

# **Non-target screening with high resolution mass spectrometry: Ready for real-world applications?**

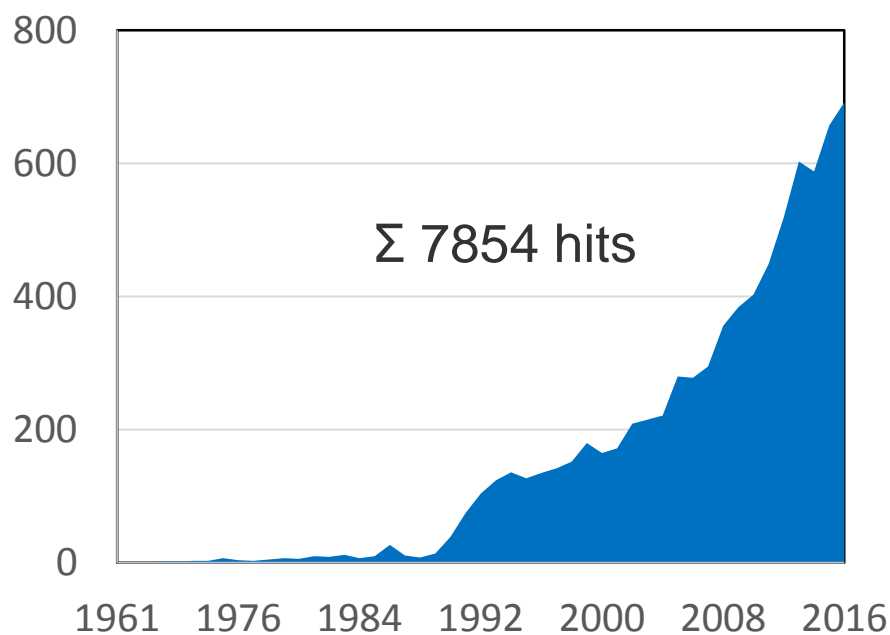
Juliane Hollender, Jennifer Schollée, Aurea Chiaia-Hernandez,  
Emma Schymanski, Heinz Singer

Eawag, Swiss Federal Institute of Aquatic Science and Technology  
Institute of Biogeochemistry and Pollutant Dynamics, ETH Zürich

# Non-target is becoming popular in science

Search for publications in web of science (May 23, 2017)

Nontarget or non-target\* & mass spectrom\*



A new approach to data evaluation in the non-target screening of organic trace substances in water analysis

Alexander M. J. K. <sup>1,3</sup>, Wolfgang Schulz <sup>2,3</sup>, Wolfgang K. Ruck <sup>3</sup>, Walter H. Weber <sup>1</sup>

<sup>1</sup> Swiss Federal Institute for Aquatic Science and Technology (Eawag), CH-8600 Dübendorf, Switzerland  
<sup>2</sup> Institute for Environmental Chemistry, University of Applied Sciences, CH-8400 Winterthur, Switzerland  
<sup>3</sup> Institute for Environmental Chemistry, University of Applied Sciences, CH-8400 Winterthur, Switzerland

# Chance to explore the iceberg of chemicals?



⇒ Confirmation with reference standards

⇒ Expected compounds which exact mass can be screened

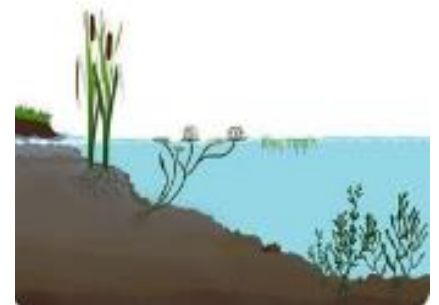
⇒ All remaining components with no prior information

## General workflow

## Three application examples

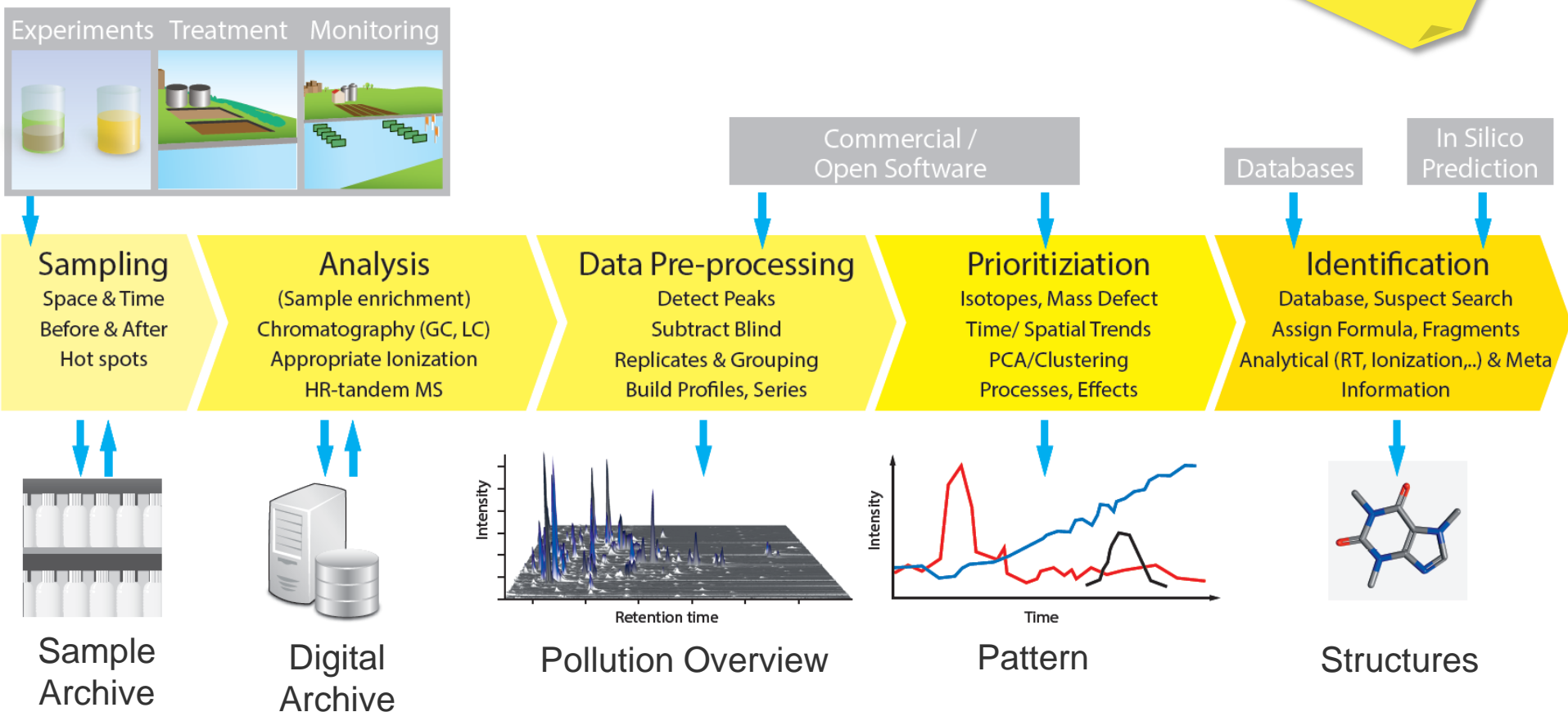
- Daily monitoring of river Rhine
- Evaluation of wastewater treatment technologies
- Assessment of (political) mitigation measures using sediment archives

## Conclusions

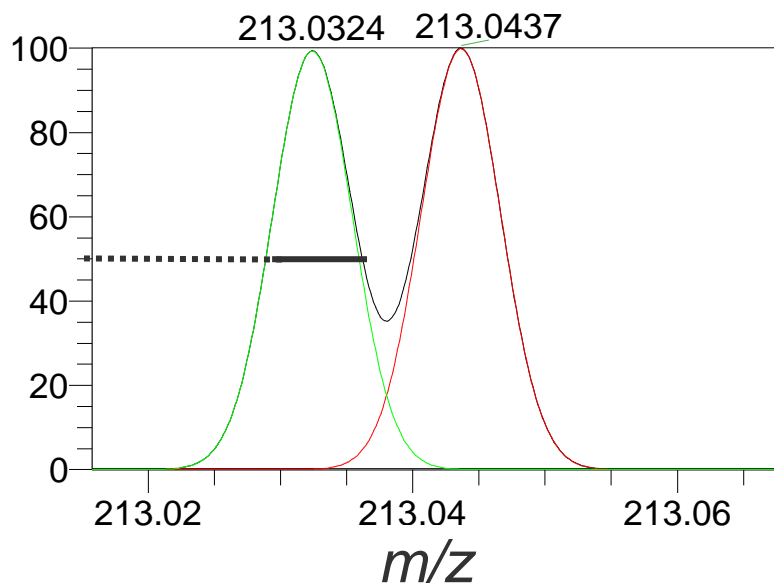
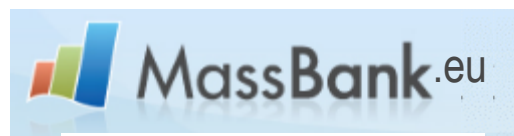


# Non-target screening workflow

Comprehensive  
target & suspect  
screening



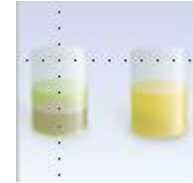
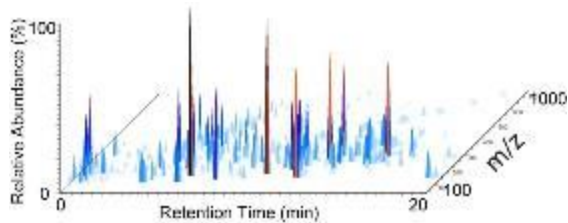
- high mass accuracy (< 5 ppm)
- high mass resolution (0.001 Da)
- high sensitivity in fullscan mode
- high stability over time
- Spectra libraries
- Compound databases
- Computational tools



**Resolution**  $R = \frac{m}{\Delta m}$

$m$  = nominal mass

$\Delta m$  = mass difference full width  
at half maximum (FWHM)



## Data-driven

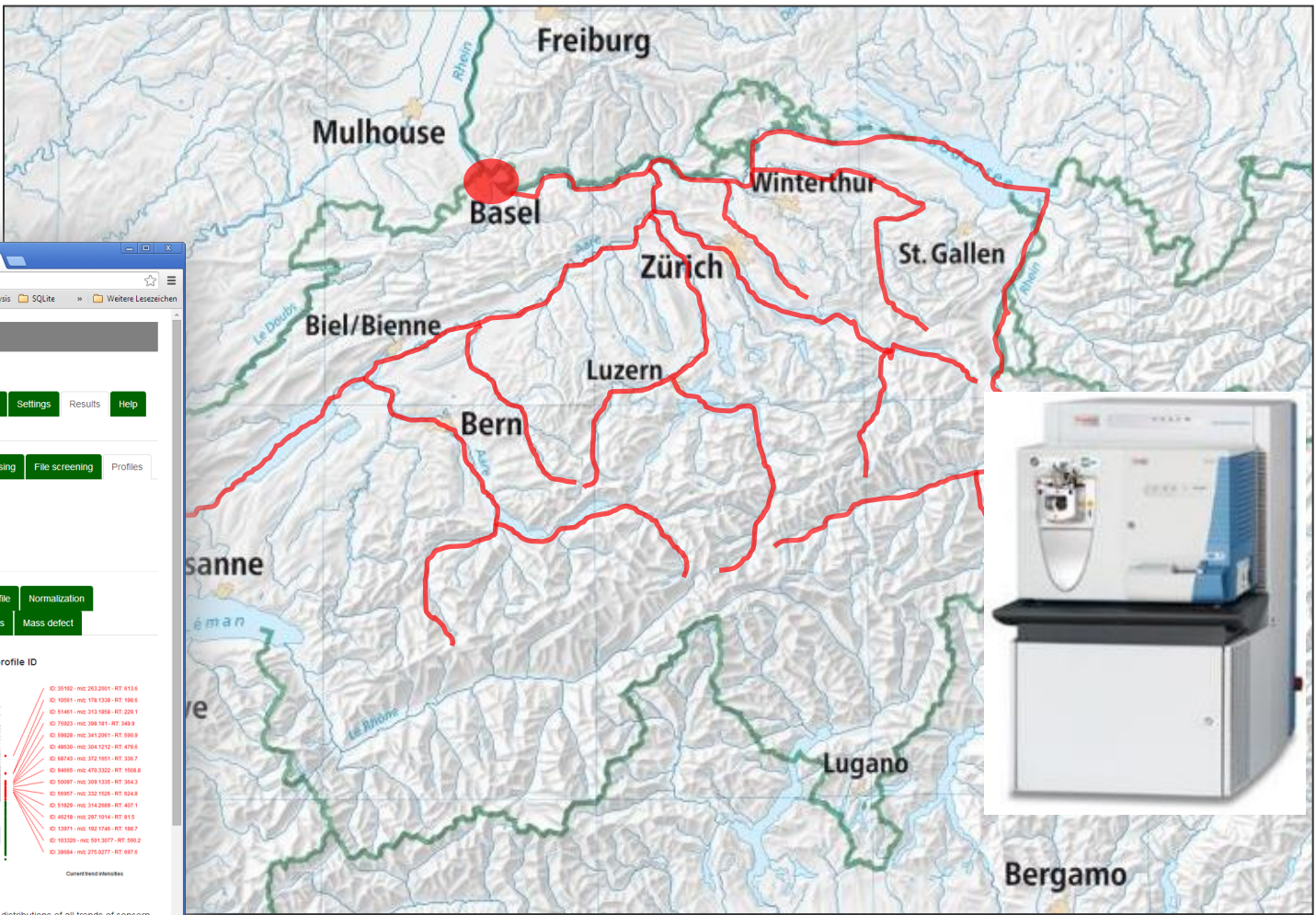
- Frequency, abundance of masses
- Component with **characteristic isotope** pattern (C, Cl, Br, N, O, S)
- Part of **homologue series** (mass difference, Kendrick mass defect)
- **Suspect** screening
- Specific functional groups (**MS/MS**, neutral loss)
- **Temporal** or **spatial profile** over several samples

