

The High Throughput Toxicokinetic (HTTK) R Package

John Wambaugh

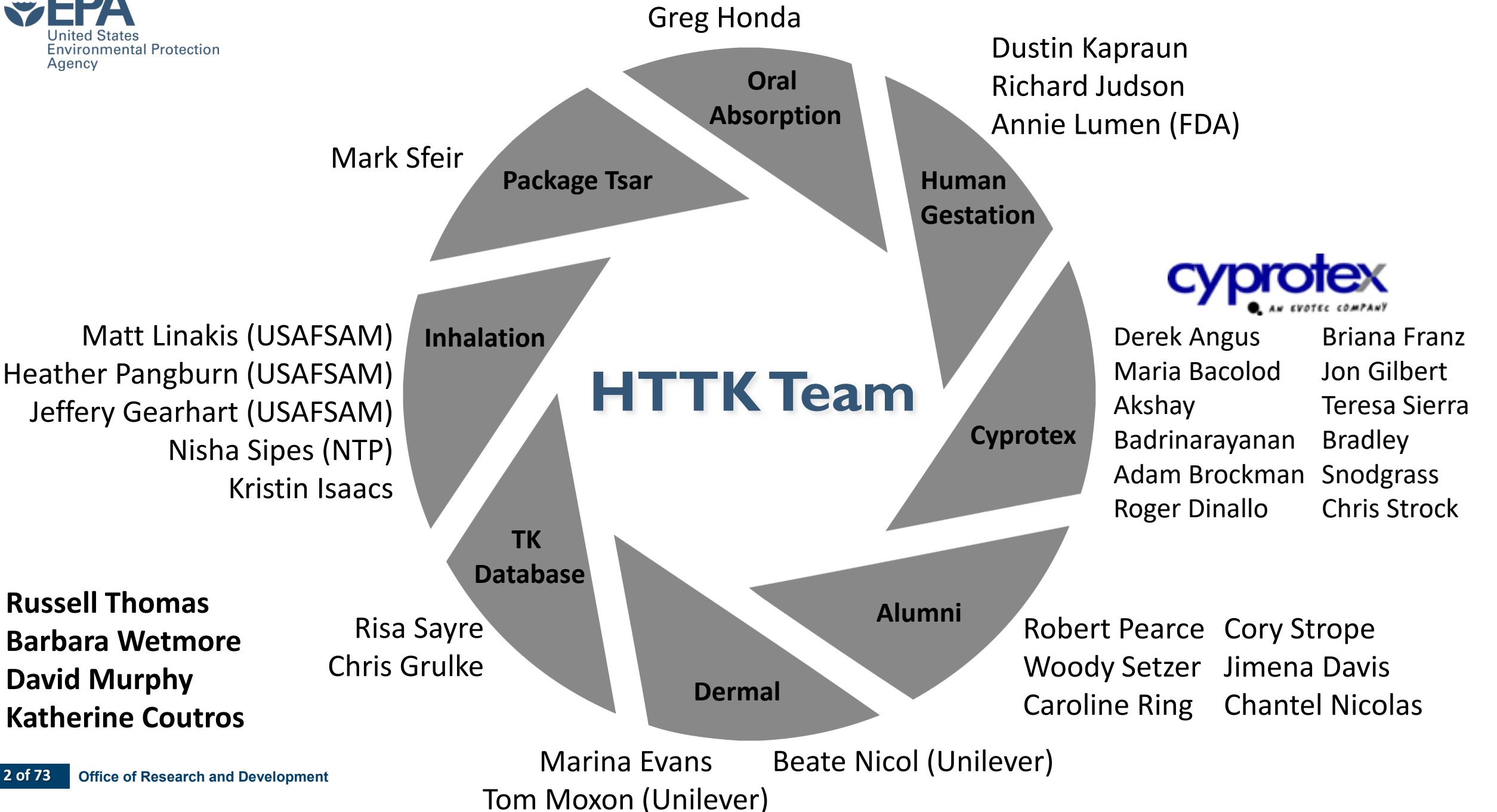
National Center for Computational Toxicology

Office of Research and Development

U.S. Environmental Protection Agency

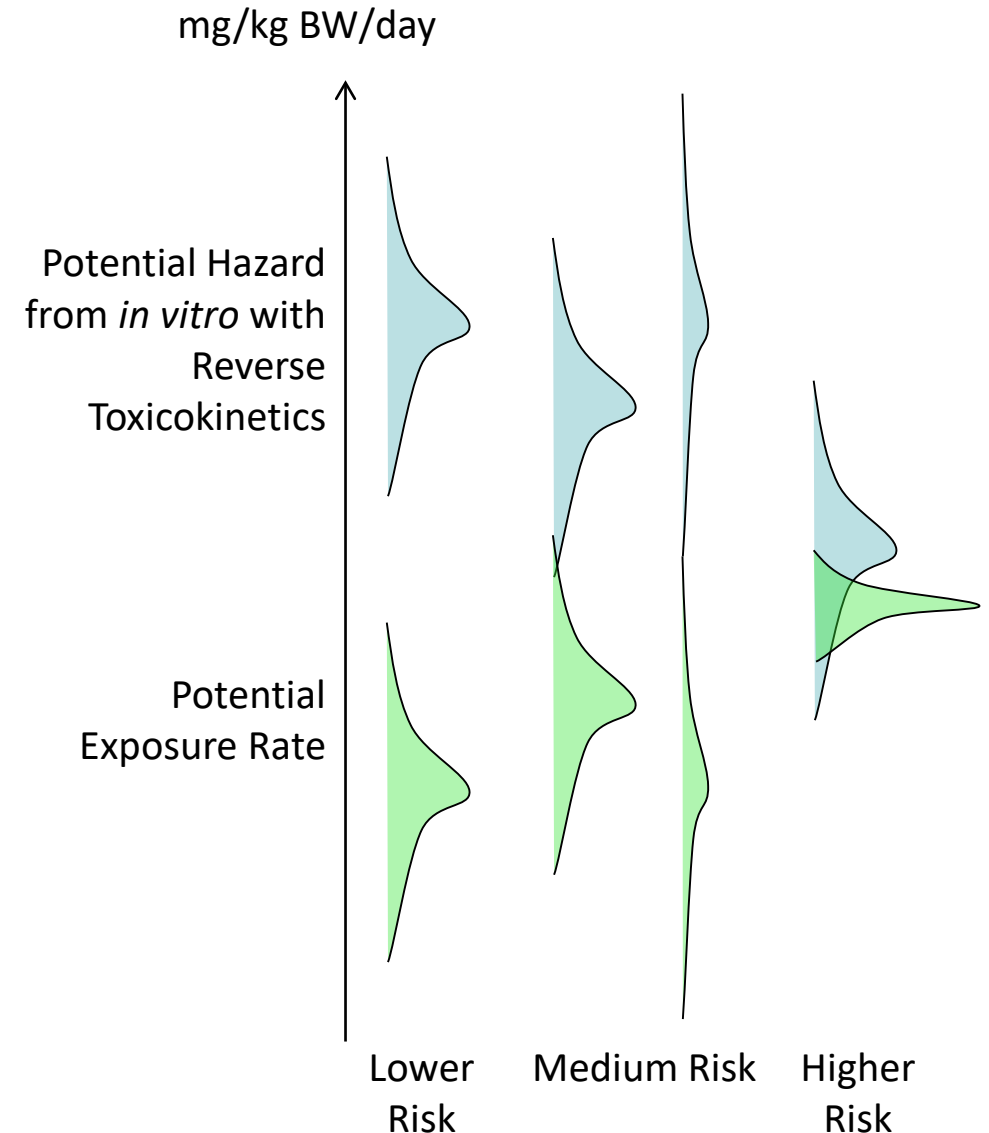
**Computational Toxicology
Community of Practice Webinar**

June 27, 2019

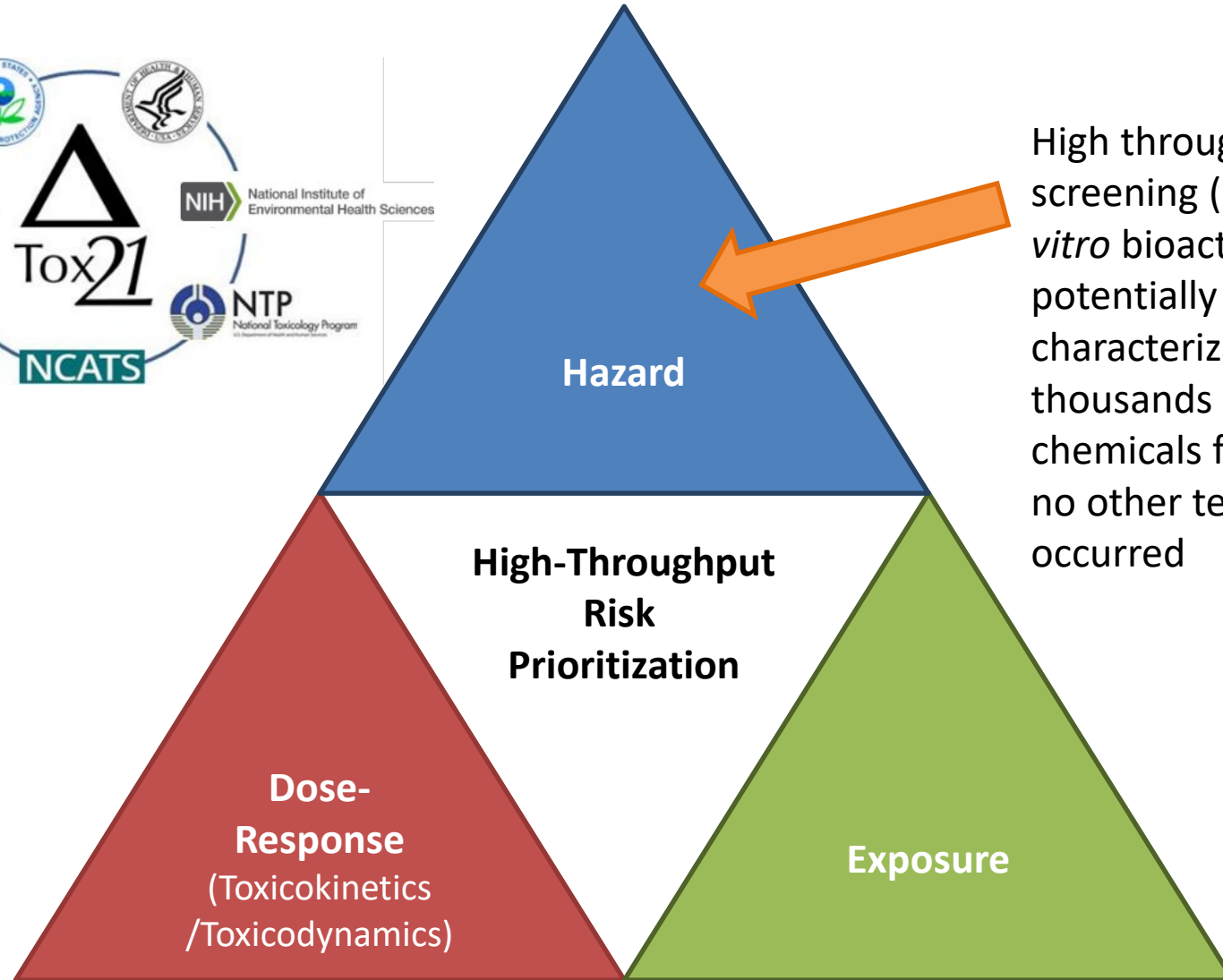


Chemical Risk = Hazard x Exposure

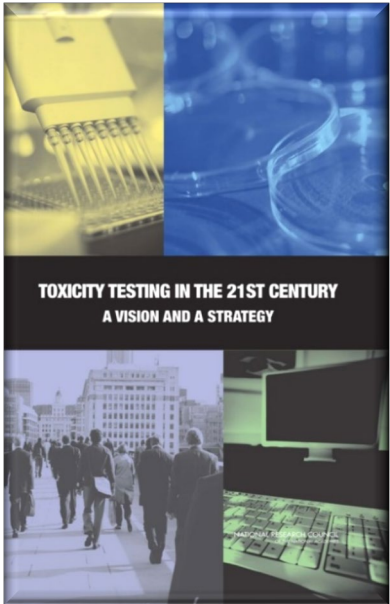
- The U.S. National Research Council (1983) identified chemical risk as a function of both inherent hazard and exposure
- To address the thousands of chemicals in commerce and the environment, we need new approach methodologies (NAMs) that can inform prioritization of chemicals most worthy of additional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

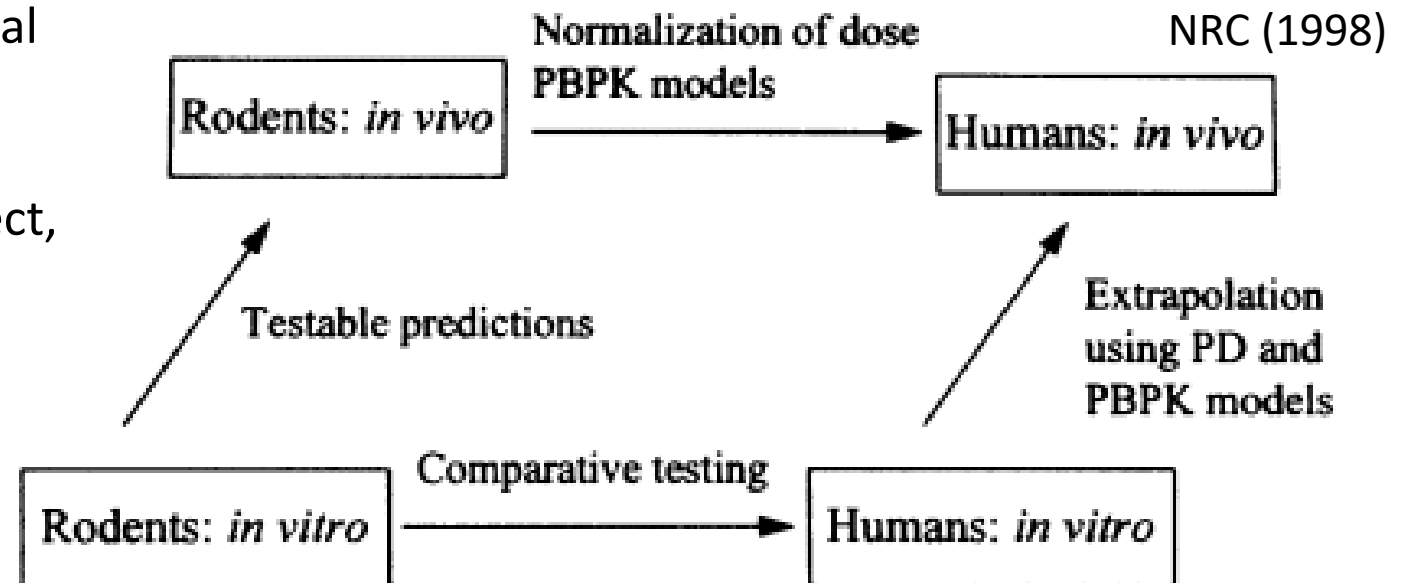
- IVIVE-PK/TK (**Pharmacokinetics/Toxicokinetics**):

- 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**):

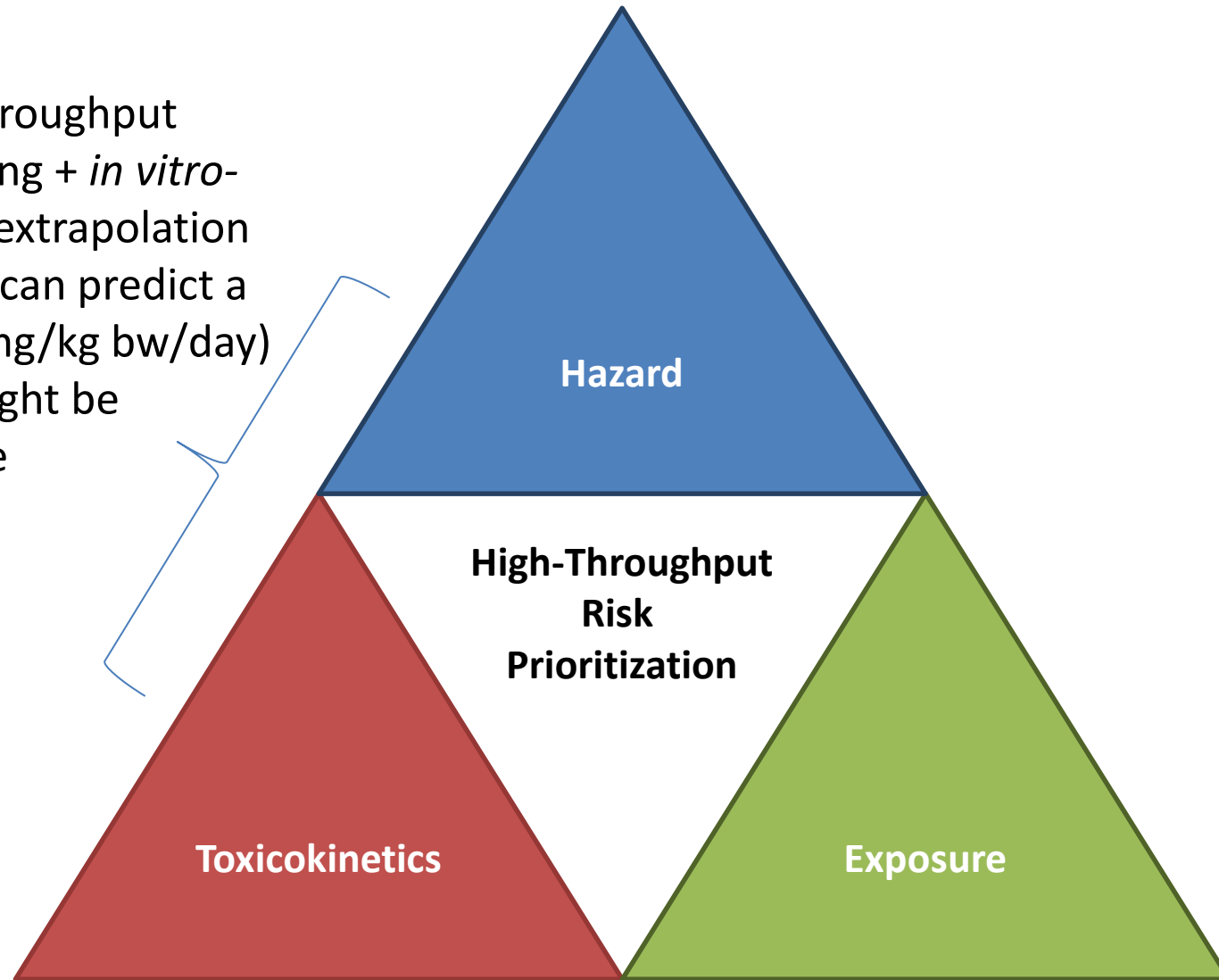
- Effect of molecules/chemicals at biological target *in vivo*
- Assay design/selection important
- Perturbation as adverse/therapeutic effect, reversible/ irreversible effects

- Both contribute to *in vivo* effect prediction



New Exposure Data and Models

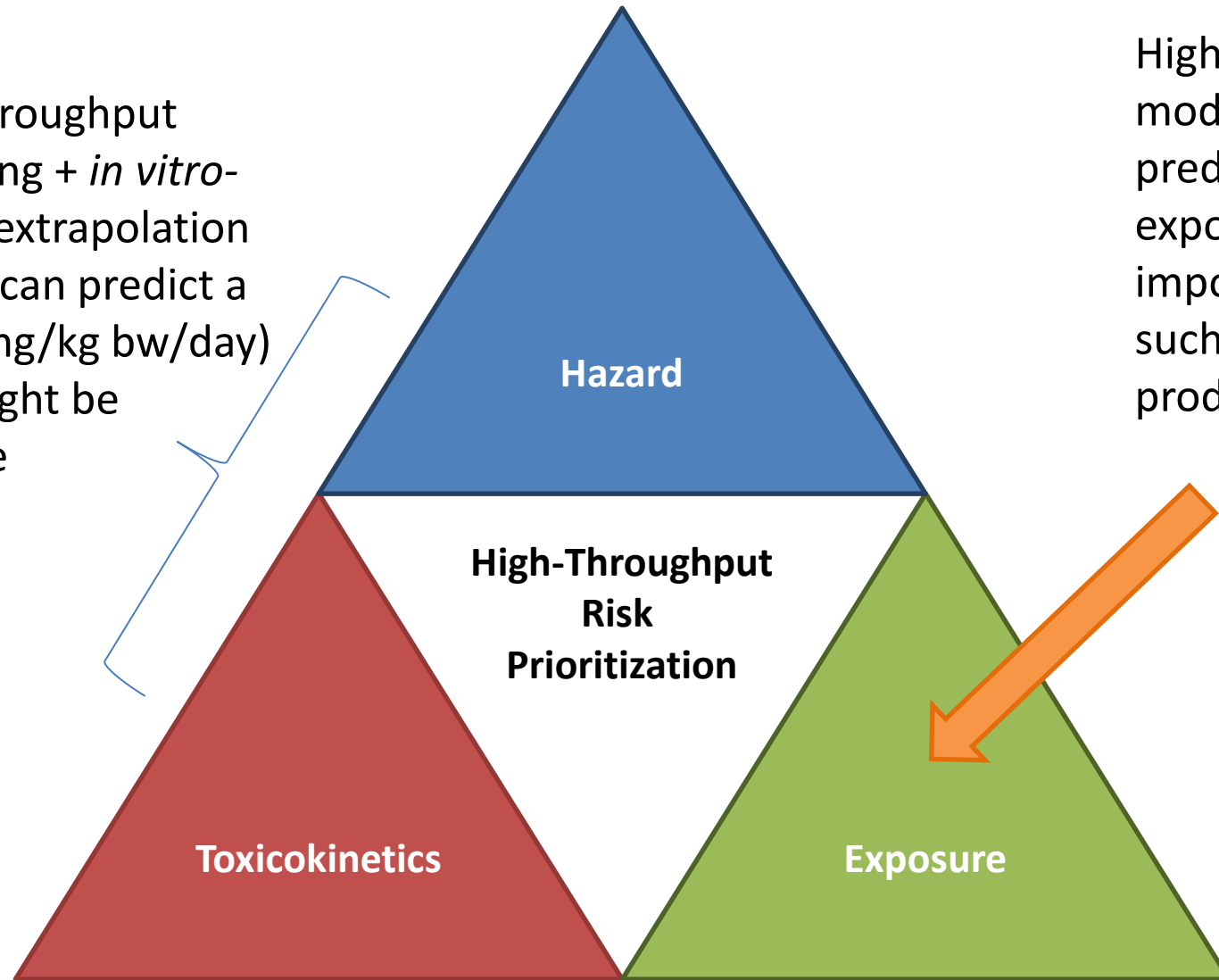
High throughput
screening + *in vitro*-
in vivo extrapolation
(IVIVE) can predict a
dose (mg/kg bw/day)
that might be
adverse



