April 13, 2021

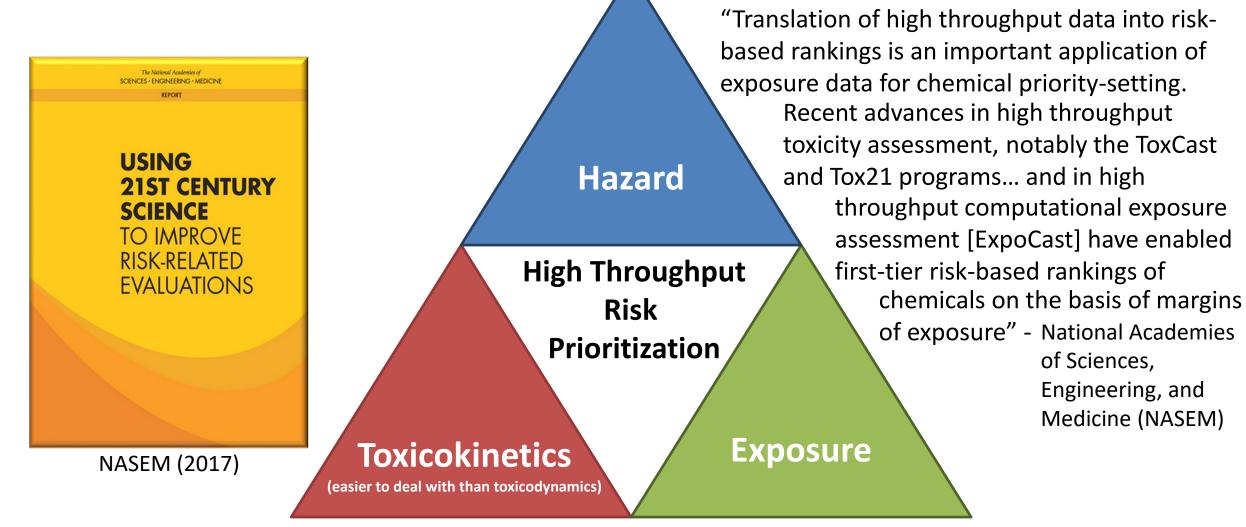
United States Environmental Protection Agency

# NAMS 101 New Approach Methodologies for Exposure

John Wambaugh Center for Computational Toxicology and Exposure Office of Research and Development U.S. Environmental Protection Agency wambaugh.john@epa.gov https://orcid.org/0000-0002-4024-534X

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





In order to perform risk-based ranking we need data on hazard, toxicokinetics, and exposure...

**Calculating Chemical Risk** 



dose-response relationship, and exposure to characterize

potential risks to public health and the environment. A chemical

with minimal toxicity might pose a risk if exposures are exten-

sive, repeated, and/or occurring during critical windows across

the human life span. Exposure assessment involves under-

standing human activity, and this activity is confounded by interindividual variability that is both biological and behavioral.

Exposures further vary between the general population and

susceptible or occupationally exposed populations. Recent

computational exposure efforts have tackled these problems

through the creation of new tools and predictive models. These

tools include machine learning to draw inferences from existing

data and computer-enhanced screening analyses to generate

new data. Mathematical models provide frameworks describing

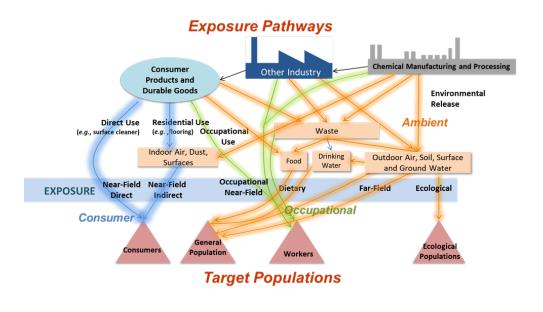
 The tools to characterize both toxicity and exposure have evolved significantly in the past decade

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**Environmental Protection** 

 NAMs for exposure science are being developed to enable risk assessors to more rapidly address public health challenges and chemical regulation





<sup>10</sup> National Health and Environmental Effects Research Laboratory, Office of Research and Development, United States Environmental Protection Agency, Research Triangle Park, NC 27711, USA

Corresponding author: Wambaugh, John F. (Wambaugh.john@epa.gov)

Current Opinion in Toxicology 2019, 15:76–92 This review comes from a themed issue on Risk Assessment in Toxicology Edited by Anne Marie Vinggaard and Richard Judson Available online 31 July 2019 For a complete overview see the Issue and the Editorial https://doi.org/10.1016/j.cotox.2019.07.001

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Exposure NAM Class	Description	Traditional Approach	Measurem	Toxicokineti	Models	Descriptors	Evaluation	Machine Lear
Measurements	New techniques including screening analyses capable of detecting hundreds of chemicals present in a sample	Targeted (chemical-specific) analyses	-	•	•	•		•
Toxicokinetics	High throughput methods using <i>in vitro</i> data to generate chemical-specific models	Analyses based on in vivo animal studies	•	-		•		•
HTE Models	Models capable of making predictions for thousands of chemicals	Models requiring detailed, chemical- and scenario-specific information	•	•	-	•		
Chemical Descriptors	Informatic approaches for organizing chemical information in a machine-readable format	Tools targeted at single chemical analyses by humans				-		•
Evaluation	Statistical approaches that use the data from many chemicals to estimate the uncertainty in a prediction for a new chemical	Comparison of model predictions to data on a per chemical basis	•	•	•	•	-	•
Machine Learning	Computer algorithms to identify patterns	Manual Inspection of the Data	•	•		•		-
Prioritization	Integration of exposure and other NAMs to identify chemicals for follow-up study	Expert decision making	•	•	•	•	•	•



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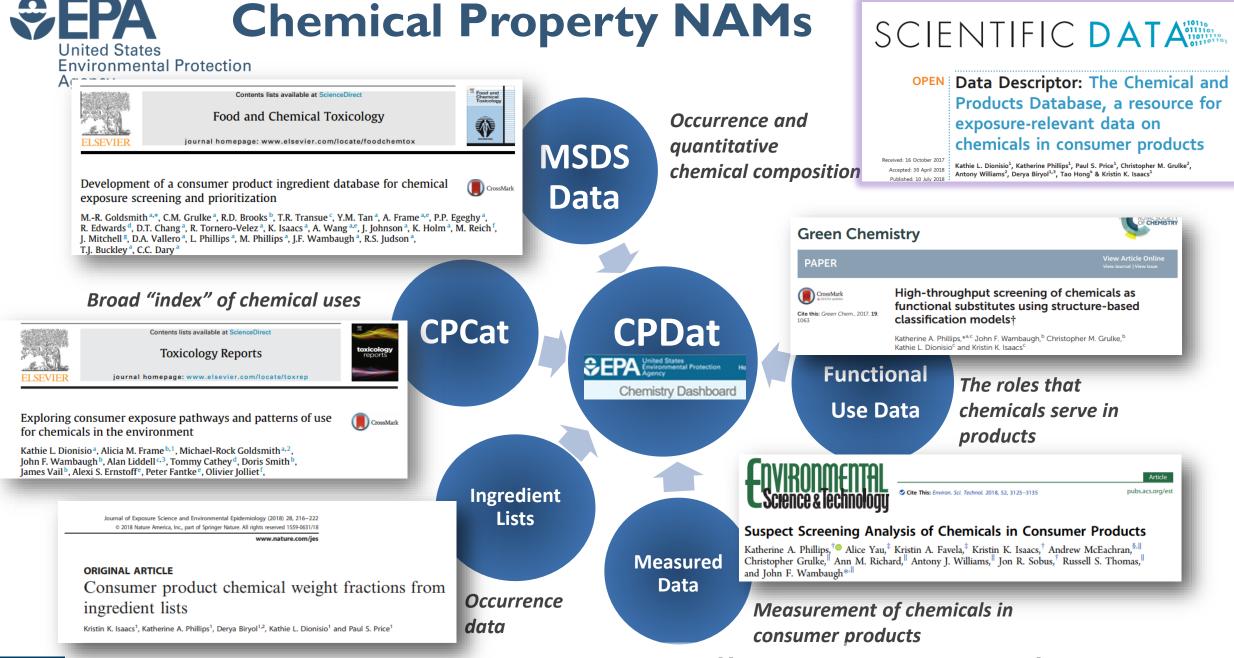
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#### https://comptox.epa.gov/dashboard