## Experiment-driven

- **Persistence**, **elimination/formation** over process
- Reaction-based search of **TPs** to link masses **before and after treatment**
- Biological, electrochemical, oxidative transformation products (TPs) formation
- Reaction with **isotopically-labelled** reagents
- **Effect-directed selection** of masses in toxic fractions

# Daily screening at the International Rhine monitoring station – River Rhine

## enviMass



enviMass v2.0

Files | Compounds | Workflow options | Settings | Results | Help

About

Quality control | EIC & Peaks | Processing | File screening | Profiles

Show profile results for ionization mode: positive

Summary | Newest trends | Single Profile | Normalization

Components | Screening | Homologues | Mass defect

Comparison of newest vs. all trends by profile ID

Global trend identifiers	Current trend identifiers
ID: 288450 - m/z: 803.543 - RT: 1270	ID: 35192 - m/z: 263.2061 - RT: 813.6
ID: 32378 - m/z: 218.2762 - RT: 315.9	ID: 10961 - m/z: 178.1328 - RT: 186.5
ID: 84805 - m/z: 160.8781 - RT: 281.9	ID: 51681 - m/z: 215.1666 - RT: 228.1
ID: 39305 - m/z: 279.0658 - RT: 822.2	ID: 75823 - m/z: 358.181 - RT: 349.9
ID: 288702 - m/z: 804.0482 - RT: 1269.5	ID: 59828 - m/z: 341.2061 - RT: 590.9
ID: 58873 - m/z: 138.3415 - RT: 1444.7	ID: 48830 - m/z: 304.1212 - RT: 478.6
ID: 80899 - m/z: 441.2074 - RT: 1390	ID: 68743 - m/z: 372.1851 - RT: 338.7
ID: 82346 - m/z: 160.8235 - RT: 1488.1	ID: 84865 - m/z: 476.3222 - RT: 1956.8
ID: 12398 - m/z: 188.2016 - RT: 1852.2	ID: 50987 - m/z: 308.1325 - RT: 344.3
ID: 18885 - m/z: 180.1517 - RT: 129.5	ID: 55957 - m/z: 332.1528 - RT: 824.4
ID: 813 - m/z: 118.0813 - RT: 73.5	ID: 51829 - m/z: 314.2688 - RT: 407.1
ID: 18011 - m/z: 200.272 - RT: 791.7	ID: 46218 - m/z: 287.1914 - RT: 81.5
ID: 28919 - m/z: 237.03 - RT: 72.8	ID: 13871 - m/z: 162.1146 - RT: 188.7
ID: 138284 - m/z: 578.261 - RT: 805.3	ID: 103300 - m/z: 541.3077 - RT: 590.2
ID: 85341 - m/z: 380.3235 - RT: 1439.4	ID: 38884 - m/z: 275.0277 - RT: 897.4

The above boxplot (grey) shows the intensity distributions of all trends of concern, listing the IDs, mean masses (m/z) and mean retention time (RT) of the profiles with





SPE followed by LC-ESI-HRMS (LTQ-Orbitrap)  
High resolved full scan spectra & MSMS fragmentation spectra



Priority substances

Site specific substances

- x calibration
- x exact quantification
- x manual data processing

- x calibration
- x semi-quant. quantification
- x automated processing

Time profiles

Statistical analysis

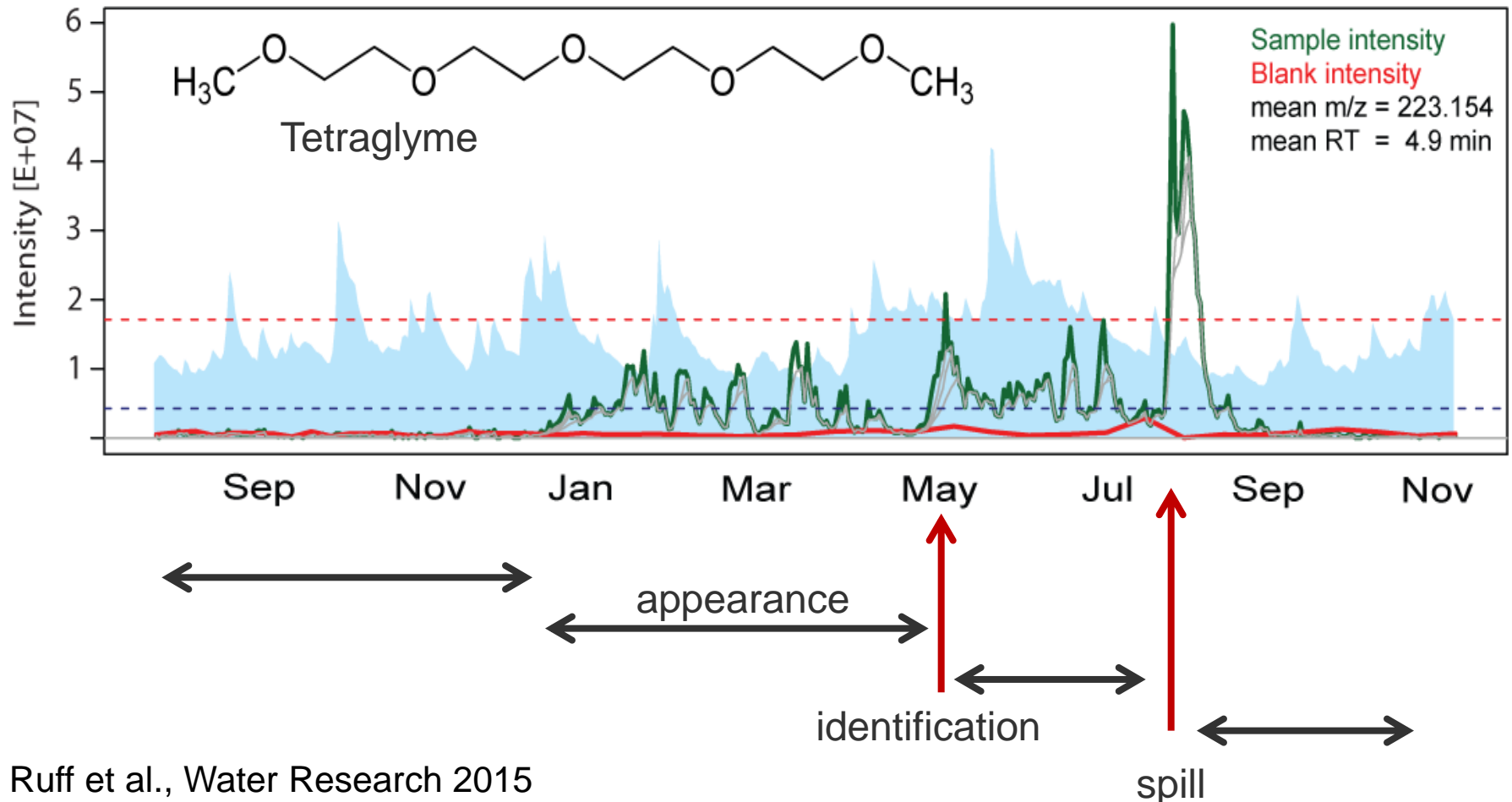
Changes / trends

Prioritization

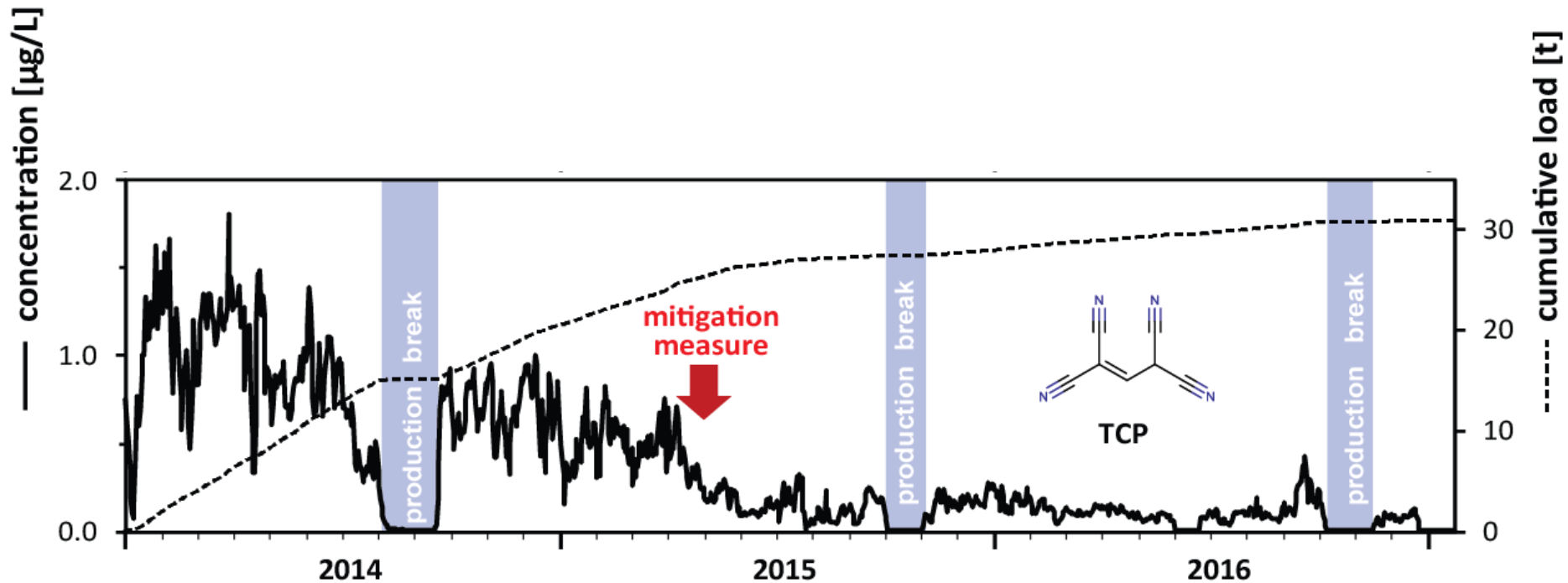
**enviMass**

# Prioritization using time profiles

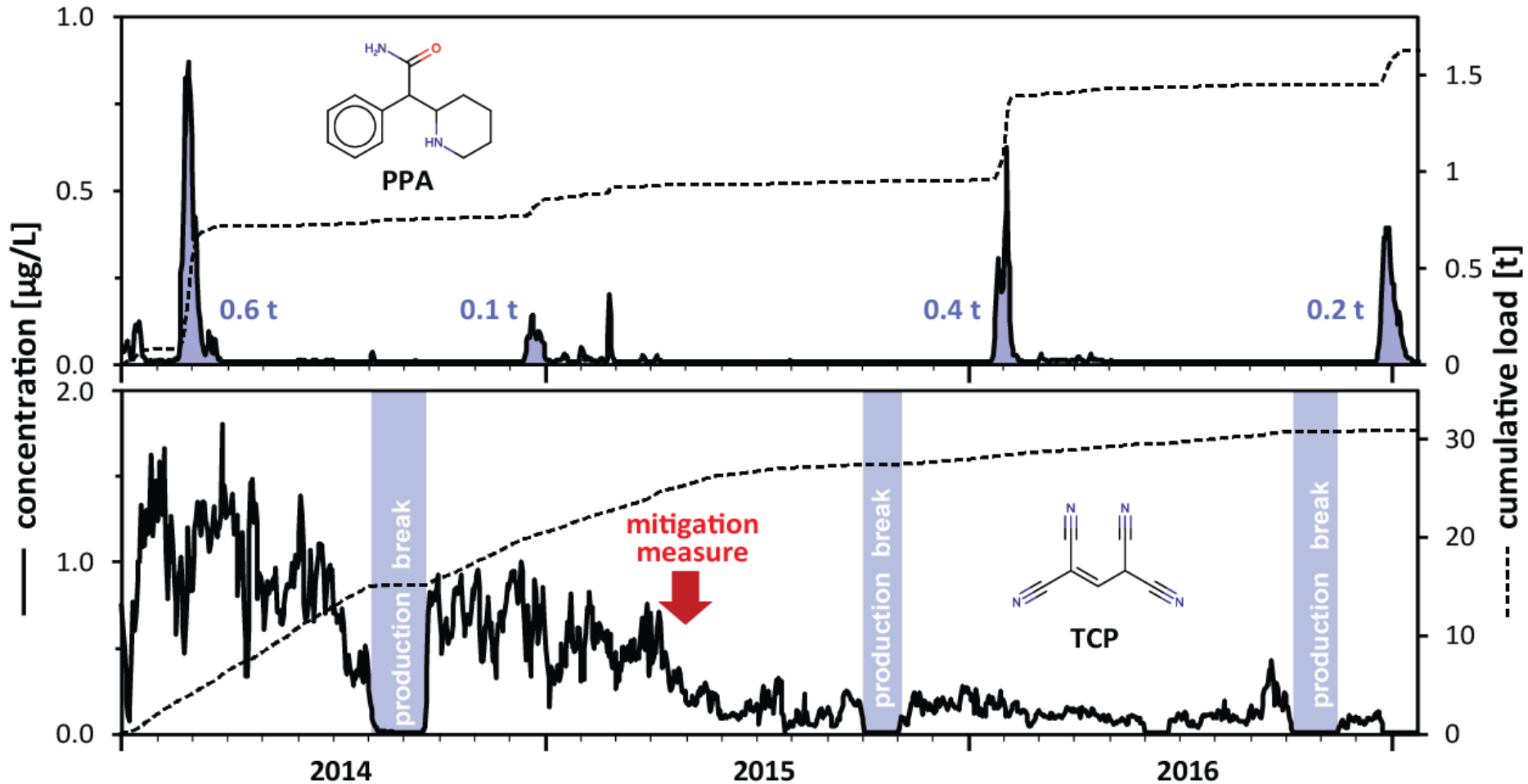
- Identification**
- Molecular formula assignment,
  - Database search
  - Prioritization of hits with information on industrial production
  - Confirmation with reference standard



