7th German Pharm-Tox Summit – Combined Online Advanced Course of the Working Groups "Regulatory Toxicology" and "Computational Toxicology" of the Society of Toxicology (GT)



Application of AOPs to support read-across assessments

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Case study - > AIM of EU ToxRisk-project

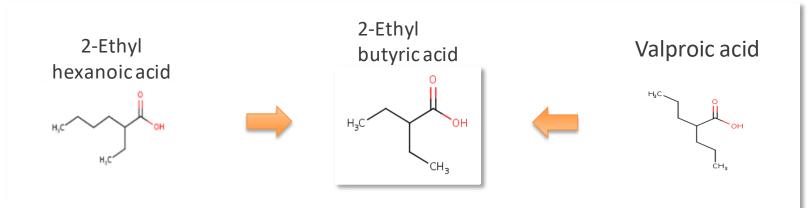
 Learn how to apply NAM in a human risk assessment for mainly two endpoints repeated dose toxicity and reprotoxicity

 Develop case studies as tool to gain experience for different aspects of regulatory decision-making

EUTOXRISK Read-Across case studies — Why?



(1) Why Read-Across Case Studies - the Regulatory Need



To date - Read-across is possible in case of similar toxicodynamic and kinetic properties (or consistent trend)

Read-Across rarely accepted by regulatory authorities

- Based often on structural & physicochemical data
- Lack of sufficient evidence to substantiate read-across justificationsfail to demonstrate toxicokinetic and toxicodynamic similarities
 - → Including lack of endpoint data on analogues provided in dossier
- Lack of scientific plausibility
 - → Disagreement with hypothesis, data not supportive of arguments presented, high uncertainty
 - → coupled with lack of evidence



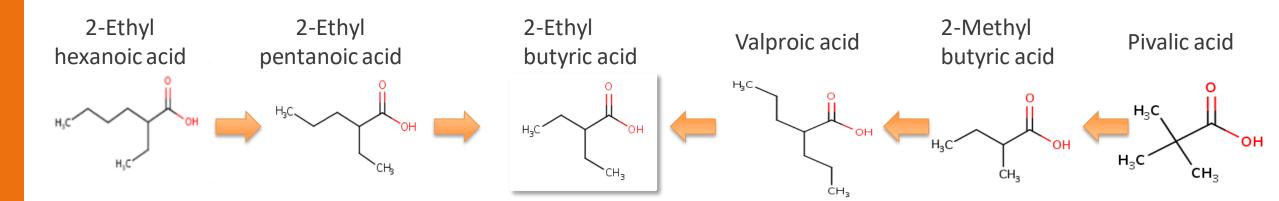
t⁴ report*

Toward Good Read-Across Practice

Nicholas Ball 1§, Mark T. D. Cronin 2§, Jie Shen 3§, Kare Mounir Bouhifd 6, Elizabeth Donley 7, Laura Eq How can NAM data Andre Kleensang⁶, Nicole Kleinstreuer⁹ Alexandra Maertens ⁶, Sue Mar<u>tv</u> a Palmer 7, David Pamies 6, Mike Penman ¹², Andrea-Nicole P , Sharon B. Stuard⁴, Grace Patlewicz ¹⁴, 2 Lhu ¹³ and Thomas Hartung ^{6,15}



(2) Why Read-across? Compare New Data to Traditional in vivo Data



NAM - Opportunities

- testing of a series of potential analogues, not limited to analogue with in vivo endpoint data
- trends can be investigated more comprehensively
- test human models
- provide mechanistic information
- help to understand kinetic properties and (dis)similarities

NAM - Challenges

- scope of in vitro testing + in silico prediction? What is good enough?
- integration of different types of information
- resulting relevance and predictivity of result
- uncertainty assessment

