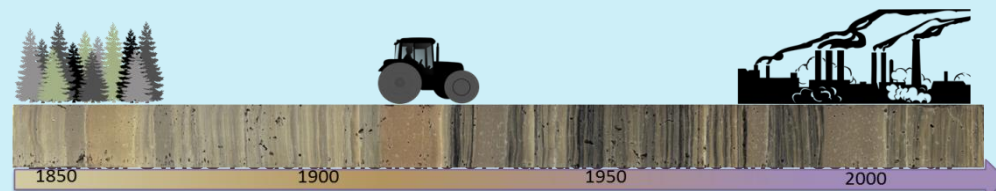


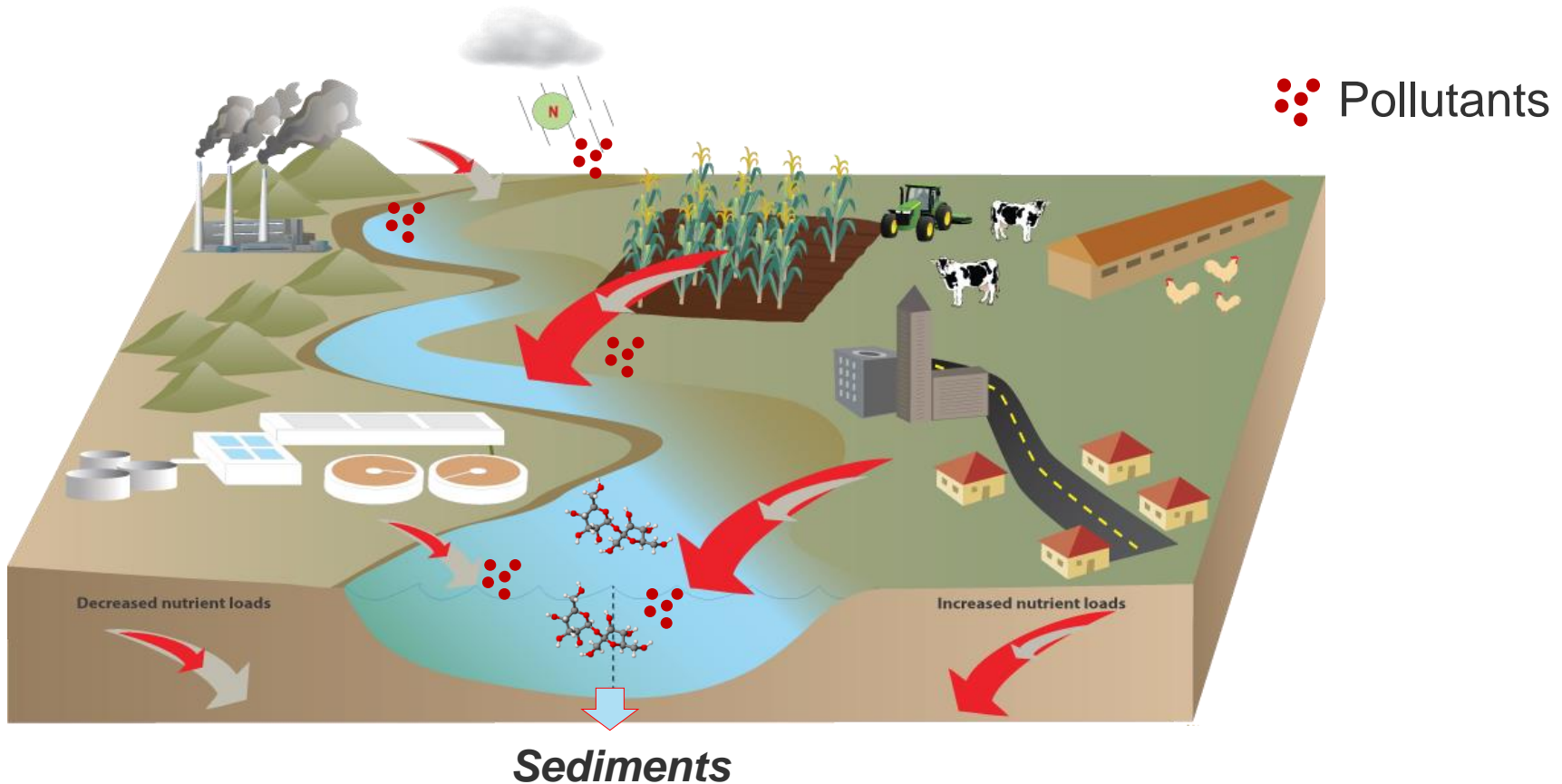
Prioritizing relevant nontarget contaminants using statistical analysis of LC-HRMS data of lake sediments

Aurea C. Chiaia-Hernandez, B. F. Günthardt, Martin P. Frey and J. Hollender

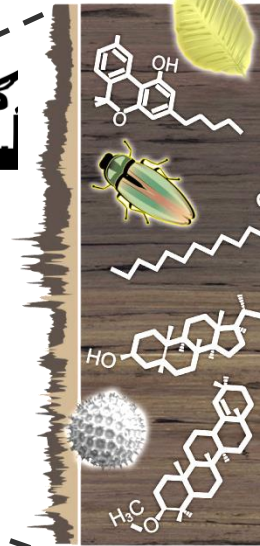
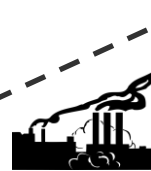
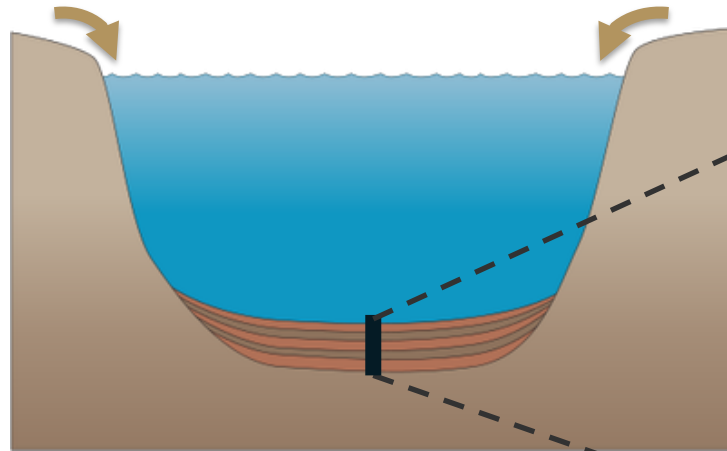


Lakes are Sensitive Barometers

- They respond to changes around them
 - Climate change (e.g. changes in rainfall)
 - Catchment change (e.g. caused by people)