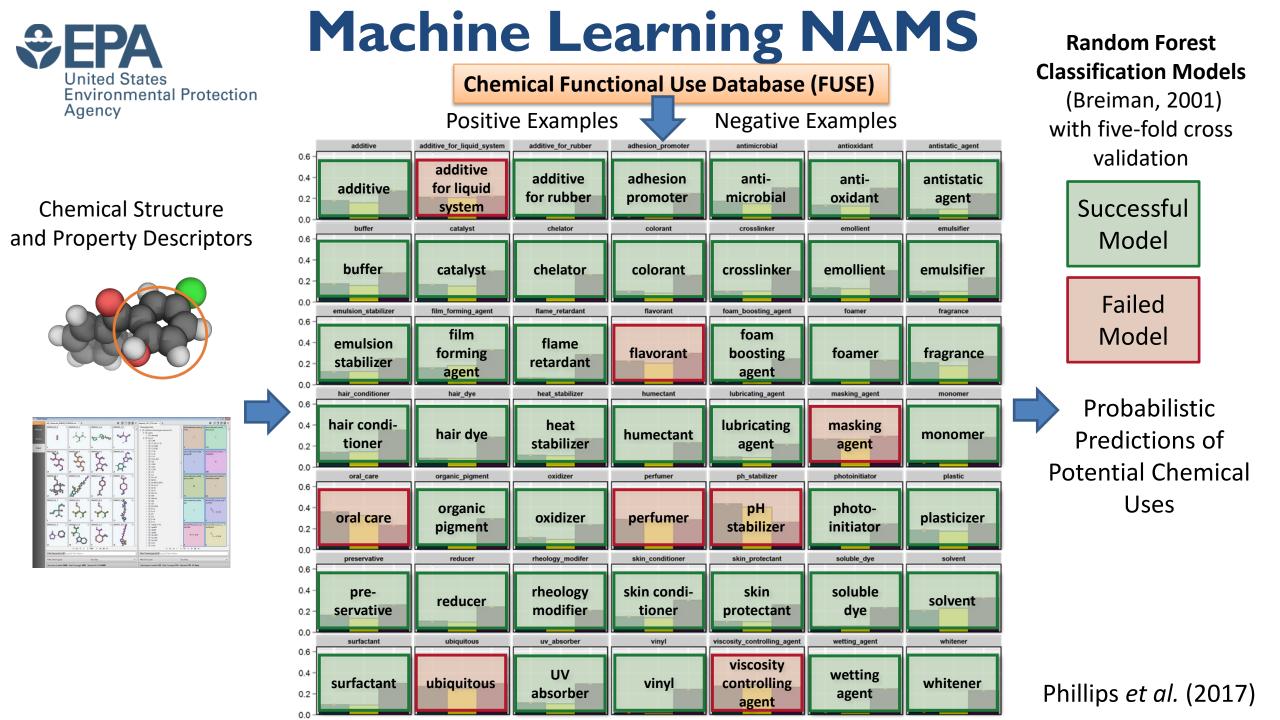
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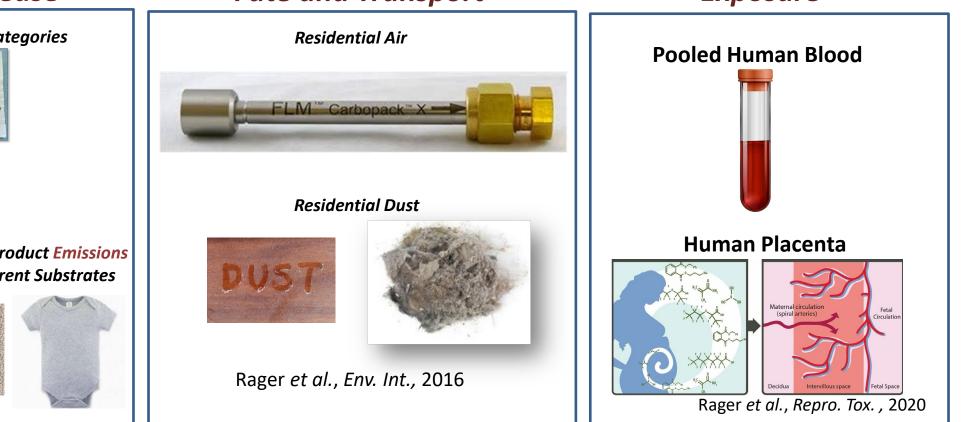
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Wambaugh et al., (2019)

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# Published and Ongoing NTA Studies in the ExpoCast Project ease Fate and Transport Exposure regories Residential Air Decled Human Bill



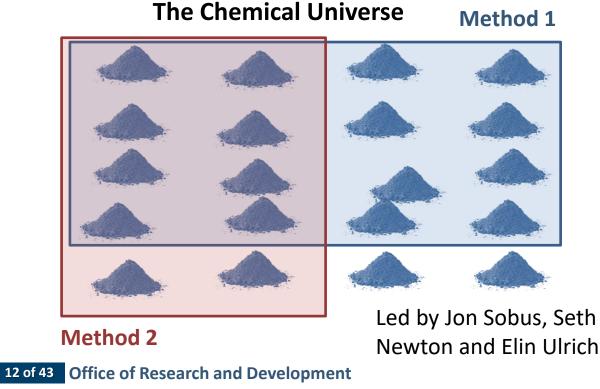
Emerging Science: How can we **quantify** concentrations of chemicals in media using NTA?

Slide from Kristin Isaacs



#### EPA's Non-Targeted Analysis Collaborative Trial (ENTACT)

- Suspect screening / Non-targeted analyses (SSA/NTA) present opportunities for new exposure data
- What NTA methods are available? What is the coverage of chemical universe and matrices? How do methods differ in their coverage?





- Phase 1:
  - Collaborators provided 10 mixtures of 100-400 ToxCast chemicals each
  - Mass spectrometry equipment vendors provided with individual chemical standards
- Phase 2: Fortified reference house dust, human serum, and silicone wristbands



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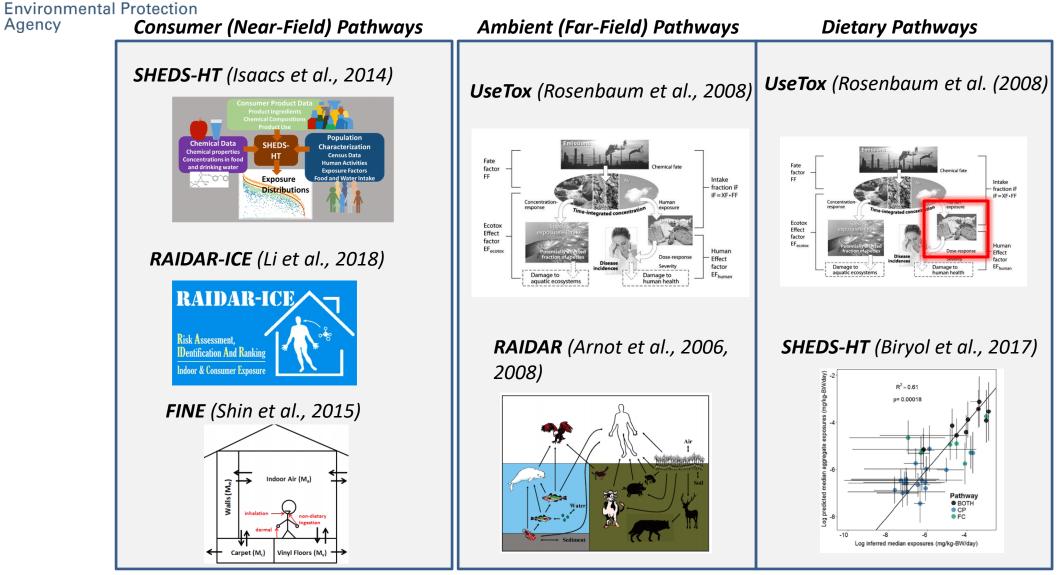
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Wambaugh et al., (2019)

