

# The High Throughput Toxicokinetic (HTTK) R Package

John Wambaugh

*National Center for Computational Toxicology*

*Office of Research and Development*

*U.S. Environmental Protection Agency*

**Computational Toxicology  
Community of Practice Webinar**

**June 27, 2019**

Greg Honda

**Oral  
Absorption**

Dustin Kapraun  
Richard Judson  
Annie Lumen (FDA)

Mark Sfeir

**Package Tsar**

**Human  
Gestation**



Derek Angus	Briana Franz
Maria Bacolod	Jon Gilbert
Akshay	Teresa Sierra
Badrinarayanan	Bradley
Adam Brockman	Snodgrass
Roger Dinallo	Chris Strock

**Cyprotex  
(lab work)**

**HTTK Team**

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**TK  
Database**

**Structure-Based  
Predictions**

**Russell Thomas**  
**Barbara Wetmore**  
David Murphy  
Katherine Coutros  
Ann Richard

Risa Sayre  
Chris Grulke

**Dermal**

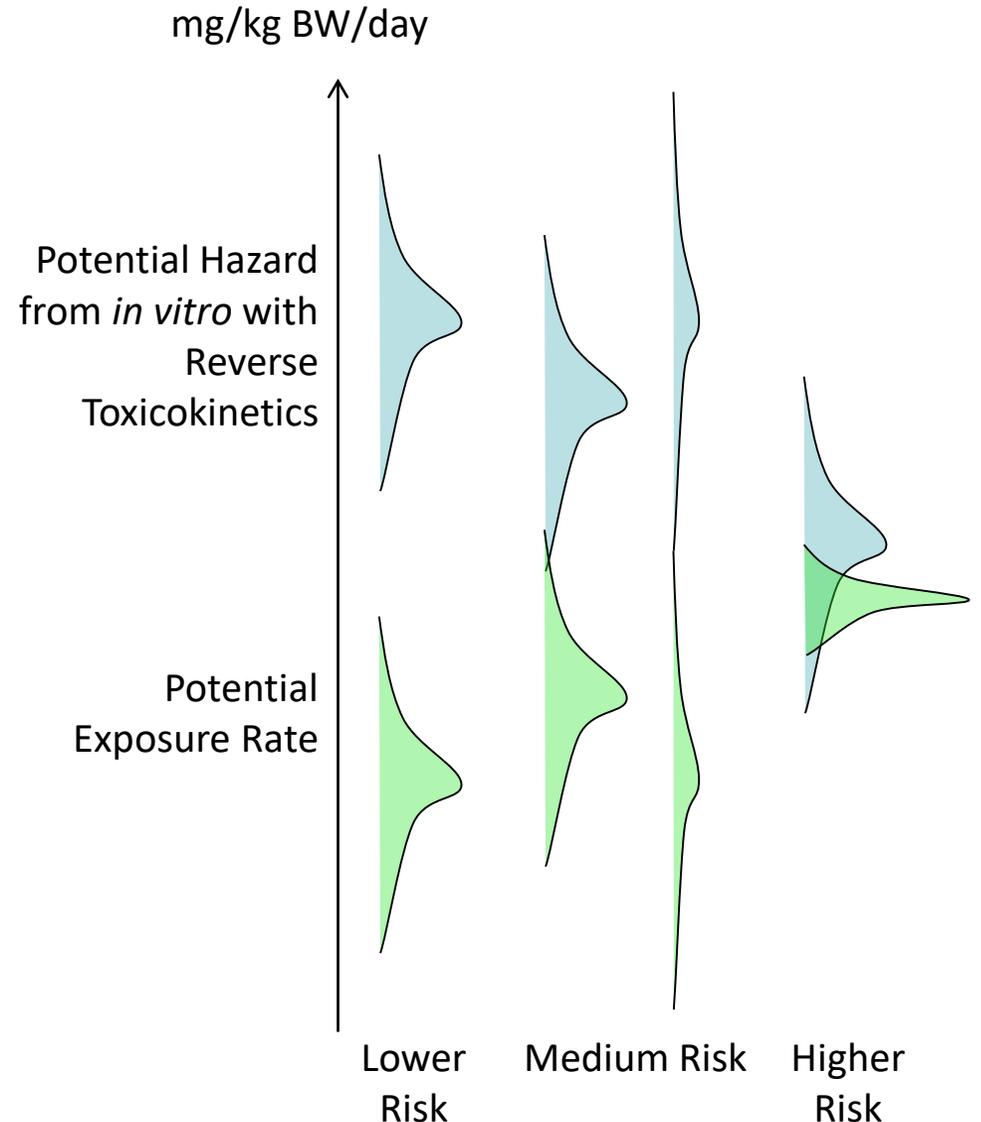
Marina Evans  
Tom Moxon (Unilever)  
Beate Nicol (Unilever)

**Alumni**

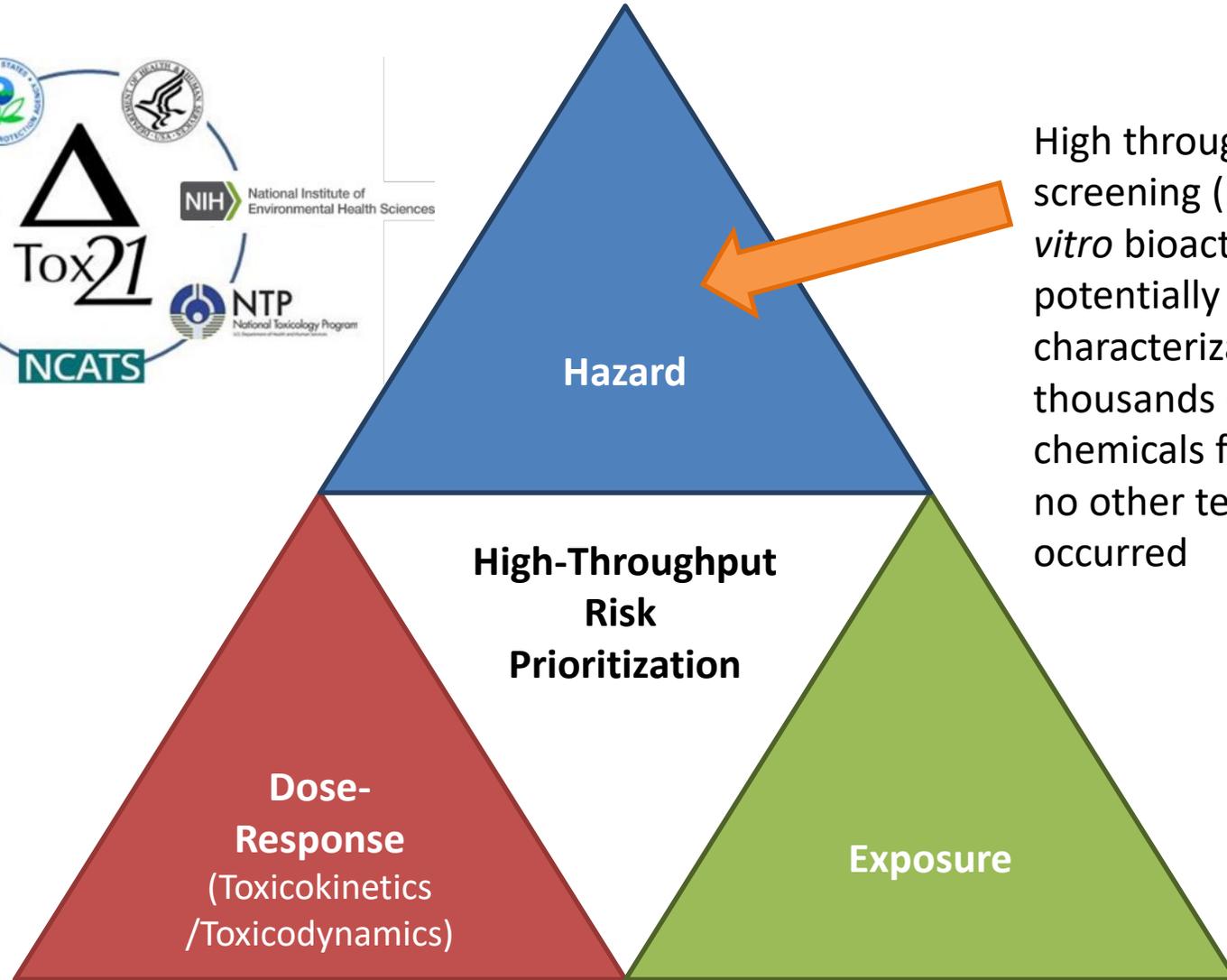
Robert Pearce	Cory Strope
Woody Setzer	Jimena Davis
Caroline Ring	Chantel Nicolas

# Chemical Risk = Hazard x Exposure

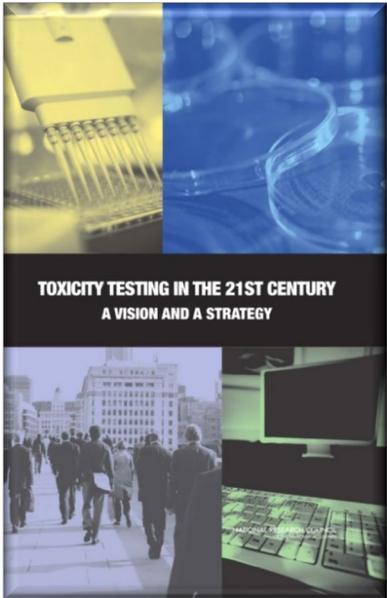
- The U.S. National Research Council (1983) identified chemical risk as a function of both inherent hazard and exposure
- To address the thousands of chemicals in commerce and the environment, we need new approach methodologies (NAMs) that can inform prioritization of chemicals most worthy of additional 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*

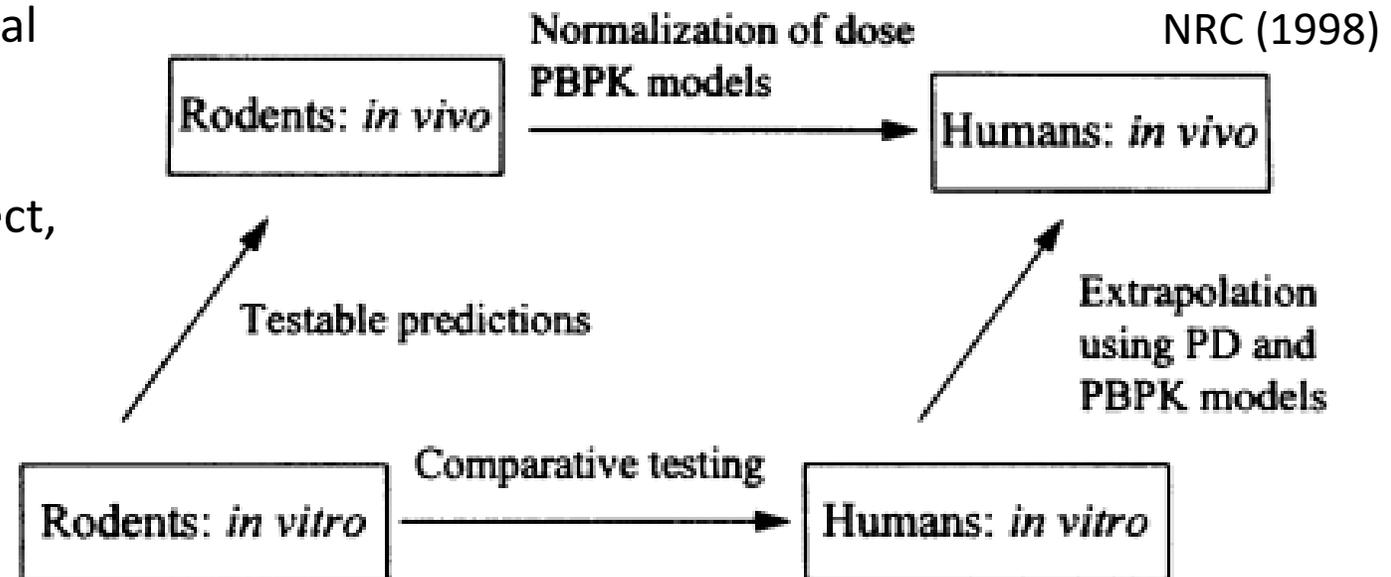
- **IVIVE-PK/TK (Pharmacokinetics/Toxicokinetics):**

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

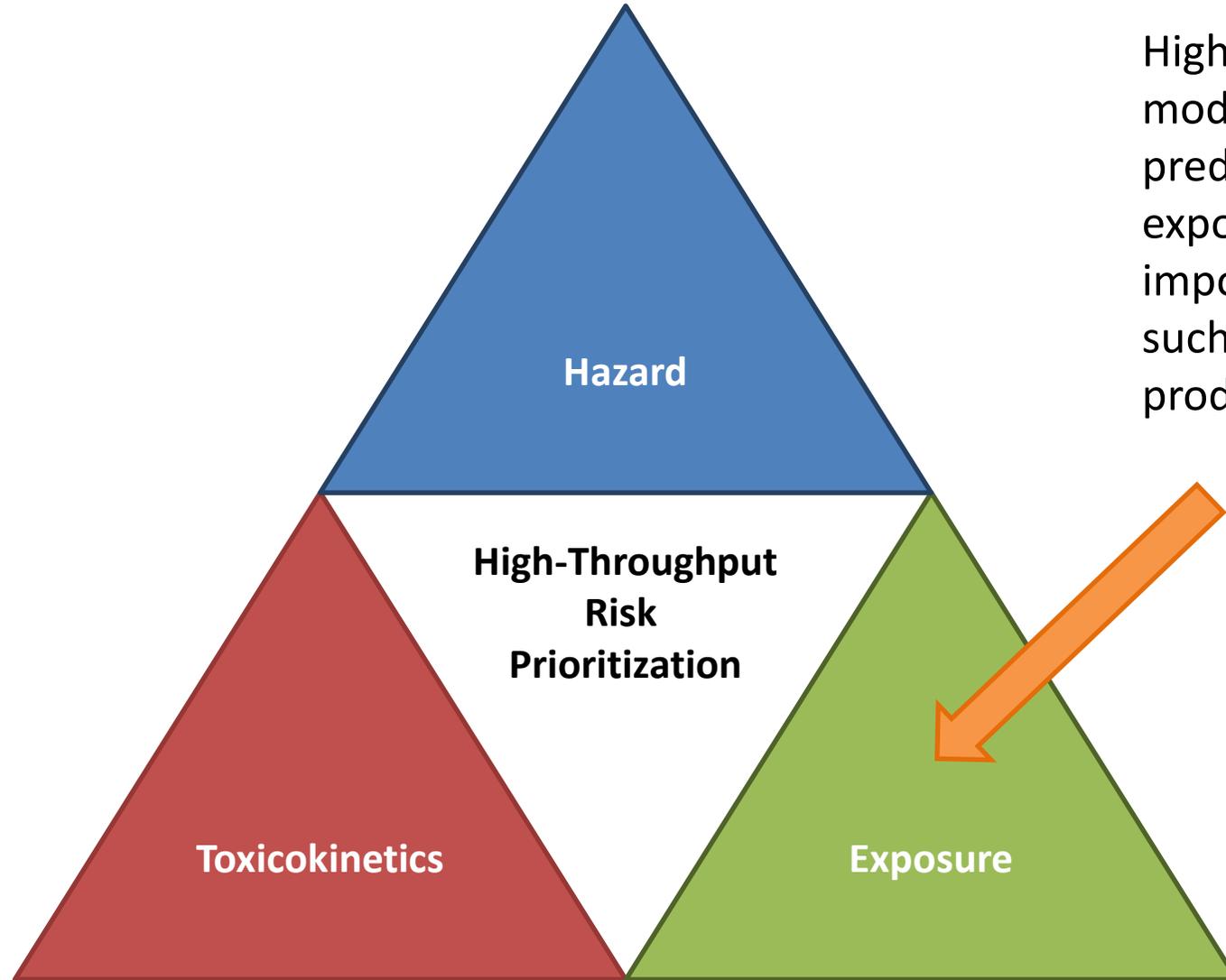
- **IVIVE-PD/TD (Pharmacodynamics/Toxicodynamics):**

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

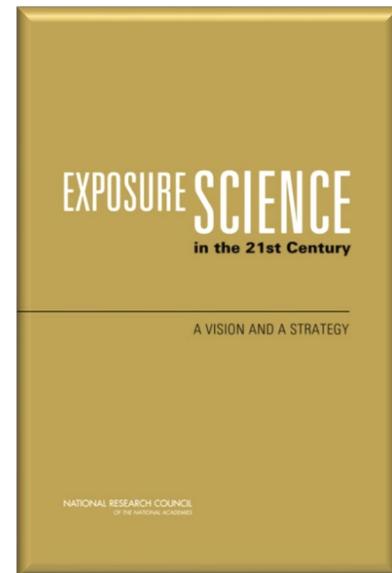
- Both contribute to *in vivo* effect prediction



# New Exposure Data and Models



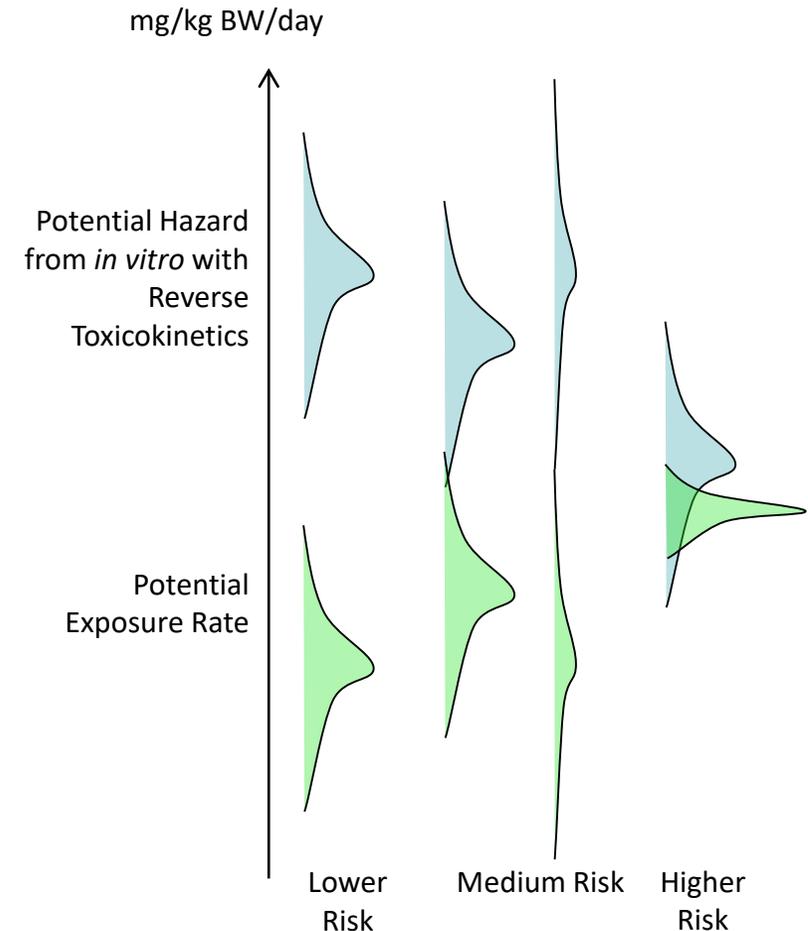
High throughput models exist to make predictions of exposure via specific, important pathways such as residential product use and diet



NRC (2012)

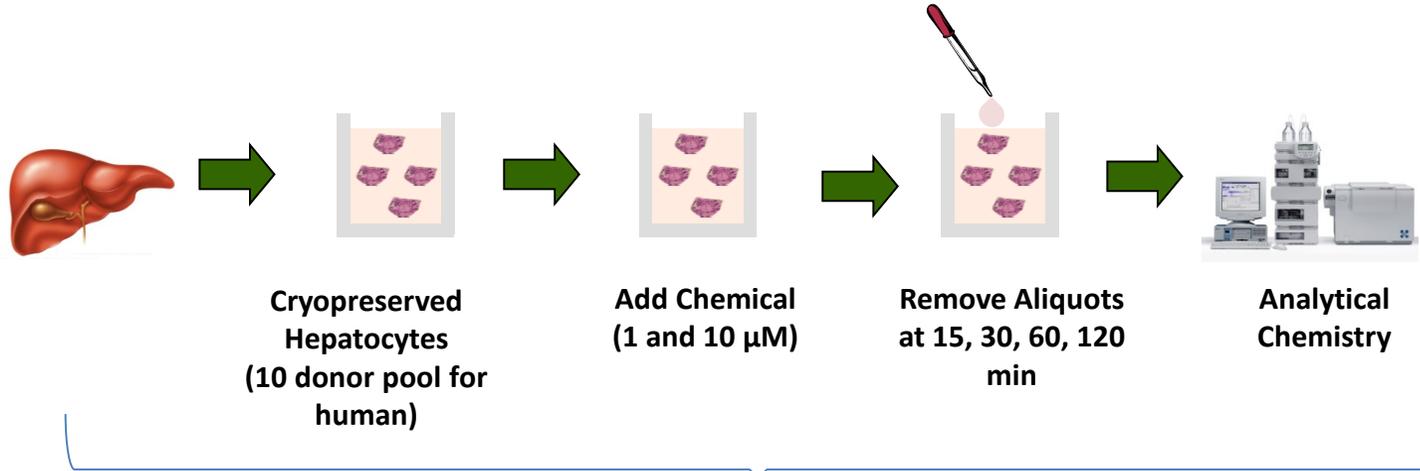
# High Throughput Toxicokinetics (HTTK)

- **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)

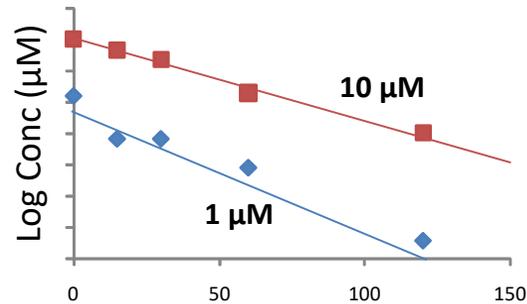


# In Vitro Data for HTTK

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



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

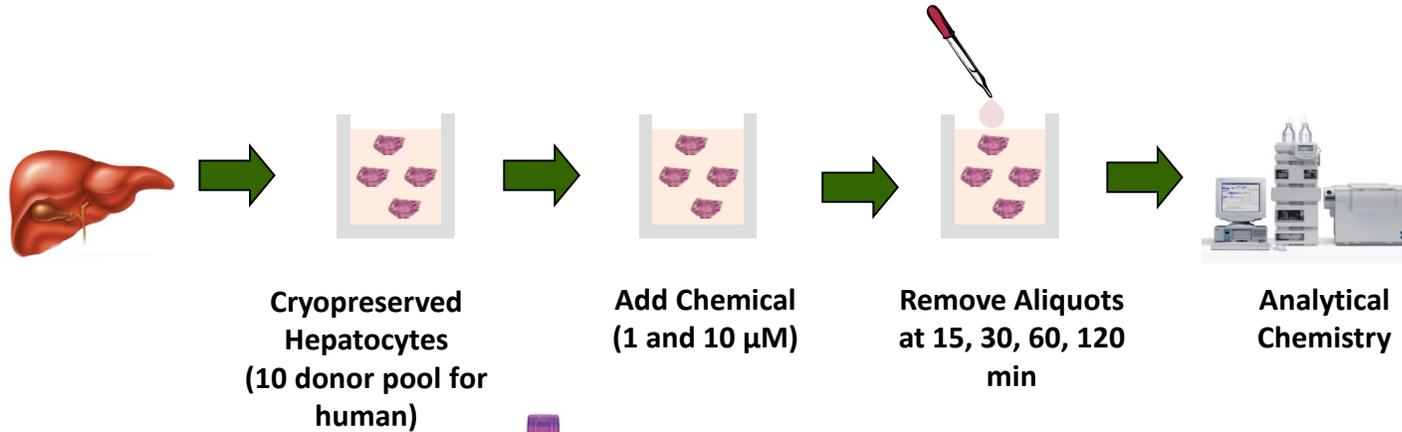


We perform the assay at 1 and 10  $\mu\text{M}$  to check for saturation of metabolizing enzymes.

