

# Placing toxicology data in the context of exposure

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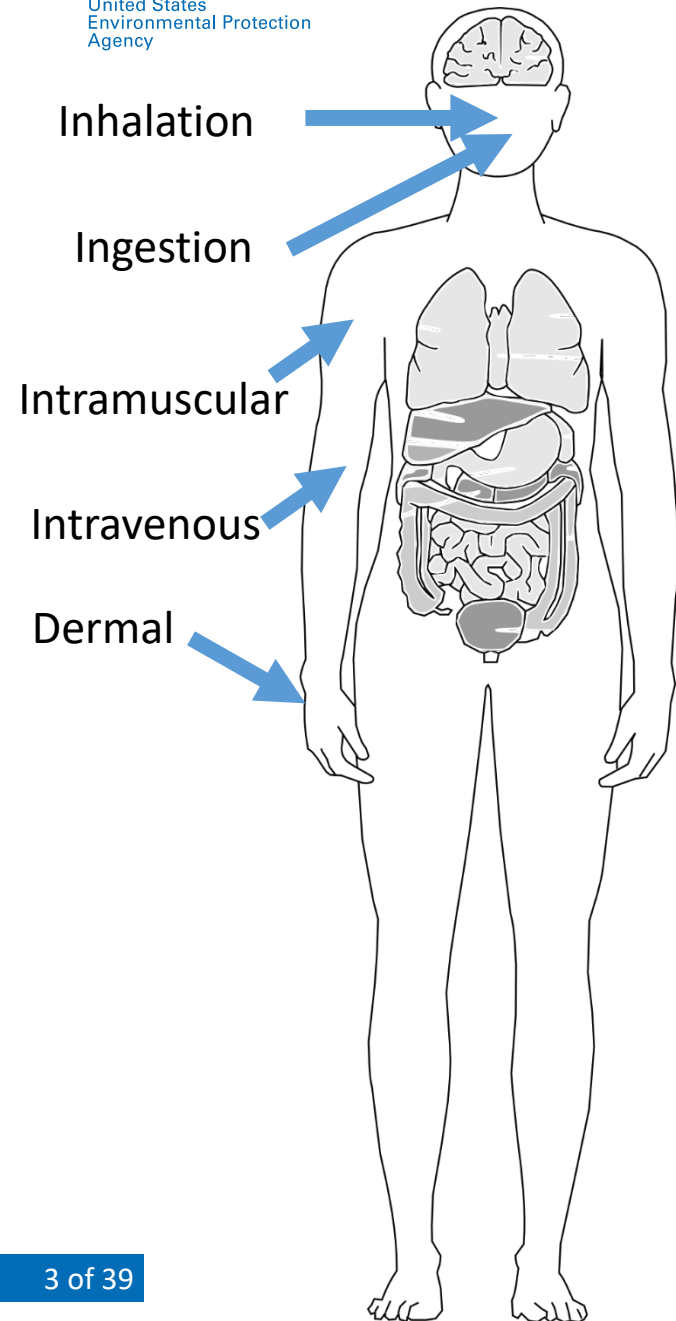


*The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA*

# Overview

- Motivation: “The dose makes the poison”
- Risk = hazard vs. exposure
- Problem: Traditional approaches insufficient to screen thousands of chemicals
- Solution: New approach methodologies (NAMs)
  - NAMs for hazard
  - NAMs for exposure
- Problem: Hazard NAMs estimate biologically active *concentrations*. How to compare to external exposure rates?
- Solution: In vitro-in vivo extrapolation using high-throughput toxicokinetic modeling

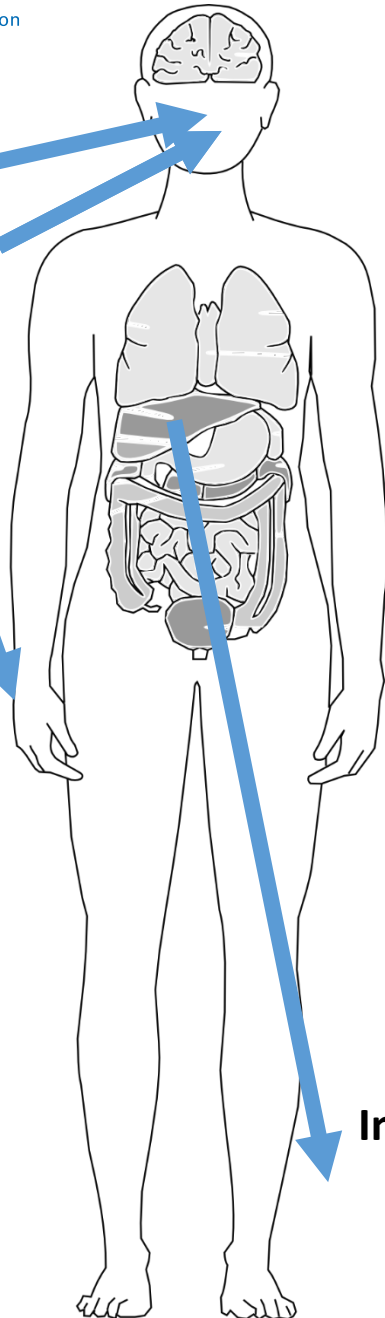
# Scenario: You are exposed to chemicals





# Scenario: You are exposed to chemicals

**External exposure**



## Things you might want to know....

- What chemicals are you exposed to? How much? How often?
- Do the chemicals get inside your body?
- If so, how much gets inside?
  - For example, what is the concentration of each chemical in your blood?
- Is that enough to cause any kind of health effect?

**Internal dose** = Amount/concentration of chemical or drug in one or more body tissues of interest

# Difficulty level: Answer these questions for thousands of environmental chemicals, and for the whole population

- Most non-food, non-drug chemicals, ranging from industrial waste to dyes to packing materials, are covered by the Toxic Substances Control Act (TSCA) and come under EPA's purview
- Currently 41,953 “active” (currently-used) chemicals on TSCA inventory, and hundreds of new ones listed every year
- Need a way to rapidly **prioritize** chemicals for more detailed evaluation

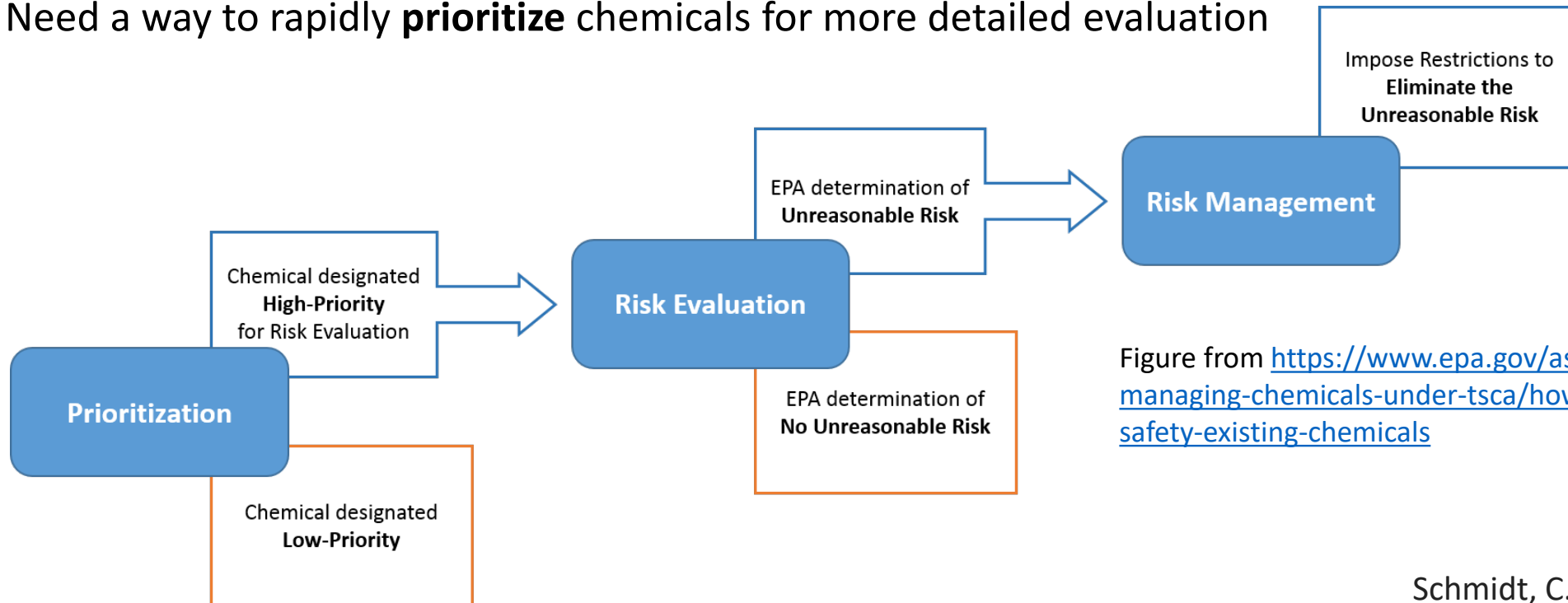


Figure from <https://www.epa.gov/assessing-and-managing-chemicals-under-tsca/how-epa-evaluates-safety-existing-chemicals>

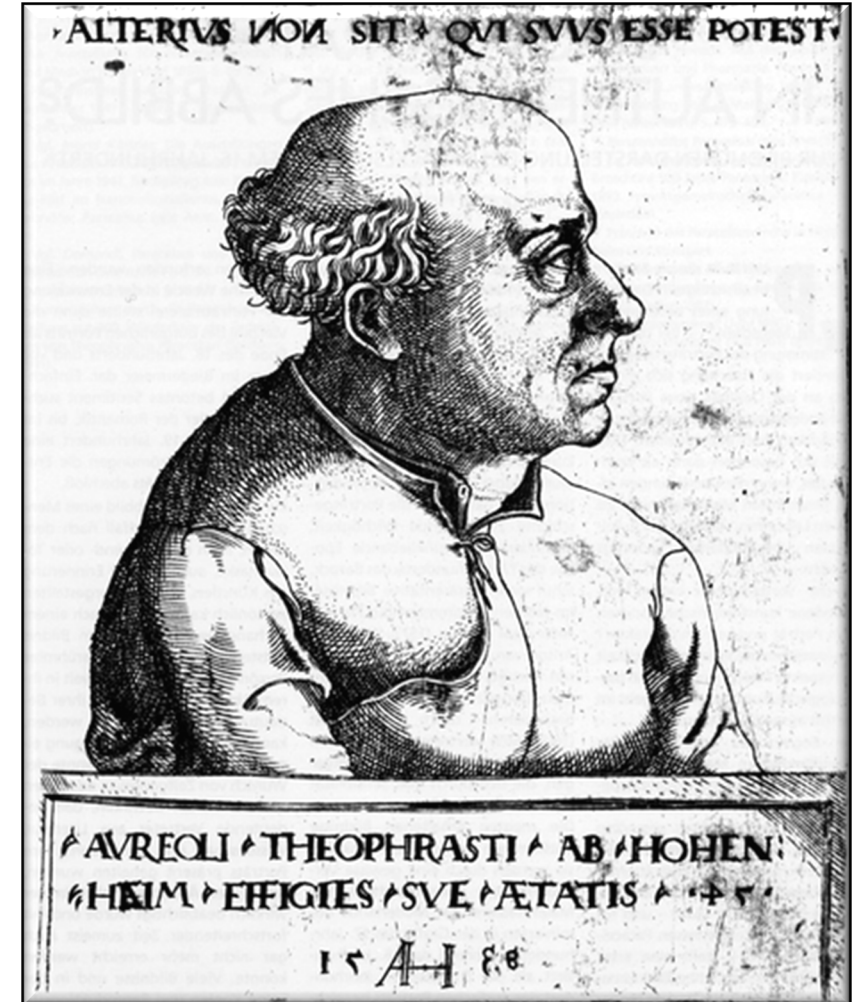
# Paracelsus: “The dose makes the poison”

*“What is there that is not poison? All things are poison and nothing is without poison. Solely the dose determines that a thing is not a poison”*  
— Paracelsus (1493-1541)

**Hazard:** What type of harm could occur, and what dose would be necessary to produce this harm

**Exposure:** What dose someone actually receives

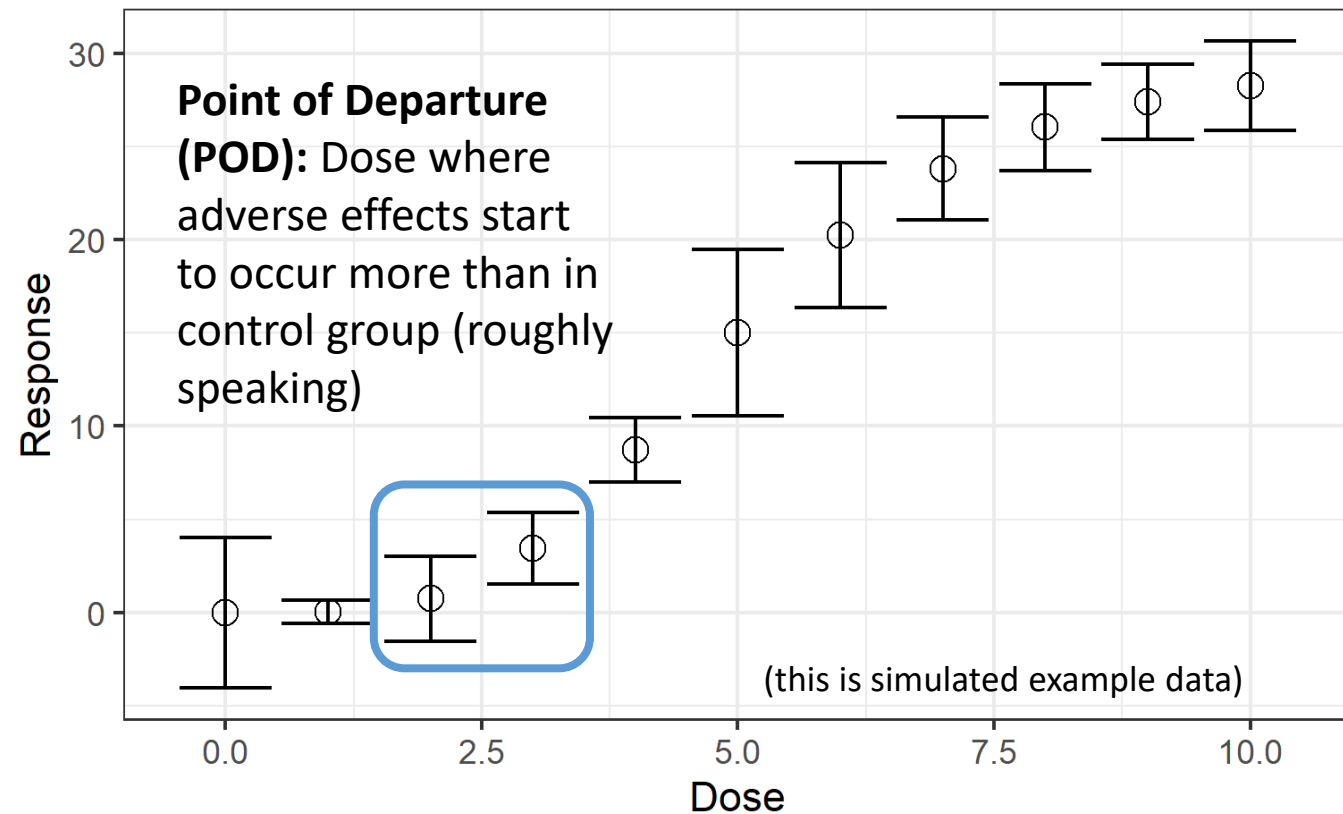
**Risk:** The likelihood that harm will occur from the exposure actually received



# Traditional hazard data comes from studies *in vivo*, one chemical at a time



[Observe adverse effects in each dose group  
after days, weeks, months, or years of dosing]



# New approach methodologies for hazard: *In vitro* high-throughput screening (HTS) assays, e.g. ToxCast/Tox21



Thousands of chemicals are screened in concentration-response across hundreds of *in vitro* assays for various kinds of biological activity (binding, signaling, viability...) – now with transcriptomics!

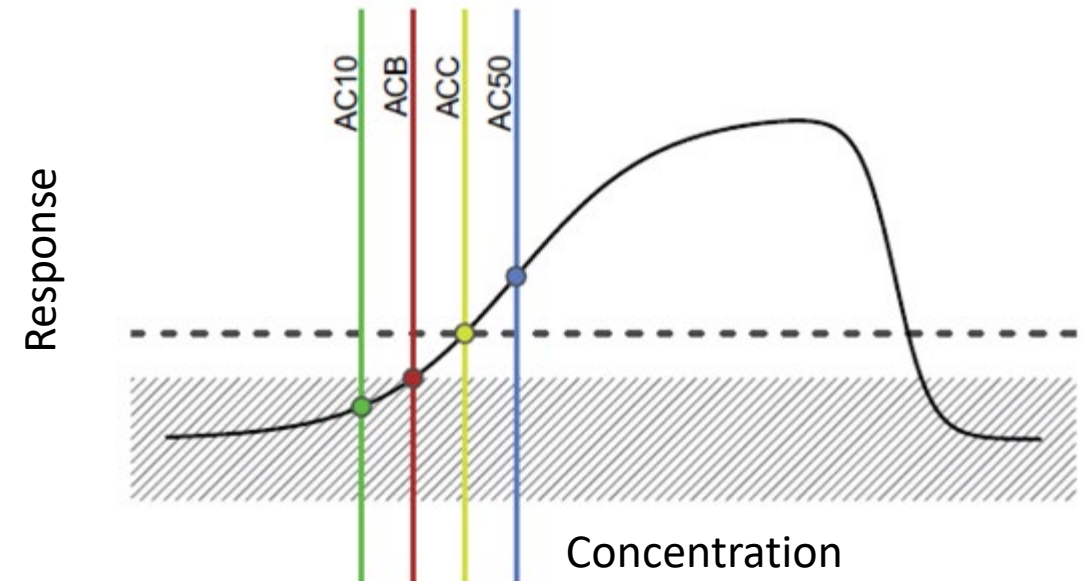
[Schmidt 2009; Dix et al. 2007; Kavlock et al. 2018; Filer *et al.*, 2016; Franzosa et al. 2021]

**Data:** For each chemical, ***in vitro* concentrations associated with bioactivity** in each assay, if any

All data are public:

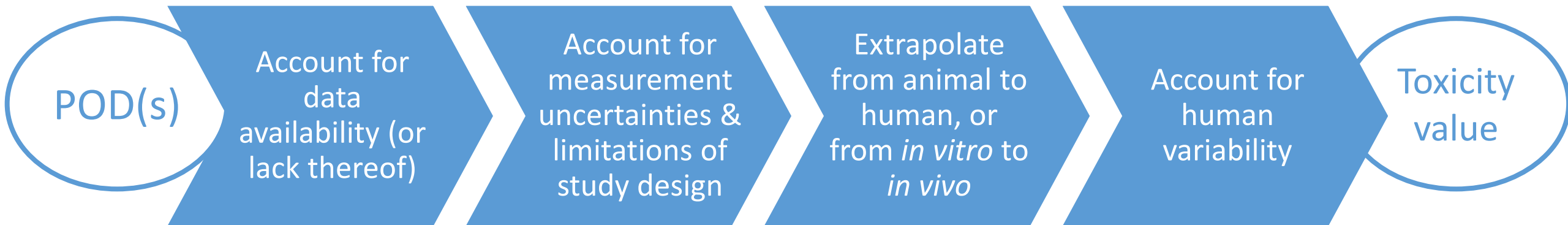
<http://comptox.epa.gov/dashboard/>

<https://www.epa.gov/chemical-research/exploring-toxcast-data-downloadable-data>





Hazard data is then extrapolated to develop a toxicity value:  
a dose below which an adverse effect is considered unlikely



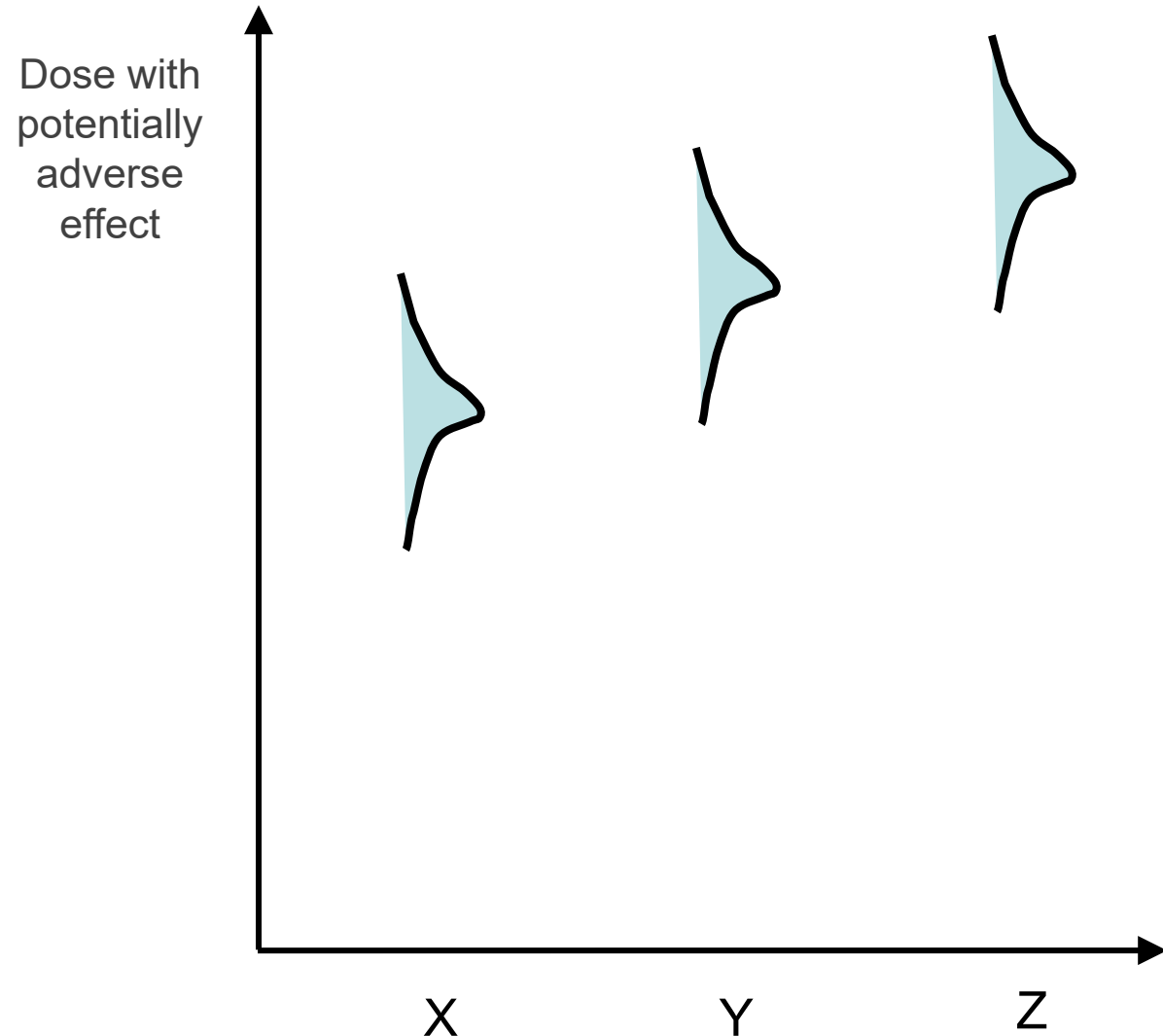
Chiu et al. (2018)  
National Academies of Science (2009)  
US EPA (2002)

# Sometimes chemicals are ranked based on hazard/toxicity data alone

Here are some fictitious toxicity values for three chemicals, shown as distributions

Poll: Which of these three chemicals poses the greatest concern for human health?