#### **High Throughput Models for Key Pathways**



#### **Slide from Kristin Isaacs**

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#### Fit-for-Purpose Exposure Modeling Frameworks

**Environmental Protection** Increasing Complexity Agency Mechanistic Full exposure description of the assessment First-tier built environment and exposure assessment/ processes, including screening temporal variability Prioritization Description of *Level of aggregation across* human behavior sources, routes, scenarios, or population chemicals

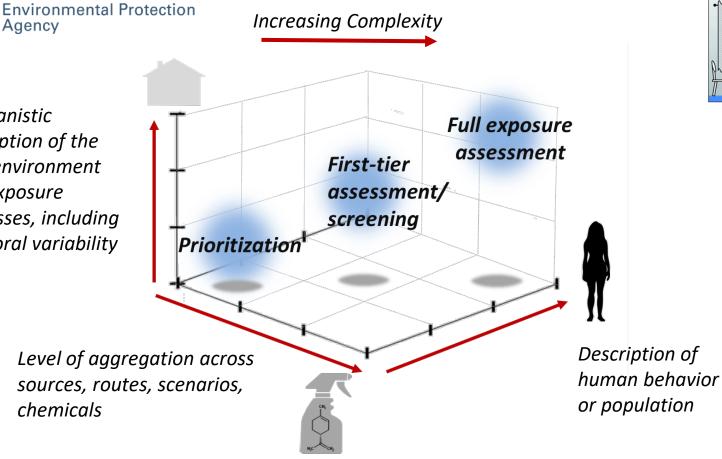
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#### Fit-for-Purpose Exposure Modeling Frameworks

*Mechanistic* description of the built environment and exposure processes, including temporal variability

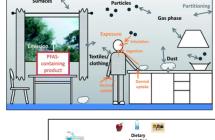
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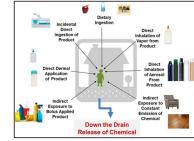
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- Models of different levels of complexity have • overlapping data needs
- They also share some universal challenges •

**Slide from Kristin Isaacs** 





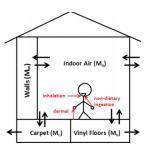
Risk Assessment, Dentification And Rank or & Consumer Expos

**RAIDAR-ICE** 

Eichler and Little, 2020

SHEDS-HT, Isaacs et al., 2014

Li et al., 2018



FINE, Shin et al., 2015



EPA, 2019



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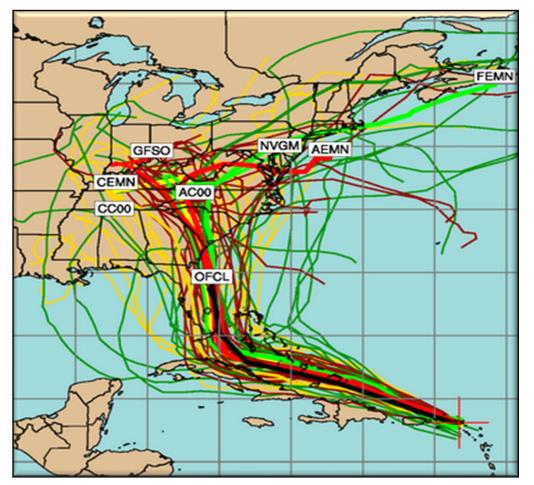
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# **Ensemble Predictions**

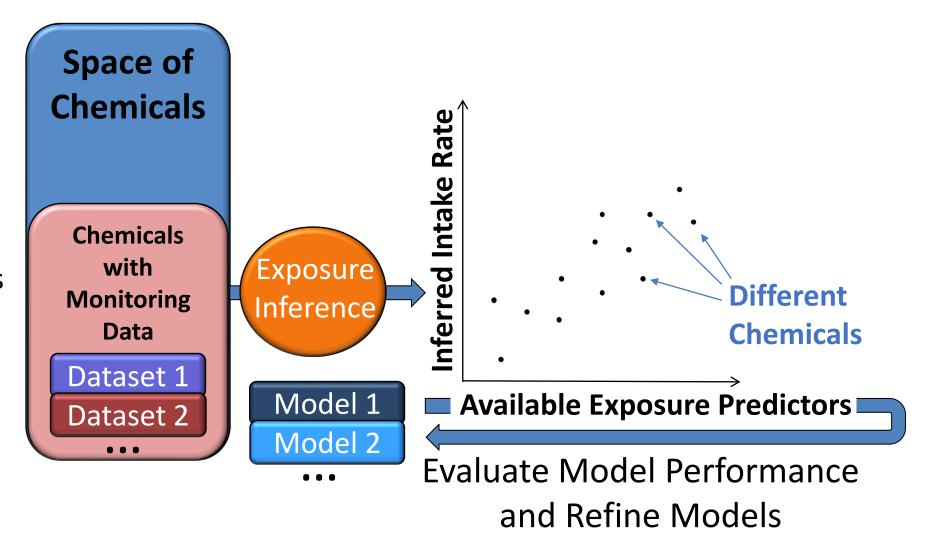
- We can use ensemble methods to make more stable models and characterize uncertainty
- "Ensemble methods are learning algorithms that construct a set of classifiers and then classify new data points by taking a (weighted) vote of their predictions." Dietterich (2000)
- Ensemble systems have proven themselves 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

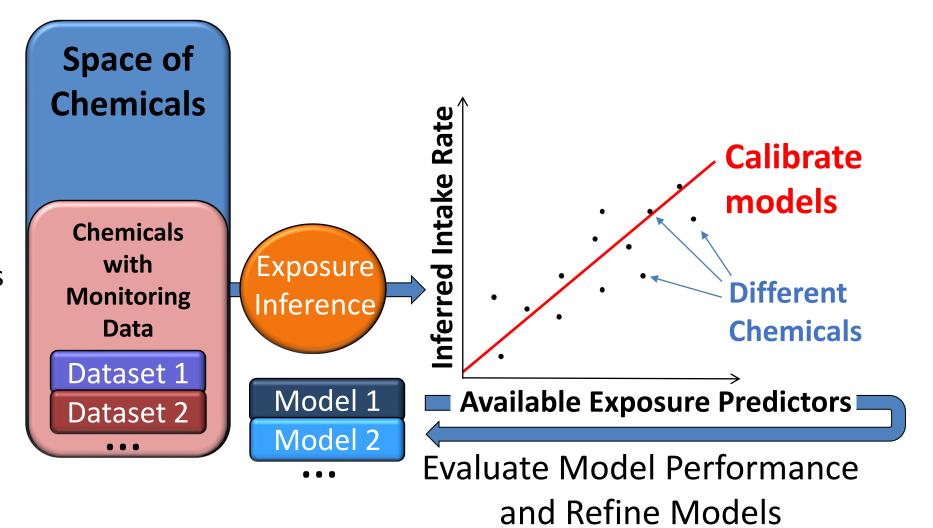


We use Bayesian methods to incorporate multiple models into consensus predictions for 1000s of chemicals within the **Systematic Empirical Evaluation** of Models (SEEM) (Wambaugh et al., 2013, 2014; Ring et al., 2018)



