

**Ascona, 29. May - 3. June, 2016**

**Non-target screening using HPLC-HRMS  
in combination with Effect-Directed Analysis  
to prioritize contaminants in the aquatic environment**

**- Non-target real world applications -**

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Laboratory for Operation Control and Research

Langenau, Germany

- **Non-Target screening in drinking water analysis**
- **Introduction to HPTLC-EDA\***
- **Reciprocal iso-inhibition volume (RIV)**
- **Linkage of HPTLC-EDA with HPLC-MS**
- **Conclusion**

\*HPTLC: High Performance Thin Layer Chromatography

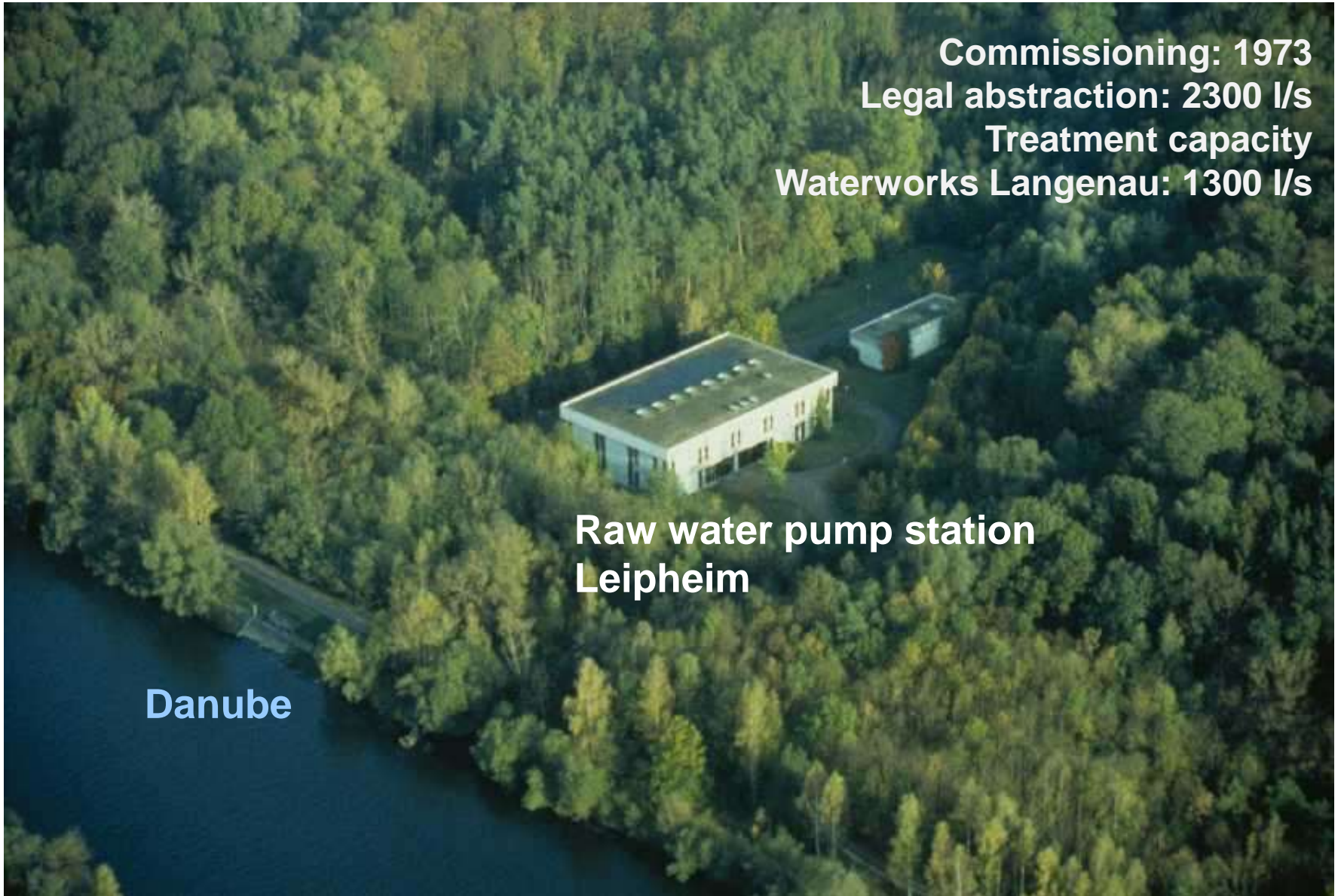
EDA: Effect-directed analysis

# River water abstracted from the Danube

Commissioning: 1973  
Legal abstraction: 2300 l/s  
Treatment capacity  
Waterworks Langenau: 1300 l/s

Raw water pump station  
Leipheim

Danube



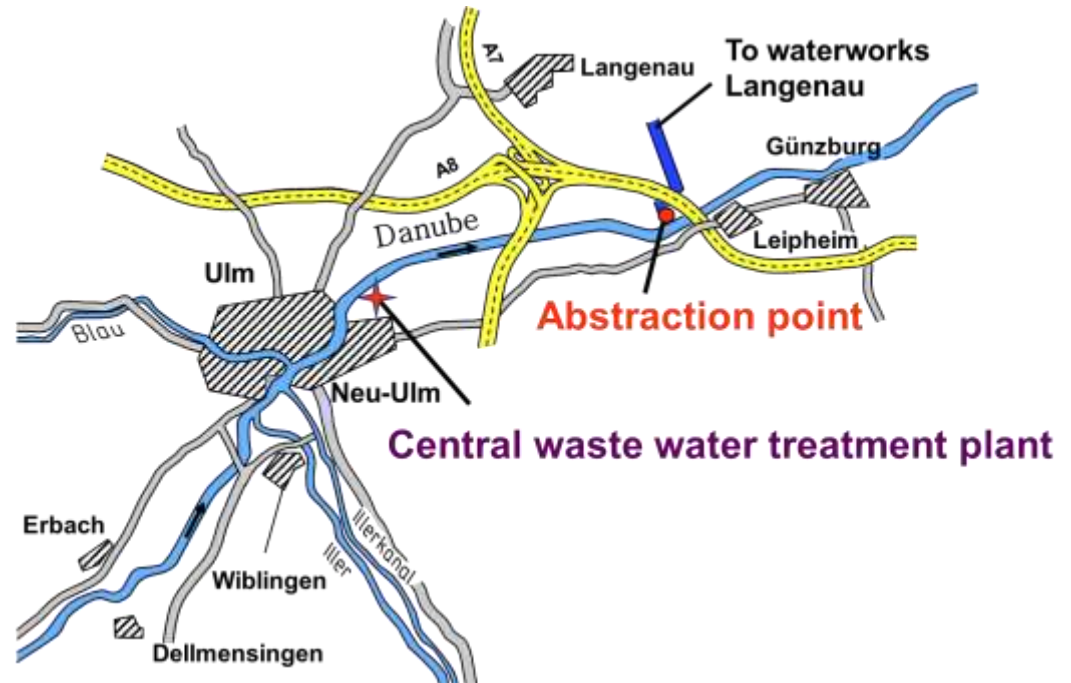
surface water  
river danube



treatment  
(process)



drinking water





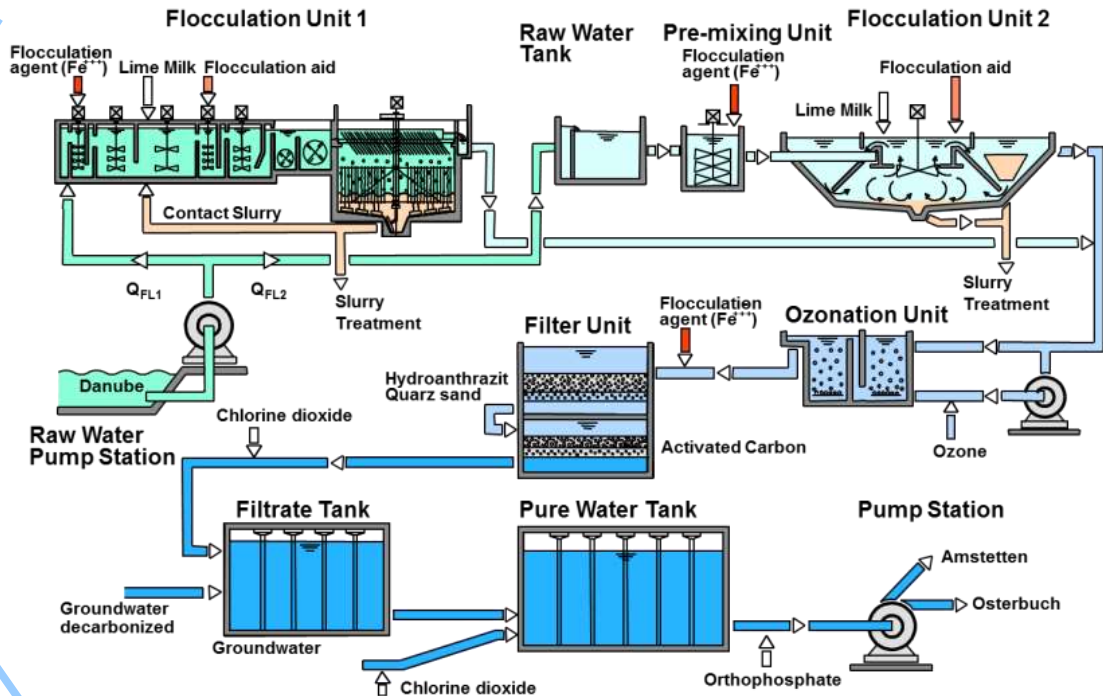
surface water  
river danube



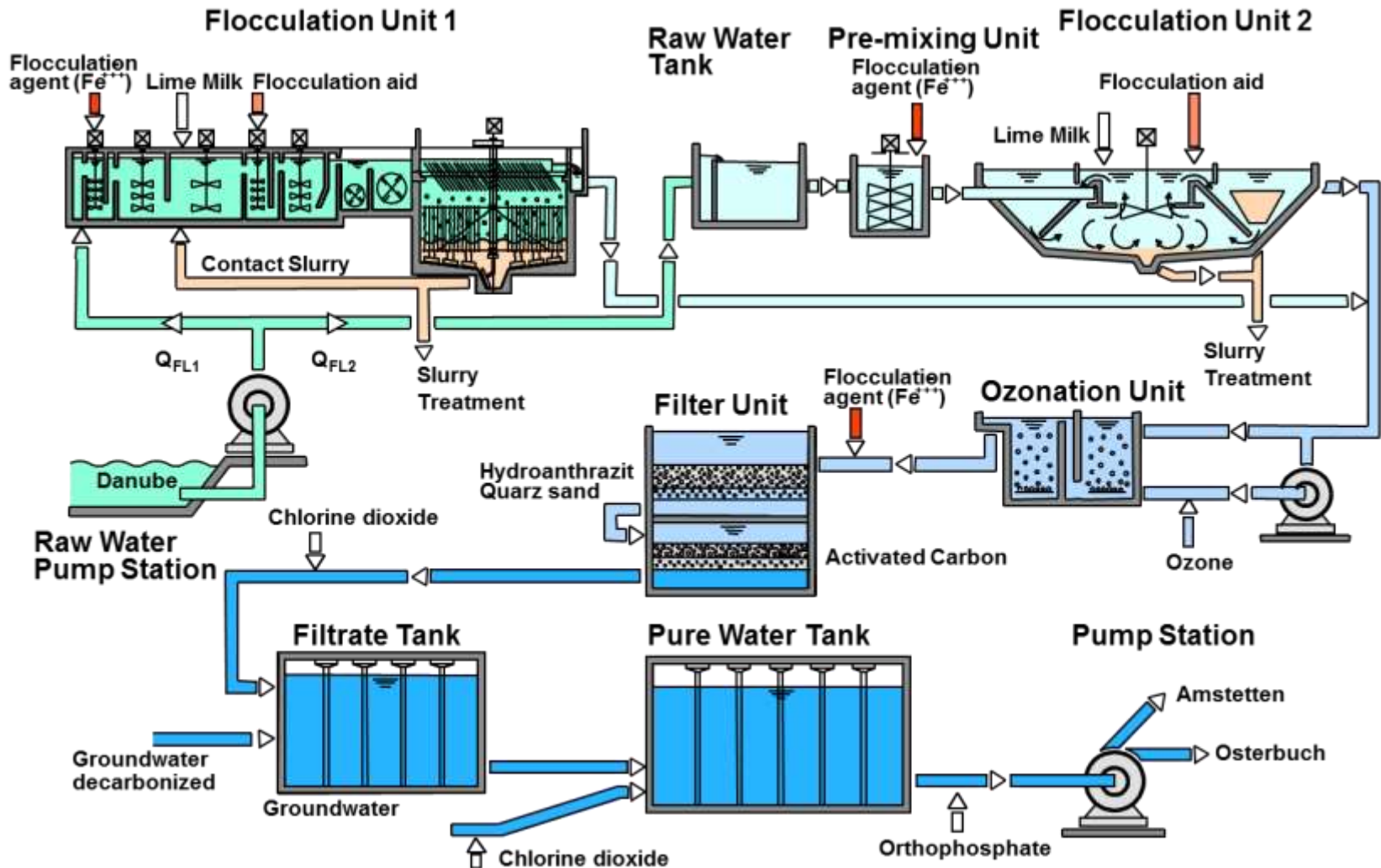
treatment  
(process)



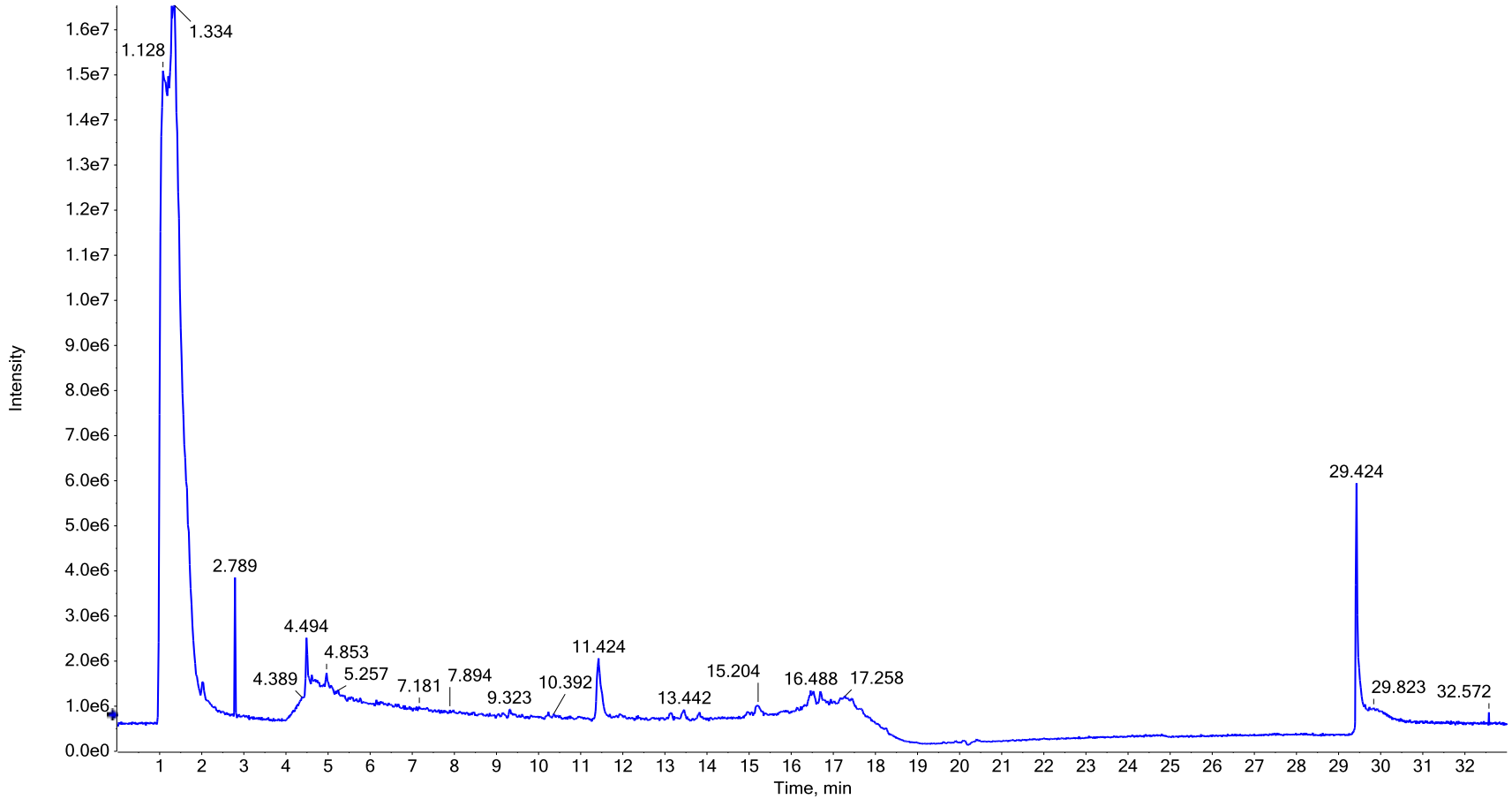
drinking water



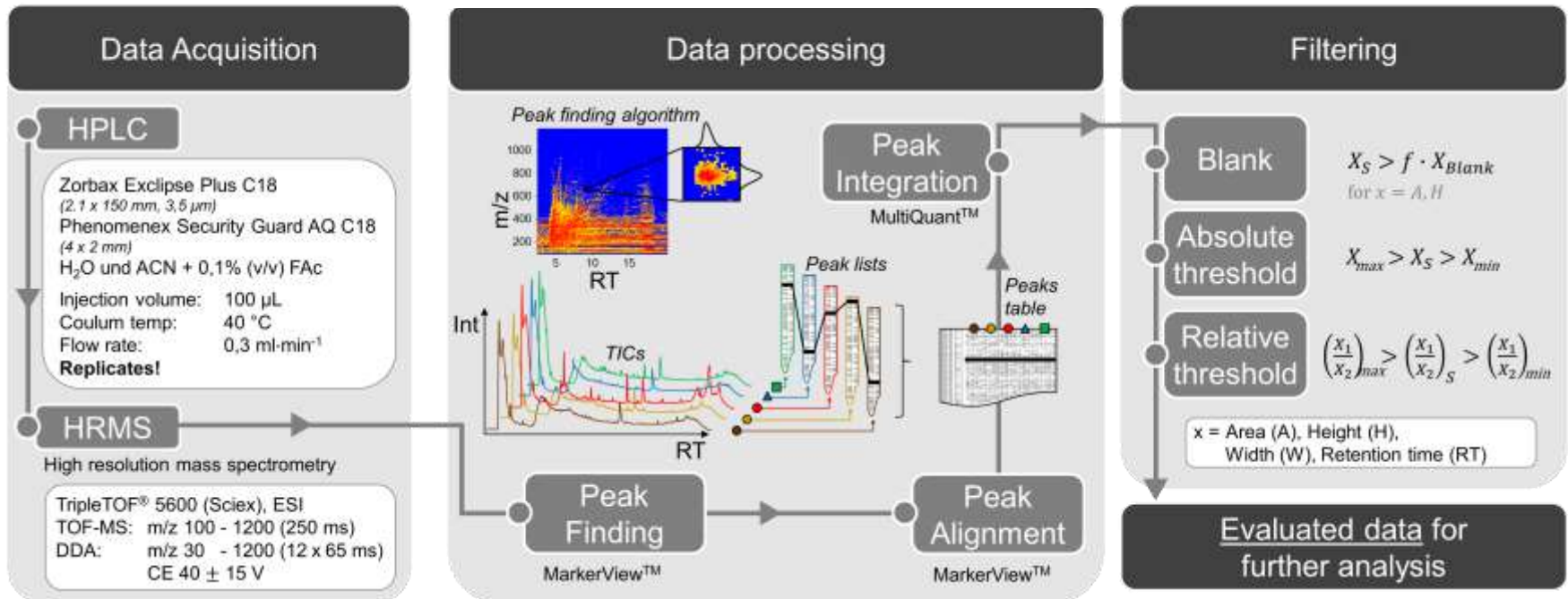
# Treatment of Danube water at Langenau waterworks



# TIC of HPLC-HRMS Chromatogram river Danube



# Non-Target-Screening Workflow



Talk Tobias Bader  
 Tuesday



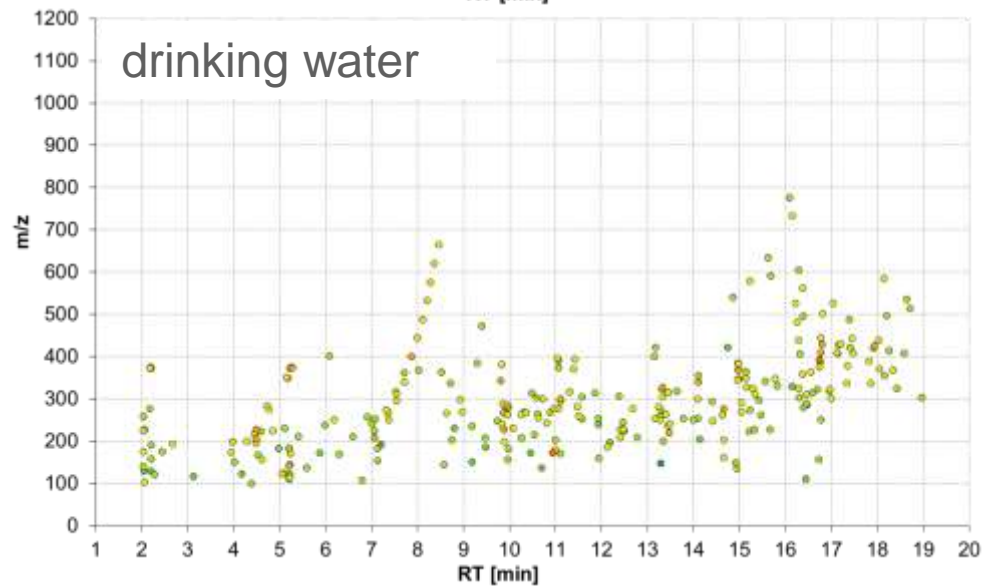
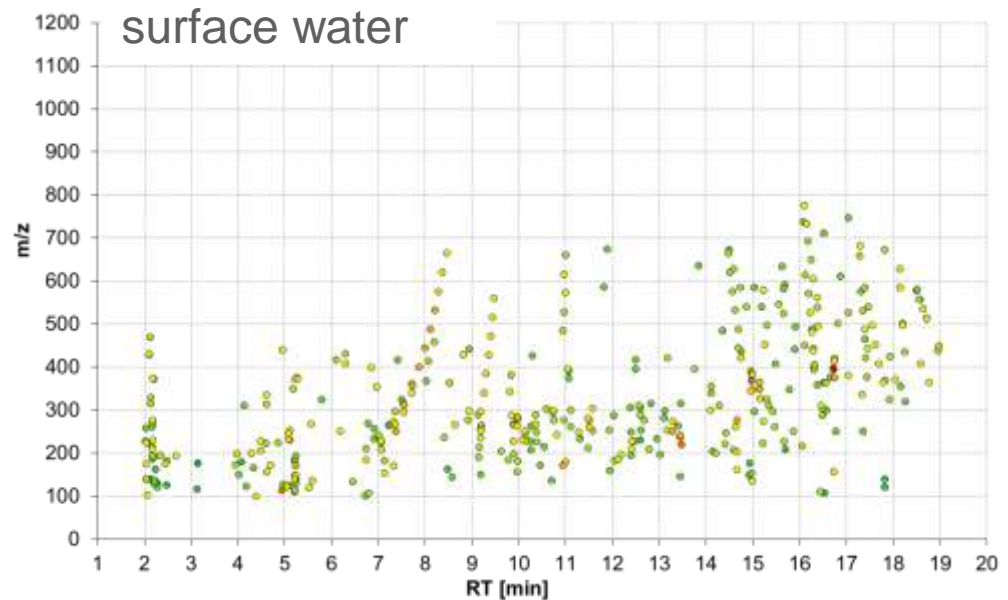
surface water  
river danube

