

# IDENTIFICATION AND QUANTIFICATION OF TRANSFORMATION PRODUCTS IN THE AQUATIC ENVIRONMENT BY HIGH RESOLUTION MASS SPECTROMETRY

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# Transformation Products (TPs)/Metabolites in European Directives and Guidelines



## **Drinking Water Directive:**

...pesticides and their **relevant metabolites** in drinking water must not exceed 0.1 µg/L.



**Council Directive 91/414/EEC** concerning the placing of plant protection products on the market

...**64 times „relevant metabolites“**



**EMA-Guideline** on the environmental risk assessment of medical products for human use (June 2006)

... **relevant metabolites** ....

# Analytical strategy to identify and quantify transformation products (TPs)



# Analytical procedure

## Challenges:

Low concentrations

Broad set of compounds

Ionization in positive and negative mode

Identification without reference standards



**Sample - ground, surface, waste water**  
- 1/0.5/0.2 liter, filtration,  
- pH 6.8, 90 isotope labeled internal standards



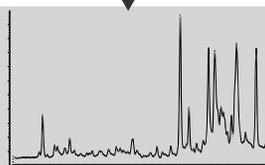
**Enrichment (SPE)**  
- mixed bed cartridge (RP, Ionic exchange mode)



**Chromatography**  
- Reversed Phase Xbridge 50 x 2mm  
- H<sub>2</sub>O/MeOH gradient with 0.1% formic acid



**LTQ-Orbitrap-MS**  
- Electrospray ionisation (positive/negative run)  
- HR scan with data dependent MSMS-scan



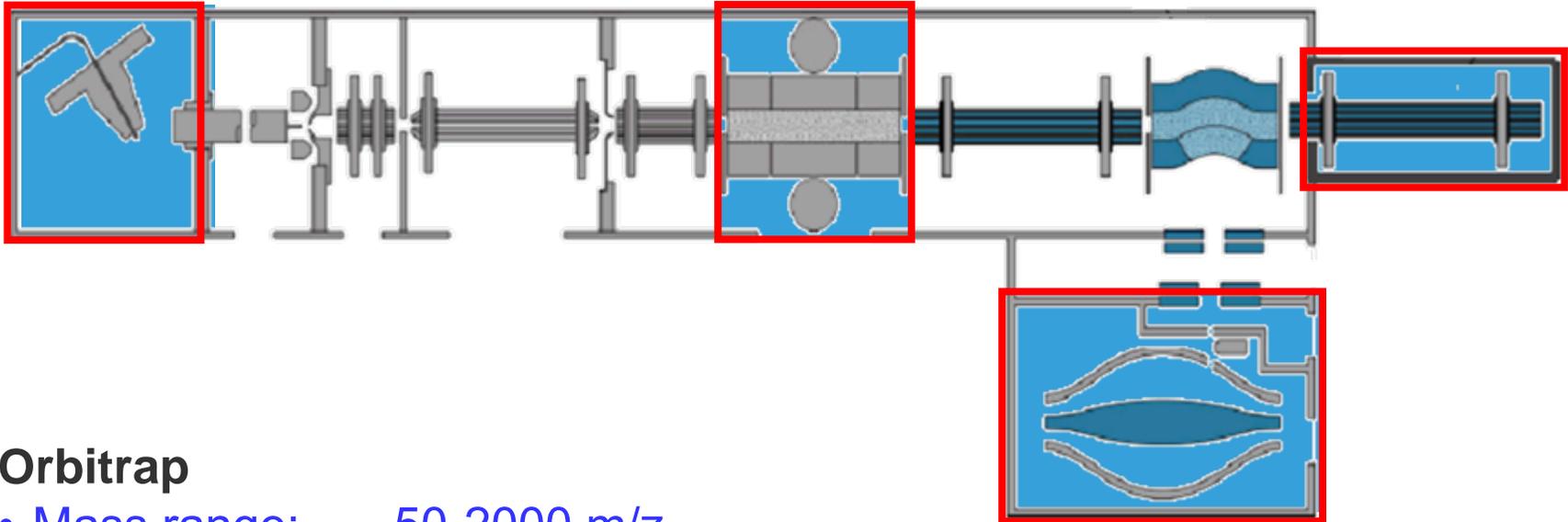
**HR Orbitrap-Chromatogram**  
- Quantification of ~240 target analytes  
- Target, suspects and non-target screening

# Hybrid mass spectrometer - Orbitrap XL

## Electrospray Interface

## Linear Ion trap LTQ

- Mass range: 50-2000 m/z
- Resolution: Unit resolution
- Fragmentation: CID



## Orbitrap

- Mass range: 50-2000 m/z
- Resolution: 100,000 (@ 400m/z)
- Accuracy: < 5 ppm

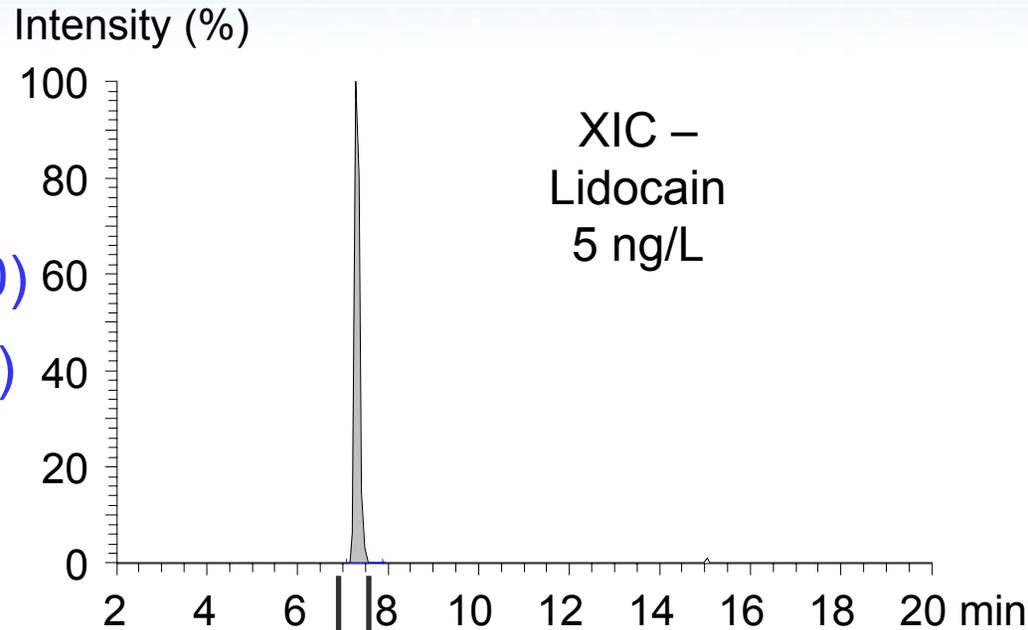
## Collision Cell

- Fragmentation: higher energy collision dissociation (HCD)

# MS settings for screening

## Per 30sec LC-peak:

- 10 MS scans (R 60,000)
- 30 HCD MSMS scans (R 7,500)
- 30 CID MSMS scans (Unit res.)



