



HELMHOLTZ  
ZENTRUM FÜR  
UMWELTFORSCHUNG  
UFZ

# How well can we predict combined effects?

Rolf Altenburger  
21.01.2022



SWEDISH SOCIETY  
OF TOXICOLOGY  
SINCE 1969

# Outline

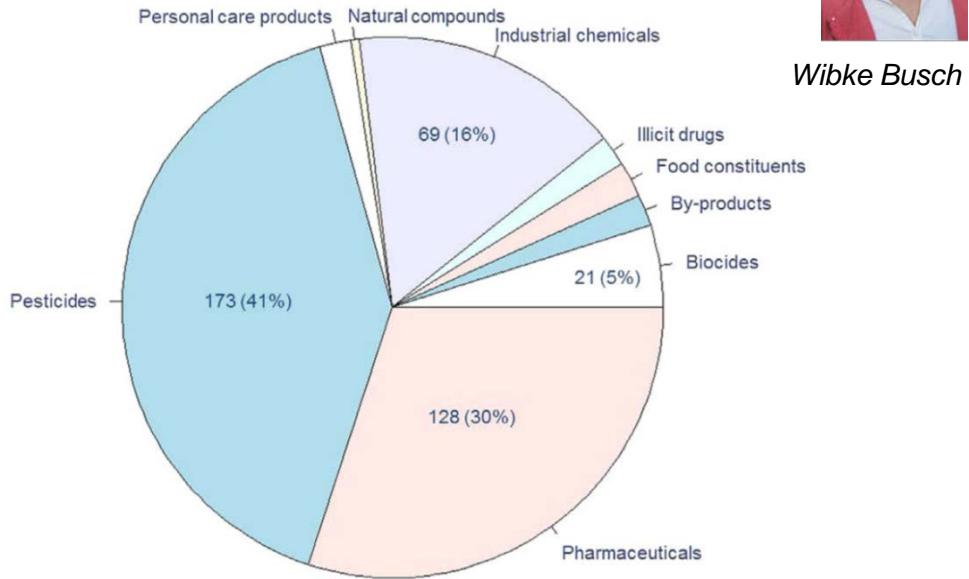
---

- The exposome challenge
- Targeted effect-based mixture detection
- Component-based combined effect prediction
- Mixture exposure and multiple responses

# Chemicals in the environment → Exposome challenge



e.g. >400 contaminants  
in European freshwaters



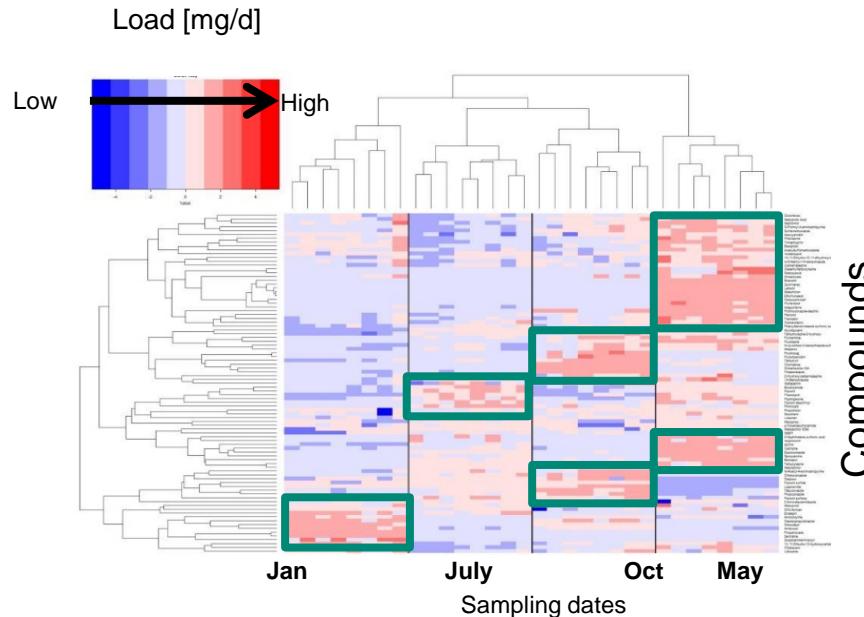
Wibke Busch

Busch, W., et al. (2016): Micropollutants in European rivers: A mode of action survey to support the development of effect-based tools for water monitoring. *Environ. Toxicol. Chem.* 35 (8), 1887 - 1899

# Chemicals in the environment → Exposome challenge



## Nontarget analytics in WWTP effluent



Liza Beckers

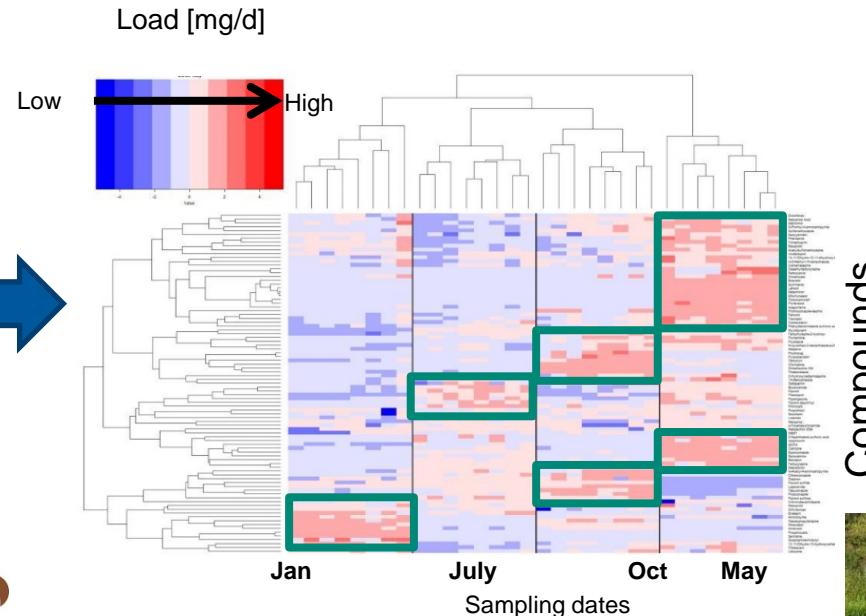
Beckers et al., 2018, *W.*, (2018): Characterization and risk assessment of weather dynamics in organic pollutants mixtures form discharge from a separate sewer system. *Water Res.* 135, 122 - 133

# Chemicals in the environment → Exposome challenge



Nontarget analytics in WWTP effluent

Exposure  
of the population



Compounds

Exposure  
of aquatic  
organisms



# Exposome challenge in humans

## Chemical universe

### Source universe

CAS universe  
163M  
(CAS RN)

Anthro-  
pogenic

Industrial  
products  
30 k (KEMI)

Non-  
anthro-  
pogenic

## Exposome

### Exposome universe

#### Proxies:

Blood exposome  
50k (Barupal)

Urine exposome  
1.4k (HMDB)

Dermal exposome

...

## Human biomonitoring

NHANES  
319 (1999-2017 campaigns)

GERES

HMB4EU  
19 (priority substances and groups)



Sebastian Huhn

Multiplied by (bio)-  
transformation and  
degradation

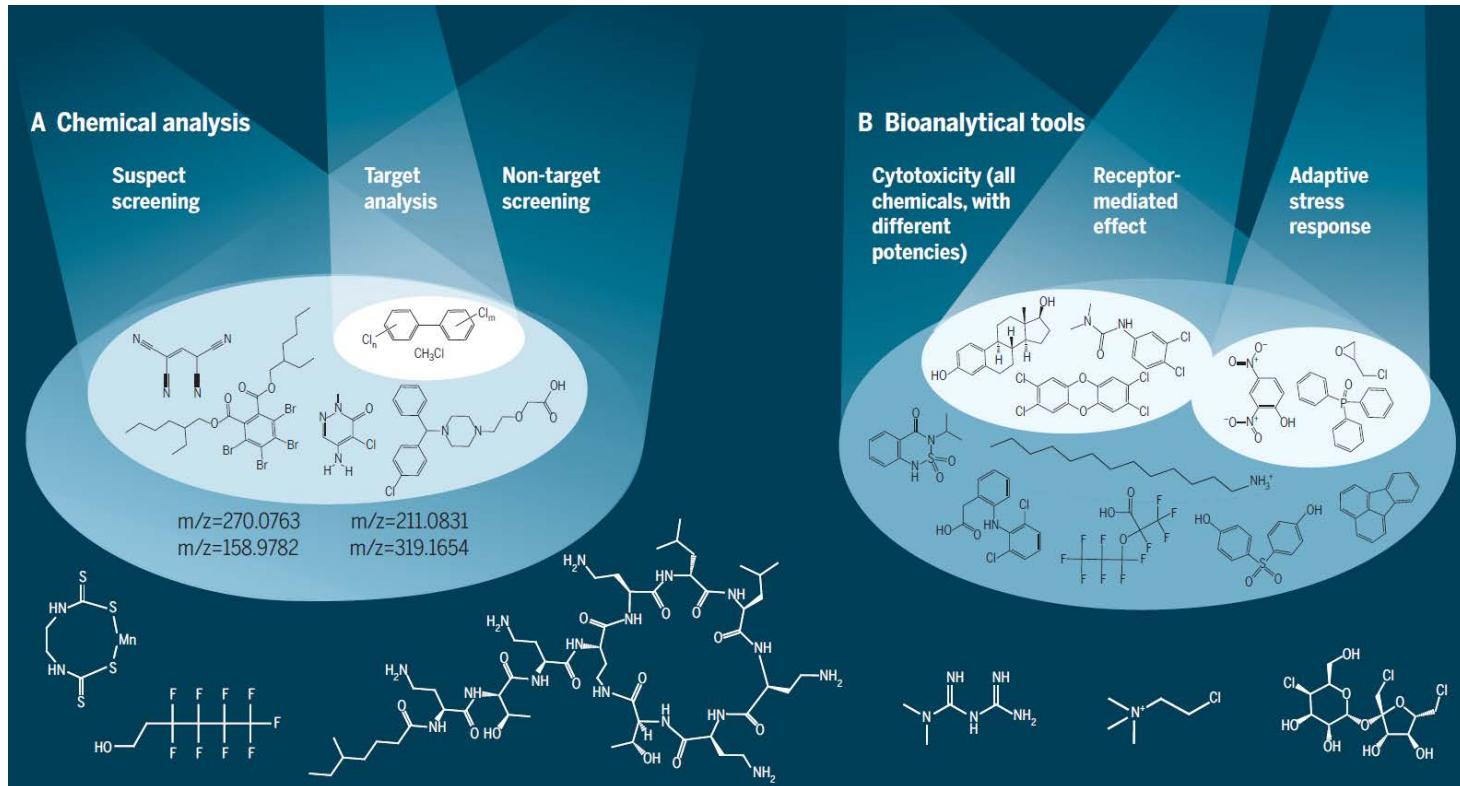
Huhn, S., et al (2021):  
*Unravelling the chemical exposome in cohort studies: routes explored and steps to become comprehensive.*  
*Environ. Sci. Eur.* 33 , art. 17

