

The High Throughput Toxicokinetic (HTTK) R Package

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Office of Research and Development

U.S. Environmental Protection Agency

**Computational Toxicology
Community of Practice Webinar**

June 27, 2019

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**Oral
Absorption**

Dustin Kapraun
Richard Judson
Annie Lumen (FDA)

Mark Sfeir

Package Tsar

**Human
Gestation**



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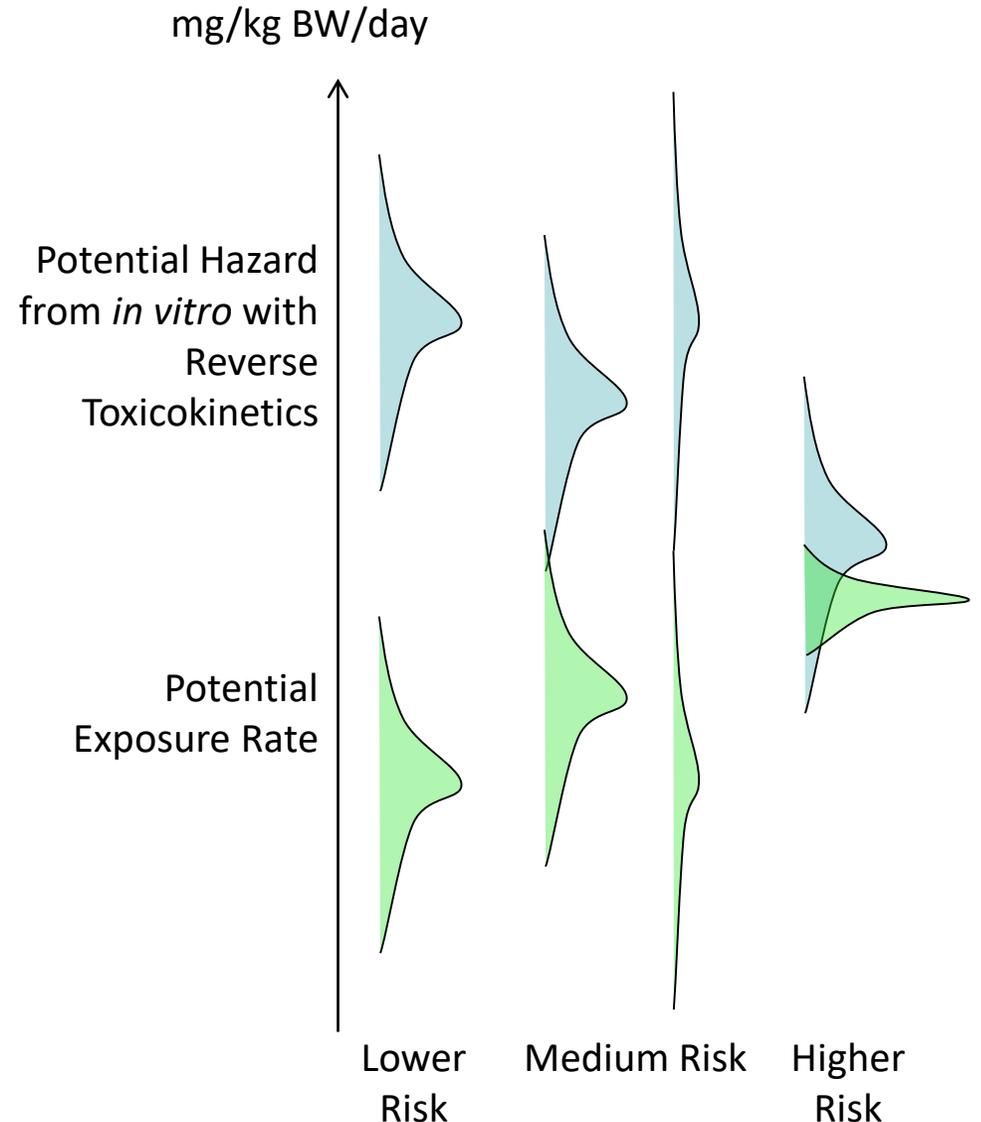
Marina Evans

Beate Nicol (Unilever)

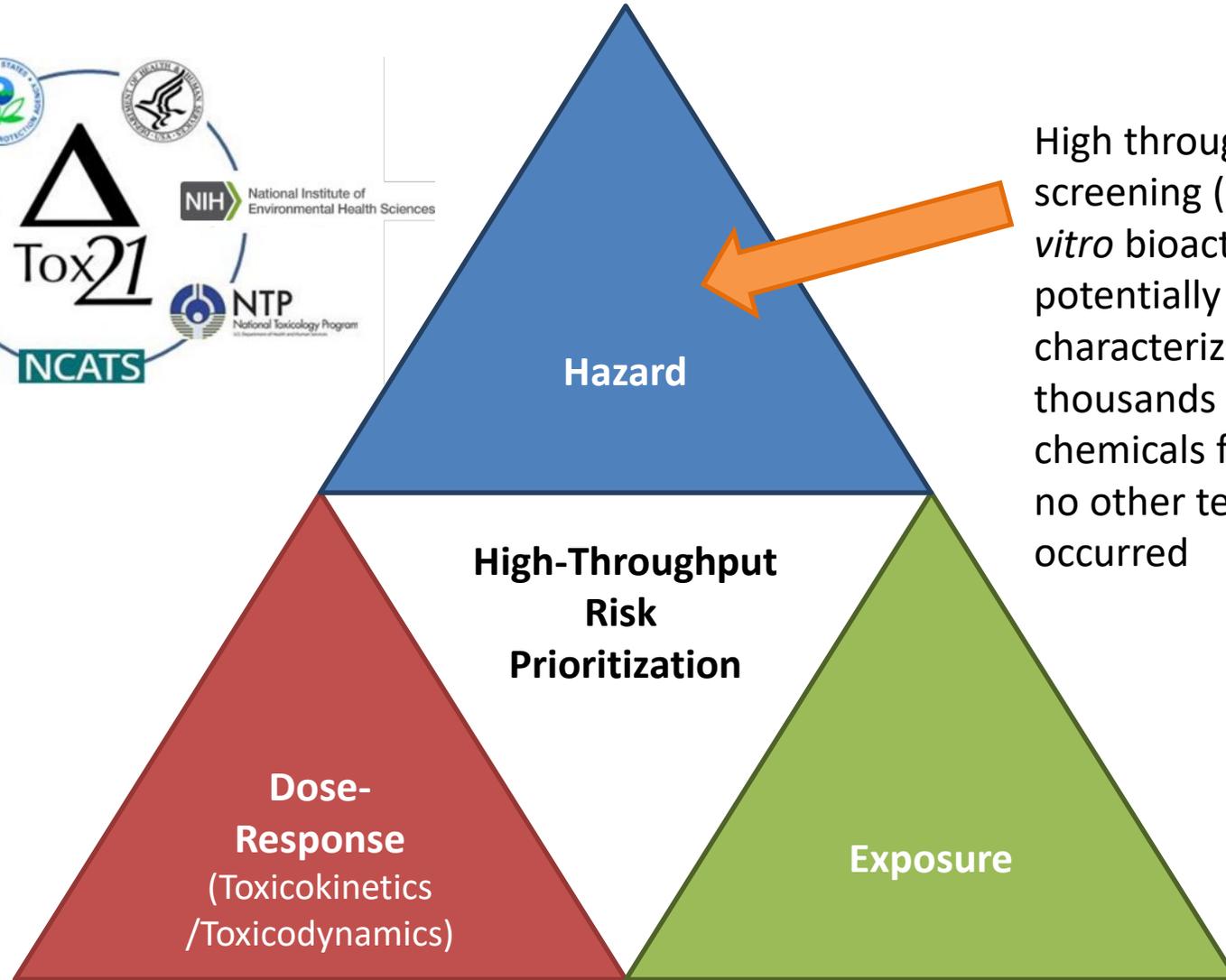
Tom Moxon (Unilever)

Chemical Risk = Hazard x Exposure

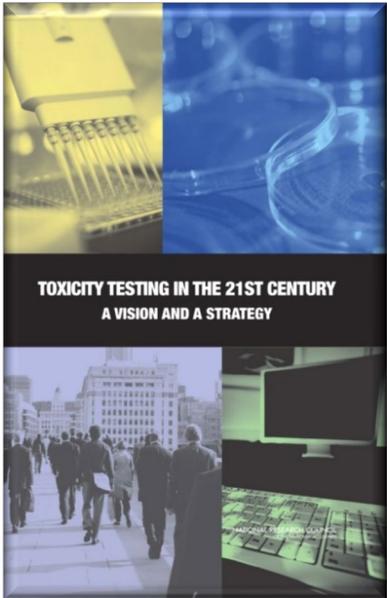
- The U.S. National Research Council (1983) identified chemical risk as a function of both inherent hazard and exposure
- To address the thousands of chemicals in commerce and the environment, we need new approach methodologies (NAMs) that can inform prioritization of chemicals most worthy of additional study
- High throughput risk prioritization needs:
 1. High throughput hazard characterization (Dix et al., 2007, Collins et al., 2008)
 2. High throughput exposure forecasts (Wambaugh et al., 2013, 2014)
 3. High throughput toxicokinetics (i.e., dose-response relationship) linking hazard and exposure (Wetmore et al., 2012, 2015)



High-Throughput Risk Prioritization



High throughput screening (HTS) for *in vitro* bioactivity potentially allows characterization of thousands of chemicals for which no other testing has occurred



NRC (2007)

In Vitro - In Vivo Extrapolation (IVIVE)

Utilization of *in vitro* experimental data to predict phenomena *in vivo*

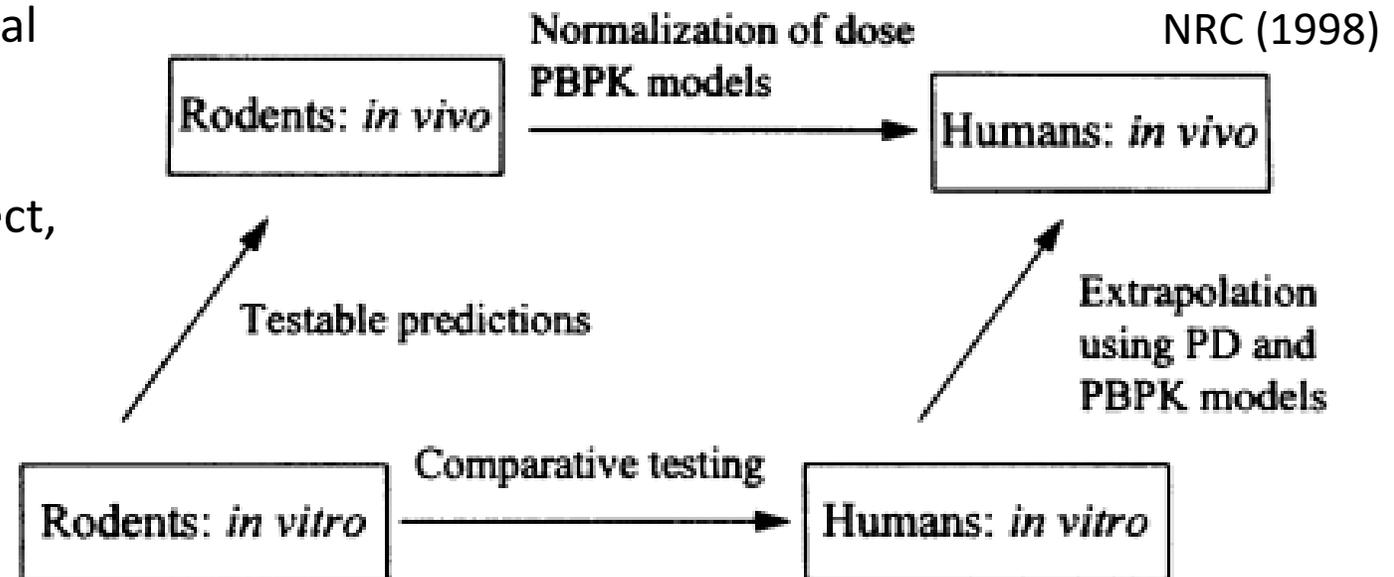
- IVIVE-PK/TK (**Pharmacokinetics/Toxicokinetics**):

- Fate of molecules/chemicals in body
- Considers absorption, distribution, metabolism, excretion (ADME)
- Uses empirical PK and physiologically-based (PBPK) modeling

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

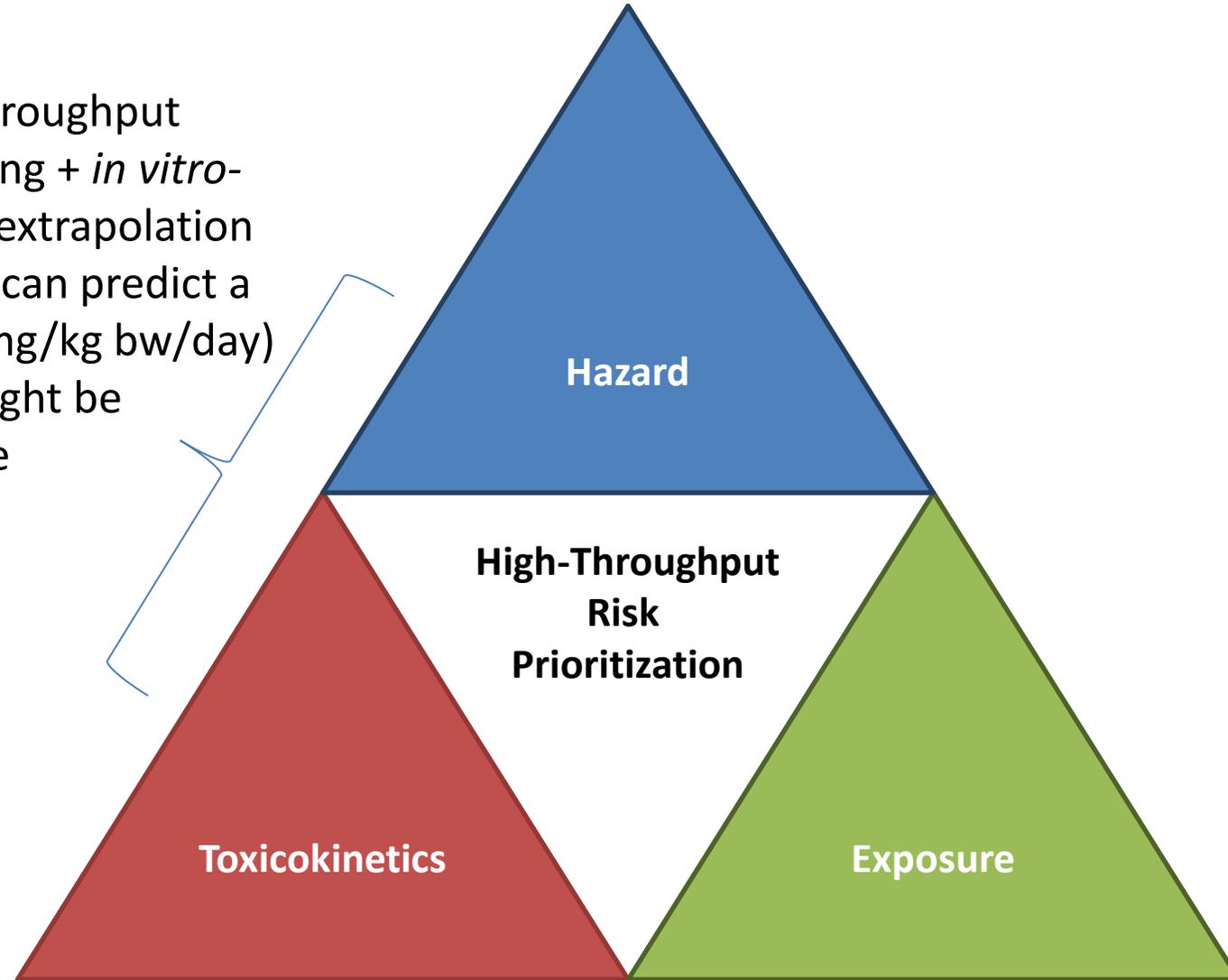
- Effect of molecules/chemicals at biological target *in vivo*
- Assay design/selection important
- Perturbation as adverse/therapeutic effect, reversible/ irreversible effects

- Both contribute to *in vivo* effect prediction



New Exposure Data and Models

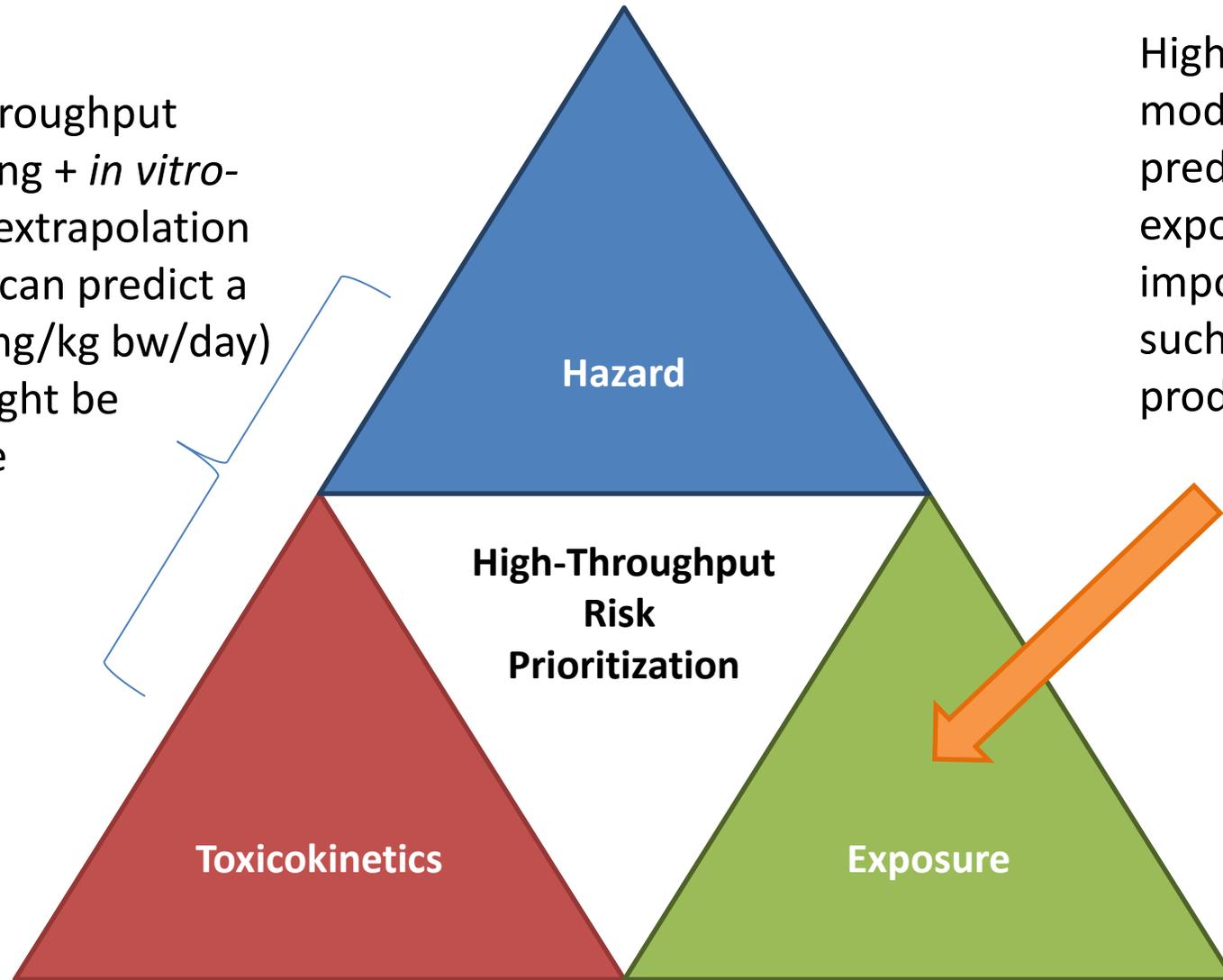
High throughput screening + *in vitro*-*in vivo* extrapolation (IVIVE) can predict a dose (mg/kg bw/day) that might be adverse



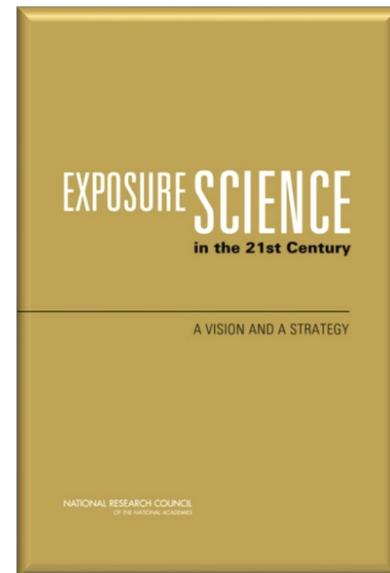
Wetmore et al. (2012, 2015)

New Exposure Data and Models

High throughput screening + *in vitro*-*in vivo* extrapolation (IVIVE) can predict a dose (mg/kg bw/day) that might be adverse



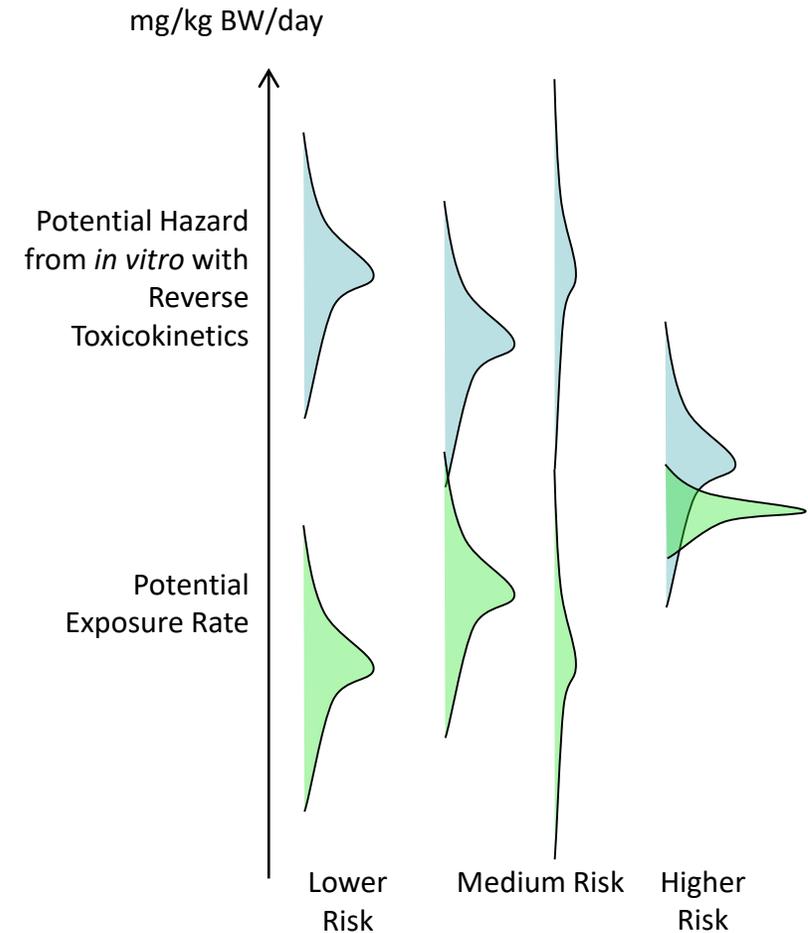
High throughput models exist to make predictions of exposure via specific, important pathways such as residential product use and diet



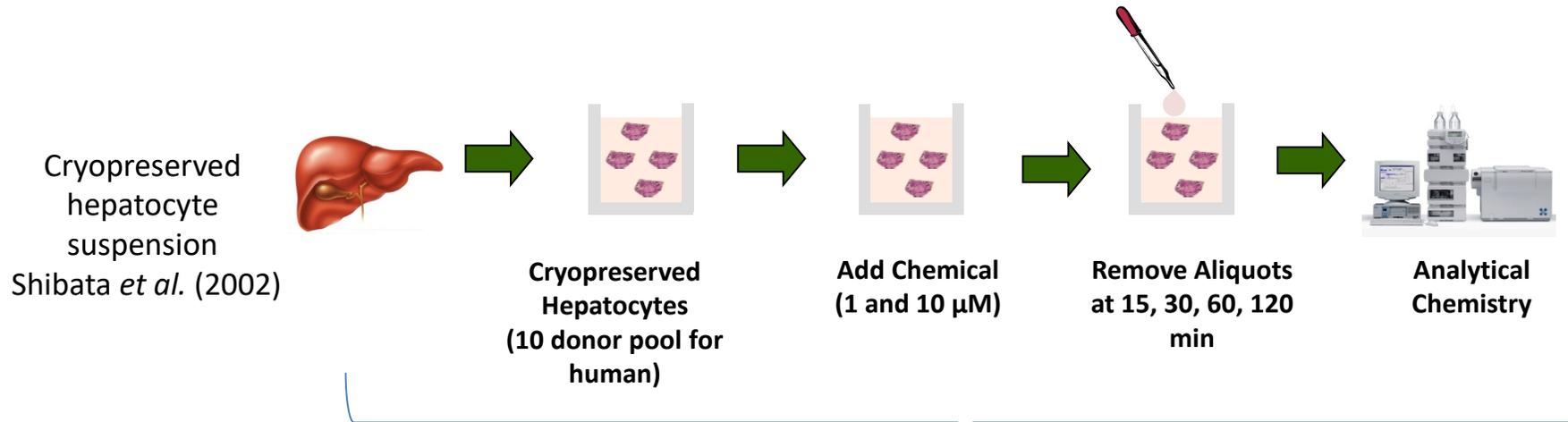
NRC (2012)

High Throughput Toxicokinetics (HTTK)

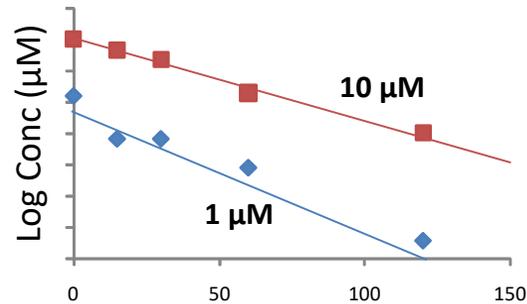
- **Most chemicals do not have TK data**
- In order to address greater numbers of chemicals we collect *in vitro*, high throughput toxicokinetic (HTTK) data (Rotroff et al., 2010, Wetmore et al., 2012, 2015)
- HTTK methods have been used by the pharmaceutical industry to determine range of efficacious 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 (*i.e.*, *in vitro-in vivo* extrapolation, or **IVIVE**) (e.g., Wetmore et al., 2015)
- **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 Data for HTTK



The rate of disappearance of parent compound (slope of line) is the **hepatic clearance** ($\mu\text{L}/\text{min}/10^6$ hepatocytes)

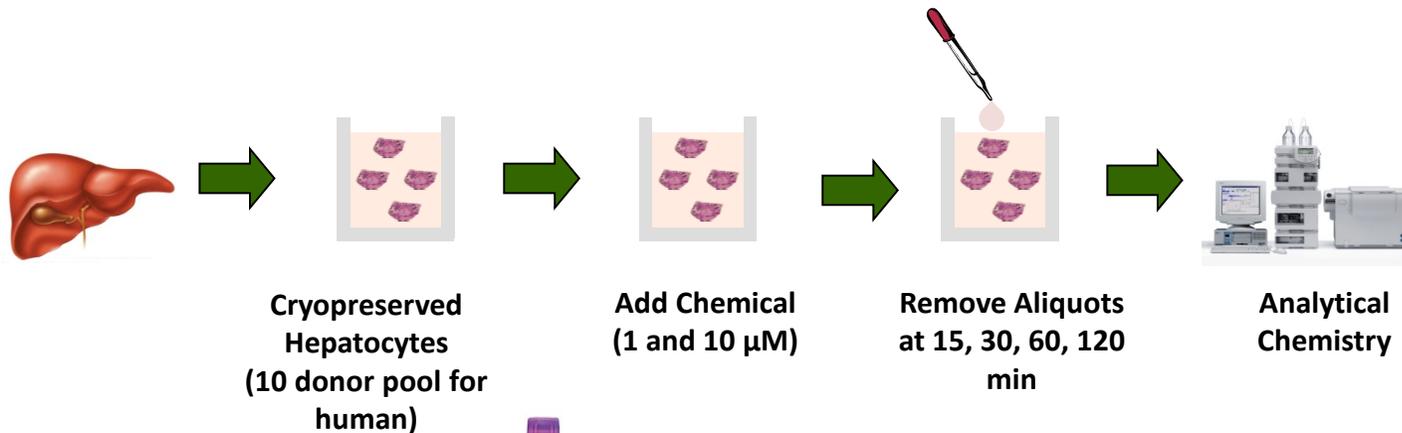


We perform the assay at 1 and 10 μM to check for saturation of metabolizing enzymes.