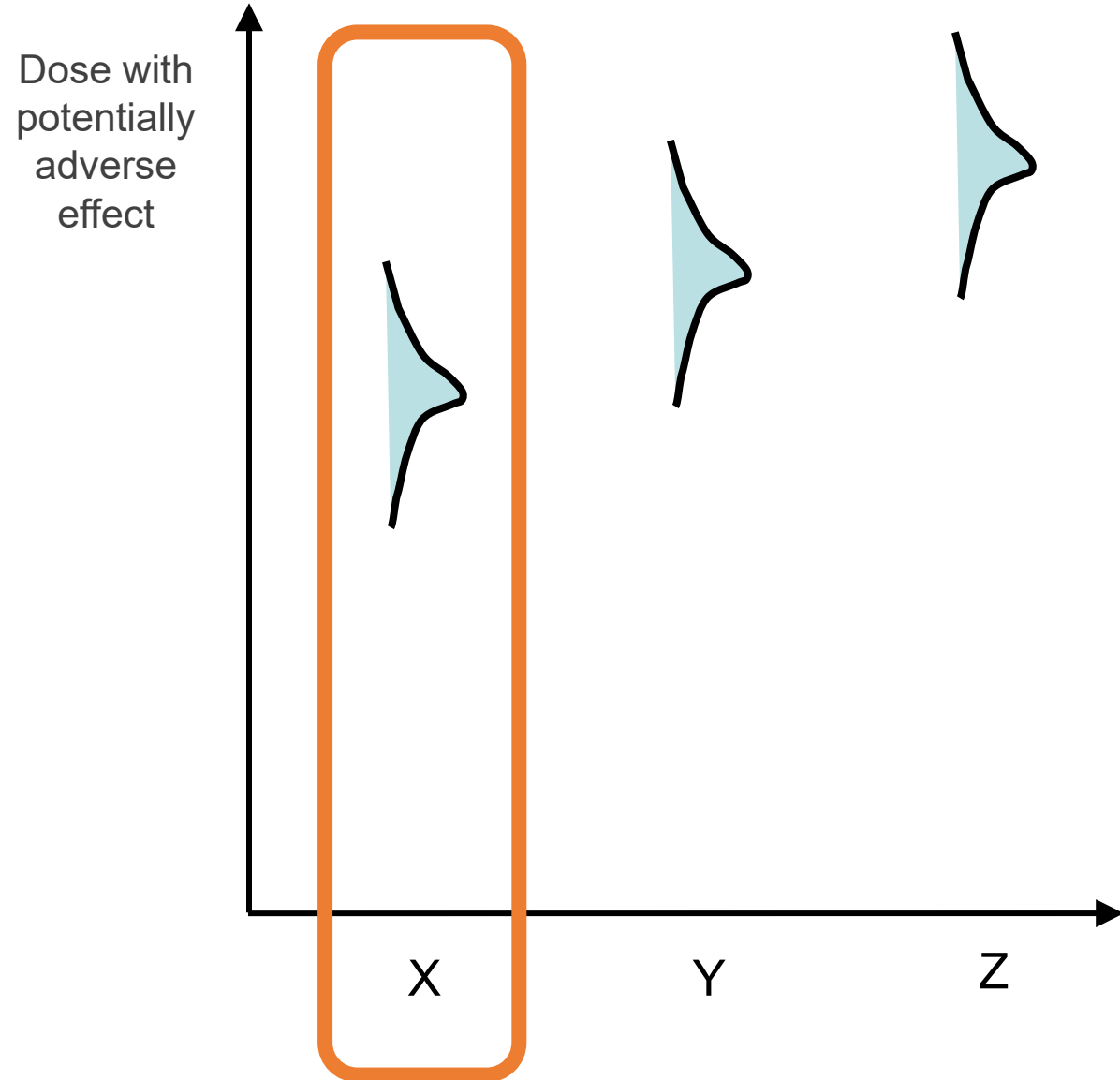
1. X
2. Y
3. Z



# Sometimes chemicals are ranked based on hazard/toxicity data alone

Chemical X has the lowest toxicity value, meaning it's the most potent (produces adverse effects at the lowest dose).

But does that make it the most concerning?



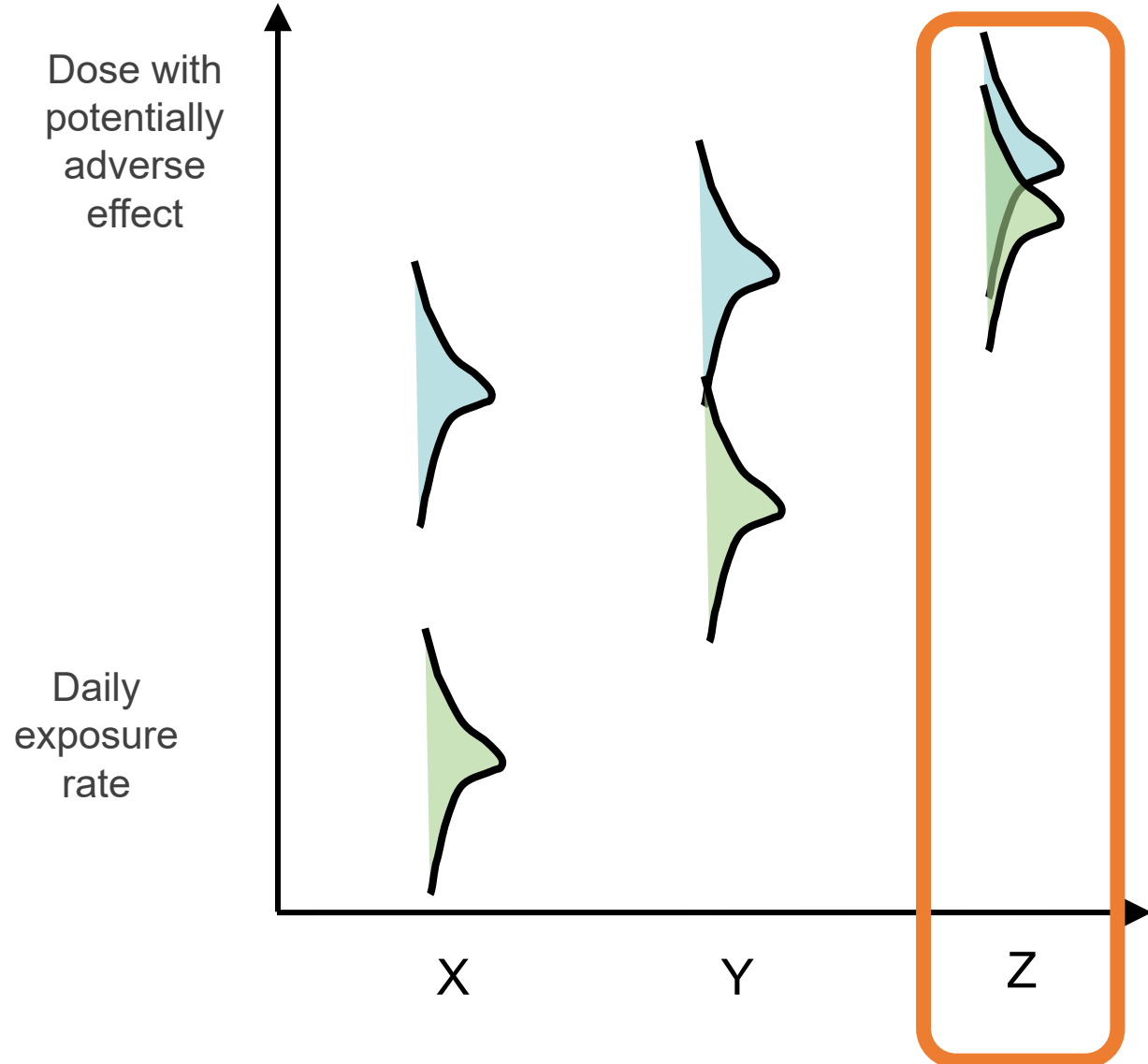
But “the dose makes the poison”: hazard/toxicity needs to be put in the context of exposure to assess risk

When we know exposure,  
**Chemical Z** is actually the most concerning!

“Margin of exposure” (MOE)  
approach:

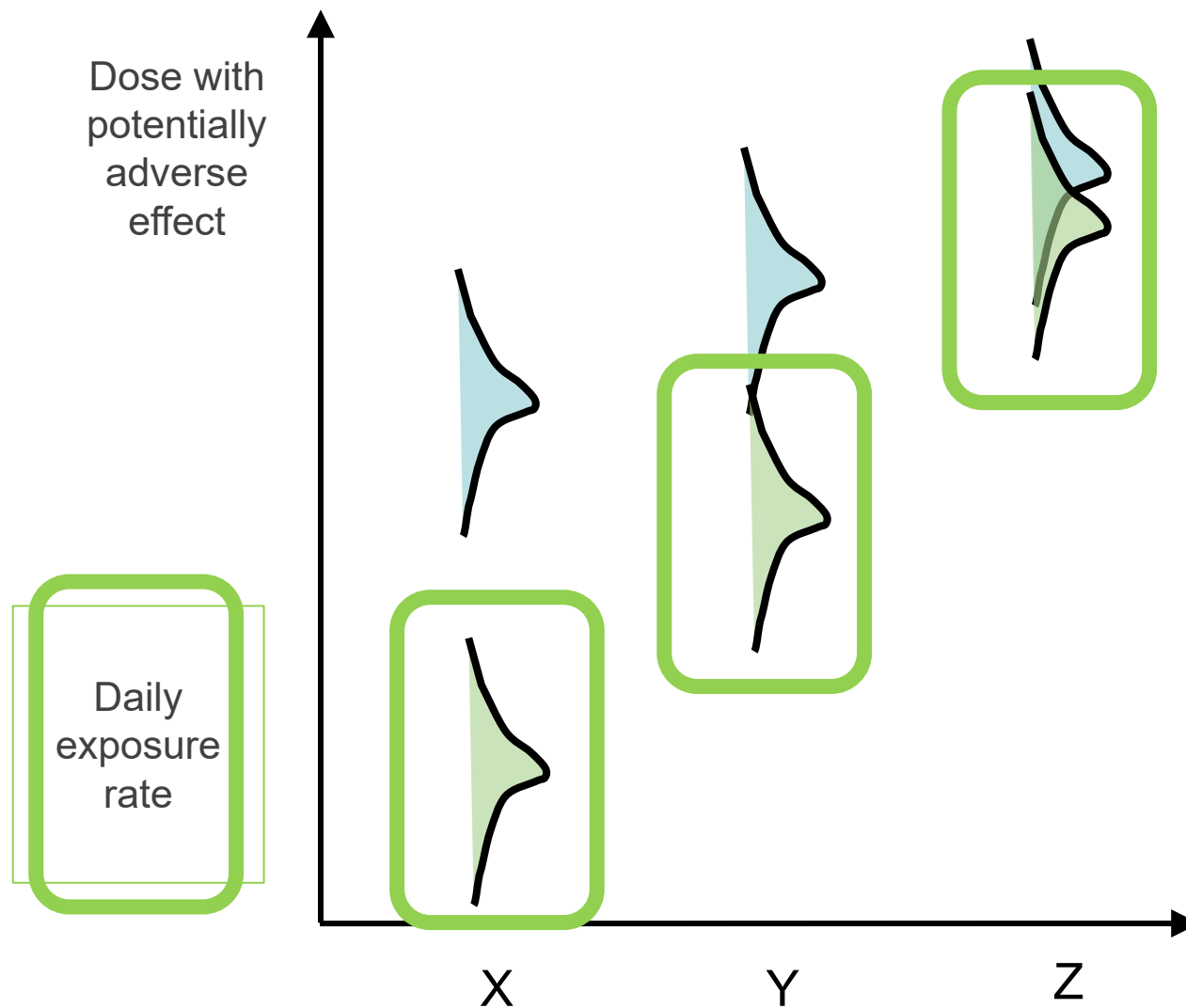
$\text{MOE} = \text{Potentially hazardous dose} / \text{Estimated exposure}$

Higher MOE = less potential risk  
(specific MOE thresholds exist for specific regulatory risk-assessment contexts!)



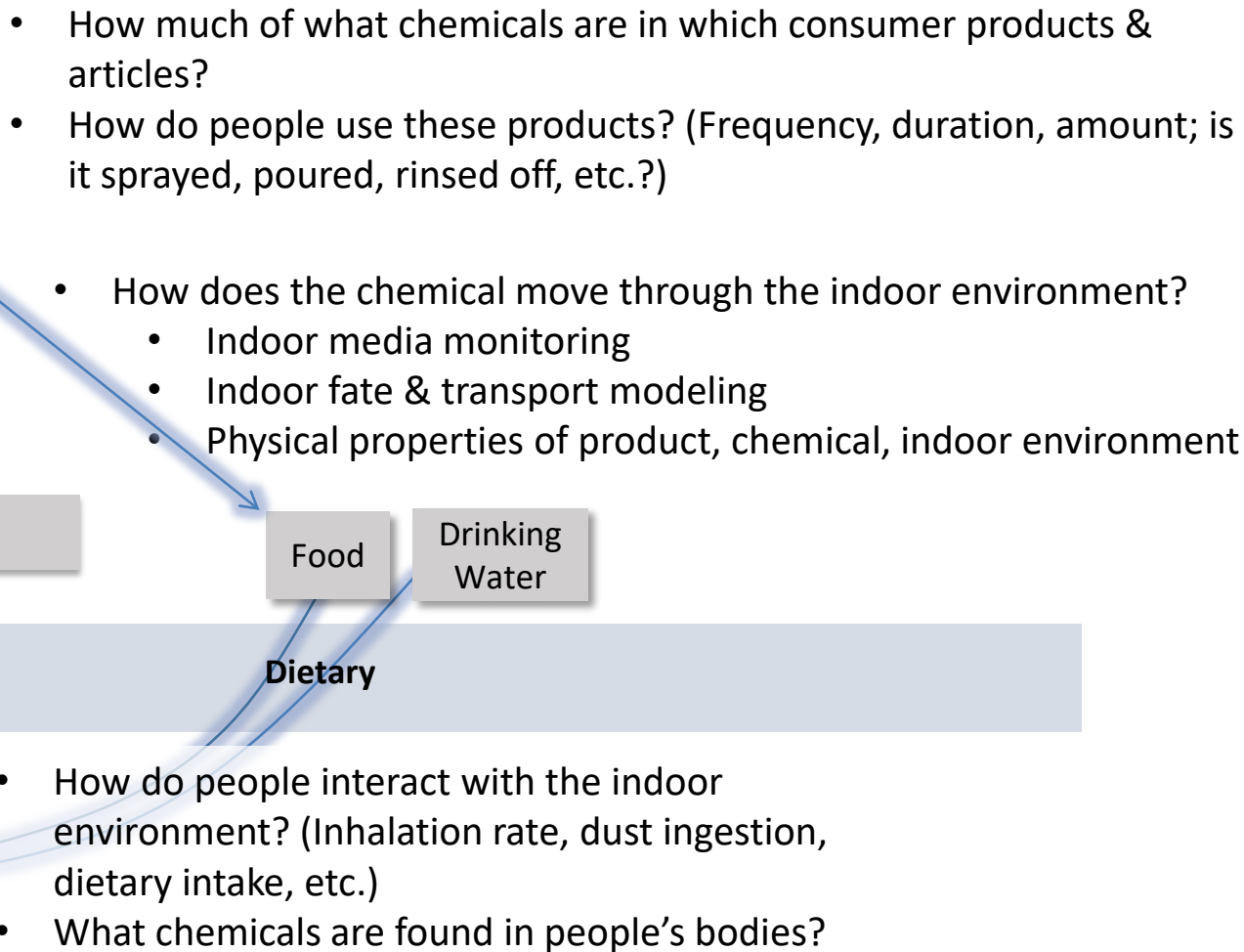


# So how do we get information about exposure?

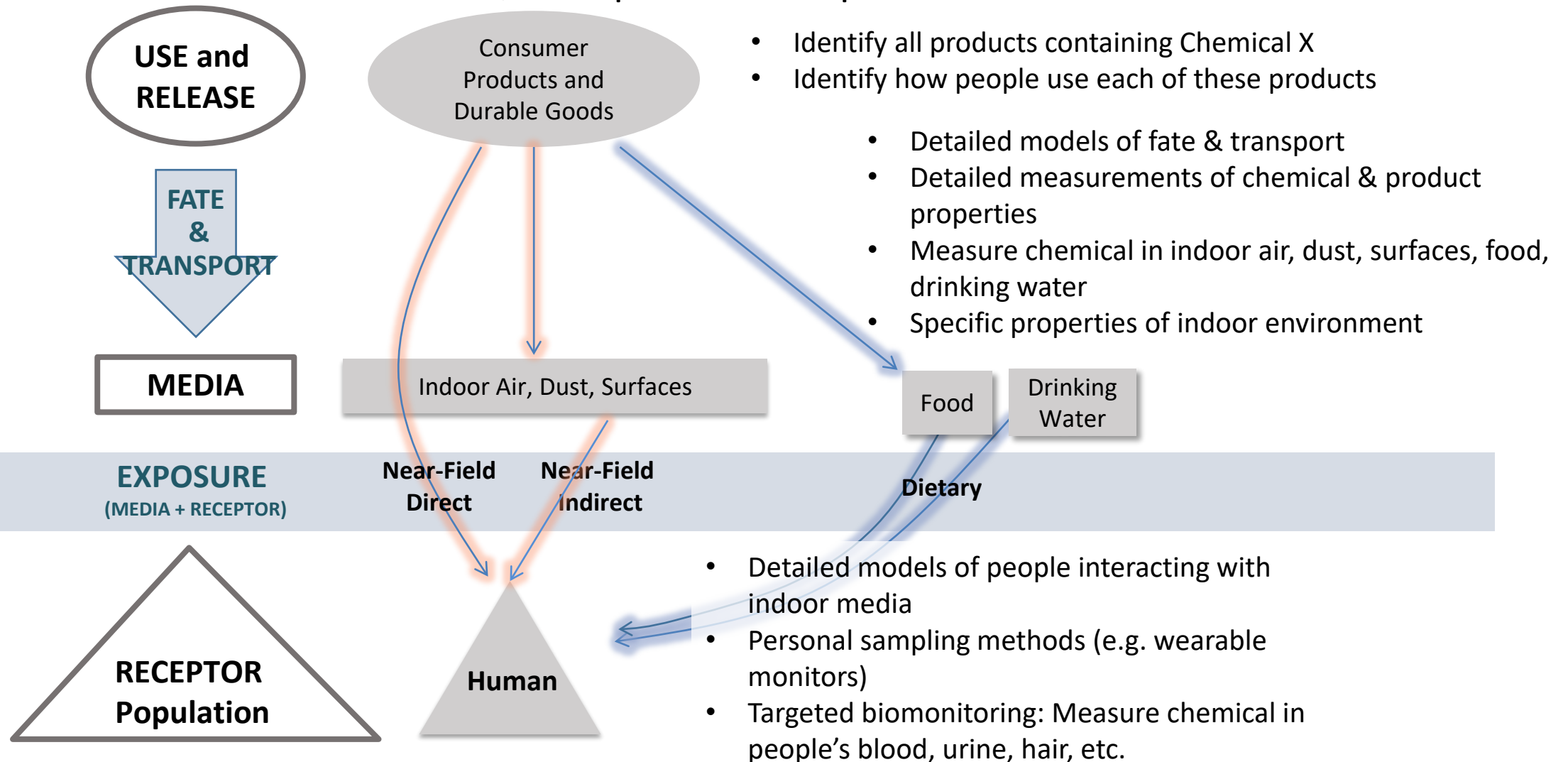


Exposure is assessed by tracing a chemical from its source (where it is released) to where a “receptor” (a person, animal, or plant) interacts with it



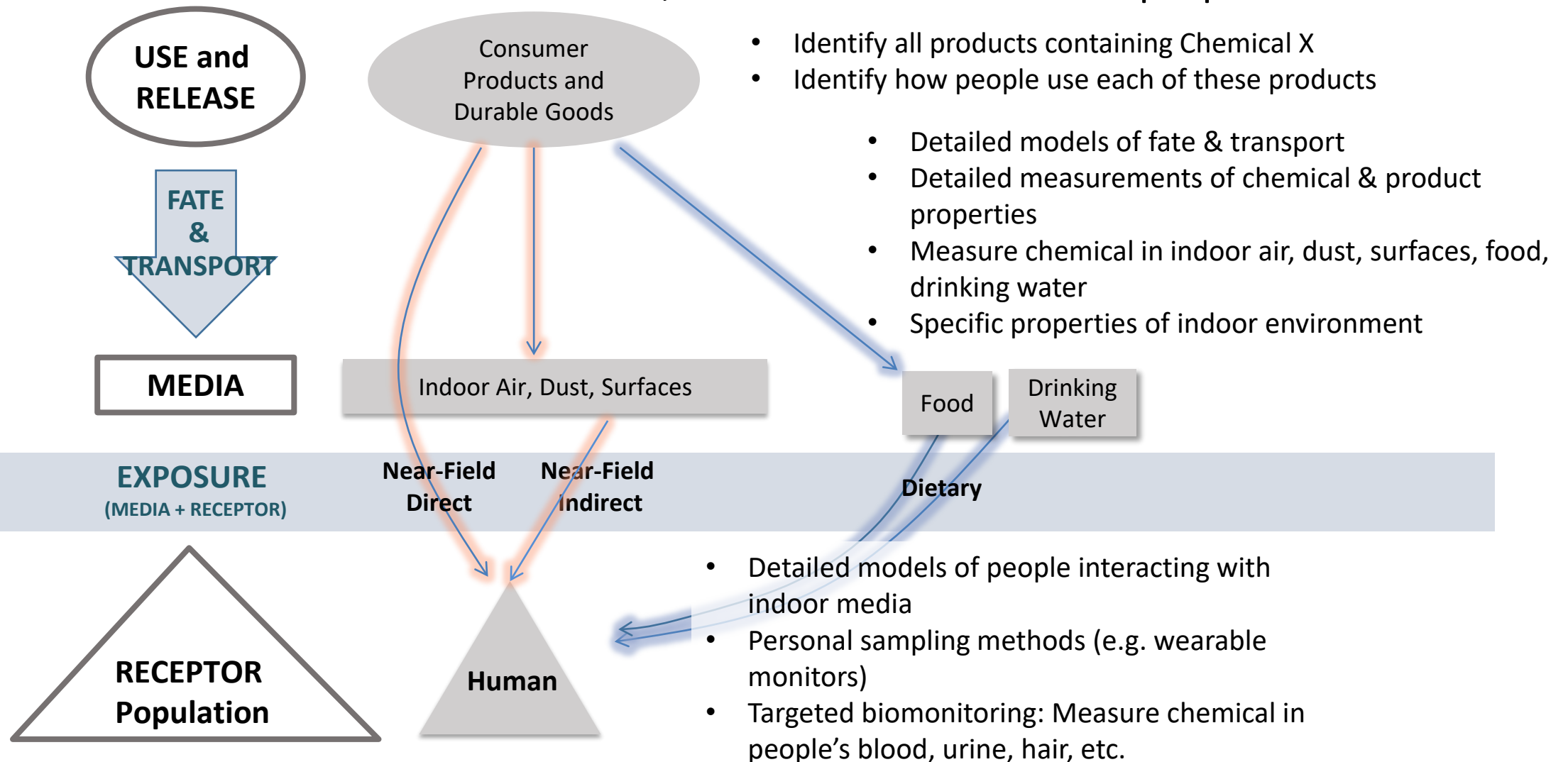


# Traditional exposure assessment: Gather detailed data for one chemical at a time, in specific exposure scenarios





# Difficulty level: Get exposure data for thousands of environmental chemicals, and for the whole population



# New Approach Methodologies (NAMs) for high-throughput exposure science: EPA's ExpoCast project



ExpoCast exposure NAMs aim to inform every part of the source-to-receptor exposure model, in ways that:

- identify and address key pathways of exposure
- can be applied rapidly, to large numbers of chemicals
- leverage existing information to make predictions for data-poor chemicals
- quantify error and uncertainty in predictions
- can be used to prioritize chemicals by potential risk



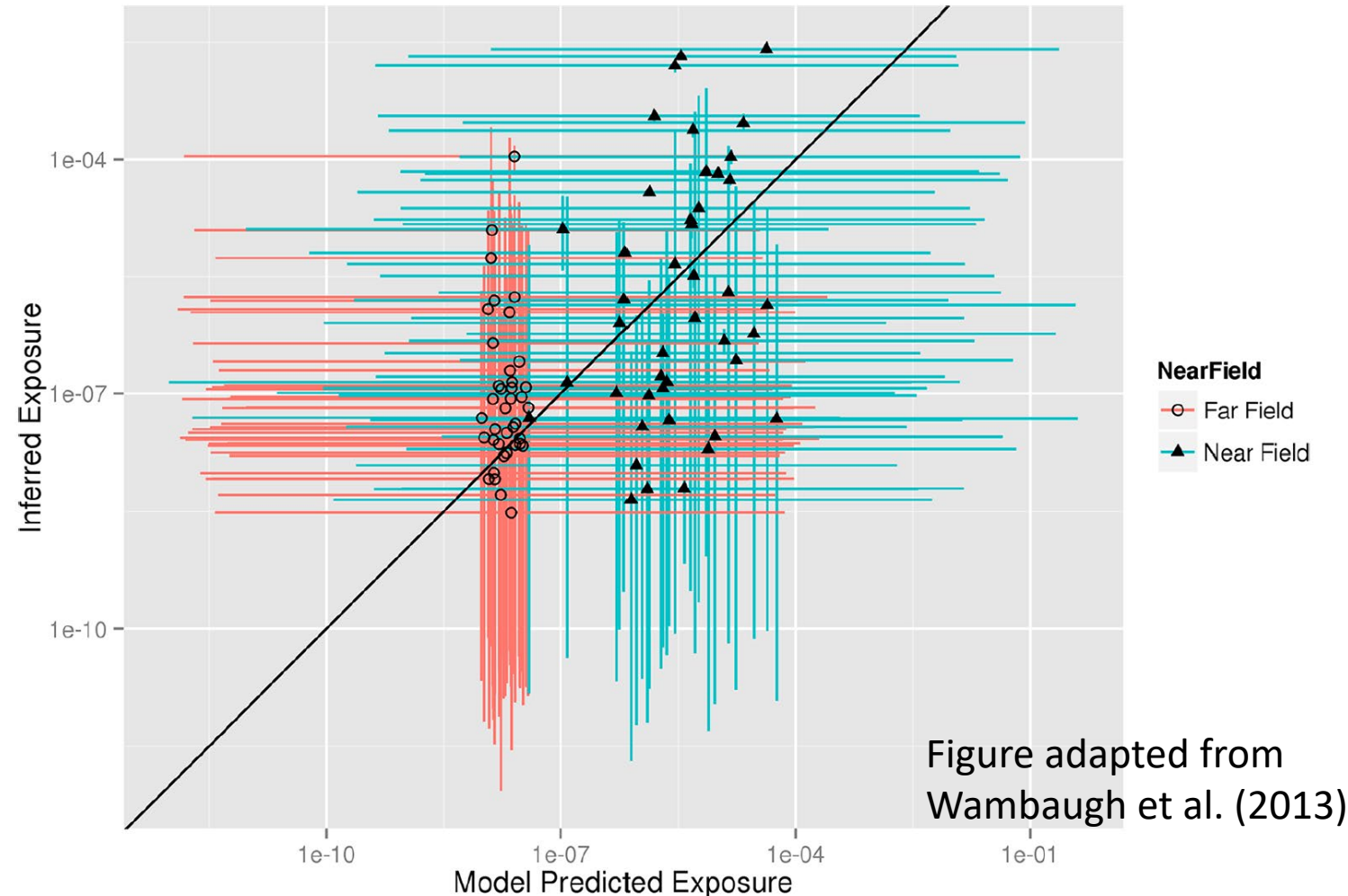


Exposure NAM Class	Description	Traditional Approach
<b>Cheminformatics</b>	Curate & organize existing exposure data for large numbers of chemicals	Tools targeted at single chemical analyses by humans
<b>Machine Learning</b>	Fill data gaps using computer algorithms to make inferences based on existing data	Manual inspection of the data
<b>Non-Targeted Measurements</b>	Screen for hundreds of unknown chemicals in environmental media using advanced analytical & computational chemistry techniques	Targeted (chemical-specific) analyses
<b>HTE Models</b>	Source-to-receptor exposure models that can make predictions rapidly for large numbers of chemicals	Exposure models requiring detailed, chemical- and scenario-specific information
<b>Consensus Modeling &amp; Evaluation</b>	Statistical approaches that use existing exposure data and model results for many chemicals to predict exposure for a new chemical (and evaluate predictive performance of specific HTE models)	Comparison of model predictions to data on a per chemical basis

# A key early ExpoCast result (2013): Consumer product exposures are an important pathway

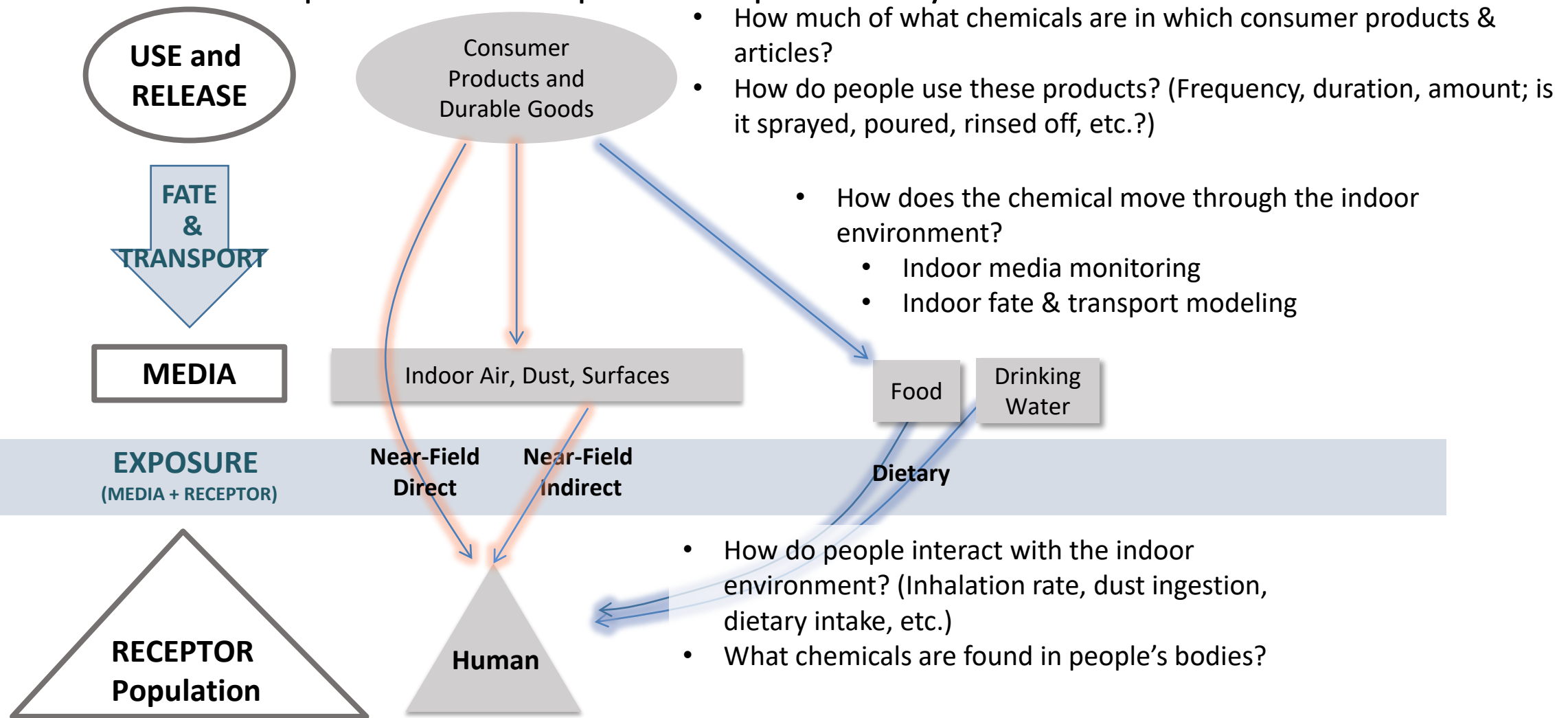
Binary indicator for indoor/consumer use — *all by itself* — explains ~10% of variability in exposure between chemicals.

And chemicals with indoor/consumer use had *higher* exposures.





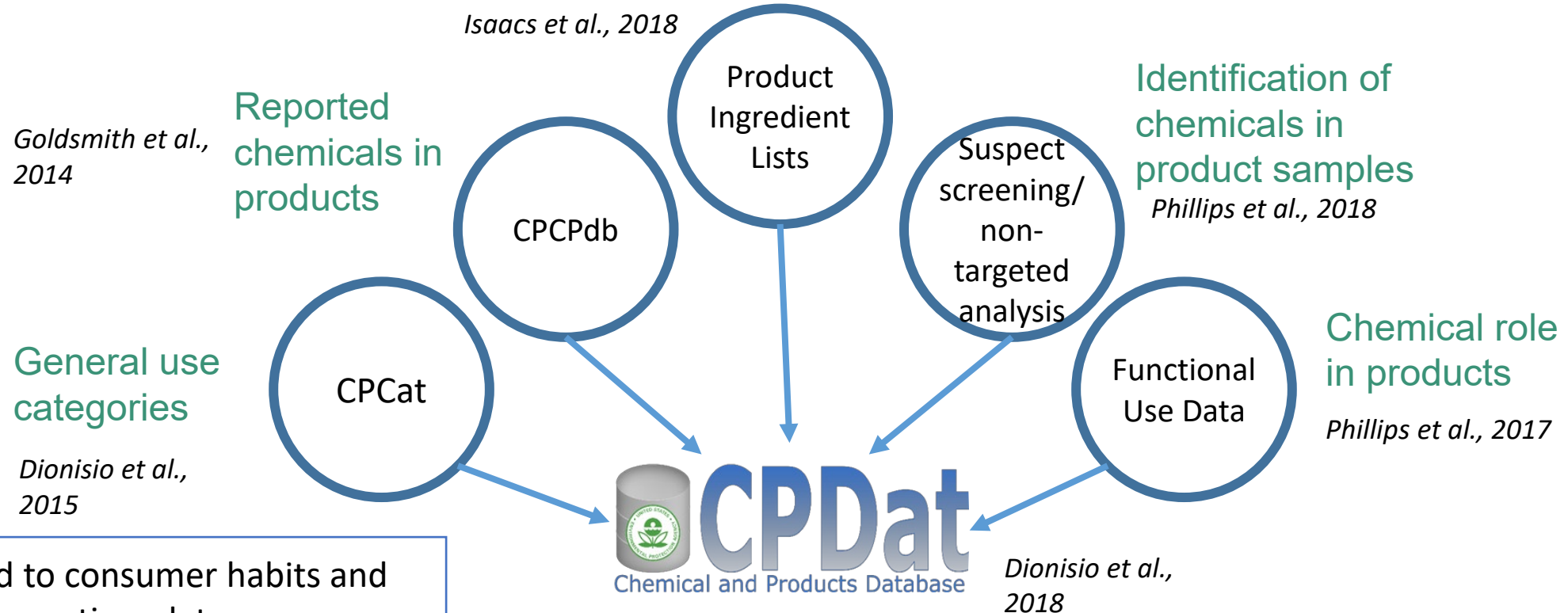
# So, many (but not all!) ExpoCast efforts have focused on consumer products exposure pathways



# Chemical use & release for consumer products: Informatics approach to organizing existing data

Reported or estimated  
weight fraction

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

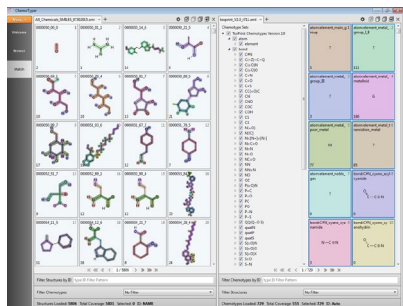
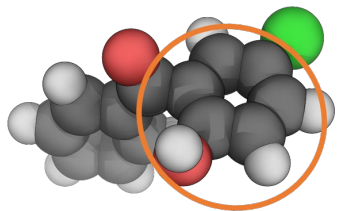


- CPDat integrates heterogeneous data on many chemicals & products from many different sources
- Makes these data machine-readable, batch-searchable
- Rapidly informs chemical use for consumer exposure scenario

# Chemical use: For chemicals without consumer product use data: predict unknown functional uses with machine learning

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

Chemical Structure  
and Property Descriptors



## Chemical Functional Use Database (FUSE)

Positive Examples

Negative Examples



Random Forest Classification  
Models  
(Breiman, 2001)  
with five-fold cross validation

Successful  
Model

Failed  
Model

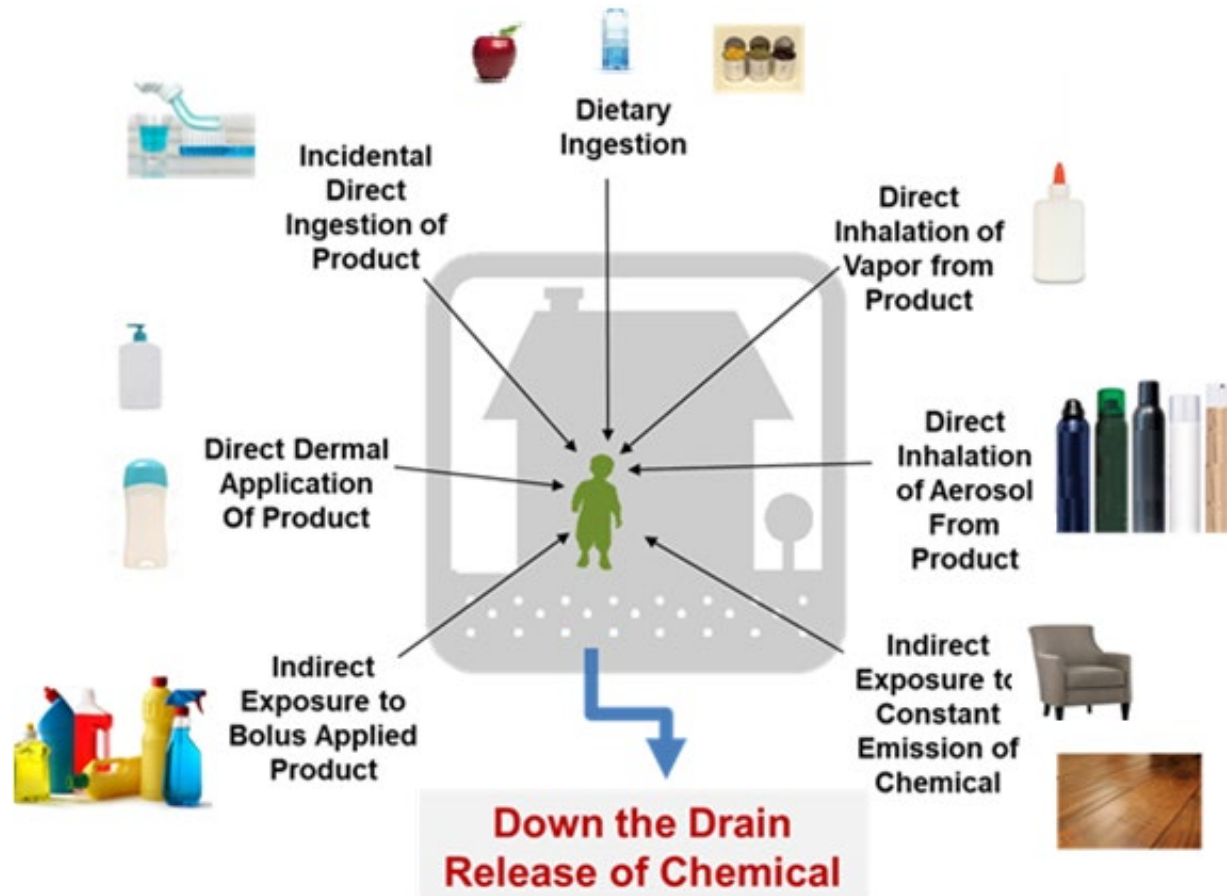


**Probabilistic  
Predictions of  
Potential Chemical  
Uses**

(including whether in  
consumer products)

Phillips *et al.* (2017)

# Modeling exposure from source to receptor: SHEDS-HT: a high-throughput population consumer exposure model (Isaacs et al., 2014)



- Sources chemical use data from CPDat
- Sources existing data on population variability in consumer habits & practices from literature
- Sources data on population variability in diet from CDC NHANES (national dietary survey data)  
(<https://www.cdc.gov/nchs/nhanes/index.htm>)
- Includes existing data on population daily activities from EPA CHAD  
(<https://www.epa.gov/fera/consolidated-human-activity-database-chad>)
- Available as R package 'ShedsHT'  
<https://github.com/HumanExposure/SHEDSHTRPackage>



# Non-Targeted Analysis: Which chemicals are found in consumer products? In indoor environmental media? In humans?

(Sobus et al., 2018; Ulrich et al., 2019)

## Source and Release

*Pilot: 20 Consumer Product Categories*



Phillips et al., *Env. Sci. Tech.* 2018

*Recycled Consumer Materials*



Lowe et al., 2018

*Consumer Product Emissions from Different Substrates*



## Fate and Transport

*Residential Air*



*Residential Dust*



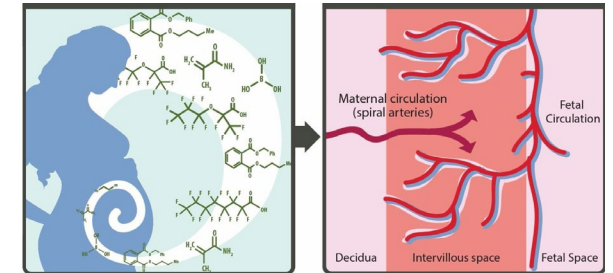
Rager et al., *Env. Int.*, 2016

## Exposure

*Pooled Human Blood*

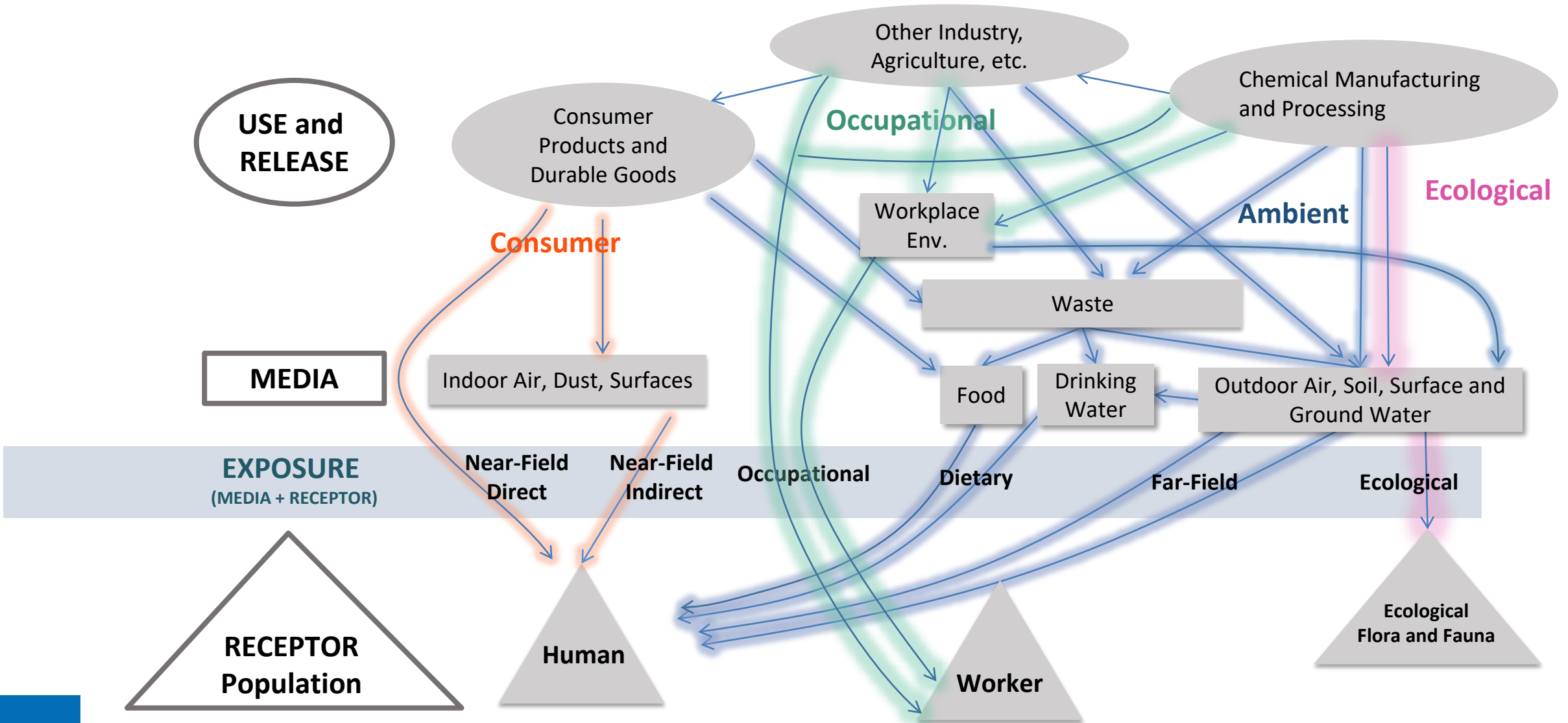


*Human Placenta*



Rager et al., *Repro. Tox.*, 2020

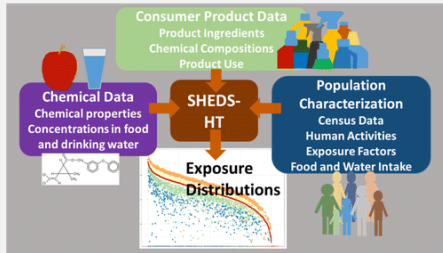
# What about exposure pathways *other* than consumer/residential?