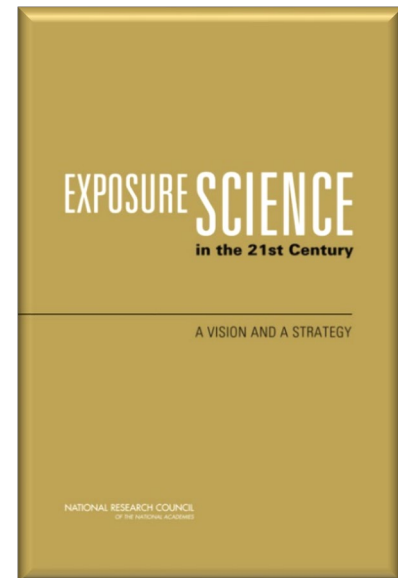
Wetmore et al. (2012, 2015)

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



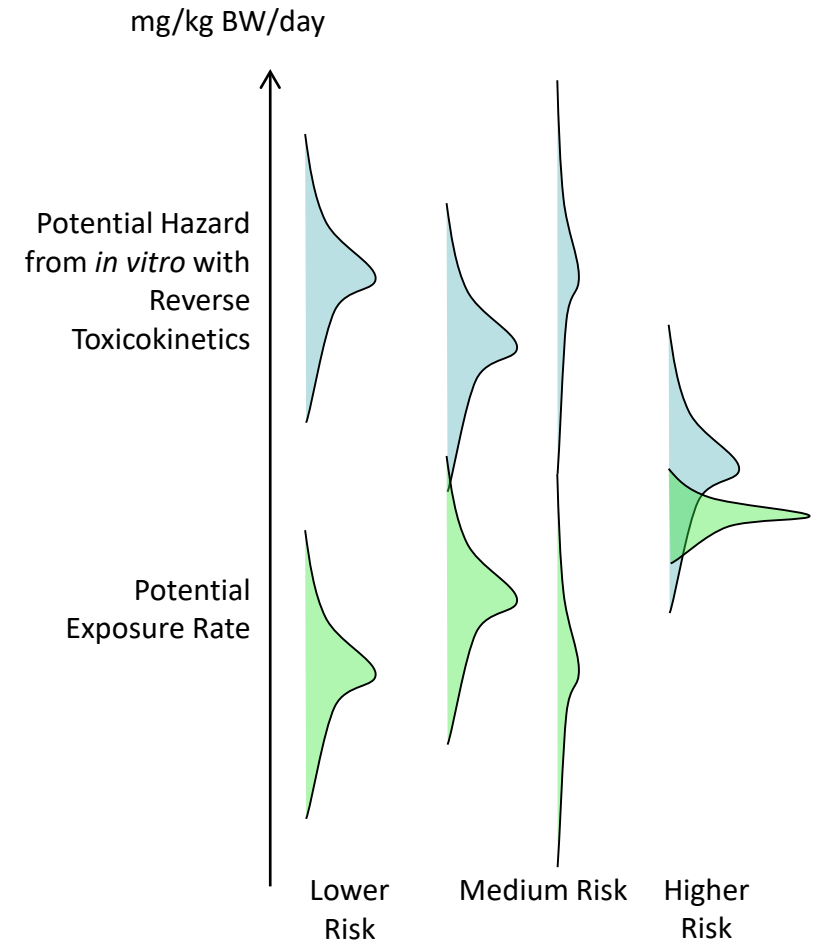
High throughput models exist to make predictions of exposure via specific, important pathways such as residential product use and diet



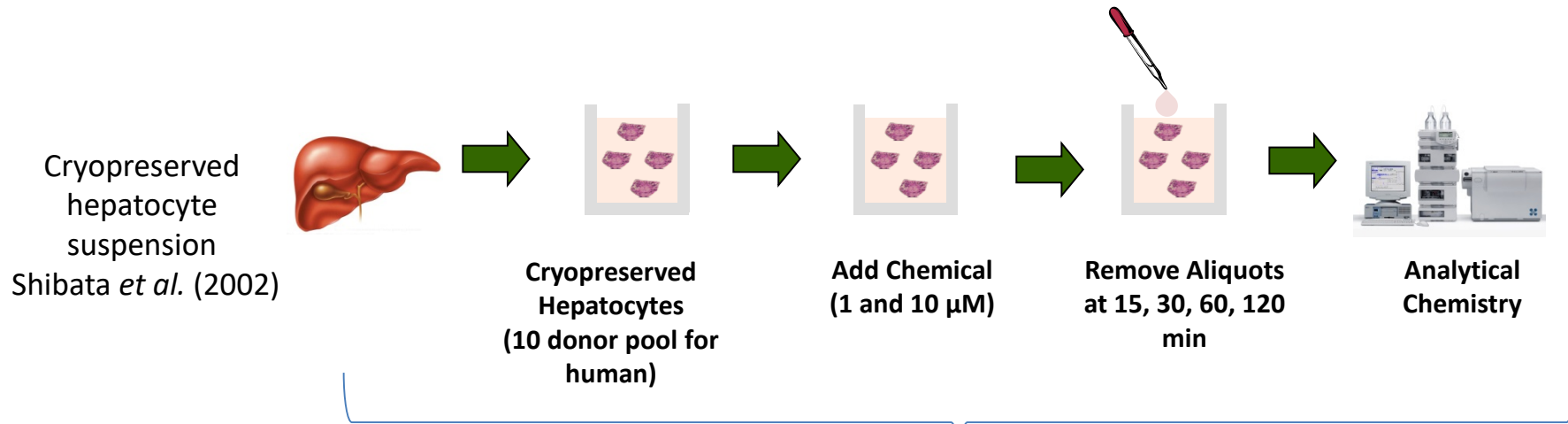
NRC (2012)

High Throughput Toxicokinetics (HTTK)

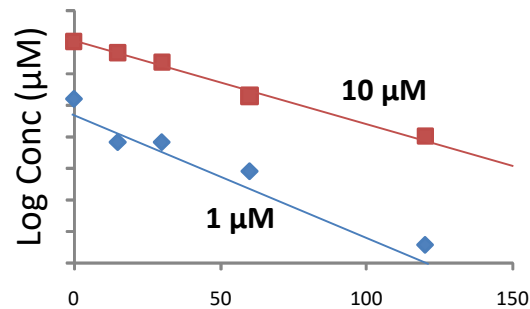
- **Most chemicals do not have TK data**
- In order to address greater numbers of chemicals we collect *in vitro*, high throughput toxicokinetic (HTTK) data (Rotroff et al., 2010, Wetmore et al., 2012, 2015)
- HTTK methods have been used by the pharmaceutical industry to determine range of efficacious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

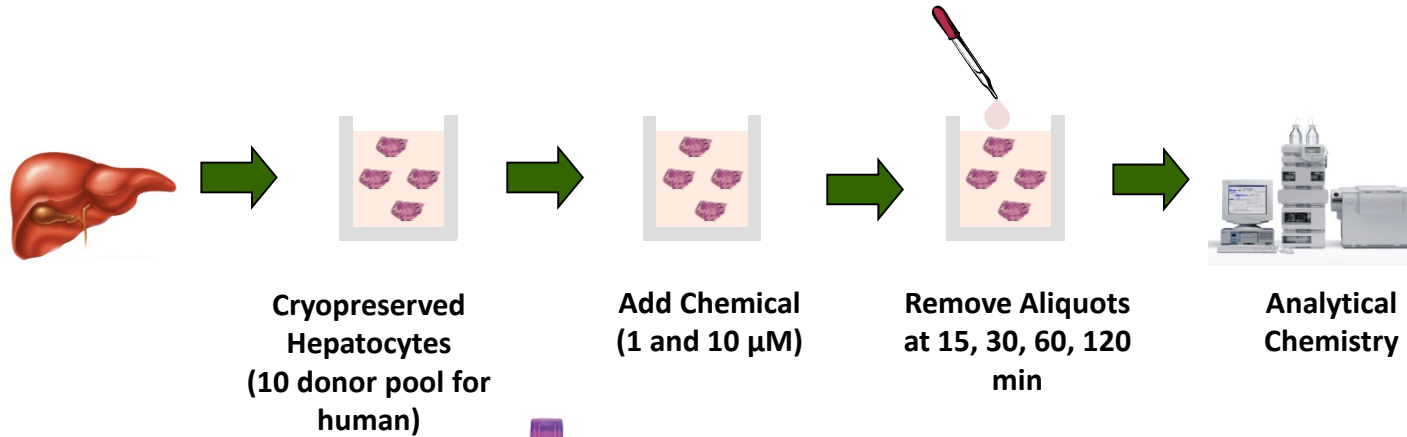


We perform the assay at 1 and 10 μM to check for saturation of metabolizing enzymes.

- **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)

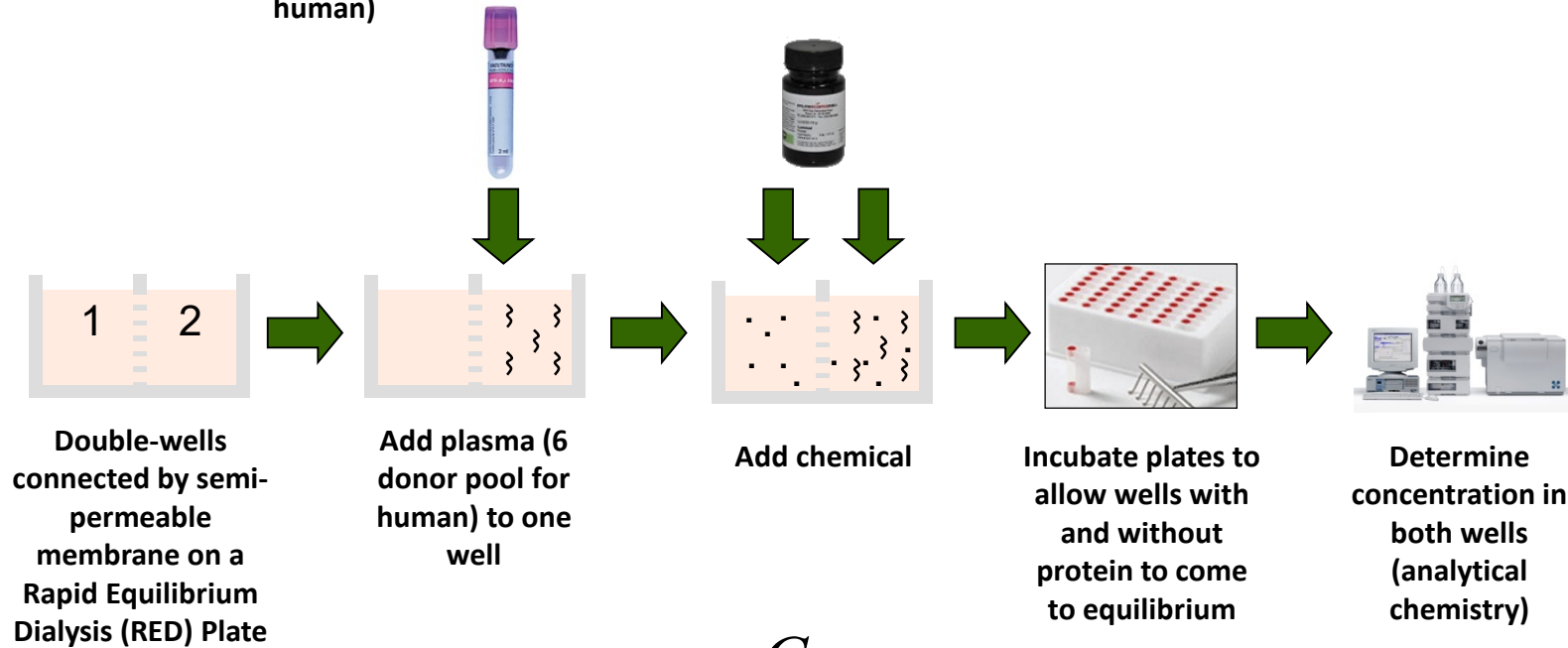
In Vitro Data for HTTK

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



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

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

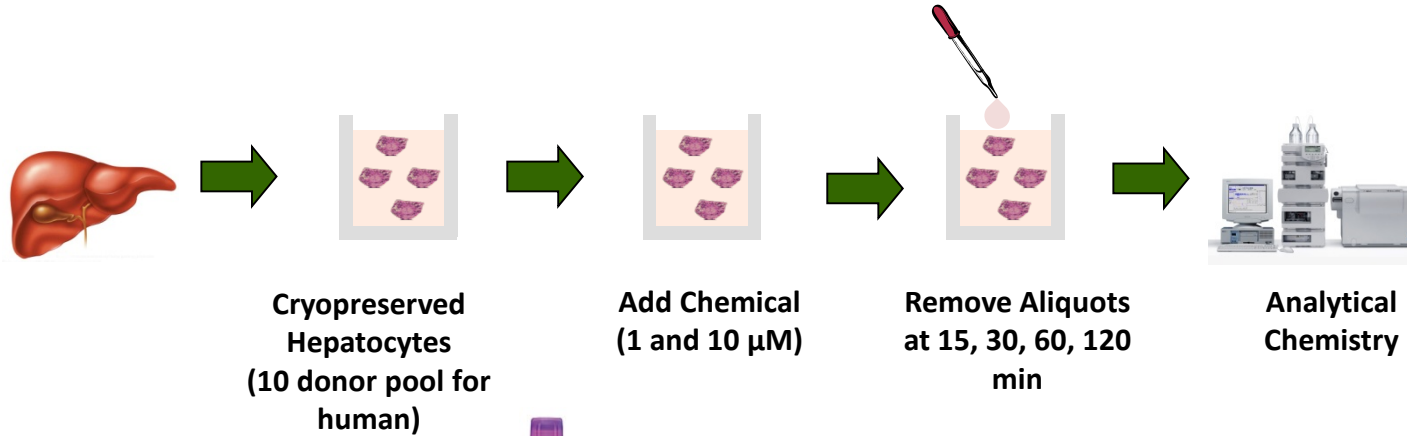


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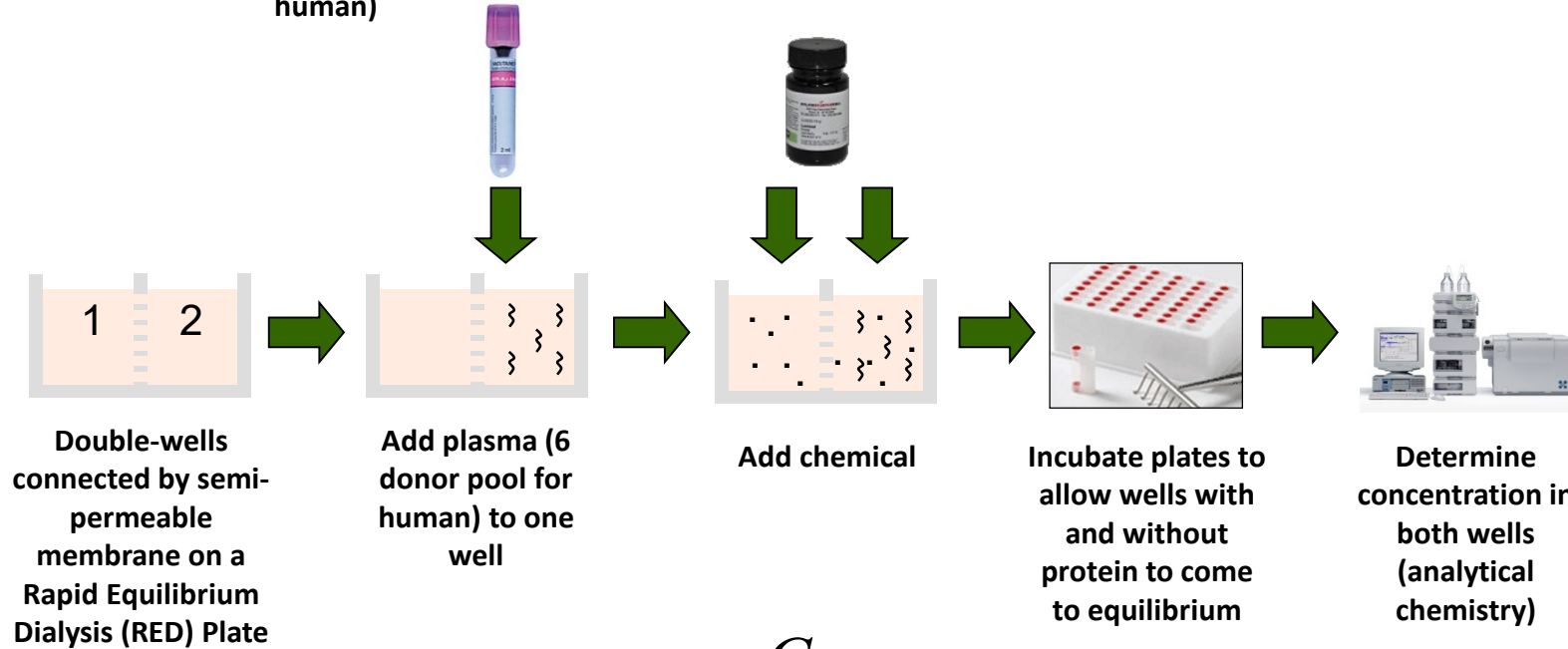
$$F_{ub,p} = \frac{C_{well1}}{C_{well2}}$$

In Vitro Data for HTTK

Cryopreserved
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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)

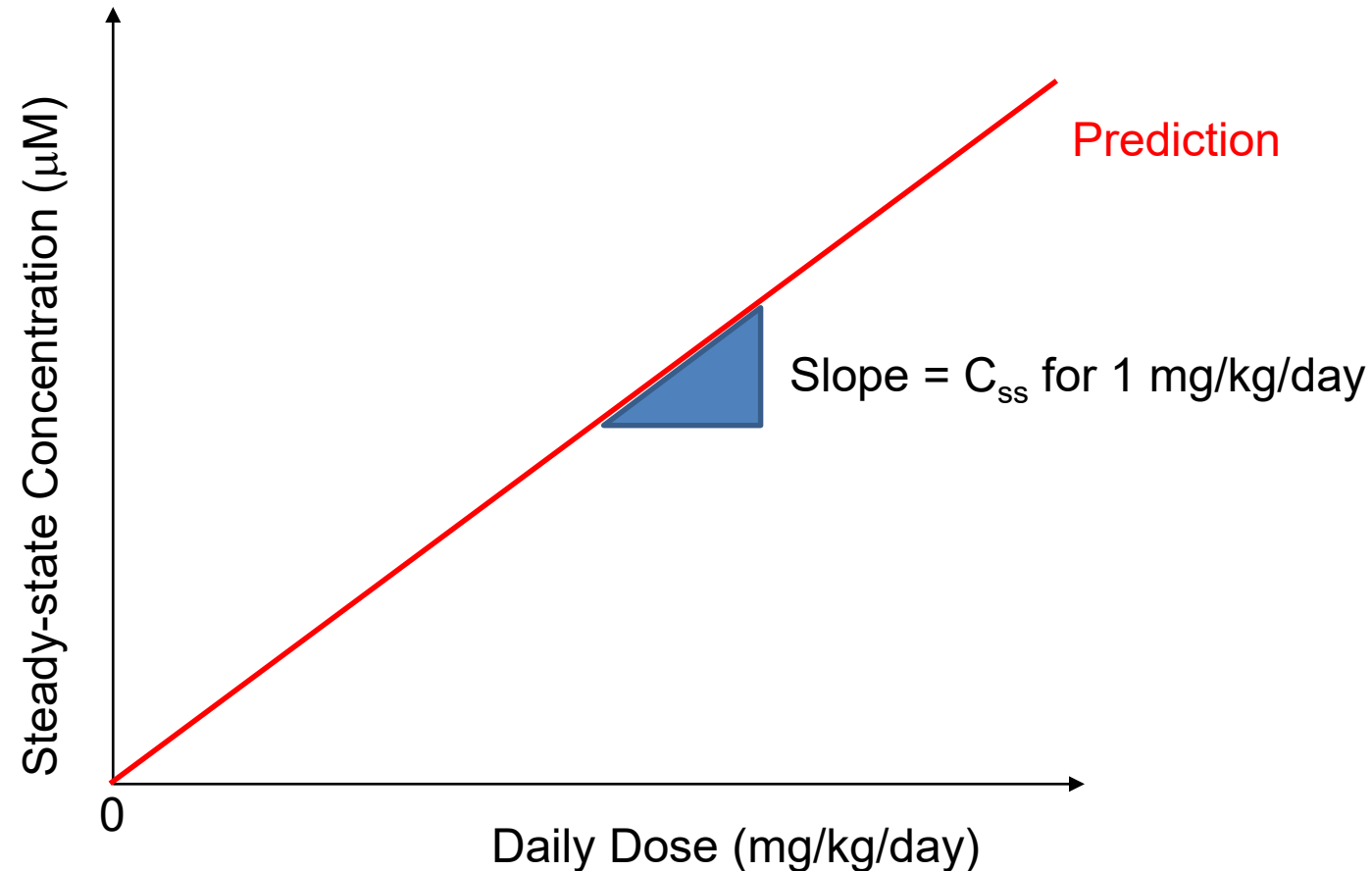
Hepatic Metabolism
(Cl_{int} : Scaled hepatic clearance
 Q_l : Blood flow to liver)