Sediments = History Books of the Environment

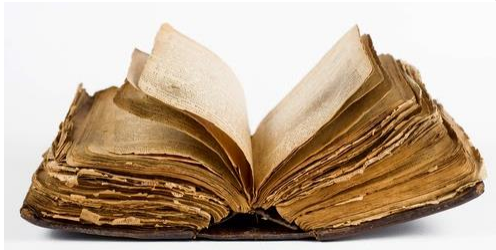


2000

1900

1800

1700



M. LAVRIEUX

Matrix



Age model

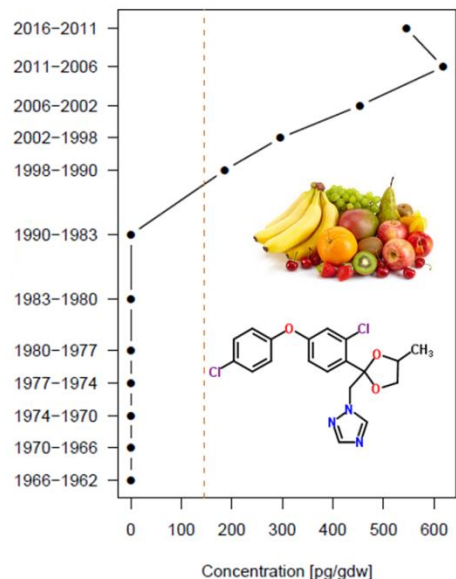
1900

1950

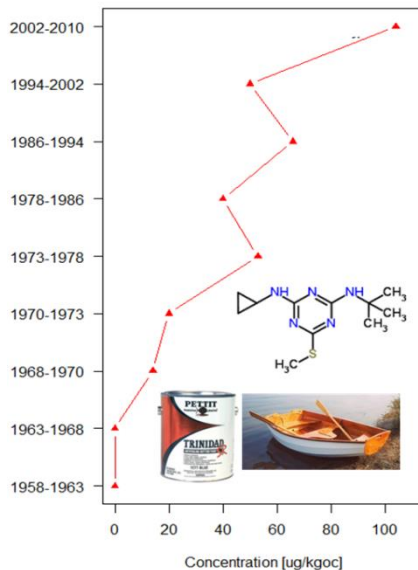
2000

Target and Suspect Screening by HRMS – Known Compounds

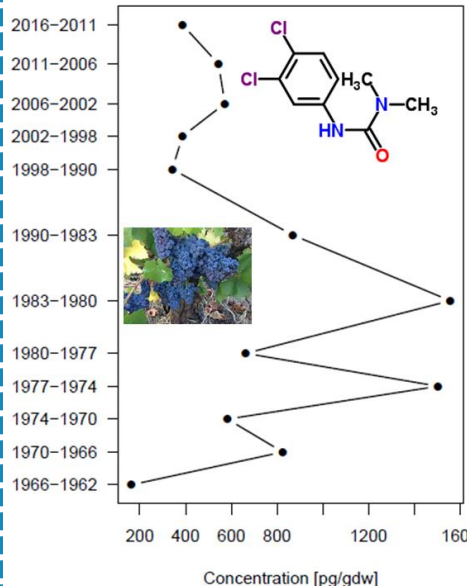
Difenoconazole



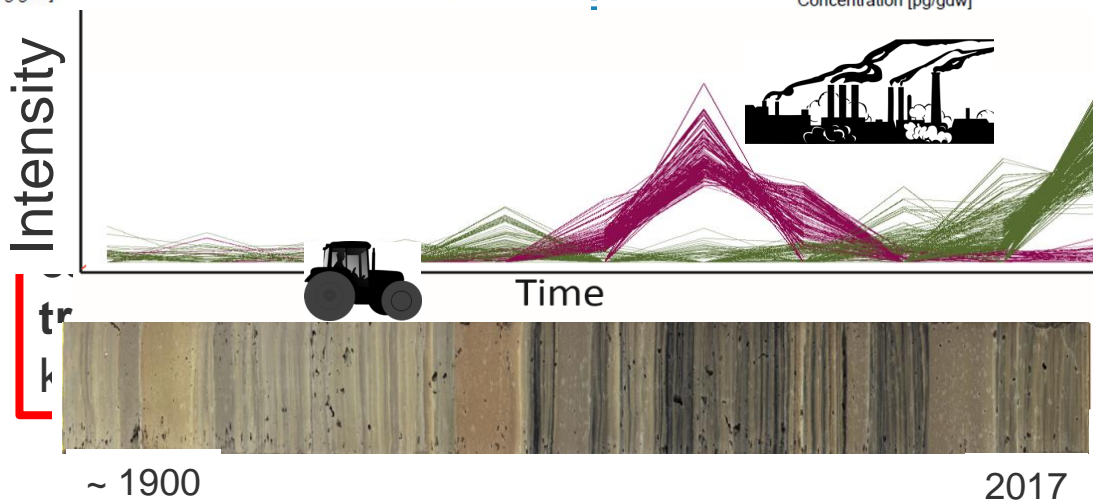
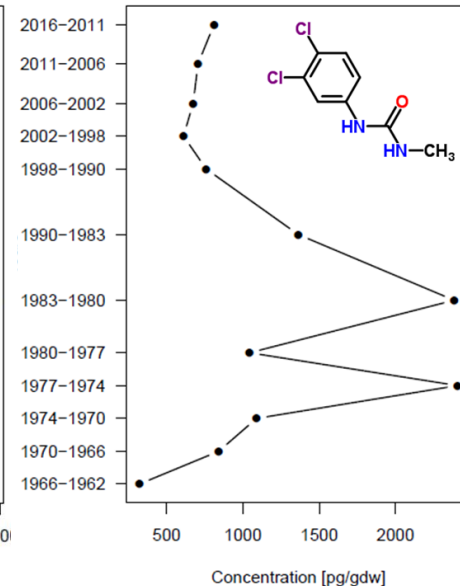
Irgarol



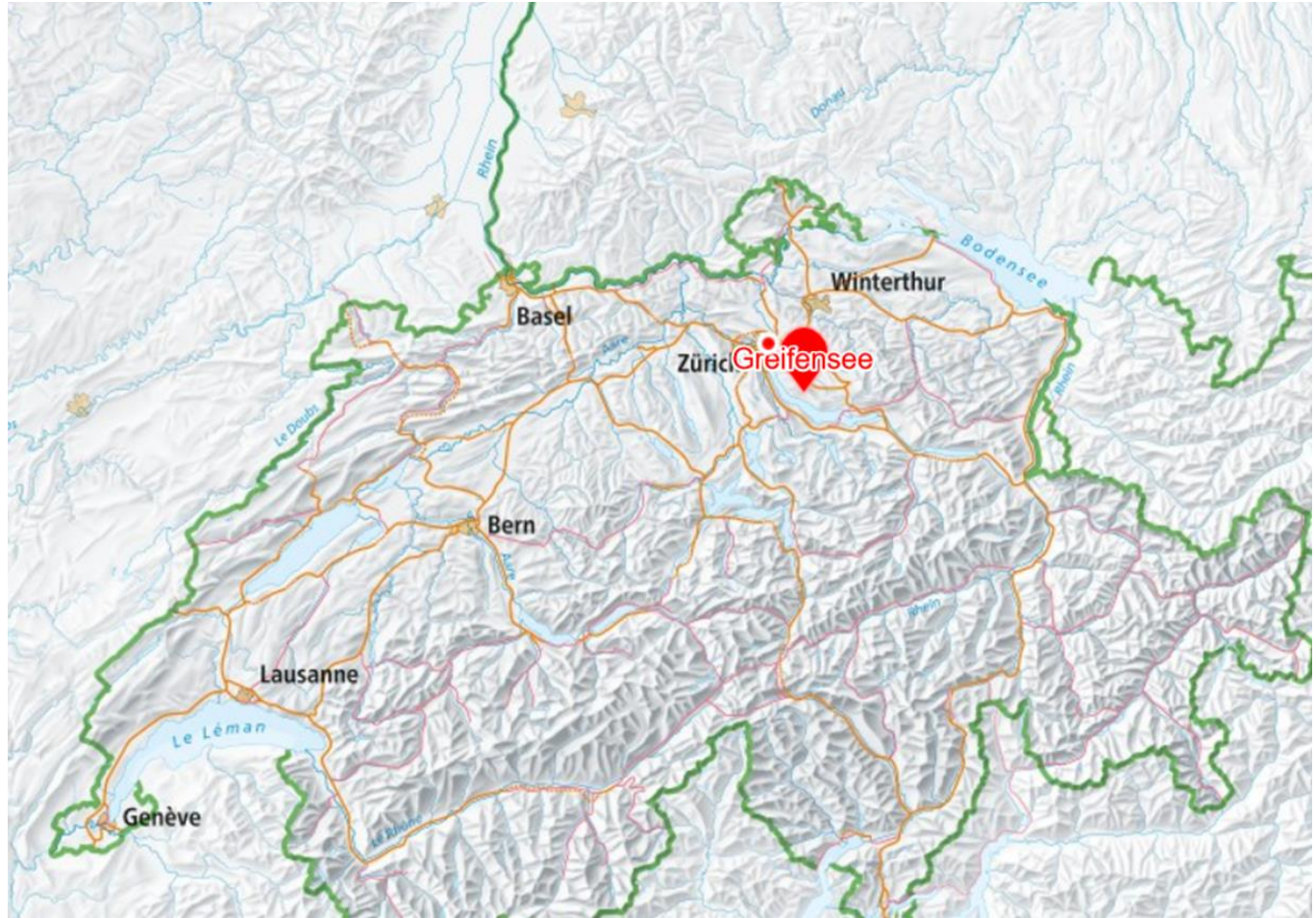
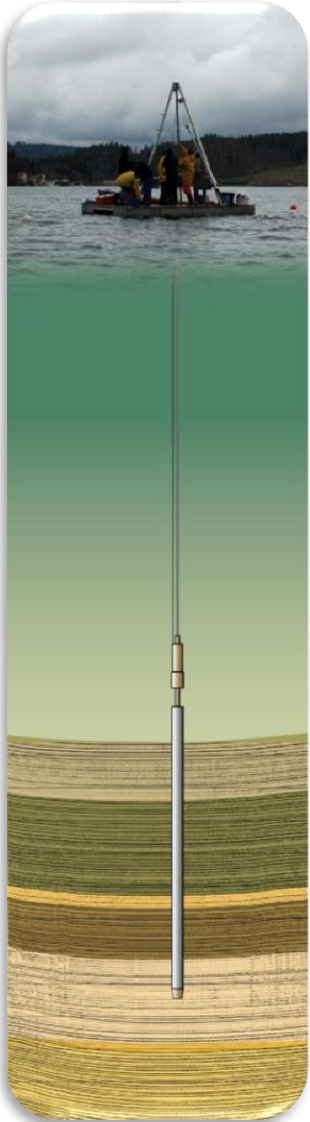
Diuron



Diuron-desmethyl



1. Collecting Lake Sediments



2. Profile Detection- enviMass

enviMass v3.2

Project folder:
E:/enviMass/Lugano_analys:

Current state:
Opened existing project

Project state:

Tasks	Status
Peak picking	✓
Compound patterns	✓
Quality control	✓
Mass recalibration	✓
Median intensity normalization	✓
align	✓
EIC_correlation	✓
Replicate filter	✓
Blind subtraction #1	✓
Profile extraction	✓
LOD interpolation	✓
Isotopol. grouping	✓
Adduct grouping	✓
Homologue detection	✓
IS screening	✓
Target screening	✓
Calibration	✓
IS-based intensity normalization	✓
Profile filtering	✓
Trend detection	✓
Quantification	✓
File componentization	✓
Recovery	✓
-	✓
excluded / not run	✗

Files | Compounds | Workflow options | Settings | Calibration | Results | Help & About

Data viewer | Quality control | Processing | Screening & quantification | Grouping | Profiles

Select ionization (switch to negative):
positive

Summary | Latest trends | Single Profile | Normalization

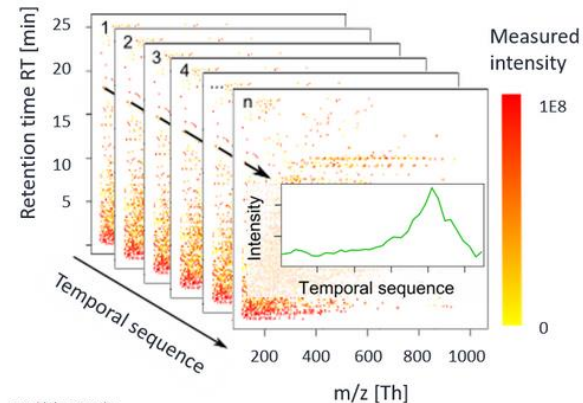
Extraction of individual profiles

Enter the ID of a profile to extract relevant information. Profile IDs are listed both in the Summary tab and the ... choose an entry number to show a listed profile.