# High Throughput Exposure (HTE) models can predict exposures via key pathways (for chemicals with enough data to parameterize models)

## Consumer (Near-Field) Pathways

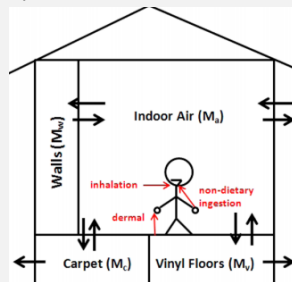
*SHEDS-HT* (Isaacs et al., 2014)



*RAIDAR-ICE* (Li et al., 2018)

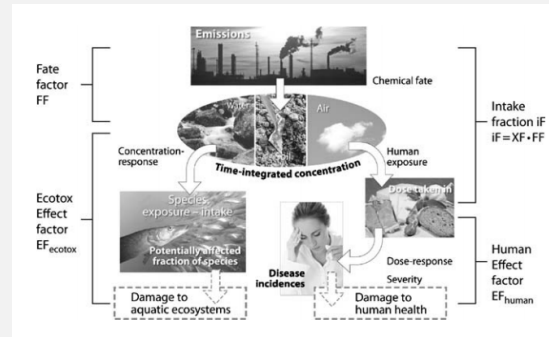


*FINE* (Shin et al., 2015)

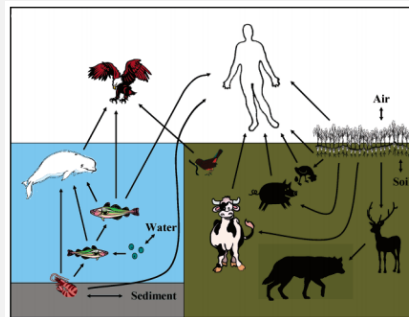


## Ambient (Far-Field) Pathways

*UseTox* (Rosenbaum et al., 2008)



*RAIDAR* (Arnot et al., 2006, 2008)

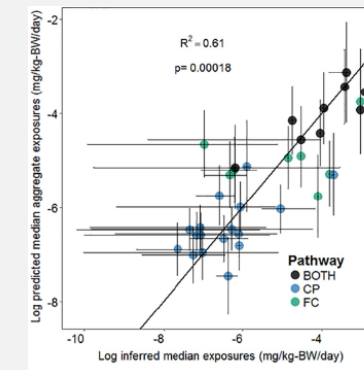


## Dietary Pathways

*UseTox* (Rosenbaum et al. (2008)



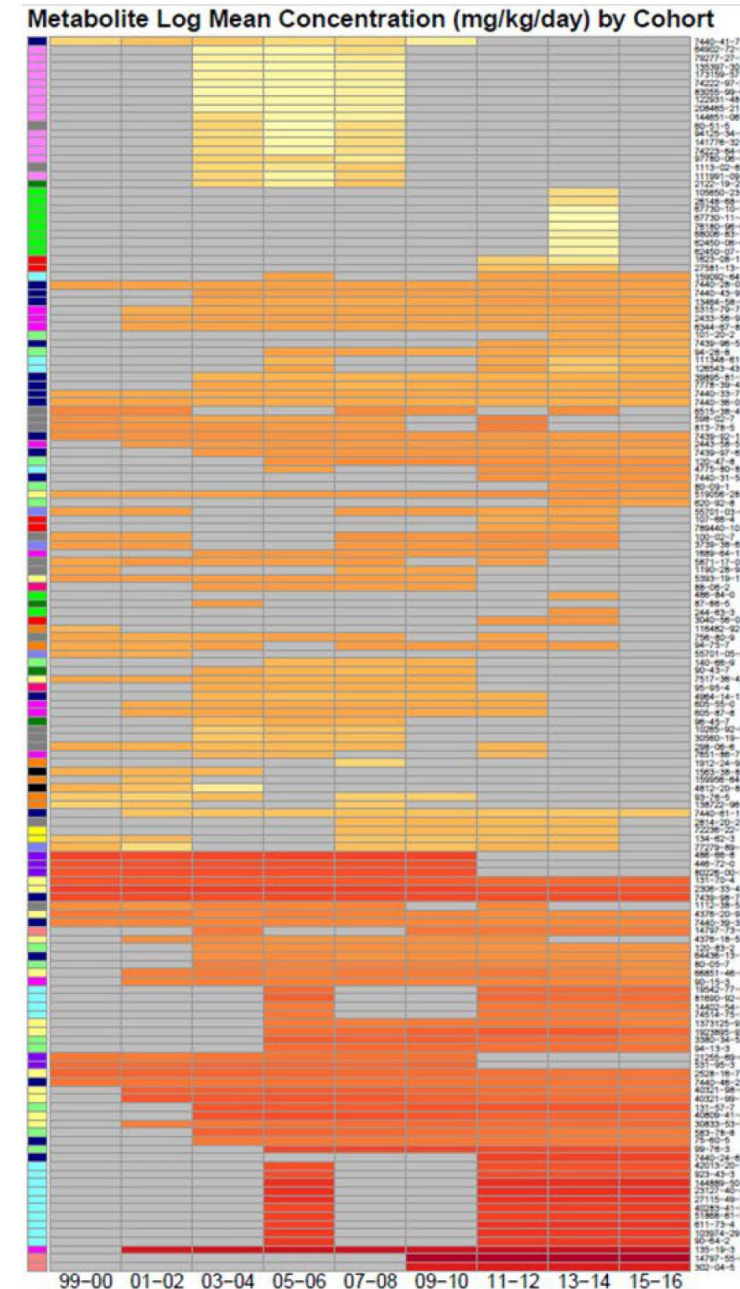
*SHEDS-HT* (Biryol et al., 2017)





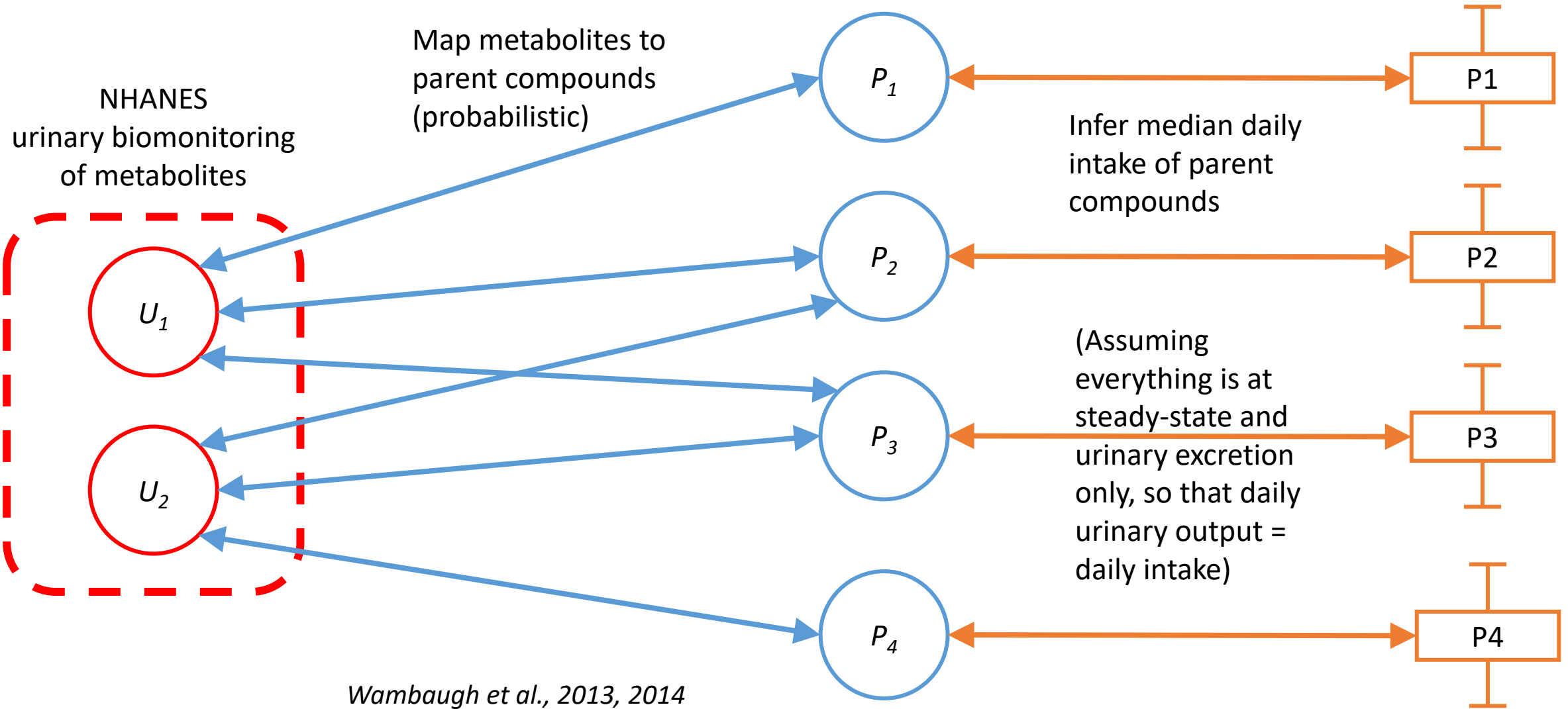
# Aggregate exposures (over *all* pathways) can be inferred from population exposure biomonitoring

- Exposure biomonitoring measures internal body levels of various chemicals of interest, or their metabolites
  - e.g. in blood, urine, hair, breastmilk, etc.
- A key source of exposure biomonitoring data is CDC NHANES (National Health & Nutrition Examination Survey)
  - Large-scale, nationally-representative survey of US population
  - 2-year cycles: starting in 1999, most recent published data 2016
- NHANES gathers various health & nutrition data
  - Previously mentioned: dietary intake survey (used in SHEDS-HT model)
- Including urine levels of 151 metabolites (mapping to 179 possible parent chemicals) [see figure at right!]
- All data publicly available (anonymized) at <https://www.cdc.gov/nchs/nhanes/index.htm>



Wambaugh et al., 2013, 2014; Stanfield et al., in prep  
Figure courtesy of Dr. Zachary Stanfield

# ExpoCast work: Bayesian inference of external exposures from internal biomonitoring data

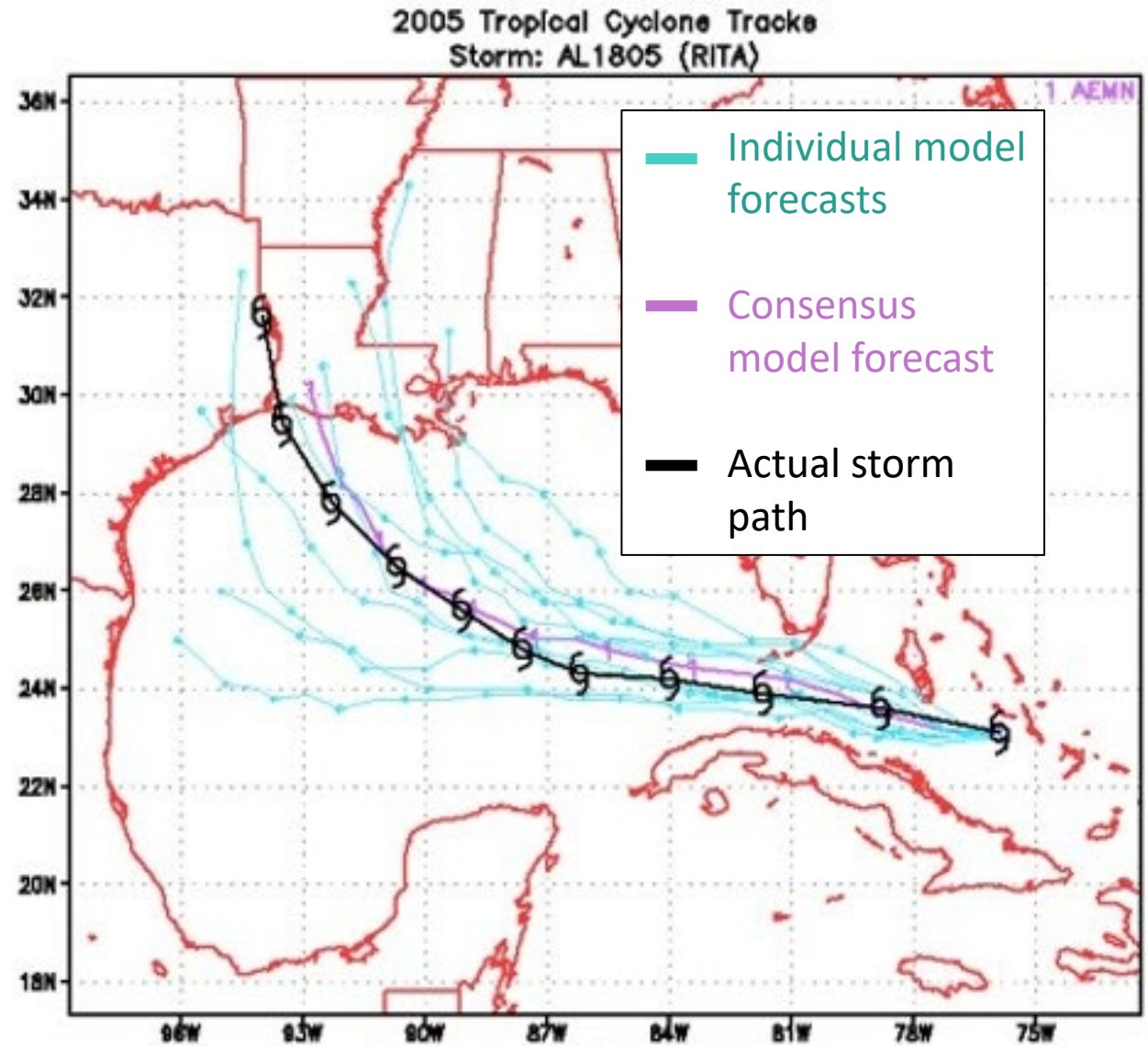


Wambaugh et al., 2013, 2014  
Stanfield et al., 2021



We can integrate all of these exposure models and data sources into a *consensus model* for aggregate exposure!

- Consensus models may be familiar from weather forecasting: e.g. predicting hurricane paths
- Average together the individual model predictions
- Individual models can be *weighted* to correct for model biases
  - e.g. a model that usually predicts a path too far west
  - e.g. a model that usually over-predicts storm intensity
- We can make an analogous consensus model for aggregate human daily intake!



# SEEM3: A consensus model for aggregate exposure

SEEM3 = Systematic Empirical Evaluation of Models, version 3

Ring et al. (2019)

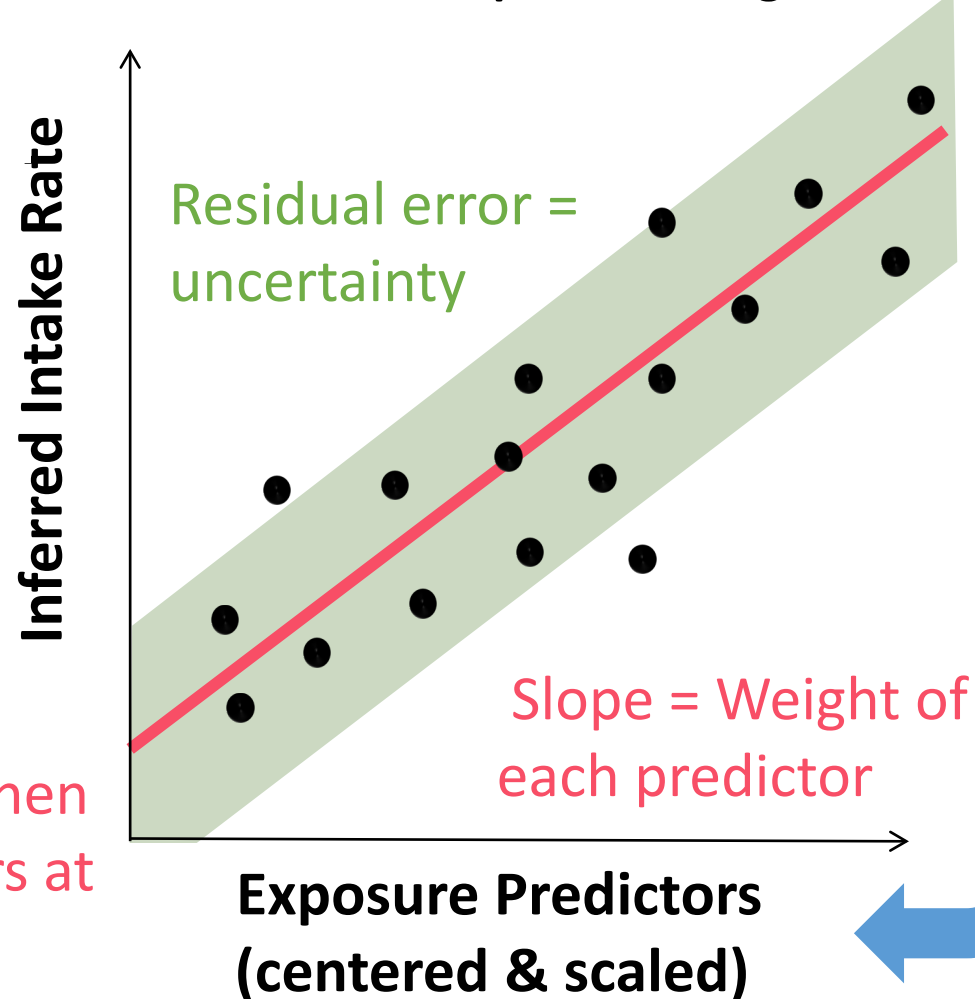
Train model on inferred  
exposures from NHANES  
biomonitoring data

$$\boxed{P1} = \bullet$$

**Bayesian inference** =  
*Probabilistic* estimates of  
**intercept**, **slopes**, and  
**uncertainty**

**Intercept** =  
Exposure when  
all predictors at  
mean value

**SEEM3 is a multiple linear regression!**



## Exposure Predictors:

- Predictions of HT exposure models (USETox, RAIDAR, FINE, SHEDS-HT...)
- Chemical production volume (U.S.)
- Existing EPA pesticide exposure assessments
- Presence on Stockholm Convention list of banned persistent organic pollutants

Missing predictor data:  
Impute mean

# SEEM3 includes pathways of exposure

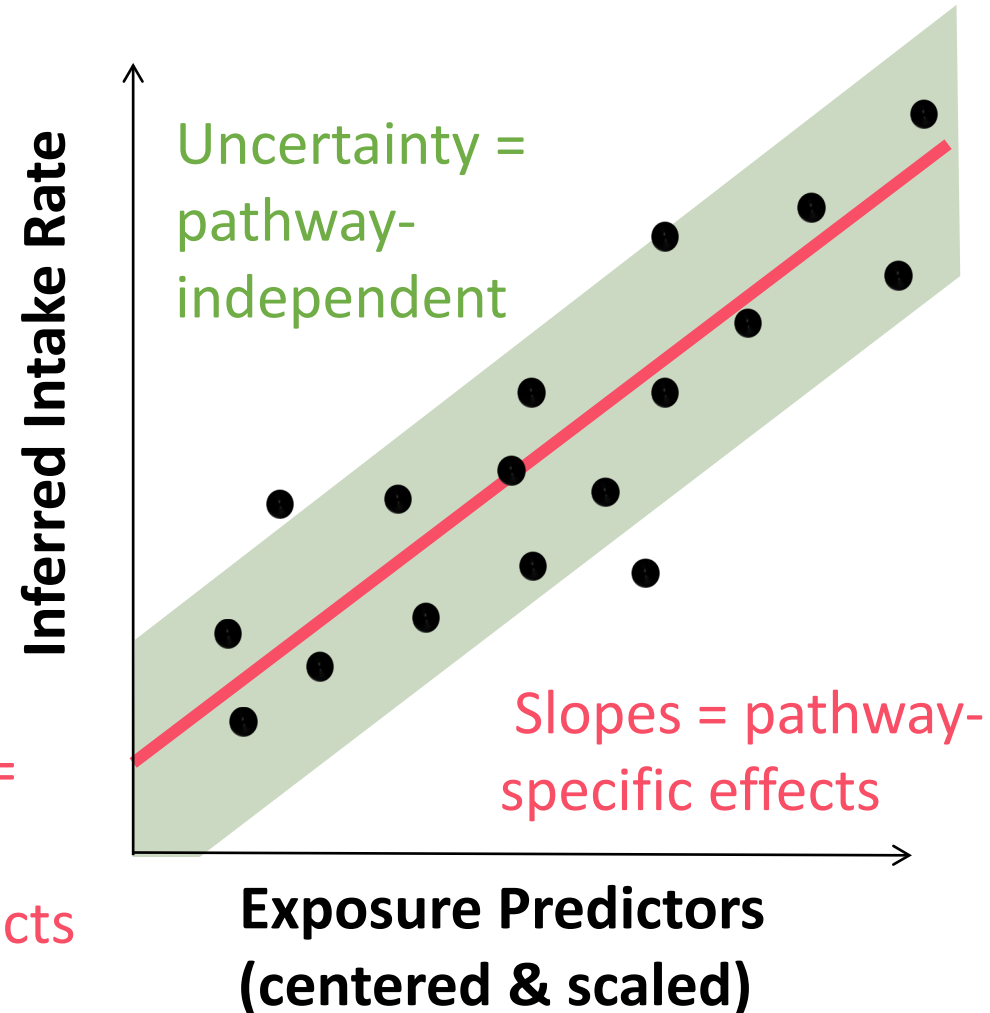
Ring et al. (2019)

Machine-learning model (random forest) predicts **exposure pathway probability** for each chemical:

