

Toxicokinetic Human Variability Simulations Help Establish a Risk Context

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**Data and Diversity:
Improving Coverage of Relevant Populations
(Toxicology Track)**

Thursday, June 9, 2022

The views expressed in this presentation are those of the author
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US EPA Office of Research and Development

- The Office of Research and Development (ORD) is the scientific research arm of EPA
 - 543 peer-reviewed journal articles in 2019
- Research is conducted by ORD's four national centers, and three offices organized to address:
 - Public health and env. assessment; comp. tox. and exposure; env. measurement and modeling; and env. solutions and emergency response.
- 13 facilities across the United States
- Research conducted by a combination of Federal scientists (including uniformed members of the **Public Health Service**); contract researchers; and postdoctoral, graduate student, and post-baccalaureate trainees

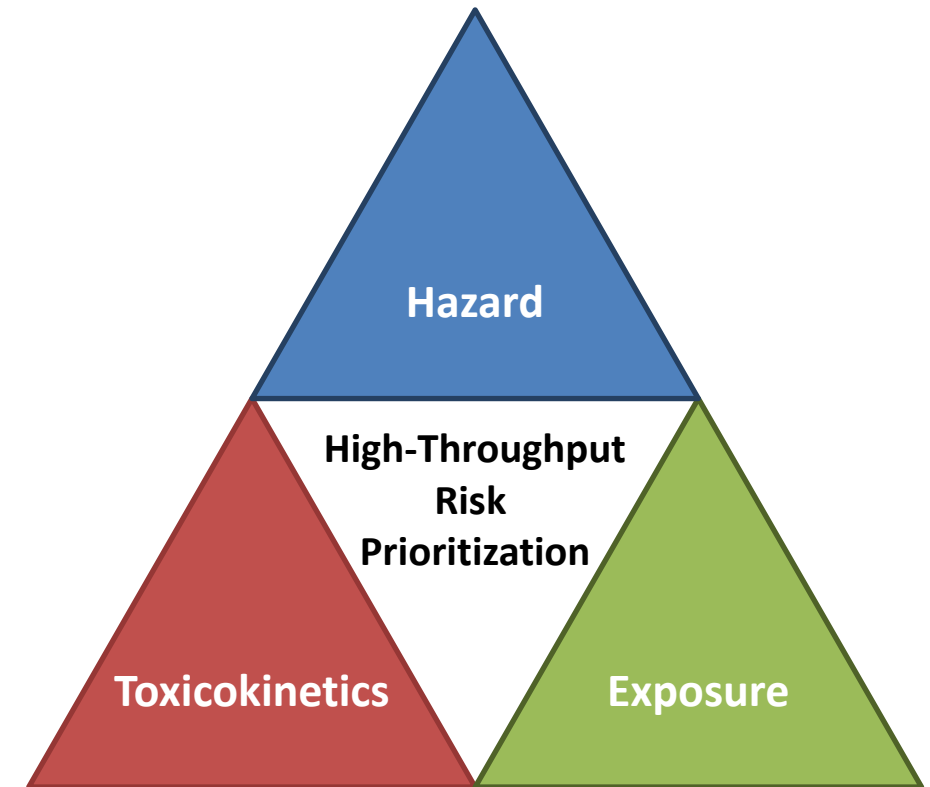


Credit: the Research Triangle Foundation

ORD Facility in
Research Triangle Park, NC

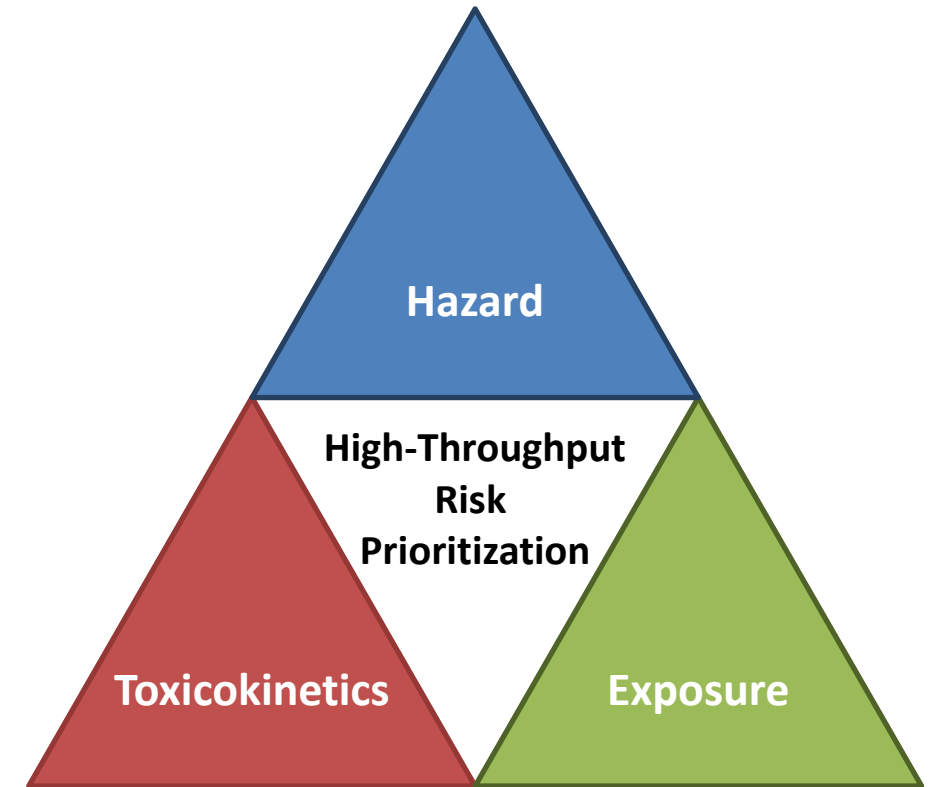
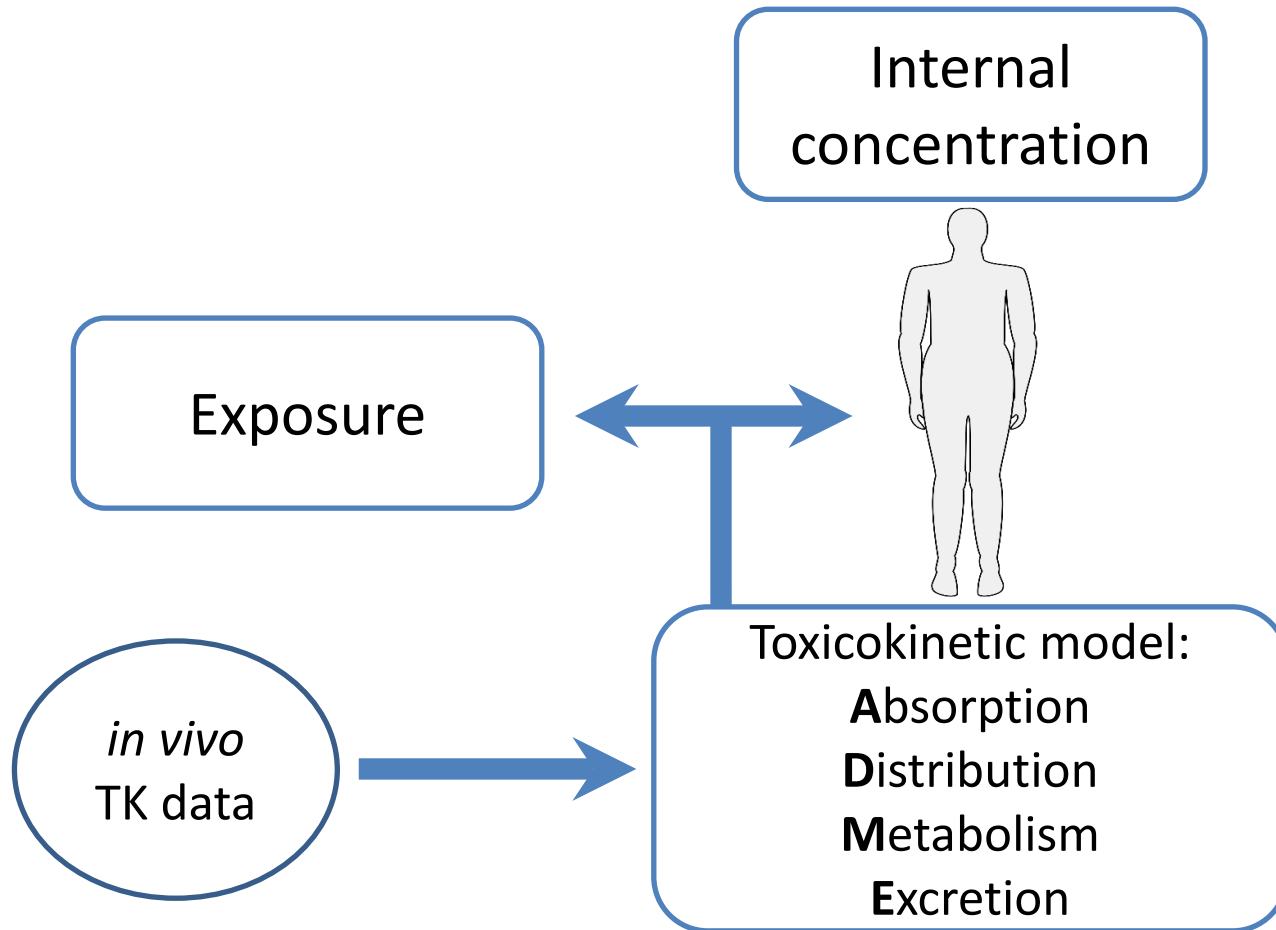
Calculating Chemical Risk

- High throughput risk prioritization based upon *in vitro* screening requires comparison to exposure (for example, NRC, 1983)
 - Data obtained *in vitro* must be placed in an *in vivo* context: *in vitro-in vivo* extrapolation (IVIVE)
- Information must be relevant to the scenario, for example, consumer, ambient, or occupational exposure.



