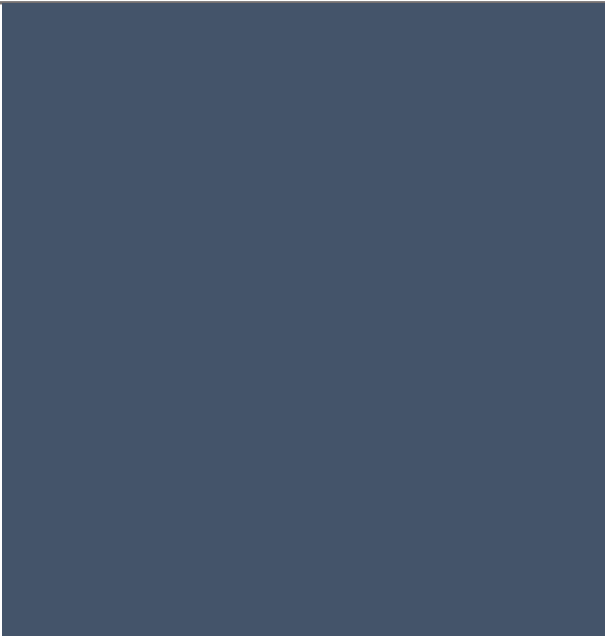


Note: this document may contain some elements that are not fully accessible to users with disabilities. If you need assistance accessing any information in this document, please contact ORD_Webmaster@epa.gov.



US EPA CSS-HERA Board of Scientific Counselors Chemical Safety Subcommittee Meeting

US EPA CSS-HERA BOSC Meeting – February 2-5, 2021

CSS Session 1B Slides



The work presented within represents US EPA Office of Research and Development research activities. Material includes both peer reviewed, published results and work-in-progress research. Please do not cite or quote slides.

Table of Contents

High-Throughput Exposure Models and the Systematic Empirical Evaluation of Models (SEEM) Framework (John Wambaugh).....	3
High-Throughput Toxicokinetic Models and <i>In Vitro-In Vivo</i> Extrapolation (IVIVE) (Barbara Wetmore).....	25
NAMs for Exposure: Non-Targeted Analysis (Jon Sobus).....	49

High Throughput Exposure Models and the Systematic Empirical Evaluation of Models (SEEM) Framework

John Wambaugh

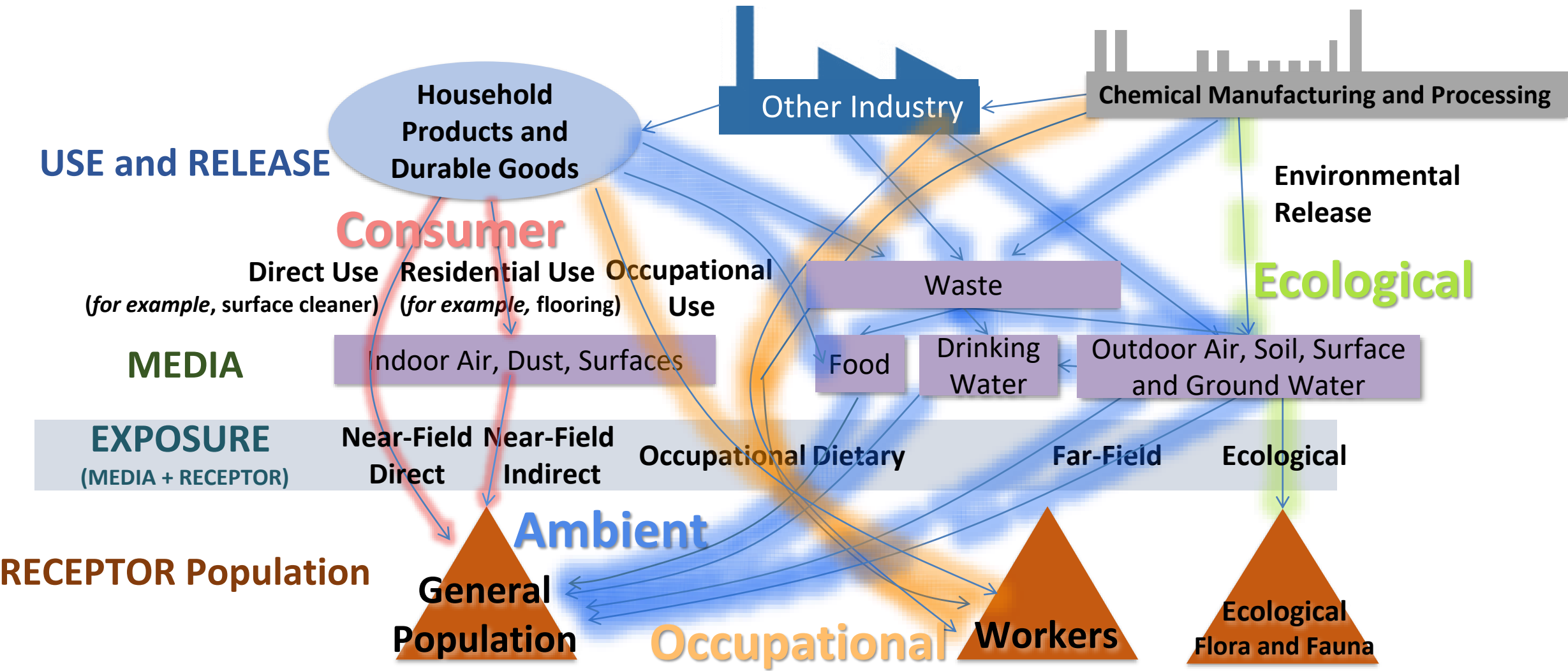
US EPA CSS-HERA Board of Scientific Counselors
Chemical Safety Subcommittee Meeting

Stakeholder Need as stated in research plan

Chemical exposure scenarios and pathways:

Chemical evaluations require information to estimate exposure via a variety of high-priority pathways, including scenario-specific data and models particular to consumer products and materials in the indoor environment, as well as occupational, ambient and ecological pathways.

Exposure Pathways



Properties of High-Throughput Exposure Models

- 1) Capable of handling **many chemicals with minimal descriptive information**
- 2) Cover **one or more relevant exposure routes**
- 3) **Allow for integration** with models for other pathways
- 4) Scientifically plausible
- 5) Allow for the assessment of interindividual and intraindividual variation in exposure
- 6) **Amenable to integration within statistical frameworks that quantify uncertainty**
- 7) **No more complicated than necessary**



Current Opinion in Toxicology

Available online 31 July 2019

In Press, Journal Pre-proof ?



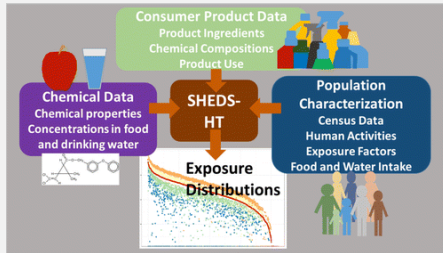
New Approach Methodologies for Exposure Science

John F. Wambaugh¹  , Jane C. Bare², Courtney C. Carignan³, Kathie L. Dionisio⁴, Robin E. Dodson^{5, 6}, Olivier Jolliet⁷, Xiaoyu Liu⁸, David E. Meyer², Seth R. Newton⁴, Katherine A. Phillips⁴, Paul S. Price⁴, Caroline L. Ring⁹, Hyeong-Moo Shin¹⁰, Jon R. Sobus⁴, Tamara Tal¹¹, Elin M. Ulrich⁴, Daniel A. Vallero⁴, Barbara A. Wetmore⁴, Kristin K. Isaacs⁴

Existing HT Models for Key Pathways

Consumer (Near-Field) Pathways

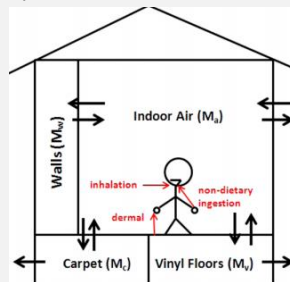
SHEDS-HT (Isaacs et al., 2014)



RAIDAR-ICE (Li et al., 2018)

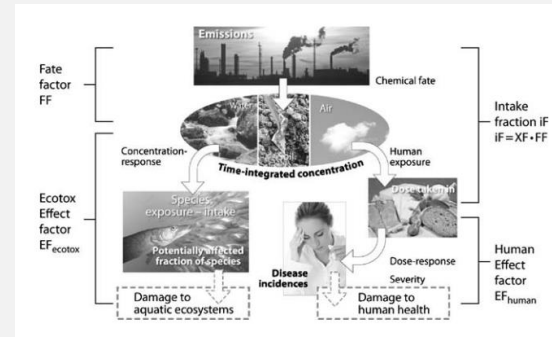


FINE (Shin et al., 2015)

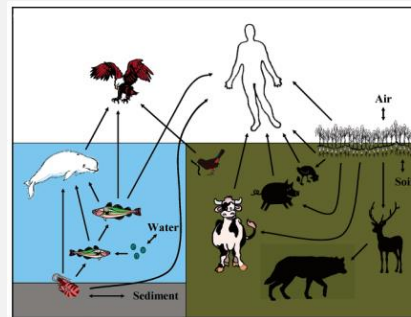


Ambient (Far-Field) Pathways

UseTox (Rosenbaum et al., 2008)



RAIDAR (Arnot et al., 2006, 2008)

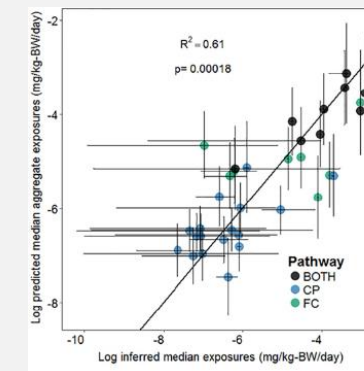


Dietary Pathways

UseTox (Rosenbaum et al. (2008)

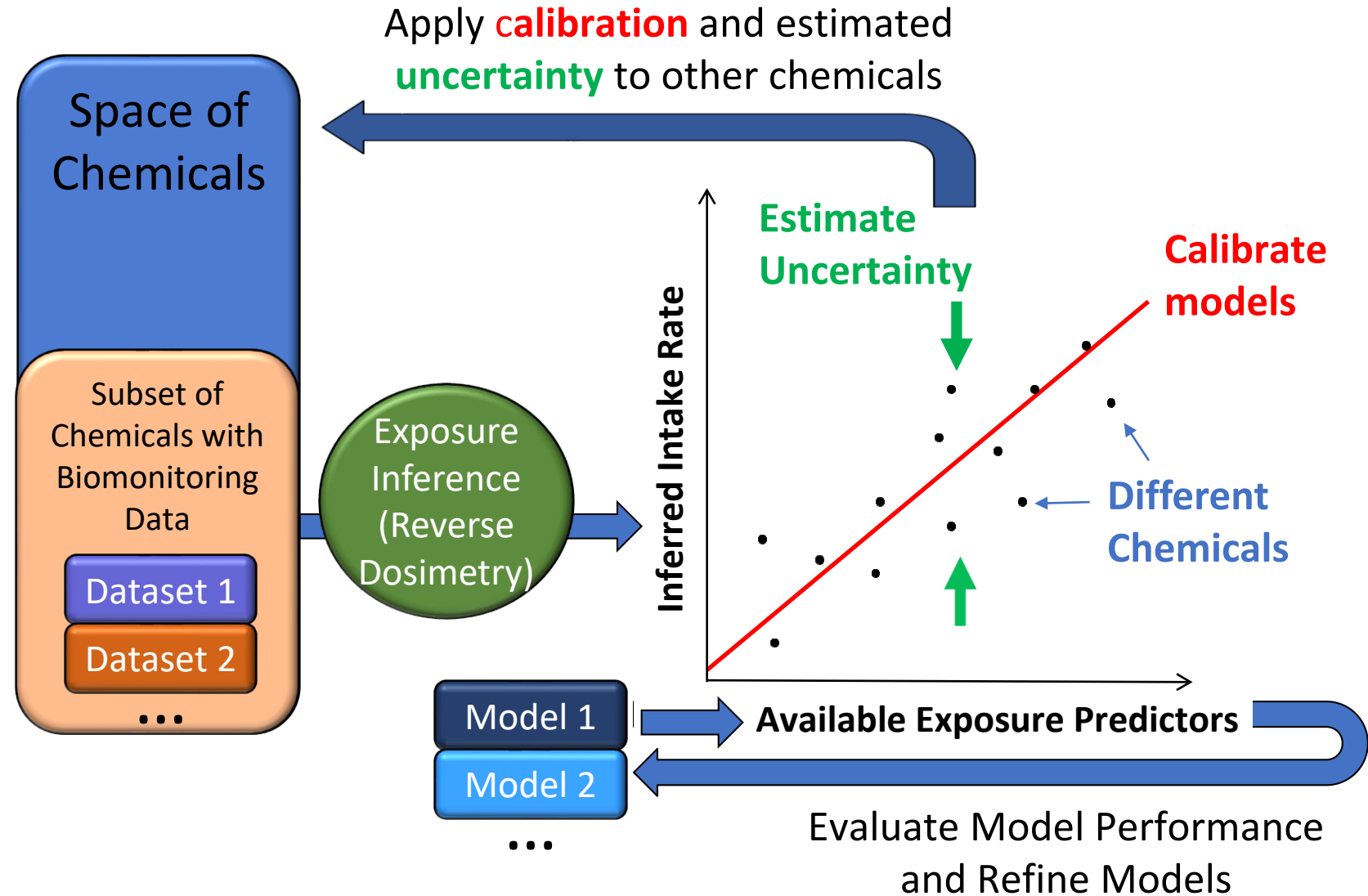


SHEDS-HT (Biryol et al., 2017)



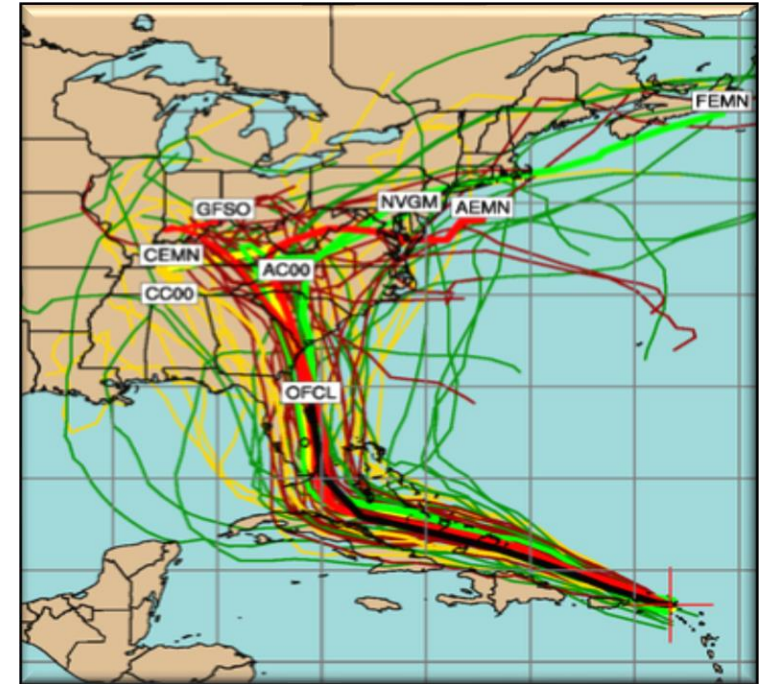
Consensus Exposure Predictions with the SEEM Framework

- Different exposure models incorporate **knowledge, assumptions, and data** (MacLeod et al., 2010)
- We incorporate multiple models (including SHEDS-HT, USEtox, RAIDAR) into consensus predictions for 1000s of chemicals within the **Systematic Empirical Evaluation of Models (SEEM)** (Wambaugh et al., 2013, 2014, Ring et al., 2019)
- Evaluation is like a sensitivity analysis: What models are working? What data are most needed?



Ensemble Predictions

- We can use ensemble methods to make more stable models and characterize uncertainty
- “Ensemble methods are learning algorithms that construct a set of classifiers and then classify new data points by taking a (weighted) vote of their predictions.”
Dietterich (2000)
- Ensemble systems have proven themselves to be very effective and extremely versatile in a broad spectrum of problem domains and real-world applications
(Polikar, 2012)
- Ensemble learning techniques in the machine learning paradigm can be used to integrate predictions from multiple tools. Pradeep (2016)



Hurricane Path Prediction is an
Example of Integrating Multiple Models
US EPA CSS-HERA BOSC Meeting – February 2-5, 2021

SEEM3 Collaboration

Jon Arnot, Deborah H. Bennett, Peter P. Egeghy, Peter Fantke, Lei Huang, Kristin K. Isaacs, Olivier Jolliet, Hyeong-Moo Shin, Katherine A. Phillips, Caroline Ring, R. Woodrow Setzer, John F. Wambaugh, Johnny Westgate



Ring et al. (2018)

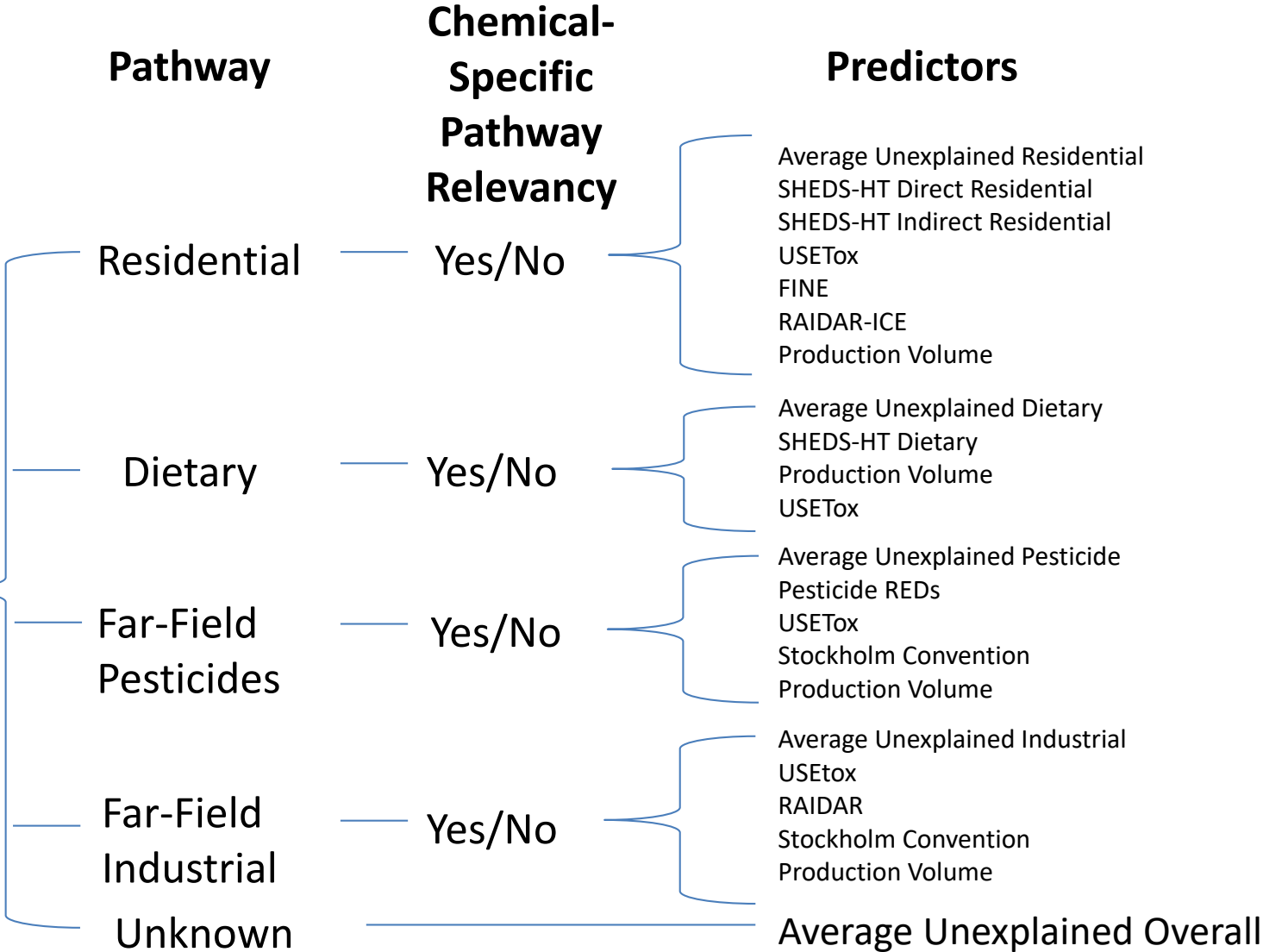
Predictor	Reference(s)	Chemicals Predicted	Pathway(s)
EPA Inventory Update Reporting and Chemical Data Reporting (CDR) (2015)	US EPA (2018)	7856	All
Stockholm Convention of Banned Persistent Organic Pollutants (2017)	Lallas (2001)	248	Far-Field Industrial and Pesticide
EPA Pesticide Reregistration Eligibility Documents (REDs) Exposure Assessments (Through 2015)	Wetmore et al. (2012, 2015)	239	Far-Field Pesticide
United Nations Environment Program and Society for Environmental Toxicology and Chemistry toxicity model (USEtox) Industrial Scenario (2.0)	Rosenbaum et al. (2008)	8167	Far-Field Industrial
USEtox Pesticide Scenario (2.0)	Fantke et al. (2011, 2012, 2016)	940	Far-Field Pesticide
Risk Assessment IDentification And Ranking (RAIDAR) Far-Field (2.02)	Arnot et al. (2008)	8167	Far-Field Pesticide
EPA Stochastic Human Exposure Dose Simulator High Throughput (SHEDS-HT) Near-Field Direct (2017)	Isaacs (2017)	7511	Far-Field Industrial and Pesticide
SHEDS-HT Near-field Indirect (2017)	Isaacs (2017)	1119	Residential
Fugacity-based INdoor Exposure (FINE) (2017)	Bennett et al. (2004), Shin et al. (2012)	645	Residential
RAIDAR-ICE Near-Field (0.803)	Arnot et al., (2014), Zhang et al. (2014)	1221	Residential
USEtox Residential Scenario (2.0)	Jolliet et al. (2015), Huang et al. (2016,2017)	615	Residential
USEtox Dietary Scenario (2.0)	Jolliet et al. (2015), Huang et al. (2016), Ernstoff et al. (2017)	8167	Dietary