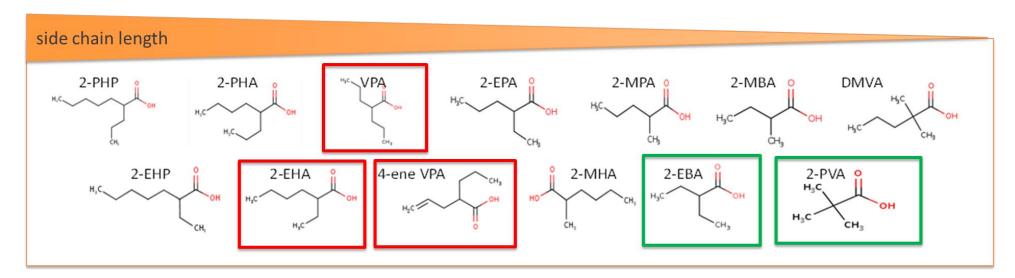


Case study 1— RAX hypothesis for systemic toxicity, lead effect liver steatosis

Core structure

$$R_1$$
 R_2
 R_3
 R_1 , R_2 = linear alkyl

 R_3 = CH_3 or H

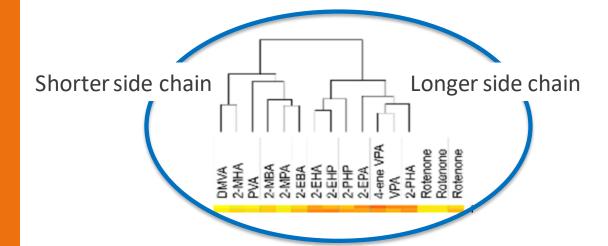


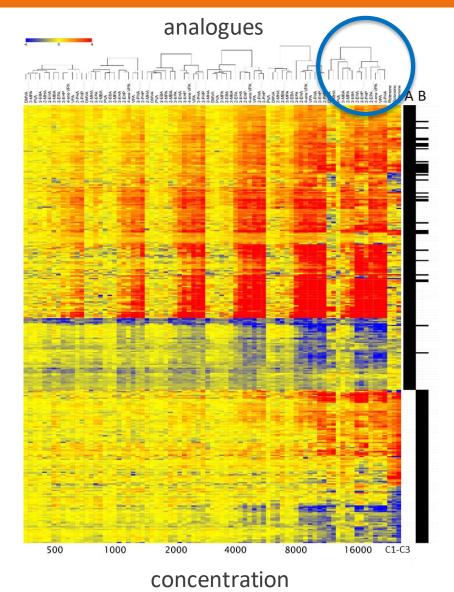
- liver steatosis, in peclinical rodent studies
- no liver effect observed up to highest in vivo tested dose



Analysis of DEGs - first indication of biological similarity

- Dose dependent testing of HepG2 cells
- TempOseq 1500⁺ panel (about 3500 genes)
- Expression profile of carboxylix acids differ from rotenone (michondrial complex I inhibitor)
- Analogues with longer side chain are more active and cluster together



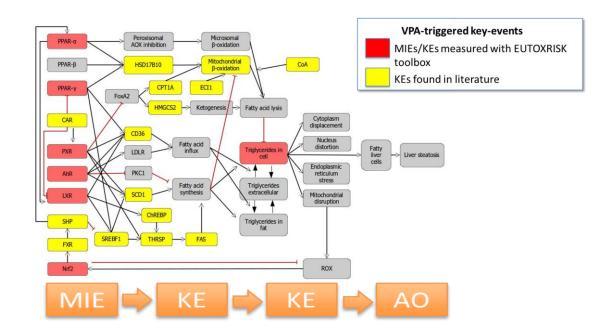


A- DEGs of carboxylic acids

B- DEGs of rotenone



Learn to use AOPs AOP network for liver steatosis



AOP Network

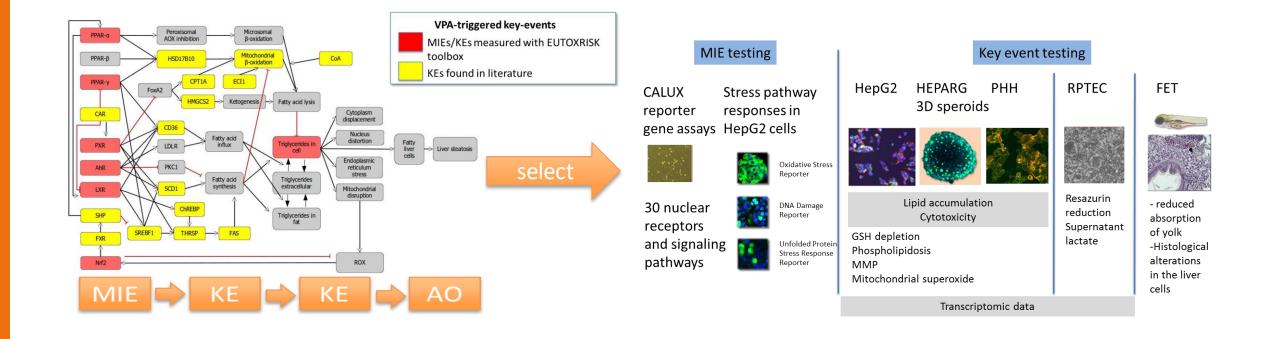
AOP network

- comprise evidence from 55 different AOPs
- Coloured boxes Evidence known for VPA
- Red- MIEs and KE tested in EUTOXRISK in vitro toolbox
- is used to infrom the testing strategy