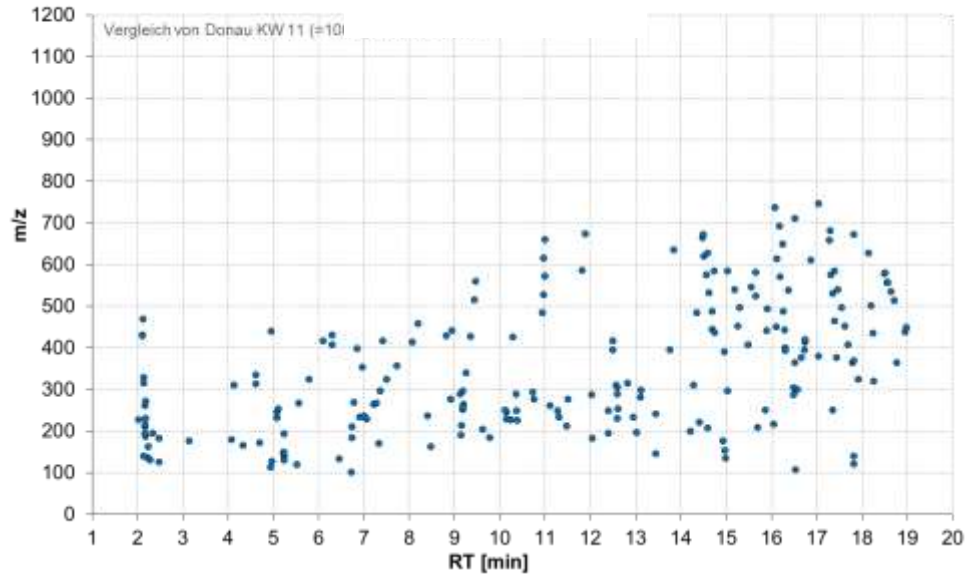


treatment  
(process)

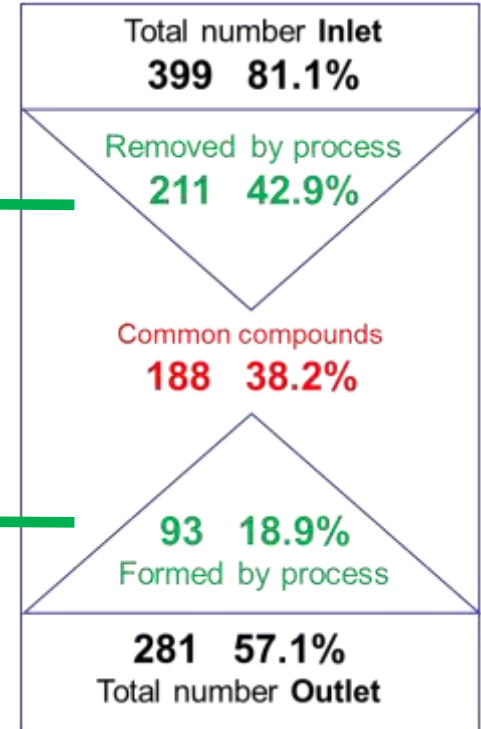
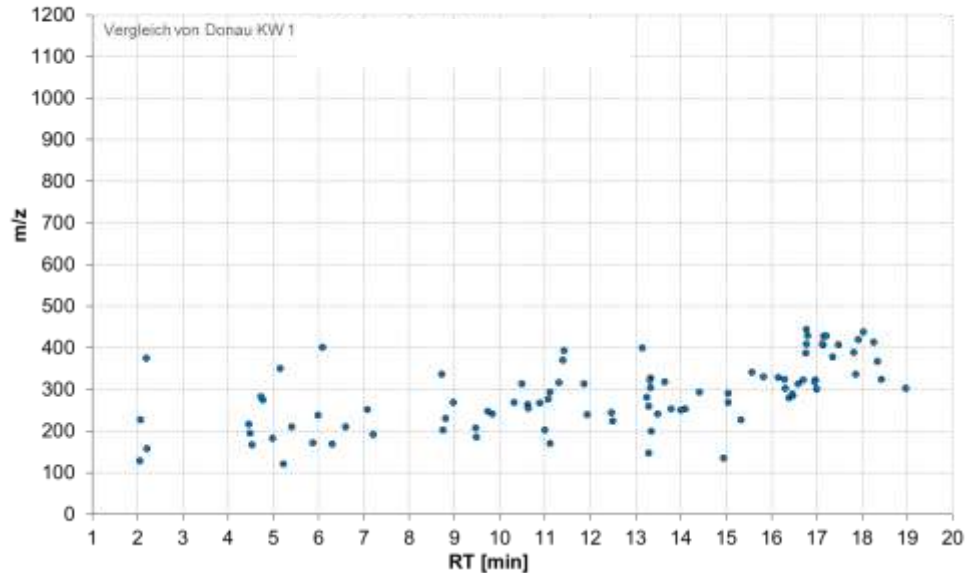


drinking water

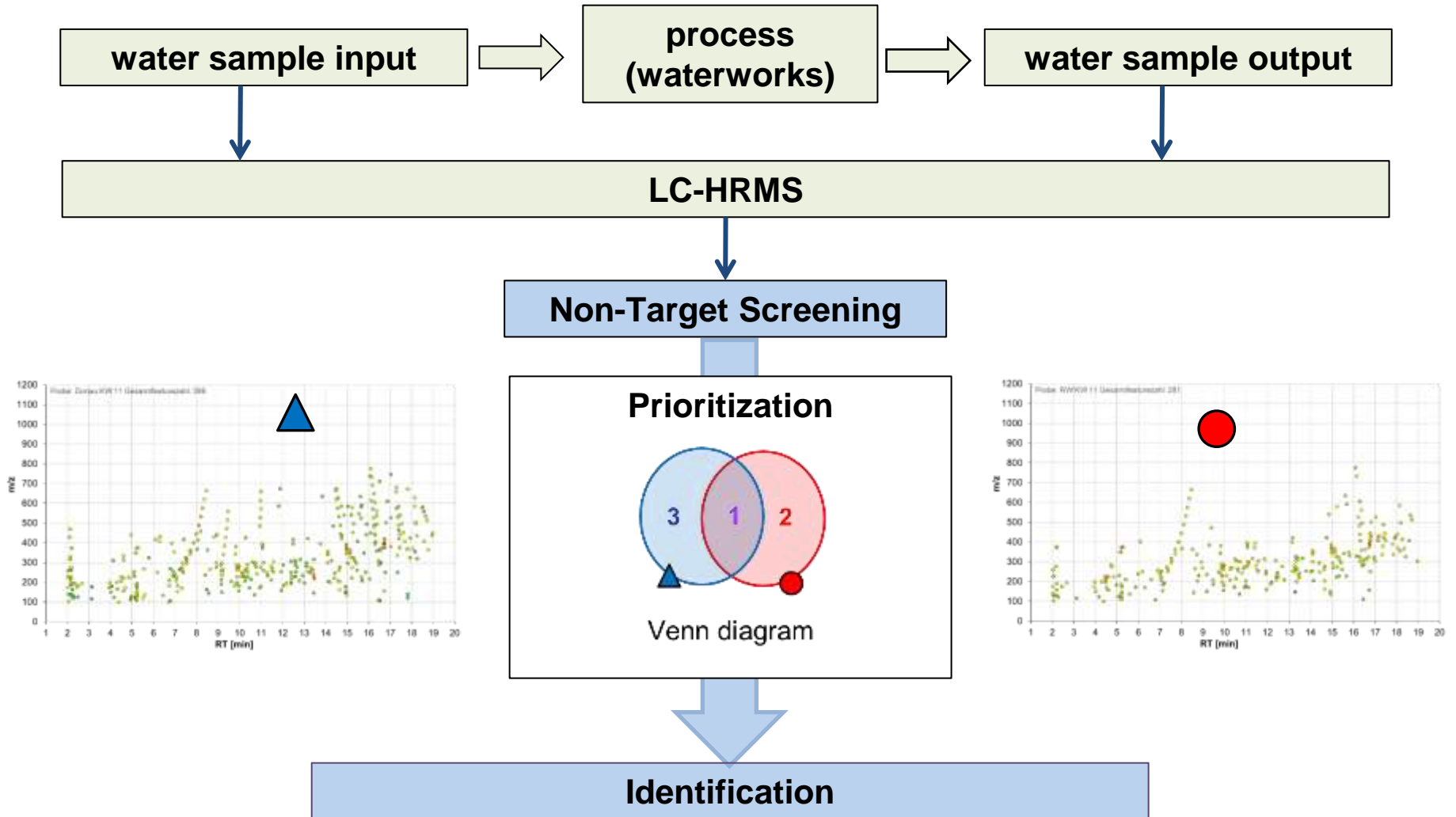




● 2 - Features die durch den Prozess selbst hinzukommen  
93



# Strategy for Prioritization



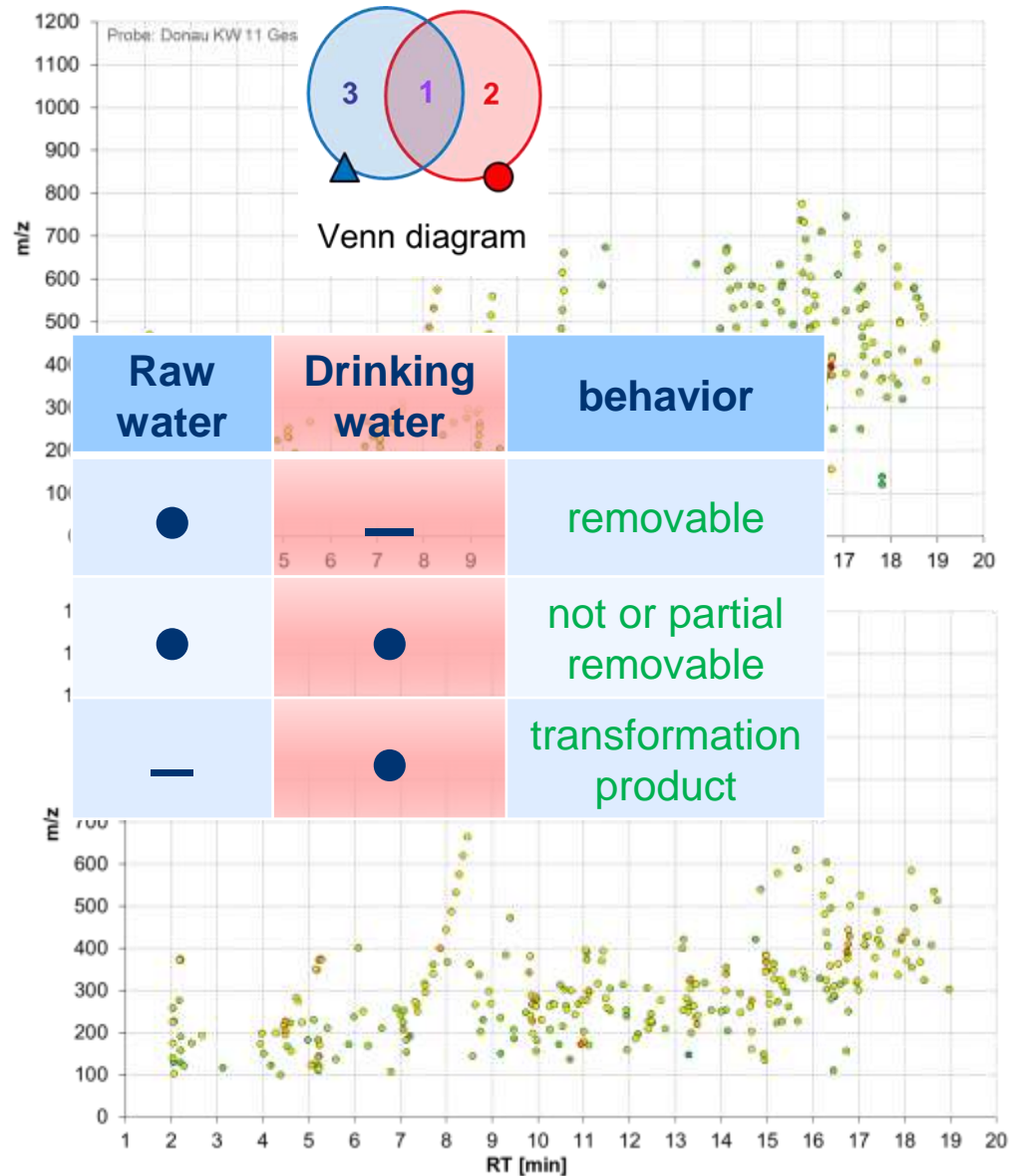
surface water  
river danube



treatment  
(process)



drinking water



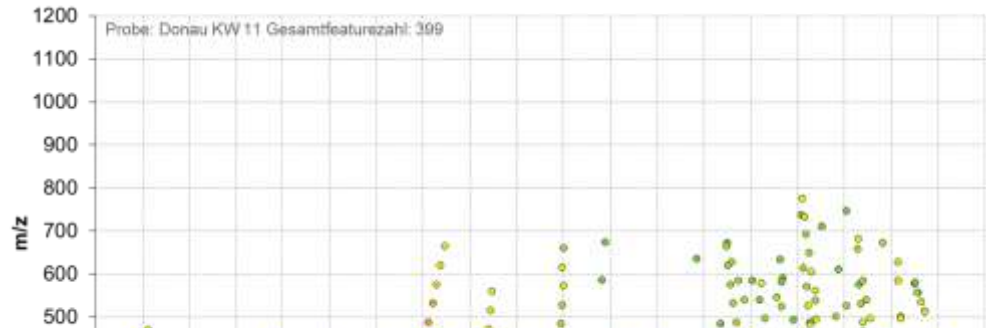
surface water  
river danube



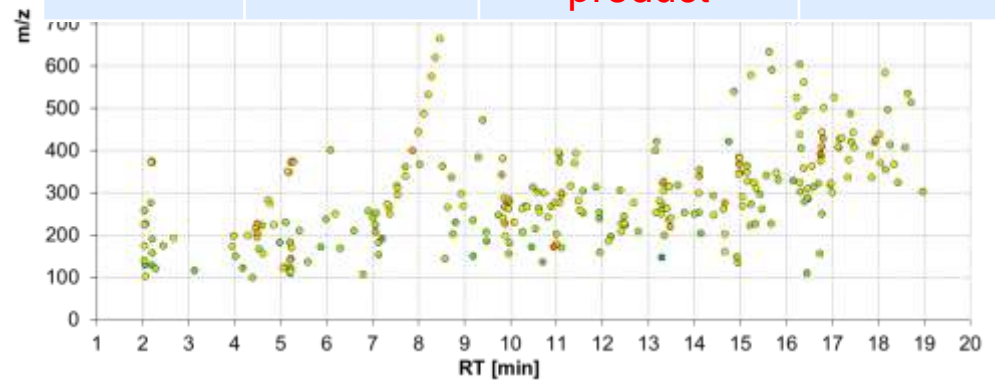
treatment  
(process)



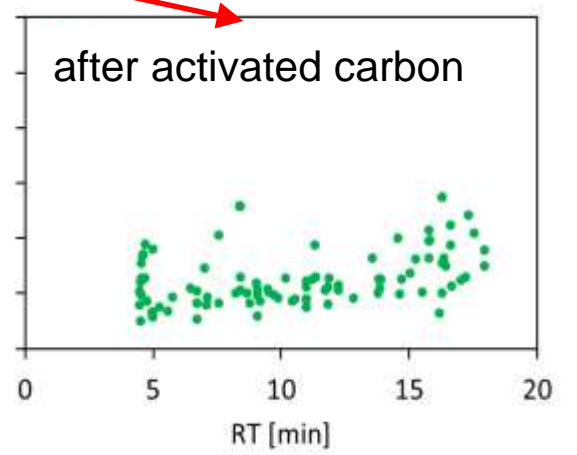
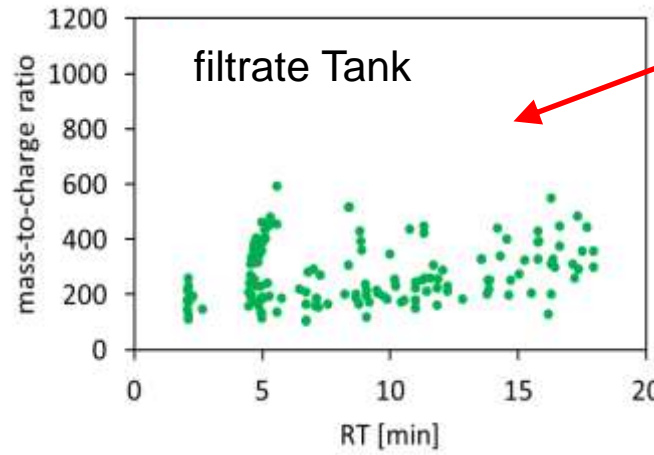
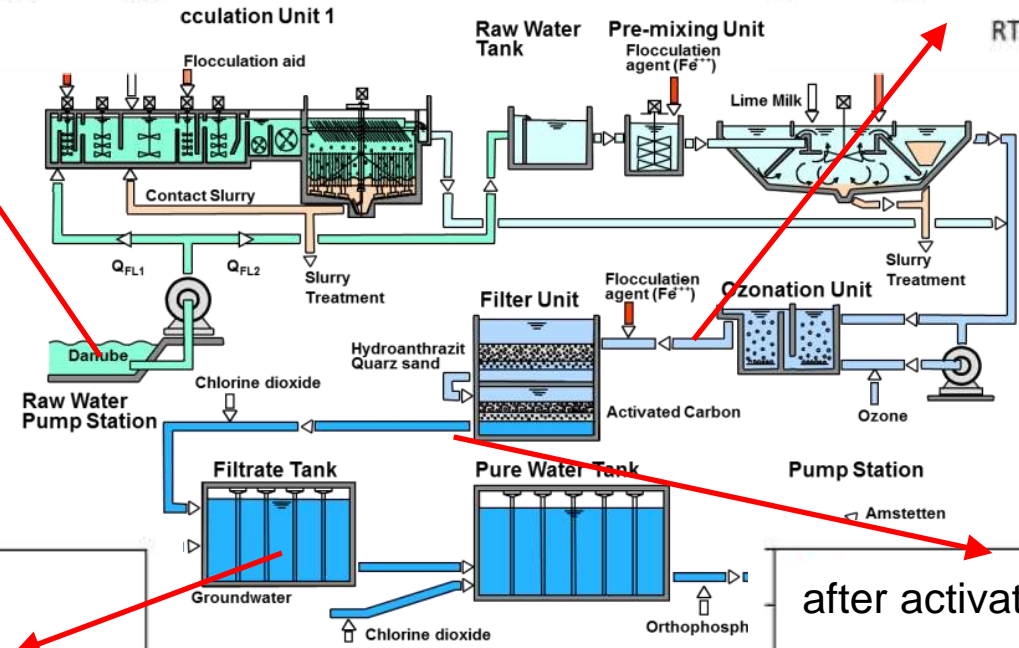
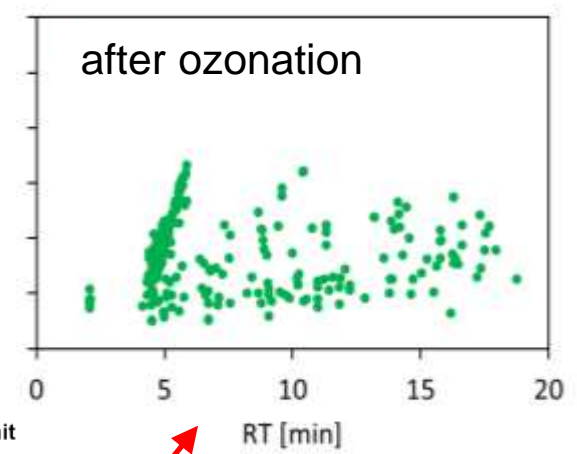
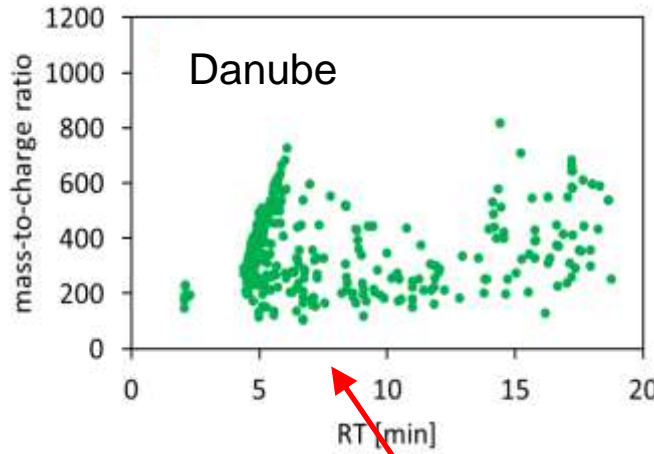
drinking water



