



3rd meeting in the OECD advisory group
for molecular screening and toxicogenomics
(Paris, Oct. 26-27)

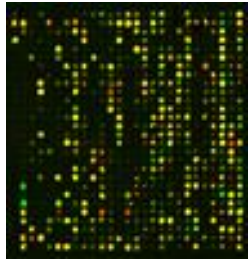


Ecotoxicogenomics in predictive risk assessment — potential use of *in vitro* and *in vivo* transcriptomics for assessing toxicological effects of chemicals

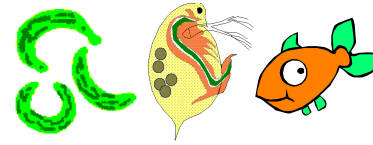
Knut Erik Tollefsen

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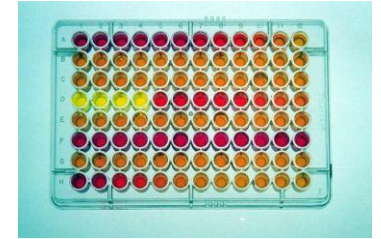
Ecotoxicology and risk assessment



**Molecular
toxicology**



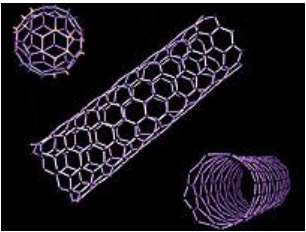
**Ecotoxicological
testing**



**Biomarker
research**

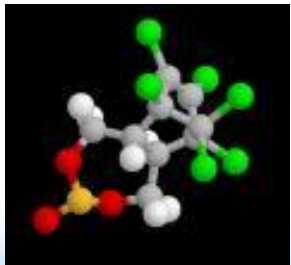


**Effluent
monitoring**



**Cellular
toxicology**

**Chemistry
&
Ecotoxicology**



**Ecotoxicological
consulting**

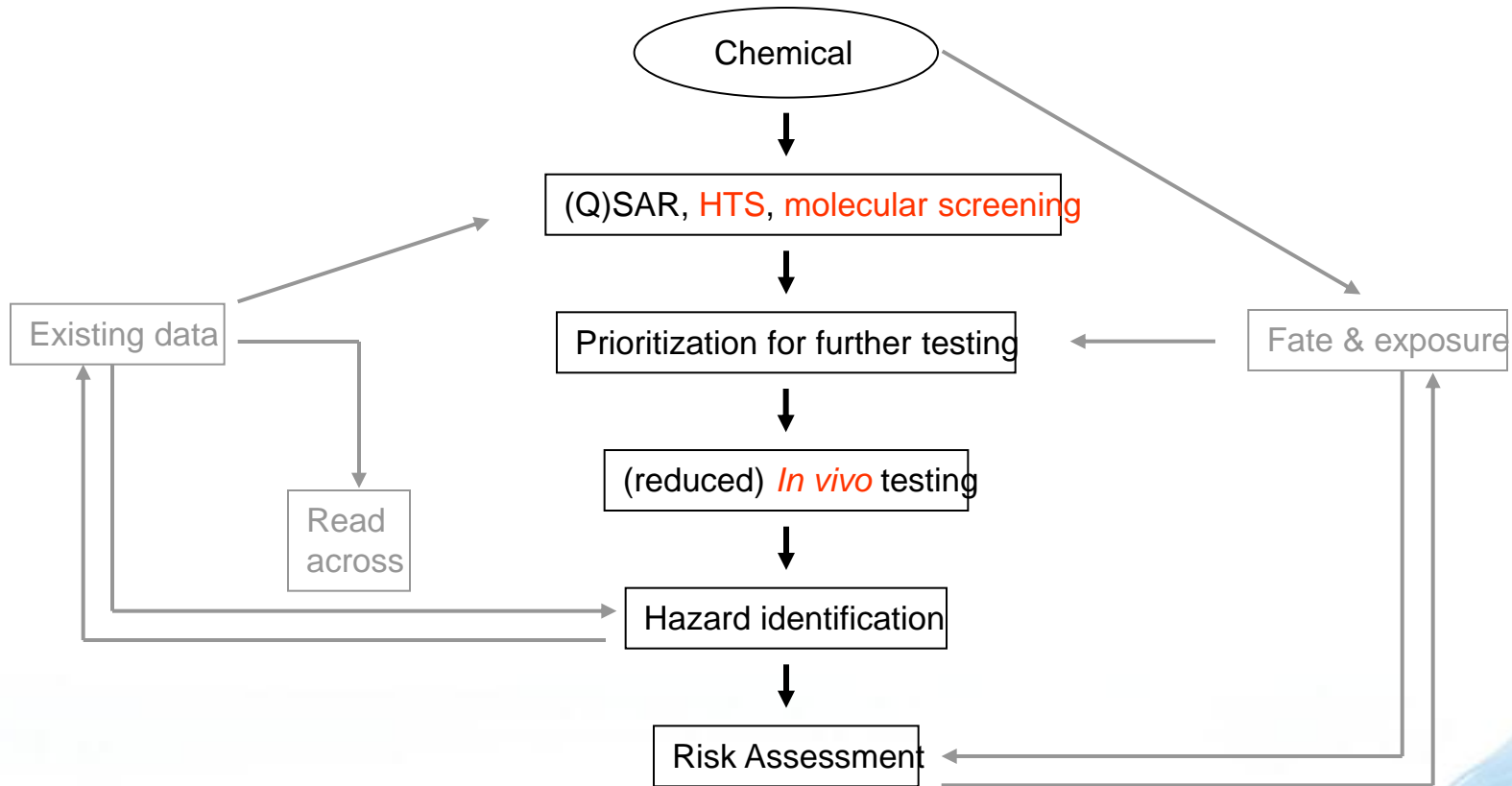
**Environmental
monitoring**



Risk assessment



Integrated testing strategies