Orbitrap

MS Scan  
R 60,000

MS/MS Scan  
HCD

MS/MS Scan  
HCD

MS/MS Scan  
HCD

MS S  
R 60,000

LTD

MS/MS Scan  
CID

MS/MS Scan  
CID

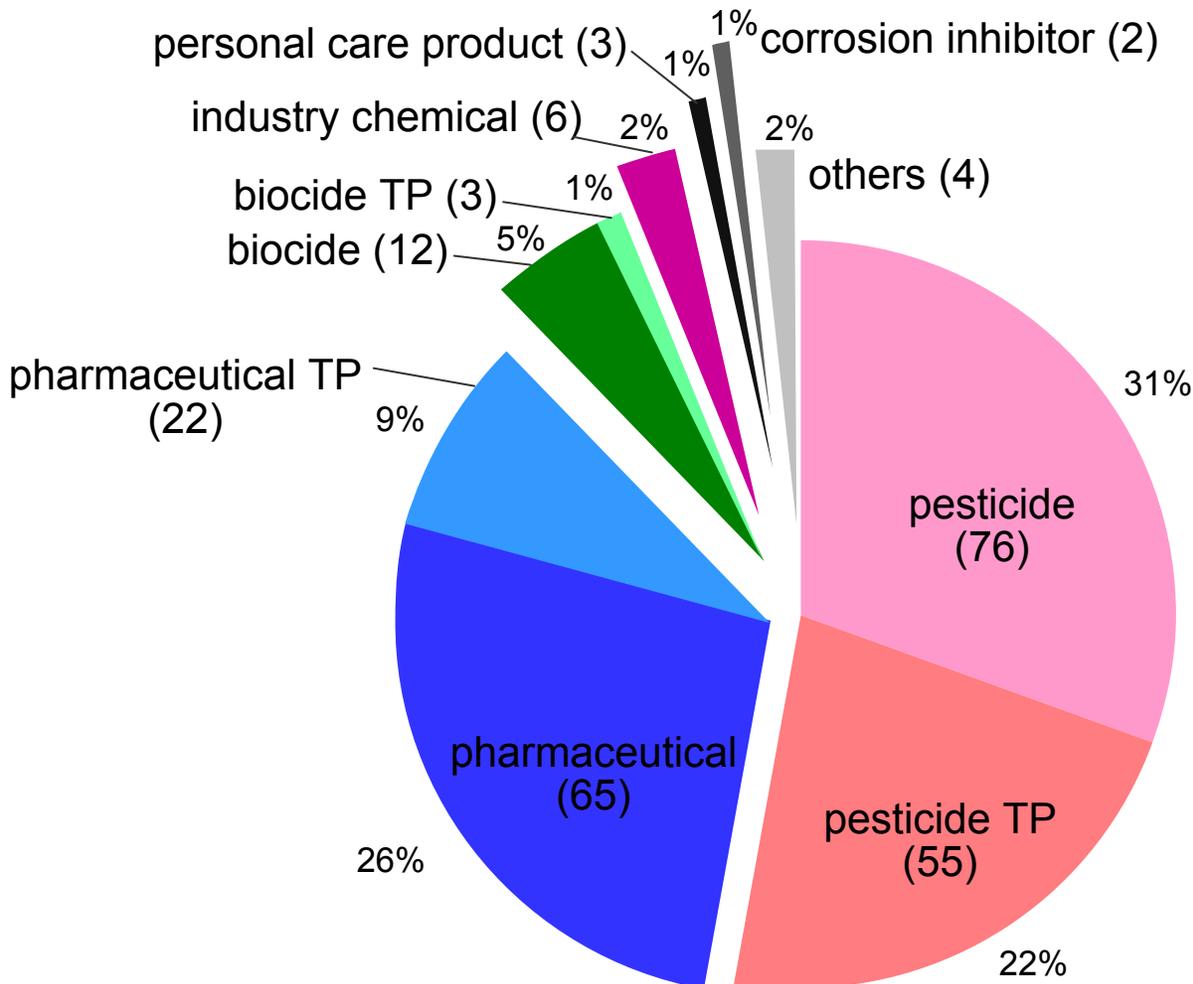
MS/MS Scan  
CID

data-dependent MS/MS with target list (170 pos/90 neg)  
and dynamic exclusion (8 sec)

3 sec cycle time

# Suitability of screening procedure

167 parent compounds, 81 transformation products (TP):  
Mr: 115 – 1000; Kow: -2.2 up to 5.7, 52 % neutral, 48 % ionic



## Method detection limits:

(surface water)

75 % ≤ 10 ng/L

20 % 10 - 100 ng/L

5 % < 1000 ng/L

## Relative recoveries:

(surface water)