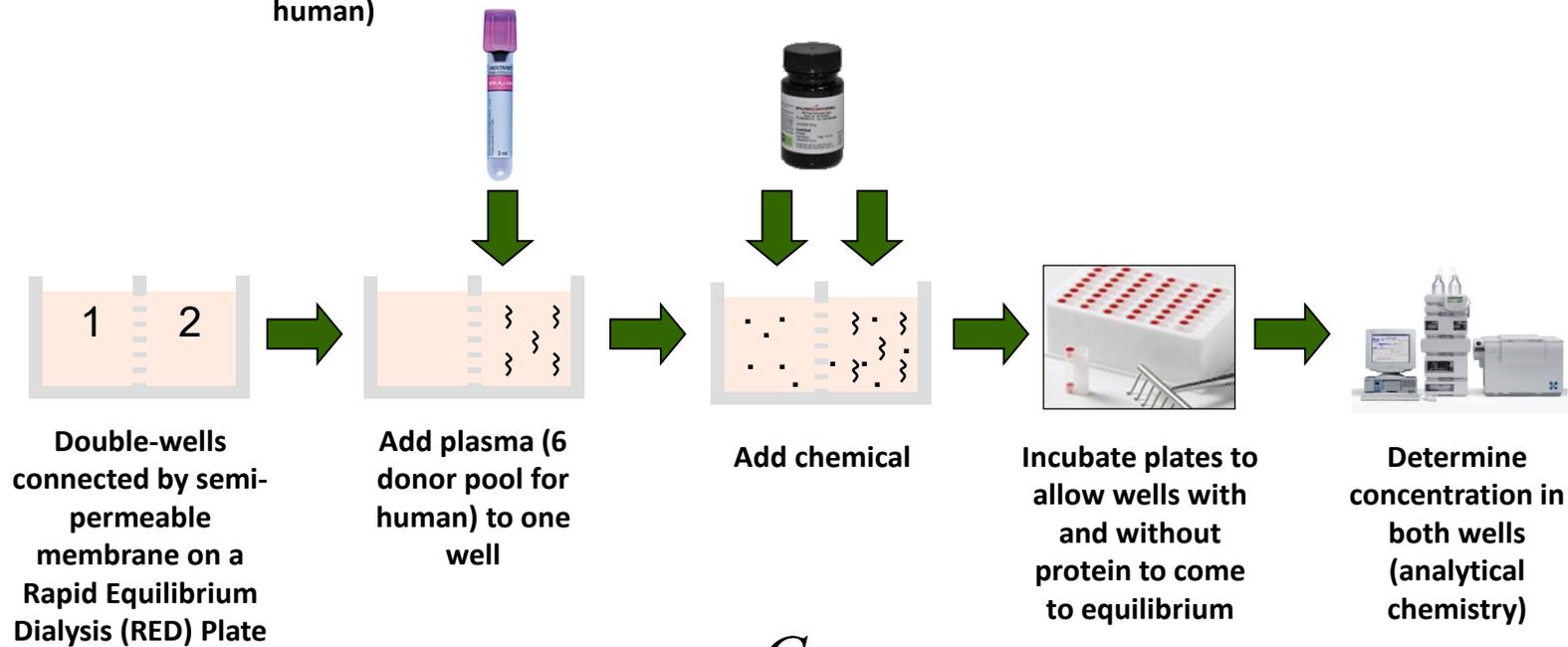
- **Most chemicals do not have TK data** – we use *in vitro* HTTK methods adapted from pharma to fill gaps
- In drug development, HTTK methods allow IVIVE to estimate therapeutic doses for clinical studies – predicted concentrations are typically on the order of values measured in clinical trials (Wang, 2010)

In Vitro Data for HTTK

Cryopreserved
hepatocyte
suspension
Shibata *et al.* (2002)



Rapid Equilibrium
Dialysis (RED)
Waters *et al.* (2008)

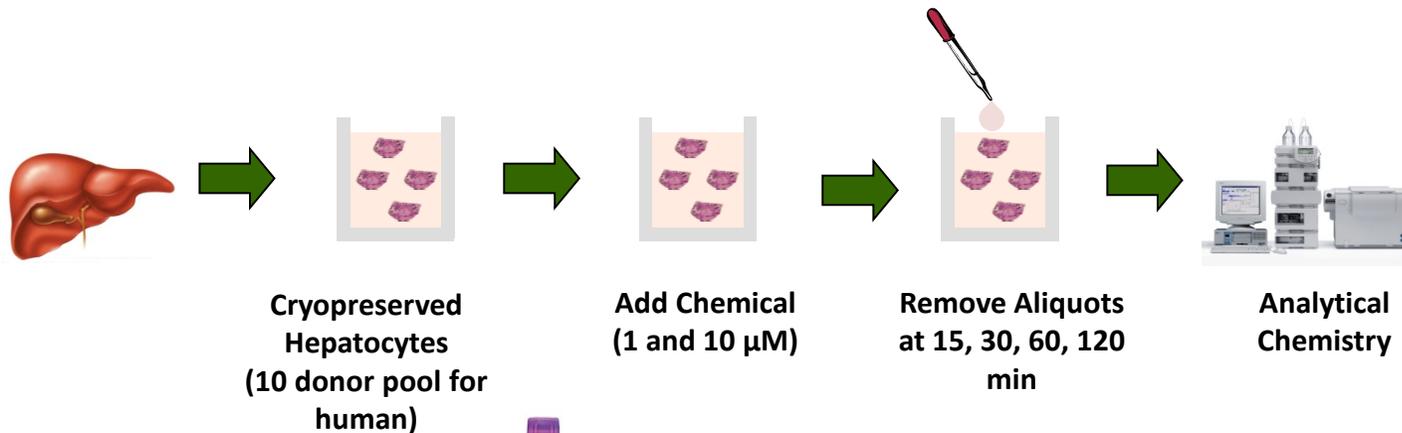


$$F_{ub,p} = \frac{C_{well1}}{C_{well2}}$$

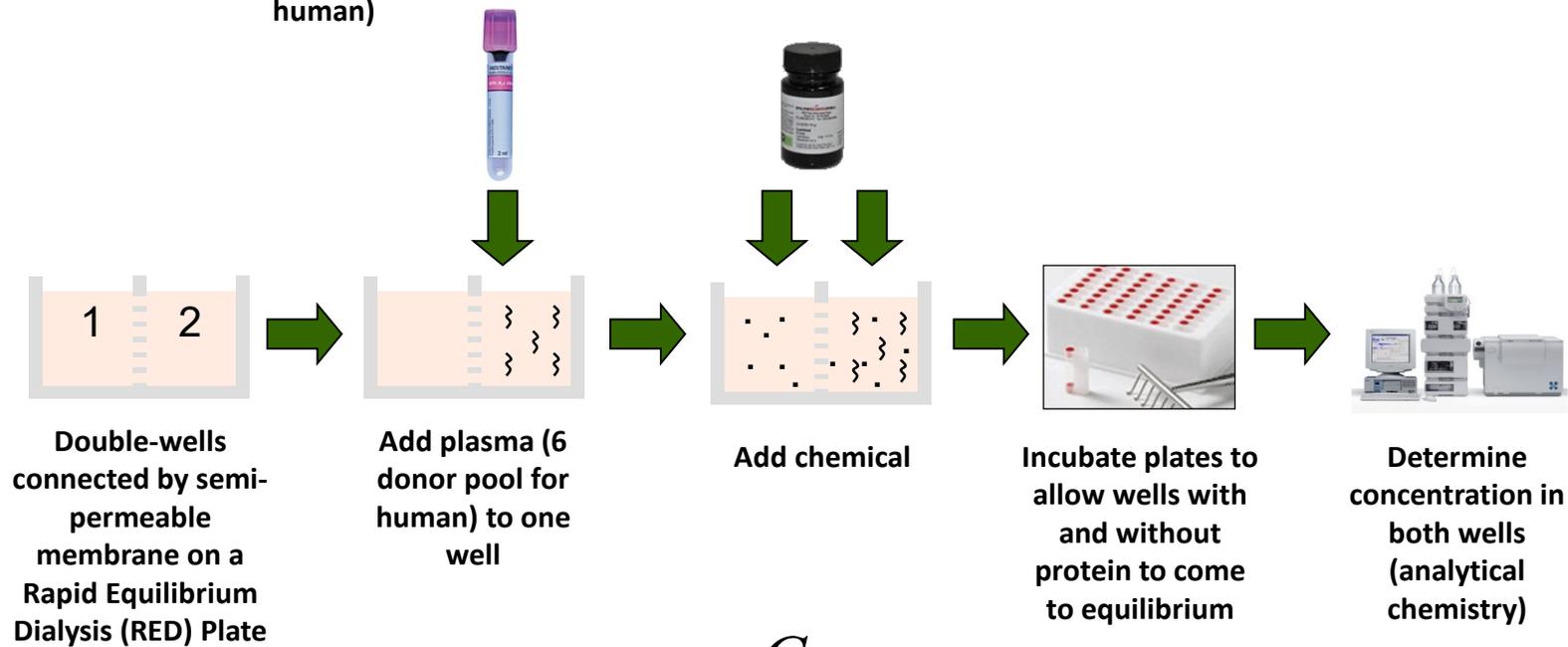
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In Vitro Data for HTTK

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Waters *et al.* (2008)



$$F_{ub,p} = \frac{C_{well1}}{C_{well2}}$$

- **Most chemicals do not have TK data** – we use *in vitro* HTTK methods adapted from pharma to fill gaps

- Environmental chemicals:

Rotroff *et al.* (2010)
35 chemicals

Wetmore *et al.* (2012)
+204 chemicals

Wetmore *et al.* (2015)
+163 chemicals

Wambaugh *et al.* (submitted)
+389 chemicals

Simple Model for Steady-State Plasma Concentration (C_{ss})

$$C_{ss} = \frac{\textit{oral dose rate}}{\underbrace{(GFR * f_{up})}_{\text{Passive Renal Clearance}} + \underbrace{\left(Q_l * f_{up} * \frac{Cl_{int}}{Q_l + f_{up} * Cl_{int}} \right)}_{\text{Hepatic Metabolism}}}$$

Wilkinson and Shand (1975)

Passive Renal Clearance
 (GFR: Glomerular filtration rate
 f_{up} : fraction unbound in plasma)

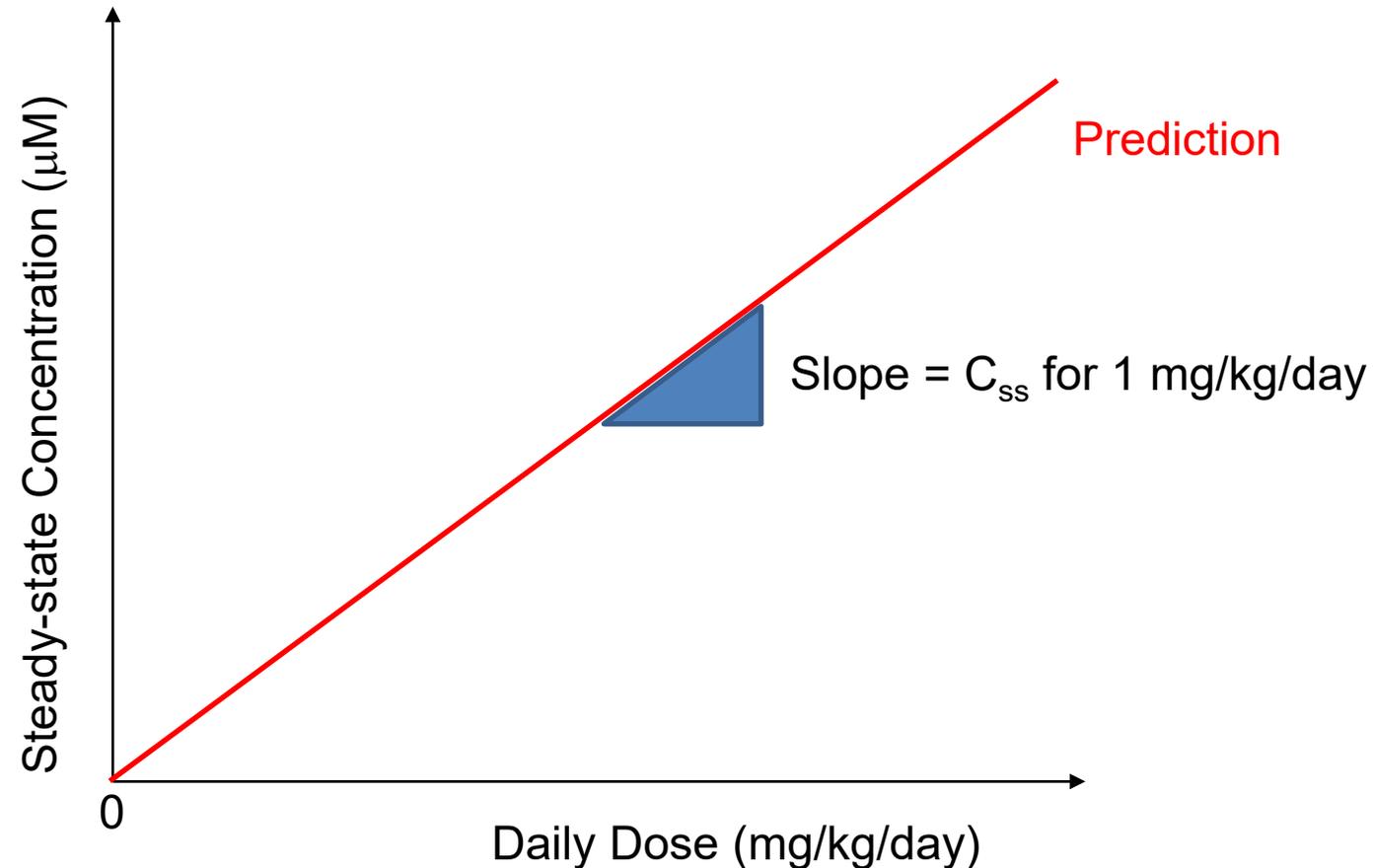
Hepatic Metabolism
 (Cl_{int} : Scaled hepatic clearance
 Q_l : Blood flow to liver)

Assume that Steady-State is Linear with Dose

$$C_{ss} = \frac{\text{oral dose rate}}{(GFR * f_{up}) + \left(Q_l * f_{up} * \frac{Cl_{int}}{Q_l + f_{up} * Cl_{int}} \right)}$$

Assume that Steady-State is Linear with Dose

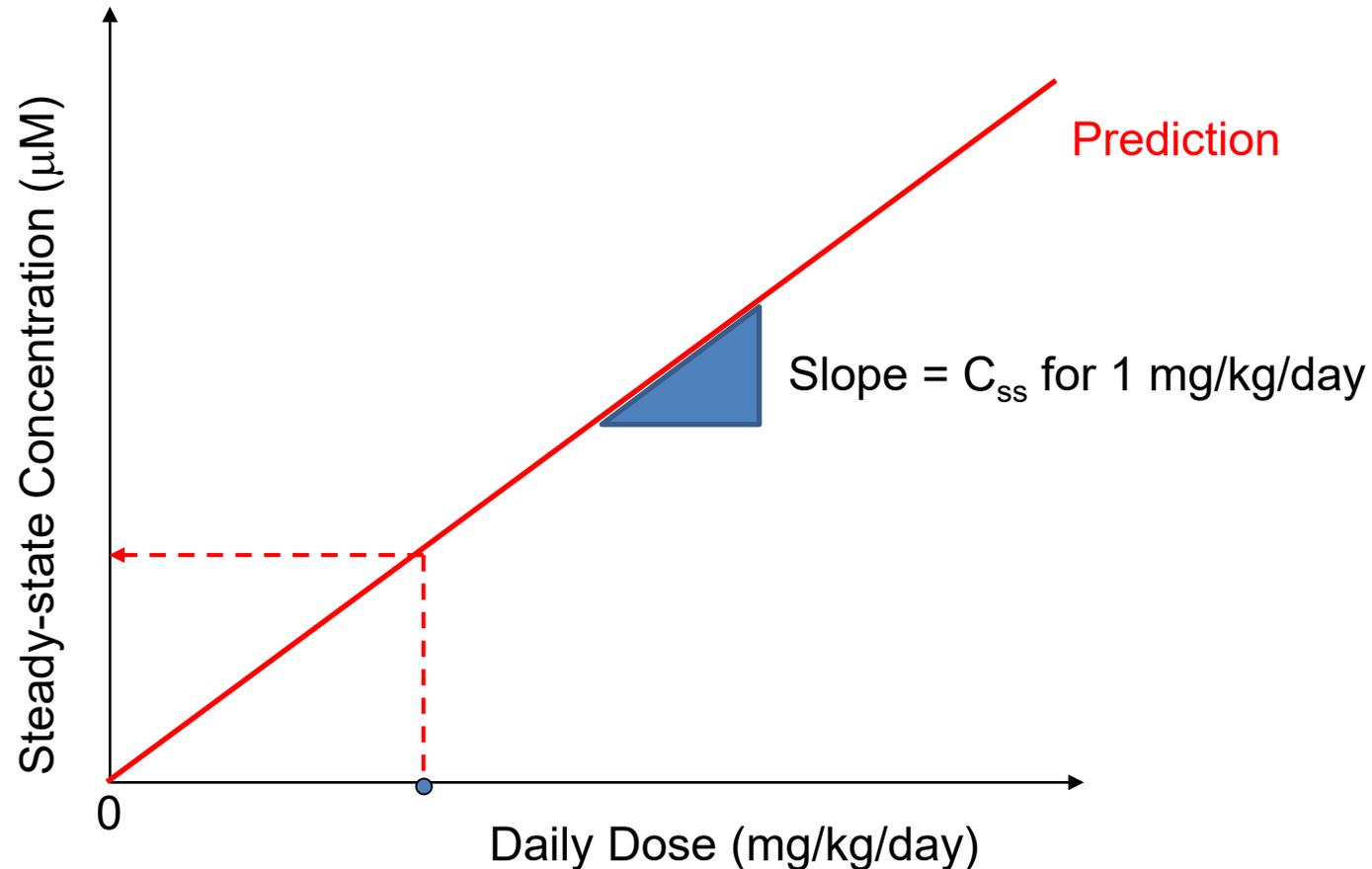
$$C_{ss} = \frac{\text{oral dose rate}}{(GFR * f_{up}) + \left(Q_l * f_{up} * \frac{Cl_{int}}{Q_l + f_{up} * Cl_{int}} \right)}$$



- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

Assume that Steady-State is Linear with Dose

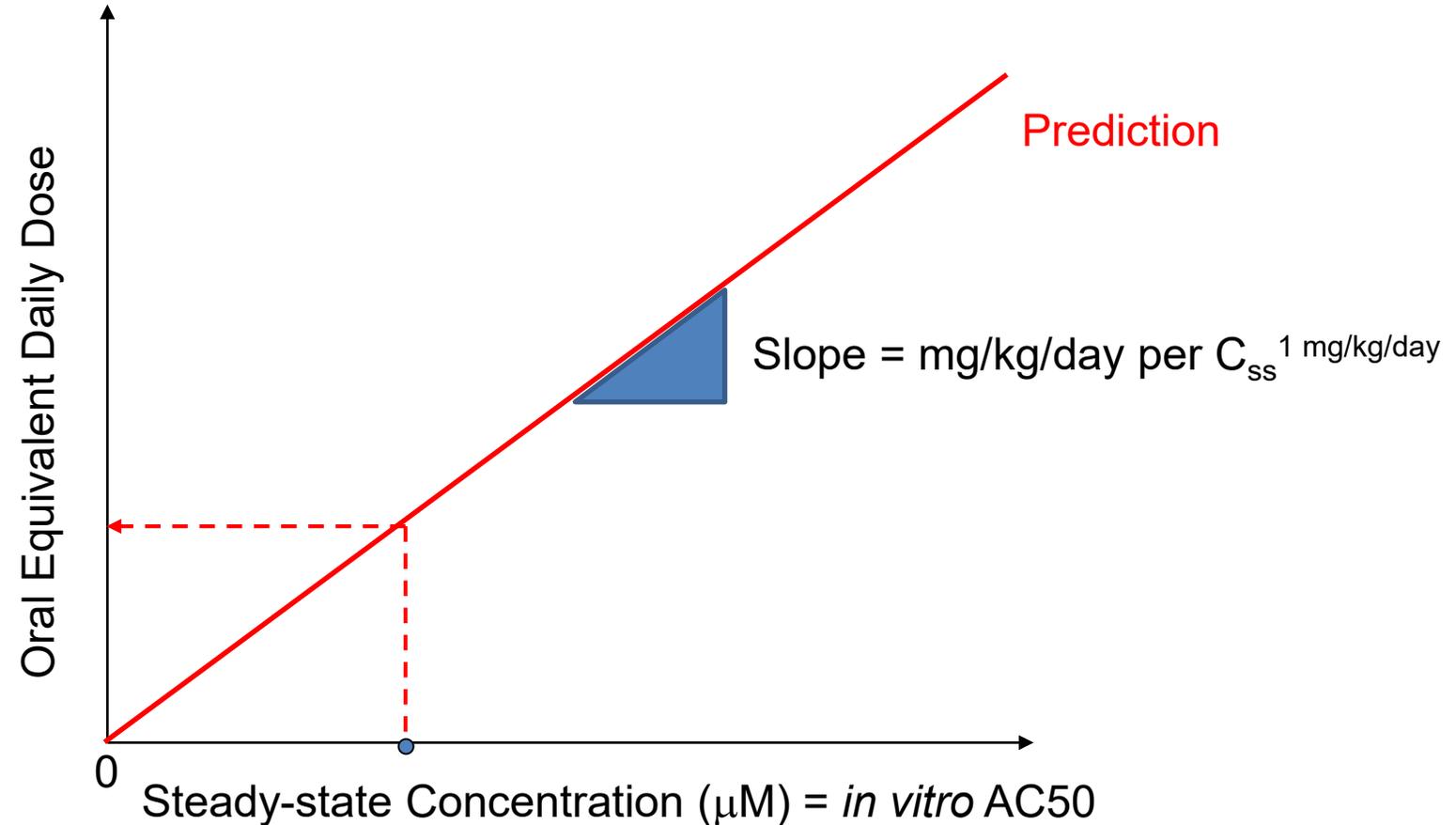
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- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

HTTK Allows Steady-State In Vitro-In Vivo Extrapolation (IVIVE)