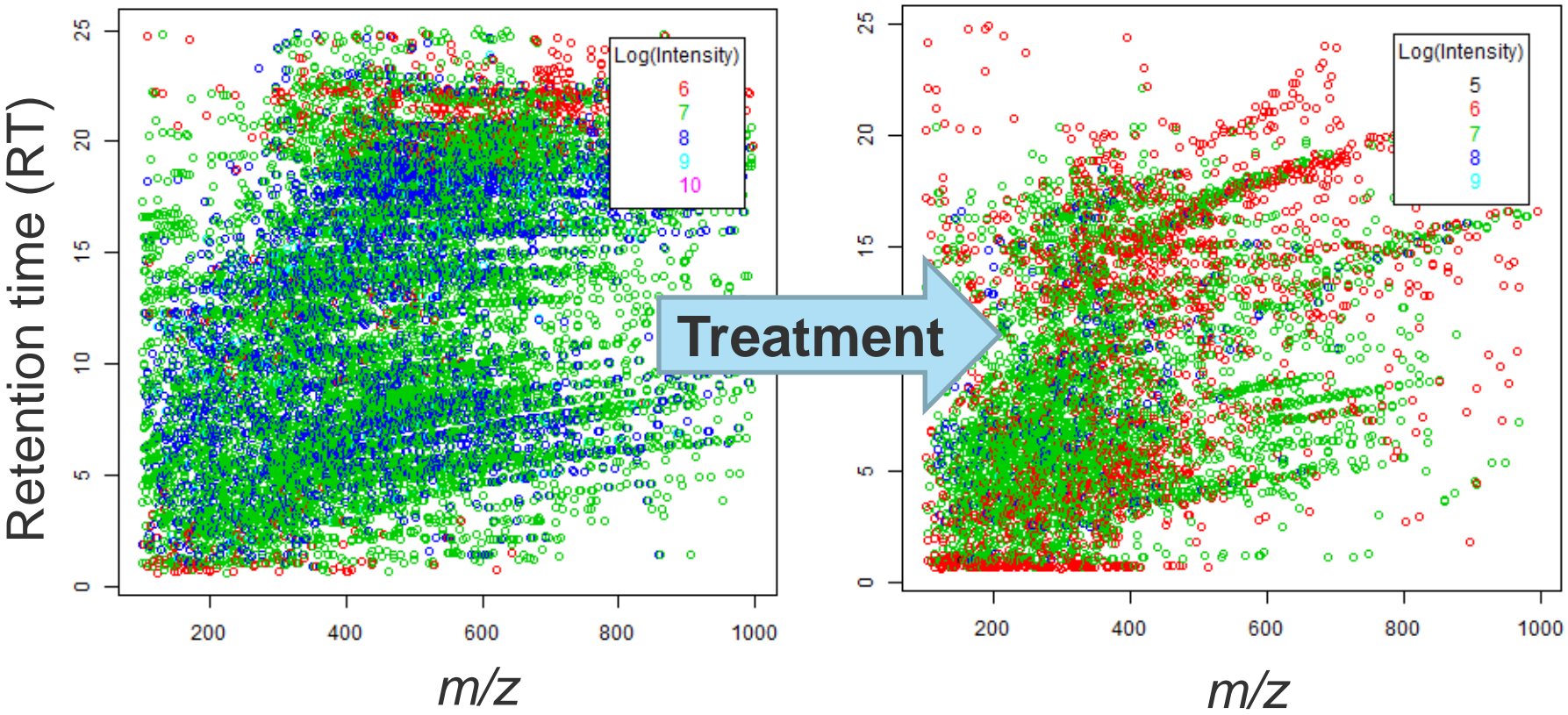
## Cumulative loads over three years



## Cumulative loads over three years

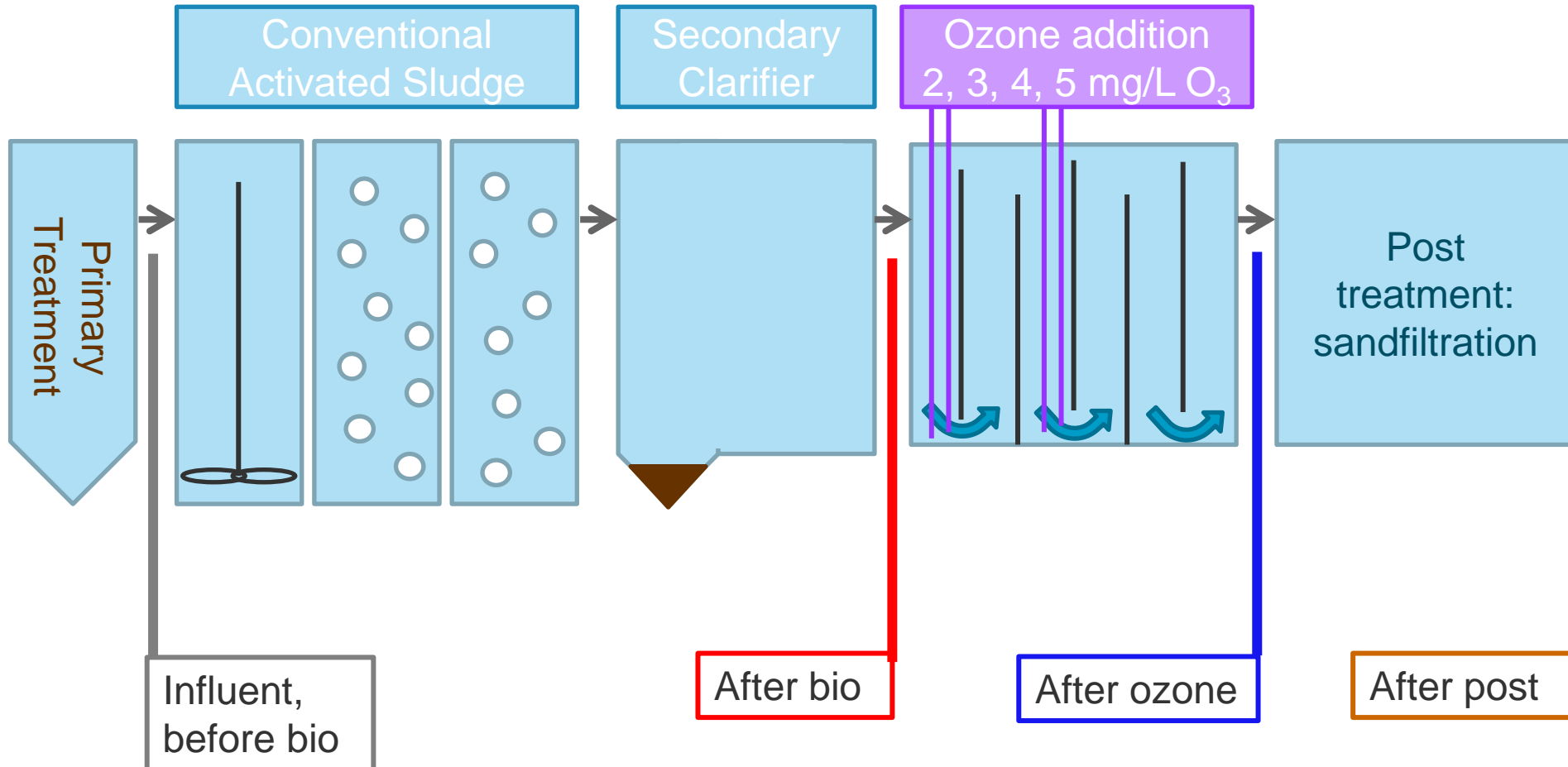


## Complex wastewater matrix with contaminants



# Wastewater treatment with ozonation

Full scale Wastewater treatment plant (WWTP) in Dübendorf, Switzerland



24-hr flow-proportional composites (3 consecutive days)

# Workflow – Suspect & non-target screening

## Data Acquisition

Online solid-phase extraction

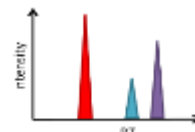


LC-Orbitrap MS/MS  
(Atlantis; +/- ESI  
R= 70,000 – 140,000)

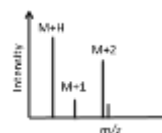


## Data Processing

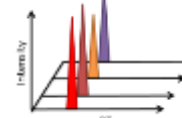
Peak detection  
enviPick package



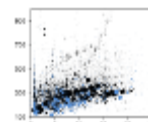
Componentization  
nontarget package



Profile building  
enviMass package

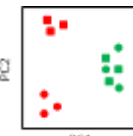


Blind subtraction  
R script



## Data Analysis

Multivariate  
Statistics – PCA, HCA



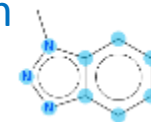
Linking Parent/TP  
R script



Suspect Screening  
R script



Structure elucidation  
MetFrag, MassBank,  
STOFF-IDENT

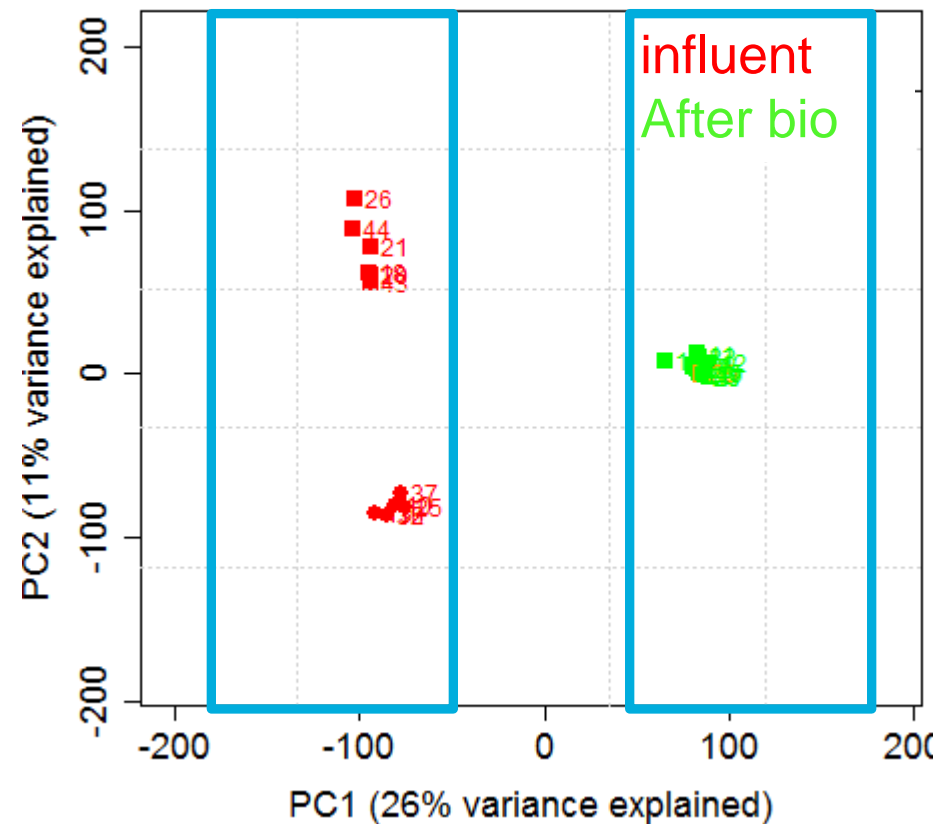


Workflow modified from: Schollée et al. 2015, Anal. Chem

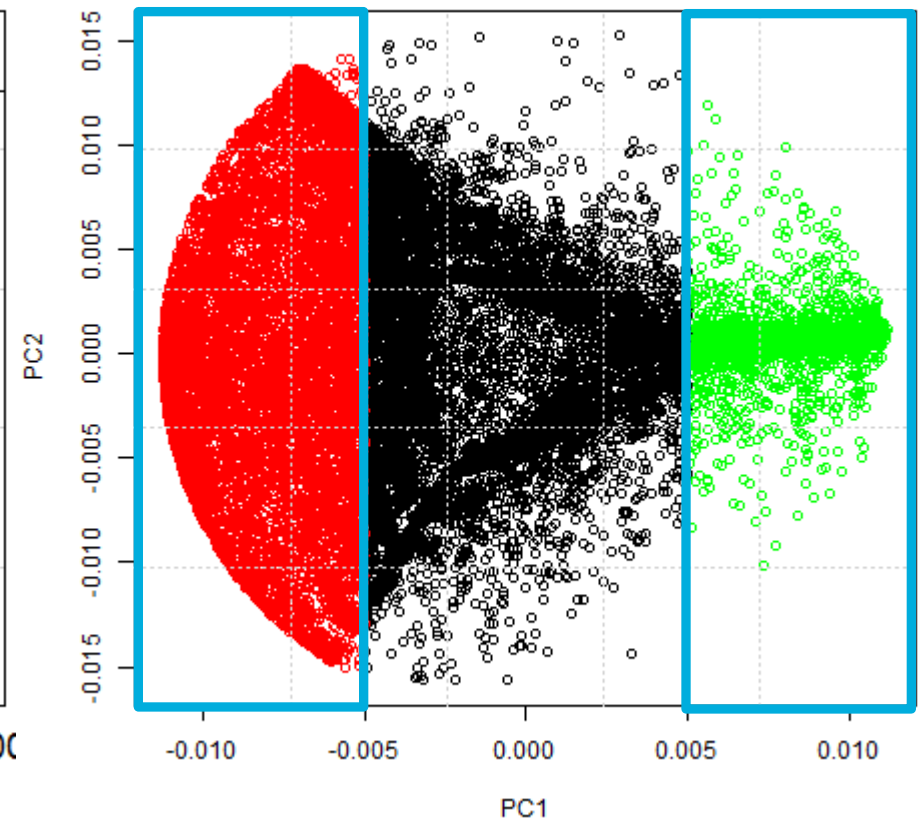
Software available at: <http://www.eawag.ch/en/departement/uchem/software/>