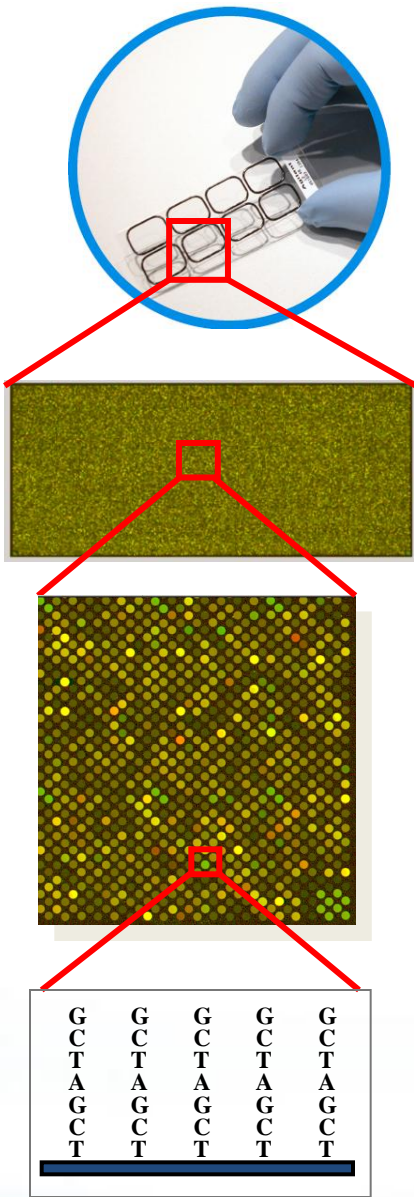


Objectives

- Develop transcriptomic tools for aquatic toxicity testing
- Apply technology to *in vitro* and *in vivo* test systems
- Screen for potential adverse effects of chemicals & mixtures
- Determine whether technology is applicable to risk assessment

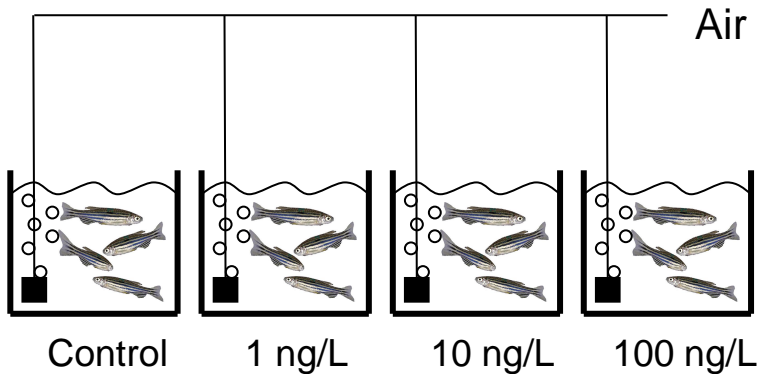
Oligo microarrays

- Flexible (custom & standard) microarrays
- 60-mer oligonucleotides printed on array (15-244k)
- Chip technology accomodates up to 1 million gene probes (features)
- Gene expression detected by laser scanner technology
- Bioinformatics translate gene expression to meaningful biological information
- Full genome sequence available for some species
- Currently applied for *in vitro* and/or *in vivo* testing (zebrafish, salmonids, cod, halibut, calanus, mussels)
- Implemented in environmental monitoring (salmonids, cod, calanus, mussels)