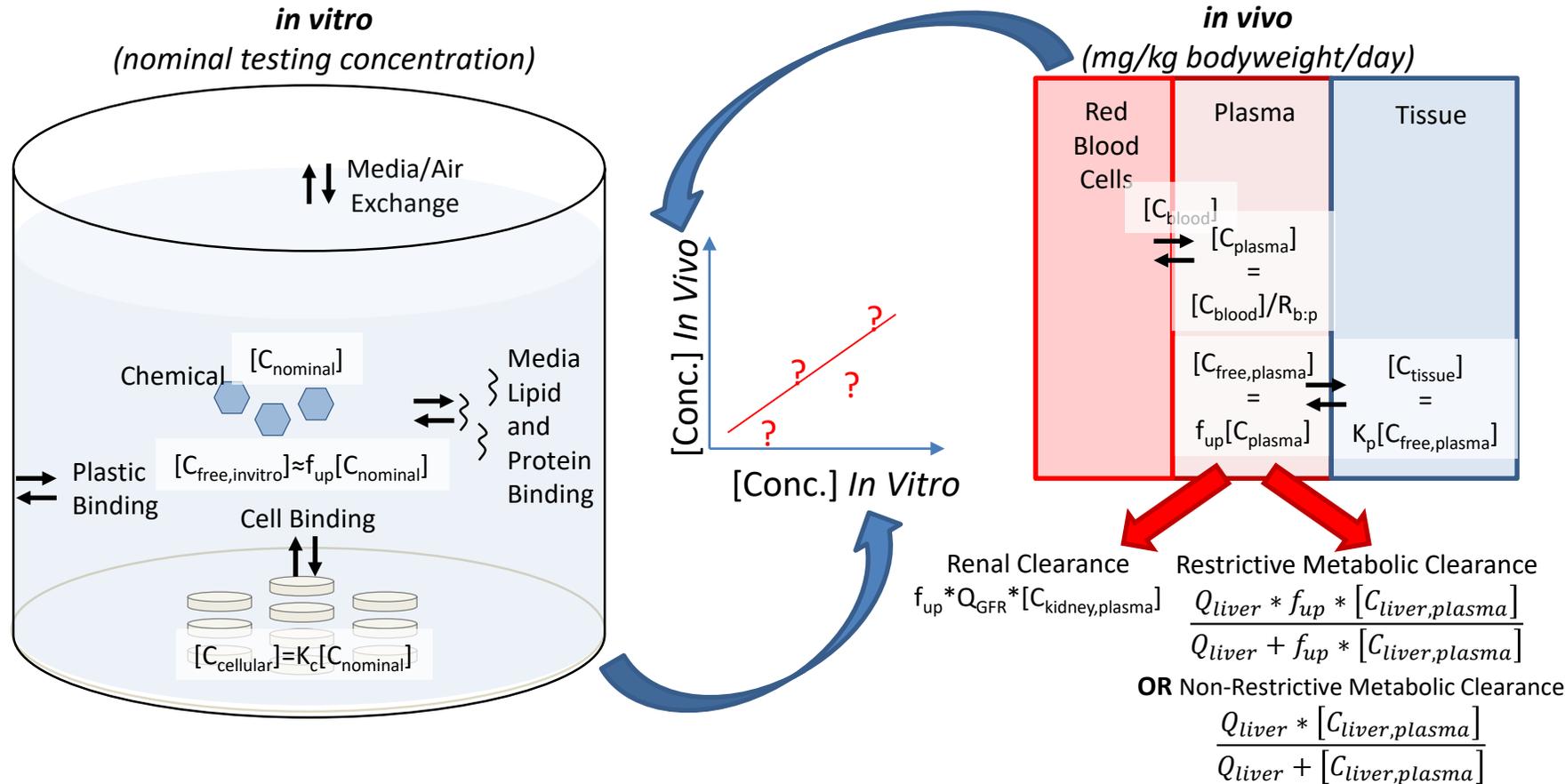
$$C_{ss} = \frac{\text{oral dose rate}}{(GFR * f_{up}) + \left(Q_l * f_{up} * \frac{Cl_{int}}{Q_l + f_{up} * Cl_{int}} \right)}$$



- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

High-Throughput Toxicokinetics (HTTK) for *In Vitro-In Vivo* Extrapolation (IVIVE)

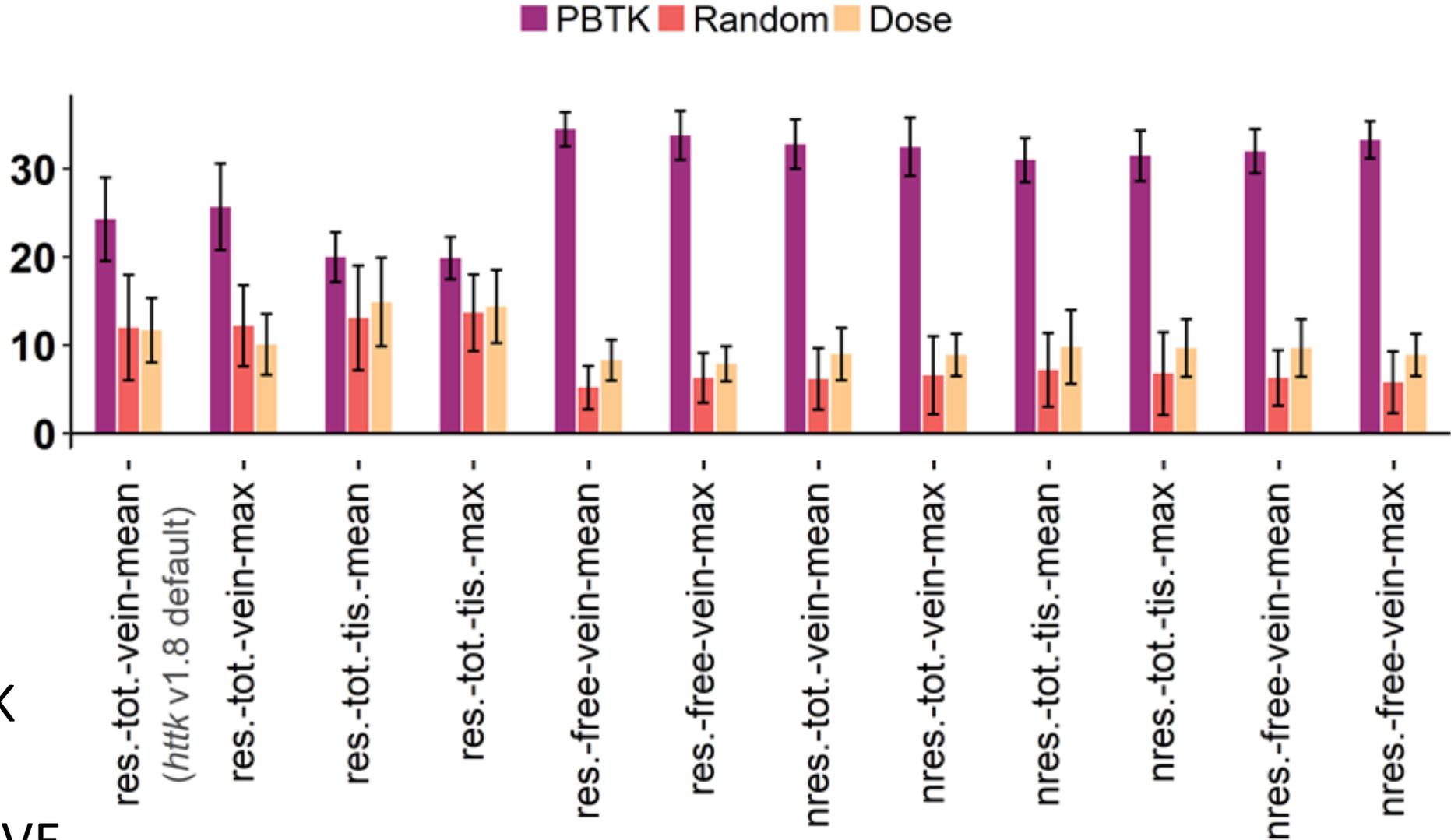
Using the generic HTTK physiologically based toxicokinetics model to inform IVIVE...



Selecting the appropriate *in vitro* and *in vivo* concentrations for extrapolation

Optimizing HTKK-based IVIVE

Number of times model selected as best for predicting *in vivo* endpoints



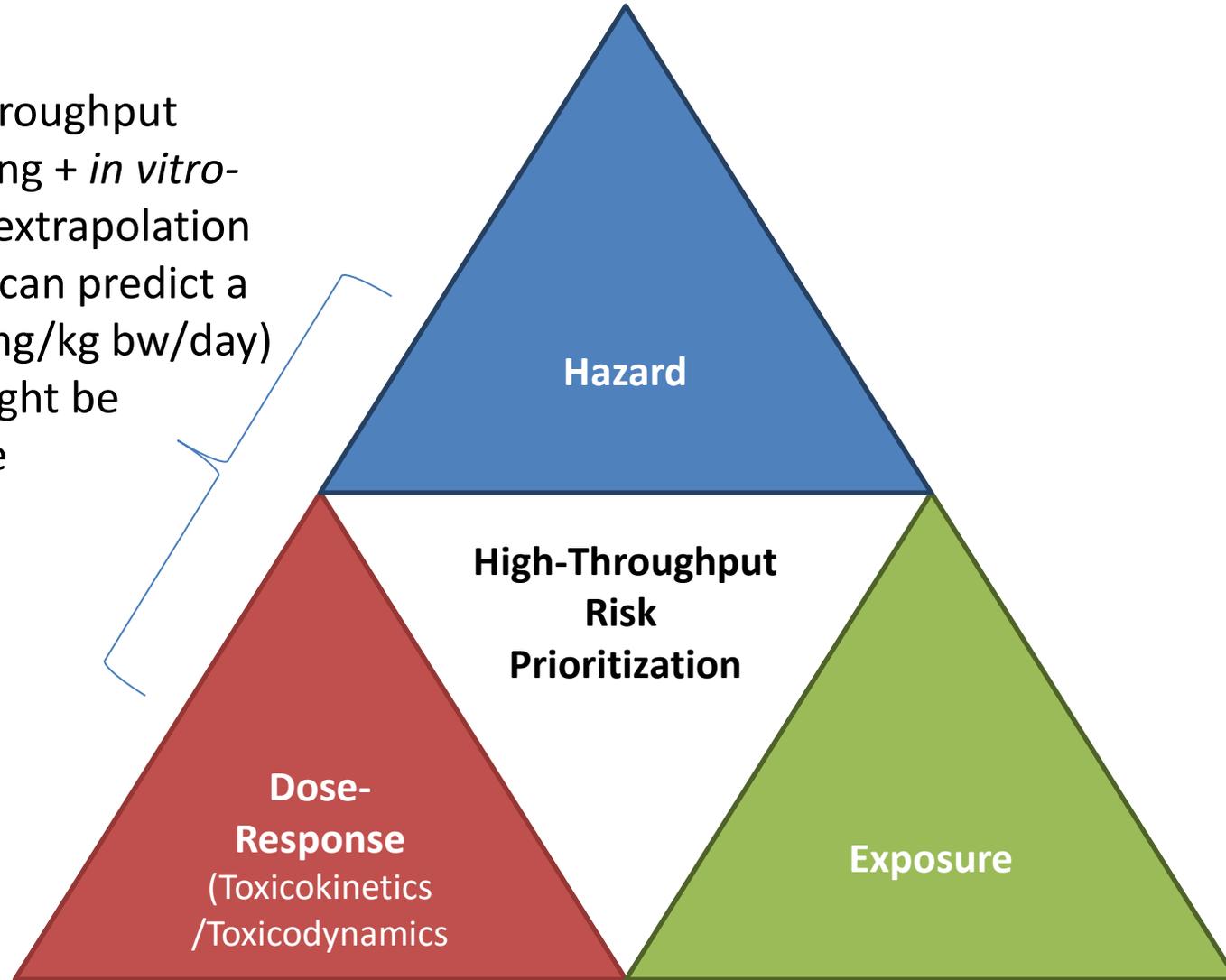
Using PBTK
Models

Improves IVIVE

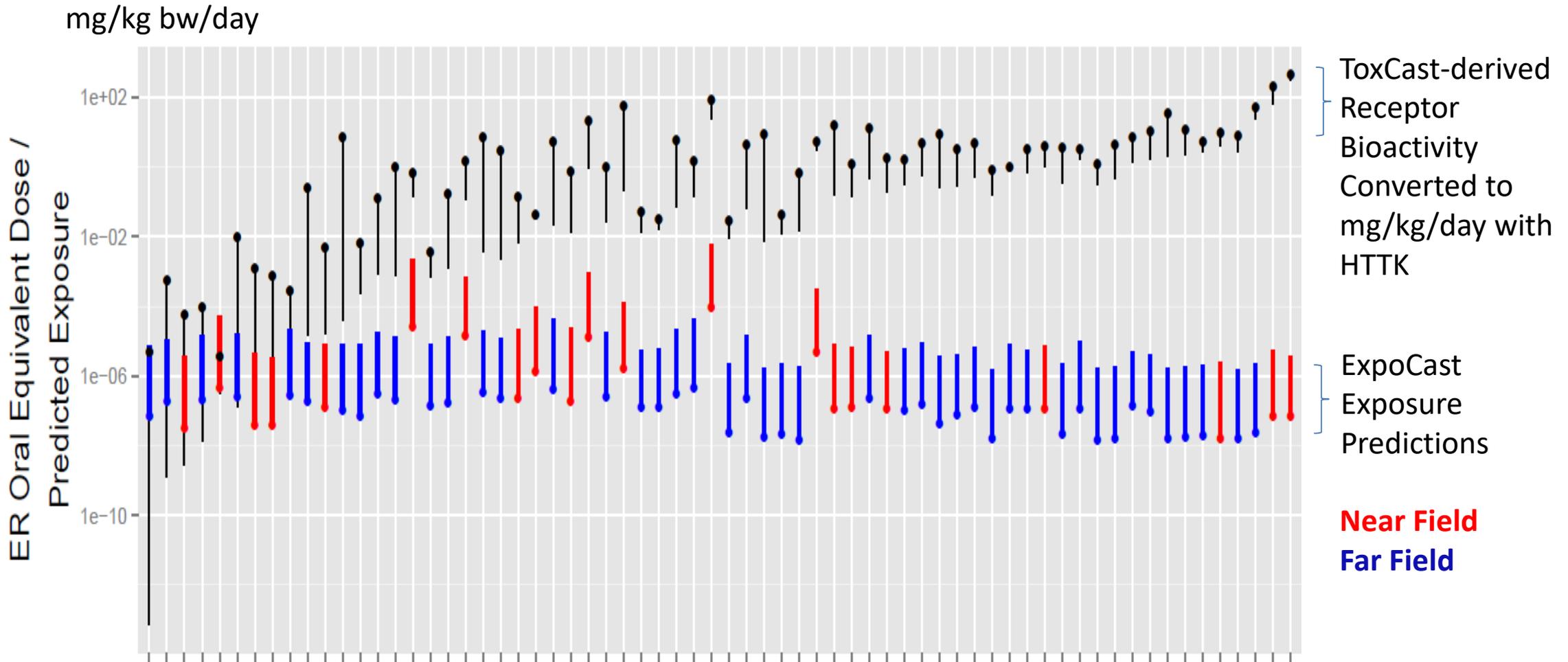
Various Combinations of IVIVE Assumptions

New Exposure Data and Models

High throughput screening + *in vitro*-*in vivo* extrapolation (IVIVE) can predict a dose (mg/kg bw/day) that might be adverse



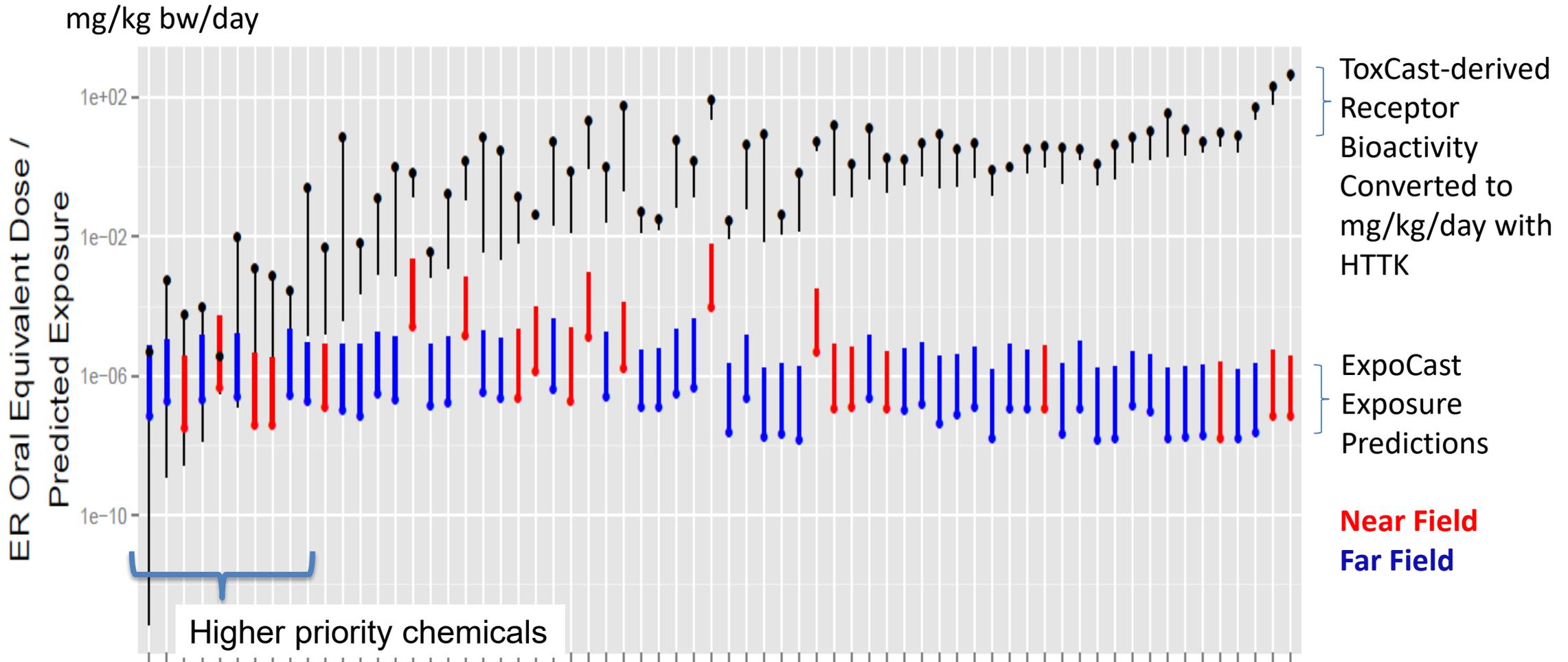
High Throughput Risk Prioritization in Practice



ToxCast Chemicals

December, 2014 Panel:
“Scientific Issues Associated with Integrated Endocrine
Bioactivity and Exposure-Based Prioritization and Screening”

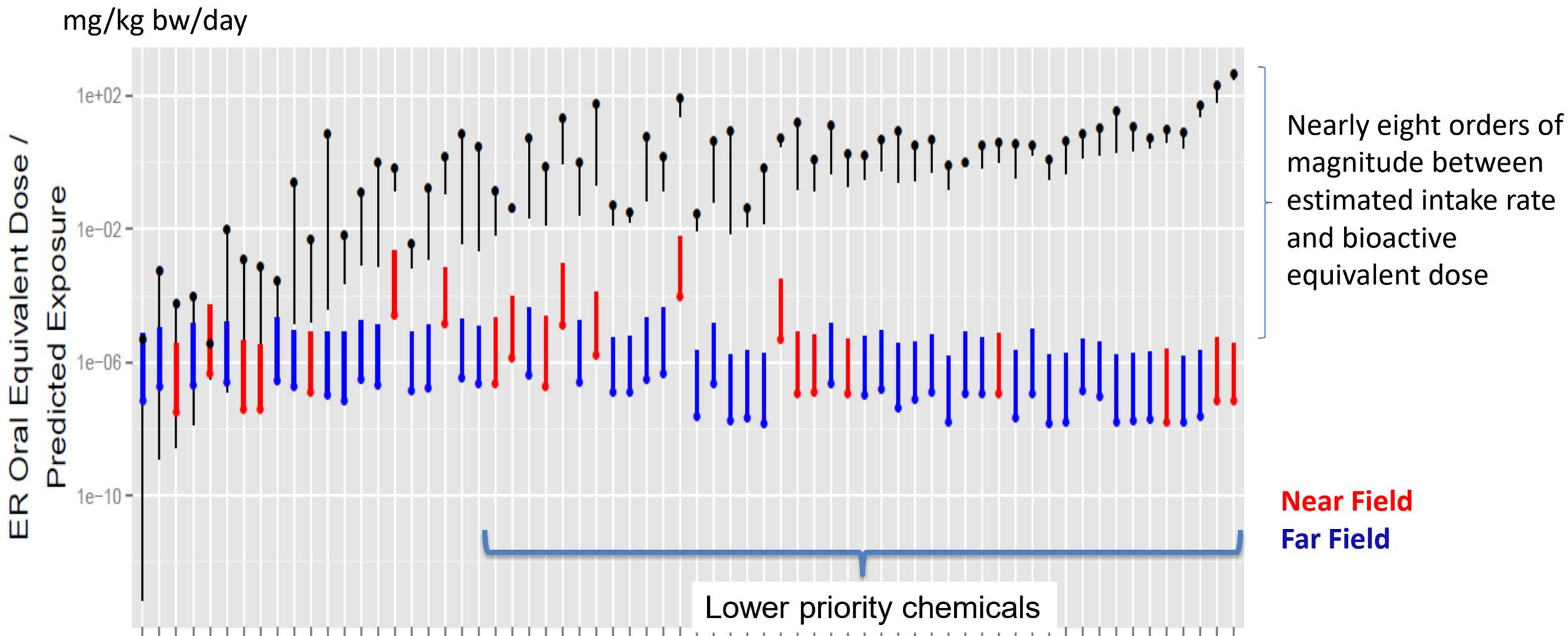
High Throughput Risk Prioritization in Practice



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December, 2014 Panel:
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High Throughput Risk Prioritization in Practice

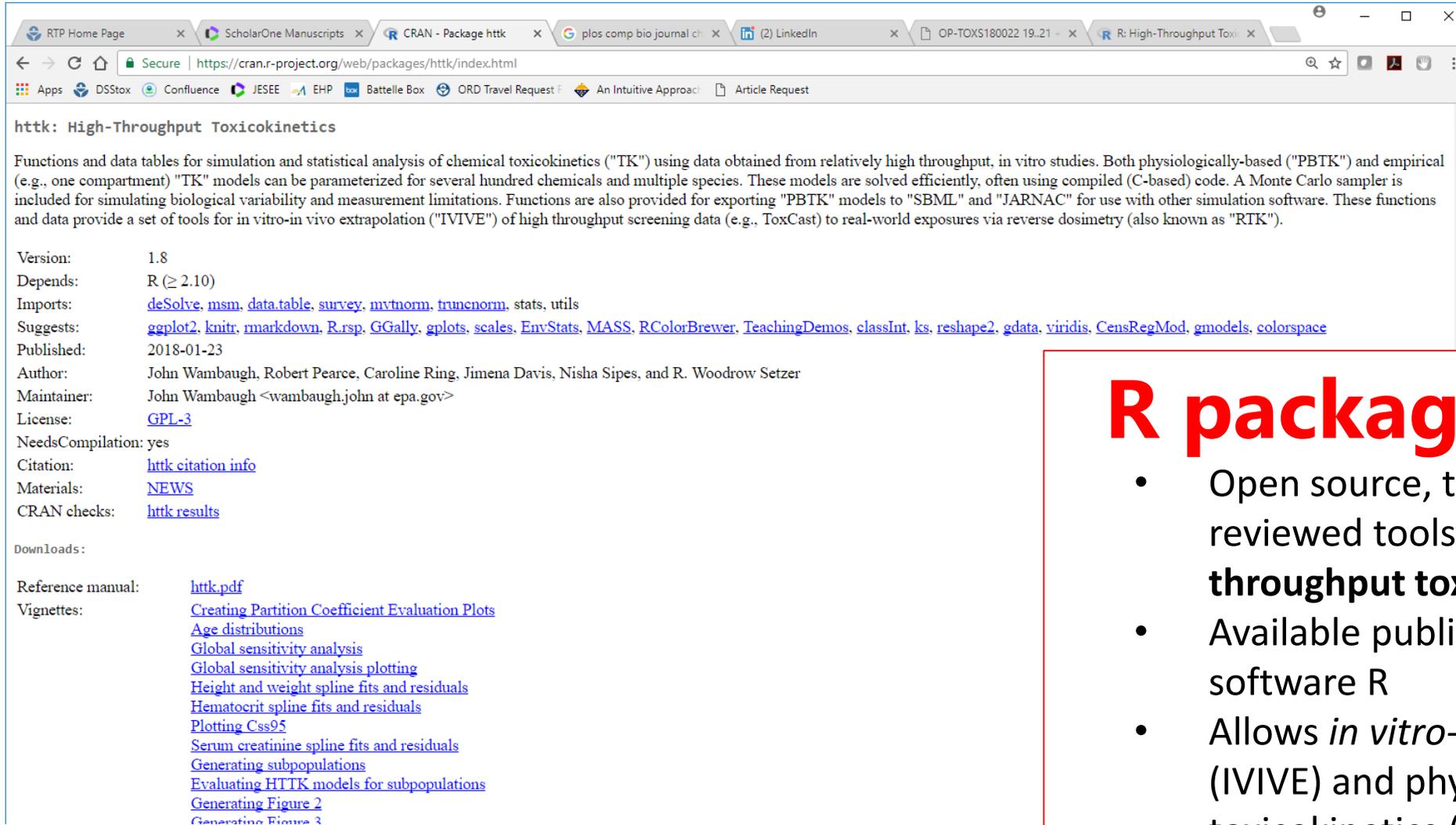


ToxCast Chemicals

December, 2014 Panel:
“Scientific Issues Associated with Integrated Endocrine
Bioactivity and Exposure-Based Prioritization and Screening”

Open Source Tools and Data for HTKK

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



The screenshot shows the CRAN package page for 'httk'. The browser tabs include 'RTP Home Page', 'ScholarOne Manuscripts', 'CRAN - Package httk', 'plos comp bio journal ch', '(2) LinkedIn', 'OP-TOXS180022 19.21', and 'R: High-Throughput Tox'. The address bar shows the URL 'https://cran.r-project.org/web/packages/httk/index.html'. The page content includes the package name 'httk: High-Throughput Toxicokinetics', a description of its functions, and various metadata fields.

httk: High-Throughput Toxicokinetics

Functions and data tables for simulation and statistical analysis of chemical toxicokinetics ("TK") using data obtained from relatively high throughput, in vitro studies. Both physiologically-based ("PBTk") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability and measurement limitations. Functions are also provided for exporting "PBTk" models to "SBML" and "JARNAC" for use with other simulation software. These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry (also known as "RTK").