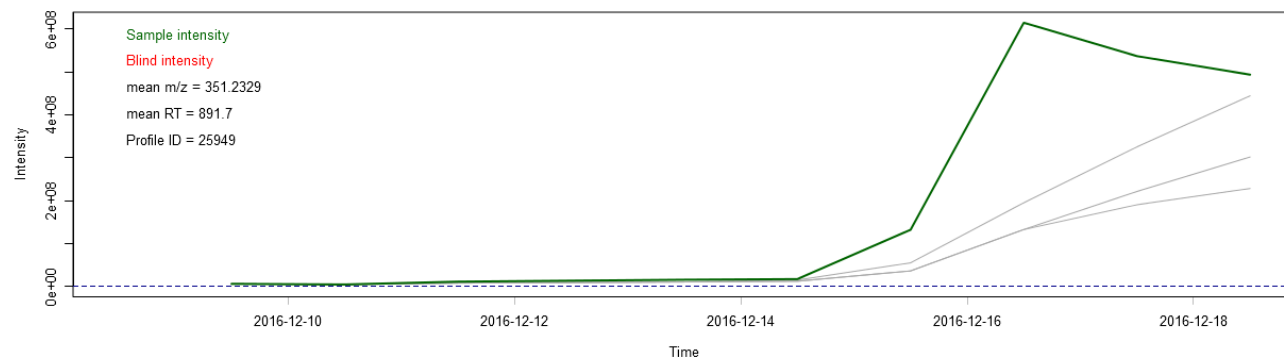
profile ID:

Entry # in (filtered, sorted) profile list:

Logarithmic intensity: no yes



Draw rectangles and double-click into them to zoom, double-click again to zoom out.



Profile Detection (enviMass) vs. Concentration Profiles

Irgarol

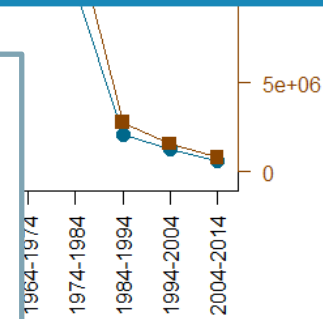
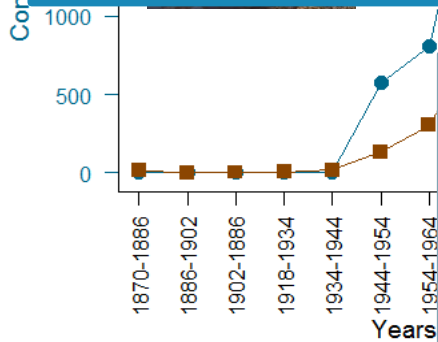
Dichlorophene

= Fully automated workflow

+ > 71,000

- > 148,000

Concentration (ng/l)



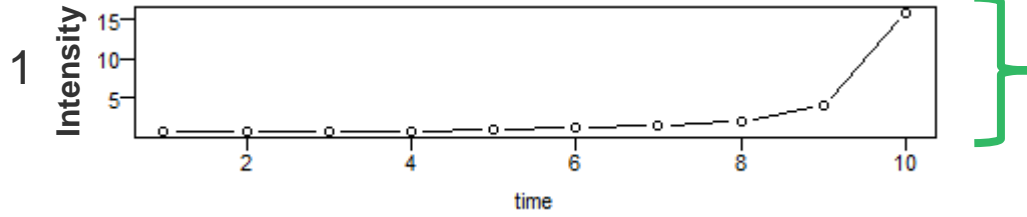
● Measured concentration

...ed using ...
...s

Matrix

M. LAVRIEUX

3. Trend Characterization: Spearman's Rank Correlation

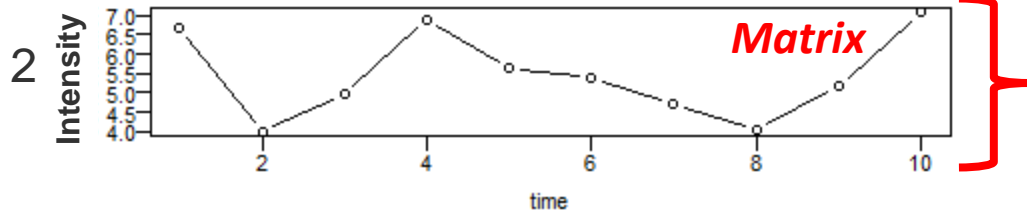


Pearson (r)

0.66

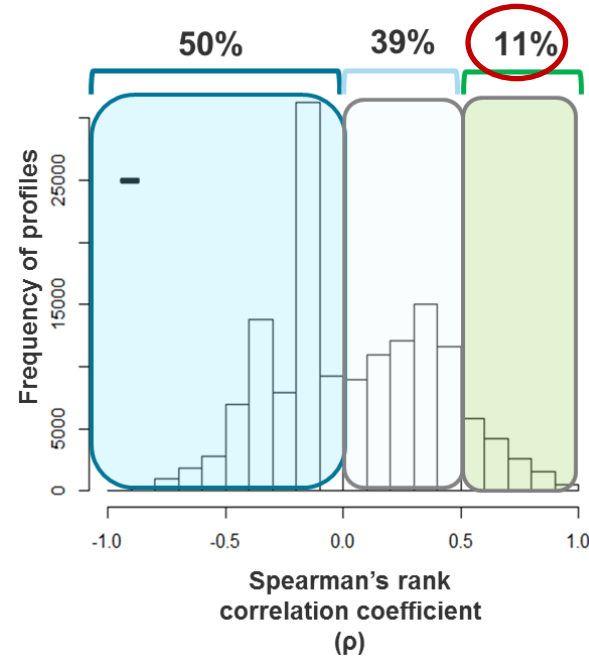
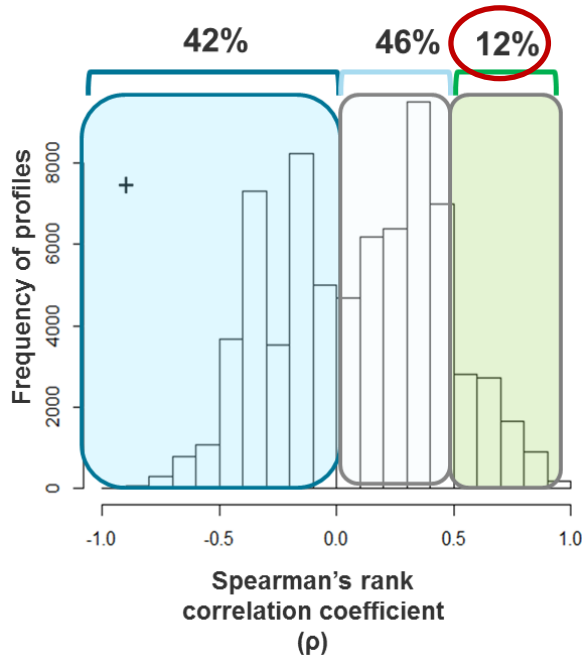
Spearman (ρ)