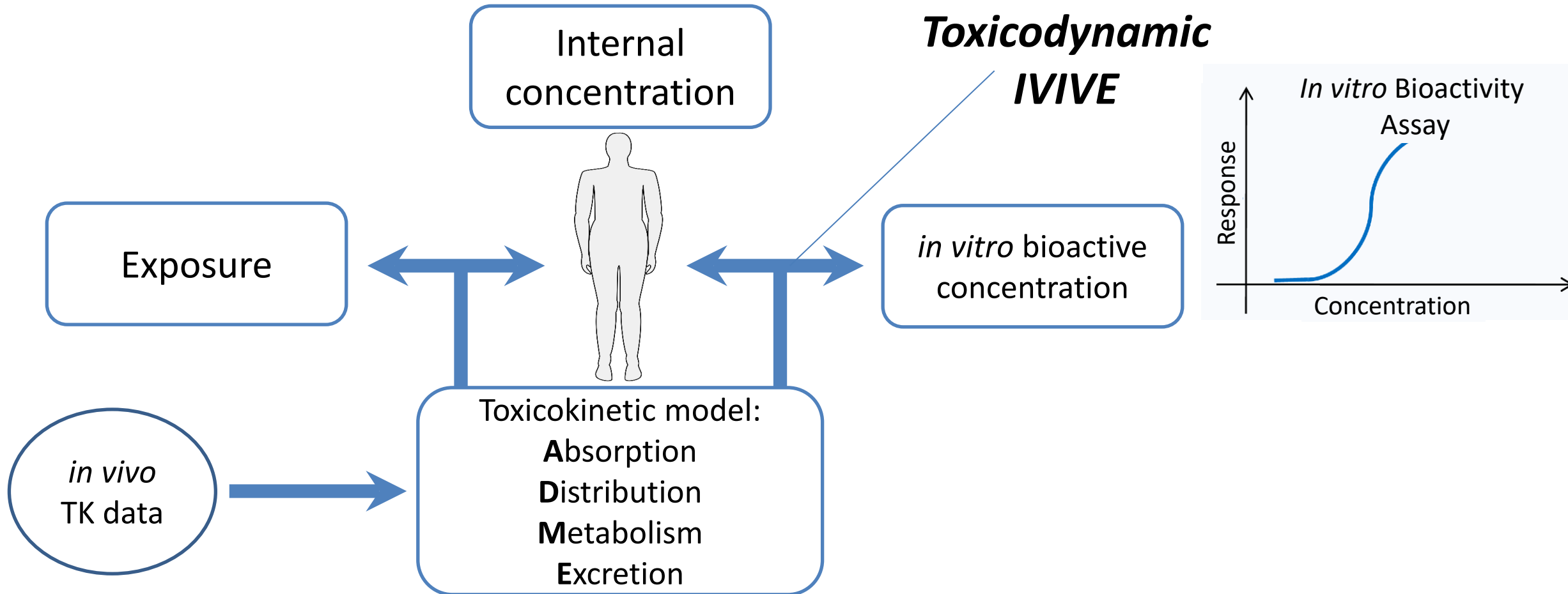
Toxicokinetics

- Toxicokinetics describes the absorption, distribution, metabolism, and excretion of a chemical by the body:
 - Chemical-specific
 - Links exposure with internal concentrations



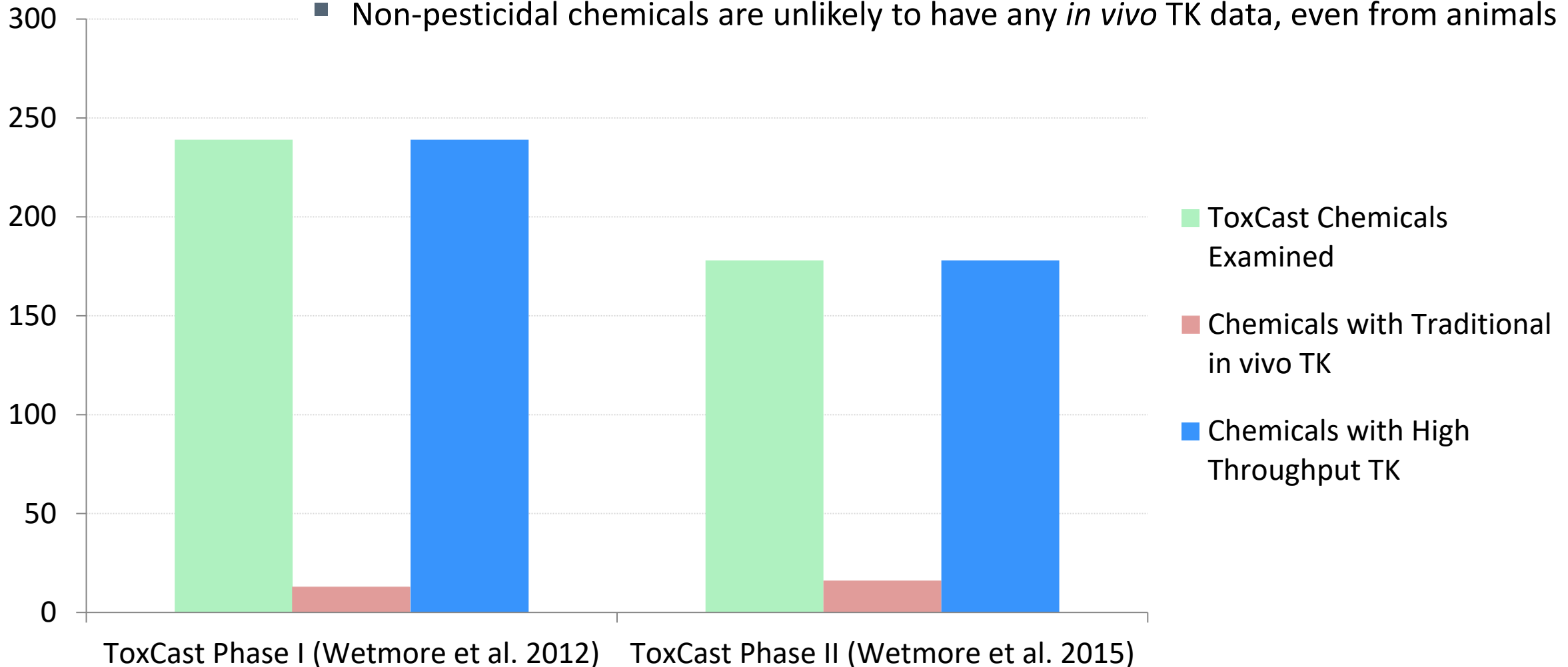
In Vitro-In Vivo Extrapolation (IVIVE)

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



Most Chemicals lack Toxicokinetic Data

- Most non-pharmaceutical chemicals – for example, flame retardants, plasticizers, pesticides, solvents – do not have human *in vivo* TK data.
- Non-pesticidal chemicals are unlikely to have any *in vivo* TK data, even from animals

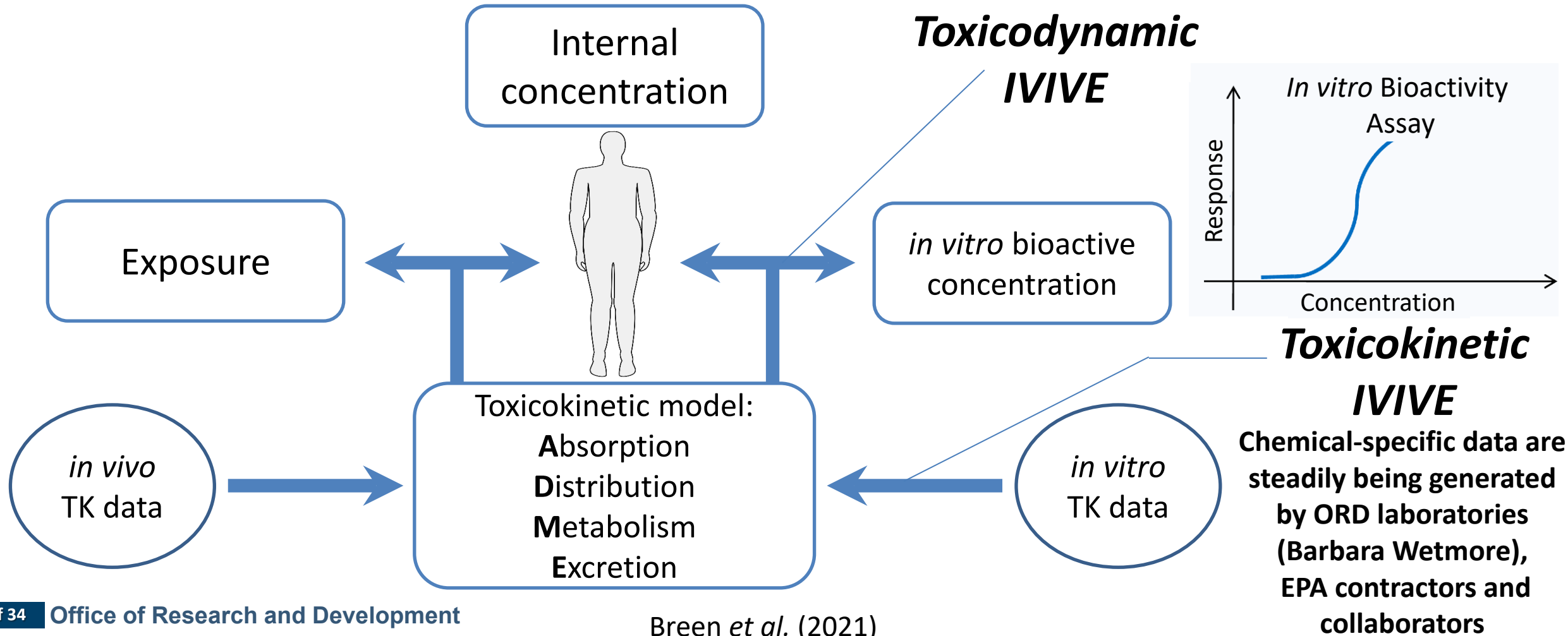


High Throughput Toxicokinetics (HTTK)