Experimental models

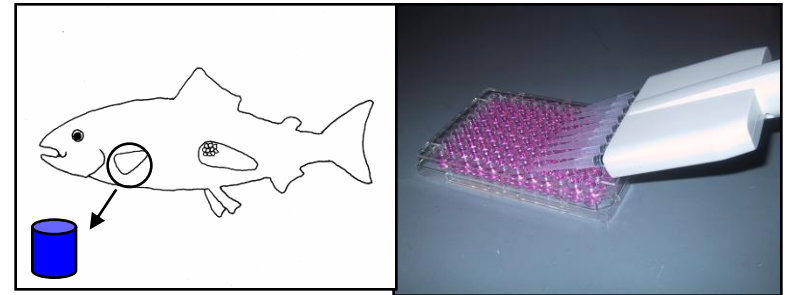
In vivo
(1-4 days exposure)



Studies

(xeno)estrogen (E2, EE2)
PAH (BaP)
Persistent compounds (HBCD)
Pharmaceuticals (Diclofenac)
Artificial sweetener (Sucralose)
Metals (Copper)
Nanoparticles (Silver NPs)
Road run-off/tunnel wash water

In vitro
(1-4 days exposure)

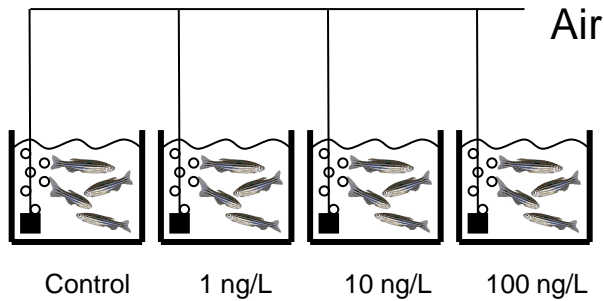
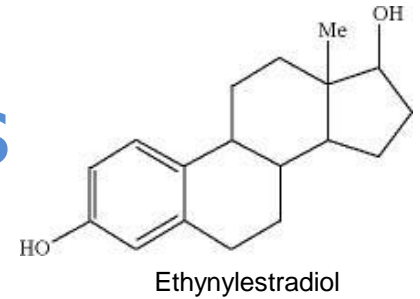


Studies

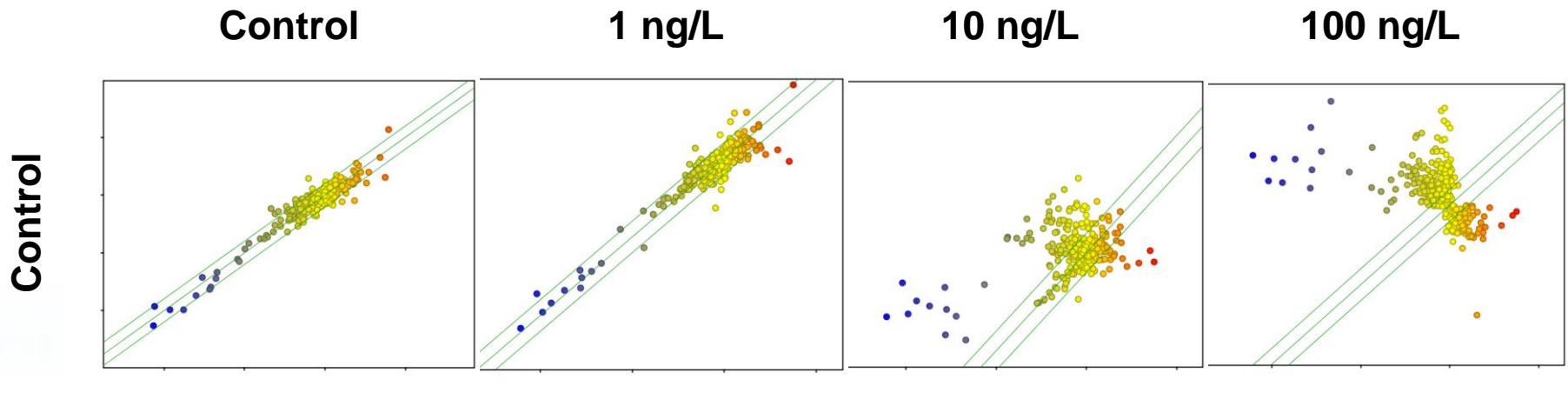
(xeno)estrogen (EE2)
Persistent compounds (TCDD)
Direct mutagens (1-NQO)
Pesticides (paraquat)
Mixtures



