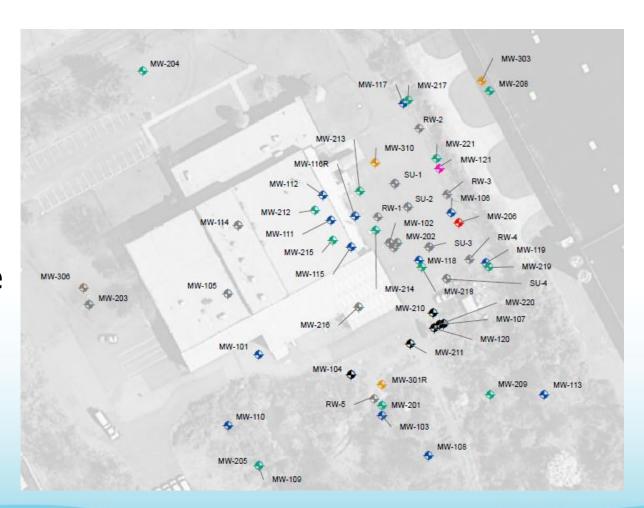
- Each sample's pattern is compared against a database
 - MI's in-house database
 - Made up of well-characterized samples with known reductive dechlorination classifications
 - Incorporation of machine learning to predict unknown samples based on the database

Thousands of features per sample



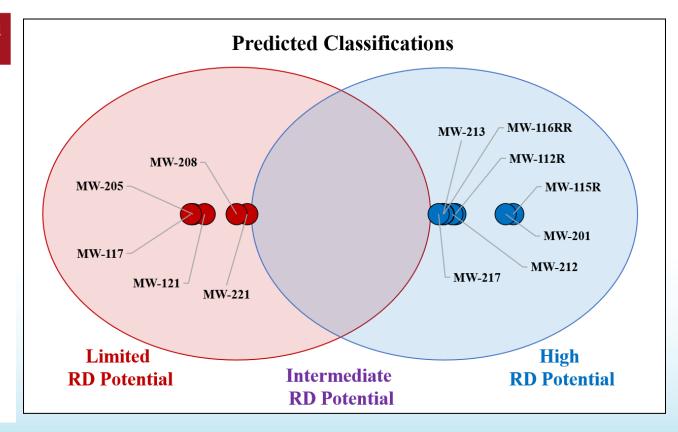


- Chlorinated solvent site
- QuantArray-Chlor and MetaArray
- Site chemistry, geochem, and history were not shared before analysis





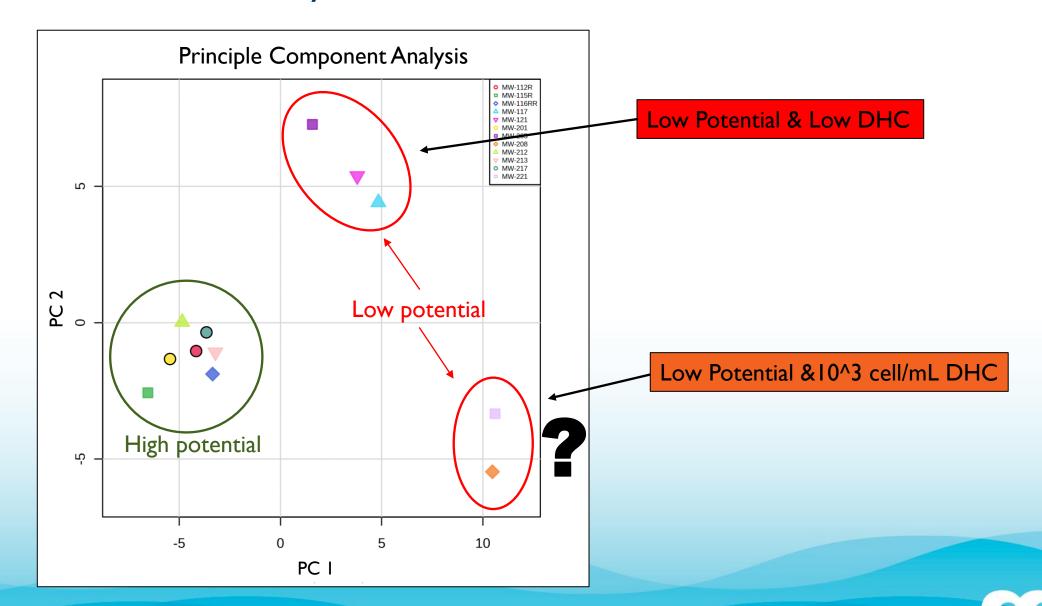
Sample Name	Predicted Class	SVM Model Accuracy ¹
MW-115R	High Reductive Dechlorination Potential	80%
MW-212	High Reductive Dechlorination Potential	80%
MW-121	Limited Reductive Dechlorination Potential	80%
MW-221	Limited Reductive Dechlorination Potential	80%
MW-116RR	High Reductive Dechlorination Potential	80%
MW-213	High Reductive Dechlorination Potential	80%
MW-117	Limited Reductive Dechlorination Potential	80%