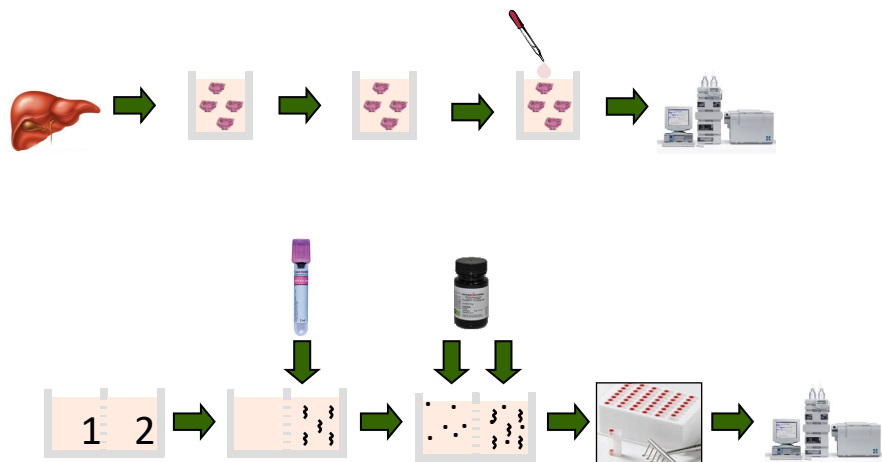
- To provide toxicokinetic data for larger numbers of chemicals collect *in vitro*, high throughput toxicokinetic (HTTK) data
(for example, Rotroff *et al.*, 2010, Wetmore *et al.*, 2012, 2015)
- HTTK methods have been used by the pharmaceutical industry to determine range of efficacious doses and to prospectively evaluate success of planned clinical trials
(Jamei, *et al.*, 2009; Wang, 2010)
- The **primary goal** of HTTK is to provide a human dose context for bioactive *in vitro* concentrations from HTS (that is, *in vitro-in vivo* extrapolation, or **IVIVE**)
(for example, Wetmore *et al.*, 2015)
- A **secondary goal** is to provide **open-source data and models** for evaluation and use by the broader scientific community (Pearce *et al.*, 2017a)

In Vitro-In Vivo Extrapolation (IVIVE)

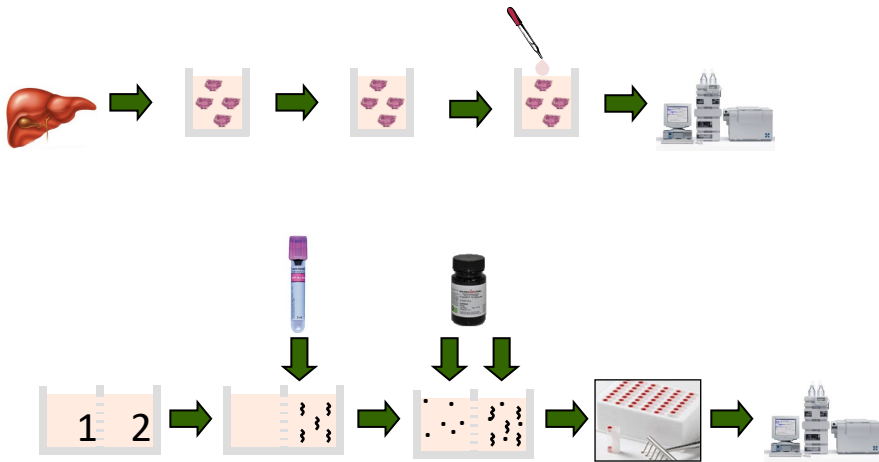
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In vitro toxicokinetic data



In vitro toxicokinetic data



Rotroff *et al.* (2010)

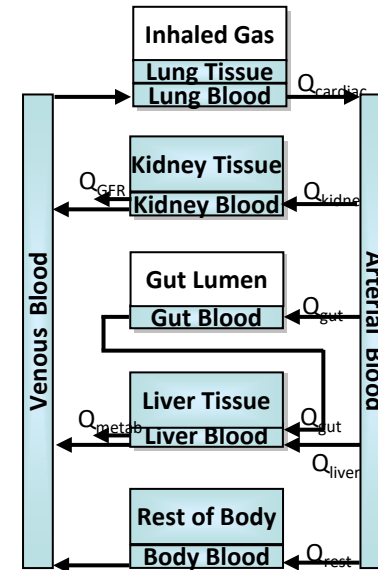
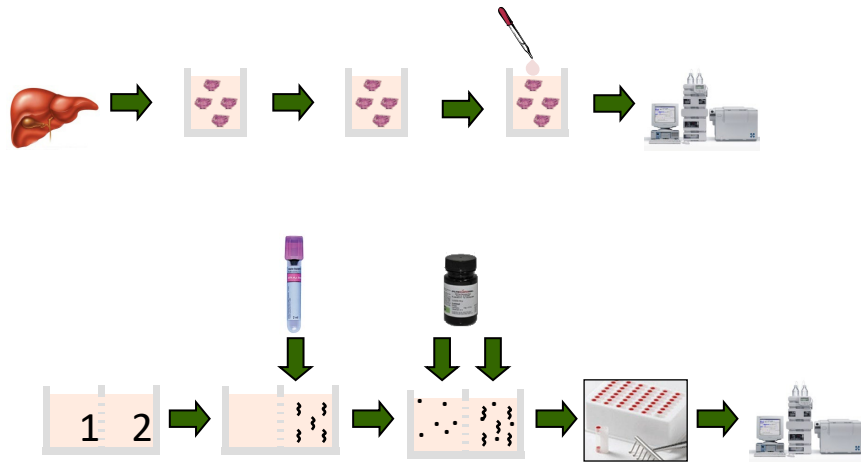
Wetmore *et al.* (2012)

Wetmore *et al.* (2015)

Wambaugh *et al.* (2019)

High Throughput Toxicokinetics (HTTK)

In vitro toxicokinetic data + generic toxicokinetic model



Rotroff *et al.* (2010)

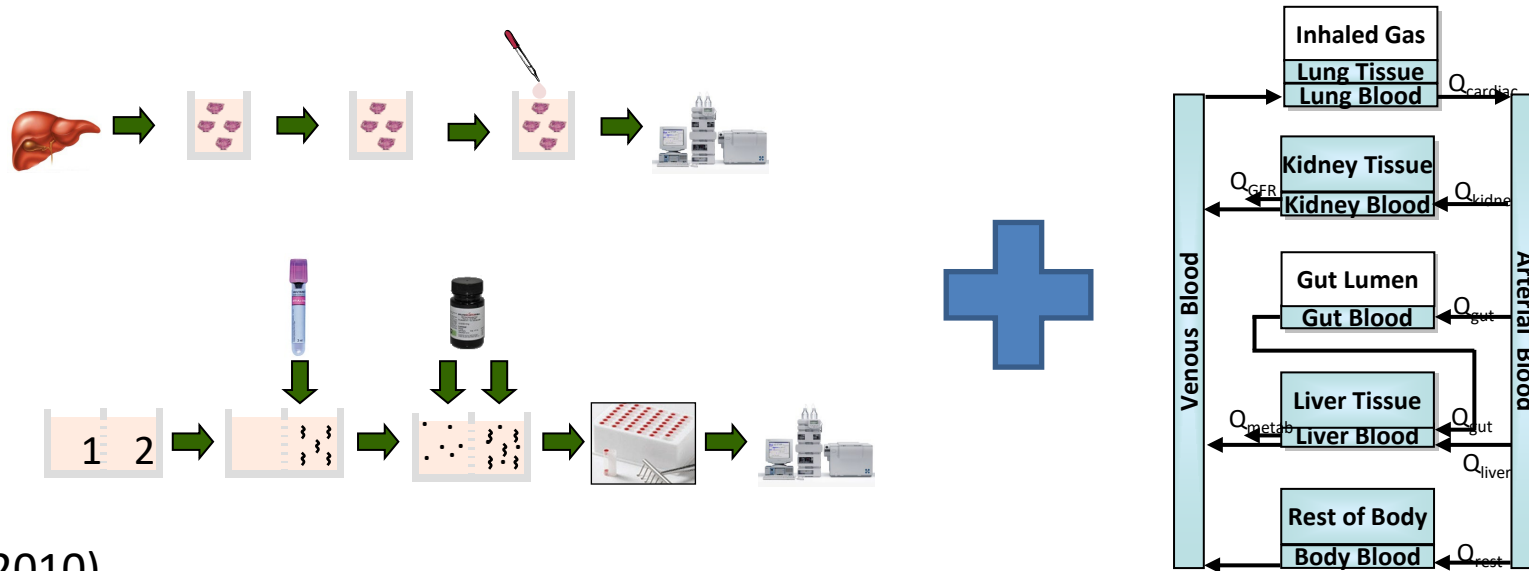
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High Throughput Toxicokinetics (HTTK)

In vitro toxicokinetic data + generic toxicokinetic model

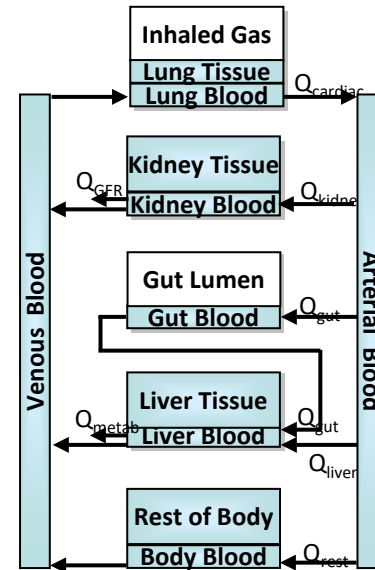
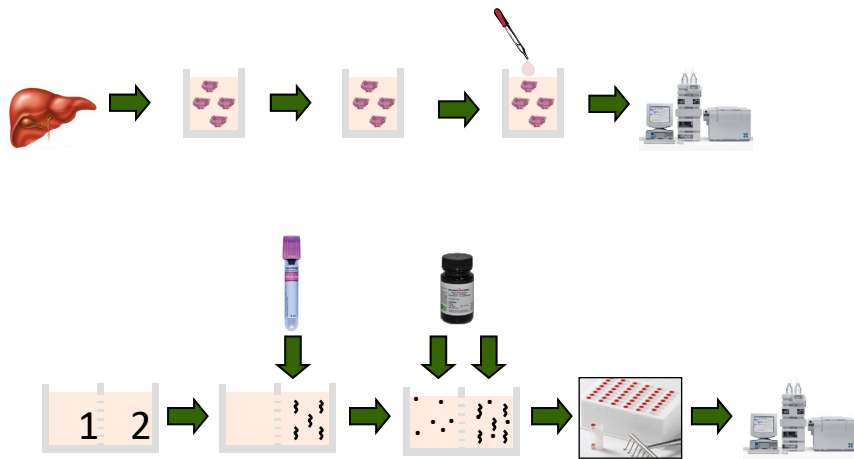


