

Monitoring Bioremediation with Molecular Tools

Hauke Smidt

Laboratory of Microbiology Wageningen University





Fate of Environmental Pollutants

- Biodegradability largely depends on
 - chemical structure
 - environmental conditions

&

Purifying Capacity of microbiota

Fate of Environmental Pollutants

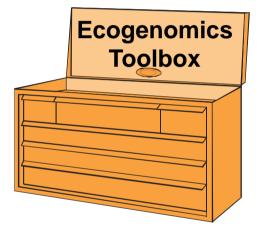


Opening up the black boxes

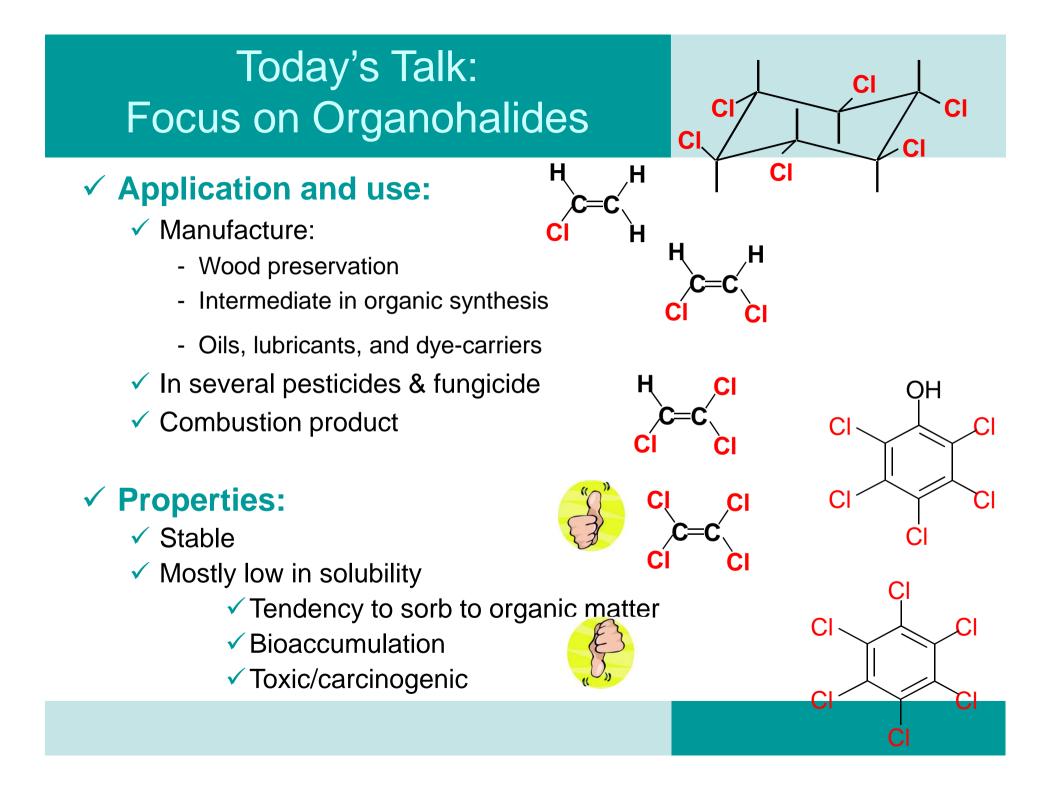
How to open the black box

Molecular Toolbox to assess and predict

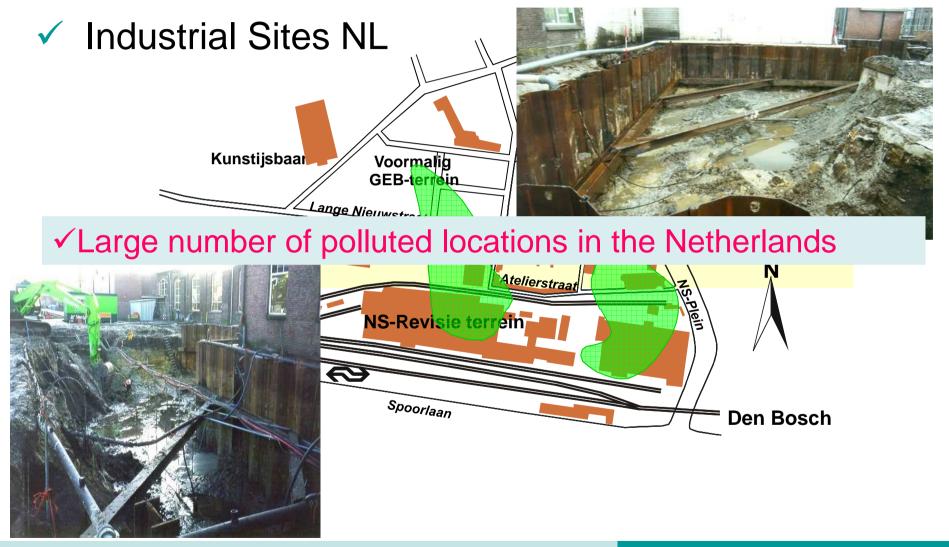
- Phylogenetic identity / complexity (WHO)
- ✓ Catabolic capacity (POTENTIAL)
- ✓ Catabolic activity (ACTION)



 Exploring and defining the self-purifying capacity of natural and man-made environments



Bioremediation of Aquifers and Sediments



Chlorinated Aliphatic Hydrocarbon (CAH) pollution

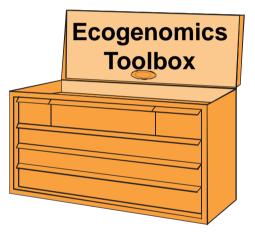
Biological cleanup of chlorinated compounds

- Biodegradability of halogenated hydrocarbons largely depends on
 - chemical structure
 - environmental conditions
 - Purifying capacity of microbiota
- Reductive dehalogenation is often crucial step for initial biodegradation in anoxic environments
 - dechlorination by specific enzymes
 + energy conservation : organohalide respiration

