

A Population-Based Human *In Vitro* Approach to Characterize Inter-Individual Variability in Responses to Chemical Mixtures

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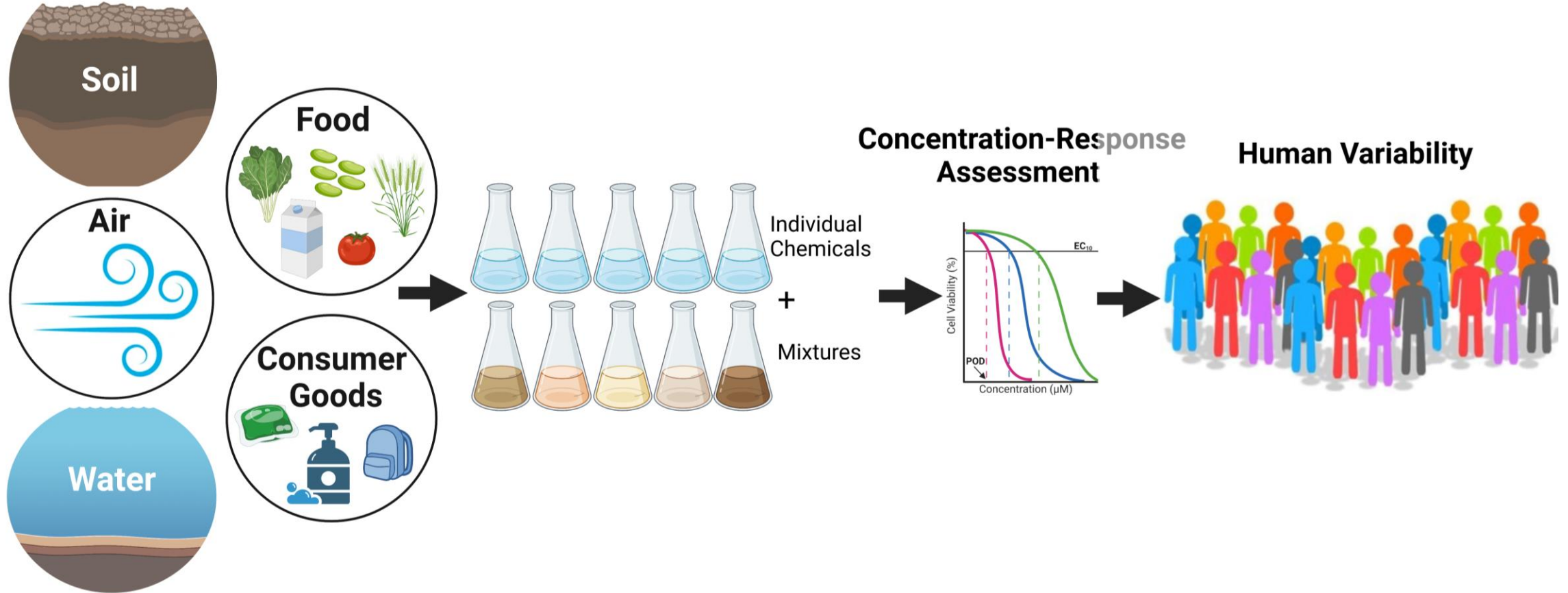
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Various Exposures and Unknown Outcomes

Sources of Exposures



How can mixtures be assessed?

Current challenges in mixtures risk assessment:

- 1) Health effects
 - 2) Unknown composition of the mixtures
 - 3) Exposure assessment
- Existing risk assessment methods rely on data from individual chemicals
 - No standardized approach to assess risk of mixtures

Two proposed methods:

- 1) Whole-mixture approach
- 2) Component-based approach

GUIDANCE

ADOPTED: 17 November 2021
doi: 10.2903/j.efsa.2021.7033

Guidance Document on Scientific criteria for grouping chemicals into assessment groups for human risk assessment of combined exposure to multiple chemicals

EFSA Scientific Committee,
Simon John More, Vasileios Bampidis, Diane Benford, Claude Bragard, Antonio Hernandez-Jerez, Susanne Hougaard Bennekou, Thorhallur Ingi Halldorsson, Konstantinos Panagiotis Koutsoumanis, Claude Lambré, Kyriaki Machera, Hanspeter Naegeli, Søren Saxmose Nielsen, Josef Rudolf Schlatter, Dieter Schrenk, Vittorio Silano, Dominique Turck, Maged Younes, Emilio Benfenati, Amélie Crépet, Jan Dirk Te Biesebeek, Emanuela Testai, Bruno Dujardin, Jean Lou CM Dorne and Christer Hogstrand

Abstract

This guidance document provides harmonised and flexible methodologies to apply scientific criteria and prioritisation methods for grouping chemicals into assessment groups for human risk assessment of combined exposure to multiple chemicals. In the context of EFSA's risk assessments, the problem formulation step defines the chemicals to be assessed in the terms of reference usually through regulatory criteria often set by risk managers based on legislative requirements. Scientific criteria such as hazard-driven criteria can be used to group these chemicals into assessment groups. In this guidance document, a framework is proposed to apply hazard-driven criteria for grouping of chemicals into assessment groups using mechanistic information on toxicity as the gold standard where available (i.e. common mode of action or adverse outcome pathway) through a structured weight of evidence approach. However, when such mechanistic data are not available, grouping may be performed using a common adverse outcome. Toxicokinetic data can also be useful for grouping, particularly when metabolism information is available for a class of compounds and common toxicologically relevant metabolites are shared. In addition, prioritisation methods provide means to identify low-priority chemicals and reduce the number of chemicals in an assessment group. Prioritisation methods include combined risk-based approaches, risk-based approaches for single chemicals and exposure-driven approaches. Case studies have been provided to illustrate the practical application of hazard-driven criteria and the use of prioritisation methods for grouping of chemicals in assessment groups. Recommendations for future work are discussed.

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Keywords: harmonised methodologies, human risk assessment, combined exposure to multiple chemicals, scientific criteria, grouping, assessment groups, dose addition

Requestor: EFSA
Question number: EFSA-Q-2019-00517
Correspondence: sc.secretariat@efsa.europa.eu

as part of EFSA work programme, part of a roadmap for action and its

Assessment of multiple Chemicals

nts received from the T Agencies, EU Member ers in the phases 1 & 2 of ess

ne implementation of human health mains of activity. Harmonised tools and efficient assessment of both . Available methods will be fit-for- islations.

onised approach for the assessment non-dietary exposure to multiple d in four work packages: methods d implementation.

vious projects to identify relevant hods will then be translated into all domains of EFSA, and necessary . In a second instance, EFSA will ossible needs for further innovation osure assessment.

k assessment to multiple chemicals will first prioritise the routine risk th a subsequent integration of non- er chemicals will also be developed of pesticides, the scope of the

implementation plan will be extended to other chemicals.

- In view of increasing the outreach towards national competent authorities, EFSA will promote the implementation of the methods and tools developed and provide adequate training, where necessary.

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Why are we interested in population-based *in vitro* methods?

Traditional Toxicity Testing Methods: *In Vivo*

- Time and labor-intensive, expensive, and low throughput
- Challenges with extrapolation to humans
- Models often overlook inter-individual variability
- Ethical concerns



New Approach Methods for Toxicity Testing: *In Vitro*

- Faster, cheaper, and higher-throughput
- Ability to look at biologically-relevant phenotypes
- Can evaluate inter-individual and inter-species variability
- Reduces use of animal testing



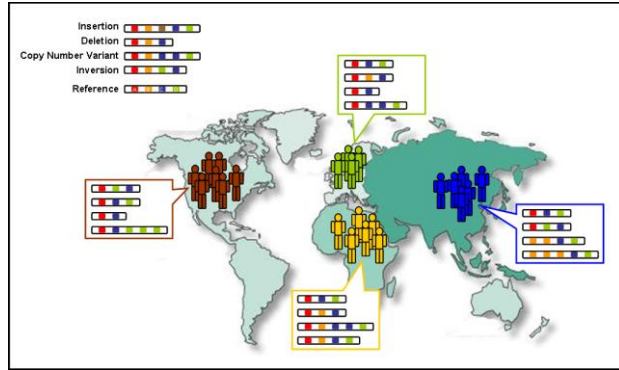
Population-Based Human *In Vitro* Models

- Human lymphoblast cell lines (1,000+ donors)
- Human induced pluripotent stem cell-derived cardiomyocytes (~43 donors)
- Assess inter-individual and chemical-specific variability
- Translation to humans