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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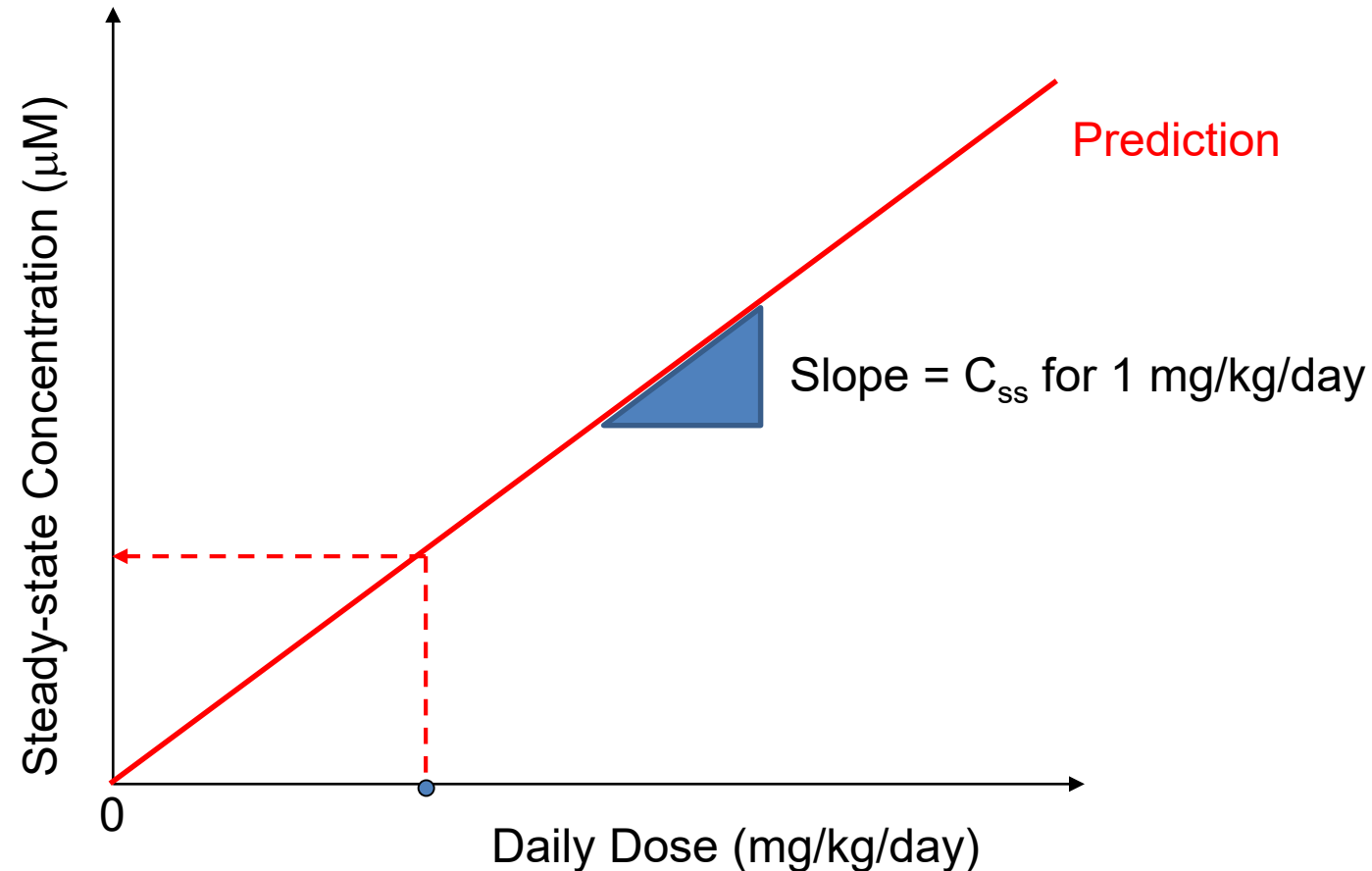


- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

Wetmore *et al.* (2012)

Assume that Steady-State is Linear with Dose

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

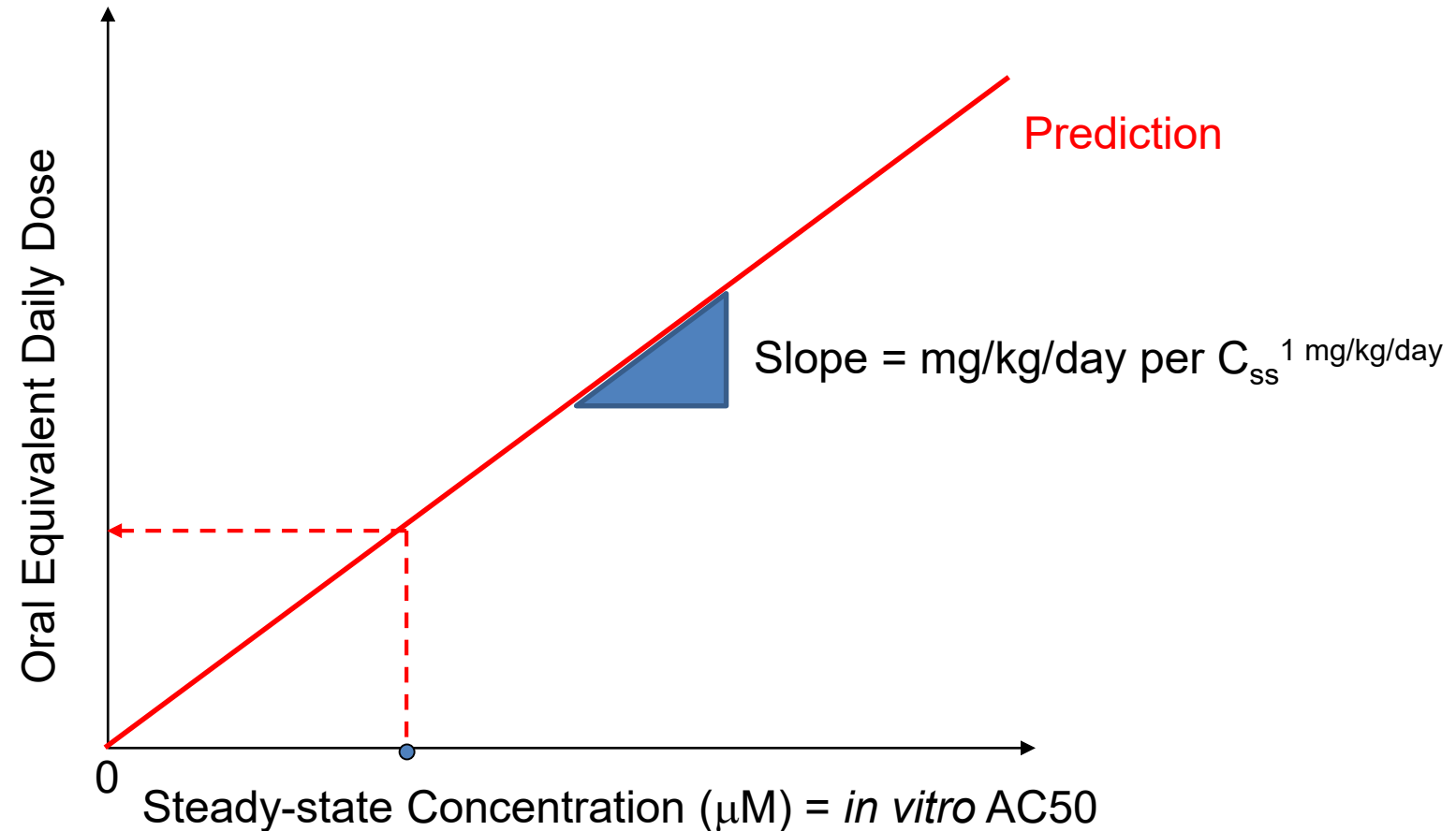


- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

Wetmore *et al.* (2012)

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

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

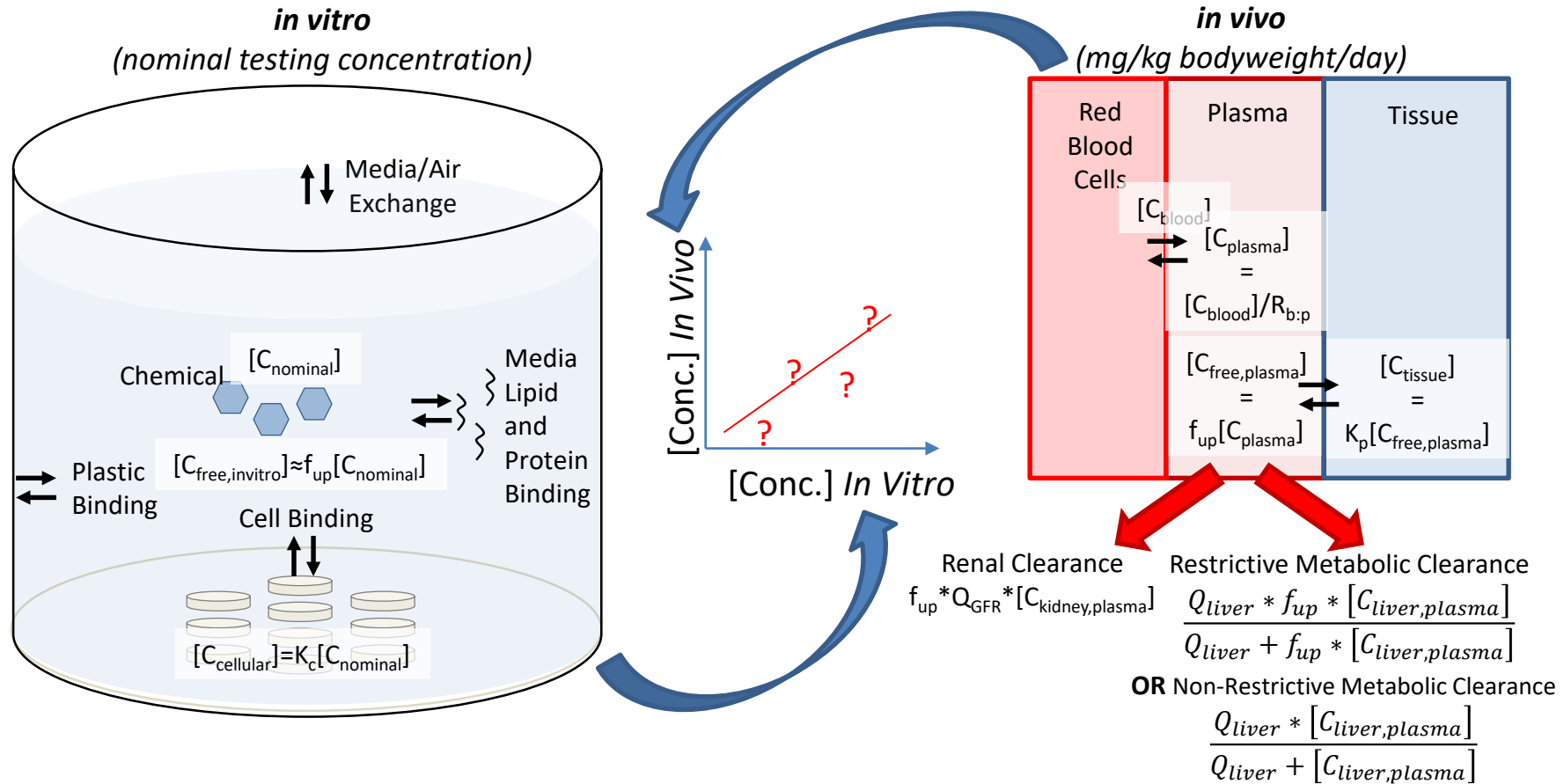


- Can calculate predicted steady-state concentration (C_{ss}) for a 1 mg/kg/day dose and multiply to get concentrations for other doses

Wetmore *et al.* (2012)

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

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



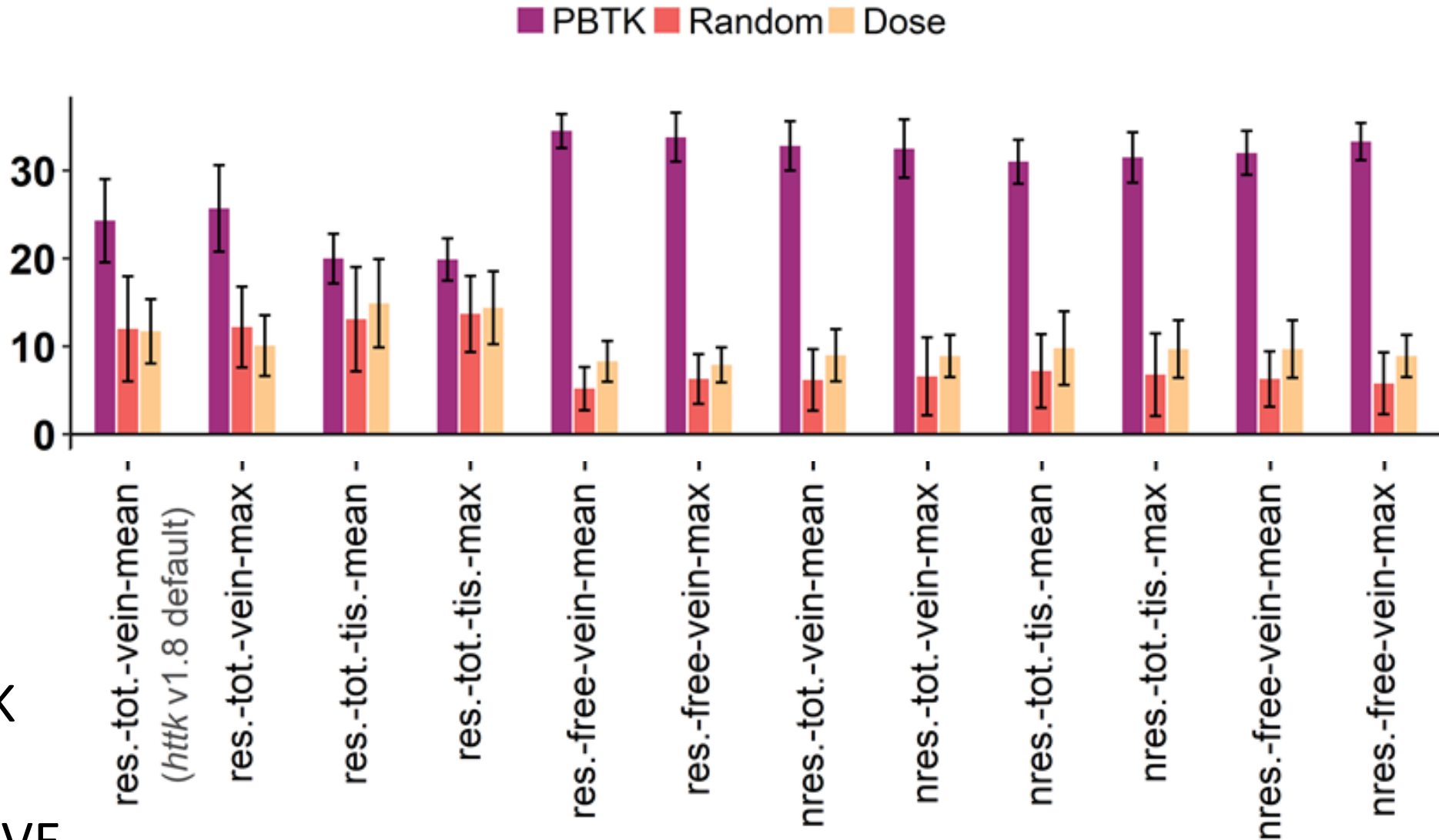
Selecting the appropriate *in vitro* and *in vivo* concentrations for extrapolation

Optimizing HTK-based IVIVE

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

Using PBTK
Models

Improves IVIVE

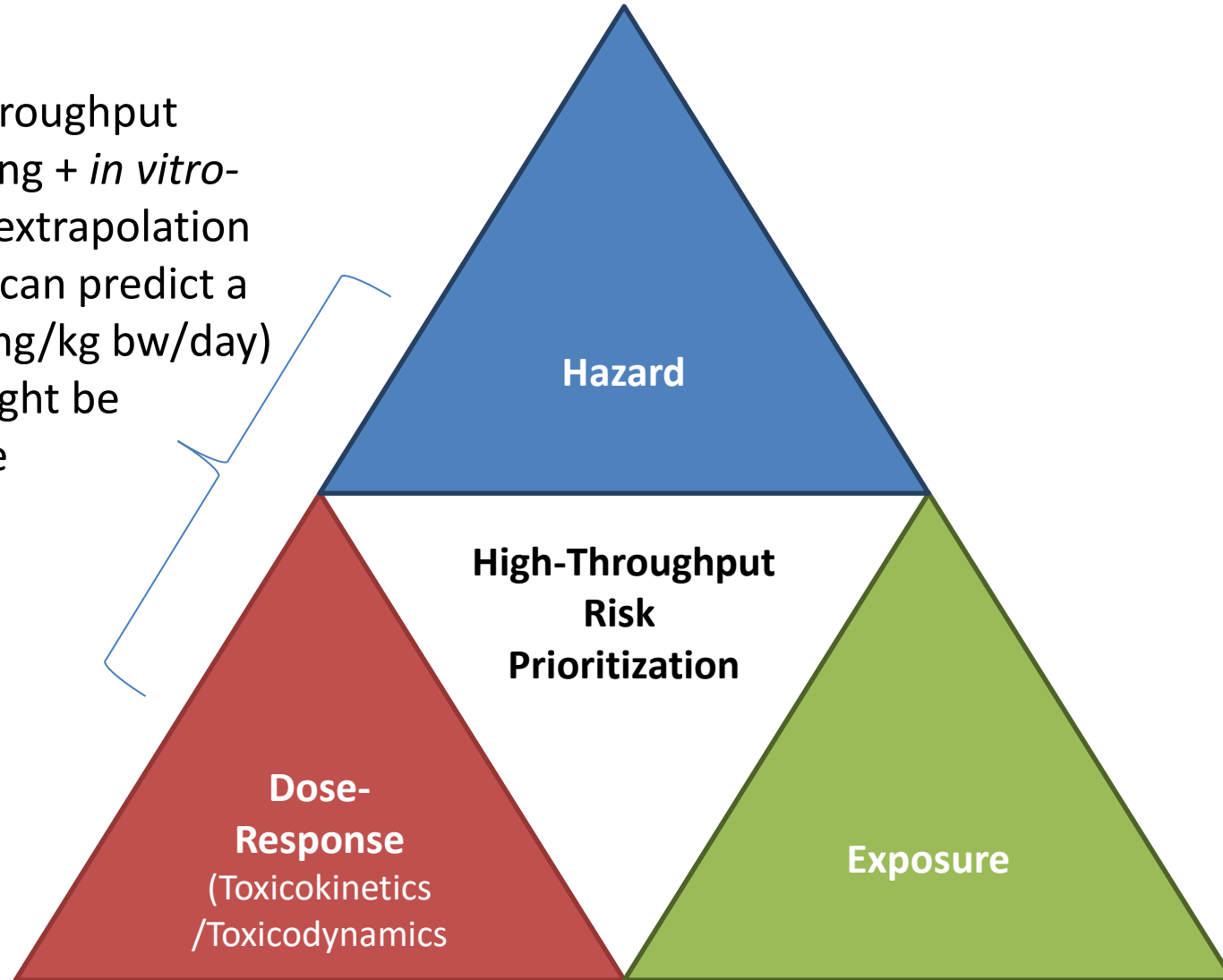


Various Combinations of IVIVE Assumptions

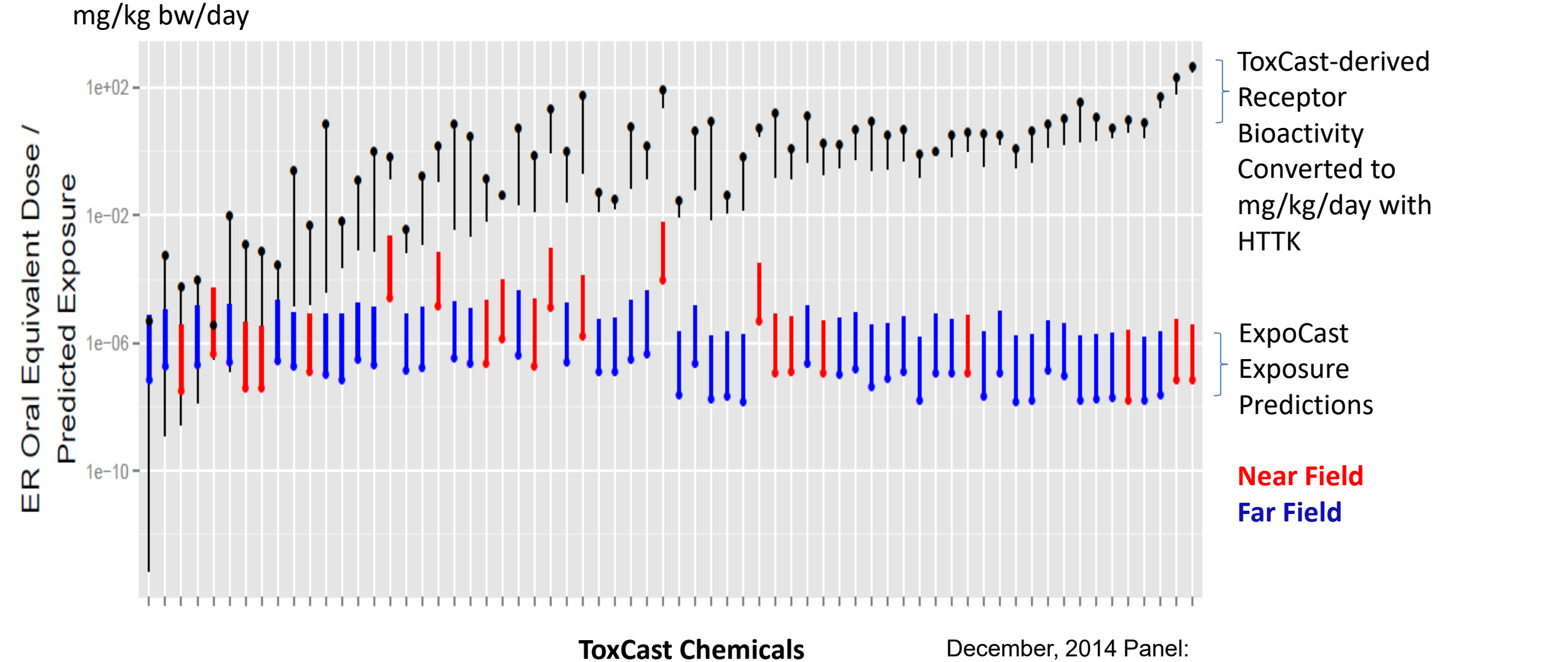
Honda et al. (2019)