SEEM3 Considers Pathway of Exposure

We organize models by the exposure pathways they cover

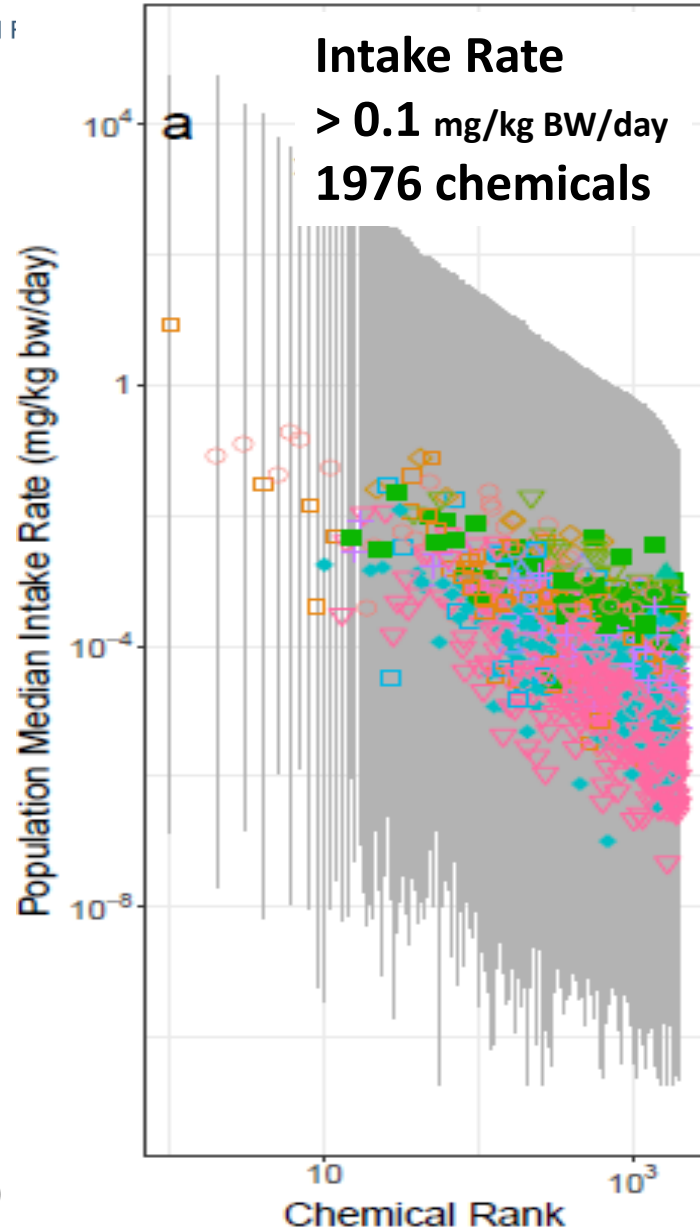
We calibrate predictors based on ability to explain median NHANES exposure rates

**General Population
Median
Chemical Exposure
(mg/kg BW/day)**



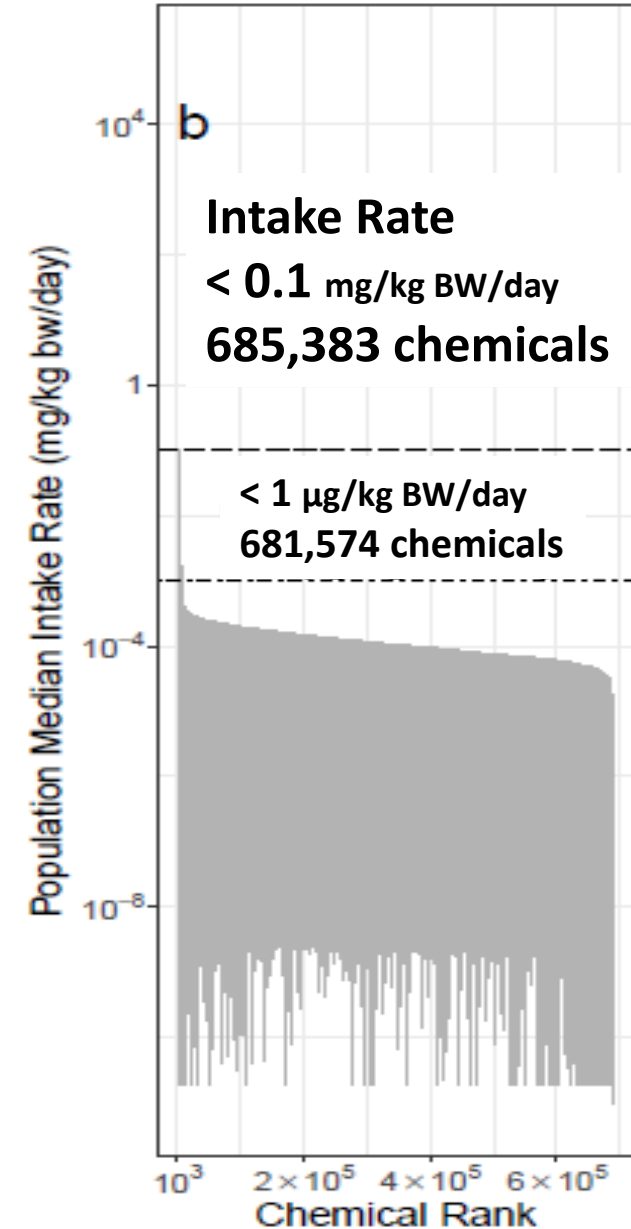
Ring et al. (2018)

Consensus Modeling of Median Chemical Intake



- Pathway(s)**
- Dietary
 - Dietary, Industrial
 - ◇ Dietary, Pesticide
 - △ Dietary, Pesticide, Industrial
 - ▽ Dietary, Residential
 - Dietary, Residential, Industrial
 - Dietary, Residential, Pesticide
 - ▲ Dietary, Residential, Pesticide, Industrial
 - ◆ Industrial
 - Pesticide
 - Pesticide, Industrial
 - △ Residential
 - + Residential, Industrial
 - × Residential, Pesticide
 - ◇ Residential, Pesticide, Industrial
 - ▽ Unknown

Of 687,359 chemicals evaluated, 30% have less than a 50% probability for exposure via any of the four pathways and are considered outside the “domain of applicability”



ExpoCast SEEM Models: Required Building Blocks for the Output

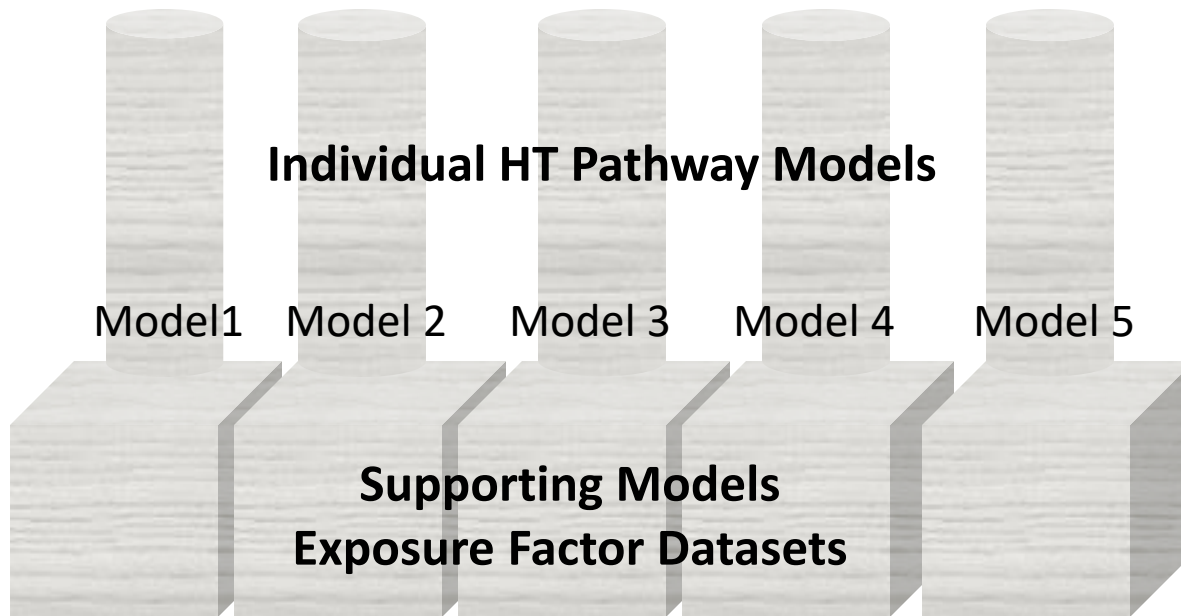


Supporting Models
Exposure Factor Datasets

Machine-learning models for filling gaps from structure when no data are available

Composition and use/release data

ExpoCast SEEM Models: Required Building Blocks for the Output

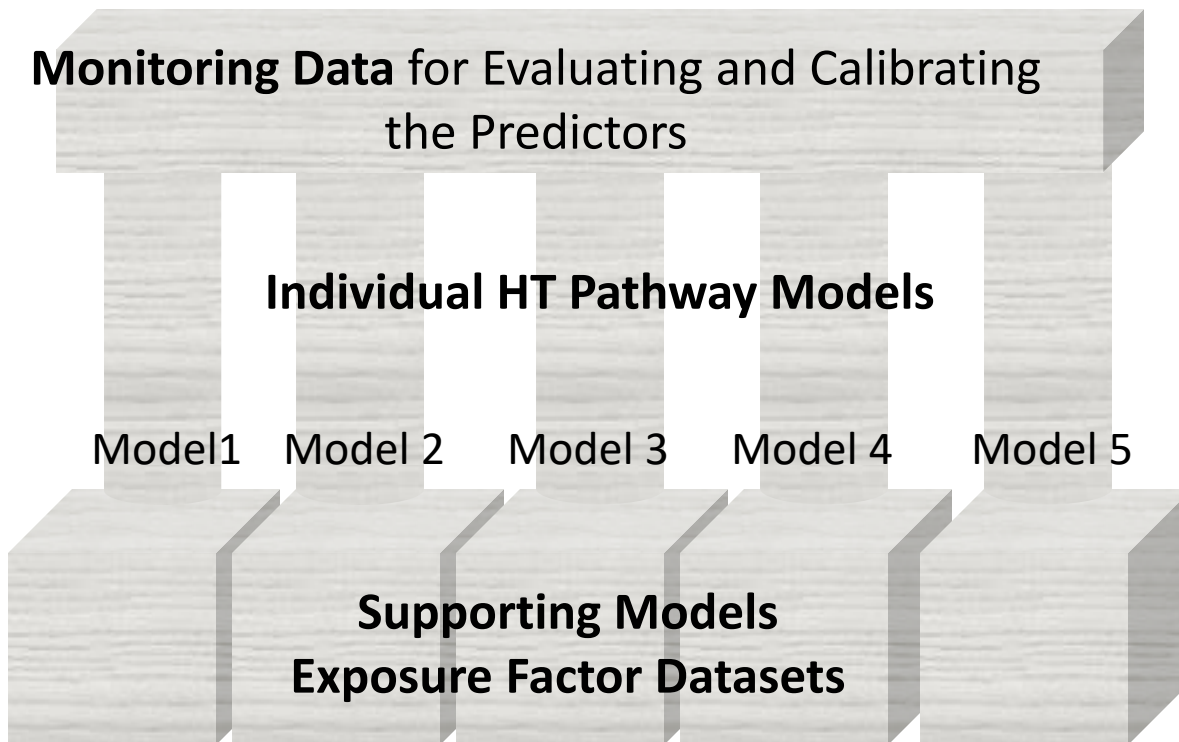


*for example, SHEDS-HT, HT ChemSteer,
external models*

*Machine-learning models for filling gaps from
structure when no data are available*

Composition and use/release data

ExpoCast SEEM Models: Required Building Blocks for the Output



Including NHANES biomonitoring and USGS water datasets

for example, SHEDS-HT, HT ChemSteer, external models

Machine-learning models for filling gaps from structure when no data are available

Composition and use/release data

ExpoCast SEEM Models: Required Building Blocks for the Output

Consensus SEEM Predictions for Receptor

Monitoring Data for Evaluating and Calibrating
the Predictors

Individual HT Pathway Models

Model 1

Model 2

Model 3

Model 4

Model 5

Supporting Models
Exposure Factor Datasets

*Including NHANES biomonitoring and
USGS water datasets*

*for example, SHEDS-HT, HT ChemSteer,
external models*

*Machine-learning models for filling gaps from
structure when no data are available*

Composition and use/release data

ExpoCast SEEM Models: Required Building Blocks for the Output

Consensus SEEM Predictions* for Receptor

Monitoring Data for Evaluating and Calibrating
the Predictors

Individual HT Pathway Models*

Model 1

Model 2

Model 3

Model 4

Model 5

Supporting Models*

Exposure Factor Datasets

*New Approach

Methodologies for Exposure:
Application to Real Decision Contexts

*Including NHANES biomonitoring and
USGS water datasets*

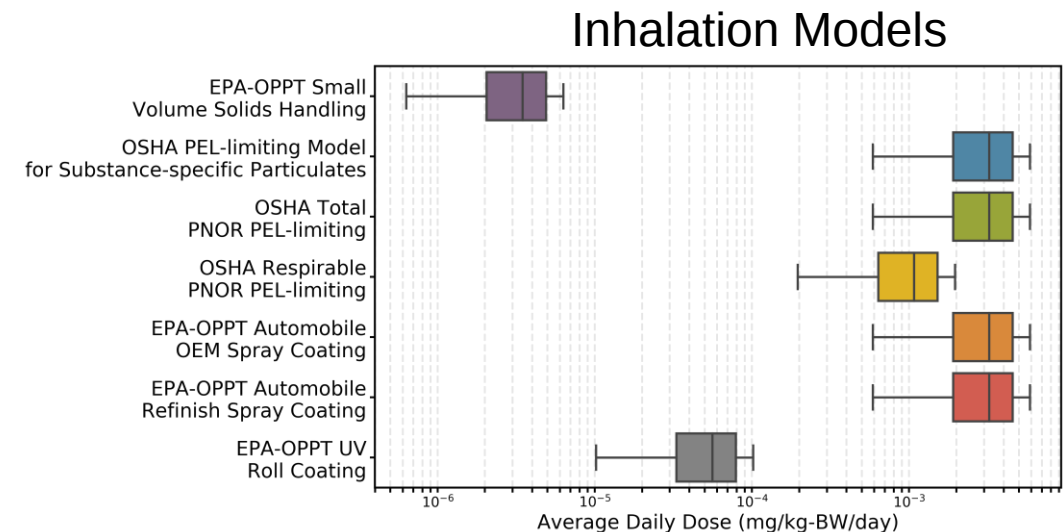
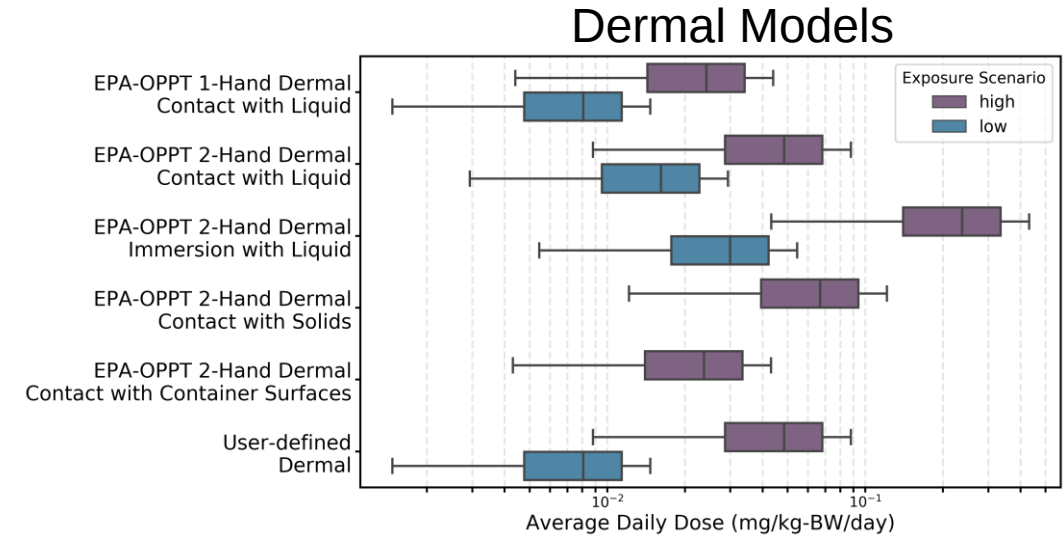
*for example, SHEDS-HT, HT ChemSteer,
external models*

*Machine-learning models for filling gaps from
structure when no data are available*

Composition and use/release data

Formatting Occupational Exposure Models for HT Use

- We have developed consensus models for consumer and some ambient pathways, but ecological and occupational consensus models are ongoing
- Many predictors for these pathways exist, but they are not typically oriented for high throughput capacity, for example EPA's ChemSTEER (Chemical Screening Tool for Exposures and Environmental Releases)
- Command Line Occupational Exposure Tool (CLOET) a command line tool that allows use of ChemSTEER v3.0 in a high throughput manner
- Multiple scenarios for each model have been run and tested against ChemSTEER GUI to test for model fidelity.



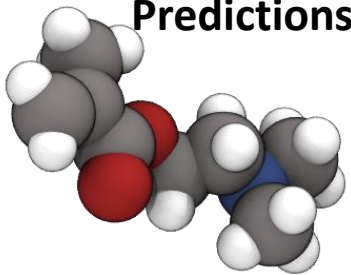
Concentrations were varied from 0.1 to 1

US EPA CSS-HERA BOSC Meeting – February 2-5, 2021

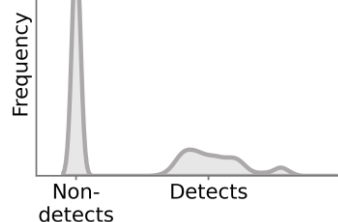
Two-Stage Occupational Exposure Model

- OSHA's chemical exposure health data set for air samples was used to build a two-stage model that predicts 1) if a chemical is likely to be detected in air and 2) what the likely concentration would be
- OPERA physicochemical property distributions across NAICS sector and subsectors are included as input distributions to the models in addition to the OSHA data
- Bayesian Hierarchical Regression allows us to organize our predictions (either detect/non-detect or concentration) by NAICS Sector and/or Subsector

OPERA Property Predictions

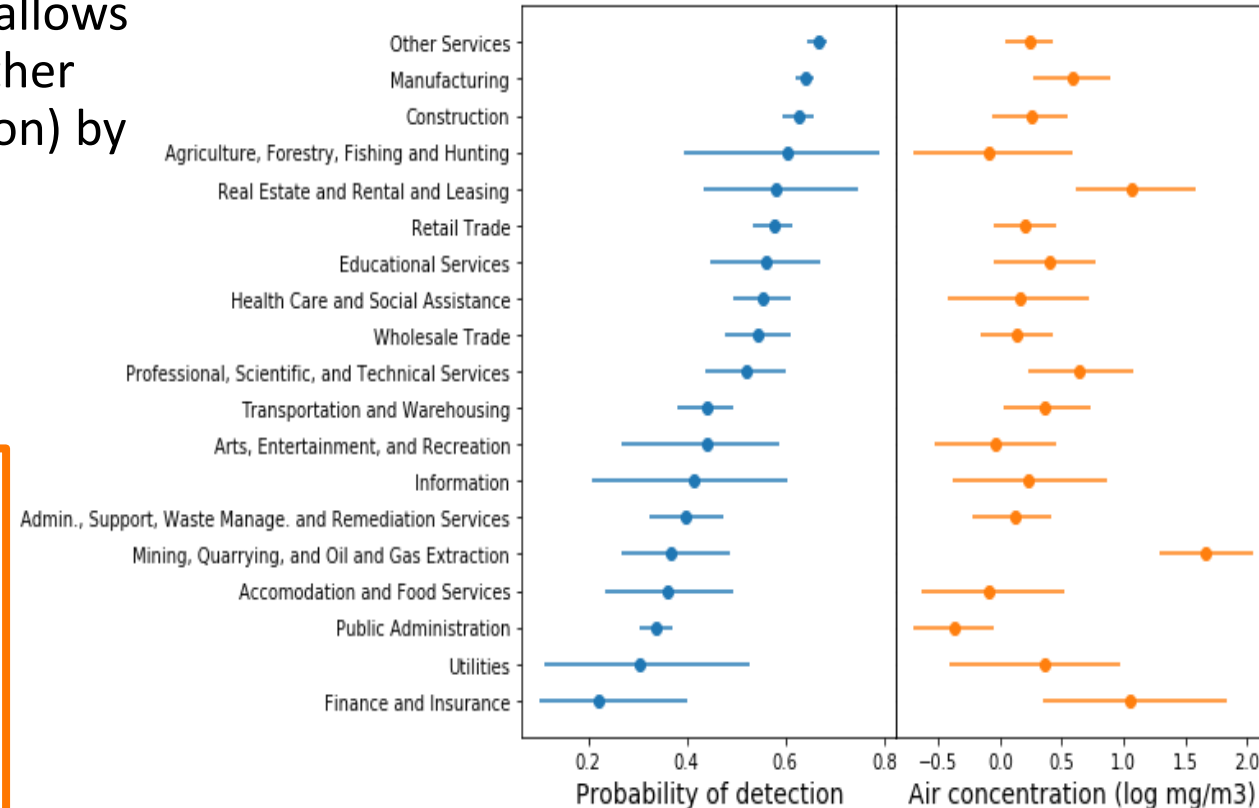
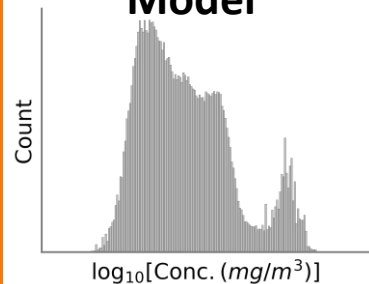