# Outline

---

- The exposome challenge
- Targeted effect-based mixture detection
- Component-based combined effect prediction
- Mixture exposure and multiple responses

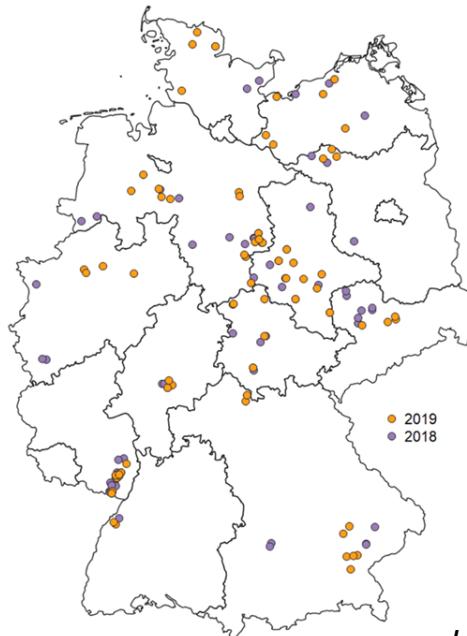
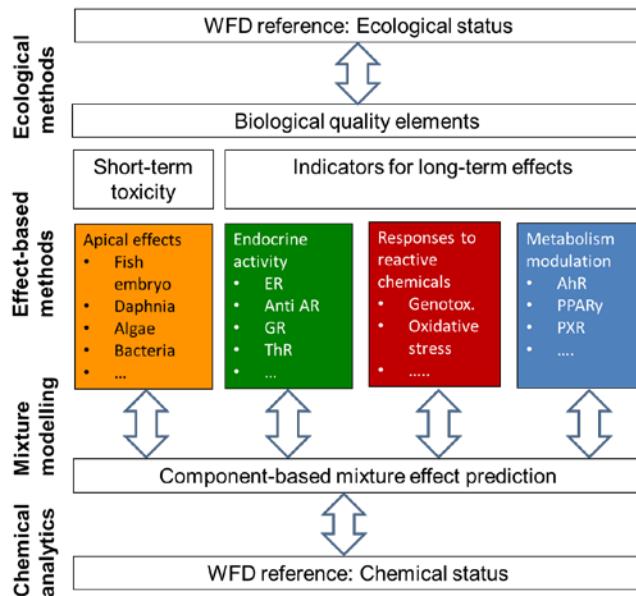
# Effect-based mixture detection → bioanalytics



Beate Escher

Escher, Stapleton,  
Schymanski 2020.  
*Tracking complex  
mixtures of  
chemicals in our  
changing  
environment.*  
Science 367: 388-  
393

# Effect-based mixture detection → bioanalytics



Matthias Liess

Altenburger et al. (2019): Future water quality monitoring: improving the balance between exposure and toxicity assessments of real-world pollutant mixtures. *Environ. Sci. Eur.* 31, art. 12

Liess, et al. (2021) Pesticides are the dominant stressor for vulnerable insects in lowland streams. *Water Res* 201:117262

# Outline

---

- The exposome challenge
- Targeted effect-based mixture detection
- Component-based combined effect prediction
- Mixture exposure and multiple responses

# Predicting combined effects → the concepts

**LOEWE Additivity -  
Dose Addition –  
Concentration Addition**

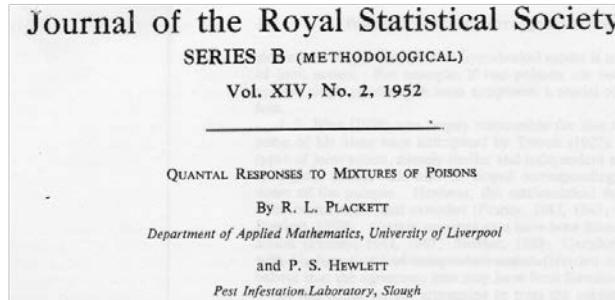
$$1 = \frac{c_1}{EC_{x,1}} + \frac{c_2}{EC_{x,2}}$$

$$ECx_{mix} = \left( \sum_{i=1}^n \frac{p_i}{F_i^{-1}(x_i)} \right)^{-1}$$

**BLISS Independence -  
Independent Action -  
Response Addition**

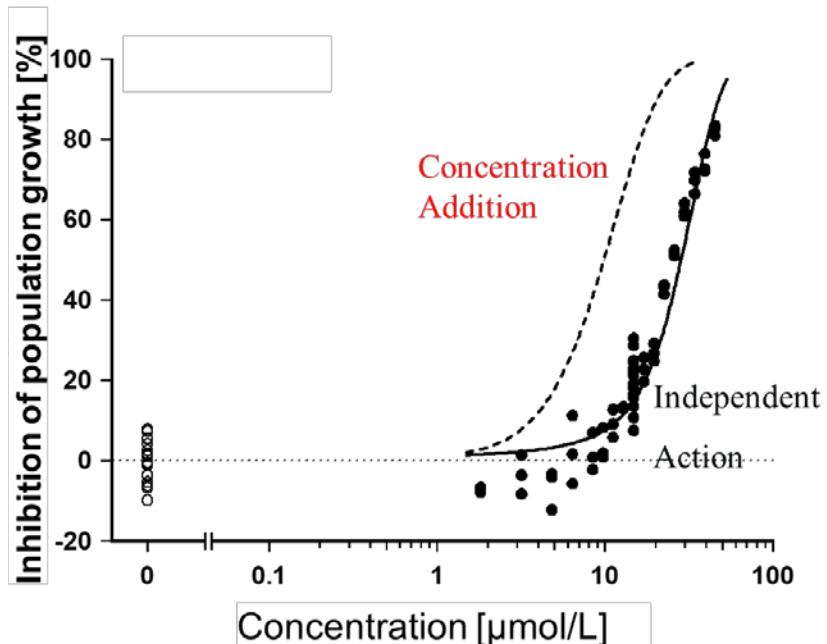
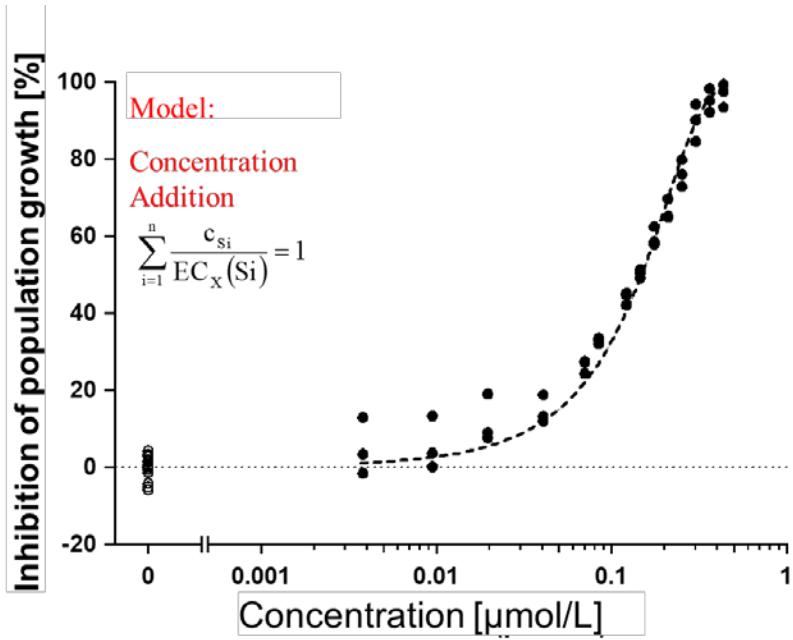
$$E_{(c1,2)} = E_{(c1)} + E_{(c2)} - E_{(c1)} \bullet E_{(c2)}$$