New Exposure Data and Models

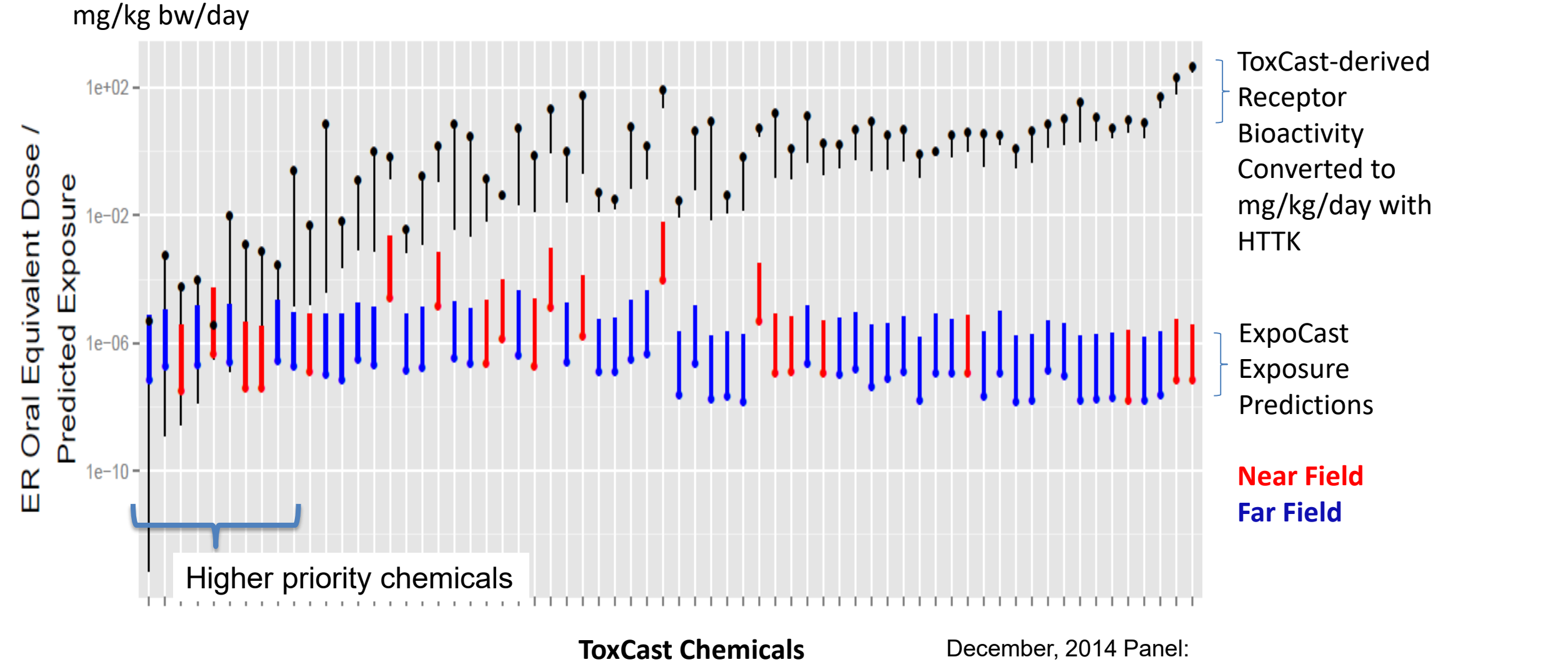
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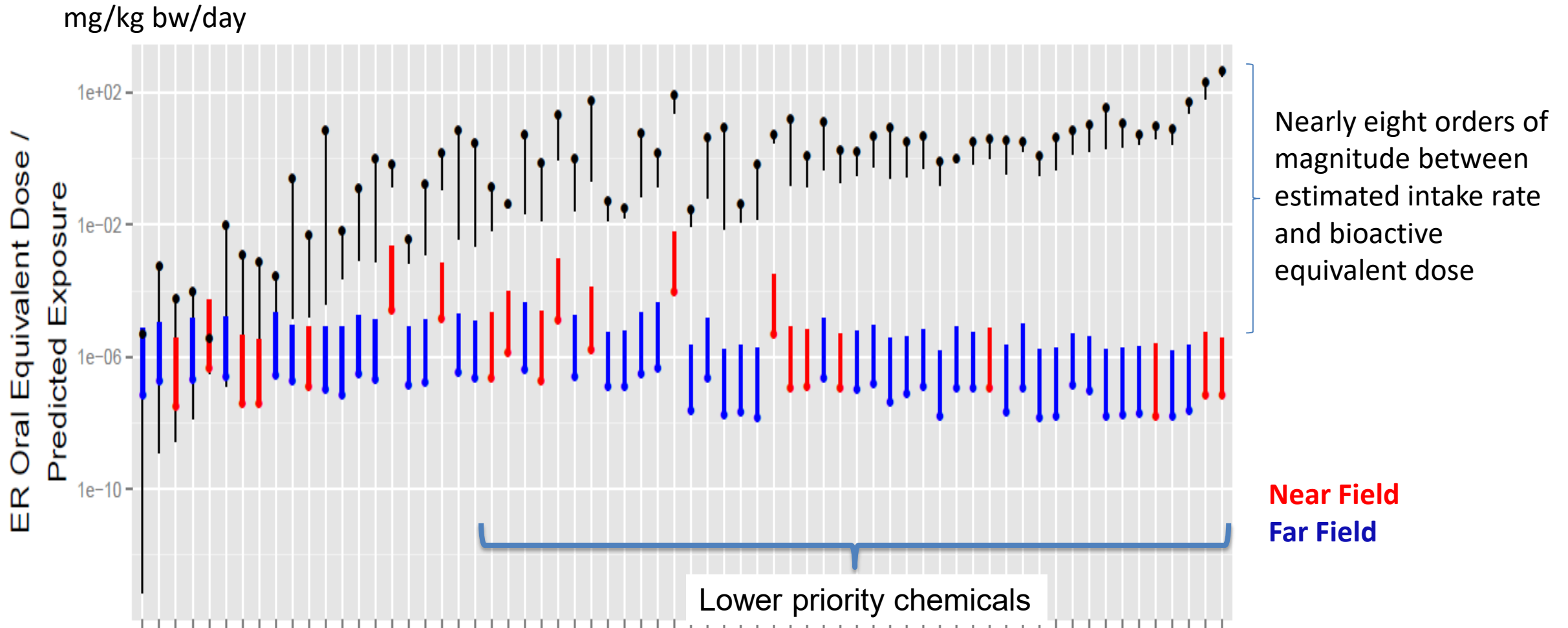
High Throughput Risk Prioritization in Practice



High Throughput Risk Prioritization in Practice



High Throughput Risk Prioritization in Practice

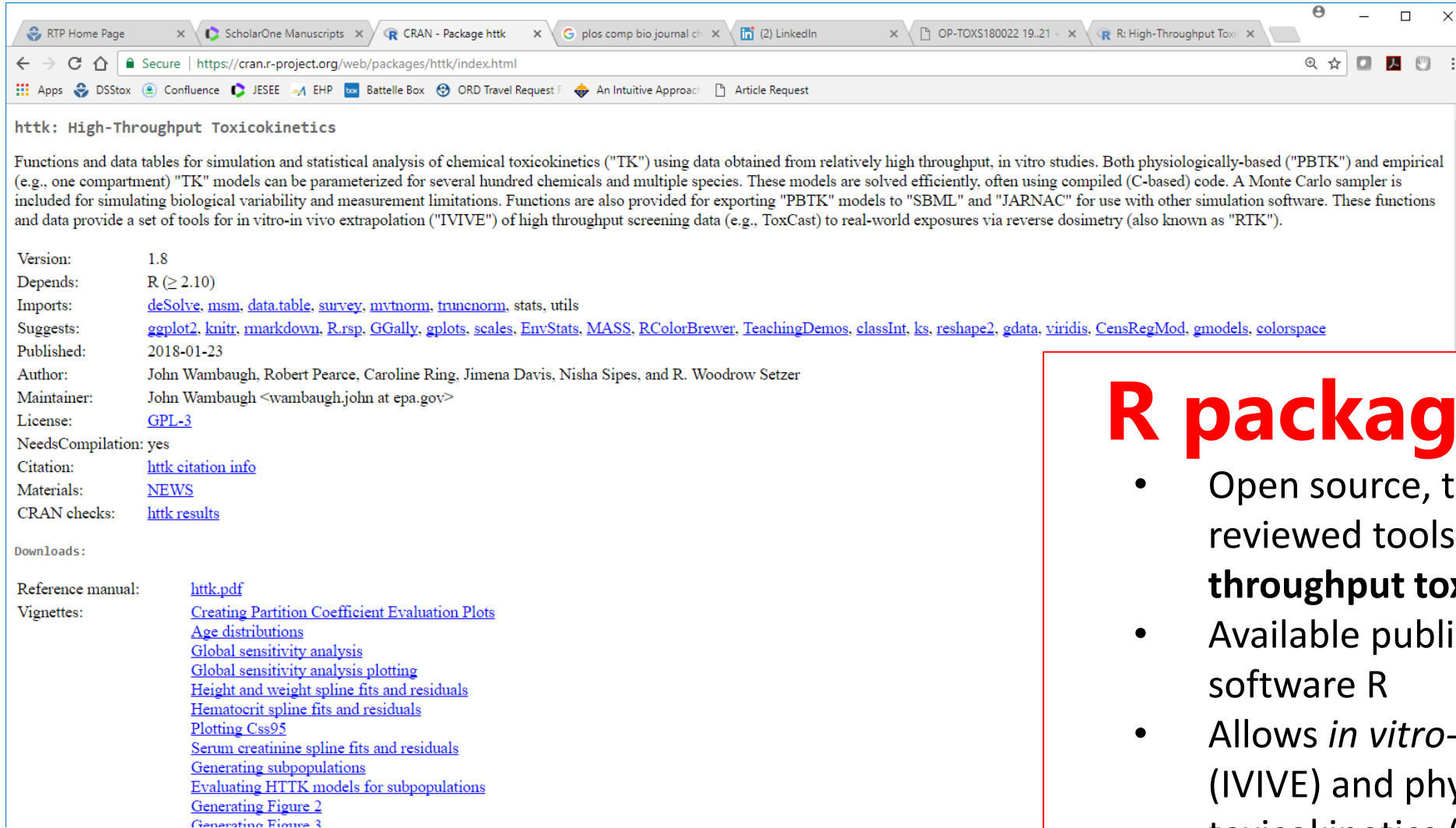


ToxCast Chemicals

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

Open Source Tools and Data for HTTK

<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', '(2) LinkedIn', 'OP-TOXS180022 19_21', and 'R: High-Throughput Toxicokinetics'. 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 (≥ 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 HTTK models for subpopulations](#), [Generating Figure 2](#), [Generating Figure 3](#)

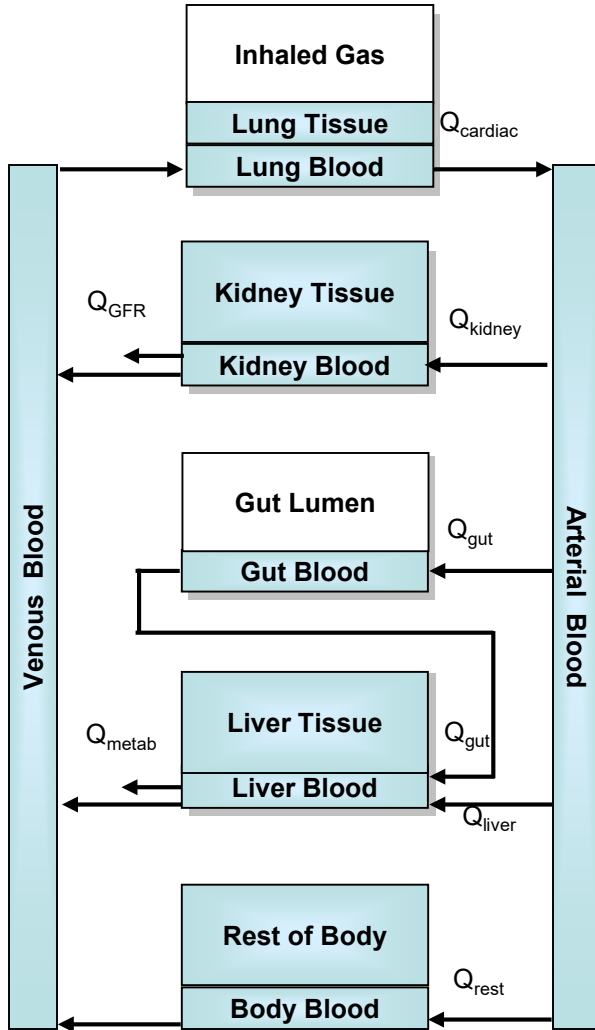
R package "httk"

- 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)
```

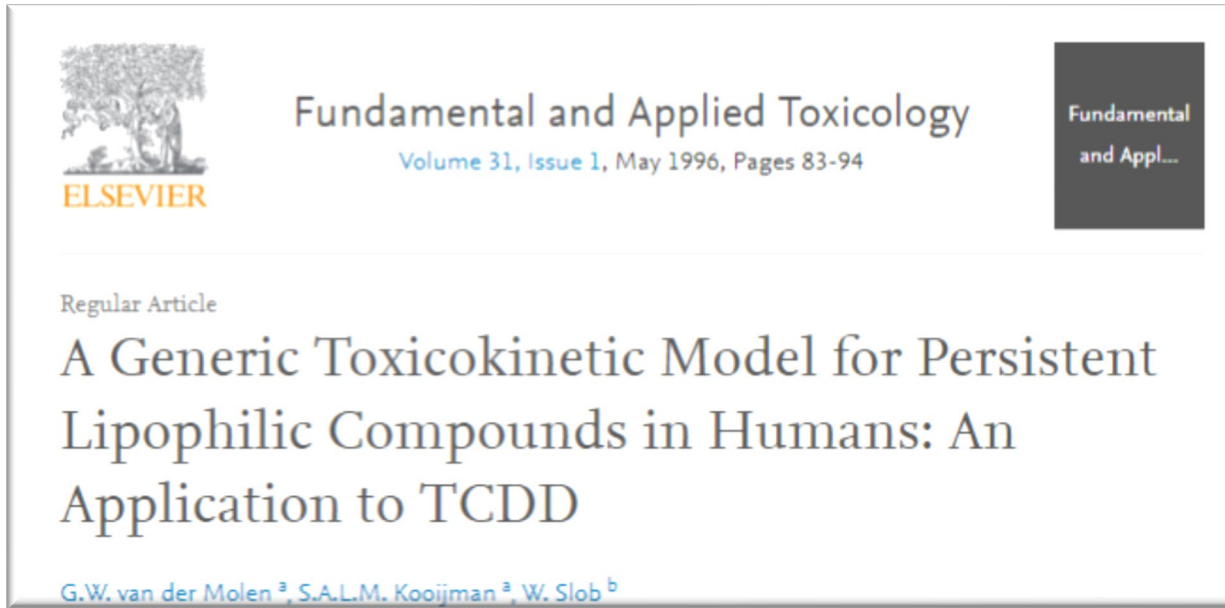
```
#Steady-state concentration (uM) for 1 mg/kg/day for 0.5 quantile for rat for Acetochlor (calculated value):  
calc_mc_css(chem.cas="34256-82-1",species="Rat",which.quantile=0.5)
```