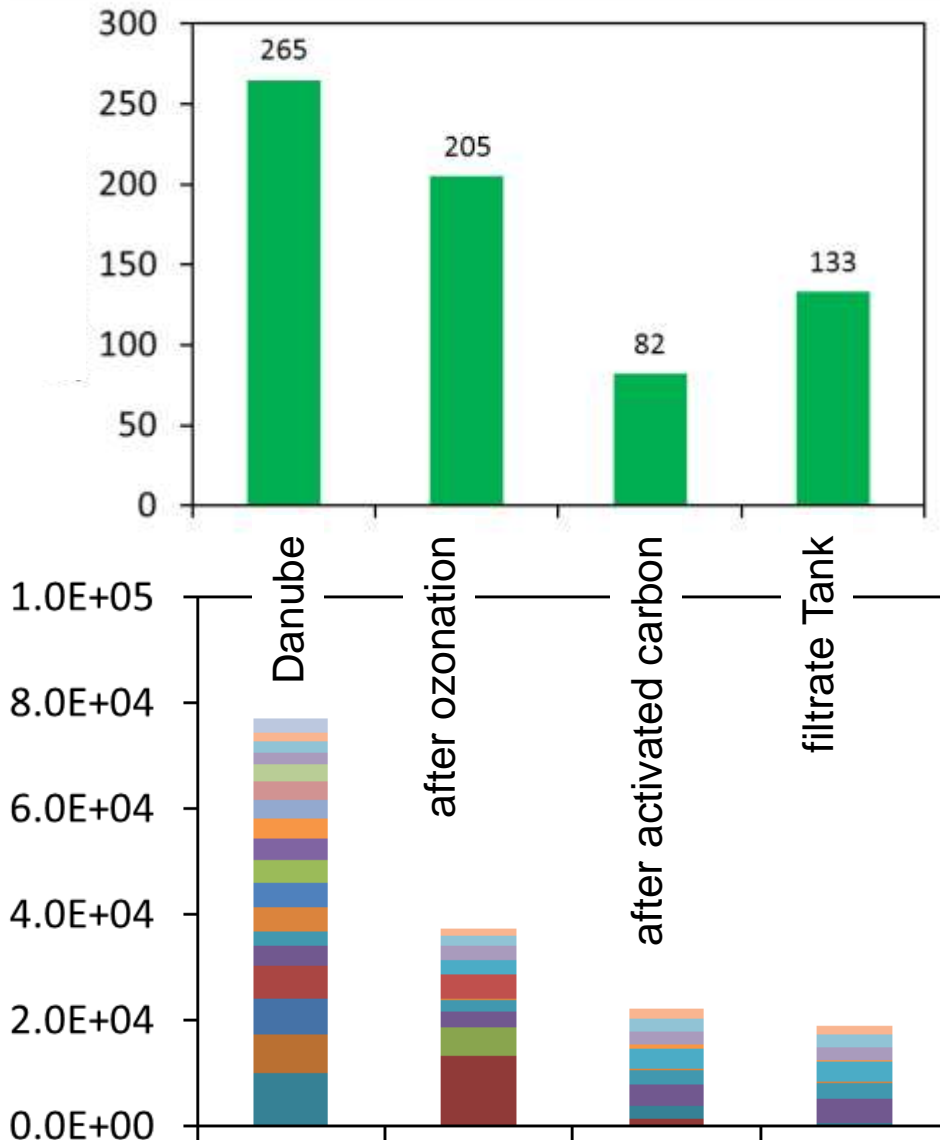
Raw water	Drinking water	behavior	Prioritization
●	—	removable	2
●	●	not or partial removable	1
—	●	transformation product	1







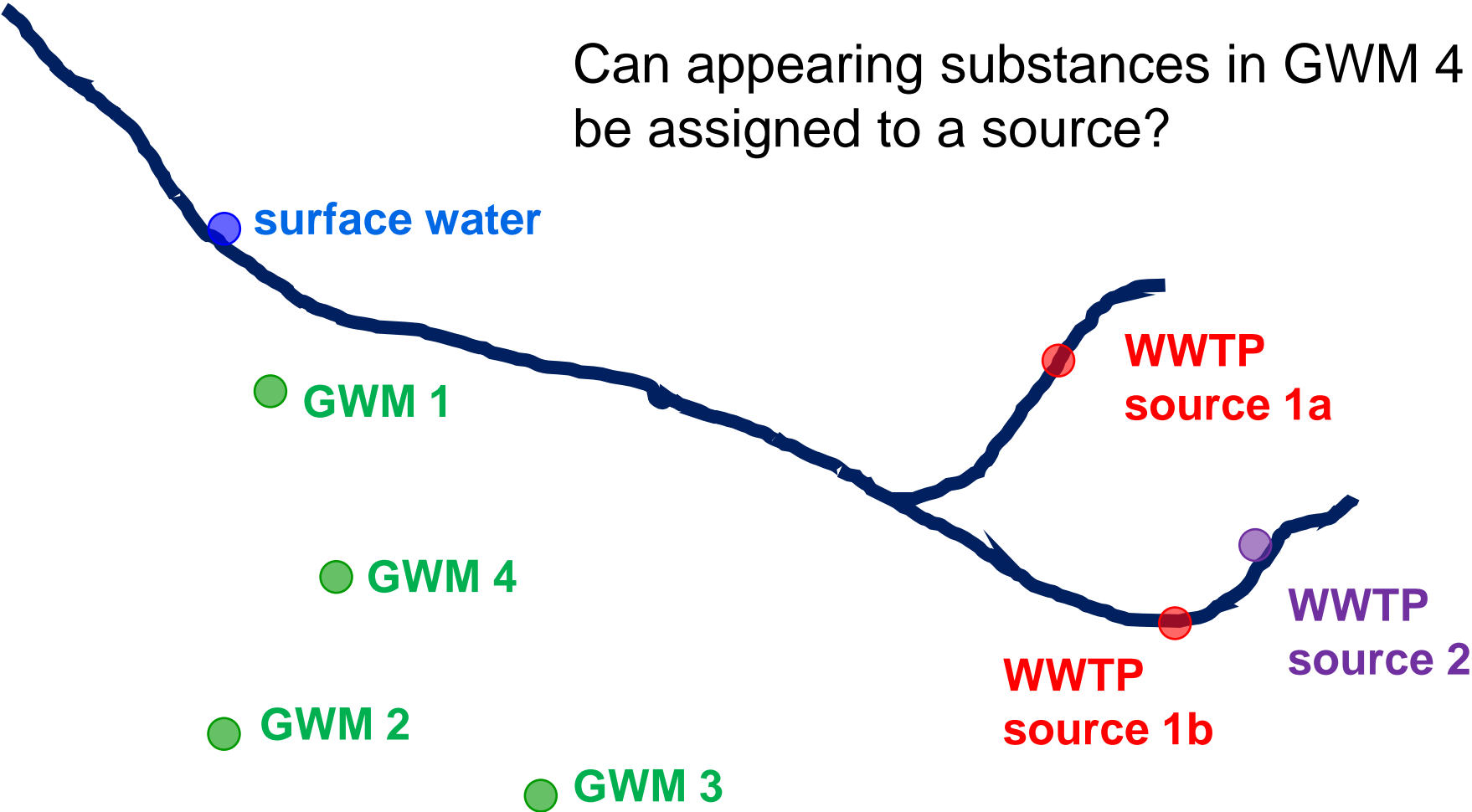
## Number of Features



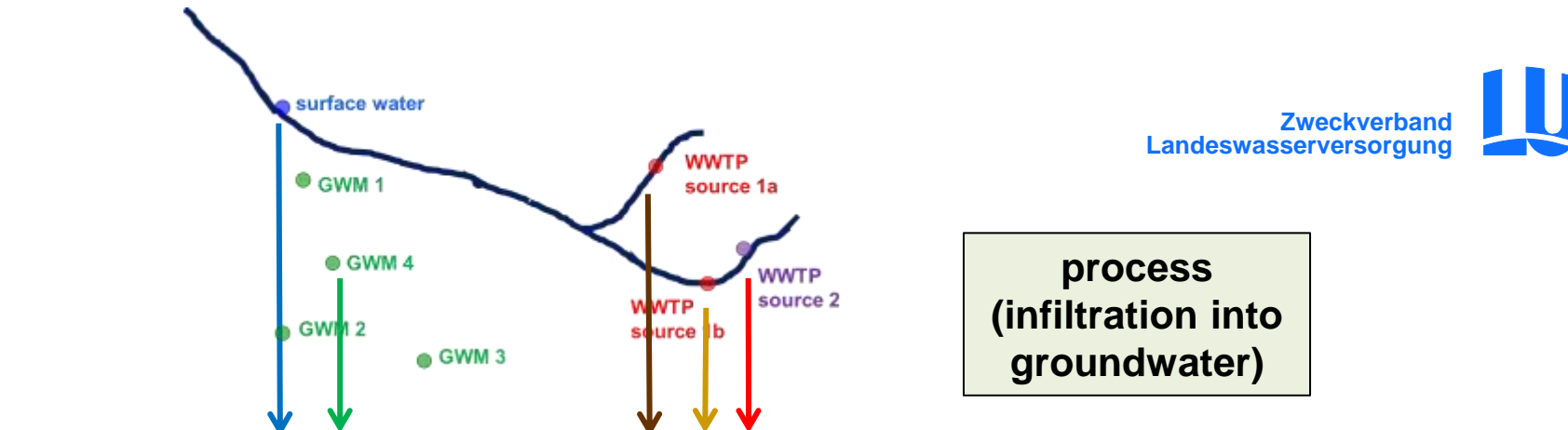
## Intensity of the TOP 25 features in the Danube

# overview of sampling points

Can appearing substances in GWM 4 be assigned to a source?



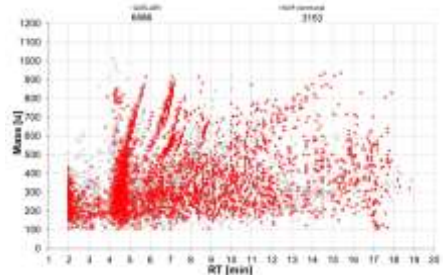
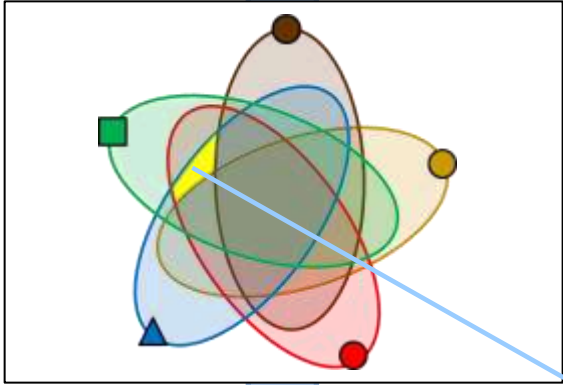
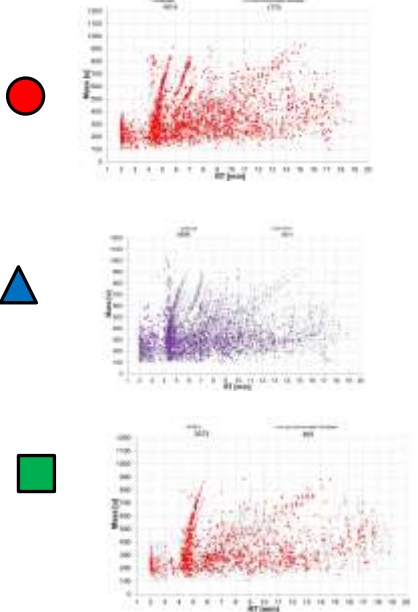
GWM: groundwater monitoring wells  
WWTP: waste water treatment plant



process  
(infiltration into  
groundwater)

LC-HRMS

Non-Target Screening



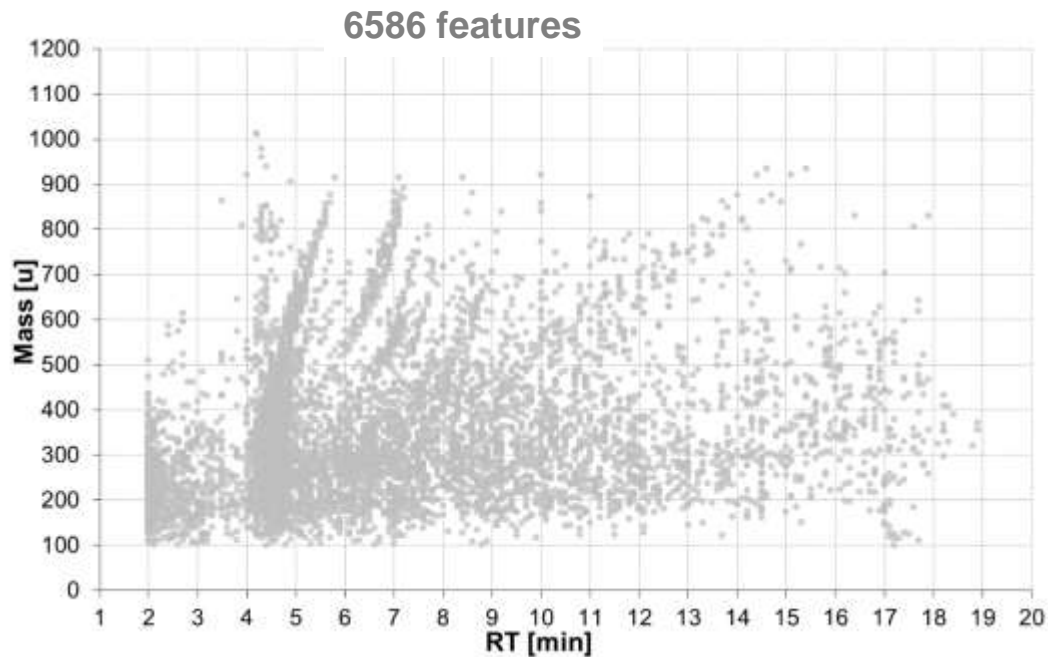
(● U ●)

$(\text{red circle} \cap \text{blue triangle} \cap \text{green square}) \setminus (\text{brown circle} \cup \text{yellow circle})$