Endocrine disruptors



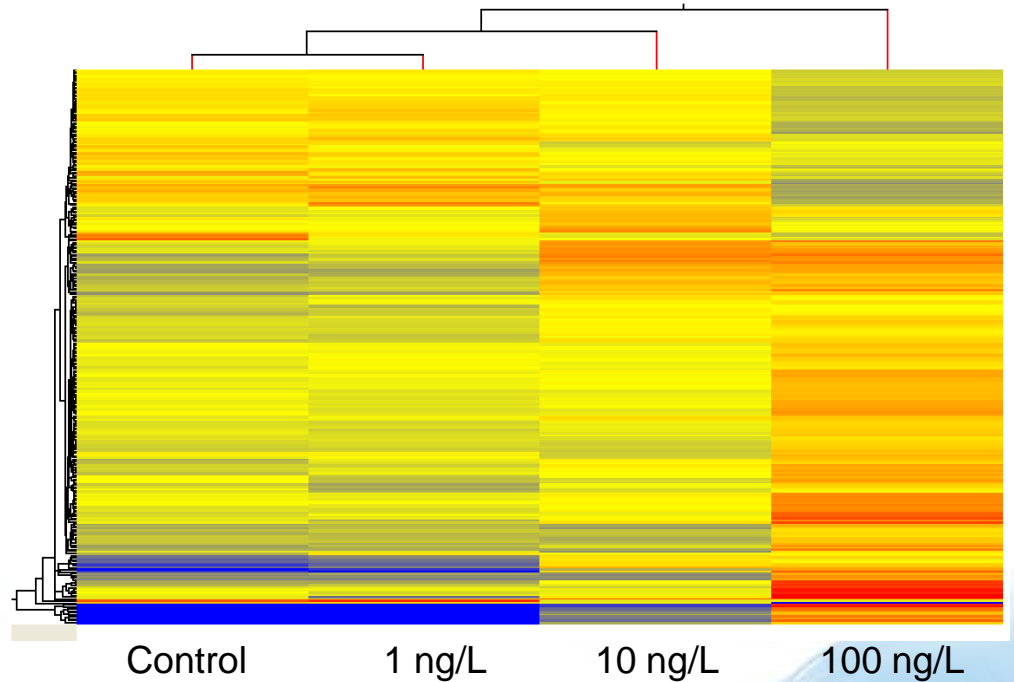
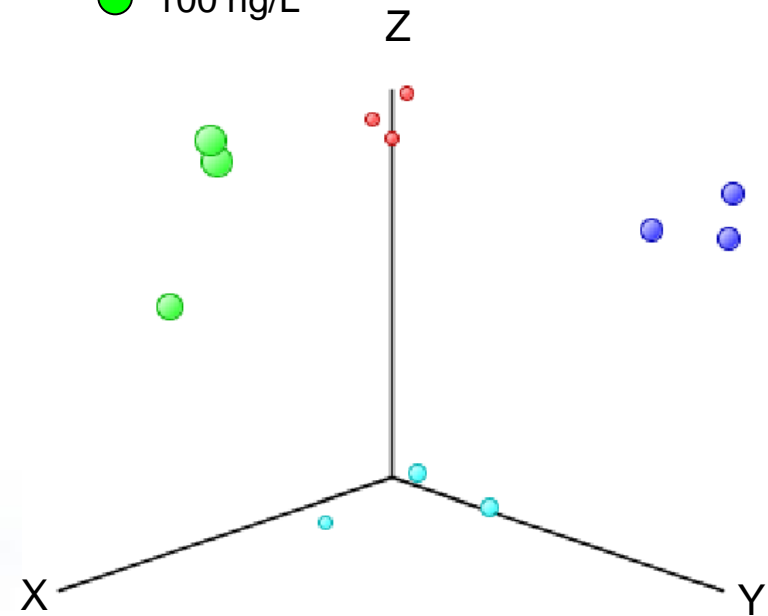
17 α -ethynylestradiol (EE2)



Differential gene expression in male zebrafish livers after 48h exposure (323 features)