$$X = 1 - \prod_{i=1}^n \left( 1 - F_i(p_i \bullet (ECx_{mix})) \right)$$



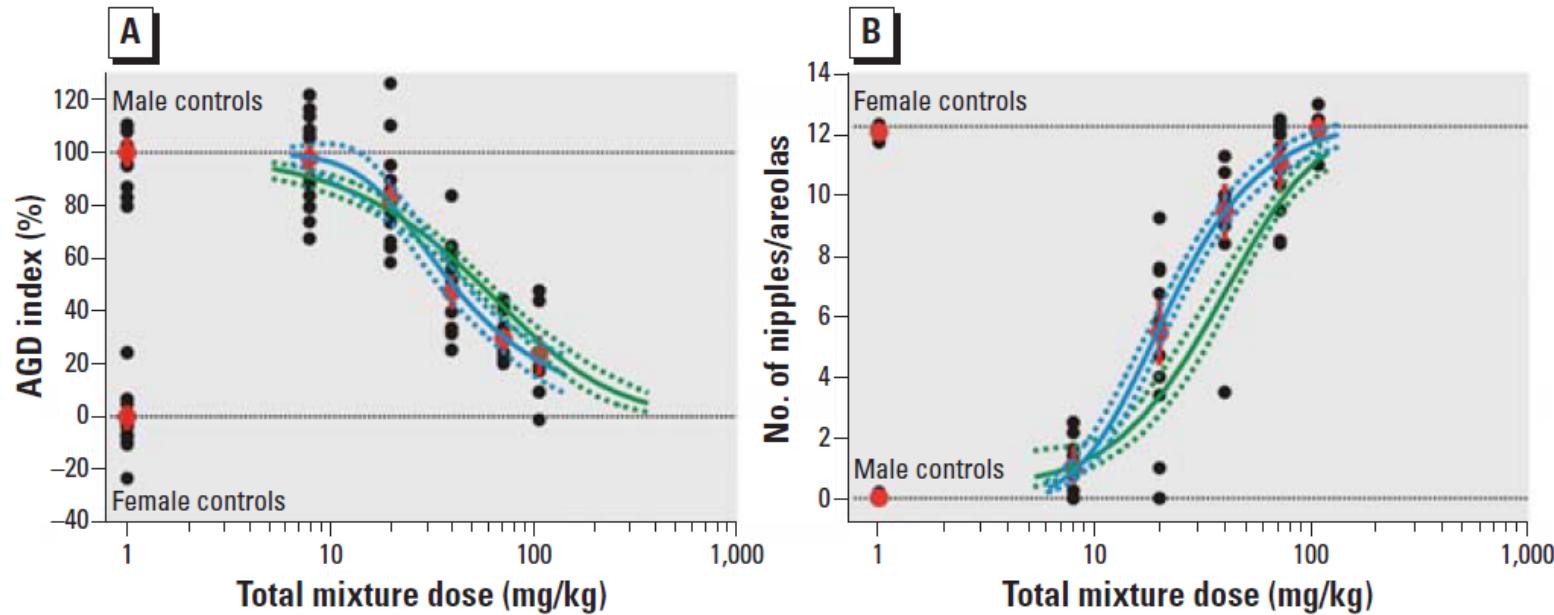
		Mode-of-action	
		Similar	Dissimilar
Effect	Non-interactive	Simple similar	Simple independent
	Interactive	Complex similar	Dependent

# Predictability of combined effects → ecotoxicology



Faust et al. 2000  
J Environ Qual 29:1063

# Predictability of combined effects → rat toxicology



**Figure 1.** Effects of mixed exposure to vinclozolin, flutamide, and procymidone on AGD (A) and NR (B).

**Combined Exposure to Anti-Androgens Exacerbates Disruption of Sexual Differentiation in the Rat**

Ulla Hass,<sup>1</sup> Martin Scholze,<sup>2</sup> Sofie Christiansen,<sup>1</sup> Majken Dalgaard,<sup>1</sup> Anne Marie Vinggaard,<sup>1</sup> Marta Axelstad,<sup>1</sup> Stine Broeng Metzdorff,<sup>1</sup> and Andreas Kortenkamp<sup>2</sup>

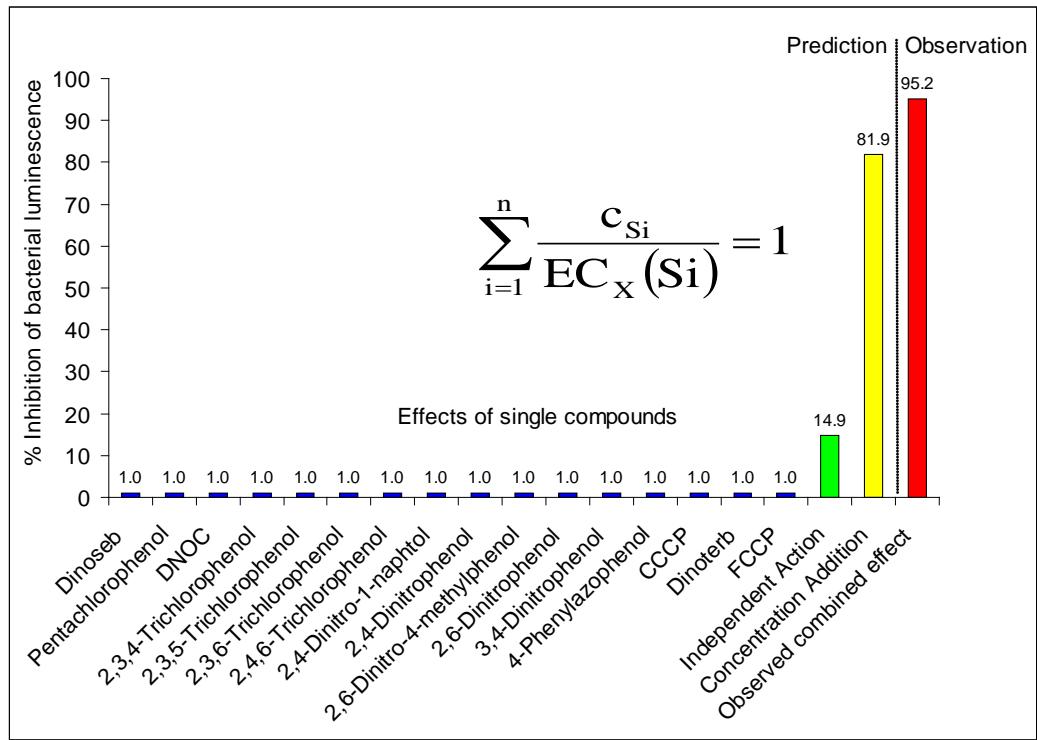
*Environ Health Perspec 115  
Suppl 1, 122-128 (2007)*

<sup>1</sup>Danish Institute for Food and Veterinary Research, Department of Toxicology and Risk Assessment, Søborg, Denmark; <sup>2</sup>The School of Pharmacy, University of London, London, United Kingdom

# Predicting combined effects at low EC



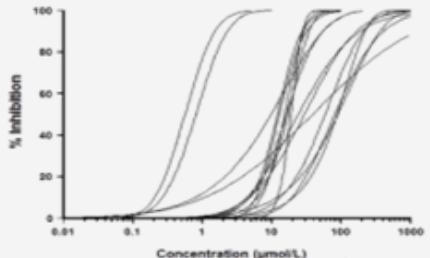
**‘Something from nothing’**



Altenburger & Greco,  
Integr Environ Assess Manag.  
5:62, 2009

# Predictability depends on good quality input data

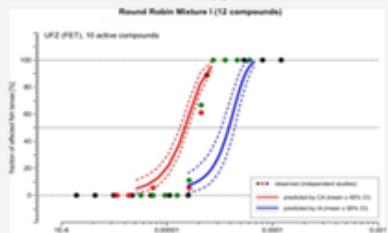
## Concentration- effect relationship



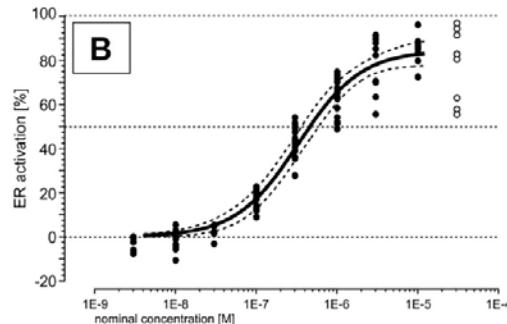
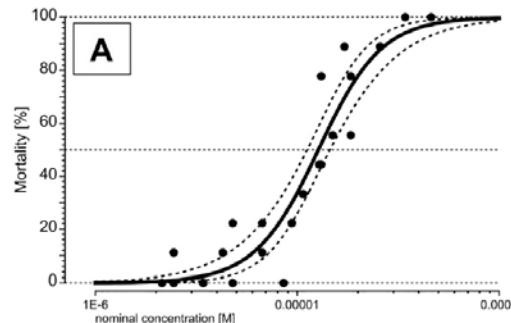
Mixture  
effect  
prediction

$$EC_X(\text{mixture}) = \left( \sum_{i=1}^n p_i EC_{X,i} \right)^{-1}$$
$$E(c_{\text{mixture}}) = \prod_{i=1}^n E(c_i)$$