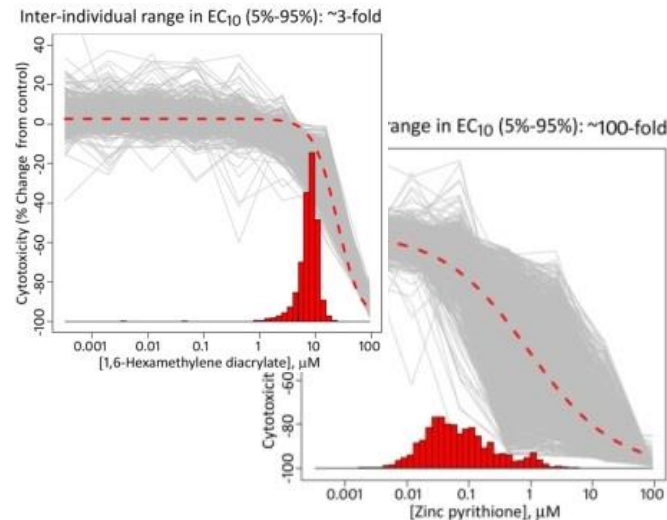


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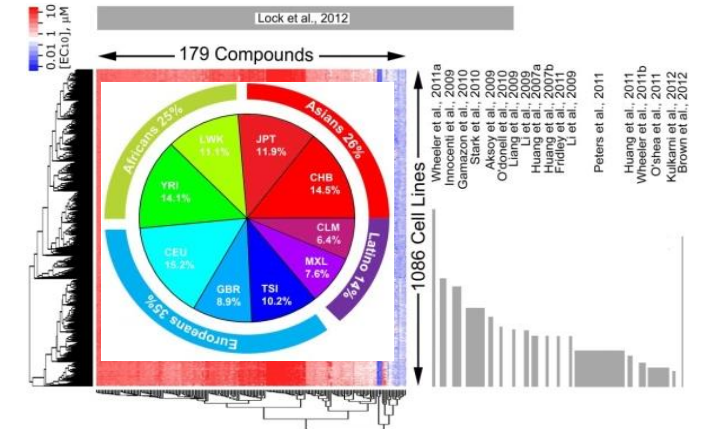
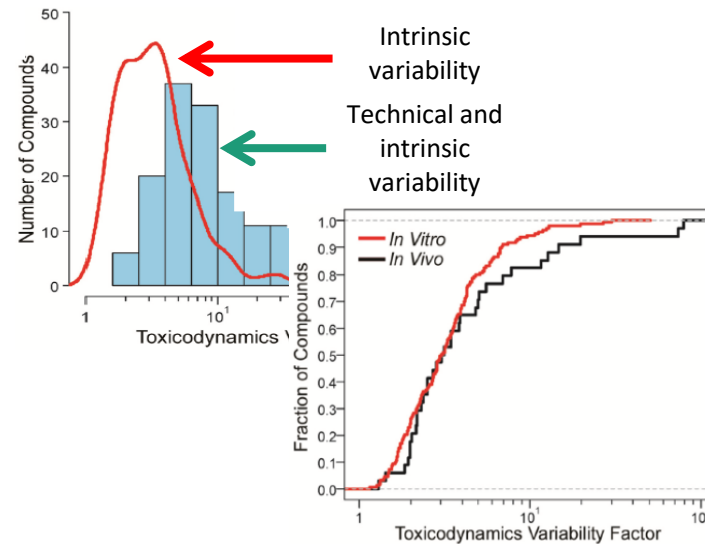
Why do toxicity testing in human lymphoblast cell lines?



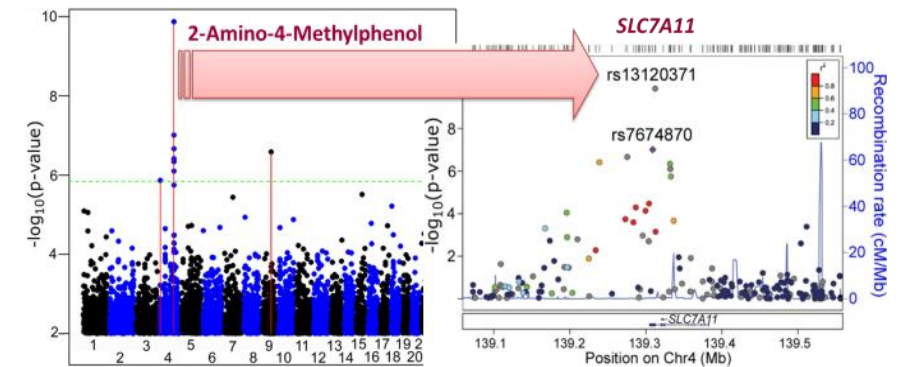
Hazard and Dose-Response Assessments



Ability to Assess Variability and Relevant to Humans



Mechanistic Hypotheses and Identification of Susceptible Genes



- Abdo et al [Environ Int](#) 85:147-55, 2015
- Abdo et al [Environ Health Perspect](#) 123(5):458-66,
- Eduati et al [Nat Biotechnol](#) 33(9):933-40, 2015
- Chiu et al [ALTEX](#) 34(3):377-388, 2017

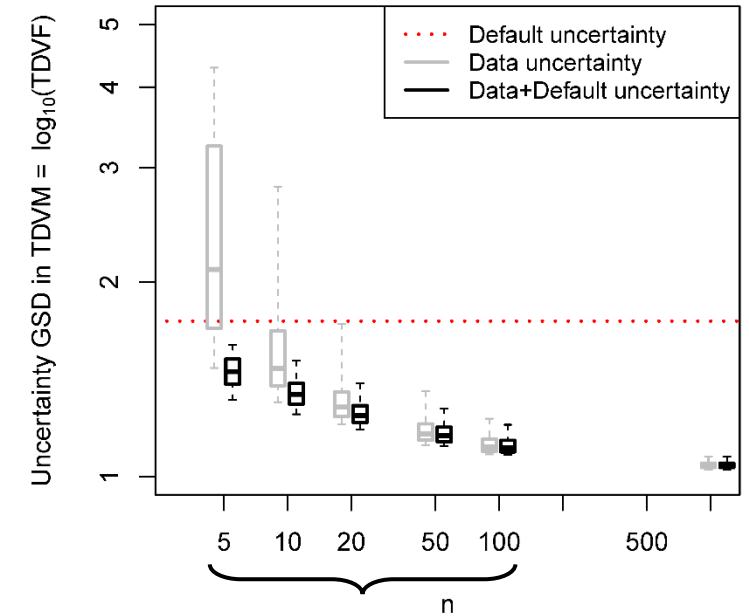
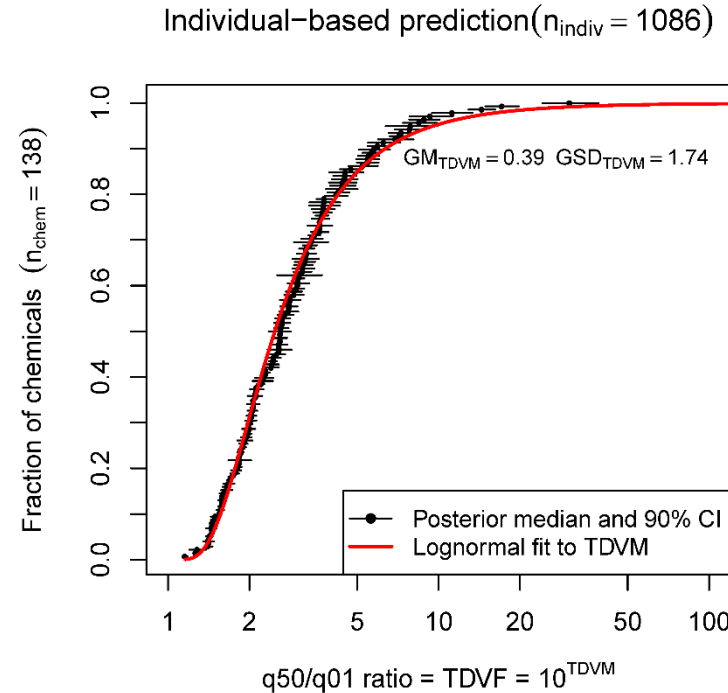


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Do we need 1000+ cell lines to study human variability?

How many individuals do we really need for *in vitro* screening?