Version: 1.8
Depends: R (\geq 2.10)
Imports: [deSolve](#), [msm](#), [data.table](#), [survey](#), [mvtnorm](#), [truncnorm](#), stats, utils
Suggests: [ggplot2](#), [knitr](#), [rmarkdown](#), [R.rsp](#), [GGally](#), [gplots](#), [scales](#), [EnvStats](#), [MASS](#), [RColorBrewer](#), [TeachingDemos](#), [classInt](#), [ks](#), [reshape2](#), [gdata](#), [viridis](#), [CensRegMod](#), [gmodels](#), [colorspace](#)
Published: 2018-01-23
Author: John Wambaugh, Robert Pearce, Caroline Ring, Jimena Davis, Nisha Sipes, and R. Woodrow Setzer
Maintainer: John Wambaugh <wambaugh.john at epa.gov>
License: [GPL-3](#)
NeedsCompilation: yes
Citation: [httk citation info](#)
Materials: [NEWS](#)
CRAN checks: [httk results](#)

Downloads:

Reference manual: [httk.pdf](#)
Vignettes: [Creating Partition Coefficient Evaluation Plots](#), [Age distributions](#), [Global sensitivity analysis](#), [Global sensitivity analysis plotting](#), [Height and weight spline fits and residuals](#), [Hematocrit spline fits and residuals](#), [Plotting C_{ss95}](#), [Serum creatinine spline fits and residuals](#), [Generating subpopulations](#), [Evaluating HTKK models for subpopulations](#), [Generating Figure 2](#), [Generating Figure 3](#)

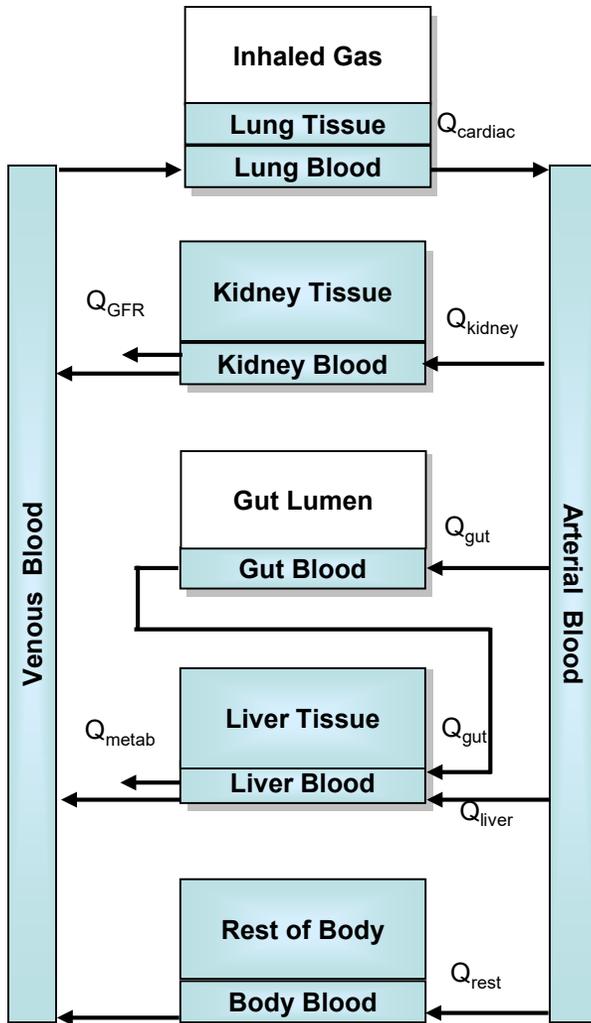
R package "httk"

- 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)

What you can do with R Package “httk”?

- Allows one compartment, three-compartment, and PBTK modeling
- Allows conversion of *in vitro* concentration to *in vivo* doses
- Allows prediction of internal tissue concentrations from dose regimen (oral and intravenous)
- A peer-reviewed paper in the Journal of Statistical software provides a how-to guide (Pearce et al., 2017a)
- You can use the built in chemical library or add more chemical information (examples provided in JSS paper)
- You can load specific (older) versions of the package
- You can use specific demographics in the population simulator (Ring et al., 2017)
- You can control the built in random number generator to reproduce the same random sequence (function `set.seed()`)

A General Physiologically-based Toxicokinetic (PBTK) Model



- “httk” includes a generic PBTK model
- Some tissues (e.g. arterial blood) are simple compartments, while others (e.g. kidney) are compound compartments consisting of separate blood and tissue sections with constant partitioning (i.e., tissue specific partition coefficients)
- Exposures are absorbed from reservoirs (gut lumen)
- Some specific tissues (lung, kidney, gut, and liver) are modeled explicitly, others (e.g. fat, brain, bones) are lumped into the “Rest of Body” compartment.
- The only ways chemicals “leave” the body are through metabolism (change into a metabolite) in the liver or excretion by glomerular filtration into the proximal tubules of the kidney (which filter into the lumen of the kidney).

Why Build Another Generic PBTK Tool?

	SimCYP	ADMET Predictor / GastroPlus	MEGen	IndusChemFate	httk
Maker	SimCYP Consortium / Certara	Simulations Plus	UK Health and Safety Laboratory	Cefic LRI	US EPA
Reference	Jamei et al. (2009)	Lukacova et al., (2009)	Loizou et al. (2011)	Jongeneelen et al., (2013)	Pearce et al. (2017a)
Availability	License, but inexpensive for research	License, but inexpensive for research	Free: http://xnet.hsl.gov.uk/megen	Free: http://cefic-lri.org/lri_toolbox/induschemfate/	Free: https://CRAN.R-project.org/package=httk
Open Source	No	No	Yes	No	Yes
Default PBPK Structure	Yes	Yes	No	Yes	Yes
Expandable PBPK Structure	No	No	Yes	No	No
Population Variability	Yes	No	No	No	Yes
Batch Mode	Yes	Yes	No	No	Yes
Graphical User Interface	Yes	Yes	Yes	Excel	No
Physiological Data	Yes	Yes	Yes	Yes	Yes
Chemical-Specific Data Library	Many Clinical Drugs	No	No	15 Environmental Compounds	543 Pharmaceutical and ToxCast Compounds
Ionizable Compounds	Yes	Yes	Potentially	No	Yes
Export Function	No	No	Matlab and AcslX	No	SBML and Jarnac
R Integration	No	No	No	No	Yes
Easy Reverse Dosimetry	Yes	Yes	No	No	Yes
Future Proof XML	No	No	Yes	No	No

We want to do a statistical analysis (using R) for as many chemicals as possible

Oral Equivalent Dose Examples

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.95  
quantile, for Acetochlor (published value):
```

```
get_wetmore_oral_equiv(0.1,chem.cas="34256-82-1")
```

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.95  
quantile, for Acetochlor (calculated value):
```

```
calc_mc_oral_equiv(0.1,chem.cas="34256-82-1")
```

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.05,  
0.5, and 0.95 quantile, for Acetochlor (published values):
```

```
get_wetmore_oral_equiv(0.1,chem.cas="34256-82-1",which.quantile=c(0.05,0.5,0.95))
```

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.05,  
0.5, and 0.95 quantiles, for Acetochlor (calculated value):
```

```
calc_mc_oral_equiv(0.1,chem.cas="34256-82-1",which.quantile=c(0.05,0.5,0.95))
```

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for rat, 0.95  
quantile, for Acetochlor (calculated value):
```

```
calc_mc_oral_equiv(0.1,chem.cas="34256-82-1",species="Rat")
```

Interspecies Extrapolation Examples

```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for human for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1")
```

```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for rat for Acetochlor (should produce errors since  
there is no published value, 0.5 quantile only):  
get_wetmore_css(chem.cas="34256-82-1",species="Rat")
```

```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for rat for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1",species="Rat")
```

```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.5 quantile for rat for Acetochlor (published value):  
get_wetmore_css(chem.cas="34256-82-1",species="Rat",which.quantile=0.5)
```

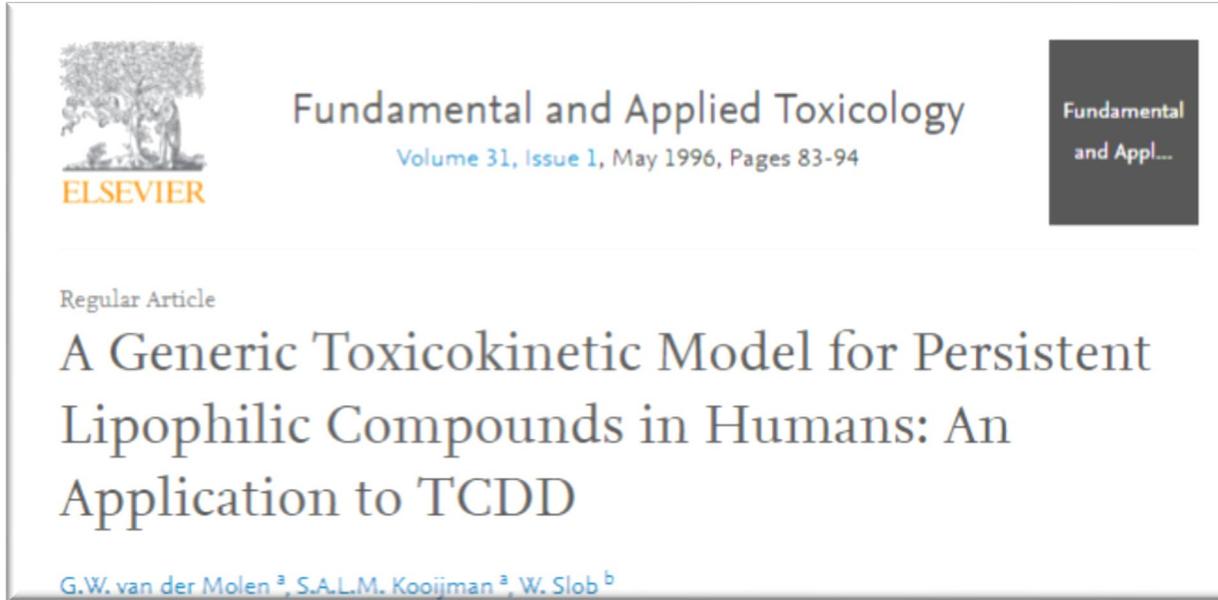
```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.5 quantile for rat for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1",species="Rat",which.quantile=0.5)
```