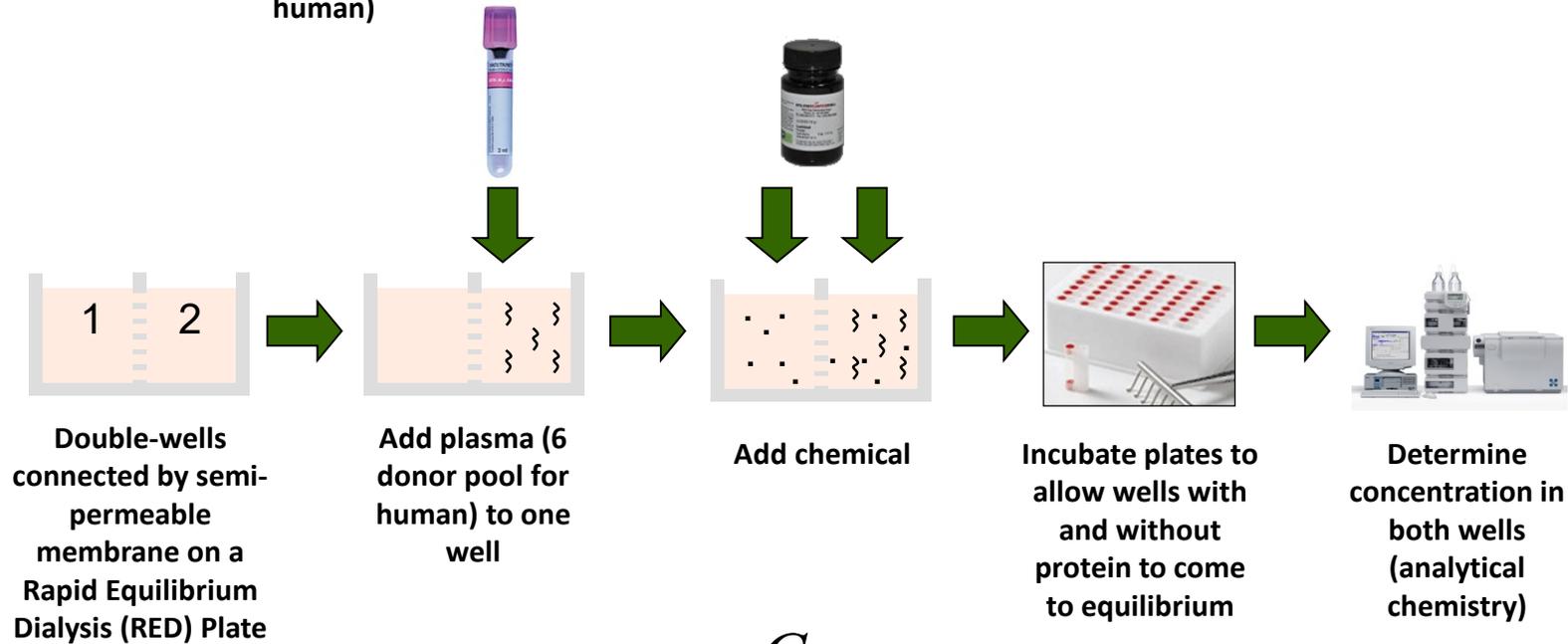
- **Most chemicals do not have TK data** – we use *in vitro* HTTK methods adapted from pharma to fill gaps
- In drug development, HTTK methods allow IVIVE to estimate therapeutic doses for clinical studies – predicted concentrations are typically on the order of values measured in clinical trials (Wang,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

# In Vitro Data for HTTK

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



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

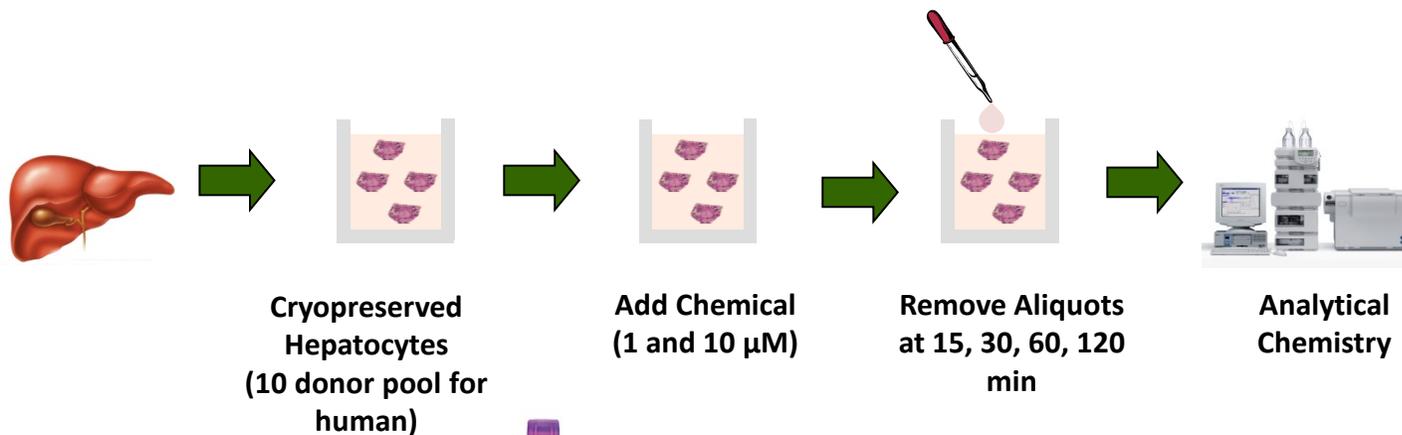


$$F_{ub,p} = \frac{C_{well1}}{C_{well2}}$$

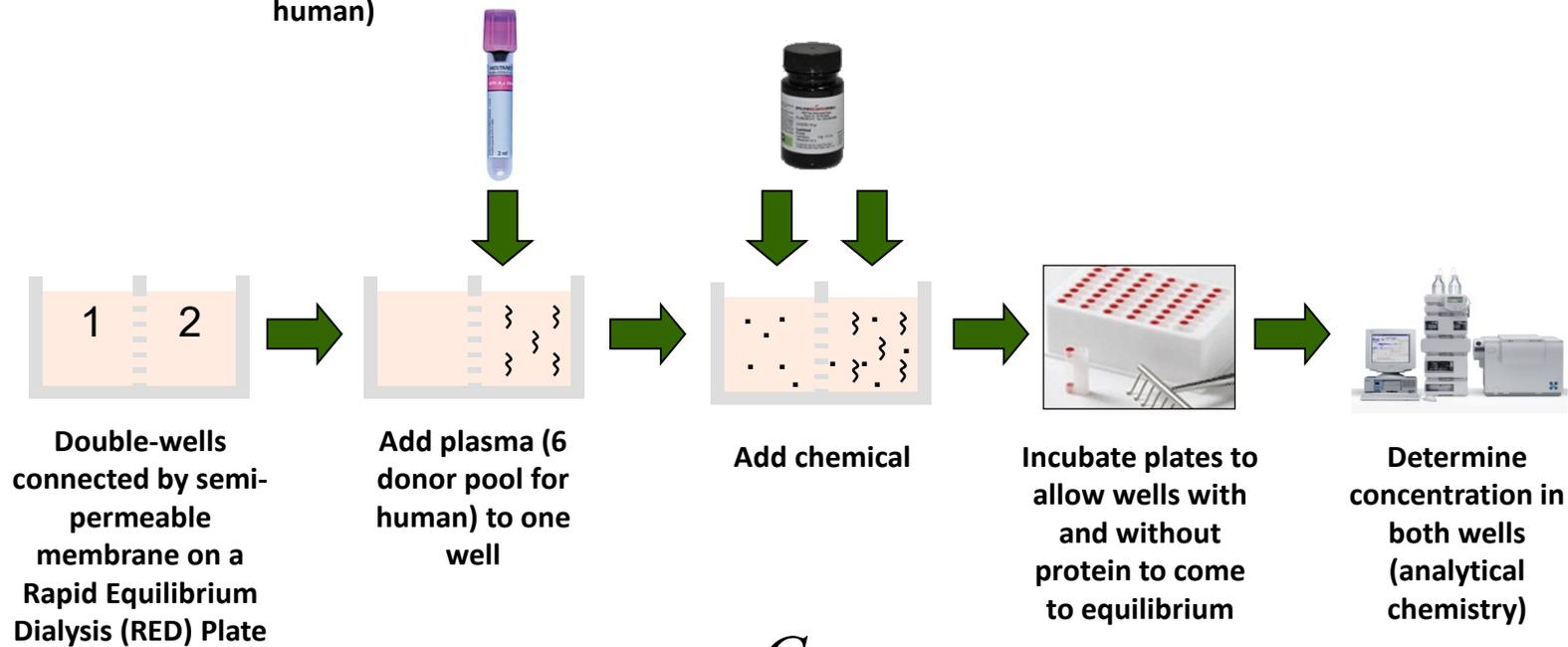
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$$F_{ub,p} = \frac{C_{well1}}{C_{well2}}$$

- 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

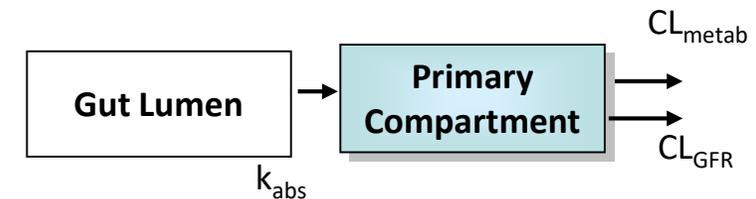
# Simple Model for Steady-State Plasma Concentration ( $C_{ss}$ )

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

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)

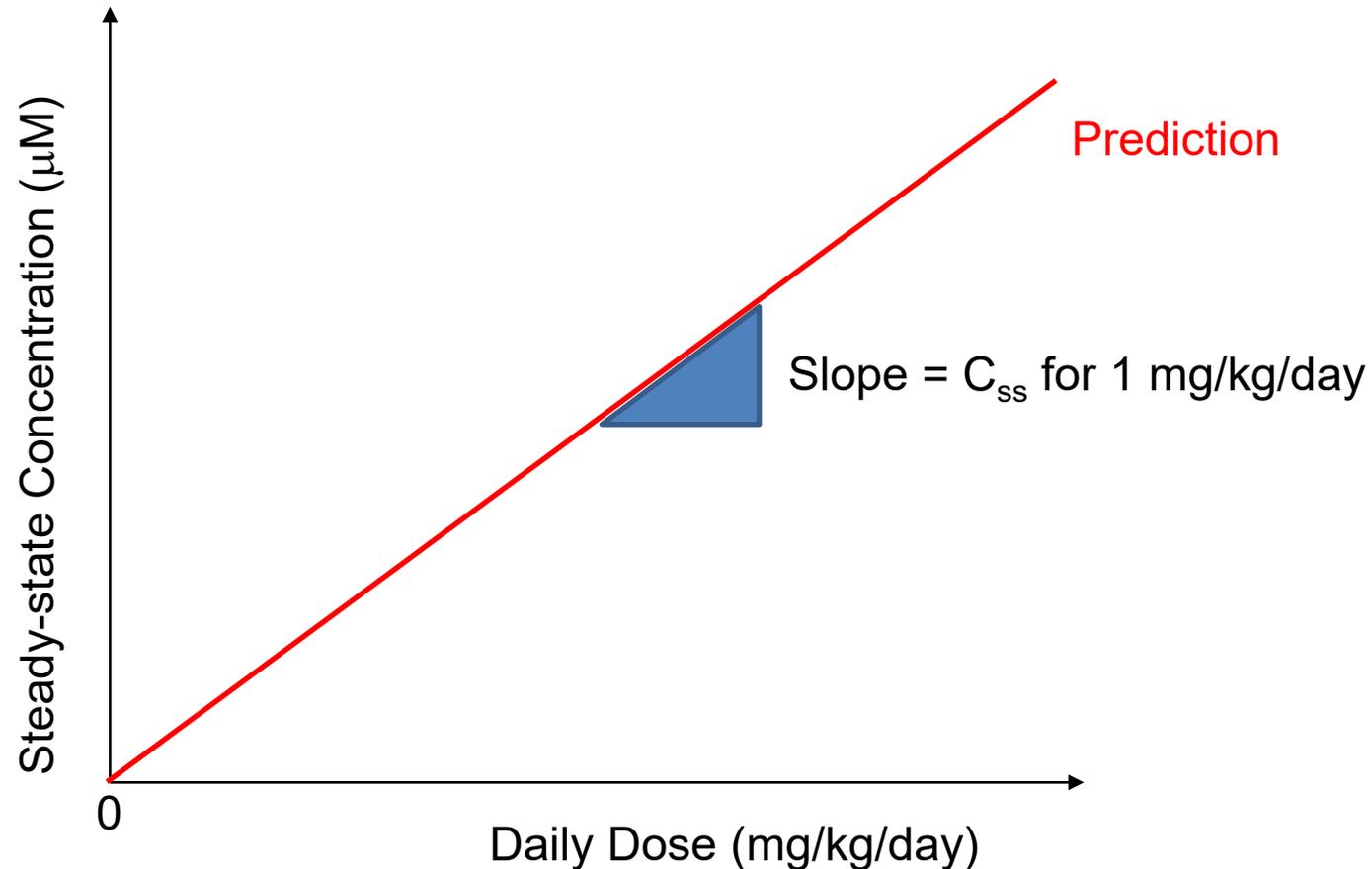


# Assume that Steady-State is Linear with Dose

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

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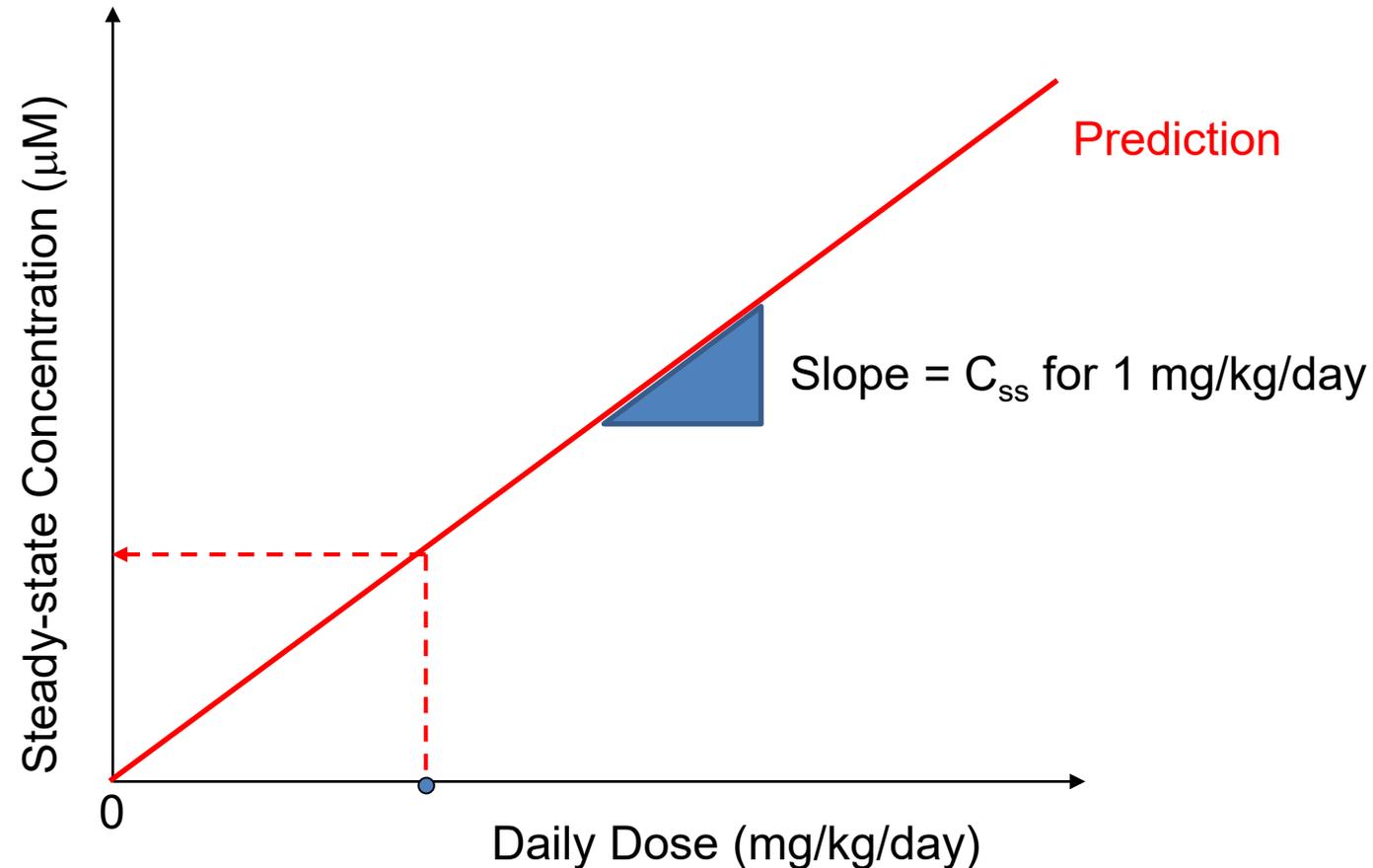
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- Can calculate predicted steady-state concentration ( $C_{ss}$ ) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

# Assume that Steady-State is Linear with Dose

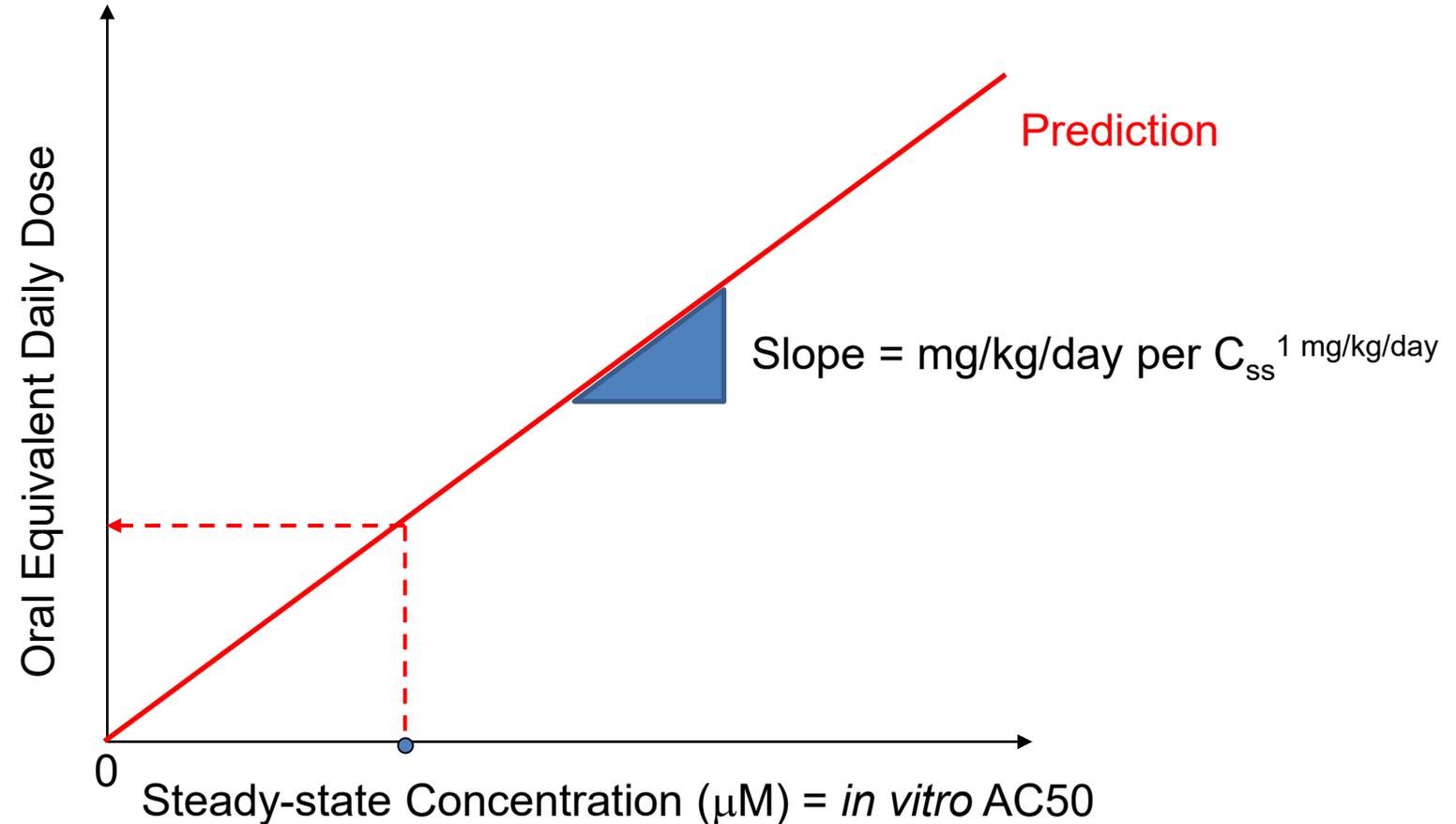
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# HTTK Allows Steady-State In Vitro-In Vivo Extrapolation (IVIVE)

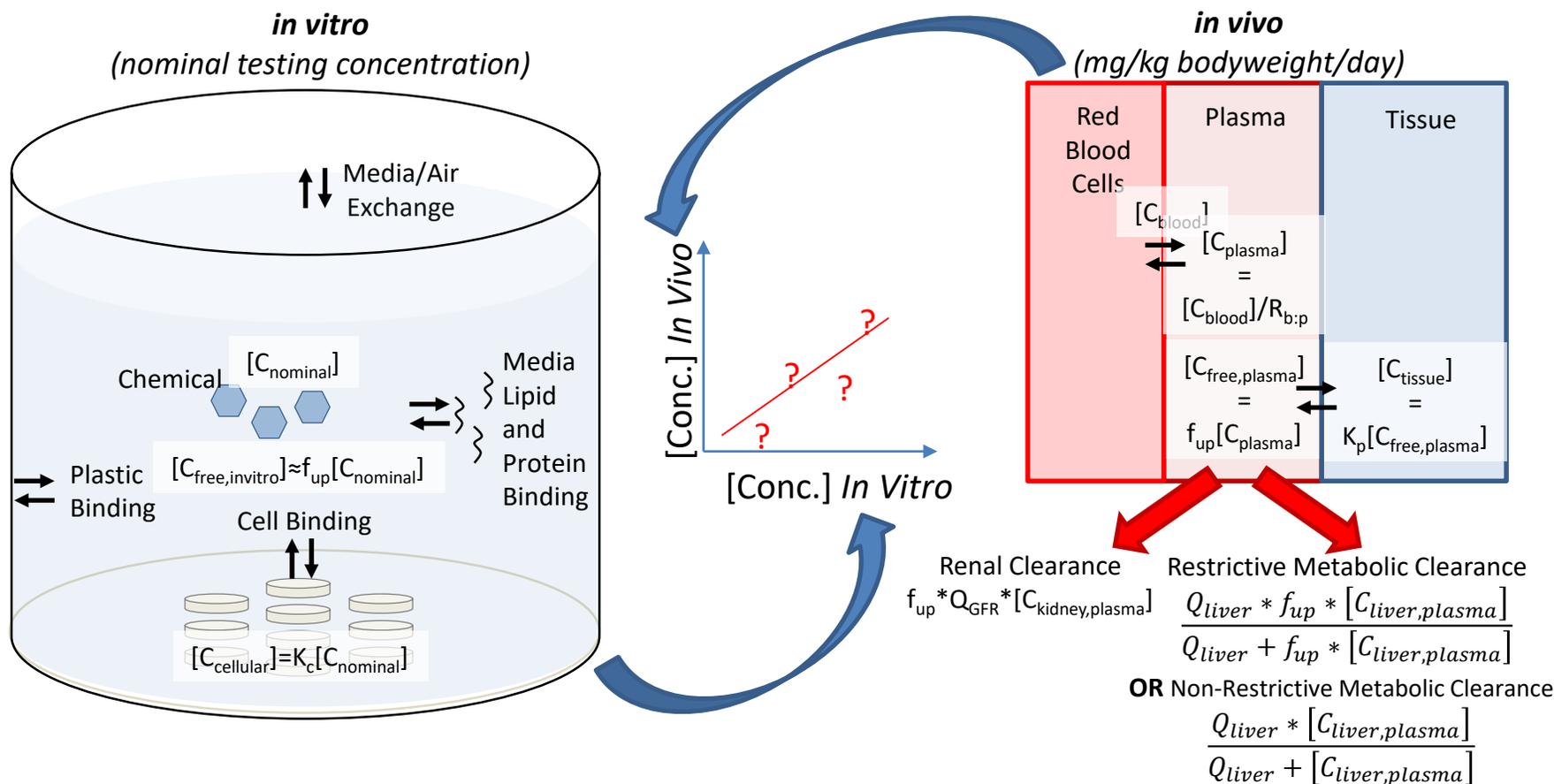
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- Can calculate predicted steady-state concentration ( $C_{ss}$ ) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

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

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



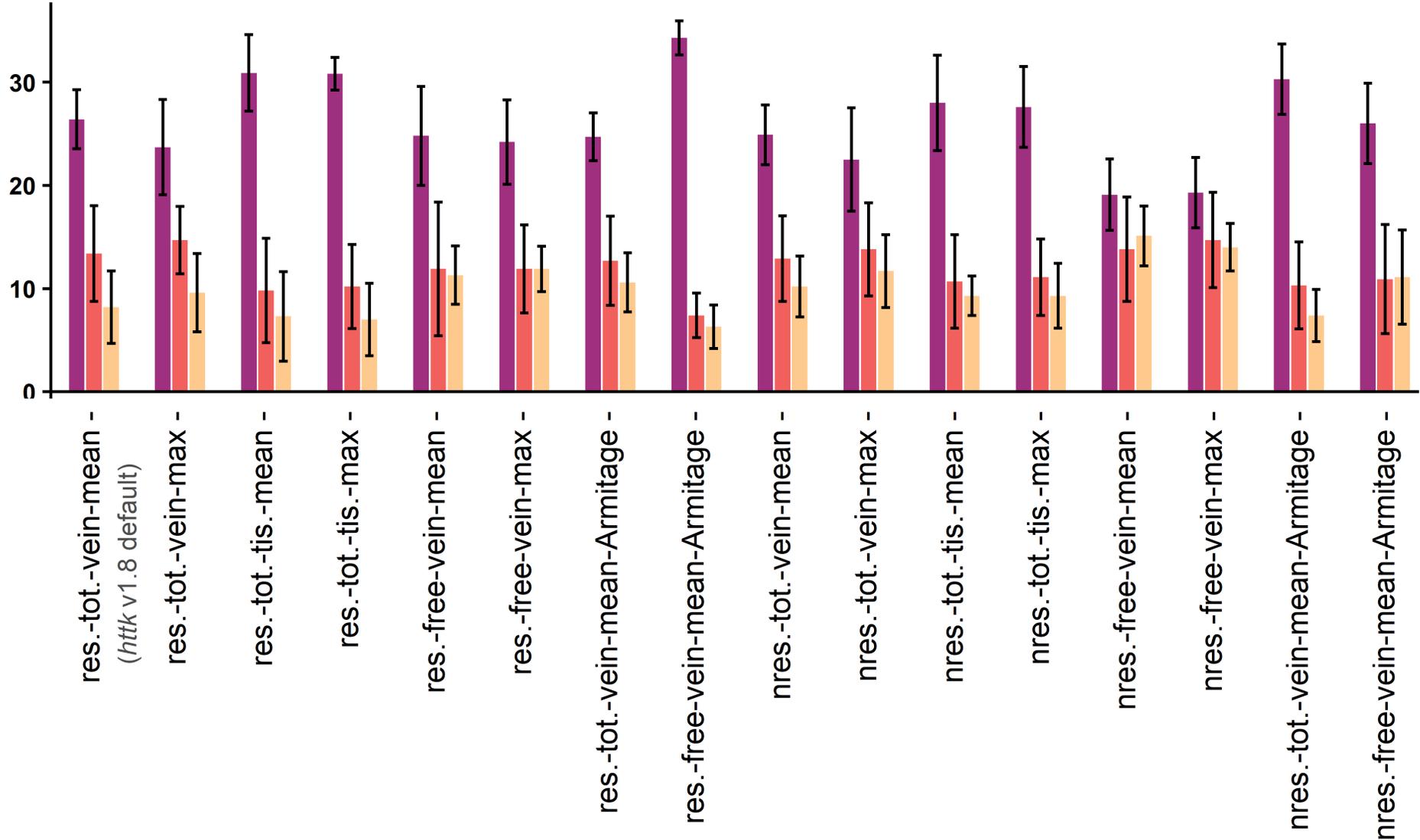
Selecting the appropriate *in vitro* and *in vivo* concentrations for extrapolation