- Consumer
- Dietary
- Industrial
- Pesticide

based on chemical structure & properties

Intercepts =  
pathway-  
specific effects

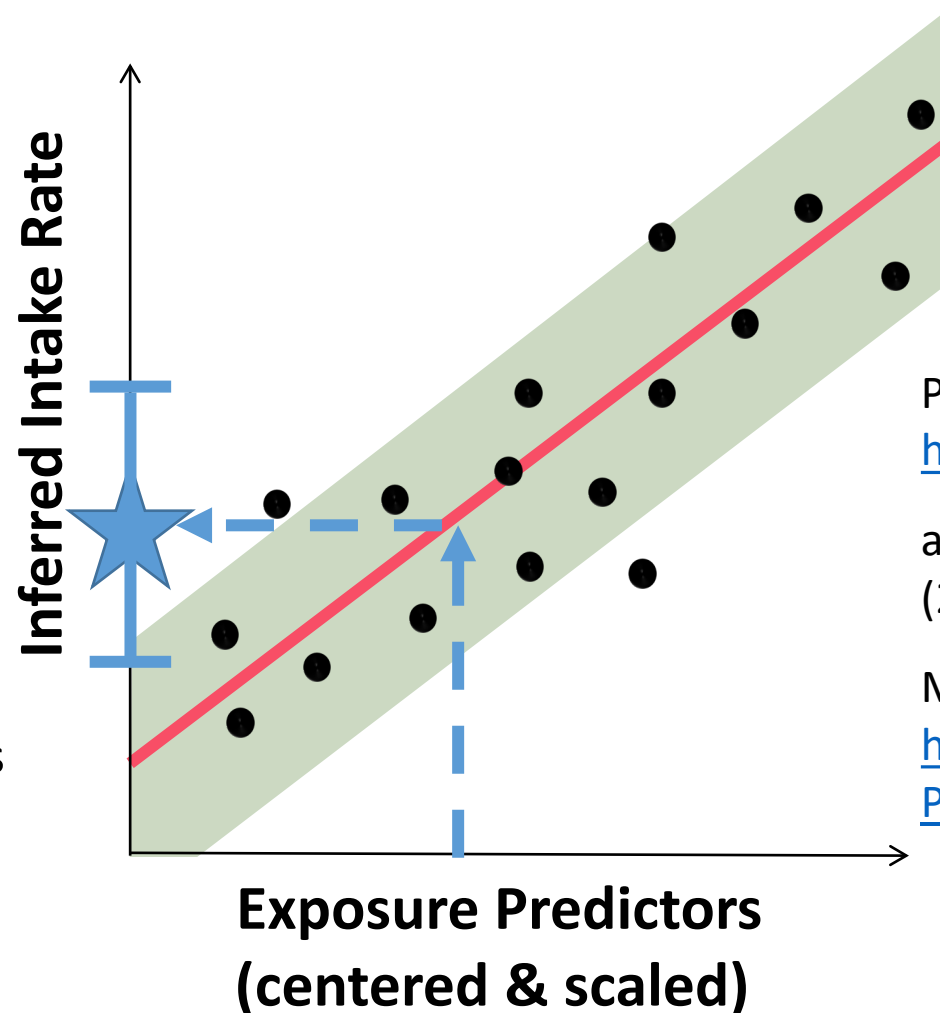


Pathway-specific weights (slopes) for each predictor = predictive strength of that predictor for that pathway

(hence the “evaluation of models” in the SEEM3 name)

# SEEM3 can *predict* median exposures for data-poor chemicals – and quantify uncertainty in the predictions

There are SEEM3 predicted median exposures for 687,359 chemicals!  
(Every compound with a structure in DSSTox library as of 2018)

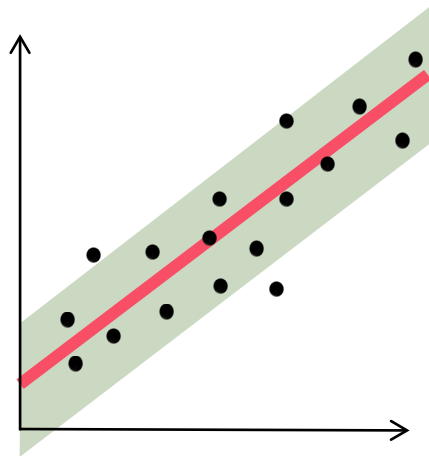
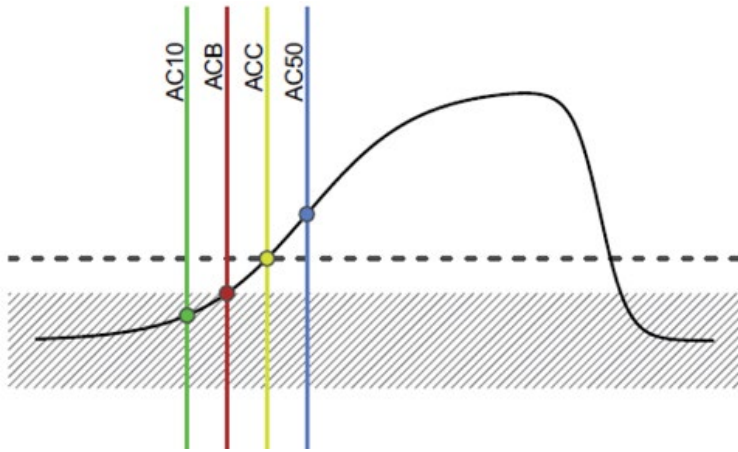


Predictions available on  
<https://comptox.epa.gov/dashboard>  
and as Supplemental Material to Ring et al. (2019)

Model available as R package:  
<https://github.com/HumanExposure/SEEM3R>  
Package

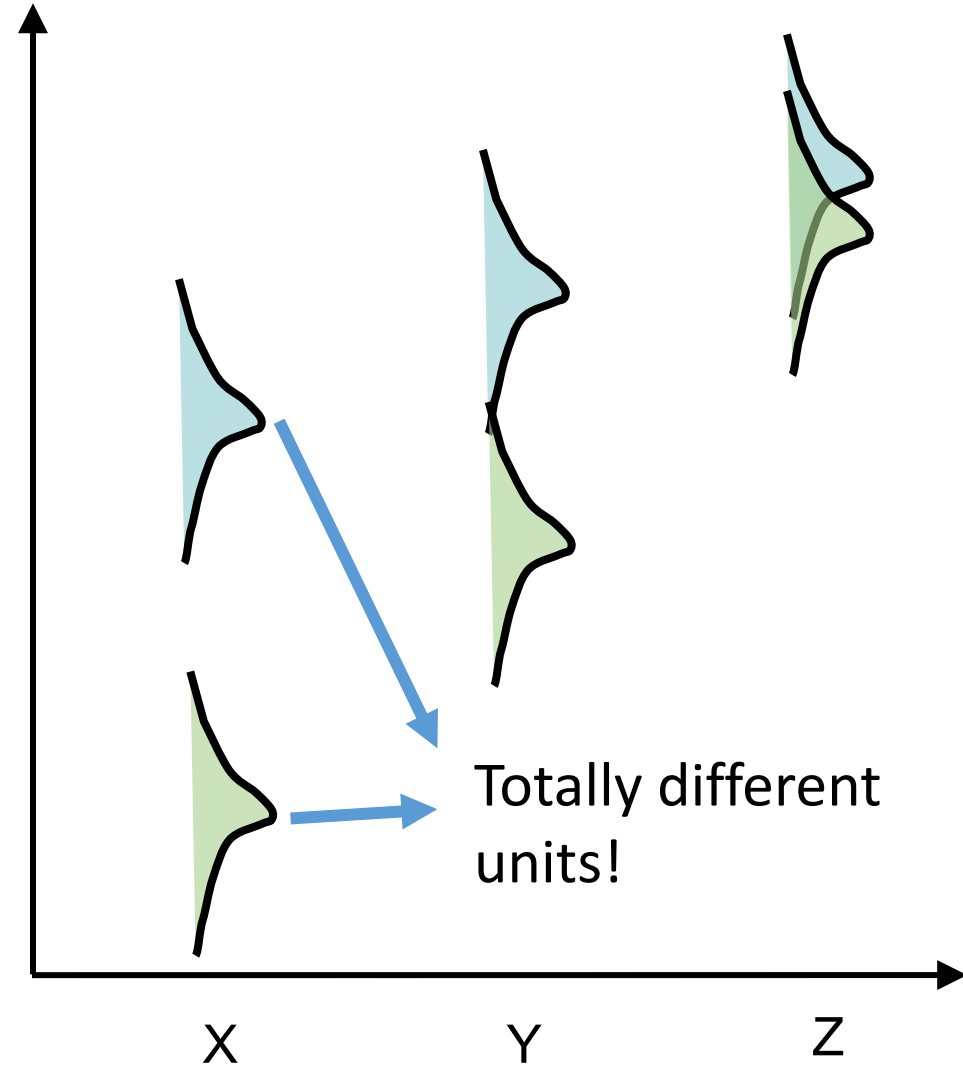
# Web demonstration: How to find exposure data and predictions on the CompTox Chemicals Dashboard

So, we can predict exposures using all of these clever computational tools. But how does that help us when we have *in vitro* hazard data only in the form of *in vitro* bioactive concentrations?



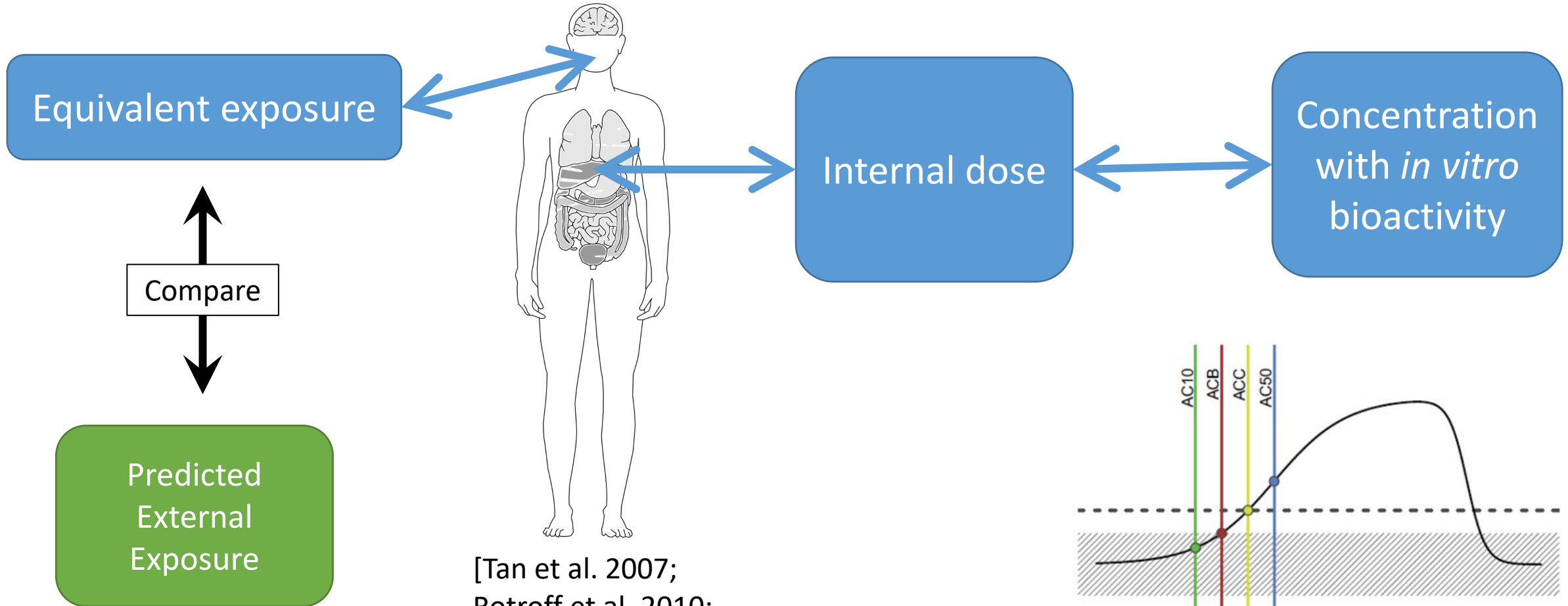
Concentration  
bioactive *in vitro*  
(uM)

Daily  
exposure  
rate  
(mg/kg/day)



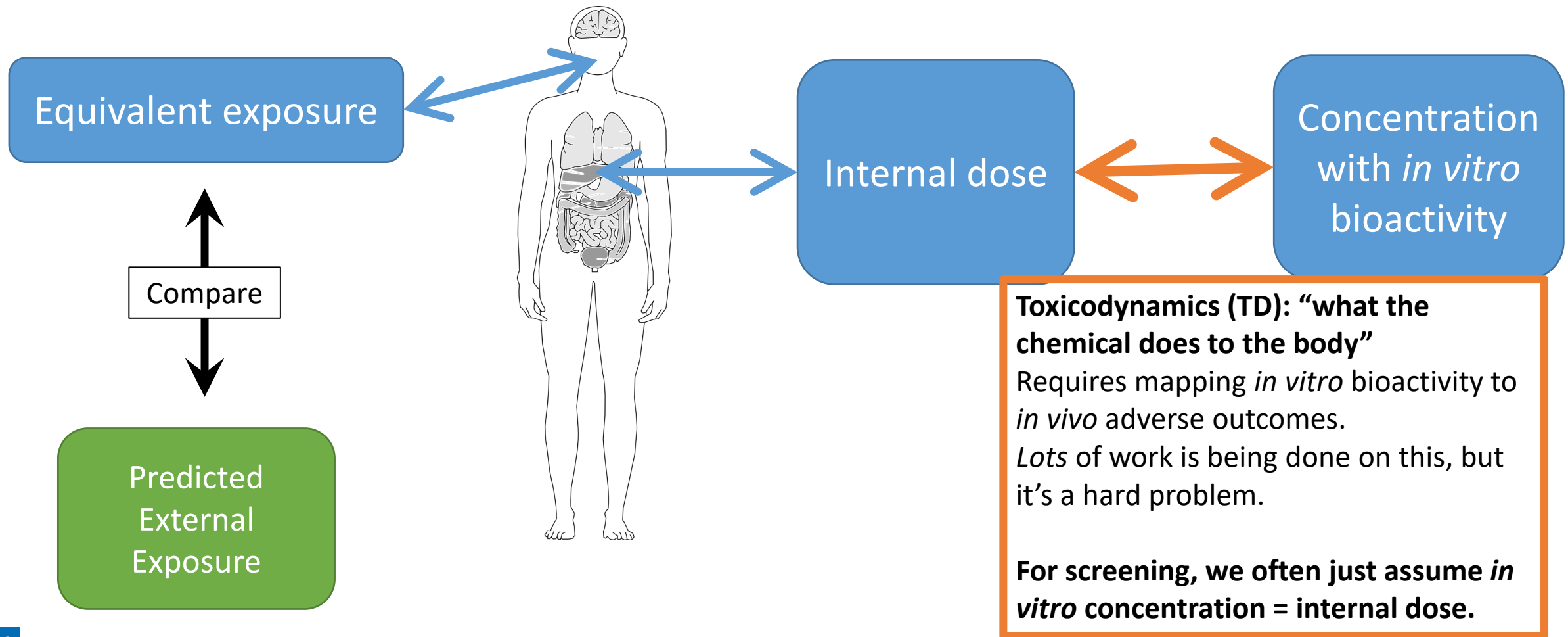


Need to link *in vitro* concentrations to *in vivo* exposures:  
*in vitro-in vivo* extrapolation (IVIVE) —  
and we need to do IVIVE for thousands of chemicals and  
the whole population!

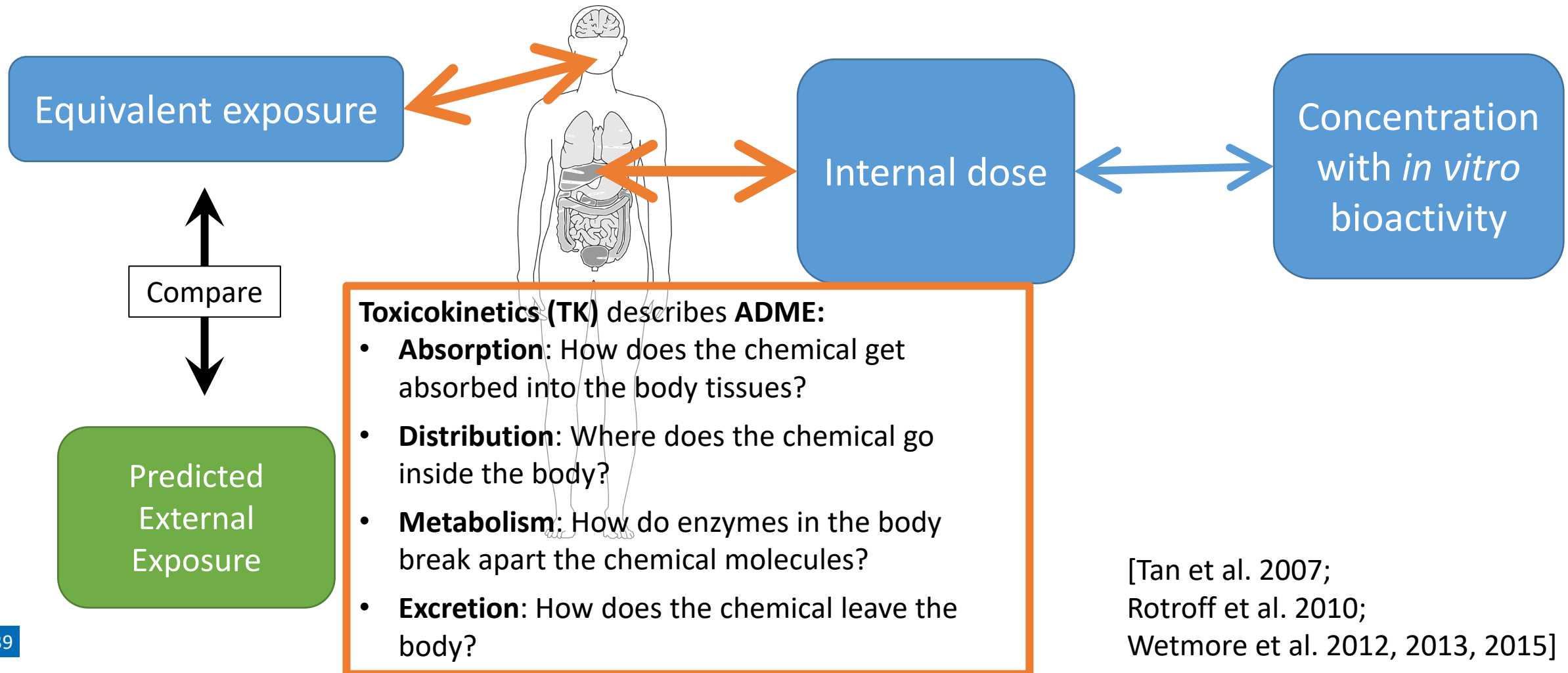


[Tan et al. 2007;  
Rotroff et al. 2010;  
Wetmore et al. 2012, 2013, 2015]

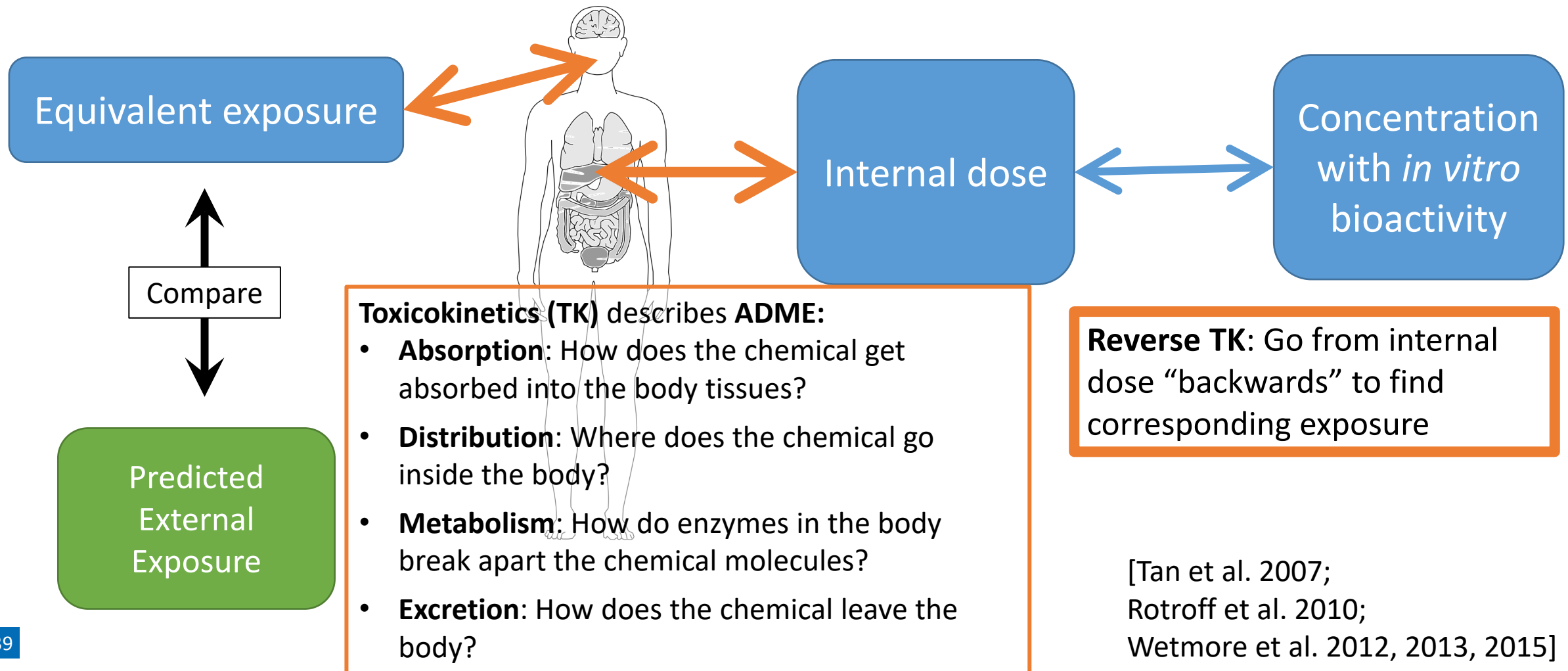
# Mapping between *in vitro* bioactive concentration and internal dose is a **toxicodynamics** problem



# Mapping between internal dose and external exposure is a **toxicokinetics** problem

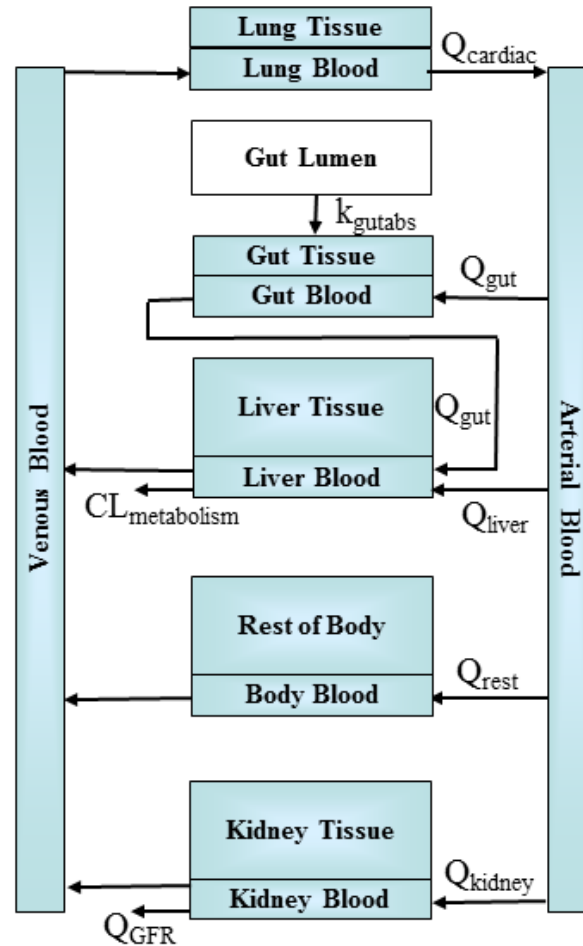
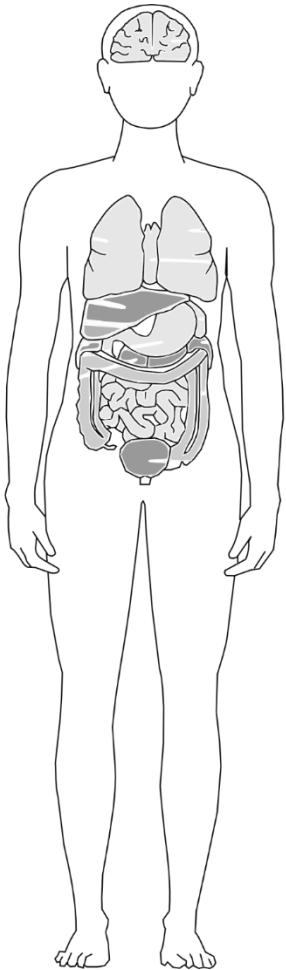