Learn to use AOPs AOP network for liver steatosis

AOP Network



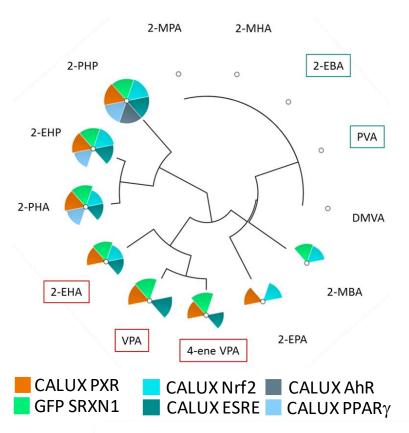
[::::] EUTOXRISK

Targeted in vitro testing battery

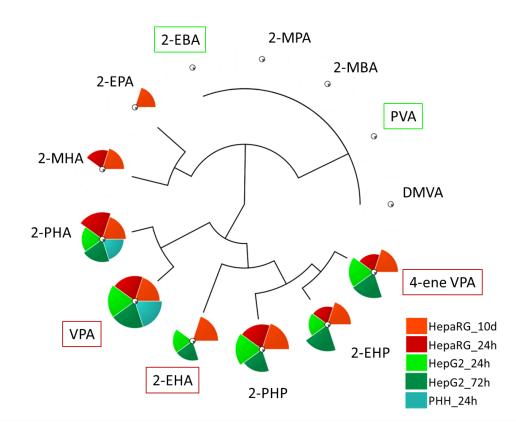
Learn to use AOPs in Read-Across Context



MIE/early KEs - reporter gene assays



Late KE – lipid accumulation

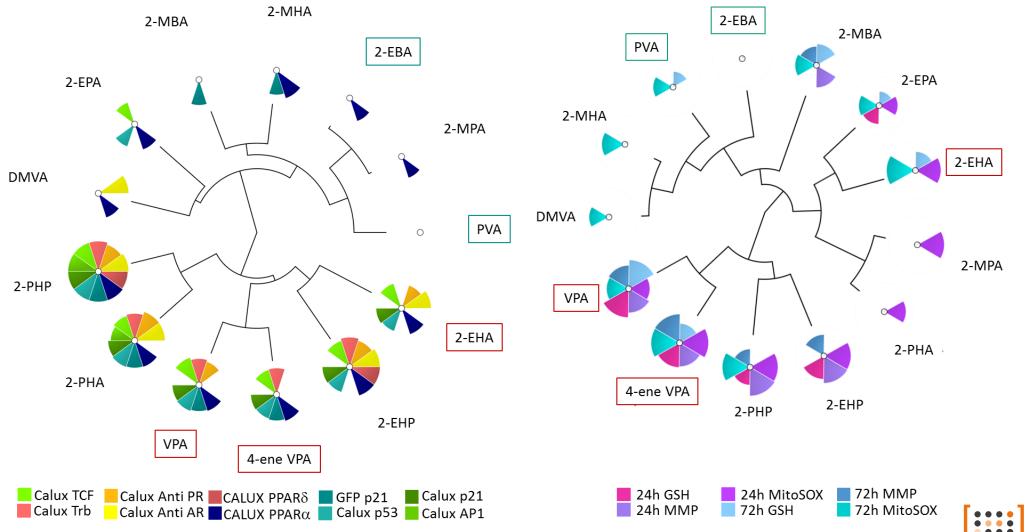


MIEs and Lipid accumulation – increased activity with increasing side chain length; in vivo pos. and neg. compounds predicted correctly