Rotroff *et al.* (2010)
Wetmore *et al.* (2012)
Wetmore *et al.* (2015)
Wambaugh *et al.* (2019)

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

High Throughput Toxicokinetics (HTTK)

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



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

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

= *httk*

Open Source Tools and Data for HTTK

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

CRAN - Package httk

cran.r-project.org/web/packages/httk/index.html

httk: High-Throughput Toxicokinetics

Generic models and chemical-specific data for simulation and statistical analysis of chemical toxicokinetics ("TK") as described by Pearce et al. (2017) <[doi:10.18637/jss.v079.i04](https://doi.org/10.18637/jss.v079.i04)>. Chemical-specific in vitro data have been obtained from relatively high-throughput experiments. Both physiologically-based ("PBTk") and empirical (for example, one compartment) "TK" models can be parameterized with the data provided for thousands of chemicals, multiple exposure routes, and various species. The models consist of systems of ordinary differential equations which are solved using compiled (C-based) code for speed. A Monte Carlo sampler is included, which allows for simulating human biological variability (Ring et al., 2017 <[doi:10.1016/j.envint.2017.06.004](https://doi.org/10.1016/j.envint.2017.06.004)>) and propagating parameter uncertainty. Calibrated methods are included for predicting tissue:plasma partition coefficient (Wetmore et al., 2015 <[doi:10.1093/toxsci/kfv171](https://doi.org/10.1093/toxsci/kfv171)>). These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high-throughput exposures via reverse dosimetry (also known as "RTK") (Wetmore et al., 2015 <[doi:10.1093/toxsci/kfv171](https://doi.org/10.1093/toxsci/kfv171)>).

Version: 2.1.0
Depends: R (≥ 2.10)
Imports: deSolve, msm, data.table, survey, mvtnorm, truncnorm, stats, graphics, utils, magrittr, purrr, met
Suggests: ggplot2, knitr, rmarkdown, R.rsp, GGally, gplots, scales, EnvStats, MASS, RColorBrewer, Teac
Published: 2022-03-26
Author: John Wambaugh, Dustin Kapra, [ctb], Nisha Sipes [id] [ctb], Barbara Wetmore [id] [ctb], Woodrow Setzer [id] [ctb]
Maintainer: John Wambaugh <wambaugh.john@epa.gov>
BugReports: <https://github.com/USEPA/CompTox-ExpoCast-httk>
License: GPL-3
Copyright: This package is primarily developed by employees of the U.S. Federal government as part of the
URL: <https://www.epa.gov/chemical-research/rapid-chemical-exposure-and-dose-research>
NeedsCompilation: yes
Citation: [httk citation info](#)
Materials: [NEWS](#)
CRAN checks: [httk results](#)

downloads 1071/month

R package "httk"

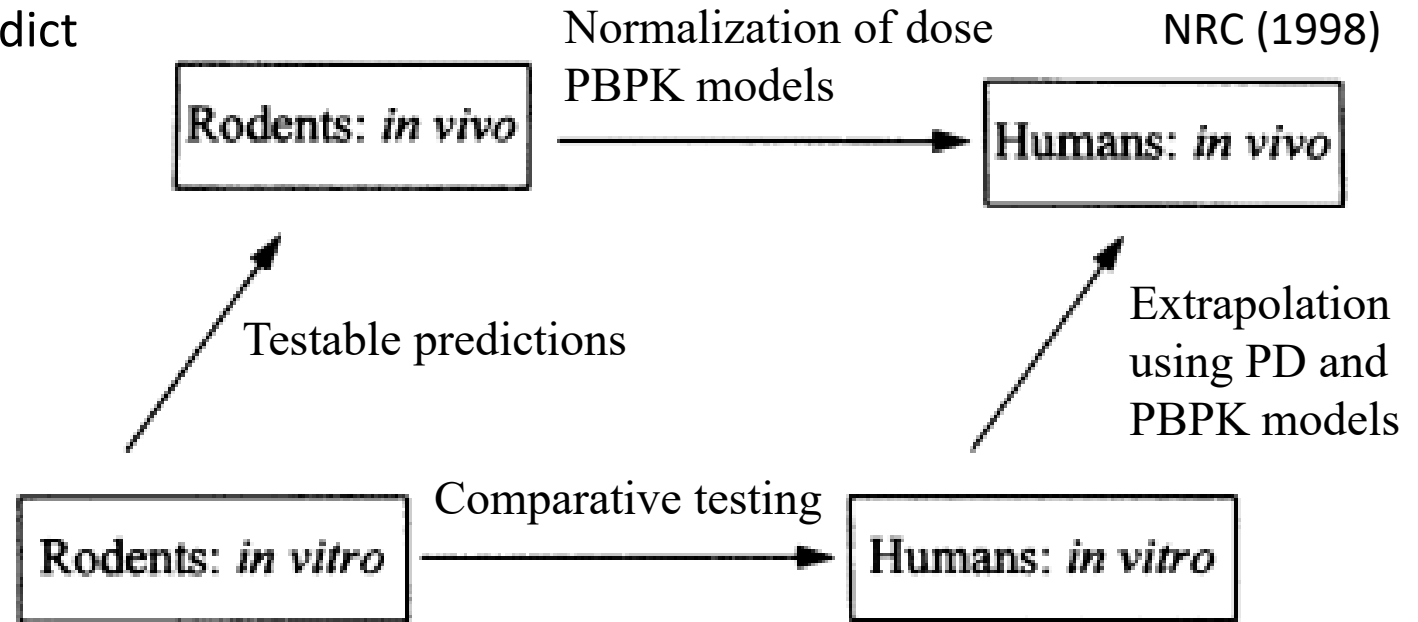
- Open source, transparent, and peer-reviewed tools and data for **high throughput toxicokinetics (httk)**
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTk)
- Human-specific data for 987 chemicals
- Described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

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

In Vitro - *In Vivo* Extrapolation (IVIVE)

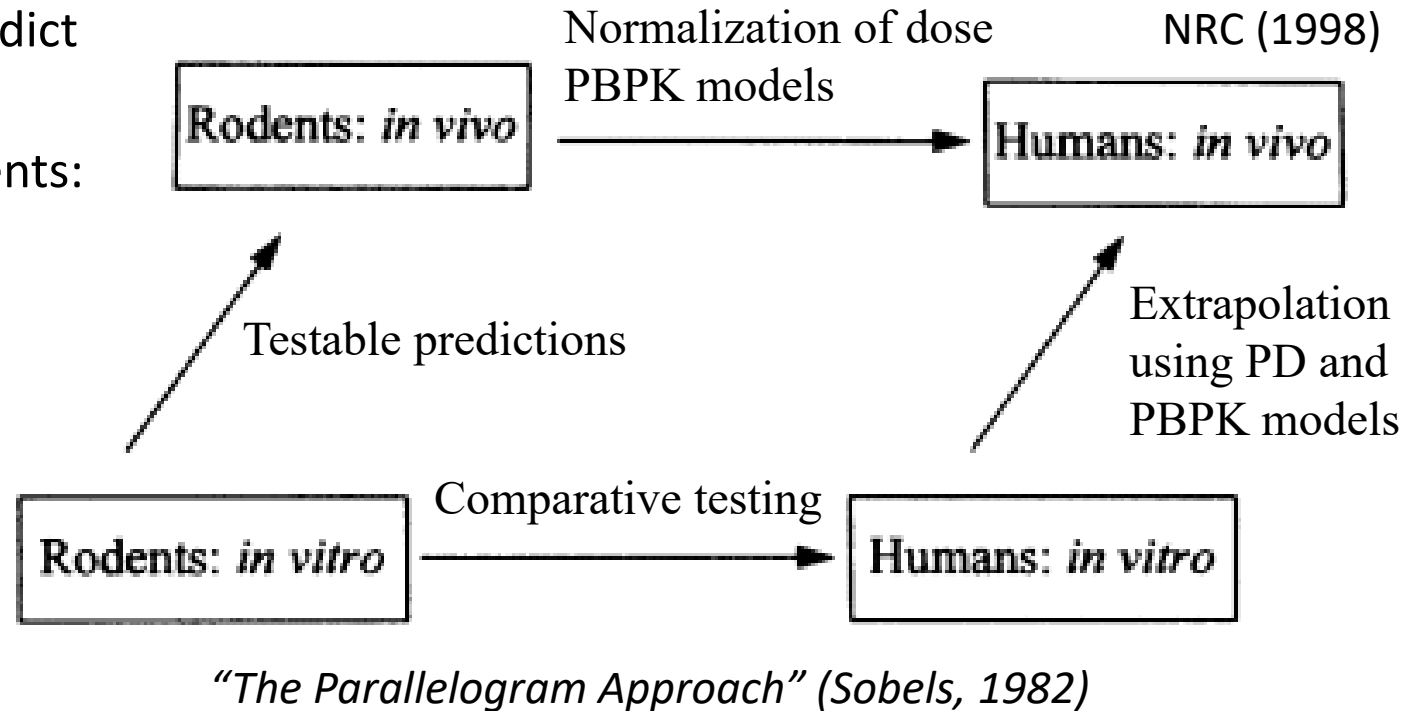
- HTTK allows *in vitro-in vivo* extrapolation (IVIVE)
 - the use of *in vitro* experimental data to predict phenomena *in vivo*.