```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for mouse for Acetochlor (should produce error since  
there is no published value, human and rat only):  
get_wetmore_css(chem.cas="34256-82-1",species="Mouse")
```

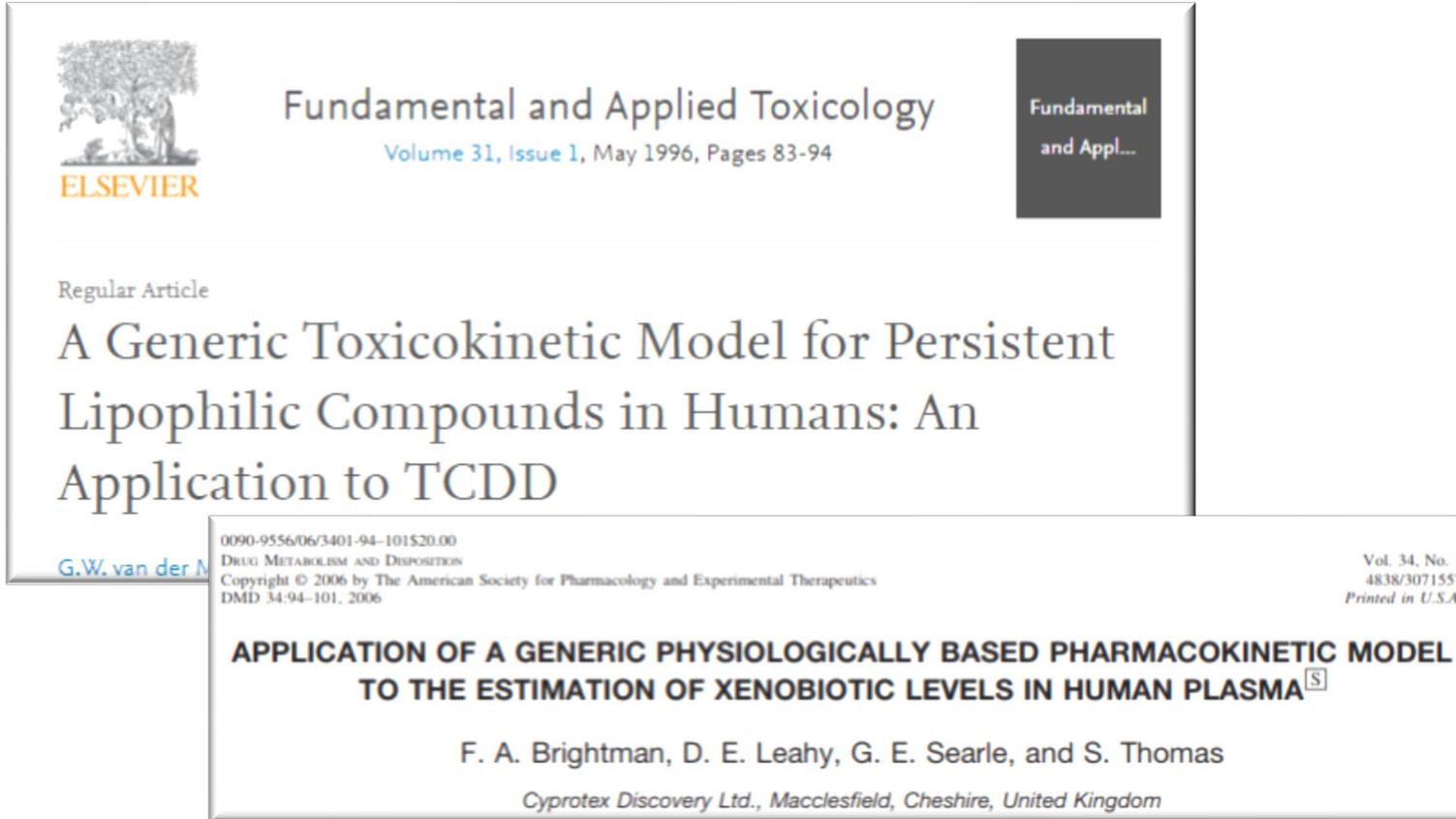
```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for mouse for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1",species="Mouse")  
calc_mc_css(chem.cas="34256-82-1",species="Mouse",default.to.human=T)
```

Generic PBTK Models

There is nothing new about the idea of generic PBTK models...



There is nothing new about the idea of generic PBTK models...



 **ELSEVIER**

Fundamental and Applied Toxicology
Volume 31, Issue 1, May 1996, Pages 83-94

Fundamental and Appl...

Regular Article

A Generic Toxicokinetic Model for Persistent Lipophilic Compounds in Humans: An Application to TCDD

G.W. van der Molen

0090-9556/06/3401-94-101\$20.00
DRUG METABOLISM AND DISPOSITION
Copyright © 2006 by The American Society for Pharmacology and Experimental Therapeutics
DMD 34:94-101, 2006

Vol. 34, No. 1
4838/3071557
Printed in U.S.A.

APPLICATION OF A GENERIC PHYSIOLOGICALLY BASED PHARMACOKINETIC MODEL TO THE ESTIMATION OF XENOBIOTIC LEVELS IN HUMAN PLASMA^S

F. A. Brightman, D. E. Leahy, G. E. Searle, and S. Thomas

Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

Generic PBTK Models

There is nothing new about the idea of generic PBTK models...



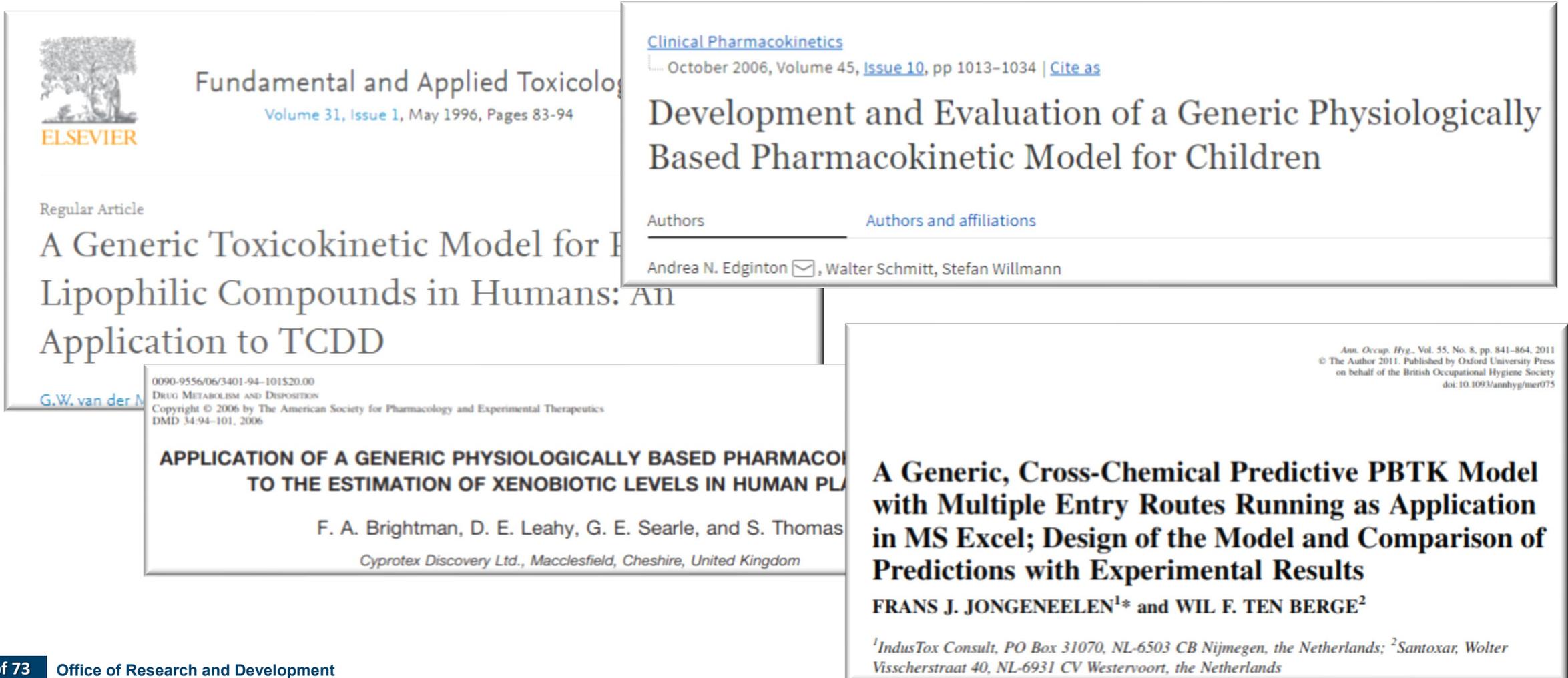
Fundamental and Applied Toxicology
Volume 31, Issue 1, May 1996, Pages 83-94
ELSEVIER
Regular Article
A Generic Toxicokinetic Model for Lipophilic Compounds in Humans: An Application to TCDD
G.W. van der ...

Clinical Pharmacokinetics
October 2006, Volume 45, Issue 10, pp 1013-1034 | [Cite as](#)
Development and Evaluation of a Generic Physiologically Based Pharmacokinetic Model for Children
Authors: [Authors and affiliations](#)
Andrea N. Edginton , Walter Schmitt, Stefan Willmann

0090-9556/06/3401-94-101\$20.00
DRUG METABOLISM AND DISPOSITION
Copyright © 2006 by The American Society for Pharmacology and Experimental Therapeutics
DMD 34:94-101, 2006
Vol. 34, No. 1
4838/3071557
Printed in U.S.A.
APPLICATION OF A GENERIC PHYSIOLOGICALLY BASED PHARMACOKINETIC MODEL TO THE ESTIMATION OF XENOBIOTIC LEVELS IN HUMAN PLASMA^S
F. A. Brightman, D. E. Leahy, G. E. Searle, and S. Thomas
Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

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ELSEVIER
Regular Article
A Generic Toxicokinetic Model for Lipophilic Compounds in Humans: An Application to TCDD
G.W. van der ...
0090-9556/06/3401-94-101\$20.00
DRUG METABOLISM AND DISPOSITION
Copyright © 2006 by The American Society for Pharmacology and Experimental Therapeutics
DMD 34:94-101, 2006

Development and Evaluation of a Generic Physiologically Based Pharmacokinetic Model for Children
Clinical Pharmacokinetics
October 2006, Volume 45, Issue 10, pp 1013-1034 | Cite as
Authors: Andrea N. Edginton, Walter Schmitt, Stefan Willmann

A Generic, Cross-Chemical Predictive PBTK Model with Multiple Entry Routes Running as Application in MS Excel; Design of the Model and Comparison of Predictions with Experimental Results
FRANS J. JONGENEELLEN^{1*} and WIL F. TEN BERGE²
¹IndusTox Consult, PO Box 31070, NL-6503 CB Nijmegen, the Netherlands; ²Santoxar, Wolter Visscherstraat 40, NL-6931 CV Westervoort, the Netherlands

APPLICATION OF A GENERIC PHYSIOLOGICALLY BASED PHARMACOKINETIC MODEL TO THE ESTIMATION OF XENOBIOTIC LEVELS IN HUMAN PLASMA
F. A. Brightman, D. E. Leahy, G. E. Searle, and S. Thomas
Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

TOXICOLOGICAL SCIENCES **126**(1), 5–15 (2012)
doi:10.1093/toxsci/kfr295
Advance Access publication November 1, 2011

Physiologically Based Pharmacokinetic Model Use in Risk Assessment—Why Being Published Is Not Enough

Eva D. McLanahan,^{*-1} Hisham A. El-Masri,[†] Lisa M. Sweeney,[‡] Leonid Y. Kopylev,^{||} Harvey J. Clewell,[§] John F. Wambaugh,[¶] and P. M. Schlosser^{||}

“Although publication of a PBPK model in a peer-reviewed journal is a mark of good science, subsequent evaluation of published models and the supporting computer code is necessary for their consideration for use in [Human Health Risk Assessments]”

The White House

Office of the Press Secretary

For Immediate Release

May 09, 2013

Executive Order -- Making Open and Machine Readable the New Default for Government Information

EXECUTIVE ORDER

MAKING OPEN AND MACHINE READABLE THE NEW DEFAULT FOR GOVERNMENT INFORMATION

By the authority vested in me as President by the Constitution and the laws of the United States of America, it is hereby ordered as follows:

Section 1. General Principles. Openness in government strengthens our democracy, promotes the delivery of efficient and effective services to the public, and contributes to economic growth. As one vital benefit of open government, making information resources easy to find, accessible, and usable

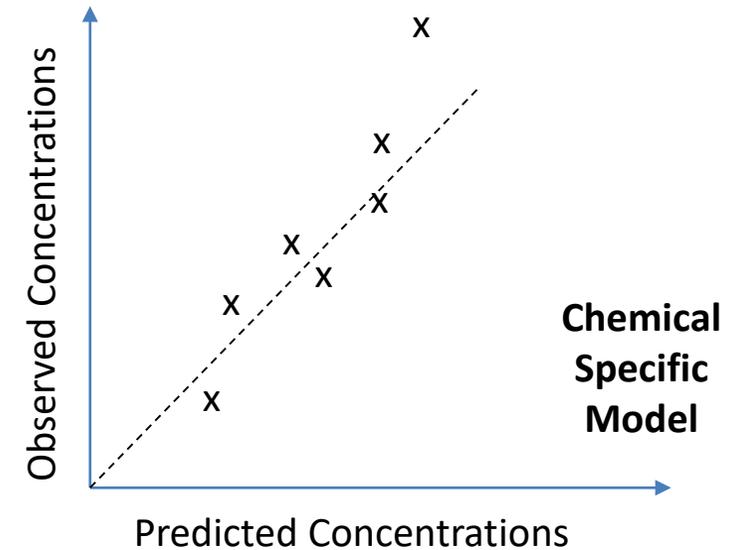
“...the default state of new and modernized Government information resources shall be open and machine readable.”

Doing Statistical Analysis with HTK

- If we are to use HTK, we need confidence in predictive ability
- In drug development, HTK methods estimate therapeutic doses for clinical studies – predicted concentrations are typically on the order of values measured in clinical trials (Wang, 2010)
 - For most compounds in the environment there will be no clinical trials
- Uncertainty must be well characterized
 - We compare to *in vivo* data to get **empirical estimates of HTK uncertainty**
 - ORD has both compiled existing (literature) TK data (Wambaugh *et al.*, 2015) and conducted new experiments in rats on chemicals with HTK *in vitro* data (Wambaugh *et al.*, 2018)
 - Any approximations, omissions, or mistakes should work to increase the estimated uncertainty when evaluated systematically across chemicals

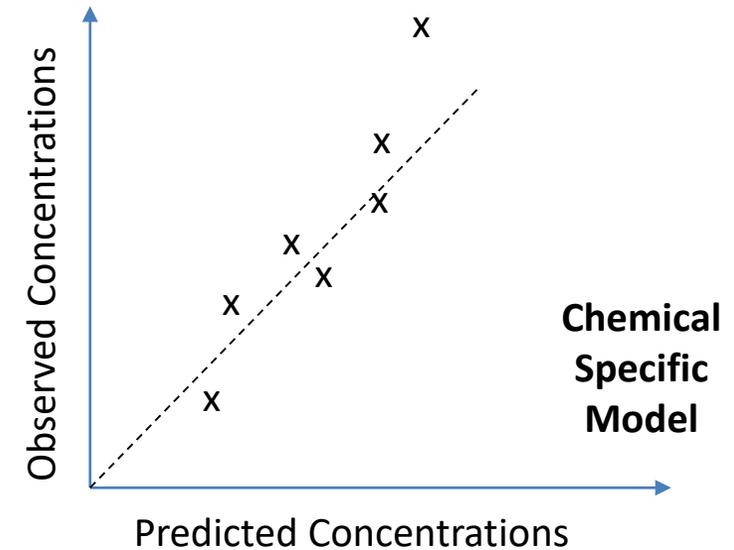
Building Confidence in TK Models

- In order to evaluate a **chemical-specific TK model** for “chemical x” you can compare the predictions to *in vivo* measured data
 - Can estimate bias
 - Can estimate uncertainty
 - Can consider using model to extrapolate to other situations (dose, route, physiology) where you don’t have data



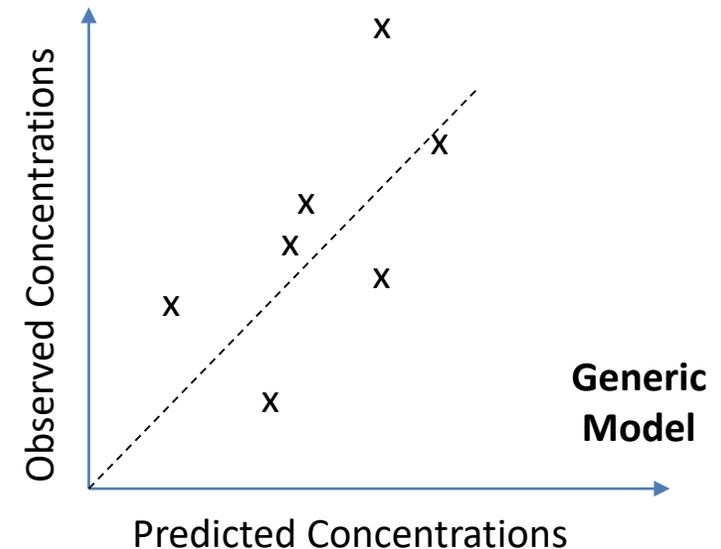
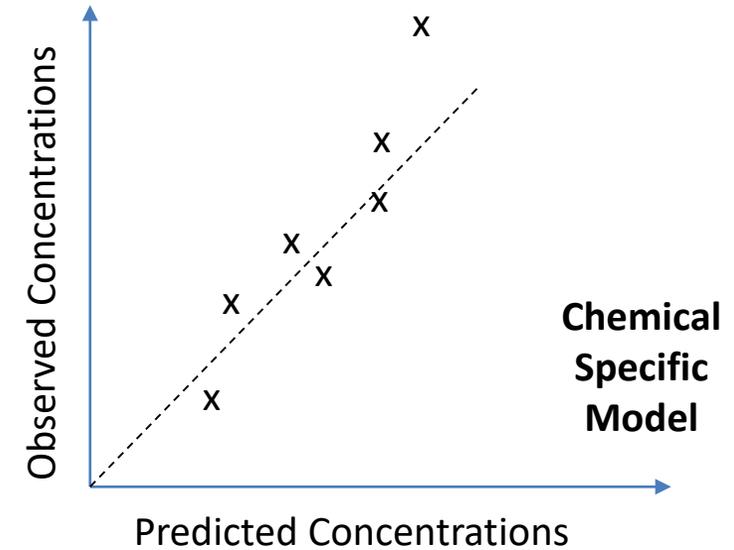
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- However, we do not typically have TK data



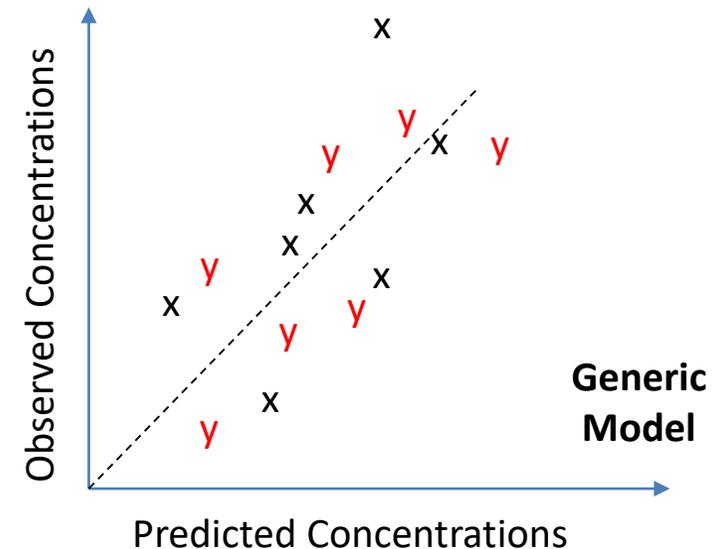
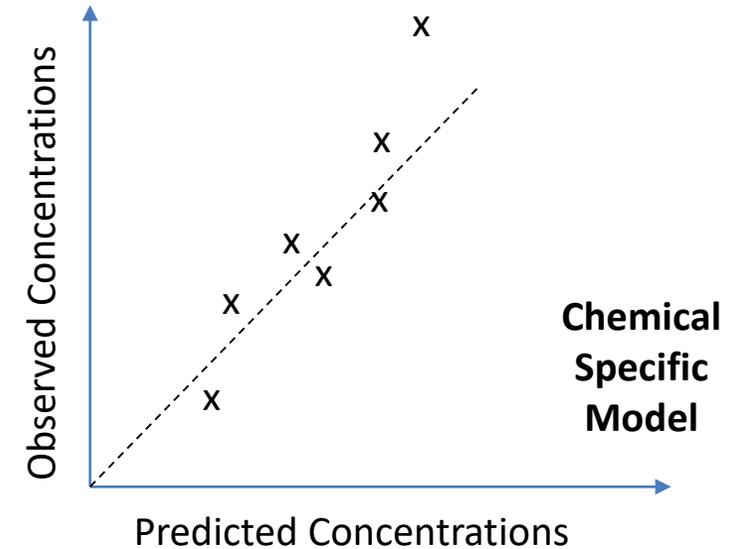
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- However, we do not typically have TK data
- We can parameterize a **generic TK model**, and evaluate that model for as many chemicals as we do have data
 - We do expect larger uncertainty, but also greater confidence in model implementation
 - Estimate bias and uncertainty, and try to correlate with chemical-specific properties



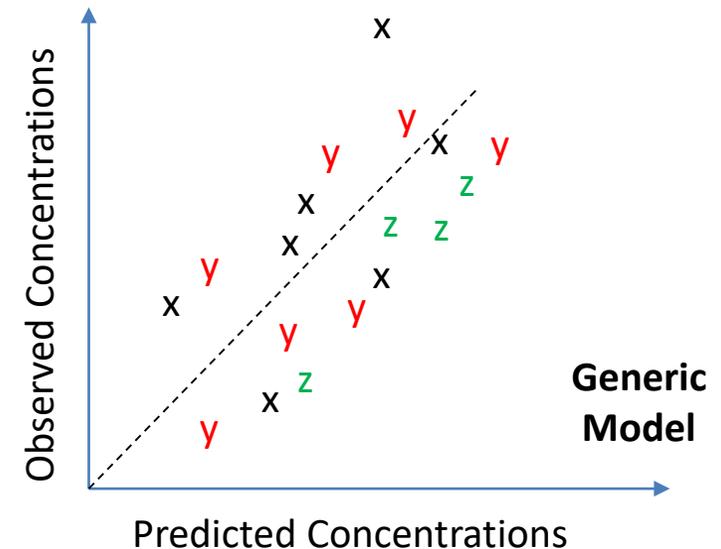
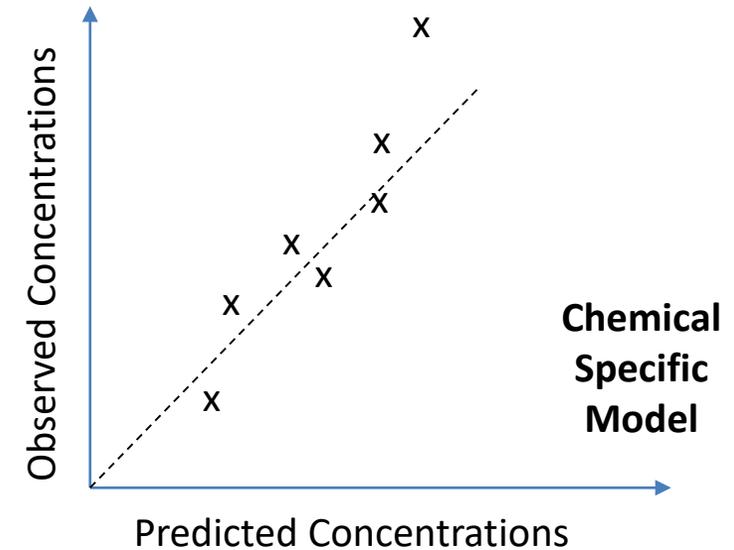
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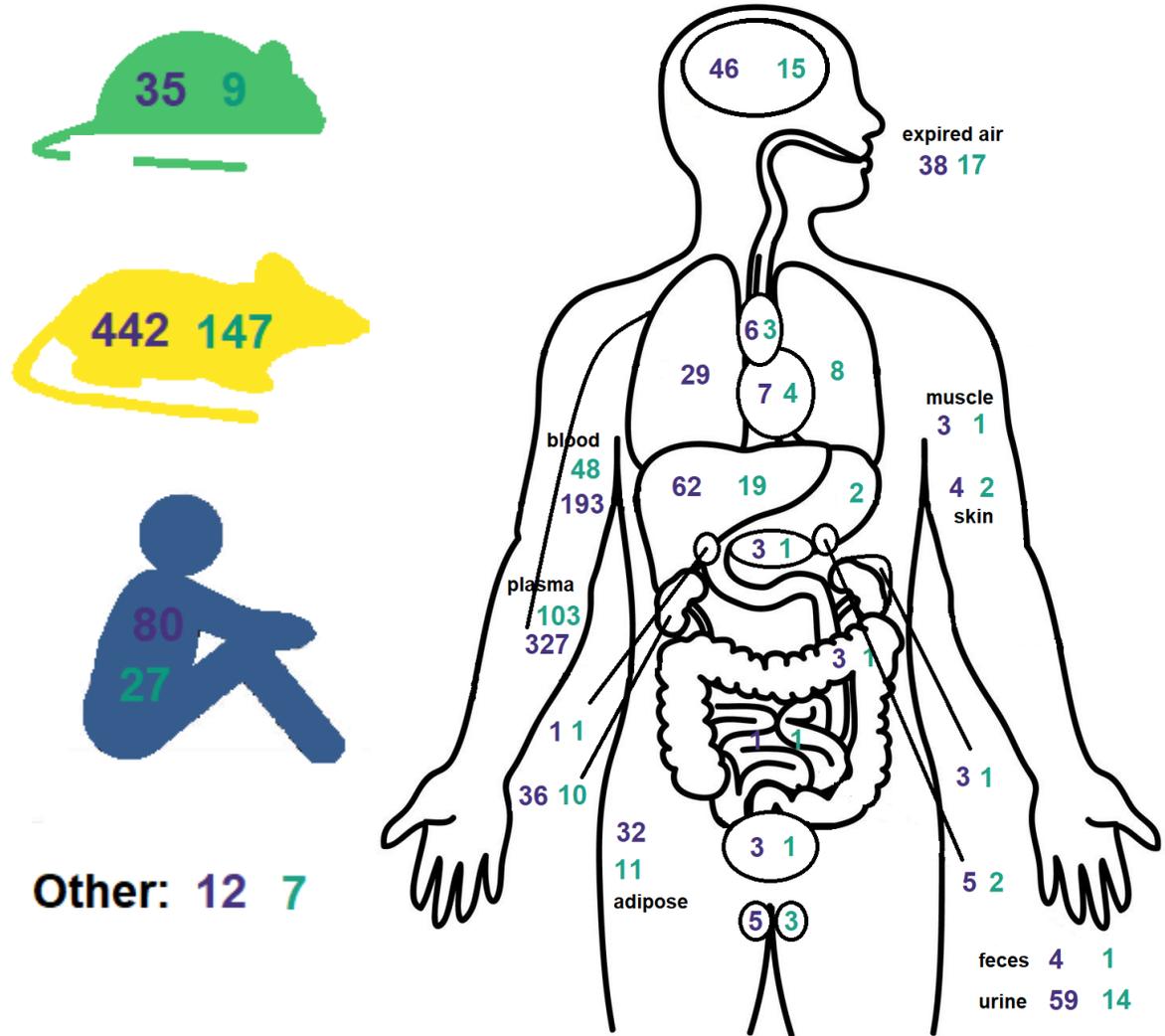
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In Vivo TK Database

- EPA is developing 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
- Database will be made available through web interface and through the “httk” R package
- Standardized, open source curve fitting software *invivoPKfit* used to calibrate models to all data:

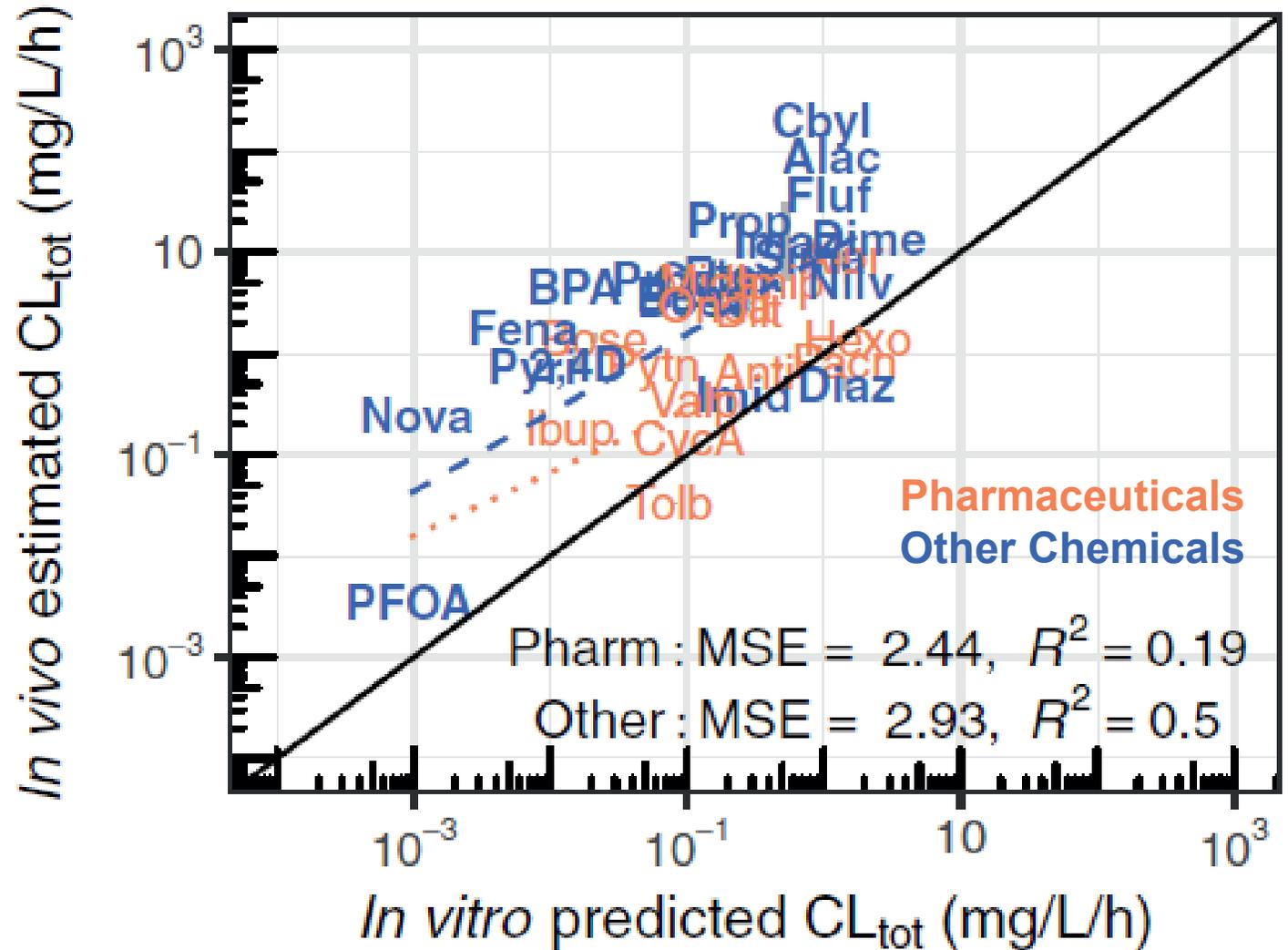


<https://github.com/USEPA/CompTox-ExpoCast-invivoPKfit>

Sayre et al. (in preparation)

Observed Total Clearance

- We estimate clearance from two processes – hepatic metabolism (liver) and passive glomerular filtration (kidney)
- This appears to work better for pharmaceuticals than other chemicals:
 - ToxCast chemicals are overestimated
- Non-pharmaceuticals may be subject to extrahepatic metabolism and/or active transport



Variability

Different crayons
have different
colors...



Variability

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



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

Sample NHANES
biometrics for
actual individuals:

Sex
Race/ethnicity
Age
Height
Weight
Serum creatinine

Population simulator for HTTK



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Sex
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Age
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Population simulator for HTTK



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.)

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

Sample NHANES biometrics for actual individuals:

- Sex
- Race/ethnicity
- Age
- Height
- Weight
- Serum creatinine

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.)

Uncertainty

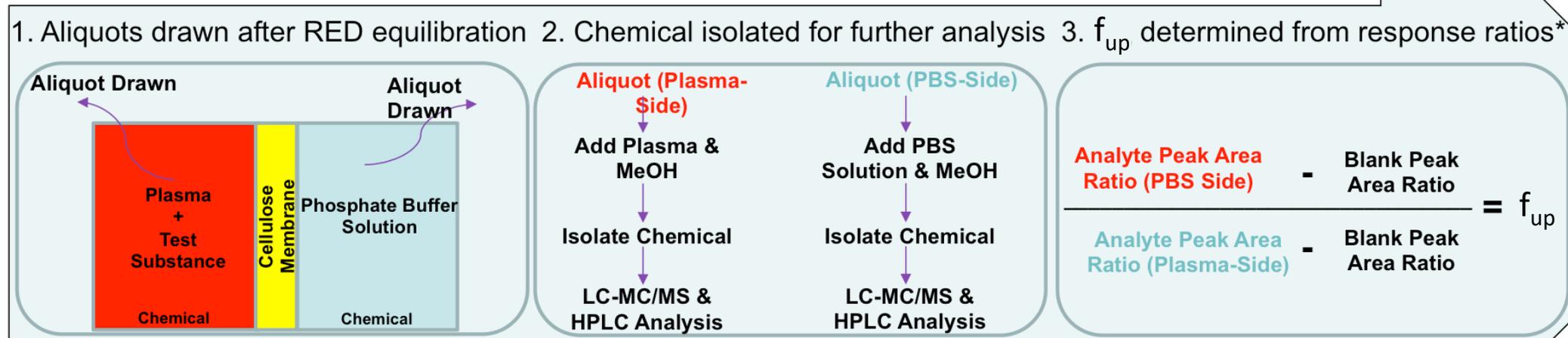
Until I open the
box, I don't know
what colors I
have...

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



Analytical Chemistry is an HTTK Bottleneck

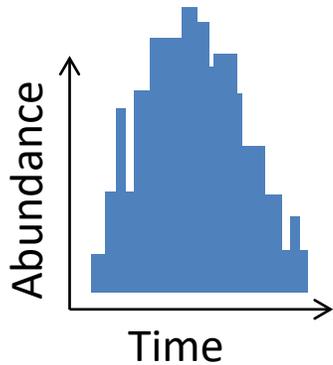
- For HTTK we always need to develop a chemical-specific method for quantitating amount of chemical *in vitro*
 - This is very different from HTS where the same readout (e.g., bioluminescence) can be used for most chemicals
- In Wetmore et al. (2012), the rapid equilibrium dialysis (RED) assay (Waters et al. 2008) failed for fraction unbound in plasma (f_{up}) 38% of the chemicals.



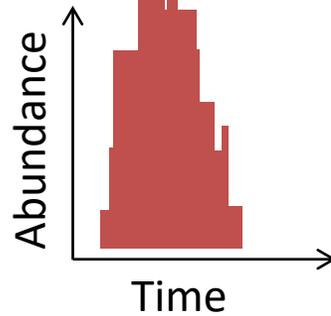
New HTTK Measurements and Uncertainty Analysis

The HTTK *in vitro* assays need to measure differences in chemical concentration

Internal Standard



Chemical Peak

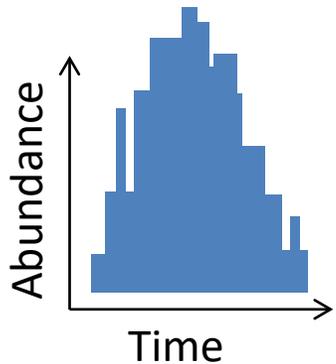


- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio R of the chemical peak to the ITSD

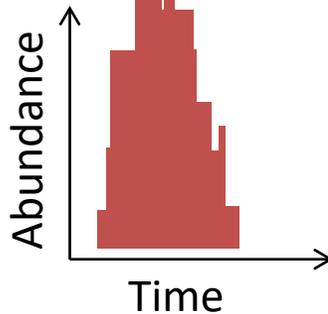
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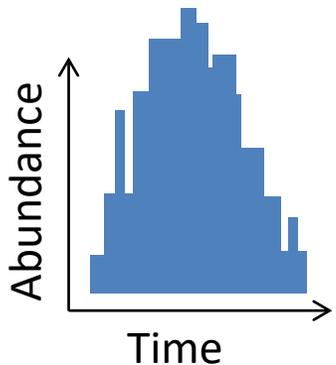
Internal Standard



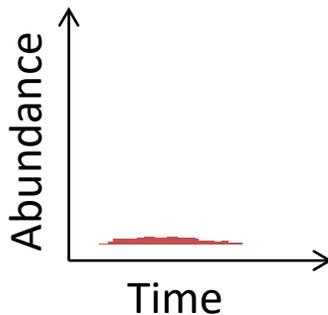
Chemical Peak



Internal Standard



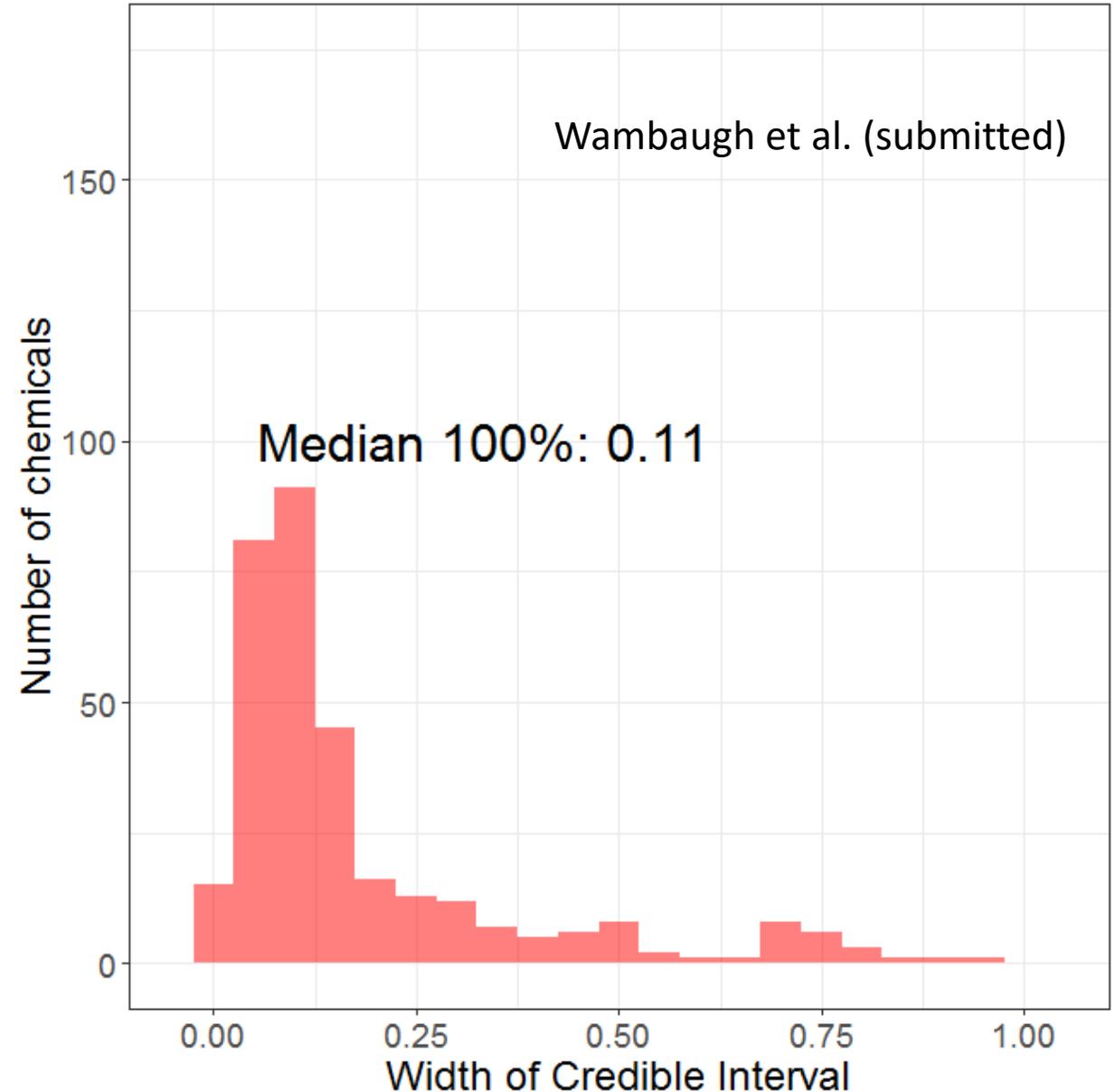
Chemical Peak



- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio R of the chemical peak to the ITSD
- For new measurements HTTK (>200 compounds to data) performed by Cyprotex, we have modified RED protocol to use a titration of plasma protein (10%, 30%, 100%) of physiological concentration
 - Keeps chemical concentration in the same range
- Analyzed data in Bayesian framework that included a model for analytical chemistry
 - Bayesian approach gives a credible interval (range of values that would be consistent with the data) – quantitative uncertainty

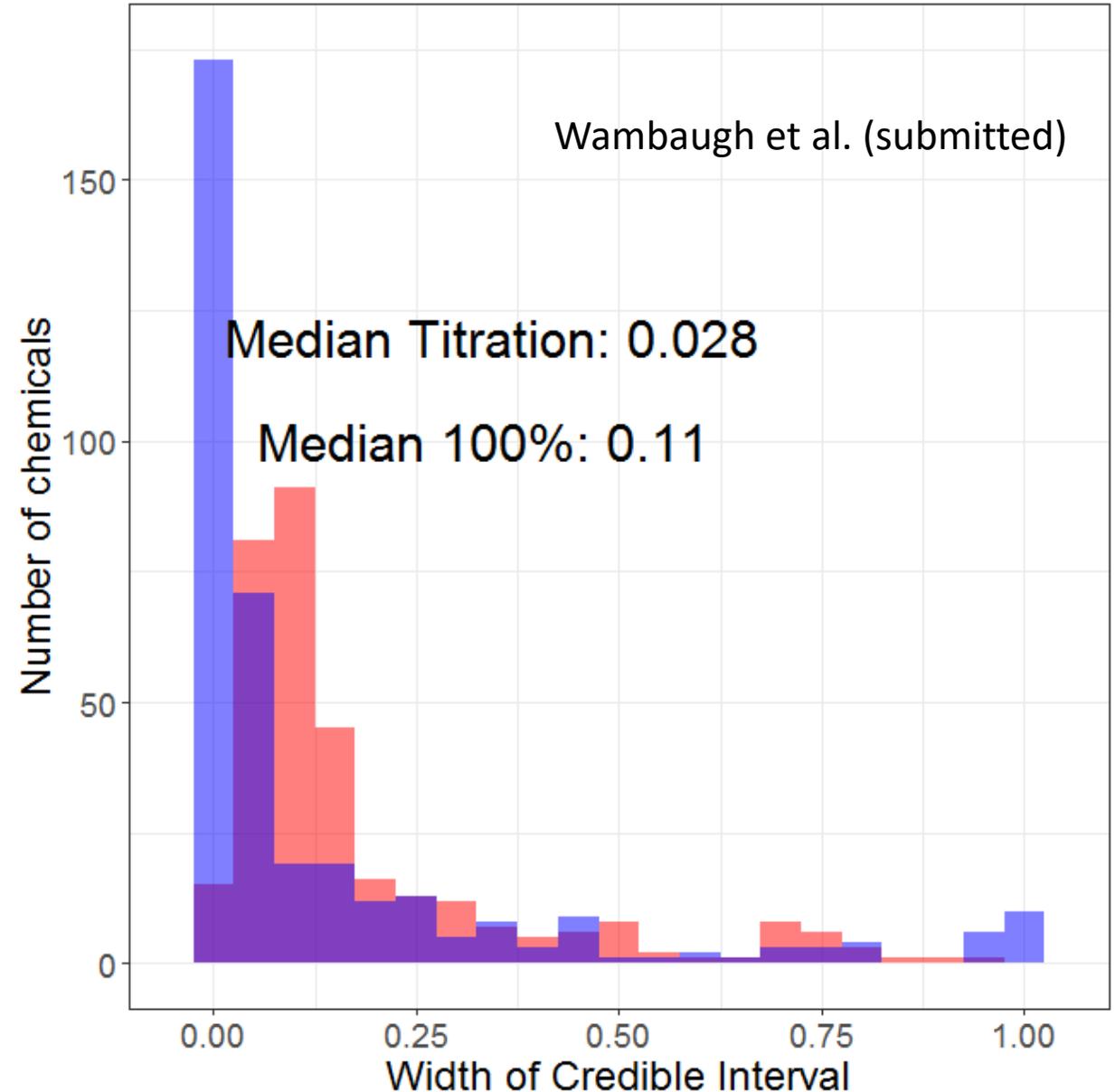
New Plasma Binding Protocol Reduces Uncertainty

- New protocol performs assay at 100%, 30%, and 10% of physiologic protein concentration
- Median uncertainty for 100% physiological concentration only: $\pm 5.5\%$



New Plasma Binding Protocol Reduces Uncertainty

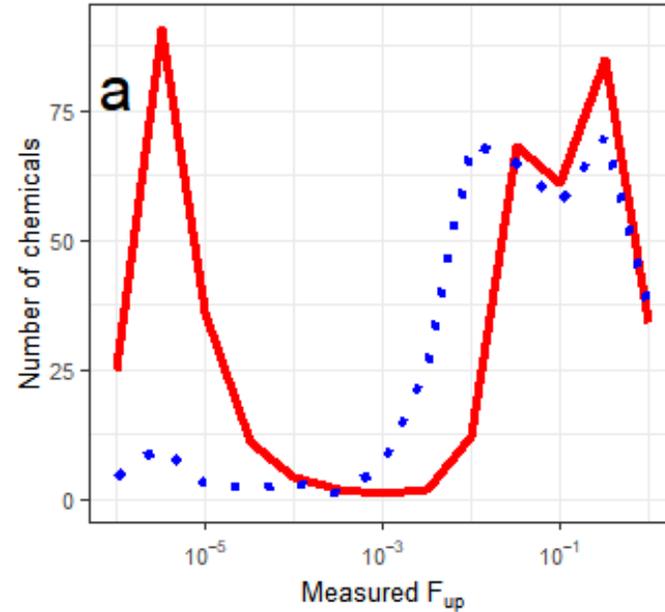
- New protocol performs assay at 100%, 30%, and 10% of physiologic protein concentration
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- Median uncertainty for three-point assay: $\pm 1.4\%$



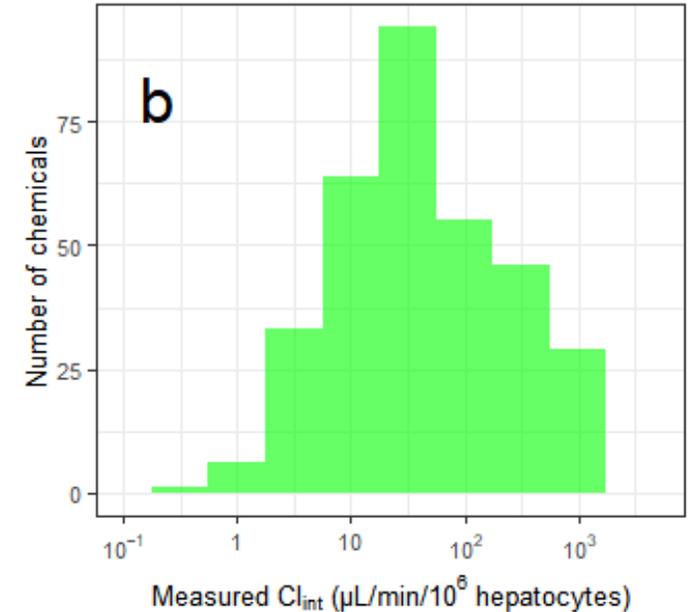
New Data!

New experimental measurements of f_{up} and Cl_{int} are reported for 418 and 467 chemicals, respectively. These data raise the HTTK chemical coverage of the ToxCast Phase I and II libraries to 57%.

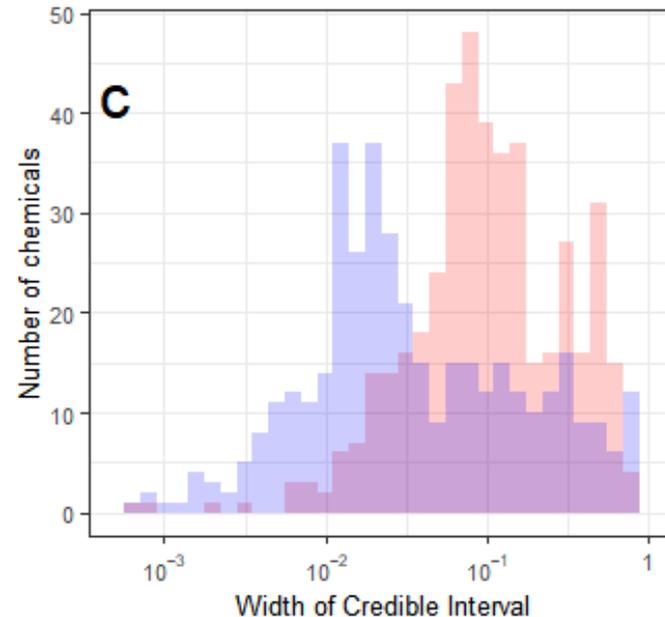