MIEs and KEs that do not belong to the AOP show no trend

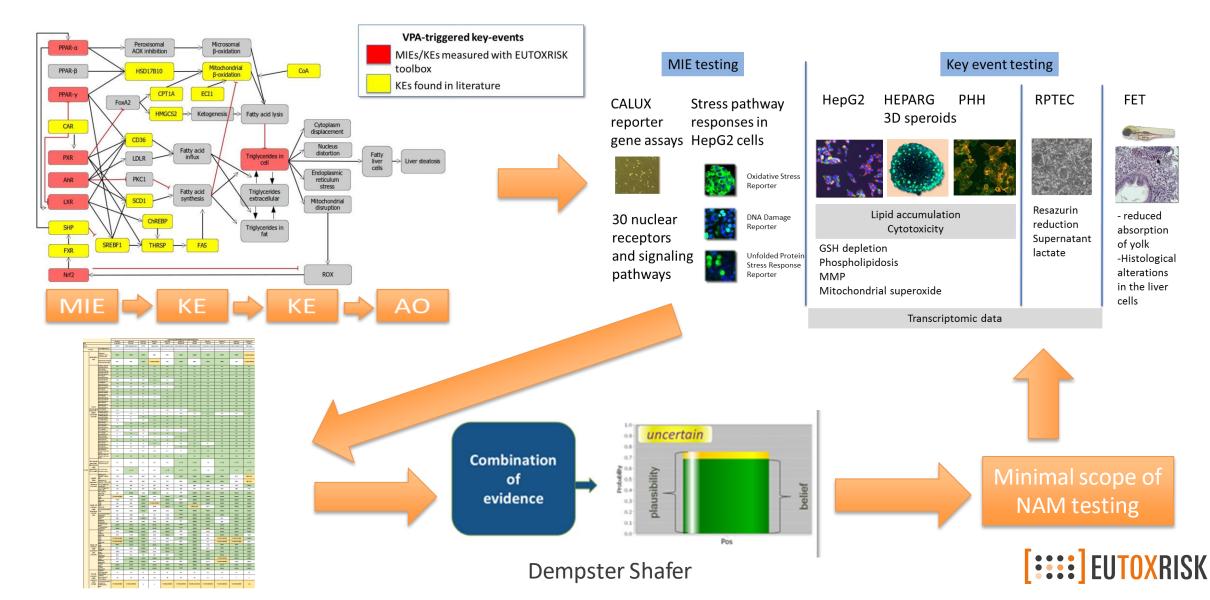
MIE/early KEs not in AOP

Mitochondrial dysfunction (HepG2 cells)



Learn to use AOPs AOP network for liver steatosis





Read-Across Supported by NAMs -Toxikodynamics

NAMs used to illustrate shared mode of action

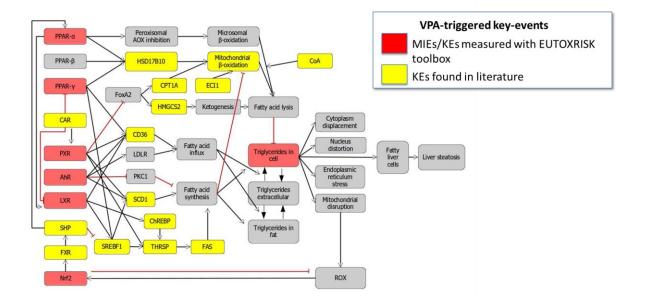
- 2-EPA show activation of MIEs and KEs belonging to AOP, induces late KE "lipid accumulation" in different liver cells
- Early MIEs/KEs can be used to prove similar mode of action



First learnings – application of AOPs in reg risk assessment

- AOP-based testing strategy
 - AOP-network -> no need to test all
 MIEs/KEs; test shared toxicological profile
 - include KE close to apical endpoint
 - data integration might be challenging

If no AOP available/ AOP weak: Describe the scientific rational of the testing in detail



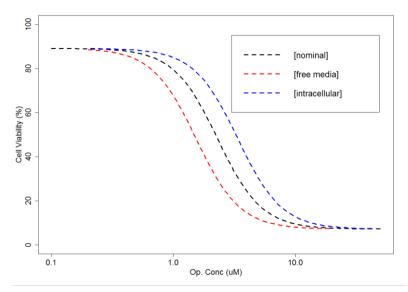


Next step – In Vitro to In Vivo Extrapolation

What is the free concentration of the test compound in the cell?