## Principal component analysis of Non-target Peaks

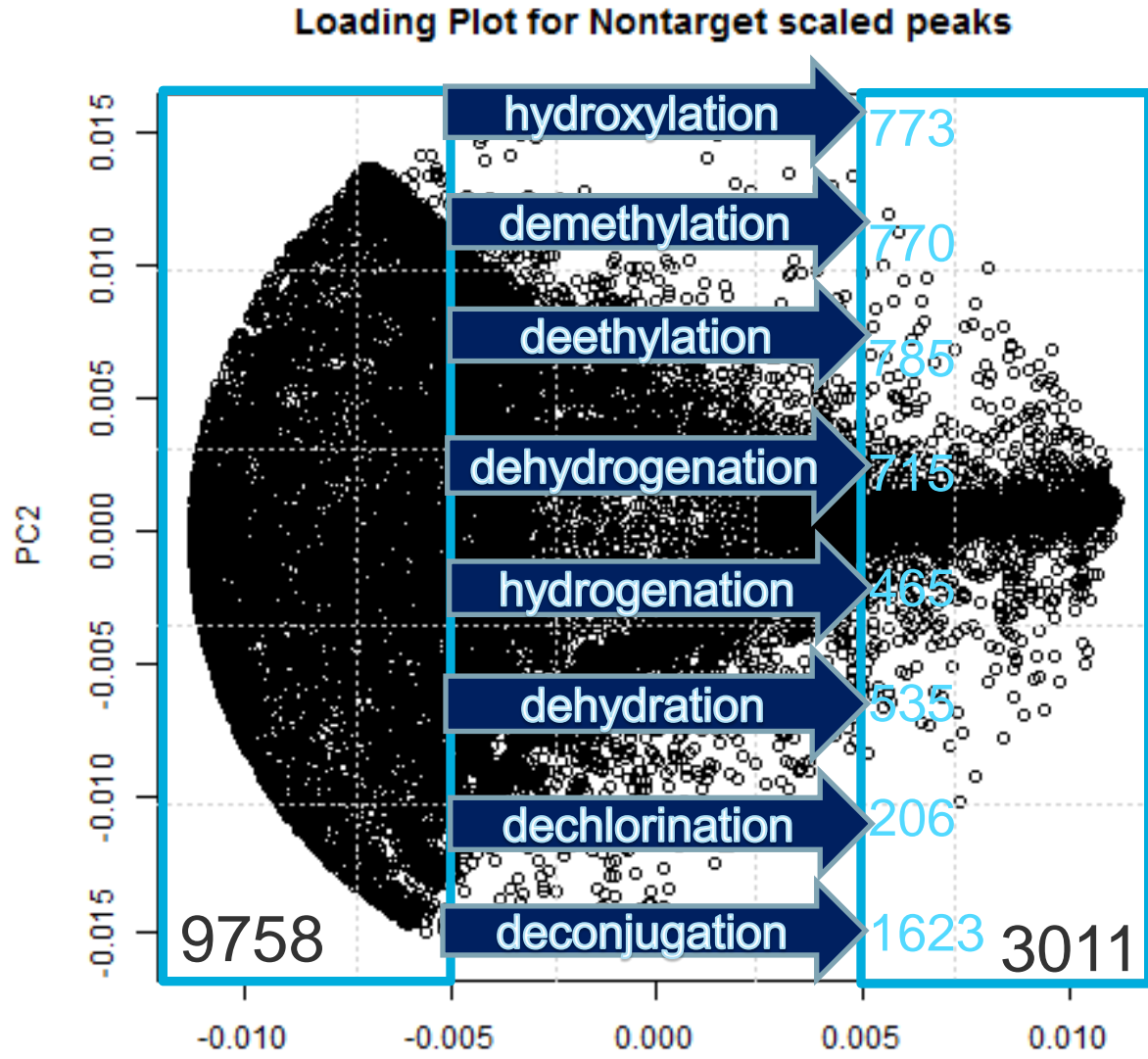
Scores Plot for Nontarget scaled peaks



Loading Plot for Nontarget scaled peaks



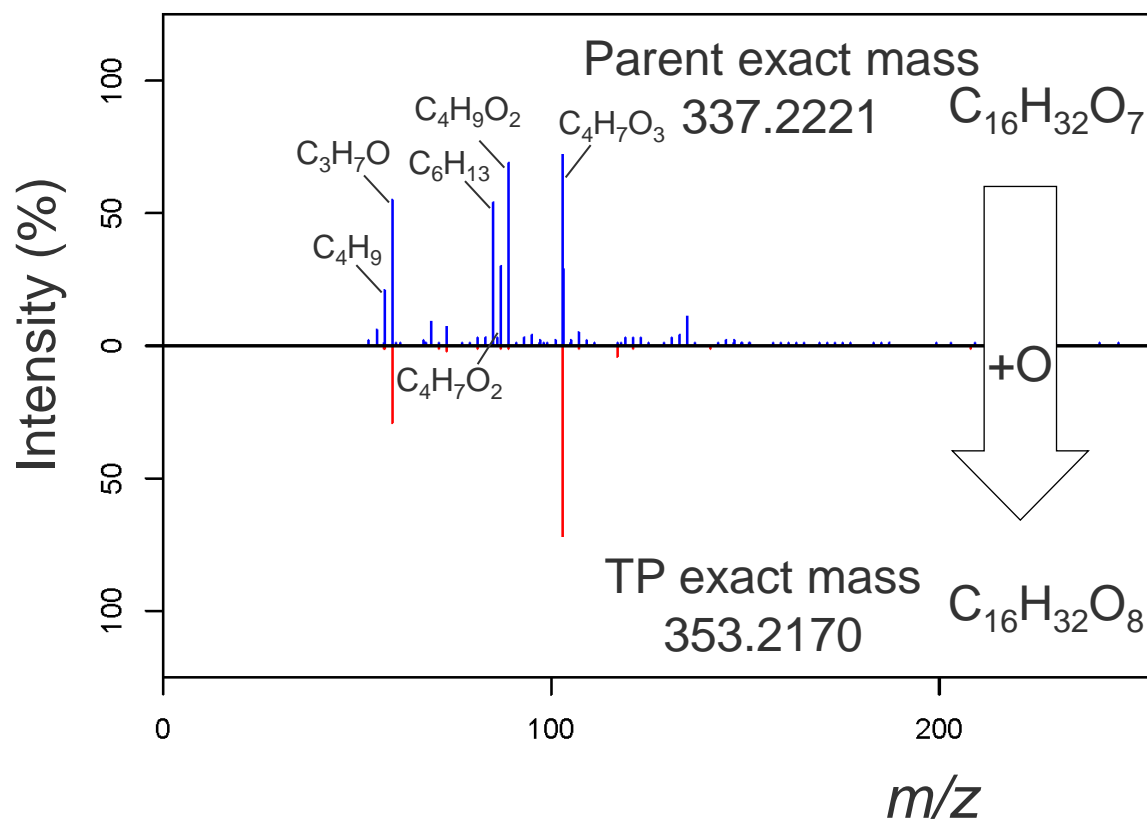




5872 linkages

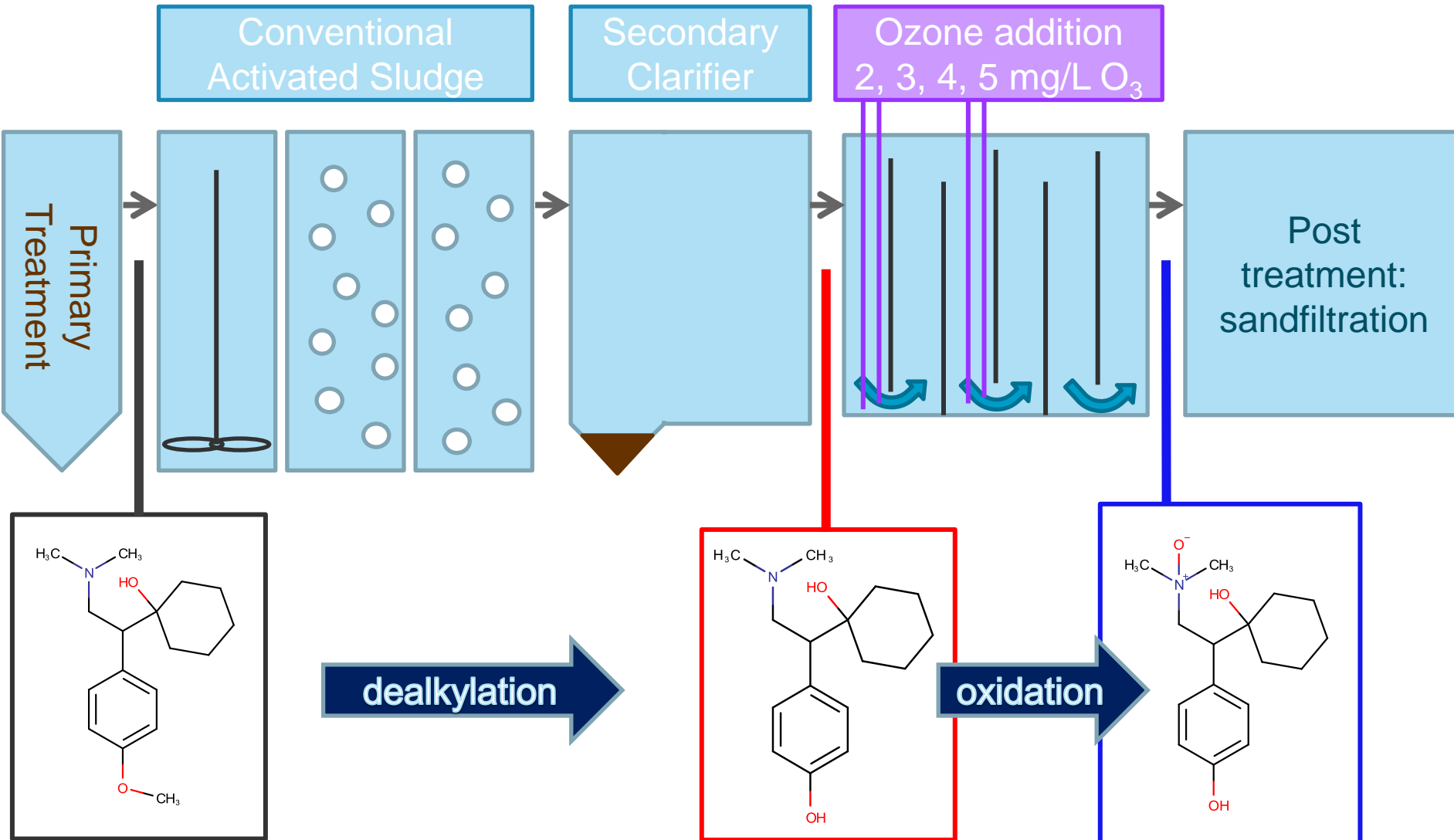
# Structure elucidation

## Example of a linkage related to a hydroxylation



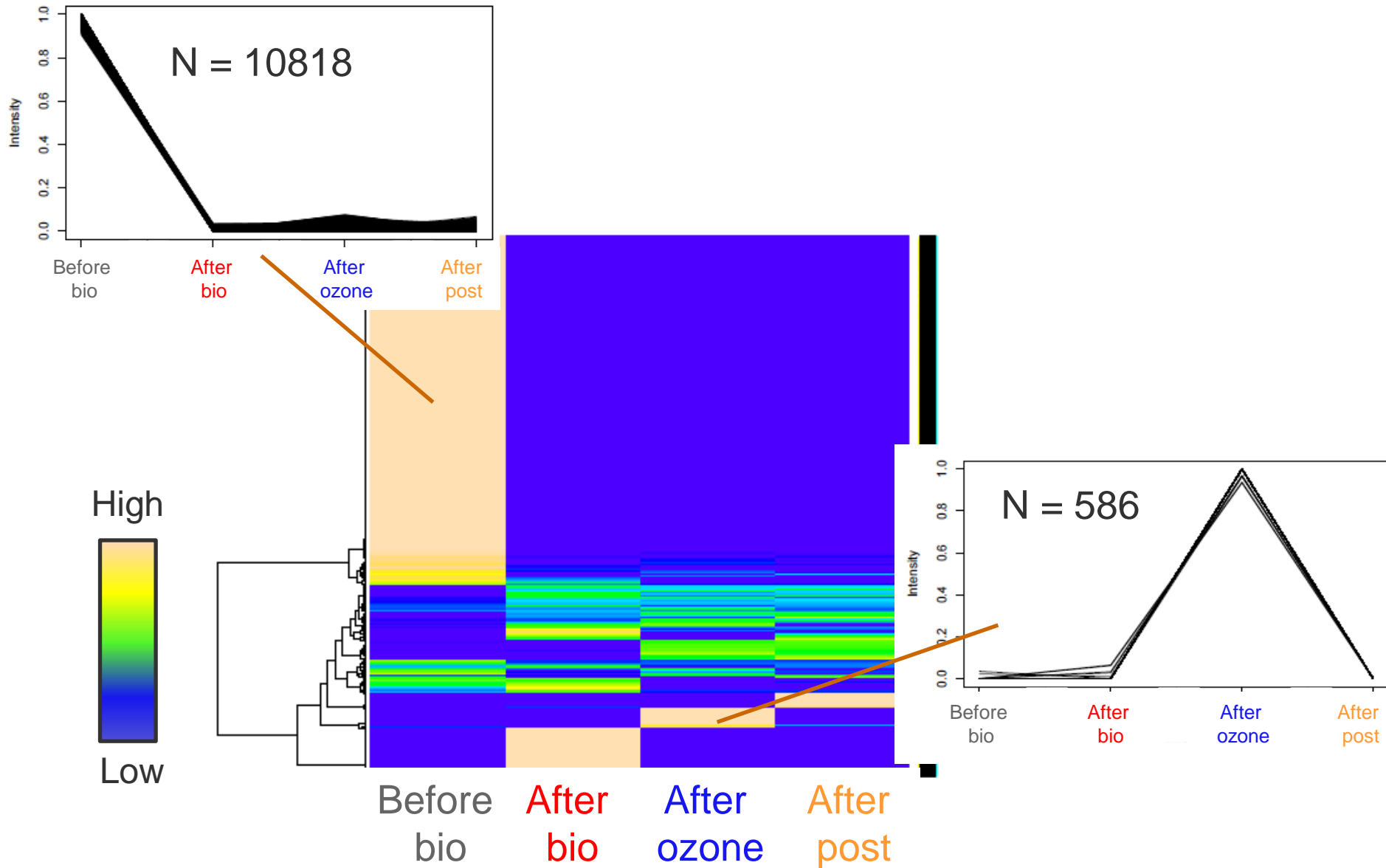
MS/MS Similarity score: 0.73

# Statistical tools and suspect screening

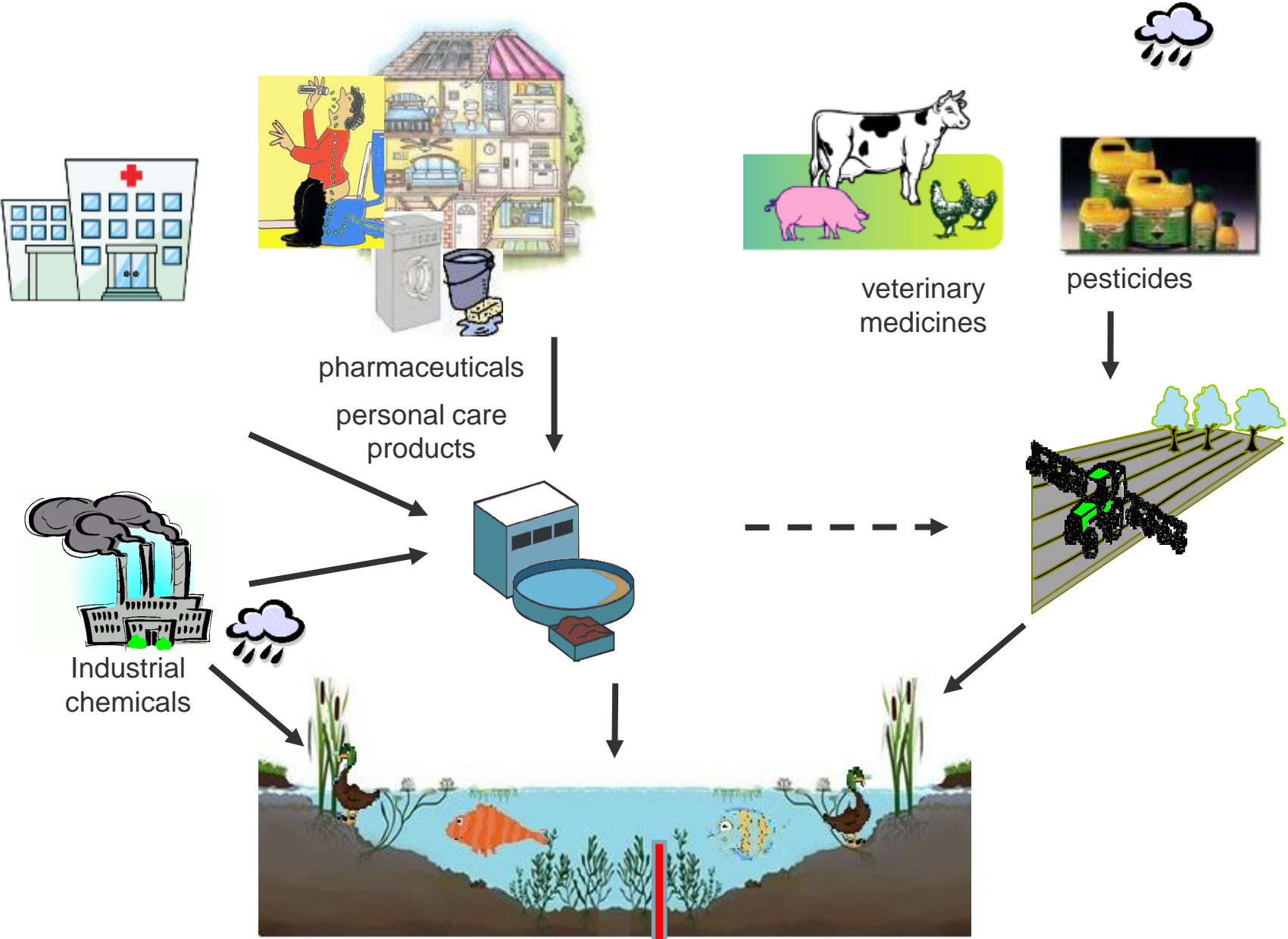


Venlafaxine degradation described in: Rúa-Gómez et al. 2012; Gulde et al. 2016

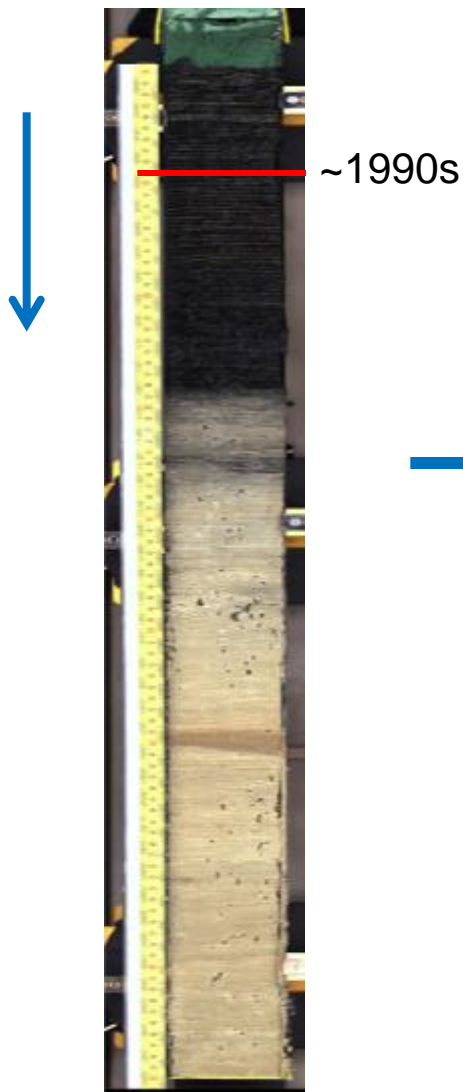