0.95



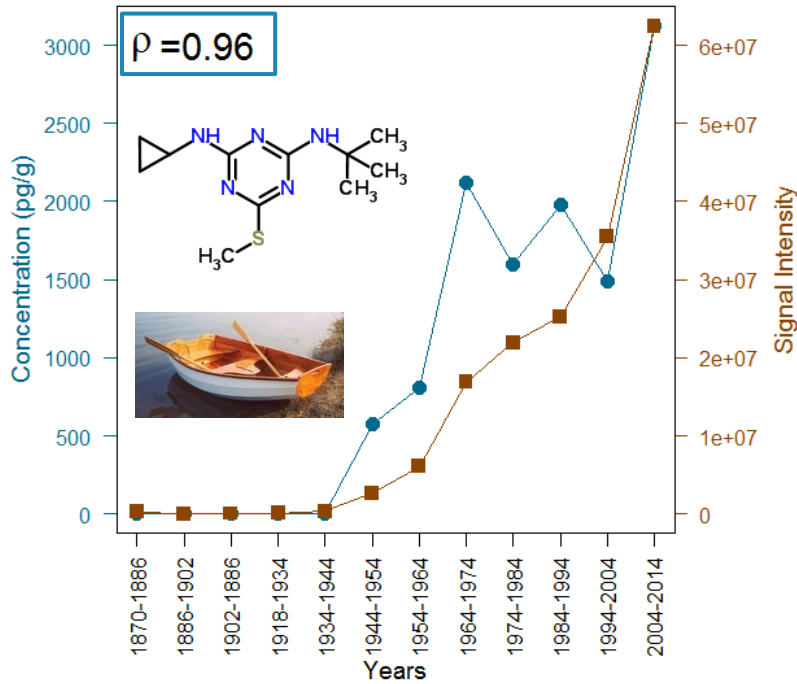
0.00

0.10

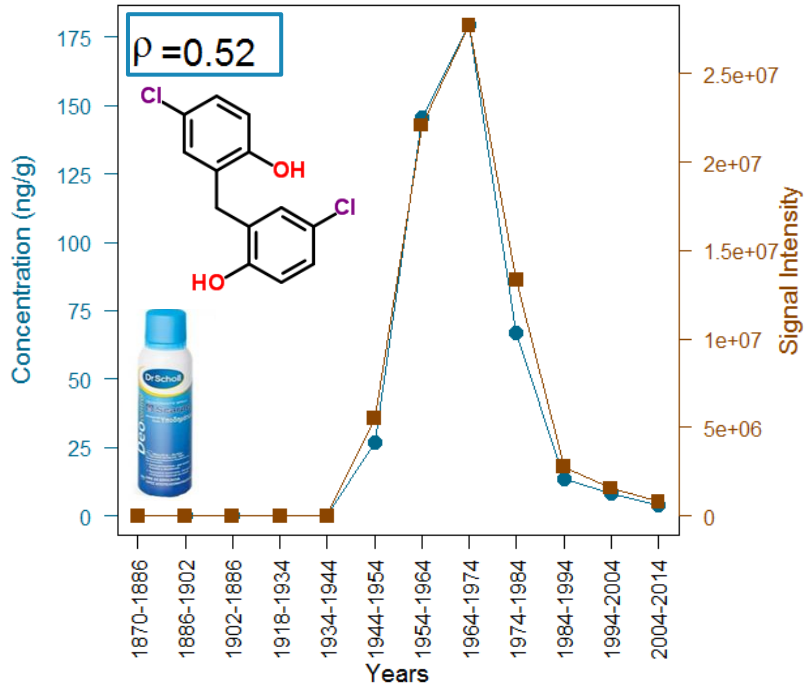


Trend characterization 1: Spearman's rank correlation

Irgarol



Dichlorophene



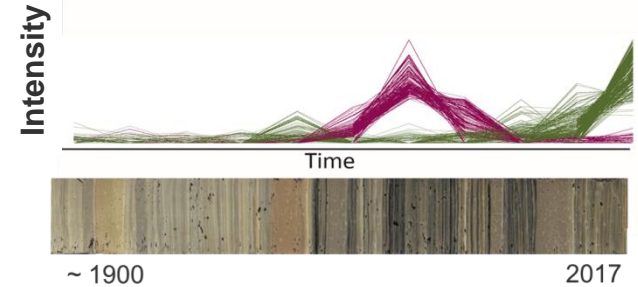
● Measured concentration

■ Profile extracted using enviMass

4. Trend Characterization: Hierarchical Cluster Analysis

Cluster analysis: Group samples that are similar

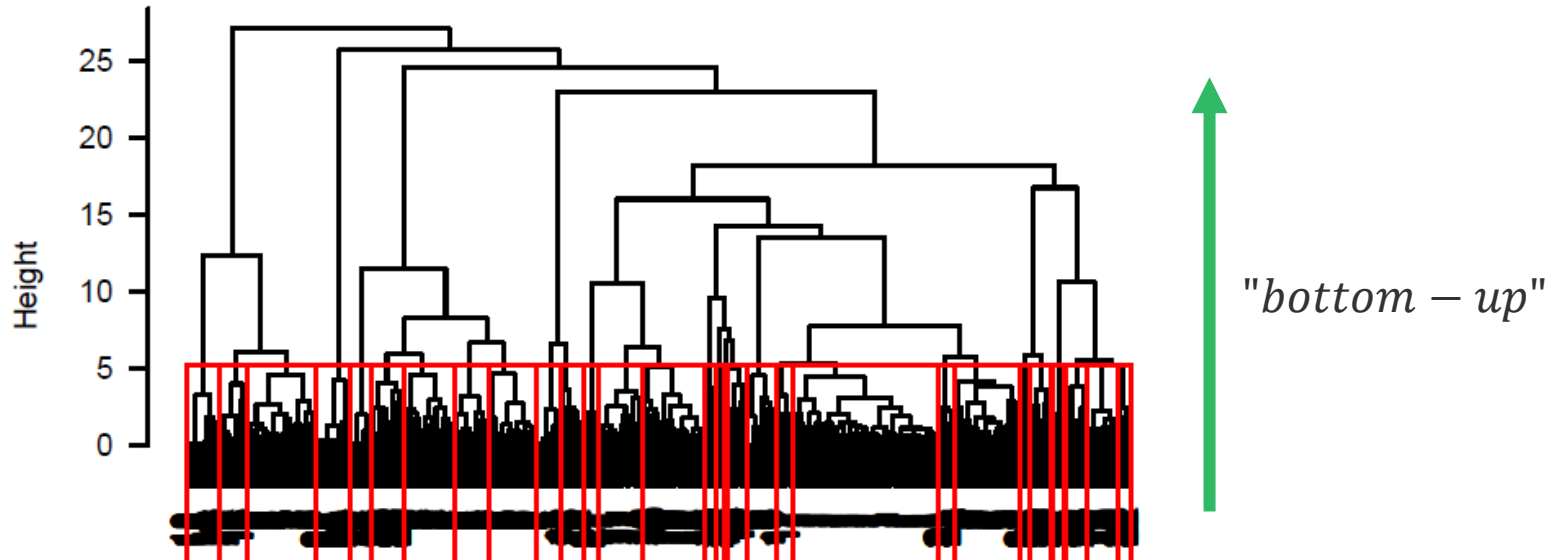
- Intensity patterns in sediments



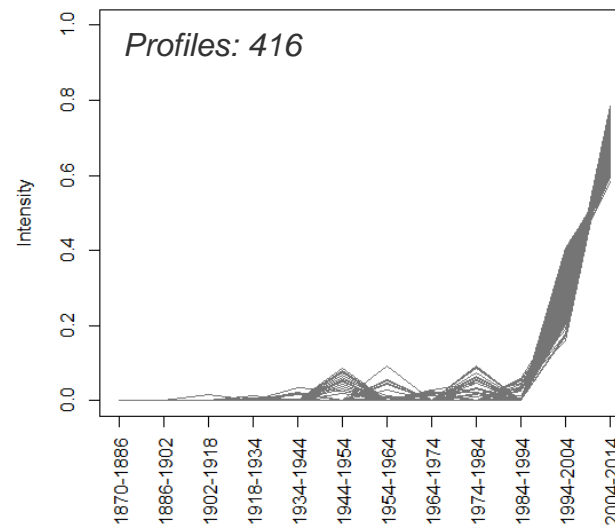
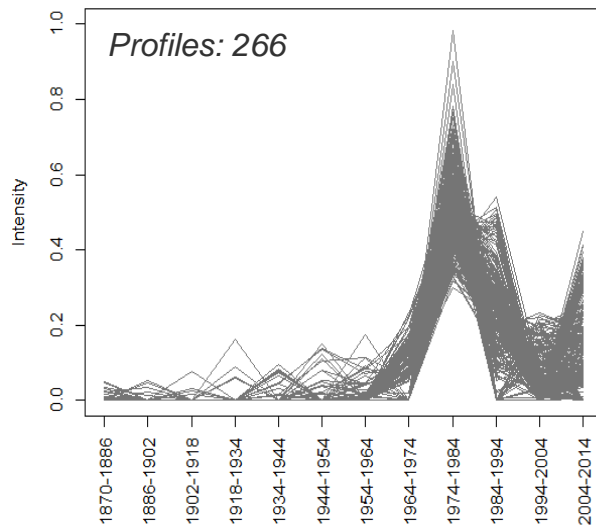
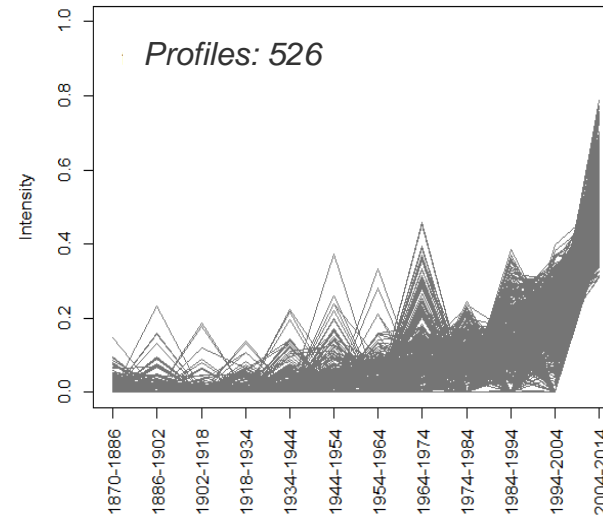
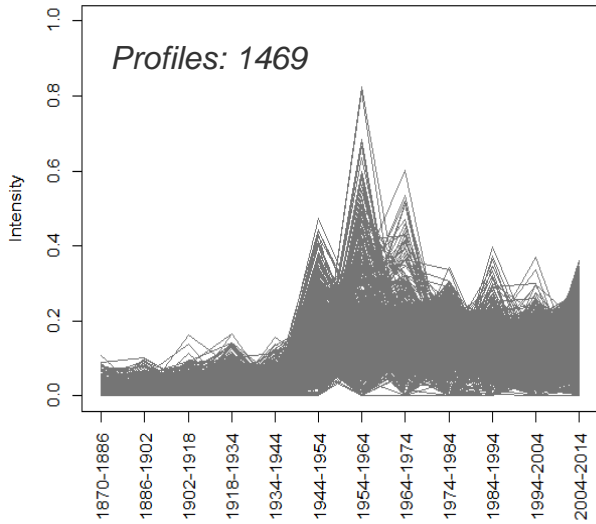
Agglomerative hierarchical clustering