"The Parallelogram Approach" (Sobels, 1982)

In Vitro - *In Vivo* Extrapolation (IVIVE)

- HTTK allows *in vitro-in vivo* extrapolation (IVIVE)
 - the use of *in vitro* experimental data to predict phenomena *in vivo*.
- IVIVE can be broken down into two components:
 - IVIVE-PK/TK
(**Pharmacokinetics/Toxicokinetics**):
 - Fate of molecules/chemicals in body
 - Considers absorption, distribution, metabolism, excretion (ADME)
 - Can use empirical PK or physiologically-based (PBPK)



In Vitro - *In Vivo* Extrapolation (IVIVE)

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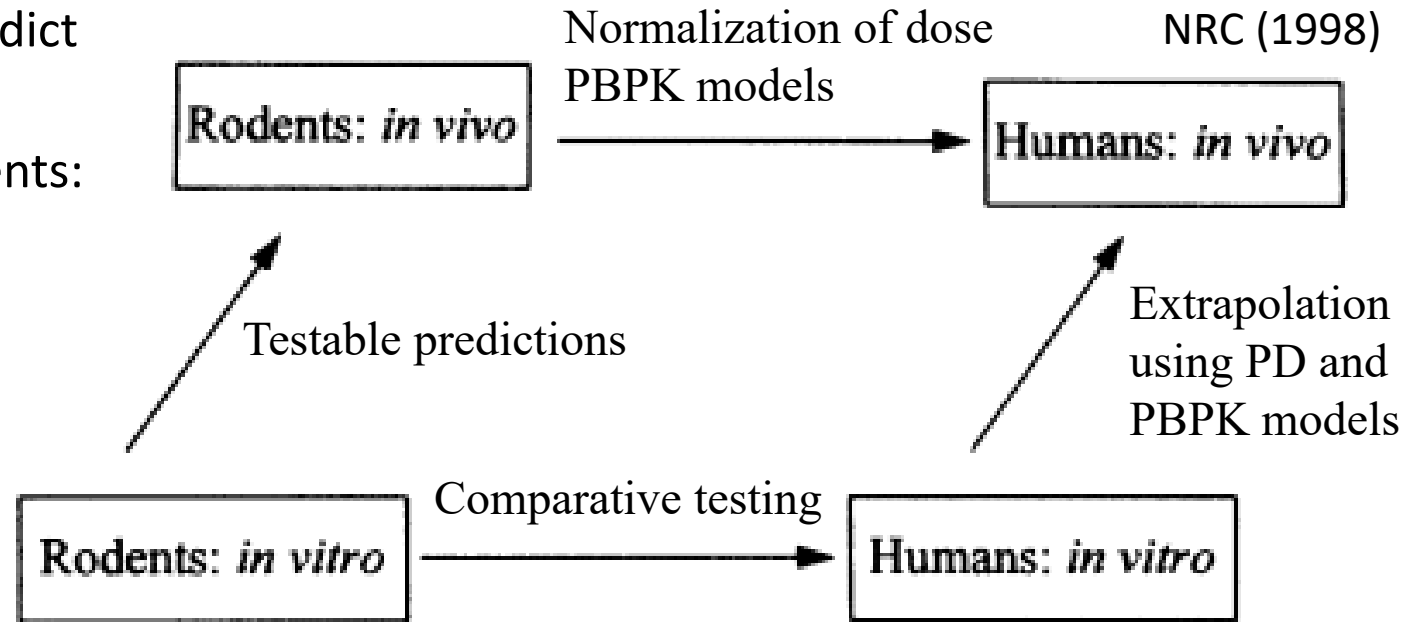
- IVIVE-PK/TK

(Pharmacokinetics/Toxicokinetics):

- Fate of molecules/chemicals in body
- Considers absorption, distribution, metabolism, excretion (ADME)
- Can use empirical PK or physiologically-based (PBPK)

- IVIVE-PD/TD (Pharmacodynamics/Toxicodynamics):

- Effect of molecules/chemicals at biological target *in vivo*
- Perturbation as adverse/therapeutic effect, reversible/ irreversible effects



"The Parallelogram Approach" (Sobels, 1982)

In Vitro - In Vivo Extrapolation (IVIVE)

- HTTK allows *in vitro-in vivo* extrapolation (IVIVE)
– the use of *in vitro* experimental data to predict phenomena *in vivo*.

- IVIVE can be broken down into two components:

- IVIVE-PK/TK

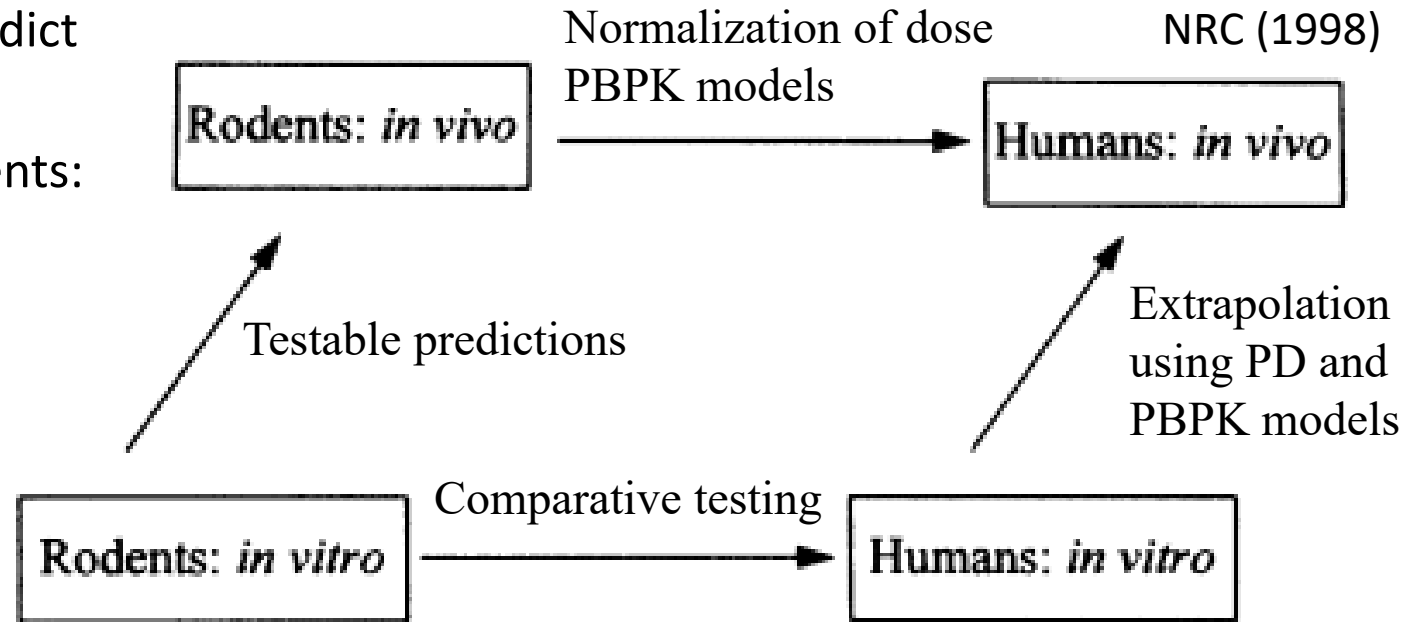
(Pharmacokinetics/Toxicokinetics):

- Fate of molecules/chemicals in body
- Considers absorption, distribution, metabolism, excretion (ADME)
- Can use empirical PK or physiologically-based (PBPK)

- IVIVE-PD/TD (Pharmacodynamics/Toxicodynamics):

- Effect of molecules/chemicals on biological systems
- Perturbation as adverse/therapeutic effect, reversible/ irreversible effects

HTTK only covers toxicokinetic extrapolation



“The Parallelogram Approach” (Sobels, 1982)

IVIVE by Scaling Factor

- There are many approaches to IVIVE, but we choose a relatively simple one:
- We make various assumptions that allow conversion of an *in vitro* concentration $[X]$ (μM) into an **administered equivalent dose** (AED) with units of mg/kg body weight/day:

$$\text{AED} = F_{IVIVE} \times [X]$$

- **AED** is the **external dose rate** that would be needed to **cause a given steady-state plasma concentration**
- F_{IVIVE} is a scaling factor that varies by chemical

IVIVE by Scaling Factor

- For a given chemical, $F_{IVIVE} = 1 / C_{ss,95}$
- $C_{ss,95}$ is the steady-state plasma concentration as the result of a 1 mg/kg/day exposure

$$AED_{95} = \frac{[X]}{C_{ss,95}}$$

- The “95” refers to the upper 95th percentile – due to human variability and measurement uncertainty there are a range of possible C_{ss} values
- All of this assumes that the individuals have enough time to come to “steady-state” with respect to their daily exposures

Don't forget:

$$\mu M = 1000 \frac{1}{MW} \frac{mg}{L}$$

Correlated Monte Carlo
sampling of physiological
model parameters built into
R “httk” package (Pearce et
al., 2017):

Sample CDC National Health
and Nutrition Examination
Survey (NHANES) biometrics
for actual individuals:

Sex
Race/ethnicity
Age
Height
Weight
Serum creatinine



Population simulator for HTTK

Correlated Monte Carlo
sampling of physiological
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R “httk” package (Pearce *et al.*, 2017):

Sample CDC National Health
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Regression equations from literature
(McNally *et al.*, 2014)
(+ residual marginal variability)

(Similar approach used in SimCYP [Jamei *et al.* 2009], GastroPlus,
PopGen [McNally *et al.* 2014], P3M [Price *et al.* 2003], physB [Bosgra *et al.* 2012], etc.)