- Power calculations based on data from Abdo et al (2015)
 - Resampled 1000+ individuals
 - Resampling across all tested chemicals



Sample sizes as small as 5 donors can be informative
Reliable estimates with 20-100 individuals



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Chiu et al [ALTEX](#) 34(3):377-388, 20172015

Slide adapted from Ivan Rusyn

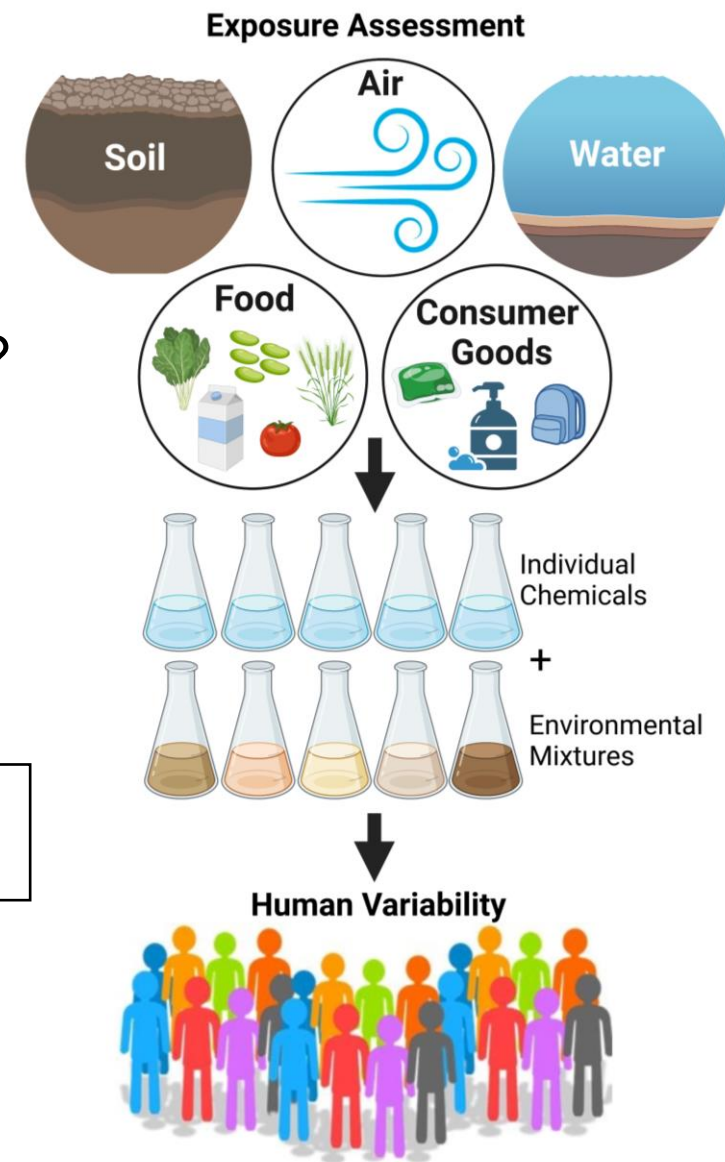
Why are we interested?

- Because there is no standard approach to test human variability in mixtures risk assessment
- Can we estimate the extent of population variability for mixtures?
 - Do we need to test both individual constituents and mixtures?
 - Is the extent of variability greater for mixtures than for chemicals?
 - Can we apply the same uncertainty factor to mixtures and chemicals?

What is our approach?

Apply population-based *in vitro* methods to assess potential toxicity of component-based mixtures

- Test defined mixtures and the individual constituents
- Use a population-based human *in vitro* model of LCLs
- Quantify toxicodynamic variability for chemicals and mixtures
- Identify potential drivers of variability through a GWAS



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Experimental Design

Lymphoblast cell lines from 4 populations (N = 146)



Testing of cytotoxicity in concentration response



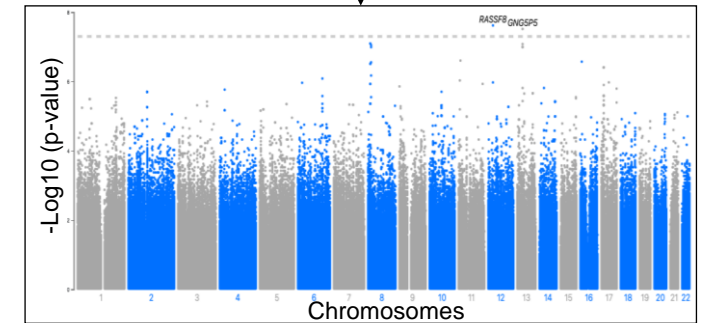
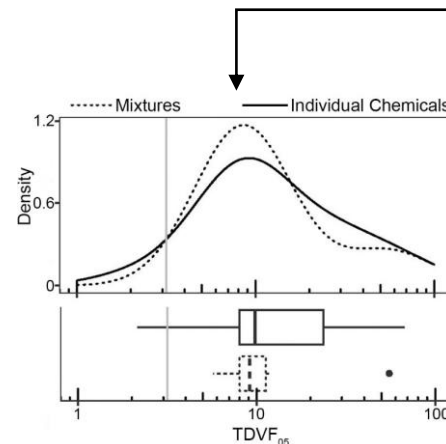
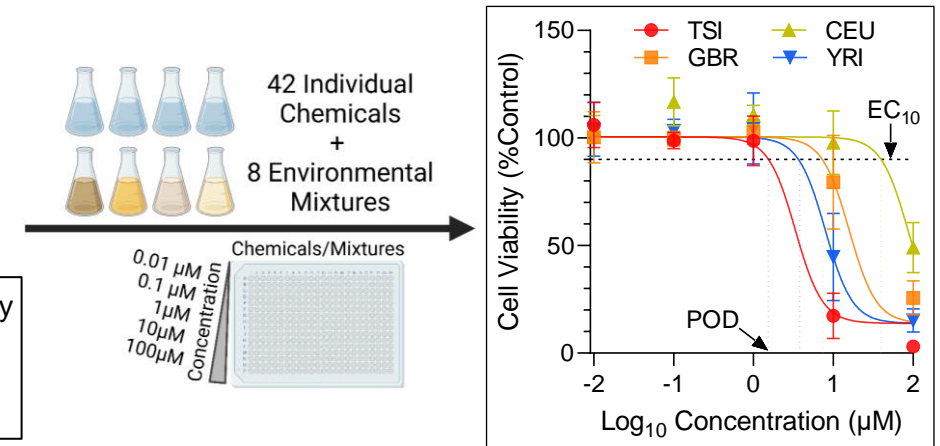
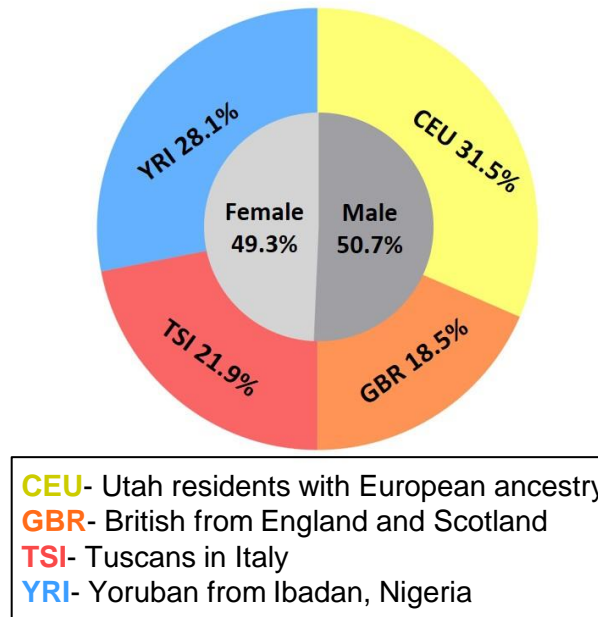
Cell line-specific concentration response modeling