# Optimizing HTTK-based IVIVE

## Reverse Dosimetry

■ PBTk ■ Random ■ AC<sub>50</sub>

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

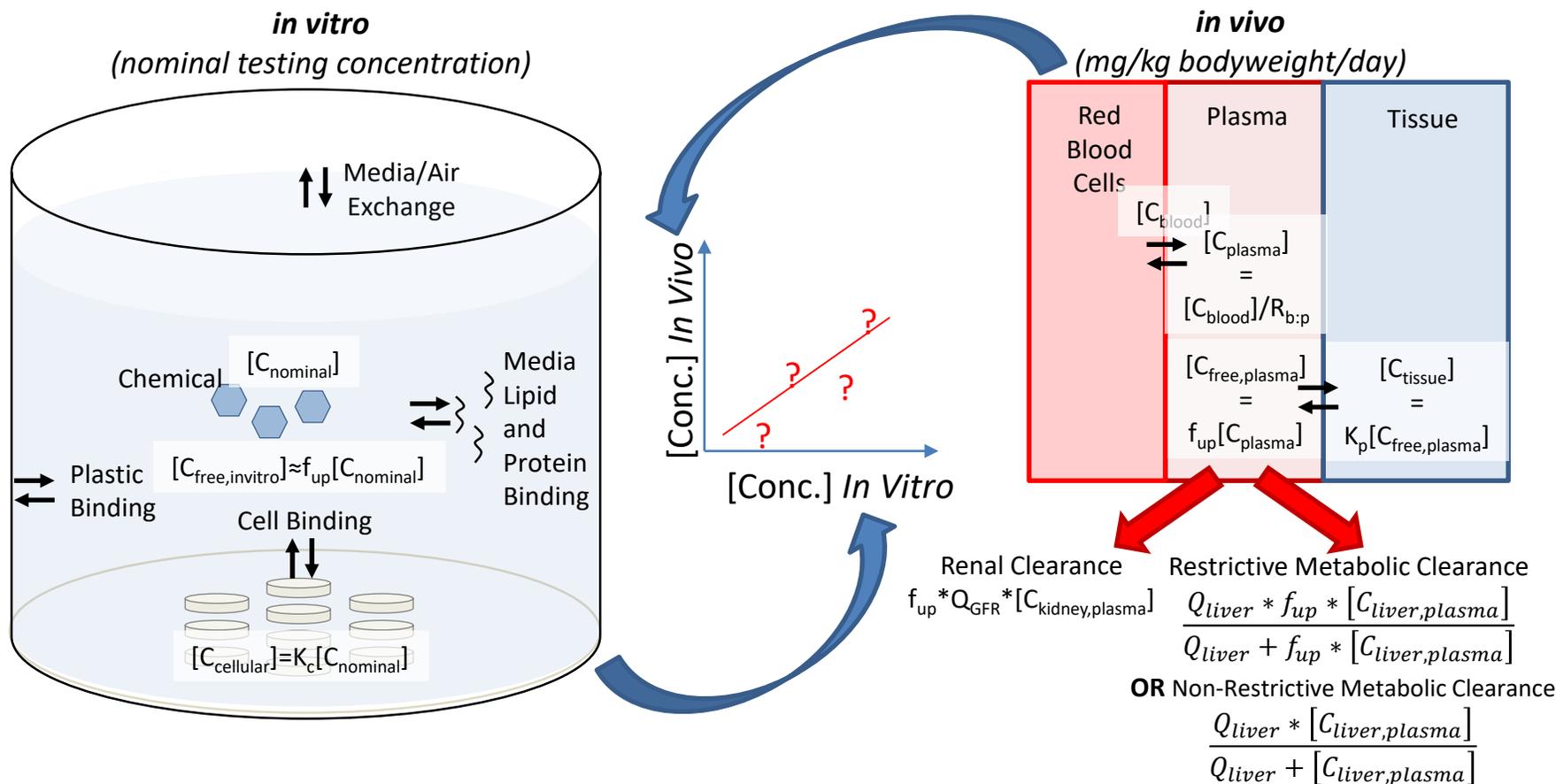


Using PBTk  
Models

Improves IVIVE

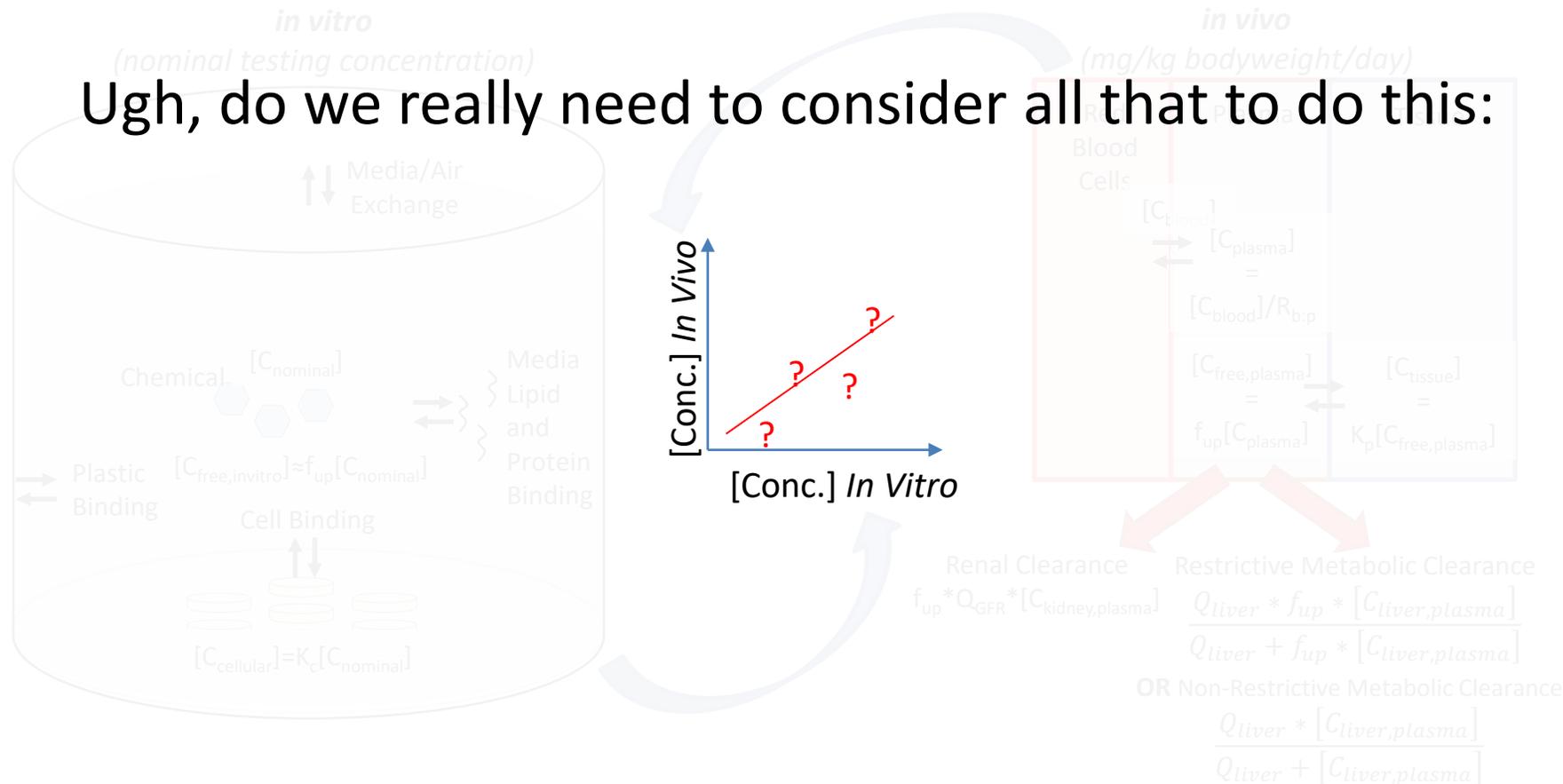
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Using the generic HTTK physiologically based toxicokinetics model to inform IVIVE...



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

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

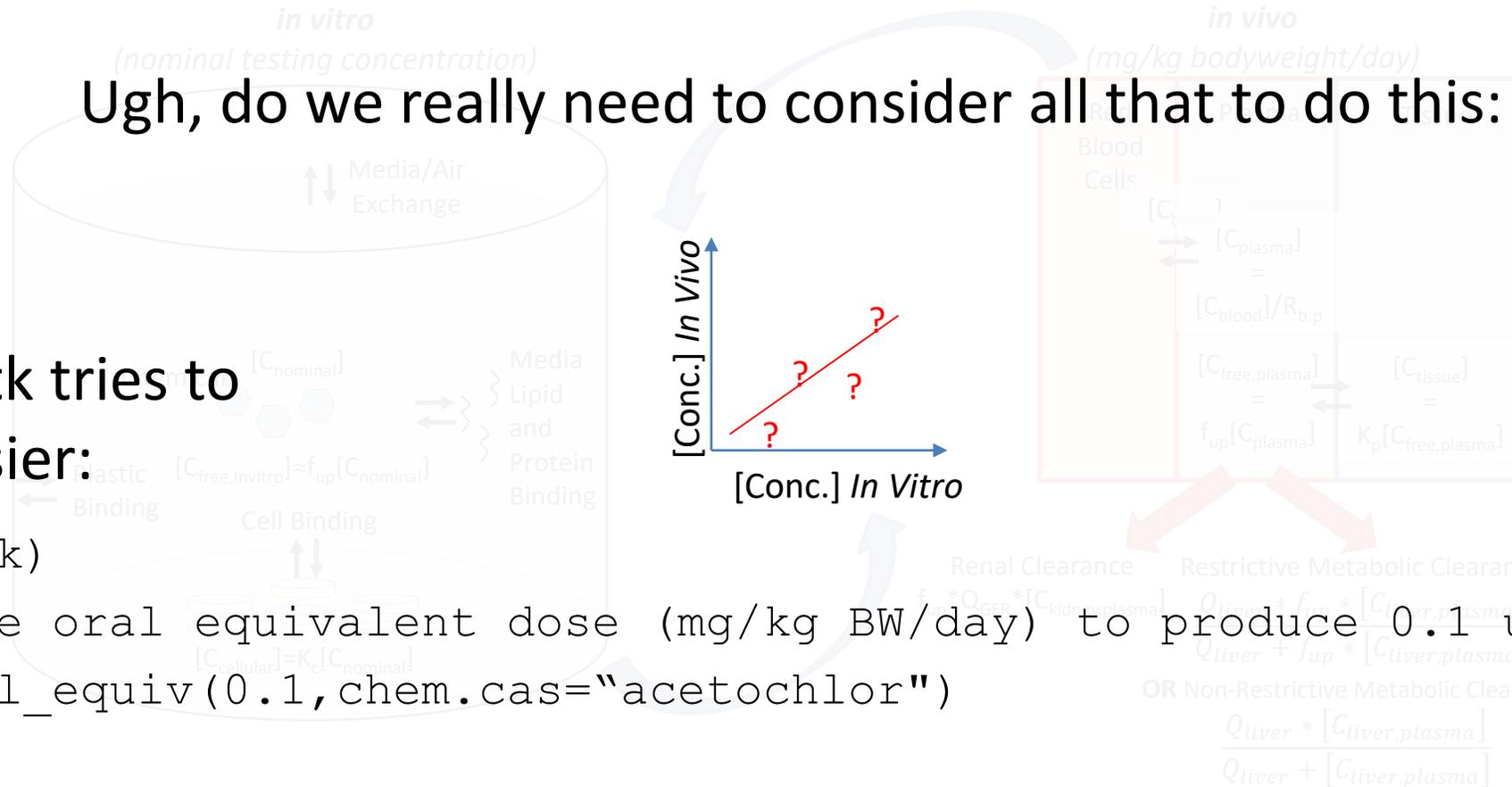
Ugh, do we really need to consider all that to do this:

Yes, but httk tries to  
make it easier:

```
library(httk)
```

```
#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM in plasma:
```

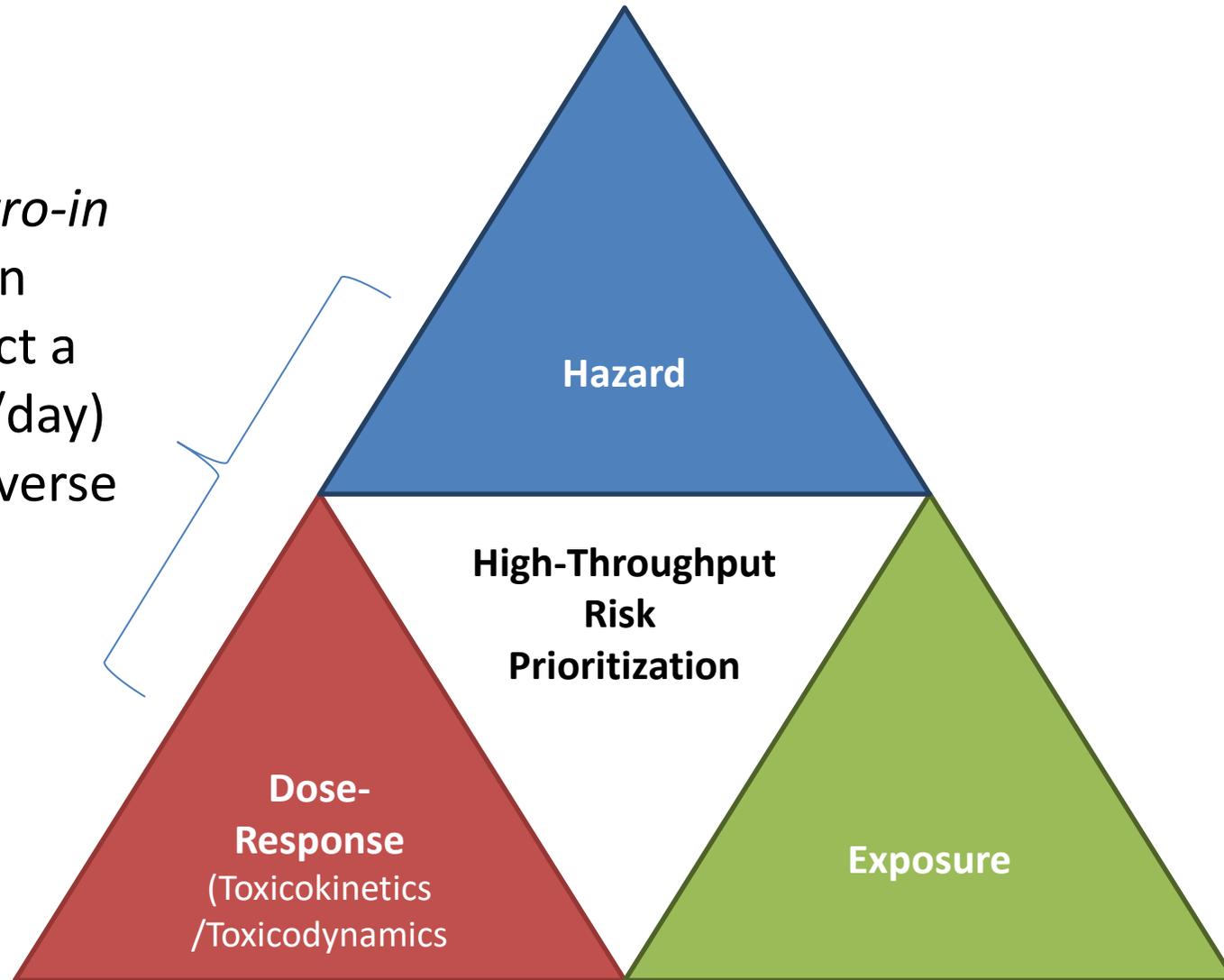
```
calc_mc_oral_equiv(0.1, chem.cas="acetochlor")
```



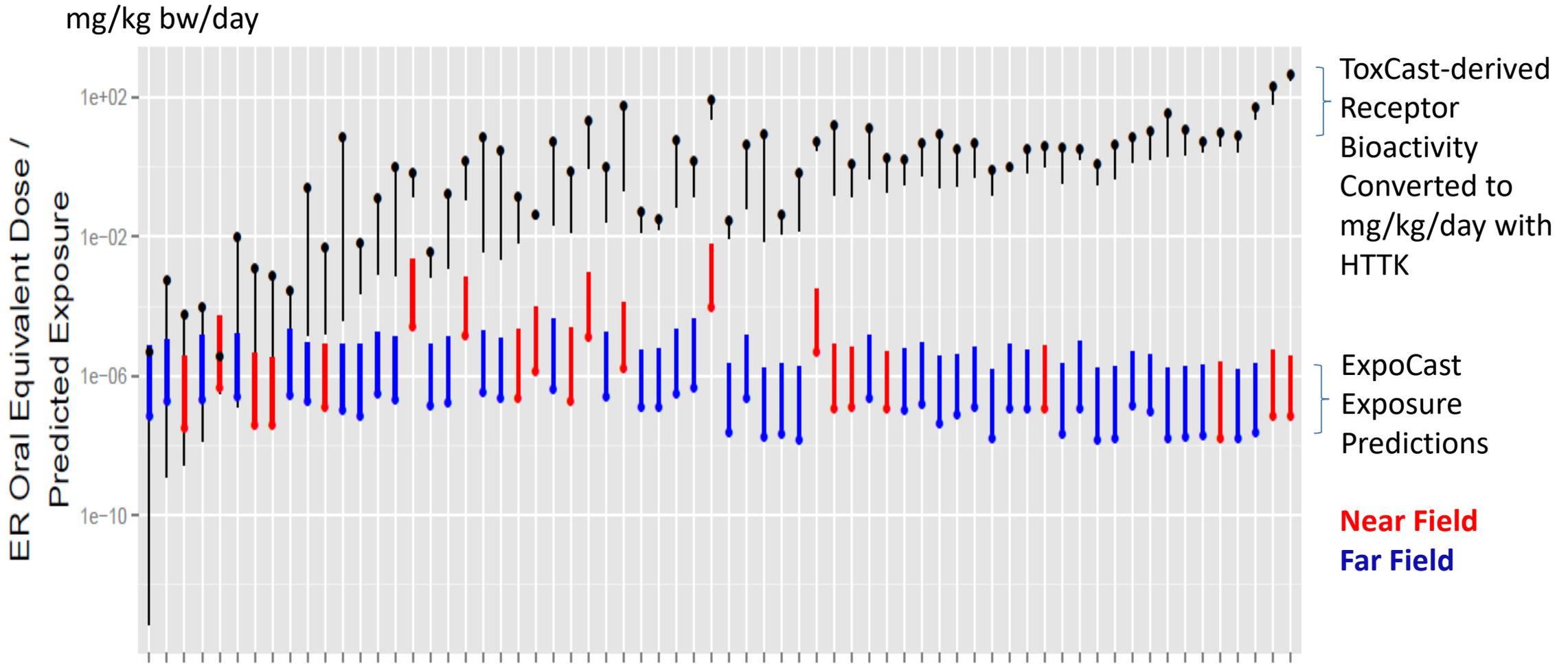
Selecting the appropriate *in vitro* and *in vivo* concentrations for extrapolation

# HTTK Facilitates IVIVE for In Vitro Screening Data

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



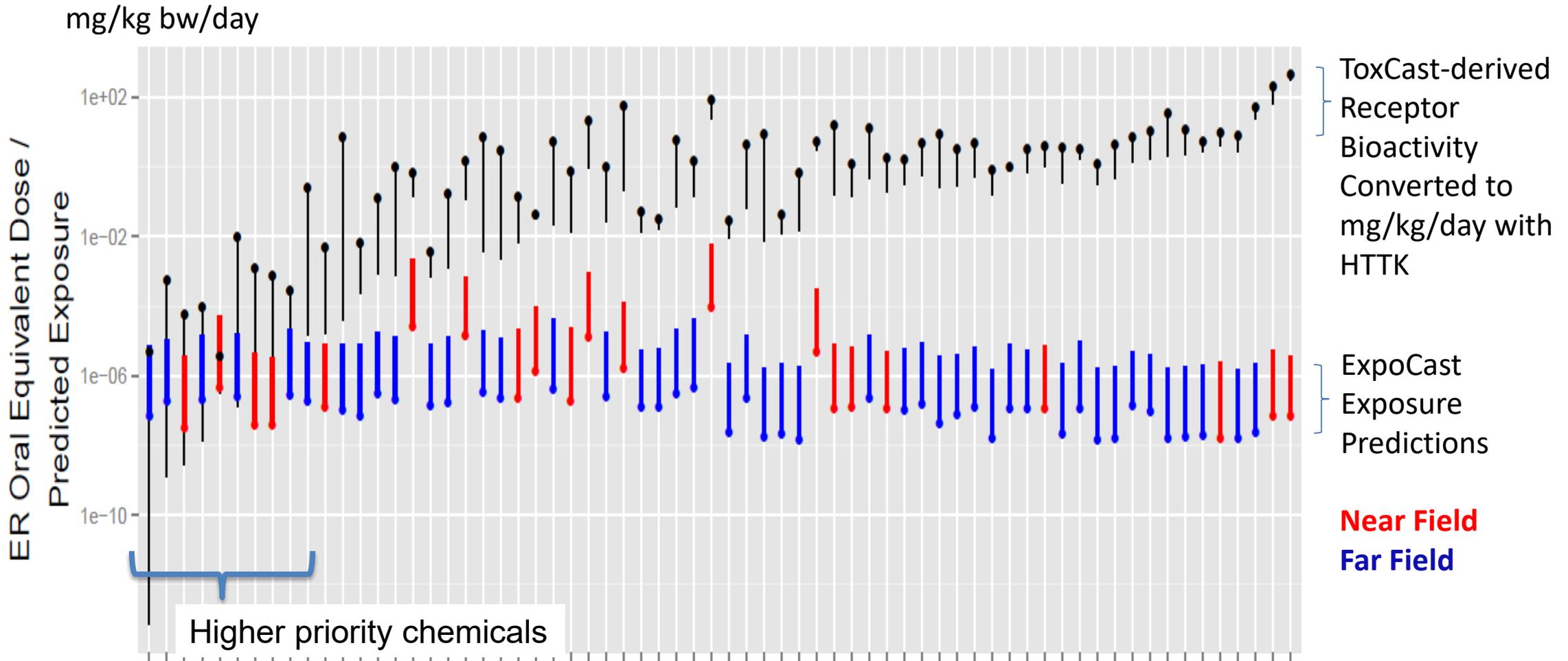
# High Throughput Risk Prioritization in Practice



**ToxCast Chemicals**

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

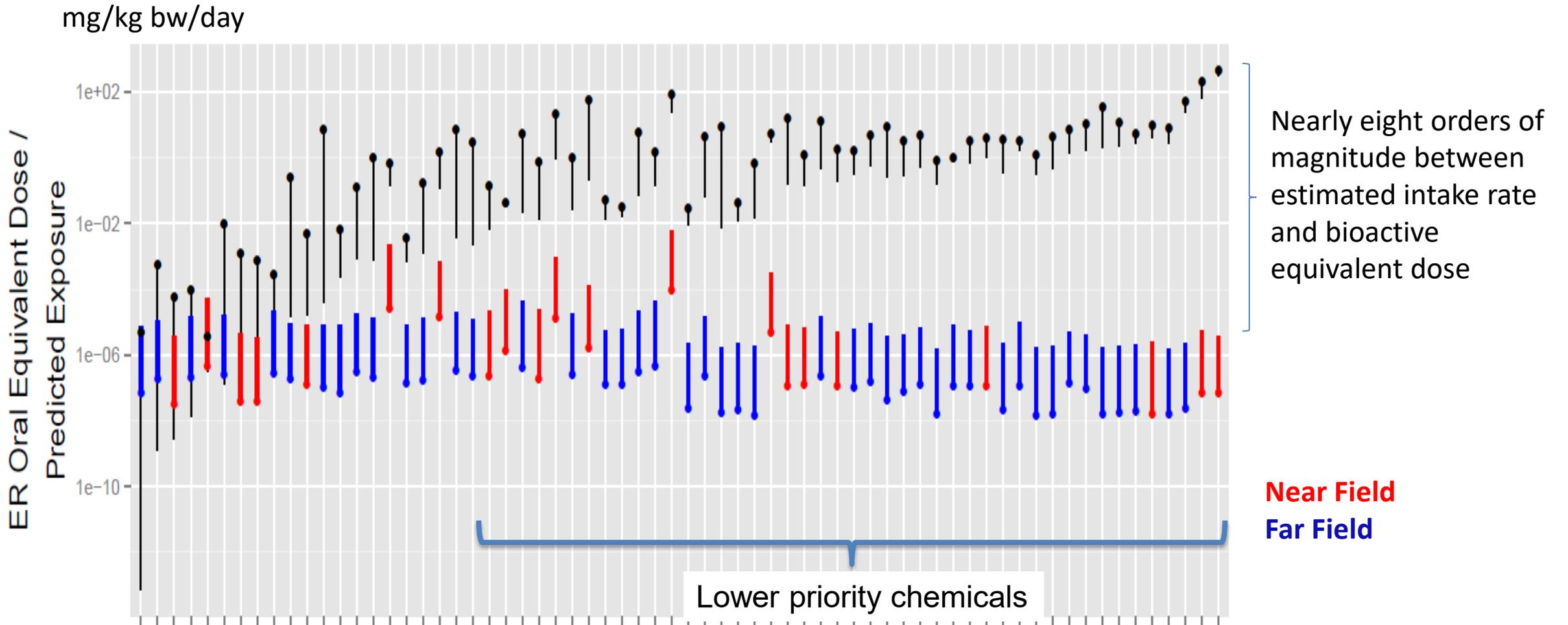
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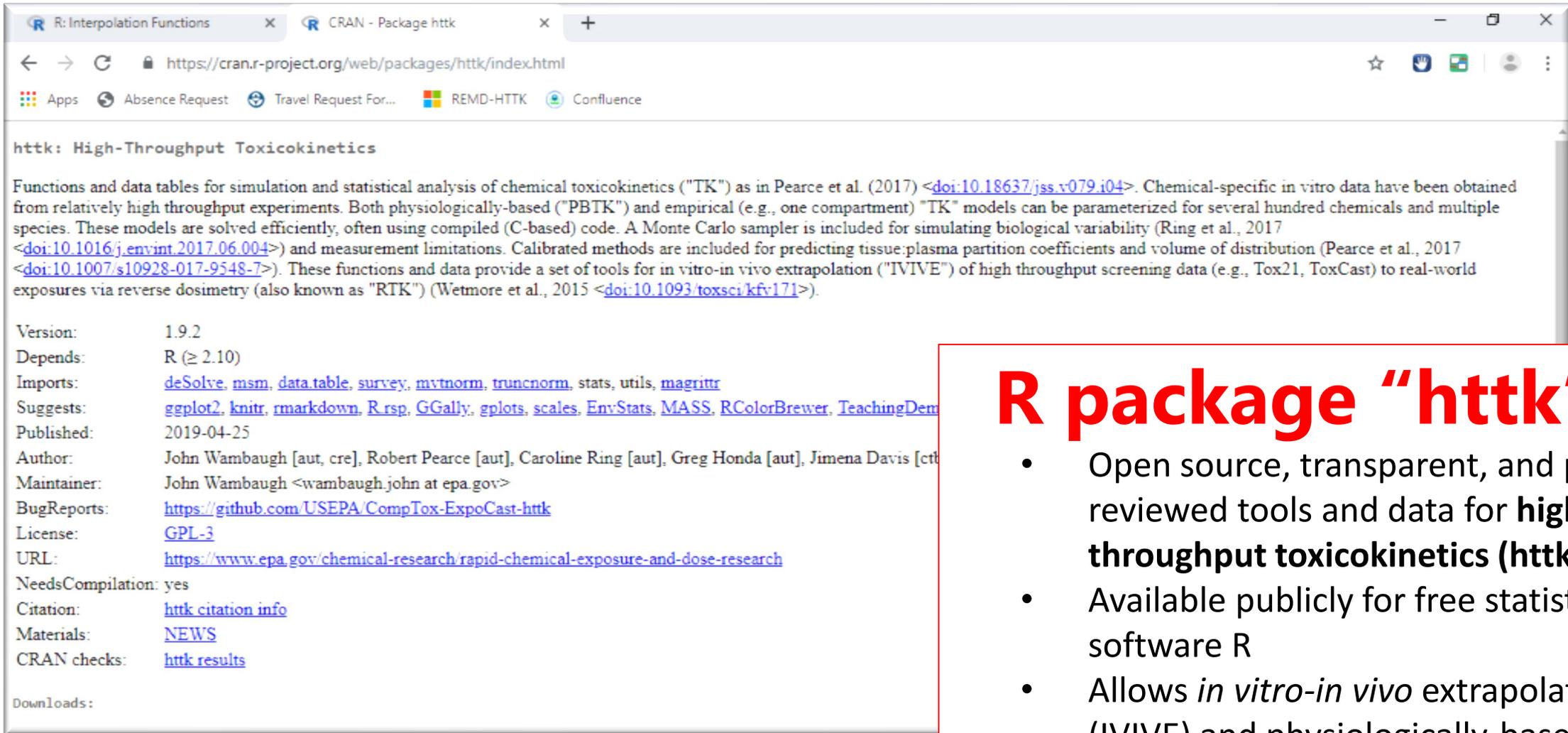
# High Throughput Risk Prioritization in Practice



December, 2014 Panel:  
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# HTTK: Open Source Tools and

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



The screenshot shows a web browser window displaying the CRAN page for the 'httk' R package. The browser tabs are 'R: Interpolation Functions' and 'CRAN - Package httk'. 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 detailed description of its functions and data tables, and a list of metadata such as version (1.9.2), dependencies (R ≥ 2.10), imports, suggests, published date (2019-04-25), author, maintainer, bug reports, license (GPL-3), and URL.