How to open the black box

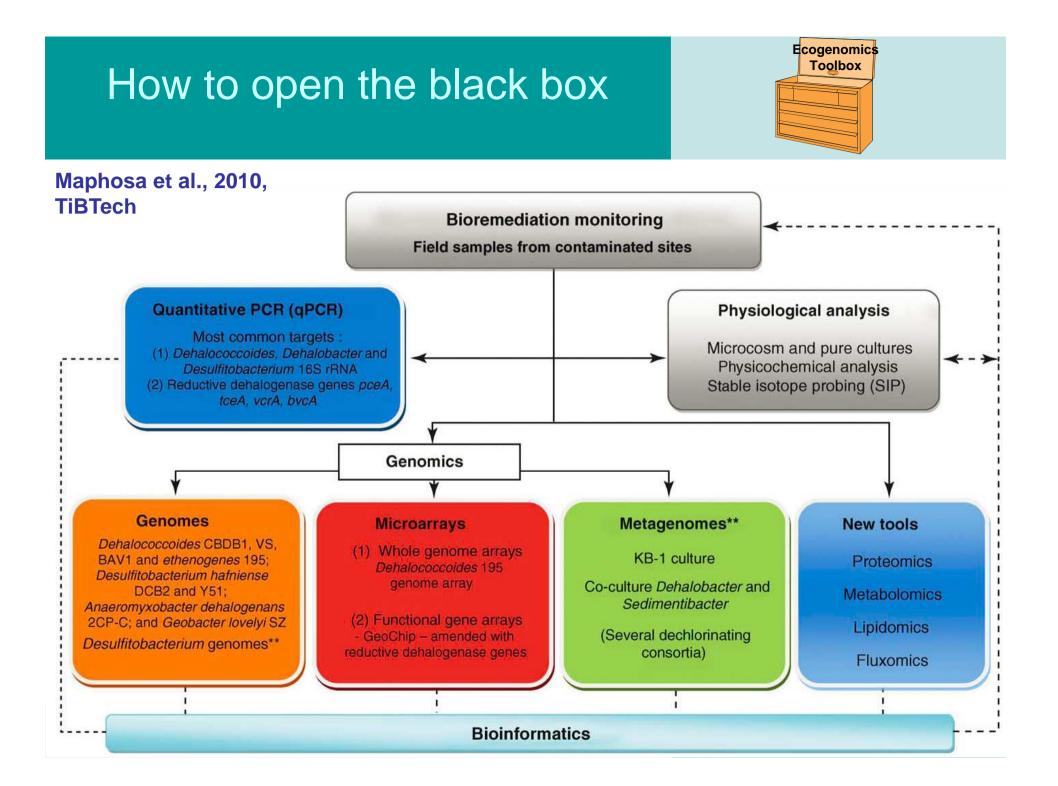
Molecular Toolbox to assess and predict

- Phylogenetic identity / complexity
- ✓ Catabolic capacity
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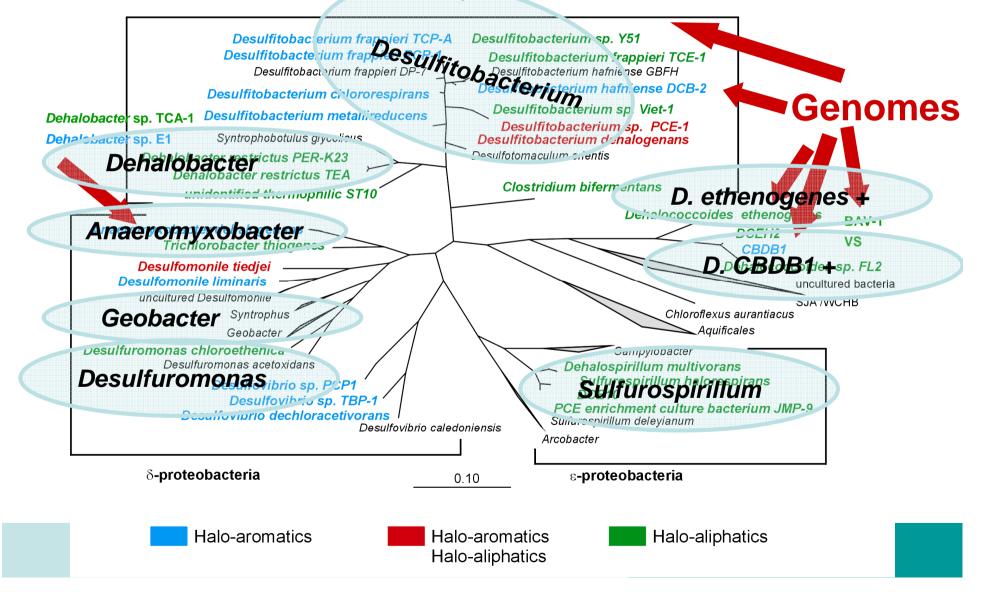
 Exploring and defining the self-purifying capacity of natural and man-made environments