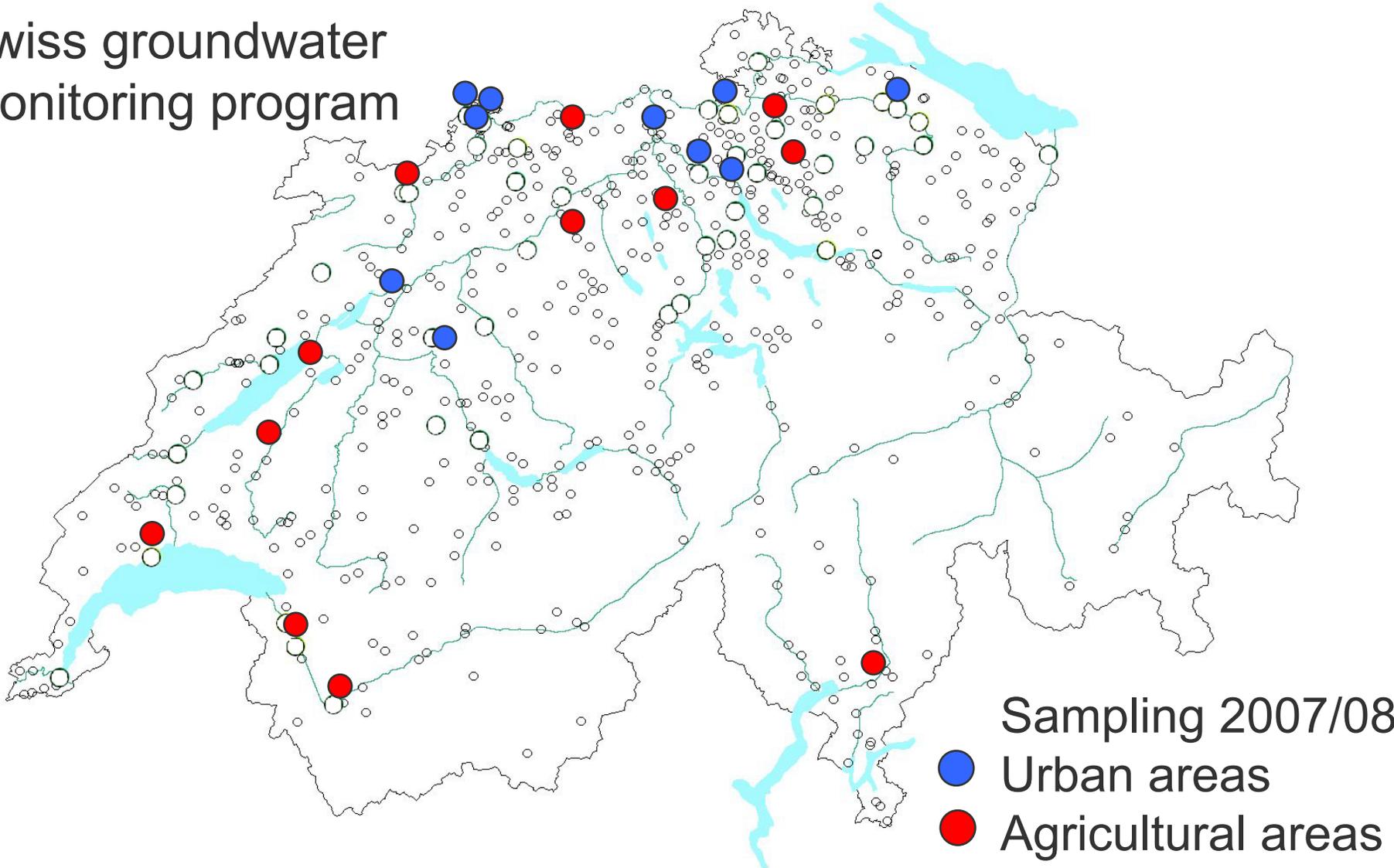
80 % > 80 %

15 % 20-80 %

5 % < 20 %

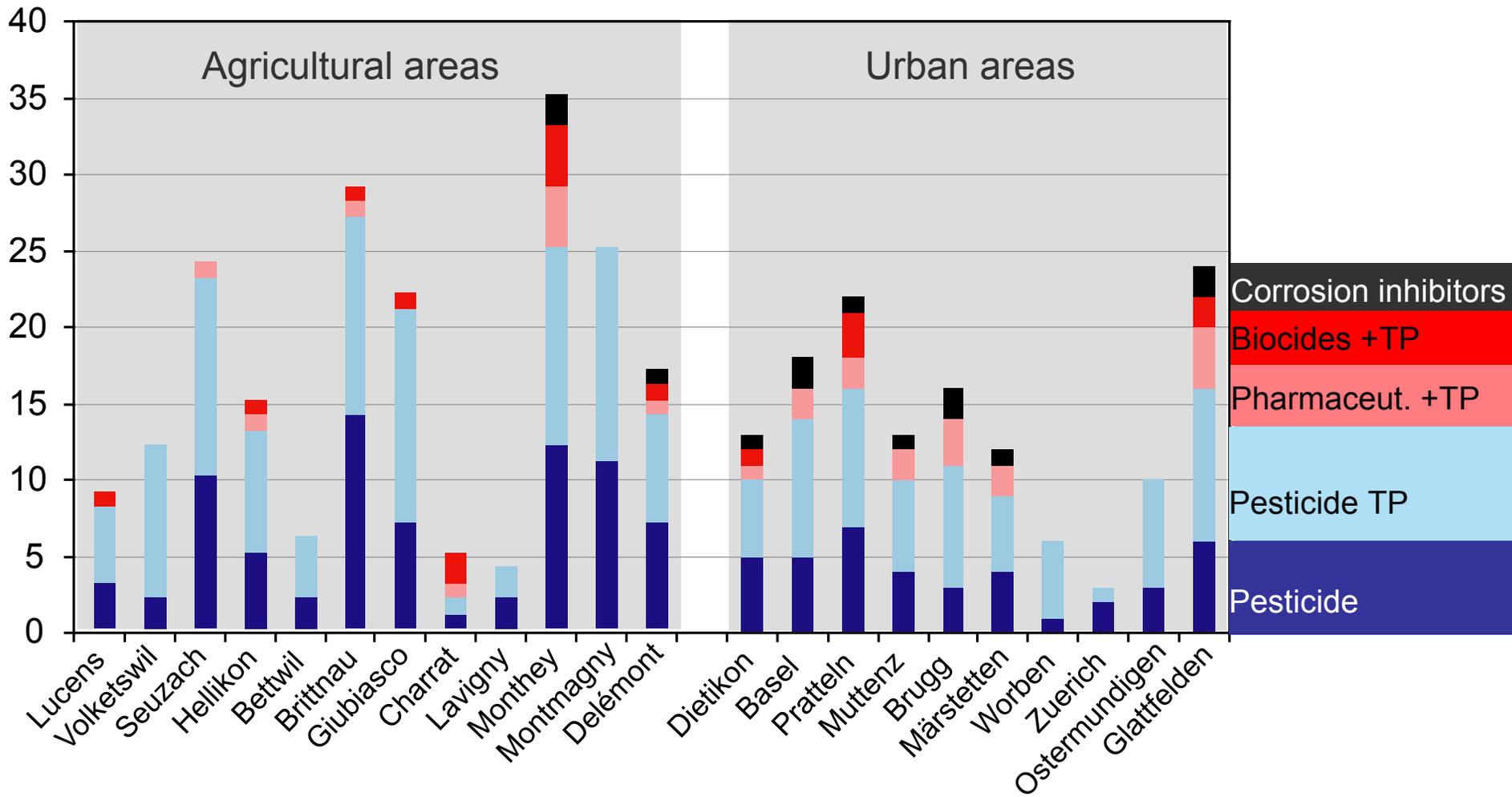
# Target screening: Case study groundwater