Mixture  
effect  
diagnosis



Mixture effects: Observed vs. Predicted

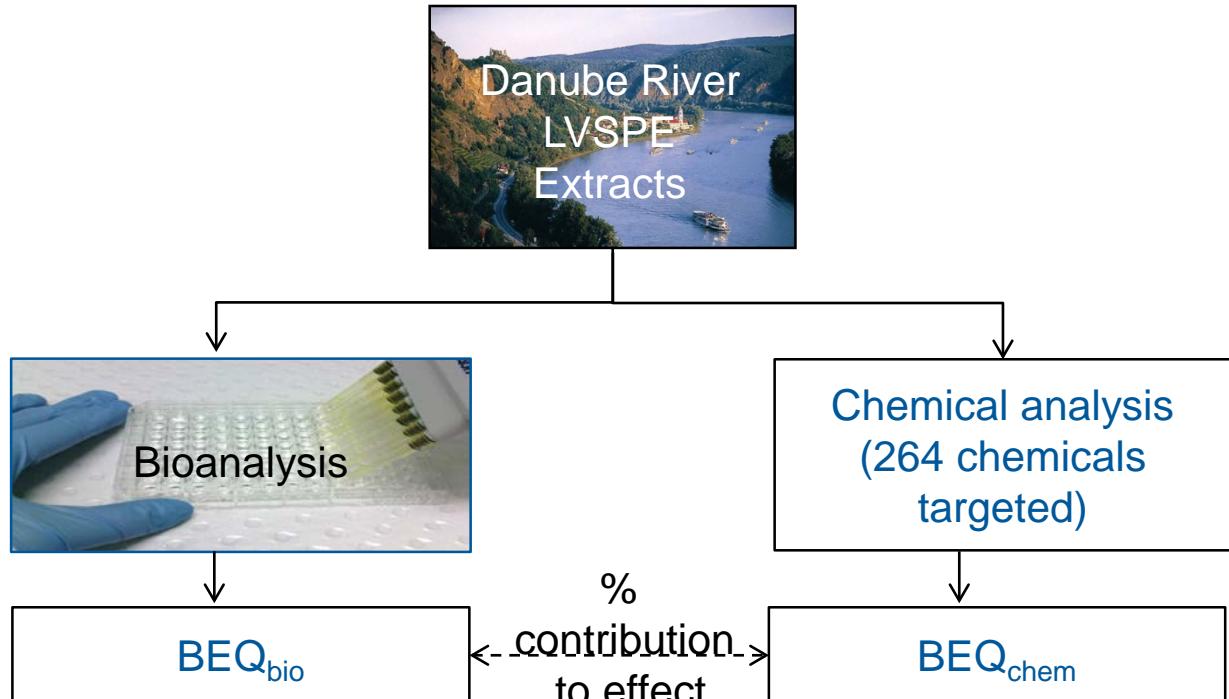


Altenburger et al. (2018): Mixture effects in samples of multiple contaminants – An inter-laboratory study with manifold bioassays. Environ. Int. 114 , 95 - 106

# Combined effects – Linking chemical & bio-analytics



„Iceberg“ modelling

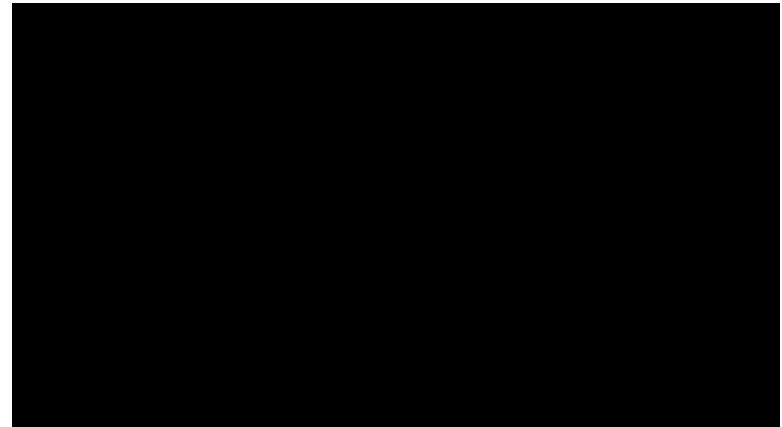


Neale, et al. (2015) ES&T 49: 14614-14624

# Outline

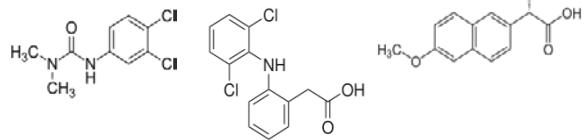
---

- The exposome challenge
- Targeted effect-based mixture detection
- Component-based combined effect prediction
- Mixture exposure and multiple responses



# → towards comprehensive combined effect predictions

Zebrafish eggs exposed to



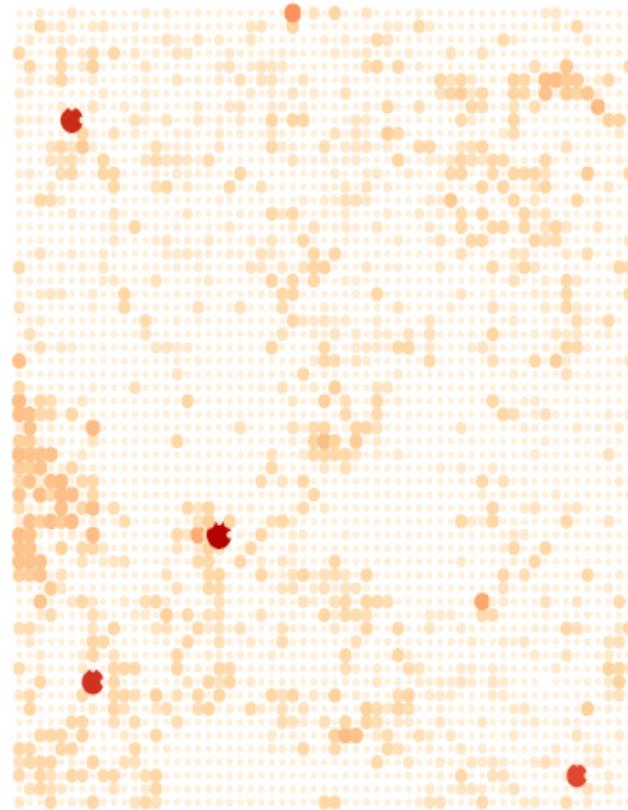
Diuron

Diclofenac

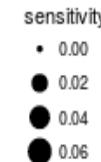
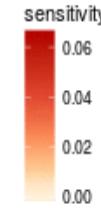
Naproxen



hours under exposure: 3

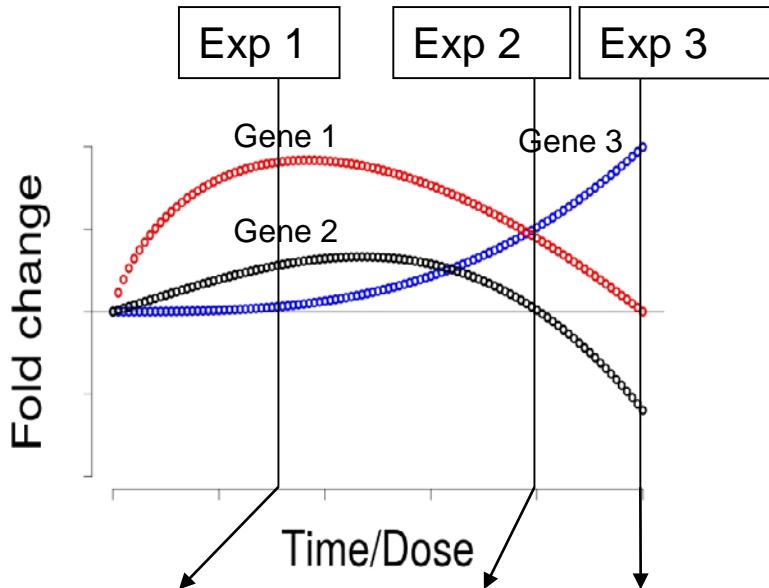


Transcriptome  
response



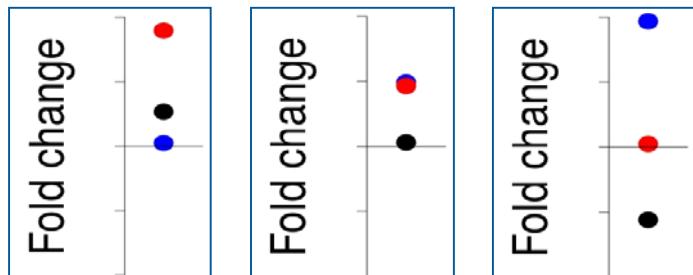
Andreas Schüttler

# → towards comprehensive combined effect predictions



→ dense sampling design

Dose	0 Control	LC25 /x <sup>4</sup>	LC25 /x <sup>3</sup>	LC25 /x <sup>2</sup>	LC25 /x	LC25
Time age ZFE (h of exposure)						
24 hpf (0) – Exposure Start						
27 hpf (3)	XX	X	X	X	X	XX
30 hpf (6)	XX	X	X	X	X	XX
36 hpf (12)	XX	X	X	X	X	XX
48 hpf (24)	XX	X	X	X	X	XX
72 hpf (48)	XX	XX	XX	XX	XX	XX
96 hpf (72)	XX	XX	XX	XX	XX	XX