418 Chemicals Measured



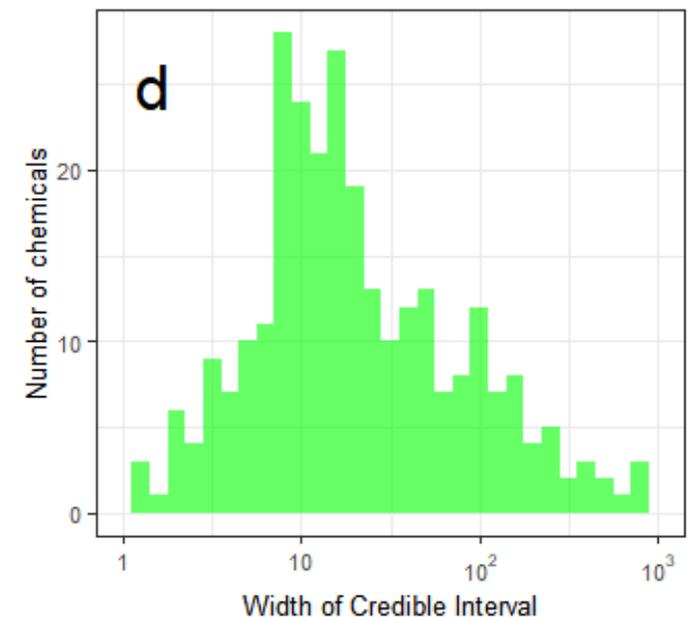
467 Chemicals Measured



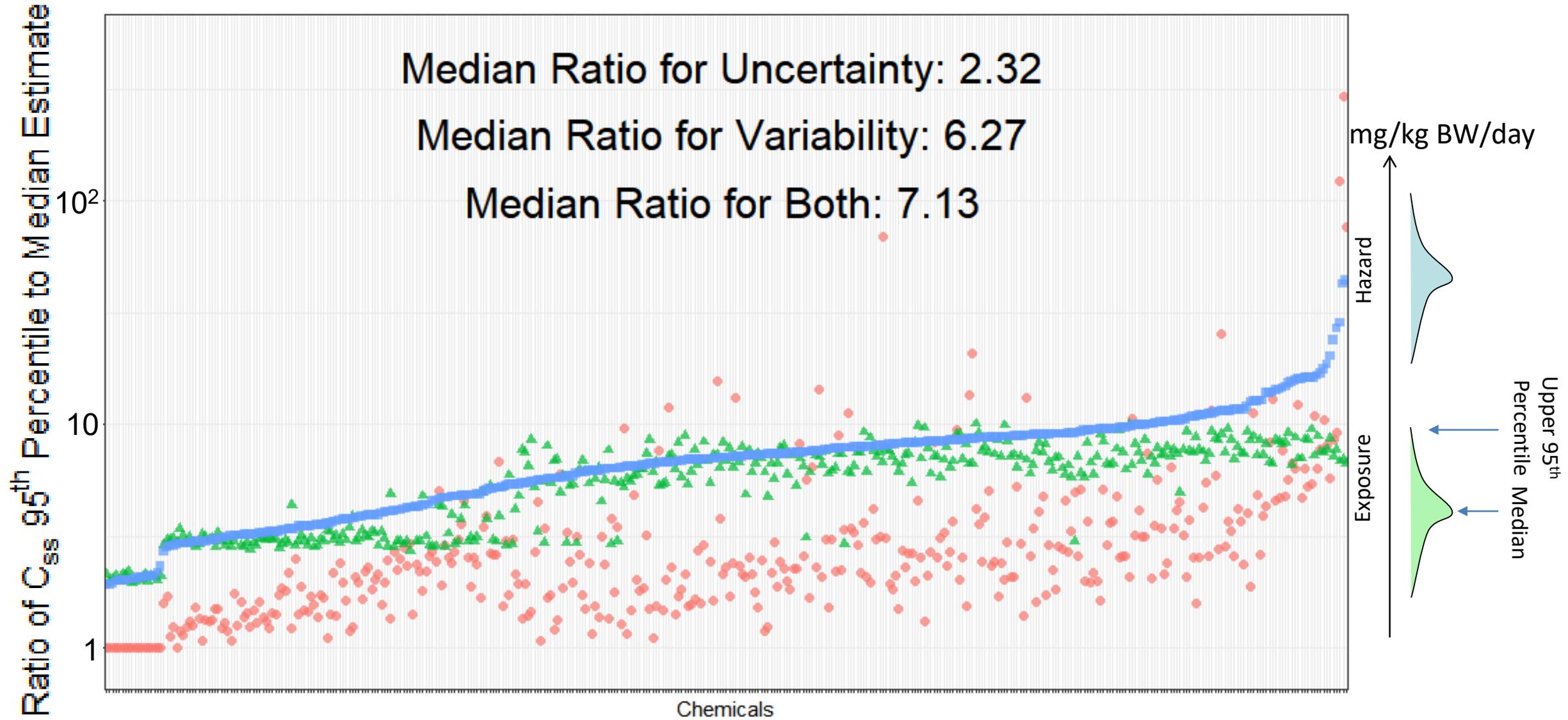
Cl: 0.1 CV: 0.4 / Cl: 0.03 CV: 0.1



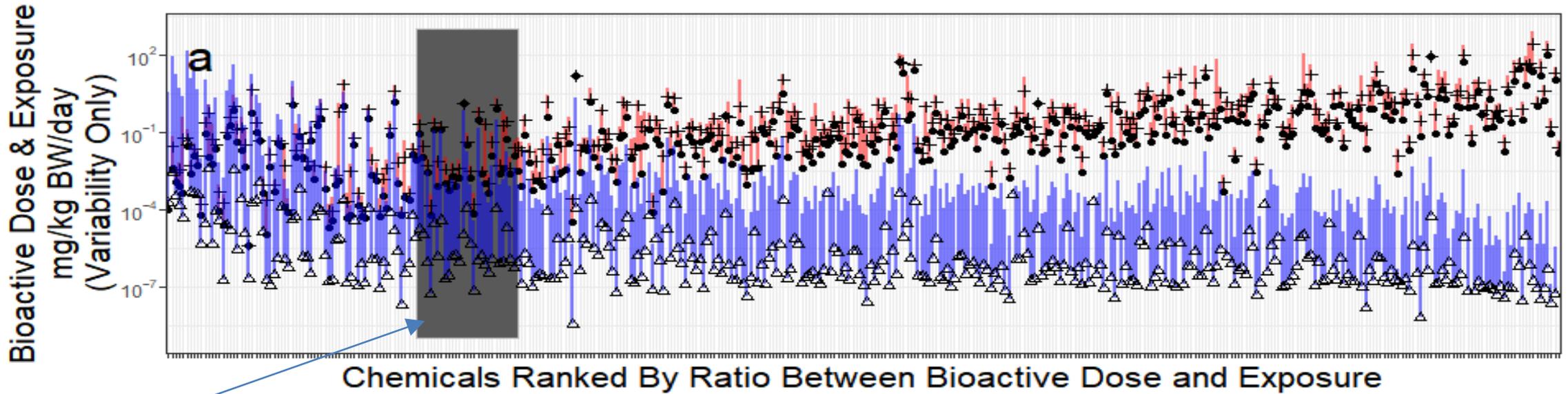
Median Cl: 14 CV: 0.31



Quantifying the Impact of Uncertainty

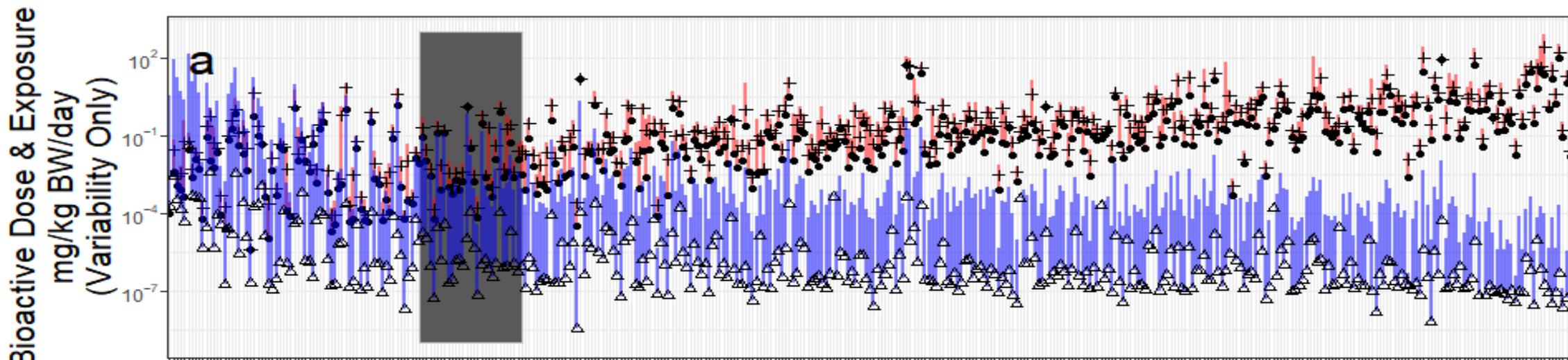


New IVIVE For 393 ToxCast Chemicals

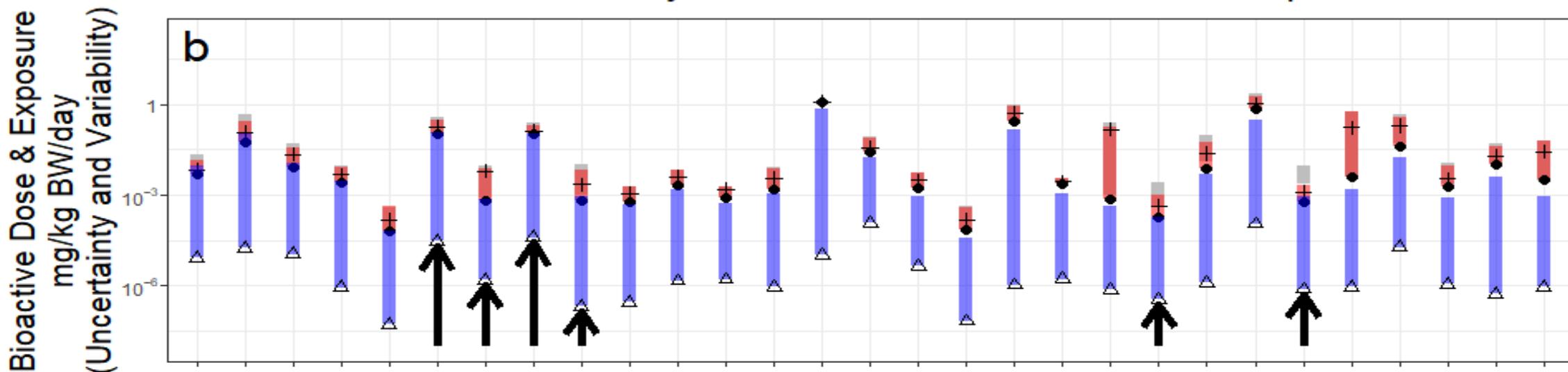


Including chemical-specific uncertainty only caused changes in whether or not exposure and bioactivity overlapped in a small region

The Impact of Measurement Uncertainty



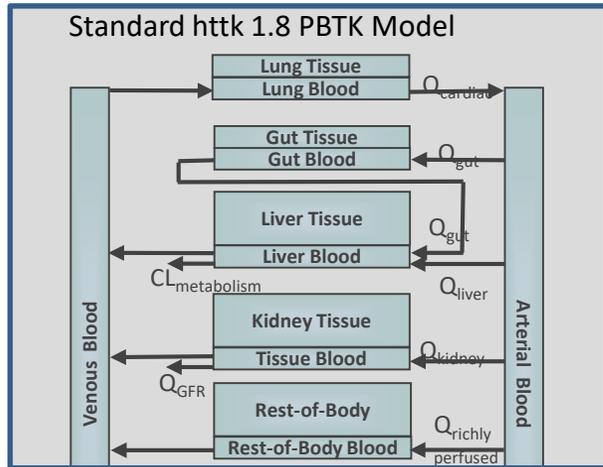
Chemicals Ranked By Ratio Between Bioactive Dose and Exposure



Only six more chemicals overlap

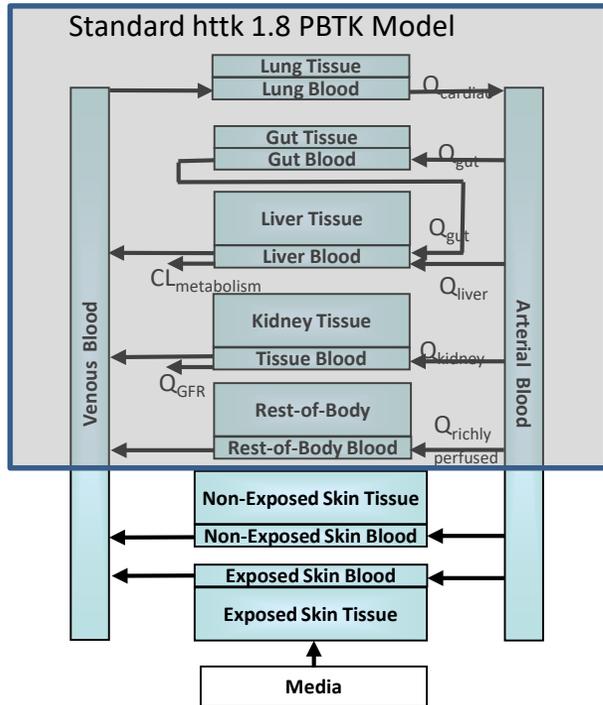
Wambaugh et al. (submitted)

New HT-PBTK Models



- We are working to augment the basic HT-PBTK model with new PBTK models
 - For example, inhalation PBTK will allow for calculation of “inhalation equivalent doses” instead of oral equivalents
- Each model will be released publicly upon peer-reviewed publication
- Pre-publication models can be shared under a MTA
- We assume there will be coding errors and over-simplifications, so each publication involves curation of evaluation data from the scientific literature and through statistical analysis

New HT-PBTK Models

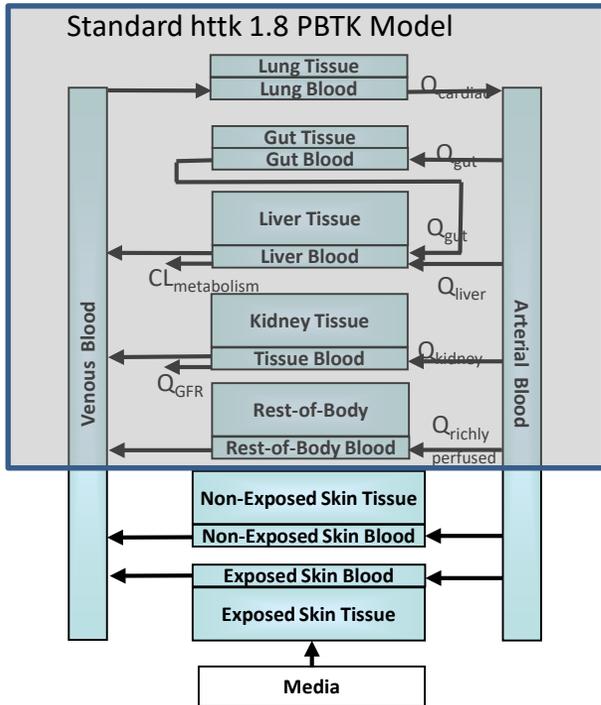
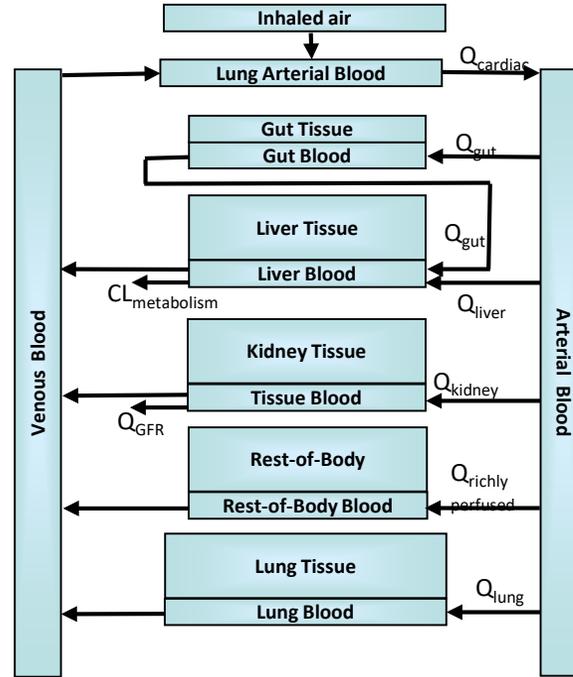


Dermal Exposure Route

EPA, Unilever, INERIS

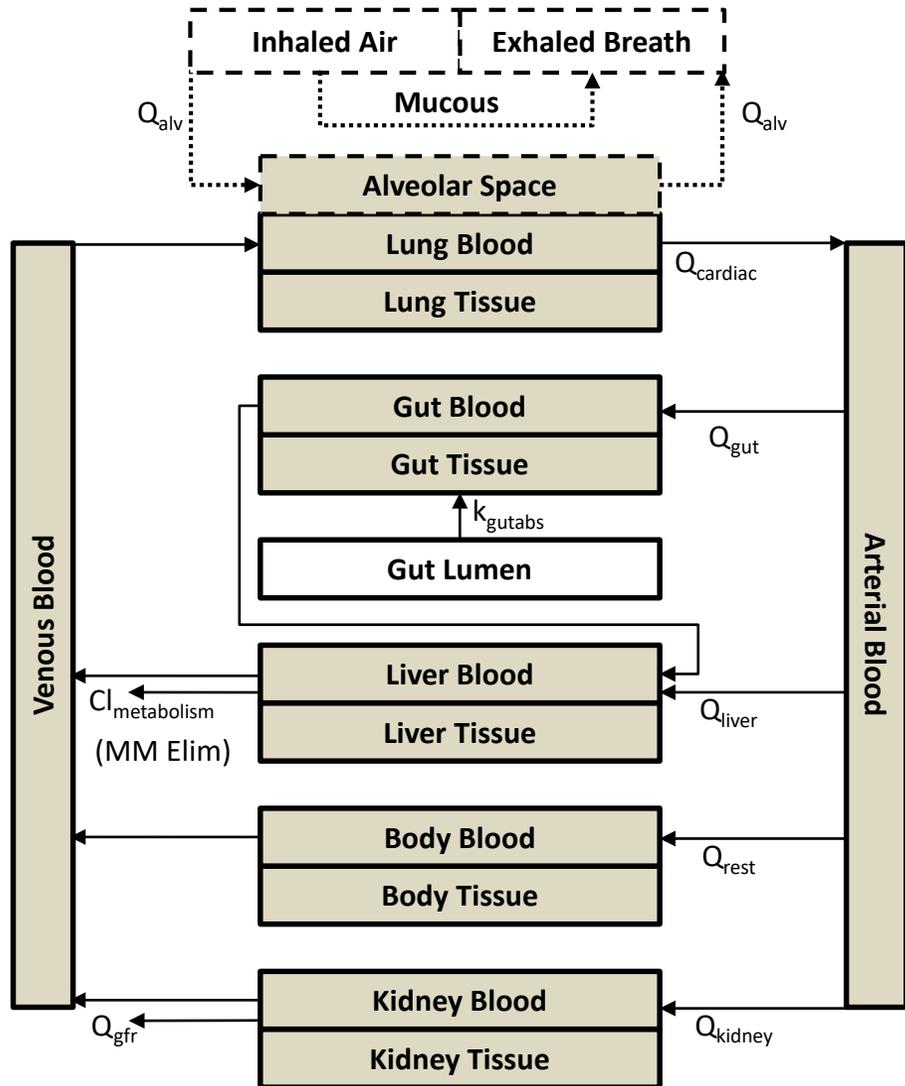
New HT-PBTK Models

Gas Inhalation Exposure Route EPA, USAF

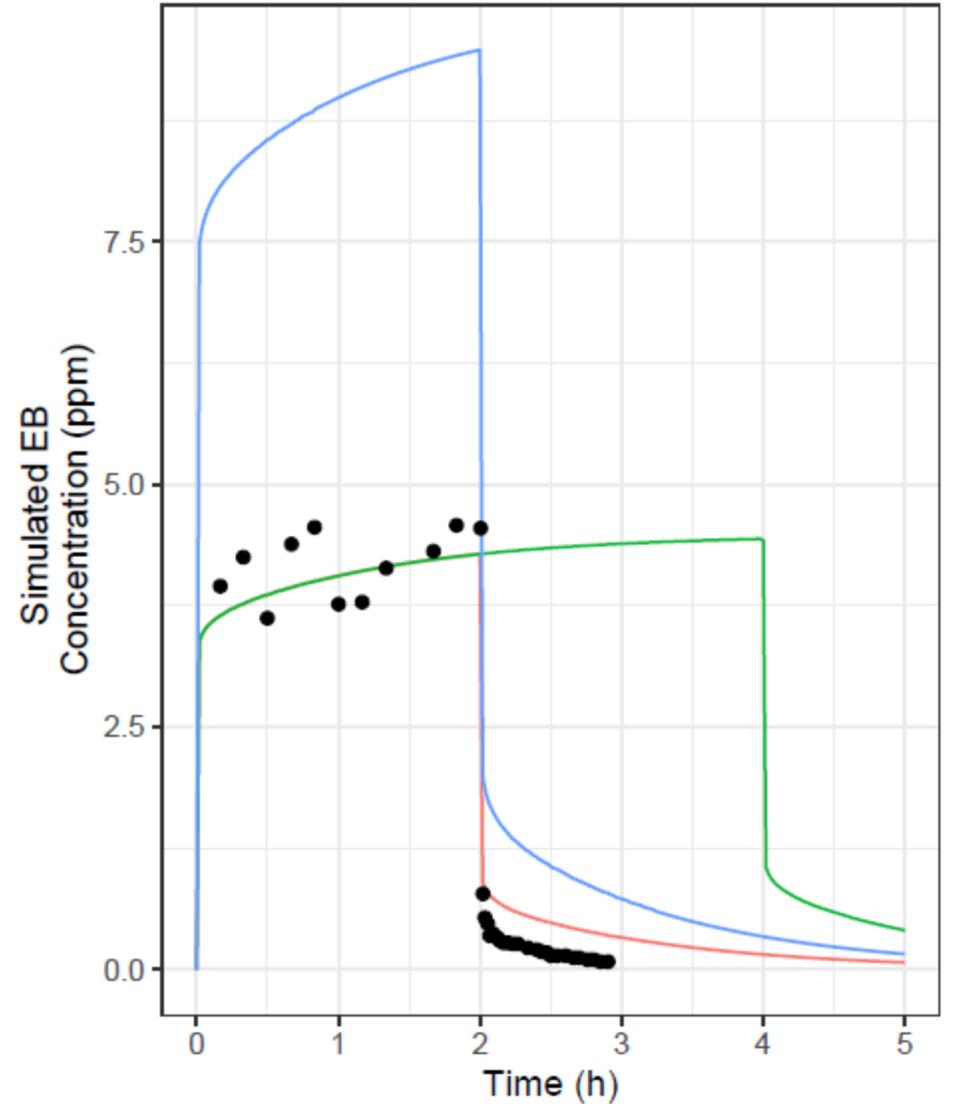


Dermal Exposure Route EPA, Unilever, INERIS

Generic Gas Inhalation Model



1,3-Butadiene (Human, 5ppm for 2h in EB)

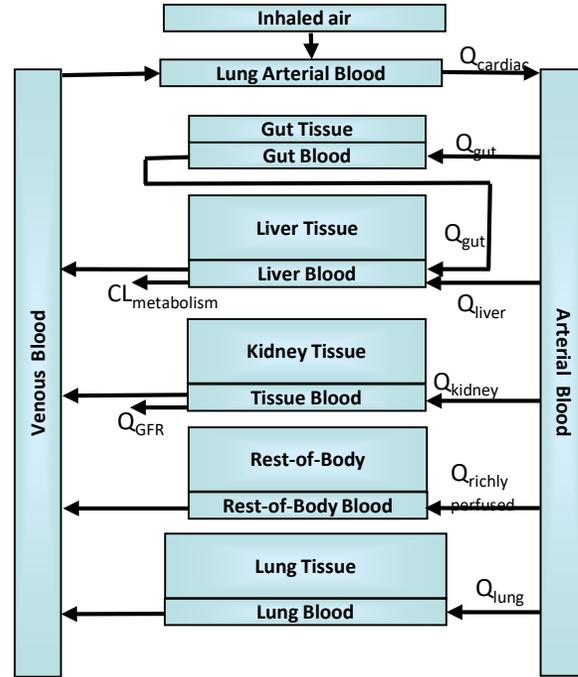


- Correct
- Used 4h exposure instead of 2h
- Used mg/m³ dose units instead of ppm

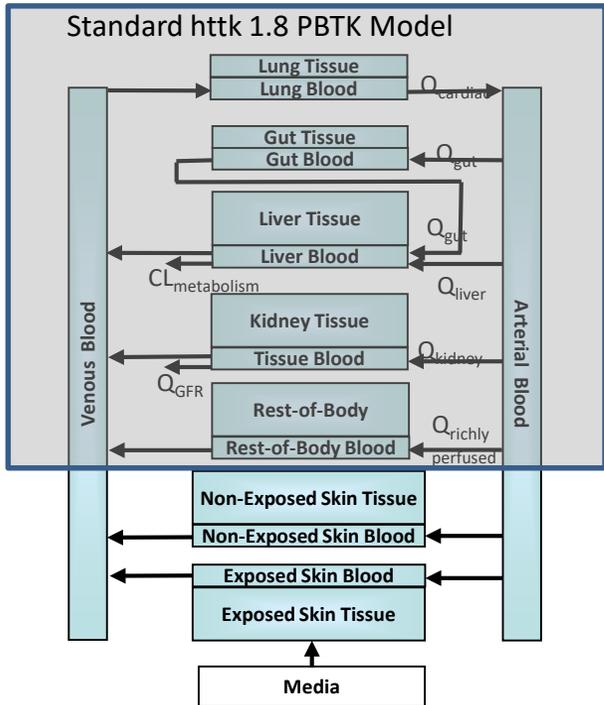
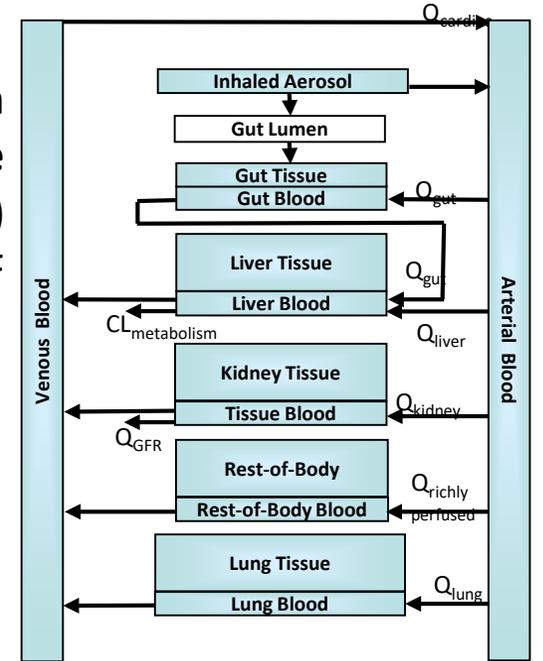
Figure from Matt Linakis (USAFSAM)

New HT-PBTK Models

Gas Inhalation Exposure Route EPA, USAF



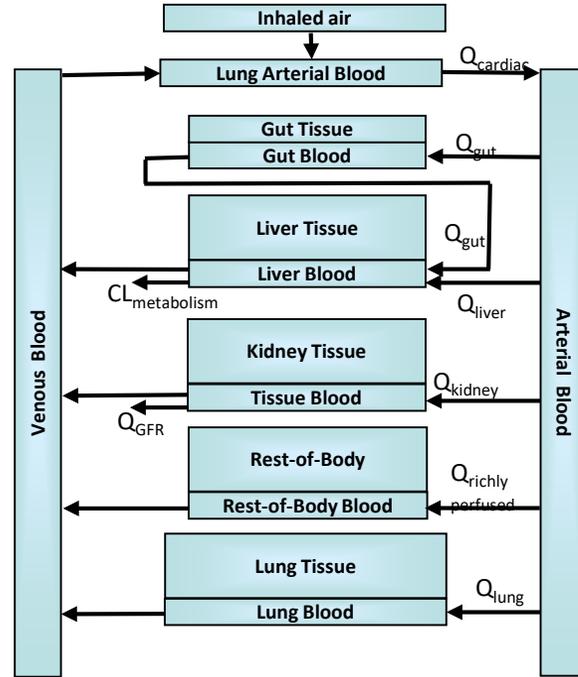
Aerosol Inhalation Exposure Route (with APEX model) EPA, USAF



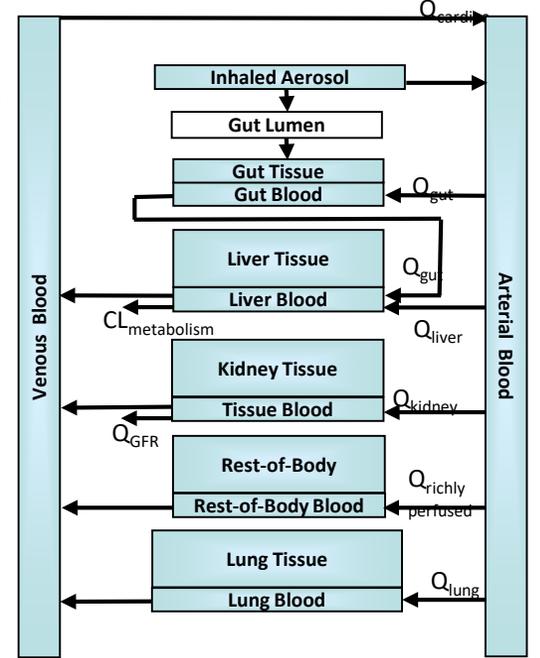
Dermal Exposure Route EPA, Unilever, INERIS

New HT-PBTK Models

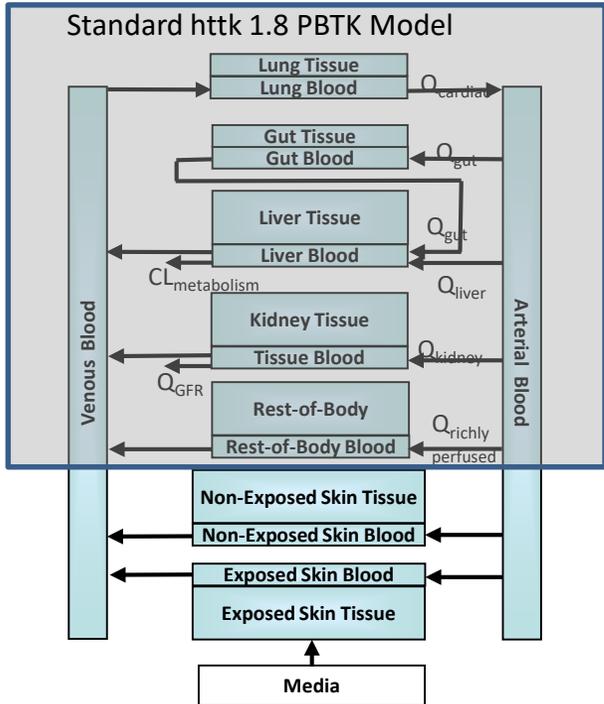
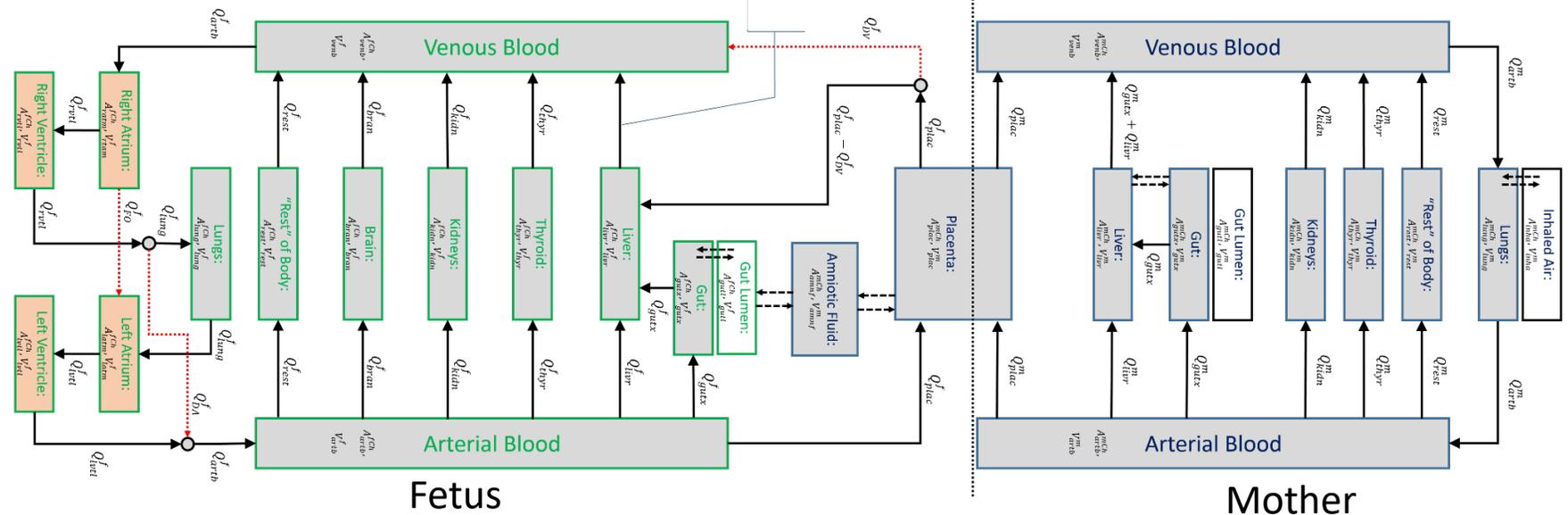
**Gas Inhalation
Exposure Route**
EPA, USAFSAM



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



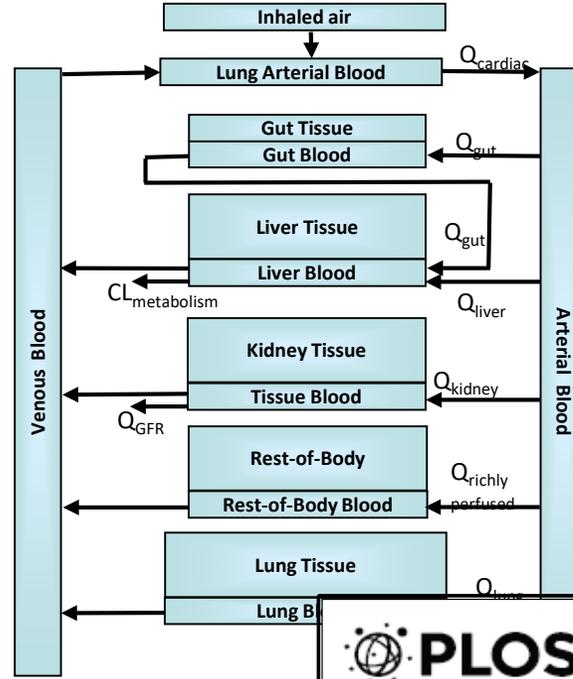
Human Gestational Model
EPA, FDA



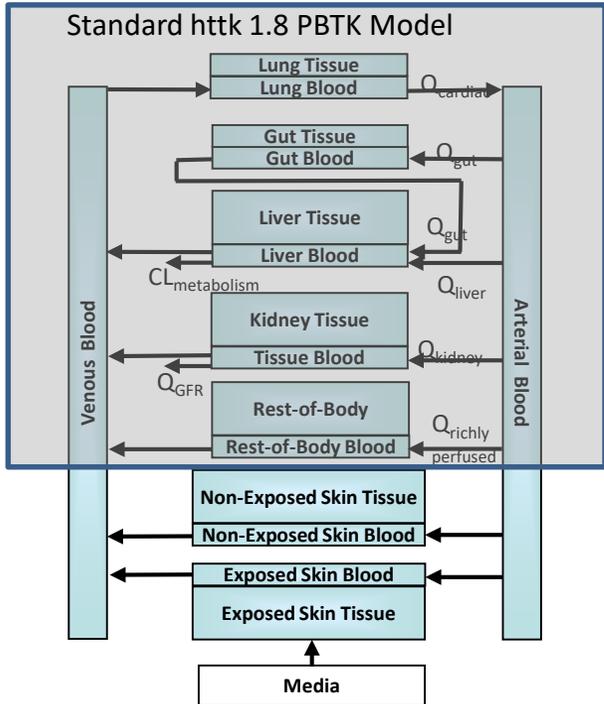
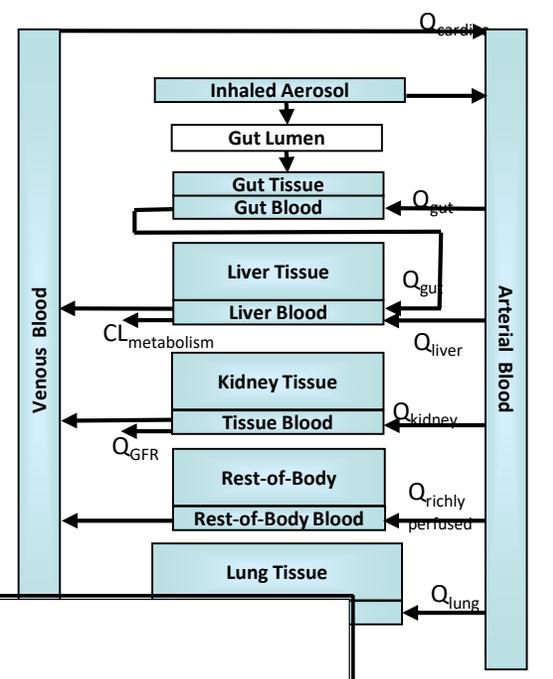
Dermal Exposure Route
EPA, Unilever, INERIS

New HT-PBTK Models

Gas Inhalation Exposure Route EPA, USAFSAM



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



Dermal Exposure Route EPA, Unilever, INERIS

PLOS ONE

RESEARCH ARTICLE

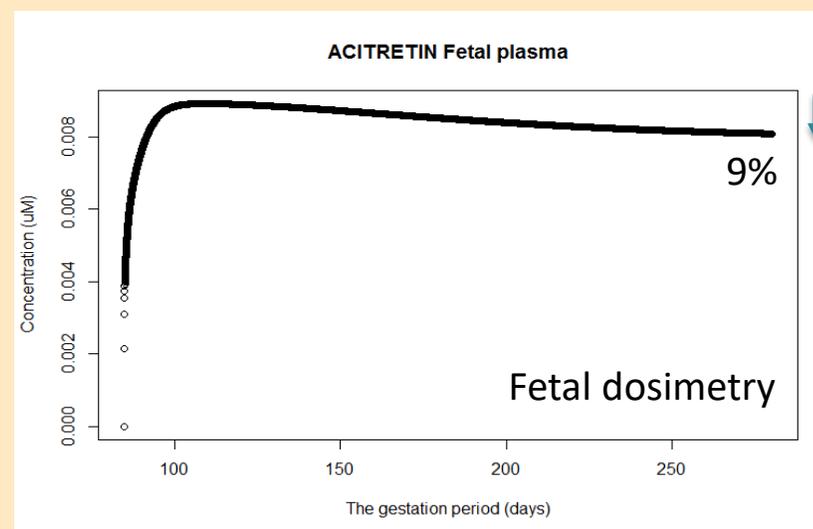
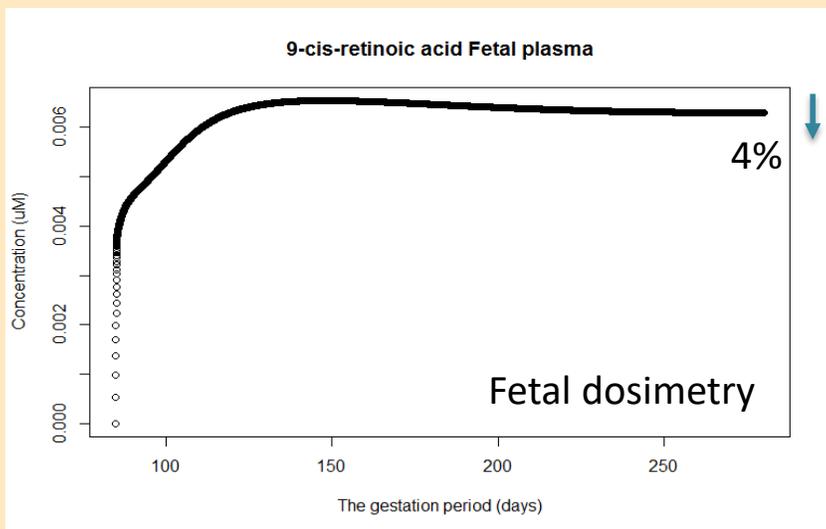
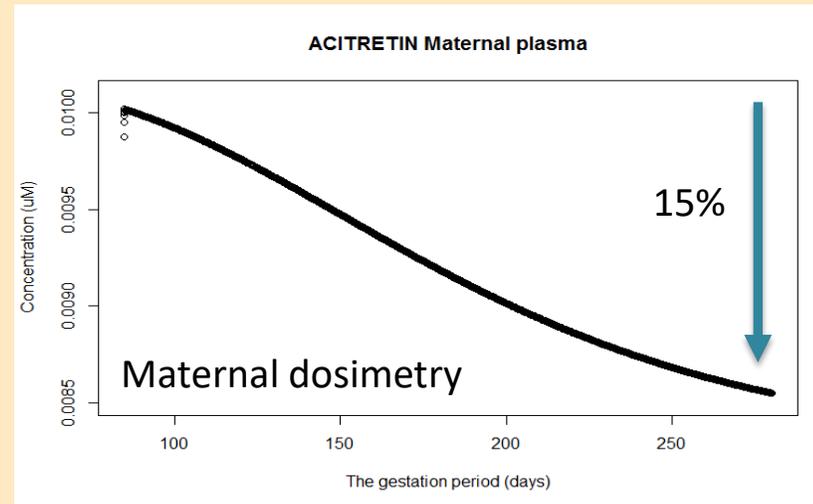
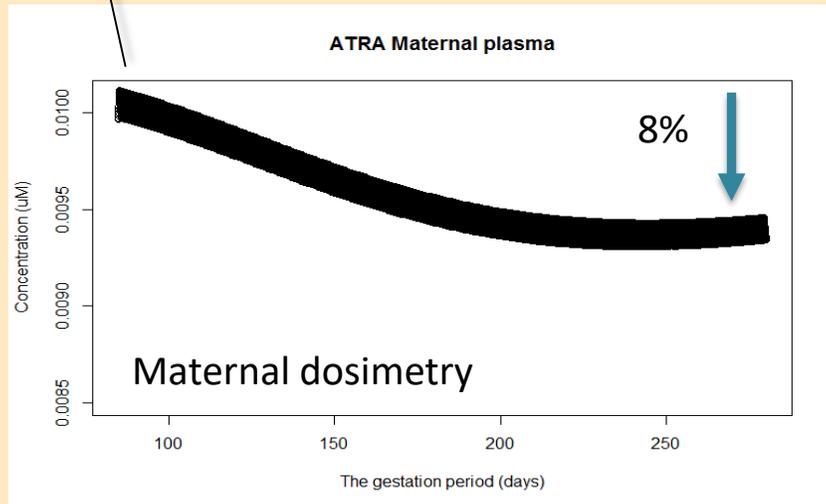
Empirical models for anatomical and physiological changes in a human mother and fetus during pregnancy and gestation

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

Normalized initial plasma concentration for each retinoid analogue

Maternal/Fetal HTTK Model Predictions for Retinoid Analogues:

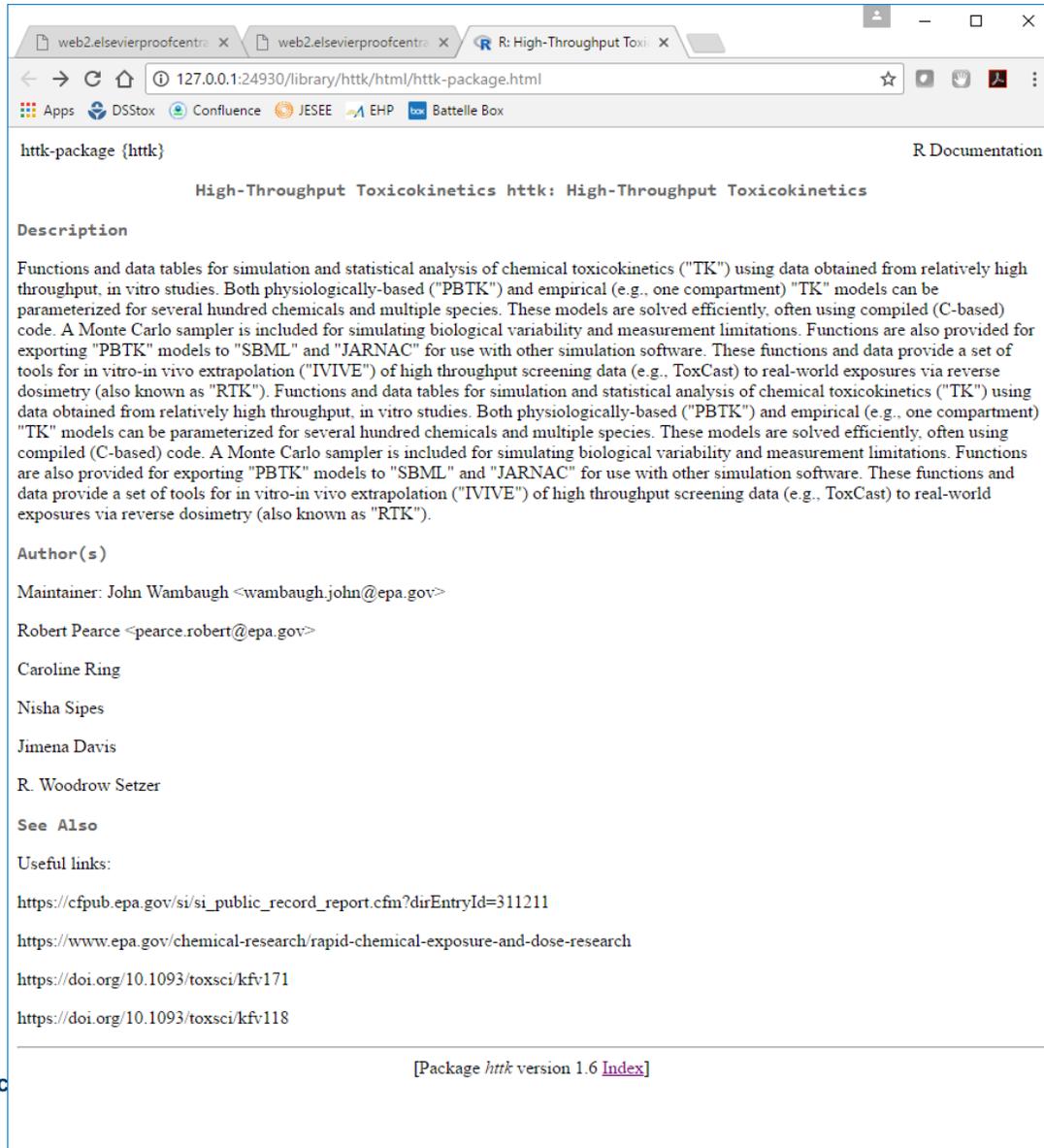


- Decrease in maternal plasma concentrations for retinoid analogues ranged from 8-15%

- Decrease in Fetal plasma concentrations for retinoid analogues ranged from 4-9%

HTTK is (mostly) Documented

Within R: type “help(httk)”



The screenshot shows a web browser window displaying the R documentation for the 'httk' package. The browser tabs include 'web2.elsevierproofcentr...' and 'R: High-Throughput Toxic...'. The address bar shows '127.0.0.1:24930/library/httk/html/httk-package.html'. The page title is 'httk-package {httk}' and it is categorized as 'R Documentation'. The main heading is 'High-Throughput Toxicokinetics httk: High-Throughput Toxicokinetics'. The 'Description' section explains that the package provides functions and data tables for simulation and statistical analysis of chemical toxicokinetics (TK) using data from high-throughput, in vitro studies. It mentions both physiologically-based (PBTK) and empirical (one-compartment) TK models, parameterized for hundreds of chemicals and multiple species. The package includes a Monte Carlo sampler for biological variability and measurement limitations, and functions for exporting PBTK models to SBML and JARNAC. It also provides tools for in vitro-in vivo extrapolation (IVIVE) of high-throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry (RTK). The 'Author(s)' section lists: Maintainer: John Wambaugh <wambaugh.john@epa.gov>, Robert Pearce <pearce.robert@epa.gov>, Caroline Ring, Nisha Sipes, Jimena Davis, and R. Woodrow Setzer. The 'See Also' section is empty. The 'Useful links' section contains: https://cfpub.epa.gov/si/si_public_record_report.cfm?dirEntryId=311211, <https://www.epa.gov/chemical-research/rapid-chemical-exposure-and-dose-research>, <https://doi.org/10.1093/toxsci/kfv171>, and <https://doi.org/10.1093/toxsci/kfv118>. At the bottom, it says '[Package httk version 1.6 [Index](#)]'. The browser's taskbar shows 'Apps', 'DSStox', 'Confluence', 'JESSE', 'EHP', and 'Battelle Box'.

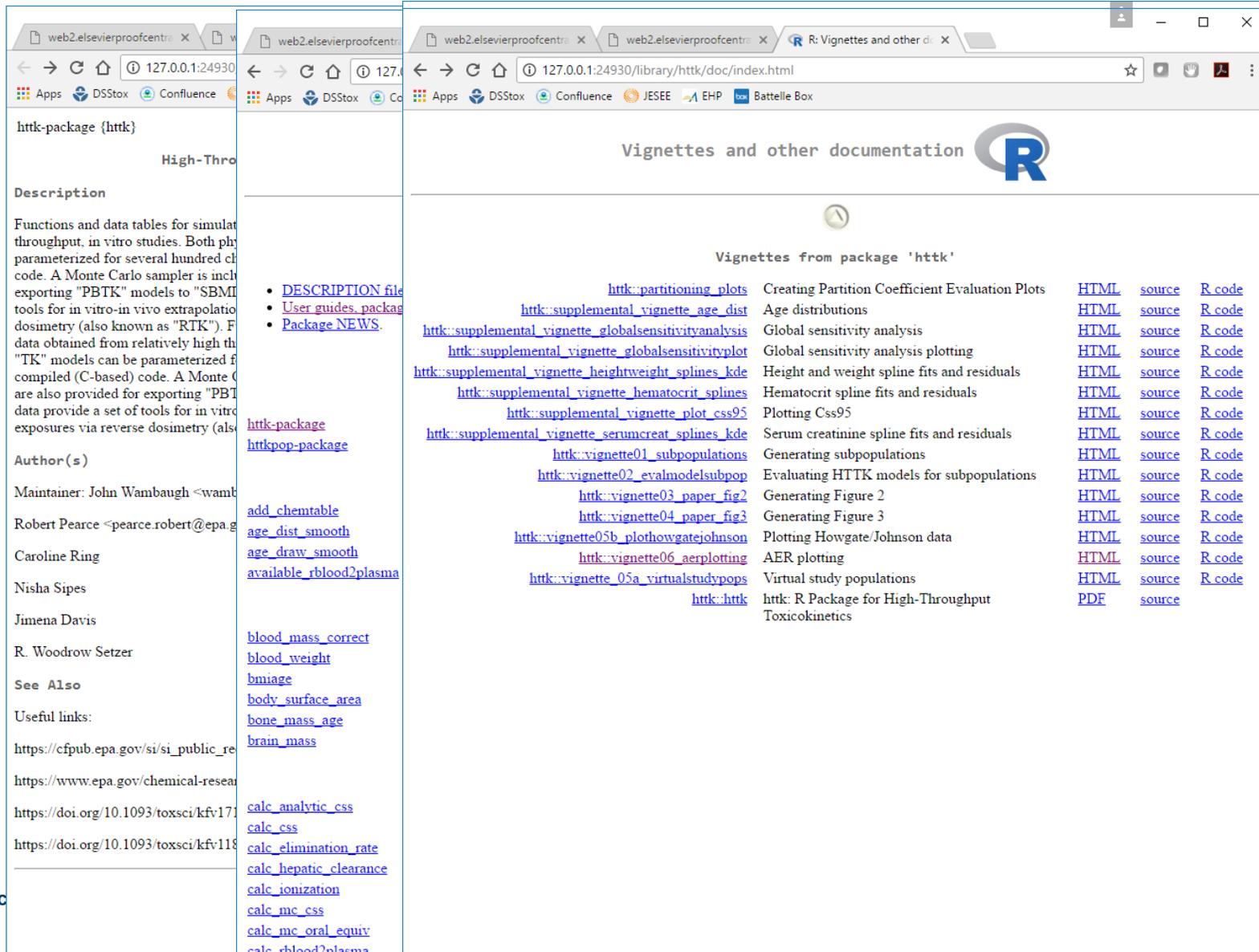
HTTK is (mostly) Documented

Within R: type "help(httk)"

The screenshot shows a web browser window displaying the documentation for the R package 'httk'. The browser address bar shows the URL '127.0.0.1:24930/library/httk/html/00Index.html'. The page title is 'High-Throughput Toxicokinetics' with the R logo. Below the title, it says 'Documentation for package 'httk' version 1.6'. There are three bullet points: 'DESCRIPTION file', 'User guides, package vignettes and other documentation', and 'Package NEWS'. A 'Help Pages' section lists letters A through W. Under 'A', there are two entries: 'httk-package' (High-Throughput Toxicokinetics httk: High-Throughput Toxicokinetics) and 'httkpop-package' (Virtual population generator for HTTK). Under 'B', there are four entries: 'add_chemtable' (Add a table of chemical information for use in making httk predictions), 'age_dist_smooth' (Smoothed age distributions by race and gender), 'age_draw_smooth' (Draws ages from a smoothed distribution for a given gender/race combination), and 'available_rblood2plasma' (Find the best available ratio of the blood to plasma concentration constant). Under 'C', there are six entries: 'blood_mass_correct' (Find average blood masses by age), 'blood_weight' (Predict blood mass), 'bmiage' (CDC BMI-for-age charts), 'body_surface_area' (Predict body surface area), 'bone_mass_age' (Predict bone mass), and 'brain_mass' (Predict brain mass).

HTTK is (mostly) Documented

Within R: type “help(httk)”



The screenshot shows a web browser window with the URL `127.0.0.1:24930/library/httk/doc/index.html`. The page content includes:

httk-package {httk}
High-Throughput Toxicokinetics

Description
Functions and data tables for simulating high-throughput, in vitro studies. Both phylogenetically parameterized for several hundred of code. A Monte Carlo sampler is included for exporting "PBTK" models to "SBMI" models for in vitro-in vivo extrapolation dosimetry (also known as "RTK"). For data obtained from relatively high throughput "TK" models can be parameterized for compiled (C-based) code. A Monte Carlo sampler is also provided for exporting "PBT" data provide a set of tools for in vitro exposures via reverse dosimetry (also known as "RTK").

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Maintainer: John Wambaugh <wambaugh.john@epa.gov>
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Caroline Ring
Nisha Sipes
Jimena Davis
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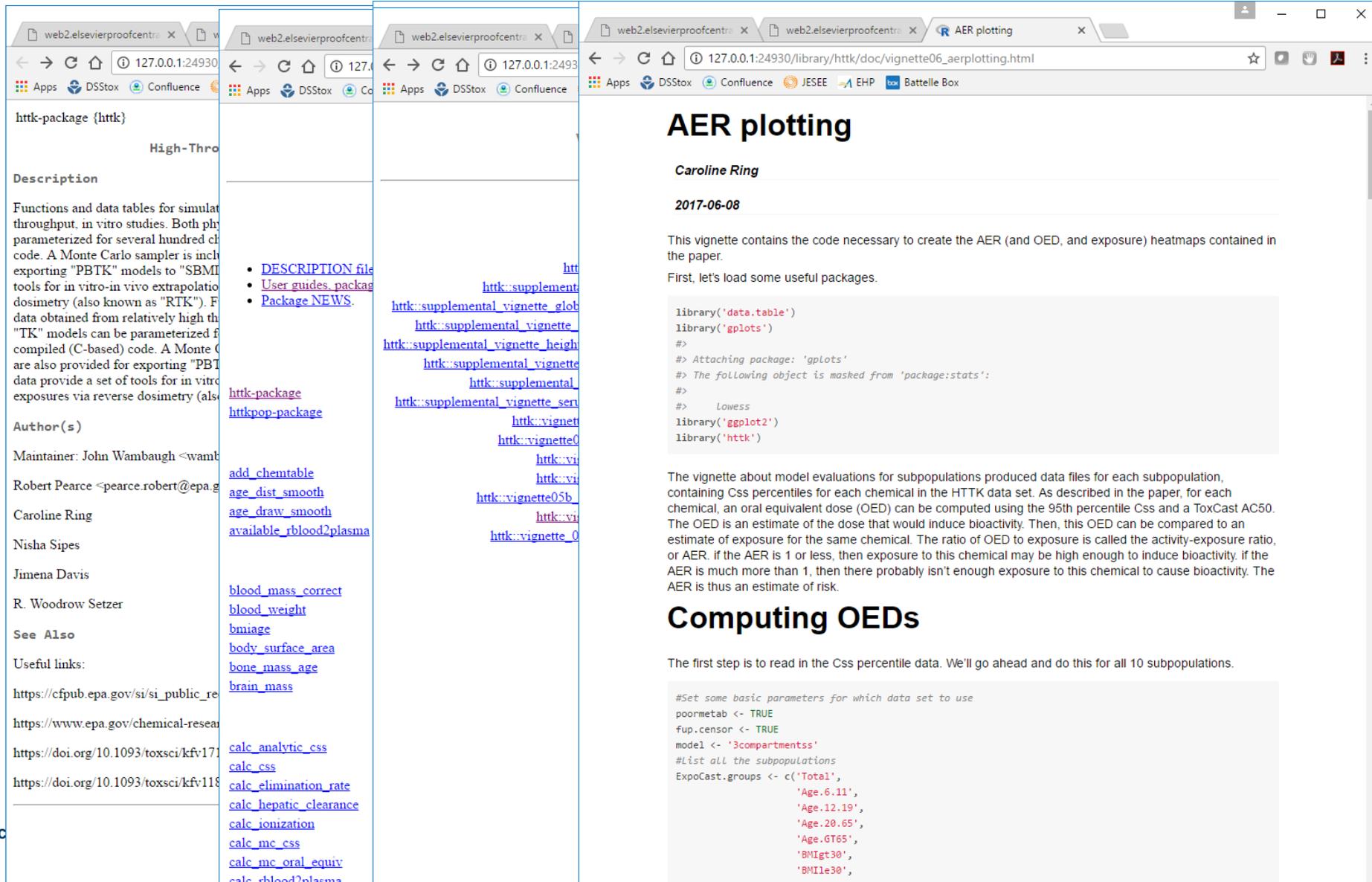
See Also
Useful links:
https://cfpub.epa.gov/si/si_public_research/
<https://www.epa.gov/chemical-research/>
<https://doi.org/10.1093/toxsci/kfv171>
<https://doi.org/10.1093/toxsci/kfv118>

Vignettes from package 'httk'

Vignette Name	Description	HTML	source	R code
http://partitioning_plots	Creating Partition Coefficient Evaluation Plots	HTML	source	R code
http://supplemental_vignette_age_dist	Age distributions	HTML	source	R code
http://supplemental_vignette_globalsensitivityanalysis	Global sensitivity analysis	HTML	source	R code
http://supplemental_vignette_globalsensitivityplot	Global sensitivity analysis plotting	HTML	source	R code
http://supplemental_vignette_heightweight_splines_kde	Height and weight spline fits and residuals	HTML	source	R code
http://supplemental_vignette_hematocrit_splines	Hematocrit spline fits and residuals	HTML	source	R code
http://supplemental_vignette_plot_css95	Plotting Css95	HTML	source	R code
http://supplemental_vignette_serumcreat_splines_kde	Serum creatinine spline fits and residuals	HTML	source	R code
http://vignette01_subpopulations	Generating subpopulations	HTML	source	R code
http://vignette02_evalmodelsubpop	Evaluating HTTK models for subpopulations	HTML	source	R code
http://vignette03_paper_fig2	Generating Figure 2	HTML	source	R code
http://vignette04_paper_fig3	Generating Figure 3	HTML	source	R code
http://vignette05b_plothowgatejohnson	Plotting Howgate/Johnson data	HTML	source	R code
http://vignette06_aerplotting	AER plotting	HTML	source	R code
http://vignette_05a_virtualstudypops	Virtual study populations	HTML	source	R code
http://httk	httk: R Package for High-Throughput Toxicokinetics	PDF	source	

HTTK is (mostly) Documented

Within R: type "help(httk)"



The image shows a screenshot of an R help environment with two windows open. The left window displays the help page for the 'httk' package, and the right window displays the 'AER plotting' vignette.

httk-package {httk}

High-Throughput

Description

Functions and data tables for simulating high-throughput, in vitro studies. Both phylogenetically parameterized for several hundred of code. A Monte Carlo sampler is included for exporting "PBTK" models to "SBMI" models for in vitro-in vivo extrapolation dosimetry (also known as "RTK"). Functions for data obtained from relatively high throughput "TK" models can be parameterized for compiled (C-based) code. A Monte Carlo sampler is also provided for exporting "PBT" data provide a set of tools for in vitro exposures via reverse dosimetry (also known as "RTK").

Author(s)

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See Also

Useful links:

- https://cfpub.epa.gov/si/si_public_research.cfm
- <https://www.epa.gov/chemical-research>
- <https://doi.org/10.1093/toxsci/kfv171>
- <https://doi.org/10.1093/toxsci/kfv118>

Additional links:

- [DESCRIPTION file](#)
- [User guides, packages, and vignettes](#)
- [Package NEWS](#)
- [httk-package](#)
- [httkpop-package](#)
- [add_chemtable](#)
- [age_dist_smooth](#)
- [age_draw_smooth](#)
- [available_rblood2plasma](#)
- [blood_mass_correct](#)
- [blood_weight](#)
- [bmiage](#)
- [body_surface_area](#)
- [bone_mass_age](#)
- [brain_mass](#)
- [calc_analytic_css](#)
- [calc_css](#)
- [calc_elimination_rate](#)
- [calc_hepatic_clearance](#)
- [calc_ionization](#)
- [calc_mc_css](#)
- [calc_mc_oral_equiv](#)
- [calc_rblood2plasma](#)

AER plotting

Caroline Ring

2017-06-08

This vignette contains the code necessary to create the AER (and OED, and exposure) heatmaps contained in the paper.

First, let's load some useful packages.