Detect / Non-detect Model



Non-detects

Air Concentration Model

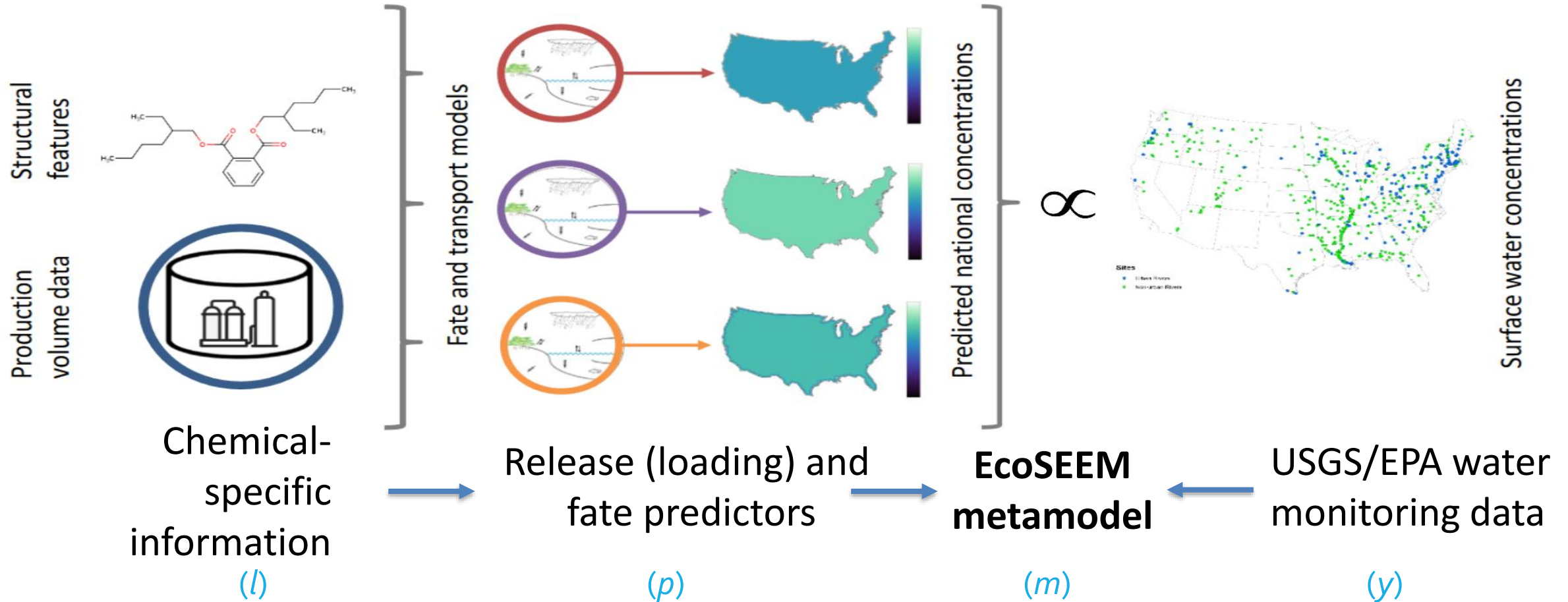


Minucci et al,
in preparation

Slide from Katherine Phillips

US EPA CSS-HERA BOSC Meeting – February 2-5, 2021

EcoSEEM Metamodel for Surface Water Chemical Concentrations



Sayre et al,
in preparation

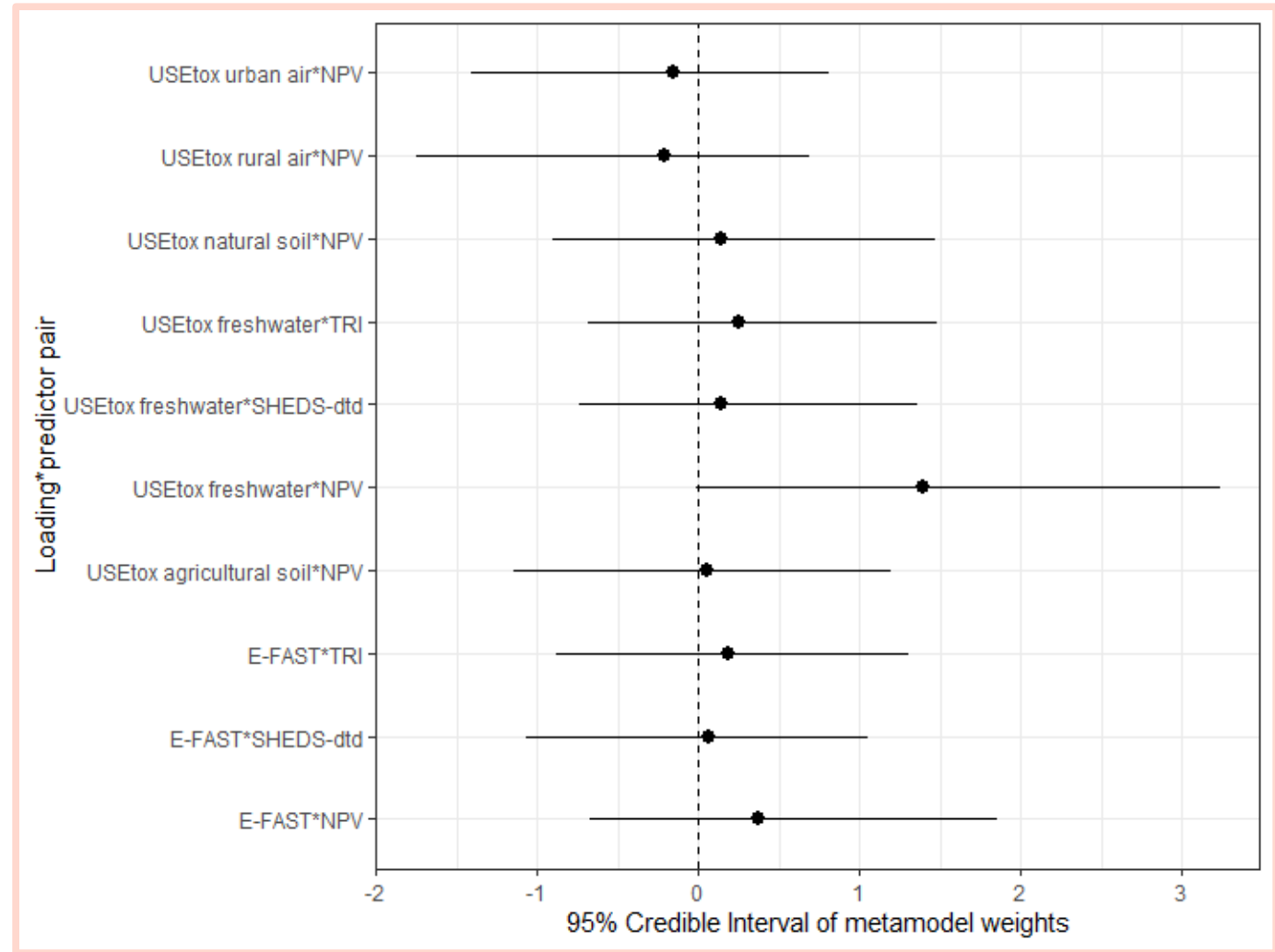
$$\ln y_i = m_0 + \sum_{j=1}^{n_j} \sum_{k=1}^{n_{kj}} m_{jk} \ln(l_{ji} p_{ki})$$

Slide from Risa Sayre

EcoSEEM Evaluating Predictive Ability of HT Surface Water Models

- The strength of the correlation between each combination of release and fate model predictions and the observed water concentrations allows model calibration
- The most informative pair for bulk concentrations was USEtox freshwater model using loadings from NPV

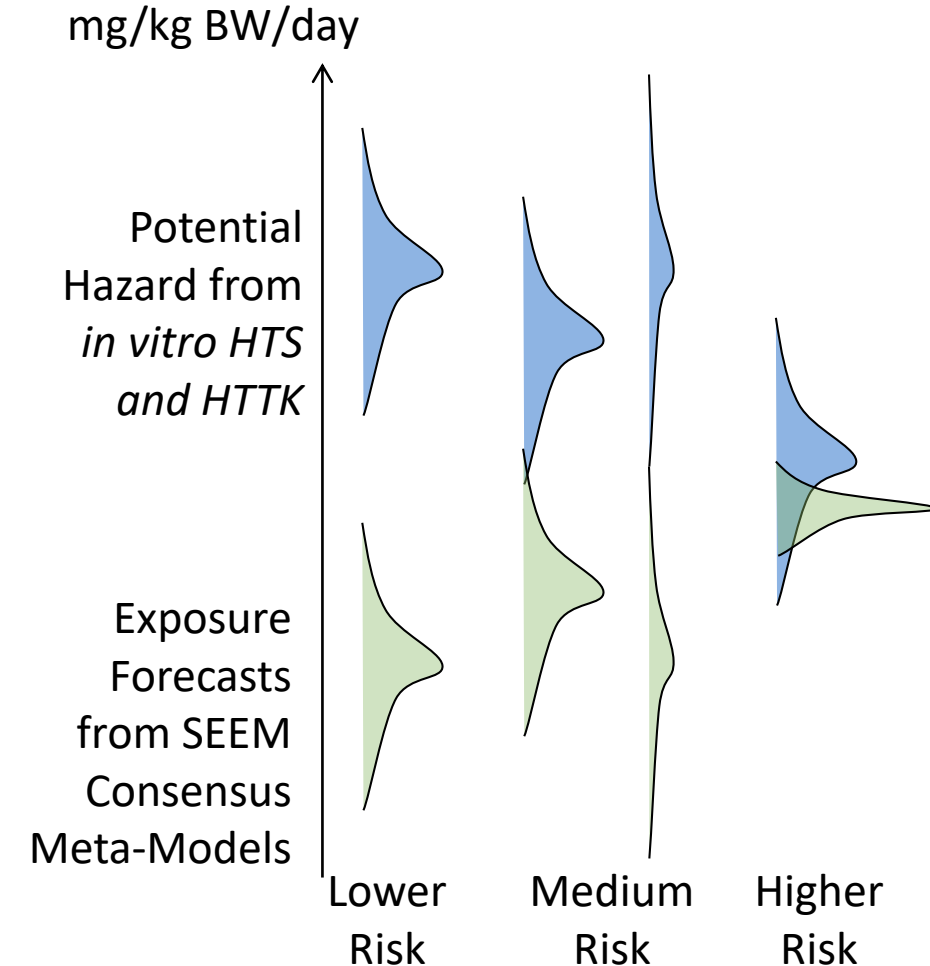
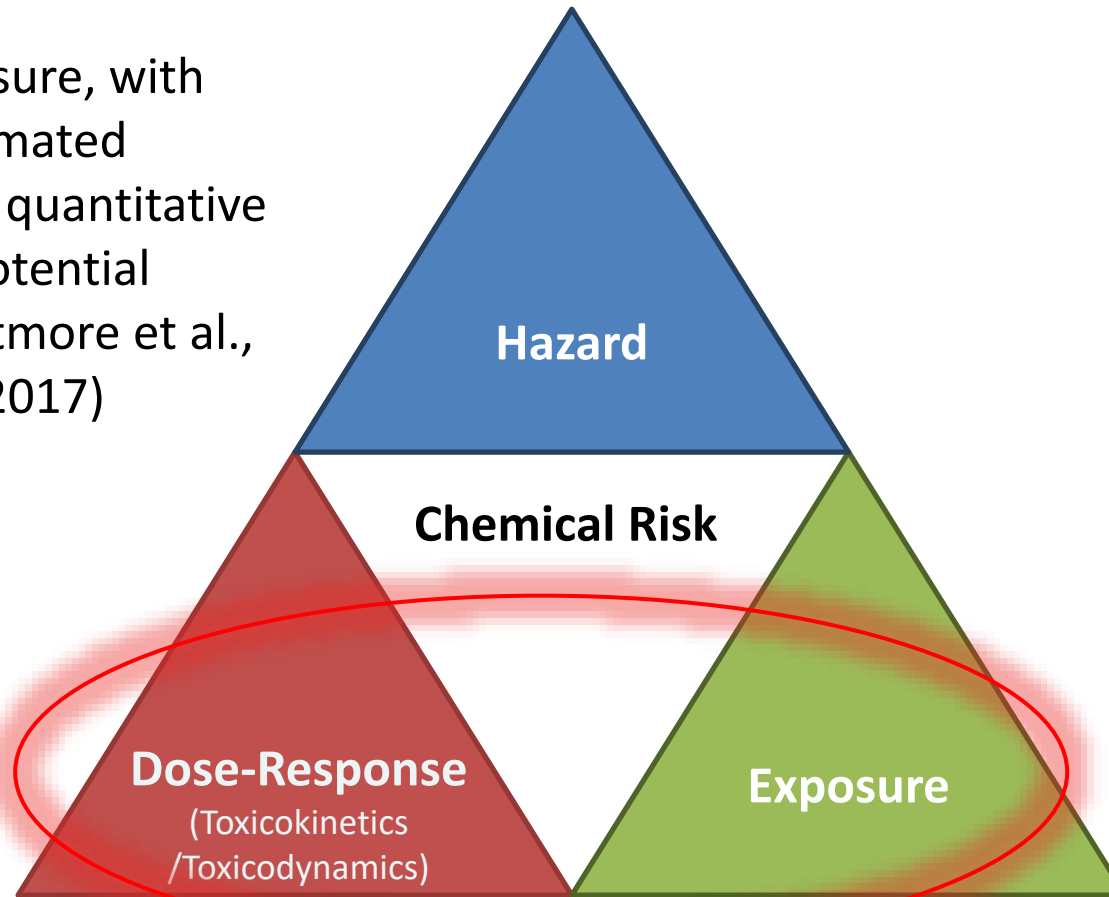
Sayre et al,
in preparation



Outlook

- SEEM metamodels have been developed for consumer and some ambient pathways (Ring et al., 2018) and ecological and occupational consensus models are in development
- Estimates of exposure, with appropriately estimated uncertainty, allow quantitative prioritization of potential chemical risk (Wetmore et al., 2015; Ring et al., 2017)

EPA's ExpoCast Project



ExpoCast Project (Exposure Forecasting)

Center for Computational Toxicology and Exposure

Linda Adams
Lucas Albrecht*
Matthew Boyce*
Miyuki Breen*
Alex Chao
Daniel Dawson*
Mike Devito
Alex East*
Lindsay Eddy*
Christopher Eklund
Peter Egeghy
Marina Evans
Alex Fisher*
Rocky Goldsmith
Louis Groff*
Chris Grulke

Colin Guider*
Mike Hughes
Victoria Hull*
Kristin Isaacs
Richard Judson
Jen Korol-Bexell*
Anna Kreutz*
Charles Lowe*
Seth Newton
Alli Phillips
Katherine Phillips
Paul Price
Tom Purucker
Ann Richard
Caroline Ring

Risa Sayre
Mark Sfeir*
Marci Smeltz*
Jon Sobus
Zach Stanfield*
Mike Tornero-Velez
Rusty Thomas
Elin Ulrich
Dan Vallero
Barbara Wetmore
John Wambaugh
Antony Williams

CEMM

Hongwan Li*
Xiaoyu Liu
Zachary Robbins*
Mark Strynar

***Trainees**

Collaborators

Arnot Research and Consulting
Jon Arnot
Johnny Westgate
Integrated Laboratory Systems
Xiaoqing Chang
Shannon Bell
National Toxicology Program
Steve Ferguson
Kamel Mansouri
Ramboll
Harvey Clewell
Silent Spring Institute
Robin Dodson
Simulations Plus
Michael Lawless
Southwest Research Institute
Alice Yau
Kristin Favela
Summit Toxicology
Lesla Aylward
Technical University of Denmark
Peter Fantke
Unilever
Beate Nicol
Cecilie Rendal
Ian Sorrell
United States Air Force
Heather Pangburn
Matt Linakis
University of California, Davis
Deborah Bennett
University of Michigan
Olivier Jolliet
University of Texas, Arlington
Hyeong-Moo Shin
University of Nevada
Li Li
University of North Carolina, Chapel Hill
Julia Rager
Marc Serre

References

EPA ORD Publications in Bold

- Annot, J.A., et al. 2006. Screening level risk assessment model for chemical fate and effects in the environment. *Environ. Sci. Technol.* 40: 2316-2323
- Annot, J.A.; Mackay D. 2008. Policies for chemical hazard and risk priority setting: Can persistence, bioaccumulation, toxicity and quantity information be combined? *Environ. Sci. Technol.* 42: 4648-4654. DOI: 10.1021/es800106g
- Barber MC, et al (2017). "Developing and applying metamodels of high resolution process-based simulations for high throughput exposure assessment of organic chemicals in riverine ecosystems." *Science of the Total Environment*, 605, 471-481.**
- Bennett, D. H.; Furtaw, E. J., Fugacity-based indoor residential pesticide fate model. *Environmental Science & Technology* 2004, 38, (7), 2142-2152.**
- Biryol D, et al.. High-throughput dietary exposure predictions for chemical migrants from food contact substances for use in chemical prioritization. *Environment international*. 2017 Nov 1;108:185-94.**
- Dietterich, Thomas G. "Ensemble methods in machine learning." *International workshop on multiple classifier systems*. Springer, Berlin, Heidelberg, 2000. **Ernstoff, A. S , et al., High-throughput migration modelling for estimating exposure to chemicals in food packaging in screening and prioritization tools. *Food and Chemical Toxicology* 2017, 109, 428-438.**
- Fantke P, Jolliet O. Life cycle human health impacts of 875 pesticides. *The International Journal of Life Cycle Assessment*. 2016 May 1;21(5):722-33.
- Fantke P , et al.. Dynamic toxicity modelling based on the USEtox matrix framework. In SETAC Europe 25th Annual Meeting 2015 (pp. 33-34). SETAC Europe.
- Huang, L , et al., A review of models for near-field exposure pathways of chemicals in consumer products. *Science of The Total Environment* 2017, 574, 1182-1208.
- Huang, L.; Jolliet, O., A parsimonious model for the release of volatile organic compounds (VOCs) encapsulated in products. *Atmospheric Environment* 2016, 127, 223-235.
- Isaacs KK , et al.. SHEDS-HT: an integrated probabilistic exposure model for prioritizing exposures to chemicals with near-field and dietary sources. *Environmental science & technology*. 2014 Nov 4;48(21):12750-9.**
- Jolliet, O , et al., P., Defining Product Intake Fraction to Quantify and Compare Exposure to Consumer Products. *Environmental Science & Technology* 2015, 49, (15), 8924-8931.
- Lallas PL. The Stockholm Convention on persistent organic pollutants. *The American Journal of International Law*. 2001 Jul 1;95(3):692-708.
- Li L, et al.. A model for risk-based screening and prioritization of human exposure to chemicals from near-field sources. *Environmental science & technology*. 2018 Nov 8;52(24):14235-44.
- MacLeod, Matthew, et al. "The state of multimedia mass-balance modeling in environmental science and decision-making." (2010): 8360-8364. Minucci et al, *in preparation*
- Minucci, Jeff et al. "High Throughput Model for Occupational Exposure"**
- Polikar, Robi. "Ensemble learning." *Ensemble machine learning*. Springer, Boston, MA, 2012. 1-34.
- Pradeep, Prachi, et al. "An ensemble model of QSAR tools for regulatory risk assessment." *Journal of cheminformatics* 8.1 (2016): 1-9.
- Rosenbaum RK, et al. USEtox—the UNEP-SETAC toxicity model: recommended characterisation factors for human toxicity and freshwater ecotoxicity in life cycle impact assessment. *The International Journal of Life Cycle Assessment*. 2008 Nov 1;13(7):532.
- Ring, Caroline L., et al. "Identifying populations sensitive to environmental chemicals by simulating toxicokinetic variability." *Environment international* 106 (2017): 105-118.**
- Ring CL, et al. Consensus modeling of median chemical intake for the US population based on predictions of exposure pathways. *Environmental science & technology*. 2018 Dec 5;53(2):719-32.**
- Sayre et al, "Consensus Model for Predicting Chemical Surface Water Concentrations"**
- Shin, H.-M., et al., Intake Fraction for the Indoor Environment: A Tool for Prioritizing Indoor Chemical Sources. *Environmental Science & Technology* 2012, 46, (18), 10063-10072.
- Shin, Hyeong-Moo, et al. "Risk-based high-throughput chemical screening and prioritization using exposure models and in vitro bioactivity assays." *Environmental science & technology* 49.11 (2015): 6760-6771
- Wambaugh, John F., et al. "High-throughput models for exposure-based chemical prioritization in the ExpoCast project." *Environmental science & technology* 47.15 (2013): 8479-848.**
- Wambaugh, John F., et al. "High Throughput Heuristics for Prioritizing Human Exposure to Environmental Chemicals." *Environmental science & technology* (2014).**
- Wambaugh JF, et al. New approach methodologies for exposure science. *Current Opinion in Toxicology*. 2019 Jun 1;15:76-92.**
- Wetmore, B. A., et al. Incorporating High-Throughput Exposure Predictions With Dosimetry-Adjusted In Vitro Bioactivity to Inform Chemical Toxicity Testing. *Toxicol. Sci.* 2015, 148 (1), 121–36.**
- Zhang, X.; Annot, J. A.; Wania, F., Model for screening-level assessment of near-field human exposure to neutral organic chemicals released indoors. *Environmental Science & Technology* 2014, 48, (20), 12312-12319.

High-Throughput Toxicokinetic Models and *In Vitro-In Vivo* Extrapolation (IVIVE)

Barbara A. Wetmore

US EPA CSS-HERA Board of Scientific Counselors
Chemical Safety Subcommittee Meeting

February 2, 2021

*The views expressed in this presentation are those of the author
and do not necessarily reflect the views or policies of the US EPA.*

NAMs for Exposure Toxicokinetics

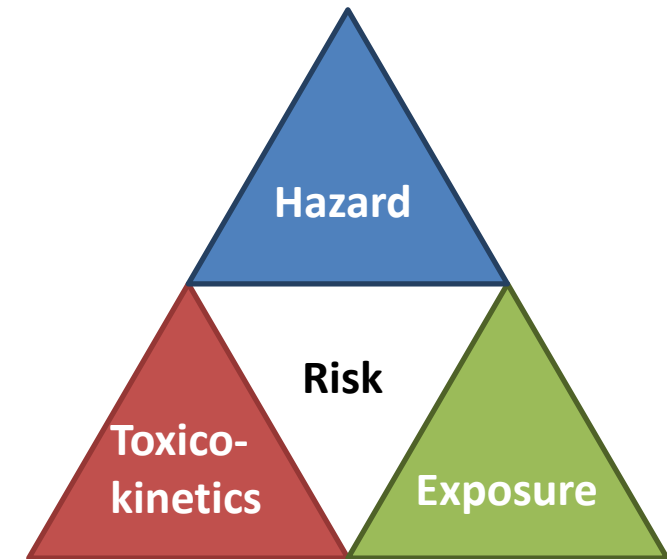
Acceptance and use of in vitro data for hazard identification is limited by uncertainties associated with exposure characterization and metabolism

Many *in vitro* systems:

- lack consideration of biotransformation capabilities
 - Overestimation of hazard for chemicals rapidly cleared *in vivo*
 - Underestimation of hazard for chemicals bioactivated *in vivo*
- lack consideration of exposure route
- lack consideration of susceptible populations / life stages
- *In vitro* potency estimates are often not adjusted for chemical availability in the *in vitro* system (ie, *in vitro* disposition)

Recent Agency Case Study Finding:

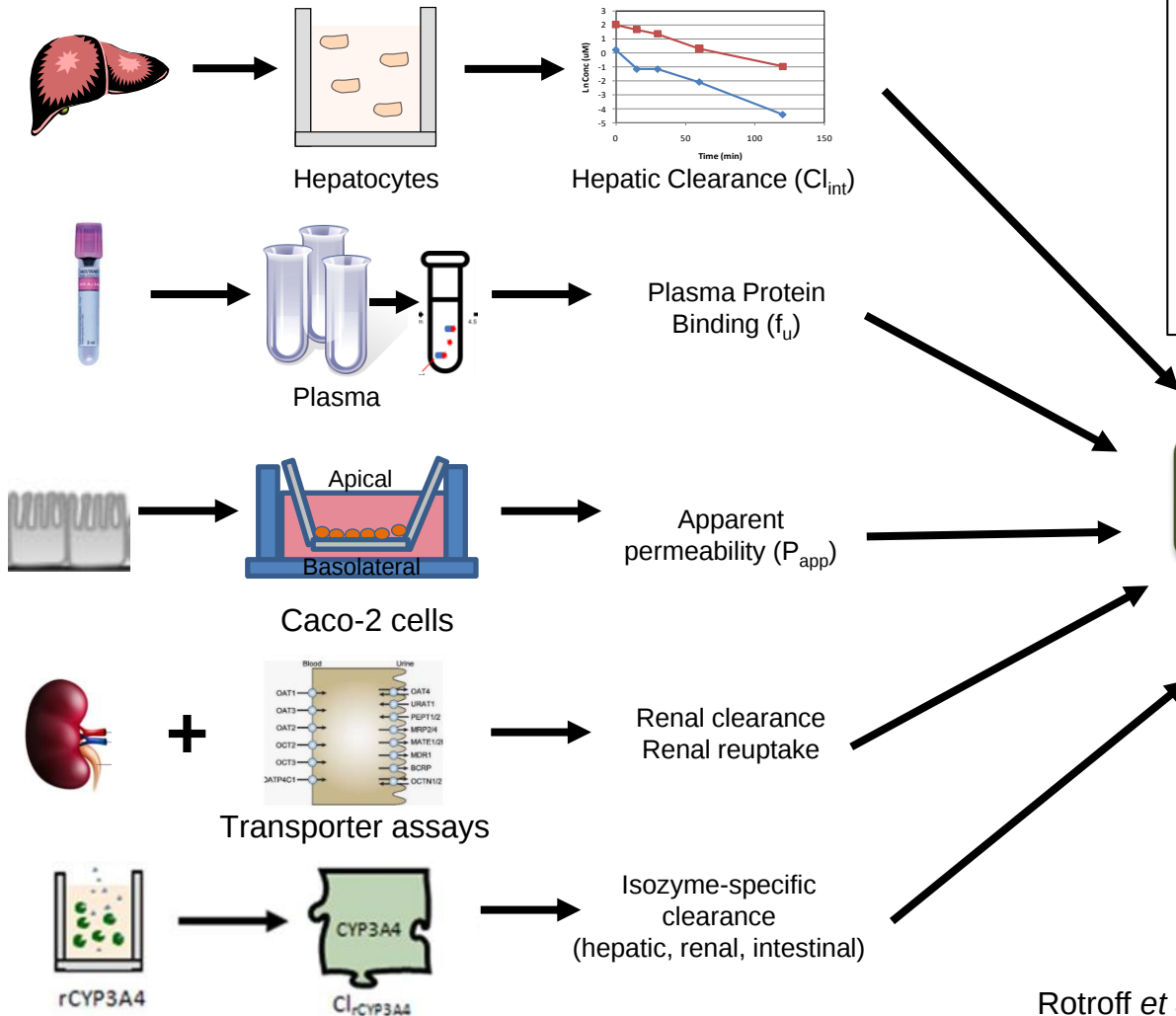
→ TK data availability rate limiting factor in TSCA screening for chemical prioritization