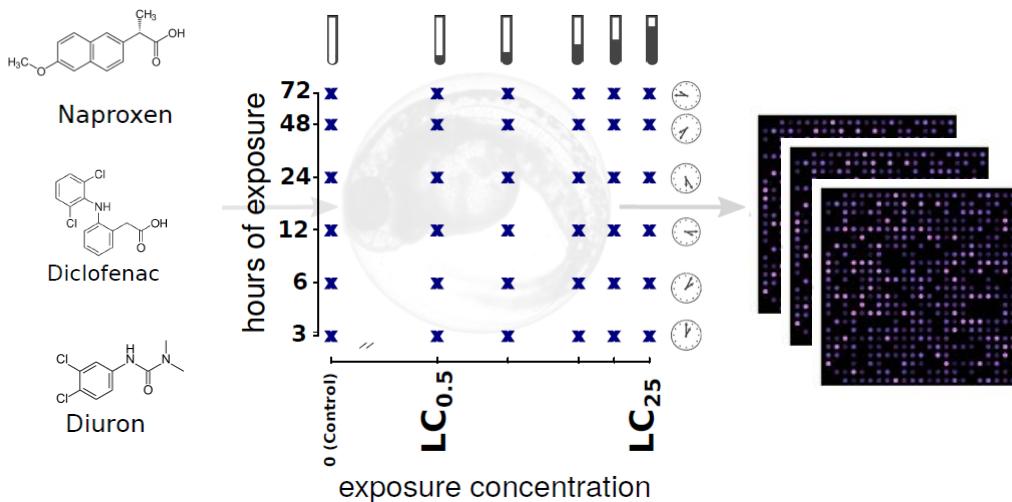


Schüttler et al. 2017,  
Toxicological Sciences 157(2): 291–304

# Questions and Approach

- How to see the complete picture?
- How to compare toxicogenomic effects of different substances?
- How to describe those effects mathematically to enable mixture calculations?

## Experiment

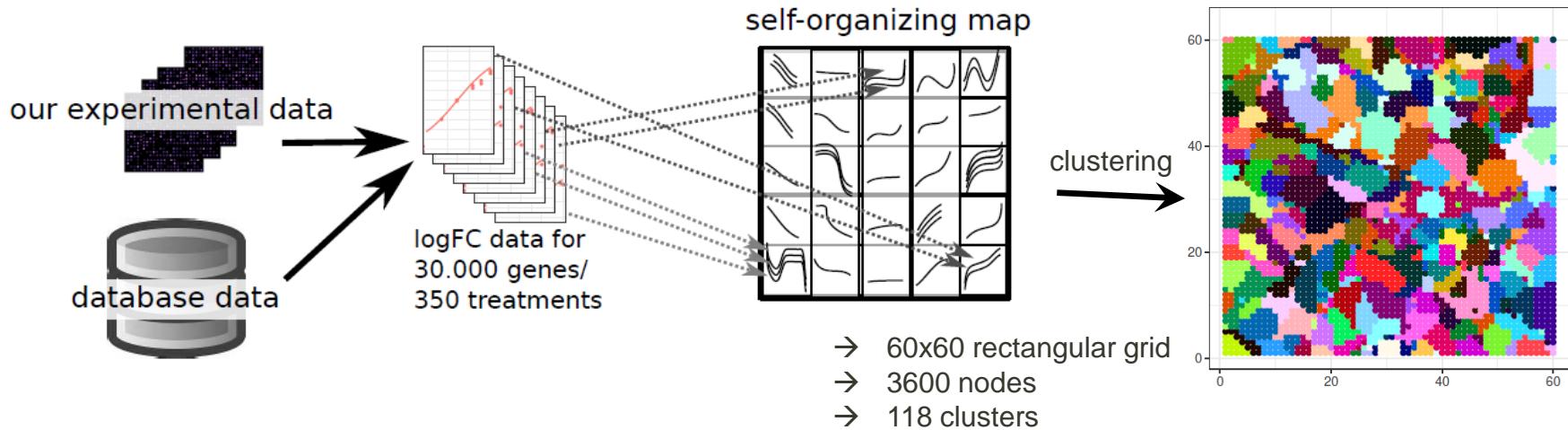


## Data analysis

- *integration of previous data*
- *aggregation of fingerprints*
- *modeling of responses*

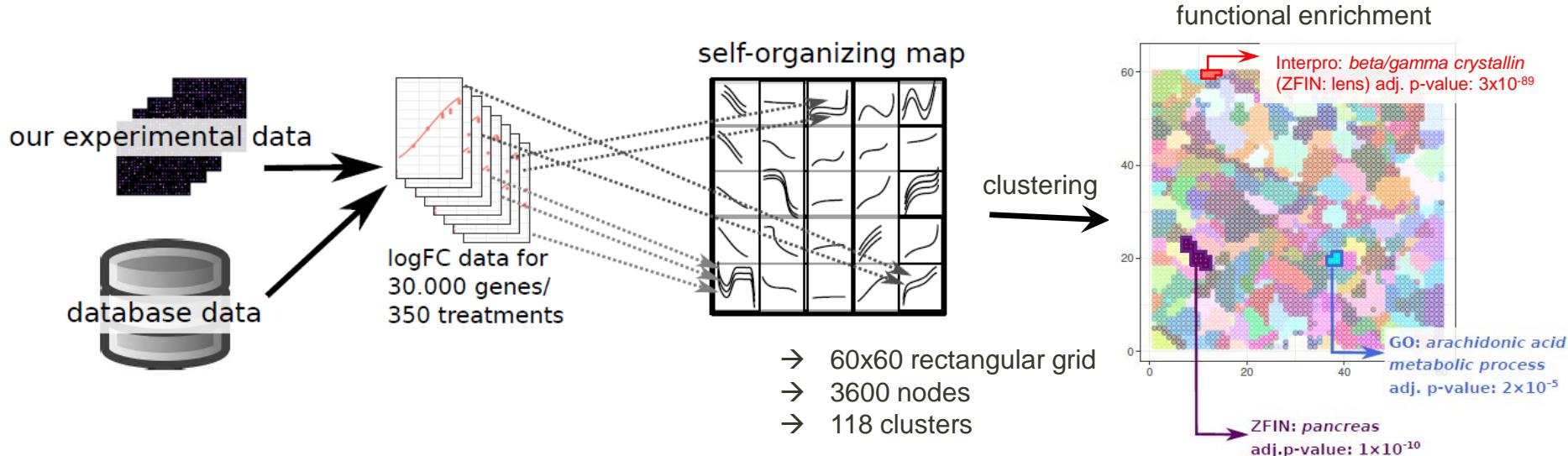
Schüttler et al., *Map and model—moving from observation to prediction in toxicogenomics*. *GigaScience* 8 (6), giz057, 2019