**httk: High-Throughput Toxicokinetics**

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

Version: 1.9.2  
Depends: R (≥ 2.10)  
Imports: [deSolve](#), [msm](#), [data.table](#), [survey](#), [mvtnorm](#), [truncnorm](#), stats, utils, [magrittr](#)  
Suggests: [ggplot2](#), [knitr](#), [rmarkdown](#), [R.rsp](#), [GGally](#), [gplots](#), [scales](#), [EnvStats](#), [MASS](#), [RColorBrewer](#), [TeachingDemos](#)  
Published: 2019-04-25  
Author: John Wambaugh [aut, cre], Robert Pearce [aut], Caroline Ring [aut], Greg Honda [aut], Jimena Davis [ctb]  
Maintainer: John Wambaugh <wambaugh.john at epa.gov>  
BugReports: <https://github.com/USEPA/CompTox-ExpoCast-httk>  
License: [GPL-3](#)  
URL: <https://www.epa.gov/chemical-research/rapid-chemical-exposure-and-dose-research>  
NeedsCompilation: yes  
Citation: [httk citation info](#)  
Materials: [NEWS](#)  
CRAN checks: [httk results](#)

Downloads:

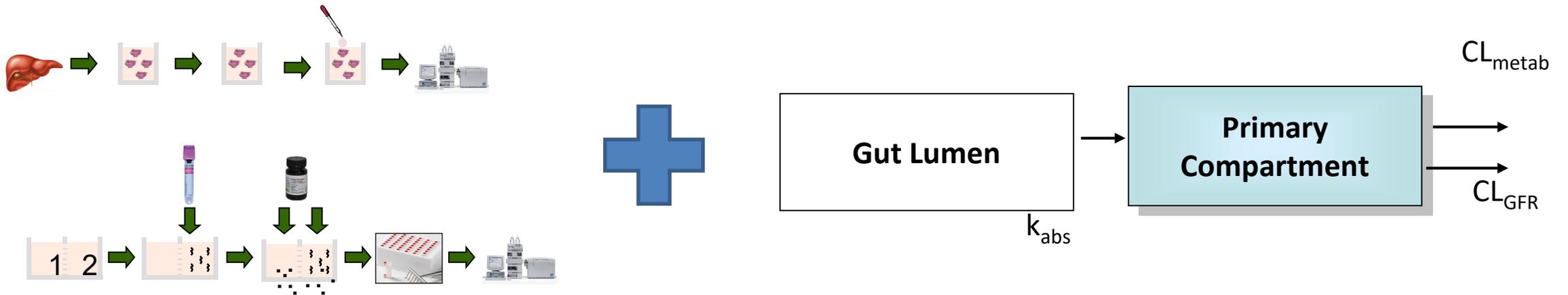
## R package "httk"

- 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”?

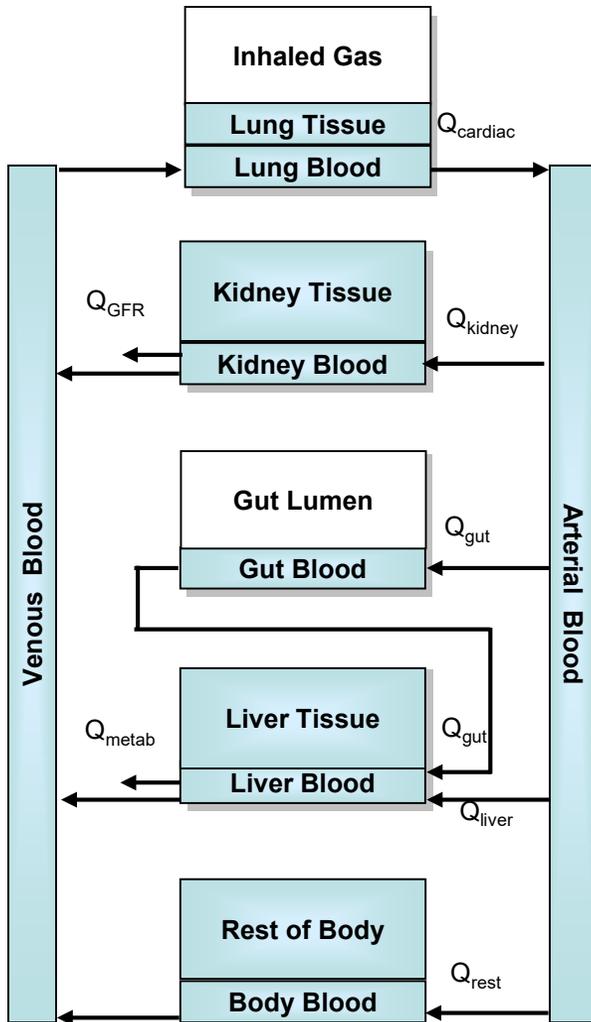
- Allows prediction of internal tissue concentrations from dose regimen (oral and intravenous)
- Allows conversion of *in vitro* concentration to *in vivo* doses
- A peer-reviewed paper in the Journal of Statistical software provides a how-to guide (Pearce et al., 2017a)
- You can use the built in chemical library or add more chemical information (examples provided in JSS paper)
- You can 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()`)

## *In vitro* toxicokinetic data + generic toxicokinetic model = high throughput toxicokinetics



= *httk*

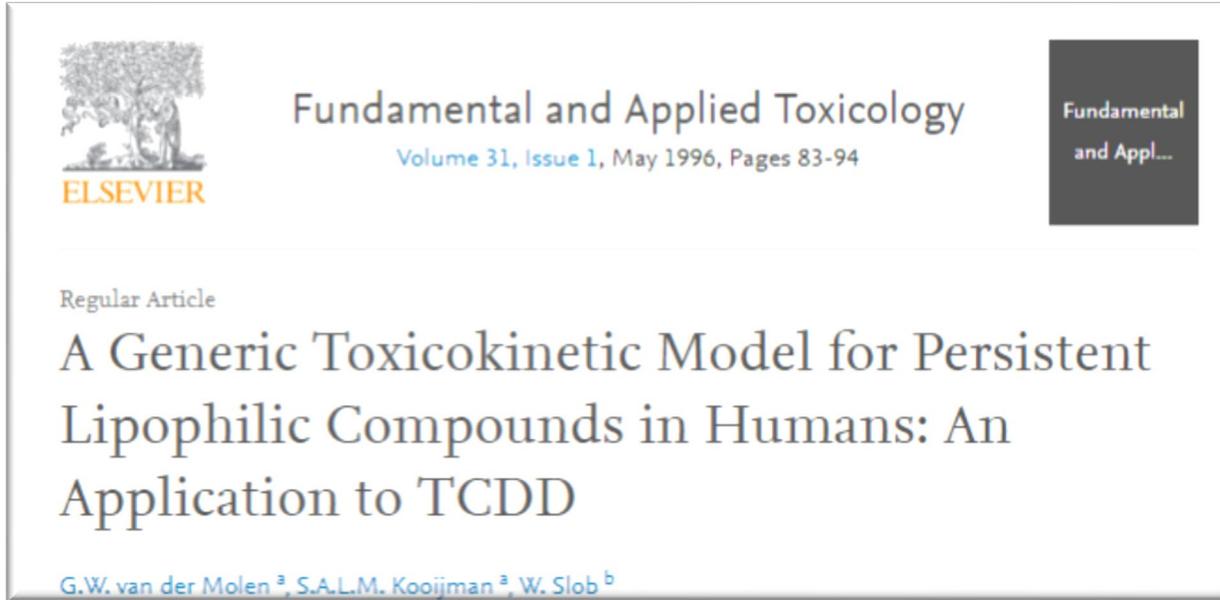
# A General Physiologically-based Toxicokinetic (PBTK) Model



- “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)
- 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).

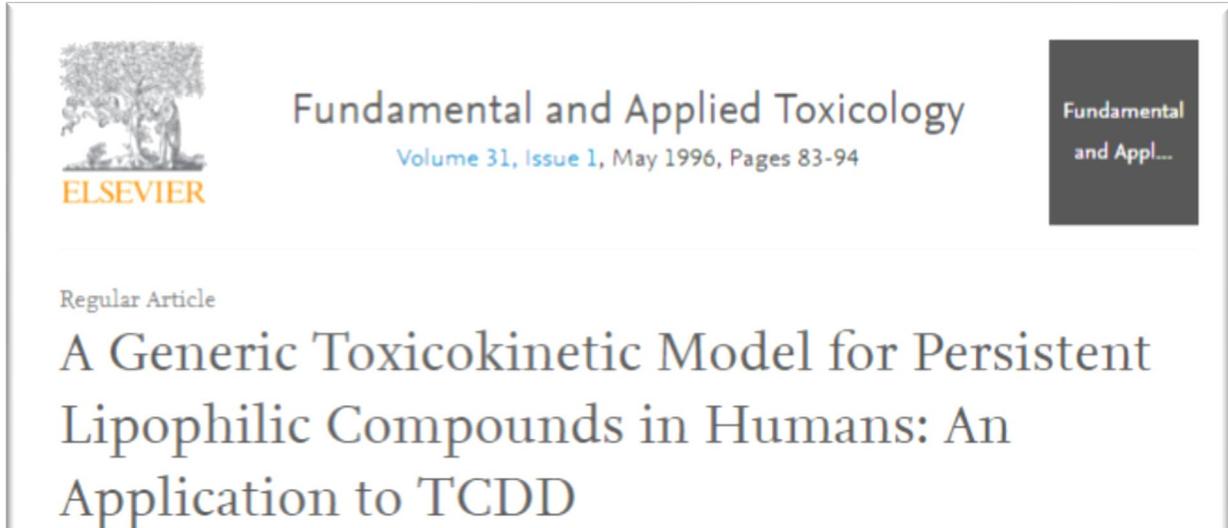
# Generic PBTK Models

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



# Generic PBTK Models

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

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<sup>S</sup>**

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

*Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom*

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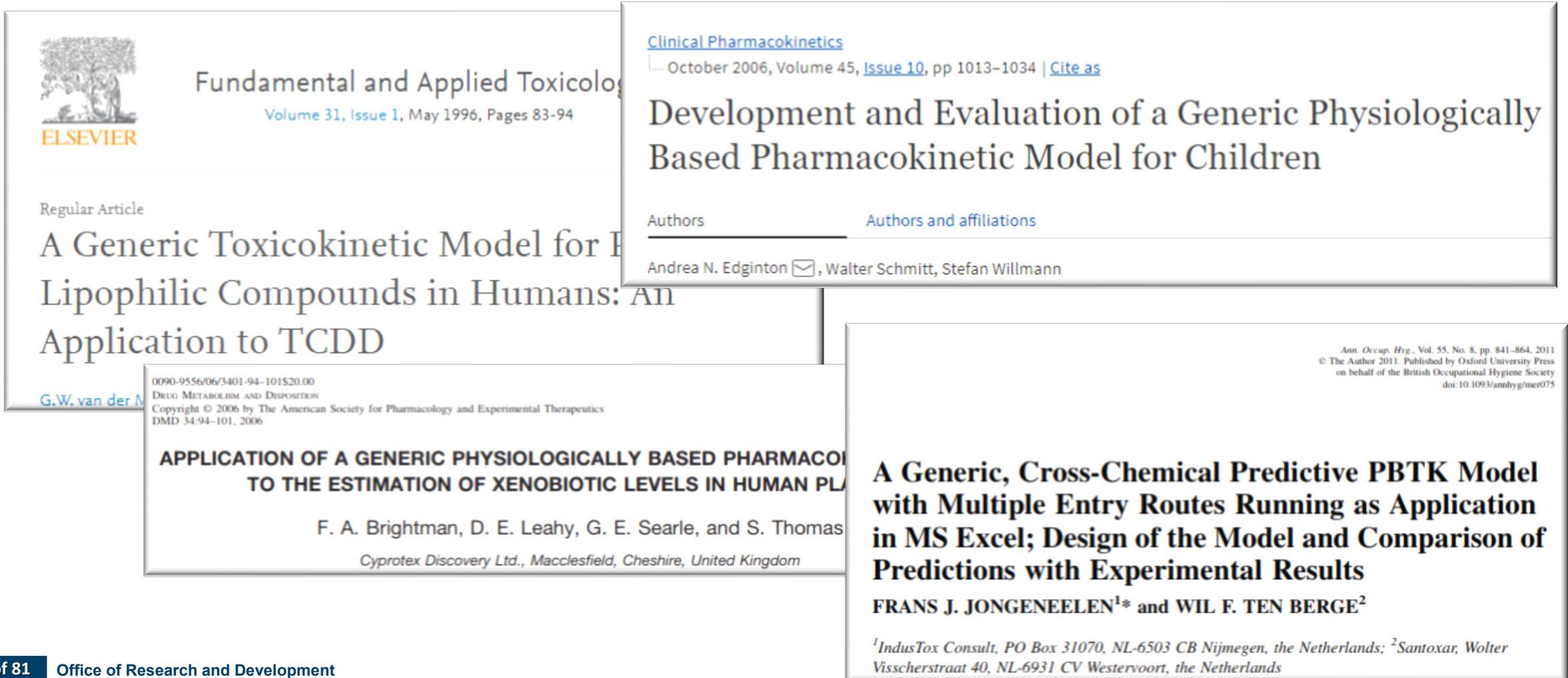
**Fundamental and Applied Toxicology**  
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ELSEVIER  
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A Generic Toxicokinetic Model for Lipophilic Compounds in Humans: An Application to TCDD  
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**Clinical 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: 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  
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Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

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The collage features three overlapping article snippets. The top-left snippet is from *Fundamental and Applied Toxicology*, Elsevier, Volume 31, Issue 1, May 1996, Pages 83-94. It is a regular article titled "A Generic Toxicokinetic Model for Lipophilic Compounds in Humans: An Application to TCDD" by G.W. van der N. The top-right snippet is from *Clinical Pharmacokinetics*, October 2006, Volume 45, Issue 10, pp 1013-1034. It is titled "Development and Evaluation of a Generic Physiologically Based Pharmacokinetic Model for Children" by Andrea N. Edginton, Walter Schmitt, and Stefan Willmann. The bottom snippet is from *Ann. Occup. Hyg.*, Vol. 55, No. 8, pp. 841-864, 2011. It is titled "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" by Frans J. Jongeneelen and Wil F. Ten Berge. The bottom snippet also includes contact information for IndusTox Consult and Santoxar.

**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  
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© The Author 2011. Published by Oxford University Press on behalf of the British Occupational Hygiene Society  
doi:10.1093/amhyg/mer075

**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<sup>1\*</sup> and WIL F. TEN BERGE<sup>2</sup>  
<sup>1</sup>IndusTox Consult, PO Box 31070, NL-6503 CB Nijmegen, the Netherlands; <sup>2</sup>Santoxar, Wolter Visscherstraat 40, NL-6931 CV Westervoort, the Netherlands

# Generic PBTK Models

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

 **Journal of Pharmaceutical Sciences**  
Volume 94, Issue 10, October 2005, Pages 2327-2343

**An Evaluation of the Utility of Physiologically Based Models of Pharmacokinetics in Early Drug Discovery**

Neil Parrott<sup>1</sup> ✉, Nicolas Paquereau<sup>1</sup>, Philippe Coassolo<sup>1</sup>, Thierry Lavé<sup>1</sup>

<sup>1</sup> F. Hoffmann-La Roche AG, Pharmaceuticals Division, CH-4070 Bl, Switzerland

Received 8 April 2005, Revised 25 May 2005, Accepted 29 May 2005, Available online 29 January 2016.

*Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom*

*Clinical Pharmacokinetics*  
2006, Volume 45, Issue 10, pp 1013-1034 | [Cite as](#)

**Development and Evaluation of a Generic Physiologically Based Pharmacokinetic Model for Children**

Authors and affiliations

Anton ✉, Walter Schmitt, Stefan Willmann

*Ann. Occup. Hyg.*, Vol. 55, No. 8, pp. 841-864, 2011  
© The Author 2011. Published by Oxford University Press  
on behalf of the British Occupational Hygiene Society  
doi:10.1093/amhyg/mer075

**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<sup>1\*</sup> and WIL F. TEN BERGE<sup>2</sup>

<sup>1</sup>IndusTox Consult, PO Box 31070, NL-6503 CB Nijmegen, the Netherlands; <sup>2</sup>Santoxar, Wolter Visscherstraat 40, NL-6931 CV Westervoort, the Netherlands

# Generic PBTK Models

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



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## An Evaluation of the Utility of Physiologically Based Models of Pharmacokinetics in Early Drug Discovery

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Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

**Expert Opinion**

## The Simcyp<sup>®</sup> Population-based ADME Simulator

Masoud Jamei<sup>†</sup>, Steve Marciniak, Kairui Feng, Adrian Barnett, Geoffrey Tucker & Amin Rostami-Hodjegan

<sup>†</sup>Modelling & Simulation Group, Simcyp Limited, Blades Enterprise Centre, John Street, Sheffield, S2 4SU, UK

The Simcyp<sup>®</sup> population-based absorption, distribution, metabolism and excretion simulator is a platform and database for 'bottom-up' mechanistic modelling and simulation of the processes of oral absorption, tissue distribution, metabolism and excretion of drugs and drug candidates in healthy and disease populations. It combines experimental data generated routinely during preclinical drug discovery and development from *in vitro* enzyme and cellular systems and relevant physicochemical attributes of compound and dosage form with demographic, physiological and genetic information on different patient populations. The mechanistic approach implemented in the Simcyp Simulator allows simulation of complex absorption, distribution, metabolism and excretion outcomes, particularly those involving multiple drug interactions, parent drug and metabolite profiles and time- and dose-dependent phenomena such as auto-induction and auto-inhibition.

## Predictions with Experimental Results

FRANS J. JONGENELEN<sup>1\*</sup> and WIL F. TEN BERGE<sup>2</sup>

<sup>1</sup>IndusTox Consult, PO Box 31070, NL-6503 CB Nijmegen, the Netherlands; <sup>2</sup>Santoxar, Wolter Visscherstraat 40, NL-6931 CV Westervoort, the Netherlands

Windows taskbar: File Explorer, Outlook, Excel, PowerPoint, Edge, Chrome, Word, Internet Explorer, PDF Reader

# 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: <a href="http://xnet.hsl.gov.uk/megen">http://xnet.hsl.gov.uk/megen</a>	Free: <a href="http://cefic-lri.org/lri_toolbox/induschemfate/">http://cefic-lri.org/lri_toolbox/induschemfate/</a>	Free: <a href="https://CRAN.R-project.org/package=httk">https://CRAN.R-project.org/package=httk</a>
Open Source	No	No	<b>Yes</b>	No	<b>Yes</b>
Default PBPK Structure	<b>Yes</b>	<b>Yes</b>	No	<b>Yes</b>	<b>Yes</b>
Expandable PBPK Structure	No	No	<b>Yes</b>	No	No
Population Variability	<b>Yes</b>	No	No	No	<b>Yes</b>
Batch Mode	<b>Yes</b>	<b>Yes</b>	No	No	<b>Yes</b>
Graphical User Interface	<b>Yes</b>	<b>Yes</b>	<b>Yes</b>	Excel	No
Physiological Data	<b>Yes</b>	<b>Yes</b>	<b>Yes</b>	<b>Yes</b>	<b>Yes</b>
Chemical-Specific Data Library	Many Clinical Drugs	No	No	15 Environmental Compounds	933 Pharmaceutical and ToxCast Compounds
Ionizable Compounds	<b>Yes</b>	<b>Yes</b>	Potentially	No	<b>Yes</b>
Export Function	No	No	Matlab and AcslX	No	SBML and Jarnac
R Integration	No	No	No	No	<b>Yes</b>
Easy Reverse Dosimetry	<b>Yes</b>	<b>Yes</b>	No	No	<b>Yes</b>
Future Proof XML	No	No	<b>Yes</b>	No	No

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,<sup>\*-1</sup> Hisham A. El-Masri,<sup>†</sup> Lisa M. Sweeney,<sup>‡</sup> Leonid Y. Kopylev,<sup>||</sup> Harvey J. Clewell,<sup>§</sup> John F. Wambaugh,<sup>¶</sup>  
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“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]”

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### 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.”

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

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# EXAMPLE: Oral Equivalent Doses

```
#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_lit_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_lit_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")
```

# EXAMPLE: Interspecies Extrapolation

```
#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_lit_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_lit_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_lit_css(chem.cas="34256-82-1",species="Mouse")
```

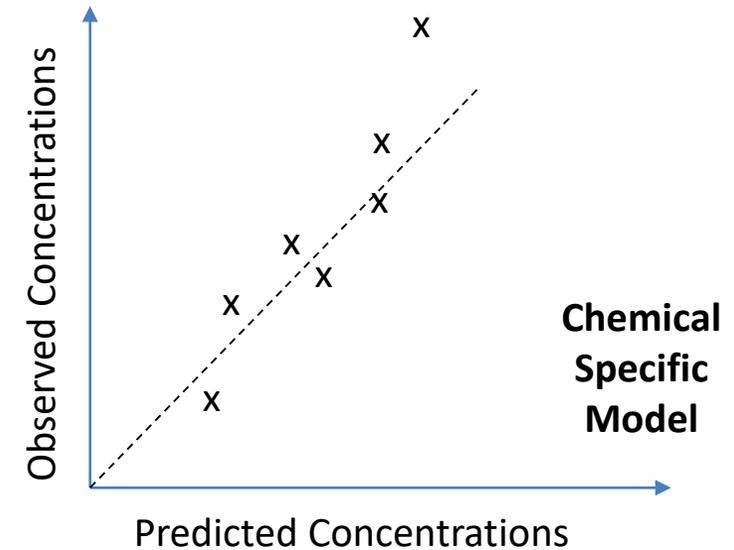
```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for mouse for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1",species="Mouse")  
calc_mc_css(chem.cas="34256-82-1",species="Mouse",default.to.human=T)
```

# 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**
  - Any approximations, omissions, or mistakes should work to increase the estimated uncertainty when evaluated systematically across chemicals

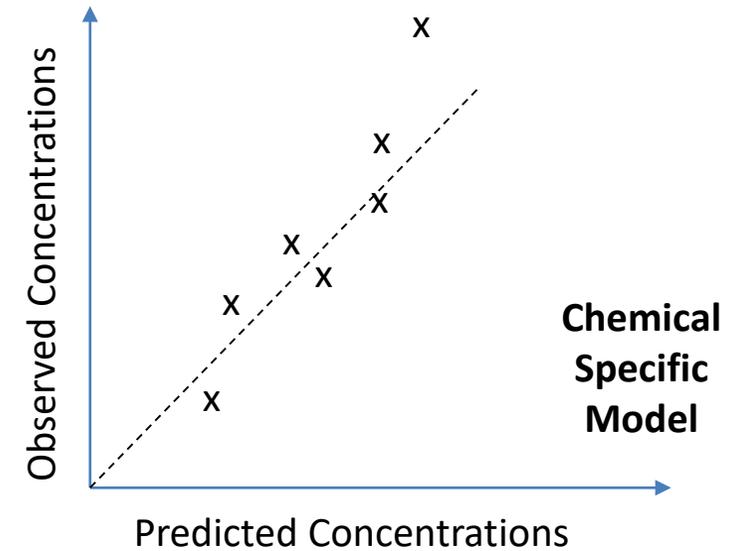
# Building Confidence in TK Models

- 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



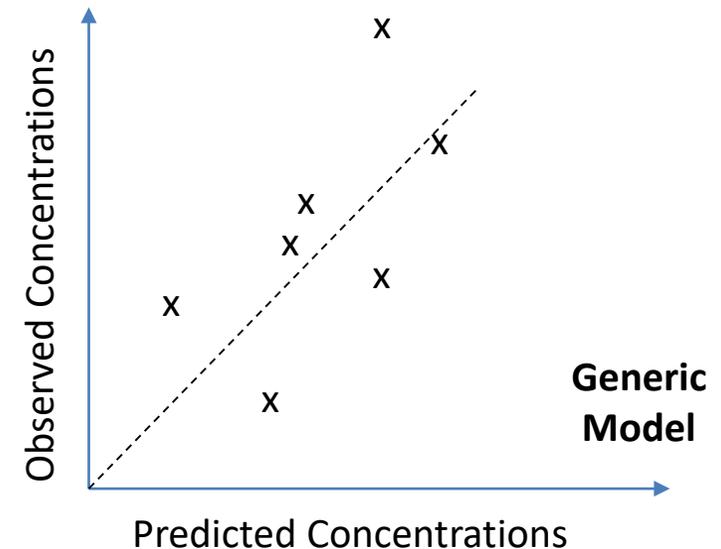
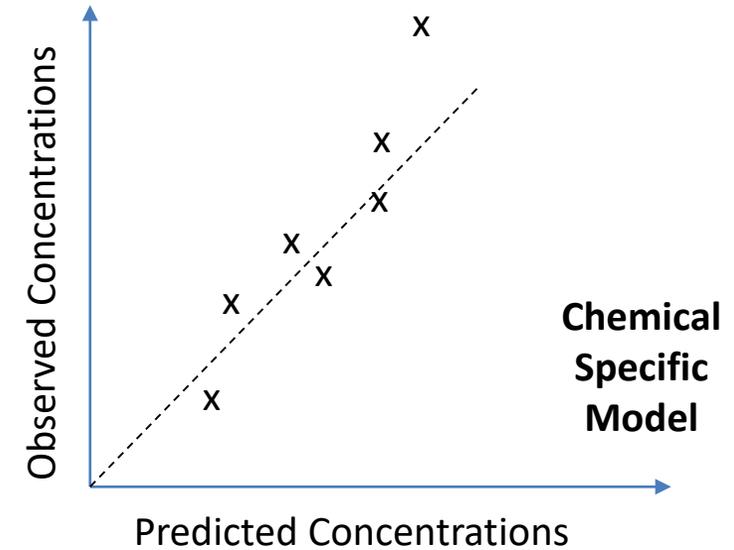
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- However, we do not typically have TK data