*“A Proof-of-Concept Case Study Integrating Publicly Available Information to Screen Candidates for Chemical Prioritization under TSCA”

In Vitro-In Vivo Extrapolation (IVIVE)

I. In Vitro Toxicokinetic Assays



IVIVE originally used and vetted in pharma applications
HT-IVIVE approach uses

- hepatic clearance
- plasma protein binding
- conservative assumptions

Predictions consistently protective of human health

IVIVE → Internal Concentration Predictions Given a Set Administered Dose

Ongoing efforts will:

- Incorporate additional TK inputs for better predictivity
- Assess impact of transporter involvement
- Evaluate extent of population variability
- Employ experimental measures to develop predictive tools

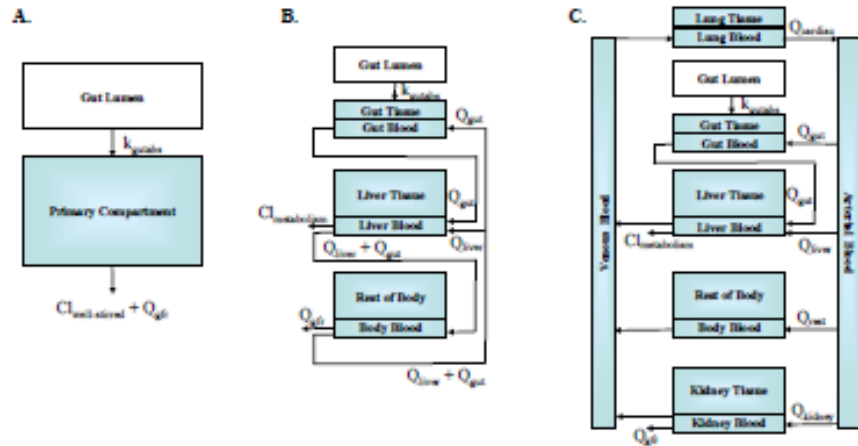
Rotroff et al., *Tox Sci.*, 2010
Wetmore et al., *Tox Sci.*, 2012
Wetmore et al., *Tox Sci.*, 2014

Wetmore et al., *Tox Sci.*, 2015
Wambaugh et al., *Tox Sci.*, 2015
Honda et al., 2019

Wambaugh et al., 2019
Smeltz et al., in preparation
Kreutz et al., in preparation

In Vitro-In Vivo Extrapolation

II. Physiologically-based Toxicokinetic Modeling



Evolving Capabilities

- Augmentation of PBTK models based on need
- Expanding to incorporate additional TK data (intestinal, renal compartments)
- Incorporating additional exposure routes
- Incorporating additional pathways (gestational)
- Incorporating demographic info to expand population-based info (variability)

“httk”: Open-source modeling package

Modeling Platform incorporates:

- chemical-specific inputs (TK data, physico-chemical)
- physiologic inputs (blood flow rates, tissue size)

into *Simulations* set up for:

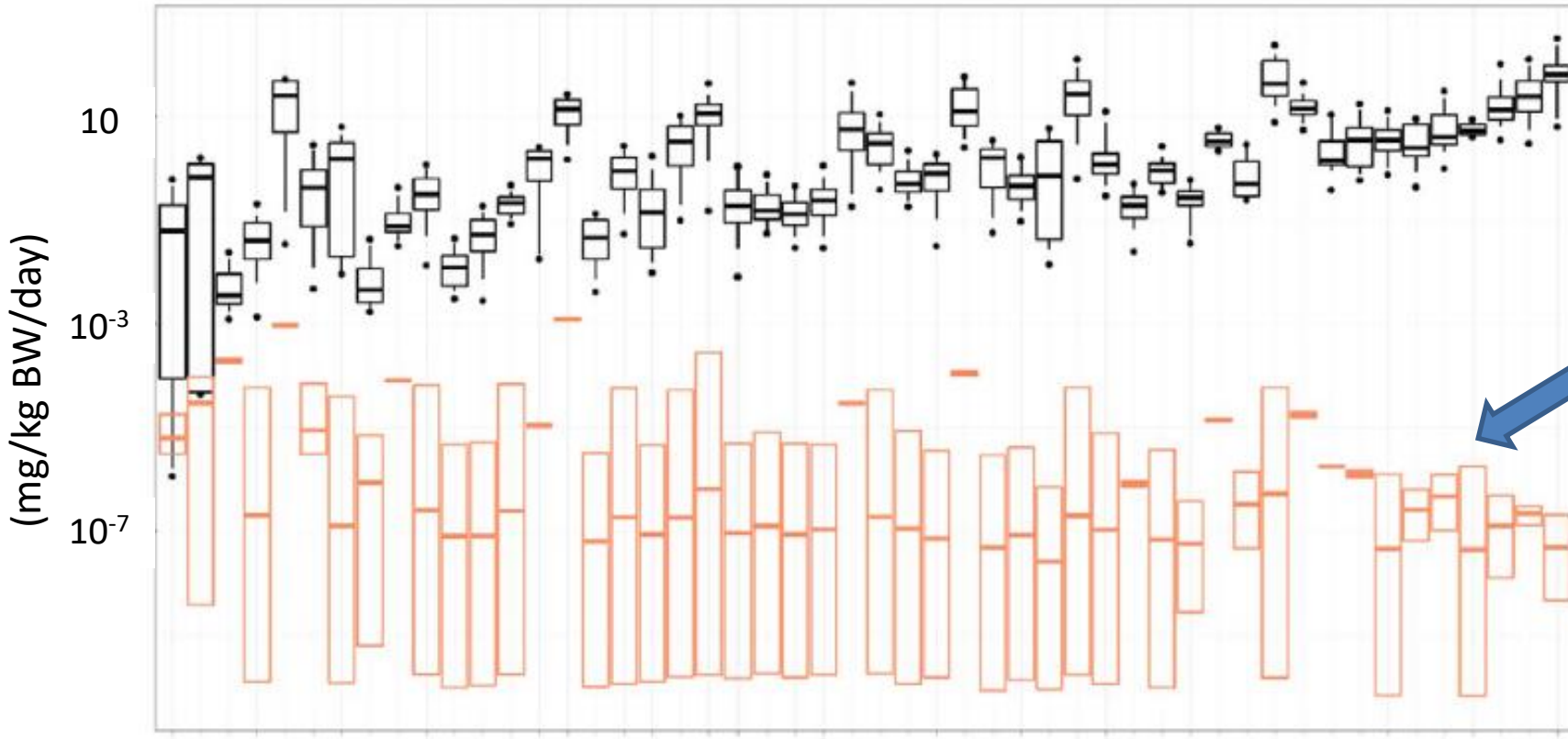
- populations of interest
 - exposures of interest
 - Capturing variability (within or across populations)
- Based on variations in the physiologic inputs (Monte Carlo)

Pearce *et al.*, 2017, *J Statistical Software*

NAMs for Prioritization

Integrating Hazard, TK, and Exposure

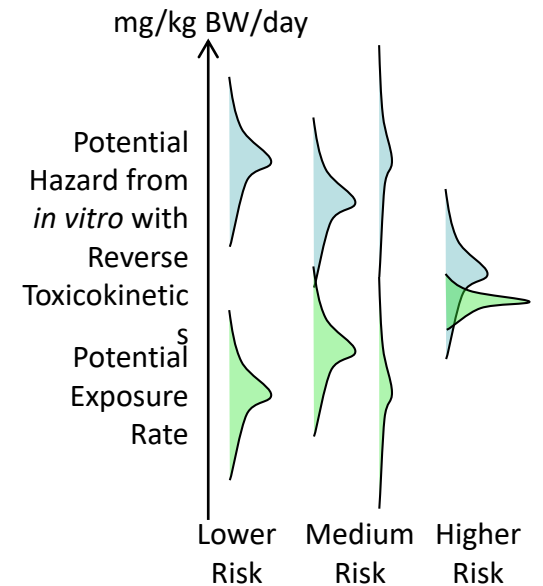
Estimated Equivalent Dose or Predicted Exposure
(mg/kg BW/day)



Chemicals Monitored by CDC NHANES

High throughput *in vitro* screening can be used to estimate doses needed to cause bioactivity

Exposure intake rates can be inferred from biomarkers



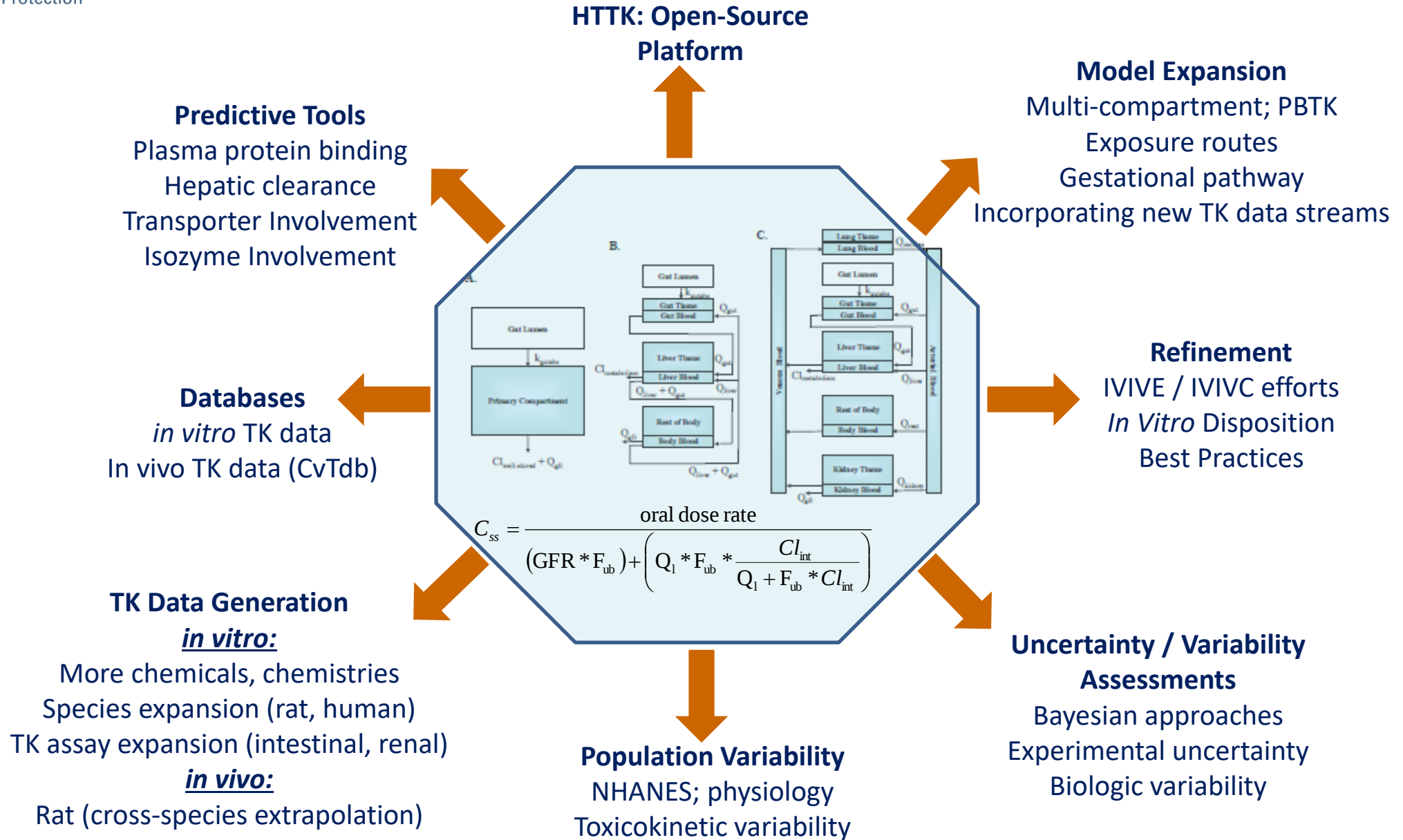
Wambaugh *et al.*, 2014
Wetmore *et al.*, 2015
Ring *et al.* (2017)
And others...

Toxicokinetics and IVIVE – Stakeholder Needs

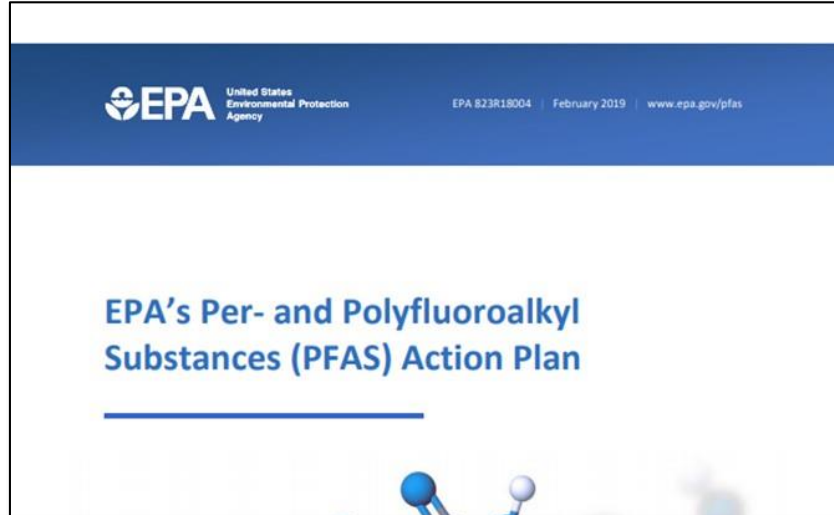
Ongoing Development of Toxicokinetic and IVIVE Tools for use in NAMs

- Primary goal: to provide a human exposure-dose context for bioactive *in vitro* concentrations from NAMs for hazard testing
 - TK Methods across TSCA landscape – including *challenging chemistries, emerging contaminants*
 - Incorporating more exposure routes and pathways
 - Tools to characterize exposures to sensitive populations and life stages
 - Characterize *in vitro* disposition across TSCA landscape
 - Tools to identify, quantitate and/or reduce sources of uncertainty
- Secondary goal: to provide **open-source data and models** for evaluation and use by the broader scientific community
 - Concomitant incorporation of above tools and data in HTK package
 - Databases with *in vitro*, *in vivo* data for use in IVIVE evaluations, *in silico* tool development

Rapid Exposure Modeling and Dosimetry



- *In Vitro* Toxicokinetic Data Generation - PFAS: Using NAMs to Fill Information Gaps

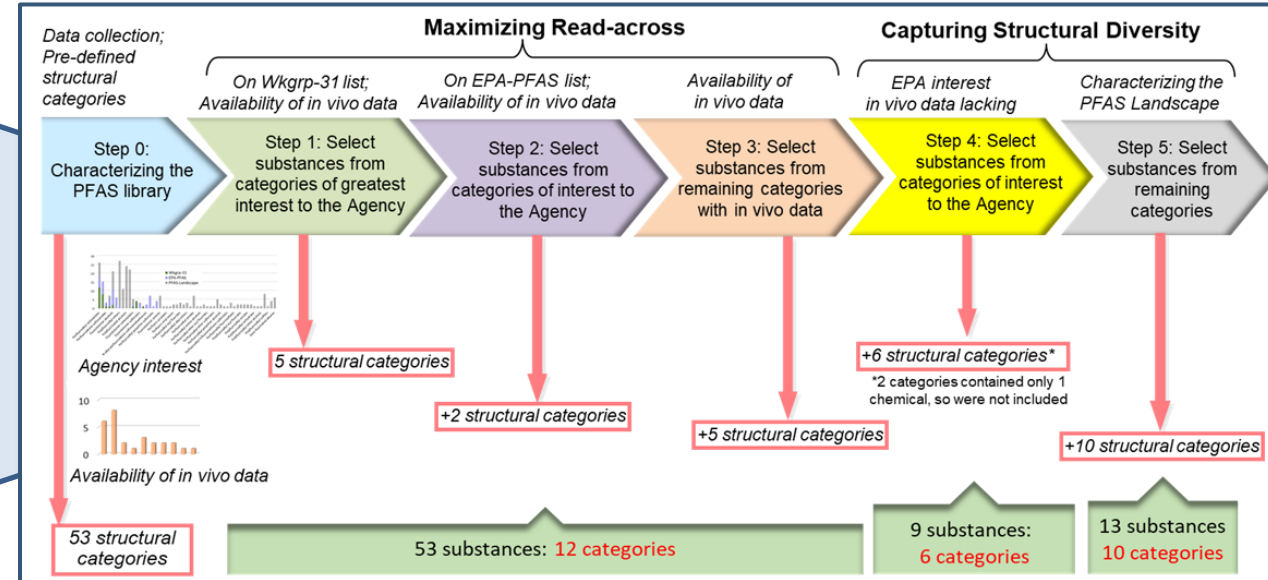


Goals:

- Generate data to support development and refinement of categories and read-across evaluation
- Incorporate substances of interest to Agency
- Characterize mechanistic and toxicokinetic properties of the broader PFAS landscape

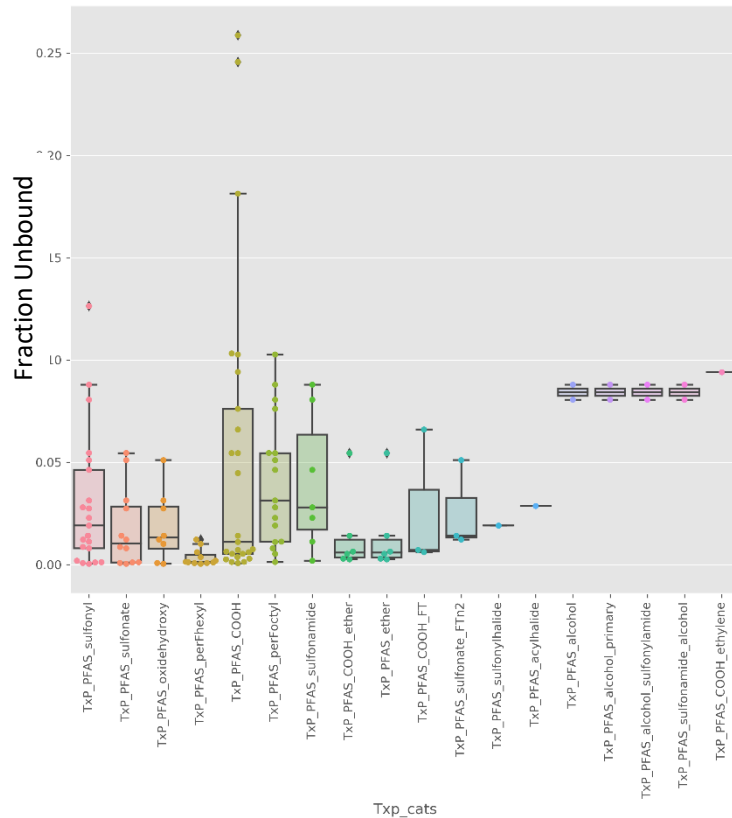
Research Area 1: What are the human health and ecological effects of exposure to PFAS?

"... the EPA plans to use new approaches such as high throughput and computational approaches to explore different chemical categories of PFAS... to inform hazard characterization, and to promote prioritization of chemicals ..."



- In Vitro Toxicokinetic Data Generation - Category-Based Analyses of Toxicokinetic Data

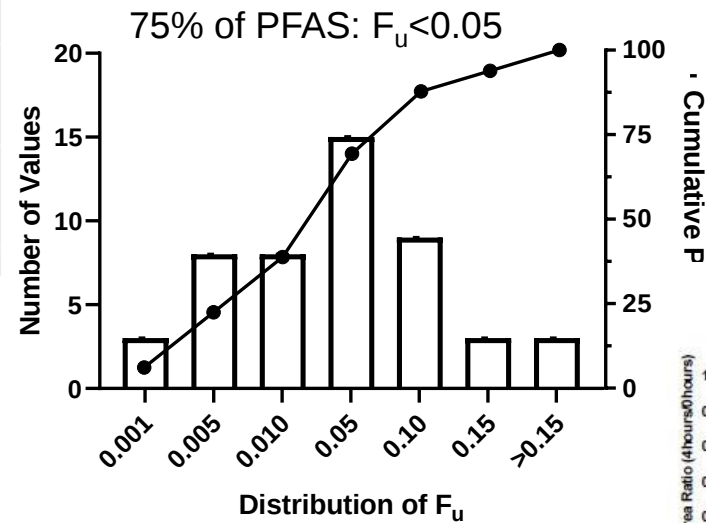
Category-Based Analysis of Plasma Protein Binding Data



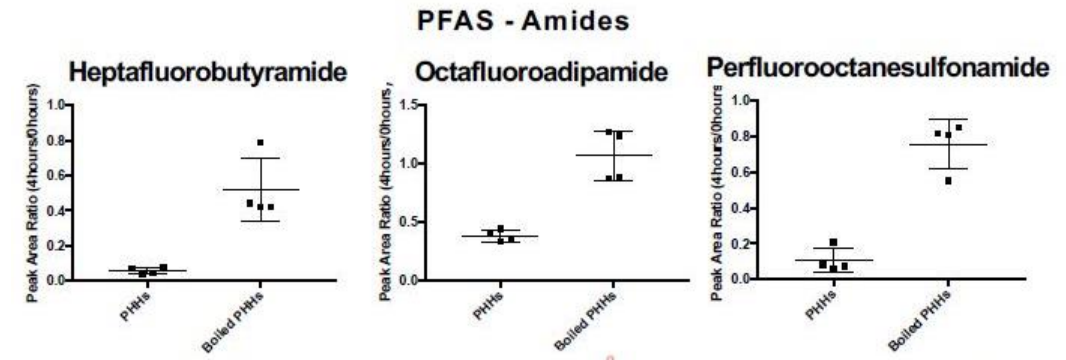
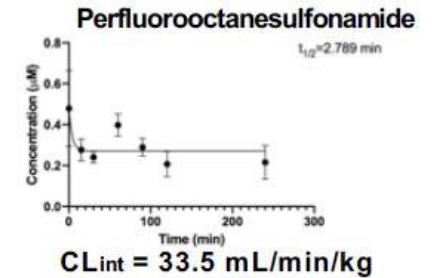
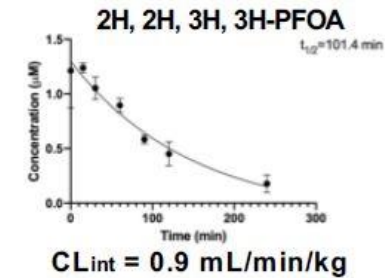
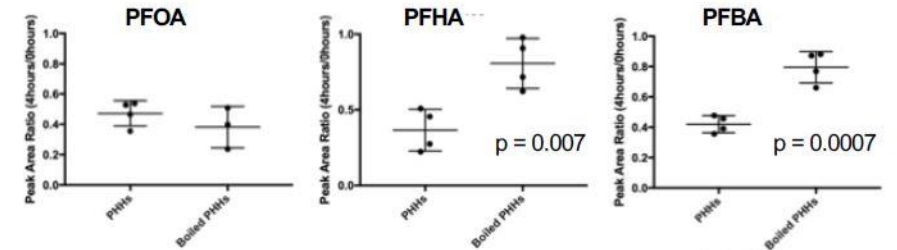
PFAS TK data: ~150 PFAS