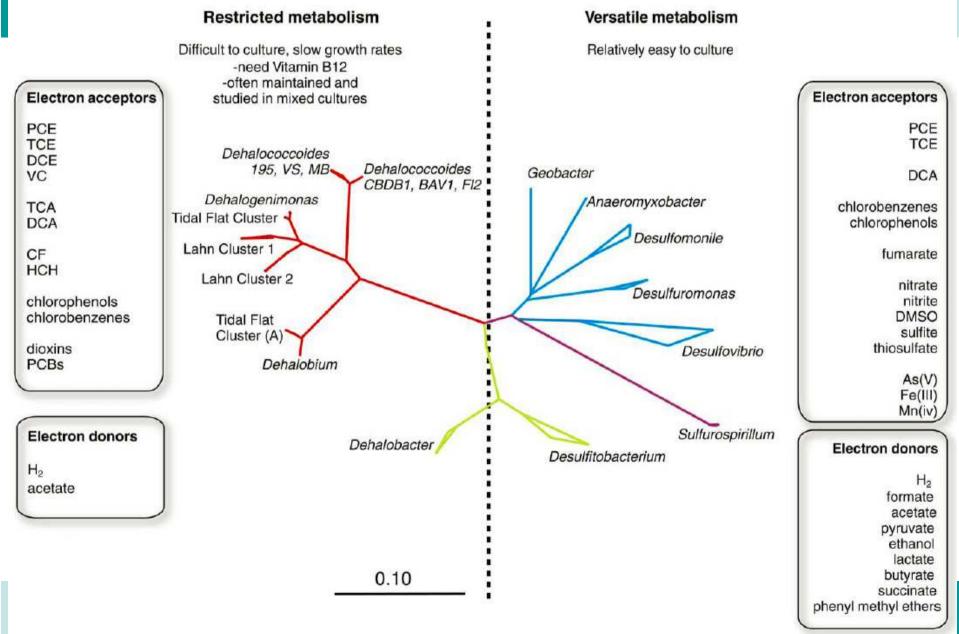


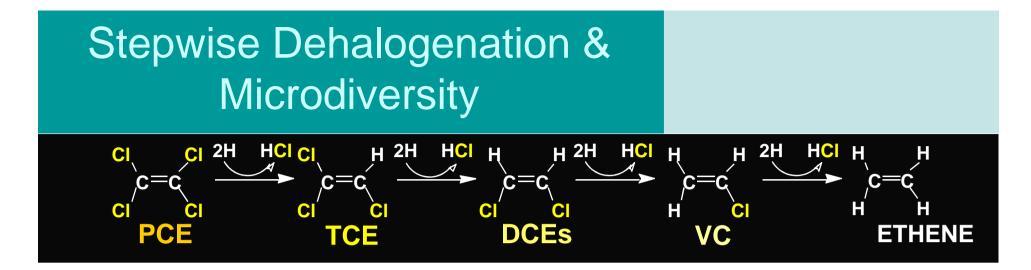
Organohalide Respiring Bacteria Diversity of isolates (16S rRNA)

Low-GC Gram-positives

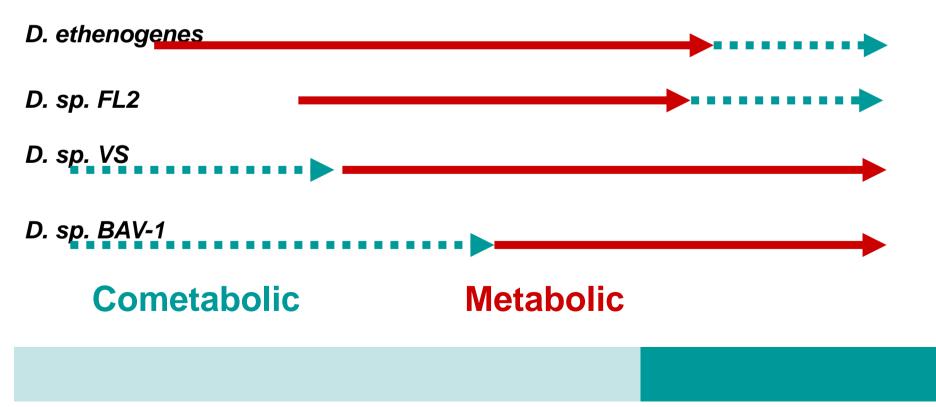


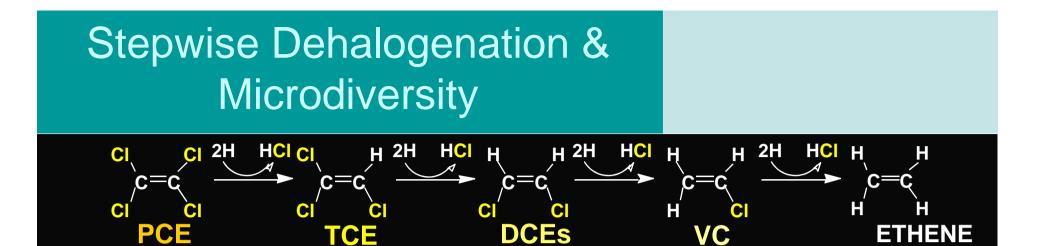
Organohalide Respiring Bacteria What do they live on

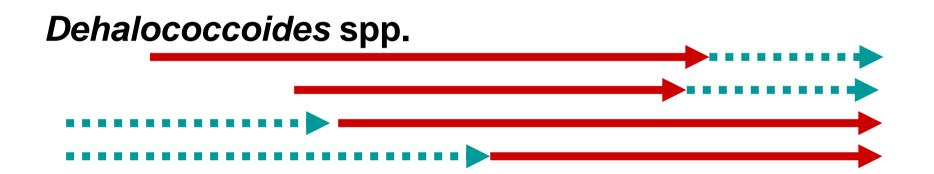




What Dehalococcoides spp. can do:







Clostridium sp. Dehalobacter spp. Desulfitobacterium spp. Desulfuromonas spp. Sulfurospirillum spp.

Lessons from Isolates

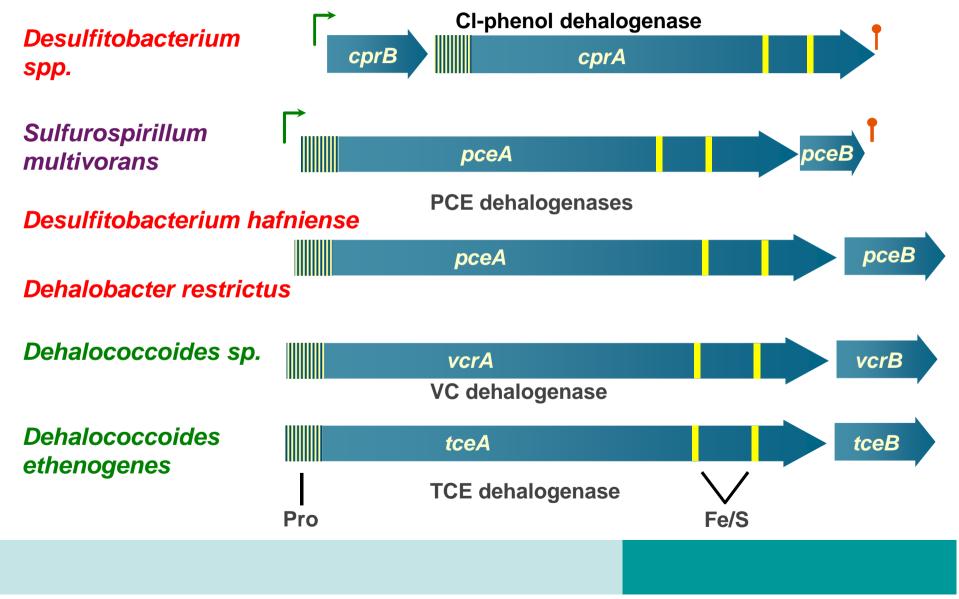
Reductive dehalogenation widespread across
 bacterial domain