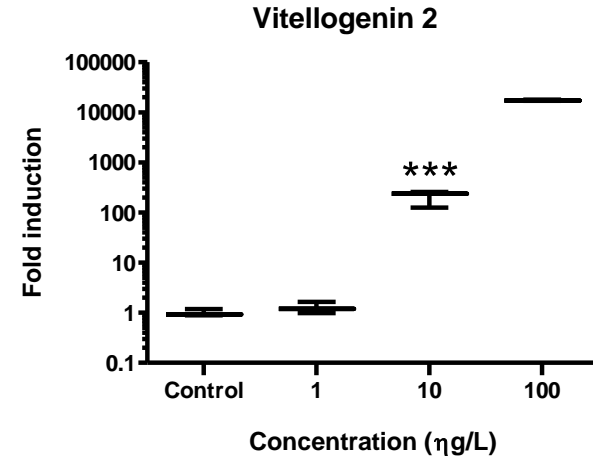
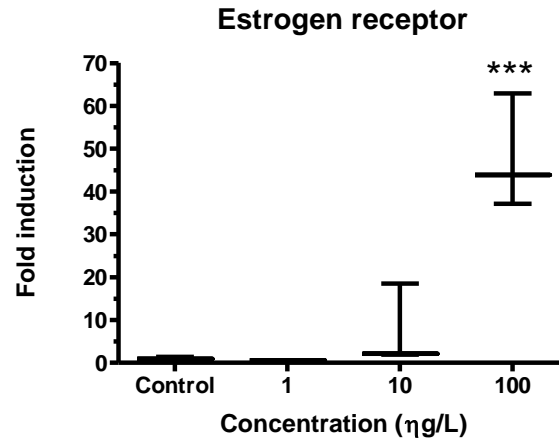
Global clustering (EE2)

- Control
- 1 ng/L
- 10 ng/L
- 100 ng/L

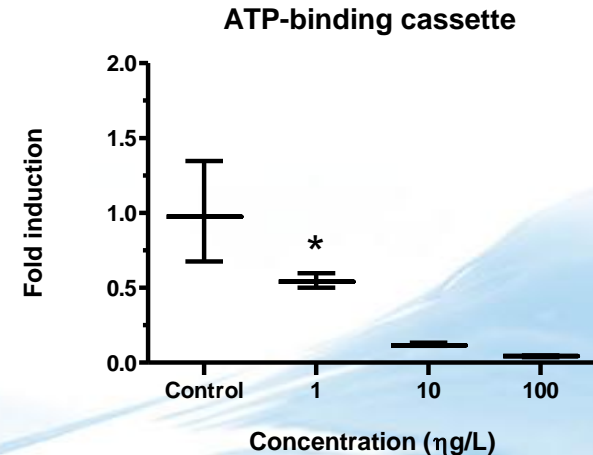
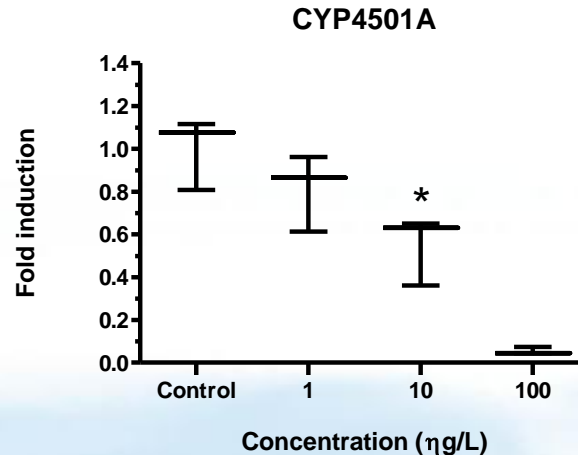


Single gene expression (EE2)

