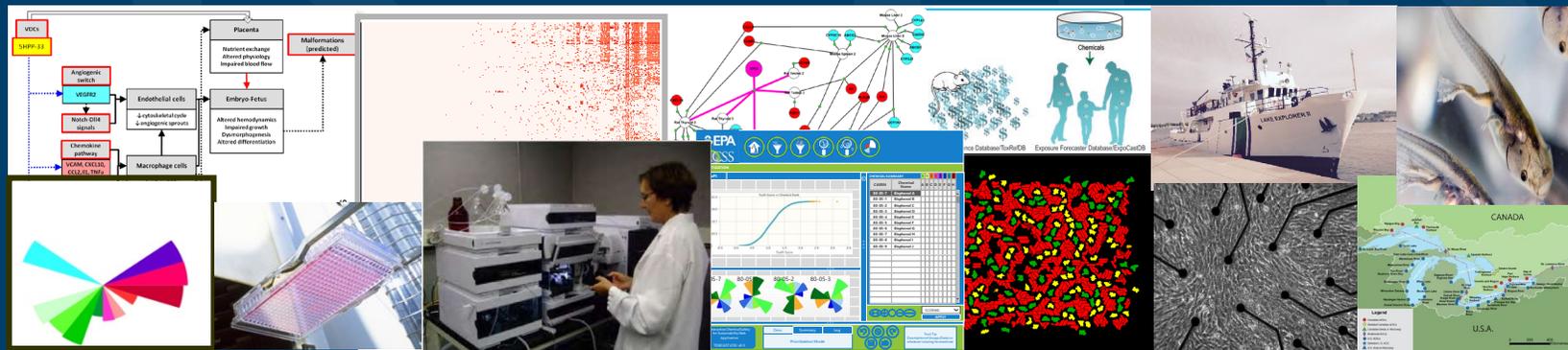


Update on Alternatives Research Activities at EPA



ICCVAM Public Forum

May 21, 2020

**Rusty Thomas
Director
Center for Computational Toxicology and Exposure**

The views expressed in this presentation are those of the presenter and do not necessarily reflect the views or policies of the U.S. EPA

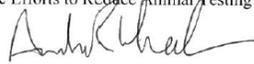
The Release of the EPA Memo Provided Clear Agency Goals for Reduction in Animal Testing



UNITED STATES ENVIRONMENTAL PROTECTION AGENCY
WASHINGTON, D.C. 20460
September 10, 2019
THE ADMINISTRATOR

MEMORANDUM

SUBJECT: Directive to Prioritize Efforts to Reduce Animal Testing

FROM: Andrew R. Wheeler 
Administrator

TO: Associate Deputy Administrator
General Counsel
Assistant Administrators
Inspector General
Chief Financial Officer
Chief of Staff
Associate Administrators
Regional Administrators

During my March 2019 all-hands address, I reiterated the U.S. Environmental Protection Agency's commitment to move away from animal testing. We are already making significant efforts to reduce, replace and refine our animal testing requirements under both statutory and strategic directives. For example, the *Toxic Substances Control Act*, amended June 22, 2016, by the Frank R. Lautenberg Chemical Safety for the 21st Century Act, requires the EPA to reduce reliance on animal testing. Also, Objective 3.3 of the *FY 2018-2022 U.S. EPA Strategic Plan* outlines a commitment to further reduce the reliance on animal testing within five years. More than 200,000 laboratory animals have been saved in recent years as a result of these collective efforts.

Scientific advancements exist today that allow us to better predict potential hazards for risk assessment purposes without the use of traditional methods that rely on animal testing. These new approach methods (NAMs), include any technologies, methodologies, approaches or combinations thereof that can be used to provide information on chemical hazard and potential human exposure that can avoid or significantly reduce the use of testing on animals. The benefits of NAMs are extensive, not only allowing us to decrease animals used while potentially evaluating more chemicals across a broader range of potential biological effects, but in a shorter timeframe with fewer resources while often achieving equal or greater biological predictivity than current animal models.

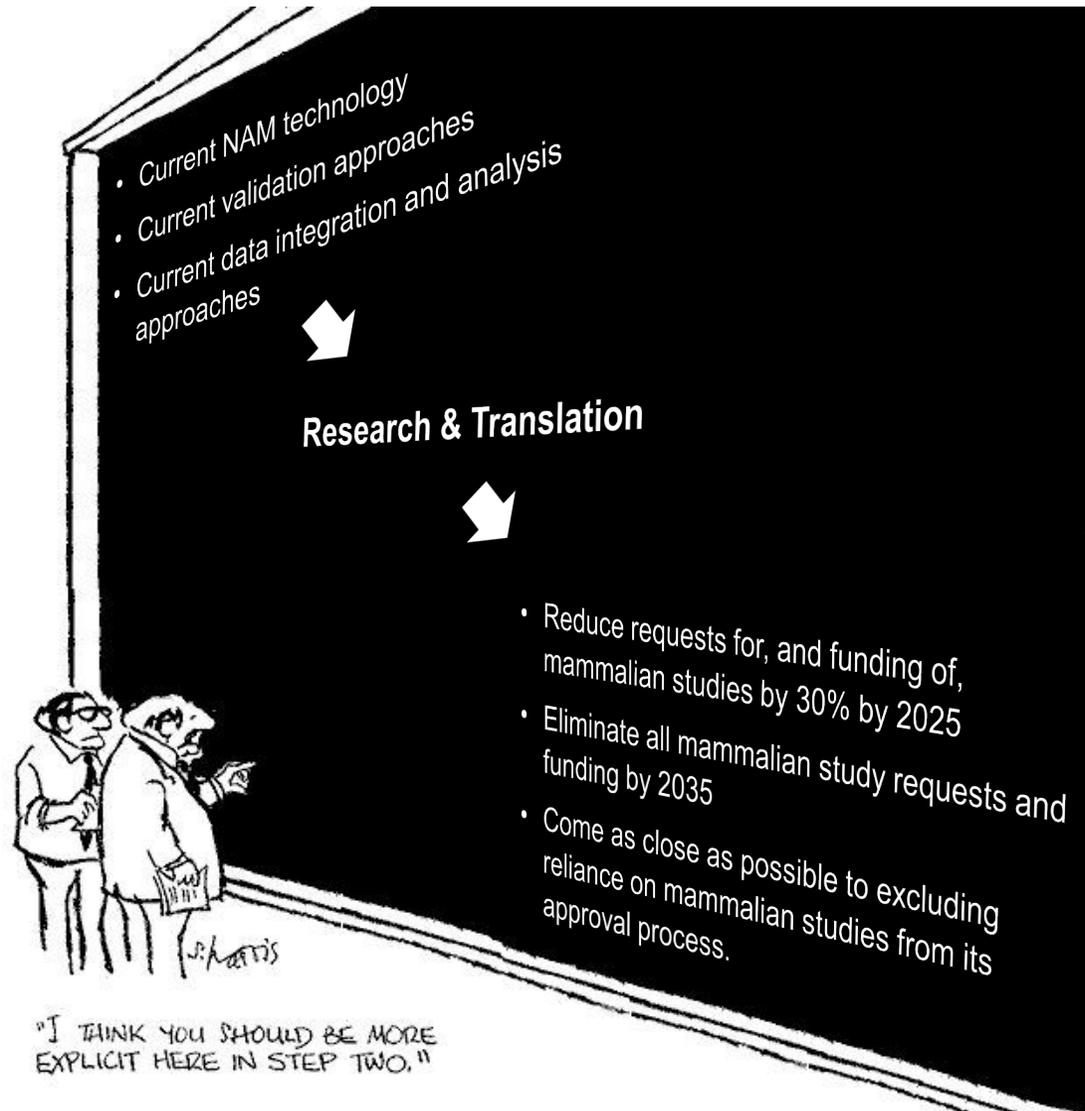
○ Goals:

- Reduce requests for, and funding of, mammalian studies by 30% by 2025
- Eliminate all mammalian study requests and funding by 2035
- Come as close as possible to excluding reliance on mammalian studies from its approval process (subject to applicable legal requirements).