Set theory

Identification

# Distinction of sources specific features for the sources



● source 1a

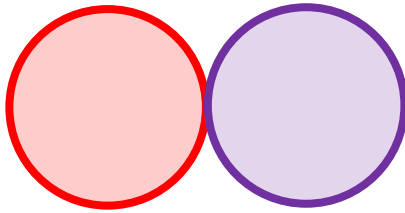
● source 1b

● source 2

Mass-RT scatterplot of all features in the sources 1a, 1b, 2



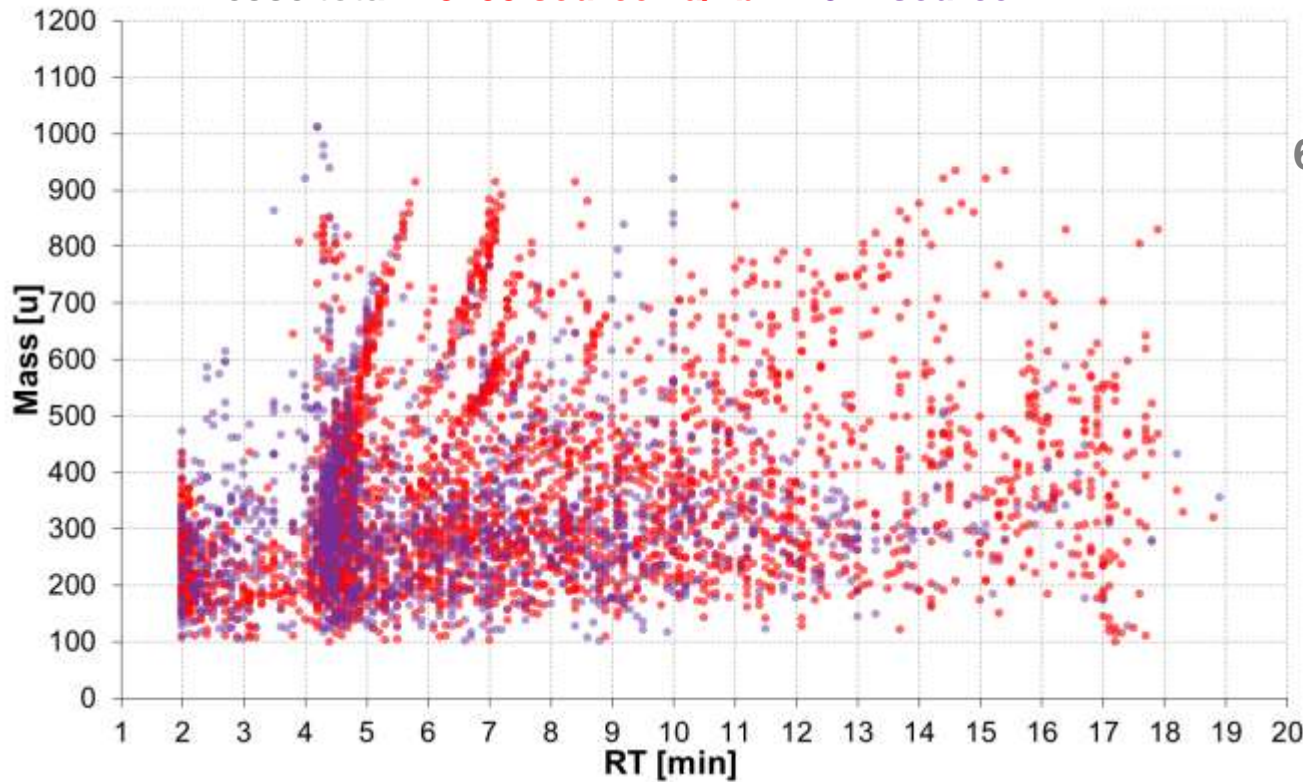
Source 1a/1b



Source 2

no interception

6586 total 3153 source 1a/1b 1611 source 2



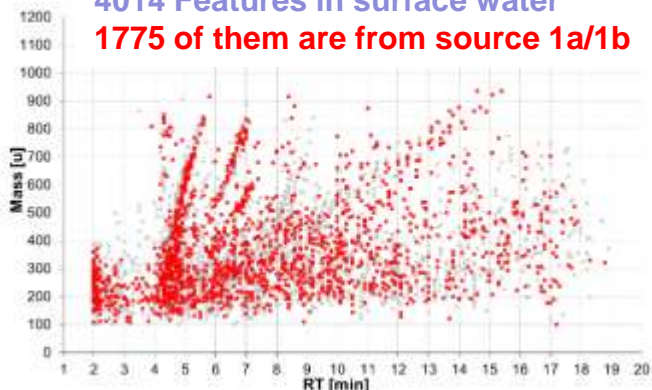
$$6586 - 3153 - 1611 = 1822$$

1822  
common features in all  
sources

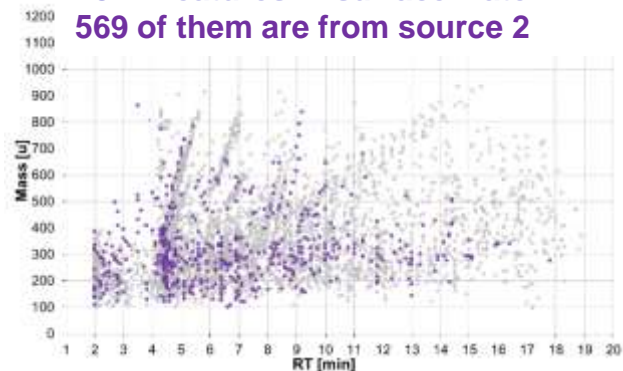
only from source 1a and 1b

only from source 2

4014 Features in surface water  
1775 of them are from source 1a/1b

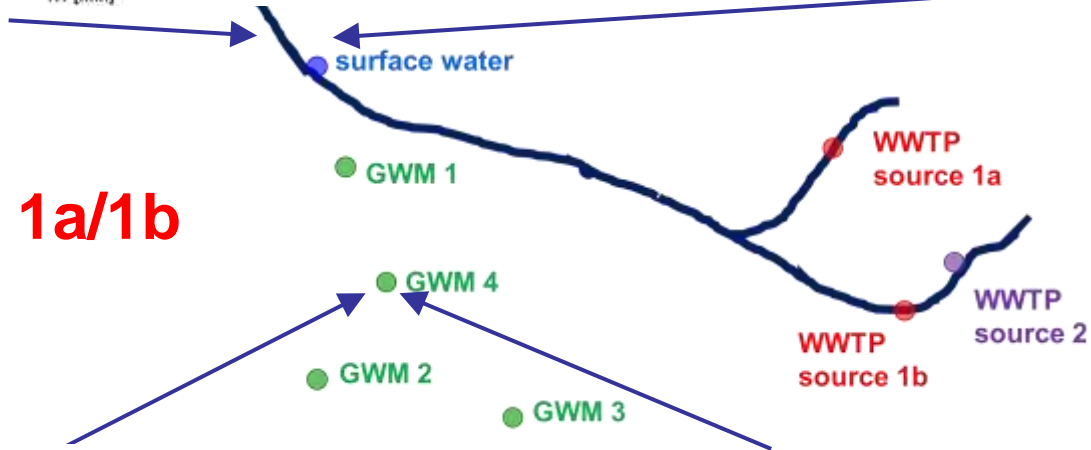


4014 Features in surface water  
569 of them are from source 2

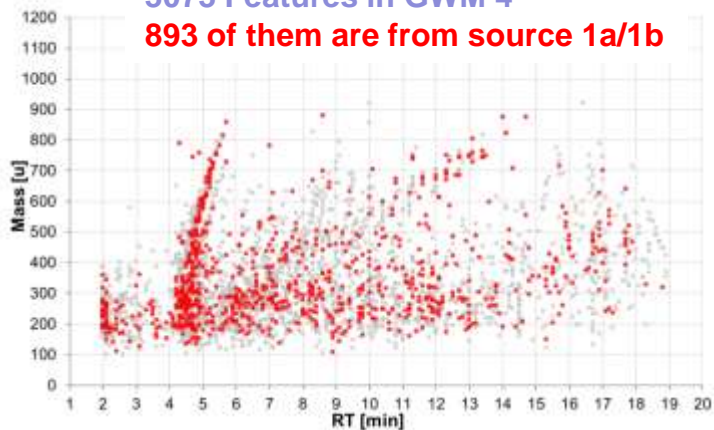


Source 1a/1b

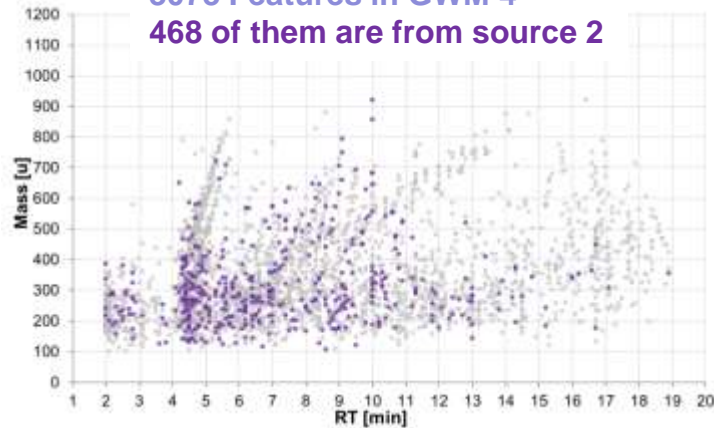
Source 2



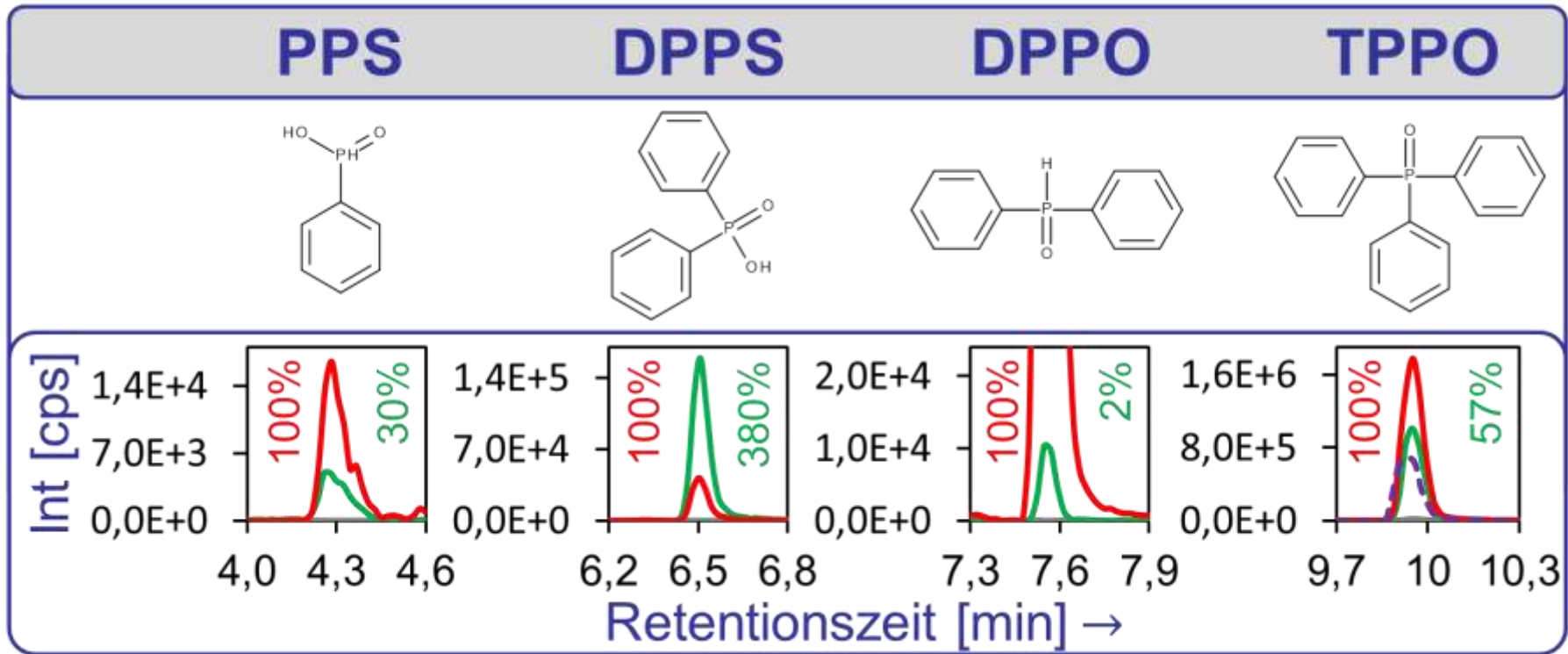
3073 Features in GWM 4  
893 of them are from source 1a/1b



3073 Features in GWM 4  
468 of them are from source 2



# Identification of some Features from source 2 present in GWM 4



Source 2

GWM 4

Blank

standard: 5  $\mu\text{g}\cdot\text{L}^{-1}$

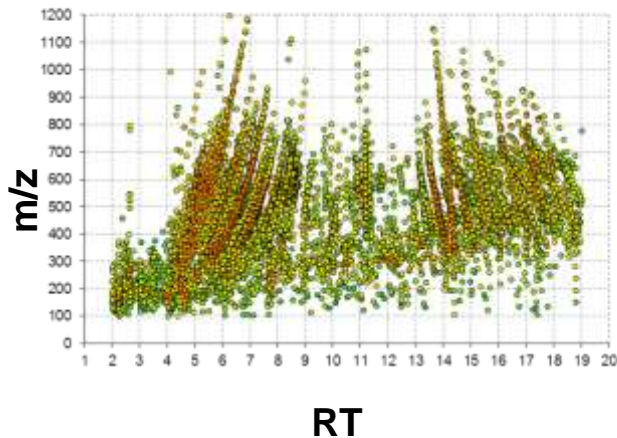
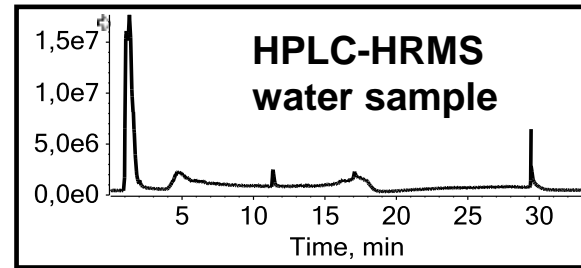
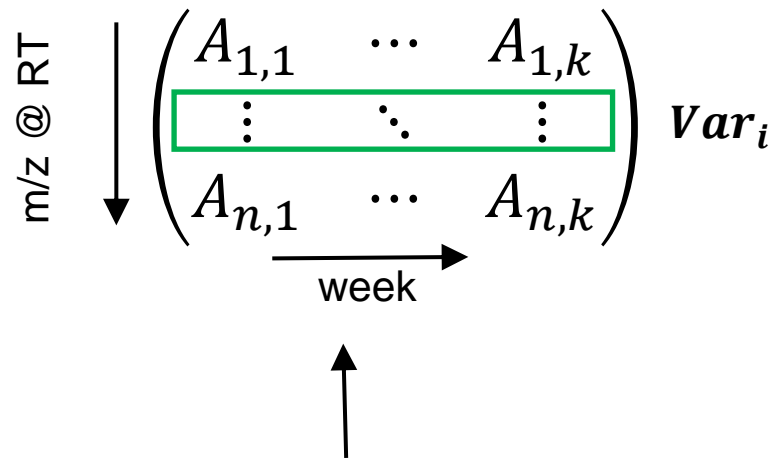
TPPO Triphenylphosphineoxid

Talk: Uwe Kunkel, Friday

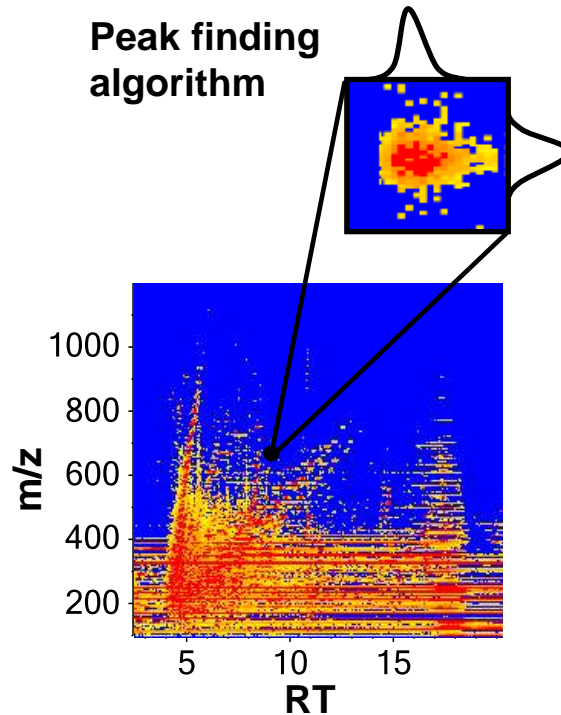
Quarternary triphenylphosphonium compounds

# Non-Target Screening

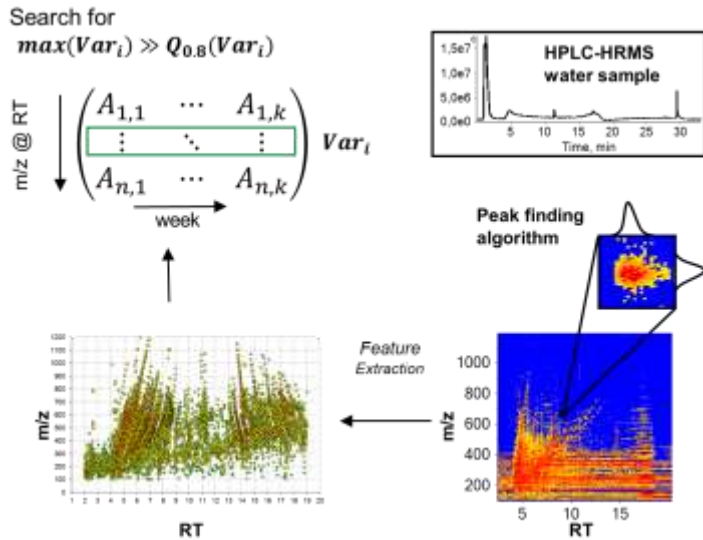
Search for  
 $\max(Var_i) \gg Q_{0.8}(Var_i)$



Feature  
Extraction



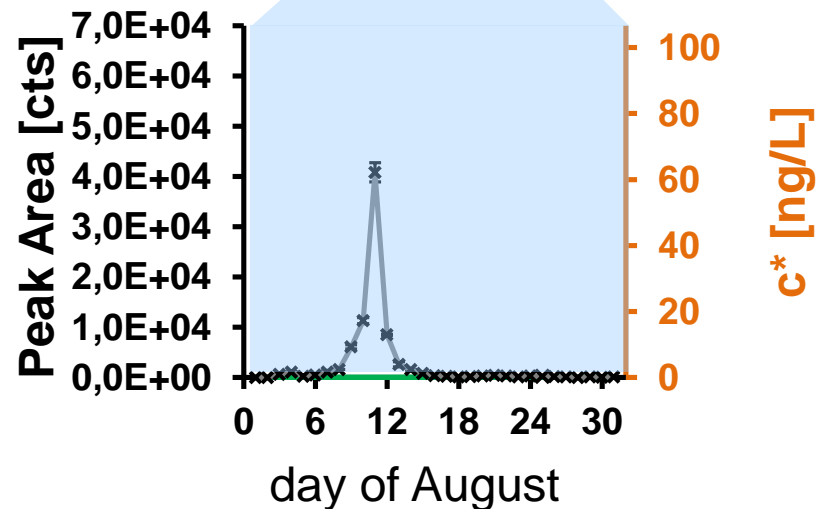
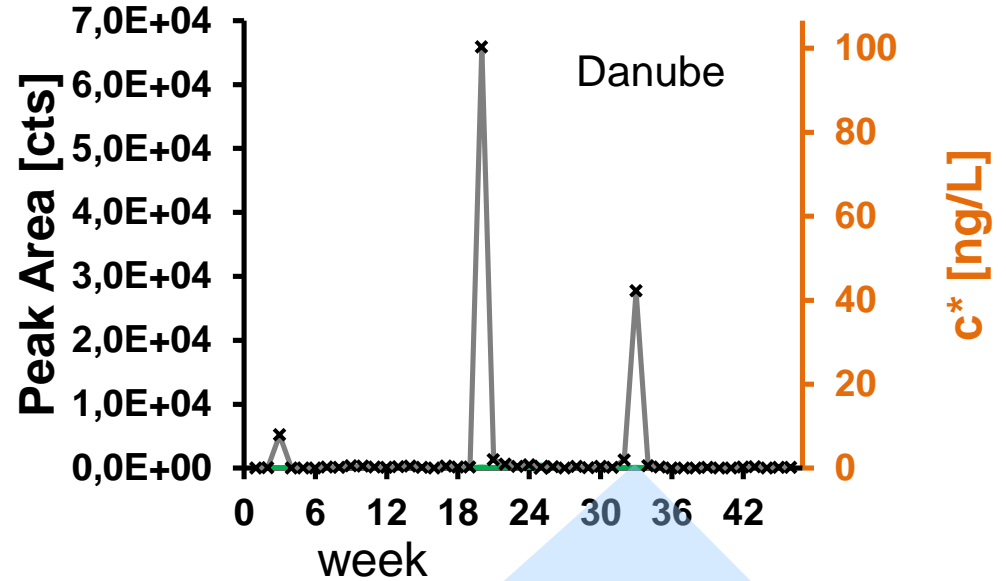
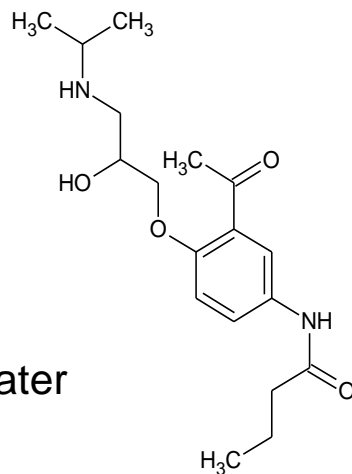
# Non-Target Screening



Acebutolol

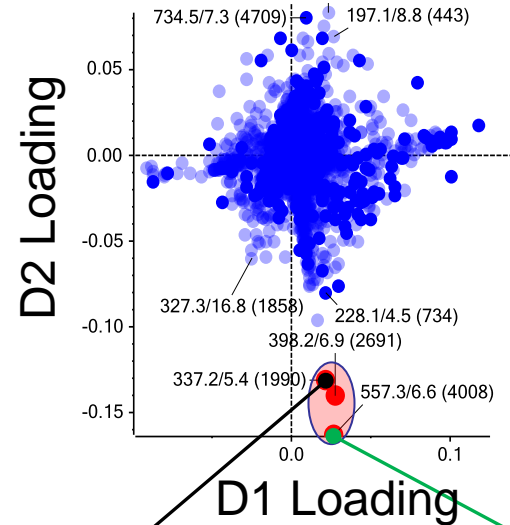
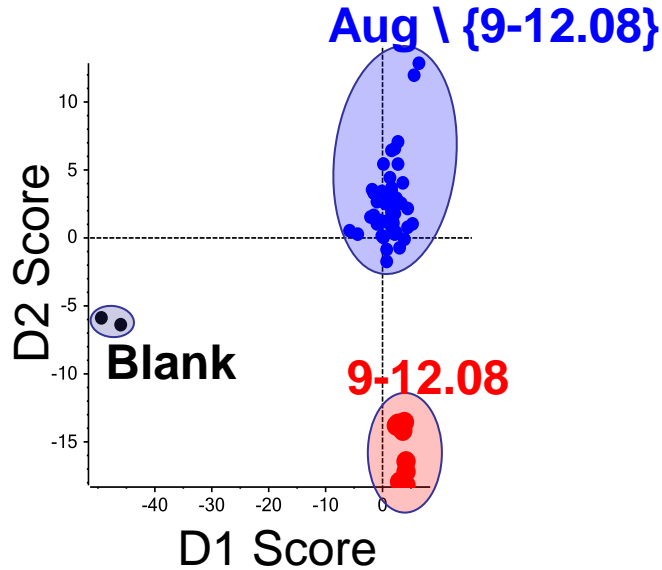
337.212@5.4min

Not found in drinking water





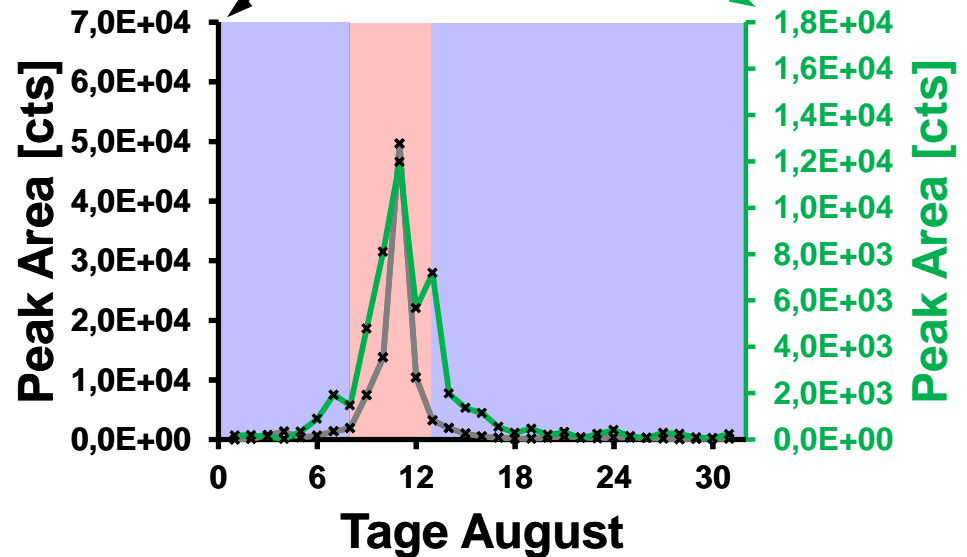
# PCA-DA

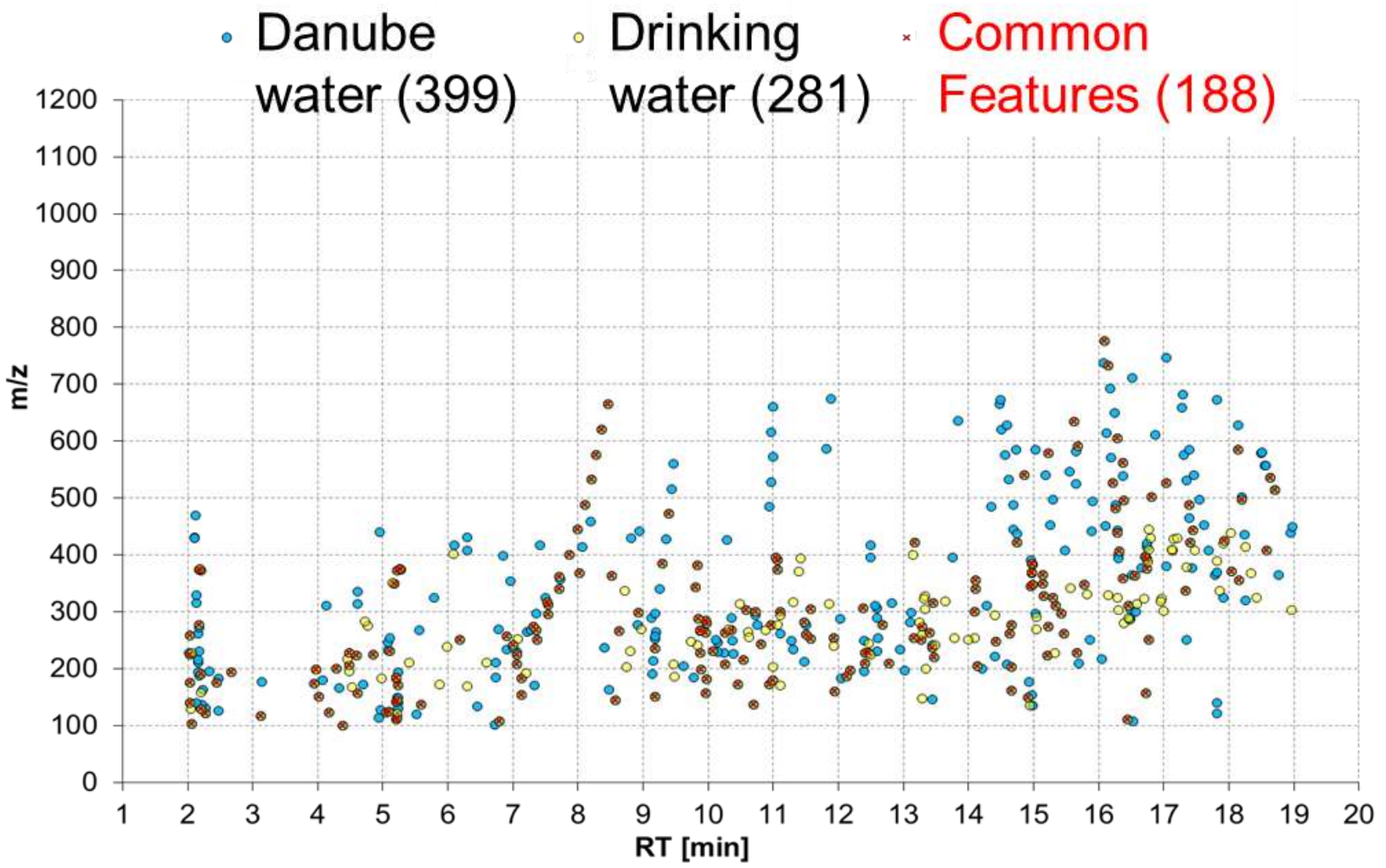


Principal component analysis and discriminant analysis (PCA-DA)