- Hepatic clearance
 - Plasma protein binding
 - Renal transporter activity
- IVIVE, modeling, TK NAMs

Preliminary set: Plasma protein binding data across 50+ PFAS



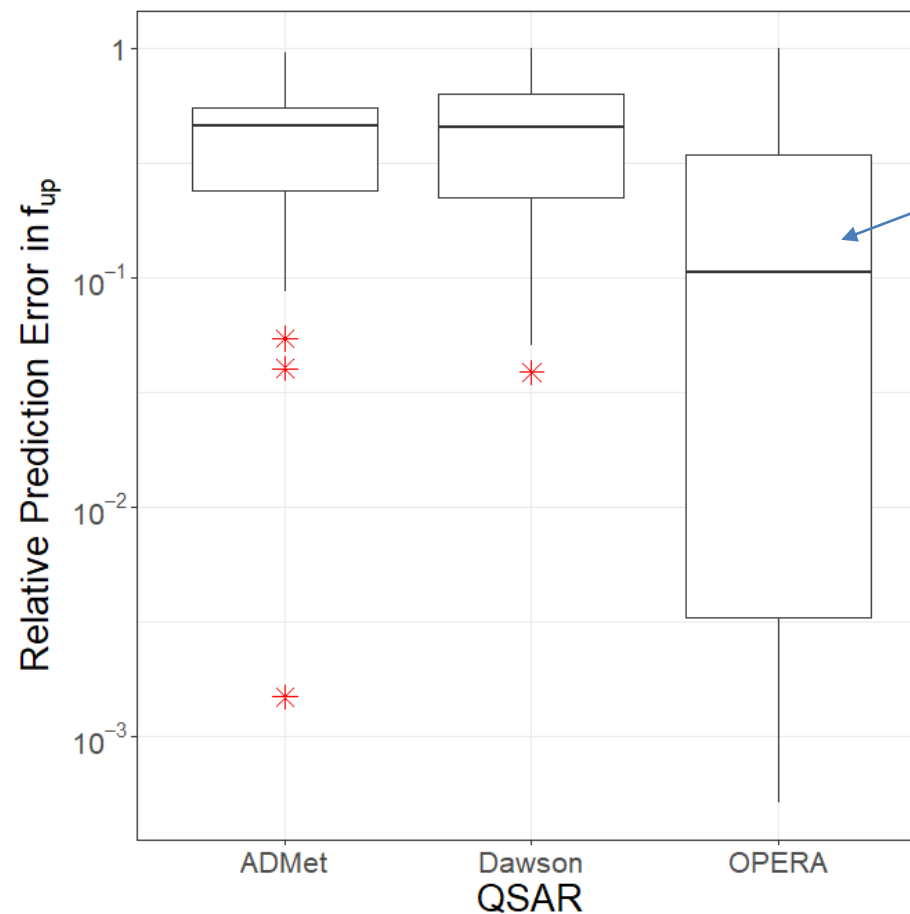
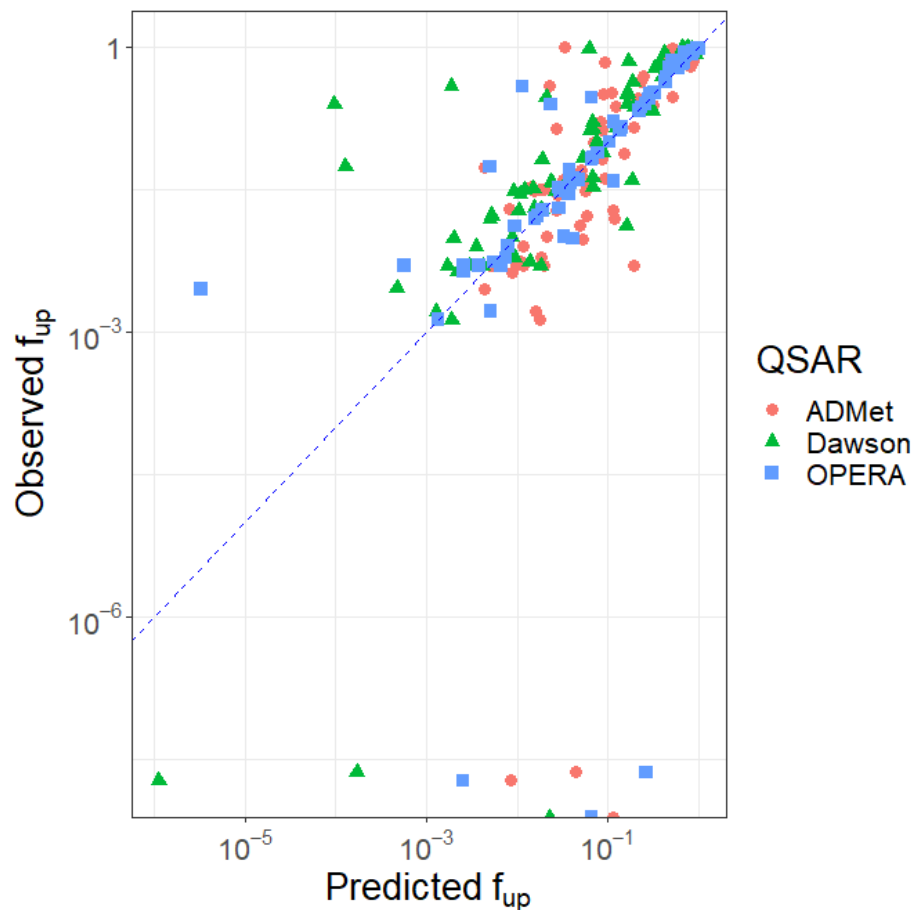
Hepatic Clearance Data



- Predictive Tool Development -

- *In vitro* TK measurements are being employed in model development and evaluation.
- Plasma protein binding (f_u); hepatic clearance (Cl_{int}) underway; others to follow.

In silico predictions for f_u (plasma protein binding)



This method uses nearest neighbors, and many evaluation chemicals are in training set

Dawson *et al.* submitted
Pradeep *et al.*, 2020
Tornero-Velez *et al.*, underway
Sipes *et al.*, 2017

- Model Expansion - Gestational Pathway

RESEARCH ARTICLE

Empirical models for anatomical and physiological changes in a human mother and fetus during pregnancy and gestation

Dustin F. Kapraun^{1*}, John F. Wambaugh², R. Woodrow Setzer², Richard S. Judson²

¹ National Center for Environmental Assessment, US Environmental Protection Agency, Research Triangle Park, North Carolina, United States of America, ² National Center for Computational Toxicology, US Environmental Protection Agency, Research Triangle Park, North Carolina, United States of America

* kapraun.dustin@epa.gov

Table 1. Itemized comparison of selected publications that contain one or more formulae related to human gestation and pregnancy.

Manuscript	[27]	[15]	[25]	[28]	[29]	[31]	[3]	[32]
Presents original data*	N	N	N	N	N	N	N	N
Presents original compiled data* set(s)	Y	N	N	Y	N	N	Y	Y
Presents original models [†] based on compiled data sets of Abduljalil et al. [28]	N	N	N	Y	Y	N	N	N
Presents original models [†] based on compiled data sets of Abduljalil et al. [33]	N	N	N	N	N	N	N	N
(+) Employs and thoroughly describes rigorous statistical methods for parameter [‡] estimation	Y	N	Y	N	N	N	Y	N
(+) Employs and thoroughly describes rigorous statistical methods for model [†] selection	N	N	N	N	N	N	Y	N
(+) Presents original models [‡] for multiple maternal compartments	N	Y	N	Y	Y	Y	Y	N
(+) Presents original models [‡] for multiple fetal compartments	N	Y	Y	N	N	Y	N	Y
(+) Presents models that reflect a biologically accurate depiction of the fetal circulatory system [§]	N	N	N	N	N	Y	N	N*
(+) Presents explicit models [‡] for “rest of body” compartments that yield feasible (e.g., non-negative) values for all relevant time points	N	N	N	N	N	N	N	Y
(+) Systematically compares original models [‡] with previously published models [‡]	N	N	N	N	N	N	N	N
(-) Presents models that contain errors or inconsistencies identified in the current manuscript	N	Y	N	N	Y	N	Y	Y

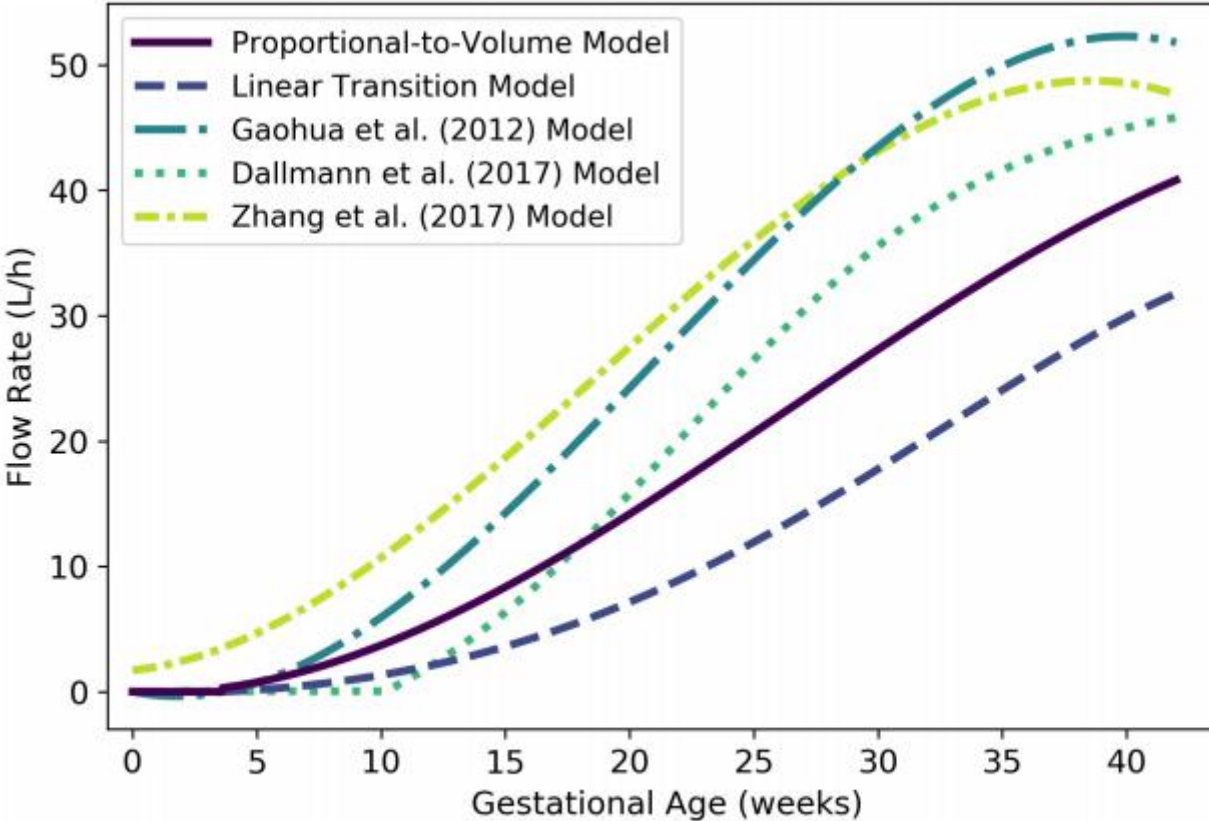
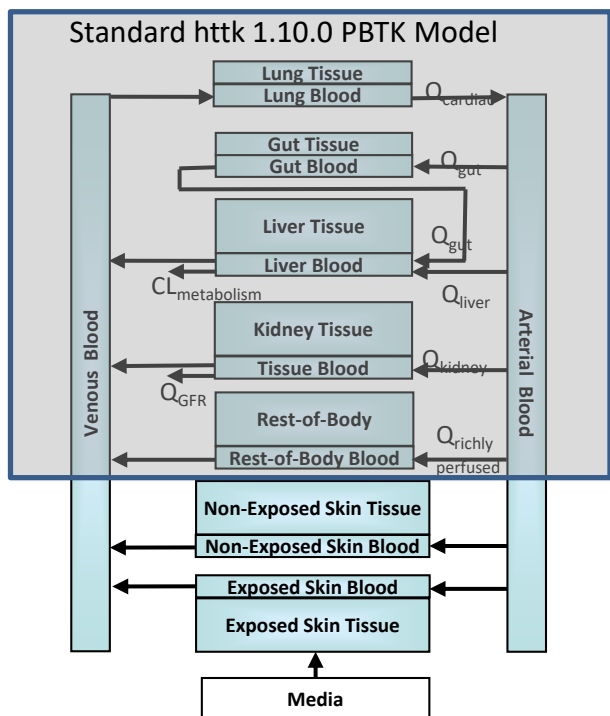


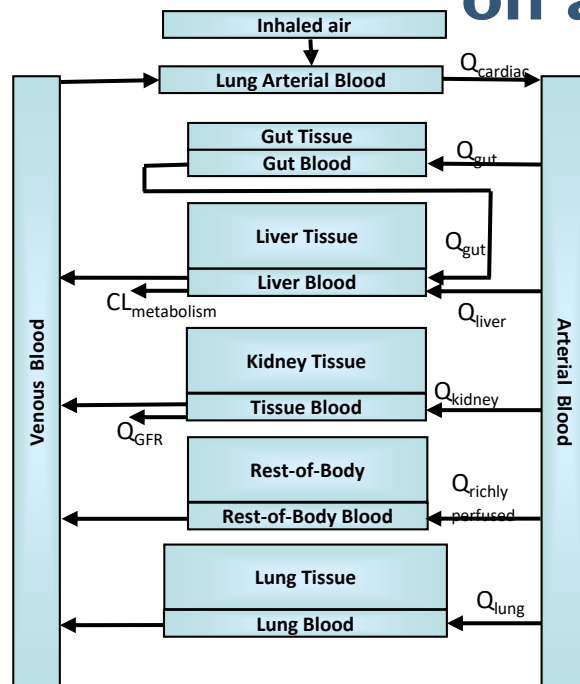
Fig 11. Maternal blood flow to the placenta vs. gestational age. The proportional-to-volume model (solid line) given by Eq 22, the linear transition model given by Eq 21, and two published models [3, 29, 32] are shown.

New HT-PBTK Models: Development is Done with “Branches” on a Git Repository

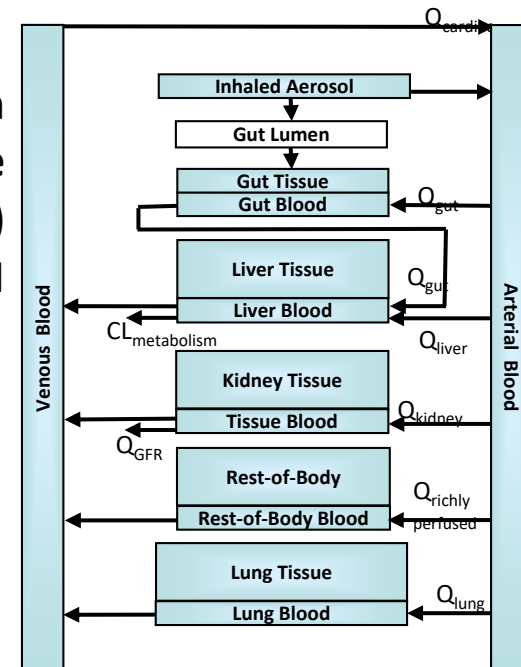
**Gas Inhalation
Exposure Route**
EPA, USAFSAM



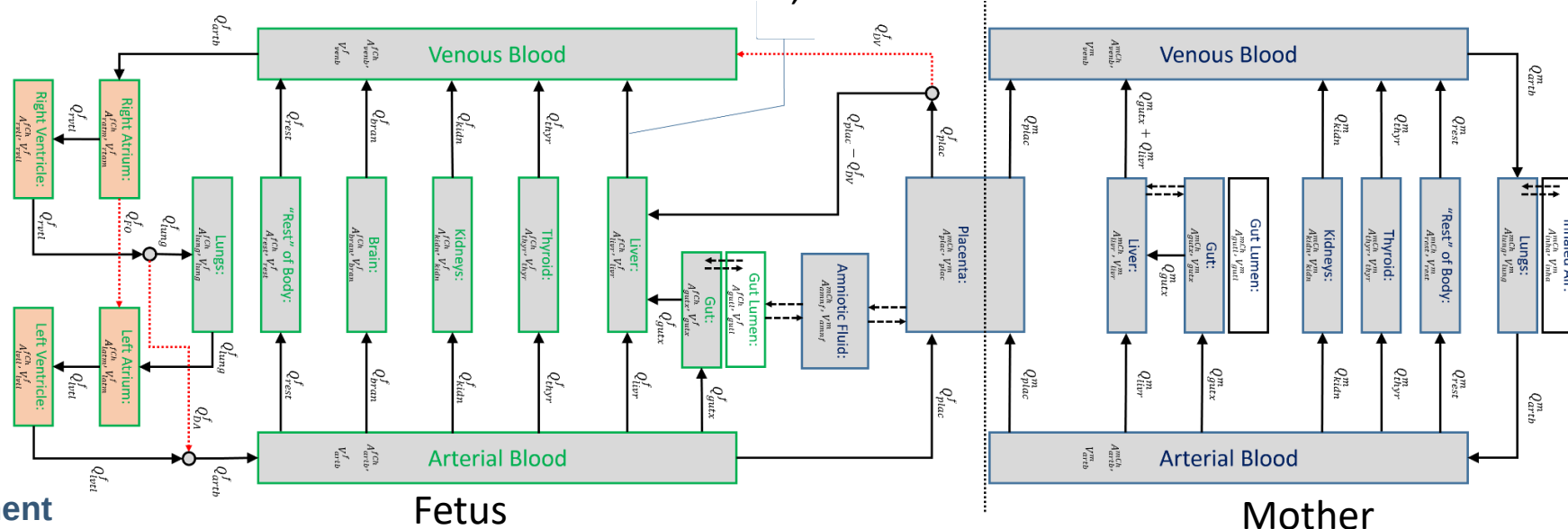
Dermal Exposure Route
EPA, Unilever, INERIS



**Aerosol Inhalation
Exposure Route
(with APEX model)**
EPA, USAFSAM



Human Gestational Model
EPA, FDA

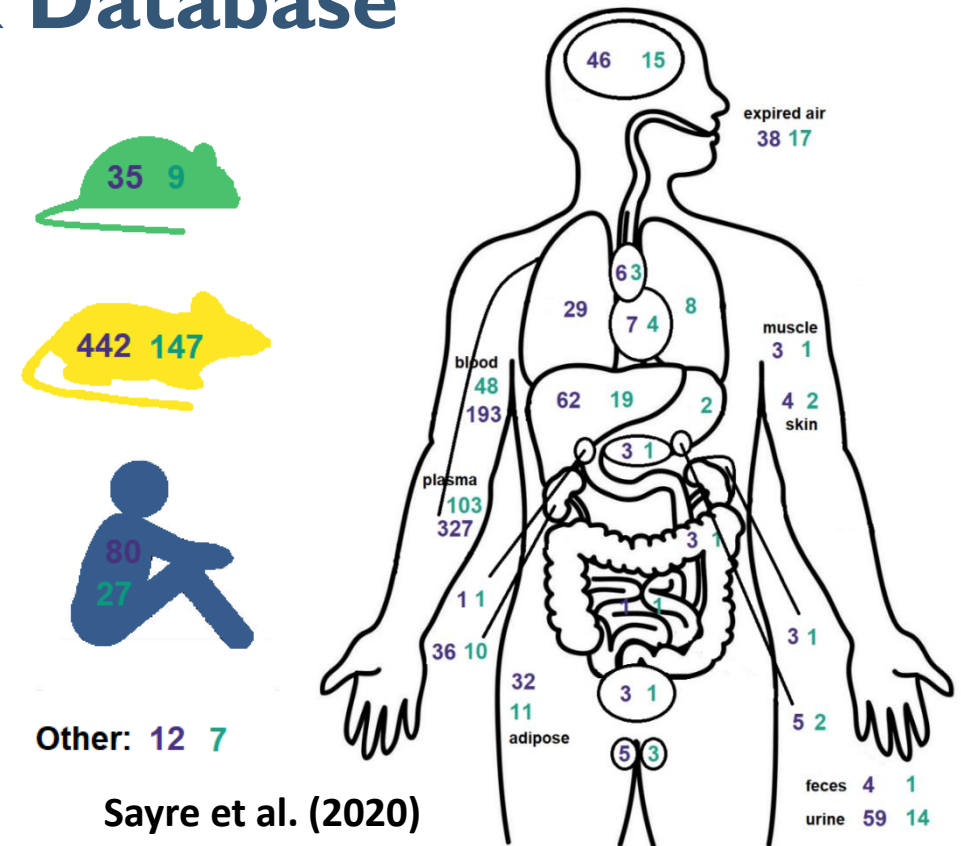


Fetus

Mother

- Database Development - CvTdb: An *In Vivo* TK Database

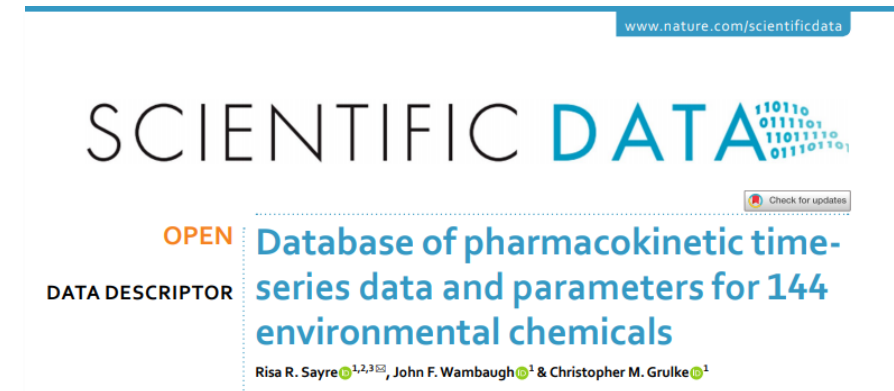
- EPA has developed a **public database of concentration vs. time data** across several species for building, calibrating, and evaluating TK models
- Effort ongoing, but to date includes:
 - 198 analytes (EPA, National Toxicology Program, literature)
 - Routes: Intravenous, dermal, oral, sub-cutaneous, and inhalation exposure
- Standardized, open-source curve fitting software invivoPKfit used to calibrate models to all data



Sayre et al. (2020)