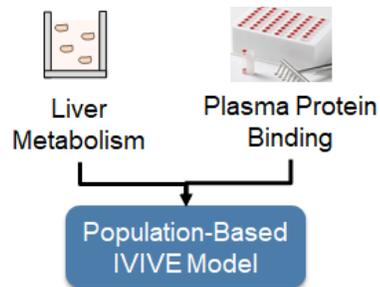
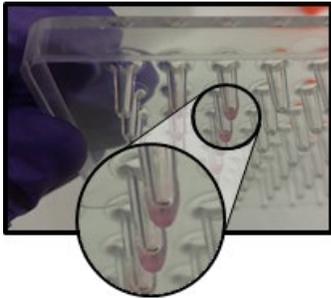
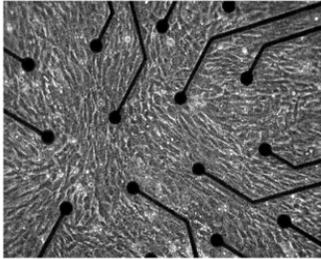
○ Objectives:

- Evaluate regulatory flexibility for accommodating the use of NAMs
- Develop baselines and metrics for assessing progress
- Validation to ensure NAMs are equivalent to or better than the animal tests
- Demonstration that NAMs are applicable for use in risk assessment and protective of human health and environment
- Engage and communicate with stakeholders

The Challenge...



ORD Research to Fill in “Step 2”



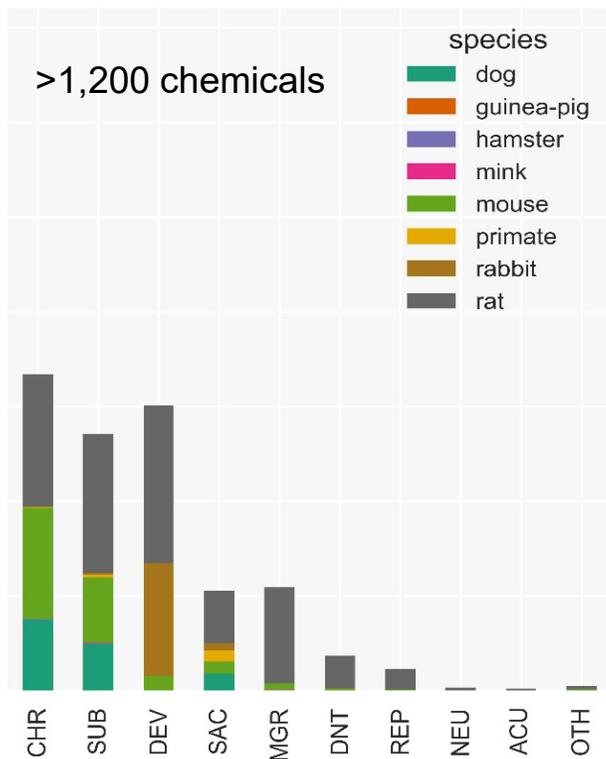
- Establish expectations on the variability of current toxicity studies
- Incorporate technological and data analysis advances to developing new alternatives
- Address limitations of *in vitro* test systems
- Build confidence through case studies

Mandate to Evaluate the Reliability and Relevance of Traditional Toxicity Testing Models

- Section 4(h) in the new TSCA legislation requires –
 - “...Administrator shall reduce and replace, to the extent practicable and scientifically justified...the use of vertebrate animals in the testing of chemical substances or mixtures...”
 - Alternative approaches need to provide “information of equivalent or better scientific quality and relevance...” than the traditional animal models

Evaluating Reproducibility of Traditional Toxicity Studies

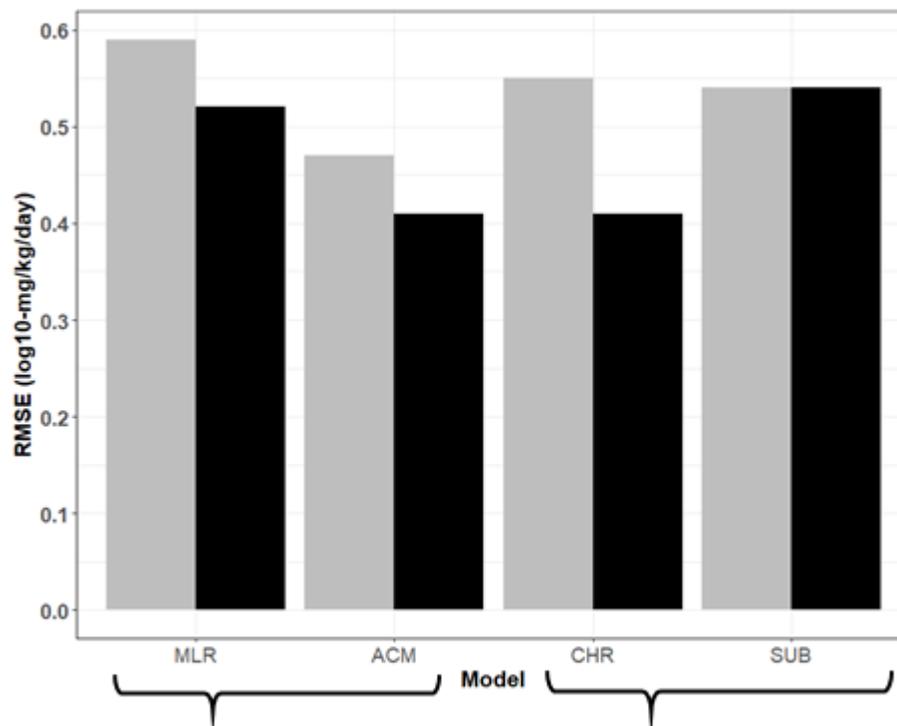
ToxRefDB Version 2.0



Study Type

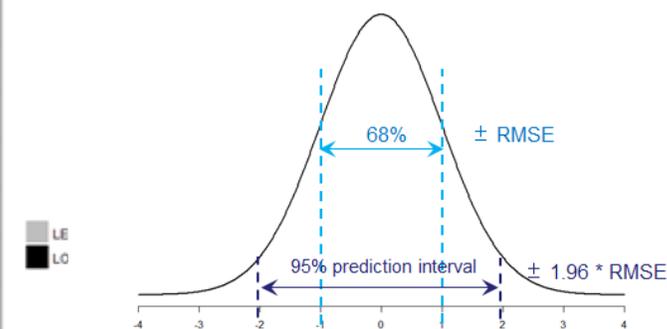
Watford et al., *Repro Toxicol*, 2019

Variability in Quantitative Effect Levels from *In Vivo* Repeat Dose Toxicity Studies



Two ways to statistically model the data across multiple study types

Variability within a specific study type



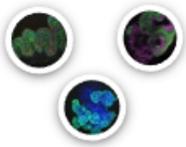
Using an RMSE=0.59, the minimum 95% PI of an LEL/LOAEL is:

1 mg/kg/day → 0.07 – 14 mg/kg/day.