## Swiss groundwater monitoring program



# Target screening: Case study groundwater

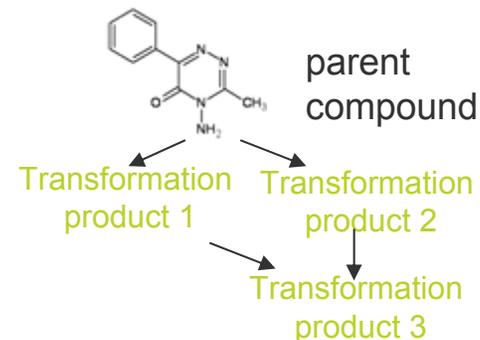
## Number of findings



# Suspects screening

## 1. Step

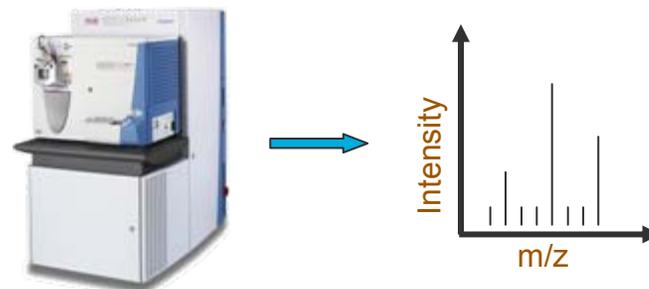
Prediction of possible transformation products using structure-biodegradation relationships & data mining



List of target transformation products

## 2. Step

Identification of transformation products in environmental samples by high resolution mass spectrometry



**Identified transformation products**

# Identification procedure without reference standards

## 1. Exact mass:

→ extracted chromatogram

## 2. Retention time:

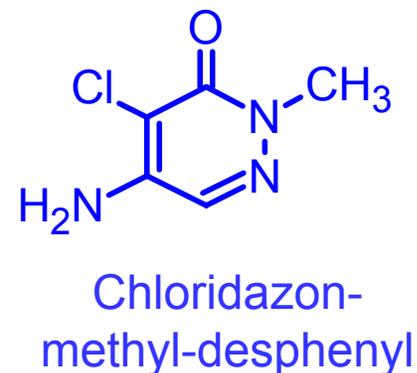
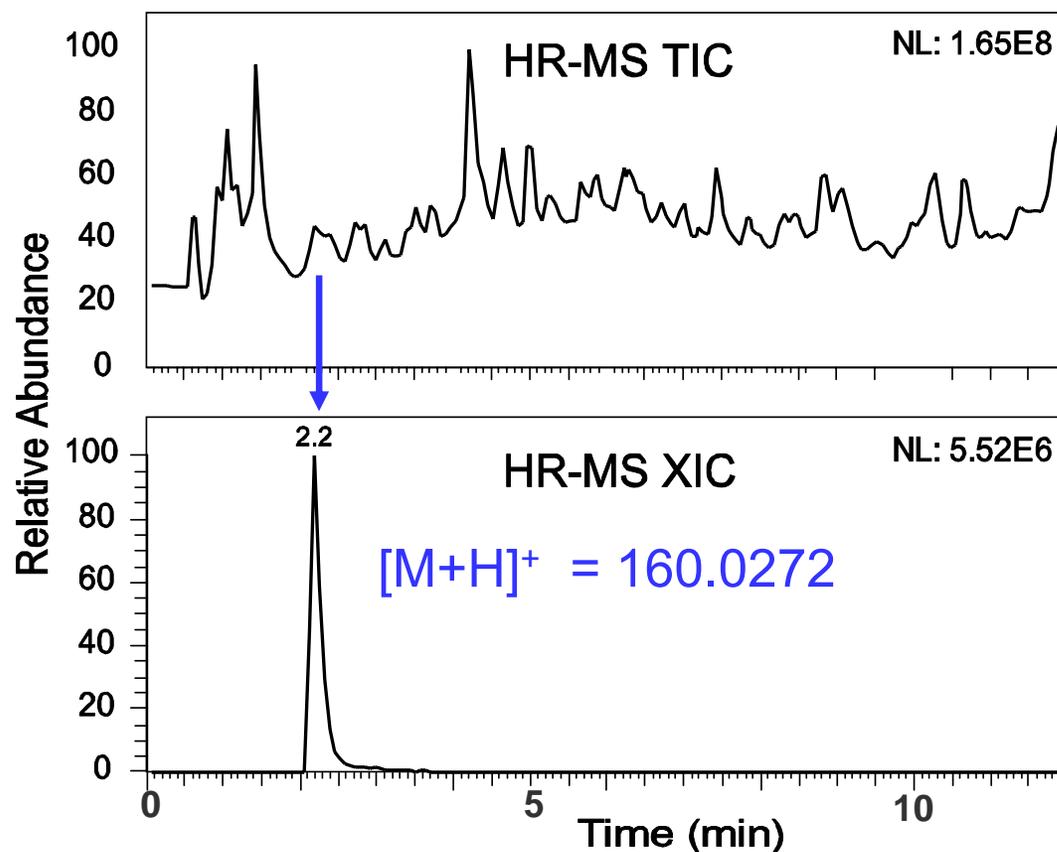
→ comparison to parent compound  
or prediction based on log K<sub>ow</sub>