- Classification
- Information retrieval

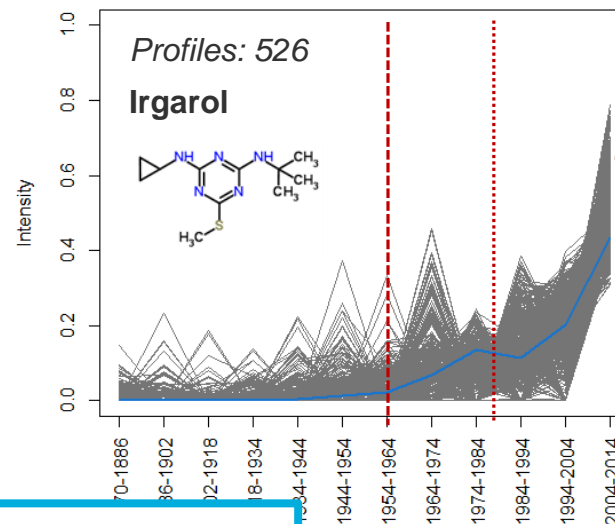
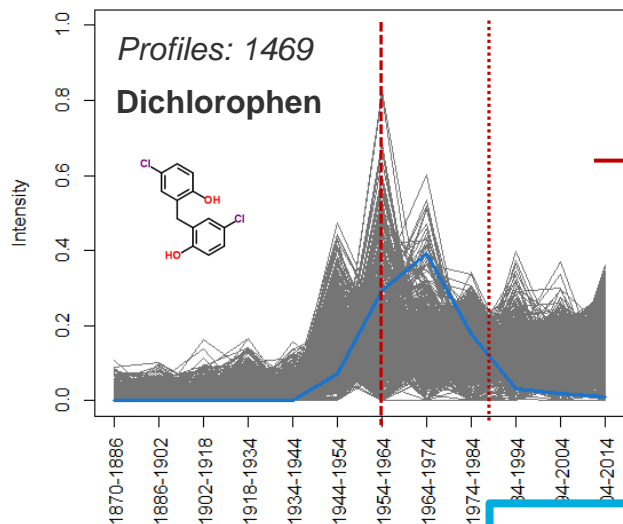
Dendrogram



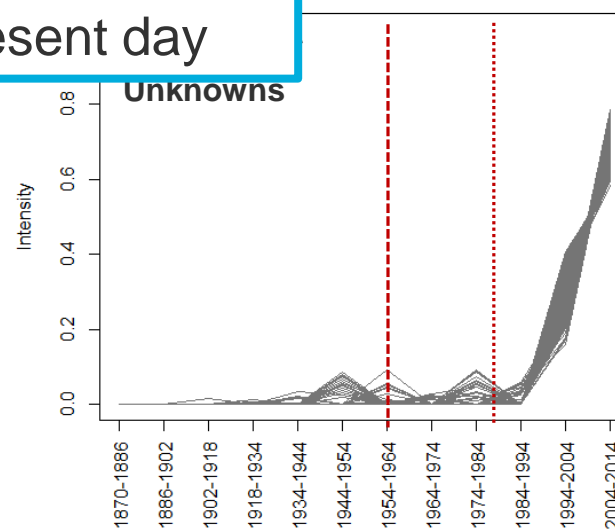
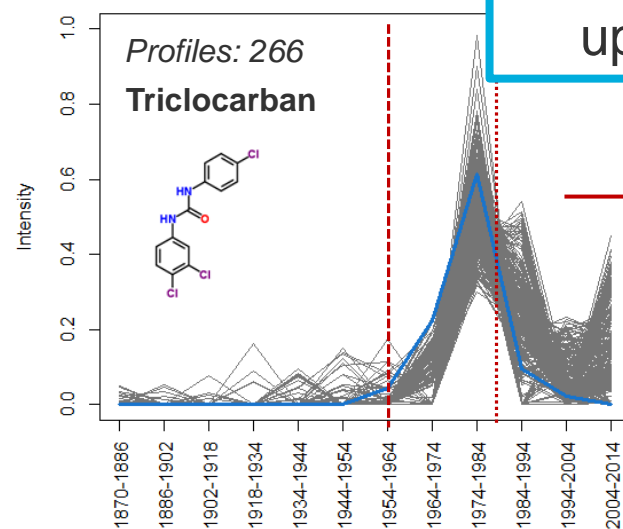
Hierarchical Clustering- Greifensee