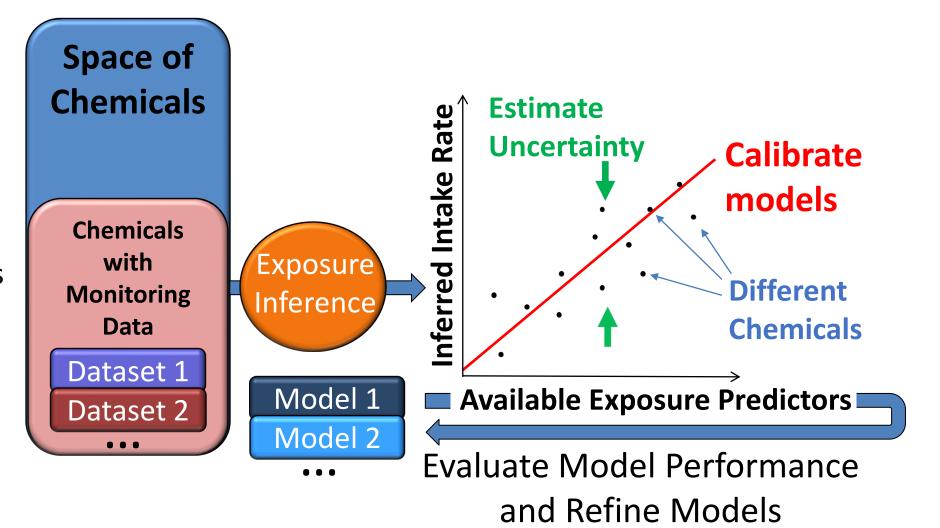


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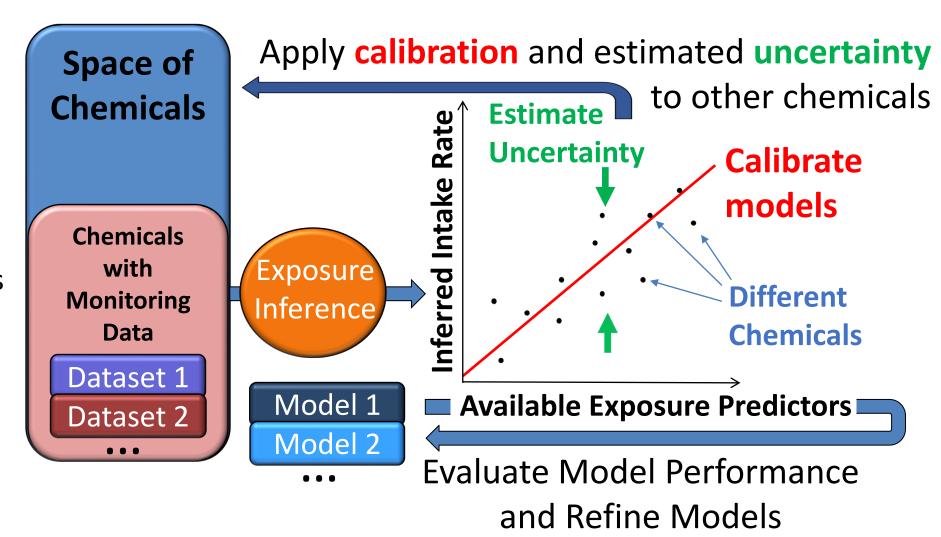


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## **SEEM3** Collaboration

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

US EPA (2018)

**Reference(s)** 

**Chemicals** 

Predicted

7856

8167

Pathway(s)

All

Dietary



Predictor

USEtox Dietary Scenario (2.0)

EPA Inventory Update Reporting and Chemical Data









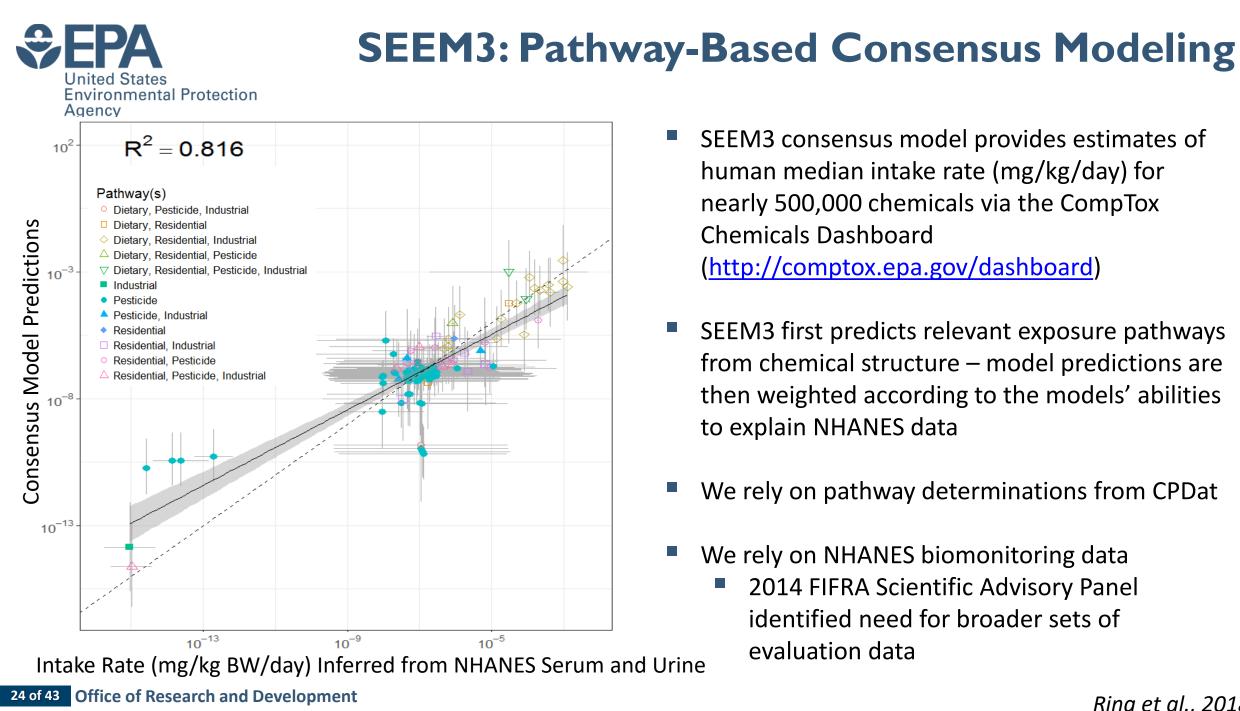


Reporting (CDR) (2015)			
Stockholm Convention of Banned Persistent Organic	Lallas (2001)	248	far field Industrial and
Pollutants (2017)			Pesticide
EPA Pesticide Reregistration Eligibility Documents	Wetmore et al. (2012, 2015)	239	far field Pesticide
(REDs) Exposure Assessments (Through 2015)			
United Nations Environment Program and Society for	Rosenbaum et al. (2008)	8167	far field Industrial
Environmental Toxicology and Chemistry toxicity model			
(USEtox) Industrial Scenario (2.0)			
USEtox Pesticide Scenario (2.0)	Fantke et al. (2011, 2012, 2016)	940	far field Pesticide
Risk Assessment IDentification And Ranking (RAIDAR)	Arnot et al. (2008)	8167	far field Pesticide
far field (2.02)			
EPA Stochastic Human Exposure Dose Simulator High	Isaacs (2017)	7511	far field Industrial and
Throughput (SHEDS-HT) near field Direct (2017)			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

Jolliet et al. (2015), Huang et al. (2016),

Ernstoff et al. (2017)