Cumulative Distribution of $TDVF_{05}$



GWAS using cell line-specific PODs



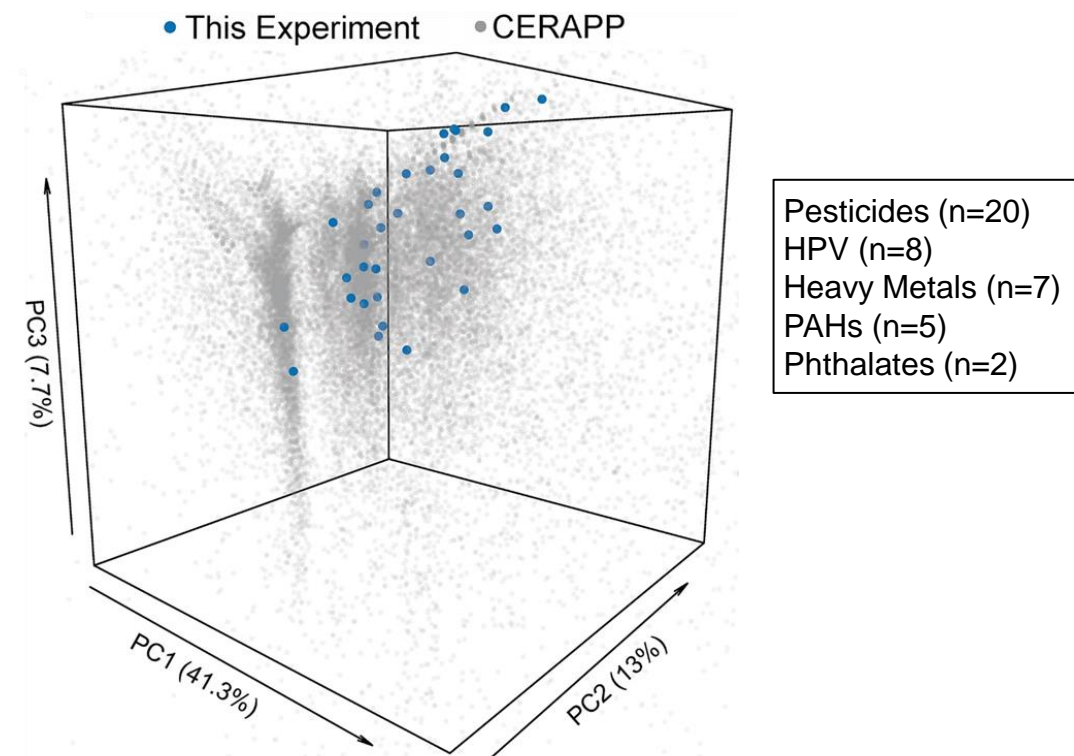
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Experimental Design

Chemical Selection and Design Mixtures

ID	SUBSTANCE NAME	CAS	Class
1	BENZO(A)ANTHRACENE	56-55-3	PAH
2	NAPHTHALENE	91-20-3	PAH
3	FLUORANTHENE	206-44-0	PAH
4	DDT, P,P'-	50-29-3	Pest
5	DIELDRIN	60-57-1	Pest
6	ALDRIN	309-00-2	Pest
7	HEPTACHLOR	76-44-8	Pest
8	LINDANE	58-89-9	Pest
9	DISULFOTON	298-04-4	Pest
10	ENDRIN	72-20-8	Pest
11	DIAZINON	333-41-5	Pest
12	HEPTACHLOR EPOXIDE	1024-57-3	Pest
13	PENTACHLOROPHENOL	87-86-5	HPV
14	DI-N-BUTYL PHTHALATE	84-74-2	Plastiz
15	CHLORPYRIFOS	2921-88-2	Pest
16	DI(2-ETHYLHEXYL)PHTHALATE	117-81-7	Plastiz
17	2,4,6-TRICHLOROPHENOL	88-06-2	HPV
18	ETHION	563-12-2	Pest
19	AZINPHOS-METHYL	86-50-0	Pest
20	2,4,5-TRICHLOROPHENOL	95-95-4	HPV
21	PARATHION	56-38-2	Pest

ID	SUBSTANCE NAME	CAS	Class
22	BENZO(B)FLUORANTHENE	205-99-2	PAH
23	TRIFLURALIN	1582-09-8	Pest
24	ACENAPHTHENE	83-32-9	PAH
25	DDD, P,P'-	72-54-8	Pest
26	BENZIDINE	92-87-5	HPV
27	ENDOSULFAN	115-29-7	Pest
28	METHOXYCHLOR	72-43-5	Pest
29	2,4-DINITROPHENOL	51-28-5	Pest
30	2,4-DINITROTOLUENE	121-14-2	HPV
31	DICOFOL	115-32-2	Pest
32	CRESOL, PARA-	106-44-5	HPV
33	DDT, O,P'-	789-02-6	Pesticide
34	4,6-DINITRO-O-CRESOL	534-52-1	HPV
35	1,2,3-TRICHLOROBENZENE	87-61-6	HPV
36	LEAD NITRATE	10099-74-8	Metal
37	CADMIUM CHLORIDE	10108-64-2	Metal
38	ZINC CHLORIDE	7646-85-7	Metal
39	MERCURIC CHLORIDE	7487-94-7	Metal
40	POTASSIUM CHROMATE	7789-00-6	Metal
41	COBALT CHLORIDE	7646-79-9	Metal
42	NICKEL CHLORIDE	7718-54-9	Metal



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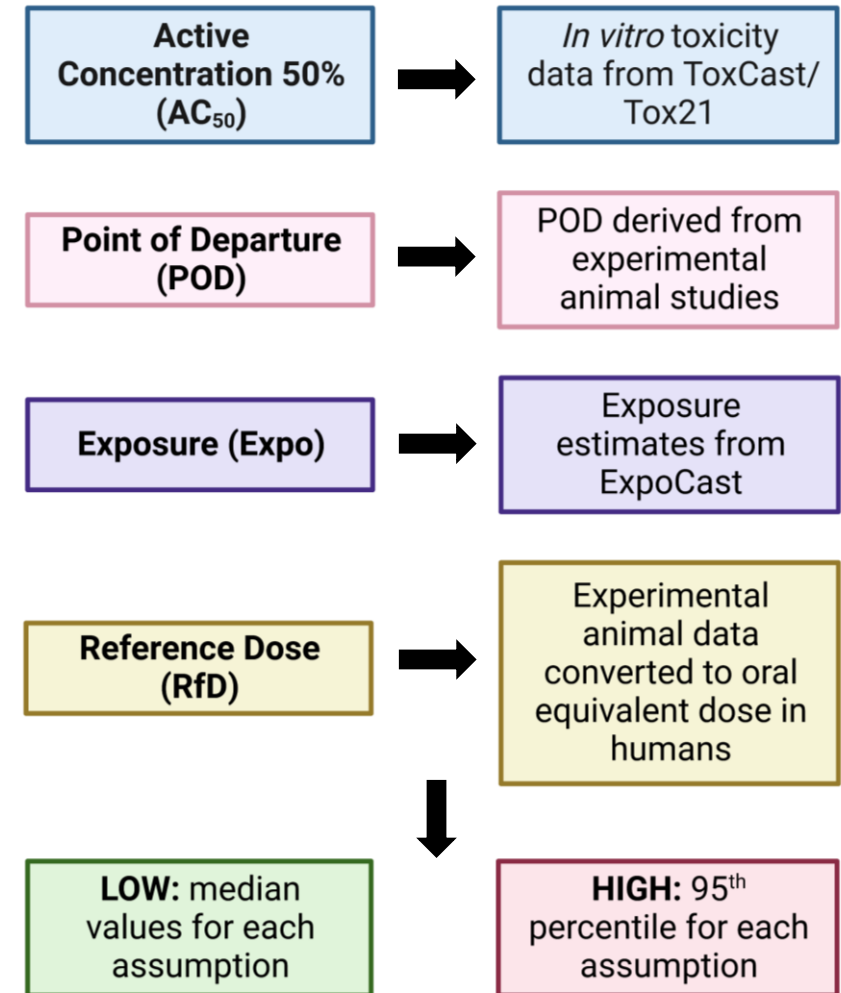
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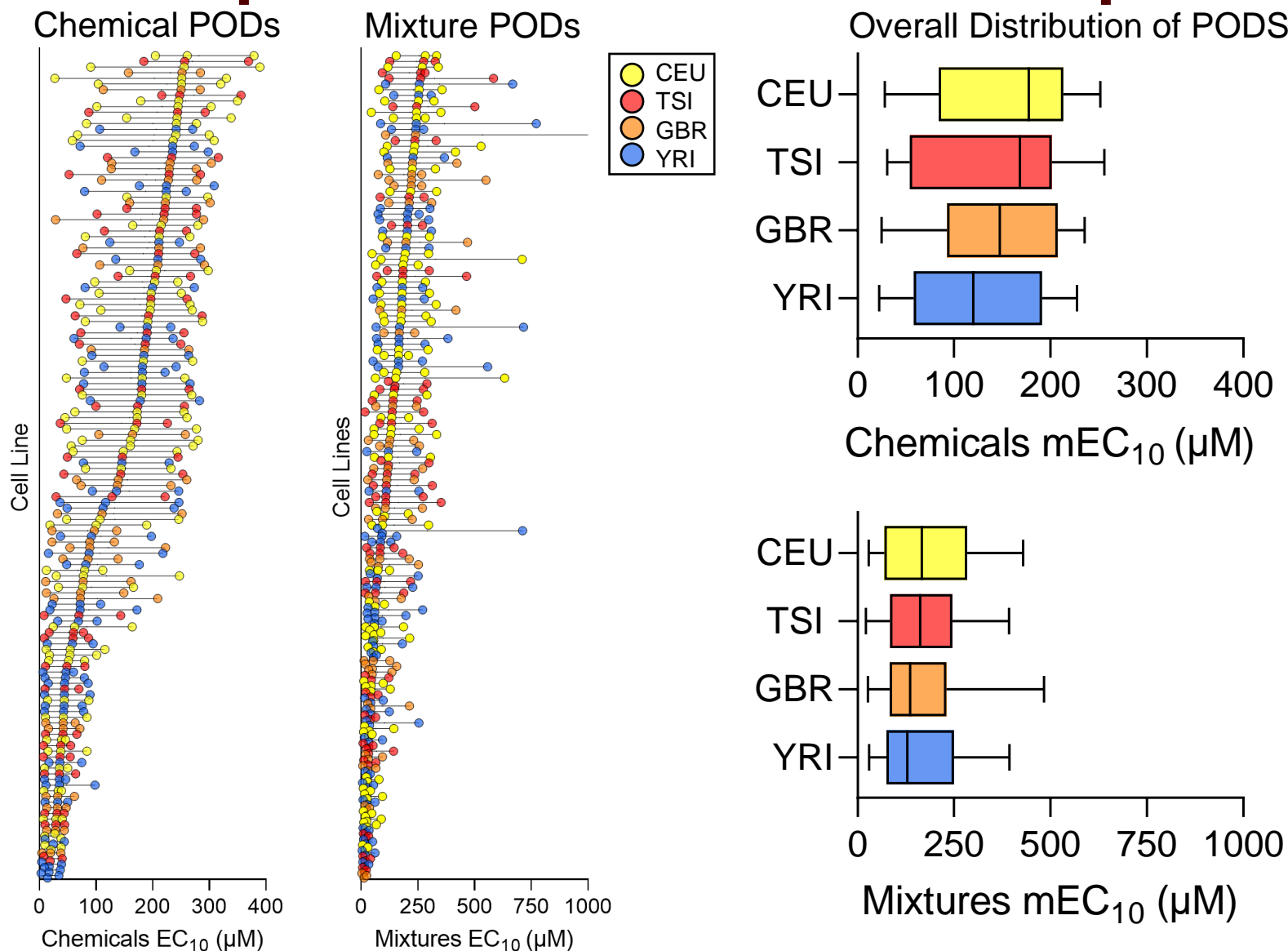
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36	LEAD NITRATE	10099-74-8	Metal
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41	COBALT CHLORIDE	7646-79-9	Metal
42	NICKEL CHLORIDE	7718-54-9	Metal