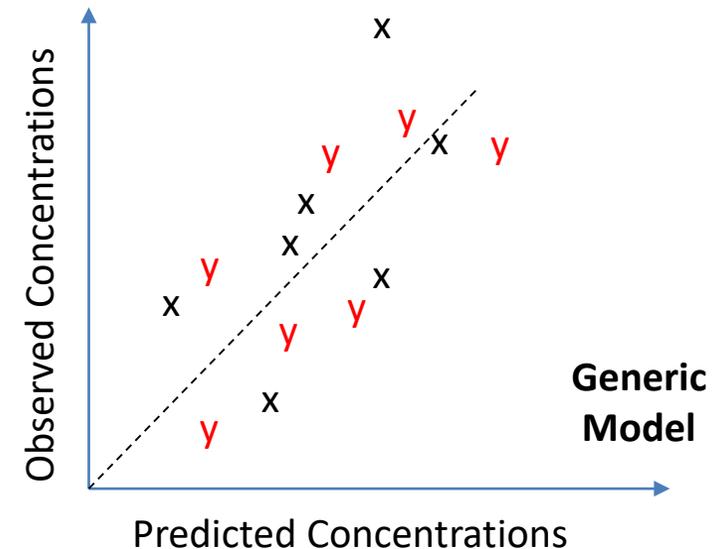
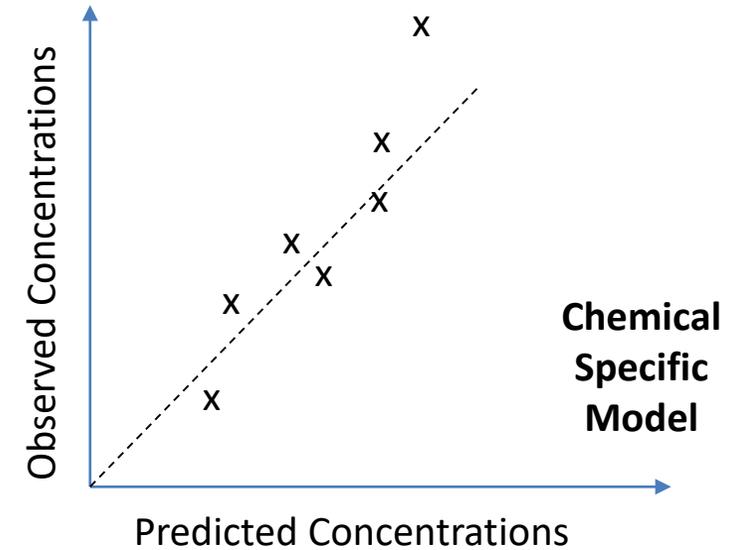
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- We can parameterize a **generic TK model**, and evaluate that model for as many chemicals as we do have data
  - We do expect larger uncertainty, but also greater confidence in model implementation
  - Estimate bias and uncertainty, and try to correlate with chemical-specific properties



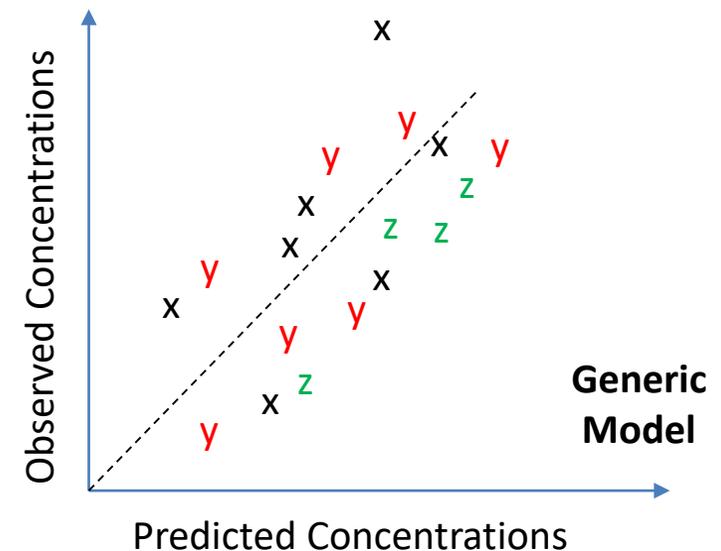
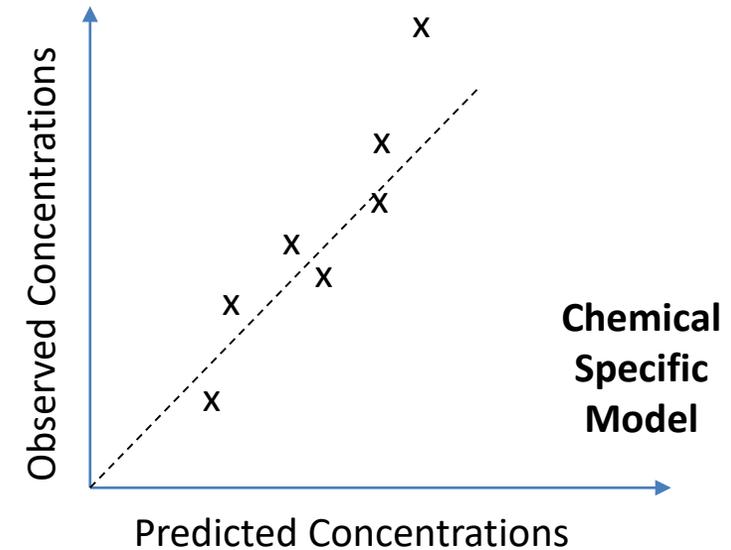
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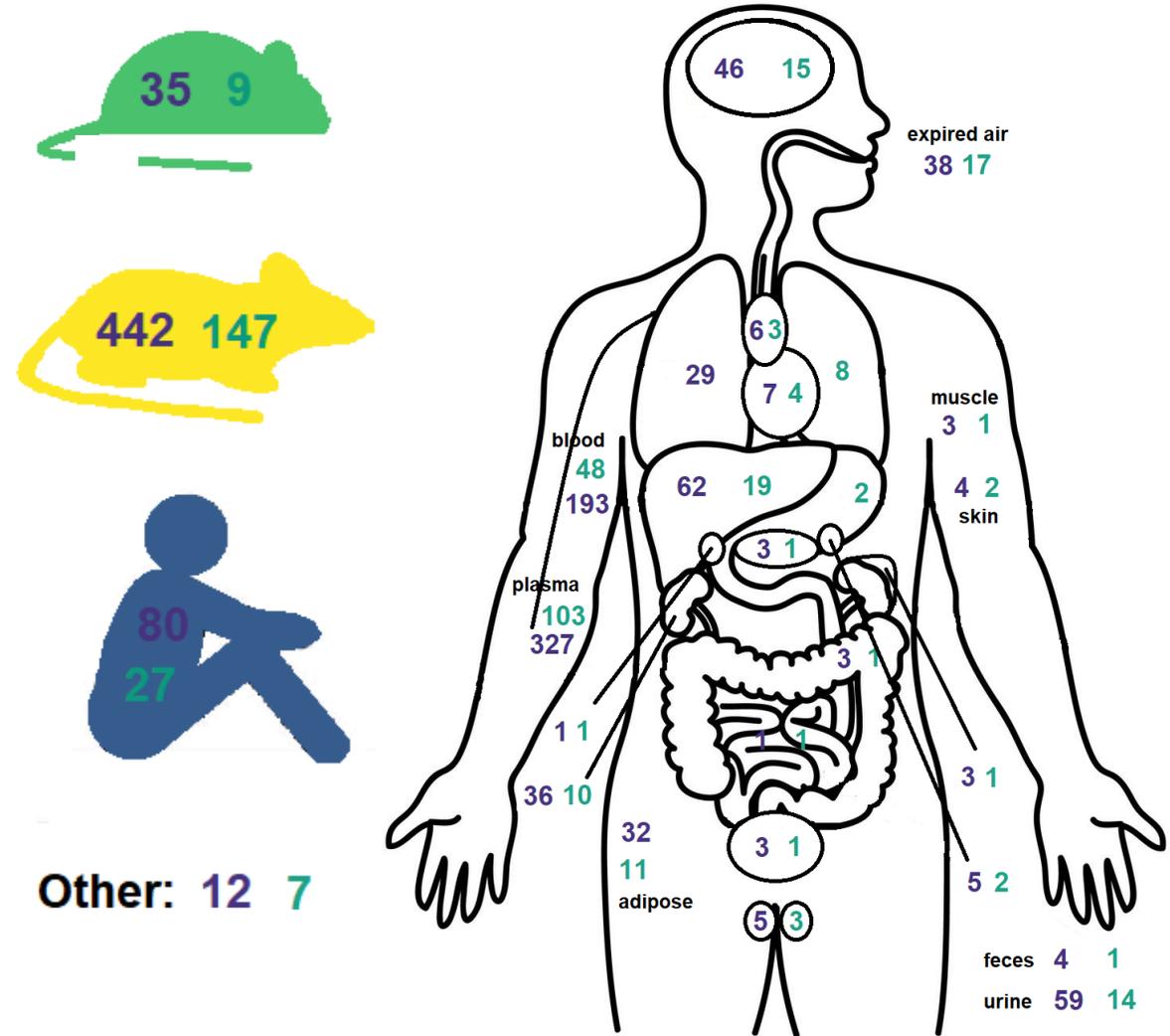
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# In Vivo TK Database

- 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
  - Data from cross lab study (EPA: Michael Hughes, Jane Ellen Simmons, Denise MacMillan, Jermaine Ford, RTI: Timothy Fennell, Rodney Snyder, Sherry Black)
- Standardized, open source curve fitting software *invivoPKfit* used to calibrate models to all data:

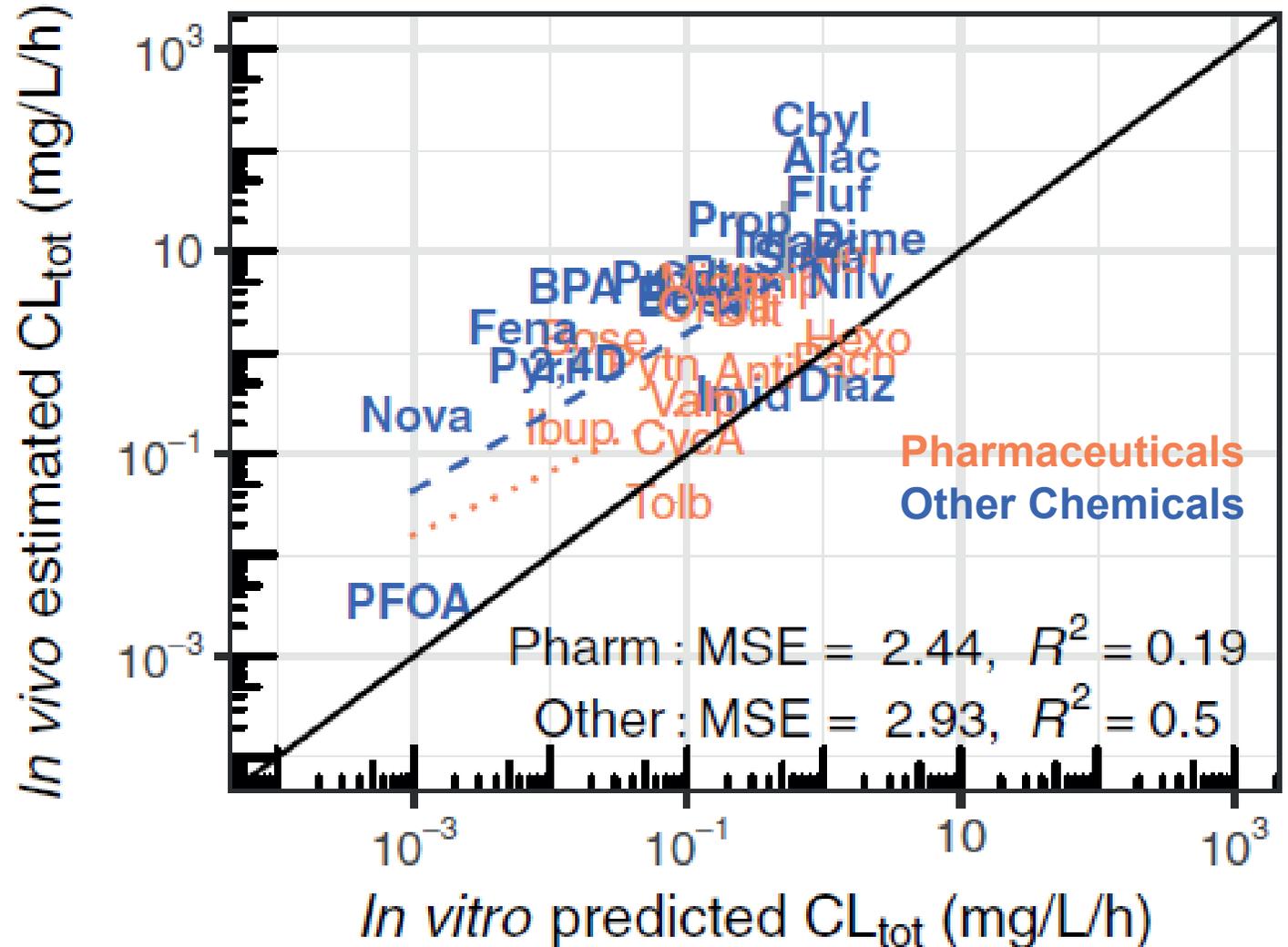


<https://github.com/USEPA/CompTox-ExpoCast-invivoPKfit>

Sayre et al. (in preparation)

# Predicted vs. 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



# Population simulator for HHTK



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

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

*Sample* NHANES biometrics for actual individuals:

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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.)

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(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.)

*Predict* physiological quantities

- Tissue masses
- Tissue blood flows
- GFR (kidney function)
- Hepatocellularity

# Uncertainty

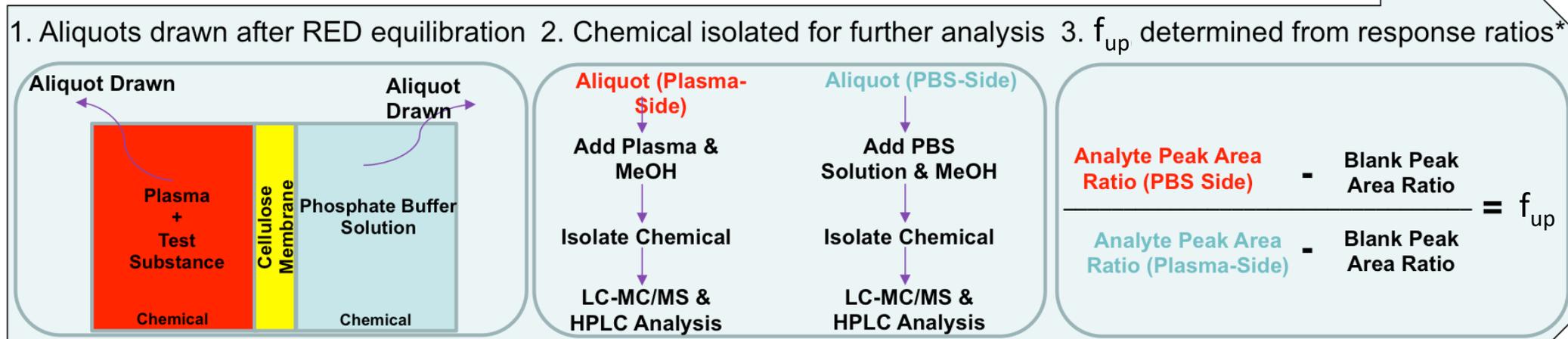
Until I open the  
box, I don't know  
what colors I  
have...

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



# Analytical Chemistry is an HTTK Bottleneck

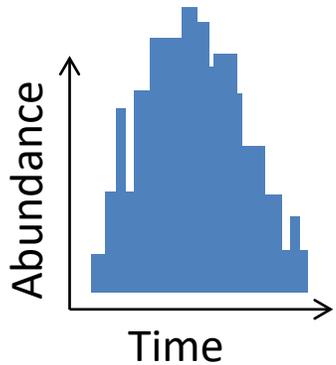
- For HTTK we always need to develop a chemical-specific method for quantitating amount of chemical *in vitro*
  - This is very different from HTS where the same readout (e.g., bioluminescence) can be used for most chemicals
- In Wetmore et al. (2012), the rapid equilibrium dialysis (RED) assay (Waters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



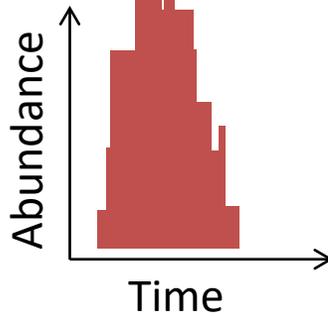
# New HTTK Measurements and Uncertainty Analysis

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

Internal Standard



Chemical Peak

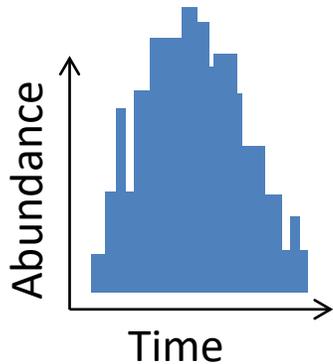


- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio  $R$  of the chemical peak to the ITSD

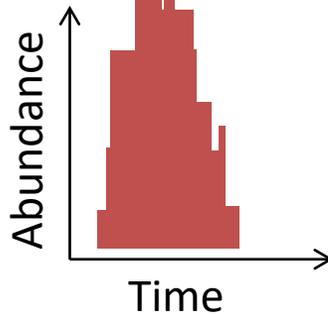
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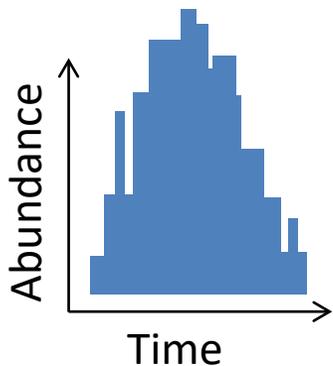


Chemical Peak

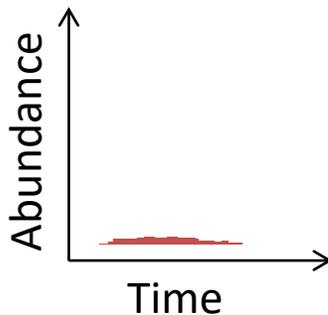


- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio  $R$  of the chemical peak to the ITSD
- For new measurements HTTK (>200 compounds to data) performed by Cyprotex, we have modified RED protocol to use a titration of plasma protein (10%, 30%, 100%) of physiological concentration

Internal Standard



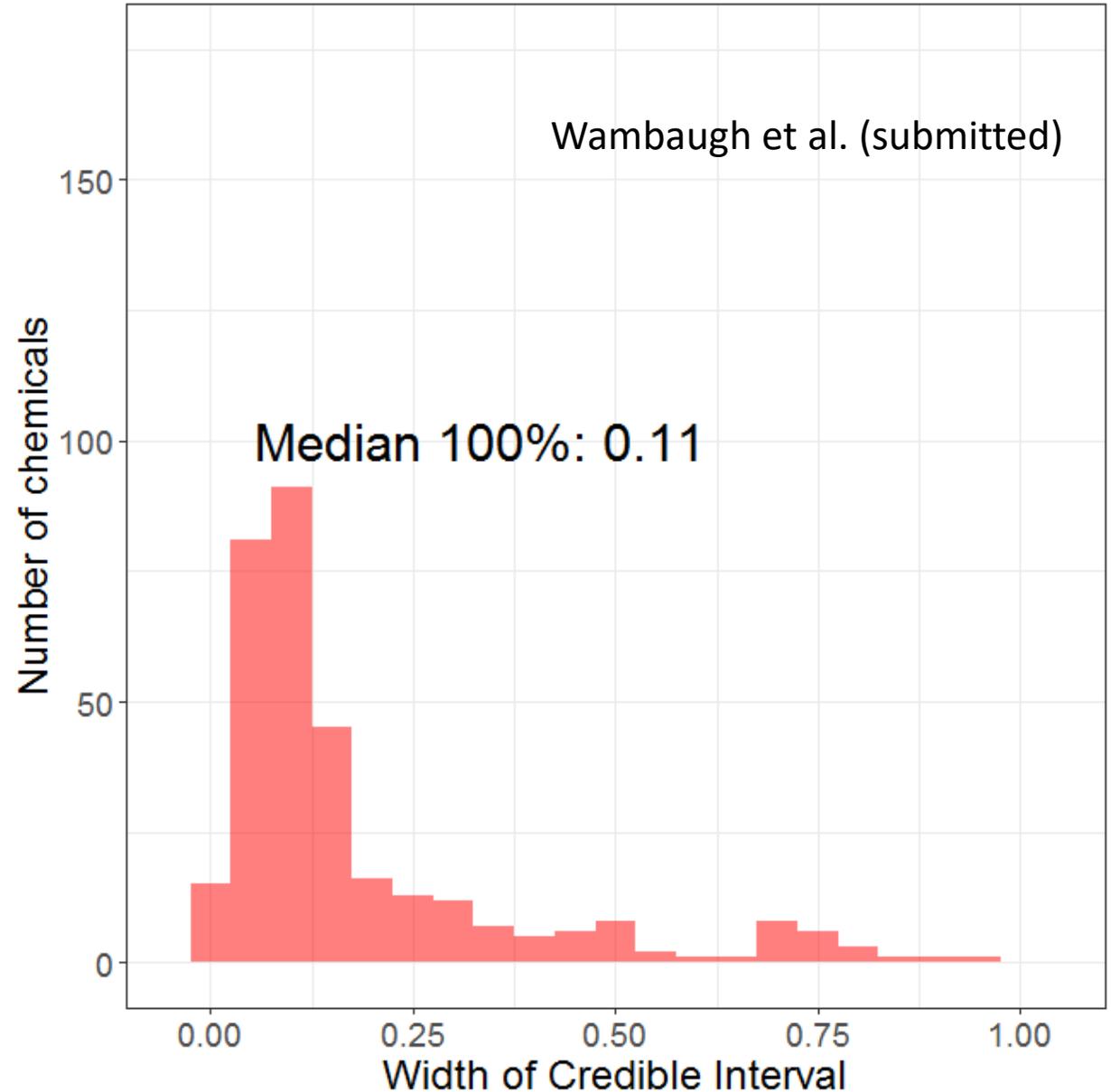
Chemical Peak



- 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

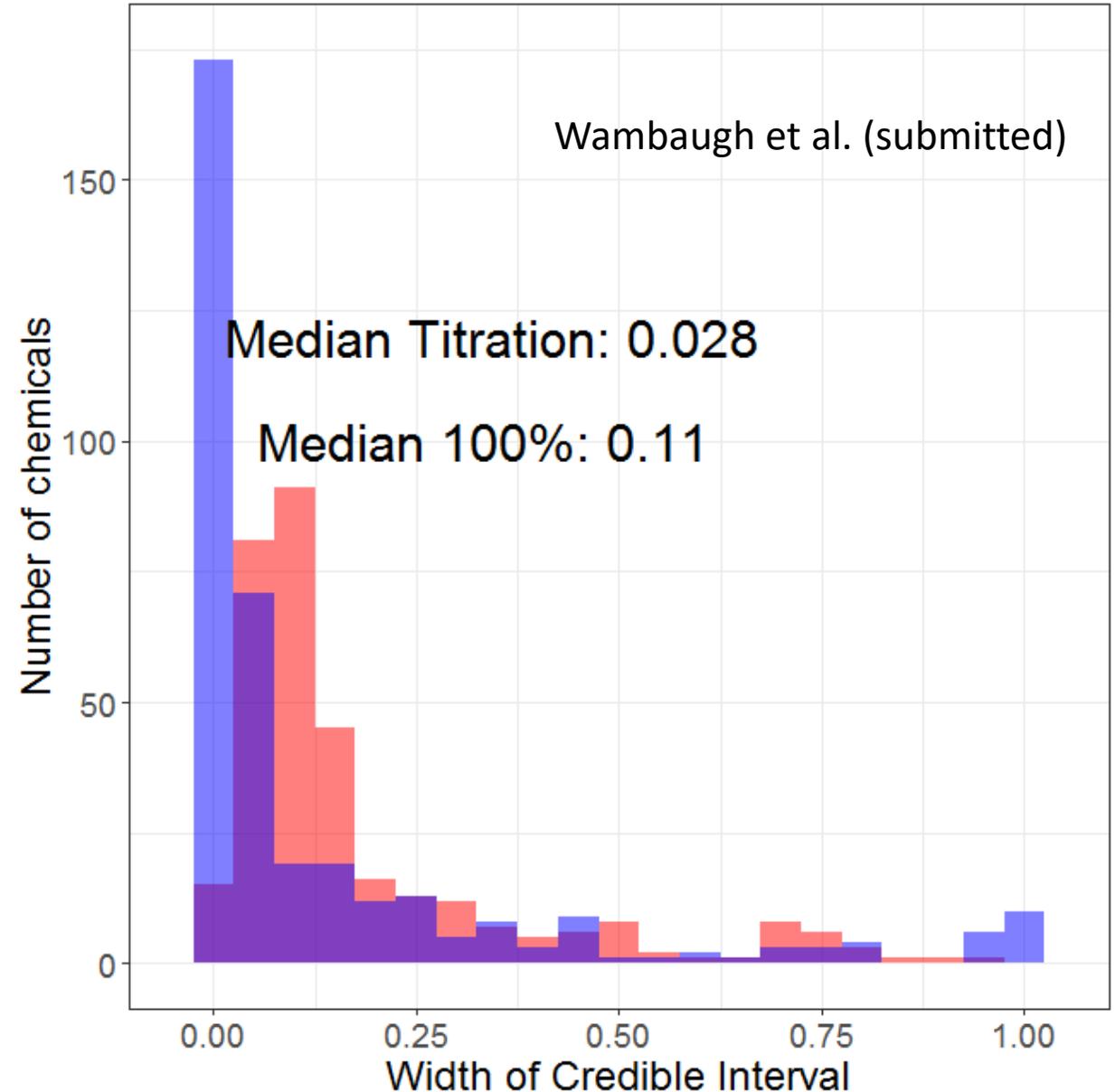
# New Plasma Binding Protocol Reduces Uncertainty

- 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

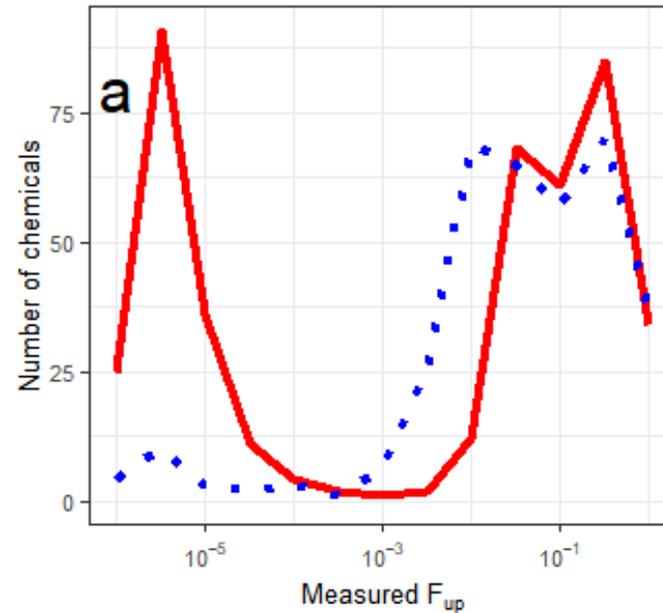
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- Median uncertainty for three-point assay:  $\pm 1.4\%$



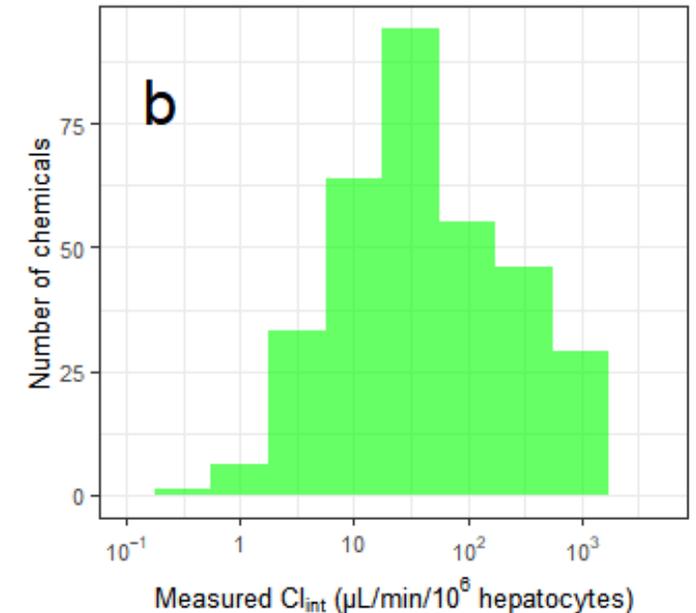
# New Data!