Metabolomics Case Study

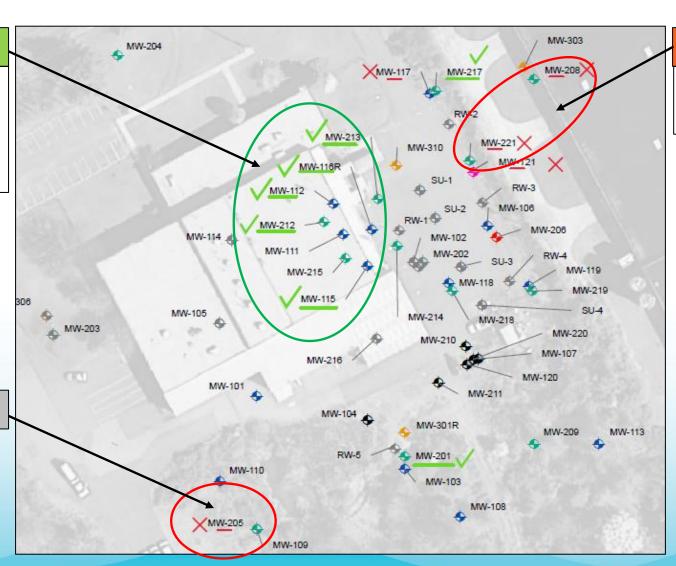




High Potential

Heavy treatment Area
Excavation
EVO Soil Mixing
ZVI/Hydraulic Fracking

Background



Low Potential &10³ cell/mL DHC

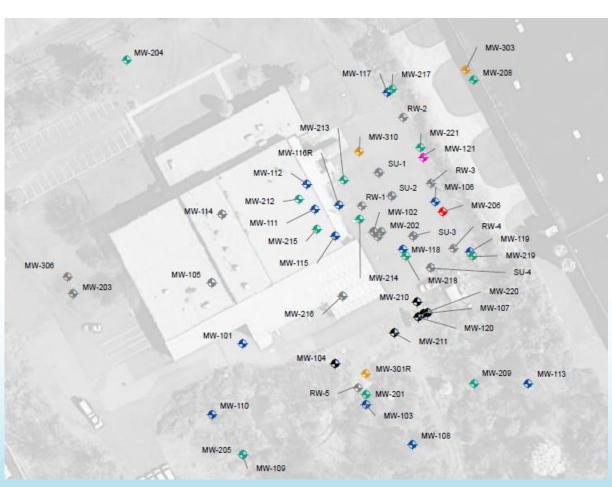
Bedrock wells

Share the same fracture line





- Next steps
 - Second sampling event
 - Compare results
 - Has the DHC and functional gene abundance changed
 - Does MetaArray indicate in the lower potential wells
 - Use the MBT results to identify the locations with issues
 - Reassess strategy in these locations





CONCLUSIONS

- Machine learning will continue to grow our databases and understanding of the biological metabolome.
- Metabolomics has the ability to help us get closer to rates of degradation.
- Key sentinel metabolites likely exist to help us define degradation potential and predict processes or community health.