## 3. Molecular structure:

→ interpretation of MS/MS fragments

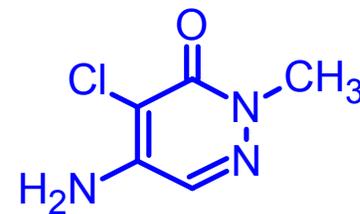
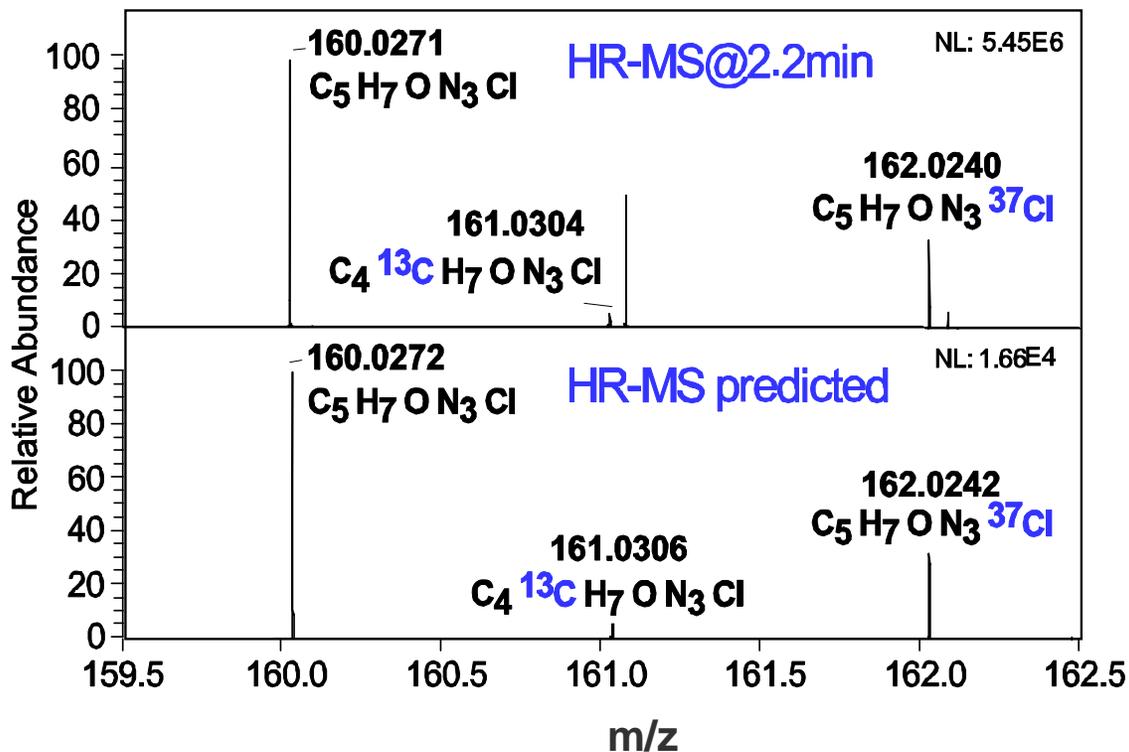
# Suspects screening: 1. Exact mass & isotope pattern

Compound detection: filtering with 5 ppm extraction window



# Suspects screening: 1. Exact mass & isotope pattern

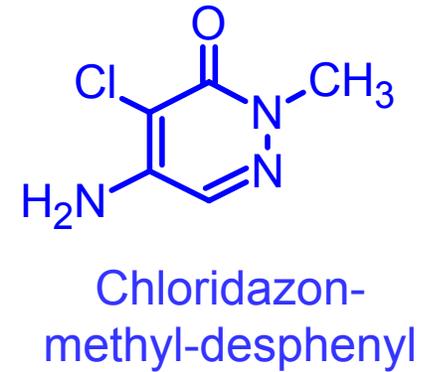
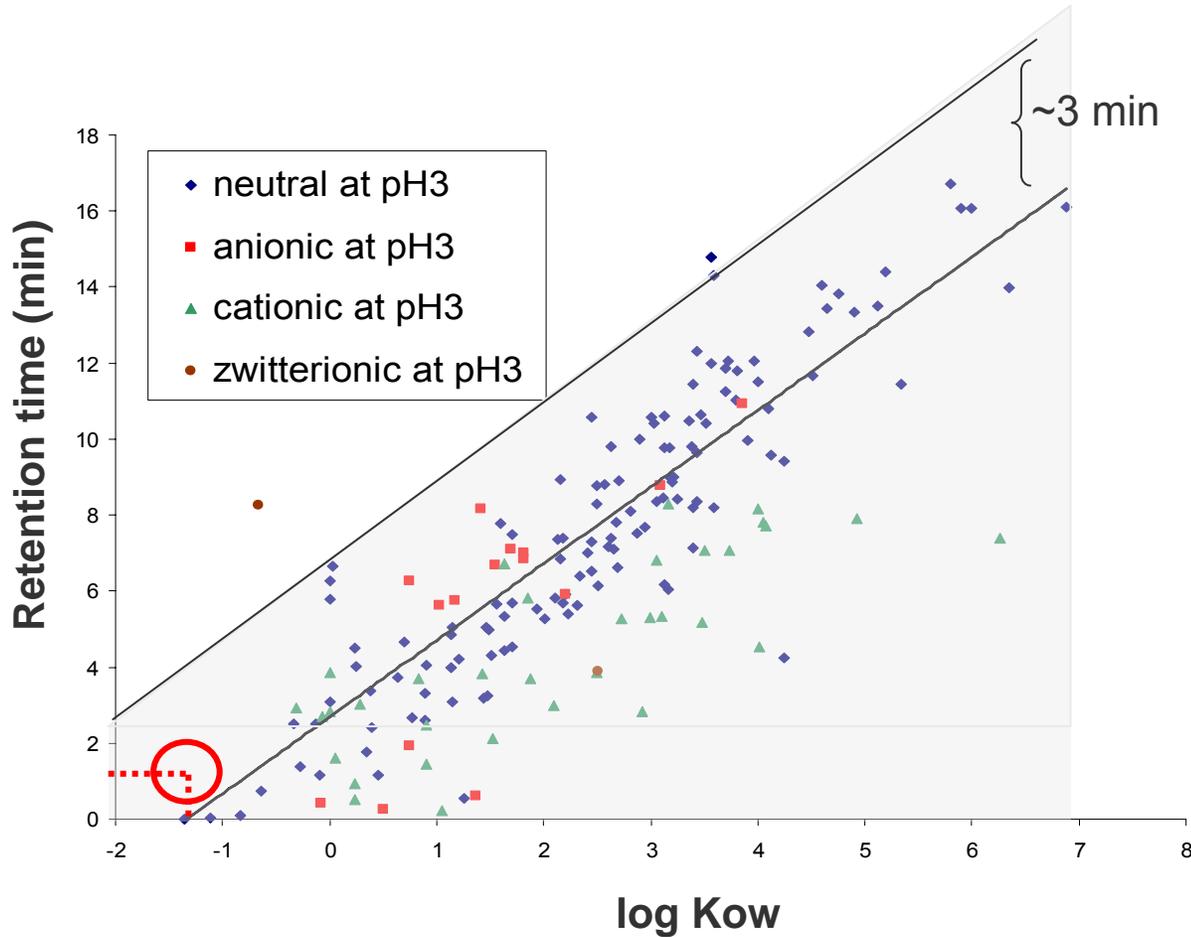
Isotope pattern:  $C_5H_6ClN_3O$