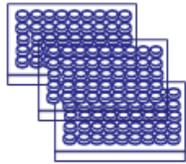
10 mg/kg/day → 0.7 – 143 mg/kg/day.

Comparing 'Cellular Pathology' With *In Vivo* Pathology Responses

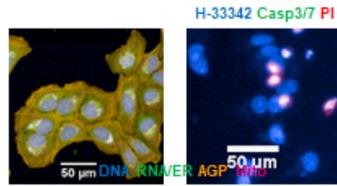
Multiple Cell Types



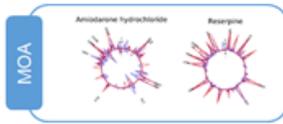
Concentration Response Screening



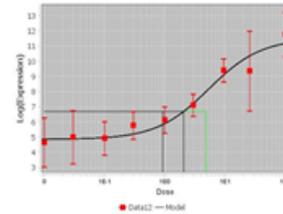
Multi-Parameter Cellular Phenotypic Profiling (HTPP)



Mode-of-Action Identification

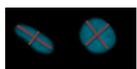
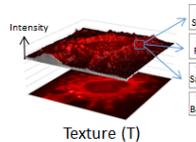
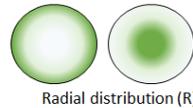
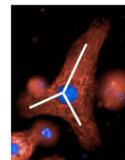
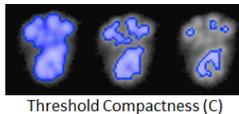
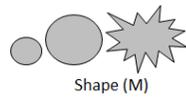


Concentration Response Modeling

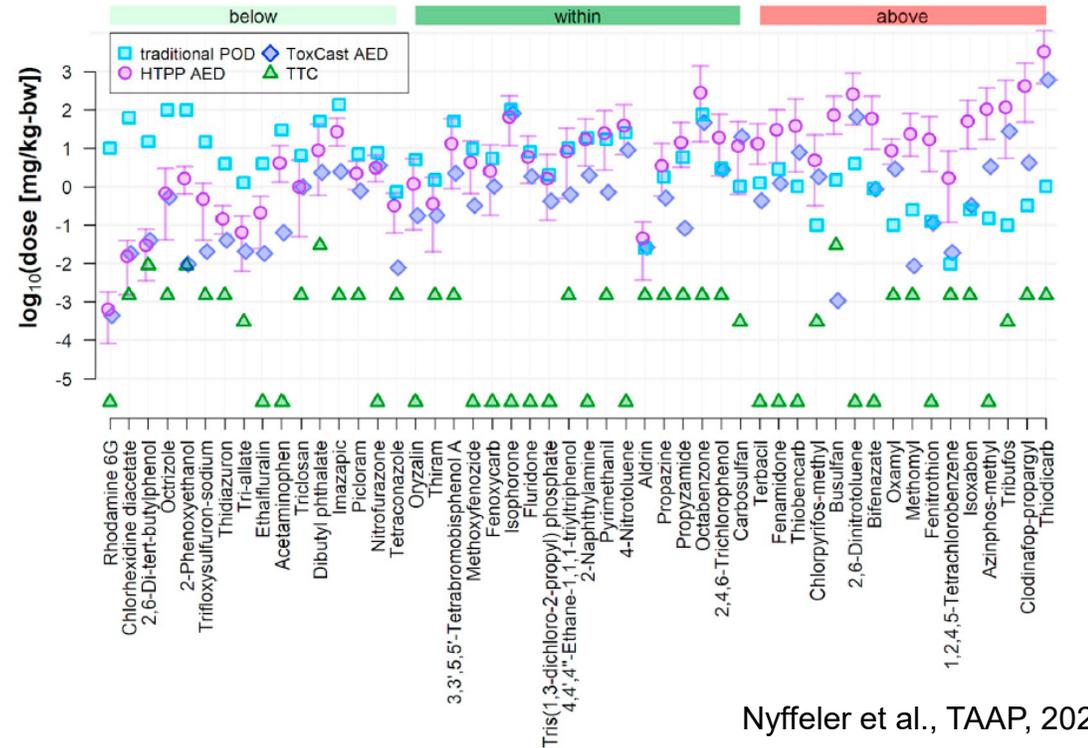
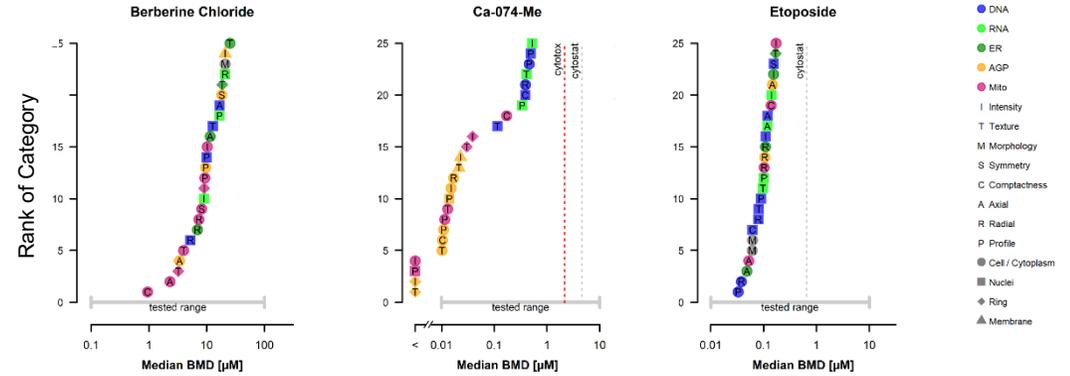