```
#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)
```

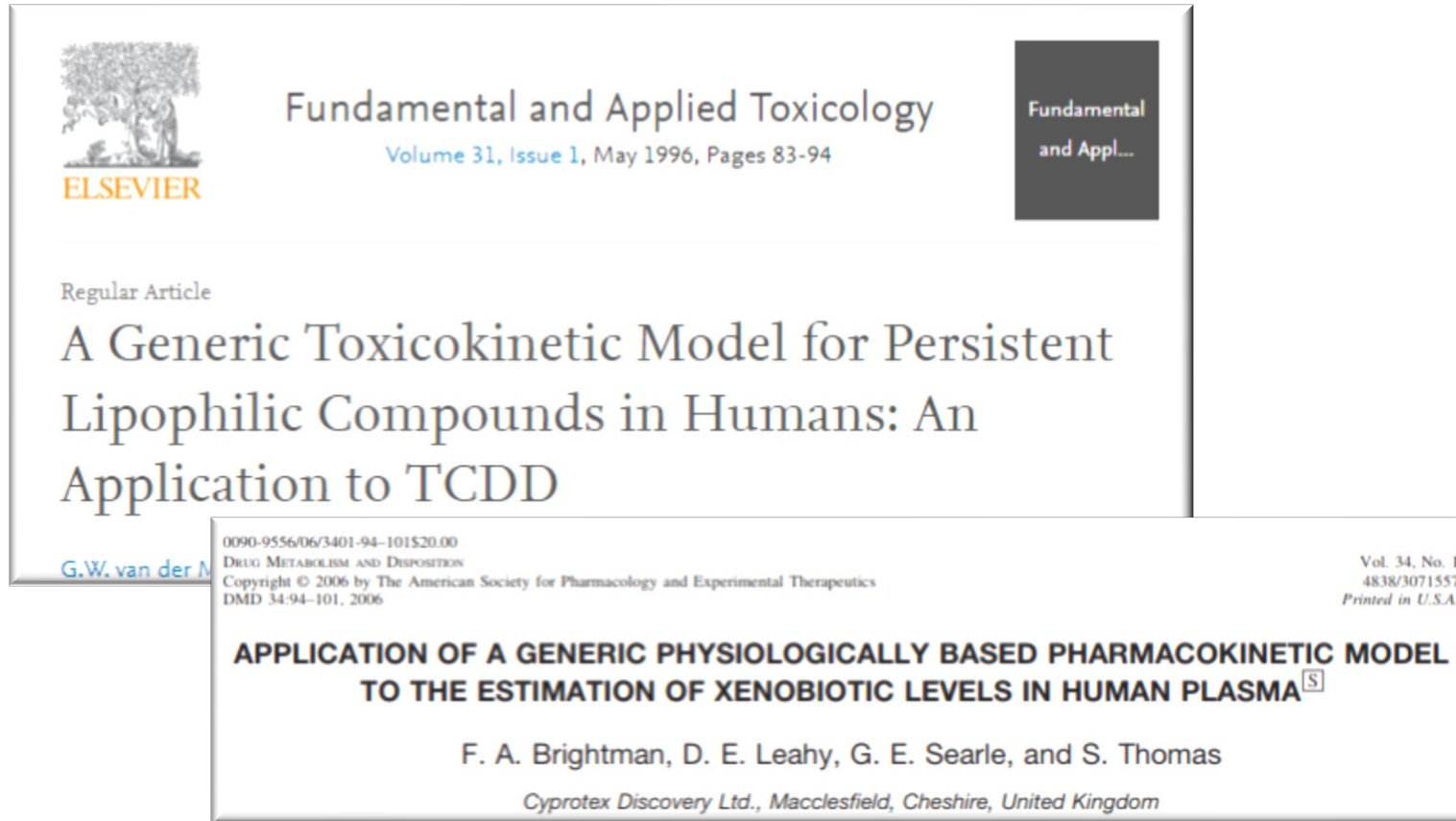
Generic PBTK Models

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




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ELSEVIER

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

Regular Article

A Generic Toxicokinetic Model for 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
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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

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Cyprotex Discovery Ltd., Macclesfield, Cheshire, United Kingdom

[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](#)

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on behalf of the British Occupational Hygiene Society
doi:10.1093/annhyg/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^{1*} and WIL F. TEN BERGE²

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

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

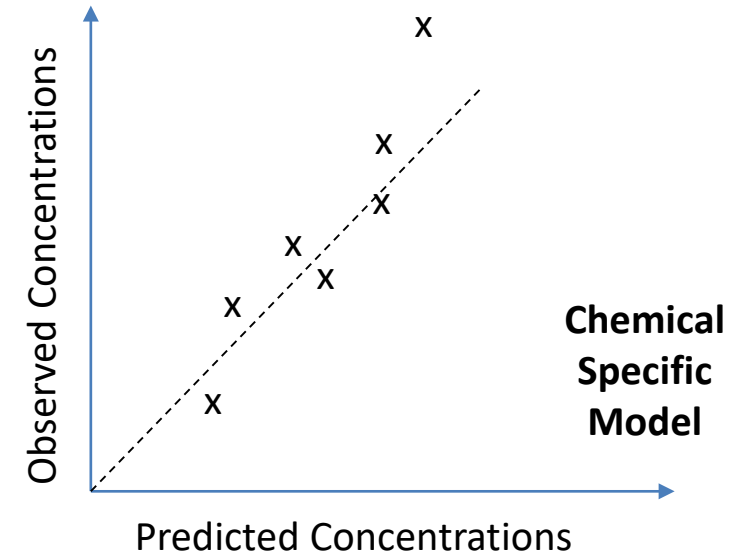
“...the default state of new and modernized Government information resources shall be open and machine readable.”

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

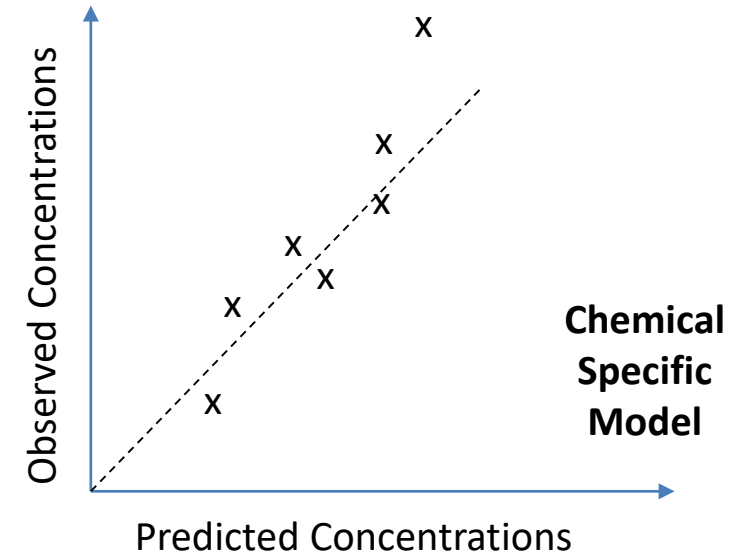
Building Confidence in TK Models

- 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



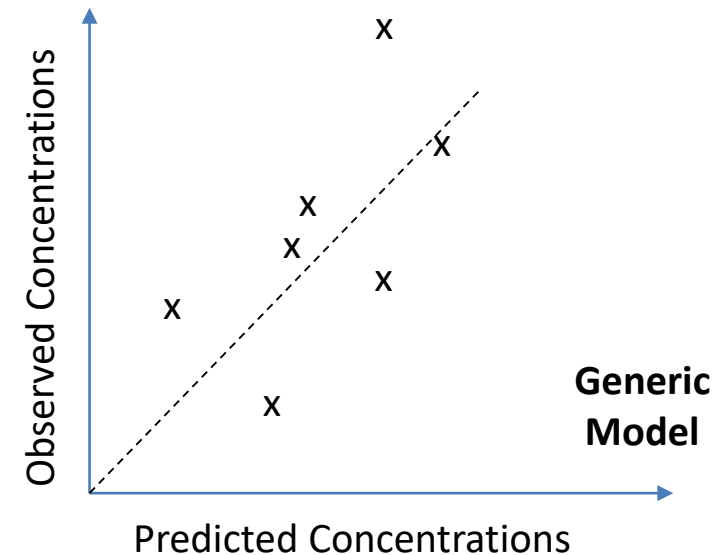
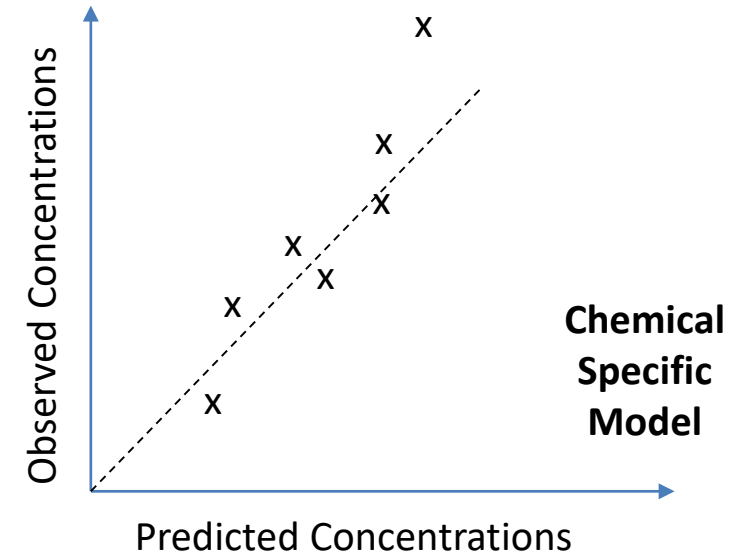
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- However, we do not typically have TK data



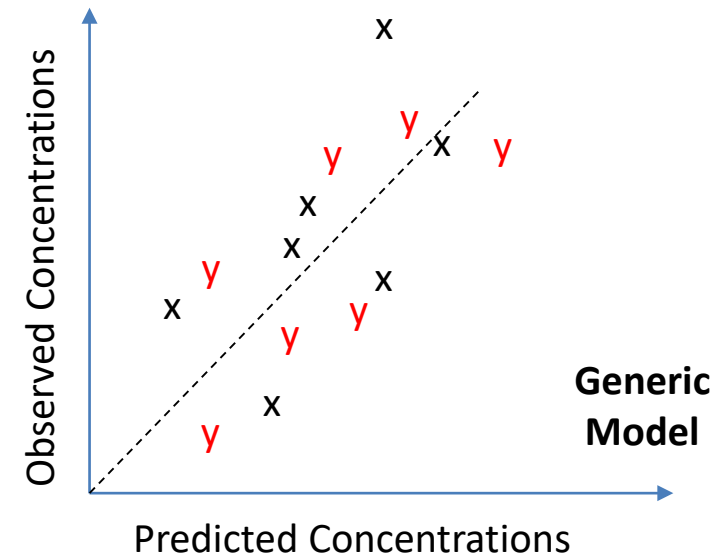
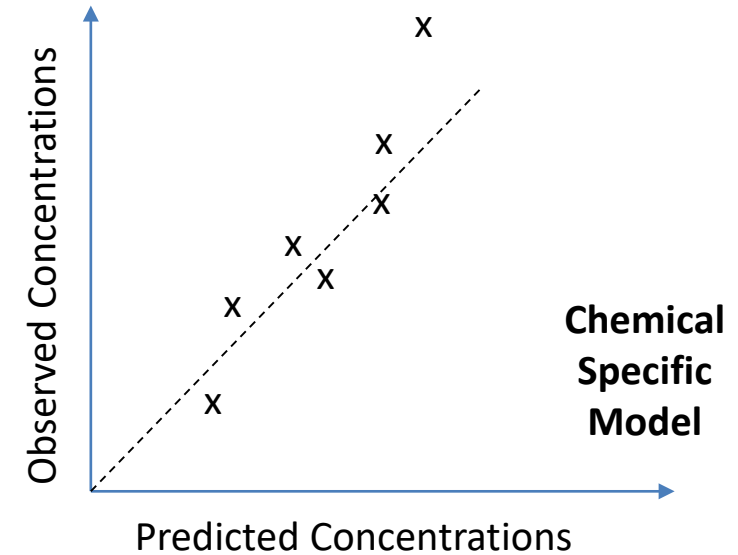
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- We can parameterize a **generic TK model**, and evaluate that model for as many chemicals as we do have data
 - We do expect larger uncertainty, but also greater confidence in model implementation
 - Estimate bias and uncertainty, and try to correlate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



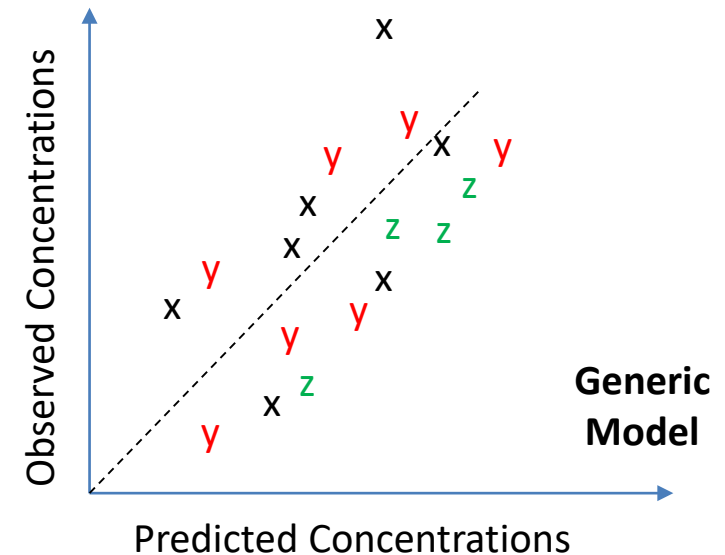
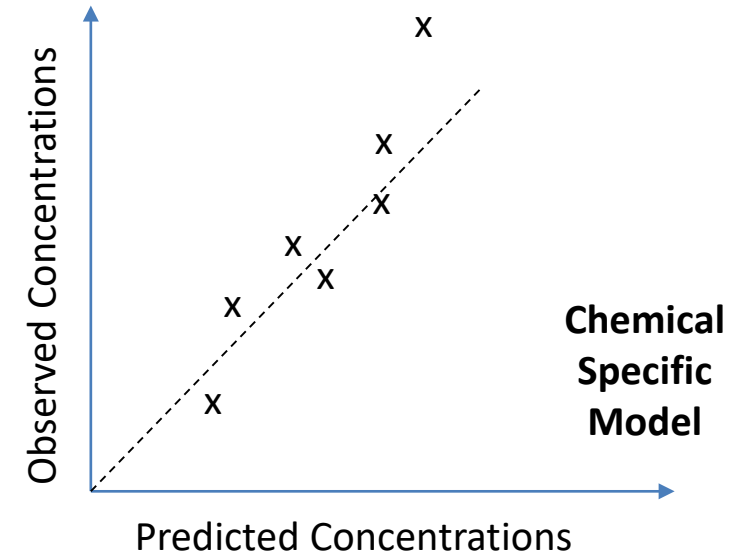
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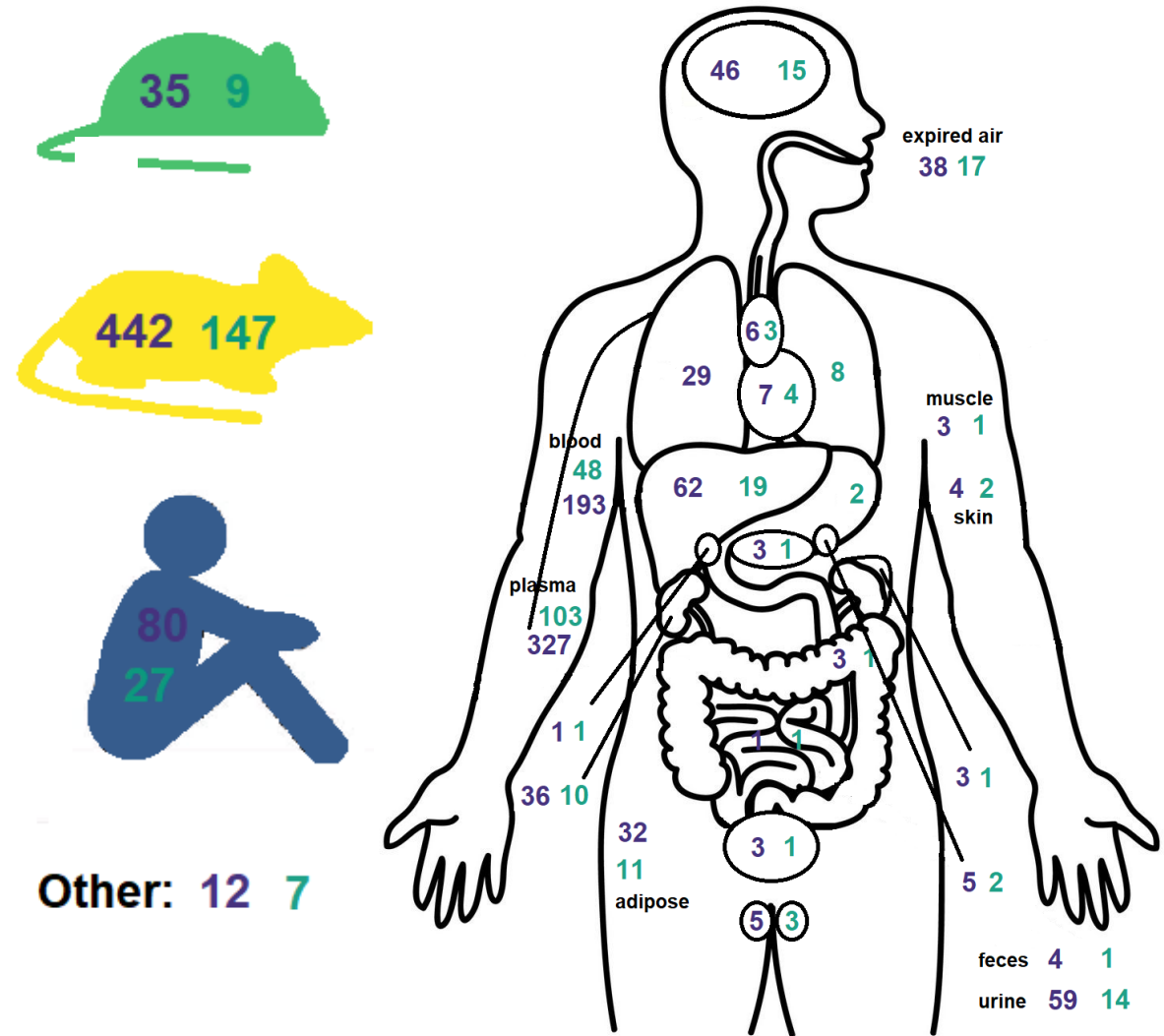
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In Vivo TK Database

- 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:

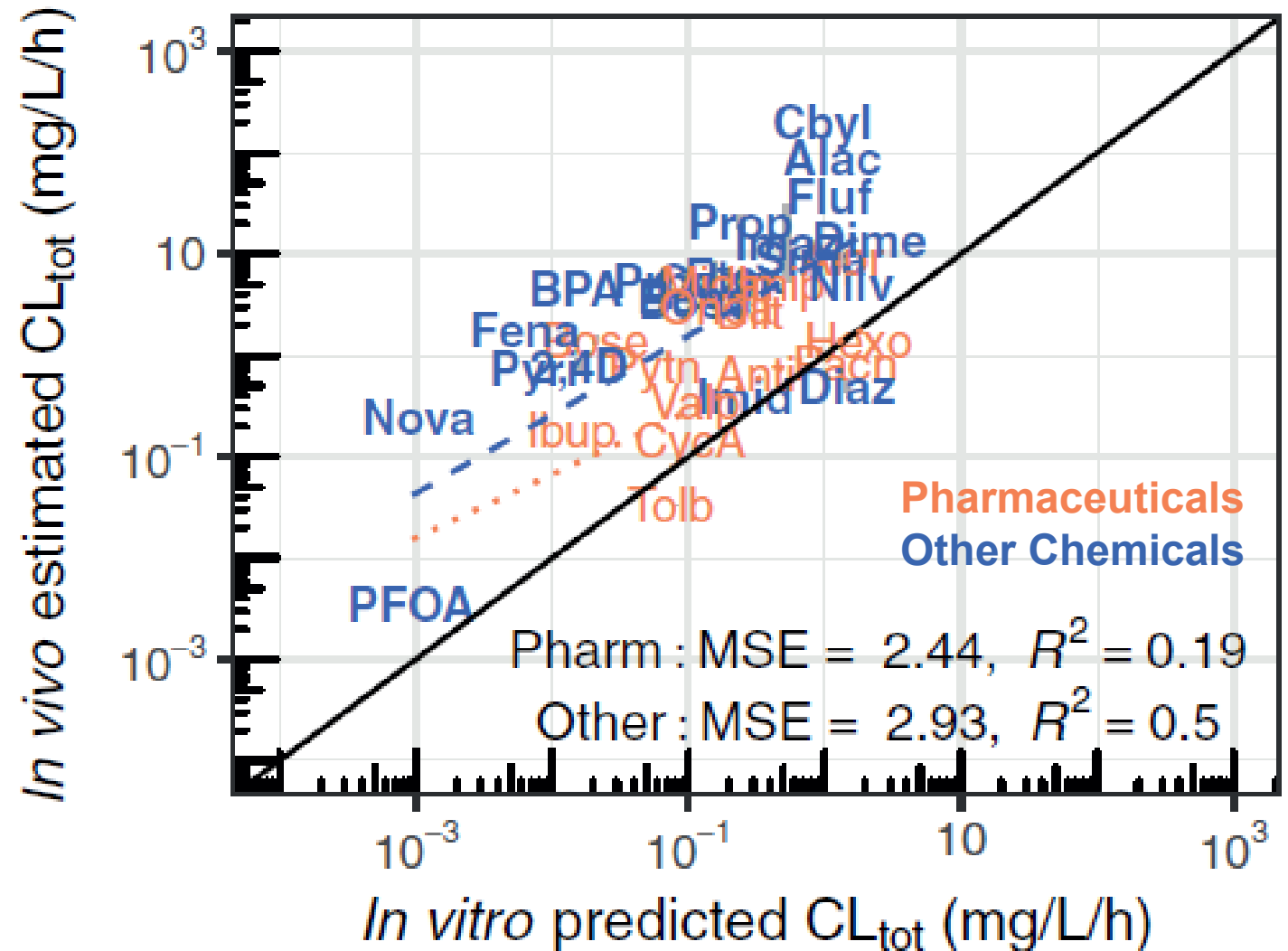
<https://github.com/USEPA/CompTox-ExpoCast-invivoPKfit>



Sayre et al. (in preparation)

- 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

Observed Total Clearance



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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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.)

Slide from Caroline Ring (ToxStrategies)

Ring *et al.* (2017)

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Sample NHANES
biometrics for
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Sex
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Population simulator for HTTK

Predict physiological
quantities

Tissue masses
Tissue blood flows
GFR (kidney function)
Hepatocellularity

Regression equations from literature
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Ring *et al.* (2017)

Uncertainty

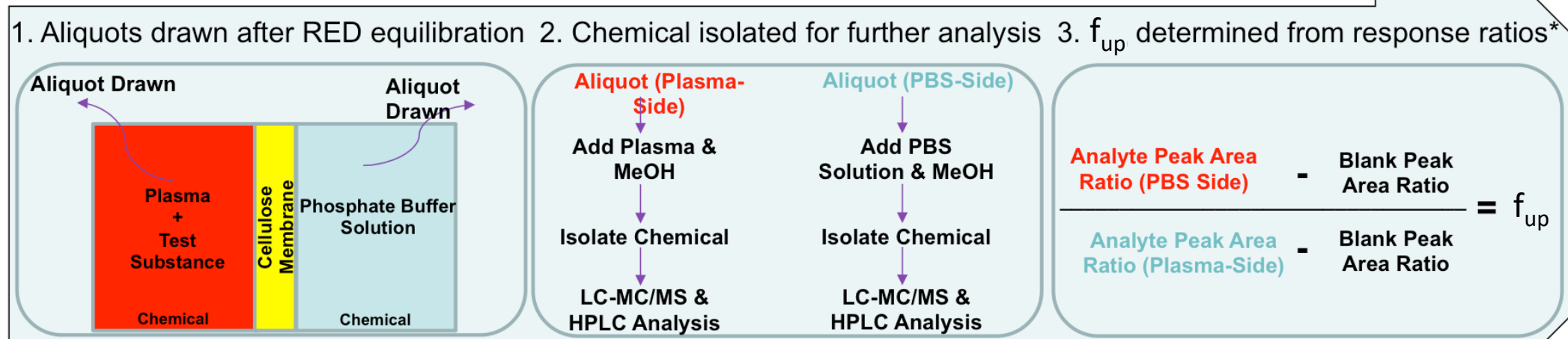
Until I open the
box, I don't know
what colors I
have...

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



Analytical Chemistry is an HTTK Bottleneck

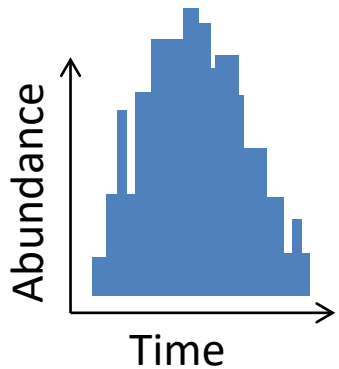
- For HTTK we always need to develop a chemical-specific method for quantitating amount of chemical *in vitro*
 - This is very different from HTS where the same readout (e.g., bioluminescence) can be used for most chemicals
- In Wetmore et al. (2012), the rapid equilibrium dialysis (RED) assay (Waters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



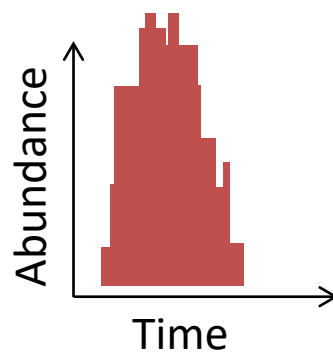
New HTTK Measurements and Uncertainty Analysis

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