- Is the effective concentration in vitro translated correctly to the in vivo concentration?
- Do we have confounding factors? Like volatility? Binding properties (to protein/serum; to plastics...)?
- ADME properties are dependent on ionization status, pH of medium.
- RAX- do we have large differences between the grouped compounds?

In Vitro Biokinetics







Predicting In Vitro Distribution – VIVD Model

$$C_{media,dissolved,u} = \frac{C_{nominal} \cdot fu_{diluted} \cdot V_{media} \cdot 1e^{-3}}{V_{bulk} + k_{air}f_{ui}V_{air} + k_{cell,u}V_{cell} + k_{plastic}SA_{media} \cdot 1e^{3}}$$



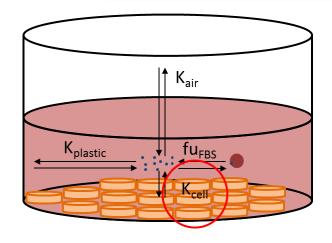
Intracellular Concentration (M) logP_{ow} VIVD: Virtual *in vitro* distribution model for the mechanistic prediction of intracellular concentrations of chemicals in *in vitro* toxicity assays

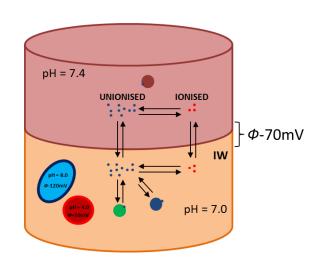


Fisher C. a, Siméon S. b, Jamei M. a, Gardner I. Bois Y.F. b

a Certara UK Limited, Simcyp Division, Acero, 1 Concourse Way, Sheffield S1 2BJ, UK

b INERIS, METO Unit, Verneuil en Halatte, France

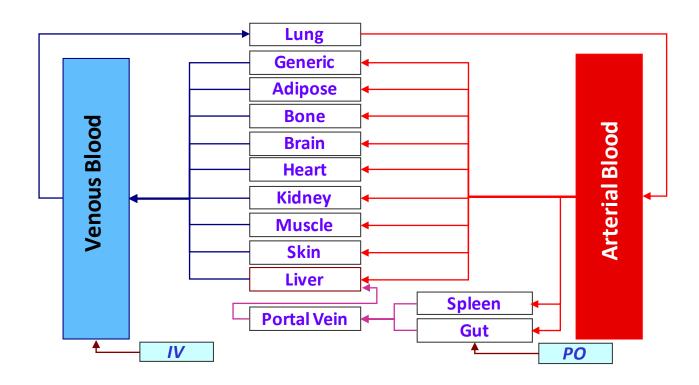








Physiologically-based Pharmacokinetic Modelling

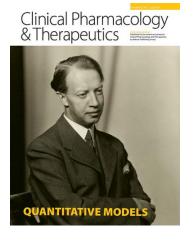


Nothing New:

Teorell, T. Studies on the diffusion effect upon ionic distribution: II. experiments on ionic accumulation. J. Gen. Physiol. 21, 107–122

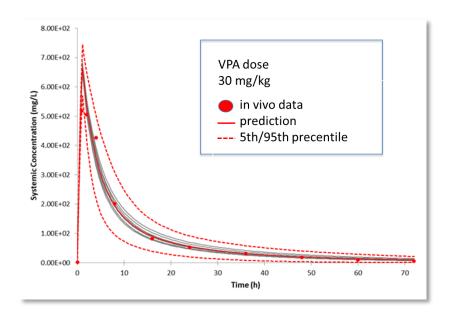
(1937)

- octanol:water partition coefficient (logP_{ow})
- pKa
- PSA(Ų), HBD
- blood to plasma ratio (B:P)
- plasma protein binding (fu)
- fraction absorbed (fa)
- first-order absorption rate constant (Ka)
- steady-state volume of distribution (V_{ss})
- intrinsic hepatic clearance (CL_{int,Hep})





VPA – data rich compound with in vivo ADME data



PBPK model predicts in vivo human data for one analogue well

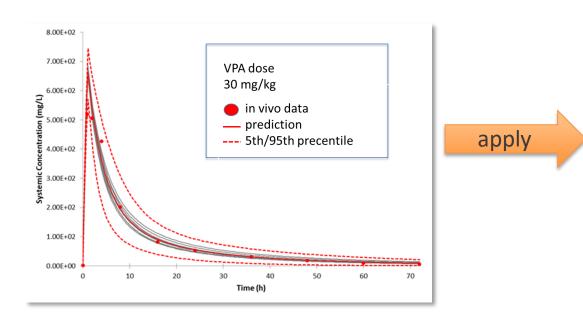
NAM derived – ppb (in silico) and intr. hep clearance in PHH (in vitro)