 Functional microdiversity – Different activities in closely related strains

Non-dehalogenating strains within organohalide respiring genera

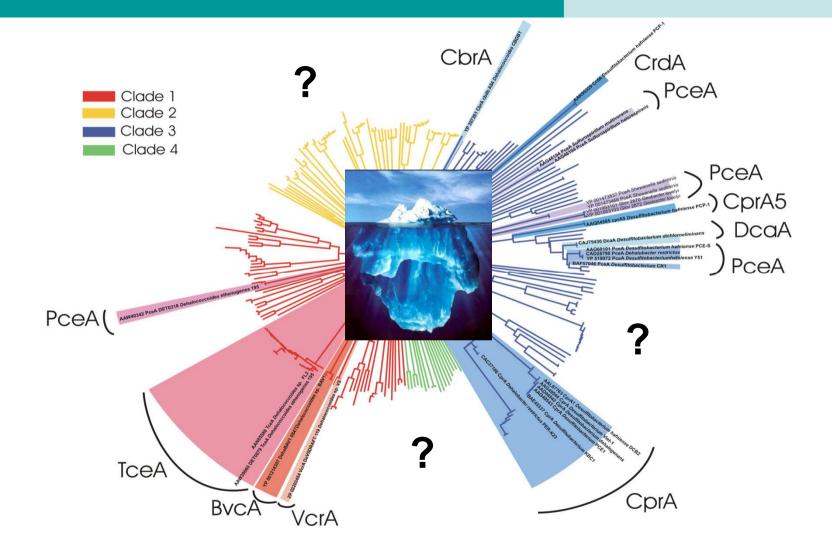
Catabolic Gene Markers Reductive dehalogenase genes

Conserved motifs



Organohalide Respiring Bacteria Some Current Diversity of Dehalogenases Many

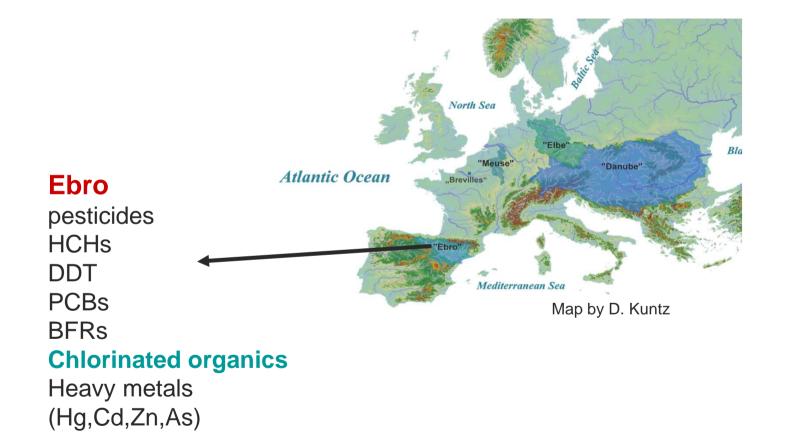
Some known Many unknown



Maphosa et al., TiBTech, 2010

Assessing Activity of OHRB Study Site: The Ebro Basin

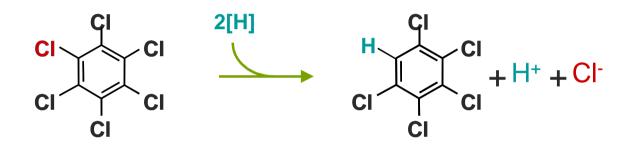




Tas et al., AEM, 2009

Focus on hexachlorobenzene

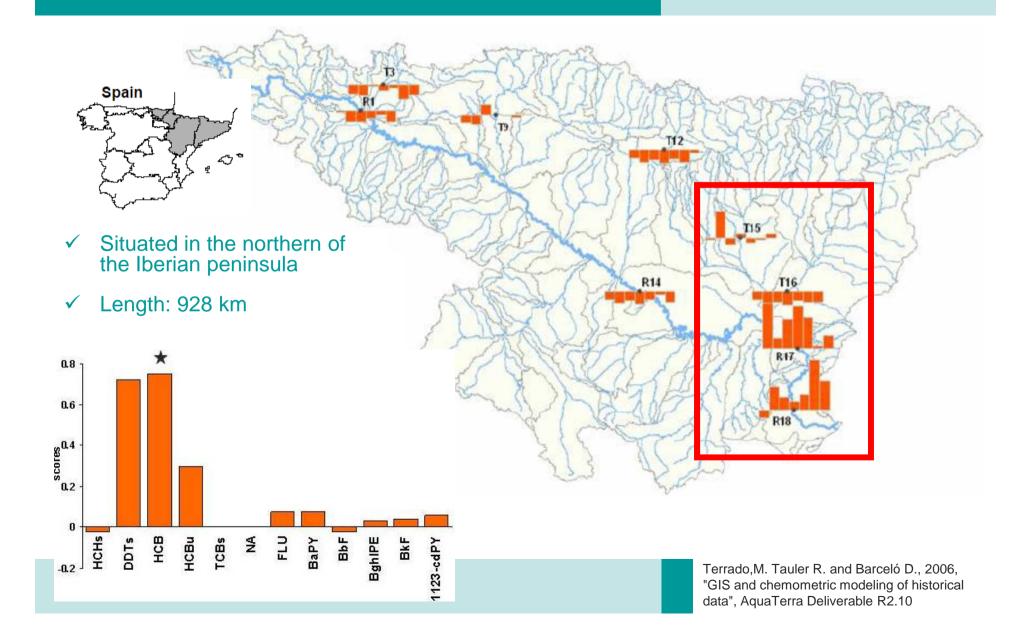
Hexachlorobenzene (HCB)



- ✓ Also here: stepwise dehalogenation
- ✓ Functional microdiversity in closely related strains of *Dehalococcoides*