Internal Standard



Chemical Peak

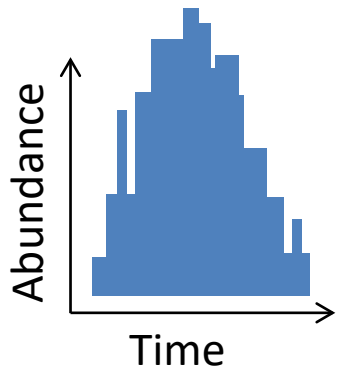


- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio R of the chemical peak to the ITSD

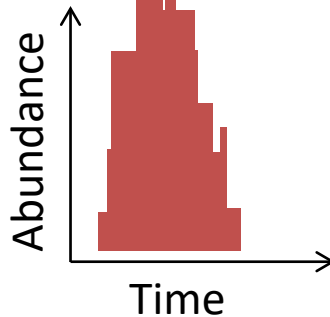
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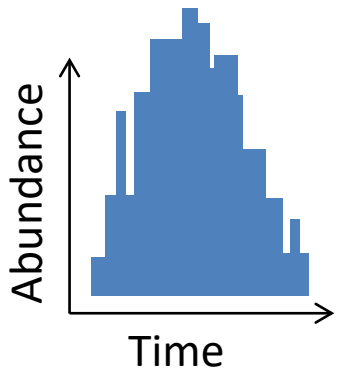


Chemical Peak

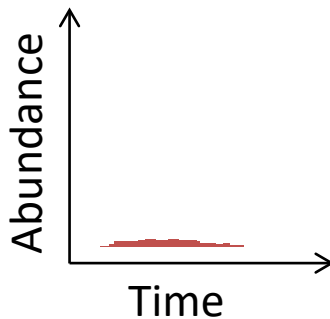


- Area of the internal standard (ITSD) at a known, fixed concentration fluctuates with time
- Find a peak that corresponds to chemical of interest, and then follow the ratio R of the chemical peak to the ITSD
- For new measurements HTTK (>200 compounds to data) performed by Cyprotex, we have modified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

Internal Standard



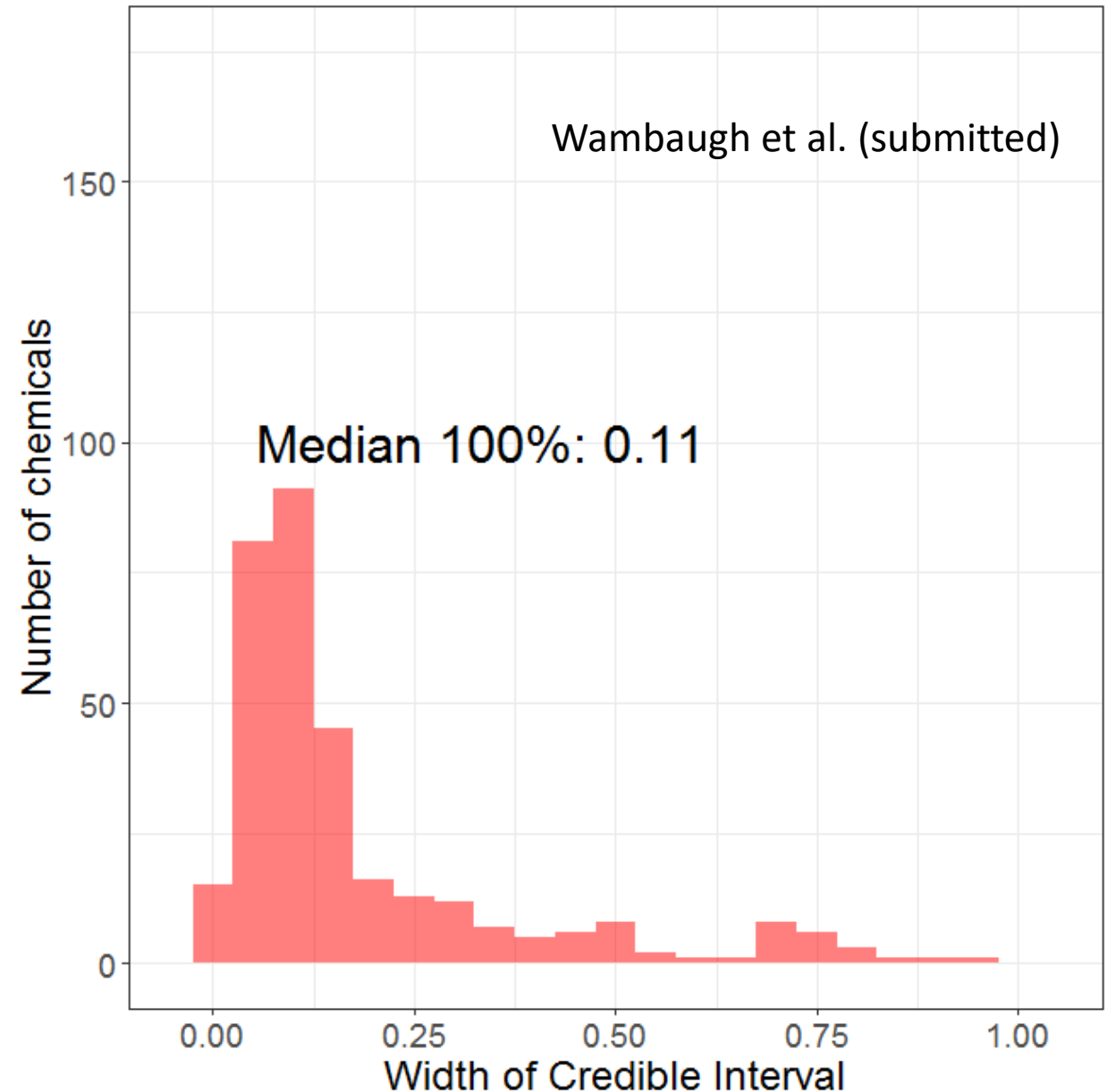
Chemical Peak



- 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

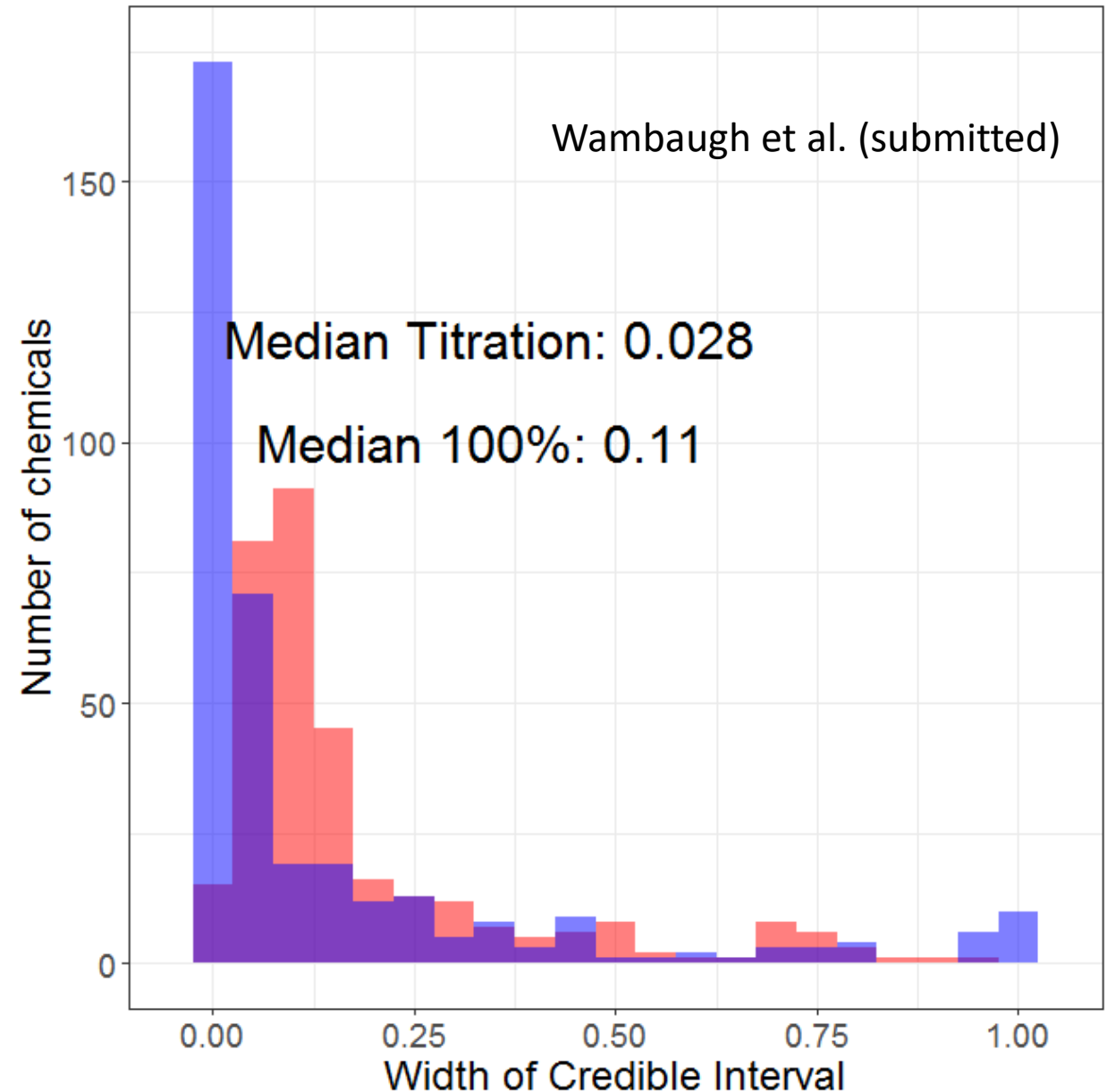
New Plasma Binding Protocol Reduces Uncertainty

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



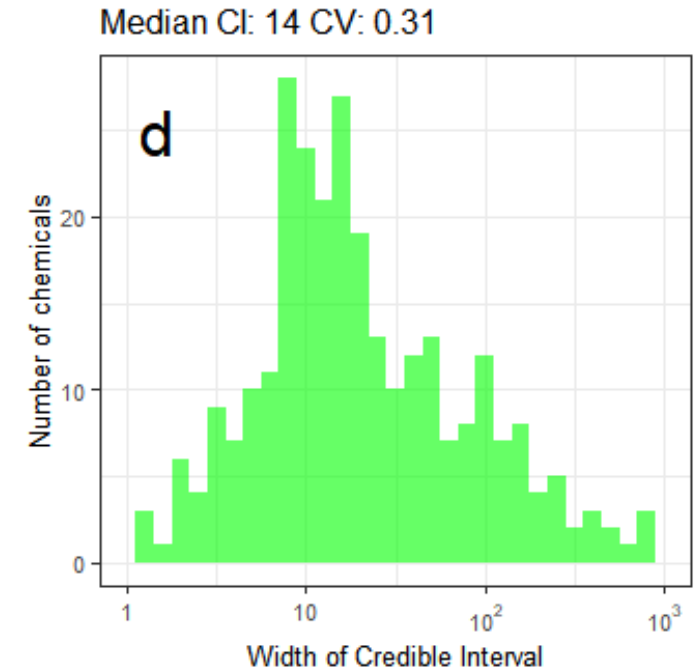
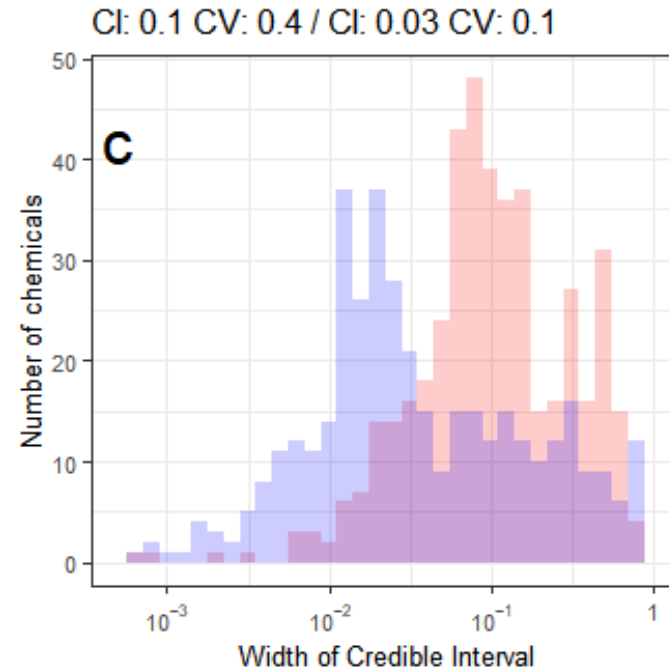
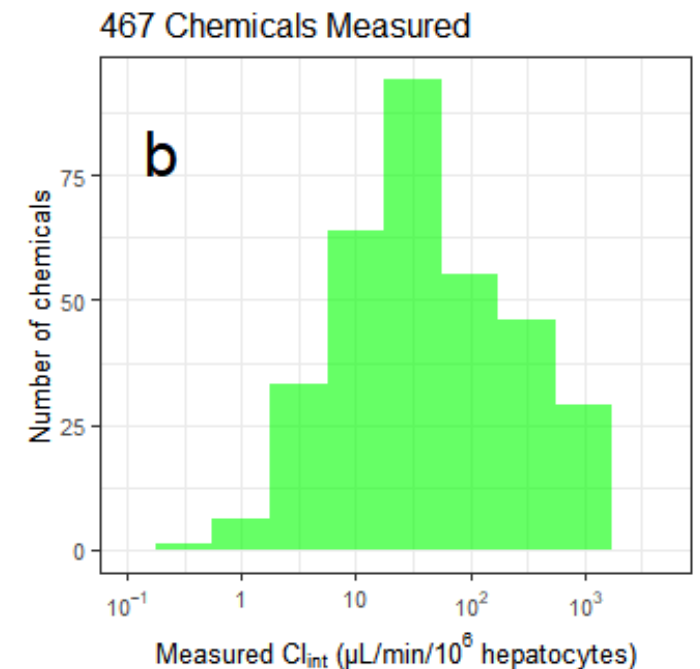
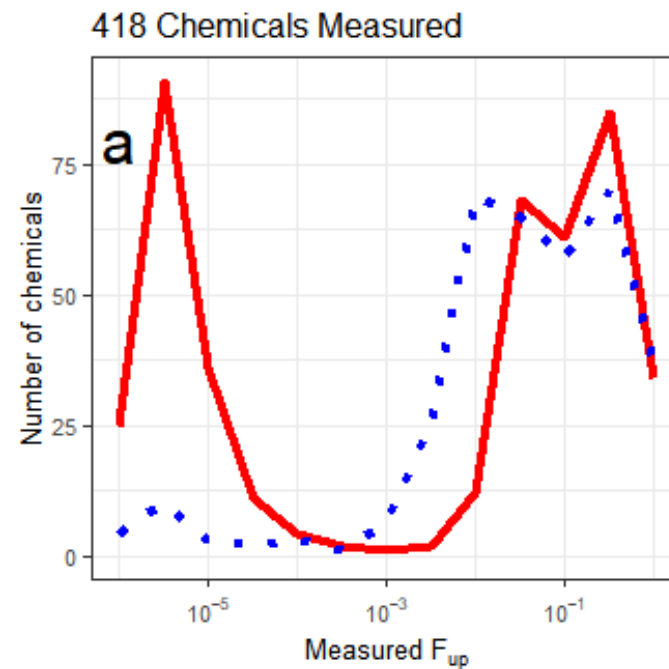
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- Median uncertainty for three-point assay: $\pm 1.4\%$

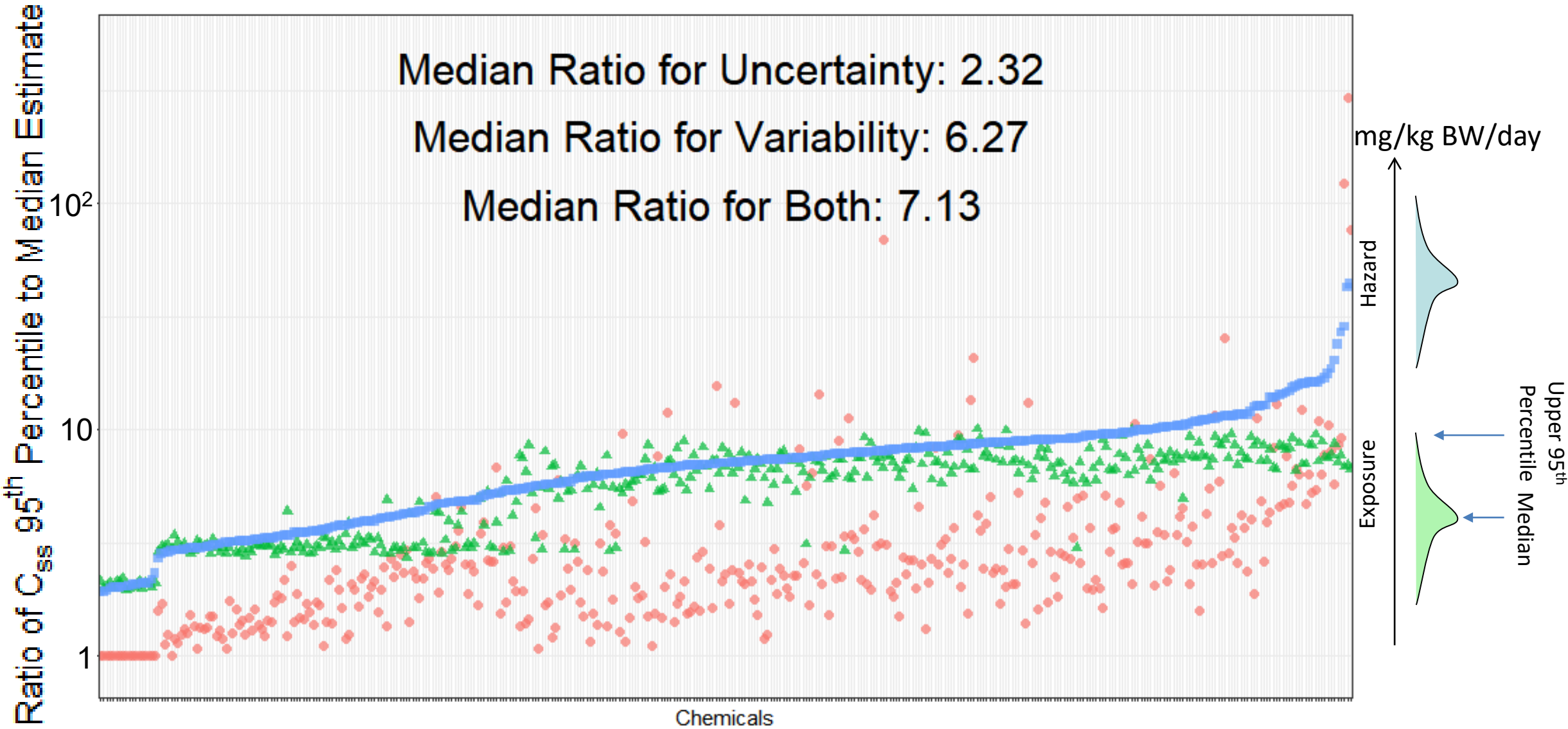


New Data!

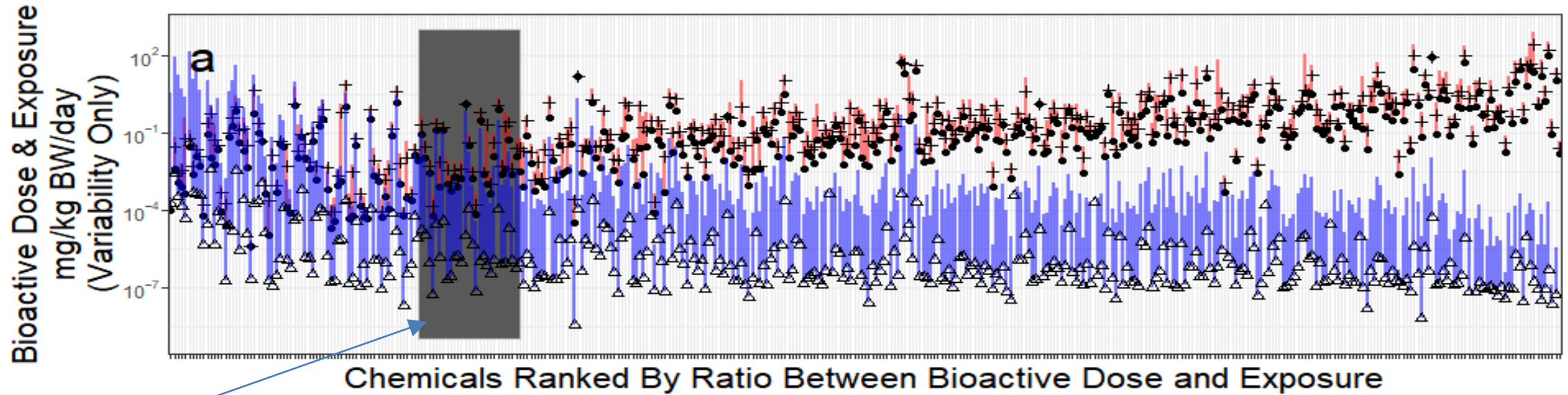
New experimental measurements of f_{up} and Cl_{int} are reported for 418 and 467 chemicals, respectively. These data raise the HTTK chemical coverage of the ToxCast Phase I and II libraries to 57%.



Quantifying the Impact of Uncertainty

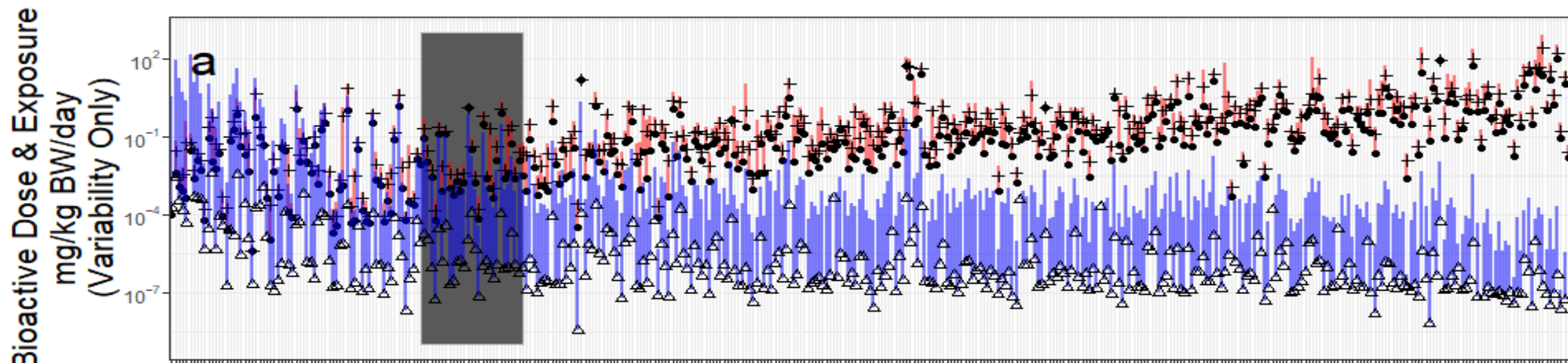


New IVIVE For 393 ToxCast Chemicals

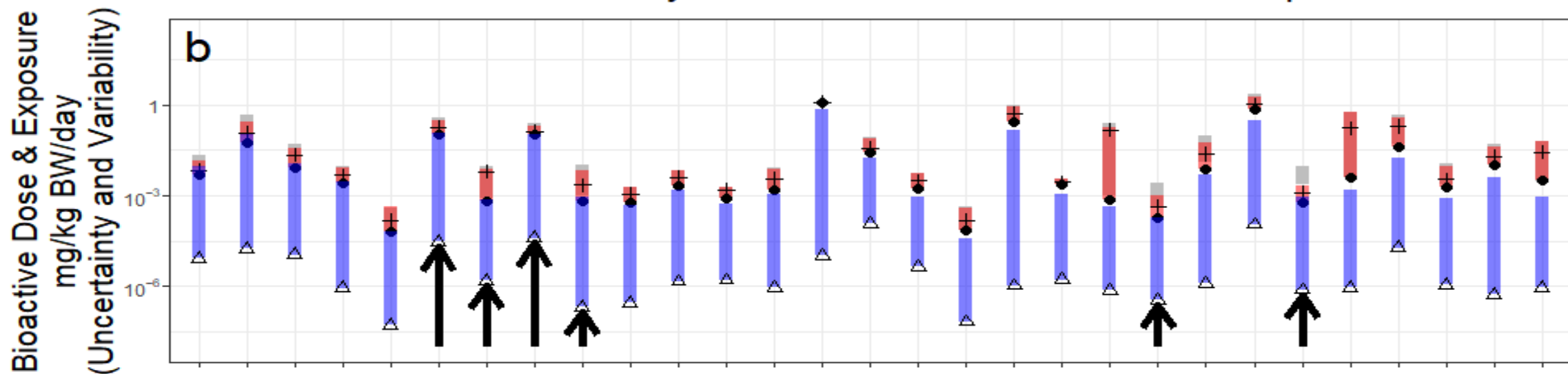


Including chemical-specific uncertainty only caused changes in whether or not exposure and bioactivity overlapped in a small region

The Impact of Measurement Uncertainty



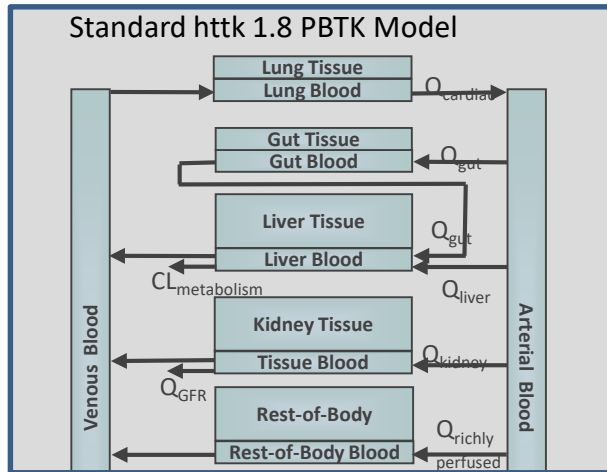
Chemicals Ranked By Ratio Between Bioactive Dose and Exposure



Only six more chemicals overlap

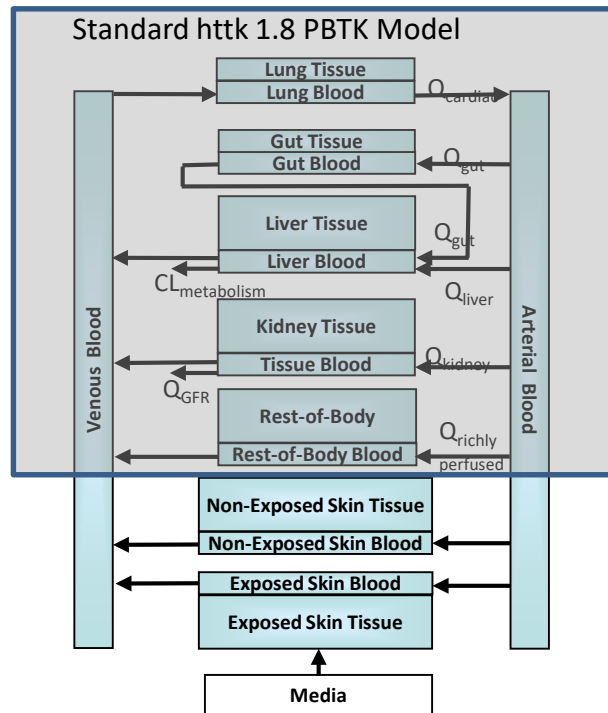
Wambaugh et al. (submitted)

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

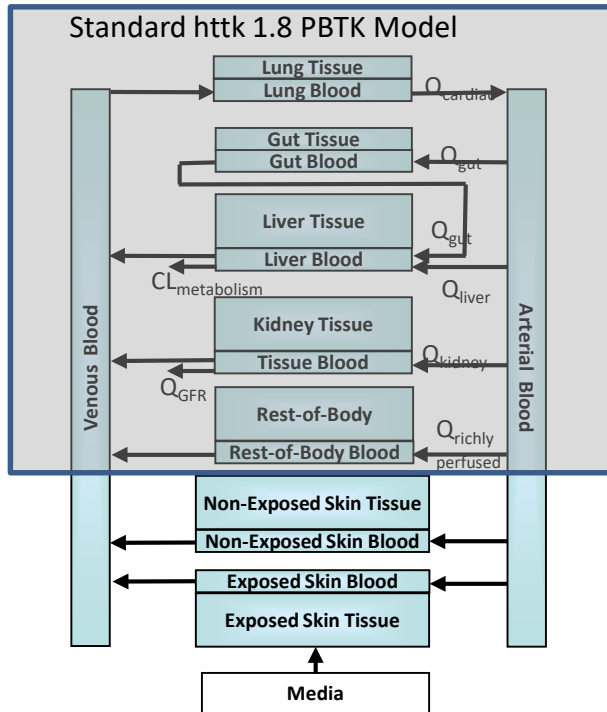
New HT-PBTK Models



Dermal Exposure Route

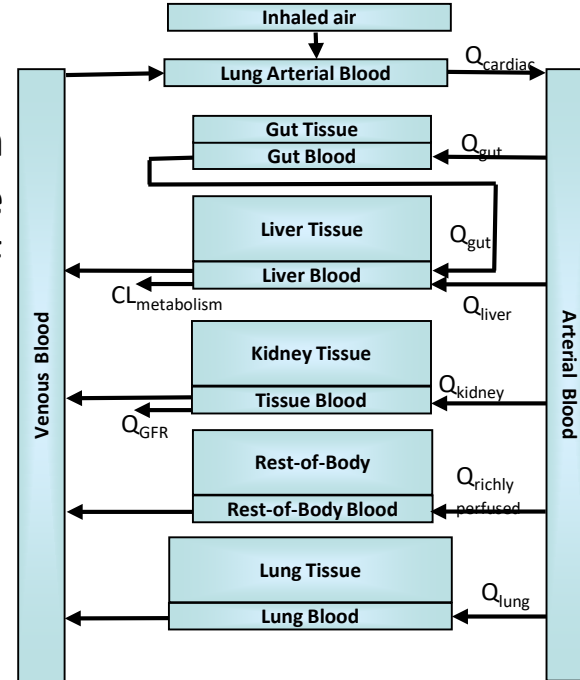
EPA, Unilever, INERIS

Gas Inhalation Exposure Route EPA, USAF



Dermal Exposure Route EPA, Unilever, INERIS

New HT-PBTK Models



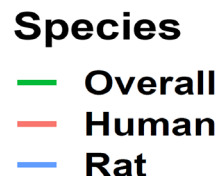
The diagram illustrates the flow of air, blood, and substances between the environment, lungs, gut, liver, body, and kidneys. It is organized into a central column of organs/tissues, flanked by a large vertical bar on the left representing the Venous Blood reservoir and a large vertical bar on the right representing the Arterial Blood reservoir.

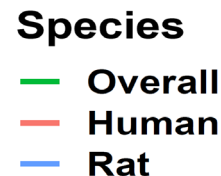
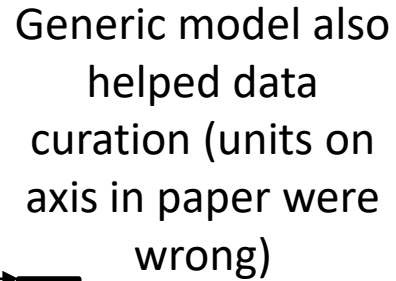
Central Column (Organs/Tissues):

- Lung:** A dashed box at the top contains "Inhaled Air" and "Exhaled Breath". Below it is a box for "Mucous". The "Alveolar Space" is a box below the mucous. The "Lung Blood" and "Lung Tissue" are stacked boxes below the alveolar space.
- Gut:** The "Gut Blood" and "Gut Tissue" are stacked boxes. Below them is the "Gut Lumen" box.
- Liver:** The "Liver Blood" and "Liver Tissue" are stacked boxes.
- Body:** The "Body Blood" and "Body Tissue" are stacked boxes.
- Kidney:** The "Kidney Blood" and "Kidney Tissue" are stacked boxes at the bottom.

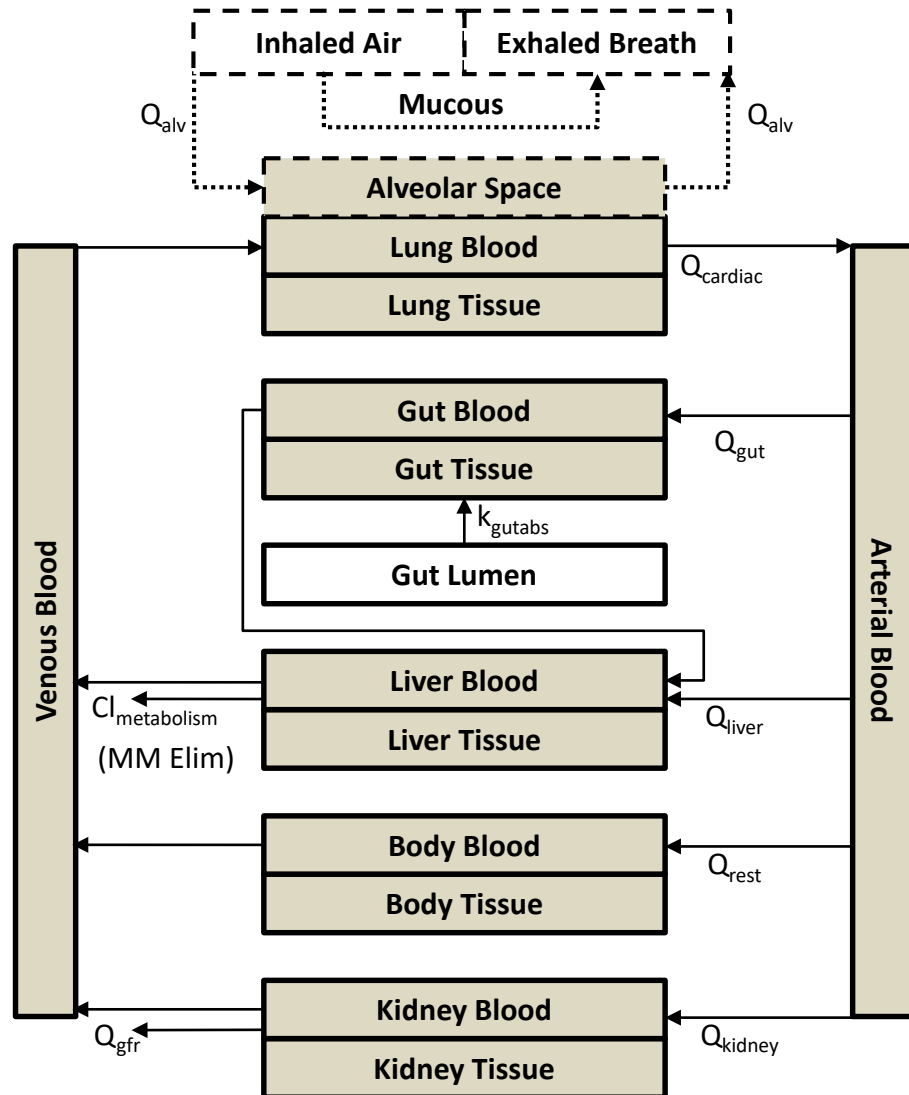
Flows and Parameters:

- Air Flow:** Q_{alv} (Alveolar ventilation) is shown entering the Alveolar Space from the Inhaled Air and leaving as Exhaled Breath.
- Blood Flow:**
 - $Q_{cardiac}$ (Cardiac output) flows from the Arterial Blood reservoir to the Lung Blood.
 - Q_{gut} (Gut blood flow) flows from the Arterial Blood reservoir to the Gut Blood.