Population simulator for HTTK

Correlated Monte Carlo
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Sample CDC National Health
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Sex
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Population simulator for HTTK

Predict physiological
quantities

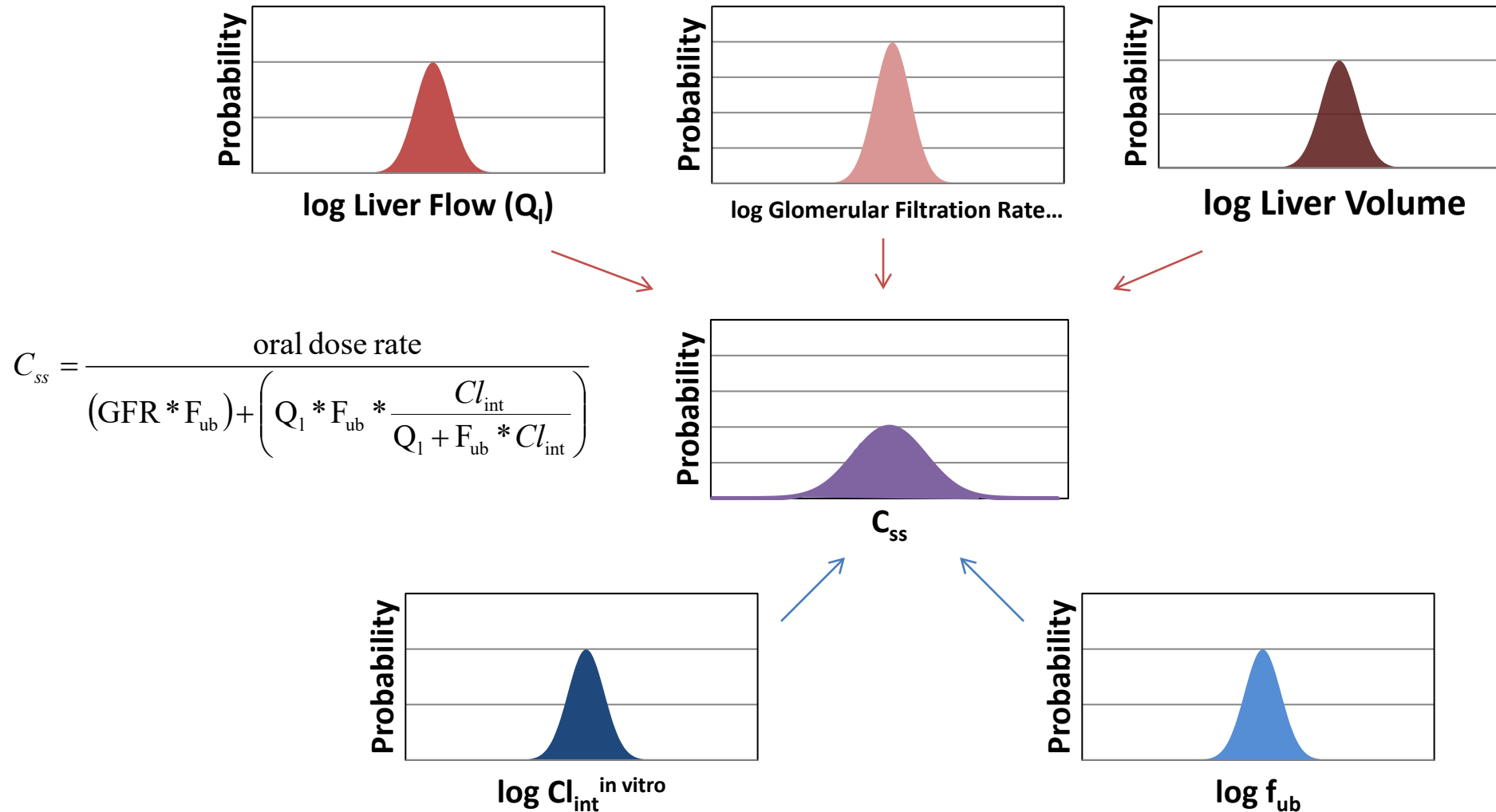
Tissue masses
Tissue blood flows
GFR (kidney function)
Hepatocellularity

Regression equations from literature
(McNally *et al.*, 2014)
(+ residual marginal variability)

(Similar approach used in SimCYP [Jamei *et al.* 2009], GastroPlus,
PopGen [McNally *et al.* 2014], P3M [Price *et al.* 2003], physB [Bosgra *et al.* 2012], etc.)

Monte Carlo Sampling

- Can be used for variability and uncertainty



Variability

Different crayons
have different
colors...



Variability

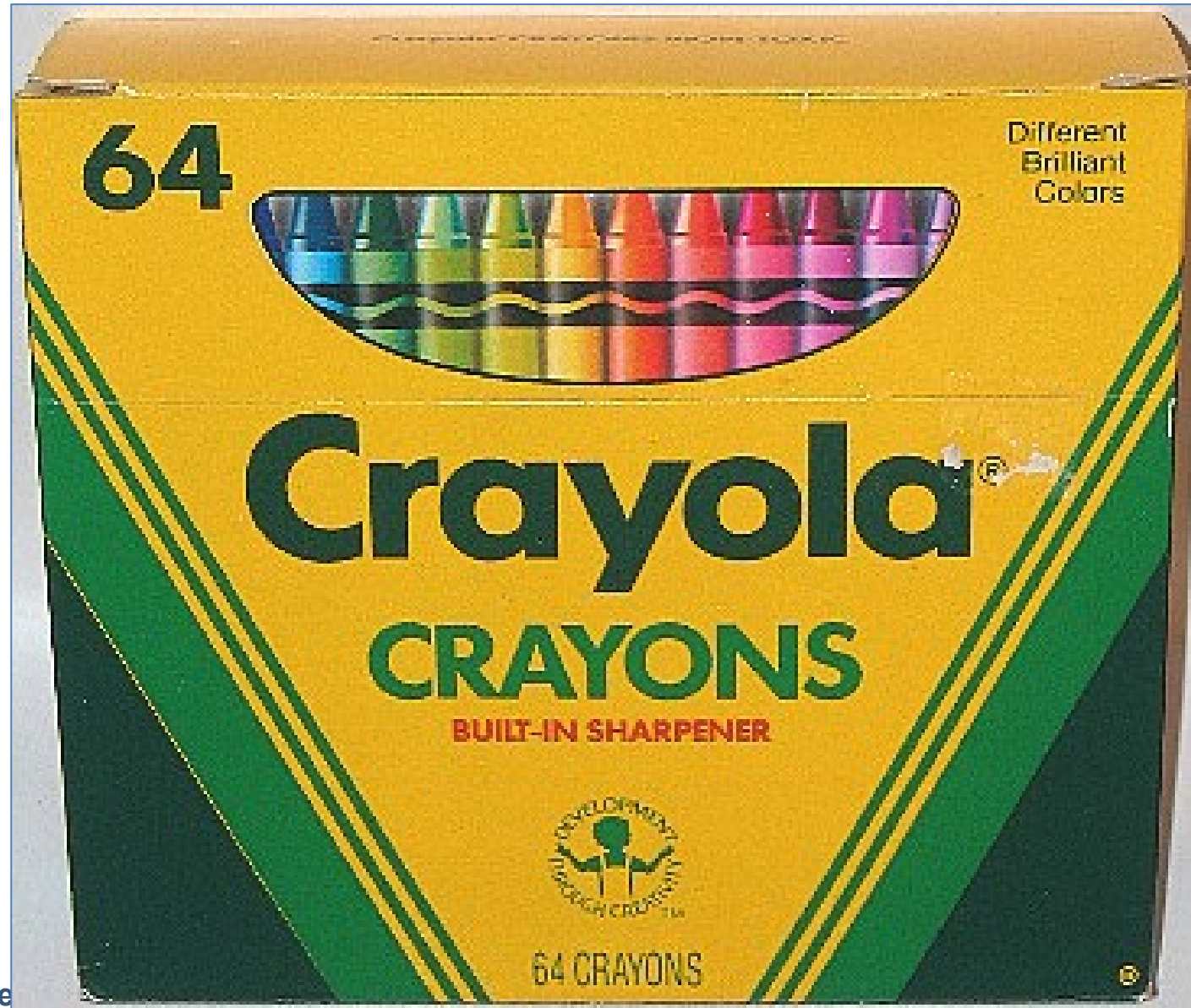
Different crayons
have different
colors, and none
of them are the
“average” color



Uncertainty

Until I open the
box, I don't know
what colors I
have...

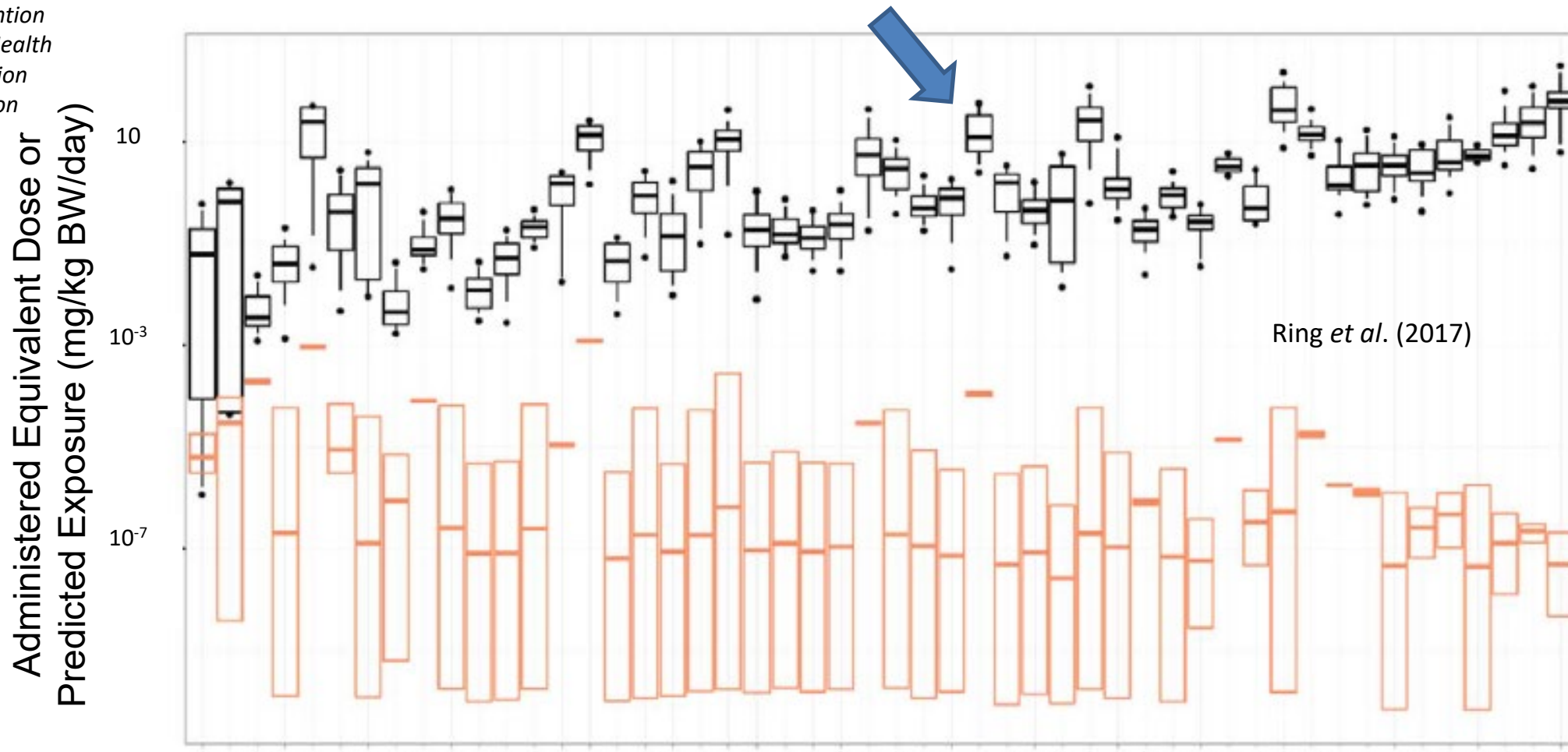
...especially if my
nine-year-old has
been around.