Preparation of 8 Defined Mixtures



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Donor-Specific Concentration-Response Profiling

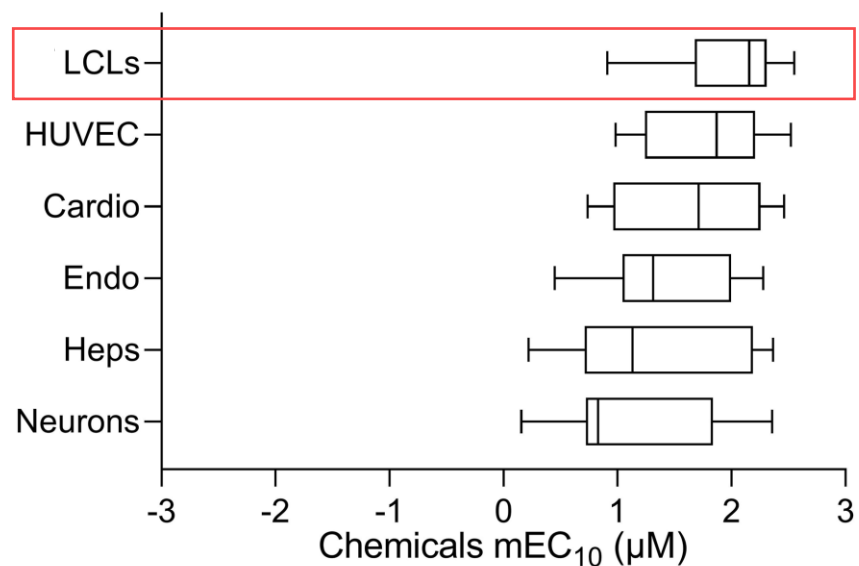


- Comparison of PODs across populations
- No significant differences across the 4 subpopulations
- YRI (subpopulation from African descent) lowest median PODs
- None of the subpopulations significantly more or less susceptible

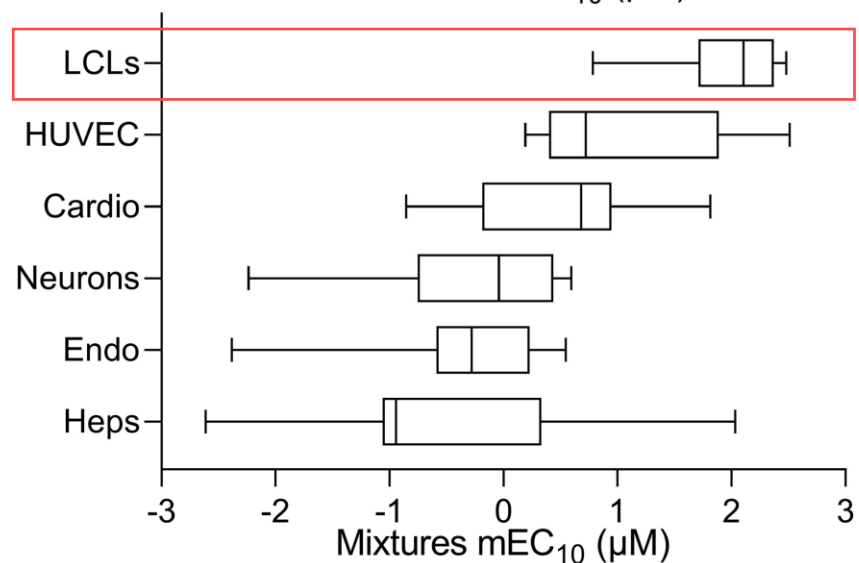


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Comparison Across Various *In Vitro* Models

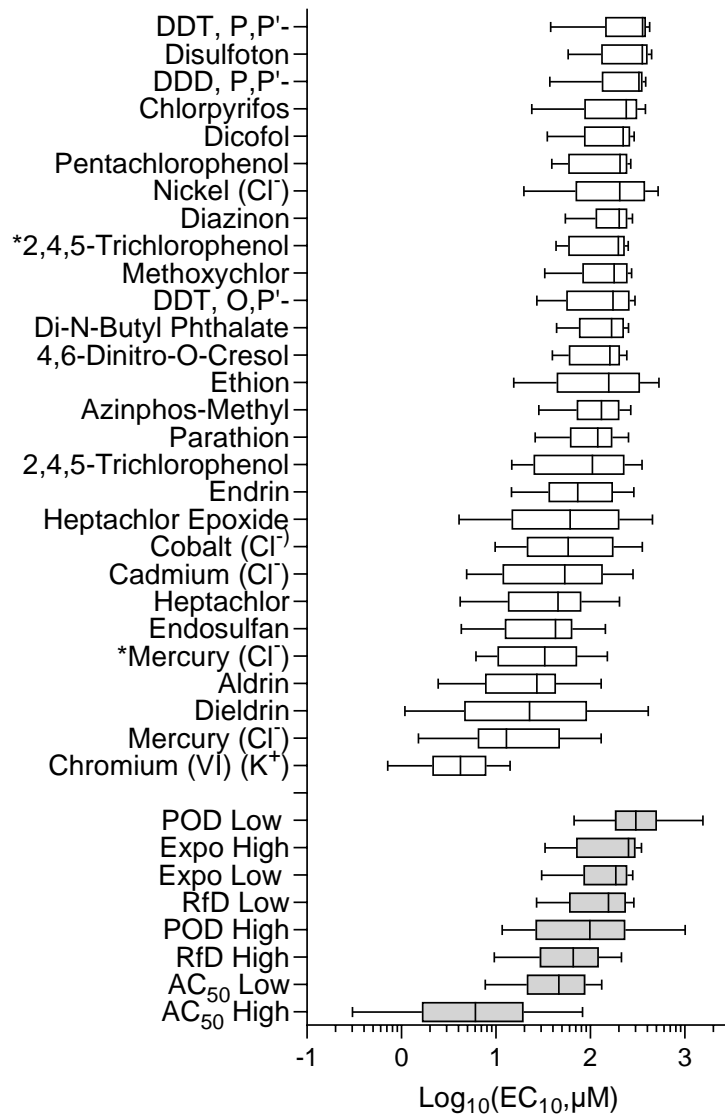


- Comparison of cytotoxic phenotypes for all models
- Chemicals and mixtures previously screened using 5 human *in vitro* models (PMID: 33395322)
- LCL within range of other *in vitro* models