# Mapping between internal dose and external exposure is a **toxicokinetics** problem

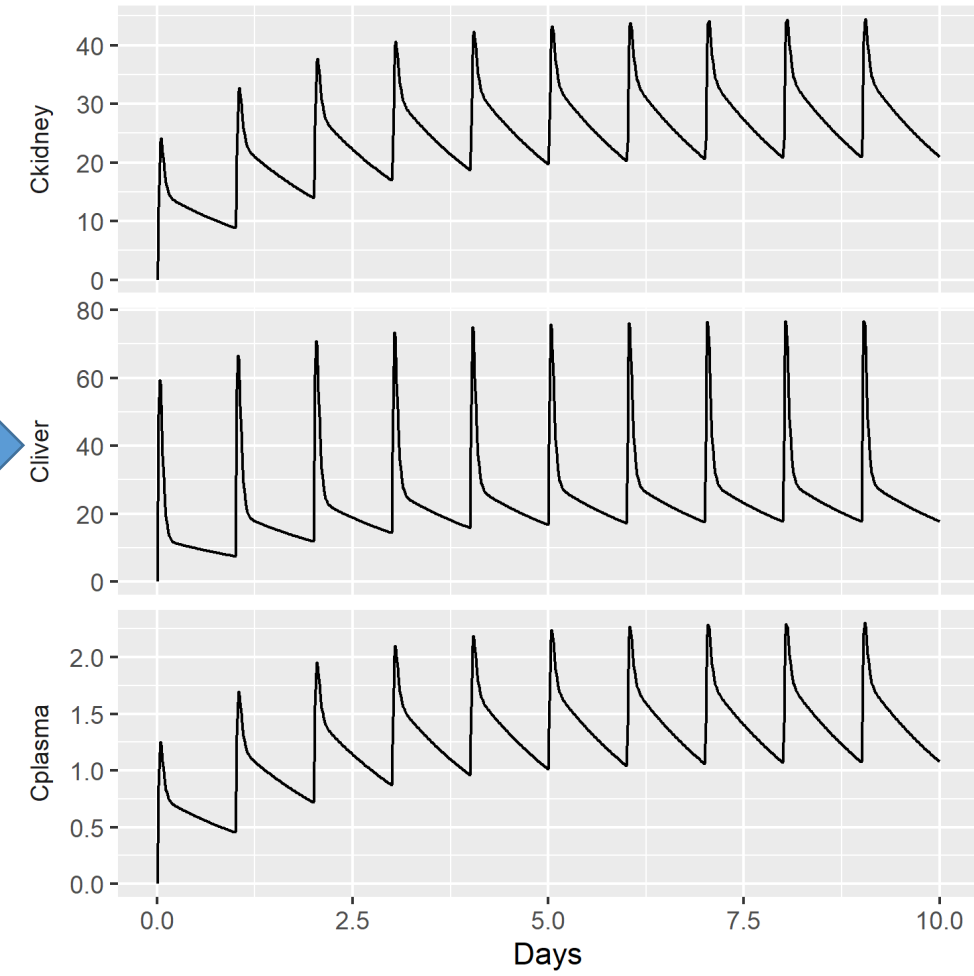


# TK models describe ADME mathematically

Body as mass-balance system  
Defined by parameters describing ADME



Concentration vs. time in each compartment



*High-throughput* IVIVE (rapid, for thousands of chemicals) requires *high-throughput* TK (HTTK)

### Characteristics of HTTK model:

- **Generic:** same model structure can be applied to all chemicals
- **Minimal chemical-specific TK parameters**
  - Only describe the most important chemical-specific ADME processes
  - Can only run model for chemicals where we know these parameters — so the fewer chemical-specific parameters, the more chemicals we can run
- **Chemical-specific TK parameters that can be measured *in vitro* or predicted *in silico***, rather than having to be measured *in vivo*
  - Use existing *in vitro* experimental methods to measure TK parameters – pharmaceutical industry has been working on this for years
- **Not too computationally intensive:** Feasible to solve rapidly for thousands of chemicals
- **Allows quantification of uncertainty & variability** in its predictions

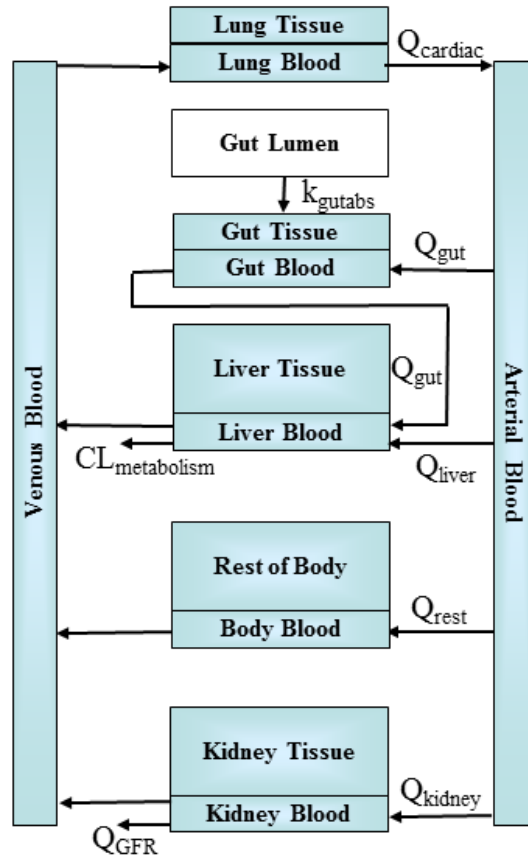


# High-throughput TK (HTTK)

## Generic physiologically-based TK (PBTk) model

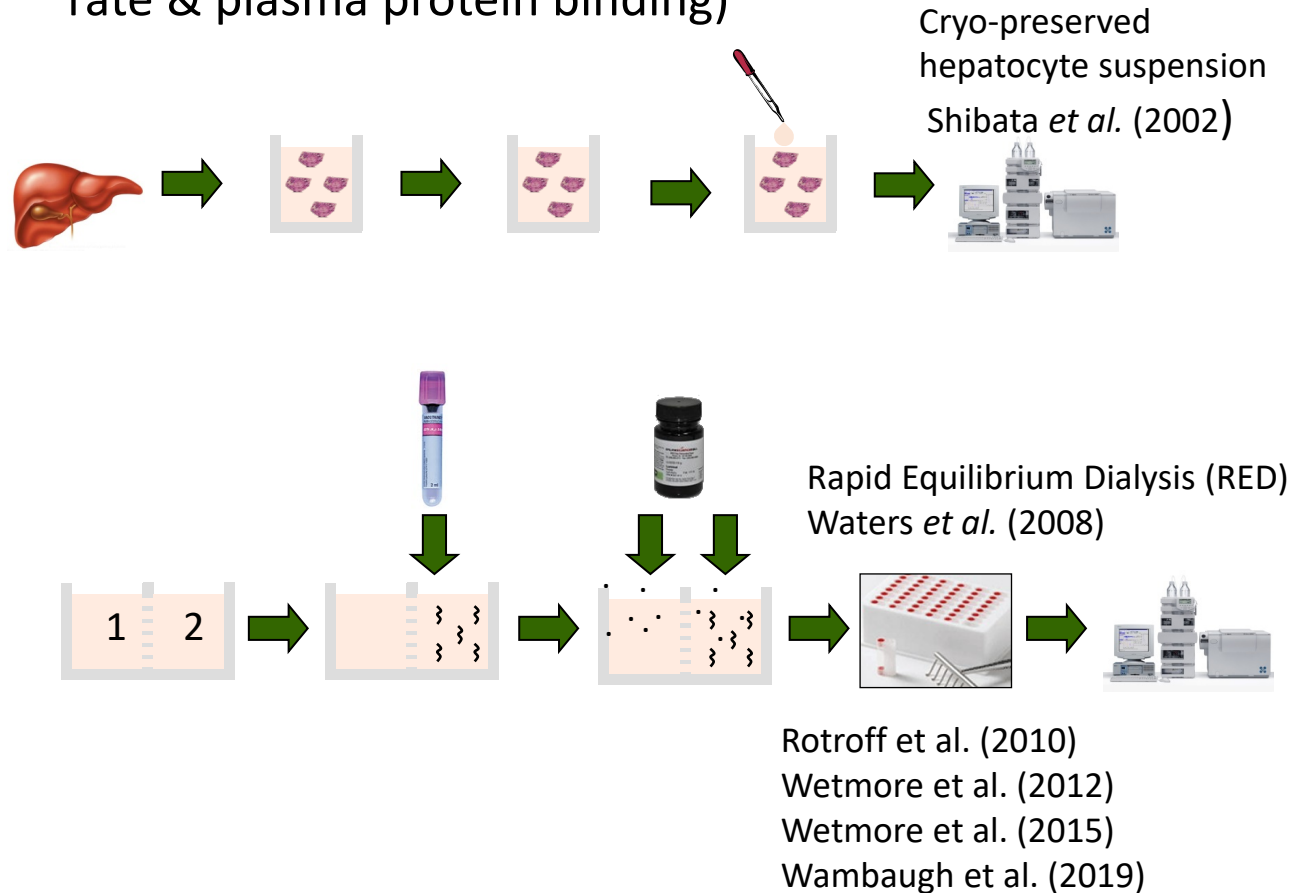
Assume clearance  
via first-order  
hepatic metabolism  
& passive renal  
filtration

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



+

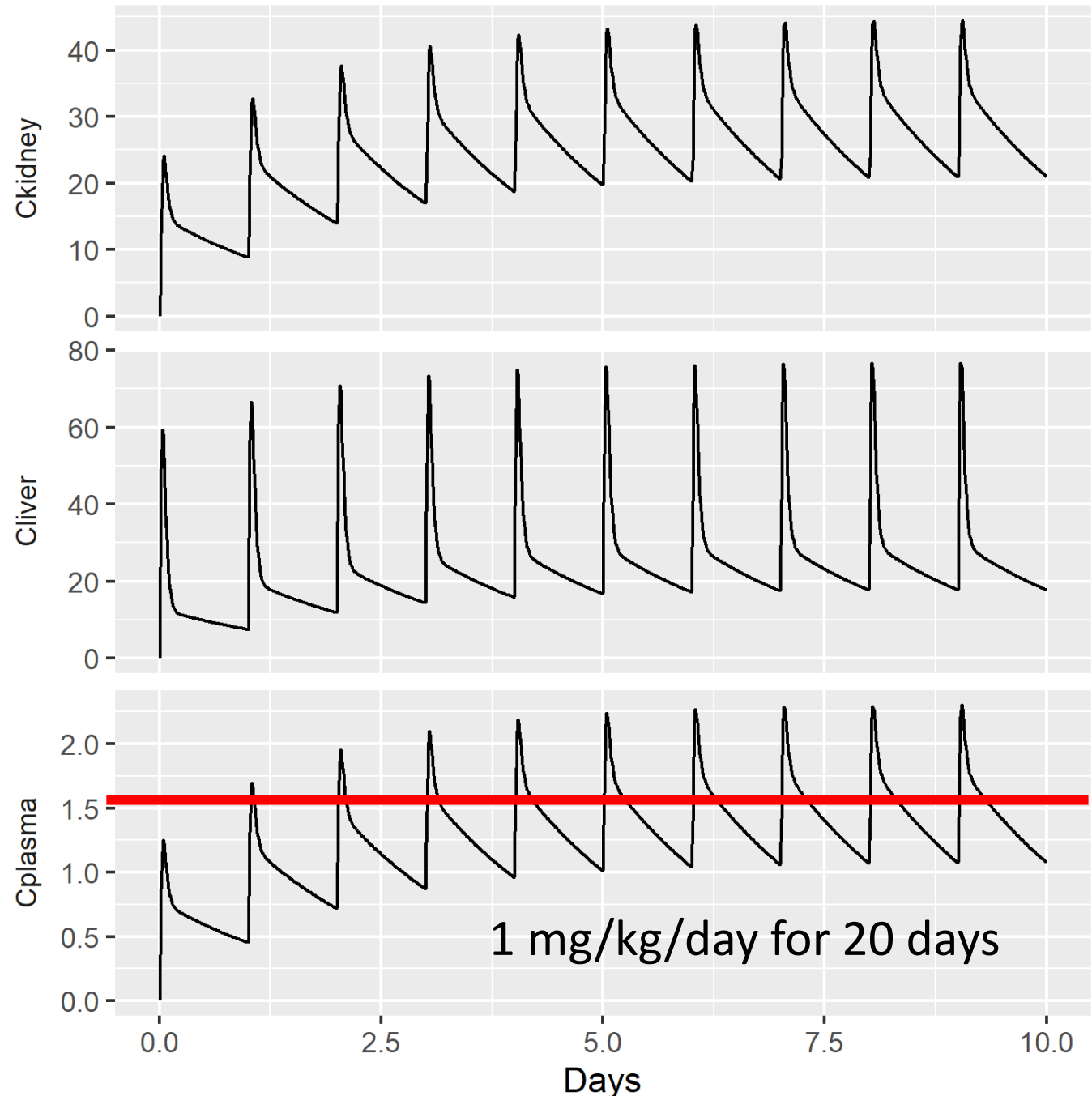
*In vitro* measurements of the minimal chemical-specific TK model parameters (hepatic clearance rate & plasma protein binding)



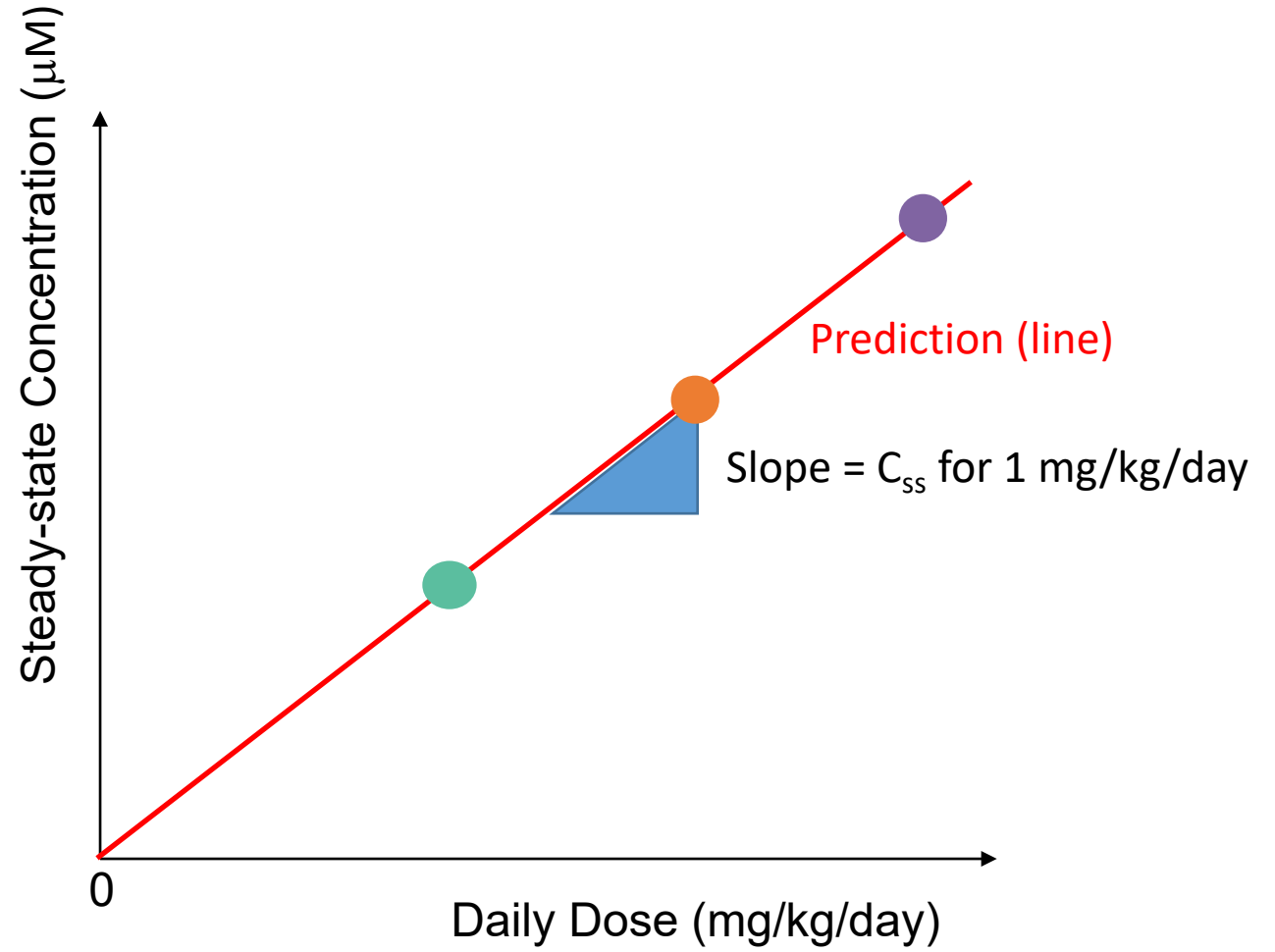
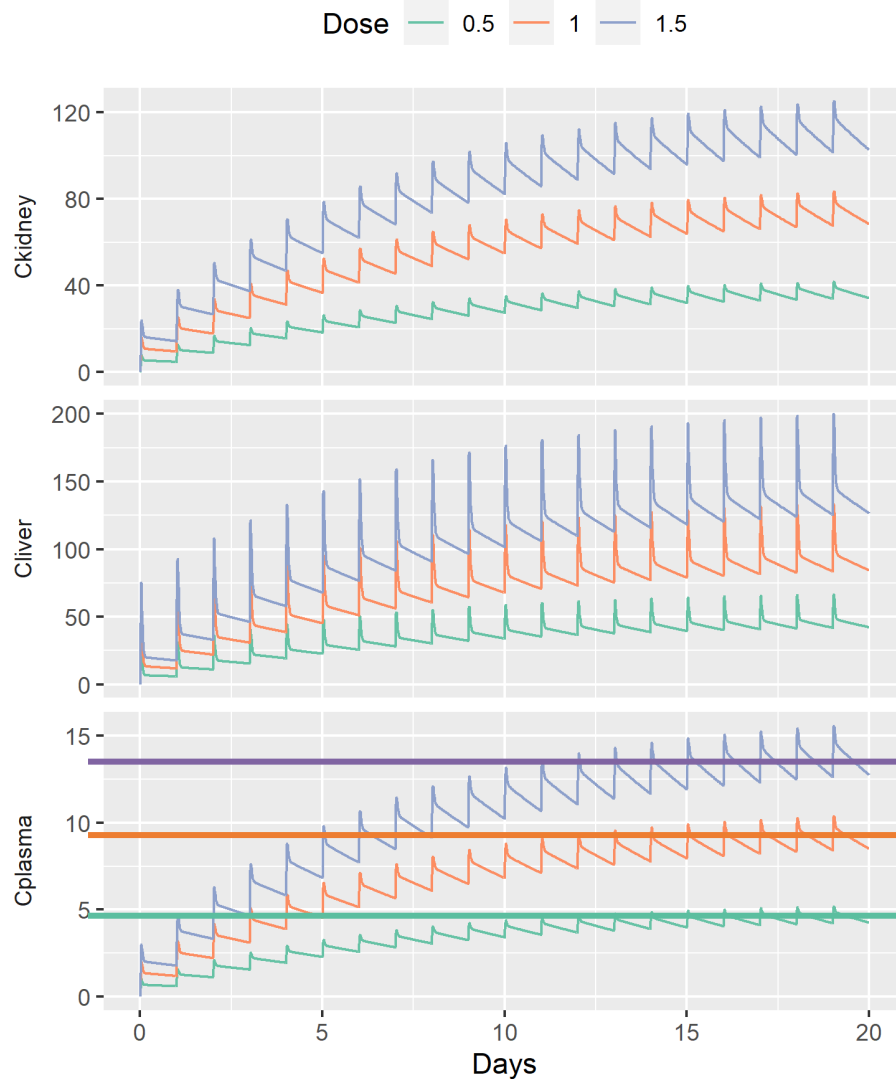
Full concentration vs. time simulations in all compartment are still too computationally intensive — need to simplify further

For chemical screening purposes, we are usually interested in what happens with long-term, low-level exposures

So we focus on the steady-state plasma concentration ( $C_{ss}$ )