Cell Compartments

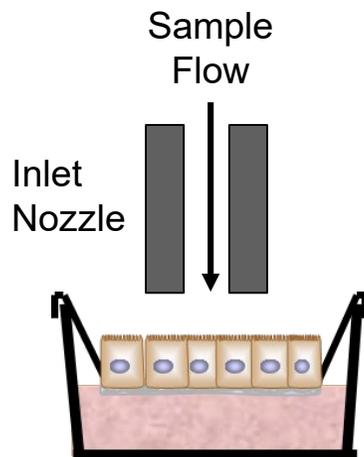
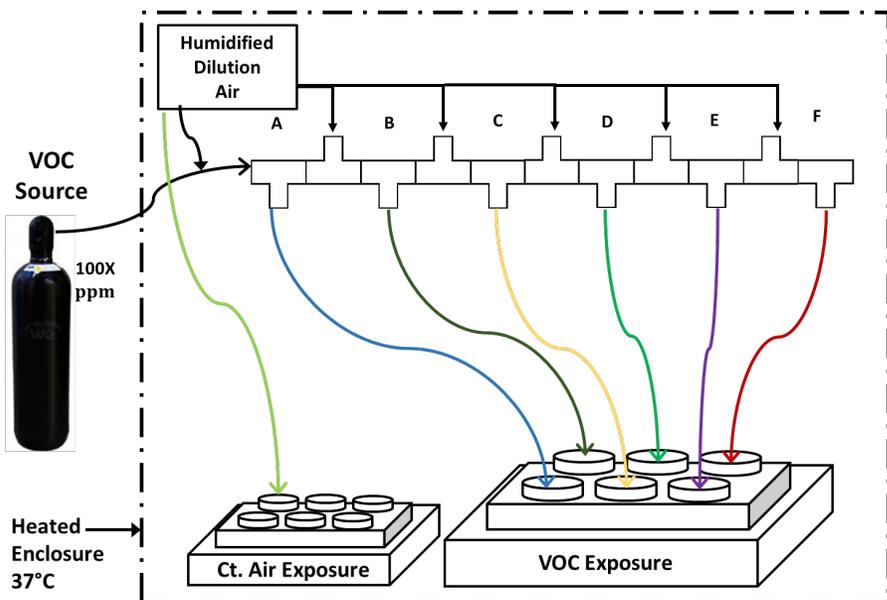
	NUCLEUS	RING	CYTOPLASM	MEMBRANE	CELL
DNA	S,C,A,R, P,I,T,M	--	--	--	S,C,A,R, P,M
RNA	S,C,A,R, P,I,T	--	--	--	S,C,A,R, P
ER	S,C,A,R, P,I,T	I,T	I,T	I	S,C,A,R, P
AGP	S,C,A,R, P,I,T	I,T	I,T	I,T	S,C,A,R, P
MITO	S,C,A,R, P,I,T	I,T	I,T	I	S,C,A,R, P



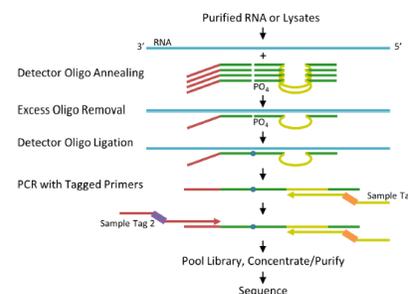
Images from PerkinElmer



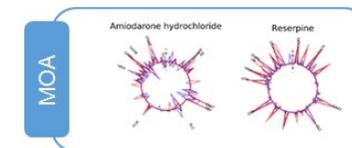
Adapting *In Vitro* Assays to Test Volatile Chemicals



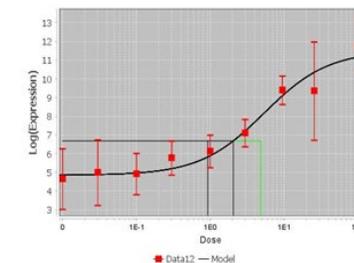
Whole Genome Transcriptomics (HTTr)



Mode-of-Action Identification



Concentration Response Modeling



Chemical Name	Gene Set Collection	BEAS-2B, BMC of most sensitive gene set (ppm)	HBEC, BMC of most sensitive gene set (ppm)	ACGIH TLV (ppm)
1-Bromopropane	MSigDB_C2	2.49302	9.93639	0.1
1-Bromopropane	MSigDB_H	2.97983	NA	
1-Bromopropane	Reactome	2.664425	NA	
Carbon Tetrachloride	MSigDB_C2	9.23691	NA	10
Carbon Tetrachloride	MSigDB_H	16.91345	NA	
Carbon Tetrachloride	Reactome	11.0172	NA	
Trichloroethylene	MSigDB_C2	48.9539	27.9907	50
Trichloroethylene	MSigDB_H	NA	36.4984	
Trichloroethylene	Reactome	69.6447	32.0725	
Dichloromethane	MSigDB_C2	136.124	269.865	100
Dichloromethane	MSigDB_H	231.7465	394.894	
Dichloromethane	Reactome	136.124	355	

A.Speen (CPHEA), M. Higuchi (CPHEA), and J. Harrill, Unpublished

Integrating *In Vitro* Assays to Predict Developmental Toxicity

Augmented DevTox prediction model uses Stemina + ToxCast assays



Profiling the ToxCast Library With a Pluripotent Human (H9) Stem Cell Line-Based Biomarker Assay for Developmental Toxicity

Todd J. Zurlinden, Katerine S. Sali, Nathaniel Rush, Parth Kothiyar, Richard S. Judson, Keith A. Houck, E. Sidney Hunter, Nancy C. Baker, Jessica A. Palmer, Russell S. Thomas, and Thomas B. Knudsen

National Center for Computational Toxicology (NCCT) and National Health and Environmental Effects Research Laboratory (NHEERL), Office of Research and Development (ORD), U.S. Environmental Protection Agency (USEPA), Research Triangle Park, North Carolina 27711; Leidos, Research Triangle Park, North Carolina 27711; and Stemina Biomarker Discovery, Inc, Madison, Wisconsin 53719

ABSTRACT

The Stemina devTOX quickPredict platform is a human pluripotent stem cell-based assay that predicts the developmental toxicity potential based on changes in cellular metabolism following chemical exposure [Palmer, J. A., Smith, A. M., Egnash, L. A., Conard, K. R., West, P. R., Burnier, R. E., Donlay, F. L. R., and Kitchener, P. R. (2013). Establishment and assessment of a new human embryonic stem cell-based biomarker assay for developmental toxicity screening. *Birth Defects Res. B Dev. Reprod. Toxicol.* 98, 343-361]. Using this assay, we screened 1065 ToxCast phase I and II chemicals in single-concentration or concentration-response for the targeted biomarker (ratio of ornithine to cystine secreted or consumed from the media). The dataset from the Stemina (STM) assay is annotated in the ToxCast portfolio as STM. Major findings from the analysis of ToxCast STM dataset include (1) 19% of 1065 chemicals yielded a prediction of developmental toxicity, (2) assay performance reached 79%-82% accuracy with high specificity (> 84%) but modest sensitivity (< 67%) when compared with *in vivo* animal models of human prenatal developmental toxicity, (3) sensitivity improved as more stringent chemical hits or evidence requirements were applied to the animal studies, and (4) statistical analysis of the most potent chemical hits on specific biochemical targets in ToxCast revealed positive and negative associations with the STM response, providing insights into the mechanistic underpinning of the targeted endpoint and its biological domain. The results of this study will be useful to improve our ability to predict *in vivo* developmental toxicants based on *in vitro* data and *in silico* models.

Key words: predictive toxicology; developmental toxicity; embryonic stem cells.

