

A New Approach Method for Characterizing Inter-Species Toxicodynamic Variability

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Presentation Outline

- Introduction:
 - Problem I: Too many chemicals (too little toxicity data)
 - Problem II: Toxicodynamic variability
 - Motivation to shift from *in vivo* approaches to new approach methodologies (NAM) *in vitro*
- Objective: Demonstrate the utility of an *in vitro* model for hazard and toxicodynamic variability assessments
 - Characterization of both inter-individual and inter-species variability
- Significance, limitations, and future directions of this work

Problem I: Too Many Chemicals





¹Natural Resources Defense Council

Images from <u>https://www.niehs.nih.gov/health/topics/agents/index.cfm</u> and <u>https://now.tufts.edu/articles/consequences-spraying-fire-retardants-wildfires</u>

90% of the 75,000 chemicals approved for use in the United States remain **inadequately tested** for potential toxicity¹

Traditional safety evaluation: in vivo

- Expensive, time-consuming, low-throughput
- Inter-species and inter-individual differences
- Unfeasible for characterizing safety of thousands of compounds

Problem I: Too Many Chemicals





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Alternative approach: in vitro

- Cheaper, faster, higher-throughput
- Biologically relevant can recapitulate characteristics of species/individuals
- Opportunity to evaluate variability
- Attractive alternative method for chemical safety evaluation

Problem II: Toxicodynamic Variability

Inter-species variability

- Testing in rodents, dogs, and/or nonhuman primates ≠ humans
- Often 1 isogenic animal strain •

Inter-individual variability

Genetic heterogeneity in the human population = sensitive subpopulations¹





RfD

&V

(UF = 3.16)

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NOAEL/LOAEL

UFs

Toxicity Testing in the 21st Century



TOXICITY TESTING IN THE 21ST CENTURY A VISION AND A STRATEGY



• Due to these challenges, there is strong motivation to decrease reliance on animal testing

- A paradigm shift toward the advancement and integration of alternative approaches for safety evaluation
- New approach methodologies (NAM) *in vitro*
- Increased efficiency in toxicity testing enables high-throughput, biology-driven evaluation

Assessing Inter-Species Variability In Vitro

- Inter-species toxicodynamic variability is poorly understood¹
- Risk assessment relies on allometric scaling² to extrapolate dose equivalencies between species (BW³)

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- Thought to account for metabolic differences in body size³
- Does not account for all observed toxicodynamic variability



¹Price *et al.*, 2008 (PMID: 18514247), ²Schneider *et al.*, 2004 (PMID: 15135212), ³Ungvari *et al.*, 2011 (PMID: 21059837)

Research Objective

Long-term goal:

- Improve *in vitro* approaches for chemical safety evaluation
- Better characterize both chemical hazard and variability
- Ultimately, advance hazard and risk assessment

Objective:

• To characterize inter-species and inter-individual variability in toxicodynamic response and investigate factors driving *in vitro* relationships

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Hypothesis:

An *in vitro* model using fibroblast cells from many species and individuals can be used to assess inter-species and inter-individual variability.

Approach and Study Design

Fibroblasts from 68 individuals of 54 diverse species



Concentration-response screening of 40 chemicals





Concentration (µM)

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TEXAS A&M

Chemical-Specific Differences in Variability



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TY

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Species Differences in Extent of <u>Inter-Individual</u> Variability



Default inter-individual UF = 3.16



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- General trend for inter-individual variability: human > monkey > dog > rat
 - Median human TDVF₀₅ ~2.5x greater than monkey, ~5x greater than rat
- 71% of chemicals had TDVF₀₅ higher than the default UF for inter-individual variation

Human Inter-Individual Variability



- Across studies, TDVF₀₅ estimates generally close to the default UF of 3.16 (1.87 5.01)
- Observable chemical-specific differences, with some chemicals exceeding the default UF

Inter-Species Variability



- Overall, median UF_{A,TD} estimate is close to the default UF
- Observable chemical-specific differences, with over half of chemicals exceeding the default UF
- Little to no relationship between body weight or lifespan and toxicodynamic sensitivity → toxicokinetics primarily drives allometric scaling *in vivo*
 - But overall allometric power may be chemical-specific



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EXA



Summary and Conclusions

 Both the inter-individual and inter-species components of variability contribute to inter-species toxicodynamic differences

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- On average, *in vitro*-derived estimates of inter-individual and inter-species variability are similar to the default UFs
- Chemical-specific differences → supports incorporation of CSAFs into risk assessment

This research demonstrates the utility of using an *in vitro* model to characterize both inter-individual and inter-species variability.



Significance, Limitations, and Future Directions

Significance of This Work

- This research demonstrates the utility of an *in vitro* model for hazard and toxicodynamic variability assessments
 - Both inter-individual and inter-species variability
- This approach is feasible for chemical screening in a manner unfeasible using traditional methods

This work contributes to the paradigm shift from traditional animal models of toxicity toward alternative approaches to inform hazard and risk assessment.

Limitations



- Need to optimize sample size for accurate hazard and variability assessments
- For cardiotoxicity, central estimate of population-wide potency in vitro is feasible with 5 donors, but ~20 donors are needed for accurate variability estimates

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- Clinical relevance of some *in vitro* phenotypes is unclear
 - Careful selection and biological understanding of phenotype of interest
- Differences in *in vitro* and *in vivo* dose metrics
 - In vitro-to-in vivo extrapolation to inform risk assessment^{2,3}
 - Need to understand chemical free fraction, clearance, and *in vivo* exposure levels

¹Blanchette *et al.*, 2021 (in press), ²Wetmore *et al.*, 2012 (PMID: 21948869), ³Blanchette *et al.*, 2019 (PMID: 30346629)

Future Directions



- Automated, high-throughput screening pipeline¹
- Mass production of cells²
- Characterized chemical library³
- Addressing metabolism in *in vitro* models⁴
 - Bioactivation vs. bioinactivation
 - Incorporating hepatic S9 fraction (individual- or species-specific)
 - Retrofitting *in vitro*-derived data with metabolism → more relevant effect predictions



Thank you!

Questions?

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