AQUATERRA sites

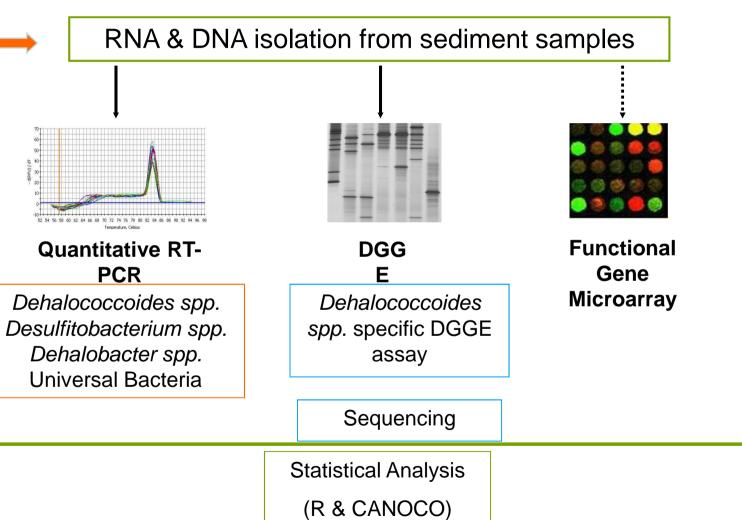
EBRO



Genomics Toolbox @ Work

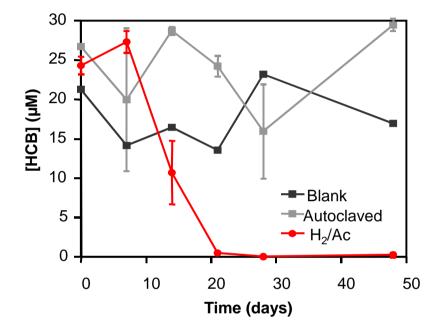




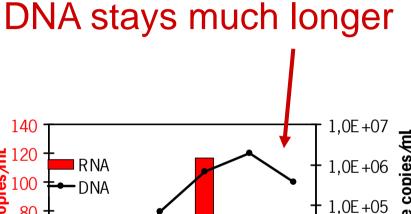


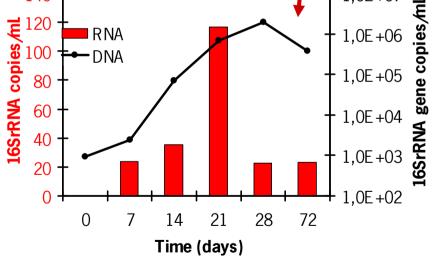
Combination of physiological and molecular ecology data

Flix sediment



 Profile of rRNA copy numbers concomitant with degradation

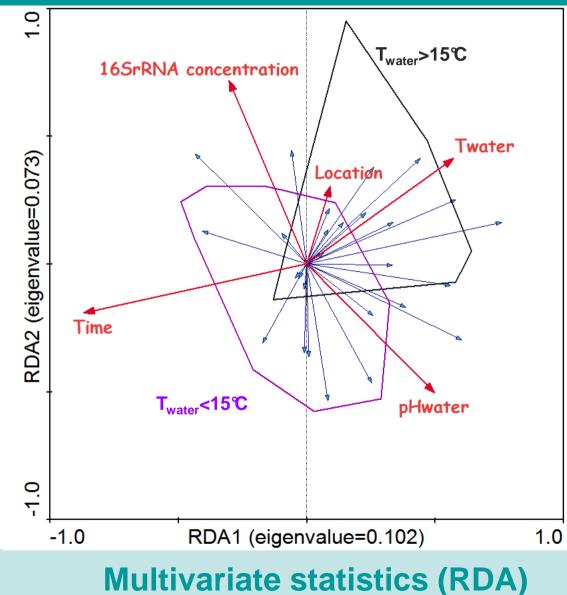




Dehalococcoides sp. specific qPCR

EBRO Dehalococcoides Diversity

DGGE fingerprinting



Conditional Effects

Variable	Р	
Time	0.002	
16SrRNA Conc.	0.004	
T _{water}	0.092	
рН _{water}	0.054	
Location	0.190	
Depth	0.146	
SPECIES (or DGGE	E Bands)	
SAMPLES		
◯ — Twater>	>15°C	
Twater⊲	Twater<15°C	
ENV. VARIABLES		
•		

rRNA Quantitative PCR profiling **Bacteria & OHRB** ✓ Ratio of 16SrRNA of OHRB to 1.2 Dcoc/UB 16SrRNA bacteria as a measure of 1.0 activity 0.8 2004 SUMMER 2005 WINTER % 0.6 2006 WINTER ✓ *Dehalococcoides* spp. appear 0.4 as most active OHRB 0.2 0.0 ✓ Activity of *Desulfitobacterium* **Rice Field** Delta Lleida Flix U5 Flix D5 Tortosa spp. in upstream locations Eff. Estr. 0.000 0.005 0.010 % 0.015 0.020 0.025 Dsulf/UB 16SrRNA 0.030

Routine HTP Monitoring of Bioremediation

✓ Multiplex assays:

✓ Allow detection of multiple targets in single experiment

✓ Reduce testing time and costs

✓ Challenges

- ✓ High number of different targets
- ✓ Very specific detection required
- ✓ Targets might be present in a wide concentration range
- Low concentration target DNA in high concentration background DNA

Routine HTP Monitoring of Bioremediation

Multiplex assays:

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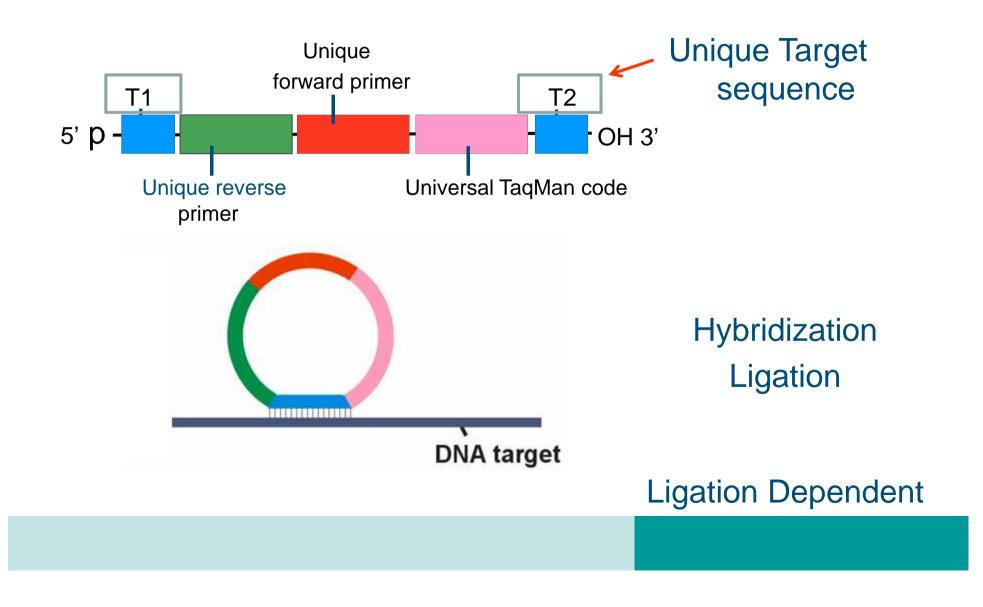
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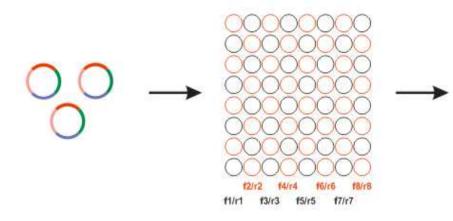
Technology

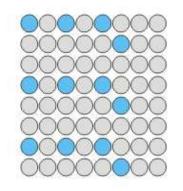
Circularization probe-based detection (Ligation probes)

PRI-Lock probe principle



Quantitative multiplex platform

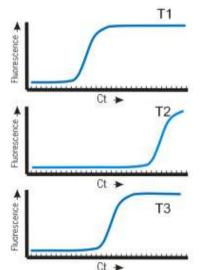


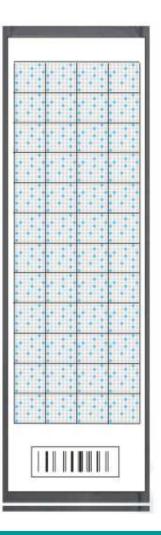


Multiplex PRI-lock ligation

singleplex probe amplification

Biotrove "33 nL Real-Time PCR array"

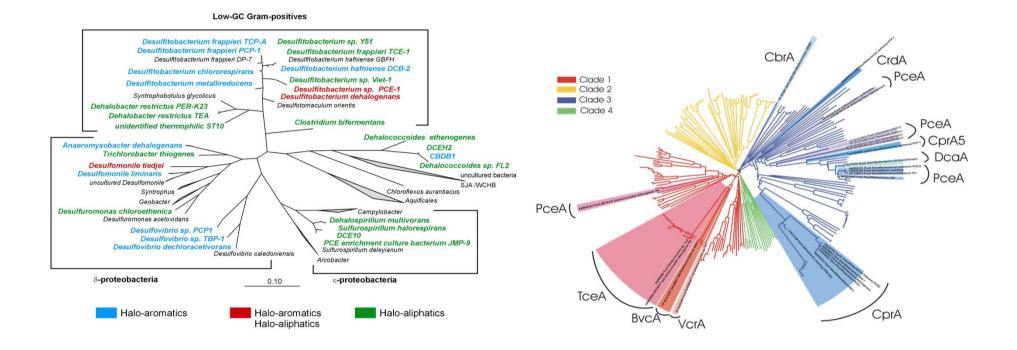




van Doorn et al., 2007

Quantitative multiplex platform

Targets

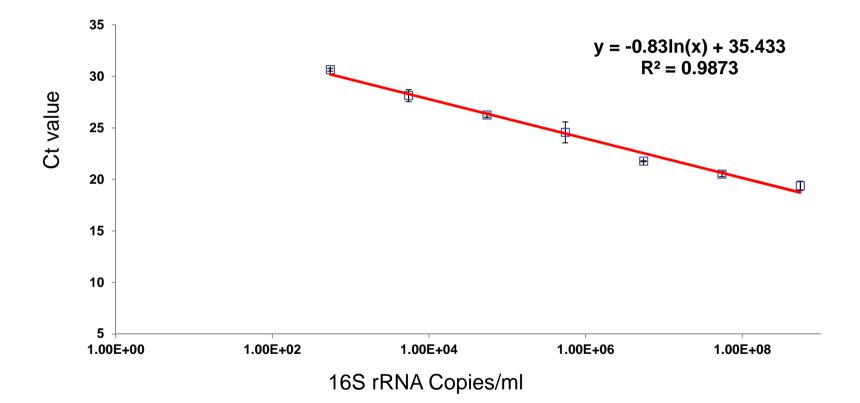


•Multiple PRI-lock ligation probes targeting representative

- •16S rRNA genes
- •reductive dehalogenase genes

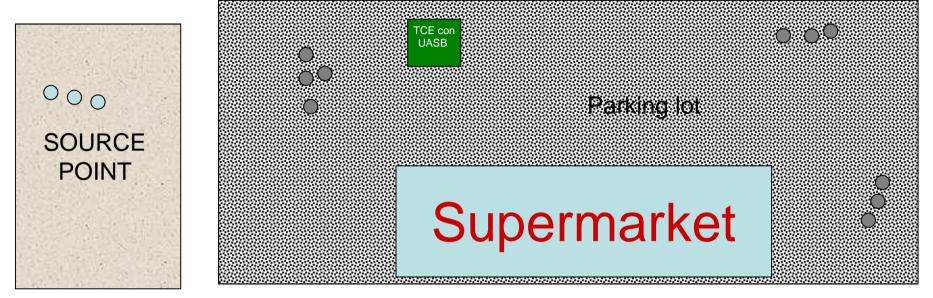
Probe Dynamic Range

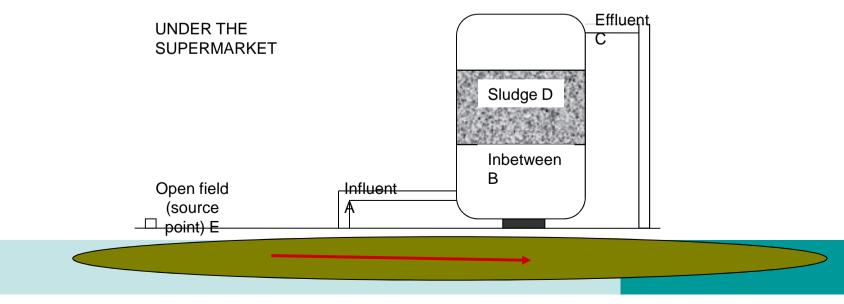
Representative Example: Standard Curve for Dehalococcoides Probe