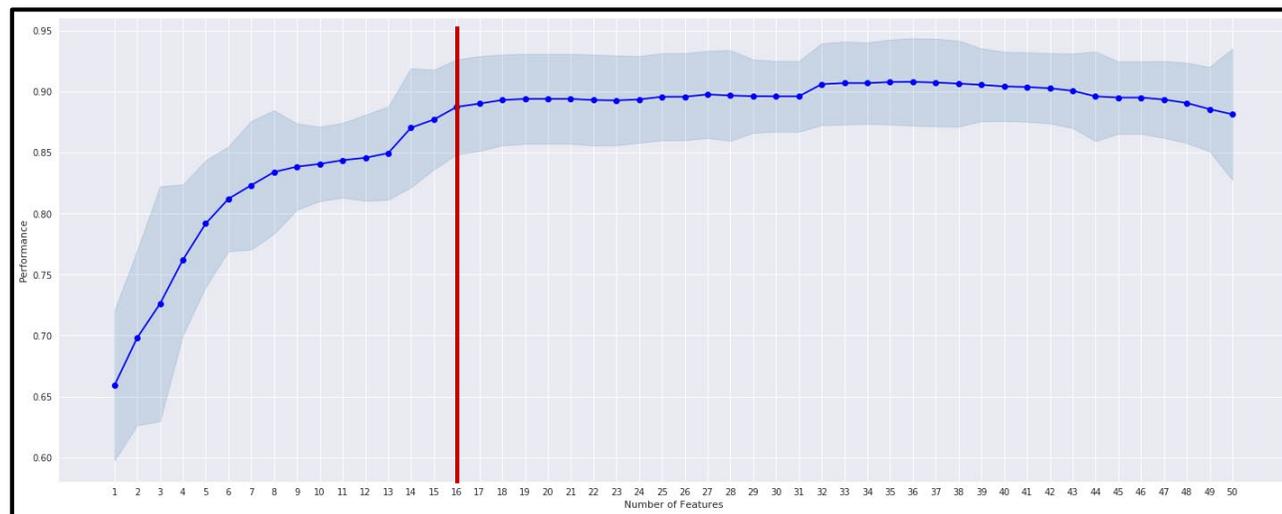
TOXICOLOGICAL SCIENCES, 174(2), 2020, 189-209

doi: 10.1093/toxsci/kfz014

Advance Access Publication Date: February 20, 2020

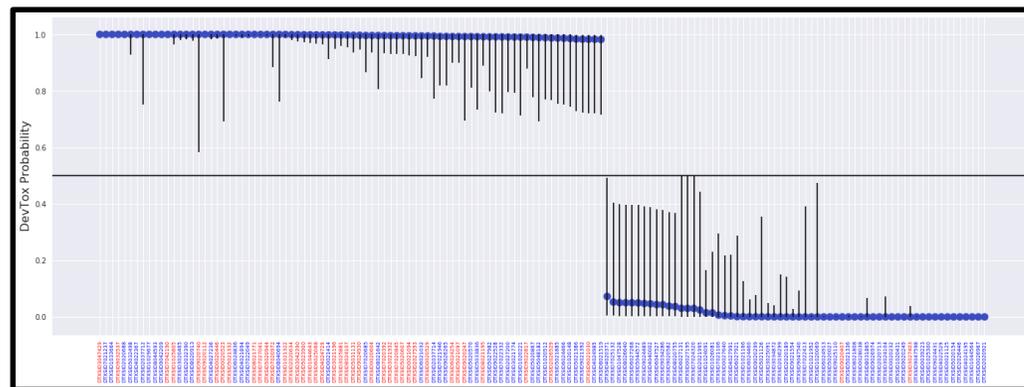
Research Article

Metric*	mean +/- sdev
ROC_AUC	0.91 +/- 0.03
Balanced Accuracy	0.82 +/- 0.04
NPV	0.80 +/- 0.05
PPV	0.90 +/- 0.08



*80/20 split (train/test) of the "Med_plus" data set (CLEAR rat OR rabbit, NO rat AND rabbit)

- Bayesian logistic regression to determine probabilistic model for DevTox
- Capability to tune model for increased sensitivity OR specificity

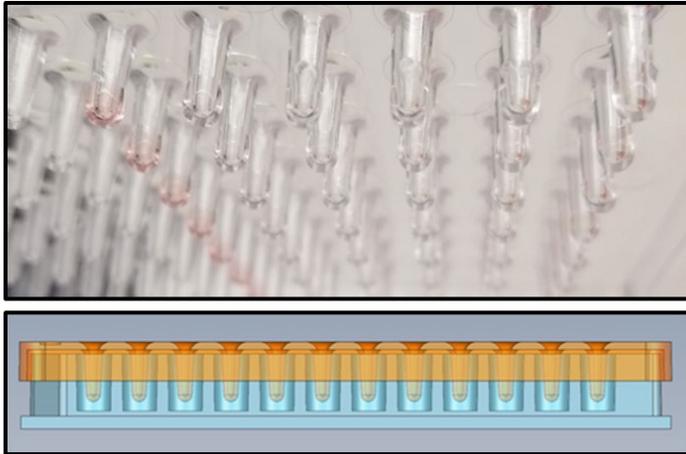


- Application of the "high specificity" model to ~580 chemicals on TSCA non-confidential inventory
- 144 chemicals predicted with confidence to fall into DevTox positive or negative domains

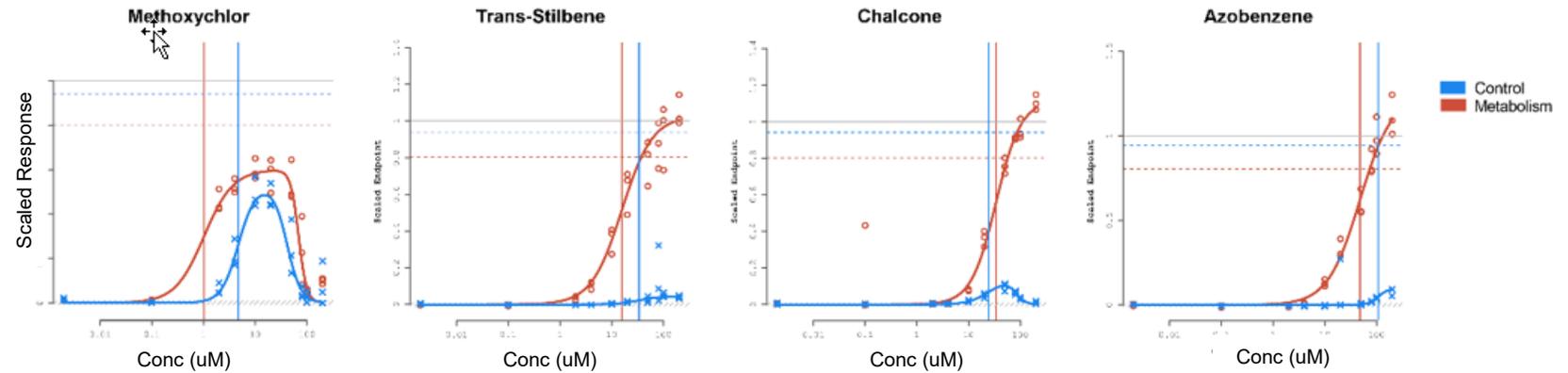
Zurlinden *et al.*, *Toxicol Sci.*, 2020
T. Zurlinden, T. Knudsen, Unpublished