In generic PBTK model,  $C_{ss}$  has a *linear* relationship with dose

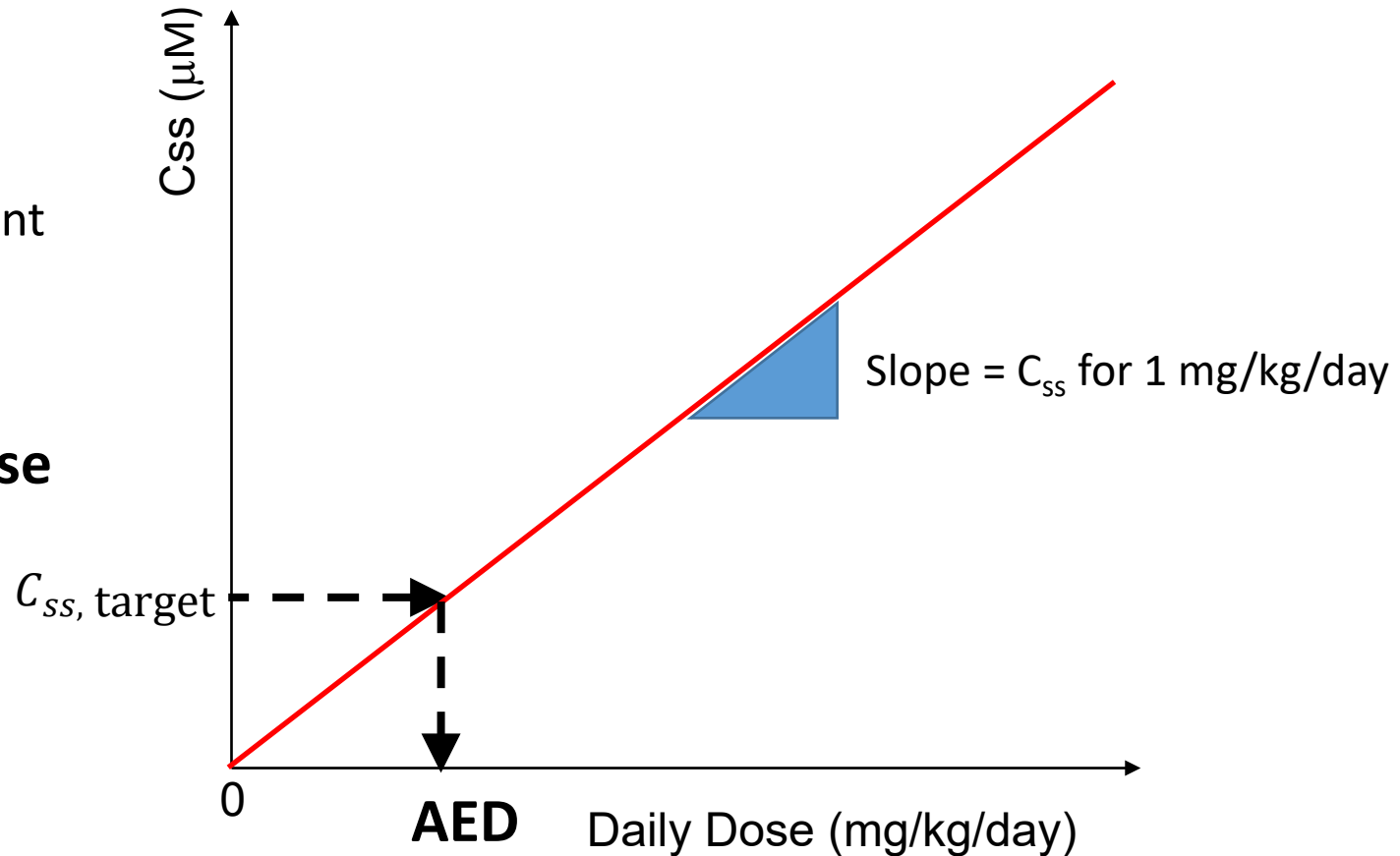


# Linear C<sub>ss</sub>-dose relationship makes reverse TK quick & easy

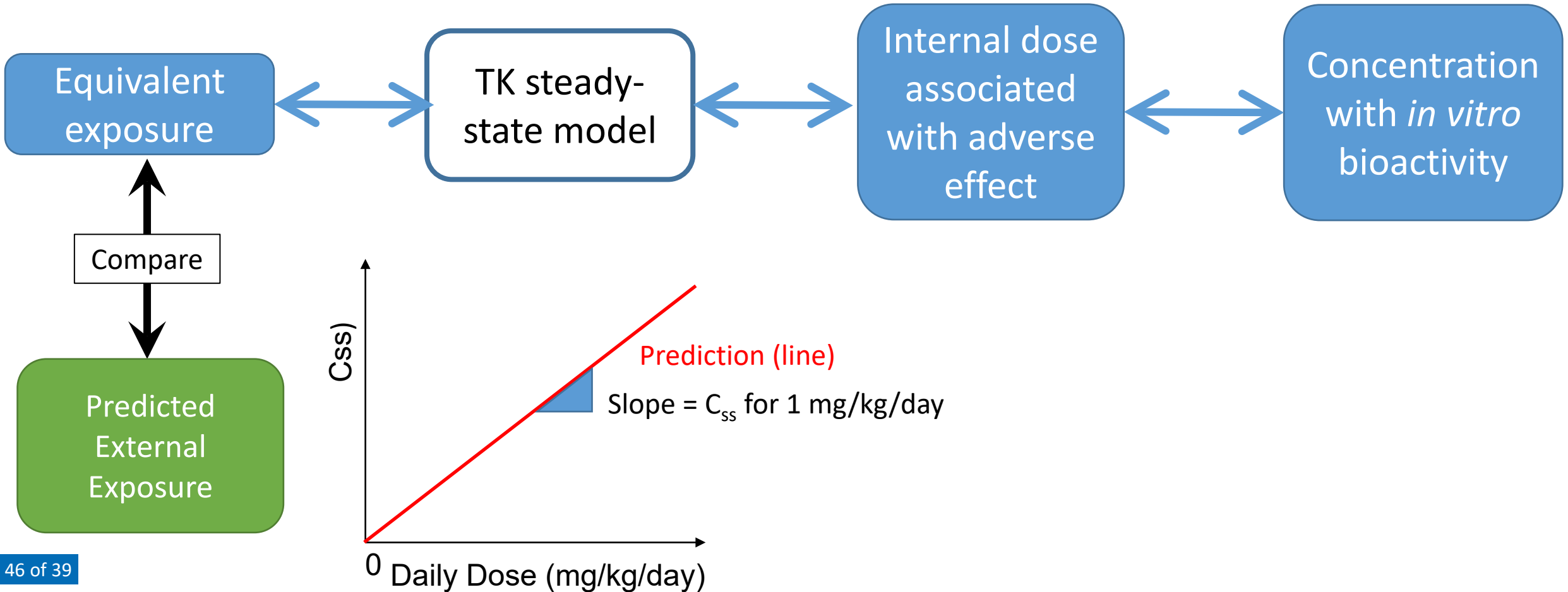
- Graphically:
  - start with the “target” concentration on the y-axis (*in vitro* bioactive concentration  $C_{ss, \text{target}}$ )
  - go over to the C<sub>ss</sub>-dose line
  - drop down to the x-axis
  - then read off the “administered equivalent dose” (AED) on the x-axis.

- Mathematically: 
$$\text{AED} = \frac{C_{ss, \text{target}}}{\text{slope}}$$

- Interpretation: **AED = the *external* dose that would produce an *internal* body concentration equal to the *in vitro* bioactive concentration**



So, we can do IVIVE rapidly for large numbers of chemicals — *if we can get the slope of the  $C_{ss}$ -dose line for each chemical*



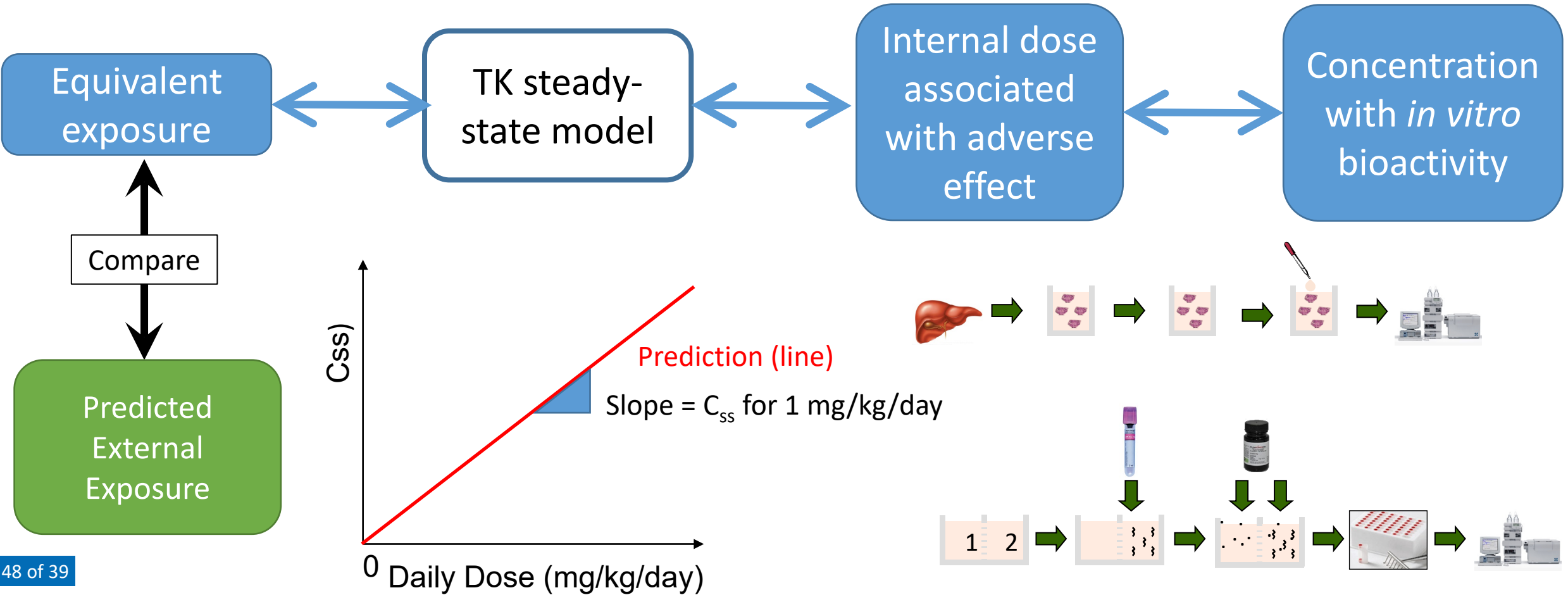
Q: What determines the slope of the line?

A: The TK model parameters that describe ADME.

Chemical-specific parameters	How do we get the parameter values?
Intrinsic hepatic clearance rate (metabolism)	Measured in HT <i>in vitro</i> assays (Rotroff <i>et al.</i> 2010; Wetmore <i>et al.</i> 2012, 2014, 2015; Wambaugh <i>et al.</i> 2019)
Fraction unbound to plasma protein	
Tissue partition coefficients (ratio of conc. in tissue to conc. in plasma)	Predict <i>in silico</i> from phys-chem properties and tissue properties (Pearce <i>et al.</i> , 2017b)
Physiological parameters (chemical-independent)	
Tissue masses (including body weight)	Gathered from data available in the published literature [Wambaugh <i>et al.</i> 2015; Pearce <i>et al.</i> 2017a]
Tissue blood flows	
Glomerular filtration rate (passive renal clearance)	
Hepatocellularity	



So to do high-throughput IVIVE for thousands of chemicals, all we need is the *in vitro* measured chemical-specific TK parameters!



CRAN - Package httpk

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

Apps Absence Request Travel Request Form REMD-HTTK Confluence Bitbucket CompTox Dashboard EHP Change Password

httpk: 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) and extrapolation ("IVIVE") of in vitro data to real-world exposures via PBTK models.

Version: 2.0.1  
Depends: R (≥ 2.10)  
Imports: deSolve, msm, data.table, survey, mvtnorm, truncnorm, statmod, ggplot2, knitr, rmarkdown, R.rsp, GGally, gplots, scales, Enrichr, ggrepel, dplyr, forcats, smatr, gttools, gridExtra  
Published: 2020-03-02  
Author: John Wambaugh [aut, cre], Robert Pearce [aut], Catherine J. Sipes [ctb], Barbara Wetmore [ctb], Woodrow Setzer [ctb]  
Maintainer: John Wambaugh <wambaugh.john@epa.gov>  
BugReports: https://github.com/EPA/CompTox-ExpoCast-httpk  
License: GPL-3  
URL: https://www.epa.gov/chemical-research/rapid-chemical-exposure-assessment  
NeedsCompilation: yes  
Citation: httpk citation info  
Materials: NEWS  
CRAN checks: httpk results

Downloads: downloads 806/month

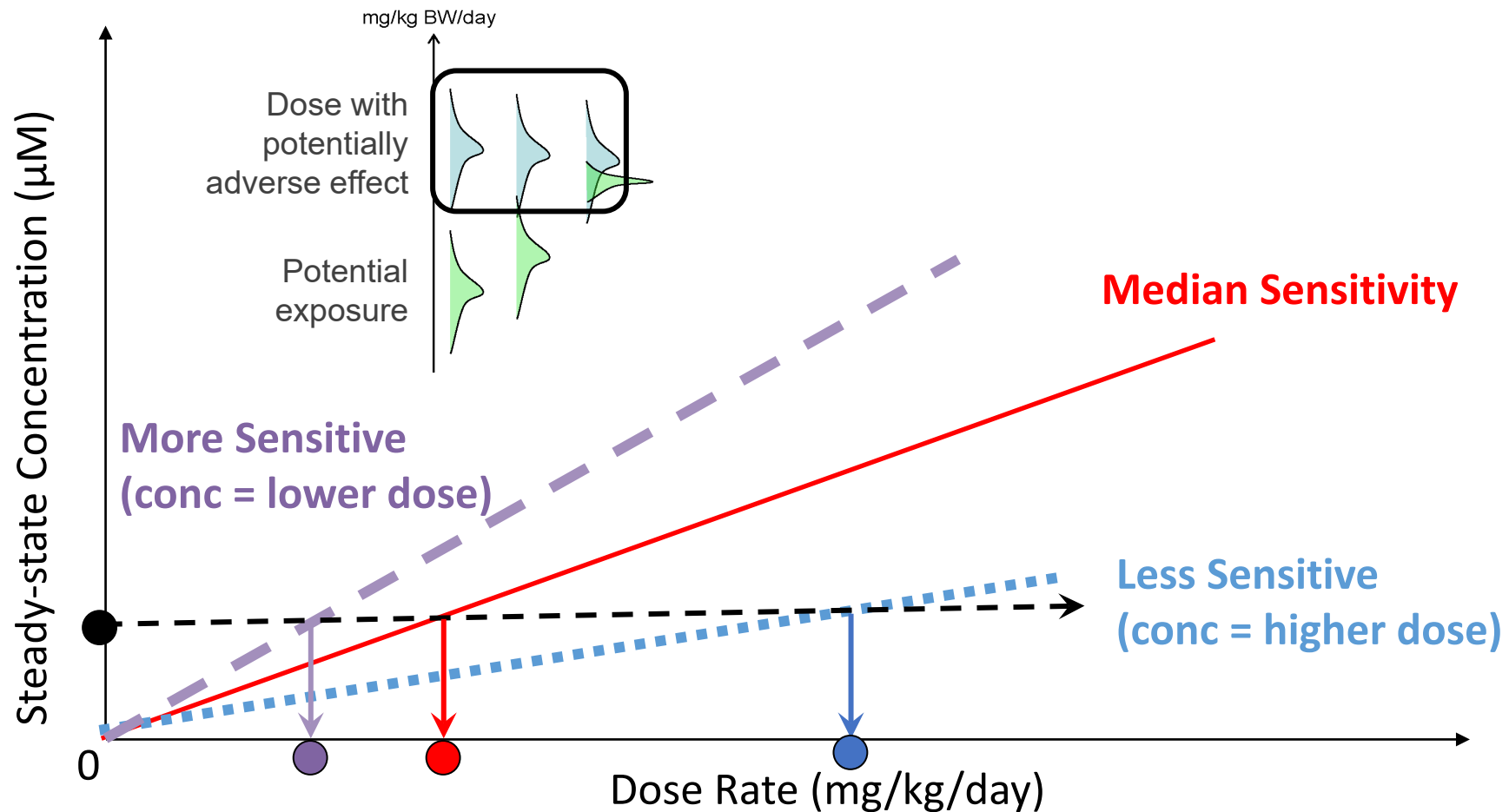
Reference manual: httpk.pdf  
Vignettes: Frank et al. (2018): Creating IVIVE Figure (Fig. 6)  
Honda et al. (2019): Updated Armitage et al. (2014) Model  
Linakis et al. (Submitted): Analysis and Figure Generation  
Pearce et al. (2017): Creating Partition Coefficient Evaluation

R package httpk

- Open source, transparent, and peer-reviewed tools and data for high throughput toxicokinetics (HTTK)
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific TK data for 987 chemicals

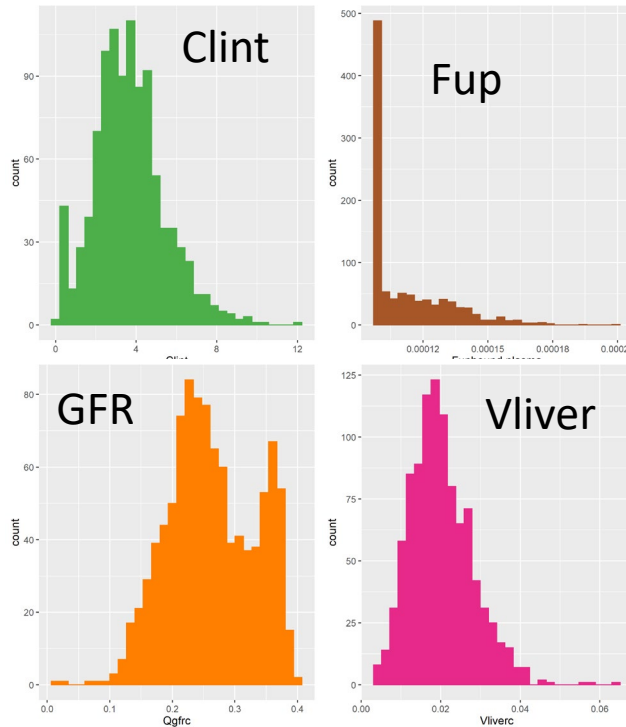
- Open source, transparent, and peer-reviewed tools and data for **high throughput toxicokinetics (HTTK)**
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific TK data for 987 chemicals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

Complication: Population biological variability in TK means that there is a *distribution* of  $C_{ss}$ -dose slopes — and thus a *distribution* of equivalent doses for any given *in vitro* bioactive concentration

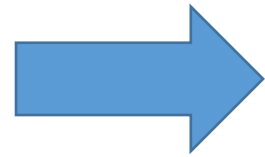


# Population variability in IVIVE can be quantified using a Monte Carlo approach: “HTTK-Pop” (Ring et al., 2017)

Sample from population distribution of TK parameters based on CDC NHANES data

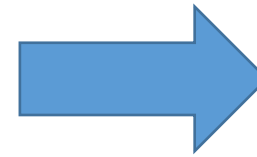


(+ other params)

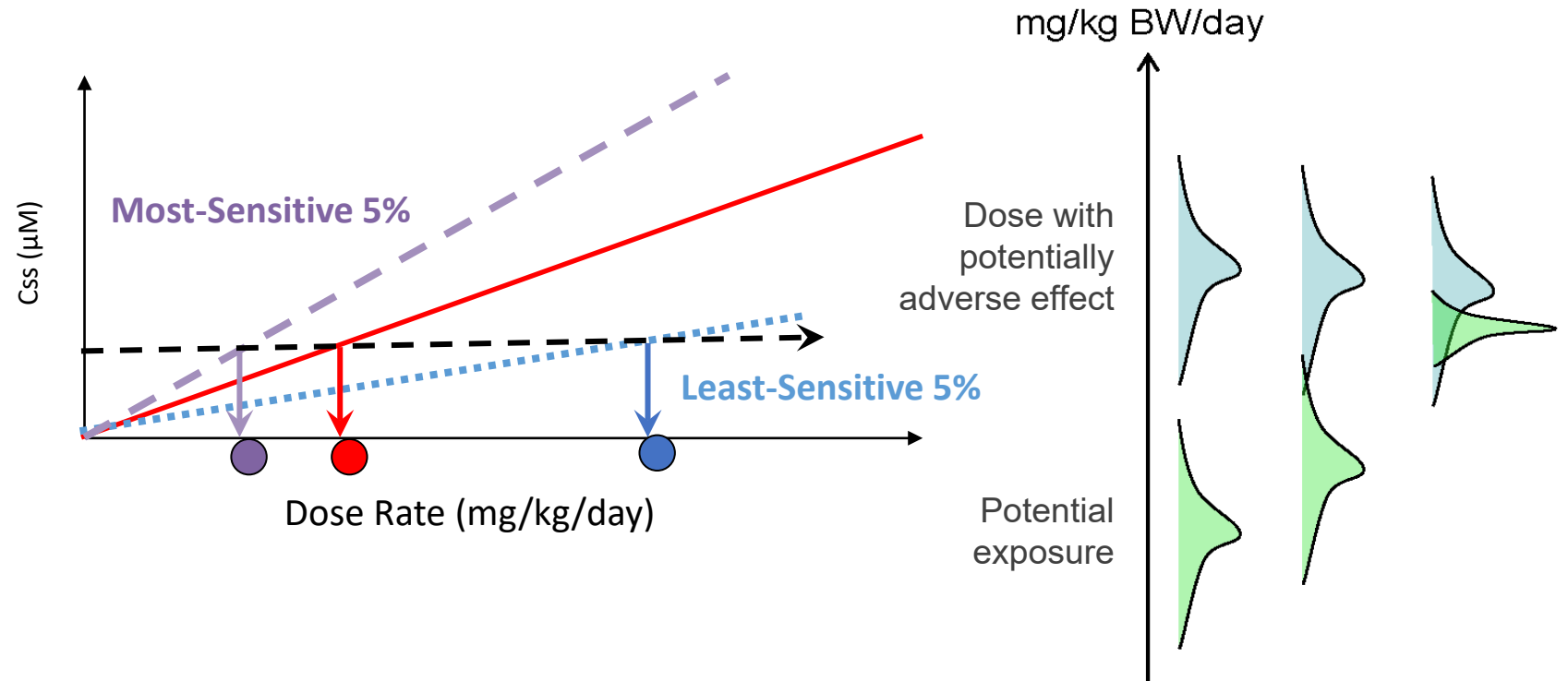


Calculate  $C_{ss}$ -dose slope for each sampled set of TK model parameters

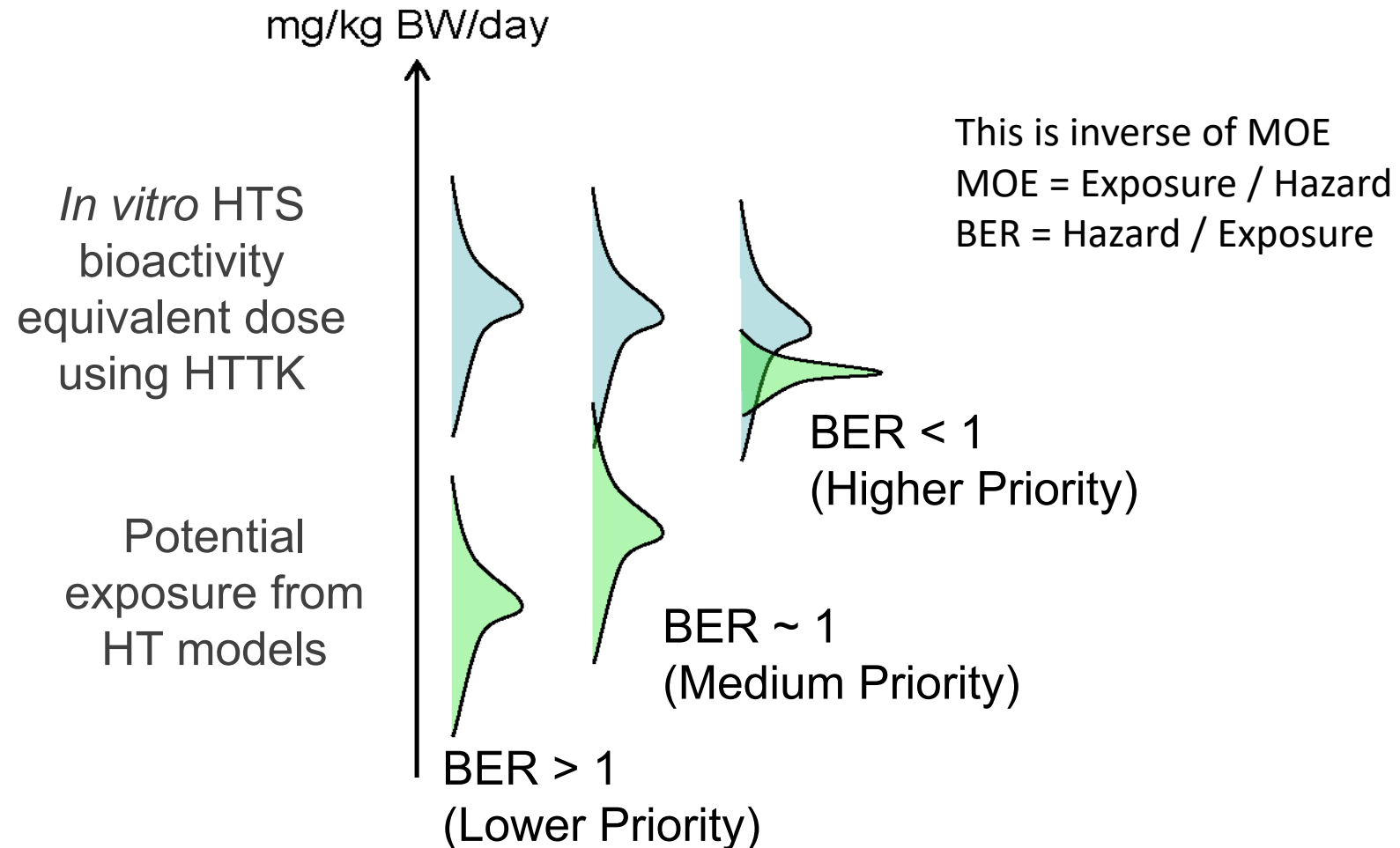
Get resulting distribution of equivalent doses



Compare equivalent dose distribution to potential exposure distribution to calculate potential risk

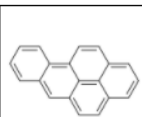


Compare the low-end equivalent dose to the high-end potential exposure to calculate “Bioactivity-Exposure Ratio” (BER).