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%.

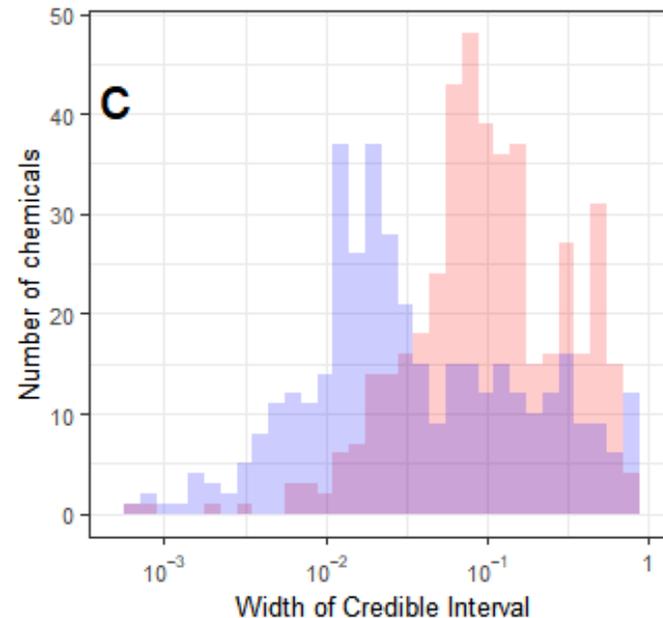
418 Chemicals Measured



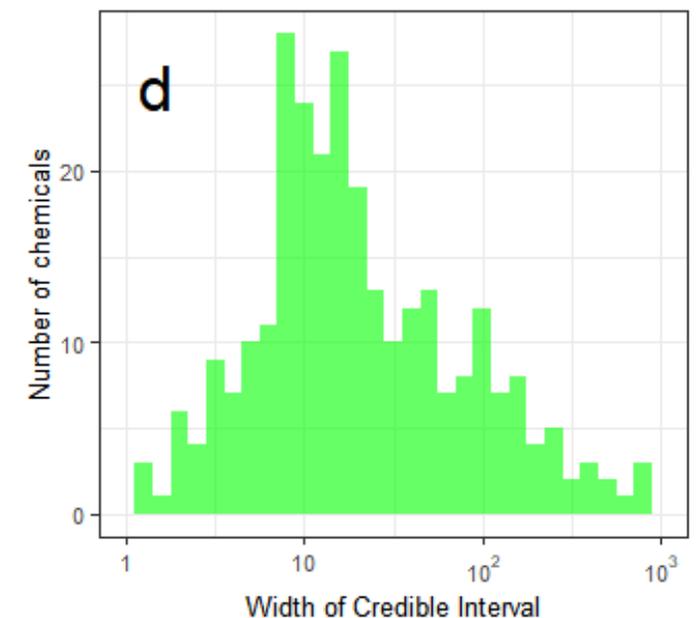
467 Chemicals Measured



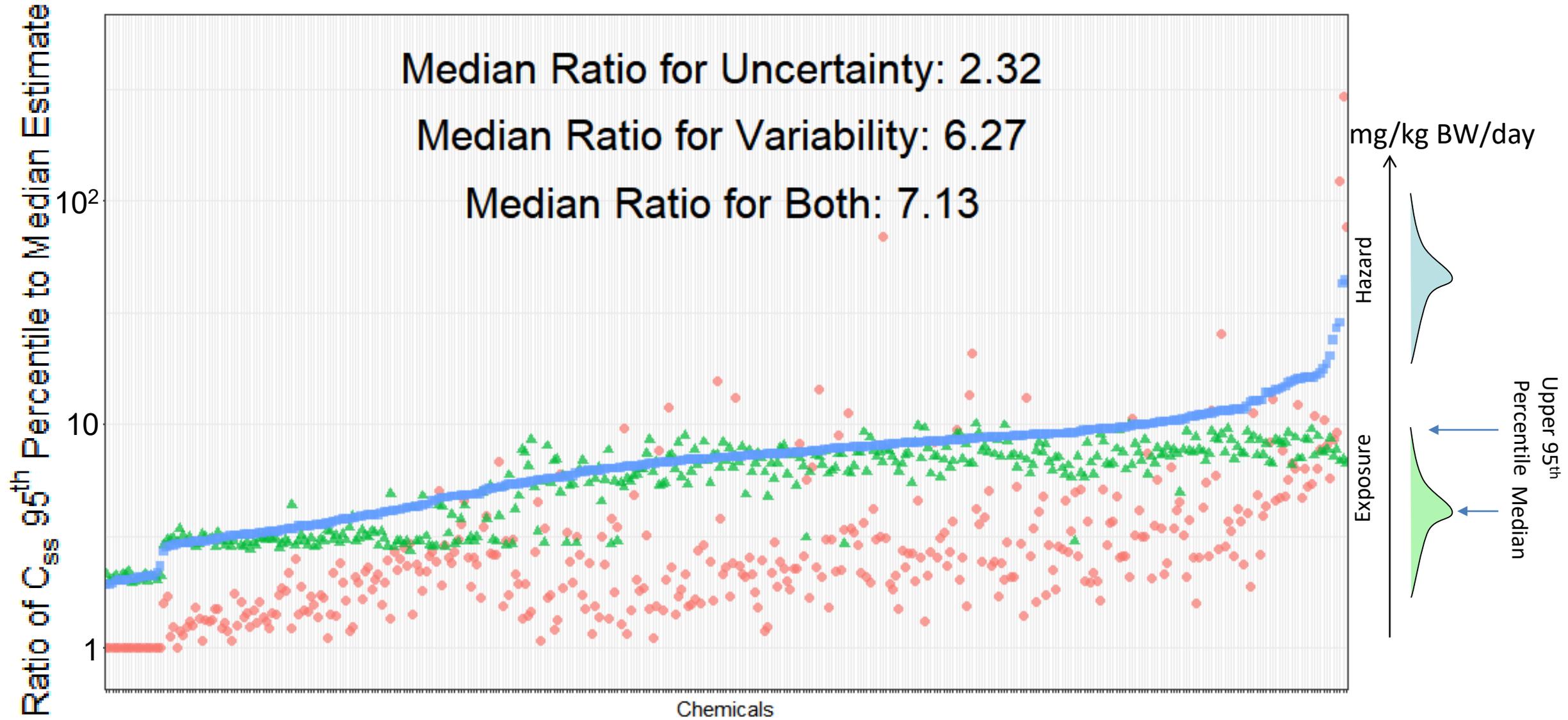
Cl: 0.1 CV: 0.4 / Cl: 0.03 CV: 0.1



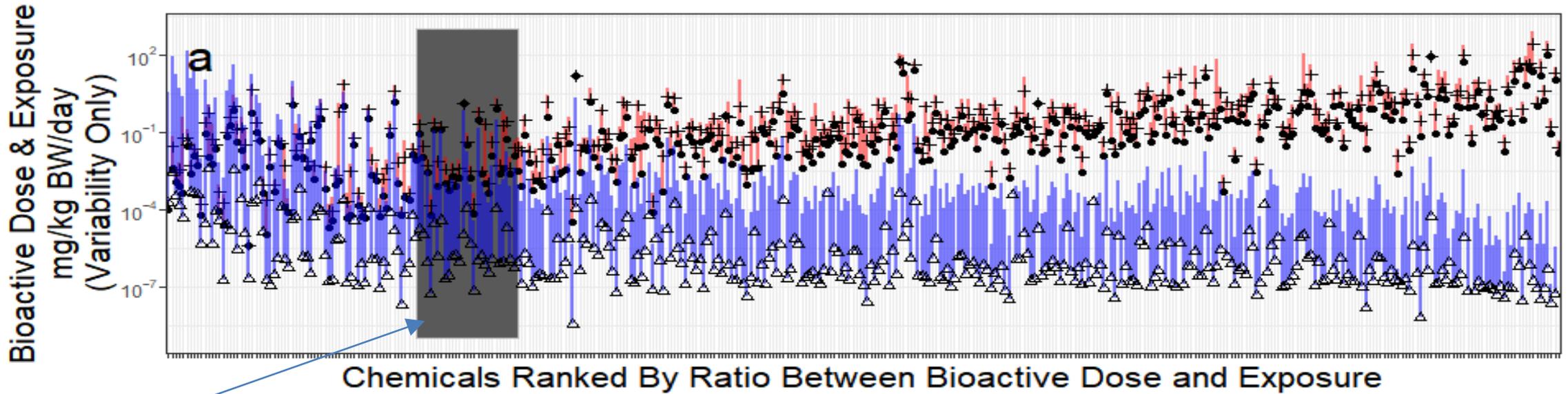
Median Cl: 14 CV: 0.31



# Quantifying the Impact of Uncertainty

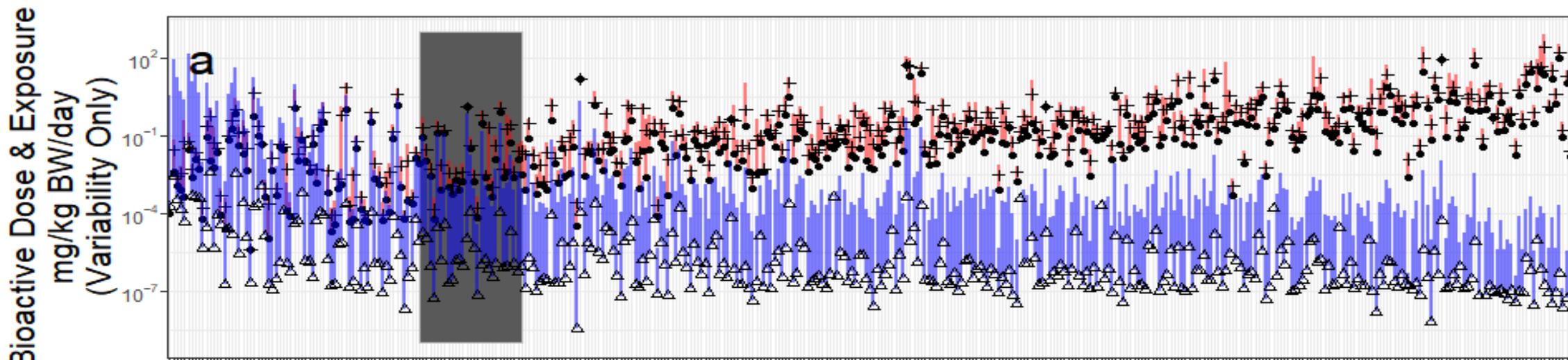


# New IVIVE For 393 ToxCast Chemicals

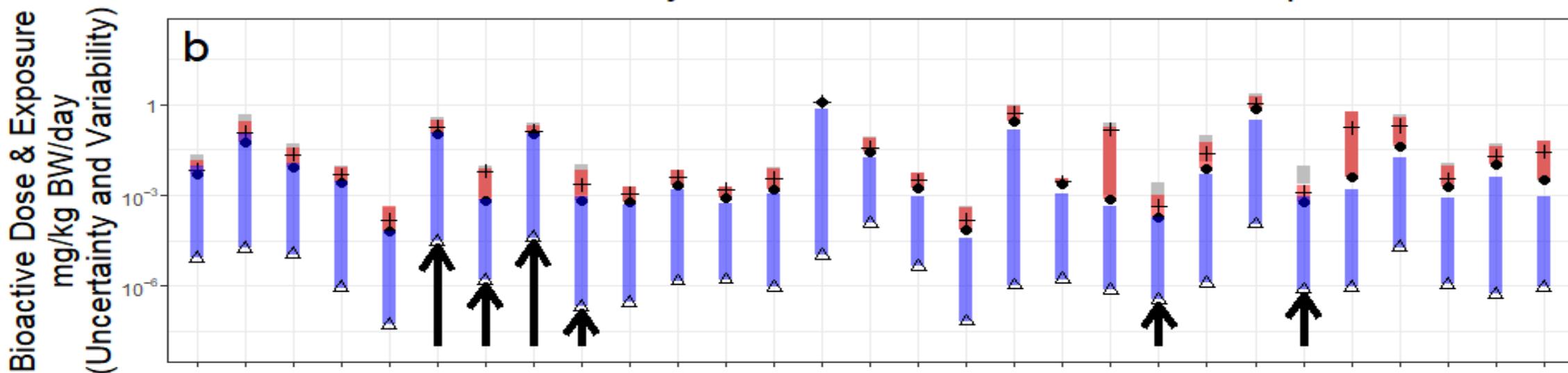


Including chemical-specific uncertainty only caused changes in whether or not exposure and bioactivity overlapped in a small region

# The Impact of Measurement Uncertainty



Chemicals Ranked By Ratio Between Bioactive Dose and Exposure

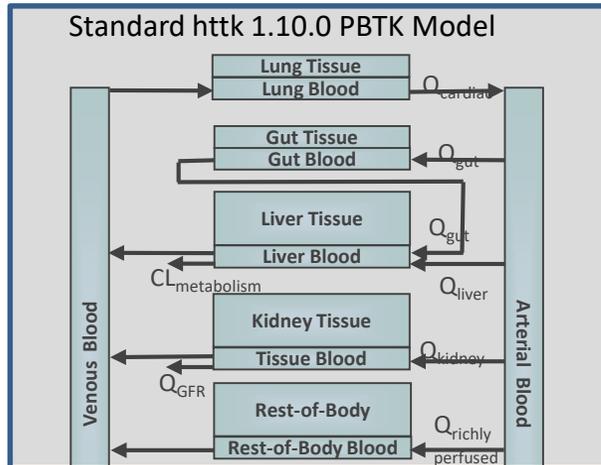


Only six more chemicals overlap

Wambaugh et al. (submitted)

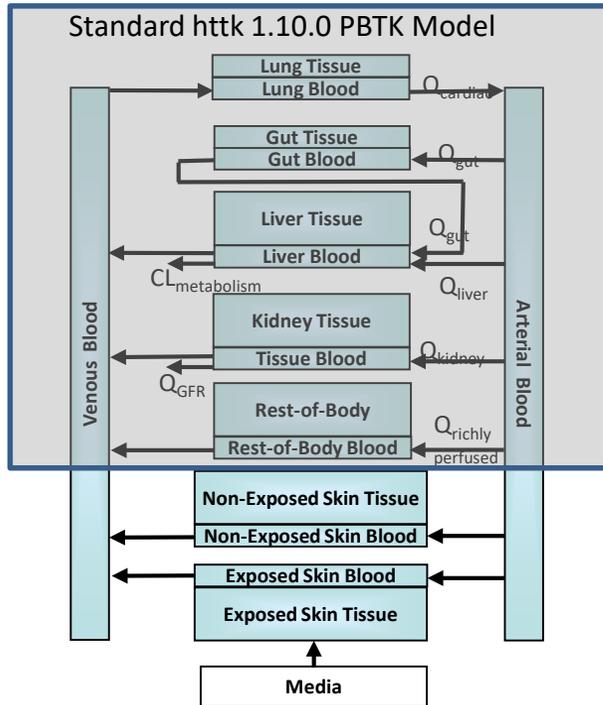
## New HT-PBTK Models

- 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 material transfer agreement (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

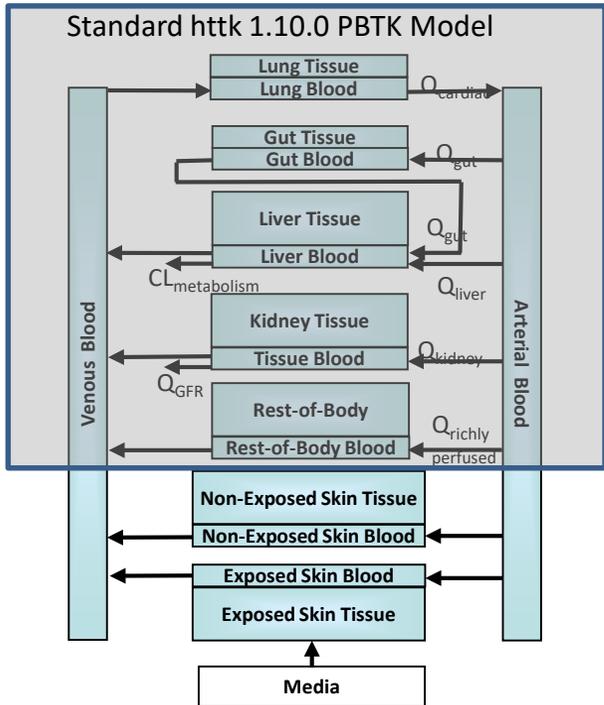
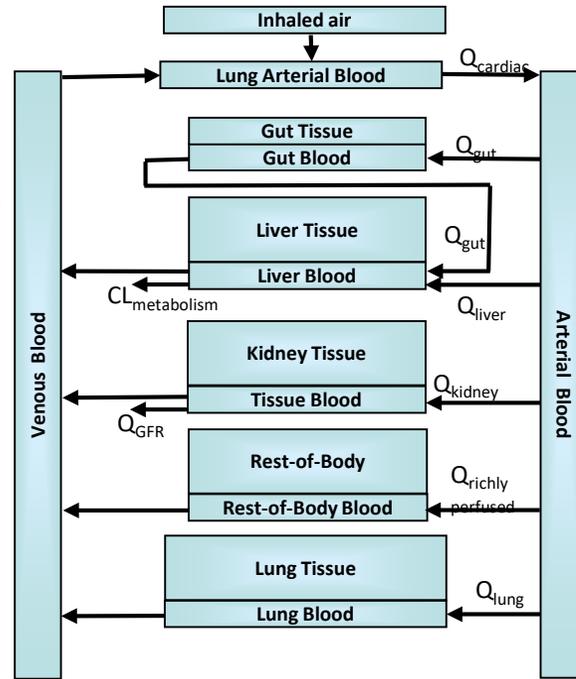


## Dermal Exposure Route

EPA, Unilever, INERIS

# New HT-PBTK Models

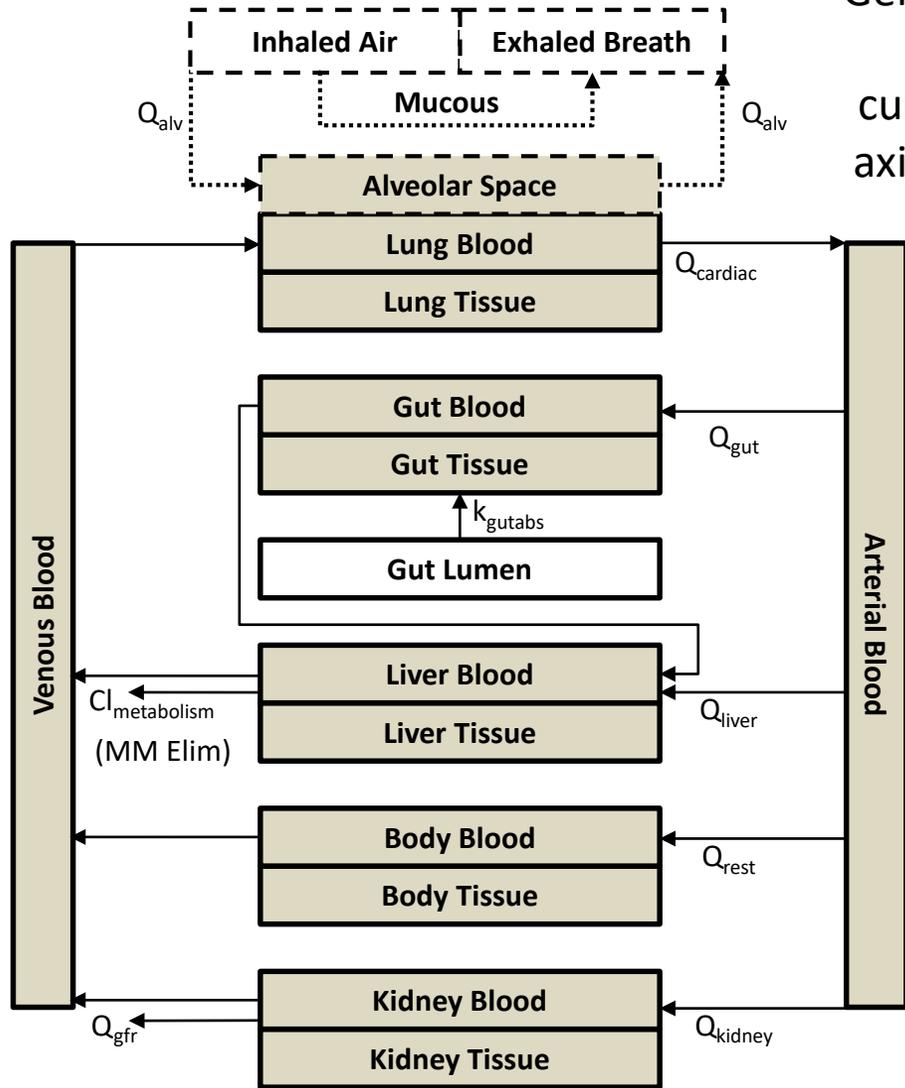
## Gas Inhalation Exposure Route EPA, USAFSAM



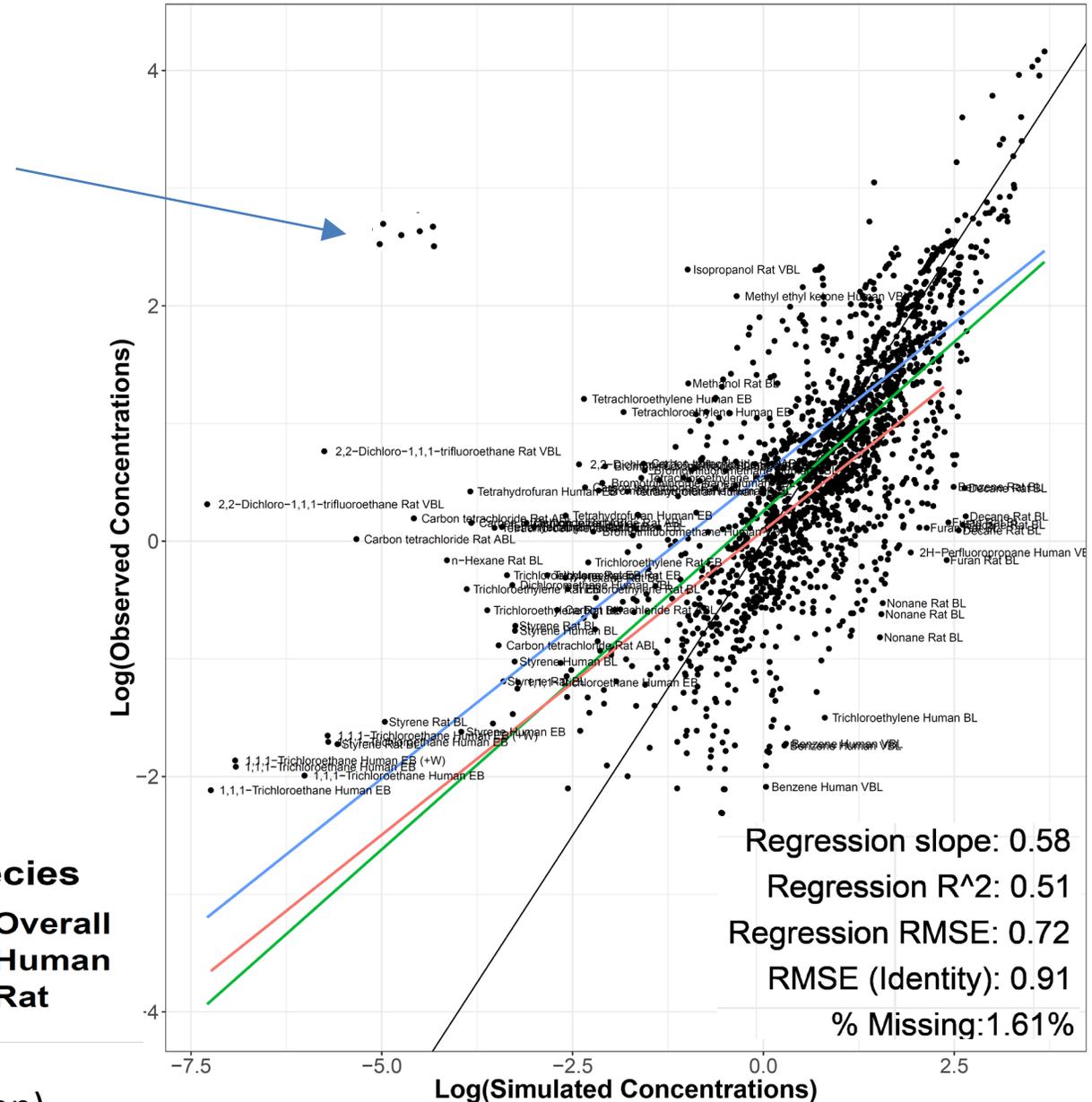
## Dermal Exposure Route EPA, Unilever, INERIS