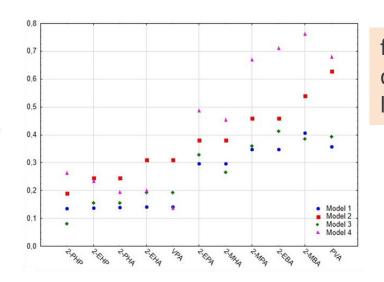


VPA model used for all analgues in RAX group



Toxikokinetic properties within grouped compounds





fu increase with decreasing side chain length

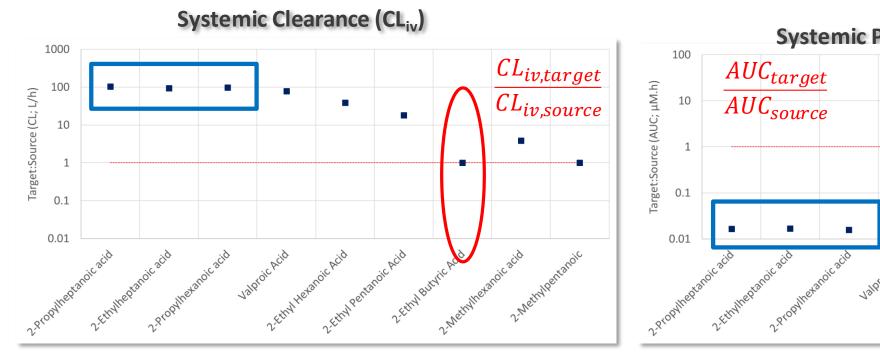
Hepatic clearance decreases with side chain length

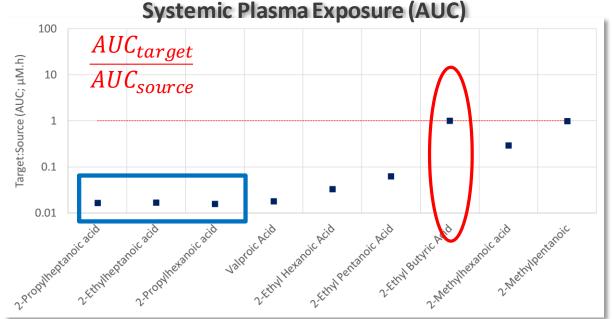
	99-66-
	VPA
CL _{int,H} (μl/min/10 ⁶ ; HμREL co-culture)	0.22

149-57-5	20225-24-5	88-09-5	97-61-0	4536-23-6
2-EHA	2-EPA	2-EBA	2-MPA	2-MHA
0.55	0.78	9.62	10.2	3.95



RAX Case Study - PBPK Predictions Across Analogues

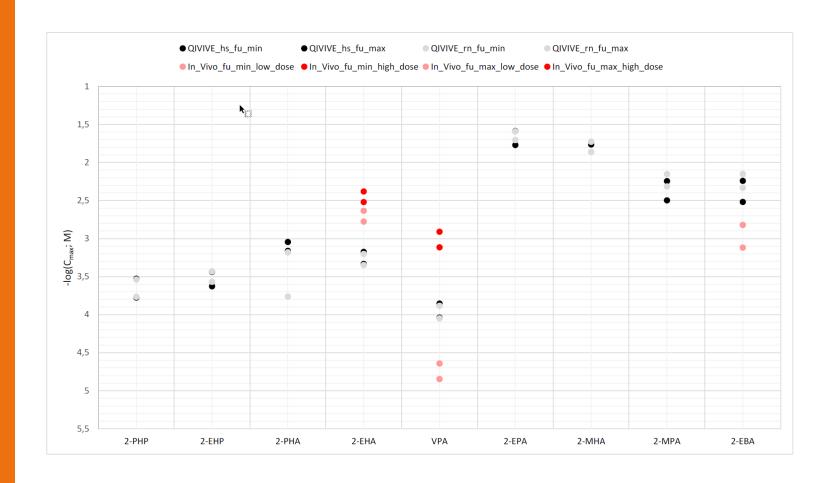




- Three source compounds show intrinsic hepatic clearance (CL_{int}) below limit of detection of *in vitro* assay (Hurel co-culture system); CL_{int} (μl/min/10⁶ hepatocytes) assumed to be 2-fold lower than VPA
- Trend observed short chain target compound 2-EBA shows comparable predicted exposure and clearance to 2-MHA and 2-MPA