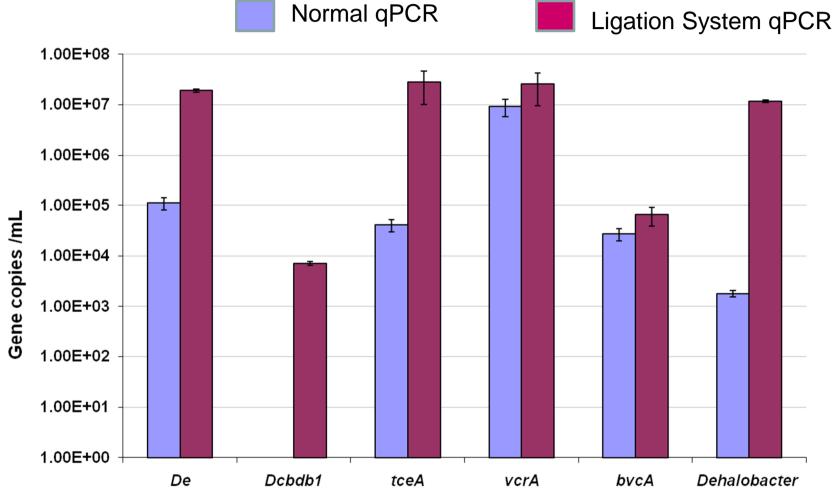
Detection Dynamic Range = 6 orders of magnitude

Cleaning Under The Supermarket...

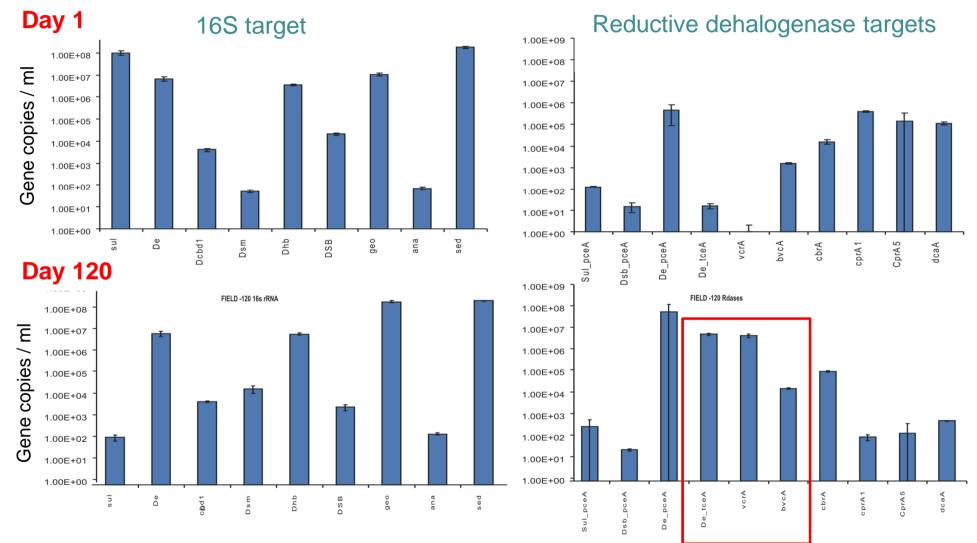




Profiling the Bioreactor SYBR qPCR vs Ligation System



Monitoring Field Samples

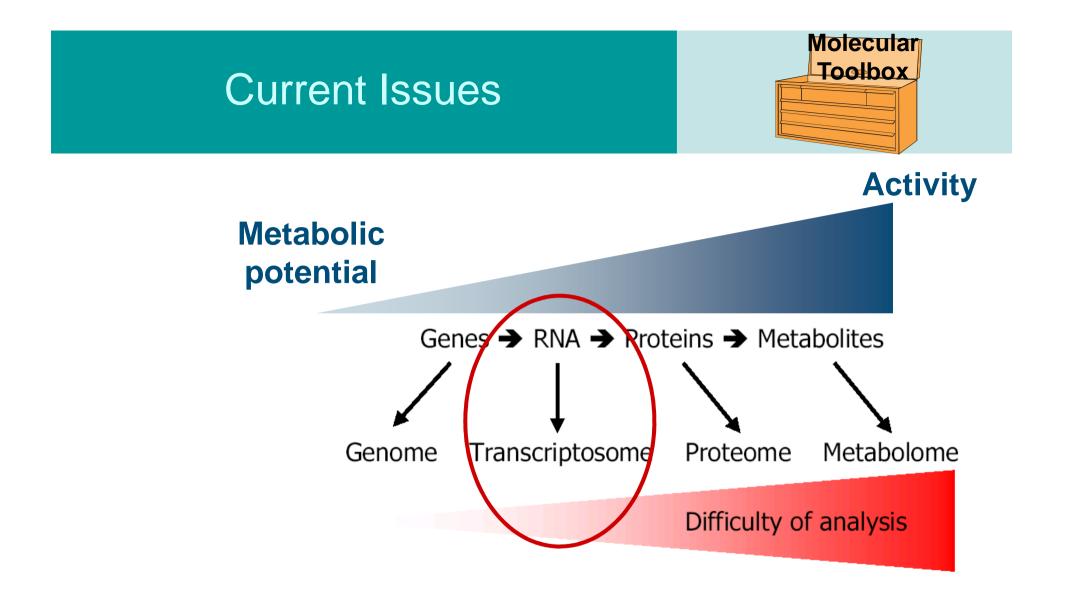


- Bioaugmentation with Dechlorinating culture :
- Increase in Reductive dehalogenases for Vinyl Chloride degradation