Chloridazon-  
methyl-desphenyl

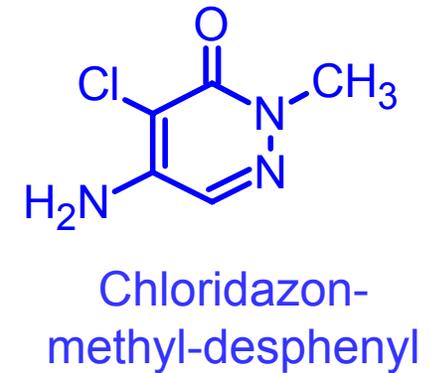
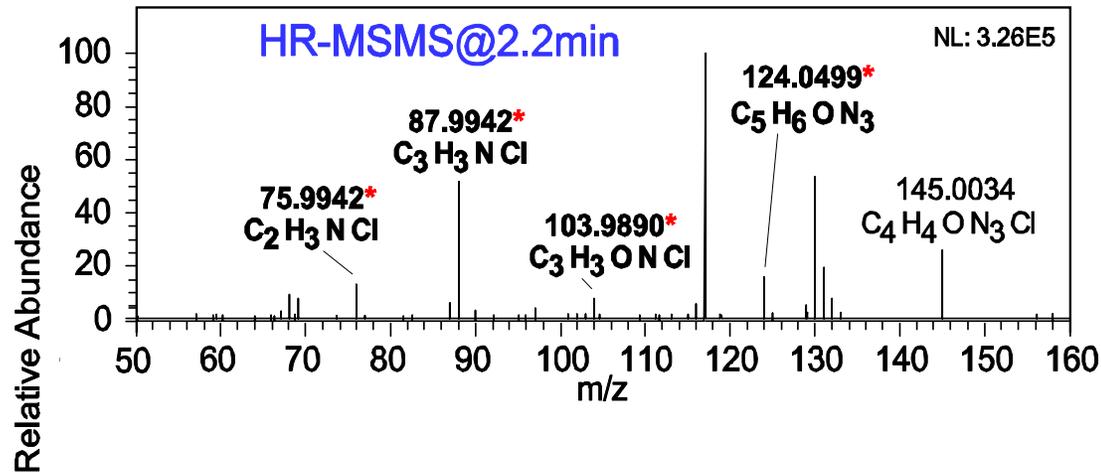
→ ~100 possible structures in Pubchem/Scifinder data bases

# Suspects screening: 2. Retention time



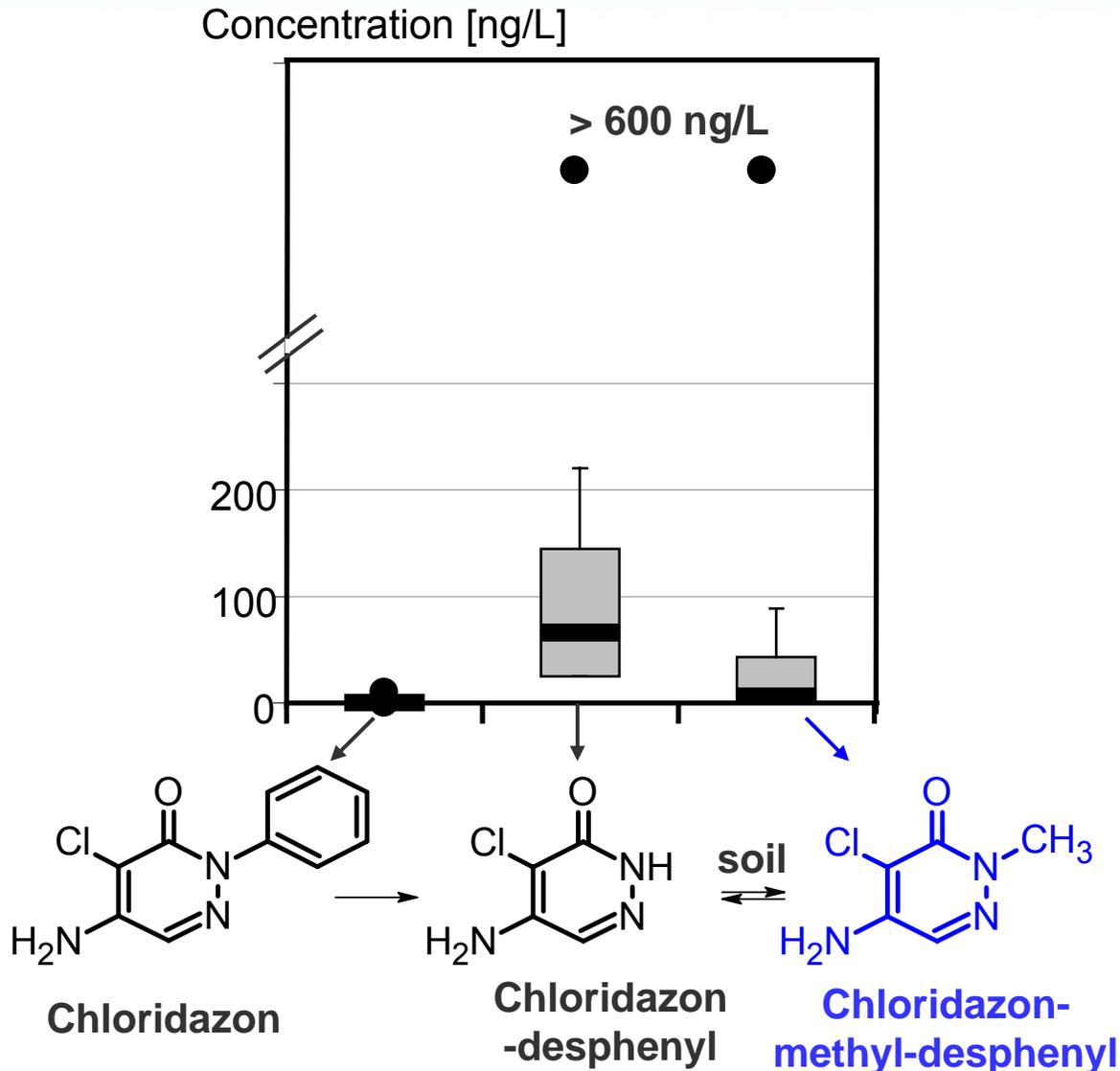
# Suspects screening:

## 3. Molecular structure - Interpretation of MS/MS fragments



\*predicted by  
fragmentation software  
(Mass frontiers)

# Chloridazon TPs in Swiss groundwater samples



# Suspects screening: Case study surface waters

7 water samples (agricultural areas, downstream wastewater treatment plants)

24 pesticides, 7 biocides, 21 pharmaceuticals

- University of Minnesota Pathway Prediction System\*
- Literature search

Predicted transformation products

1800

640

440

19

Identified transformation products

Show distinguishable peak with accurate mass

Retention time correlation

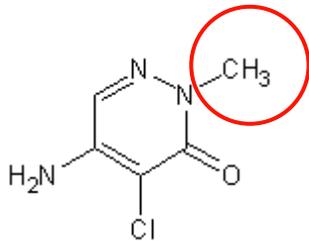
Isotope pattern, fragmentation prediction

+ 10 low concentrated transformation products identified by reference compounds

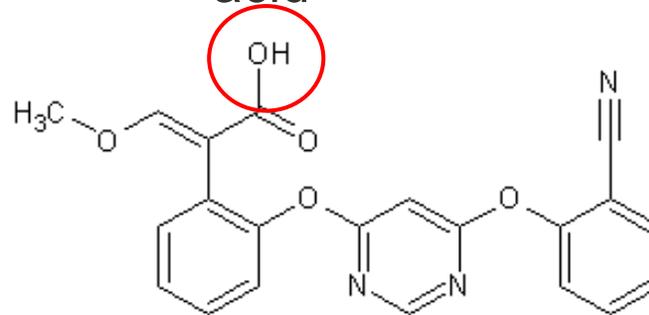
# Examples for identified transformation products

## Pesticide transformations products:

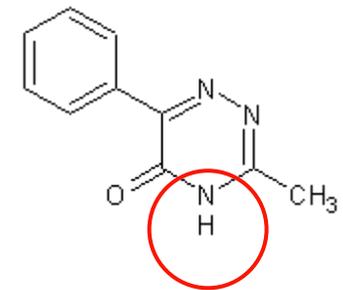
Chloridazon-methyl  
-desphenyl



Azoxystrobin  
acid



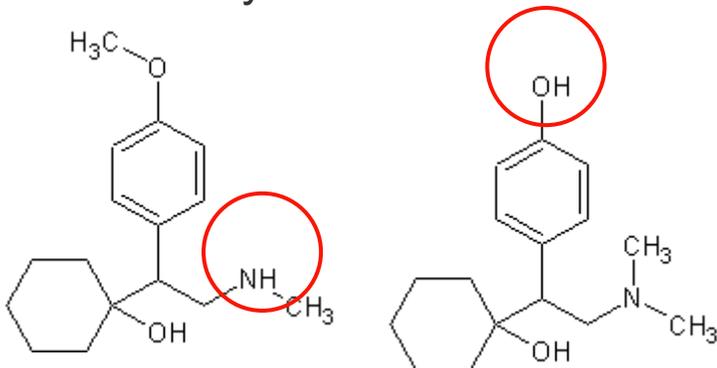
Metamitron-  
desamino



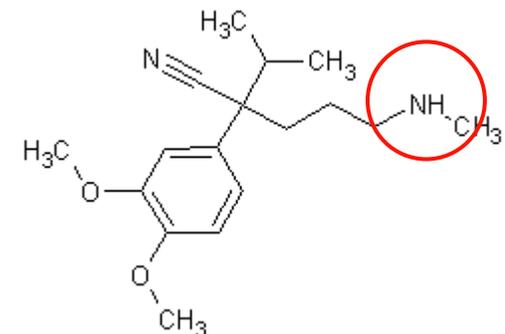
## Pharmaceutical transformation products:

N-Desmethylvenlafaxin

O-Desmethylvenlafaxin

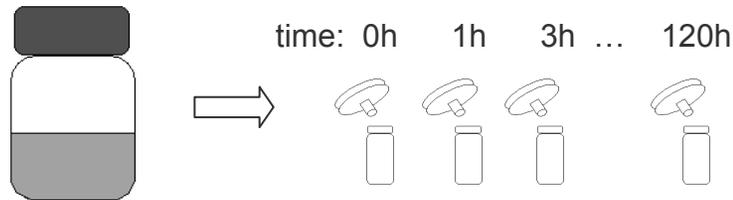


D617 (Verapamil-TP)



# Suspects screening

## Strategy for identification of relevant pharmaceutical TPs



50 mL,  $X_{SS}: 3 \text{ g}_{SS}\text{L}^{-1}$ ,  
dark,  $\text{O}_2 : >0.5 \text{ mg/L}$ ,  
 $T : 19^\circ$ , autoclaved  
controls

Qualitative  
agreement?