 - Q_{liver} (Liver blood flow) flows from the Arterial Blood reservoir to the Liver Blood.
 - Q_{rest} (Rest of body blood flow) flows from the Arterial Blood reservoir to the Body Blood.
 - Q_{kidney} (Kidney blood flow) flows from the Arterial Blood reservoir to the Kidney Blood.
- Substance Flow:**
 - Substances flow from the Gut Lumen into the Gut Blood, indicated by k_{gutabs} (gut absorption coefficient).
 - Substances flow from the Liver Blood into the Venous Blood reservoir, indicated by $Cl_{metabolism}$ (MM Elim) (metabolic clearance).
 - Substances flow from the Kidney Blood into the Venous Blood reservoir, indicated by Q_{gfr} (glomerular filtration rate).





Generic Gas Inhalation Model



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

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

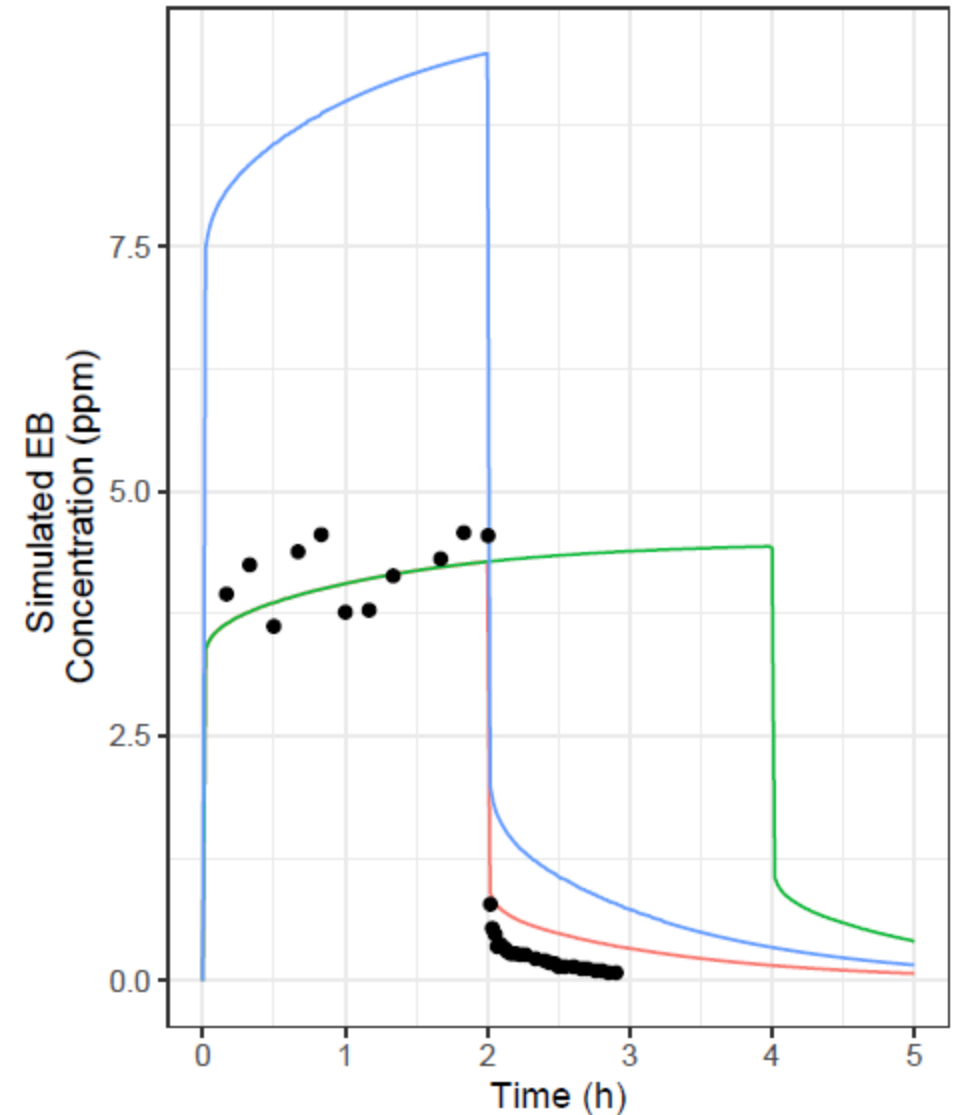
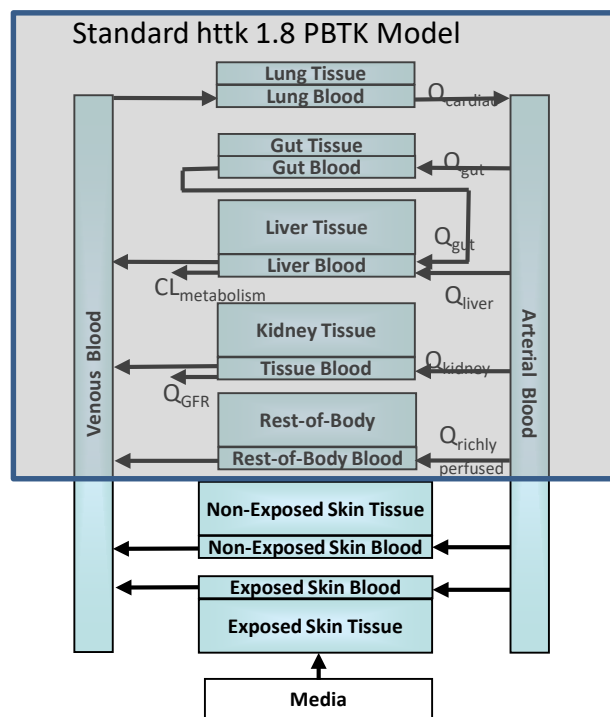


Figure from Matt Linakis (USAFSAM)

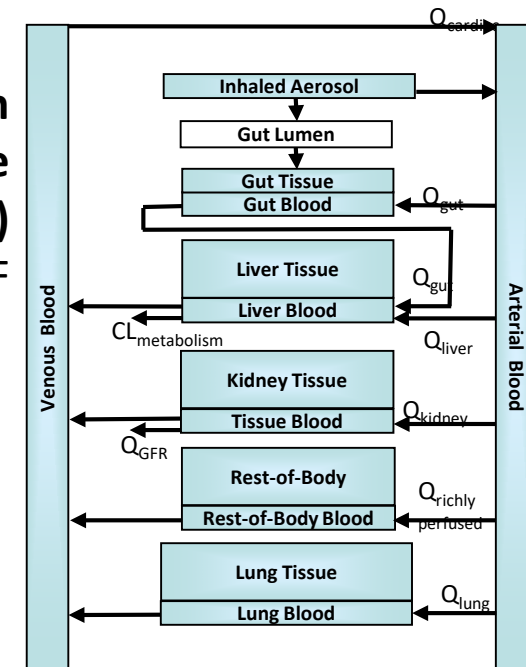
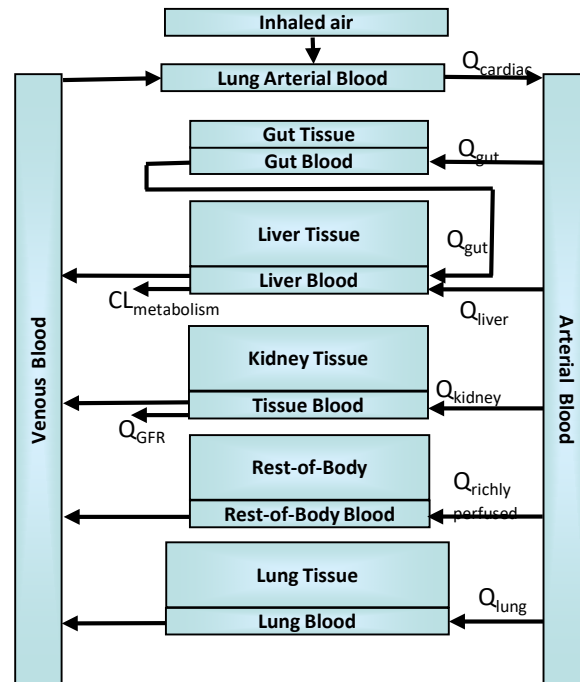
Gas Inhalation Exposure Route EPA, USAF



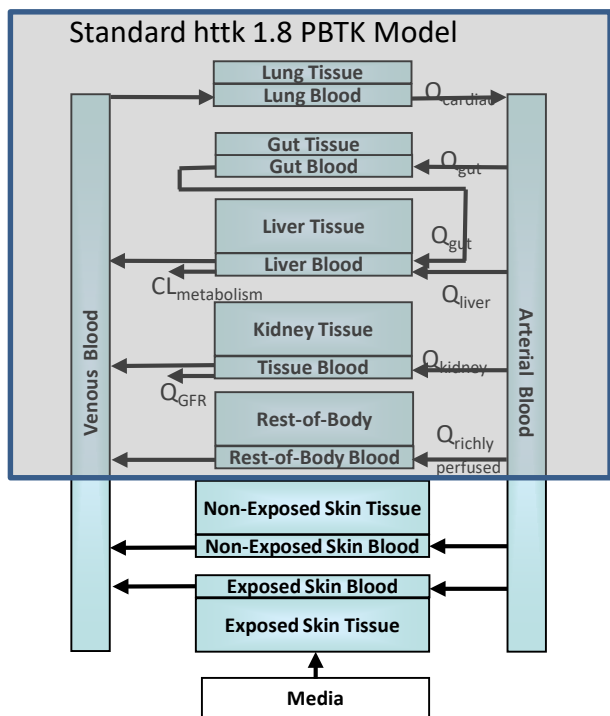
Dermal Exposure Route EPA, Unilever, INERIS

New HT-PBTK Models

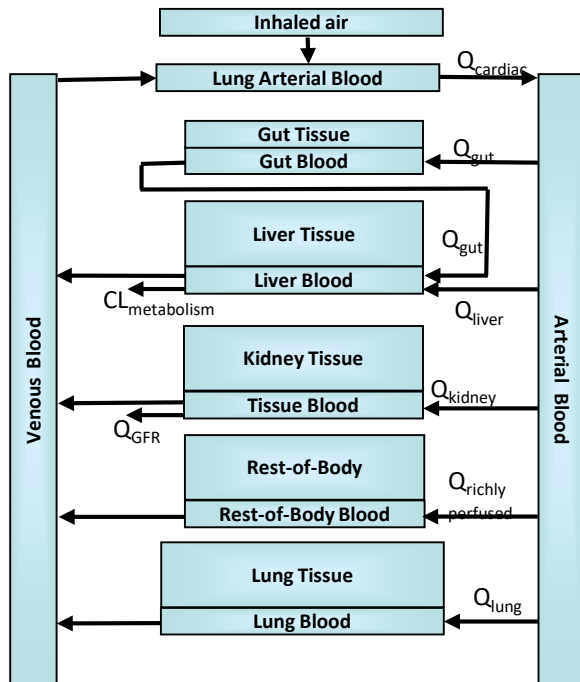
Aerosol Inhalation Exposure Route (with APEX model) EPA, USAF



Gas Inhalation Exposure Route EPA, USAFSAM

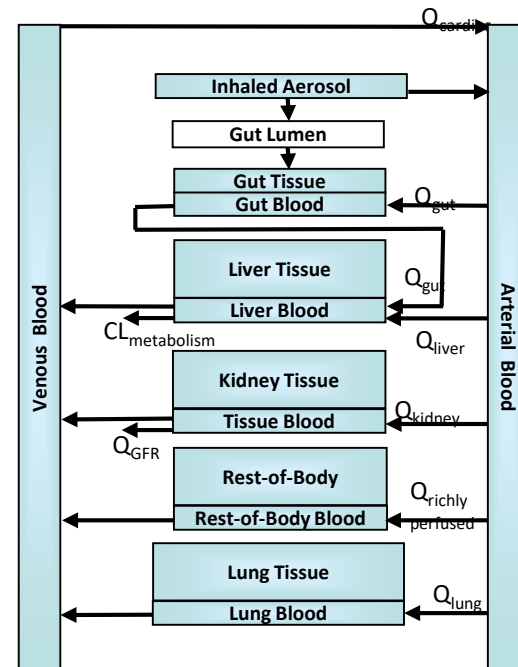


Dermal Exposure Route EPA, Unilever, INERIS

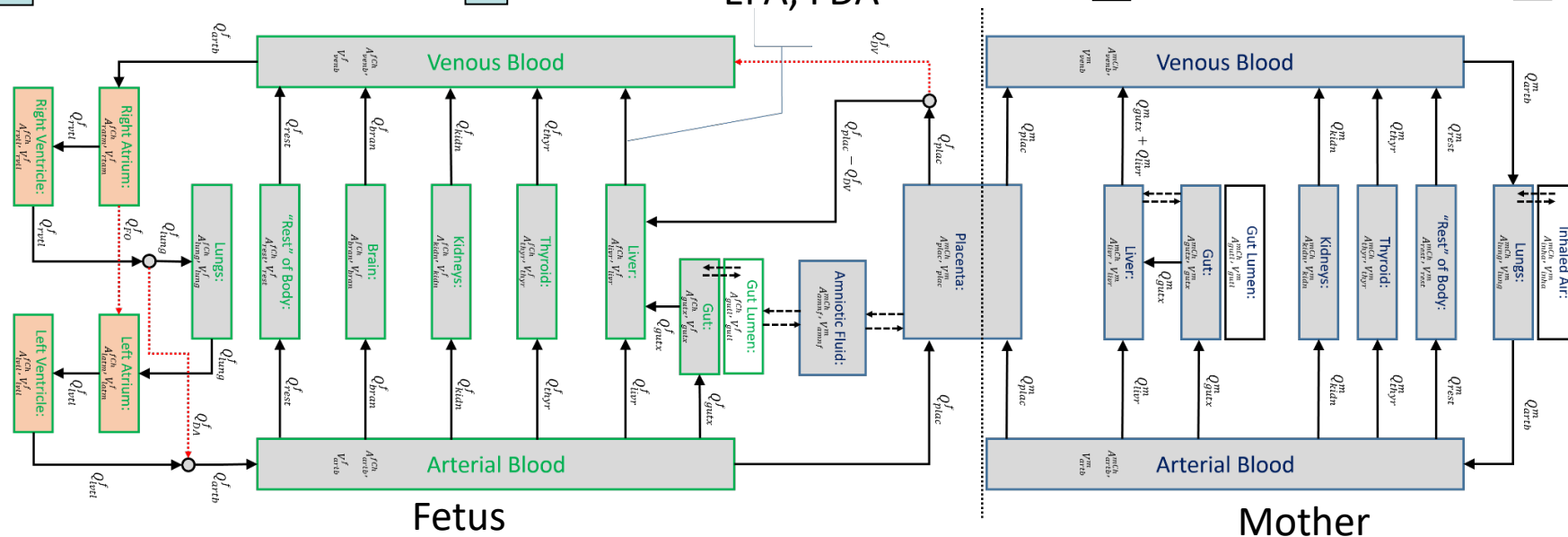


New HT-PBTK Models

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



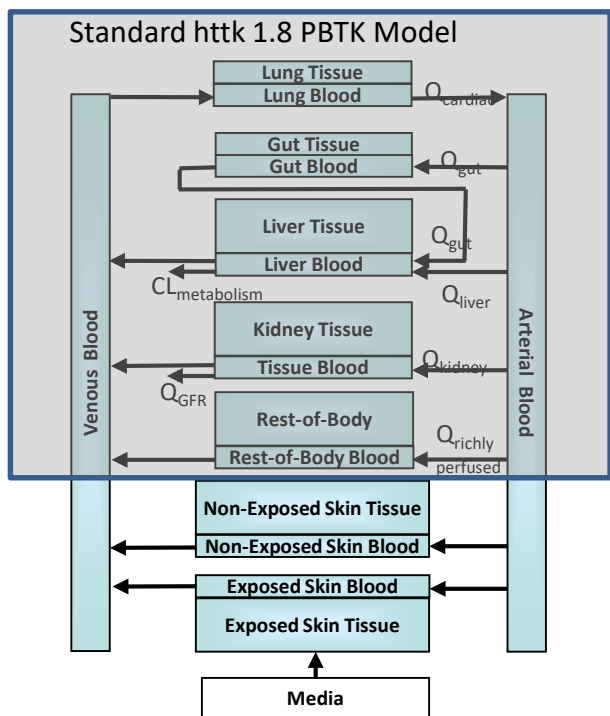
Human Gestational Model EPA, FDA



Fetus

Mother

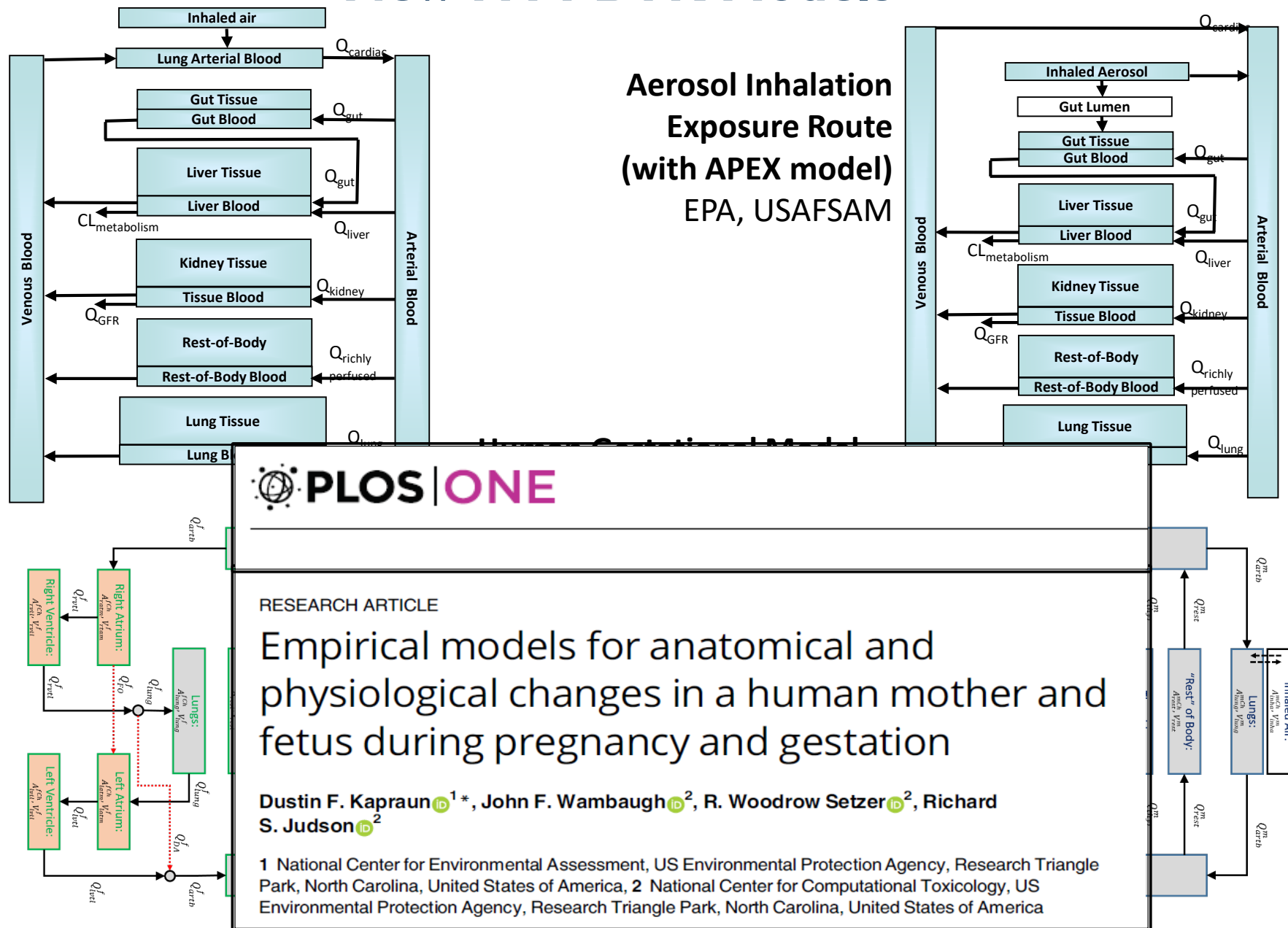
Gas Inhalation Exposure Route EPA, USAFSAM



Dermal Exposure Route EPA, Unilever, INERIS

New HT-PBTK Models

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



PLOS ONE

RESEARCH ARTICLE

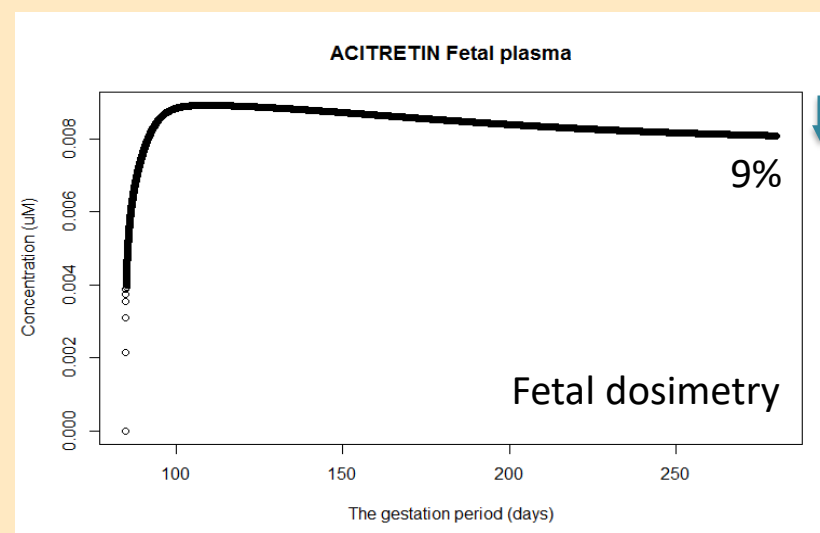
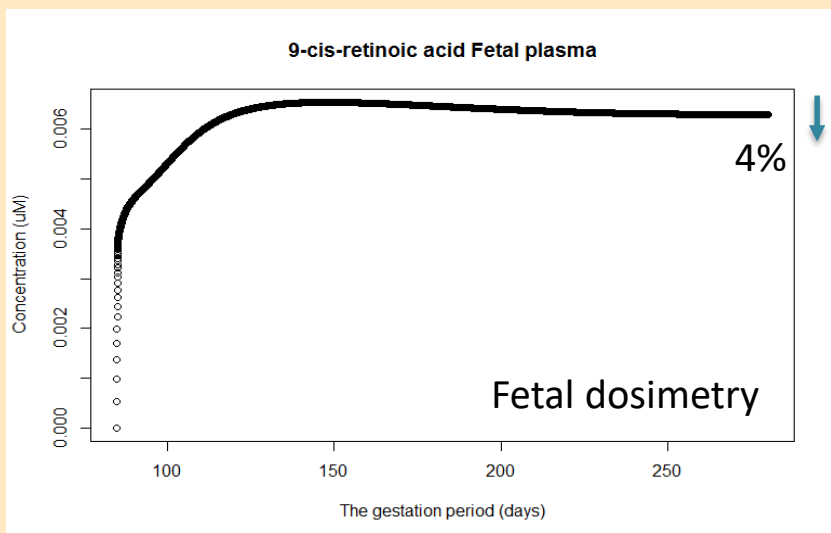
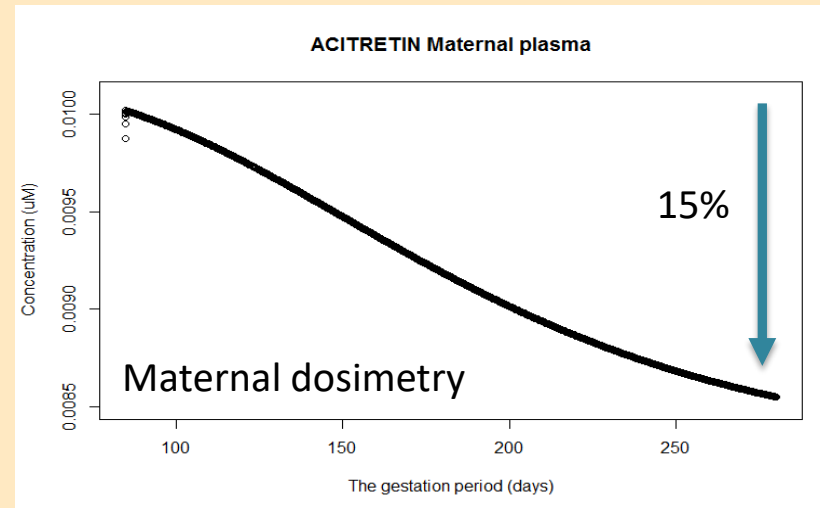
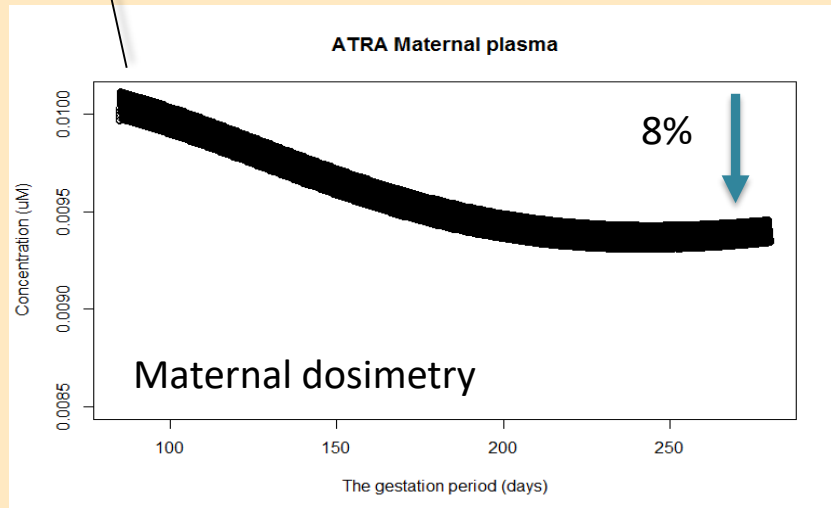
Empirical models for anatomical and physiological changes in a human mother and fetus during pregnancy and gestation

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

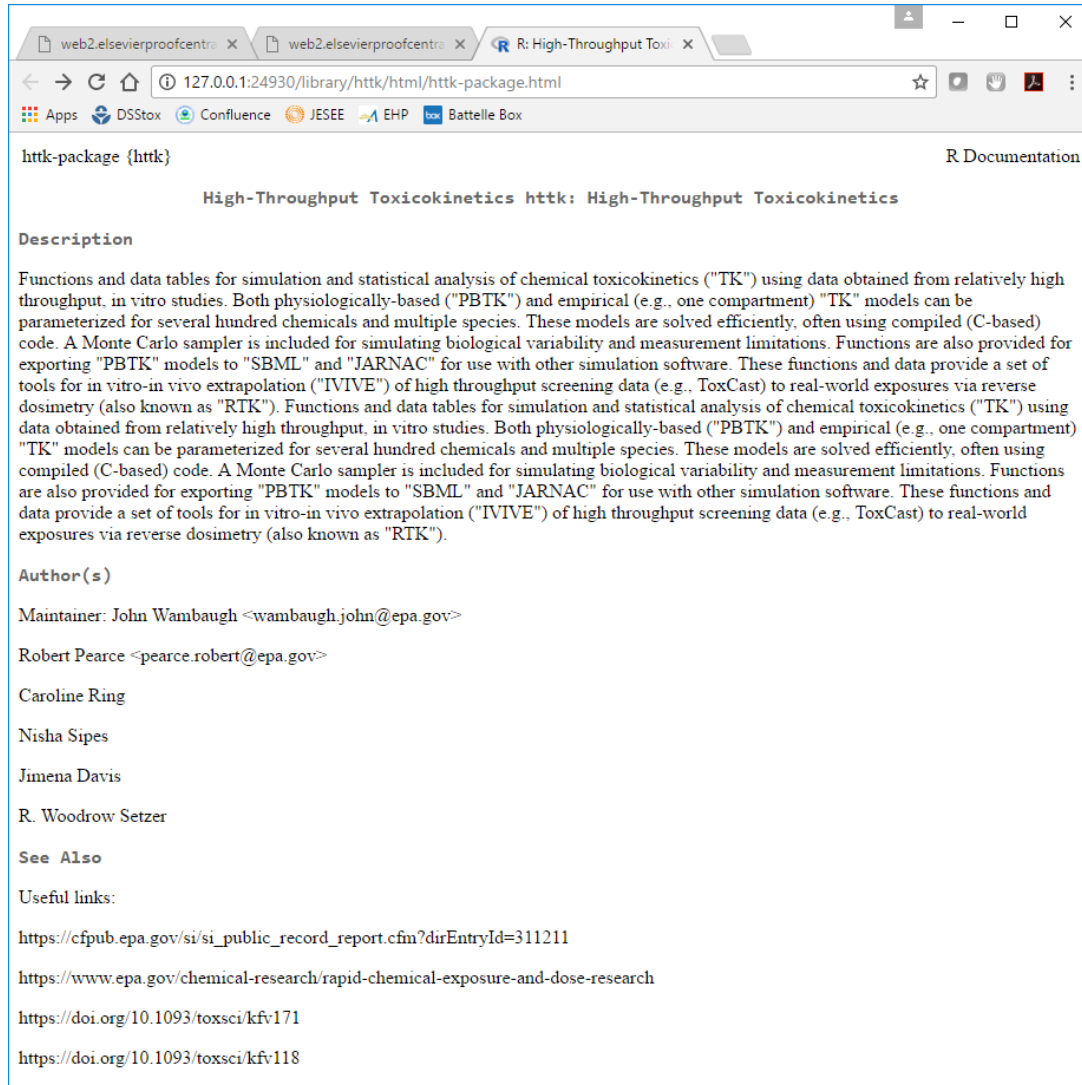
Maternal/Fetal HTTK Model Predictions for Retinoid Analogues:



- 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)”



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").

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See Also

Useful links:

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 shows the R help environment for the 'httk' package. The left sidebar contains the following information:

- httk-package {httk}**
- High-Thro**
- Description**
Functions and data tables for simulat throughput, in vitro studies. Both phy parameterized for several hundred cl 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 (are also provided for exporting "PBT data provide a set of tools for in vitro exposures via reverse dosimetry (als
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- See Also**
- Useful links:**
- 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>

The main window displays the online documentation for package 'httk' version 1.6. It includes a list of links: DESCRIPTION file, User guides, package vignettes and other documentation, and Package NEWS. Below this is a section titled 'Help Pages' with a list of functions: A, B, and C. The functions listed are:

- 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.
- 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.
- 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**: Find 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.

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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

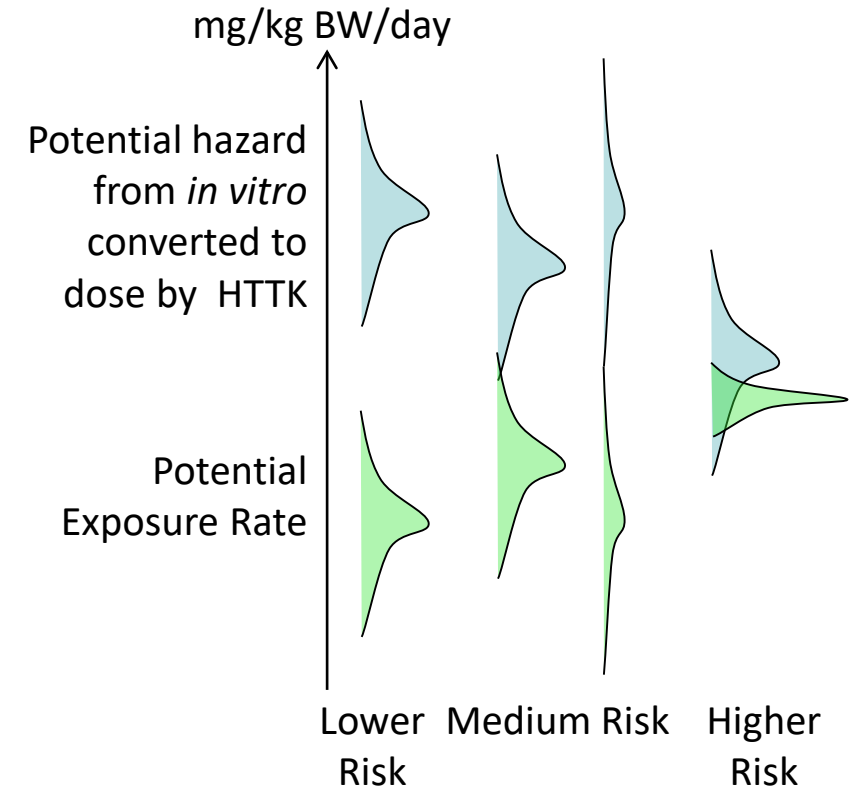
```
subset(get_cheminfo(in
fo="all"), Compound%in%
c("A", "B", "C"))
```