# Integration and aggregation of data



Andreas Schüttler et al.,  
*GigaScience*, 2019

# Integration and aggregation of data



Andreas Schüttler et al.,  
*GigaScience*, 2019

# Integration and aggregation of data

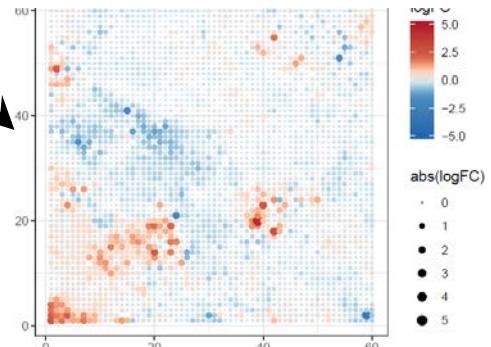
our experimental data



logFC data for  
30.000 genes/  
350 treatments

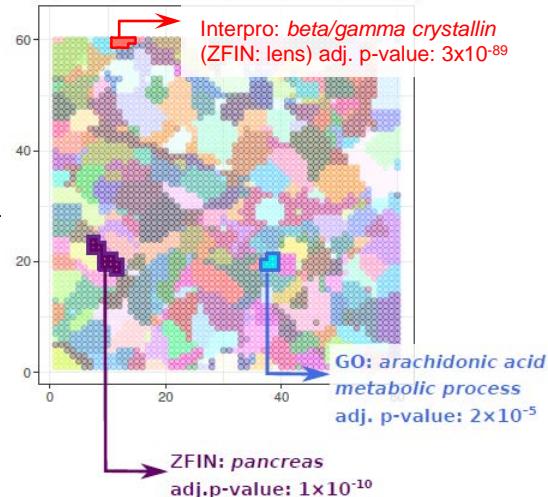
self-organizing map

- 60x60 rectangular grid
- 3600 nodes
- 118 clusters



clustering

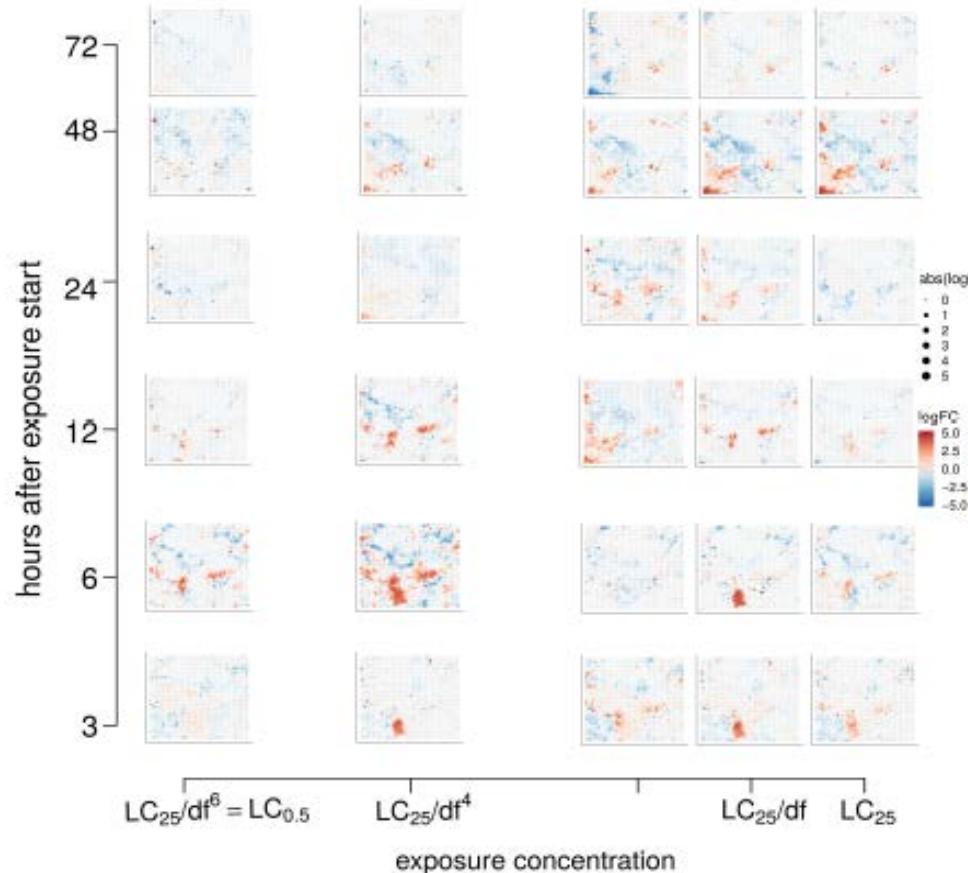
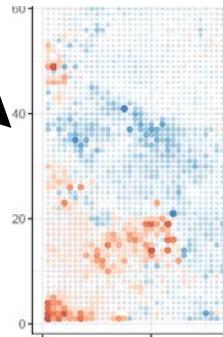
functional enrichment



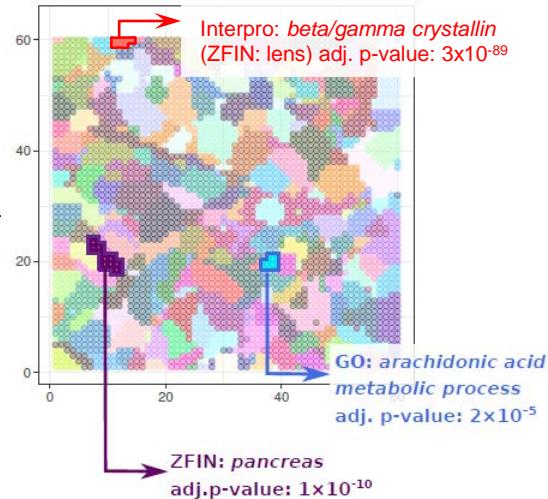
Andreas Schüttler et al.,  
*GigaScience*, 2019

# Integration and aggregation of data

our experimental

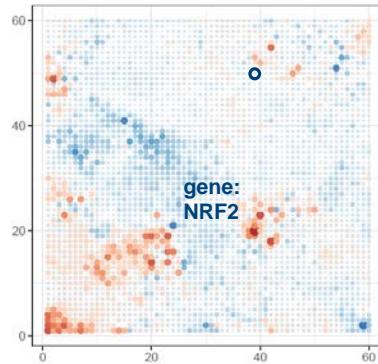
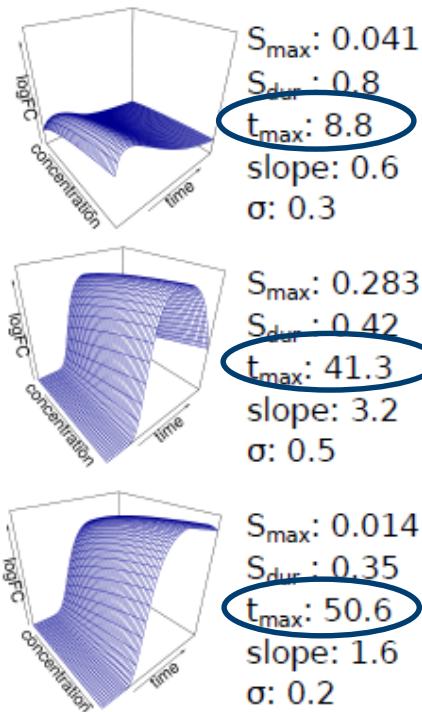
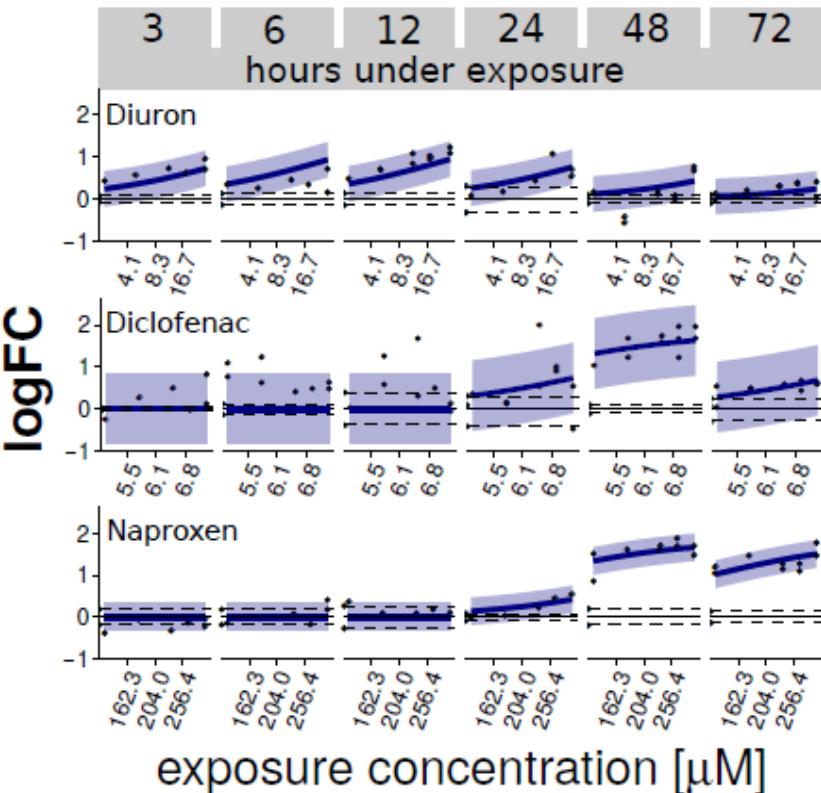


functional enrichment



Andreas Schüttler et al.,  
GigaScience, 2019

# Regression modelling



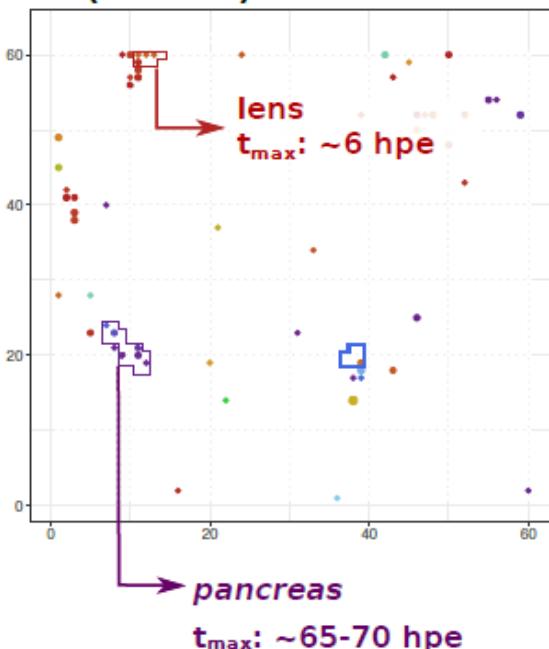
$$\log FC(c) = \frac{\log FC_{\max}}{1 + e^{-slope * (\log(c) - \log(X_{50}))}}$$

$$sensitivity(t) = \frac{1}{X_{50}(t)} = S_{\max} * e^{-0.5 * \left( \frac{\log(t) - \log(t_{\max})}{S_{dur}} \right)^2}$$