IVIVE Allows Chemical Prioritization

CDC NHANES:
U.S. Centers for
Disease Control
and Prevention
National Health
and Nutrition
Examination
Survey

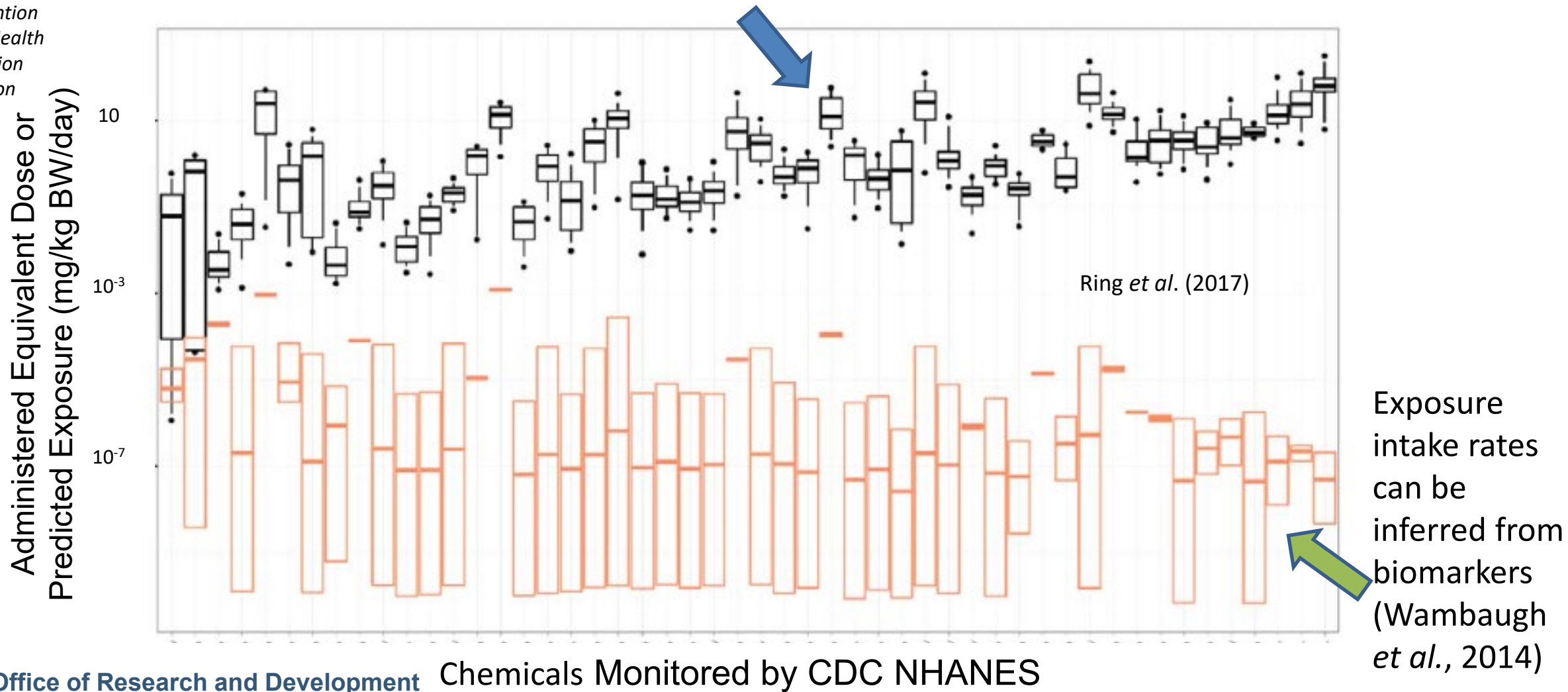
In Vitro Screening + IVIVE can estimate doses needed to cause bioactivity (Wetmore *et al.*, 2015)



IVIVE Allows Chemical Prioritization

CDC NHANES:
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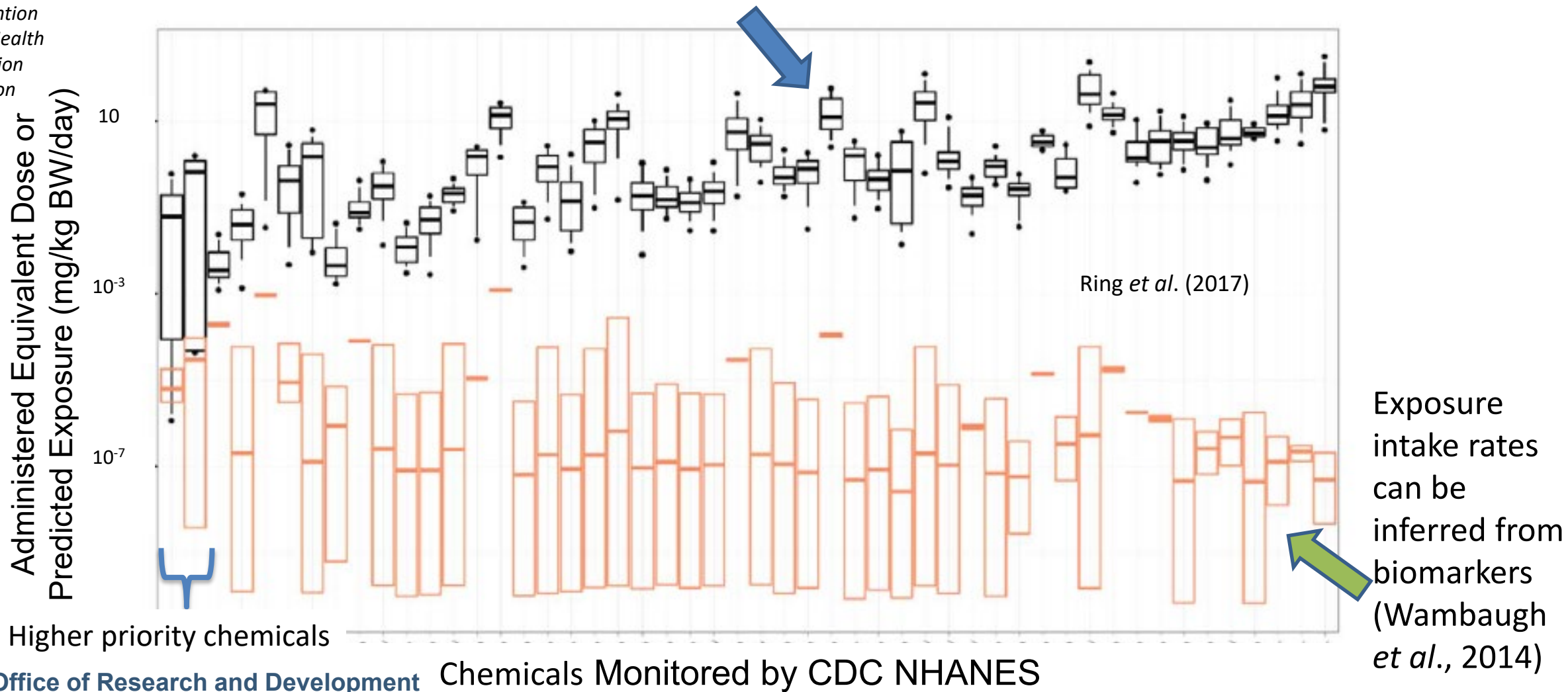
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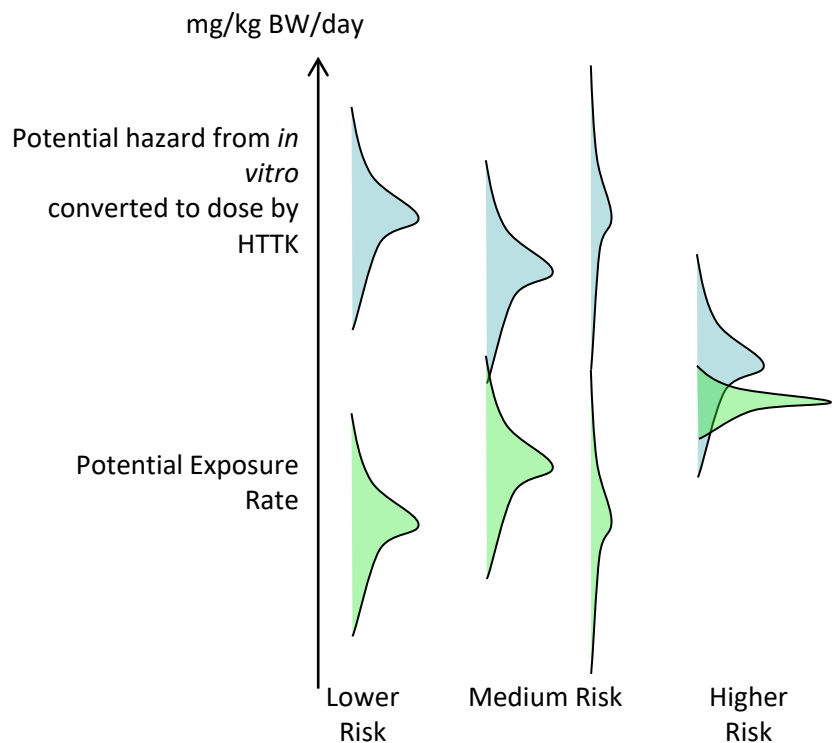
IVIVE Allows Chemical Prioritization

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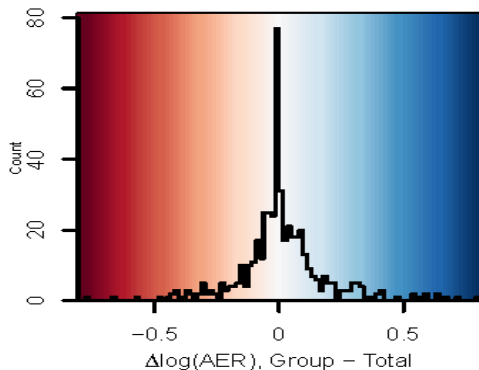
In Vitro Screening + IVIVE can estimate doses needed to cause bioactivity (Wetmore *et al.*, 2015)