Estimate human equivalant dose –bridge back to AOP



Comparison of max. plasma concentration

- QIVIVE (red)
- Reverse dosimetry from rodent study (black)

Late KE "triglyceride accumulation corresponded best to in vivo situation"

MIEs/early KE – lower heD



Read-Across Supported by NAMs –Toxikodynamics + Toxikokinetics

AOP used to illustrate shared mode of action

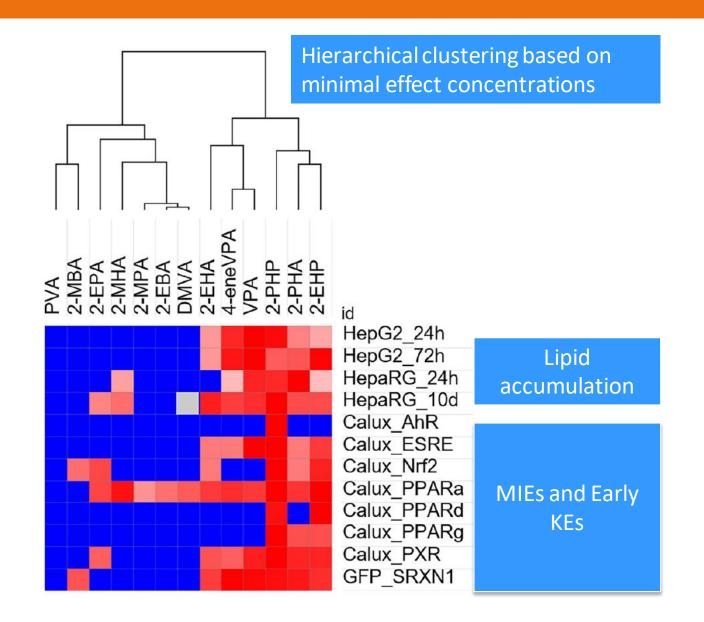
- 2-EPA show activation of MIEs and KEs belonging to AOP, induces late KE "lipid accumulation" in different liver cells
- Early MIEs/KEs can be used to prove similar mode of action

In vitro ADME assays as well as PBK modelling used to show trend in toxikokinetics

- PBK simulations for all analogues identify a trend for increasing clearance and so decreasing systemic exposure with decreasing side chain length
- Late KE shows best predictivity compared to early MIEs and KEs



Read-across - options



Option 1

 NAM can be used to idnetify nearest neighbours; in vivo data from them to read-across