# Hierarchical clustering of profiles along treatment chain



# Lake sediments as archive of pollution



## Lake sediment cores



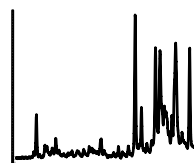
Freeze-drying & homogenization



Pressurized liquid extraction & purification using liquid-liquid extraction (Quechers)



RP chromatography (Xbridge)  
ESI +/-  
Orbitrap-HRMS/MS



- 180 Targets
- 840 Suspects
- Non-targets

# Non-target screening of m/z 450.9619 (ESI-) with negative mass defect & clear isotopic pattern



Chlorine  
adduct

Exact mass (m/z)  
**414.9846**

MS & MS/MS

**MOLGEN** : C,H,O,N,Cl,F

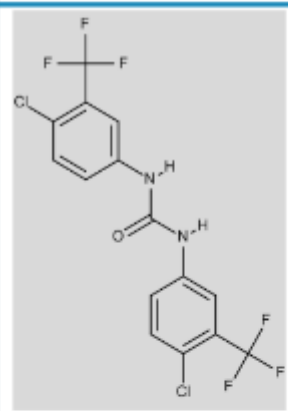
**C<sub>15</sub>H<sub>8</sub>Cl<sub>2</sub>F<sub>6</sub>N<sub>2</sub>O**