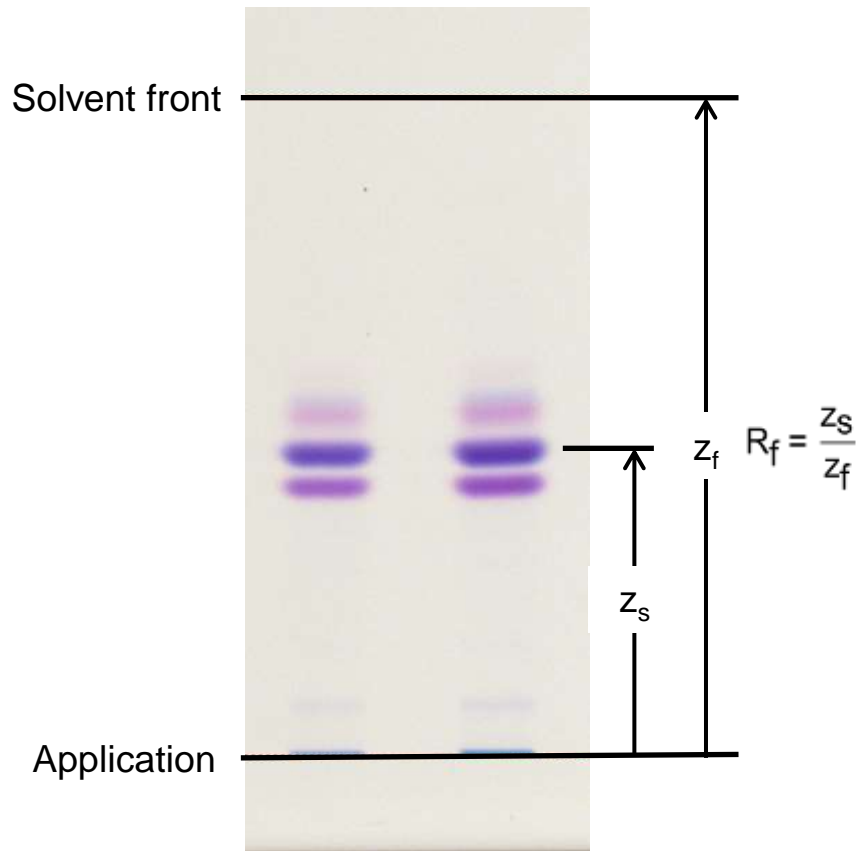
337.212@5.4min

557.255@6.6min





# Thin-layer chromatography (TLC)



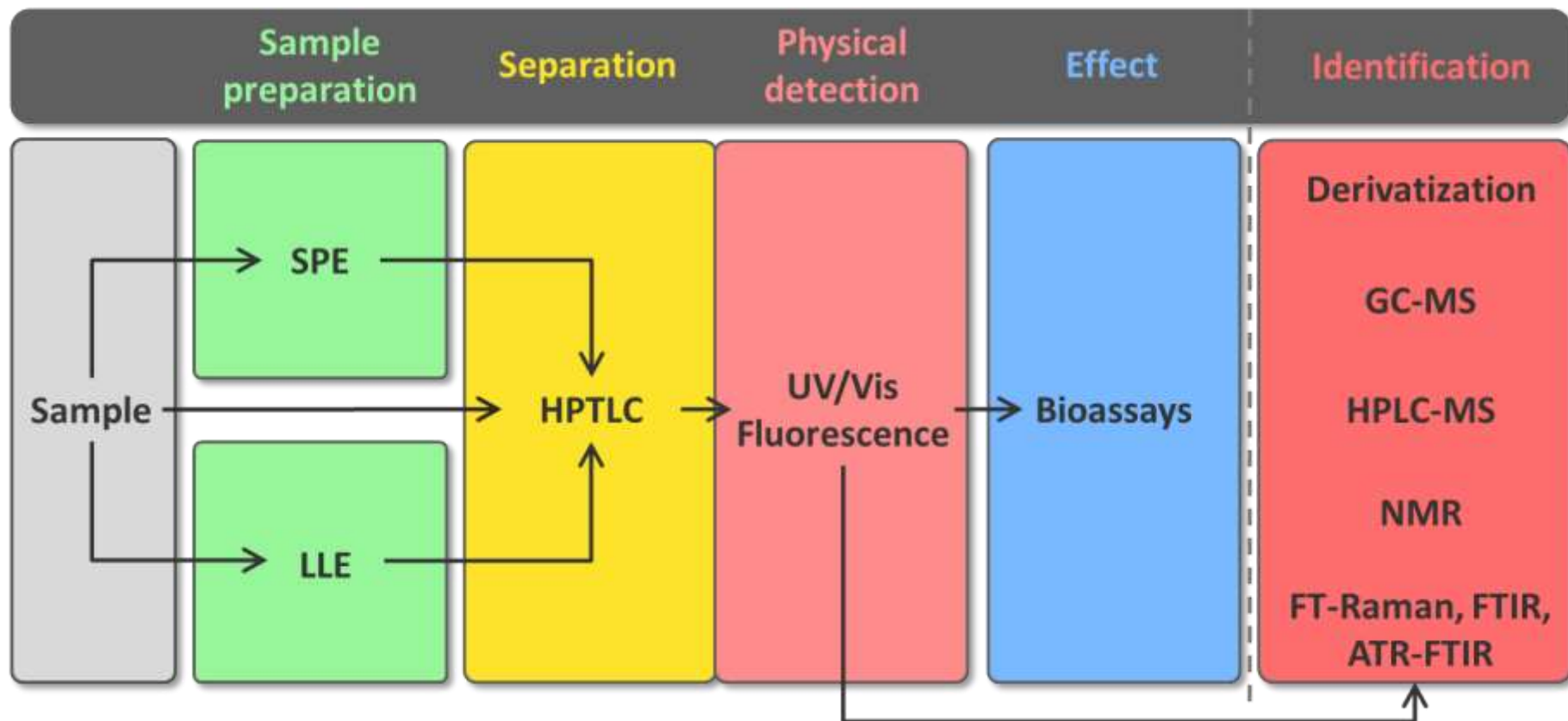
## Advantages:

- Substances which remain on the application zone were gathered
- Solvents have no influence on the biological tests
- Large choice of solvents

## Disadvantages:

- Lower separation efficiency as e.g. HPLC
- Not fully automated
- Limited separation distance

# Overview of effect-directed analysis (EDA) with thin-layer chromatography (TLC)



# Procedure of EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay

application



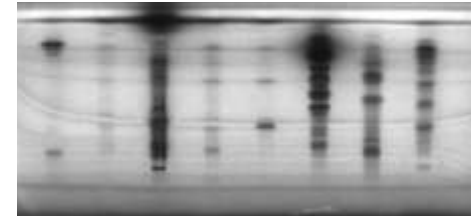
separation



immersion

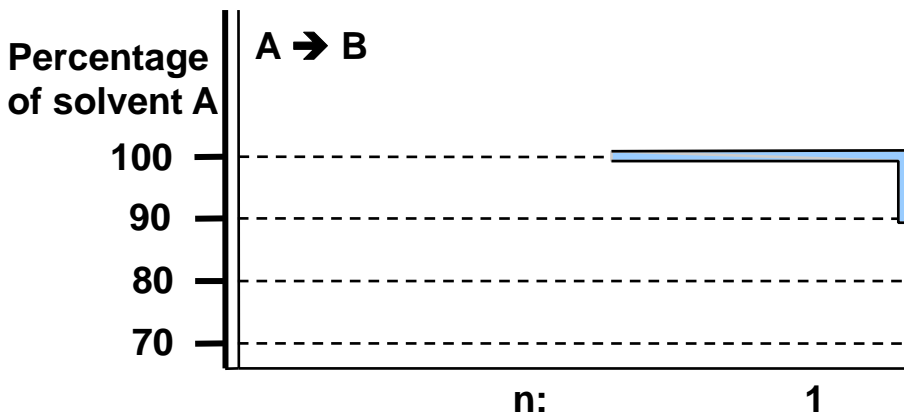
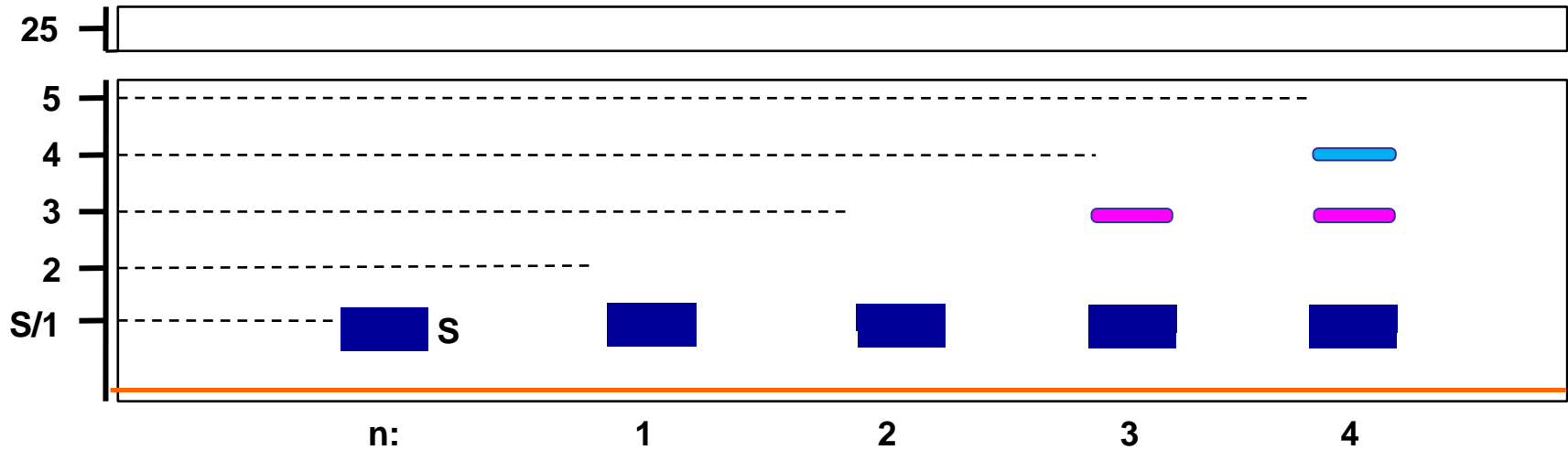


detection



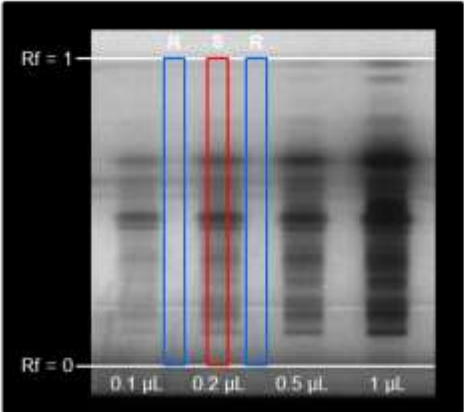


# Procedure of HPTLC-analysis with automated multiple development (AMD)

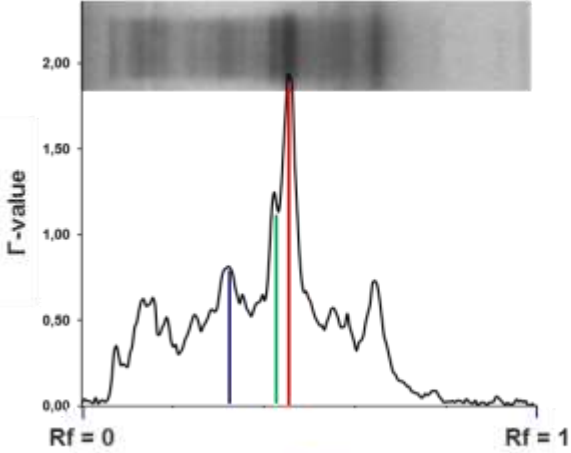


# Reciprocal iso-inhibition volume (RIV)

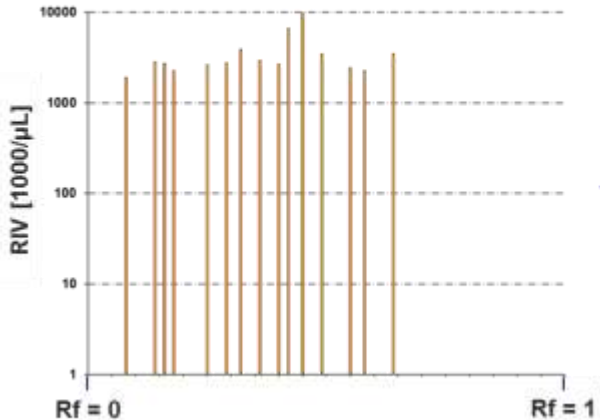
A) Bioluminescence detection



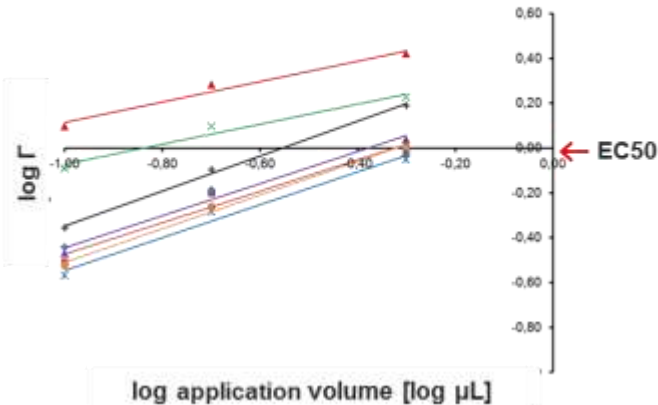
B)  $\Gamma$  value chromatogram



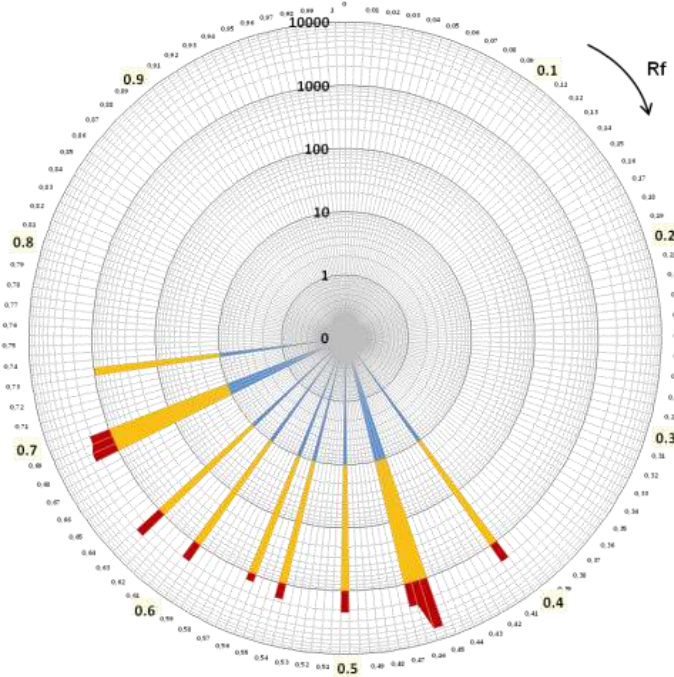
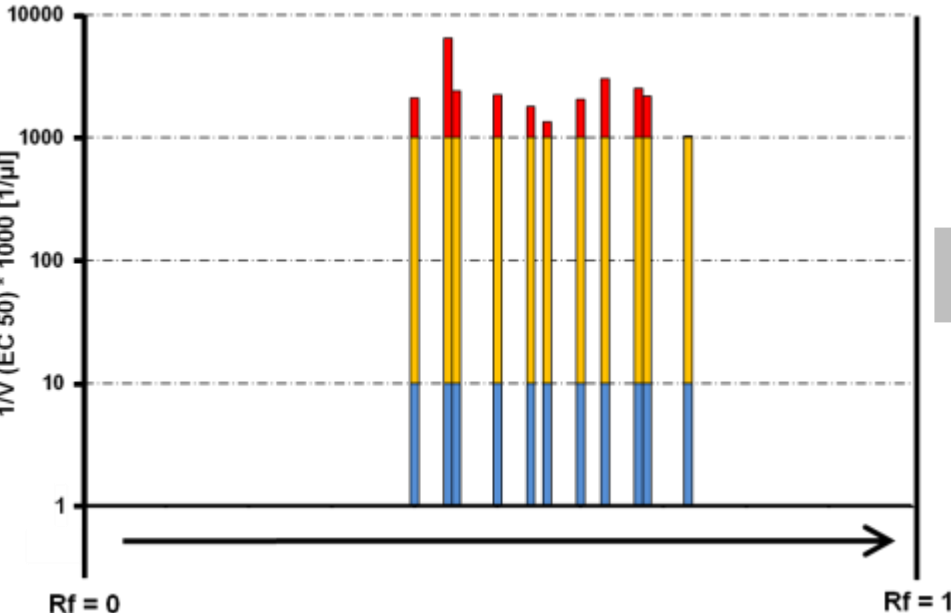
D) Calculation of the reciprocal iso-inhibition volume (RIV)



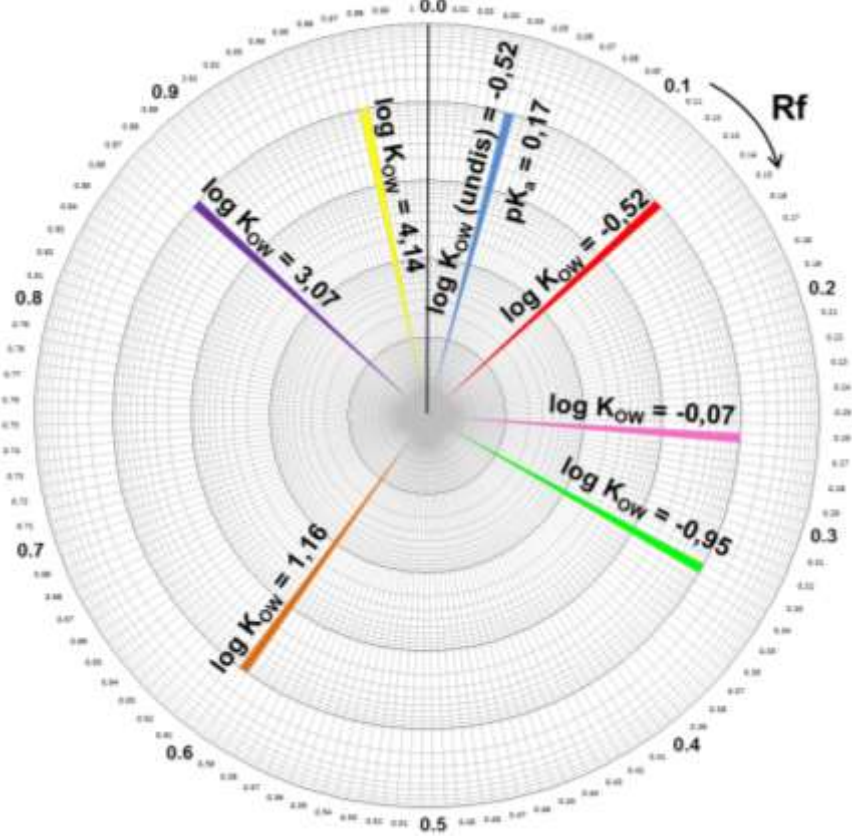
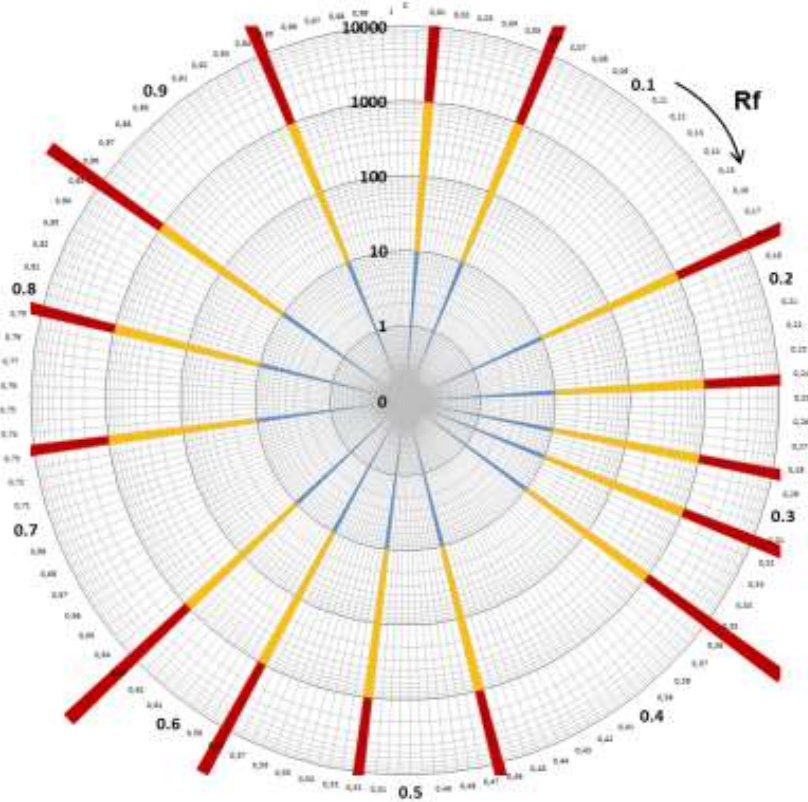
C) Dose-response relationship for the separated bands



# Reciprocal iso-inhibition volume (RIV) Polar diagram

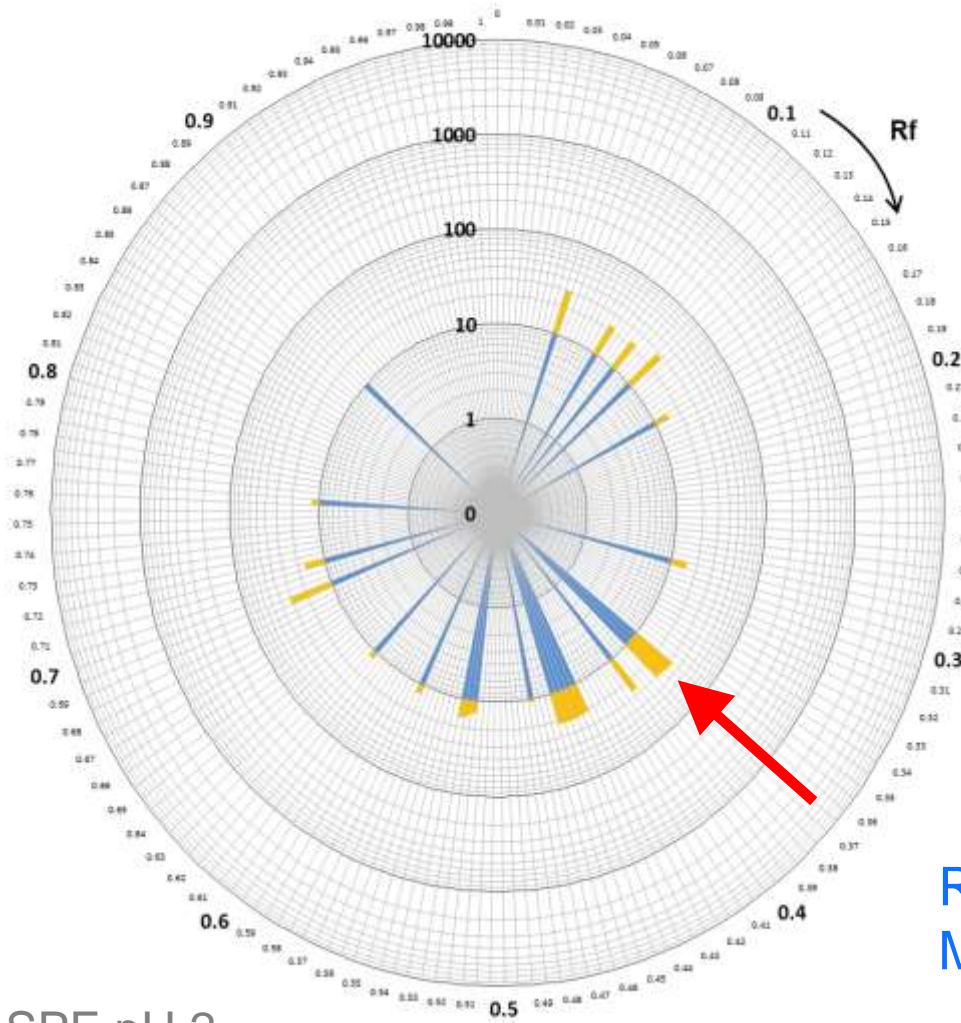


# Estimation of polarity



- 1 - Naphthalene sulfonic acid
- Theobromine
- Caffeine
- Thiourea
- Acetanilide
- Benzanilide
- Methyl yellow

# EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay

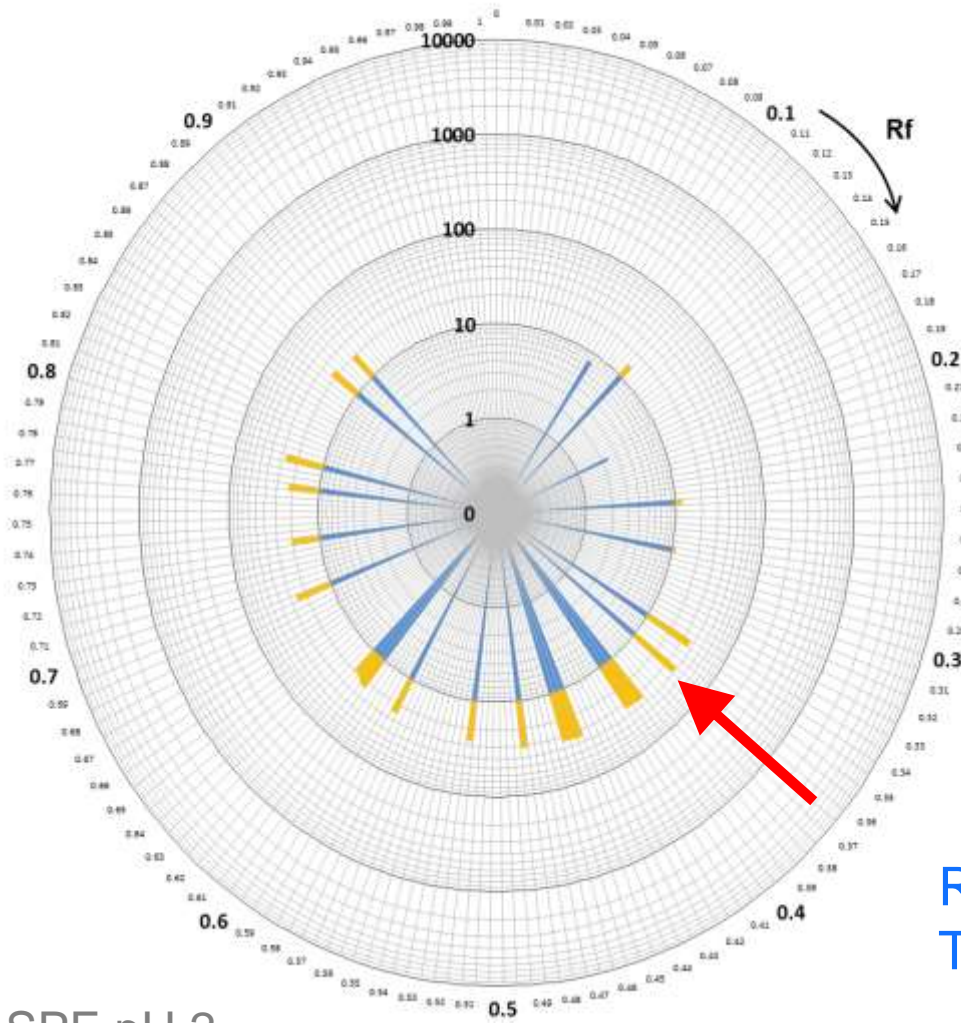


River Danube  
Monday, March 7<sup>th</sup>, 2016

SPE pH 2



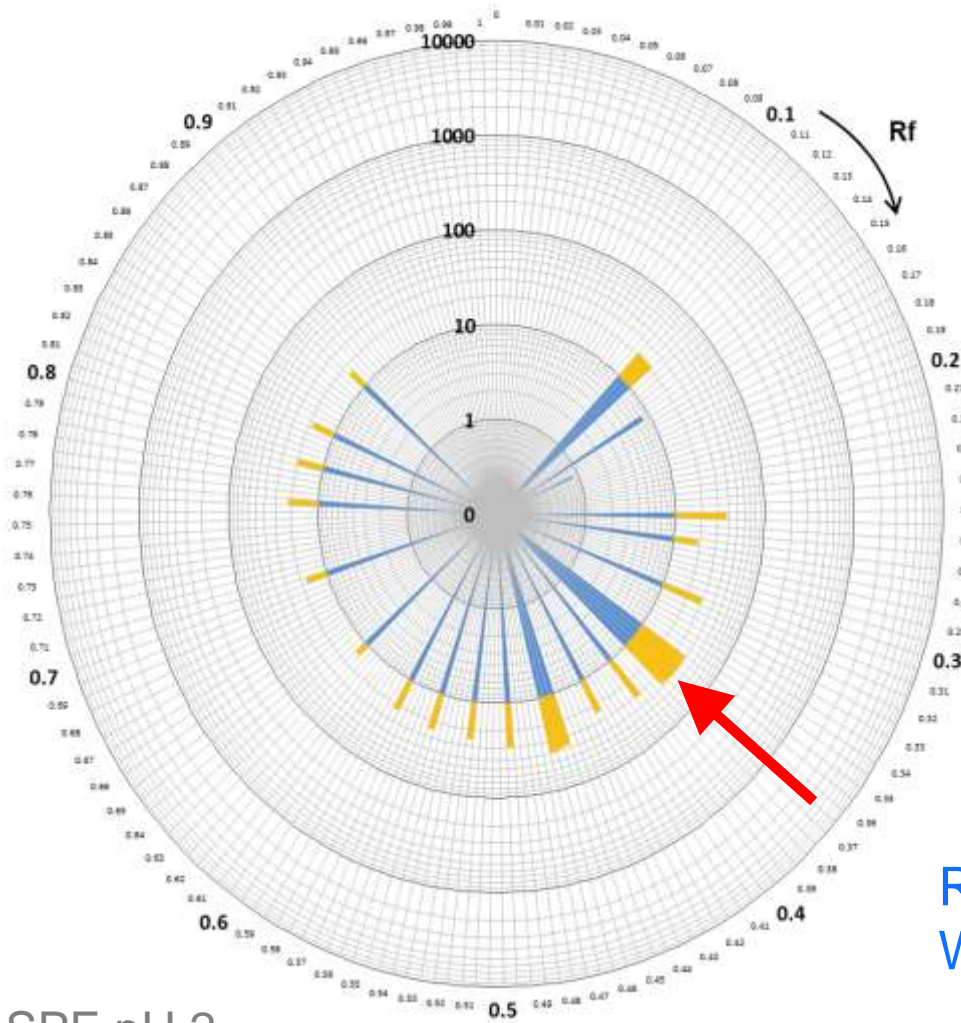
# EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay



River Danube  
Tuesday, March 8<sup>th</sup>, 2016

SPE pH 2

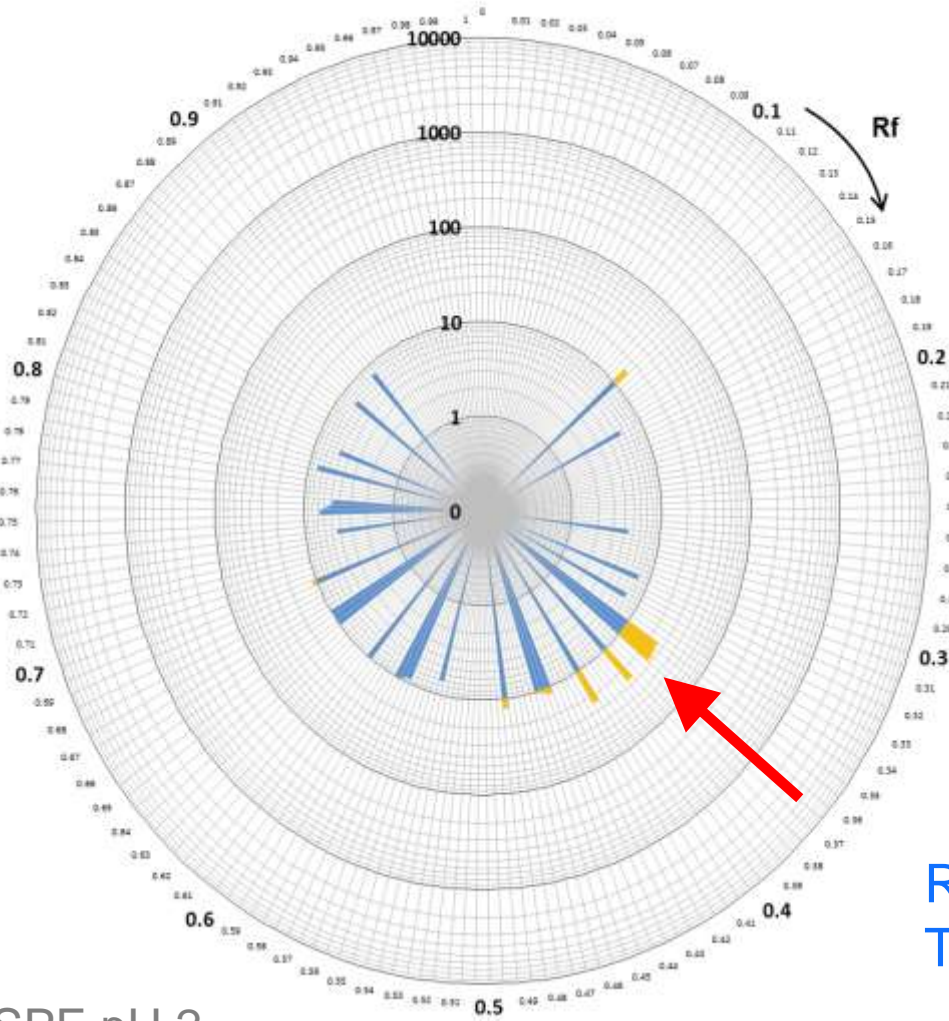
# EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay



River Danube  
Wednesday, March 9<sup>th</sup>, 2016

SPE pH 2

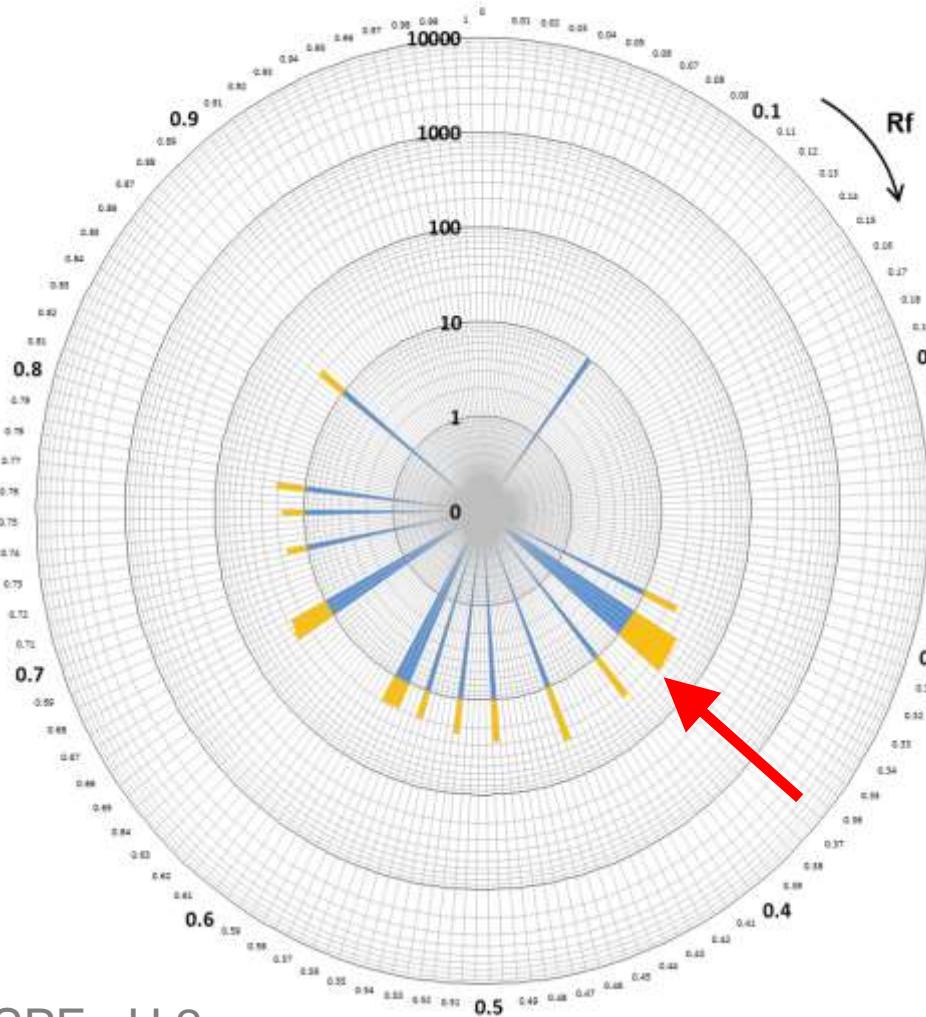
# EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay



River Danube  
Thursday, March 10<sup>th</sup>, 2016

SPE pH 2

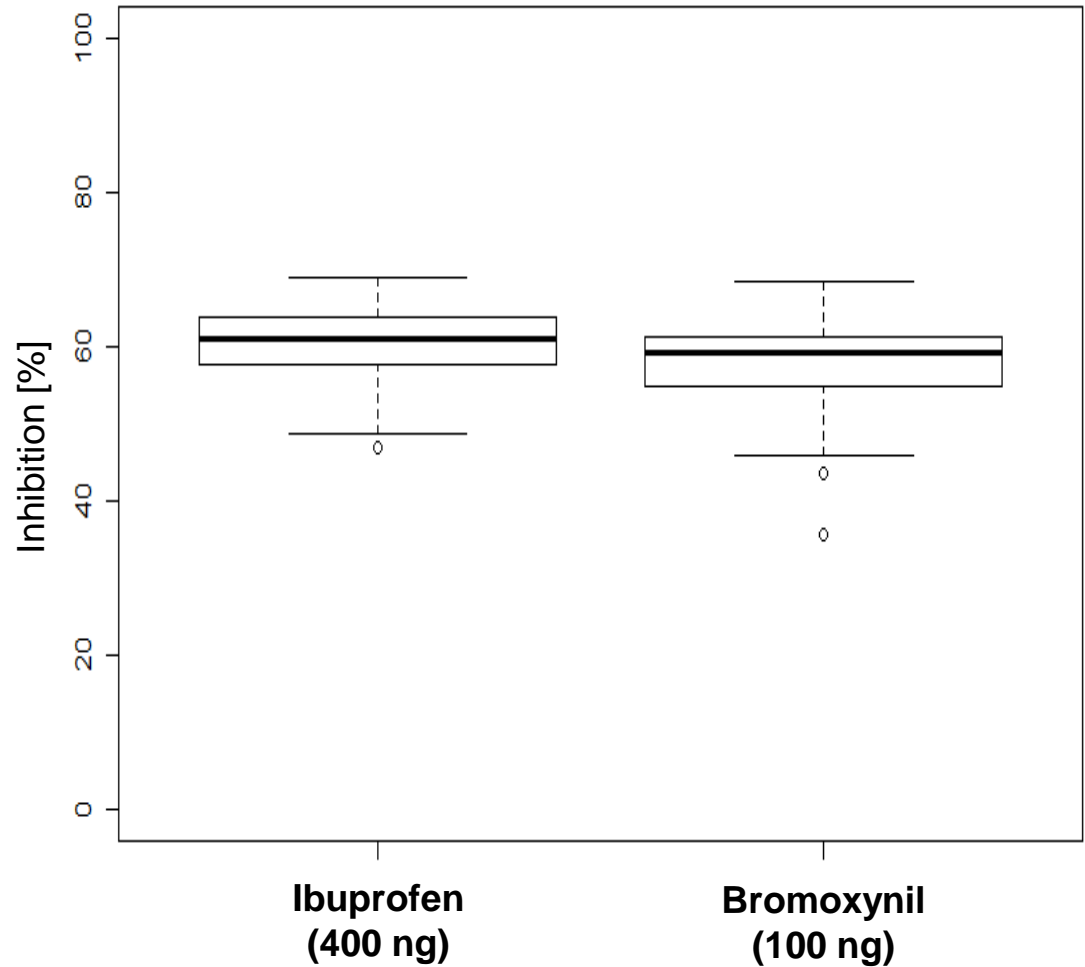
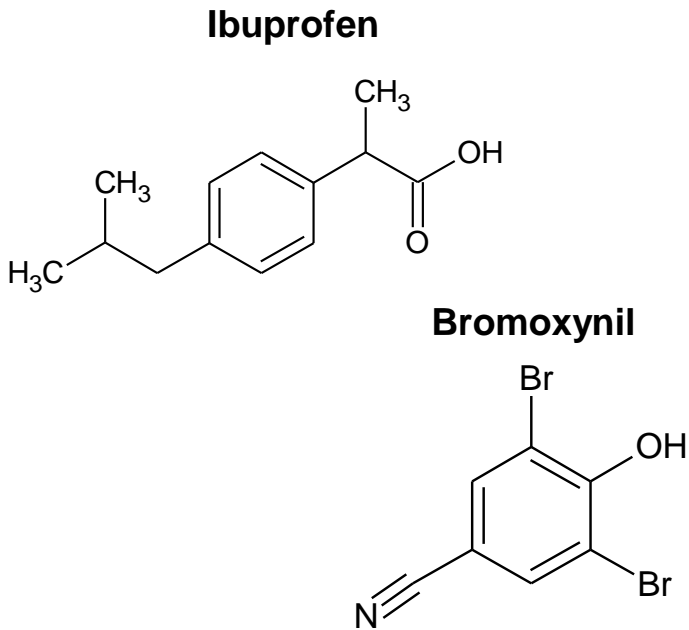
# EDA/HPTLC-analysis with *Aliivibrio fischeri* inhibition assay



River Danube  
Friday, March 11<sup>th</sup>, 2016

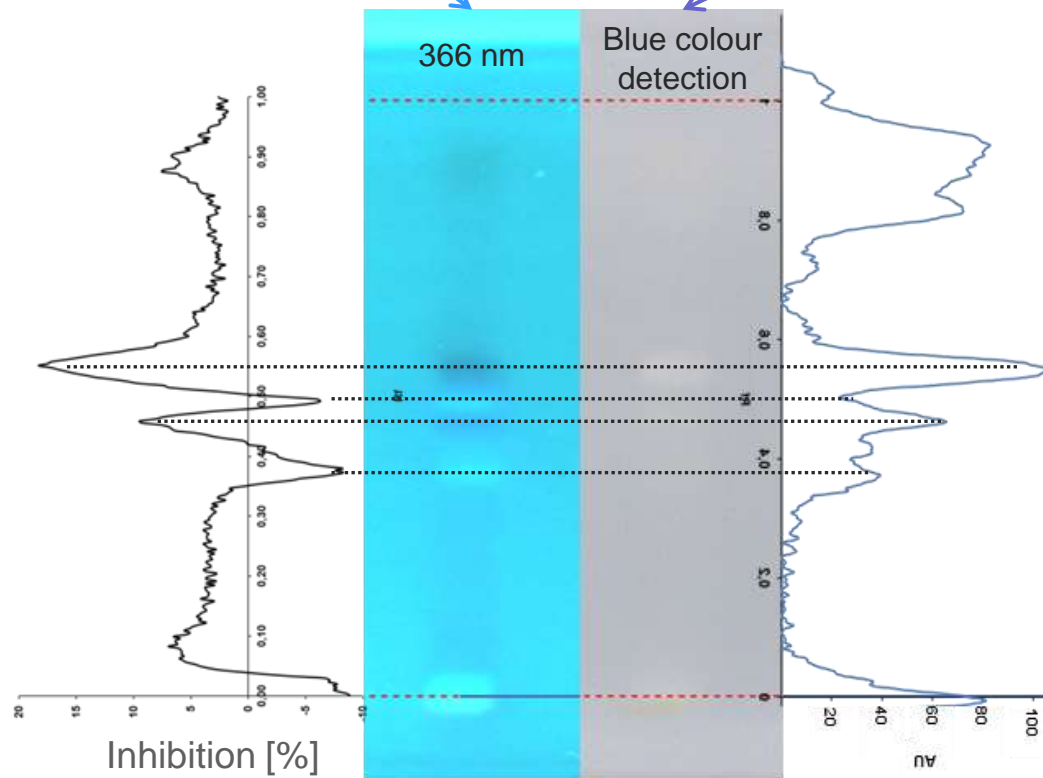
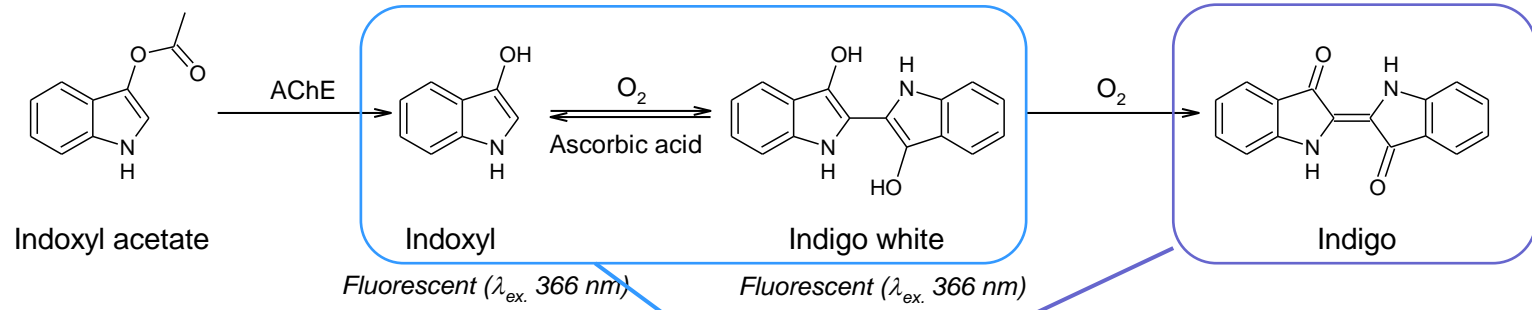
SPE pH 2