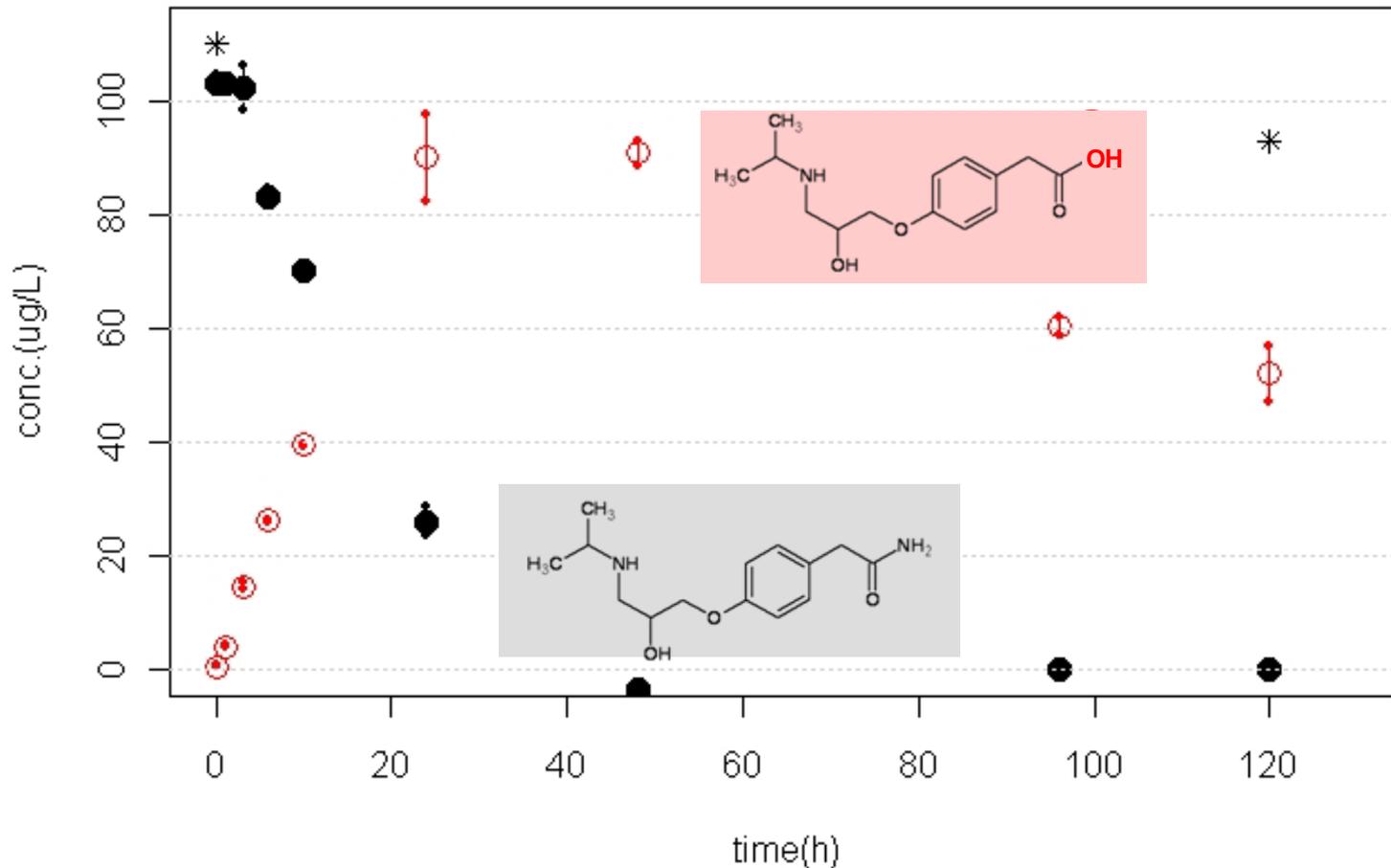
TP formed in **batch experiments** with sewage sludge from operating sewage treatment plant (STP)

**Influent / effluent** concentrations in corresponding STP

# Strategy for identification of relevant pharmaceutical TPs

Illustrative results from batch experiments

Duplicate batches for atenolol ( $\beta$ -blocker):

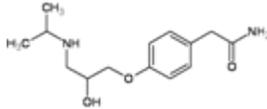
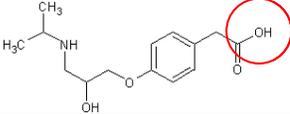
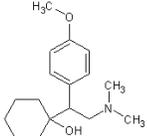
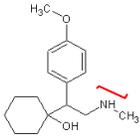
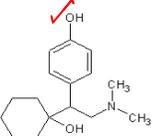


● : atenolol, ○ : atenolol acid (TP), \* : control

# Strategy for identification of relevant pharmaceutical TPs

TPs in batch experiments and STP influent/effluent

TPs with reference standards

parent compound	transformation product	structure	influent	effluent
Atenolol			✓	✓ ↓
	Atenolol acid		✓	✓ ↓
Venlafaxine			✓	✓ ↓
	N-desvenlafaxine		-	✓
	O-desvenlafaxine		✓	✓ ↑

# Conclusions

## Chemical analysis

- Combination of target and suspects screening enables evaluating the exposure to transformation products in the aquatic environment
- SPE-HPLC-ESI-MS method is necessary to enrich, separate, ionize and detect mostly polar transformation products
- High resolution mass spectrometry is indispensable to identify polar transformation products without reference standards
- Combination of laboratory batch experiments and screening of STP samples is suitable to identify new transformation products

## Exposure to transformation products

- confirmed that especially pesticide transformation products are important in groundwater
- For half of the parent compounds 1-2 transformation products were detected in surface and groundwater

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Groundwater Survey**



# Environmental transformation of organic compounds:

Towards a joint perspective on the importance of transformation products as environmental contaminants

12-17 September, 2010  
Monte Verità, Ascona, Switzerland

Organizers: Kathrin Fenner, John Sumpter, Juliane Hollender