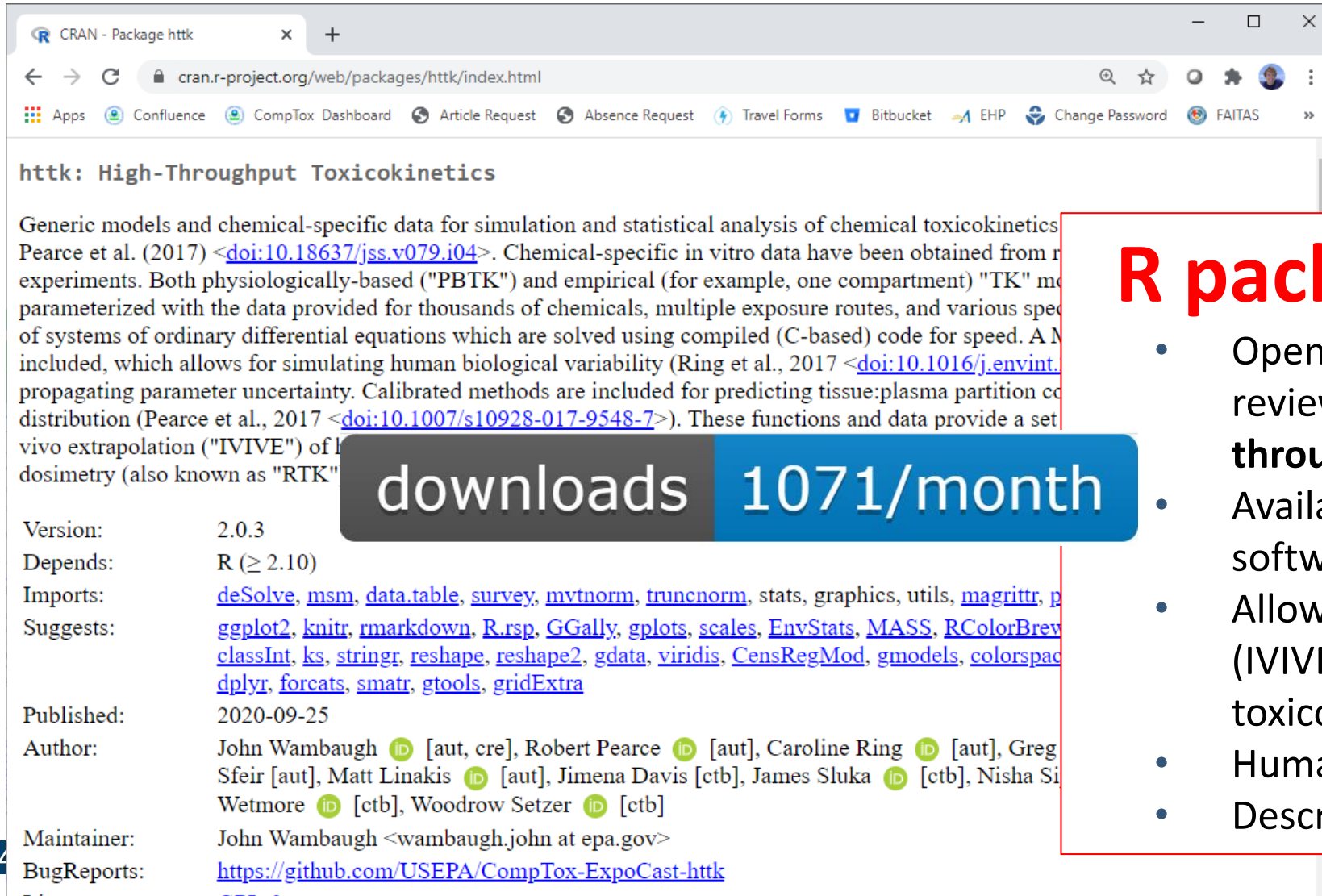
<https://github.com/USEPA/CompTox-ExpoCast-invivoPKfit>

CvTdb Link: <https://github.com/USEPA/CompTox-PK-CvTdb>



- HTTK Platform - Open-Source Tools and Data for HTTK

<https://CRAN.R-project.org/package=httk>



The screenshot shows the CRAN package page for 'httk'. The browser address bar displays 'cran.r-project.org/web/packages/httk/index.html'. The package name 'httk: High-Throughput Toxicokinetics' is prominently displayed. A large blue box indicates 'downloads 1071/month'. The description states: 'Generic models and chemical-specific data for simulation and statistical analysis of chemical toxicokinetics (Pearce et al. (2017) <doi:10.18637/jss.v079.i04>). Chemical-specific in vitro data have been obtained from experiments. Both physiologically-based ("PBTK") and empirical (for example, one compartment) "TK" models are parameterized with the data provided for thousands of chemicals, multiple exposure routes, and various species of systems of ordinary differential equations which are solved using compiled (C-based) code for speed. A Monte Carlo approach is included, which allows for simulating human biological variability (Ring et al., 2017 <doi:10.1016/j.envint.2017.05.012>), propagating parameter uncertainty. Calibrated methods are included for predicting tissue:plasma partition coefficients and distribution (Pearce et al., 2017 <doi:10.1007/s10928-017-9548-7>). These functions and data provide a set of tools for in vivo extrapolation ("IVIVE") of low-dose toxicokinetics (also known as "RTK").'

Version: 2.0.3
Depends: R (≥ 2.10)
Imports: deSolve, msm, data.table, survey, mvtnorm, truncnorm, stats, graphics, utils, magrittr, plotly, ggplot2, knitr, rmarkdown, R.spc, GGally, gplots, scales, EnvStats, MASS, RColorBrewer, classInt, ks, stringr, reshape, reshape2, gdata, viridis, CensRegMod, gmodels, colorspace, dplyr, forcats, smatr, gtools, gridExtra
Published: 2020-09-25
Author: John Wambaugh [aut, cre], Robert Pearce [aut], Caroline Ring [aut], Greg Sfeir [aut], Matt Linakis [aut], Jimena Davis [ctb], James Sluka [ctb], Nisha Siwetmore [ctb], Woodrow Setzer [ctb]
Maintainer: John Wambaugh <wambaugh.john at epa.gov>
BugReports: <https://github.com/USEPA/CompTox-ExpoCast-httk>

R package "httk"

- Open source, transparent, and peer-reviewed tools and data for **high throughput toxicokinetics (httk)**
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific data for 987 chemicals
- Described in Pearce et al. (2017a)

- HHTK Platform -

Modules within R Package “httk”

Feature	Description	Reference
Chemical Specific <i>In Vitro</i> Measurements	Metabolism and protein binding for ~1000 chemicals in human and ~200 in rat	Wetmore et al. (2012, 2013, 2015), plus others
Chemical-Specific <i>In Silico</i> Predictions	Metabolism and protein binding for ~8000 Tox21 chemicals	Sipes et al. (2017)
Generic toxicokinetic models	One compartment, three compartment, physiologically-based oral, intravenous, and inhalation (PBTK)	Pearce et al. (2017a), Linakis et al. (2020)
Tissue partition coefficient predictors	Modified Schmitt (2008) method	Pearce et al. (2017b)
Variability Simulator	Based on NHANES biometrics	Ring et al. (2017)
<i>In Vitro</i> Disposition	Armitage et al. (2014) model	Honda et al. (2019)
Uncertainty Propagation	Model parameters can be described by distributions reflecting uncertainty	Wambaugh et al. (2019)

- *In Vitro* Disposition – A Tox21 Cross Partner Project (EPA, NTP, FDA)

An Experimental Evaluation of Mass Balance Models

describing *in vitro* partitioning and disposition

- Pilot study completed
- 20 chemical case study underway
- Chemical levels quantitated across 5 *in vitro* compartments

Armitage et al. 2014 PMID 25014875

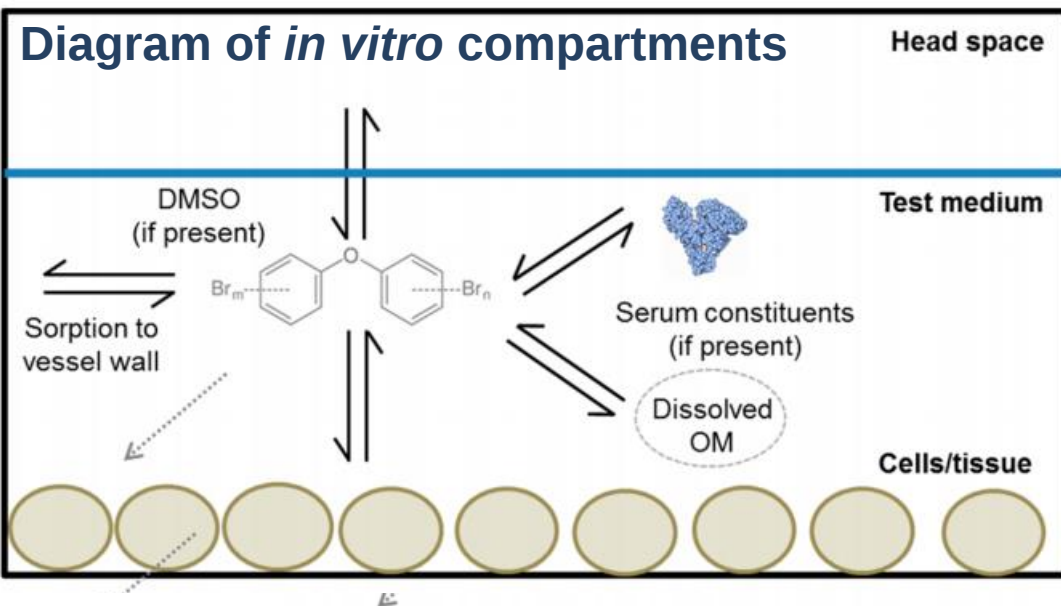
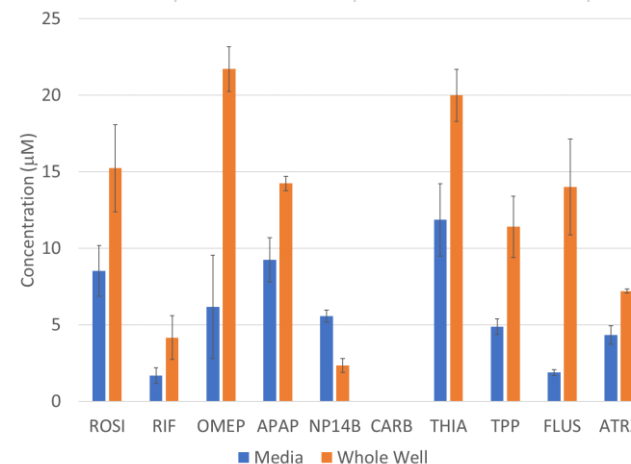


Figure 1. Conceptual representation of an *in vitro* test system.

Preliminary Design and Data

Test Plate	Test Plate Barcode	Plating Condition	Exposure Duration (hr)	Measured Compartment
A	TC00000013	Medium - cells	1	Medium
		Medium - cells	1	Plastic
B	TC00000014	Medium + cells	1	Medium
		Medium + cells	1	Plastic + Cells
C	TC00000015	Medium + cells	1	Whole Well Crash
D	TC00000016	Medium - cells	6	Medium
		Medium - cells	6	Plastic
E	TC00000017	Medium + cells	6	Medium
		Medium + cells	6	Plastic + Cells
F	TC00000018	Medium + cells	6	Whole Well Crash
G	TC00000019	Medium - cells	24	Medium
		Medium - cells	24	Plastic
H	TC00000020	Medium + cells	24	Medium
		Medium + cells	24	Plastic + Cells
I	TC00000021	Medium + cells	24	Whole Well Crash



Providing the Pieces for Prioritization

Informing TSCA

Target Population

Consumers

General Population

ToxCast

ToxCast

HTTK
Oral
Route

Dermal
Route
Needed

HTTK
Oral
Route

Aerosol
Route
Needed

Evaluation Data:
NHANES

Human

ExpoCast/SEEM

Many Exposure
Predictors

Evaluation Data:
NHANES

Human

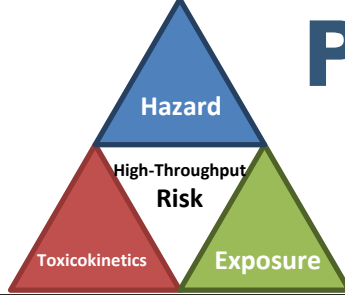
ExpoCast/SEEM

Many Exposure
Predictors

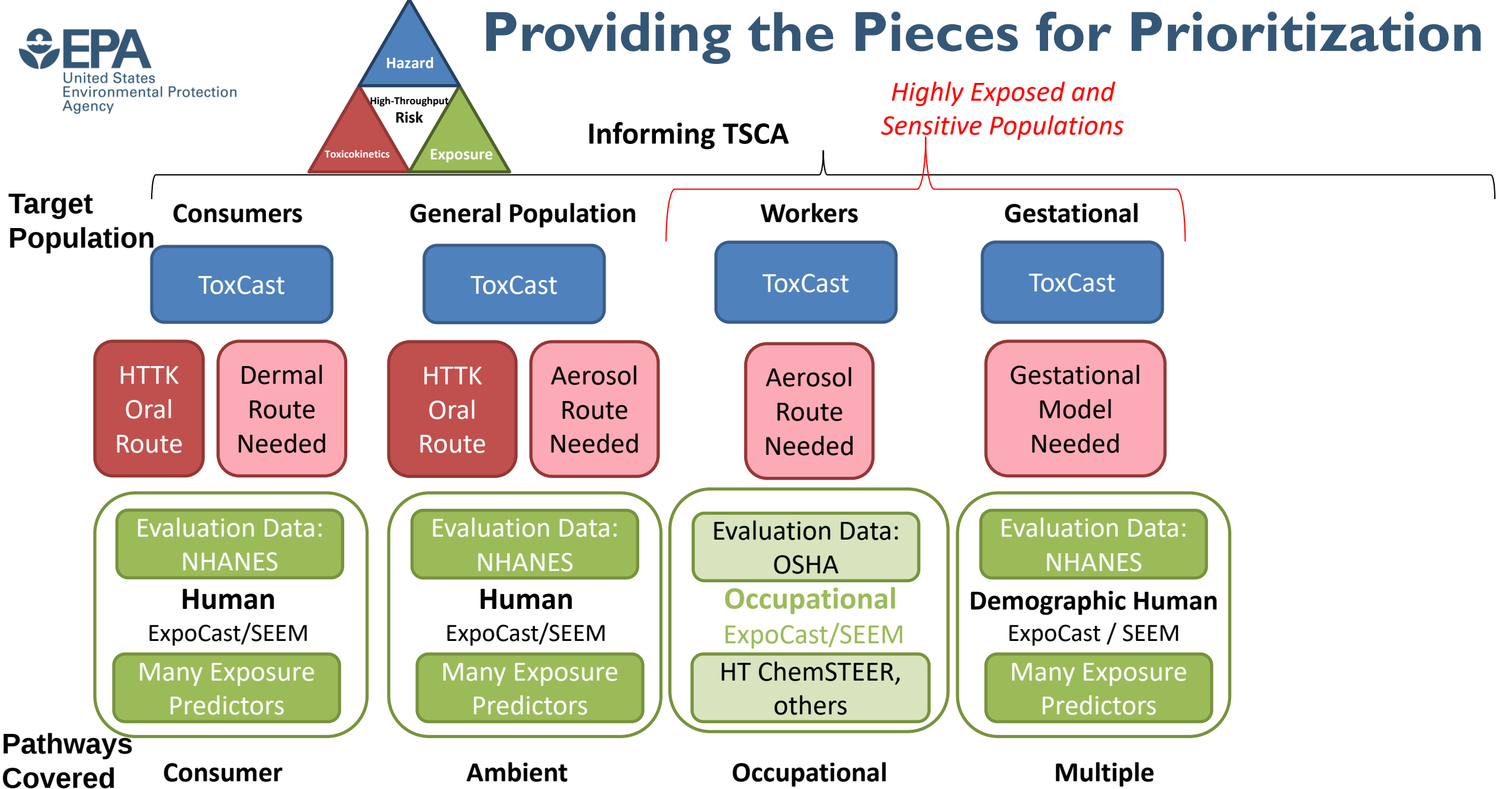
Pathways Covered

Consumer

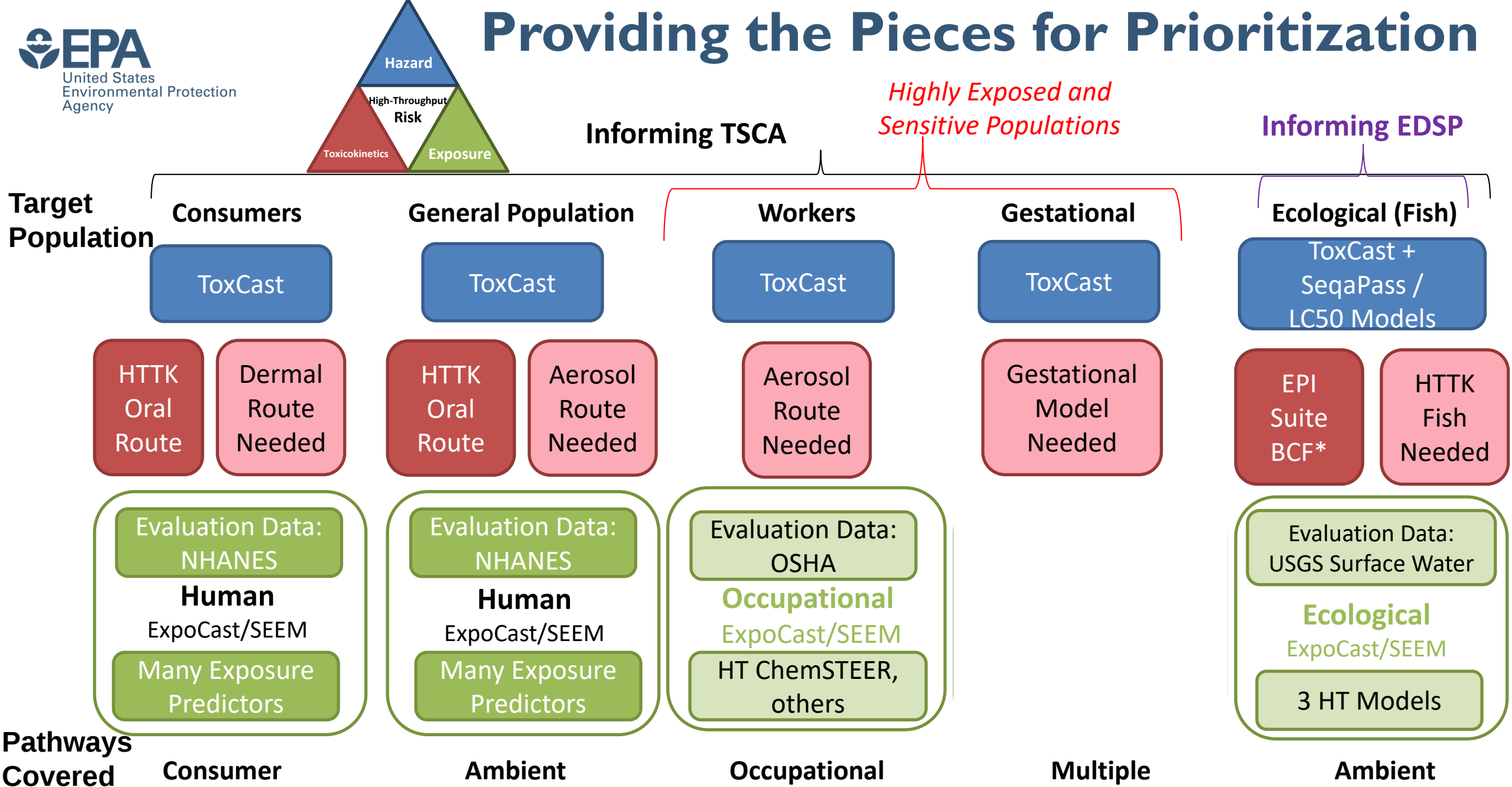
Ambient



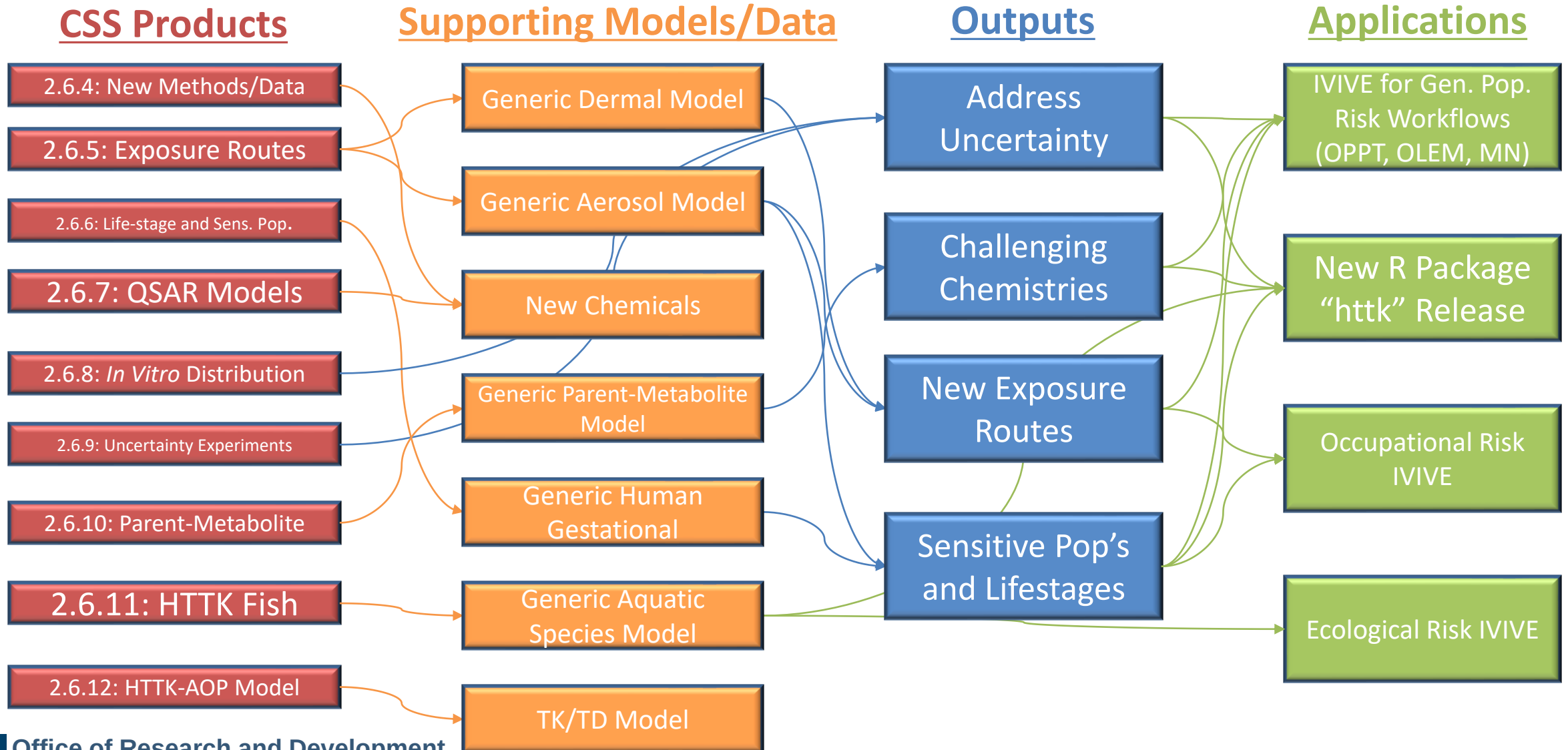
Providing the Pieces for Prioritization



Providing the Pieces for Prioritization



TK and IVIVE Projects and Relationships



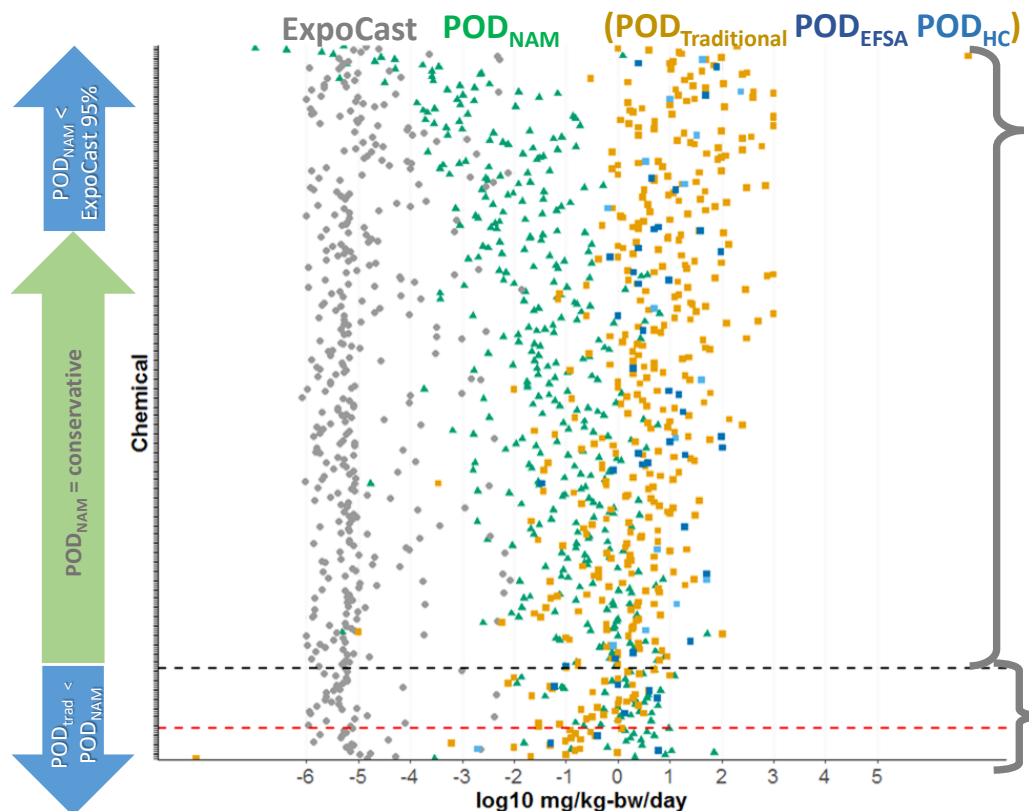
International Collaborations

- Accelerating the Pace of Chemical Risk Assessment (APCRA) -

In Vitro Bioactivity, HTTK, and *In Vivo* Toxic Doses



International case study with EPA, ASTAR, ECHA, Health Canada, and EFSA



For ~89% of the chemicals, POD_{NAM} was conservative. (~100-fold on average), but less conservative than a TTC