Up-regulation

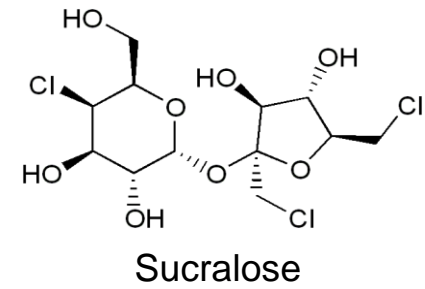


Down-regulation

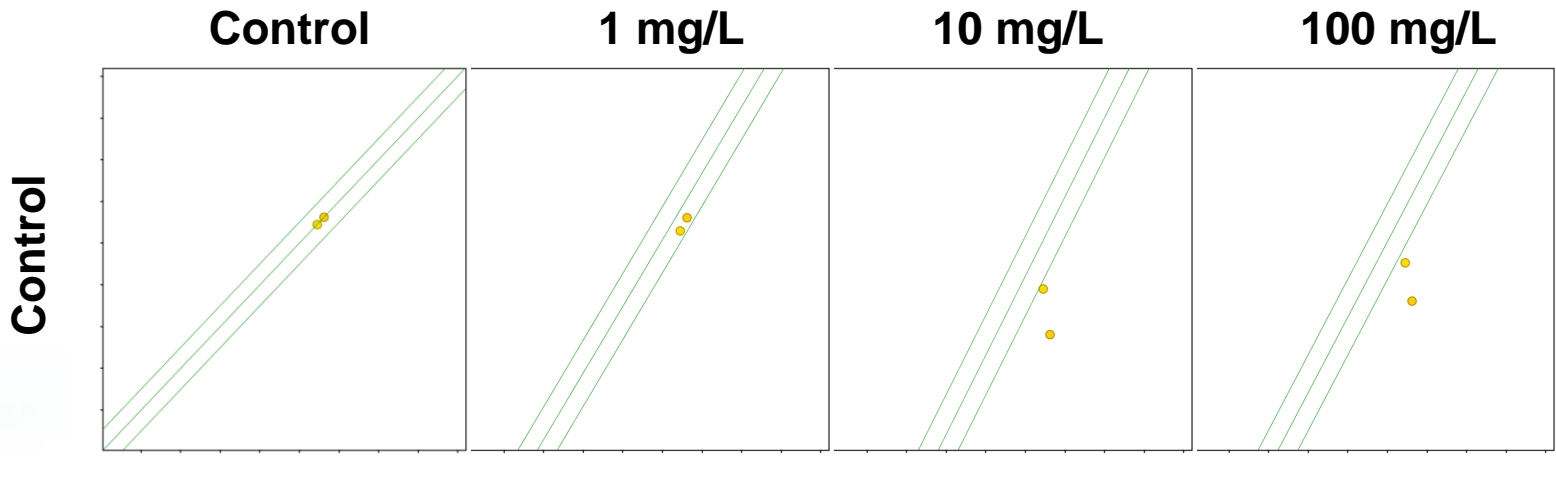




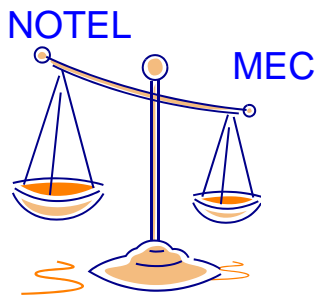
Intense sweetener



Sucralose



Differential gene expression in female zebrafish livers after 48h exposure (2 features)



Effects to be expected?

17 α -ethynylestradiol (EE2)

Environment (MEC)= 1-15 ng/L

NOEC_{acute} for fish $\geq 1 \mu\text{g/L}$

NOTEL: 1-10 ng/L

MEC/NOTEL ratio: 1-15

Sucralose

Environment (MEC)= $\sim 50 \text{ ng/L}$
(effluent: 3 $\mu\text{g/L}$)

NOEC_{acute} for fish $\geq 1800 \text{ mg/L}$

NOTEL: 10-100 mg/L

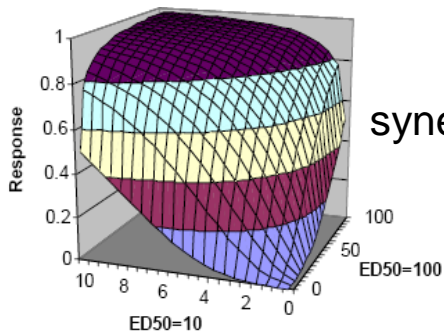
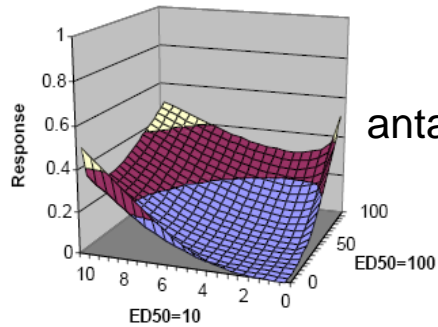
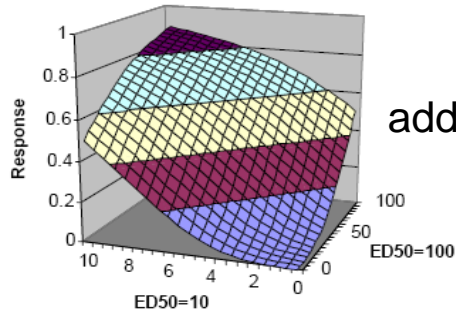
MEC/NOTEL ratio: $0.5-5 \times 10^{-7}$
(Effluent: $0.3-3 \times 10^{-4}$)