```
> 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")
```

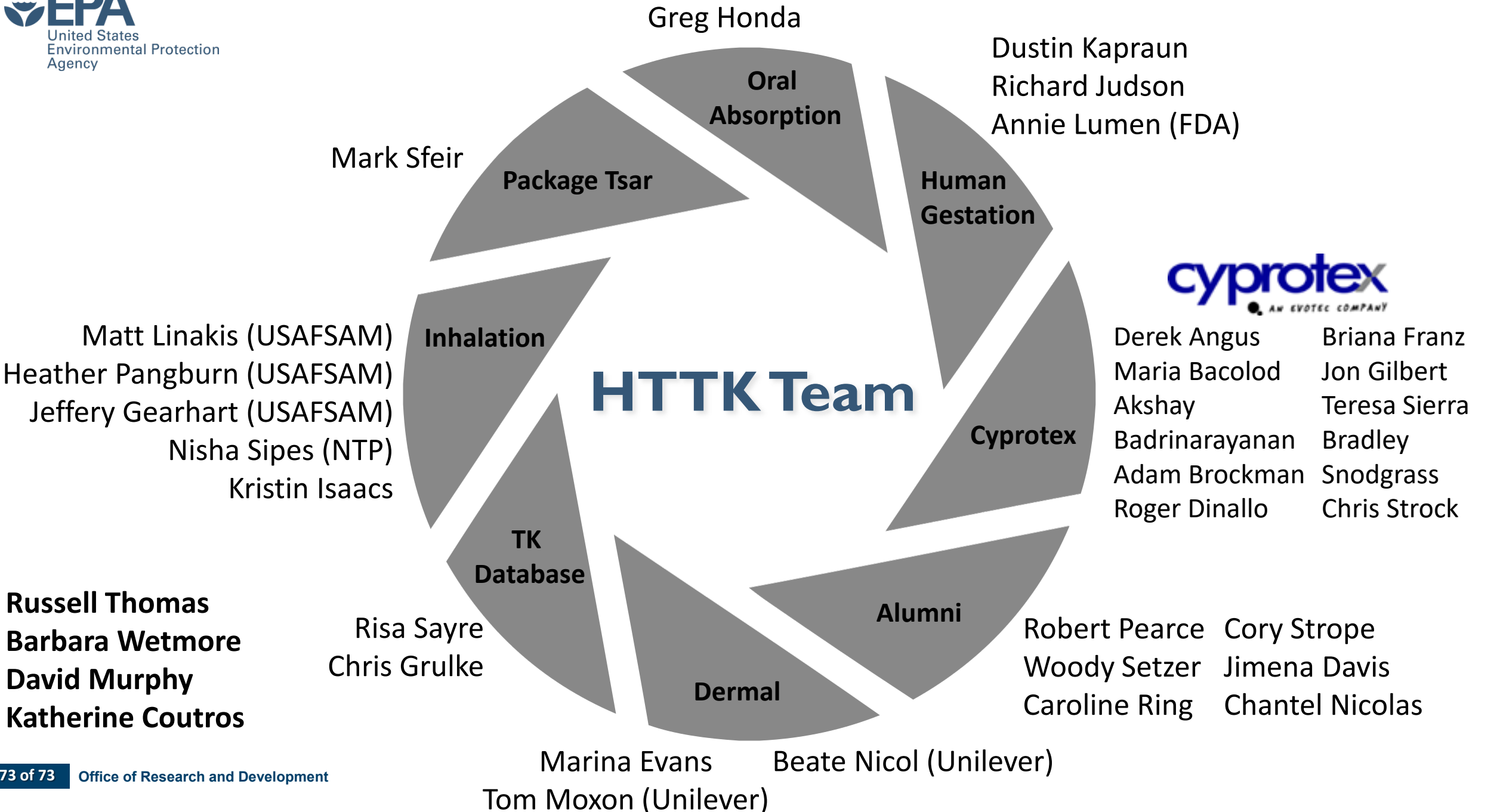
Compound	CAS	logP	pKa_Accept	pKa_Donor	MW	Human.Clint	Human.Clint.p Value	Human.Funboud.plasma	DSSTox_Substance_ID	Structure_Formula	Substance_Type
2,4-d	94-75-7	2.81	<NA>	2.81	221.03	0	0.149	0.04	DTXSID0020442	C8H6Cl2O3	Single Compound
2,4-db	94-82-6	3.53	<NA>	4.5	249.09	0	0.104	0.01	DTXSID7024035	C10H10Cl2O3	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

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





ExpoCast Project (Exposure Forecasting)

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