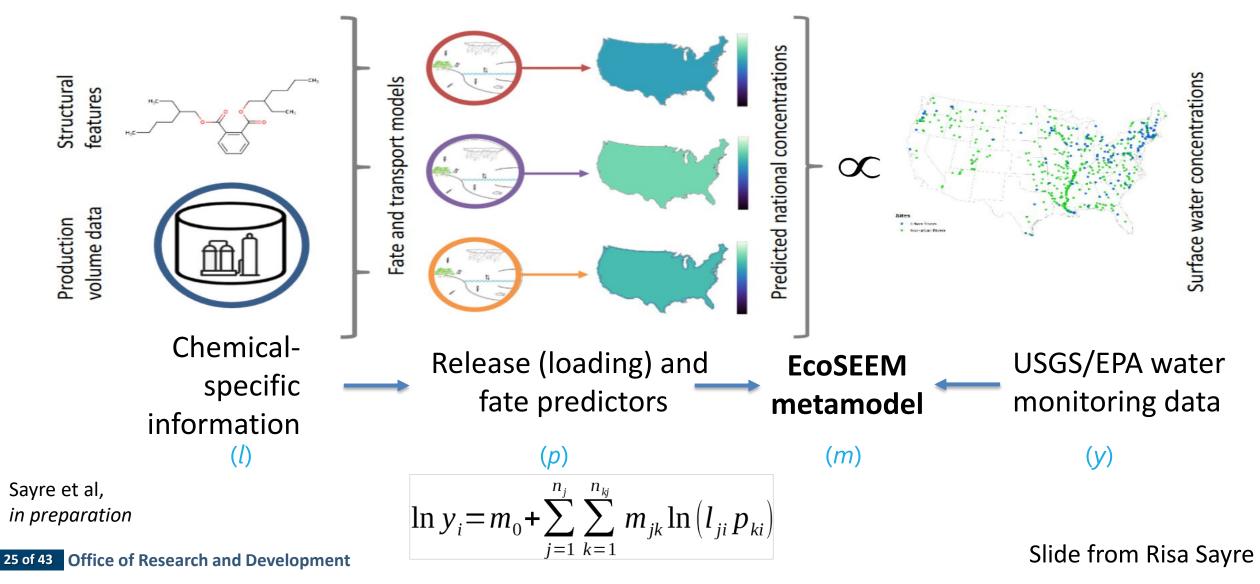
#### *Ring et al., 2018*



- SEEM3 consensus model provides estimates of human median intake rate (mg/kg/day) for nearly 500,000 chemicals via the CompTox Chemicals Dashboard (http://comptox.epa.gov/dashboard)
- SEEM3 first predicts relevant exposure pathways from chemical structure – model predictions are then weighted according to the models' abilities to explain NHANES data
- We rely on pathway determinations from CPDat
- We rely on NHANES biomonitoring data
  - 2014 FIFRA Scientific Advisory Panel identified need for broader sets of evaluation data



# **EcoSEEM Metamodel for Surface** Water Chemical Concentrations





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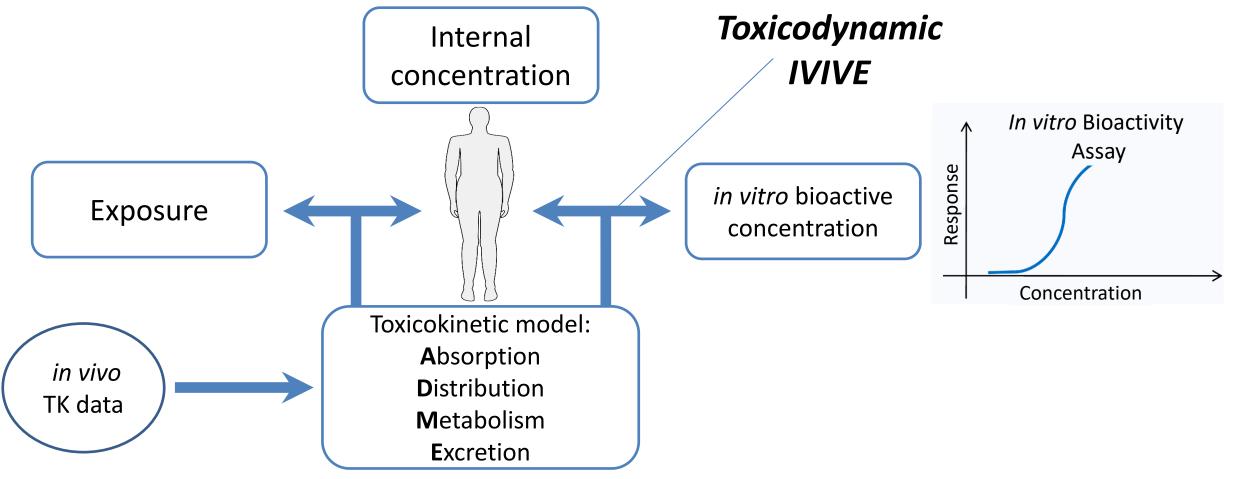
Wambaugh et al., (2019)

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# In Vitro-In Vivo Extrapolation (IVIVE)

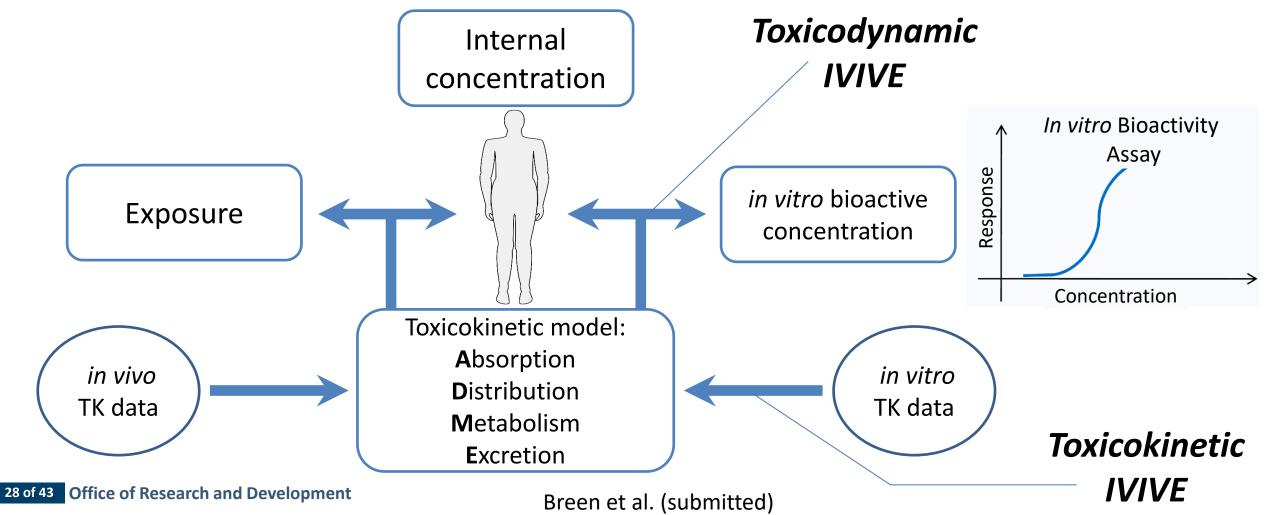
- Translation of *in vitro* high throughput screening requires chemical-specific toxicokinetic models
  - Needed for anywhere from dozens to thousands of chemicals