Incorporating Xenobiotic Metabolism Into *In Vitro* Assays

AIME Method: S9 Fraction Immobilization in Alginate Microspheres on 96- or 384-well peg

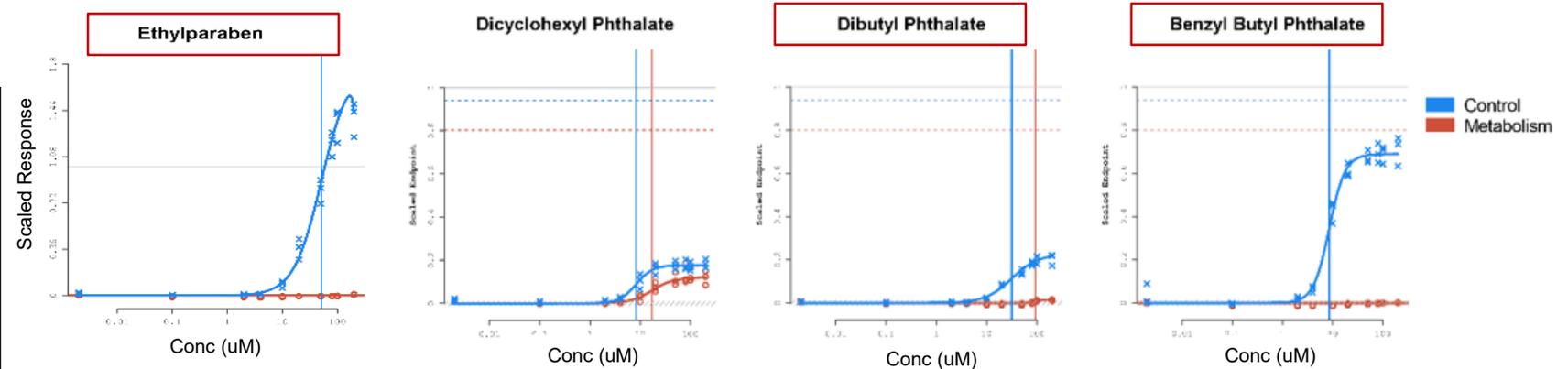


**Application to ER Transactivation Assay (ERTA)
Pilot Screening Results of Pinto et al., 2016 Library**



**Screening Window of VM7 (formerly BG1)
ER Transactivation Assay**

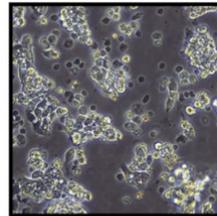
		Metabolism	
		Neg	Pos
NRS	Neg	0.91	0.89
	Pos	0.91	0.71
		Z'	



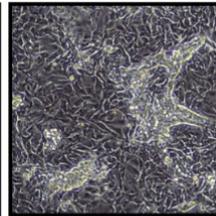
C. Deisenroth, In Review
Collaboration with Unilever