Example: Using httk to find an equivalent dose & BER for a low-end ToxCast AC50 for benzo(a)pyrene



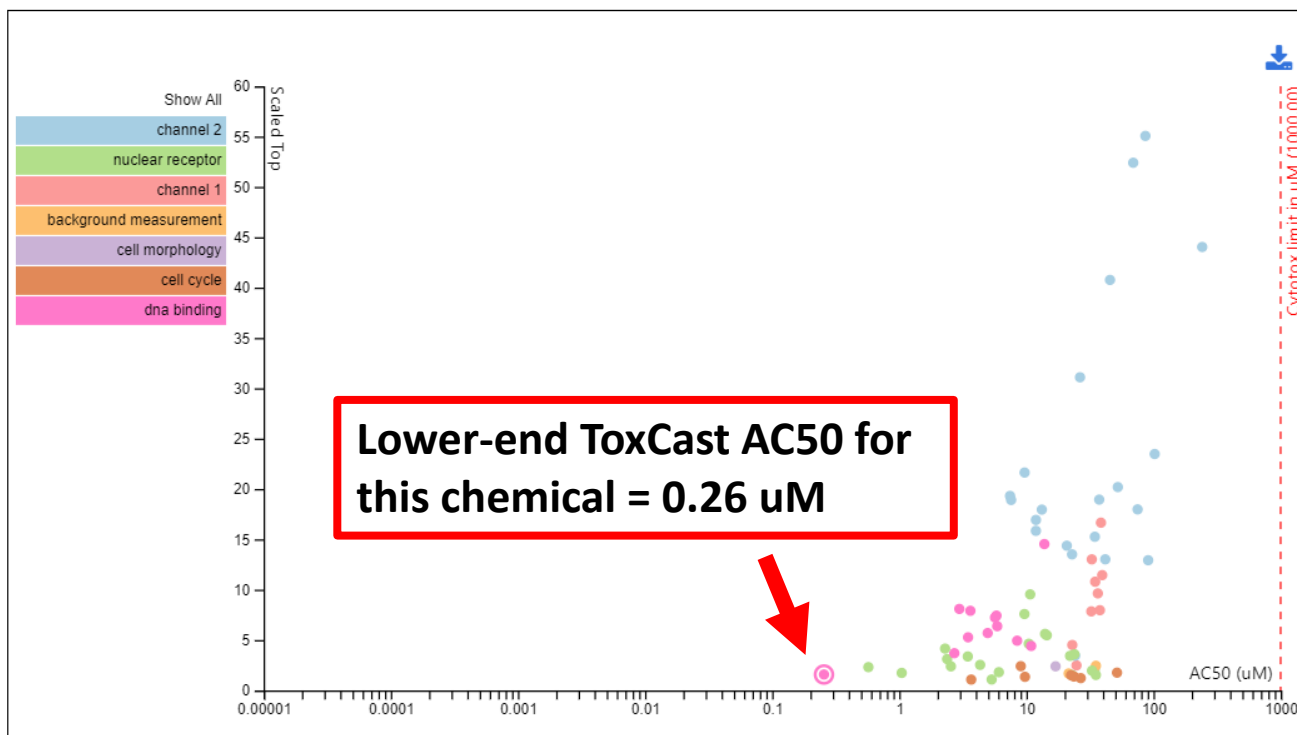
Benzo(a)pyrene  
50-32-8 | DTXSID2020139  
Searched by DSSTox Substance Id.

First: Get AC50 value. ToxCast AC50s can be found on the CompTox Chemicals Dashboard.

## Chemical Activity Summary

### TOXCAST DATA

### ASSAY DETAILS



AC50 (uM): 0.26  
Scaled top: 1.58  
Assay Endpoint Name: TOX21\_SSH\_3T3\_GLI3\_Antagonist  
Gene Symbol:  
Organism: mouse  
Tissue: embryo  
Assay Format Type: cell-based  
Biological Process Target: regulation of transcription factor activity  
Detection Technology: Luciferase-coupled ATP quantitation  
Analysis Direction: positive  
Intended Target Family: dna binding  
Description: Data from the assay component TOX21\_SSH\_3T3\_GLI3\_Antagonist was analyzed into 1 assay endpoint. This assay endpoint, TOX21\_SSH\_3T3\_GLI3\_Antagonist, was analyzed in the positive fitting direction relative to DMSO as the negative control and baseline of activity. Using a type of inducible reporter, loss-of-signal activity can be used to understand changes in the reporter gene as they relate to the gene GLI3. Furthermore, this assay endpoint can be referred to as a primary readout, because this assay has produced multiple assay endpoints where this one serves a reporter gene function.

DETAILS

EXECUTIVE SUMMARY

PROPERTIES

ENV. FATE/TRANSPORT

HAZARD

SAFETY

ADME

EXPOSURE

BIOACTIVITY

TOXCAST: SUMMARY

EDSP21

TOXCAST/TOX21

PUBCHEM

TOXCAST: MODELS

SIMILAR COMPOUNDS

To calculate population equivalent dose, use httk function  
`calc_mc_oral_equiv()`

```
> library(httk)
> set.seed(42)
> #Steady-state equivalent dose (mg/kg BW/day) to produce 0.26 uM in plasma:
  calc_mc_oral_equiv(conc=0.26,
                     chem.name="benzo(a)pyrene",
                     which.quantile = c(0.95, 0.5, 0.05),
                     input.units = "uM",
                     output.units = "mgpkgpday")
```

uM concentration converted to mgpkgpday dose for 0.95 0.5 0.05 quantile.

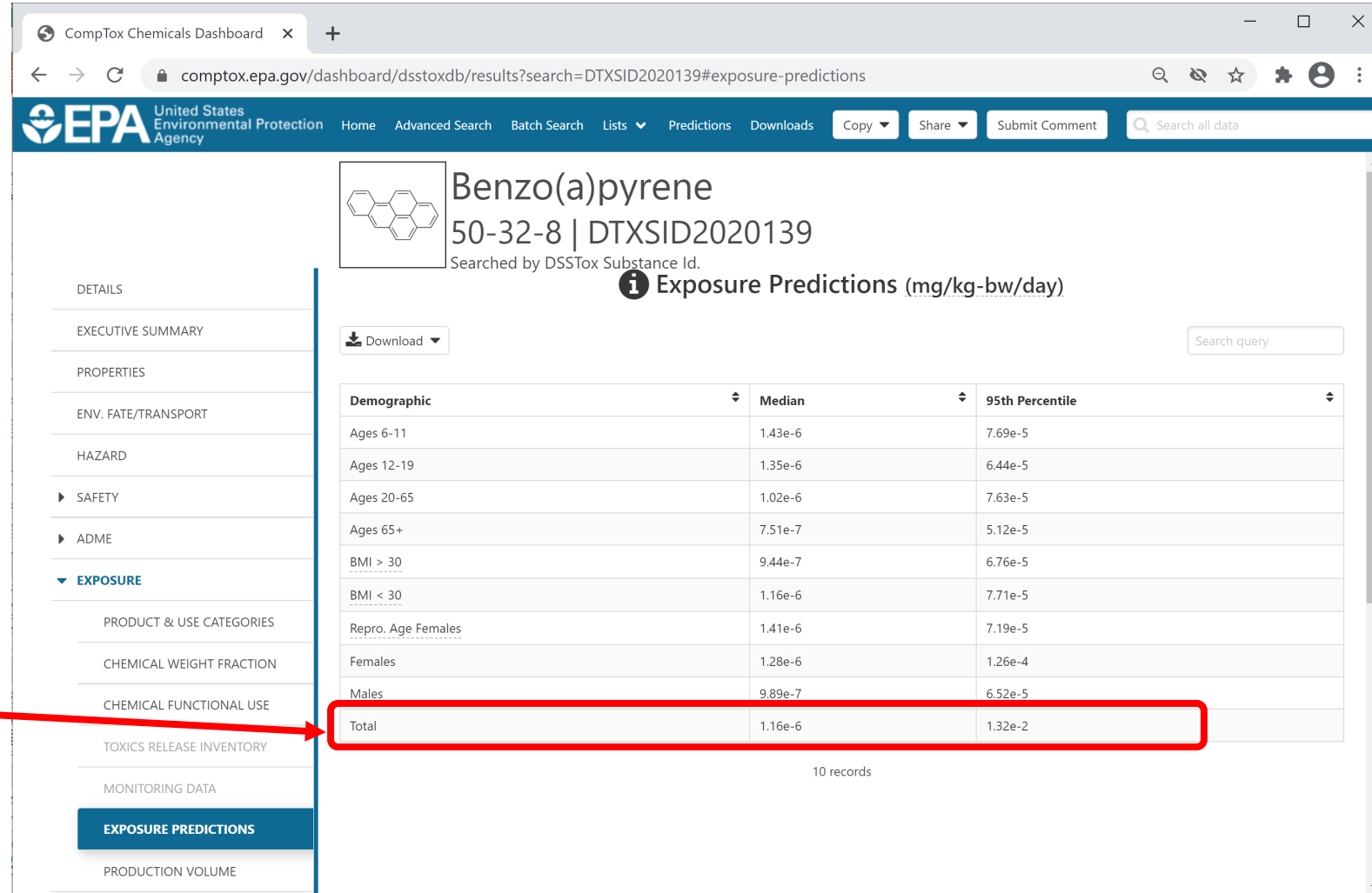
	95%	50%	5%
	0.003821	0.019090	0.067080

# Compare equivalent dose to HT exposure predictions available on EPA CompTox Chemicals Dashboard

Monte Carlo equivalent dose from  
`httpk::calc_mc_oral_equiv():`  
 uM concentration converted to  
 mg/kg/day dose for 0.95 0.5  
 0.05 quantile.

	95%	50%	5%
	0.003821	0.019090	0.067080

HT exposure predictions from Dashboard:  
 median = 1.16e-6;  
 upper bound on median = 1.32e-2  
 mg/kg/day



CompTox Chemicals Dashboard

comp tox.epa.gov/dashboard/dsstoxdb/results?search=DTXSID2020139#exposure-predictions

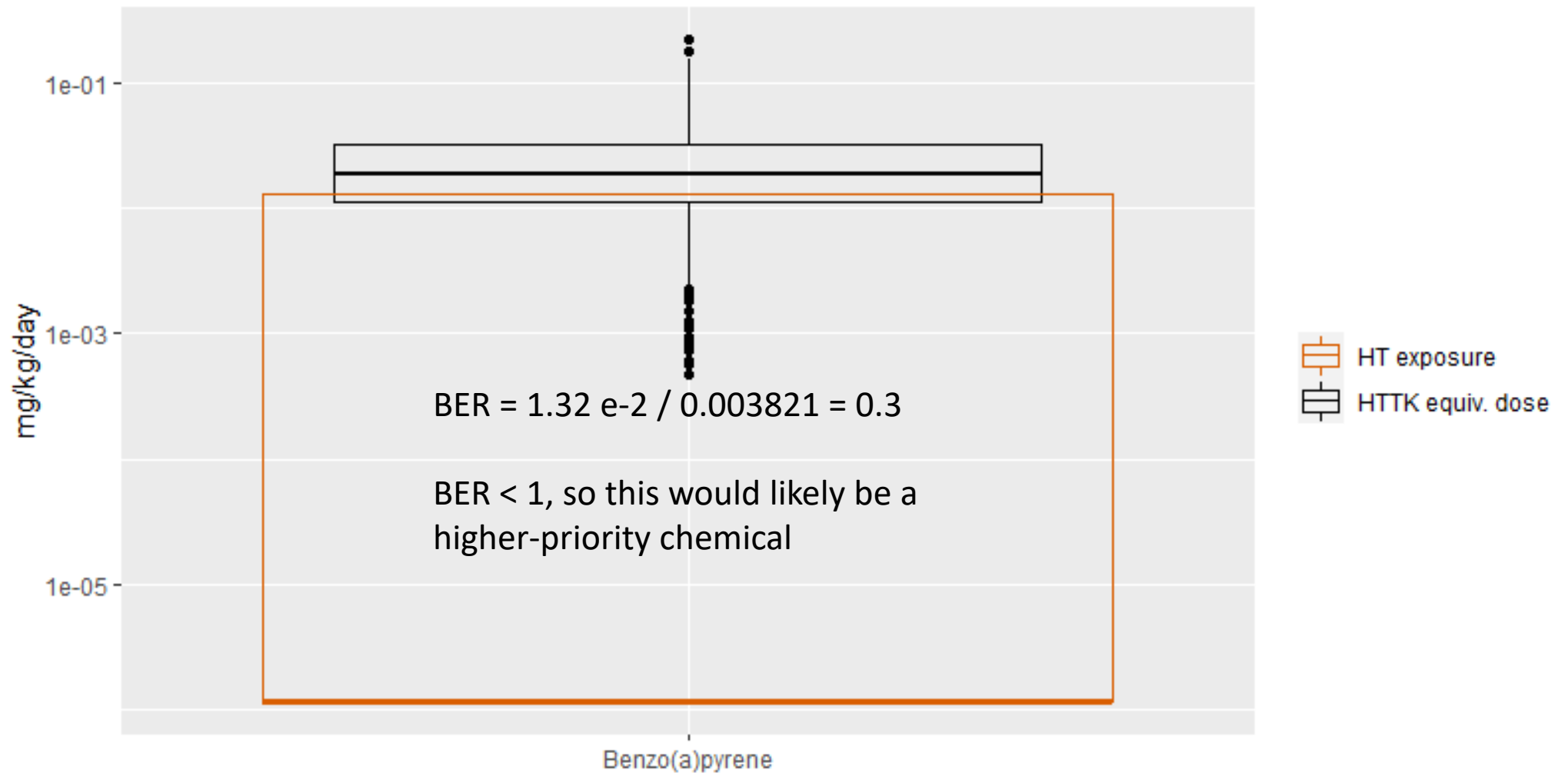
**Benzo(a)pyrene**  
 50-32-8 | DTXSID2020139  
 Searched by DSSTox Substance Id.

**Exposure Predictions (mg/kg-bw/day)**

Demographic	Median	95th Percentile
Ages 6-11	1.43e-6	7.69e-5
Ages 12-19	1.35e-6	6.44e-5
Ages 20-65	1.02e-6	7.63e-5
Ages 65+	7.51e-7	5.12e-5
BMI > 30	9.44e-7	6.76e-5
BMI < 30	1.16e-6	7.71e-5
Repro. Age Females	1.41e-6	7.19e-5
Females	1.28e-6	1.26e-4
Males	9.89e-7	6.52e-5
<b>Total</b>	<b>1.16e-6</b>	<b>1.32e-2</b>

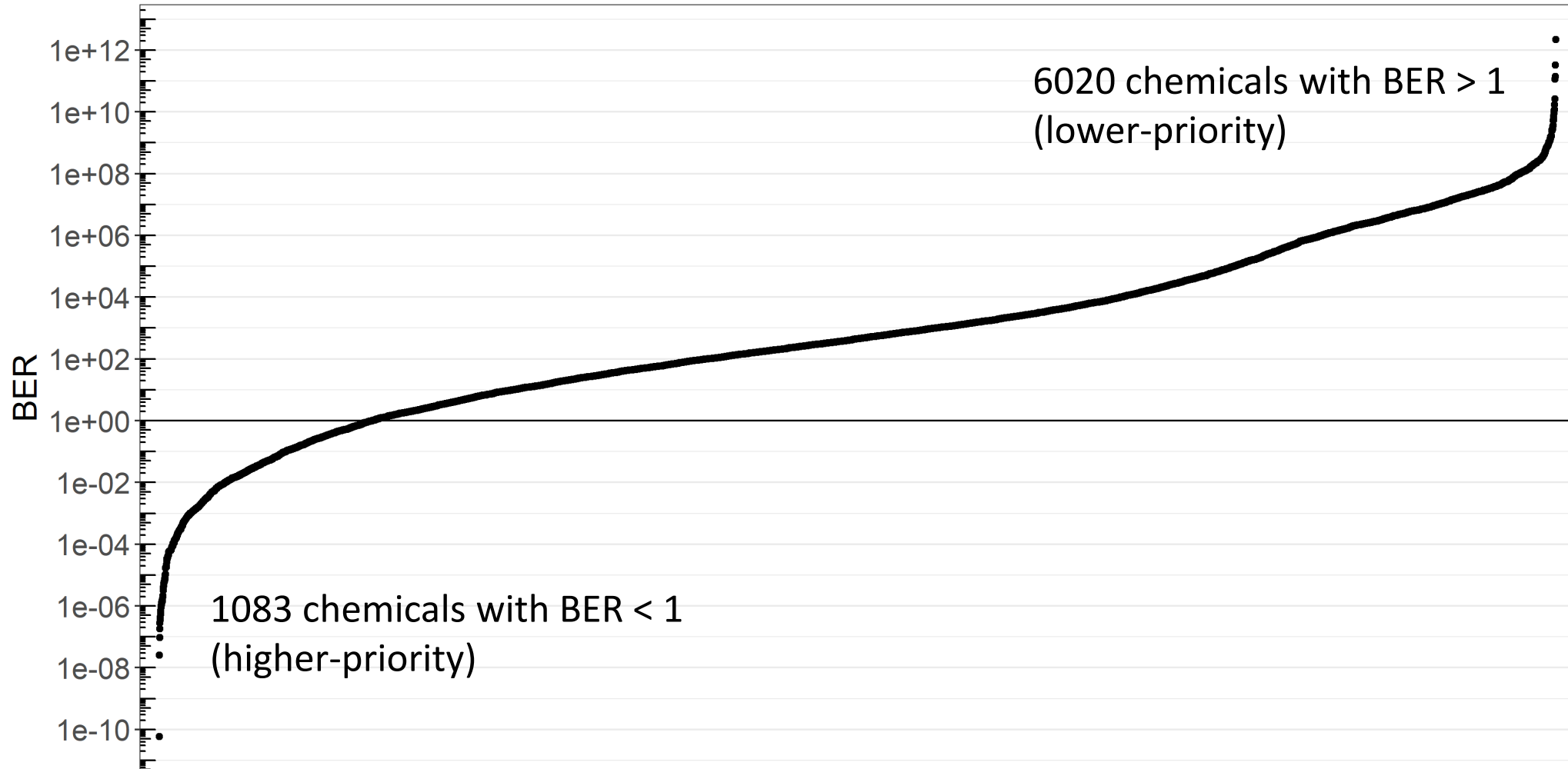
10 records

# BER: Graphical comparison of HTTK-predicted equivalent dose for ToxCast AC50, vs. HT exposure prediction





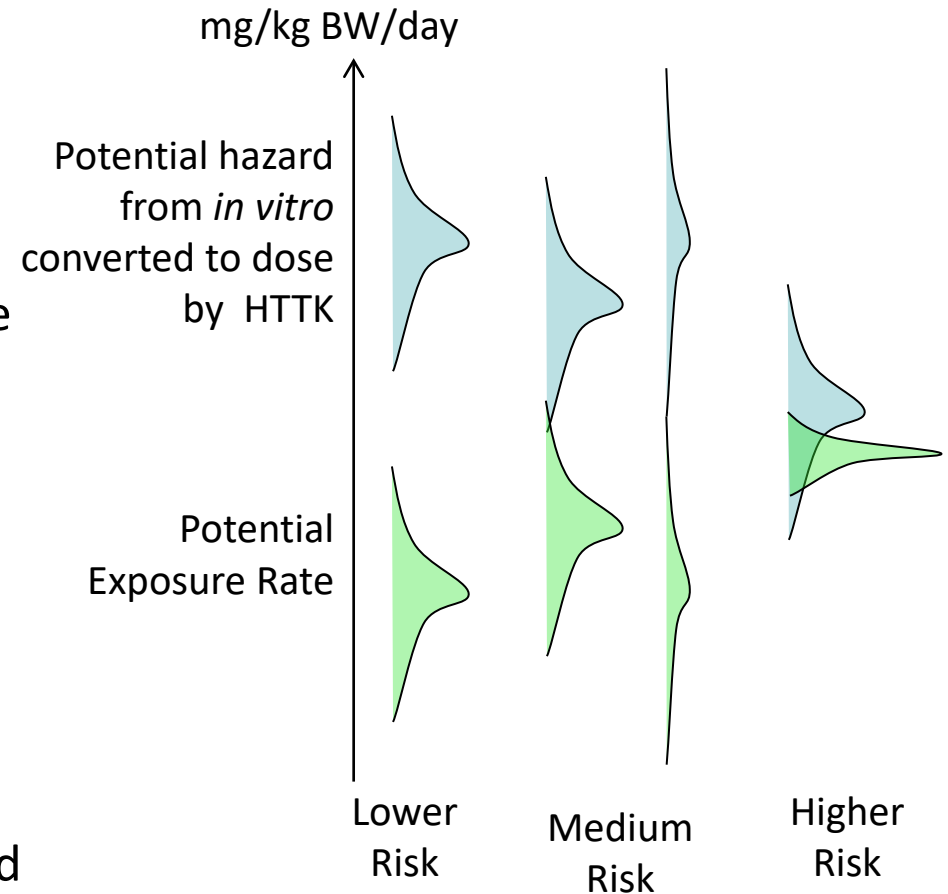
An even-more high-throughput application: BER prioritization of 7104 chemicals based on HTTK-Pop IVIVE of ToxCast AC50s and HT exposure predictions from SEEM3 model





# Summary

- “The dose makes the poison”: risk is a function of both hazard and exposure
- Hazard: When *in vivo* hazard data are not available, we can use *in vitro* high-throughput screening (HTS) assays
- Exposure: estimation requires tracing chemical from source to receptor
- When detailed chemical-specific exposure data are not available, we can use exposure NAMs to fill data gaps and make exposure predictions
- To compare *in vitro* HTS data to *in vivo* exposure estimates, we use high-throughput toxicokinetics (HTTK) -- generic model that can be parameterized with *in vitro* data
- The bioactivity-exposure ratio (BER) framework allows rapid risk-based chemical prioritization
- Hazard, exposure, and TK data and models are publicly available through the CompTox Chemicals Dashboard and as R packages



*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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