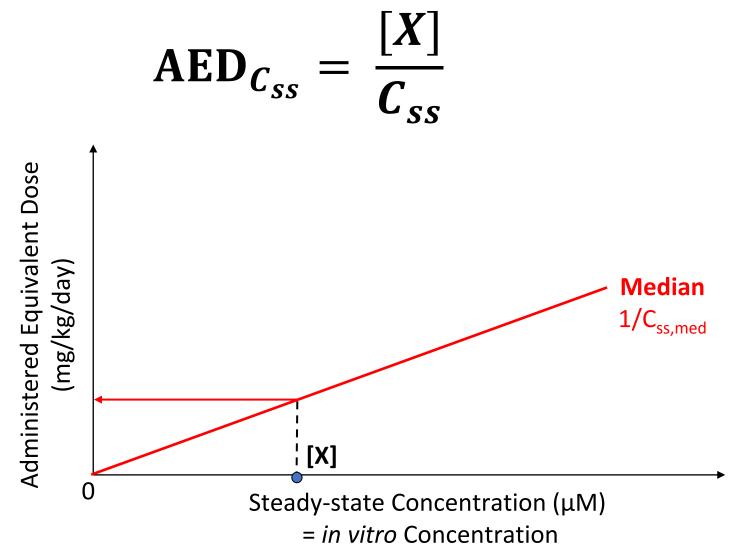
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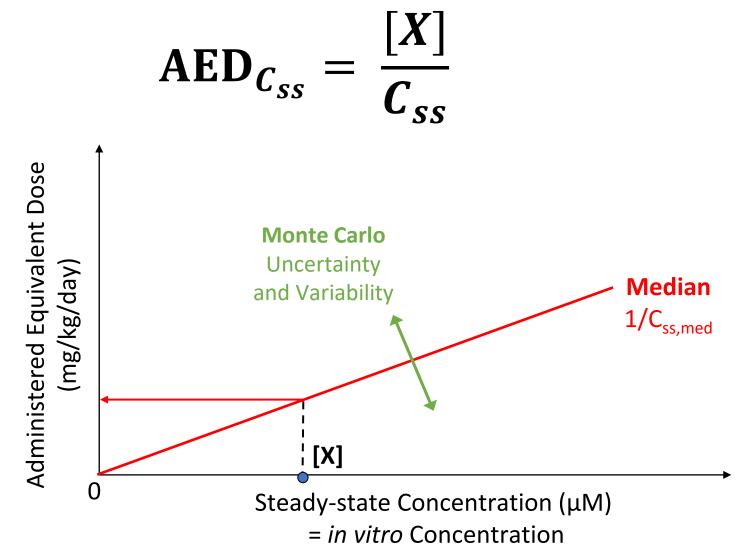


## **Reverse Dosimetry (IVIVE)**



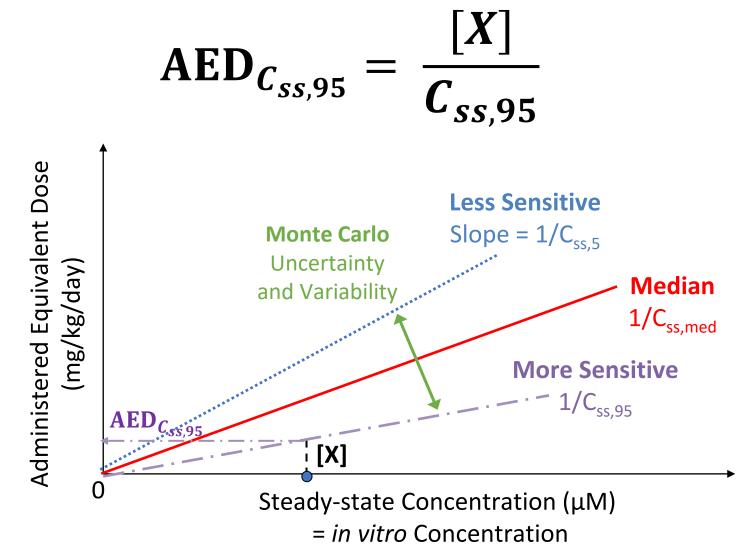


## **Reverse Dosimetry (IVIVE)**



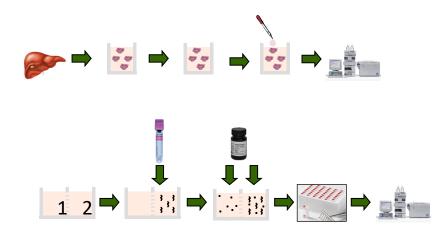


## **Reverse Dosimetry (IVIVE)**



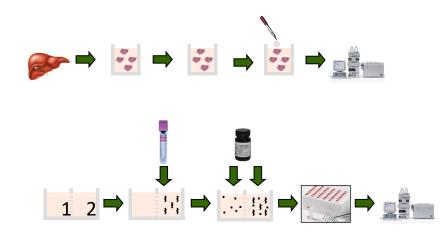


#### In vitro toxicokinetic data





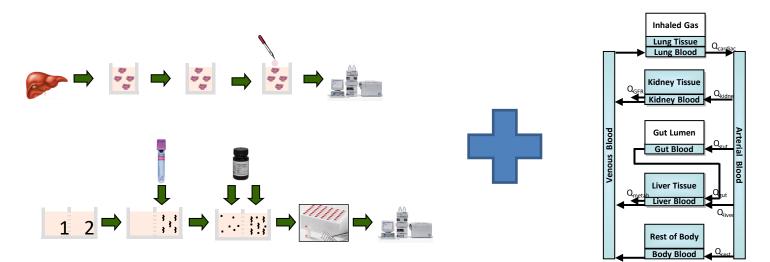
#### In vitro toxicokinetic data



Rotroff et al. (2010) Wetmore et al. (2012) Wetmore et al. (2015) Wambaugh et al. (2019)



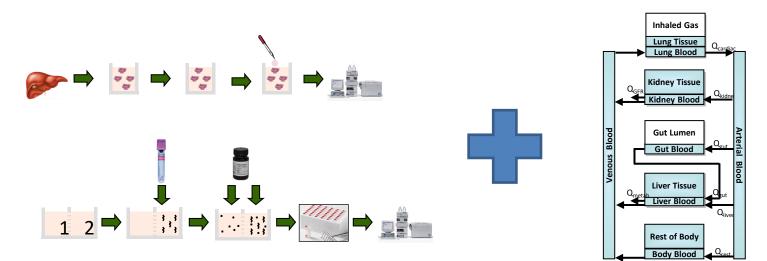
#### *In vitro* toxicokinetic data + generic toxicokinetic model



Rotroff et al. (2010) Wetmore et al. (2012) Wetmore et al. (2015) Wambaugh et al. (2019)



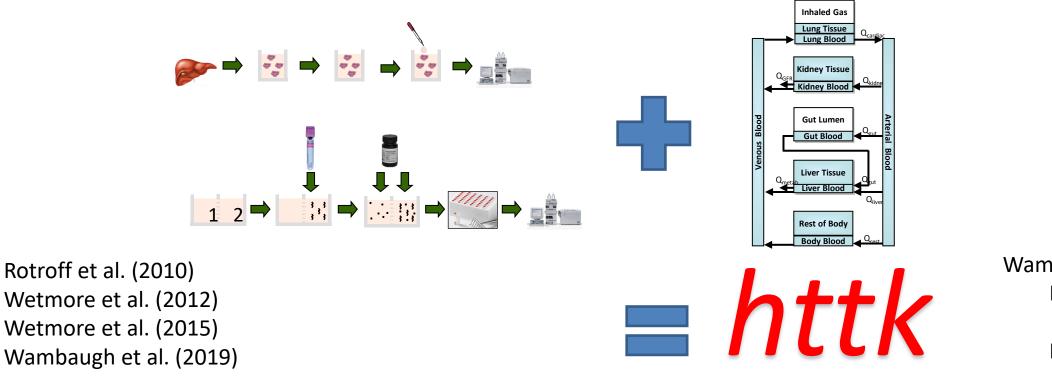
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Rotroff et al. (2010) Wetmore et al. (2012) Wetmore et al. (2015) Wambaugh et al. (2019) Wambaugh et al. (2015) Pearce et al. (2017) Ring et al. (2017) Linakis et al. (2020)