Top score of 0.96

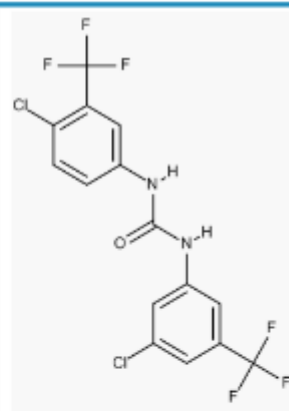
**MetFrag** scoring

Number of references

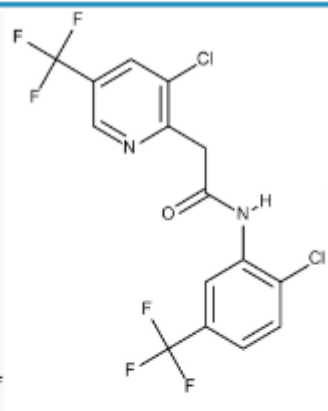
## 5 Structures



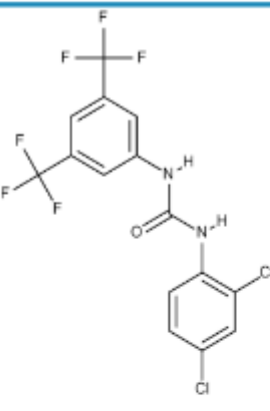
Score: 1  
References: 13



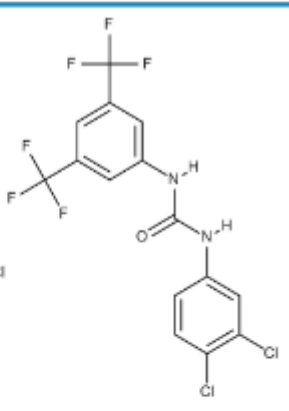
Score: 1  
References: 1



Score: 0.802  
References: 8



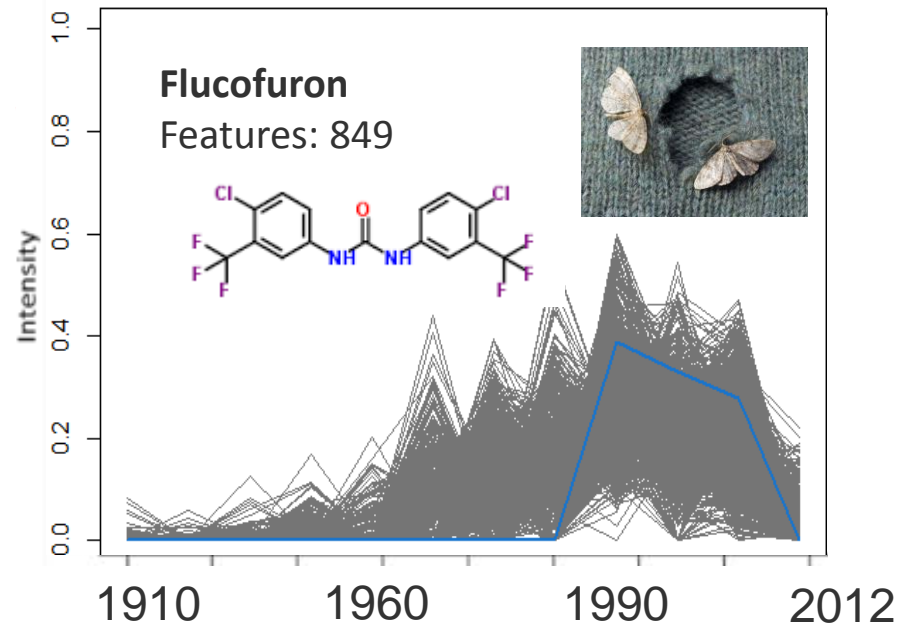
Score: 0  
References: 5



Score: 0  
References: 20

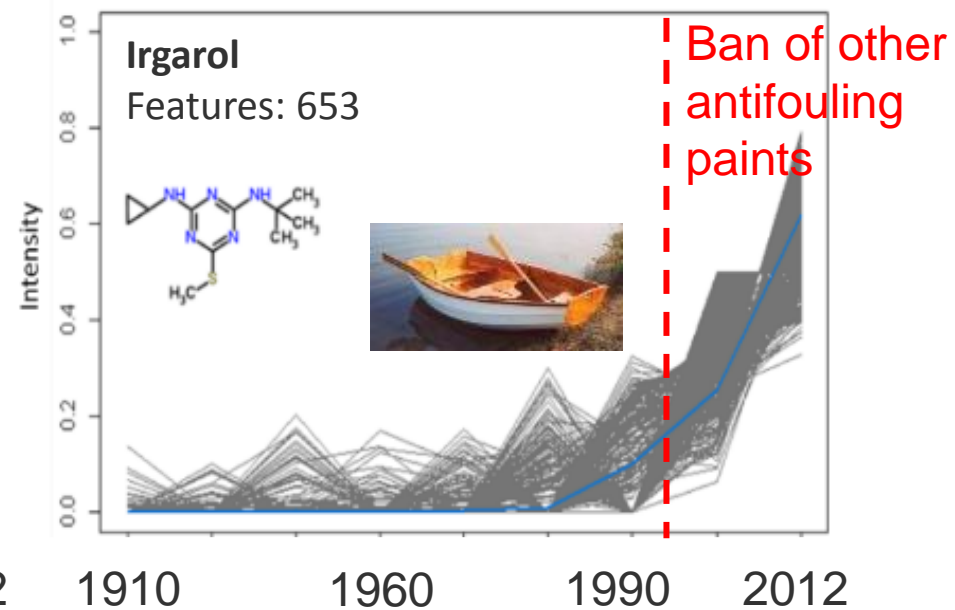
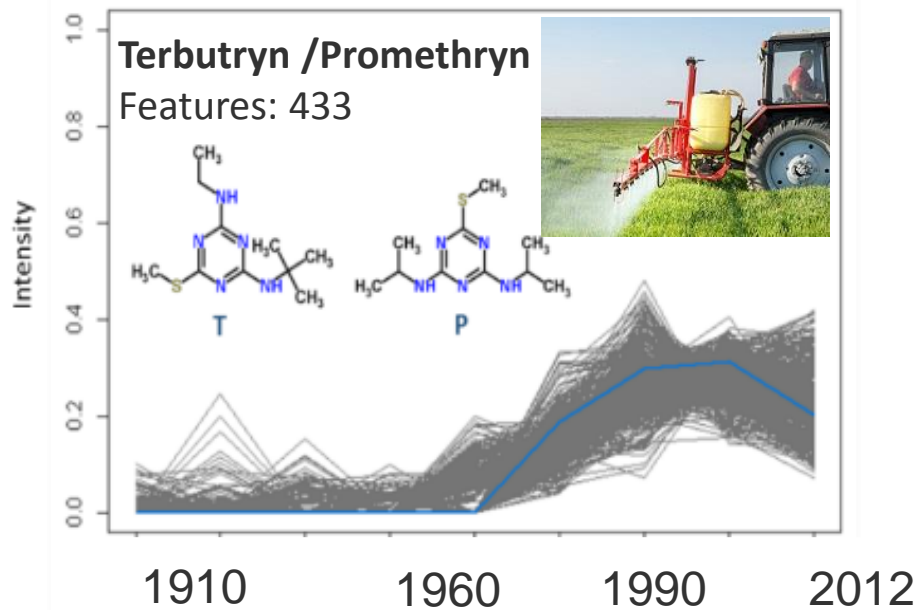
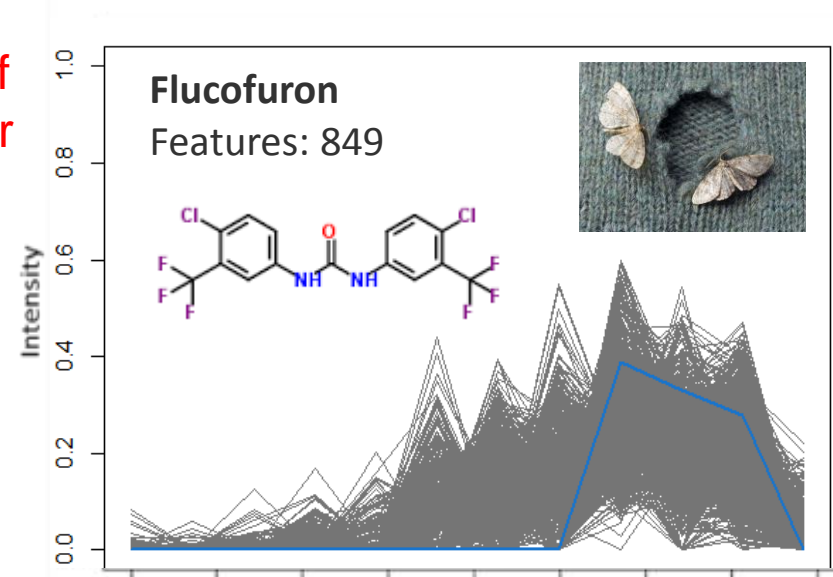
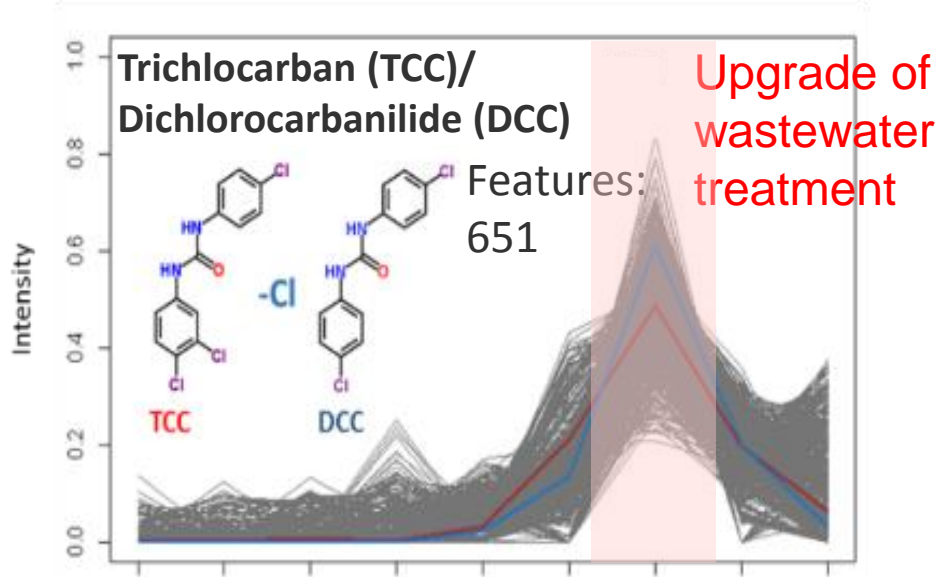
**Flucofuron**

# Characteristic trends using hierarchical clustering – Lake Lugano





# Characteristic trends using hierarchical clustering – Lake Lugano



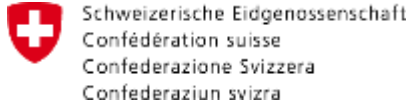
# Is non-target screening ready for real-world applications?

## Yes almost, but...

- excellent instrumentation, as well as proper data analysis tools and expert knowledge is needed
- Prioritization is mandatory to master the number of features
- Currently more characterization than identification
- Not every peak can be identified with mass spectrometry

## Future challenges

- application to other ionization techniques (e.g., APPI, APCI)
- Inclusion of other information into identification workflow (e.g., ion mobility, meta information)
- Implementation in practice labs



- Swiss Federal Office for the Environment FOEN
- Marie Curie ITN EDA-Emerge
- FP7 EU Project SOLUTIONS

- Steffen Ruppe, Reto Dolf, and Jan Mazacek, AUE Basel-City
- Matthias Ruff, Martin Loos, Barbara Günthardt, Eawag
- Martin Loos, R package developer, enviMass
- Alois Zwysig, Alfred Lück, Flavio Anselmetti, Nathalie Dubois, Eawag

Thank you for your attention



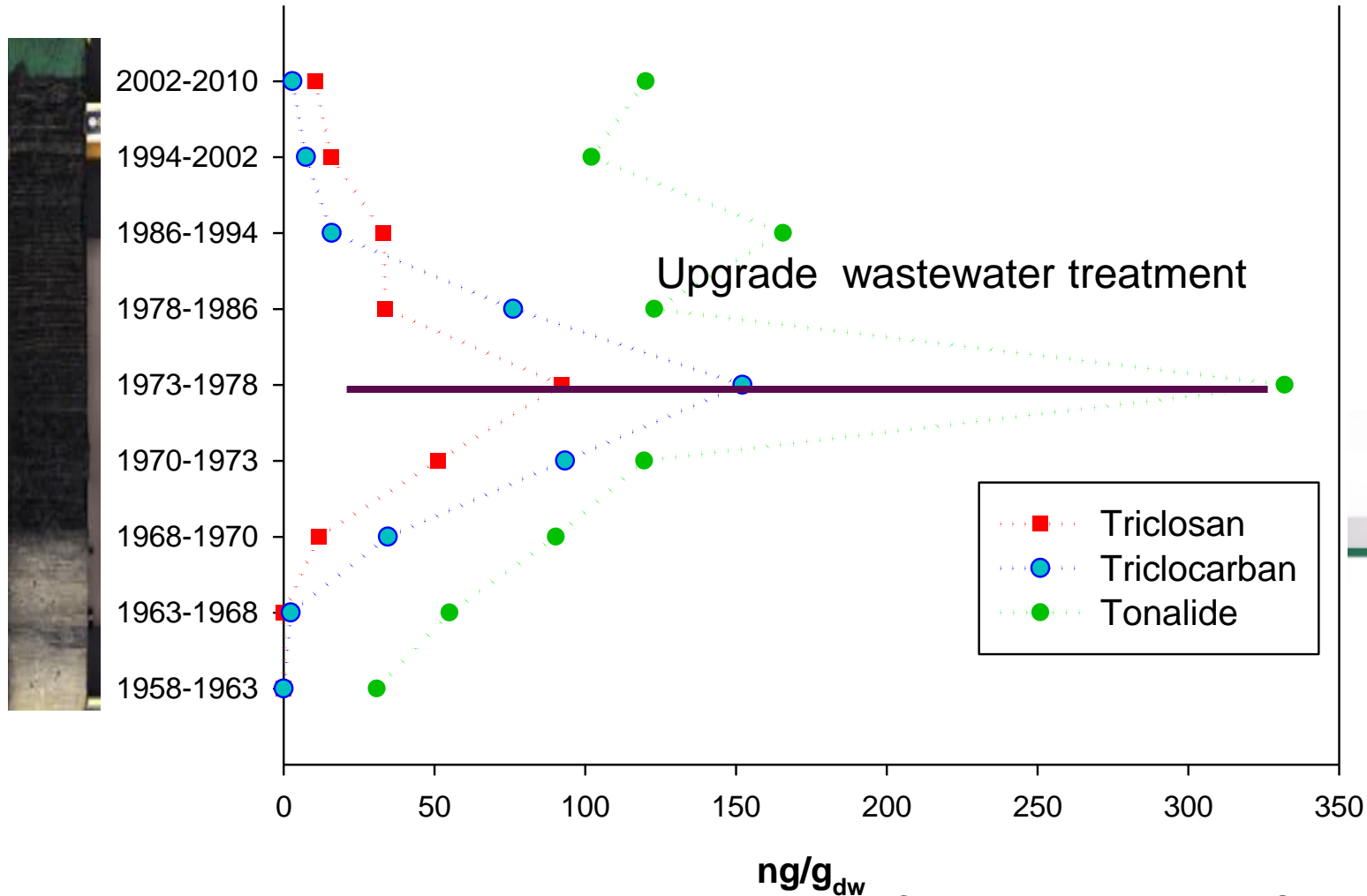
- 300 relevant compounds included in target screening
- captured load increased from 7 to 100 t per year (+ ca. 56 t complexing agents)
- Non-target analysis using enviMass has triggered several Rhine alarms