# Mapping model parameters

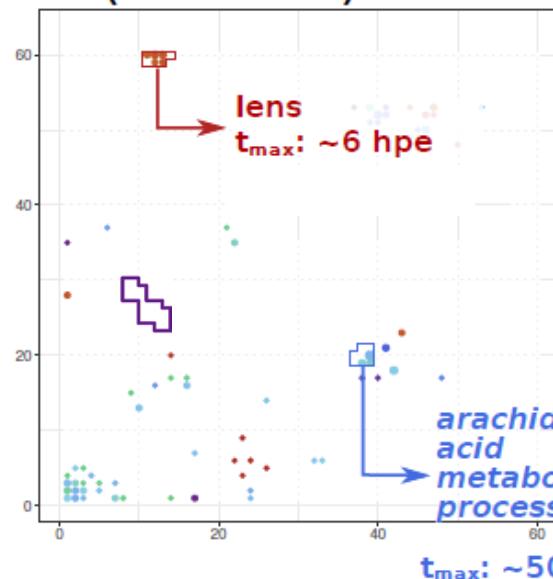
Herbicide (PSII Inhibitor)

**A (diuron)**

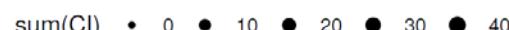
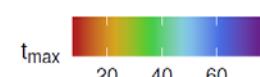
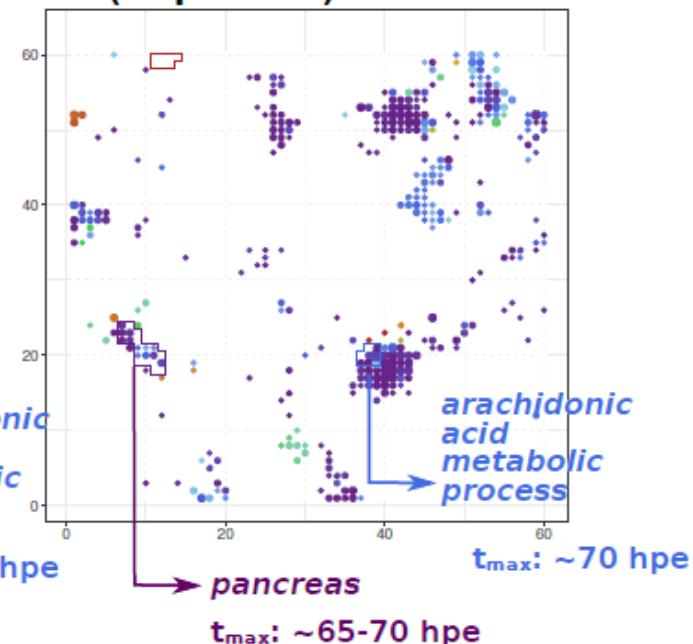


Pharmaceuticals (Cox-inhibitors)

**B (diclofenac)**

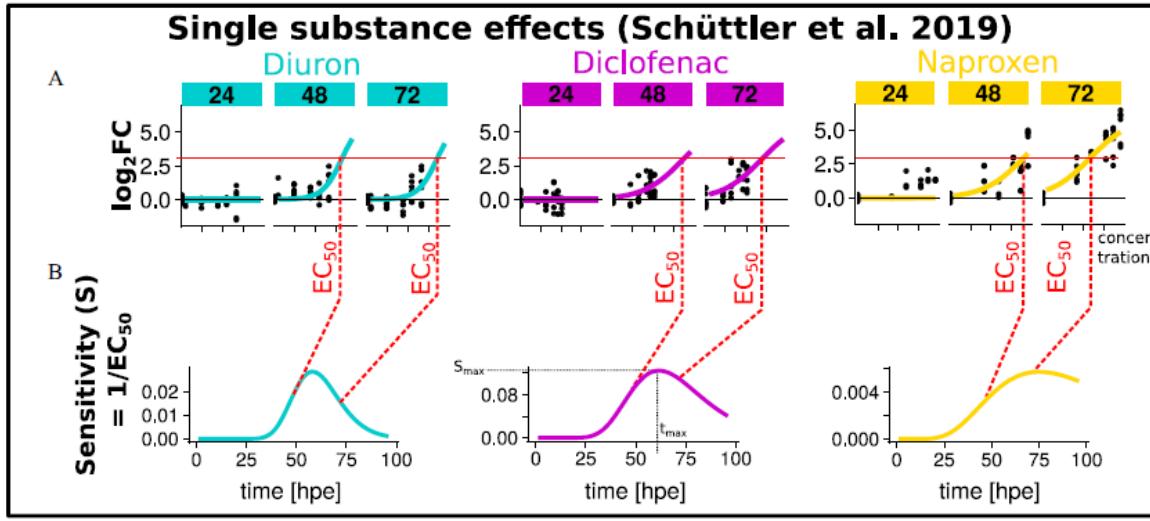


**C (naproxen)**

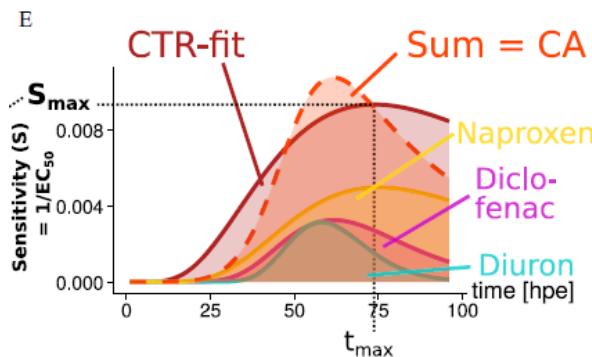


Adapted from: Schüttler et al., *GigaScience*, 2019

# Mixture effect prediction

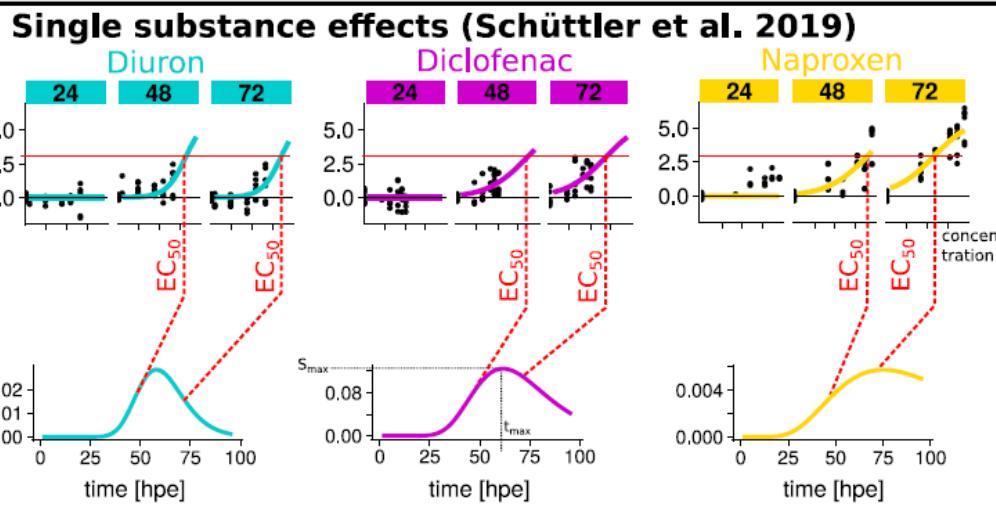
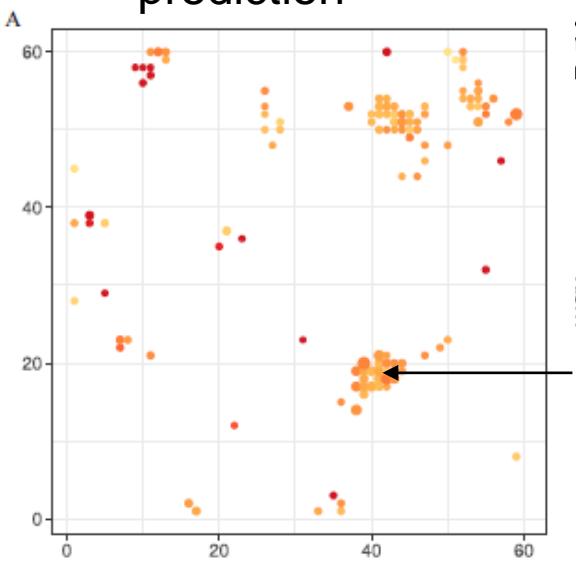