MEC: measured environmental concentration, NOTEL: No Observed Transcriptional Effect Level

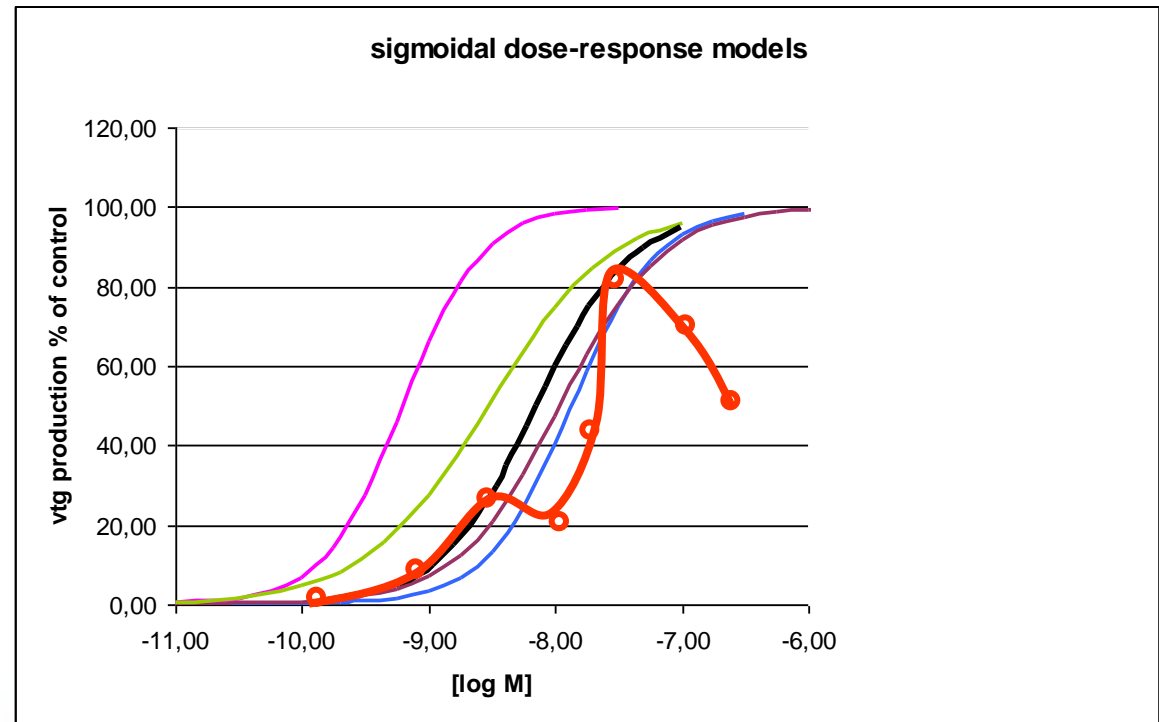
Conclusion

- MEC/NOTEL ratio for EE2 indicating transcriptional responses possible
- MEC/NOTEL ratio for Sucralose indicating large margin of safety

In vitro - mixture toxicity

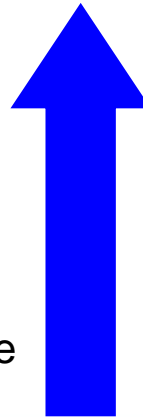
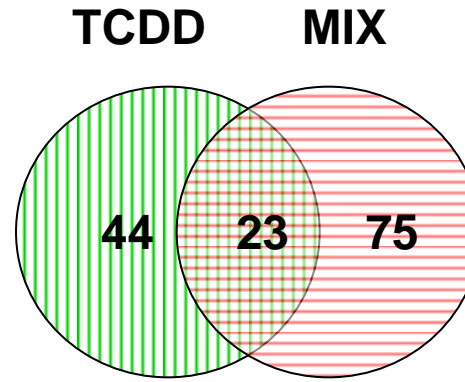
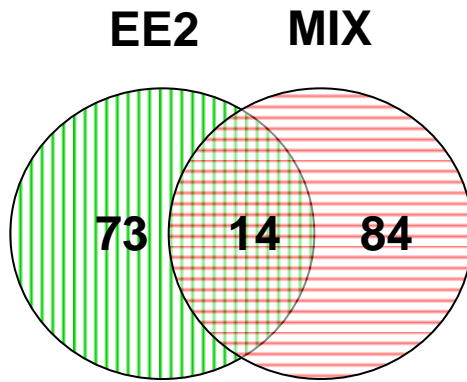
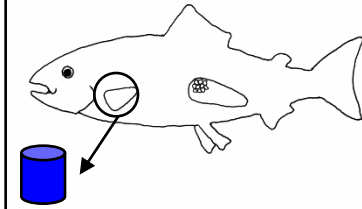