Daily trend monitoring  
and spill detection with  
enviMass since 2012

Eawag

# Temporal pollution pattern

Time series of PCPs in sediment cores from Lake Greifensee



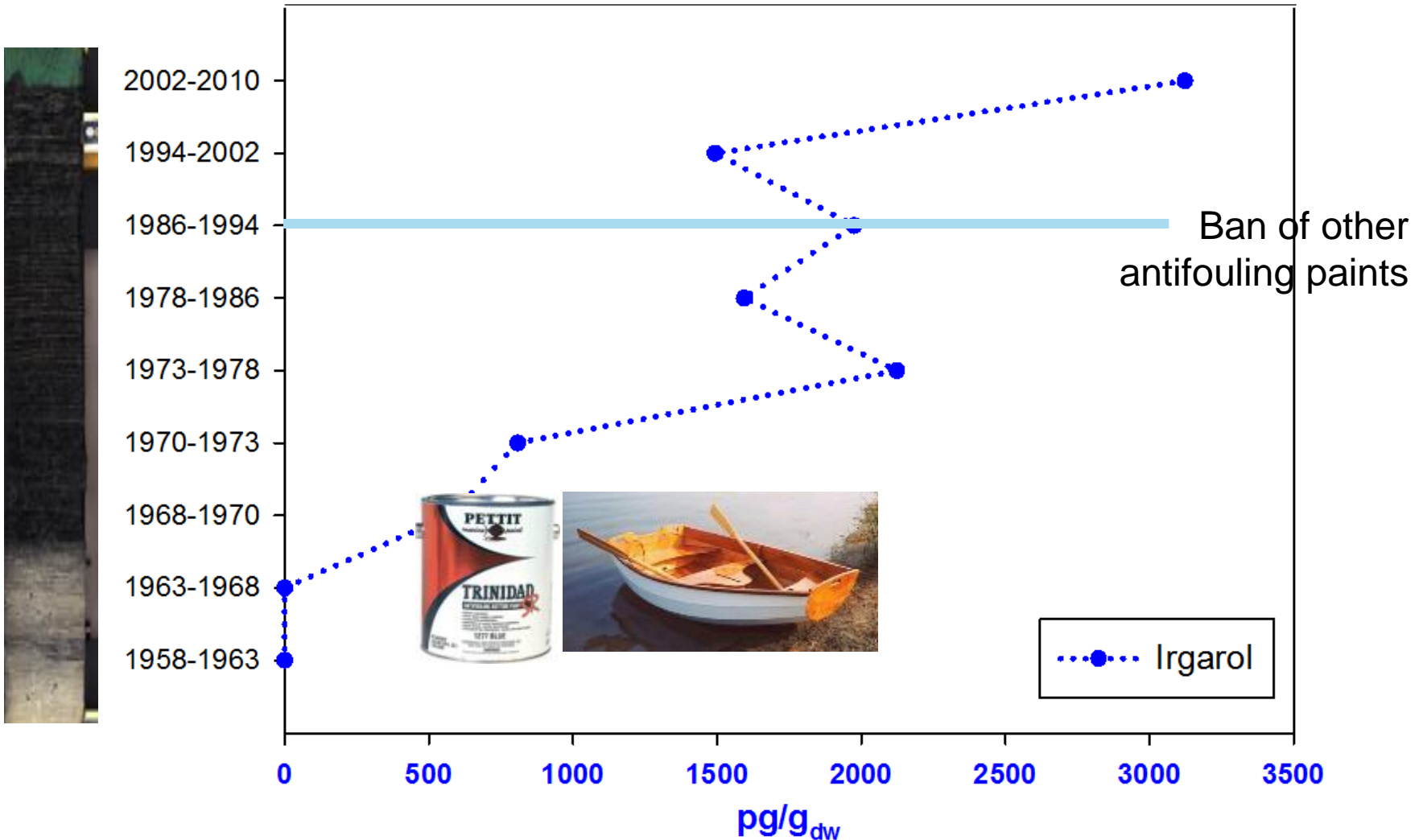
Triclosan /  
Triclocarban



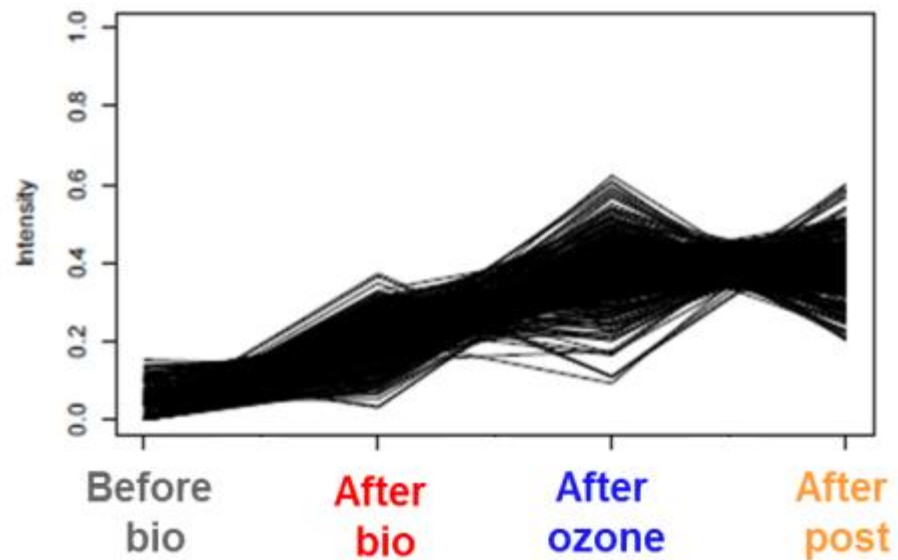
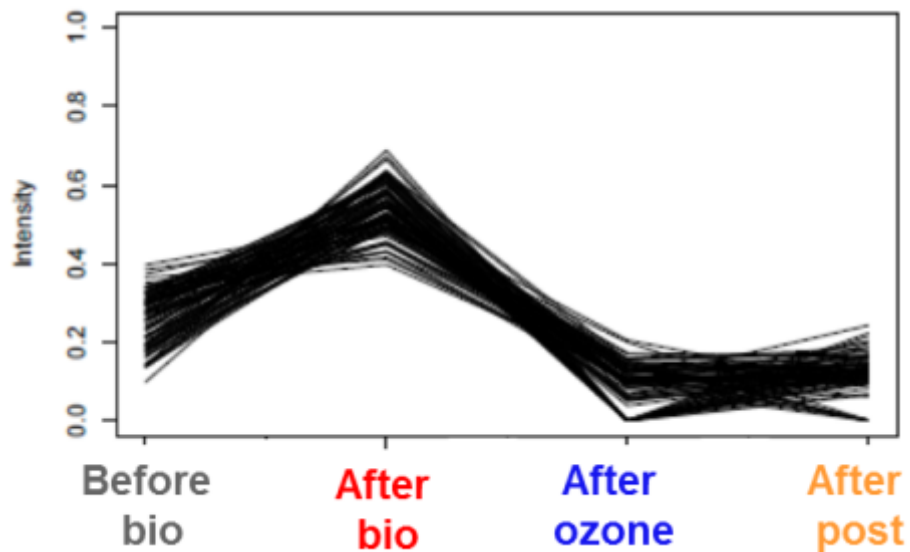
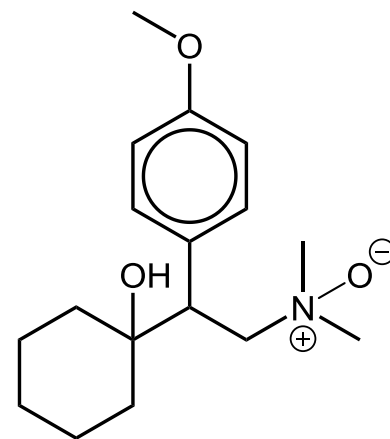
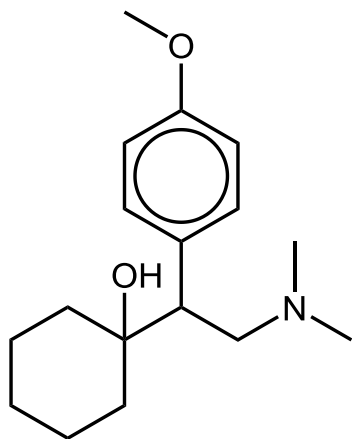
Tonalide

# Temporal pollution pattern

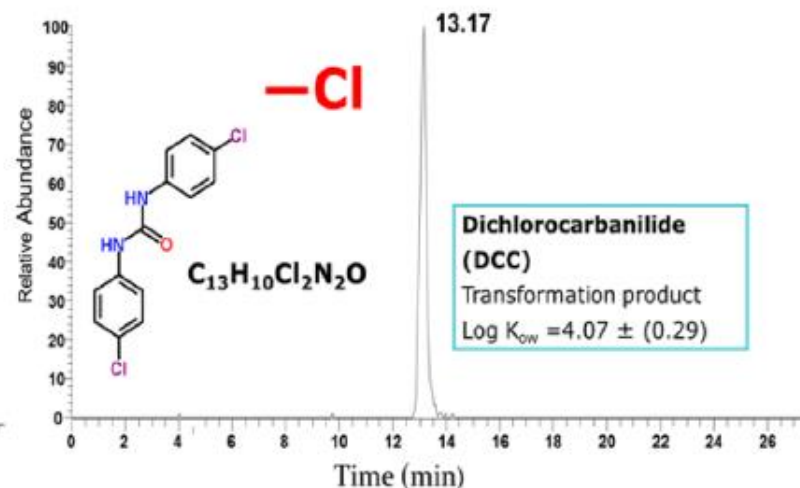
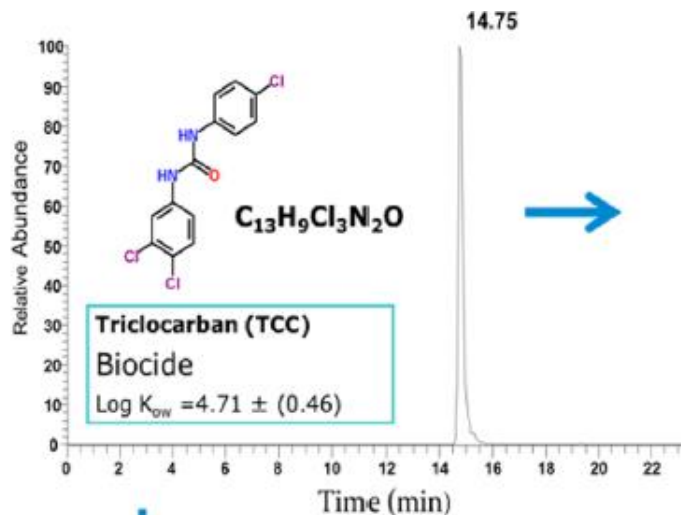
Time series of the pesticide irgarol in sediment cores from Lake Greifensee