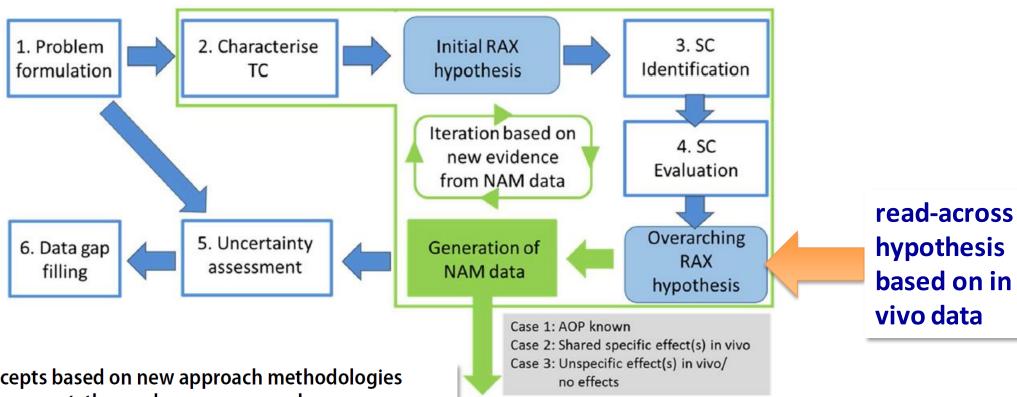
Option 2

 QIVIVE can be used to directly predict the human threshold



Result used to develope a Read-Across Assessment Schema

From several case studies



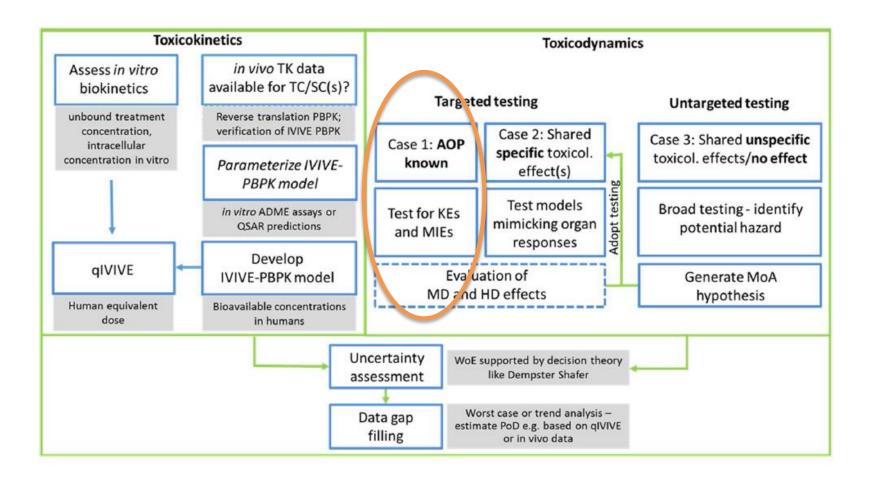
Towards grouping concepts based on new approach methodologies in chemical hazard assessment: the read-across approach of the EU-ToxRisk project

Sylvia E. Escher¹ · Hennicke Kamp² · Susanne H. Bennekou³ · Annette Bitsch¹ · Ciarán Fisher⁴ · Rabea Graepel⁵ · Jan G. Hengstler⁶ · Matthias Herzler⁷ · Derek Knight⁸ · Marcel Leist⁹ · Ulf Norinder¹⁰ · Gladys Ouédraogo¹¹ · Manuel Pastor¹² · Sharon Stuard¹³ · Andrew White¹⁴ · Barbara Zdrazil¹⁵ · Bob van de Water⁵ · Dinant Kroese¹⁶



Generation of NAMs based on Read-Across Hypothesis –

NAM used as supporting evidence to proof shared toxicodynamic and kinetic properties





Summary

- Read across case studies can be used to gain confidence in the use of non-formally standardised new approachs such as AOPs.
- In RAX in vivo endpoint data compared to the NAM based predictions -> in situ validation of formally non standardised NAM
- AOPs are extremely useful tools to inform the testing strategy
- Not all MIEs and KEs need to be tested
- In our case late KE was most appropriate to predict the human equivalent dose

Vrijenhoek NG et al. (2022) ALTEX, doi:10.14573/altex.2107261 Escher SE et al. (2022) Toxicology in Vitro 79, 105269 Kamp et al. Read across advisory document, under preparation Fischer C et al., qIVIVE, under preparation

Read-across team **AOP**

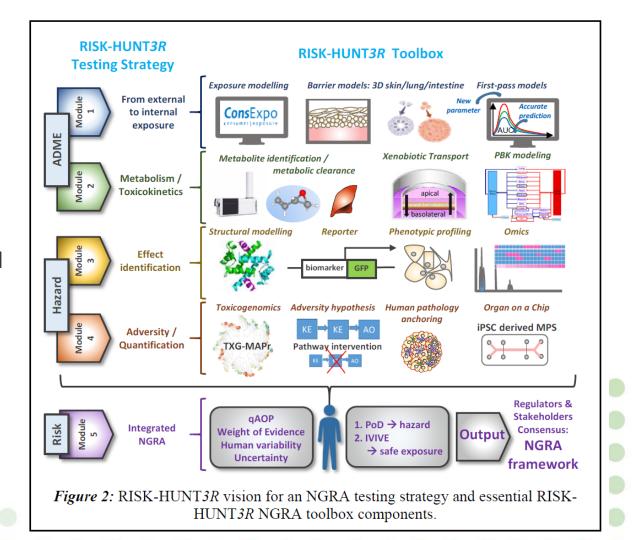


End of story?



Objectives

- Include metabolism in barrrier organs and better models for in vitro ADME
- Complete AOP landscape
- Move towards quantitative AOPs
-





Thanks for your attention