Chemicals where POD_{NAM} was not conservative enriched in OPs/carbamates

Additional Efforts and Outreach

Additional Efforts

- **In vitro TK data generation:** Ongoing, internal (>400 TSCA, incl. 150 PFAS) and external (>215); as needed on program office-initiated efforts (Office of Chemical Safety and Pollution Prevention, Office of Water)
- **In vivo TK:** rat *in vivo* studies for comparative assessments and IVIVE evaluation (Hughes *et al.*, underway)
- **Dermal Route:** permeability/partitioning models completed (Evans *et al.*), integration with HTTK begun
- **Bioavailability:** incorporation of Caco-2 data in IVIVE (Honda *et al.*, 2019; Honda *et al.*, in preparation)
- **Transporters:** TK renal transporter data generation for PFAS IVIVE modeling (Smeltz *et al.*, underway)
- **Sensitive Populations/Variability:** Isozyme-specific chemical evaluations to evaluate TK variability and supply *in silico* predictive efforts (Kreutz *et al.*, underway); Correlated Monte Carlo approach to incorporate physiologic variability (Ring *et al.*, 2017)
- **Parent-Metabolite HTTK:** NTA data for metabolism of ToxCast chemicals generated by contractor and being analyzed (Boyce *et al.* underway)

Stakeholder Outreach and Collaborations

- CompTox Chemicals Dashboard: Contains ADME data for >1000 chemicals.
- 2020 SOT: “New Data and Tools for Understanding Chemical Distribution *In Vitro*” - Nynke Kramer and John Wambaugh
- FIFRA SAP “The use of new approach methodologies (NAMs) to derive extrapolation factors and evaluate developmental neurotoxicity for human health risk assessment” - Incorporation of *in vitro* TK / HTTK
- Integration of high throughput hazard, exposure, and TK NAMs into proposed TSCA workflows (white paper, peer review)
- APCRA Collaborations – HTTK case study (underway) and NAM prospective case study (underway)
- Ongoing collaborations with Health Canada, US Geological Survey, and MN Department of Health

References

- Armitage J.M. *et al.* "Application of mass balance models and the chemical activity concept to facilitate the use of *in vitro* toxicity data for risk assessment." *Environmental Science & Technology* (2014) 48, 9770-9779.
- Bell, S.M., *et al.* "In vitro to in vivo extrapolation for high throughput prioritization and decision making." *Toxicology In Vitro* (2018) 47: 213-227.
- Honda, G.S., *et al.* "Using the concordance of *in vitro* and *in vivo* data to evaluate extrapolation assumptions." *PloS One* (2019) 14(5): e0217564.
- Jamei, M., *et al.* "The Simcyp® population-based ADME simulator." *Expert Opinion on Drug Metabolism & Toxicology* (2009) 5(2):211-223.
- Kapraun, D., *et al.* "Empirical Models for anatomical and physiological changes in a human mother and fetus during pregnancy and gestation." *PloS One* (2019). 14(5):e0215906.
- Linakis, M. *et al.* "Development and Evaluation of a high throughput inhalation model for organic chemicals." *Journal of Exposure Science & Environmental Epidemiology* (2020) 30:866-877.
- Lukacova, V. *et al.* "Prediction of modified release pharmacokinetics and pharmacodynamics from in vitro, immediate release, and intravenous data." *The AAPS journal* (2009) 11(2): 323-334.
- Mansouri, K., *et al.* "OPERA models for predicting physicochemical properties and environmental fate endpoints." *Journal of Cheminformatics* (2018). 10(1): 10.
- National Research Council. Risk Assessment in the Federal Government: Managing the Process Working Papers. (1983). National Academies Press.
- National Research Council. Issues in Potable Reuse: The viability of augmenting drinking water supplies with reclaimed water. (1998). National Academies Press.
- Paul-Friedman, K. *et al.* "Utility of *In Vitro* Bioactivity as a Lower Bound Estimate of In Vivo Adverse Effect Levels and in Risk-Based Prioritization." *Toxicological Sciences* (2020) 173(1):202-225.
- Pearce, R.G., *et al.* "httk: R Package for High-Throughput Toxicokinetics." *Journal of Statistical Software*, (2017a) 79(4):1-26.
- Pearce, R.G., *et al.* "Evaluation and calibration of high-throughput predictions of chemical distribution to tissues." *Journal of Pharmacokinetics and Pharmacodynamics* 44.6 (2017b): 549-565.
- Pradeep, P., *et al.* "Using chemical structure information to develop predictive models for *in vitro* toxicokinetic parameters to inform high-throughput risk assessment." *Computational Toxicology* (2020) Epub.
- Ring, C.L., *et al.* "Identifying populations sensitive to environmental chemicals by simulating toxicokinetic variability." *Environment International* 106 (2017): 105-118.
- Rotroff, D.M., *et al.* "Incorporating human dosimetry and exposure into high-throughput *in vitro* toxicity screening." *Toxicological Sciences* (2010) 117(2): 348-358.
- Sayre, R. *et al.*, "Database of pharmacokinetic time-series data and parameters for 144 environmental chemicals", *Scientific Data* (2020) 7(1): 1-10.
- Schmitt, W. "General approach for the calculation of tissue to plasma partition coefficients." *Toxicology in Vitro* (2008) 22(2): 457-467.
- Shibata, Y., *et al.* Prediction of hepatic clearance and availability by cryopreserved human hepatocytes: an application of serum incubation method. *Drug Metabolism and Disposition* (2002), 30(8), 892-896
- Sipes, N.S., *et al.* "An intuitive approach for predicting potential human health risk with the Tox21 10k library." *Environmental Science & Technology* (2017) 51(18): 10786-10796.
- Sobels, F. H. "The parallelogram; an indirect approach for the assessment of genetic risks from chemical mutagens." In: *Progress in Mutation Research*, Vol. 3. 1982. (K. C. Bora, G. R. Douglas, and E. R. Nestman, Eds.), Elsevier, Amsterdam, pp. 233-327.
- Tan, Y.-M., *et al.* "Reverse dosimetry: interpreting trihalomethanes biomonitoring data using physiologically based pharmacokinetic modeling." *Journal of Exposure Science & Environmental Epidemiology* (2007) 17(7):591-603.
- Thomas, Russell S., *et al.* "Incorporating Monte Carlo simulation into physiologically based pharmacokinetic models using advanced continuous simulation language (ACSL): a computational method." *Toxicological Sciences* (1996) 31(1):19-28.
- Wambaugh, J.F., *et al.* "Evaluating *In Vitro-In Vivo* Extrapolation of Toxicokinetics." *Toxicological Sciences* (2018) 163(1): 152-169.
- Wambaugh, J.F., *et al.* "Assessing Toxicokinetic Uncertainty and Variability in Risk Prioritization" *Toxicological Sciences*, 172(2), 235-251.
- Wang, Y.-H. "Confidence assessment of the Simcyp time-based approach and a static mathematical model in predicting clinical drug-drug interactions for mechanism-based CYP3A inhibitors." *Drug Metabolism and Disposition* (2010) 38(7):1094-1104.
- Wetmore, B.A., *et al.* "Integration of dosimetry, exposure and high-throughput screening data in chemical toxicity assessment." *Toxicological Sciences* (2012) 125(1):157-174.
- Wetmore, B.A., *et al.* "Relative impact of incorporating pharmacokinetics on predicting in vivo hazard and mode of action from high-throughput in vitro toxicity assays." *Toxicological Sciences* (2013) 132(2) 327-346.
- Wetmore, B.A., *et al.* "Incorporating high-throughput exposure predictions with dosimetry-adjusted in vitro bioactivity to inform chemical toxicity testing." *Toxicological Sciences* (2015) 148(1): 121-136.

ExpoCast Project (Exposure Forecasting)

Center for Computational Toxicology and Exposure

Linda Adams
Lucas Albrecht*
Matthew Boyce*
Miyuki Breen*
Alex Chao
Daniel Dawson*
Mike Devito
Alex East*
Lindsay Eddy*
Christopher Eklund
Peter Egeghy
Marina Evans
Alex Fisher*
Rocky Goldsmith
Louis Groff*
Chris Grulke

Colin Guider*
Mike Hughes
Victoria Hull*
Kristin Isaacs
Richard Judson
Jen Korol-Bexell*
Anna Kreutz*
Charles Lowe*
Seth Newton
Alli Phillips
Katherine Phillips
Tom Purucker
Ann Richard
Caroline Ring

Risa Sayre
Mark Sfeir*
Marci Smeltz*
Jon Sobus
Zach Stanfield*
Mike Tornero-Velez
Rusty Thomas
Elin Ulrich
Dan Vallero
Barbara Wetmore
John Wambaugh
Antony Williams

CEMM

Hongwan Li*
Xiaoyu Liu
Zachary Robbins*
Mark Strynar

CPHEA

Paul Price

***Trainees**

Collaborators

Arnot Research and Consulting
Jon Arnot
Johnny Westgate
Integrated Laboratory Systems
Kamel Mansouri
Xiaoqing Chang
Shannon Bell
National Toxicology Program
Steve Ferguson
Nisha Sipes
Ramboll
Harvey Clewell
Silent Spring Institute
Robin Dodson
Simulations Plus
Michael Lawless
Southwest Research Institute
Alice Yau
Kristin Favela
Summit Toxicology
Lesla Aylward
Technical University of Denmark
Peter Fantke
Unilever
Beate Nicol
Cecilie Rendal
Ian Sorrell
United States Air Force
Heather Pangburn
Matt Linakis
University of California, Davis
Deborah Bennett
University of Michigan
Olivier Jolliet
University of Texas, Arlington
Hyeong-Moo Shin
University of Nevada
Li Li
University of North Carolina, Chapel Hill
Julia Rager
Marc Serre

NAMs for Exposure: Non-Targeted Analysis

Jon Sobus, Ph.D.

Stakeholder Needs (OCSPP; EPA Regions)

Chemical safety evaluations require an improved understanding of chemical exposure scenarios and pathways

High-priority exposure data needs → consumer products,
indoor environments, occupational settings,
ambient environments, ecological pathways

Challenges

- High-quality exposure data are unavailable for many chemicals
- Measurement data traditionally generated using “targeted” methods
- Targeted analytical methods:
 - Require *a priori* knowledge of chemicals of interest
 - Produce data for few selected analytes (10s-100s)
 - Require standards for method development & compound quantitation
 - Are blind to emerging contaminants
 - Can't keep pace with the needs of 21st century chemical safety evaluations

Research Objective

Rapid Exposure Modeling and Dosimetry Output 2.7:

Develop, evaluate, and apply *non-targeted analysis (NTA)* methods, alongside targeted monitoring methods, to identify critical sources and pathways of human and ecological exposures

Key Question:

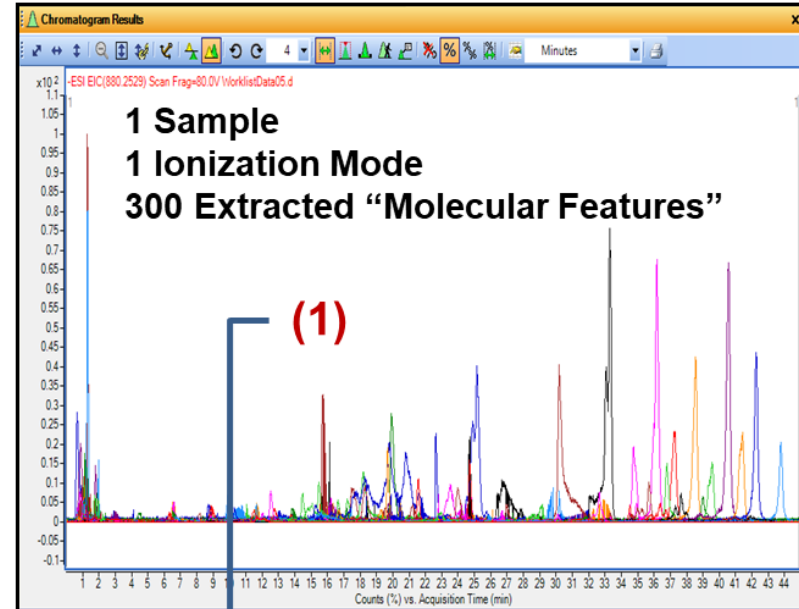
**Are NTA methods suitable to meet the needs of
21st century chemical safety evaluations?**

General NTA Workflow Steps

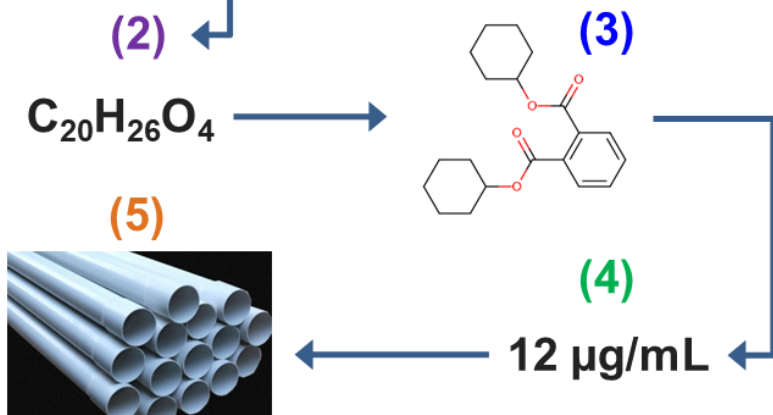
Samples



High-Resolution MS



- 1) Prioritize "molecular features"
- 2) Correctly assign formulas
- 3) Correctly assign structures
- 4) Predict chemical concentrations
- 5) Determine chemical sources

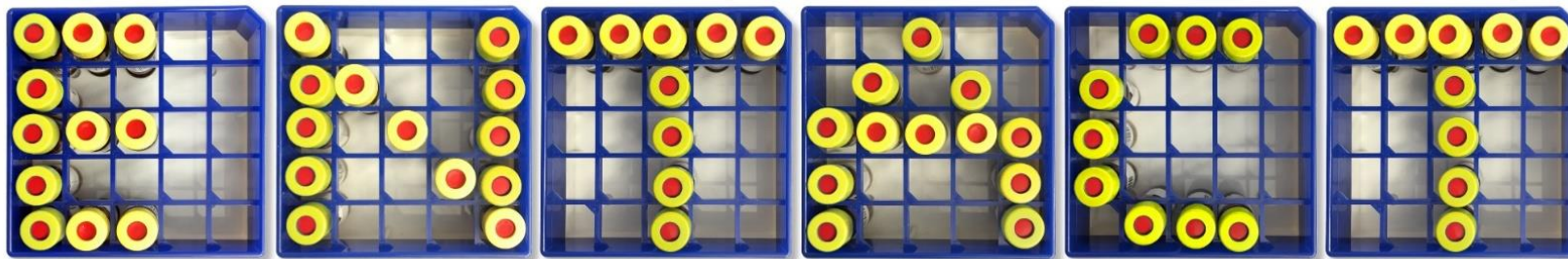
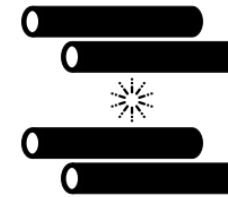
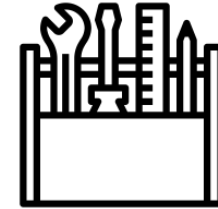


Ongoing Research Activities

- **Evaluate NTA State-of-the-Science**
 - EPA's Non-Targeted Analysis Collaborative Trial (ENTACT)
- **Develop and Disseminate Guidance Materials**
 - Benchmarking and Publications for NTA (BP4NTA)
- **Build Tools to Ensure Transparency & Reproducibility**
 - NTA Study Reporting Tool (NTA SRT)
 - EPA NTA Web Application (NTA WebApp)
- **Address Priority Data Needs with Proof-of-Concept Applications**

Evaluating NTA Science-of-the-Science

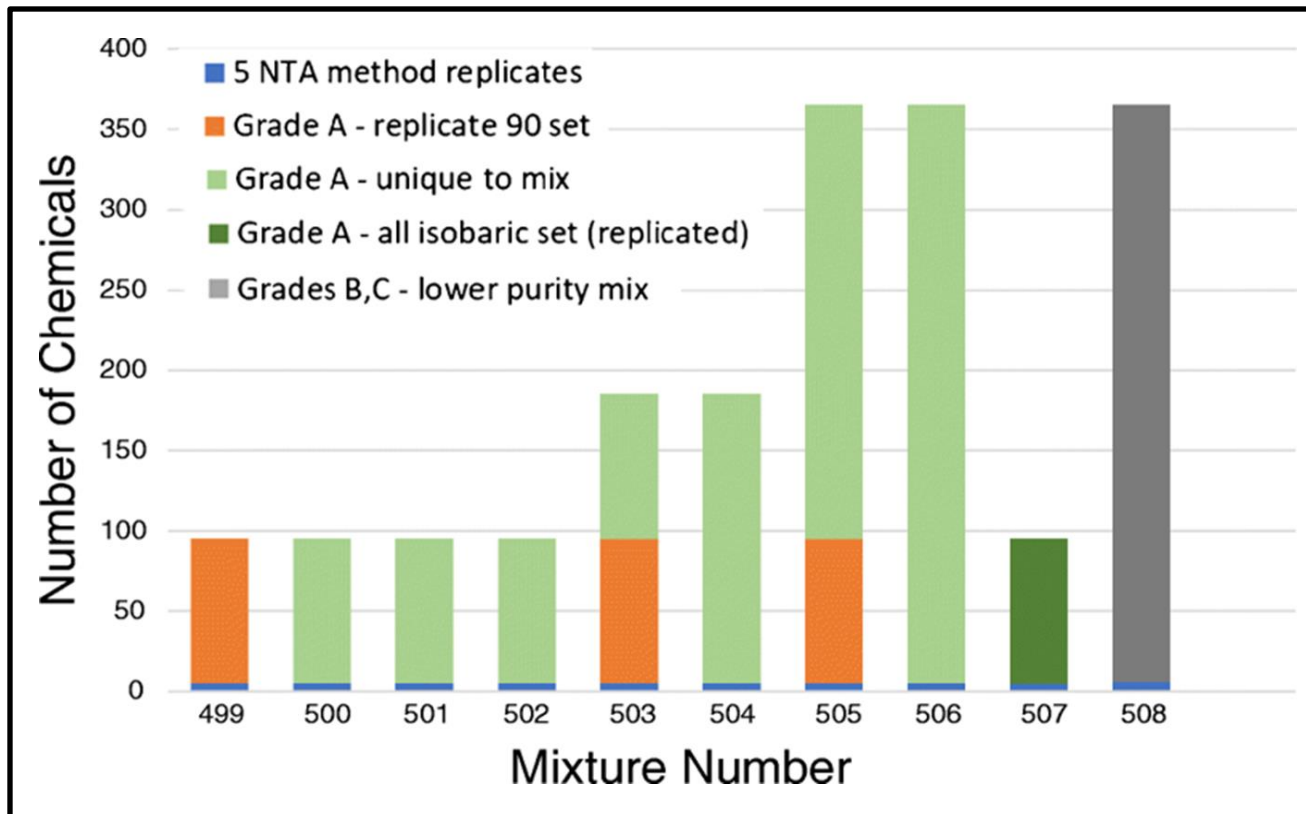
- How variable are tools and results from lab to lab?
- Are some methods/workflows better than others?
- How does sample complexity affect performance?
- What chemical space does a given method cover?
- How sensitive are specific instruments/methods?



EPA's Non-Targeted Analysis Collaborative Trial

ENTACT Study Design (Part I)

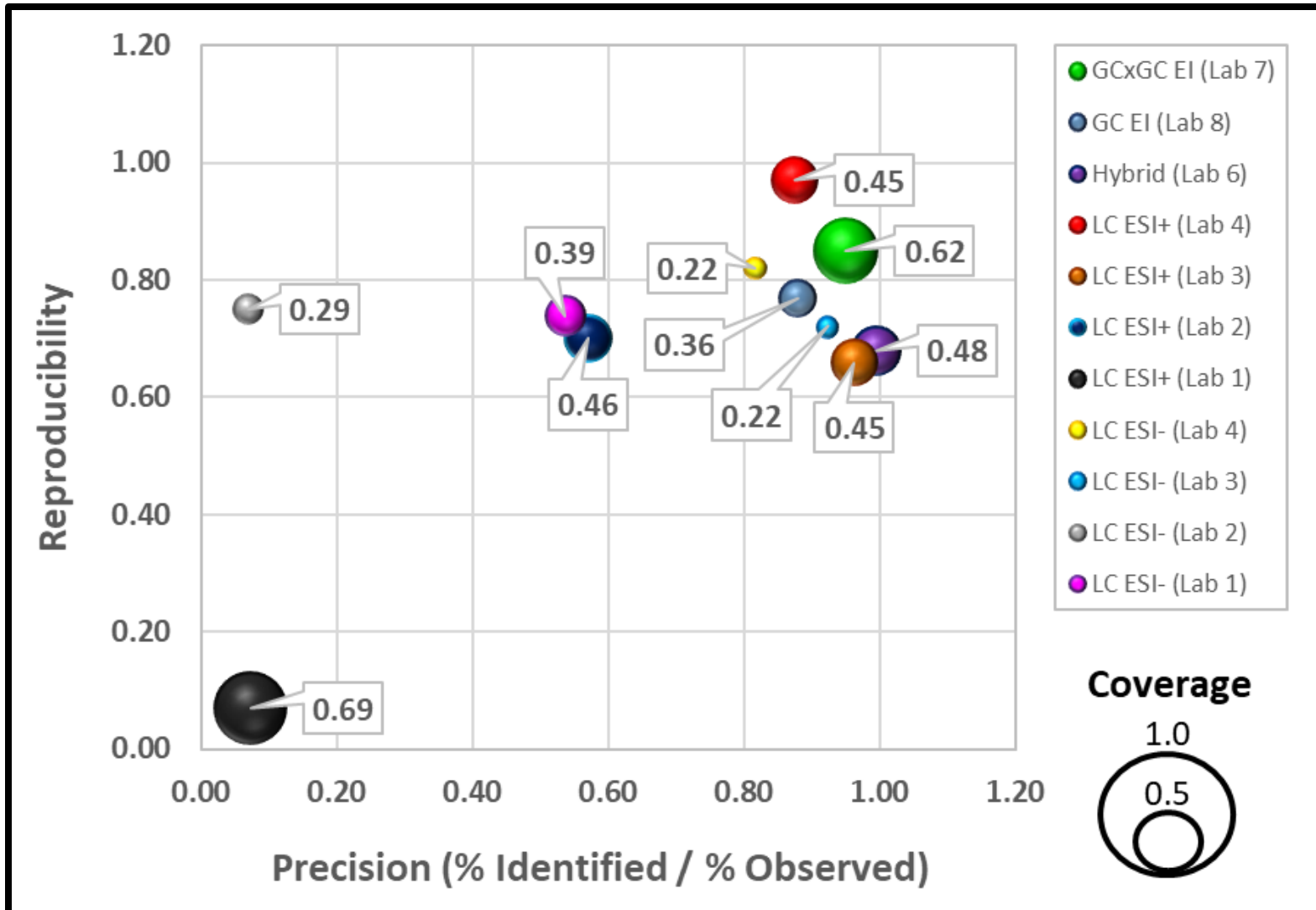
- ~30 global participants, 19 results submitted to date
- 10 synthetic mixtures of ToxCast substances (n=1269)



**Replication in
substance spikes
offers a unique
means to assess
NTA method
reproducibility!**

Ulrich et al. 2019. doi: 10.1007/s00216-018-1435-6

Performance Comparison Across Methods



Metrics:

Bubble Size →
What % observed?
(of those spiked)

X-Axis →
What % correct?
(of those observed)

Y-Axis →
What % consistent?
(of those correct)

Take-Away Messages from ENTACT (to date...)