# In vitro toxicokinetic data + generic toxicokinetic model = high(er) throughput toxicokinetics

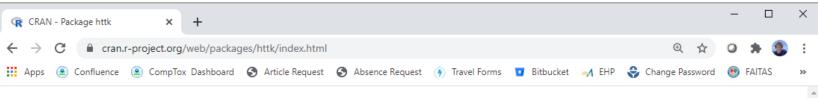


Wambaugh et al. (2015) Pearce et al. (2017) Ring et al. (2017) Linakis et al. (2020)



# **Open-Source Tools and Data for HTTK**

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



#### httk: High-Throughput Toxicokinetics

Generic models and chemical-specific data for simulation and statistical analysis of chemical toxicokinetics Pearce et al. (2017) <<u>doi:10.18637/jss.v079.i04</u>>. Chemical-specific in vitro data have been obtained from r experiments. Both physiologically-based ("PBTK") and empirical (for example, one compartment) "TK" me parameterized with the data provided for thousands of chemicals, multiple exposure routes, and various spec of systems of ordinary differential equations which are solved using compiled (C-based) code for speed. A N included, which allows for simulating human biological variability (Ring et al., 2017 <<u>doi:10.1016/j.envint.</u> propagating parameter uncertainty. Calibrated methods are included for predicting tissue:plasma partition co distribution (Pearce et al., 2017 <<u>doi:10.1007/s10928-017-9548-7</u>>). These functions and data provide a set vivo extrapolation ("IVIVE") of high throughput screening data (for example, Tox21, ToxCast) to real-world dosimetry (also known as "RTK") (Wetmore et al. 2015 <<u>doi:10.1093/toxeci/bfu171></u>)

Version: Depends:	$\frac{2.0.3}{R(\geq 2.10)}$ downloads 1071/month	
Imports:	deSolve, msm, data.table, survey, mythorm, trunchorm, stats, graphics, utils, <u>magrittr, p</u>	
Suggests:	<u>ggplot2, knitr, rmarkdown, R.rsp, GGally, gplots, scales, EnvStats, MASS, RColorBrev</u> <u>classInt, ks, stringr, reshape, reshape2, gdata, viridis, CensRegMod, gmodels, colorspac</u> <u>dplyr, forcats, smatr, gtools, gridExtra</u>	
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 Si Wetmore ( [ctb], Woodrow Setzer ( [ctb])	,
Maintainer:	John Wambaugh <wambaugh.john at="" epa.gov=""></wambaugh.john>	
RugRenorts:	https://github.com/USEPA/CompTox-ExpoCast-httk	

# R package "httk"

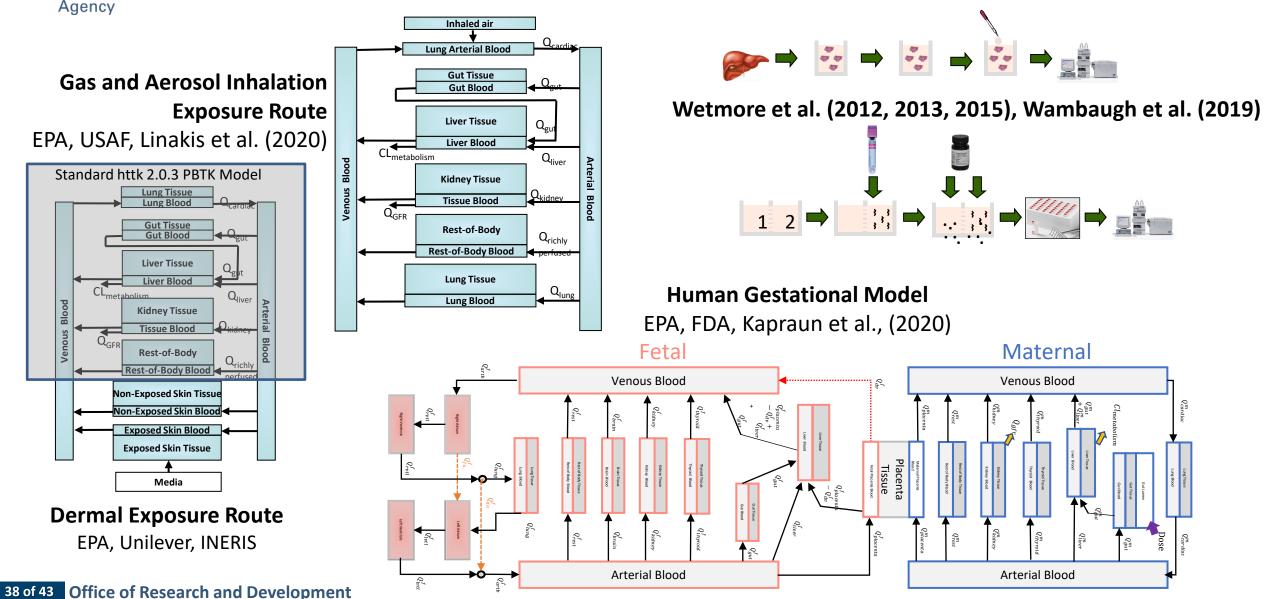
- Open source, transparent, and peerreviewed tools and data for high throughput toxicokinetics (httk)
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific data for 987 chemicals
- Described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



Blood

# **Toxicokinetics NAMs:** In Vitro Measurements and Generic

**PBTK Models** 



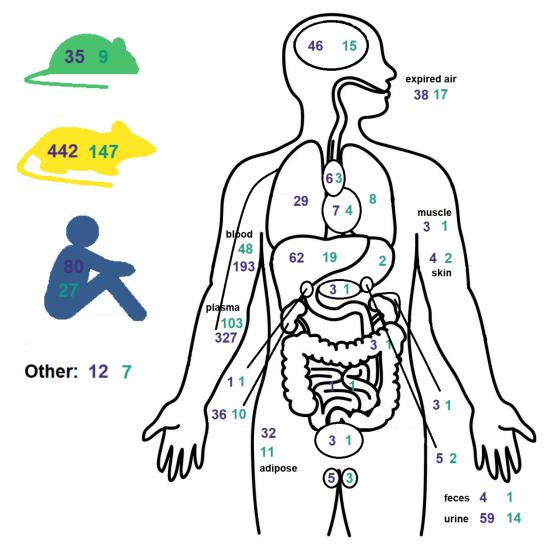


## In Vivo TK Database

#### https://github.com/USEPA/CompTox-PK-CvTdb

- EPA has developed a public database of concentration
   vs. time data for building, calibrating, and evaluating TK models
- Curation and development 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:

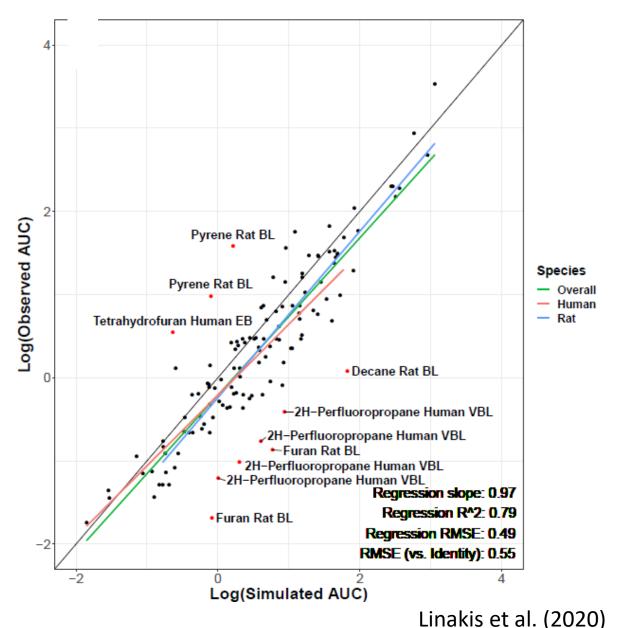
https://github.com/USEPA/CompTox-ExpoCast-invivoPKfit