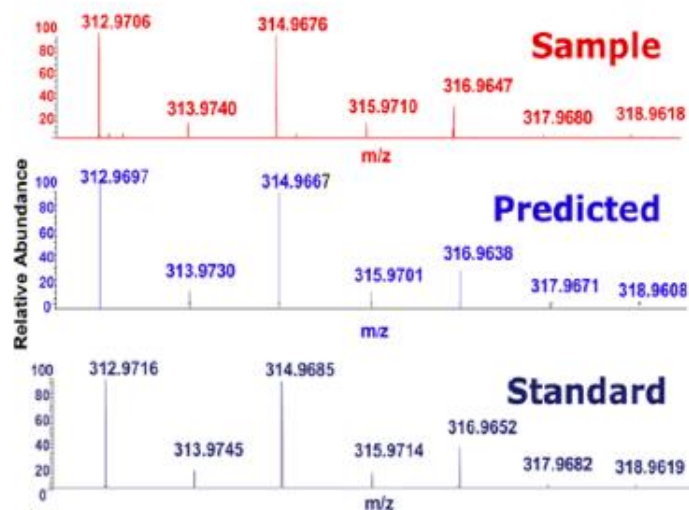
# Validation with Target Compounds



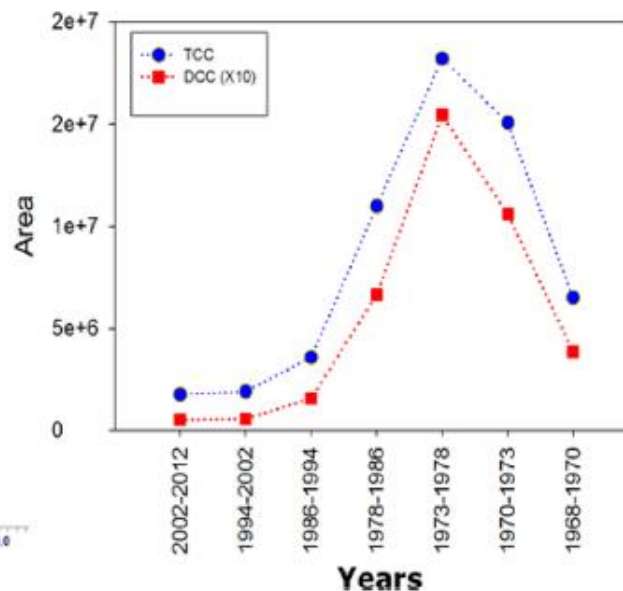
# Screening of transformation products



## Isotope pattern

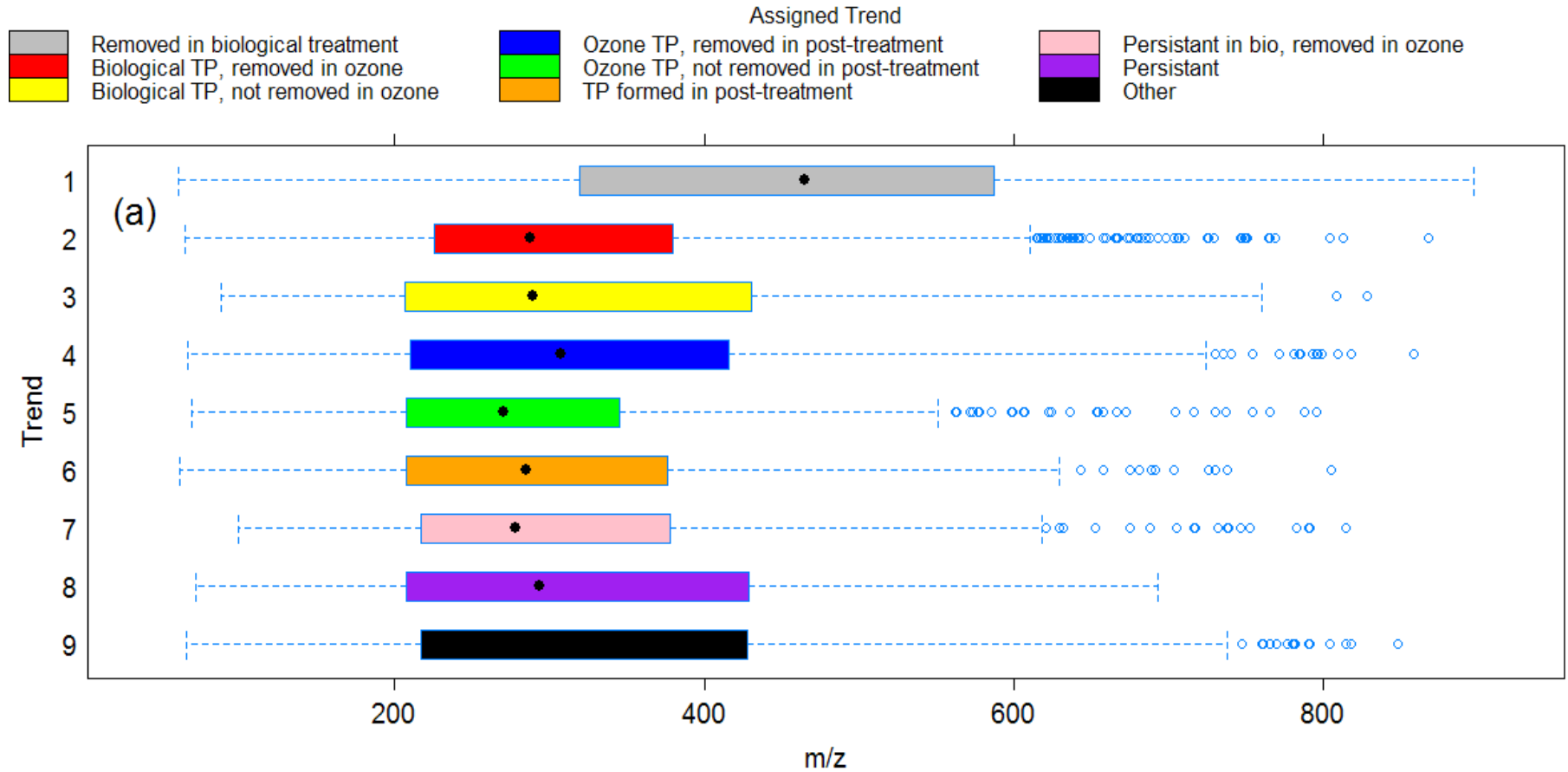


## Temporal resolution of TCC and DCC based on peak areas



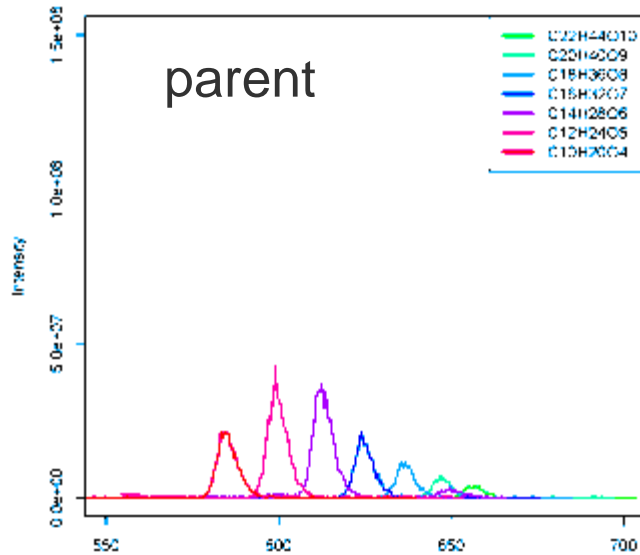


## Sand Filter Post-treatment

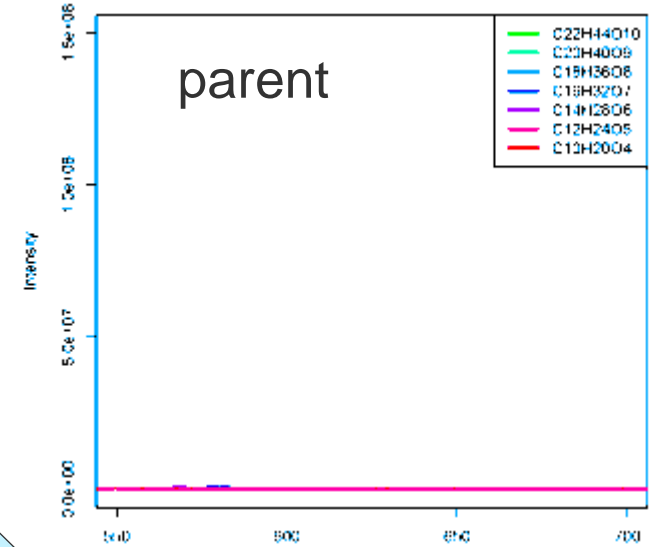


# Transformation Products of Homologue series ( $\Delta C_2H_4O$ )

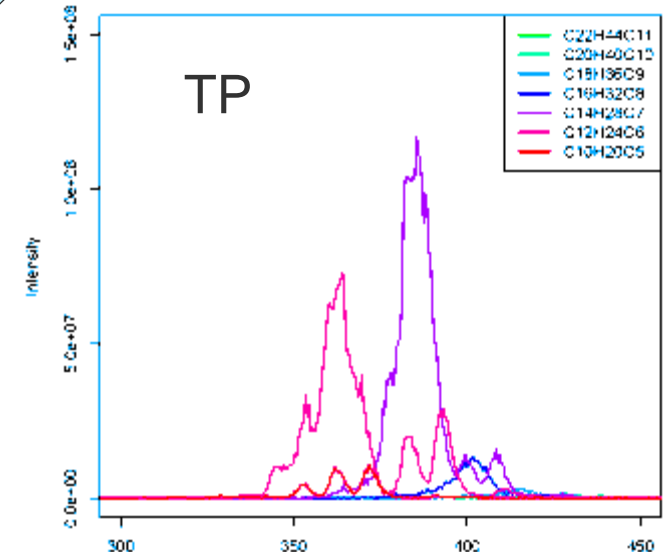
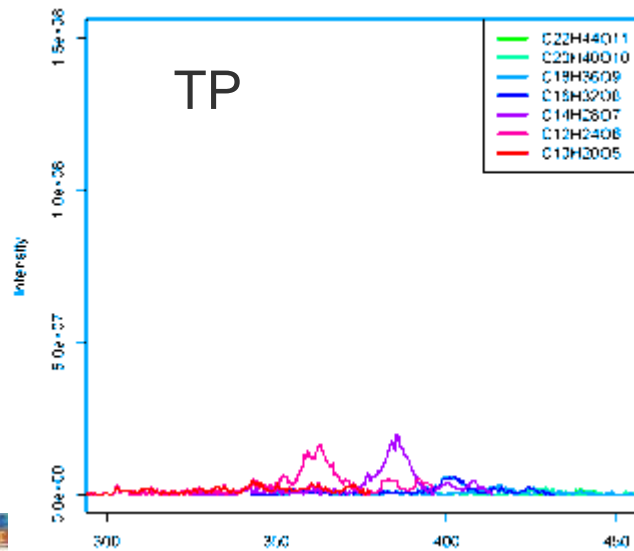
influent



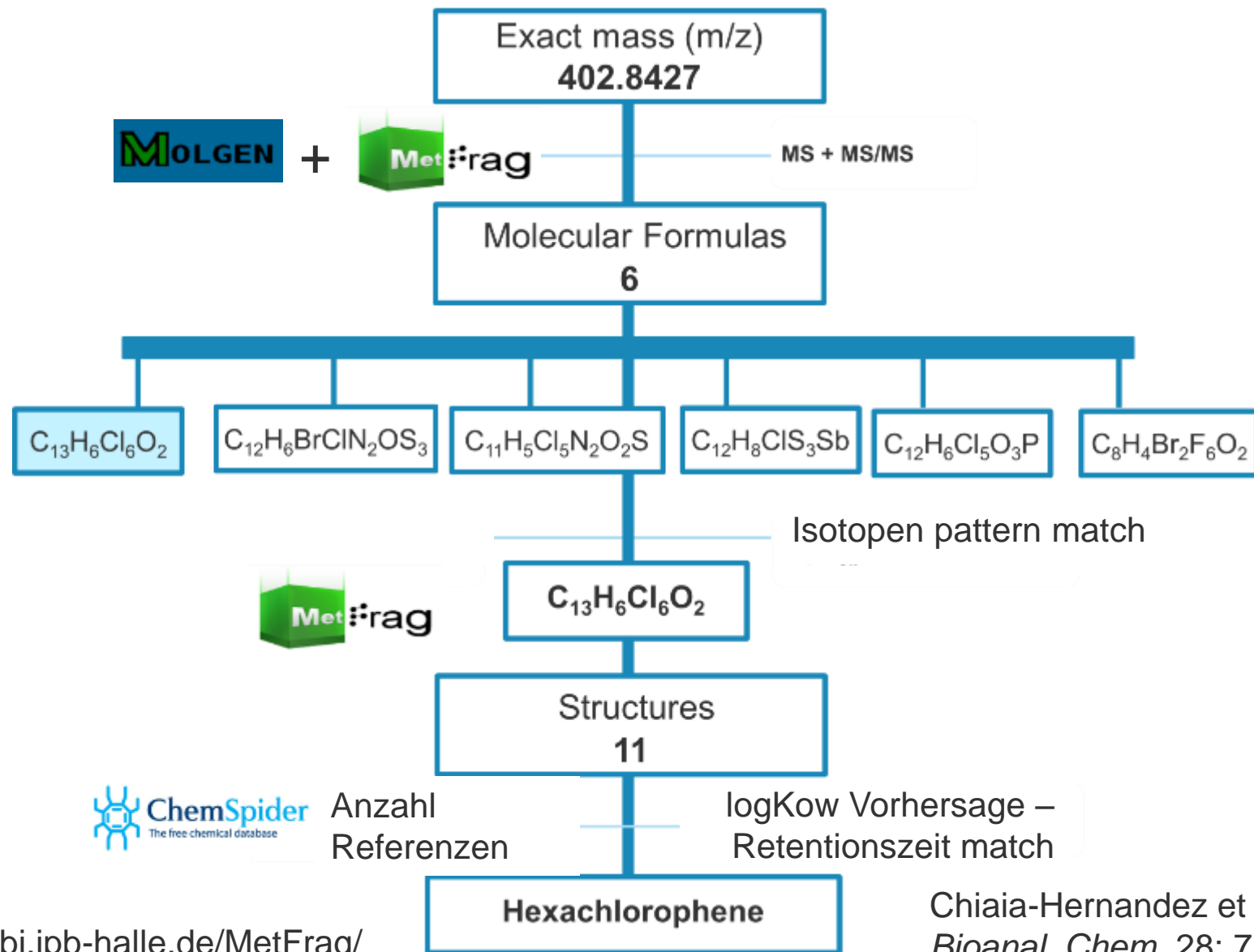
After bio



Hydroxylation

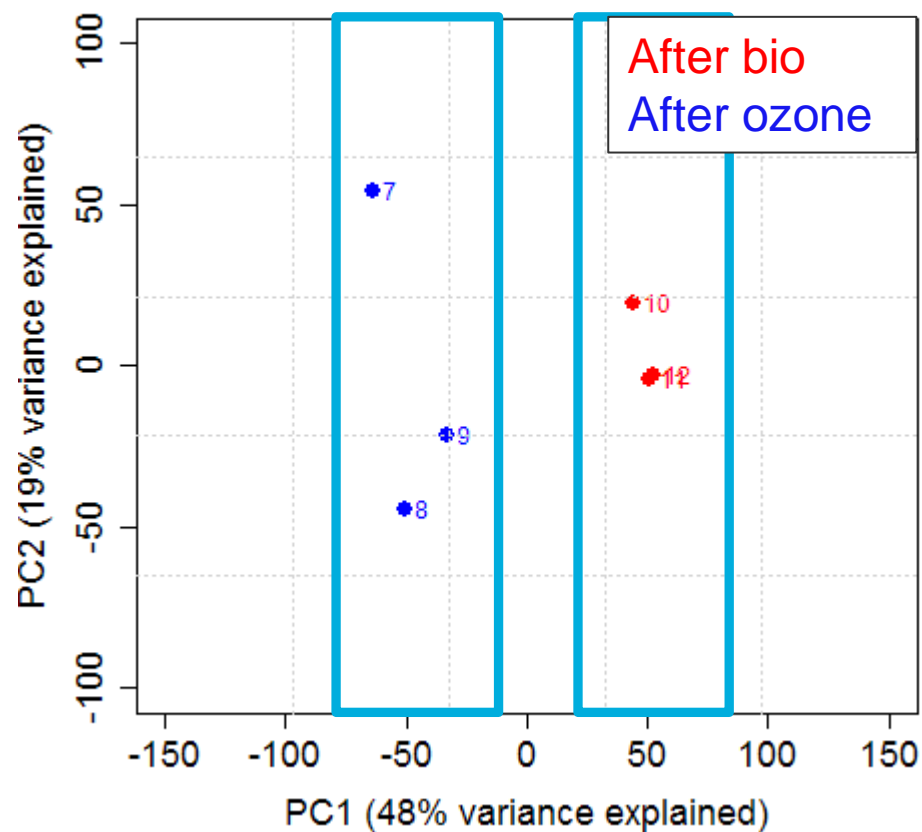


# Non-Target Screening von intensiven Peaks mit negativem Massendefekt und Chlorisotopenpattern

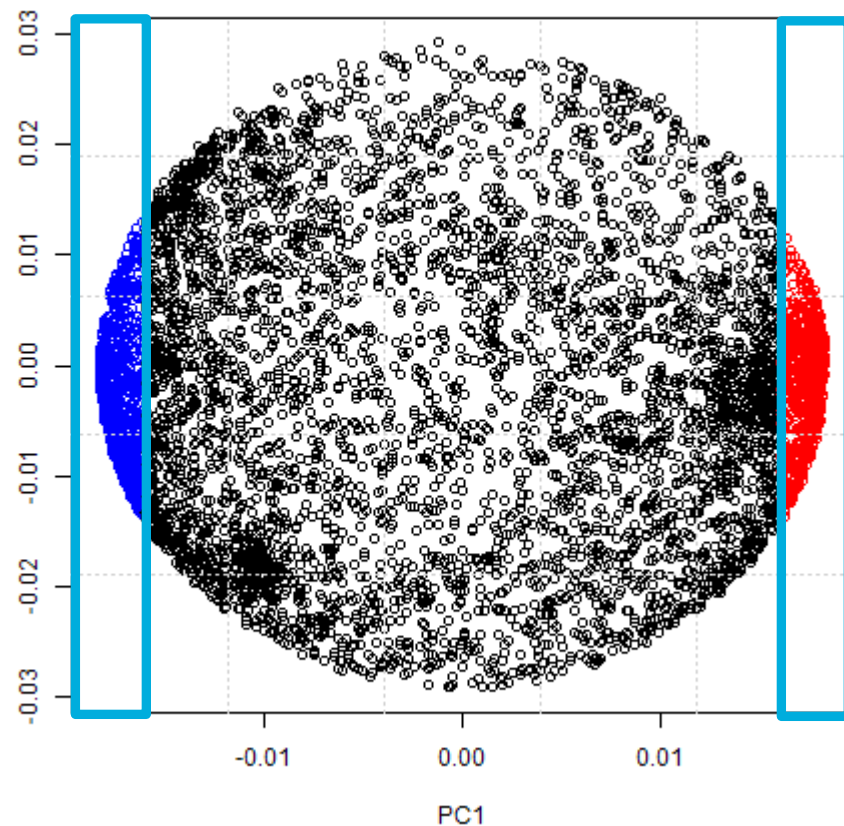


Example of 3 mg/L Ozone

Scores Plot for Nontarget scaled peaks



Loading Plot for Nontarget scaled peaks



# Loads along the river Rhine: example lamotrigin