Developing Organotypic Culture Models to Identify Tissue/Organ Effects

Normal Human
Thyroid Gland



Harvest Follicle
Fragments



Attachment and
Outgrowth of Cells

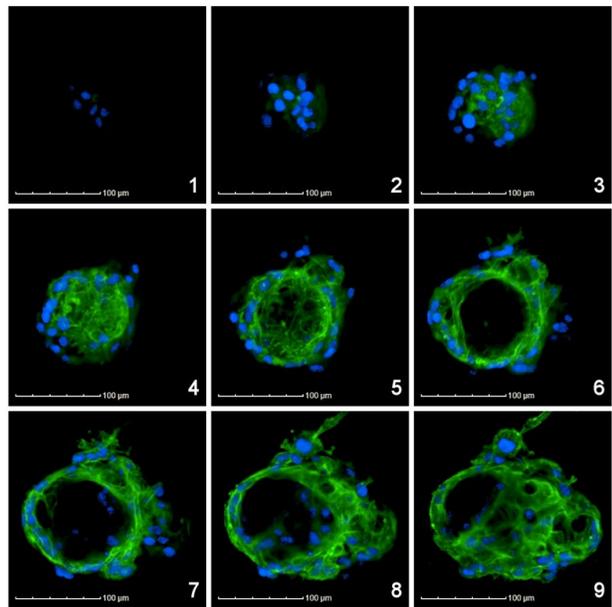
2D Cell Expansion



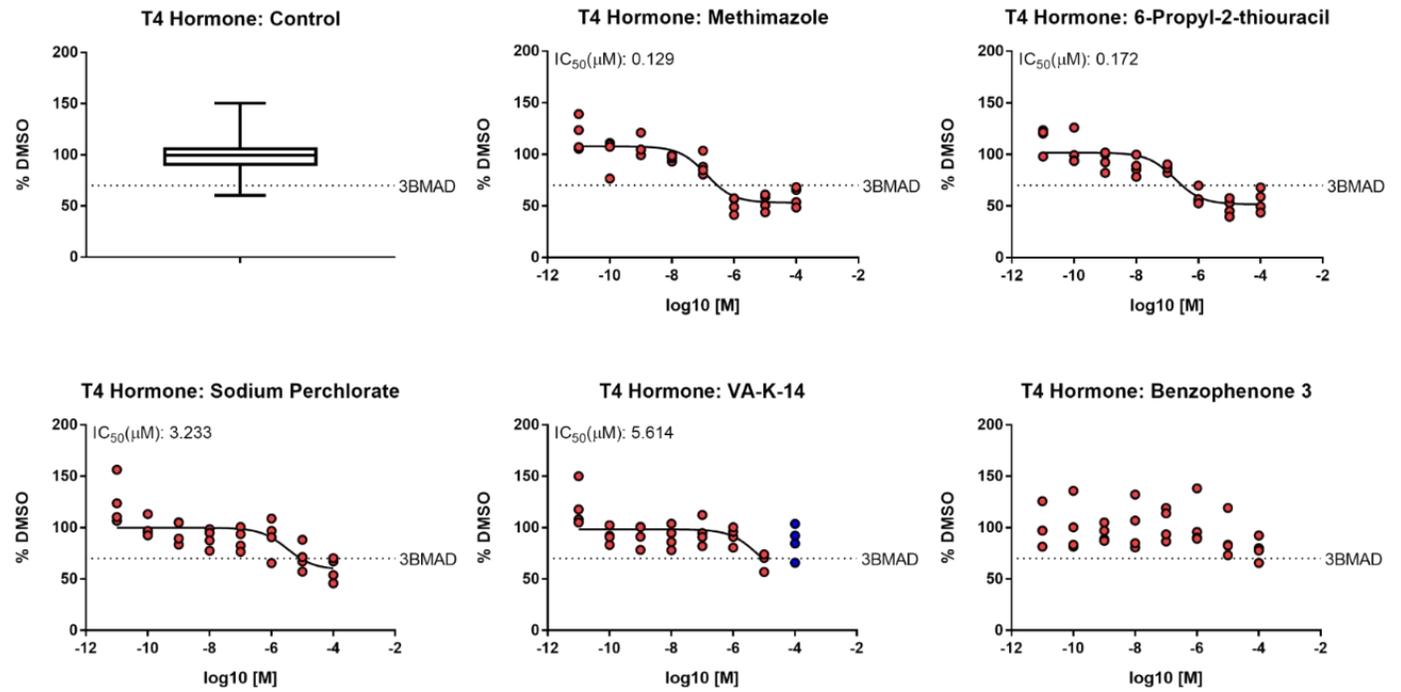
2D Monolayer
Culture



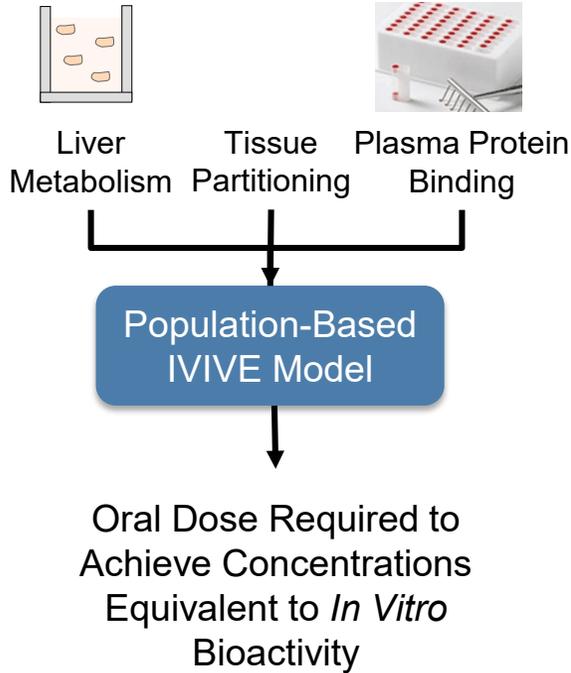
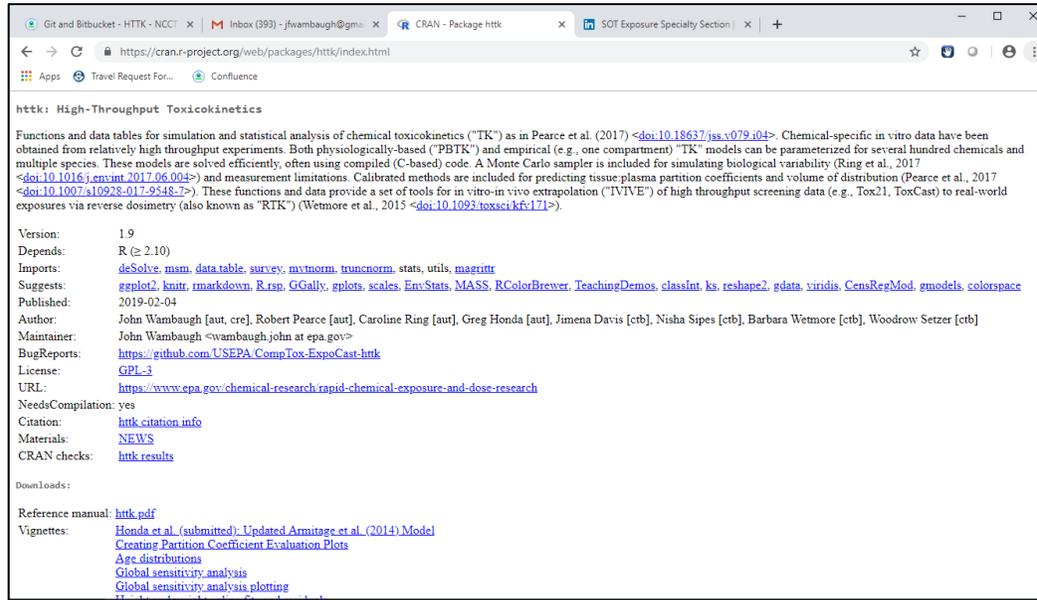
3D Sandwich
Culture



Blue, Hoechst 33342 /DNA
Green, Phalloidin/Actin



Putting Alternative Test Results in a Dose Context

```

httk: High-Throughput Toxicokinetics

Functions and data tables for simulation and statistical analysis of chemical toxicokinetics ("TK") as in Pearce et al. (2017) <doi:10.18637/jss.v079.i04>. Chemical-specific in vitro data have been obtained from relatively high throughput experiments. Both physiologically-based ("PBTK") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability (Ring et al., 2017 <doi:10.1016/j.envint.2017.06.004>) and measurement limitations. Calibrated methods are included for predicting tissue:plasma partition coefficients and volume of distribution (Pearce et al., 2017 <doi:10.1007/s10928-017-9548-7>). These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., Tox21, ToxCast) to real-world exposures via reverse dosimetry (also known as "RTK") (Wetmore et al., 2015 <doi:10.1093/toxsci/kfv171>).

Version: 1.9
Depends: R (>= 2.10)
Imports: deSolve, msm, data.table, survey, mvtnorm, truncnorm, stats, utils, magrittr
Suggests: ggplot2, knitr, rmarkdown, R.rsp, GGally, gplots, scales, EnvStats, MASS, RColorBrewer, TeachingDemos, classInt, ks, reshape2, gdata, viridis, CensRegMod, gmodels, colorspace
Published: 2019-02-04
Author: John Wambaugh [aut, cre], Robert Pearce [aut], Caroline Ring [aut], Greg Honda [aut], Jimena Davis [ctb], Nisha Sipes [ctb], Barbara Wetmore [ctb], Woodrow Setzer [ctb]
Maintainer: John Wambaugh <wambaugh.john@epa.gov>
BugReports: https://github.com/USEPA/CompTox-ExpoCast-httk
License: GPL-3
URL: https://www.epa.gov/chemical-research/rapid-chemical-exposure-and-dose-research
NeedsCompilation: yes
Citation: httk citation info
Materials: NEWS
CRAN checks: httk results

Downloads:

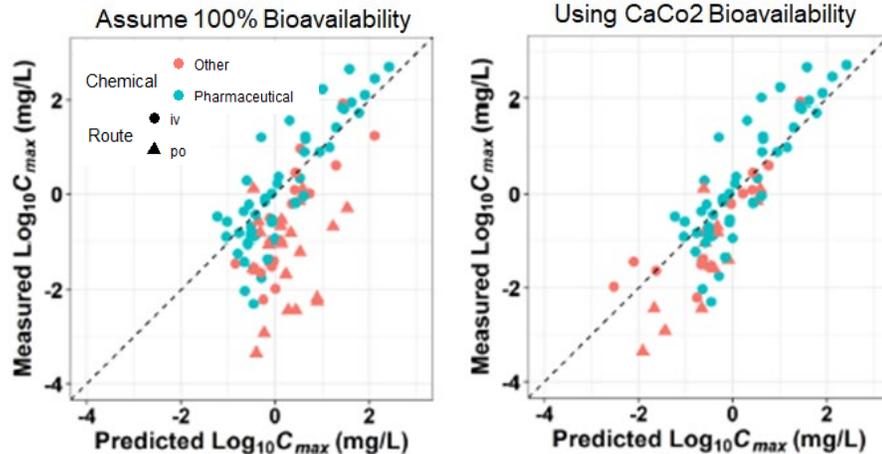
Reference manual: httk.pdf
Vignettes: Honda et al. (submitted), Updated Armitage et al. (2014) Model, Creating Partition Coefficient Evaluation Plots, Age distributions, Global sensitivity analysis, Global sensitivity analysis plotting
  