```
library('data.table')
library('gplots')
#> Attaching package: 'gplots'
#> The following object is masked from 'package:stats':
#>
#> lowess
library('ggplot2')
library('httk')
```

The vignette about model evaluations for subpopulations produced data files for each subpopulation, containing Css percentiles for each chemical in the HTTK data set. As described in the paper, for each chemical, an oral equivalent dose (OED) can be computed using the 95th percentile Css and a ToxCast AC50. The OED is an estimate of the dose that would induce bioactivity. Then, this OED can be compared to an estimate of exposure for the same chemical. The ratio of OED to exposure is called the activity-exposure ratio, or AER. If the AER is 1 or less, then exposure to this chemical may be high enough to induce bioactivity; if the AER is much more than 1, then there probably isn't enough exposure to this chemical to cause bioactivity. The AER is thus an estimate of risk.

Computing OEDs

The first step is to read in the Css percentile data. We'll go ahead and do this for all 10 subpopulations.

```
#Set some basic parameters for which data set to use
poormetab <- TRUE
fup.censur <- TRUE
model <- '3compartmentss'
#List all the subpopulations
ExpoCast.groups <- c('Total',
                    'Age.6.11',
                    'Age.12.19',
                    'Age.20.65',
                    'Age.GT65',
                    'BMIgt30',
                    'BMIle30',
```

Does My Chemical Have HTTK Data?

Is a chemical available?