# Generic Gas Inhalation Model

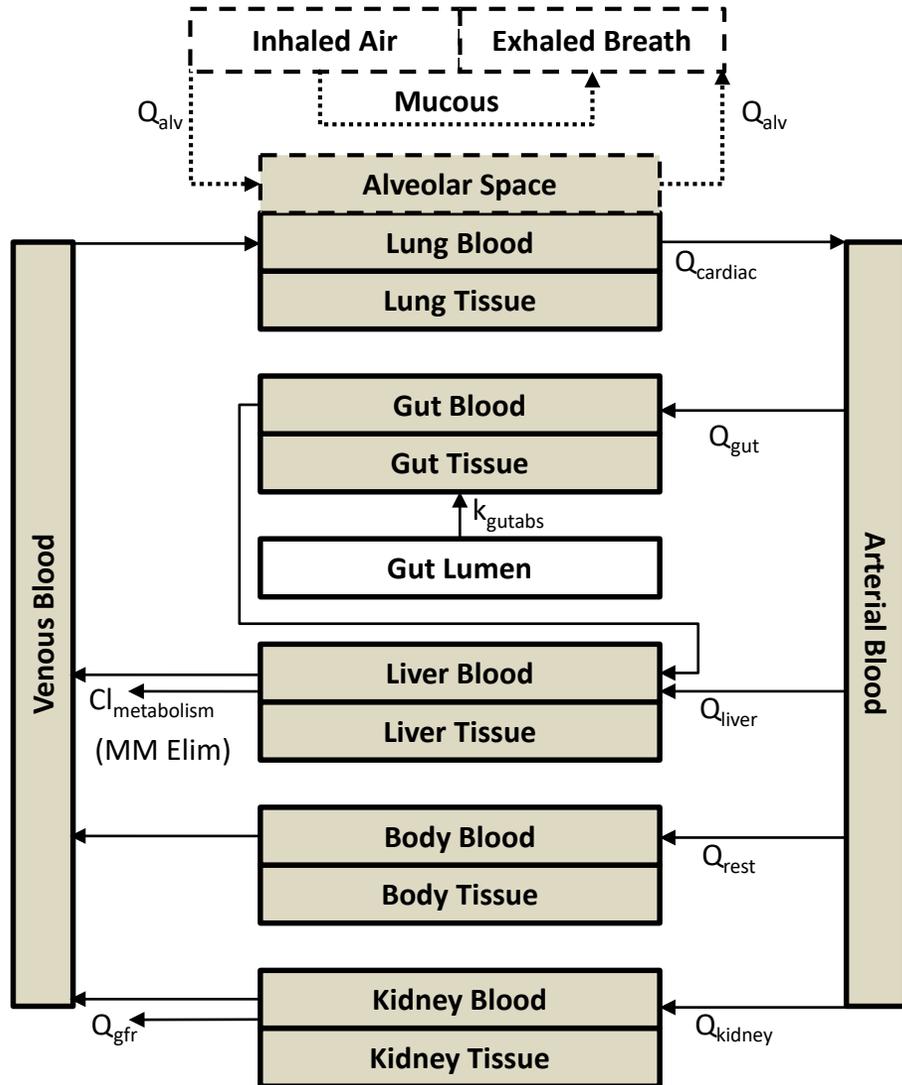


Generic model also helped data curation (units on axis in paper were wrong)

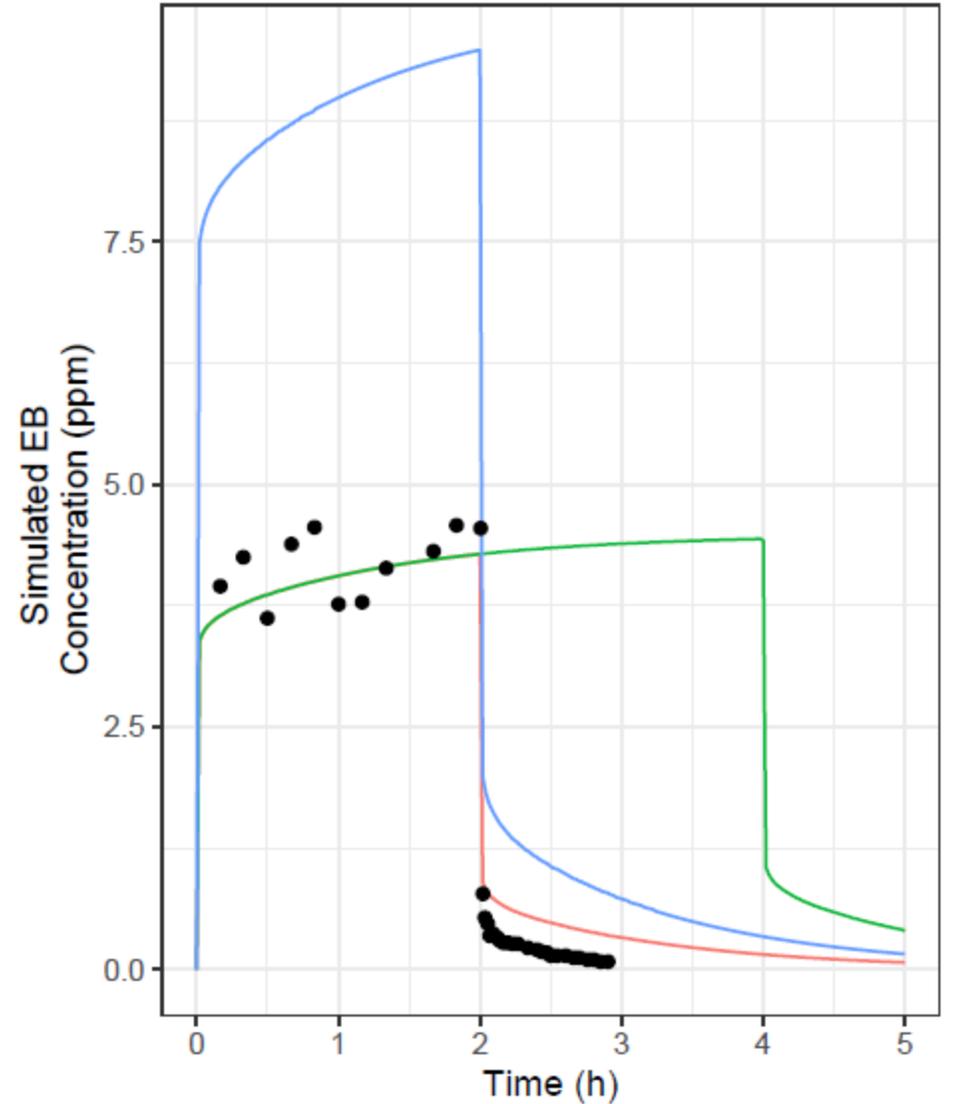


Linakis et al. (in preparation)

# Generic Gas Inhalation Model



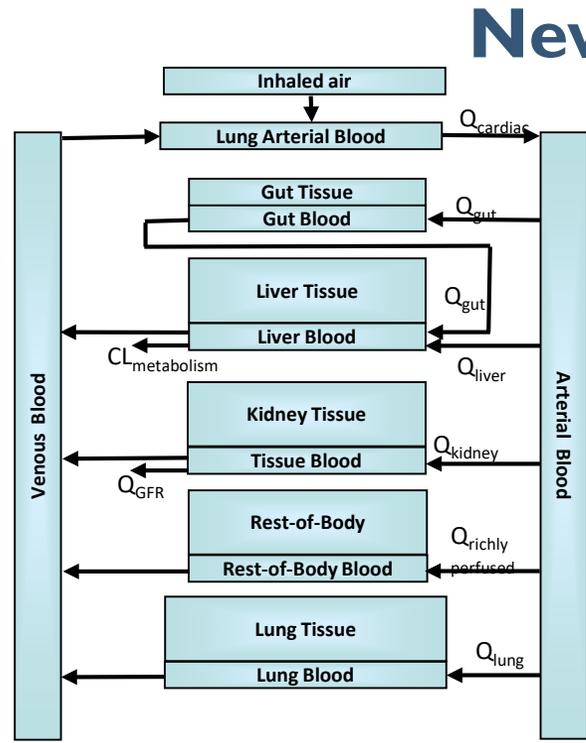
1,3-Butadiene (Human, 5ppm for 2h in EB)



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

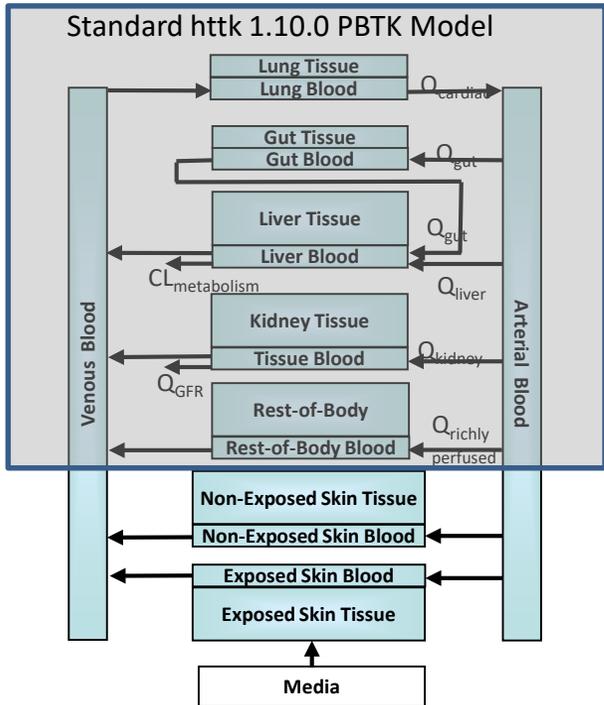
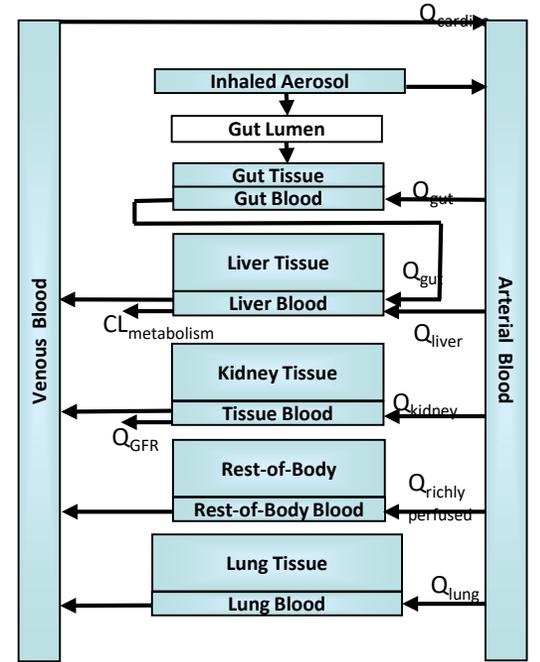
Figure from Matt Linakis (USAFSAM)

**Gas Inhalation  
Exposure Route**  
EPA, USAFSAM



**New HT-PBTK Models**

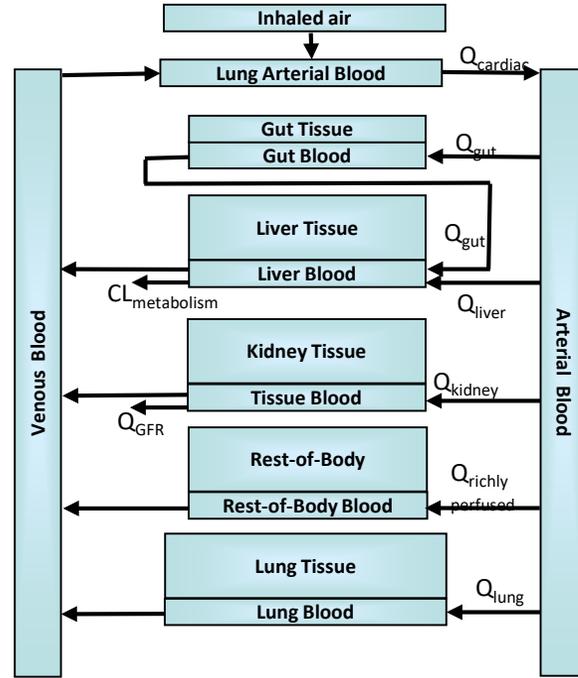
**Aerosol Inhalation  
Exposure Route  
(with APEX model)**  
EPA, USAFSAM



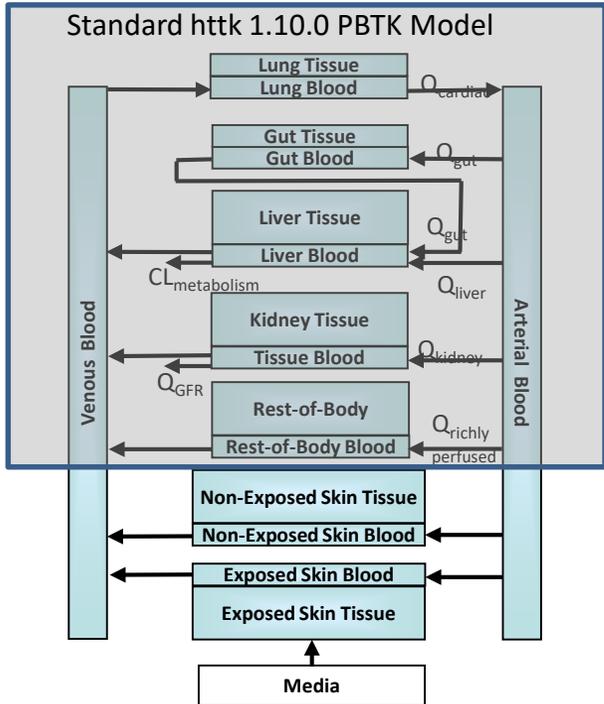
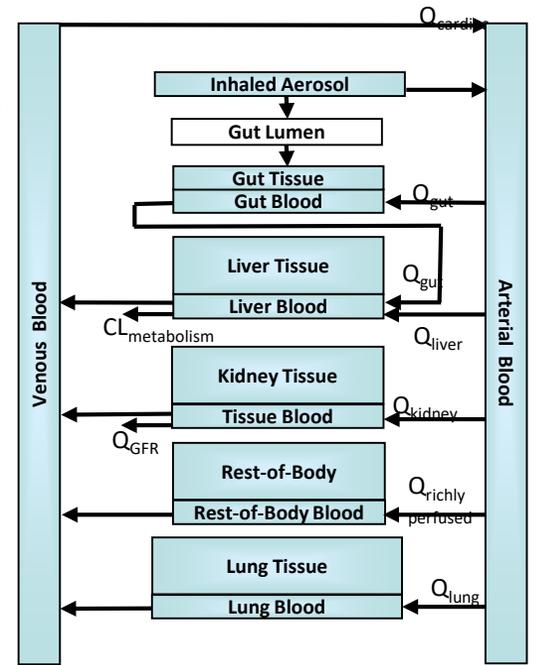
**Dermal Exposure Route**  
EPA, Unilever, INERIS

# New HT-PBTK Models

**Gas Inhalation  
Exposure Route**  
EPA, USAFSAM

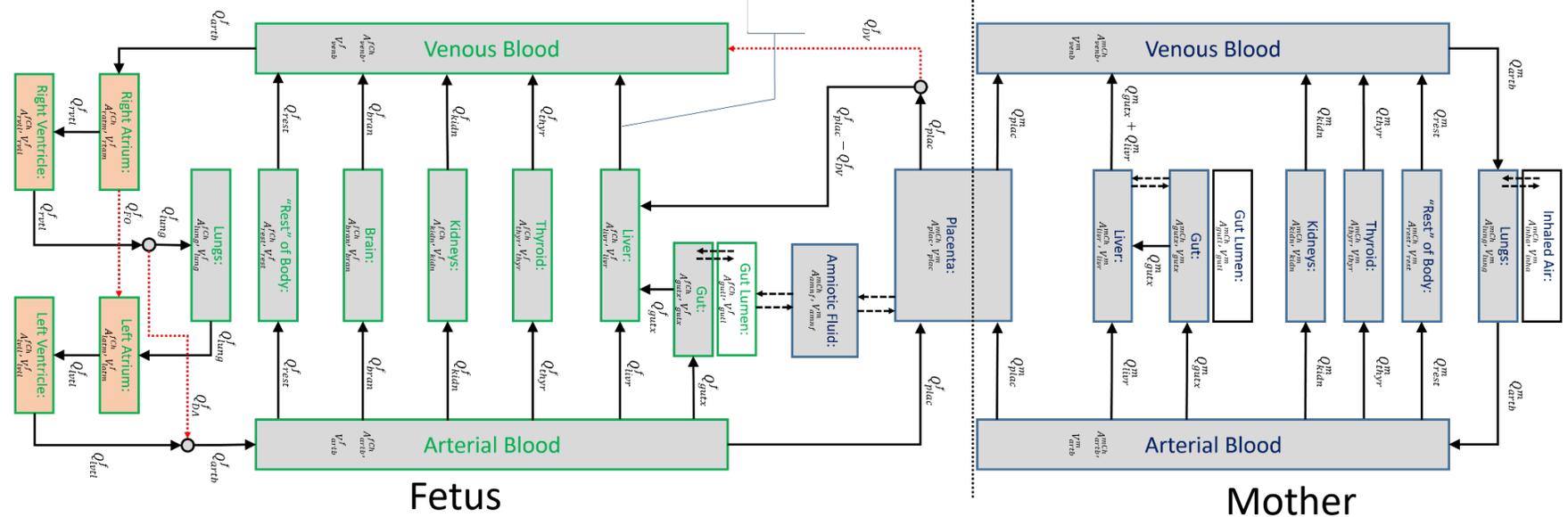


**Aerosol Inhalation  
Exposure Route  
(with APEX model)**  
EPA, USAFSAM



**Dermal Exposure Route**  
EPA, Unilever, INERIS

**Human Gestational Model**  
EPA, FDA

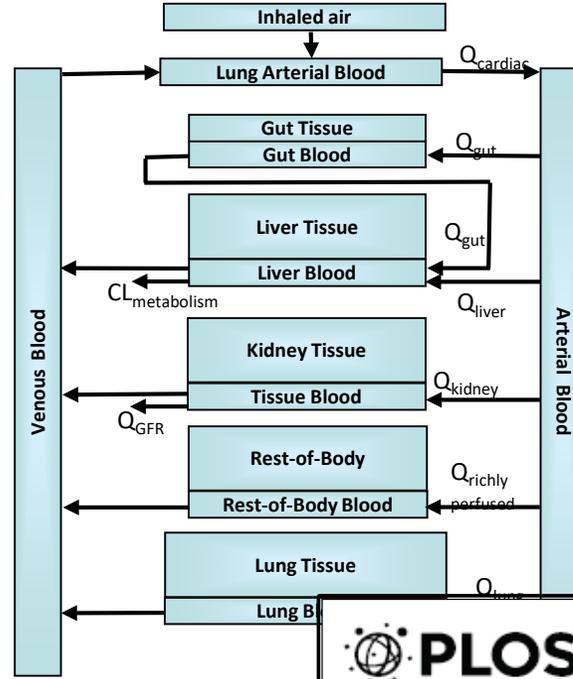


Fetus

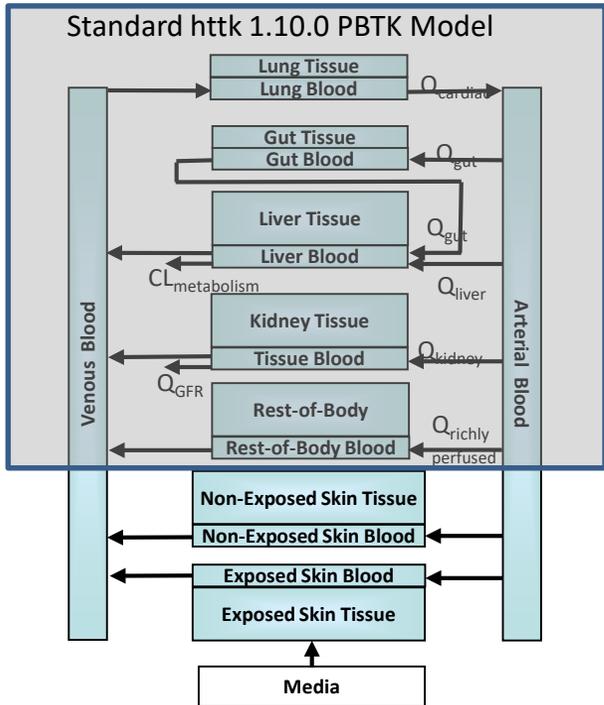
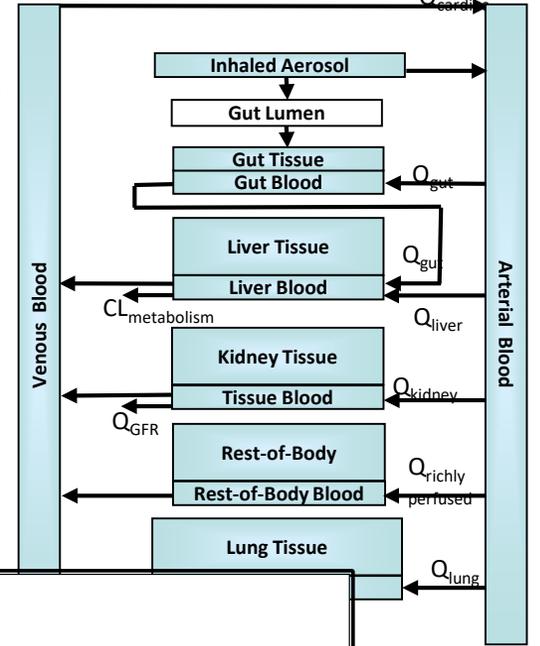
Mother

# 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

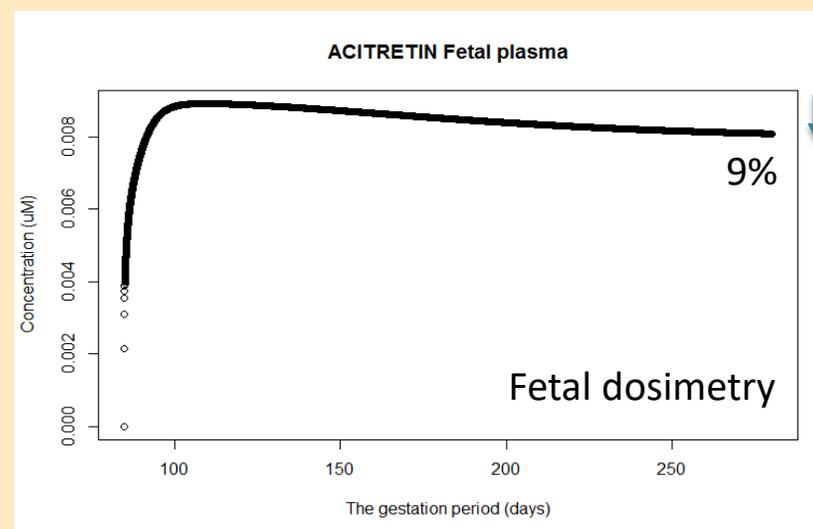
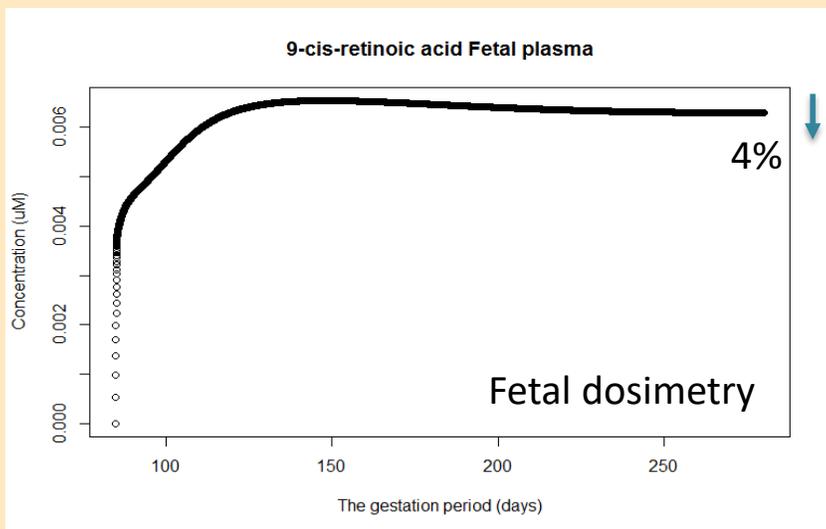
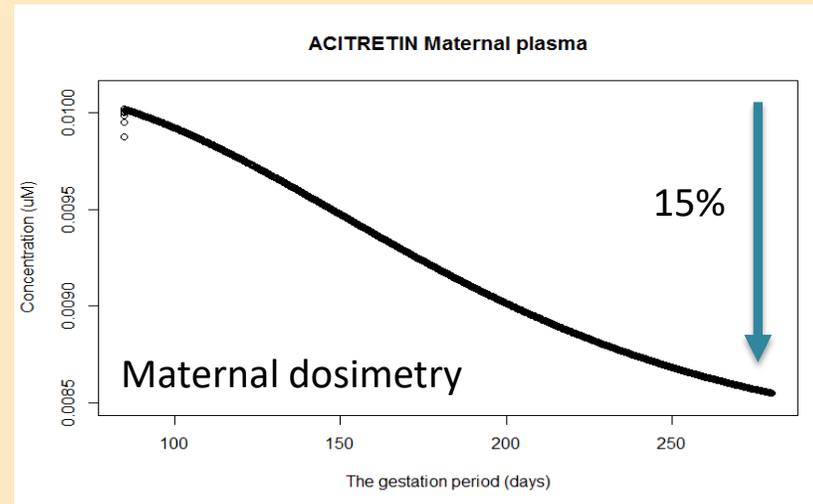
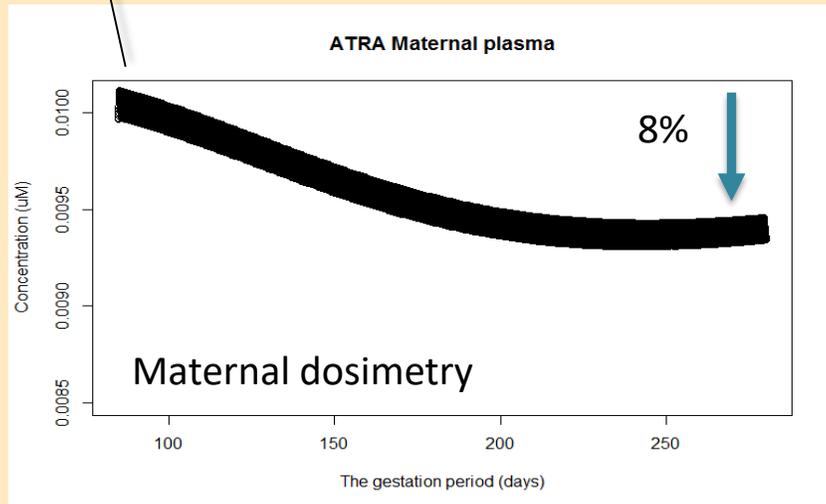
Dustin F. Kapraun<sup>1\*</sup>, John F. Wambaugh<sup>2</sup>, R. Woodrow Setzer<sup>2</sup>, Richard S. Judson<sup>2</sup>

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

This diagram shows a human circulatory system with the following components: 'Right Ventricle', 'Right Atrium', 'Left Ventricle', 'Left Atrium', 'Lungs', 'Rest of Body', and 'Inhaled Air'. Blood flow is indicated by arrows and labeled with  $Q_{cardiac}$ ,  $Q_{lung}$ ,  $Q_{rest}$ ,  $Q_{inhalation}$ , and  $Q_{exhalation}$ . The diagram illustrates the flow of blood from the heart to the lungs and then to the rest of the body, and back.