## Mixture prediction - Concentration Addition (CA)

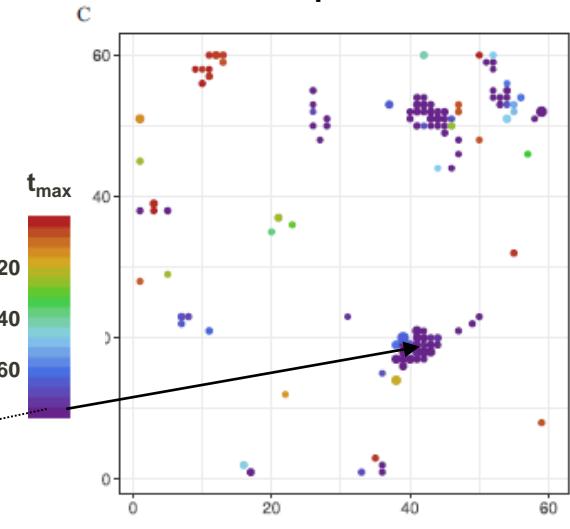
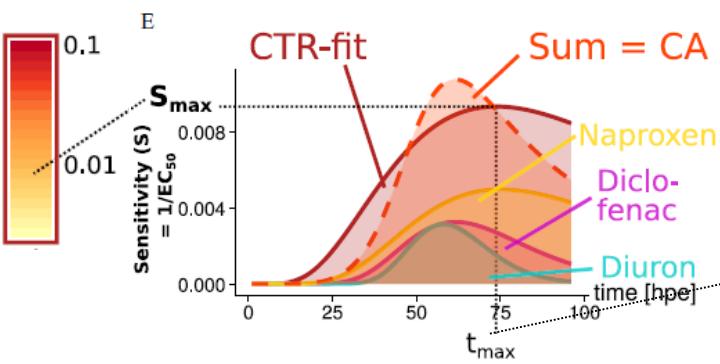


# Mixture effect prediction

$S_{\max}$  map – prediction

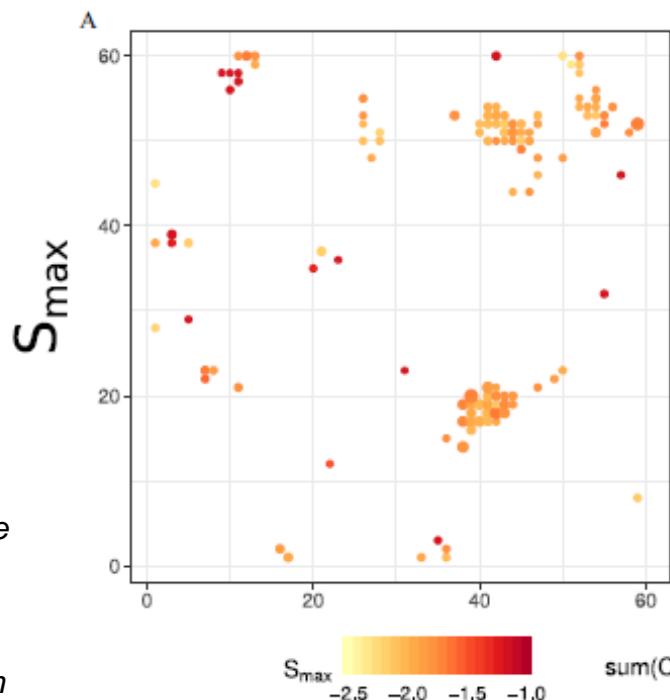


$t_{\max}$  map – prediction

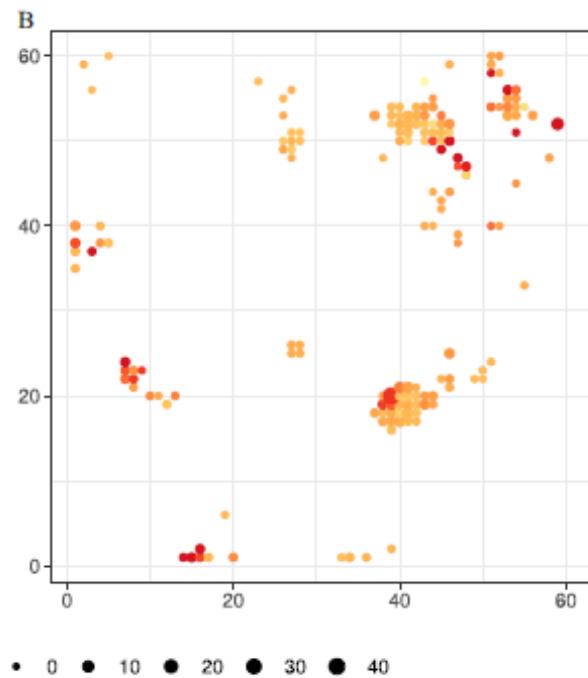


# Prediction versus Observation

**Prediction (CA)**

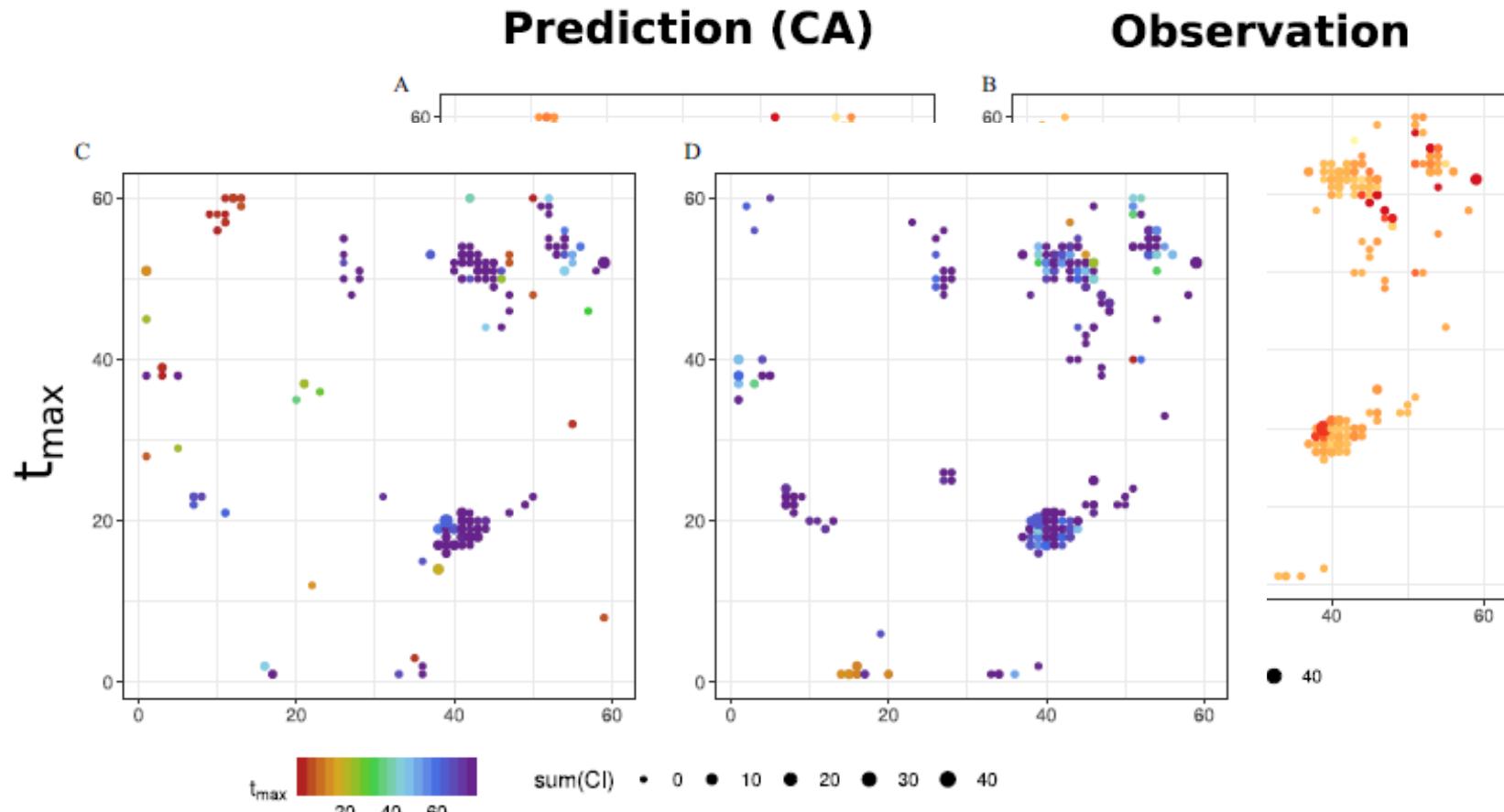


**Observation**



Schüttler et al., Transcriptome-wide prediction and measurement of combined effects induced by chemical mixture exposure in zebrafish embryos. *Environ. Health Perspect.* 2021, 129 (4), art. 047006

# Prediction versus Observation



# Summary (for combined effects in a multiple response system)

---

- Recovery of single substance effects? **YES**
- Occurrence of combined effects on the transcriptome? **YES**
  - Combined effects observed also for anticipated dissimilar acting compounds
- Can we predict the effects of a mixture on the transcriptome? **YES**
  - CA outperforms other mixture concepts (Schüttler et al. 2021, EHP)
- More-than-additive combined effects occurred rarely,
  - for the pairs of similar and dissimilar acting compounds;
  - for induced and repressed genes and nodes; but
  - can be identified by comparison of observation with CA prediction.

# Conclusions and Outlook

---

- Exposome studies will specify multiple exposure of organisms;
- Analytical characterisation of contamination and mixture exposure can be complemented by bioanalytical approaches;
- Concepts for component-based calculation of combined effects show reasonable predictability across bioassays;
- Including process knowledge allows access to more complex response patterns.

*Drakvik et al. (2020): Statement on advancing the assessment of chemical mixtures and their risks for human health and the environment. Environ. Int. 134 , art. 105267*

# Thank you for your attention

---



@UFZ\_CITE