```
> "80-05-7" %in% get_cheminfo()
[1] TRUE
```

All data on chemicals A, B, C

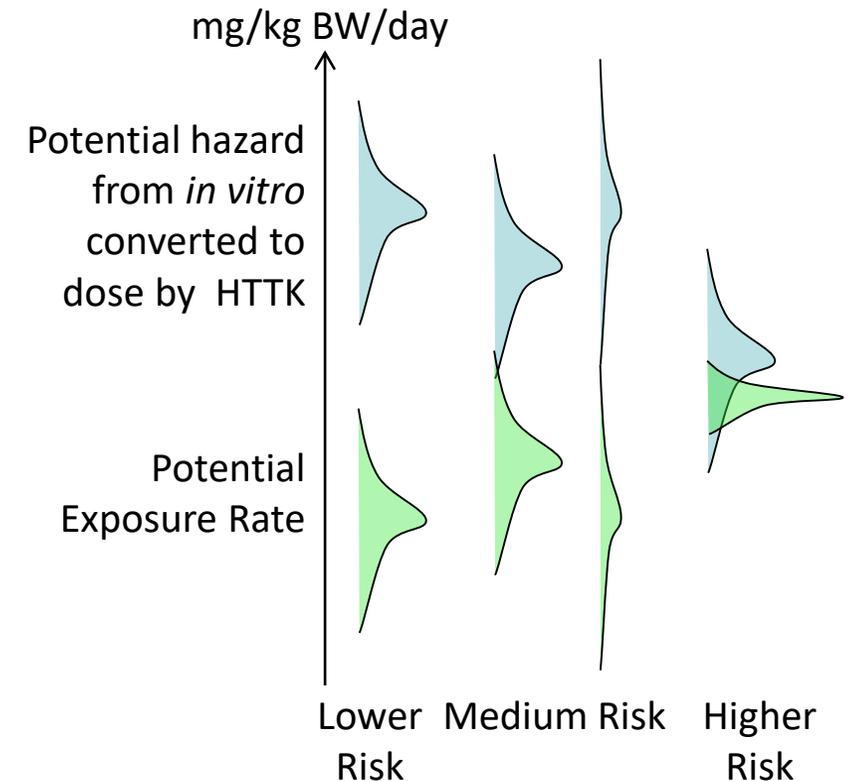
```
subset(get_cheminfo(in
fo="all"), Compound%in%
c("A", "B", "C"))
```

```
> library(httk)
> get_cheminfo()
 [1] "2971-36-0"    "94-75-7"      "94-82-6"      "90-43-7"      "1007-28-9"
 [6] "71751-41-2"  "30560-19-1"  "135410-20-7"  "34256-82-1"   "50594-66-6"
[11] "15972-60-8"  "116-06-3"    "834-12-8"     "33089-61-1"   "101-05-3"
[16] "1912-24-9"   "86-50-0"     "131860-33-8"  "22781-23-3"   "1861-40-1" ...
> get_cheminfo(info="all")
```

Compound	CAS	logP	pKa_Accept	pKa_Donor	MW	Human.Clint	Human.Clint.p Value	Human.Funbou nd.plasma	DSSTox_Substance_I d	Structure_Formula	Substance_Type
2,4-d	94-75-7	2.81	<NA>	2.81	221.03	0	0.149	0.04	DTXSID0020442	C8H6Cl2O3	Single Compound
2,4-db	94-82-6	3.53	<NA>	4.5	249.09	0	0.104	0.01	DTXSID7024035	C10H10Cl2O3	Single Compound
2-phenylphenol	90-43-7	3.09	<NA>	10.6	170.211	2.08	0.164	0.04	DTXSID2021151	C12H10O	Single Compound
6-desisopropylatrazine	1007-28-9	1.15	1.59	<NA>	173.6	0	0.539	0.46	DTXSID0037495	C5H8ClN5	Single Compound

Conclusions

- HTTK allows dosimetric adjustment of high-throughput screening (HTS) data across thousands of chemicals.
- New, chemical-specific *in vitro* experiments have been conducted by Cyprotex, using a revised protocol for measuring protein binding
- Overall, variability contributed more significantly to C_{ss} estimations of the 95th percentile
- Comparison between high throughput toxicokinetics (HTTK) predicted concentrations and *in vivo* data is a valuable approach for evaluation and establishing confidence
- Recent analyses indicate that some properties (e.g. average and maximum concentration) can be predicted with confidence.
 - A new database of *in vivo* concentration vs. time data is being developed (Sayre, in preparation)



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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