(Eide, 2004)



Multi-compound concentration-addition prediction model applied to experimental data from rainbow trout hepatocytes

In vitro gene expression

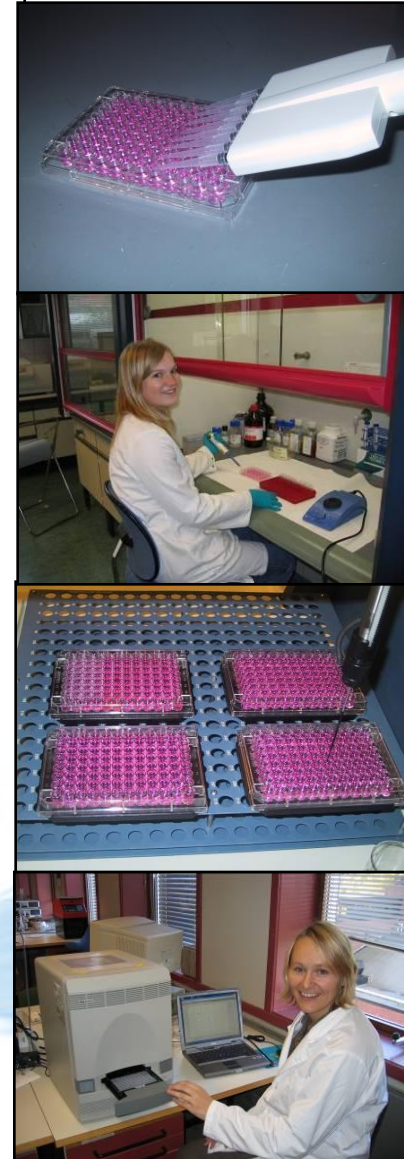


Single & Mix (equi-potent)

10 nM EE2
0.75 nM TCDD
0.75 μ M 4-Nitroquinoline-1-oxide
100 μ M Paraquat

Single chemical & combined toxicity assessment may require genome-wide response assessment

Finne, E.F. et al. (2007). *Aquat. Toxicol.* 81: 293-303



Regulatory positions (NO)

The Norwegian State Pollution Authority (SFT) support the EU (REACH) position on alternative testing methods:

*The Commission, Member States, industry and other stakeholders should continue to contribute to the promotion of alternative test methods on an international and national level including **computer supported methodologies, in vitro methodologies**, as appropriate, those based on **toxicogenomics**, and other relevant methodologies.*

(Source: REACH 40 introductory speech [fortale] for IR & CSA)

How useful are transcriptomics?

Advantages:

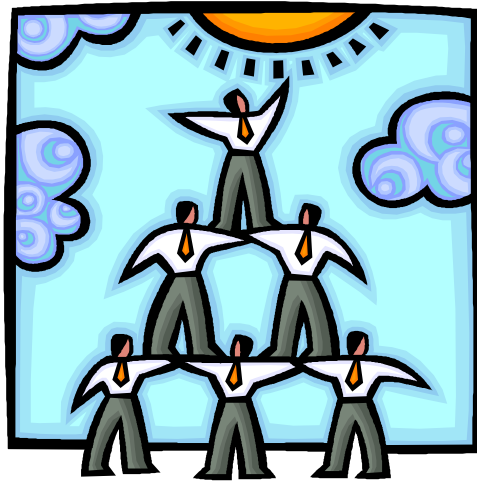
- High-throughput screening
- Hypothesis-generating
- Provide MoA based knowledge
- *in vivo* & *in vitro* applications
- Single chemical toxicity
- Mixture toxicity assessment
- Diagnostic/predictive tools
- Direct in-depth studies?

Disadvantages:

- Few tools for non-traditional species
- Few standardized guidelines & protocols
- Biostatistics/bioinformatics challenging
- Temporal variation in transcriptome
- Inter-assay reproducibility
- Whole organism responses
- Not regulatory validated nor accepted

Are we able to link transcriptome (biomarker) effects to adverse effects in RA?

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