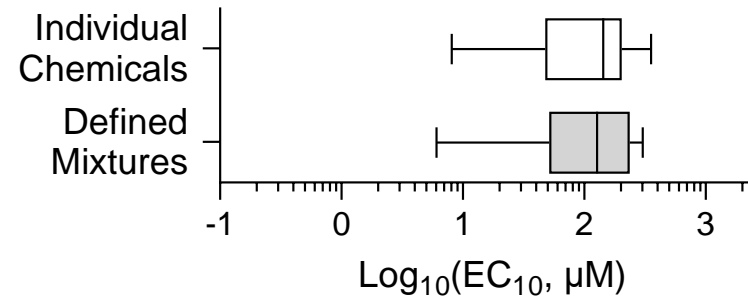


Chemical-Specific Concentration-Response Profiling

Distribution of PODs



Overall Distribution of PODs



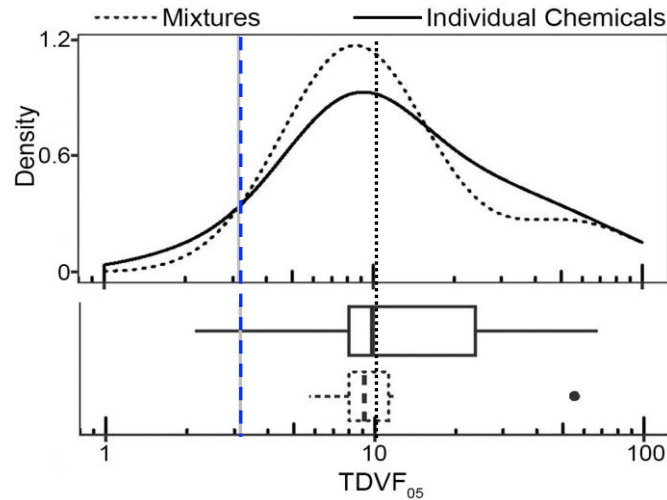
- 28 chemicals exhibited cytotoxic effects
- 17/28 chemicals were pesticides
- Heavy metals had the lowest EC₁₀
- AC₅₀ high had the lowest EC₁₀ for mixtures and the largest variability across all cells
- Median PODs for chemicals and mixtures were similar



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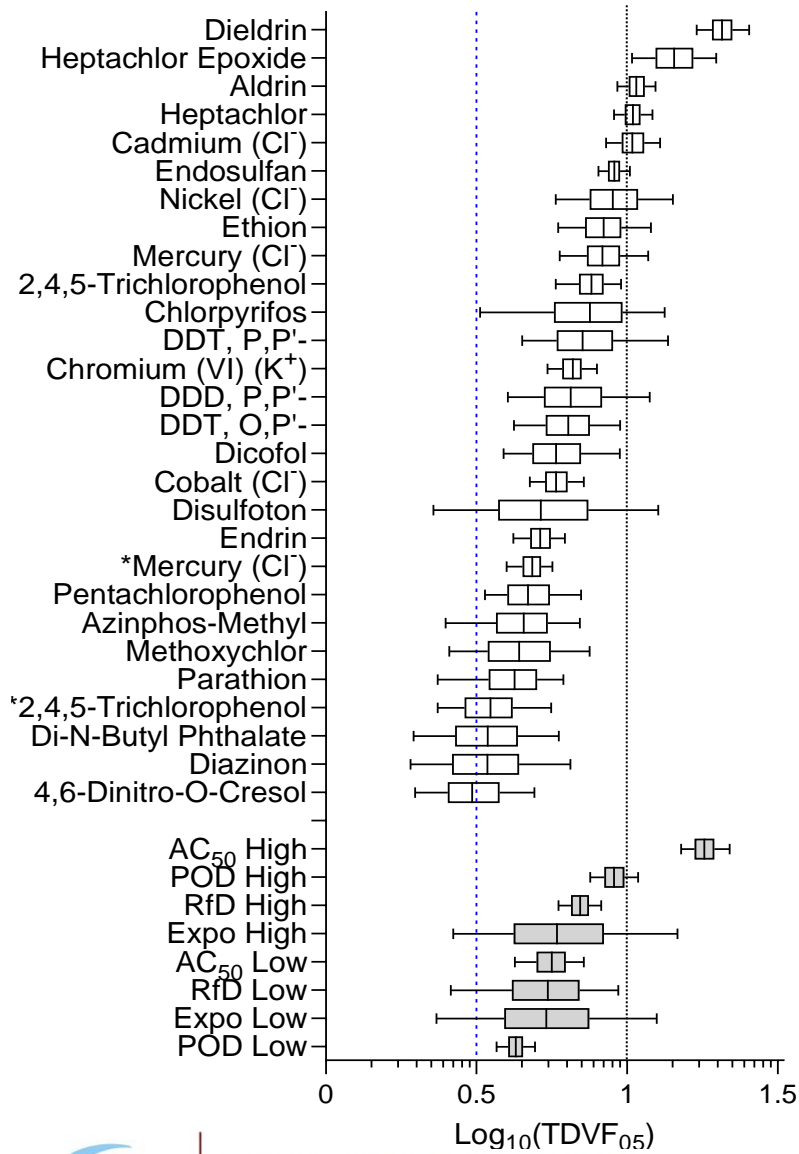
Quantifying Inter-Individual Variability

$$TDVF_{05} = \frac{EC_{10, median}}{EC_{10, 5th\ percentile}}$$



Blue Dashed line- default TDVF of 10^{1/2}
Black Dotted line- default total variability factor of 10