# Repeatability of *Aliivibrio fischeri* inhibition (N = 80)

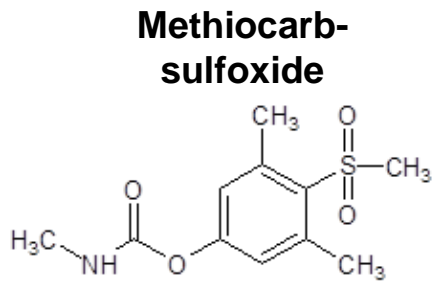




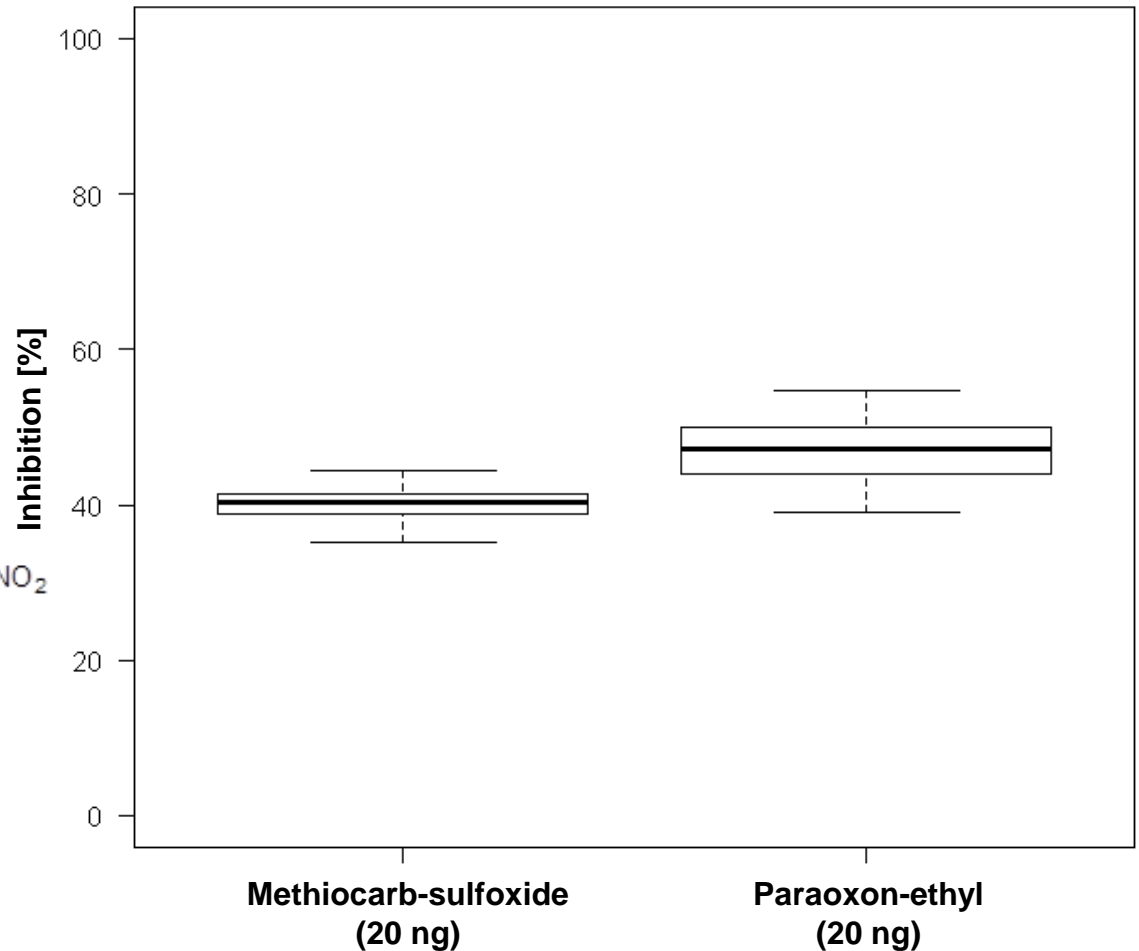
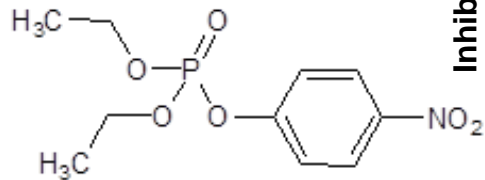
# Detection of the inhibition of AChE



# Repeatability of Acetylcholinesterase inhibition (N = 40)



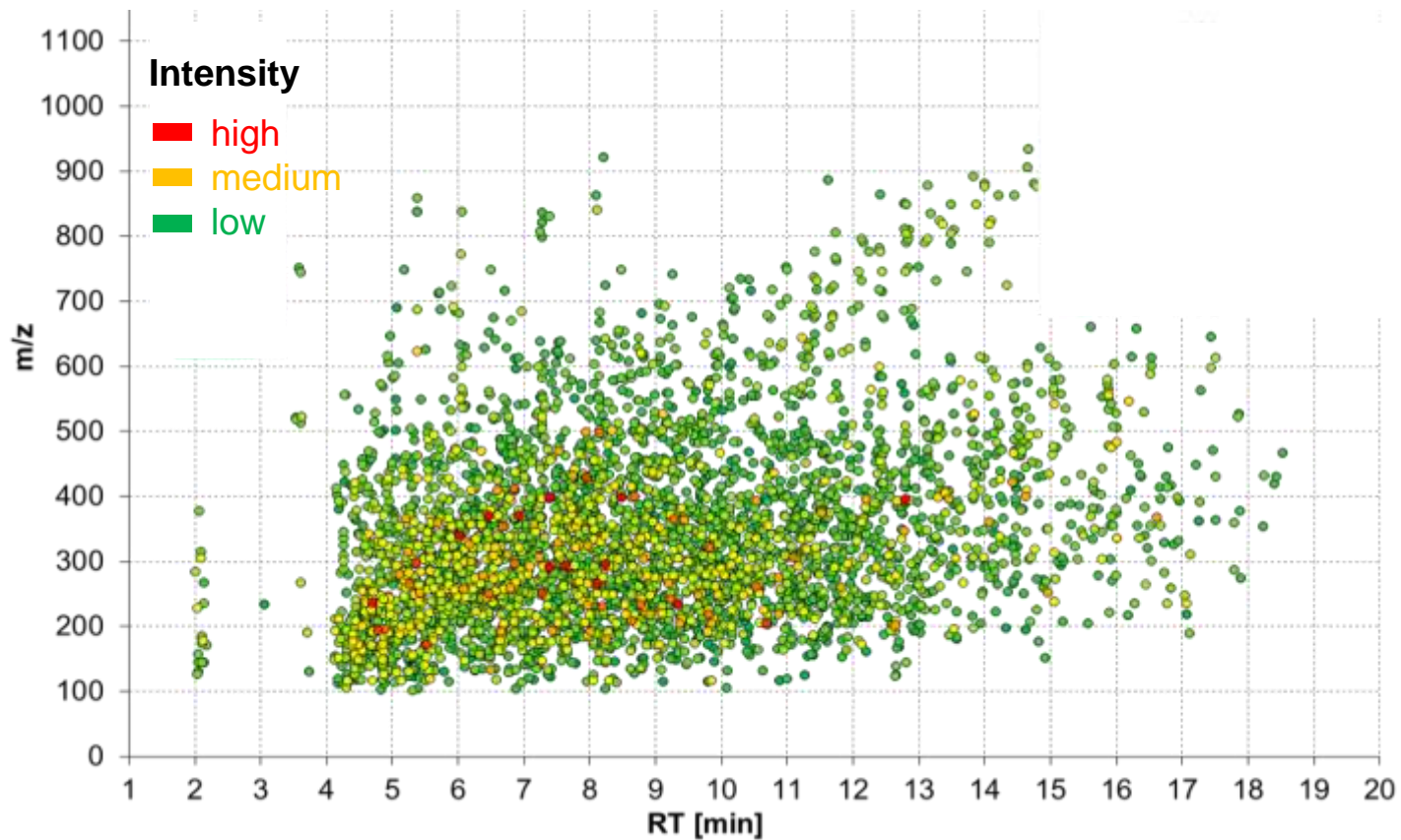
**Paraoxon-ethyl**



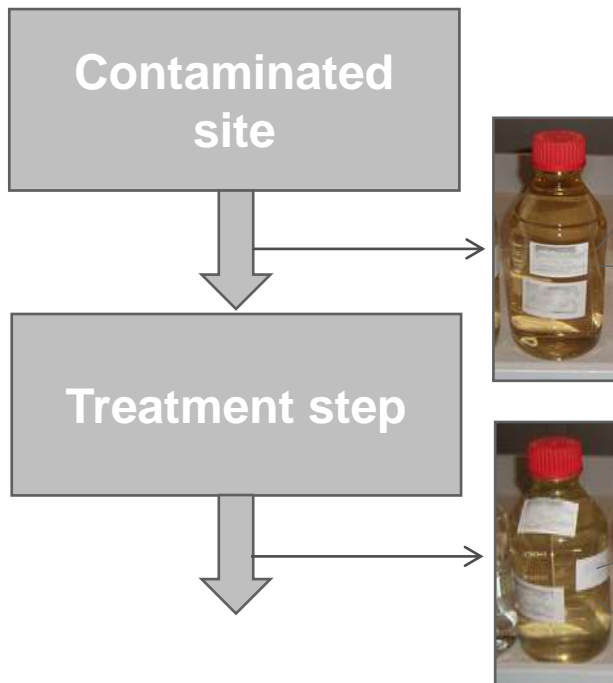
# Effluent of the industrial sewage treatment plant

Luminescence  
bacteria

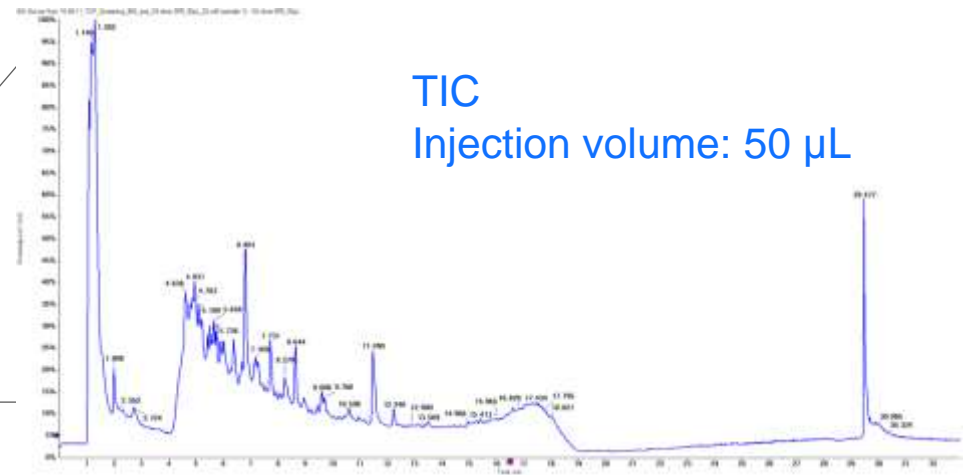
AChE



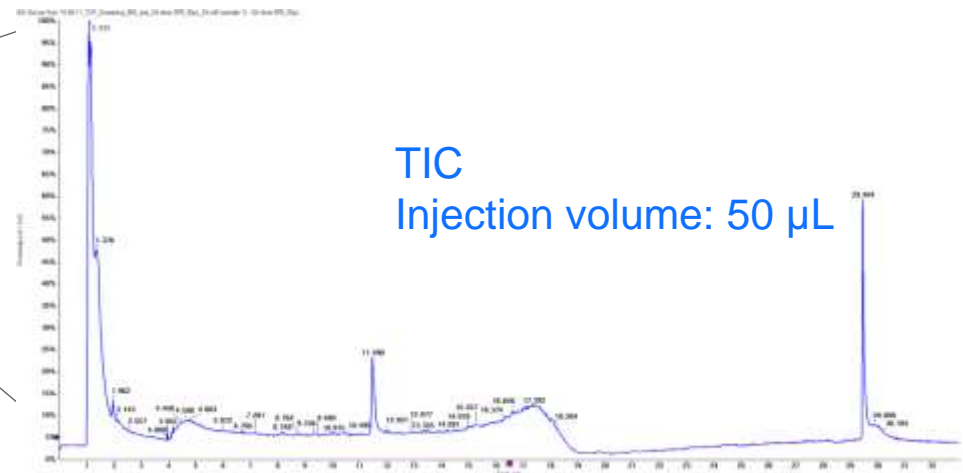
# Treatment of wastewater from a contaminated site



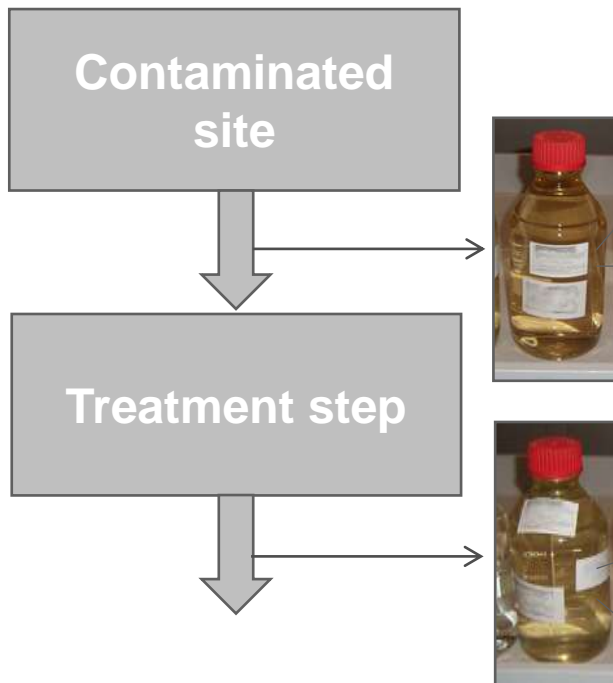
original sample before treatment



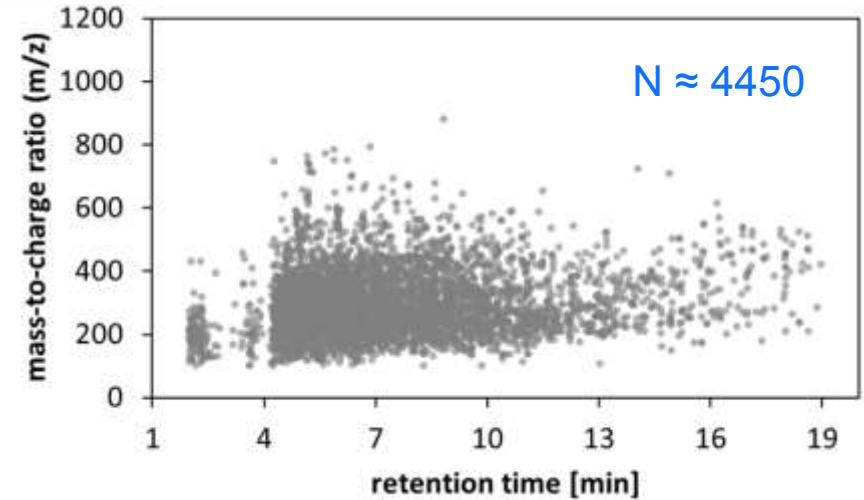
original sample after treatment



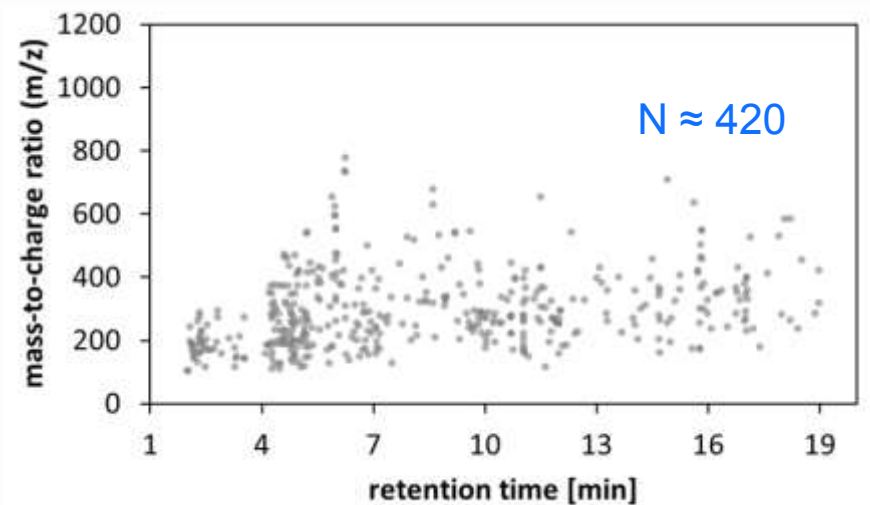
# Treatment of wastewater from a contaminated site



Mass-RT scatterplot original sample before treatment

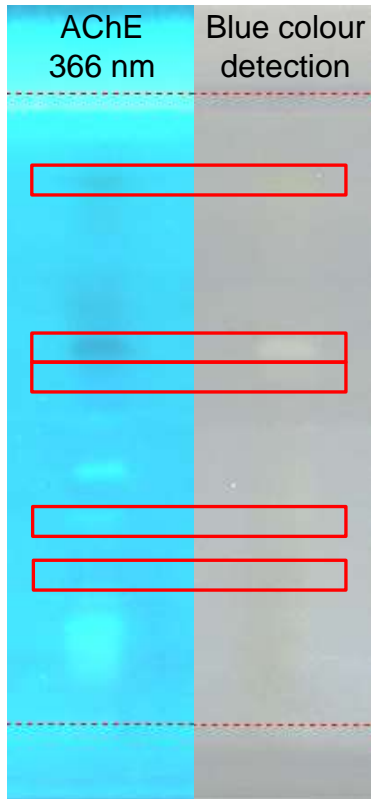


Mass-RT scatterplot original sample after treatment





# Extraction for HPLC-HRMS



Effect-directed analysis



TLC-MS Interface

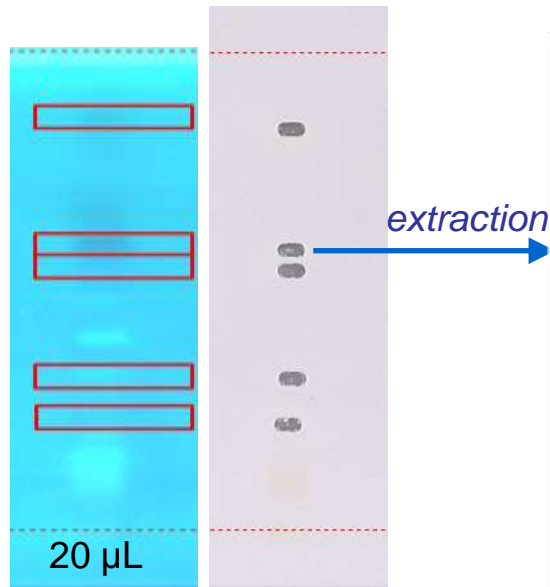


HPLC-HRMS

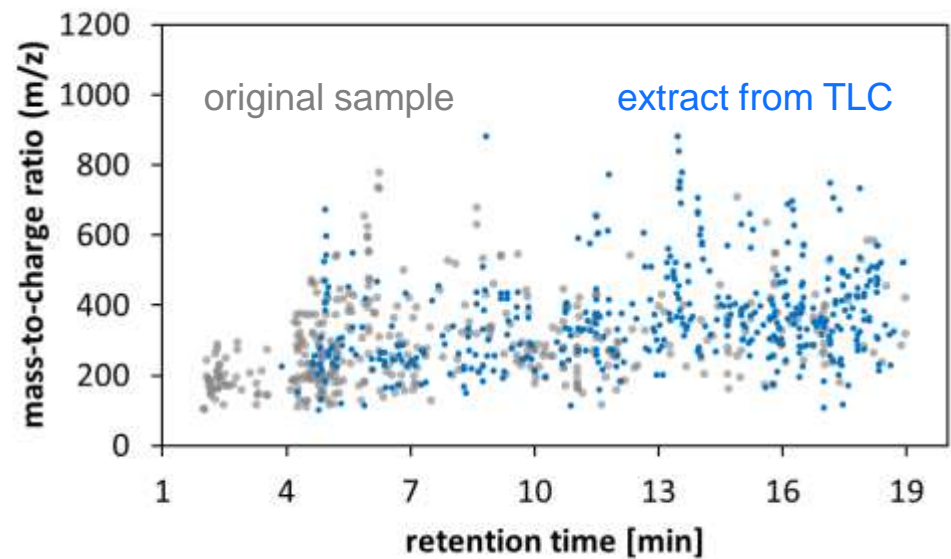
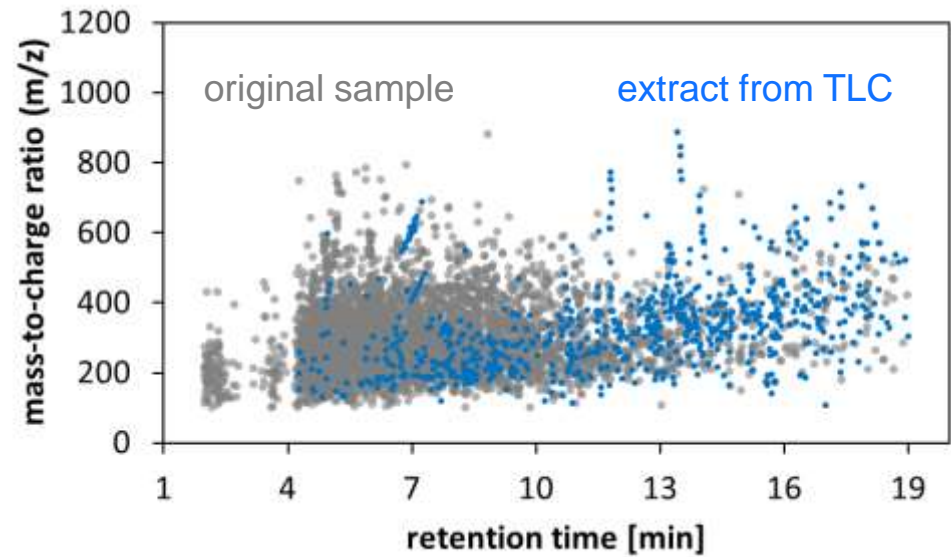
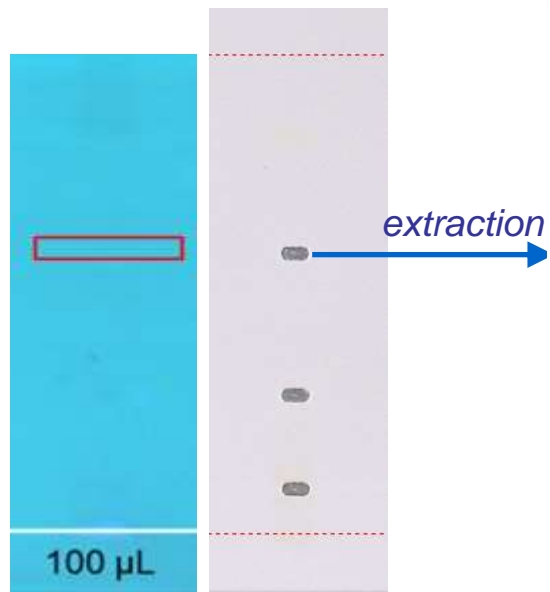
same chromatography  
for water sample  
and TLC extract