HTP Monitoring Current State of the Art

Quantitative multiplex target detection

- PRI-Lock Probes --- High specificity
- Single and multiplex target detection independent
- Detection Magnitude : 6-7 orders dynamic range
- Universal Pre- amplification improves sensitivity
- Universal TaqMan / SYBRGreen PCR conditions
- Applicable to various environmental systems and biomarkers



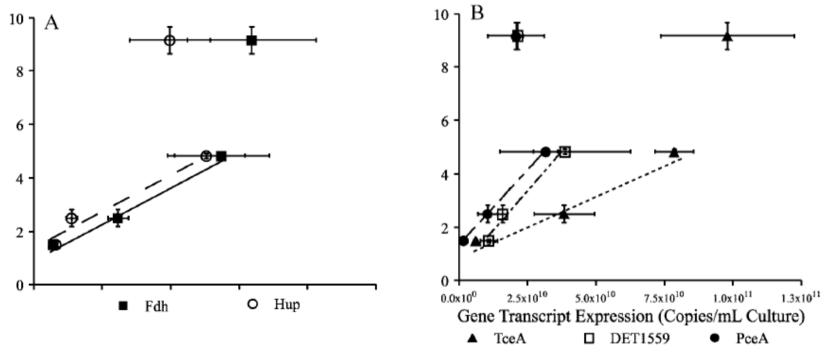
Can we actually deduce degradation rates?

Correlating gene expression Ra levels to respiration rates 200

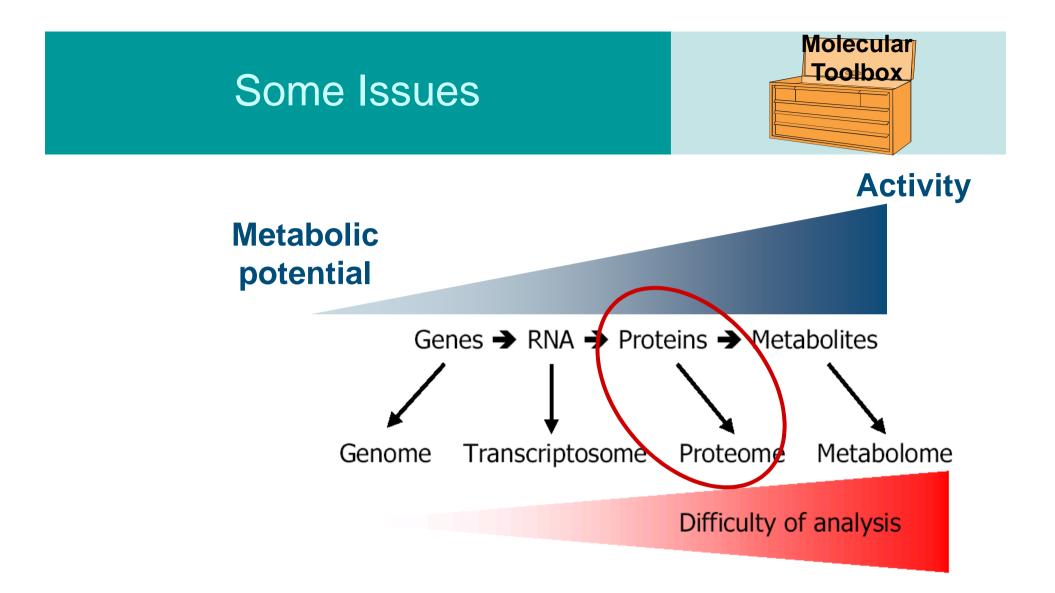
Rahm & Richardson 2008, EST

✓ Dehalococcoides ethenogenes 195

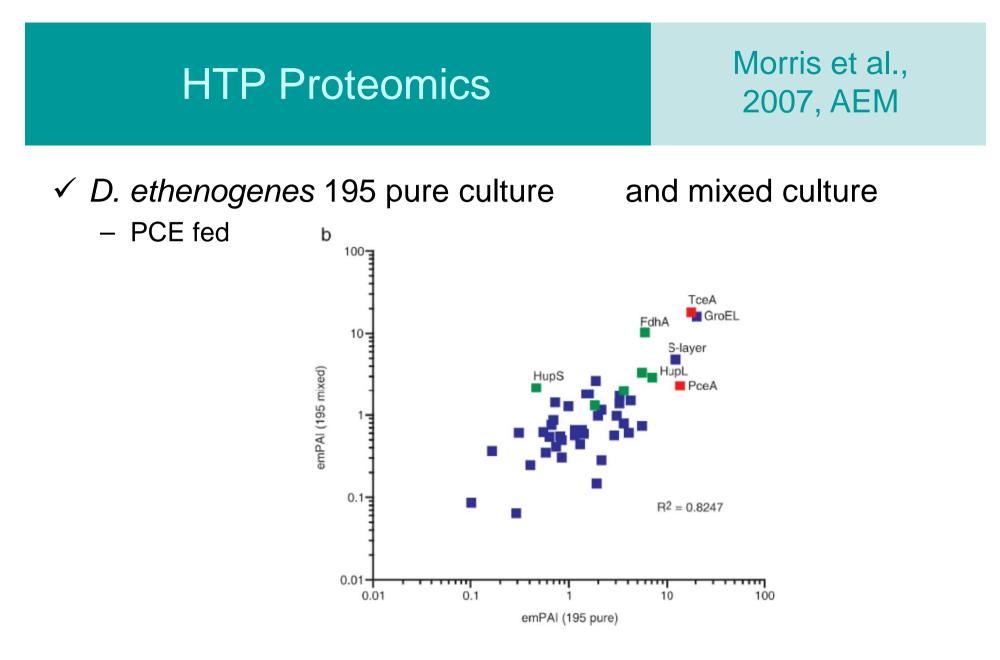




✓ Linear correlation across limited range of rates
 ✓ BUT: Rates probably strain (enzyme) specific!!



Can we look at proteins as the actual biocatalysts ?



Detection of respiratory enzymes (Rdh, H2ases, Fdh)

Special Thanks to ...

WU-Microbiology

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PRI

Ronald v. Doorn, Cor Schoen, Peter Bonants



AquaTerra

EU

BACSIN FP7 Framework Program

SEVENTH FRAMEWORK PROGRAMME