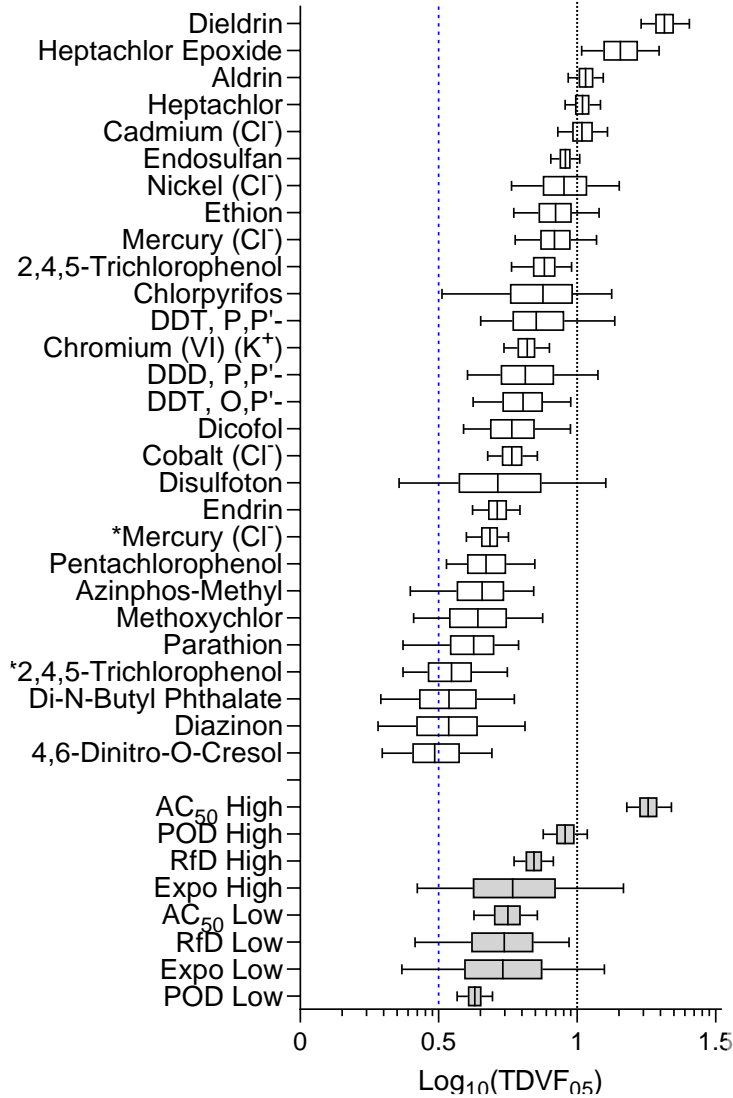
Distribution of TDVF₀₅



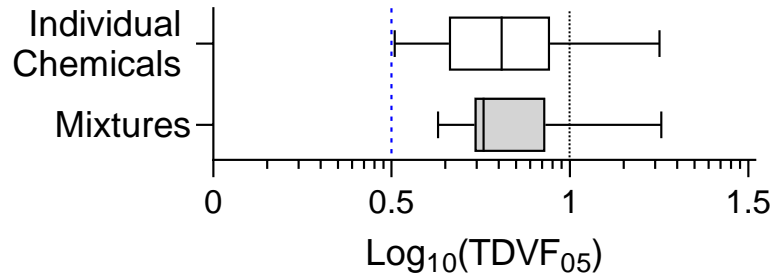
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Quantifying Inter-Individual Variability

Distribution of TDVF₀₅



Overall Distribution of TDVF₀₅



Blue Dashed line- default TDVF of $10^{1/2}$
Black Dotted line- default total variability factor of 10

- Pesticides had largest variability across all cells
- AC₅₀ had the largest variability across all cells
- Median TDVF₀₅ for chemicals and mixtures were similar
- Half a log higher than default for TDVF



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What are the potential molecular drivers of variability?

GWAS Work Flow

Run GWAS analysis for top 28 chemicals and mixtures



Identify top gene hits for each chemical/mixture



Genes functions



Gene expression in LCL



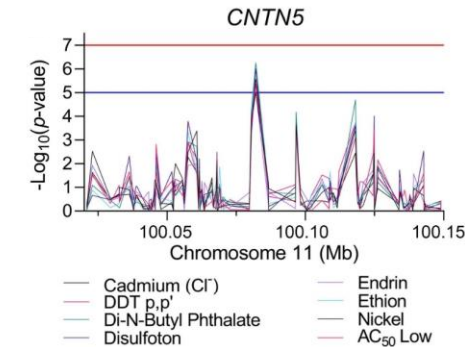
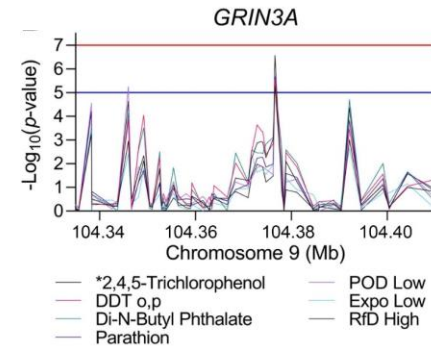
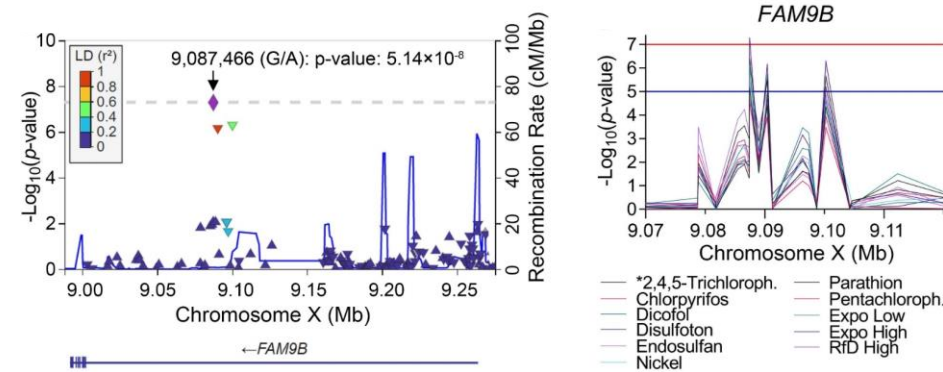
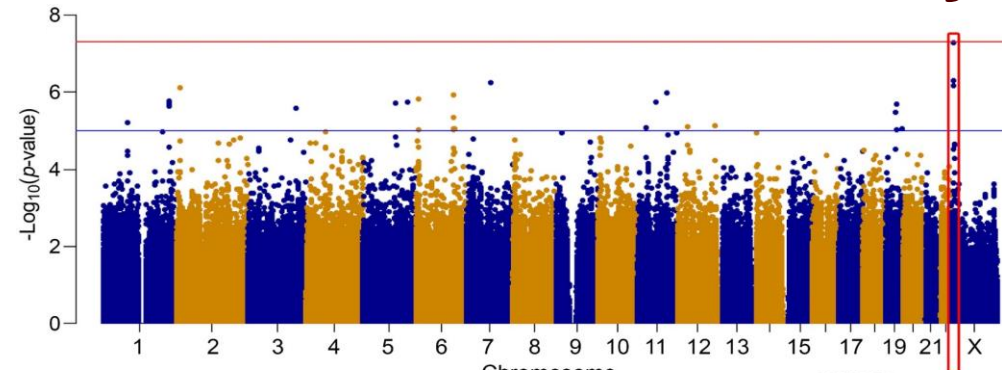
Correlation across chemicals



Gene hits in mixtures and chemicals



Research chemical exposure and gene of interest



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So what?

- Lymphoblasts were in range with human iPSC-derived models
- Quantified inter-individual variability for chemicals and mixtures
 - Population variability of mixtures does NOT exceed that of the most variable component
 - Similar TDVF₀₅ for chemicals and mixtures, BUT higher median than the default uncertainty factor of $10^{1/2}$
- Genome-wide associations among chemicals may be used to group constituents in a mixture

This model is a reasonable approach to quantify inter-individual variability and can be used to reduce uncertainties with complex exposure scenarios



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Study Limitations

Model lacks metabolic function

Reflecting acute high-dose treatments

Realistic routes of exposure

Limited chemical classes



Quantitative *in vitro* to *in vivo* extrapolation (QIVIVE)

In Vitro Data

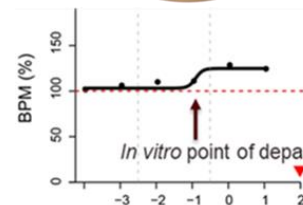
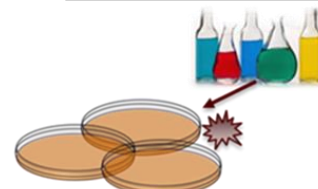
Exposure Data



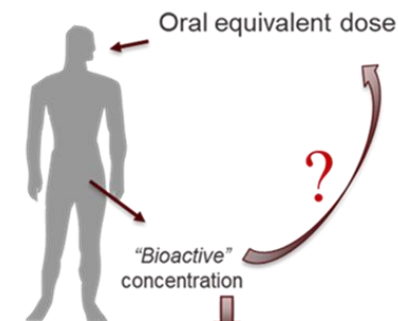
Oral Equivalent Dose in Humans

Chronic exposures

Routes of exposure



In Vitro-to-*In Vivo* Extrapolation



$$OEDs = PODs \times \frac{1 \text{ mg/kg/day}}{C_{ss}}$$



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Image adapted from Yu-Syuan Luo

Where do we go from here?