Normalized initial plasma concentration for each retinoid analogue

## Maternal/Fetal HTTK Model Predictions for Retinoid Analogues:

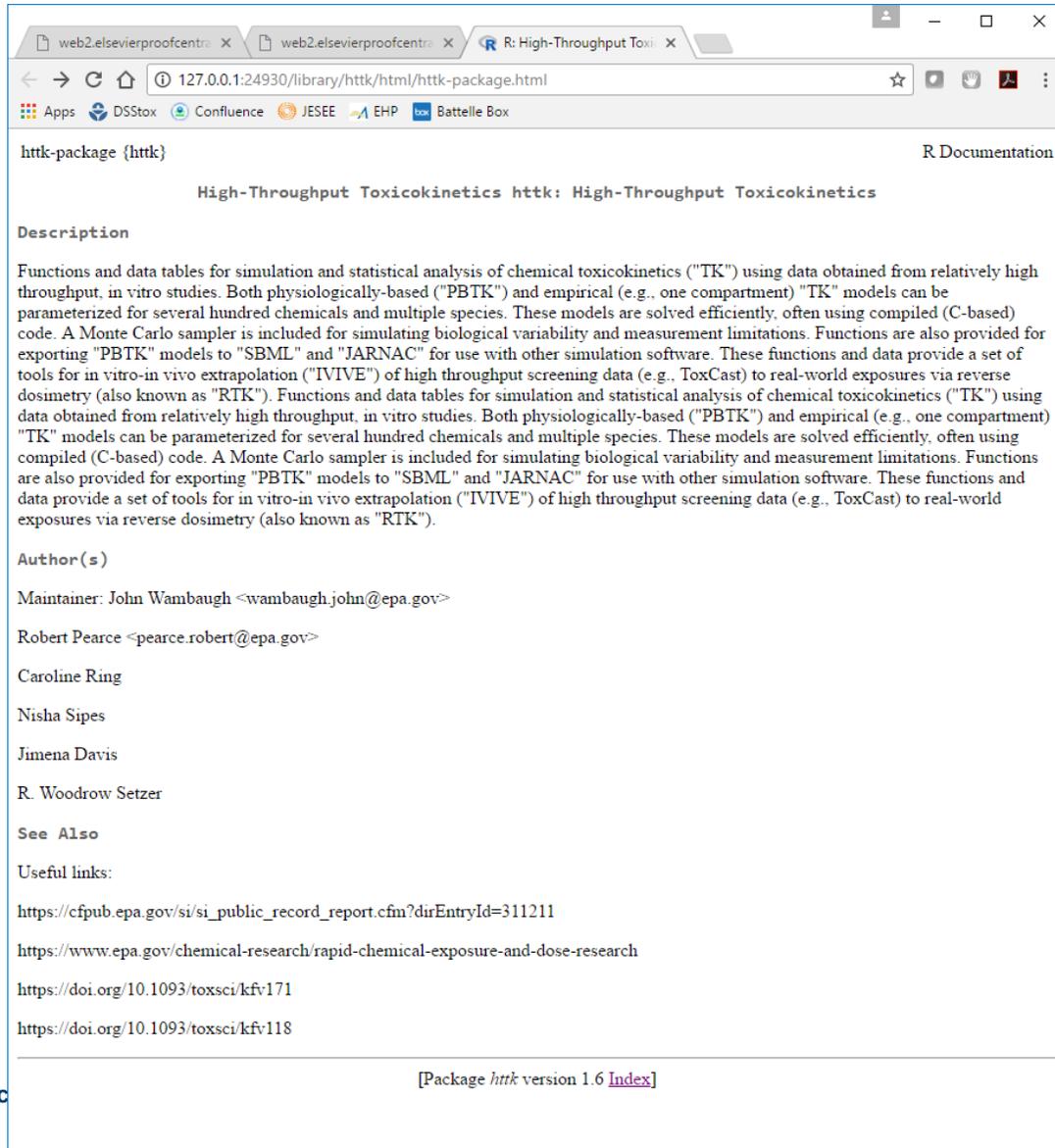


- Decrease in maternal plasma concentrations for retinoid analogues ranged from 8-15%

- 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 the URL '127.0.0.1:24930/library/httk/html/httk-package.html'. The page title is 'httk-package {httk}' and it is identified 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 the maintainer John Wambaugh and other contributors: Robert Pearce, Caroline Ring, Nisha Sipes, Jimena Davis, and R. Woodrow Setzer. A 'See Also' section and 'Useful links' are also present, with links to EPA public record reports and research pages. At the bottom, it indicates the package version is 1.6 and provides a link to the index.

httk-package {httk} R Documentation

**High-Throughput Toxicokinetics httk: High-Throughput Toxicokinetics**

**Description**

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"). 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").

**Author(s)**

Maintainer: John Wambaugh <wambaugh.john@epa.gov>  
Robert Pearce <pearce.robert@epa.gov>  
Caroline Ring  
Nisha Sipes  
Jimena Davis  
R. Woodrow Setzer

**See Also**

**Useful links:**

[https://cfpub.epa.gov/si/si\\_public\\_record\\_report.cfm?dirEntryId=311211](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>  
<https://doi.org/10.1093/toxsci/kfv118>

---

[Package *httk* version 1.6 [Index](#)]

# HTTK is (mostly) Documented

Within R: type "help(httk)"

The screenshot displays the R help environment for the 'httk' package. The left pane shows the R help text, and the right pane shows the web-based documentation.

**Left Pane (R help text):**

`httk-package {httk}`

High-Thro

**Description**

Functions and data tables for simulat  
throughput, in vitro studies. Both phy  
parameterized for several hundred of  
code. A Monte Carlo sampler is incl  
exporting "PBTk" models to "SBMI  
tools for in vitro-in vivo extrapolatio  
dosimetry (also known as "RTK"). F  
data obtained from relatively high th  
"TK" models can be parameterized f  
compiled (C-based) code. A Monte (C  
are also provided for exporting "PBT  
data provide a set of tools for in vitro  
exposures via reverse dosimetry (als

**Author(s)**

Maintainer: John Wambaugh <wamb  
Robert Pearce <pearce.robert@epa.g  
Caroline Ring  
Nisha Sipes  
Jimena Davis  
R. Woodrow Setzer

**See Also**

Useful links:  
[https://cfpub.epa.gov/si/si\\_public\\_re](https://cfpub.epa.gov/si/si_public_re)  
<https://www.epa.gov/chemical-resea>  
<https://doi.org/10.1093/toxsci/kfv171>  
<https://doi.org/10.1093/toxsci/kfv118>

**Right Pane (Web-based documentation):**

High-Throughput Toxicokinetics

Documentation for package 'httk' version 1.6

- [DESCRIPTION file.](#)
- [User guides, package vignettes and other documentation.](#)
- [Package NEWS.](#)

**Help Pages**

[A](#) [B](#) [C](#) [D](#) [E](#) [F](#) [G](#) [H](#) [I](#) [J](#) [K](#) [L](#) [M](#) [N](#) [O](#) [P](#) [R](#) [S](#) [T](#) [W](#)

[httk-package](#) High-Throughput Toxicokinetics httk: High-Throughput Toxicokinetics  
[httkpop-package](#) httkpop: Virtual population generator for HTTK.

-- A --

[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  
[available\\_rblood2plasma](#) Find the best available ratio of the blood to plasma concentration constant.

-- B --

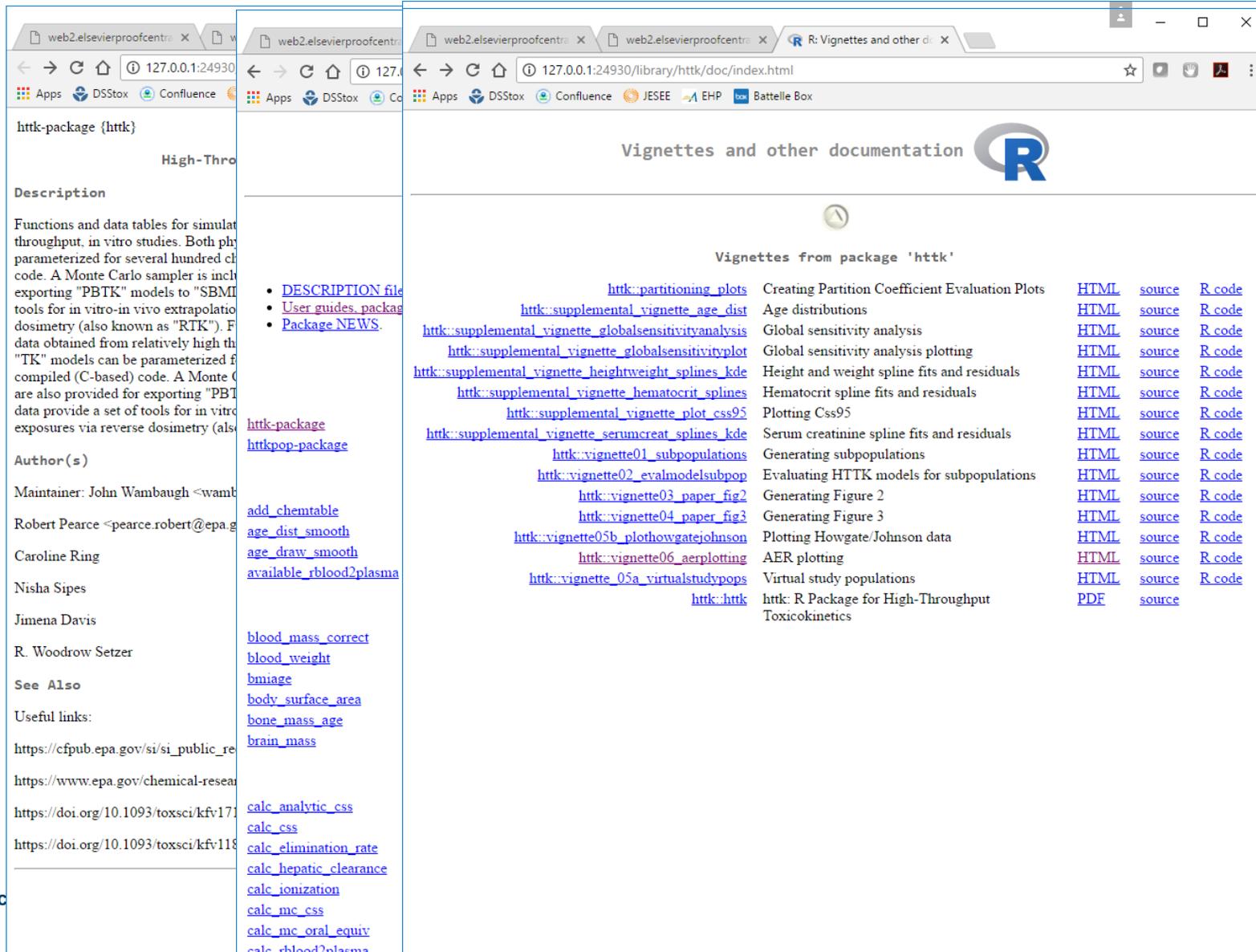
[blood\\_mass\\_correct](#) Find average blood masses by age.  
[blood\\_weight](#) Predict blood mass.  
[bmiage](#) CDC BMI-for-age charts  
[body\\_surface\\_area](#) Predict body surface area.  
[bone\\_mass\\_age](#) Predict bone mass.  
[brain\\_mass](#) Predict brain mass.

-- C --

[calc\\_analytic\\_css](#) Calculate the analytic steady state concentration.  
[calc\\_css](#) Find the steady state concentration and the day it is reached.  
[calc\\_elimination\\_rate](#) Calculate the elimination rate for a one compartment model.  
[calc\\_hepatic\\_clearance](#) Calculate the hepatic clearance.  
[calc\\_ionization](#) Calculate the ionization.  
[calc\\_mc\\_css](#) Calculate the monte carlo steady state concentration.  
[calc\\_mc\\_oral\\_equiv](#) Calculate Monte Carlo Oral Equivalent Dose  
[calc\\_rblood2plasma](#) Calculate the constant ratio of the blood concentration to the plasma concentration

# 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 is as follows:

**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"). F 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)**

Maintainer: John Wambaugh <wambaugh.john@epa.gov>  
Robert Pearce <pearce.robert@epa.gov>  
Caroline Ring  
Nisha Sipes  
Jimena Davis  
R. Woodrow Setzer

**See Also**

[add\\_chemtable](#)  
[age\\_dist\\_smooth](#)  
[age\\_draw\\_smooth](#)  
[available\\_rblood2plasma](#)  
[blood\\_mass\\_correct](#)  
[blood\\_weight](#)  
[bmiage](#)  
[body\\_surface\\_area](#)  
[bone\\_mass\\_age](#)  
[brain\\_mass](#)

**Useful links:**

[https://cfpub.epa.gov/si/si\\_public\\_research/](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
<a href="#">http://partitioning_plots</a>	Creating Partition Coefficient Evaluation Plots	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_age_dist</a>	Age distributions	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_globalsensitivityanalysis</a>	Global sensitivity analysis	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_globalsensitivityplot</a>	Global sensitivity analysis plotting	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_heightweight_splines_kde</a>	Height and weight spline fits and residuals	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_hematocrit_splines</a>	Hematocrit spline fits and residuals	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_plot_css95</a>	Plotting Css95	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://supplemental_vignette_serumcreat_splines_kde</a>	Serum creatinine spline fits and residuals	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette01_subpopulations</a>	Generating subpopulations	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette02_evalmodelsubpop</a>	Evaluating HTTK models for subpopulations	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette03_paper_fig2</a>	Generating Figure 2	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette04_paper_fig3</a>	Generating Figure 3	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette05b_plothowgatejohnson</a>	Plotting Howgate/Johnson data	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette06_aerplotting</a>	AER plotting	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://vignette_05a_virtualstudypops</a>	Virtual study populations	<a href="#">HTML</a>	<a href="#">source</a>	<a href="#">R code</a>
<a href="#">http://httk</a>	httk: R Package for High-Throughput Toxicokinetics	<a href="#">PDF</a>	<a href="#">source</a>	



# EXAMPLE: Does My Chemical Have HHTK Data?

Is a chemical available?

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

All data on chemicals A, B, C

```
subset(get_cheminfo(info="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")
```

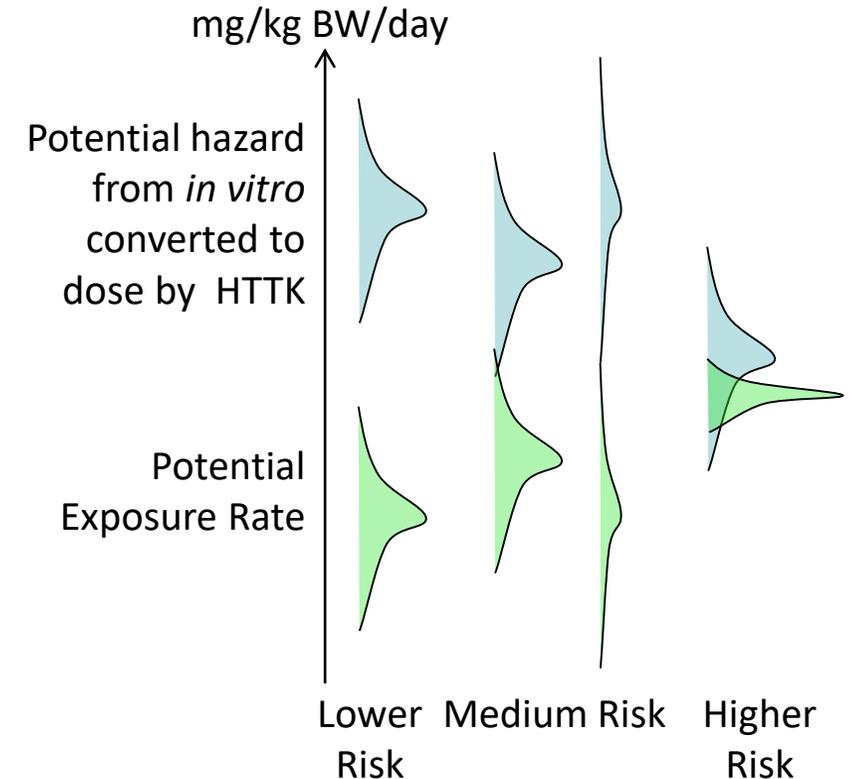
Compound	CAS	logP	pKa_Accept	pKa_Donor	MW	Human.Clint	Human.Clint.p Value	Human.Funbou nd.plasma	DSSTox_Substance_ID	Structure_Formula	Substance_Type
2,4-d	94-75-7	2.81	<NA>	2.81	221.03	0	0.149	0.04	DTXSID0020442	C8H6Cl2O3	Single Compound
2,4-db	94-82-6	3.53	<NA>	4.5	249.09	0	0.104	0.01	DTXSID7024035	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

# HTTK Limitations

- Oral absorption
  - 100% assumed, but may be very different
  - *In silico* models not necessarily appropriate for environmental chemicals
  - Honda et al. (in preparation) developing QSAR using new *in vitro* data for ToxCast Chemicals
- Hepatic Clearance ( $CL_{int}$ )
  - Not isozyme-specific (Isozyme-specific metabolism assays not HT)
  - Ten donor pool in suspension for 2-4 h misses variability and low turnover compounds
  - Isozyme abundances and activity: varies with age, ethnicity (at least) (Yasuda et al. 2008, Howgate et al. 2006, Johnson et al. 2006)
  - Parent chemical depletion only
  - *In silico* predictions of isozyme-specific metabolism? Not easy!
    - Though ADMET Predictor can do this for some isozymes, training data is mostly for pharmaceuticals
- Plasma binding assay ( $F_{up}$ )
  - Plasma protein concentration variability (Johnson et al. 2006, Israili et al. 2001)
  - Albumin or AAG binding? (Routledge 1986)
- Analytical chemistry
  - Must be able to develop method for each compound
  - Working to develop QSARs for other compounds

# Conclusions

- HTKK 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 95<sup>th</sup> percentile
- Comparison predicted concentrations and *in vivo* data is a valuable approach for evaluation and establishing confidence
- A new database of *in vivo* concentration vs. time data is being developed (Sayre, in preparation)



<https://cran.r-project.org/package=httk>

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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Absorption**

Dustin Kapraun  
Richard Judson  
Annie Lumen (FDA)

Mark Sfeir

**Package Tsar**

**Human  
Gestation**



**Inhalation**

**HTTK Team**

**Cyprotex  
(lab work)**

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Maria Bacolod	Jon Gilbert
Akshay	Teresa Sierra
Badrinarayanan	Bradley
Adam Brockman	Snodgrass
Roger Dinallo	Chris Strock

Matt Linakis (USAFSAM)  
Heather Pangburn (USAFSAM)  
Jeffery Gearhart (USAFSAM)  
Nisha Sipes (NTP)  
Kristin Isaacs

**TK  
Database**

**Structure-Based  
Predictions**

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Brandall Ingle (ICF)	Prachi Pradeep
Richard Judson	Nisha Sipes (NTP)
Rogelio Tornero-Velez	

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Chris Grulke

Marina Evans  
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# ExpoCast Project (Exposure Forecasting)

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# References

- Breyer, Stephen. *Breaking the vicious circle: Toward effective risk regulation*. Harvard University Press, 2009
- Bosgra, Sieto, et al. "An improved model to predict physiologically based model parameters and their inter-individual variability from anthropometry." *Critical reviews in toxicology* 42.9 (2012): 751-767.
- Cohen, EA Hubal, et al. "Advancing internal exposure and physiologically-based toxicokinetic modeling for 21st-century risk assessments." *Journal of exposure science & environmental epidemiology* (2018).
- Collins FS, Gray GM, Bucher JR. *Transforming environmental health protection*. Science. 2008;319:906–907. [PMC free article] [PubMed]
- Dix David, et al. "The ToxCast program for prioritizing toxicity testing of environmental chemicals." *Toxicol Sci.* 2007;95:5–12
- Honda et al., "Using the concordance of in vitro and in vivo data to evaluate extrapolation assumptions" *PLoS ONE* 14.5 (2019): e0217564.
- Jongeneelen, Frans, and Wil Ten Berge. "Simulation of urinary excretion of 1-hydroxypyrene in various scenarios of exposure to polycyclic aromatic hydrocarbons with a generic, cross-chemical predictive PBTK-model." *International archives of occupational and environmental health* 85.6 (2012): 689-702.
- Kapraun, Dustin F., et al. "Empirical models for anatomical and physiological changes in a human mother and fetus during pregnancy and gestation." *PLoS one* 14.5 (2019): e0215906.
- Linakis, et al. "Development of a Generalized Inhalation Model for use with the High-Throughput Toxicokinetics (httk) Package in R", in preparation
- Lukacova, et al.. "Prediction of modified release pharmacokinetics and pharmacodynamics from in vitro, immediate release, and intravenous data." *The AAPS journal* 11.2 (2009): 323-334.
- McNally, et al., "PopGen: a virtual human population generator." *Toxicology* (2014)
- National Research Council. (1983). *Risk Assessment in the Federal Government: Managing the Process Working Papers*. National Academies Press.
- National Research Council. (2007). *Toxicity testing in the 21st century: a vision and a strategy*. National Academies Press.
- National Research Council. *Exposure Science in the 21st Century: a Vision and a Strategy*. National Academies Press, 2012.
- Park, Youngja H., et al. "High-performance metabolic profiling of plasma from seven mammalian species for simultaneous environmental chemical surveillance and bioeffect monitoring." *Toxicology* 295.1 (2012): 47-55.
- Pearce, Robert, et al. "httk: R Package for High-Throughput Toxicokinetics." *Journal of Statistical Software*, (2017)
- Pearce, Robert G., et al. "Evaluation and calibration of high-throughput predictions of chemical distribution to tissues." *Journal of pharmacokinetics and pharmacodynamics* 44.6 (2017): 549-565.
- Peyret, Thomas, Patrick Poulin, and Kannan Krishnan. "A unified algorithm for predicting partition coefficients for PBPK modeling of drugs and environmental chemicals." *Toxicology and Applied Pharmacology* 249.3 (2010): 197-207.
- Price, Paul S., et al. "Modeling interindividual variation in physiological factors used in PBPK models of humans." *Critical reviews in toxicology* 33.5 (2003): 469-503.
- Ring, Caroline L., et al. "Identifying populations sensitive to environmental chemicals by simulating toxicokinetic variability." *Environment International* 106 (2017): 105-118.
- Ring, Caroline L., et al. "Consensus Modeling of Median Chemical Intake Based on Predictions of Exposure Pathways", *Environmental science & technology* 53.2 (2018): 719-732. Rotroff, Daniel M., et al. "Incorporating human dosimetry and exposure into high-throughput in vitro toxicity screening." *Toxicological Sciences* 117.2 (2010): 348-358.
- Sayre, Risa et al., "Database of pharmacokinetic time-series data and parameters for XX environmental chemicals" in preparation
- Schmidt, Charles W. "TOX 21: new dimensions of toxicity testing." *Environmental health perspectives* 117.8 (2009): A348.
- Schmitt, Walter. "General approach for the calculation of tissue to plasma partition coefficients." *Toxicology in Vitro* 22.2 (2008): 457-467.
- Shibata, Y., et al. (2002). Prediction of hepatic clearance and availability by cryopreserved human hepatocytes: an application of serum incubation method. *Drug Metabolism and Disposition*, 30(8), 892-896
- Sipes, Nisha S., et al. "An intuitive approach for predicting potential human health risk with the Tox21 10k library." *Environmental Science & Technology* 51.18 (2017): 10786-10796.
- Strobe, Cory L., et al. "High-throughput in-silico prediction of ionization equilibria for pharmacokinetic modeling." *Science of The Total Environment* 615 (2018): 150-160.
- Wambaugh, John F., et al. "High-throughput models for exposure-based chemical prioritization in the ExpoCast project." *Environmental science & technology* 47.15 (2013): 8479-8488.
- Wambaugh, John F., et al. "High Throughput Heuristics for Prioritizing Human Exposure to Environmental Chemicals." *Environmental science & technology* (2014).
- Wambaugh, John F., et al. "Toxicokinetic triage for environmental chemicals." *Toxicological Sciences* 147.1 (2015): 55-67.
- Wambaugh, John F., et al. "Evaluating In Vitro-In Vivo Extrapolation of Toxicokinetics." *Toxicological Sciences* 163.1 (2018): 152-169.
- Wambaugh, John F., et al. "Assessing Toxicokinetic Uncertainty and Variability in Risk Prioritization", submitted.
- Wang, Ying-Hong. "Confidence assessment of the Simcyp time-based approach and a static mathematical model in predicting clinical drug-drug interactions for mechanism-based CYP3A inhibitors." *Drug Metabolism and Disposition* 38.7 (2010): 1094-1104.
- Wetmore, Barbara A., et al. "Integration of dosimetry, exposure and high-throughput screening data in chemical toxicity assessment." *Tox. Sciences* (2012)
- Wetmore, Barbara A., et al. "Relative impact of incorporating pharmacokinetics on predicting in vivo hazard and mode of action from high-throughput in vitro toxicity assays." *toxicological sciences* 132.2 (2013): 327-346.
- Wetmore, Barbara A., et al. "Incorporating high-throughput exposure predictions with dosimetry-adjusted in vitro bioactivity to inform chemical toxicity testing." *Toxicological Sciences* 148.1 (2015): 121-136.
- Waters, Nigel J., et al. "Validation of a rapid equilibrium dialysis approach for the measurement of plasma protein binding." *Journal of pharmaceutical sciences* 97.10 (2008): 4586-4595