Transcriptomic profiles of a respiratory inhibitor and growth targeting insecticide reveal links to impaired bone mineralization and lipid homeostasis in zebrafish embryos

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SETAC Europe 2022 – Platform Presentation Speaker: Hannes Reinwald

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SETAC

Classical Toxicology



OECD 202, 203, 221, 236, ...

- Non-human animal tests
- Time & cost intensive
- Not able to detect hazardous and regulatory relevant toxic MoAs (e.g. EDCs)

Transition to understanding

Predictive Toxicology



Requires qualitative toxicological data

OMICs enable scientists to assess the responses of tens of thousands of genes and their products from a single sample.







Overall research goal





- Identification of early affected key pathways associated with the respective MoA.
- Providing high content ecotoxicogenomic data for future data mining and meta-studies. (Also public for the scientific community on ArrayExpress)





Ecotoxicogenomic fin(ger)-printing workflow setup







 Table 1: Tested insecticides with respective mode of action in the target organism and test concentrations.

 (*:96h acute fish test NOEL values median from EPA's Ecotox database; **: nominal conc.)

Substance	Mode of Action (MoA)	NOEL*	Test conc. [µg/L]**	
	(IRAC classification)		Low (LE)	High (HE)
Fenazaquin	Mitochondrial complex I electron transport	9,6 ppm	3	6
Pyriproxyfen	Juvenile hormone mimic	270 ppm	170	1700



Publicly available toxicity data from EPA's knowledgebase in cyprinidae & cypriniformes for 96h & 120h exposure compiled via *StandaRtox* (Scharmüller et. al 2020).









Reduced dimensional clustering of biological samples





t-SNE clustering for the 1500 gene transcripts with largest variance with rlog transformed DESeq2 normalized gene counts.

Clearly separated clusters of sample groups (biological replicates).









Differential expression signals for the different exposure conditions







Ecotoxicogenomic signatures in the 96 hpf zebrafish embryo





Ecotoxicogenomic signatures in the 96 hpf zebrafish embryo





Science of The Total Environment Volume 735, 15 September 2020, 139496



Pyriproxyfen induced impairment of reproductive endocrine homeostasis and gonadal histopathology in zebrafish (*Danio rerio*) by altered expression of hypothalamus-pituitary-gonadal (HPG) axis genes

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TAKE HOME MESSAGE





(?)

Although OMICs are a powerful tool to gain mechanistic insights about adverse MoAs, what are these insides worth?

- Are the observed ecotoxicogenomic signatures **MoA-specific** in the applied model?
- Are the observed profiles **reproducible among similar MoA-targeting compounds**?



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Conclusion

• Combining *omics* with OECD standardized test methods has the potential to characterize molecular profiles in aquatic non-target organisms.

Conclusion & Outlook

- Such profiles provide mechanistic insights and allow for identification of MoA-specific molecular biomarkers.
- Molecular biomarker-based screening approaches offer:
 - Cost and resource effectiveness
 - High sensitivity
 - Transition to *in vitro* cell culture based methods as alternative to non-human animal testing.



regulation of ossification

lipid homeostasis









THANK YOU FOR YOUR INTEREST!

TIME FOR QUESTIONS NOW!