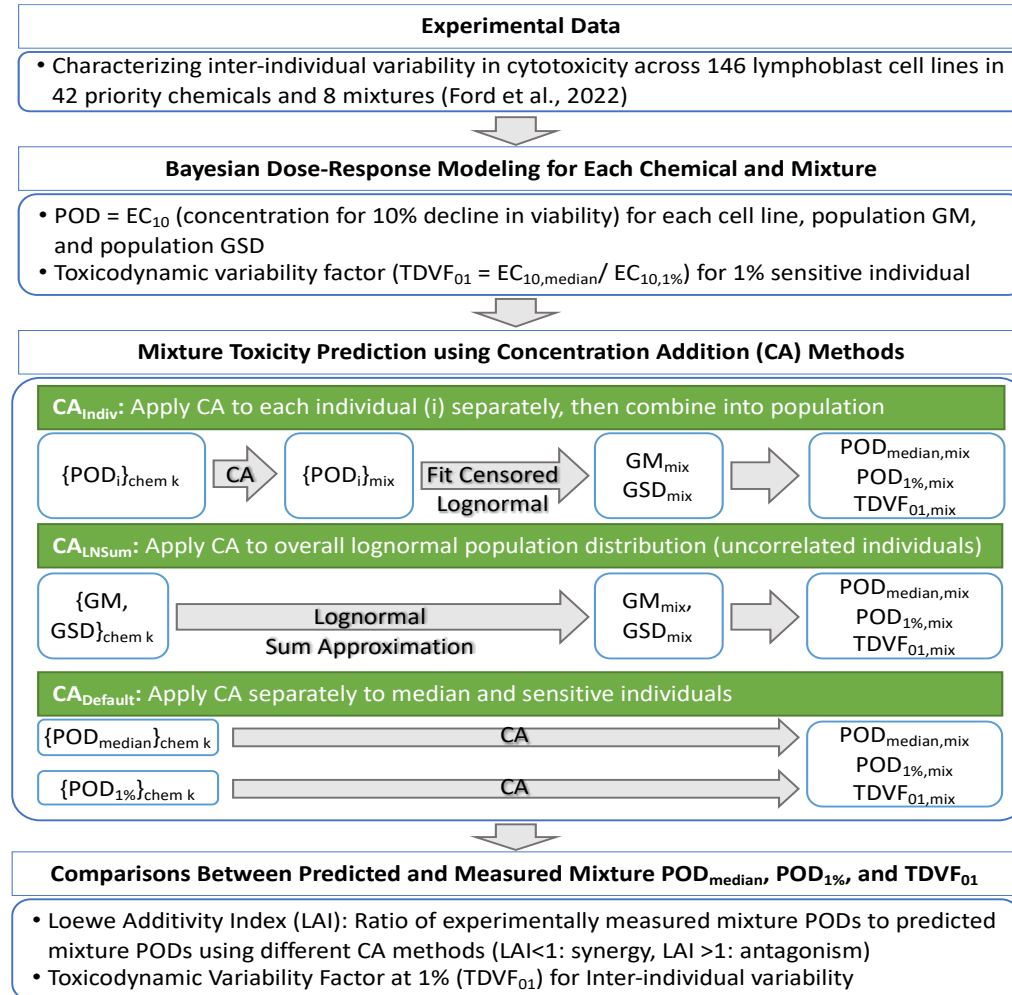
- Apply study design to evaluate toxicity of other defined and environmental mixtures
 - Screening realistic exposure scenarios using available biomonitoring data
 - Use environmental samples to conduct region-specific exposure assessments
- Complimentary work has been done with additivity models to reconstruct the variability using the chemical data (Jang et al., 2022, under review)



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Probabilistic Concentration Addition of Defined Mixture Exposures in a Population-Based Human *In Vitro* Model

Can we use concentration additivity approaches to predict inter-individual variability in responses to mixtures?



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Figures adapted from Jang *et al.*, 2022 (under review)

Probabilistic Concentration Addition of Defined Mixture Exposures in a Population-Based Human *In Vitro* Model

Can we use concentration additivity approaches to predict inter-individual variability in responses to mixtures?

Experimental Data

- Characterizing inter-individual variability in cytotoxicity across 146 lymphoblast cell lines in 42 priority chemicals and 8 mixtures (Ford et al., 2022)

Concentration additivity models may underestimate potency



Continue to use *in vitro* to test whole mixtures

OR

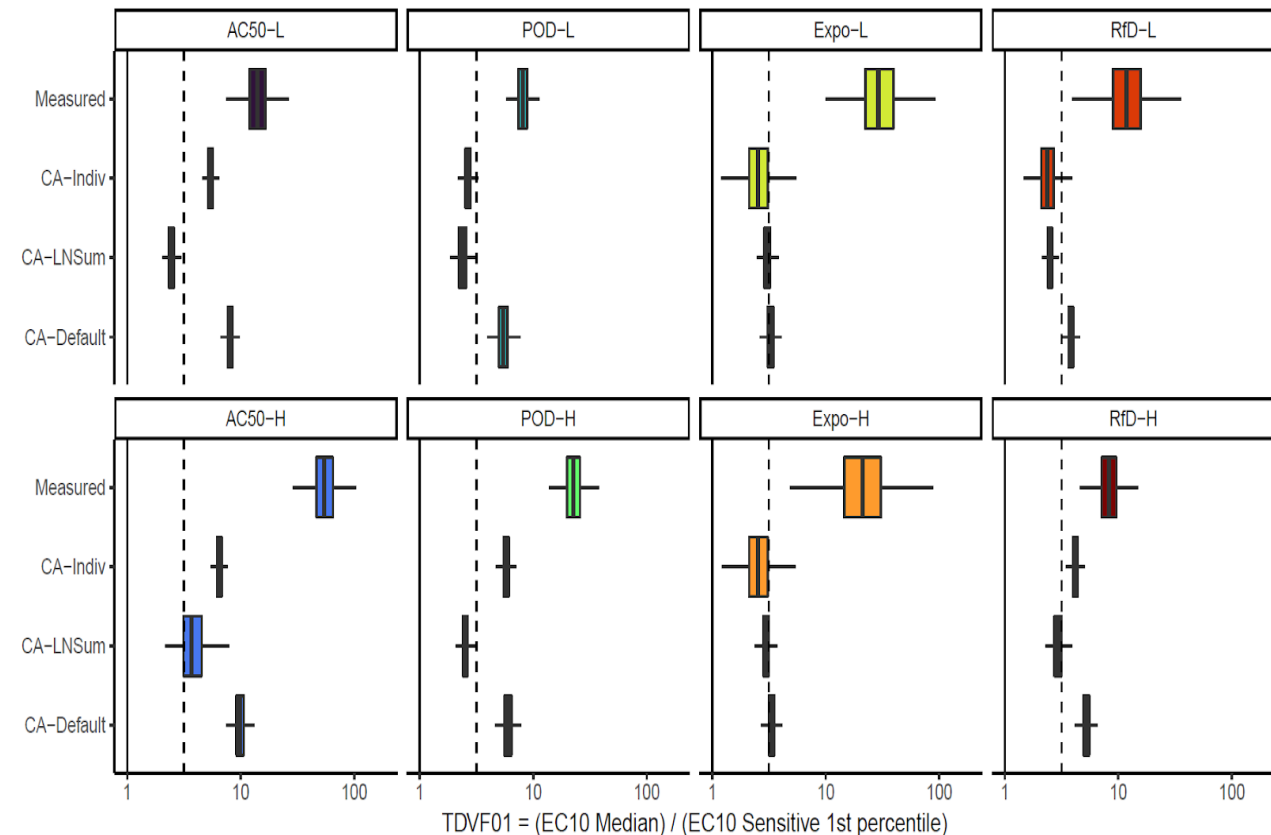
Implement more stringent risk indices
(ex: lower hazard index)



Ensure public health protection from combined exposures

Comparisons Between Predicted and Measured Mixture POD_{median} , $POD_{1\%}$, and $TDVF_{01}$

- Loewe Additivity Index (LAI): Ratio of experimentally measured mixture POD_{median} to predicted mixture POD_{median} using different CA methods (LAI < 1: synergy, LAI > 1: antagonism)
- Toxicodynamic Variability Factor at 1% ($TDVF_{01}$) for Inter-individual variability



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Figures adapted from Jang et al., 2022 (under review)

Overall Conclusions

Do we need *in vitro* systems to assess population-variability in responses to mixtures?

- Demonstrates feasibility of using population-based *in vitro* model that can be used in mixtures risk assessment
- Understand differences in inter-individual variability in responses to chemicals and mixtures
- Provides chemical and mixture-specific variability estimates that can be used to replace default assumptions
- Various concentration addition (CA) approaches demonstrate inter-individual variability, but tend to underestimate both the *in vitro* experimental POD and TDVF values
- Results from CA predictions supports continuation of *in vitro* toxicity testing for mixtures



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Thank you!

Questions?