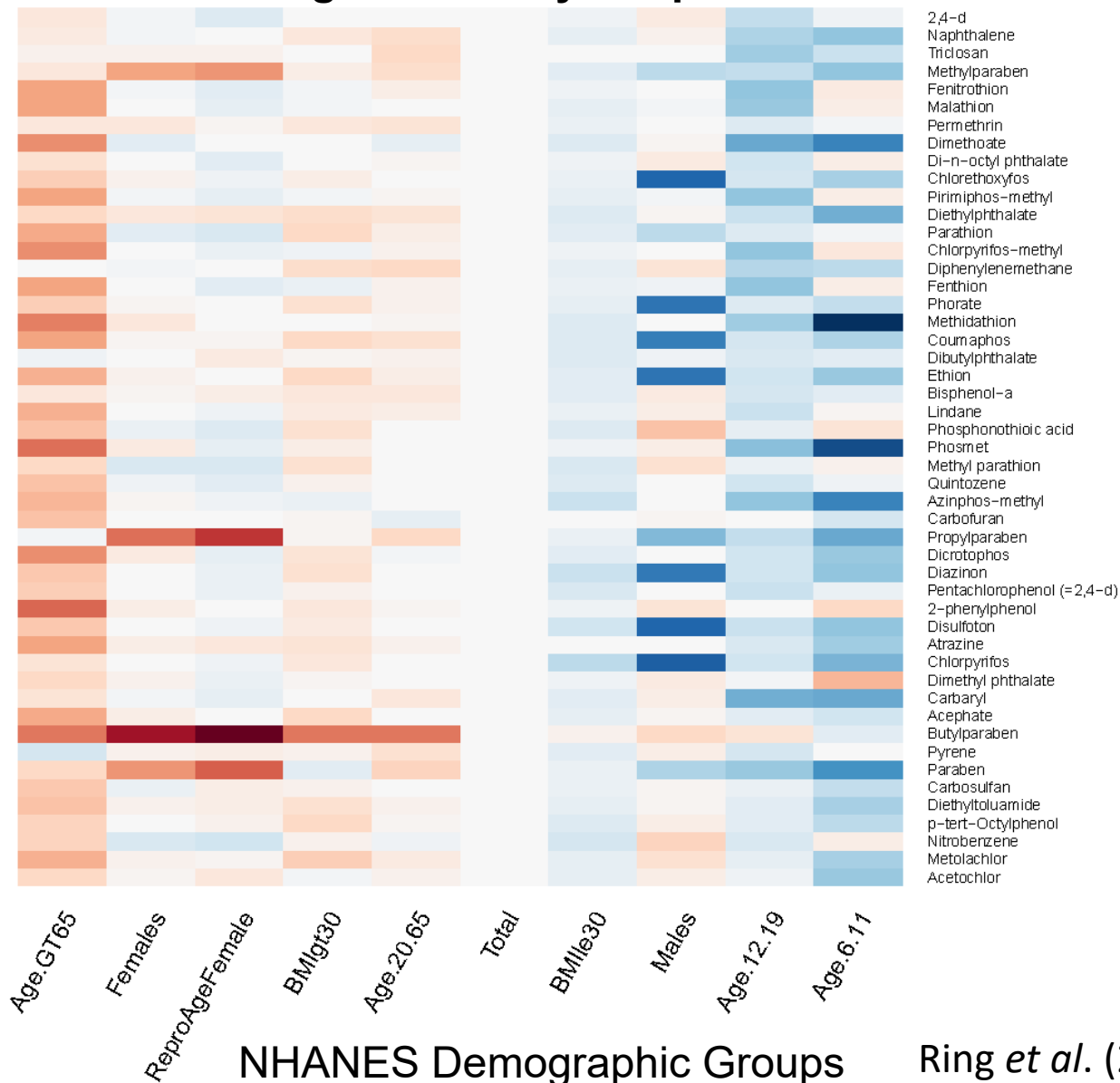
- We use HTTK to calculate margin between bioactivity and exposure for specific populations



Life-stage and Demographic Specific Predictions



Change in Activity : Exposure Ratio



NHANES Chemicals

NHANES Demographic Groups

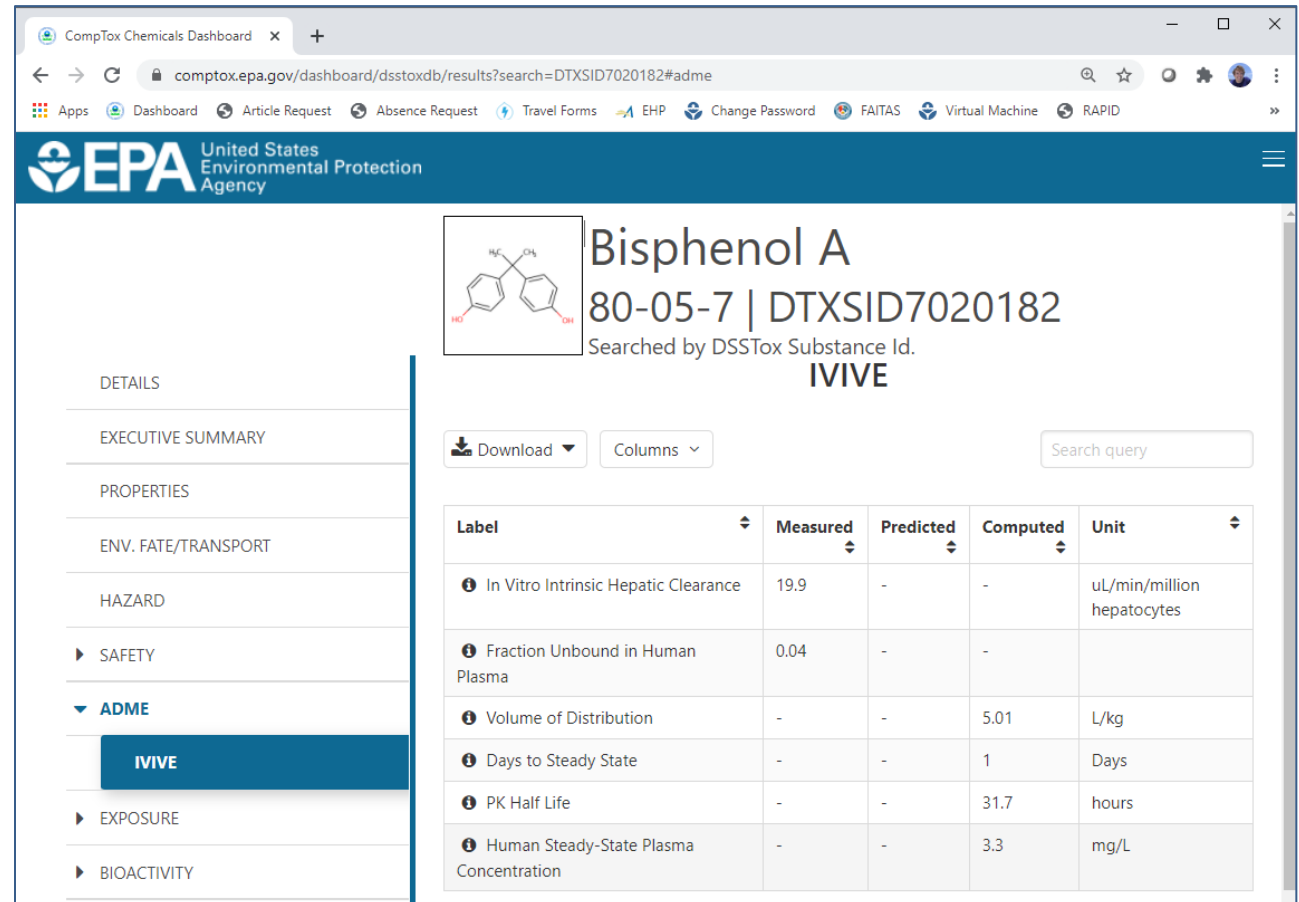
Ring *et al.* (2017)

HTTK on the CompTox Chemicals Dashboard

- The CompTox Chemicals Dashboard provides $C_{ss,95}$ values for >1000 chemicals

<https://comptox.epa.gov/dashboard/>

- We use EPA's R package "httk" to provide IVIVE predictions
- The value reported is calculated assuming a 1 mg/kg/day dose rate
- We give the upper 95th percentile of the calculated values based on a Monte Carlo simulation of human variability and uncertainty



The screenshot shows the CompTox Chemicals Dashboard for Bisphenol A (DTXSID7020182). The dashboard includes a sidebar with navigation options: DETAILS, EXECUTIVE SUMMARY, PROPERTIES, ENV. FATE/TRANSPORT, HAZARD, SAFETY, ADME (selected), EXPOSURE, and BIOACTIVITY. The main content area displays the chemical structure of Bisphenol A, its name, and its DTXSID. Below this, there is a table of IVIVE (In Vitro to In Vivo Extrapolation) results. The table has columns for Label, Measured, Predicted, Computed, and Unit. The data rows show various pharmacokinetic parameters and their predicted values.

Label	Measured	Predicted	Computed	Unit
In Vitro Intrinsic Hepatic Clearance	19.9	-	-	uL/min/million hepatocytes
Fraction Unbound in Human Plasma	0.04	-	-	
Volume of Distribution	-	-	5.01	L/kg
Days to Steady State	-	-	1	Days
PK Half Life	-	-	31.7	hours
Human Steady-State Plasma Concentration	-	-	3.3	mg/L

Conclusions

- HTTK allows dosimetric adjustment of high-throughput screening (HTS) data
 - Thousands of chemicals
 - Open source, free, and evaluated software
- HTTK accounts for human population diversity using biometrics from the CDC NHANES to predict toxicokinetic model parameters
 - Variability is simulated using a Monte Carlo approach
- Breen et al. (in preparation) updates R package “httk” to the most recent three NHANES cohorts and adds children under the age of 6
- Toxicodynamic variability is not included
- HTTK in vitro parameters are generated from pooled adult tissues



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

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Human Variability

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