# Extraction for HPLC-HRMS

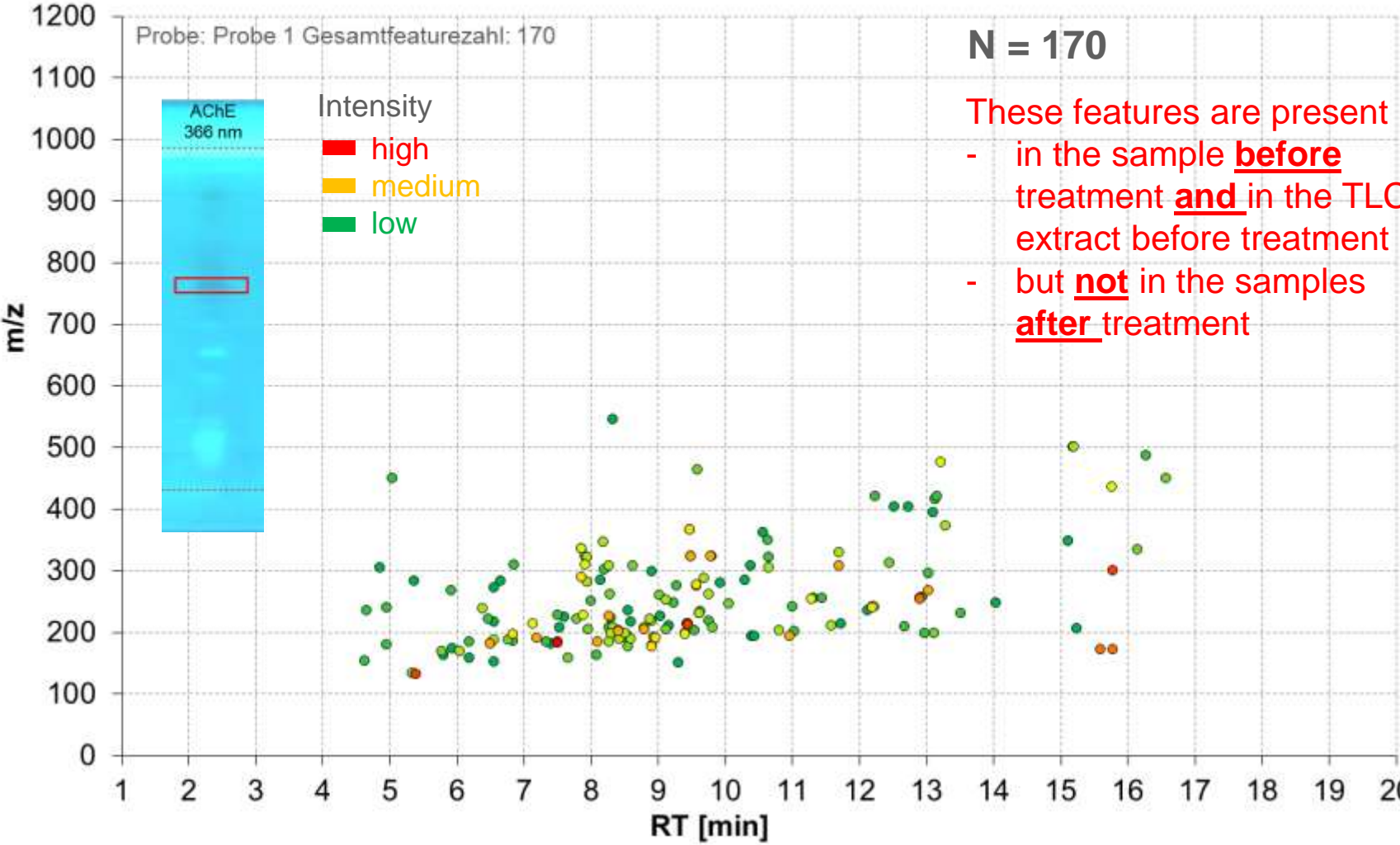
before  
treatment



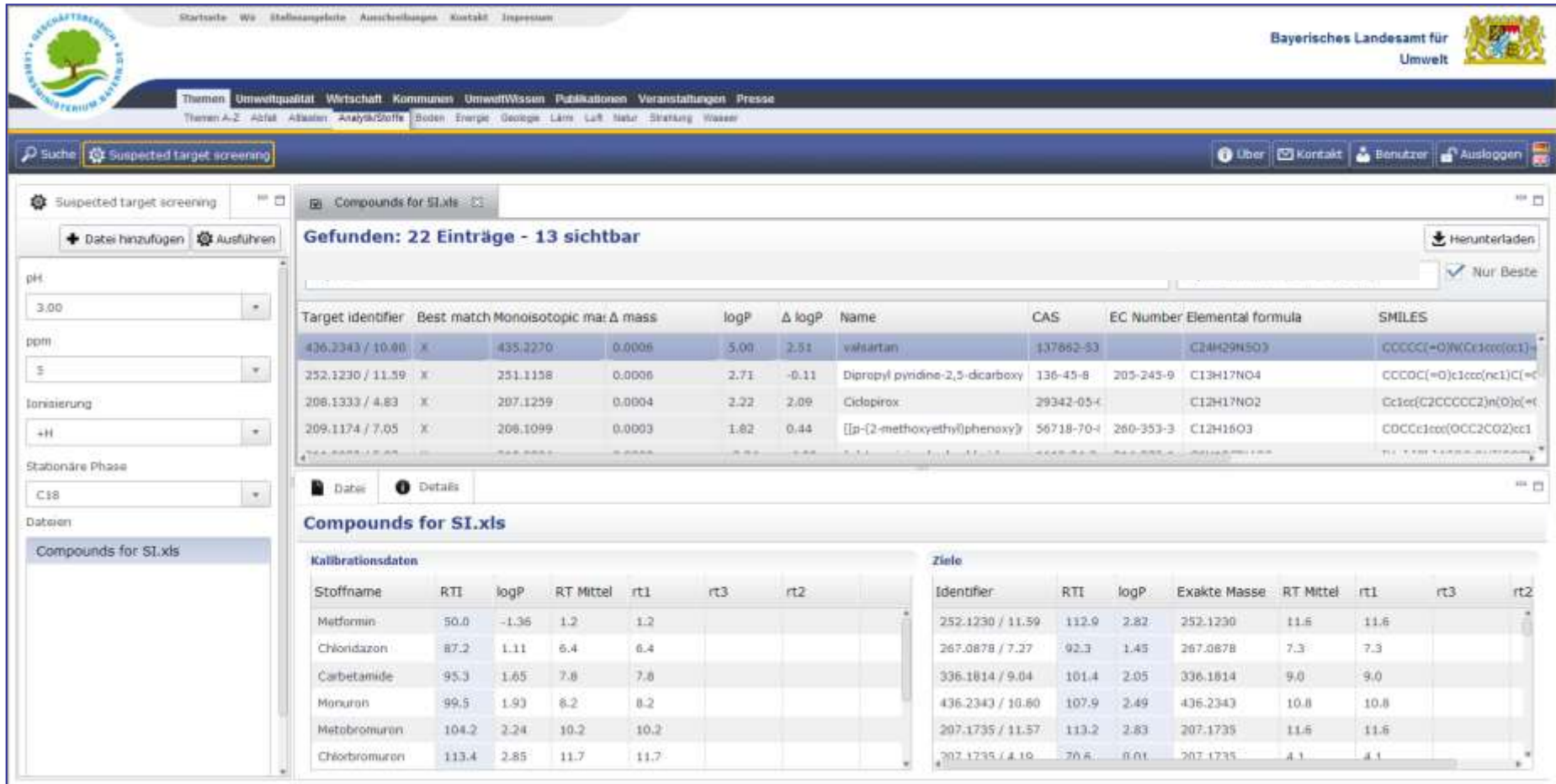
after  
treatment



# RT-m/z scatterplot of AChE inhibition zone



# Stoff-Ident Database Search



Suche **Suspected target screening** Über Kontakt Benutzer Ausloggen

Suspected target screening **Comounds for SI.xls** Herunterladen Nur Beste

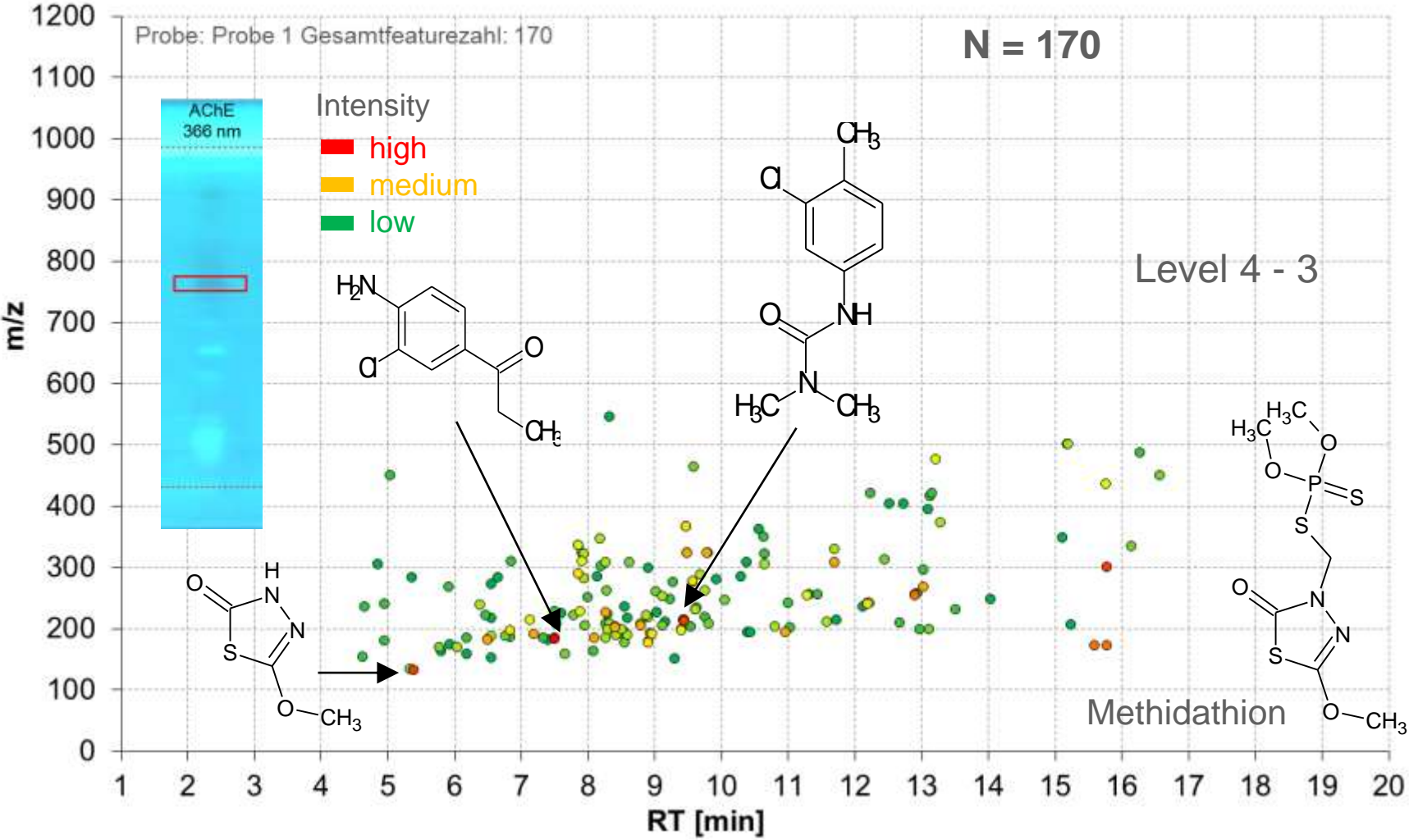
**Gefunden: 22 Einträge - 13 sichtbar**

Target Identifier	Best match	Monoisotopic m/z	Δ mass	logP	Δ logP	Name	CAS	EC Number	Elemental formula	SMILES
436.2343 / 10.00	X	435.2270	0.0006	5.00	2.51	valiartan	137862-53		C24H29H5O3	CCCCC(=O)NCC1ccc(Oc1)~
252.1230 / 11.59	X	251.1158	0.0006	2.71	-0.11	Dipropyl pyridine-2,5-dicarboxy	136-45-8	205-245-9	C13H17NO4	CCCC(=O)c1ccc(nc1)C(=O)~
208.1333 / 4.83	X	207.1259	0.0004	2.22	2.09	Ciclopirox	29342-05-4		C12H17NO2	Cc1cc(C2CCCC2)n(O)c1c~
209.1174 / 7.05	X	208.1099	0.0003	1.82	0.44	[[p-(2-methoxyethyl)phenoxy]]	56718-70-4	280-353-3	C12H16O3	COCCc1ccc(OCC2CO2)cc1~

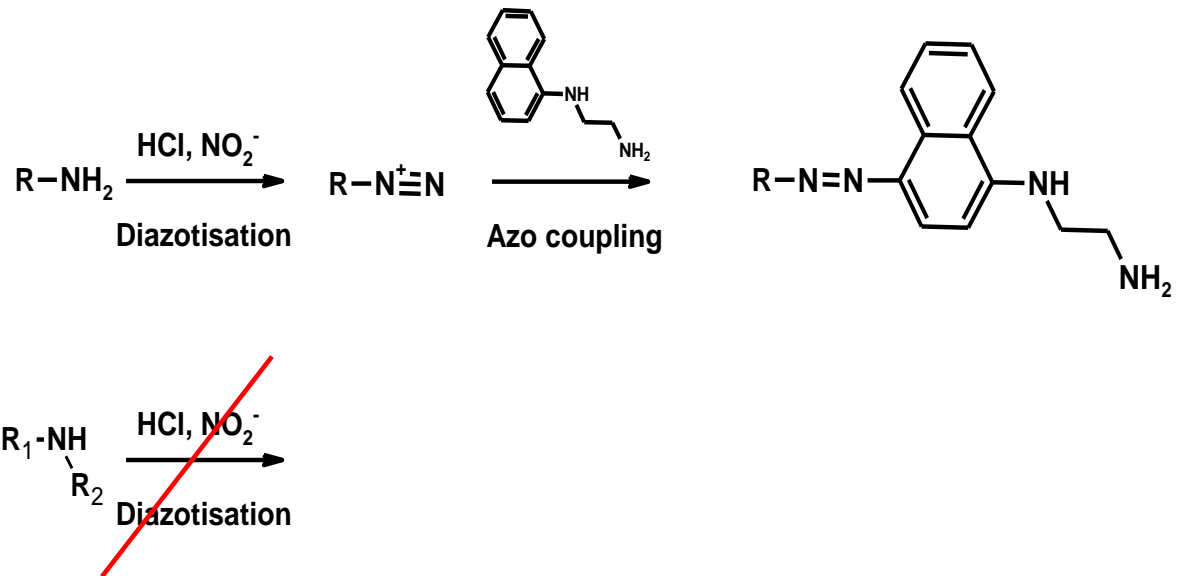
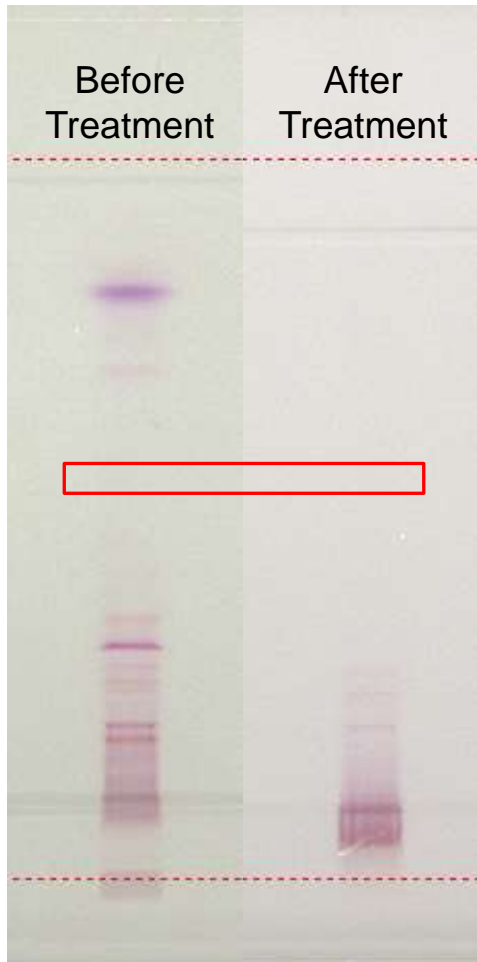
**Comounds for SI.xls**

Kalibrationsdaten							Ziele							
Stoffname	RTI	logP	RT Mittel	rt1	rt3	rt2	Identifier	RTI	logP	Exakte Masse	RT Mittel	rt1	rt3	rt2
Metformin	50.0	-1.36	1.2	1.2			252.1230 / 11.59	112.9	2.82	252.1230	11.6	11.6		
Chloridazon	87.2	1.11	6.4	6.4			267.0878 / 7.27	92.3	1.45	267.0878	7.3	7.3		
Carbetamide	95.3	1.65	7.8	7.8			336.1814 / 9.04	101.4	2.05	336.1814	9.0	9.0		
Monuron	99.5	1.93	8.2	8.2			436.2343 / 10.80	107.9	2.49	436.2343	10.8	10.8		
Metobromuron	104.2	2.24	10.2	10.2			207.1735 / 11.57	113.2	2.83	207.1735	11.6	11.6		
Chlorbromuron	113.4	2.85	11.7	11.7			207.1735 / 4.19	70.6	0.01	207.1735	4.1	4.1		

# RT-m/z scatterplot of AChE inhibition zone

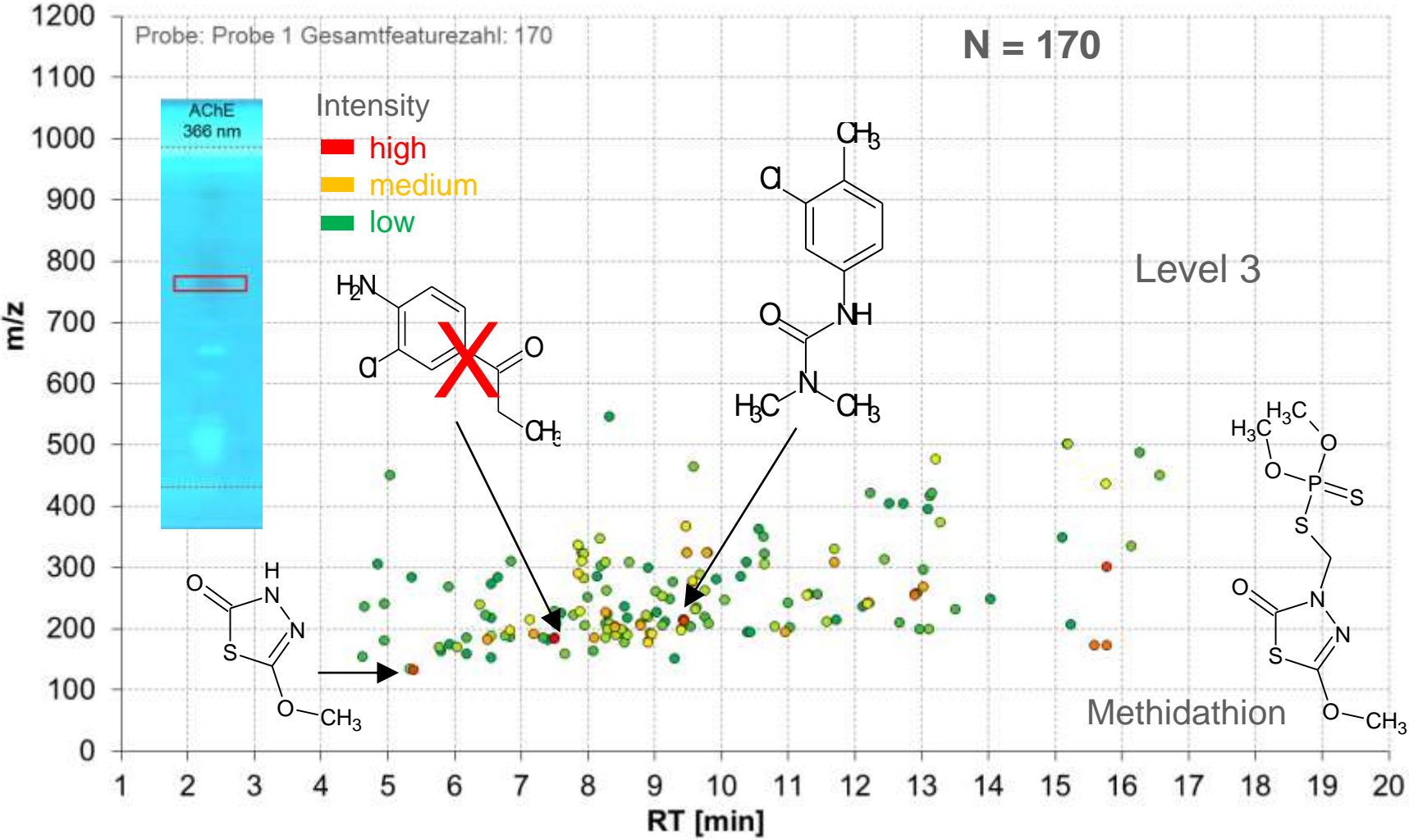


# Detection of primary amino groups with Bratton-Marshall reagent





# RT-m/z scatterplot of AChE inhibition zone



# Conclusions

- Application of set theory with the Mass-RT scatterplot is a tool for prioritization of features (i.e. Process description, identifying relevant sources of contamination)
- EDA in combination with TLC is a second dimension to distinguish samples
- EDA with TLC and different endpoints is a powerful tool for Prioritization in non-target analysis
- Non-target analysis in combination with EDA makes the gap between detection and assessment smaller

**Thank you  
for your attention!**