```

R package "httk"

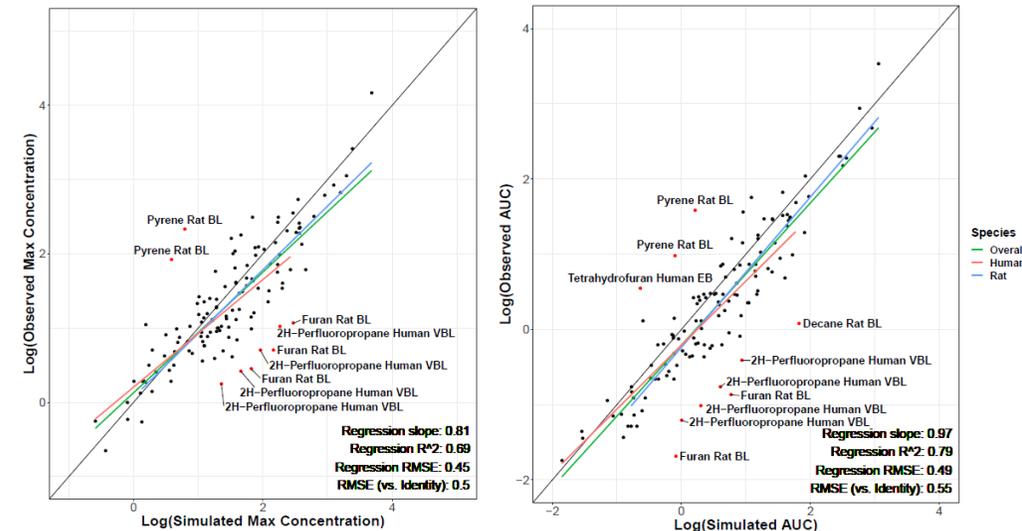
- Open source, transparent, and peer-reviewed tools and data for **high throughput toxicokinetics (httk)**
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific data for 987 chemicals
- Allows propagation of uncertainty

Rotroff *et al.*, *Tox Sci.*, 2010
 Wetmore *et al.*, *Tox Sci.*, 2012
 Wetmore *et al.*, *Tox Sci.*, 2015
 Wambaugh *et al.*, *Tox Sci.*, 2018
 Wambaugh *et al.*, *Tox Sci.*, 2019
 Linakis *et al.*, In Press.
 G. Honda and J. Wambaugh, Unpublished

Improving Oral PK Models



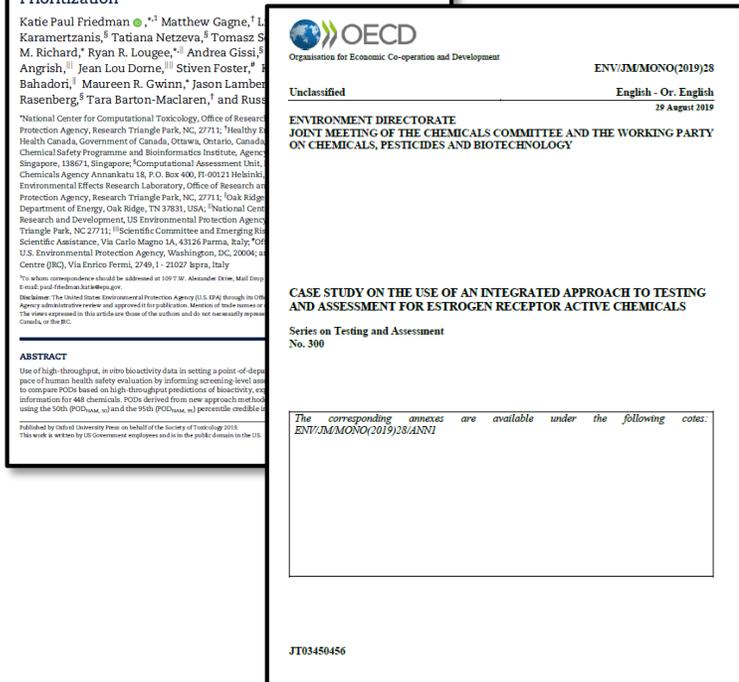
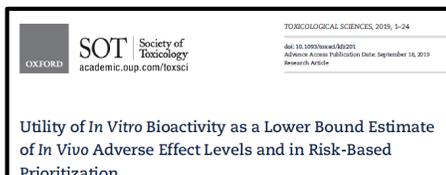
Incorporating Generic Inhalation PBPK Model



Case Studies to Build Confidence and Help Translate to Regulatory Application

Ongoing and New Case Studies

- OPP/ORD case study to use NAMs on selected pesticides with established MOAs
- OPP/ORD case study to develop a NAM for evaluating developmental neurotoxicity
- OCSP/ORD case study on integrating NAM to screen candidates for prioritization under TSCA
- OW/ORD case study on application of *in vitro* bioactivity and HTTK for screening-level assessments
- APCRA prospective case study on application of *in vitro* assays for hazard characterization
- APCRA case study on using NAMs to update chemical categories
- APCRA case study on computational approaches for rapid exposure estimates
- APCRA case study on modular integration of NAMs for identifying endocrine activity
- APCRA case study on using *in vitro* bioactivity to inform quantitative ecological hazard assessments
- APCRA case study on evaluating predictivity of HTTK methods



Recently completed case studies

Take Home Messages...

- ORD is working on a diverse portfolio of research activities to meet the Agency's animal testing reduction goals
- Characterizing the variability and relevance of existing models will aid in establishing expectations for the performance of alternative methods
- Continued development and refinement of new technologies and analysis approaches will help comprehensively evaluate potential toxicological effects
- Systematically addressing technical limitations such as a lack of metabolism, testing challenging chemicals, and identifying organ/tissue effects will enable important information gaps to be filled
- Partnering with regulators and national and international partners on case studies will increase confidence in alternatives and accelerate application for a range of decision contexts

Acknowledgements

Center for Computational Toxicology and Exposure (CCTE) Staff

Tox21 Colleagues:

NTP
FDA
NCATS

EPA Colleagues:

CEMM
CPHEA
CESER

Collaborative Partners:

Unilever
A*STAR
ECHA
EFSA
Health Canada



Research Triangle Park, NC



Cincinnati, OH



Duluth, MN



Washington, DC



Athens, GA



Gulf Breeze, FL