#### **Developing Models with the CvT Database**

- USAF and EPA developed generic gas inhalation physiologically-based toxicokinetic (PBTK) model
- Evaluated HTTK with CvTdb: 142 exposure scenarios across 41 volatile organic chemicals were modeled and compared to published *in vivo* data for humans and rat
- R<sup>2</sup> was 0.69 for predicting peak concentration
- R<sup>2</sup> was 0.79 for predicting time integrated plasma concentration (Area Under the Curve, AUC)





NAM Makes Use of

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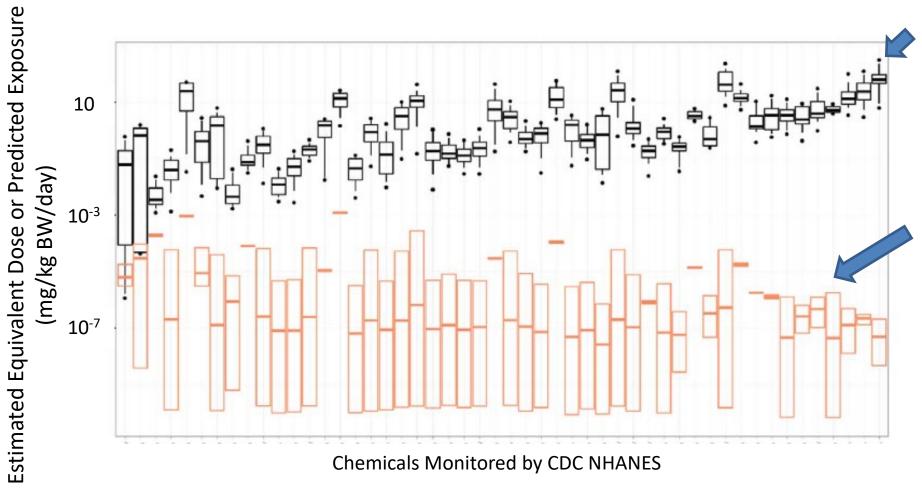
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Exposure NAM Class	Description	Traditional Approach	Measureme	Toxicokineti	Models	Descriptors	Evaluation	Machine Lear
Measurements	New techniques including screening analyses capable of detecting hundreds of chemicals present in a sample	Targeted (chemical-specific) analyses	-	•	•	•		•
Toxicokinetics	High throughput methods using <i>in vitro</i> data to generate chemical-specific models	Analyses based on <i>in vivo</i> animal studies	•	-		•		•
HTE Models	Models capable of making predictions for thousands of chemicals	Models requiring detailed, chemical- and scenario-specific information	•	•	-	•		
Chemical Descriptors	Informatic approaches for organizing chemical information in a machine-readable format	Tools targeted at single chemical analyses by humans				-		•
Evaluation	Statistical approaches that use the data from many chemicals to estimate the uncertainty in a prediction for a new chemical	Comparison of model predictions to data on a per chemical basis	•	•	•	•	-	•
Machine Learning	Computer algorithms to identify patterns	Manual Inspection of the Data	•	•		•		-
<b>Prioritization</b>	Integration of exposure and other NAMs to identify chemicals for follow-up study	Expert decision making	•	•	•	•	•	•

Wambaugh et al., (2019)

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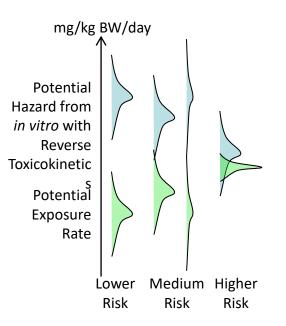


# **Chemical Prioritization NAMs**



High throughput *in vitro* screening can estimate doses needed to cause bioactivity (for example, Wetmore et al., 2015)

Exposure intake rates can be inferred from biomarkers (for example, Ring et al., 2018)

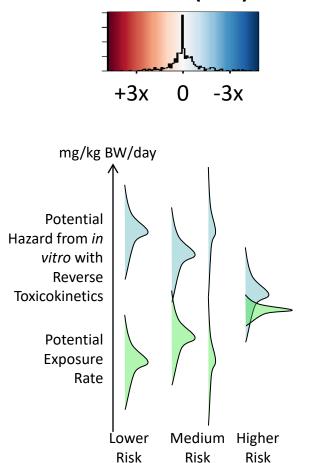


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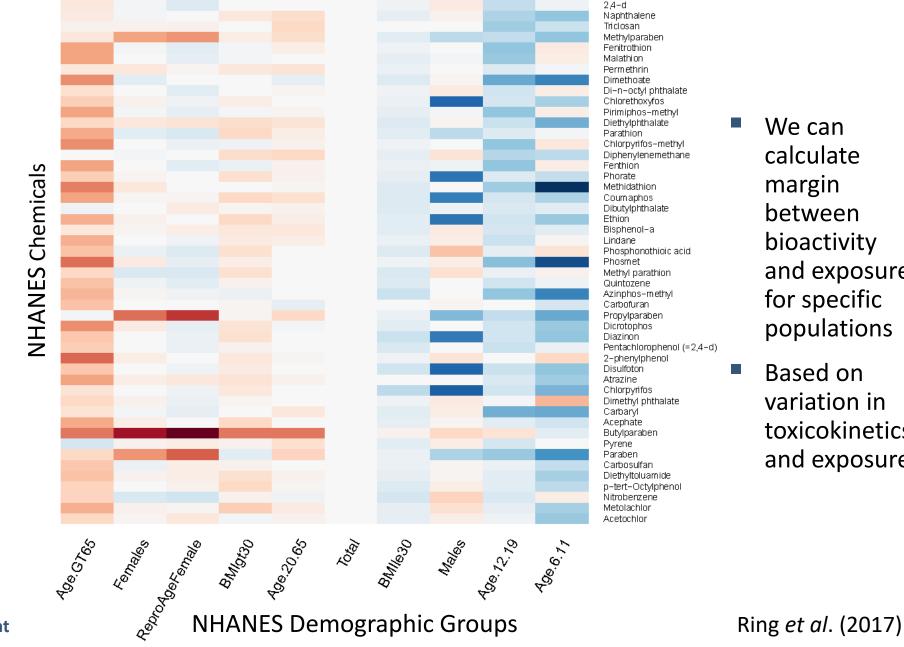
Ring *et al*. (2017)



Change in **Bioactivity : Exposure** Ratio (Risk)



#### Life-stage and Demographic Specific Predictions



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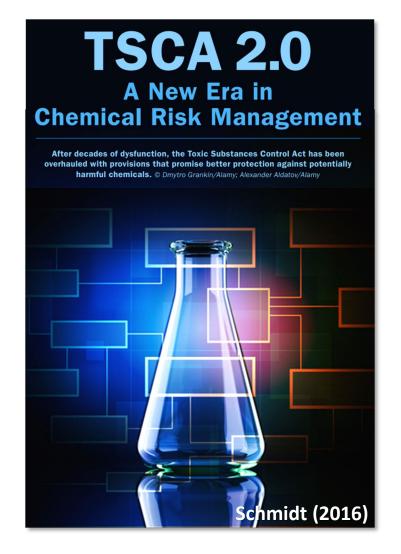
bioactivity and exposure for specific populations

variation in toxicokinetics and exposure



# Outlook

- In ExpoCast we develop models and perform experiments for both exposure and dosimetry (toxicokinetics)
- HTTK (high throughput toxicokinetics) includes a suite of peer-reviewed models for toxicokinetics that can be parameterized for nearly one thousand chemicals
  - Currently adding new models (aerosol, dermal, human gestational)
  - Adding new structure-based predictors for data that are currently measured *in vitro*
- SEEM (systematic empirical evaluation of models) is a consensus metamodeling framework for exposure
  - Trained to monitoring data (developing more)
  - Trained to chemical use data (developing more)
  - Human developed, ecological and occupational in progress



## ExpoCast Project (Exposure Forecasting)

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