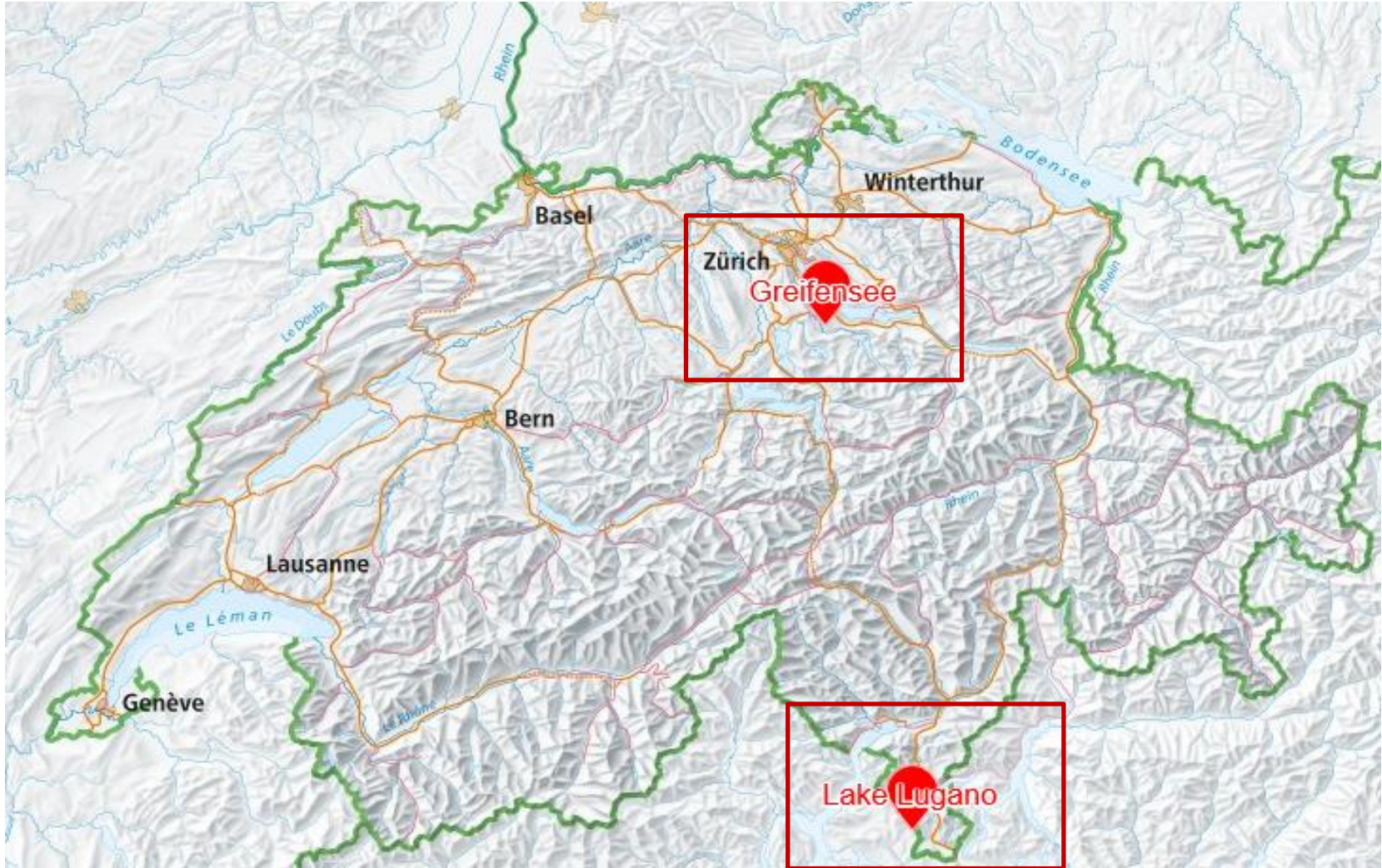
Hierarchical Clustering- Greifensee



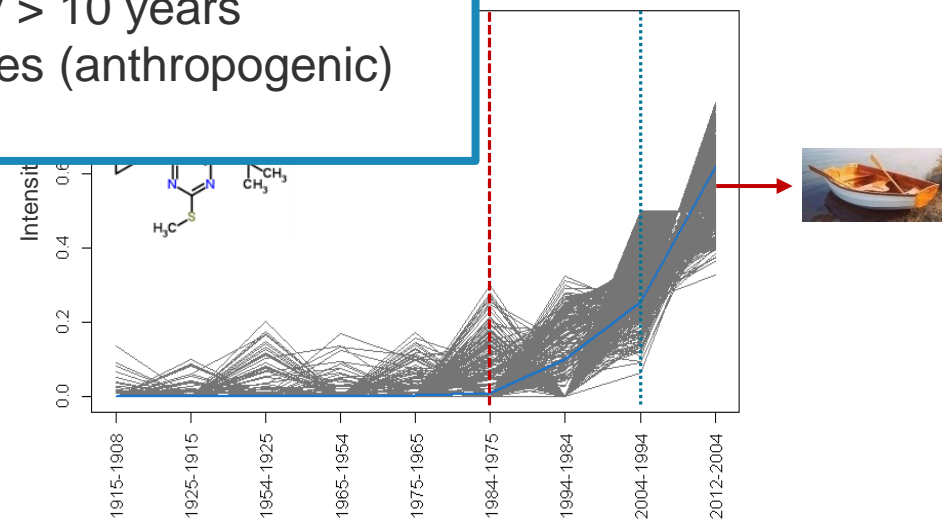
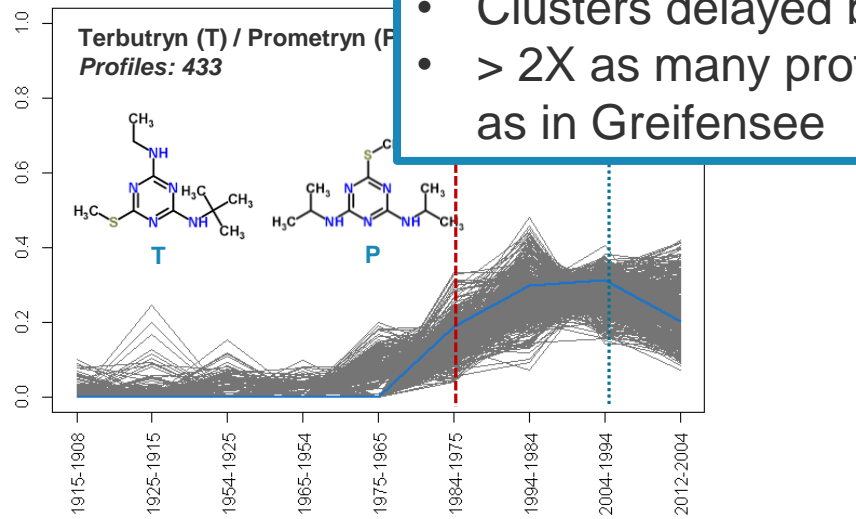
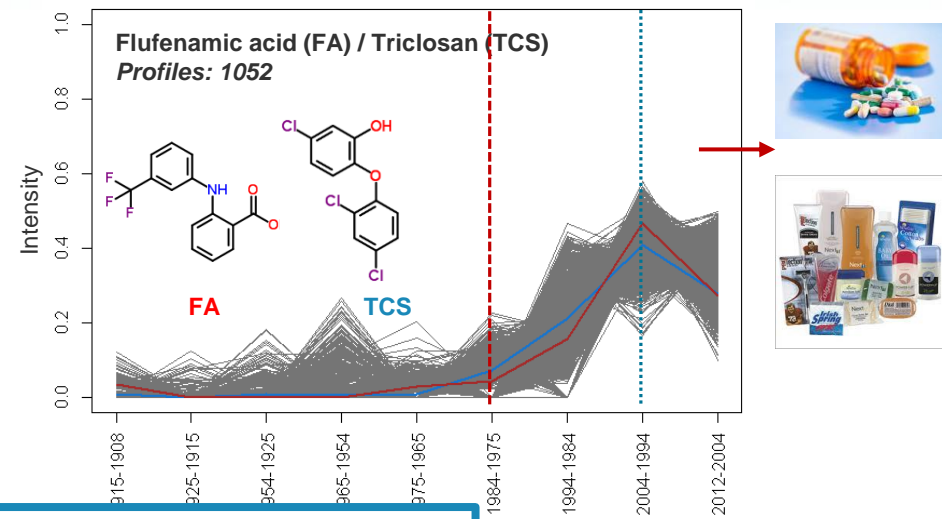
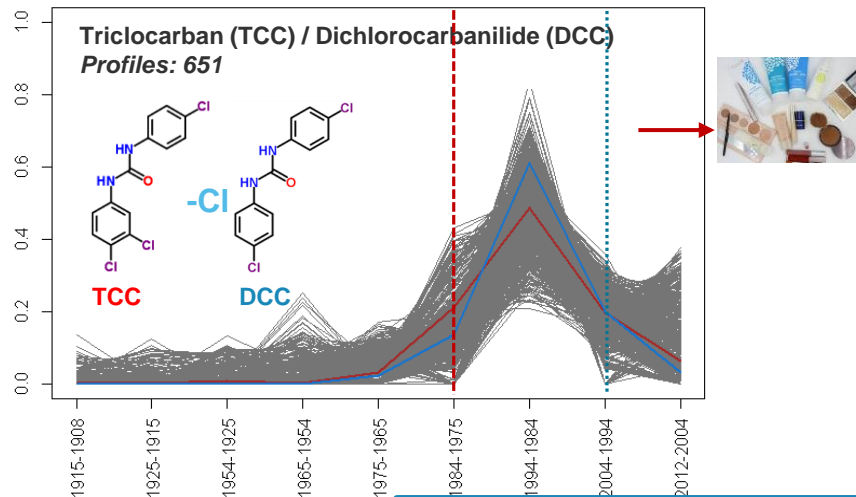
> 2,000 profiles with increasing concentrations up to the present day



Hierarchical Clustering- Lake Lugano

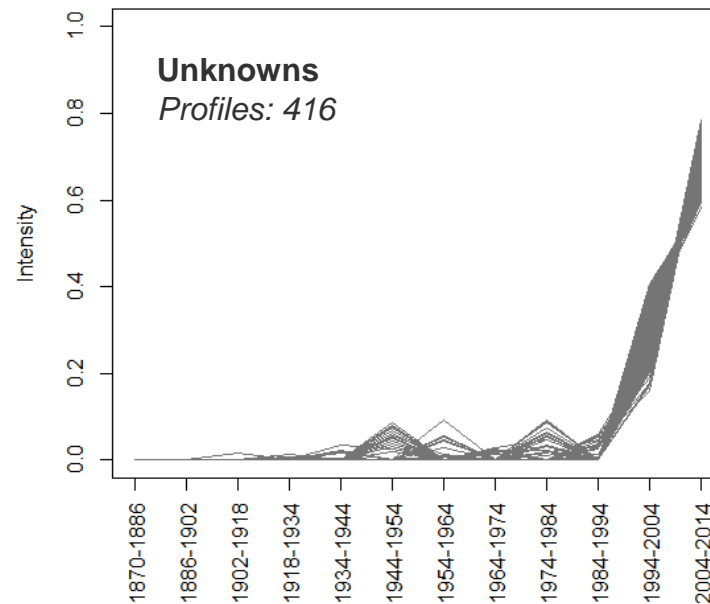
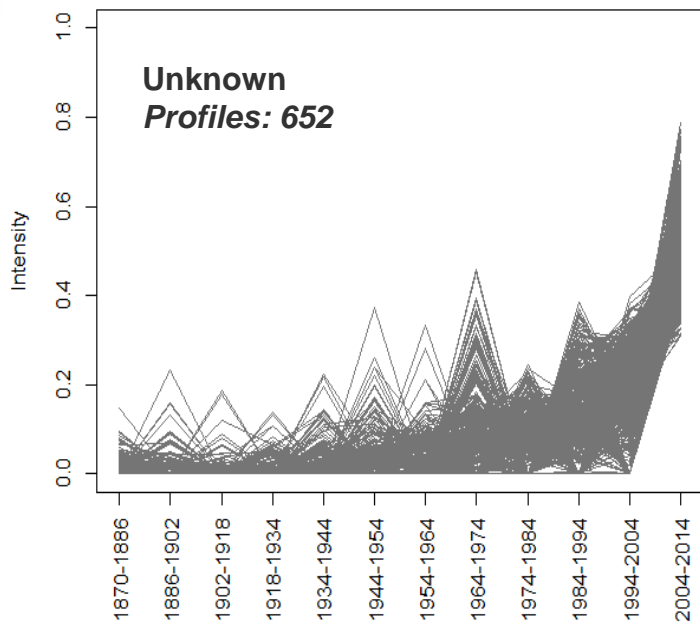


Hierarchical Clustering- Lake Lugano



- Clusters delayed by > 10 years
- > 2X as many profiles (anthropogenic) as in Greifensee

Prioritization of Relevant Nontarget Contaminants



List of candidates
Re-measured at different collision energies (HCD) of 15, 35, and 55, 75 and 90 %

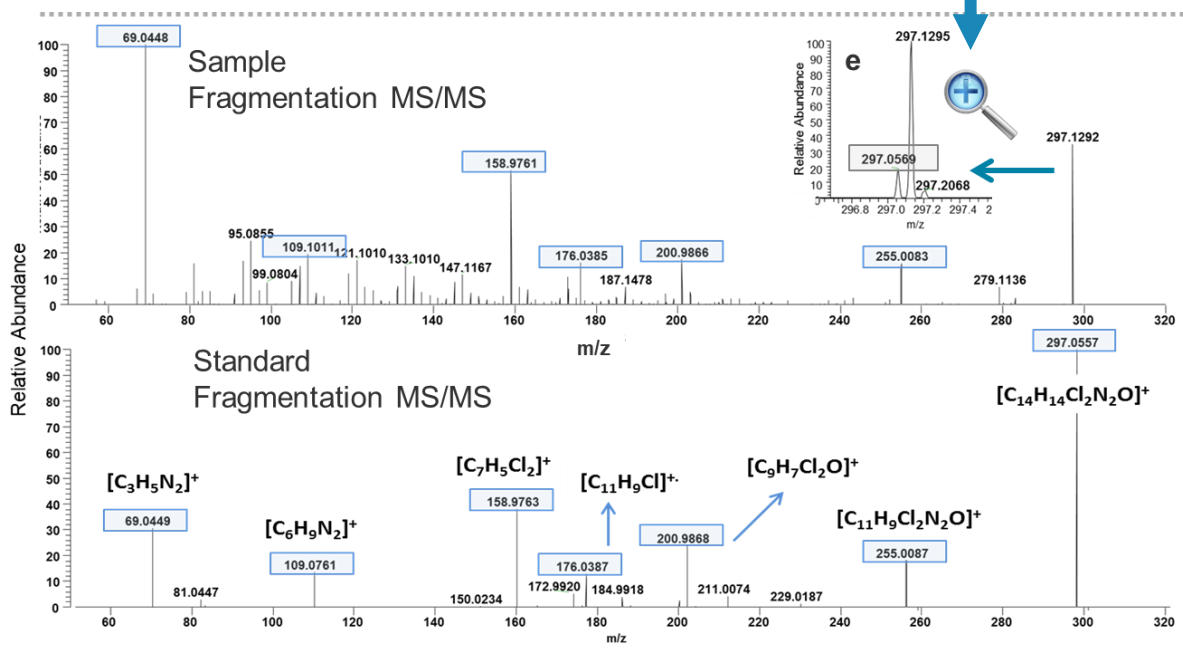
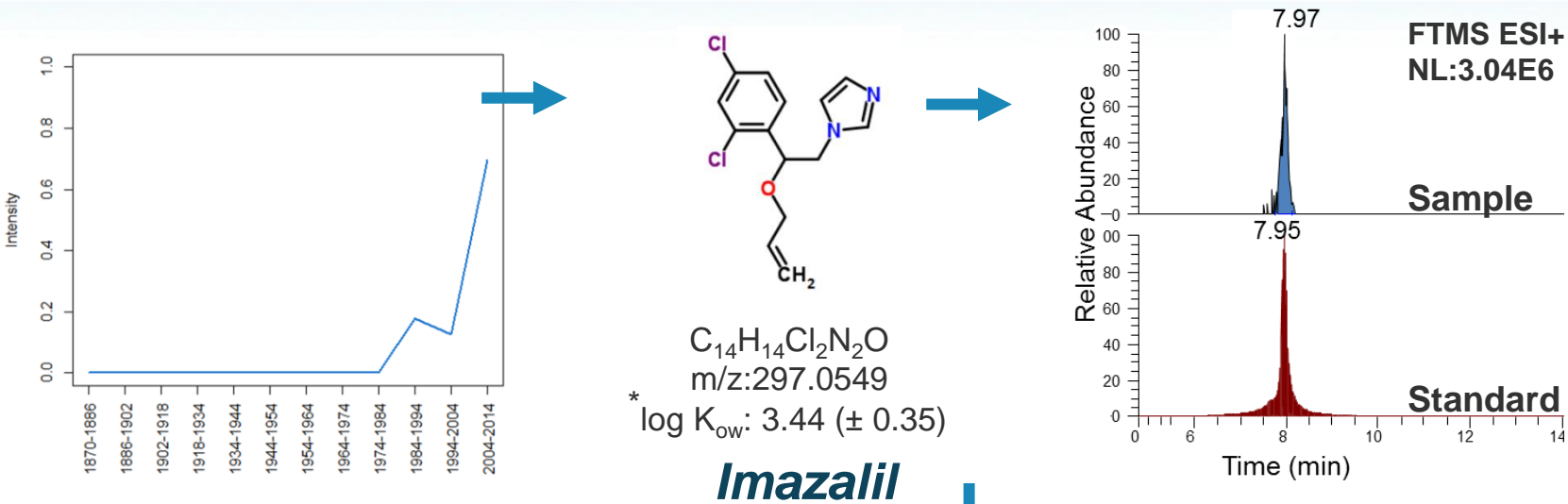
prioritized masses (+)