- Lack of transparency in methods/results reporting
- Method procedures change over short time increments
- Biased self-reporting → highlight strengths, mask weaknesses
- Blinded ToxCast mixtures allow for NTA performance assessment
- Standard performance measures highly variable across labs/methods
- Standard performance assessment methods/benchmarks must be adopted
- Benchmarks require input/consensus from NTA community
- Community focus must be on QA/QC and guidance (and innovation)

Developing and Disseminating Guidance Materials

- BP4NTA → Borne out of 2018 ENTACT workshop
- ~100 U.S. and international members
 - Government, academia, and industry
- Working Group Objectives:
 - Short term → define common NTA terms, concepts, and performance metrics
 - Short term → provide recommendations on research & reporting best practices
 - Long term → establish proficiency testing levels (ASTM/ISO)
- Products (including 3 manuscripts):
 - Website with key resources and links: <https://nontargetedanalysis.org/>
 - Guidance documents with definitions & supporting info
 - “NTA Study Reporting Tool” to standardize reporting (proposals & manuscripts)



Building Tools to Ensure Transparency & Reproducibility

The “NTA Study Reporting Tool” (NTA SRT):

- Standardized framework for reviewing quality of NTA reporting
- Aids NTA study design and review (proposals & manuscripts)
- Follows chronology of typical NTA studies with detailed examples
- Scale-based scoring (numeric & colorimetric) for individual study attributes
- HTML interactive version via BP4NTA website (hyperlinks → supporting docs.)
- Fillable PDF version available for download (via website)
- Comment box for periodic updates/revisions (via website)
- Working with journal editors for initial testing and deployment

NTA Study Reporting Tool (draft version)

NTA Study Chronology

Study Sections & Categories			Example Information to Report	Numeric & Colorimetric Scoring	Rationale/Notes
Methods	Study Design	Objectives & Scope	- Study goals and hypotheses - Scope of the study with respect to use of NTA / suspect screening - Expected chemical coverage of approach and potential limitations	1	
		Sample Information & Preparation	- Sample collection/replication, handling/storage, preparation, extraction, & clean-up methods (and related QA practices) - Intended use of samples (e.g., method development, compound identification, etc.) - Development and intended use of blanks	2	
		QC Spikes & Controls	- Development of spikes/controls (e.g., isotopically labeled standards/spikes, native standard spikes, matrix pools) - Intended use of QC or other spikes/controls (e.g., to monitor instrument performance, data normalization, etc.)	2	
	Data Acquisition	Analytical Sequence	- Sample randomization and use of replicate injections - Inclusion of blanks and QC samples in the acquisition sequence - Information about single vs. multiple analytical batches	3	
		Chromatography	- Instrument specifications - Method settings (e.g., column/guard, mobile phases, gradient, injection techniques)	3	
		Mass Spectrometry	- Instrument specifications - Instrument calibration and/or tuning procedures - Method settings (e.g., ...)	3	
		Software	- File conversion information - Software program(s) used - Workflow steps (e.g., ...) - Feature detection thresholds - Data correction or not - Software programs(s) used	2	
	Data Outputs	Statistical & Chemometric Outputs	- Basic statistical outputs (e.g., adj. p-values, standard deviations, test statistics) - Results of chemometric analyses (e.g., reported classifications/groupings of features or samples, observed trends in the data) - Visuals/plots (e.g., Venn diagrams, heatmaps, clustering dendrograms, volcano plots, network diagrams, PCA and loading plots) - New statistical metrics, algorithms, packages, and/or scripts	NA	
		Identification & Confidence Levels	- Reported identifications and associated confidence levels (e.g., levels described by Schymanski et al.) - Supporting data for annotation/identification (e.g., formula match scores, fine isotope pattern, retention time match, MS/MS match scores, source of MS/MS spectra) - For features with lower confidence IDs, (i.e., not standard-confirmed), proposed tentative structures and other annotated data - Semi-quantification or quantification data - Exported MS/MS spectra (e.g., as a library, database, or deposition into online repository)	3	
		Data Acquisition QA/QC	- Quality: Adherence to QA/QC protocols for sample preparation and data acquisition - Boundary: Description of the potential impacts of methods (sample prep, chromatographic, MS) on observable chemical space - Accuracy: Reported chromatographic and mass accuracy	1	
Results	QA/QC Metrics	Data Processing & Analysis QA/QC	- Precision: Variability of observed retention time, precursor mass error, and abundance - Quality: Outcomes of QC checks along the data processing & analysis workflow - Boundary: Impact of data processing & analysis method(s) on observed chemical space, observed limits of detection/ID - Accuracy: Performance measures (True Positive Rate, False Positive Rate, etc.) for known compounds or samples with known classification - Precision: Reproducibility/repeatability of performance measures for known compounds or samples with known classification; Calculations such as False Discovery Rate, F1 score, etc.	0	

Hyperlinked
(HTML version)
to supporting
information

3-4 bullet point examples for each of the 13 sub-categories

Not exhaustive – intended to guide reviewers; relies on reviewer expertise/discretion.

Space for reviewer to explain assigned score

Building Tools to Ensure Transparency & Reproducibility

The EPA NTA WebApp:

- Queries NTA data against DSSTox DB (~900K substances)
- Aggregates metadata to aid candidate prioritization
- Calculates match metrics to aid candidate filtering
- Provides interactive visualization of chemical candidates
- Processes data for advanced statistical analyses
- Standardizes and documents procedures for NTA data analysis
- Adheres to recommendations from BP4NTA workgroup
- Produces publication-ready output in accordance with NTA SRT

EPA's NTA WebApp

EPA United States Environmental Protection Agency

Environmental Topics Laws & Regulations About EPA Search EPA.gov

[Contact Us](#)

NTA: non-targeted analysis of MS data (beta)

Tools

- [MS1 Tool](#)
 - [Run MS1 Tool](#)
 - [MS1 Tool Algorithms](#)
 - [MS1 Tool QA/QC](#)
 - [MS1 Tool References](#)
 - [MS2 CFMID Tool](#)

Documentation

- [Source Code](#)

Run NTA MS1 Tool

Input	Value
Project name:	Example nta
Positive MPP file (csv):	Choose File No file chosen
Negative MPP file (csv):	Choose File No file chosen
Adduct mass accuracy units:	ppm
Adduct mass accuracy:	10
Adduct retention time accuracy (mins):	0.05
Tracer file (csv; optional):	Choose File No file chosen
Tracer mass accuracy units:	ppm
Tracer mass accuracy:	5
Tracer retention time accuracy (mins):	0.1
Min sample:blank cutoff:	3
Min replicate hits:	<input type="range" value="2"/>
Max replicate CV:	0.8
Parent ion mass accuracy (ppm):	<input type="range" value="5"/>
Discard features below this retention time (mins):	0.0
Search dashboard by:	mass
Save top result only?	no
DSSTox search batch size (debugging):	150

WebApp Output:

- QA/QC tracer results
- Cleaned, unannotated file for stats analysis
- Cleaned, annotated file with DSSTox chemicals
- Complete file with all chemicals & metadata

WebApp Input:

- Experimental data files
- Tracer (QA/QC) files
- Parameters for data cleaning
- Parameters for DB searching

EPA United States Environmental Protection Agency

Environmental Topics Laws & Regulations About EPA Search EPA.gov

[Contact Us](#)

NTA: non-targeted analysis of MS data (beta)

Tools

- [MS1 Tool](#)
 - [Run MS1 Tool](#)
 - [MS1 Tool Algorithms](#)
 - [MS1 Tool QA/QC](#)
 - [MS1 Tool References](#)
 - [MS2 CFMID Tool](#)

Documentation

- [Source Code](#)

NTA Output

Job ID: XIN4V113

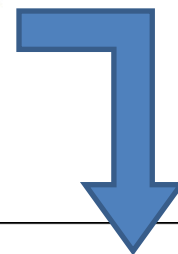
Download results:

Addressing High Priority Data Needs with Proof-of-Concept Applications

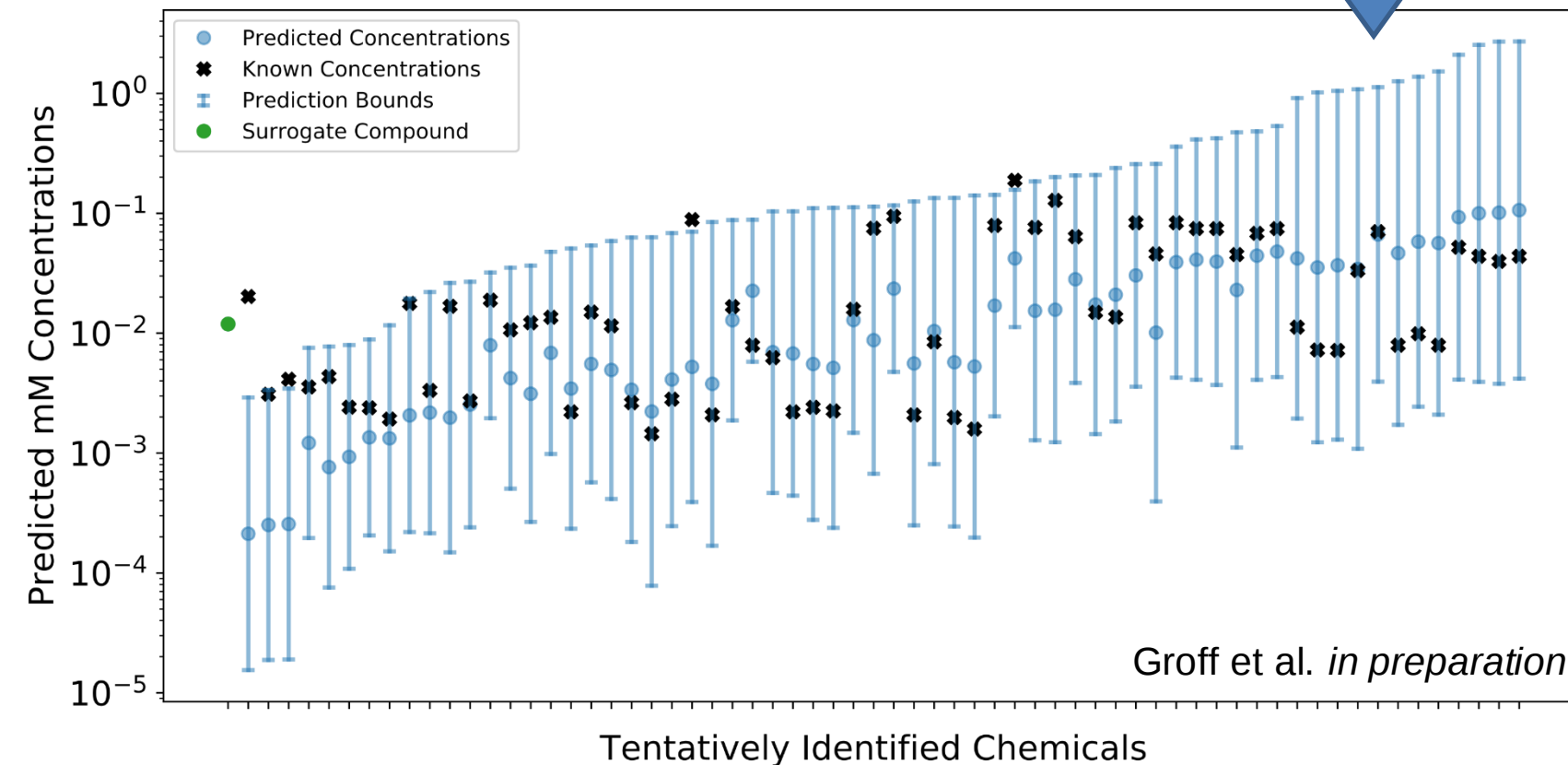
- Characterizing chemical contents of products (including UVCBs)
- Characterizing data-poor xenobiotics in biological tissues & fluids
- Identifying xenobiotic metabolites produced from *in vitro* assays
- Developing semi-quantitative (SQ) methods for risk-based interpretation
- Characterizing emerging contaminants in Brita filters (SQ proof-of-concept)
- Developing a framework for rapid response NTA



SQ NTA Proof-of-Concept



- Analysis of Brita filter extracts via GC-HRMS.
- Concentration estimates can be above or below true value.
- Prediction intervals used to bound SQ concentration estimates.
- 95% prediction intervals shown; Can use 99%, 99.9%, etc.
- Tentatively identified compounds ranked by upper bound estimates.
- Upper bound estimates compared to level-of-interest to set priorities.
- Priority compounds further examined using targeted methods.



Contributing Researchers

(EPA Affiliation Unless Otherwise Noted)

- **ENTACT:**

- ***Co-leads:*** E. Ulrich and J. Sobus
- ***Research Team:*** A. Williams, A. Chao, S. Newton, C. Lowe, C. Grulke, A. Richard, J. Grossman (ORISE)

- **BP4NTA:**

- ***Overall Co-leads:*** E. Ulrich and B. Place (NIST)
- ***Website Co-leads:*** S. Newton and S. Nason (CAES)

- **NTA SRT:**

- ***Co-leads:*** K. Peter (NIST) and A. Phillips
- ***Research Team:*** P. Gardinali (FIU), A. Knolhoff (FDA), C. Manzano (SDSU), K. Miller, M. Pristner & B. Warth (U. of Vienna), L. Sabourin & M. Sumarah (Agri-Food Canada), J. Sobus

- **NTA WebApp:**

- ***Research Team:*** J. Minucci, A. Chao, T. Purucker, A. Williams, J. McCord, H. Al-Ghoul (ORISE), M. Russell, C. Lowe, L. Groff (ORISE), J. Sobus

- **SQ NTA:**

- ***Research Team:*** L. Groff (ORISE), H. Liberatore, J. McCord, S. Newton, E. Ulrich, J. Sobus

Additional EPA Contributors



This work was supported, in part, by ORD's Pathfinder Innovation Program (PIP) and an ORD EMVL award



EPA ORD

Hussein Al-Ghoul*
Alex Chao
Jacqueline Bangma*
Matthew Boyce*
Kathie Dionisio
Louis Groff*
Jarod Grossman*
Chris Grulke
Kristin Isaacs
Sarah Laughlin*
Hannah Liberatore
Charles Lowe
Kamel Mansouri*
Aurelie Marcotte*
James McCord
Andrew McEachran*
Kelsey Miller
Jeff Minucci

EPA ORD (cont.)

Seth Newton
Grace Patlewicz
Allison Phillips
Katherine Phillips
Tom Purucker
Ann Richard
Randolph Singh*
Jon Sobus
Mark Strynar
Elin Ulrich
Ariel Wallace
John Wambaugh
Antony Williams

GDIT

Ilya Balabin
Tom Transue
Tommy Cathey

* = ORISE/ORAU

US EPA CSS-HERABOSC Meeting – February 2-5, 2021

Relevant EPA NTA Publications

- 1) Rager JE, Strynar MJ, Liang S, McMahan RL, Richard AM, Grulke CM, Wambaugh JF, Isaacs KK, Judson R, Williams AJ, Sobus JR. Linking high resolution mass spectrometry data with exposure and toxicity forecasts to advance high-throughput environmental monitoring. *Environ Int.* 2016 Mar;88:269-280.
- 2) McEachran AD, Sobus JR, Williams AJ. Identifying known unknowns using the US EPA's CompTox Chemistry Dashboard. *Anal Bioanal Chem.* 2017 Mar;409(7):1729-1735.
- 3) Newton SR, McMahan RL, Sobus JR, Mansouri K, Williams AJ, McEachran AD, Strynar MJ. Suspect screening and non-targeted analysis of drinking water using point-of-use filters. *Environ Pollut.* 2018 Mar;234:297-306.
- 4) Sobus JR, Wambaugh JF, Isaacs KK, Williams AJ, McEachran AD, Richard AM, Grulke CM, Ulrich EM, Rager JE, Strynar MJ, Newton SR. Integrating tools for non-targeted analysis research and chemical safety evaluations at the US EPA. *J Expo Sci Environ Epidemiol.* 2018 Sep;28(5):411-426.
- 5) Phillips KA, Yau A, Favela KA, Isaacs KK, McEachran A, Grulke C, Richard AM, Williams AJ, Sobus JR, Thomas RS, Wambaugh JF. Suspect screening analysis of chemicals in consumer products. *Environ Sci Technol.* 2018 Mar 6;52(5):3125-3135.
- 6) McEachran AD, Mansouri K, Newton SR, Beverly BEJ, Sobus JR, Williams AJ. A comparison of three liquid chromatography (LC) retention time prediction models. *Talanta.* 2018 May 15;182:371-379.
- 7) Ulrich EM, Sobus JR, Grulke CM, Richard AM, Newton SR, Strynar MJ, Mansouri K, Williams AJ. EPA's non-targeted analysis collaborative trial (ENTACT): genesis, design, and initial findings. *Anal Bioanal Chem.* 2019 Feb;411(4):853-866.
- 8) Sobus JR, Grossman JN, Chao A, Singh R, Williams AJ, Grulke CM, Richard AM, Newton SR, McEachran AD, Ulrich EM. Using prepared mixtures of ToxCast chemicals to evaluate non-targeted analysis (NTA) method performance. *Anal Bioanal Chem.* 2019 Feb;411(4):835-851.
- 9) Catron TR, Swank A, Wehmas LC, Phelps D, Keely SP, Brinkman NE, McCord J, Singh R, Sobus J, Wood CE, Strynar M, Wheaton E, Tal T. Microbiota alter metabolism and mediate neurodevelopmental toxicity of 17 β -estradiol. *Sci Rep.* 2019 May 8;9(1):7064.
- 10) McEachran AD, Balabin I, Cathey T, Transue TR, Al-Ghoul H, Grulke C, Sobus JR, Williams AJ. Linking in silico MS/MS spectra with chemistry data to improve identification of unknowns. *Sci Data.* 2019 Aug 2;6(1):141.
- 11) Nuñez JR, Colby SM, Thomas DG, Tfaily MM, Tolic N, Ulrich EM, Sobus JR, Metz TO, Teeguarden JG, Renslow RS. Evaluation of in silico multifeature libraries for providing evidence for the presence of small molecules in synthetic blinded samples. *J Chem Inf Model.* 2019 Sep 23;59(9):4052-4060.

Publications (cont.)

- 12) Weitekamp CA, Phelps D, Swank A, McCord J, Sobus JR, Catron T, Keely S, Brinkman N, Zurlinden T, Wheaton E, Strynar M, McQueen C, Wood CE, Tal T. Triclosan-selected host-associated microbiota perform xenobiotic biotransformations in larval zebrafish. *Toxicol Sci.* 2019 Sep 5;172(1):109–122.
- 13) Pleil JD, Wallace MAG, McCord J, Madden MC, Sobus J, Ferguson G. How do cancer-sniffing dogs sort biological samples? Exploring case-control samples with non-targeted LC-Orbitrap, GC-MS, and immunochemistry methods. *J Breath Res.* 2019 Nov 19;14(1):016006.
- 14) Chao A, Al-Ghoul H, McEachran AD, Balabin I, Transue T, Cathey T, Grossman JN, Singh RR, Ulrich EM, Williams AJ, Sobus JR. In silico MS/MS spectra for identifying unknowns: a critical examination using CFM-ID algorithms and ENTACT mixture samples. *Anal Bioanal Chem.* 2020 Feb;412(6):1303-1315.
- 15) Newton SR, Sobus JR, Ulrich EM, Singh RR, Chao A, McCord J, Laughlin-Toth S, Strynar M. Examining NTA performance and potential using fortified and reference house dust as part of EPA's Non-Targeted Analysis Collaborative Trial (ENTACT). *Anal Bioanal Chem.* 2020 Jul;412(18):4221-4233.
- 16) Singh RR, Chao A, Phillips KA, Xia XR, Shea D, Sobus JR, Schymanski EL, Ulrich EM. Expanded coverage of non-targeted LC-HRMS using atmospheric pressure chemical ionization: A case study with ENTACT mixtures. *Anal Bioanal Chem.* 2020 Aug;412(20):4931-4939.
- 17) McEachran AD, Chao A, Al-Ghoul H, Lowe C, Grulke C, Sobus JR, Williams AJ. Revisiting five years of CASMI contests with EPA identification tools. *Metabolites.* 2020 Jun 23;10(6):260.
- 18) Abrahamsson DP, Sobus JR, Ulrich EM, Isaacs K, Moschet C, Young TM, Bennett DH, Tulve NS. A quest to identify suitable organic tracers for estimating children's dust ingestion rates. *J Expo Sci Environ Epidemiol.* 2020 Jul 13.
- 19) Washington JW, Rosal CG, Ulrich EM, Jenkins TM. Use of carbon isotopic ratios in nontargeted analysis to screen for anthropogenic compounds in complex environmental matrices. *J Chromatogr A.* 2019 Jan 4;1583:73-79.
- 20) McEachran AD, Hedgespeth ML, Newton SR, McMahan R, Strynar M, Shea D, Nichols EG. Comparison of emerging contaminants in receiving waters downstream of a conventional wastewater treatment plant and a forest-water reuse system. *Environ Sci Pollut Res Int.* 2018 May;25(13):12451-12463.
- 21) Williams AJ, Sobus JR. Applications of the US EPA CompTox Chemicals Dashboard to support mass spectrometry and breath research. *Breathborne Biomarkers and the Human Volatilome.* 2019 [in press]; C. Davis, J. Beauchamp, and J. Pleil (ed), Elsevier Inc.
- 22) McEachran AD, Mansouri K, Grulke C, Schymanski EL, Ruttkies C, Williams AJ. "MS-Ready" structures for non-targeted high-resolution mass spectrometry screening studies. *J Cheminform.* 2018 Aug 30;10(1):45.

Questions?

sobus.jon@epa.gov



The views expressed in this presentation are those of the author and do not necessarily represent the views or policies of the U.S. Environmental Protection Agency.