- $>10^6$
- very characteristic isotopic pattern



Match to different DB
MetFrag 2.3 (R version)

Identifying Nontarget Contaminants



- < 50 kg/year in agriculture
- Veterinary product (enilconazole)



- "post-harvest fungicide in citrus fruit "

Conclusion:

- Spearman's rank correlation coefficient help to identify substances with increasing trends over time
- Hierarchical clustering is a very useful method to evaluate the contamination and distinguish compounds with different trend patterns- *Transfer to other matrices*
- Prioritization of non-targets using statistical tools is promising to reduce matrix interferences and focus on relevant contaminants
- Sediments are useful to identify chemicals with unrecognized input pathways



Adi Müller (eawag)
Martin Loos (eawag)
Emma Schymanski (eawag)
Heinz Singer (eawag)
Uchem Department-eawag
SURF Department-eawag

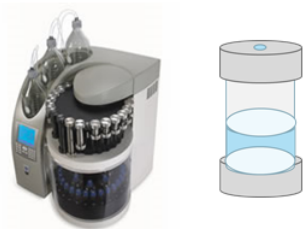


Funding:
Federal Office for the Environment (BAFU)
Academic Transition Grant (Eawag)

Analytical Methods



Preservation and Storage



Extraction and Enrichment

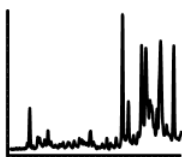
- In-cell clean up (Florisil)*



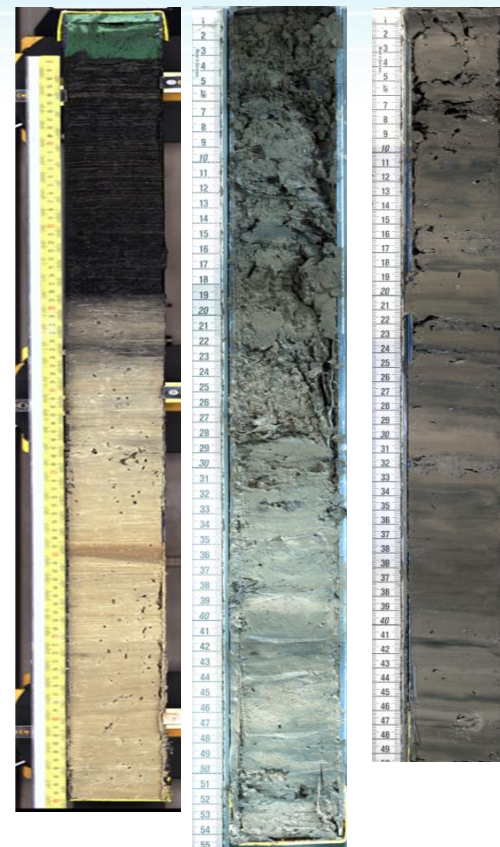
Chromatography



Orbitrap-MS
HR Chromatogram



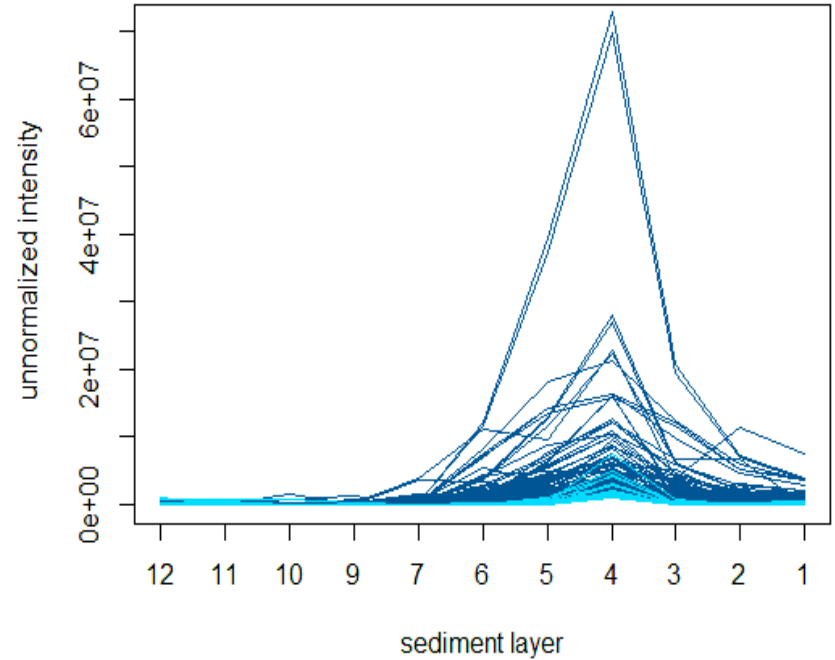
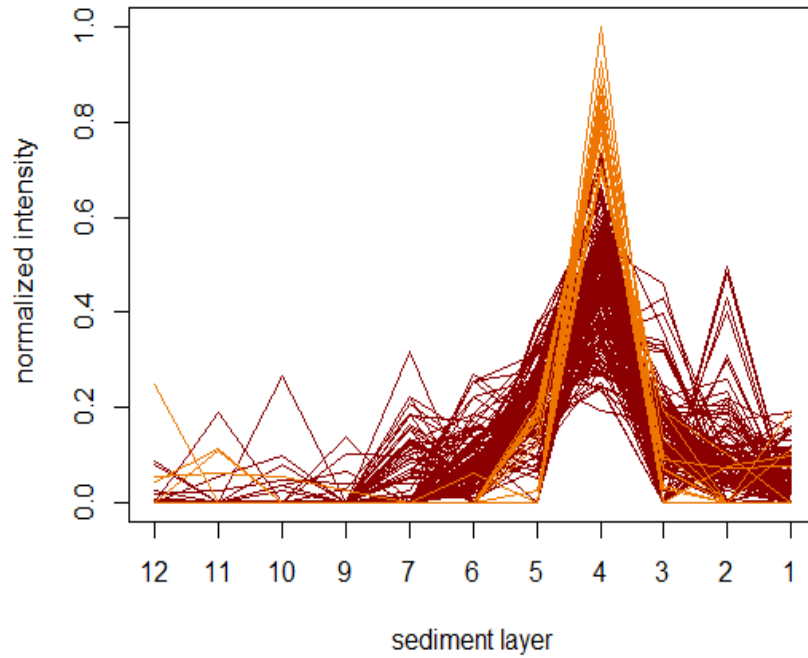
1. Target
2. Suspect
3. Non-target



- Generic method
- Non-compound class specific

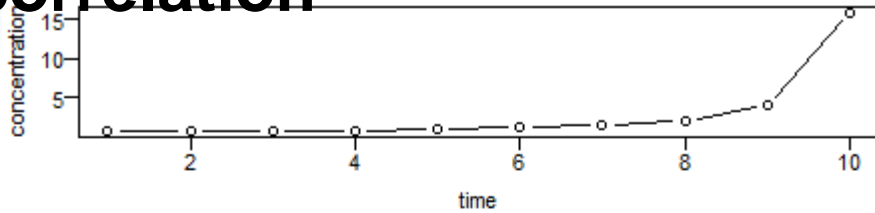
Appendix:

Normalization cluster analysis:



Trend characterization 1: Spearman's rank correlation

1 correlation



interested

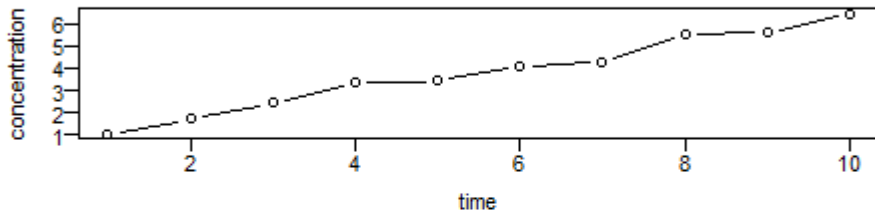
Spearman

0.95

Pearson

0.66

2

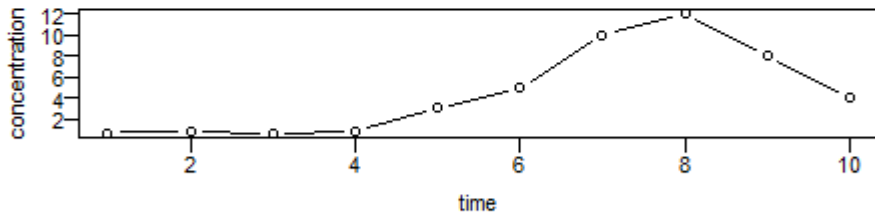


interested

1.00

0.99

3

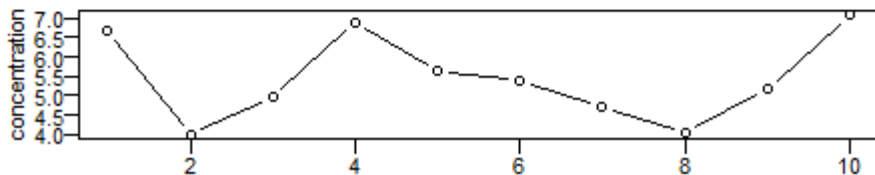


not interested

0.80

0.73

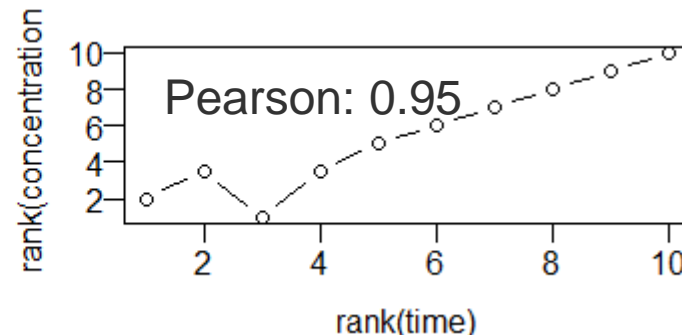
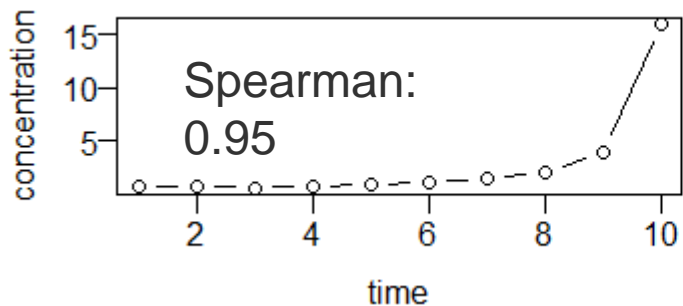
4



not interested

0.10

0.00

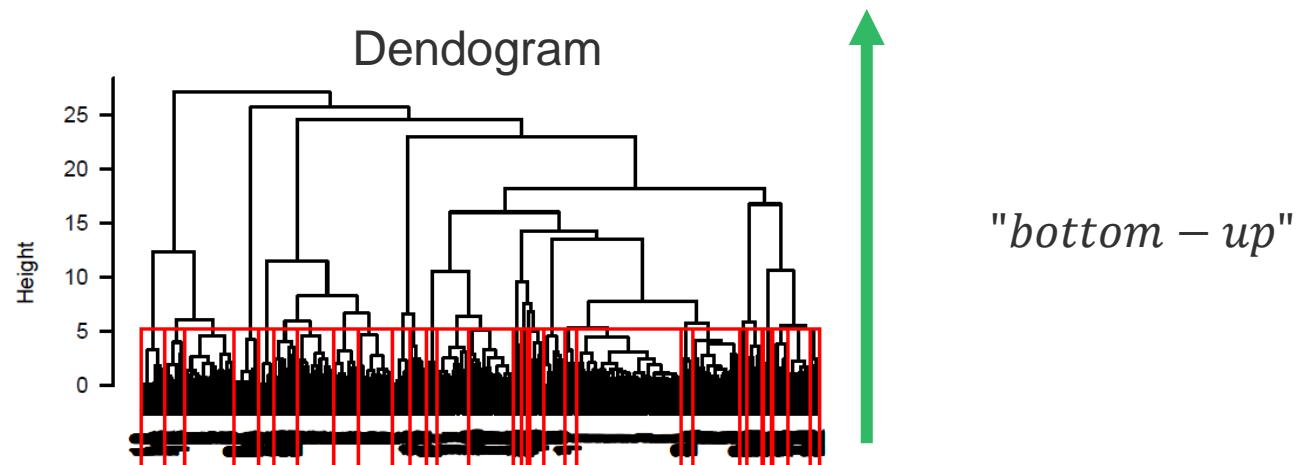


Cluster Analysis

- **Cluster Analysis (or Data segmentation)**
 - **Goals:** to group a collection of objects into subsets or "clusters"
 - Different types of cluster analysis

Hierarchical Clustering

- Classification and information retrieval
- Seeks to build a hierarchy of clusters
- Two basic paradigms: **Agglomerative (bottom up)** and divisive (top-down)

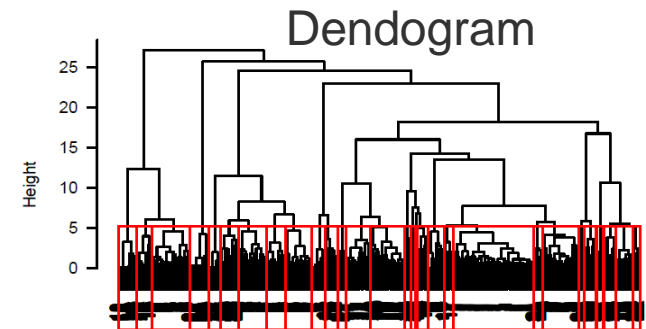


Hierarchical cluster analysis

Four main steps:

- I) Normalization (over the sum)
- II) Distance matrix (Euclidian distance)
- III) **Hierarchical clustering (Ward's minimum variance method)**
- IV) **Define number of clusters (some knowledge)**

In R
library(cluster)



Function within the package:

Daisy: Distance matrix

Arguments - methods

euclidian= root sum-of-squares of differences

Function within the package:

Hclust

Arguments - methods

ward.D2 = Ward's minimum variance method

Hierarchical cluster analysis

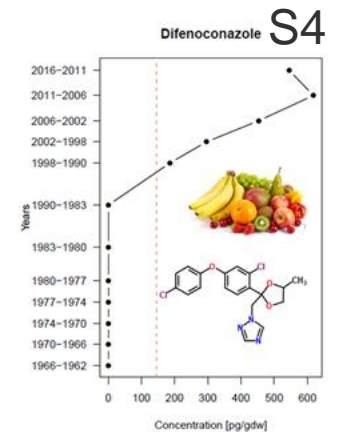
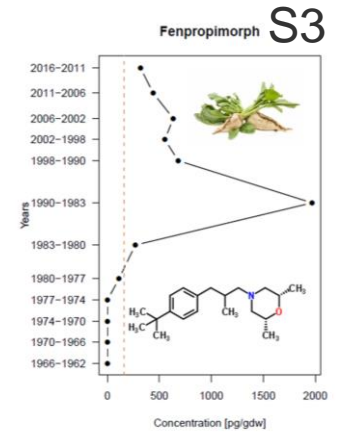
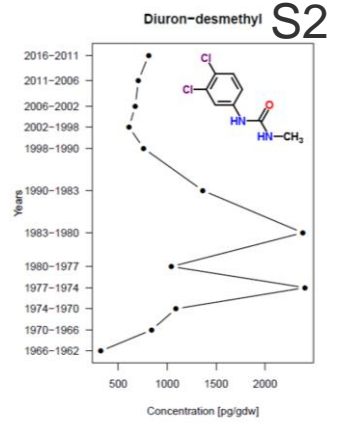
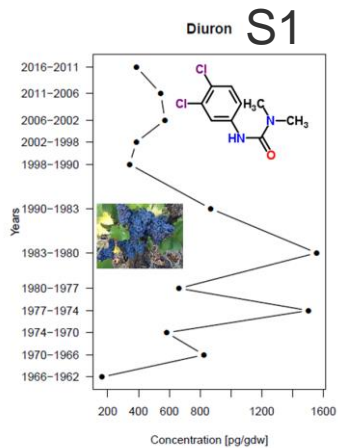
Four main steps:

I) Normalization (over the sum)

	S1	S2	S3	S4
S1	0	0.2	1.6	1.7	
S2	0.2	0	1.4	1.8	
S3	1.6	1.4	0	1.5	
S4	1.7	1.8	1.5	0	

Method)

- [



$$d(\mathbf{p}, \mathbf{q}) = d(\mathbf{q}, \mathbf{p}) = \sqrt{(q_1 - p_1)^2 + (q_2 - p_2)^2 + \dots + (q_n - p_n)^2}$$

$$= \sqrt{\sum_{i=1}^n (q_i - p_i)^2}$$

Challenges in Nontarget Screening - Unknown Compounds

- Different tools and approaches have been developed in recent years for structure elucidation
 - *In silico* fragmentation and including information on exposure, chromatographic retention, toxicity prediction
- **Pre-selection of relevant features**
 - Masses with increased identification probabilities
 - Highest intensities
 - Specific mass defects
 - Based on isotopic patterns (e.g. Cl, Br)

