

# Utility of NAMs for risk-based prioritization

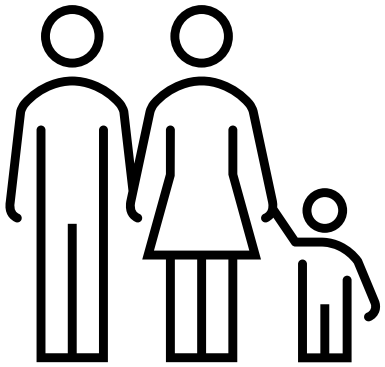
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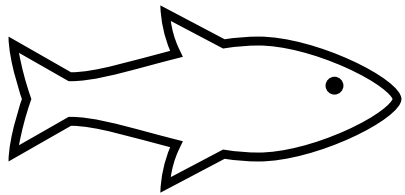
# Human & ecological populations are exposed to thousands of chemicals in the environment – which ones are highest priority?



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Schmidt, C. W. (2016)

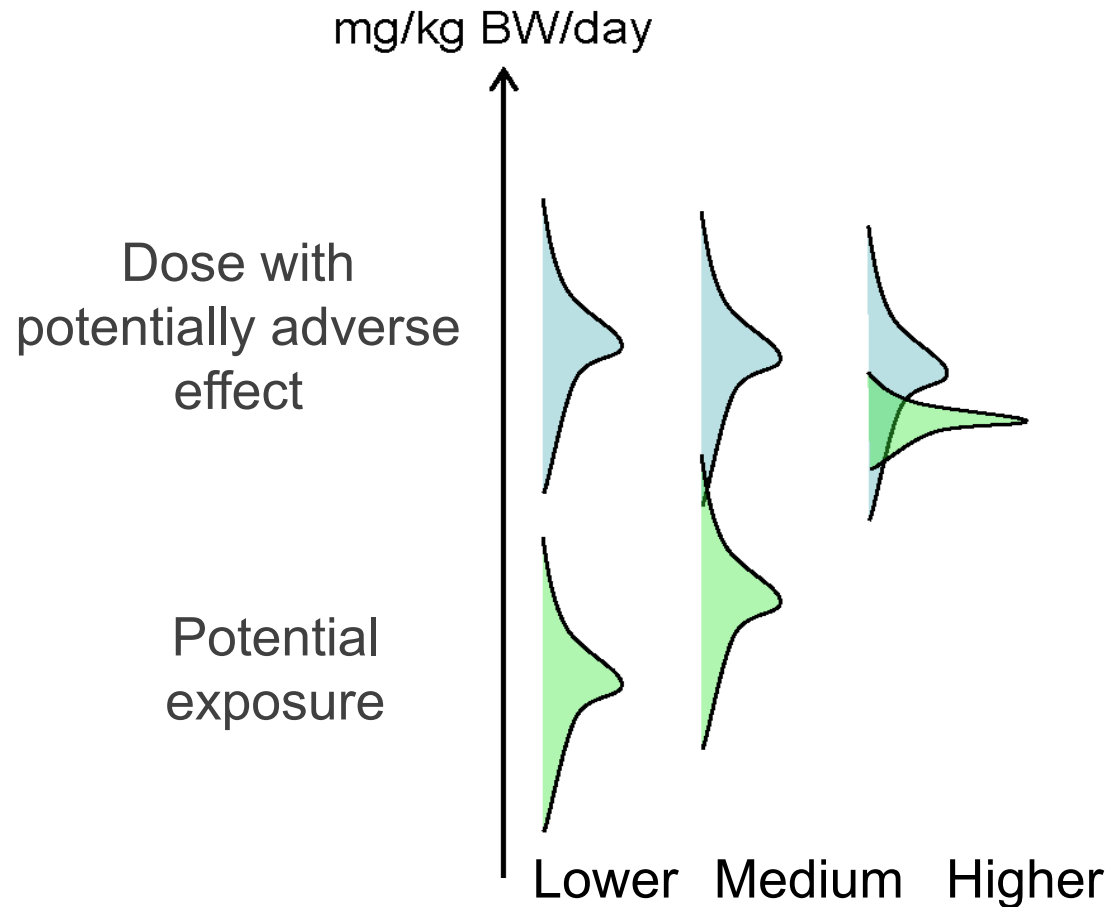


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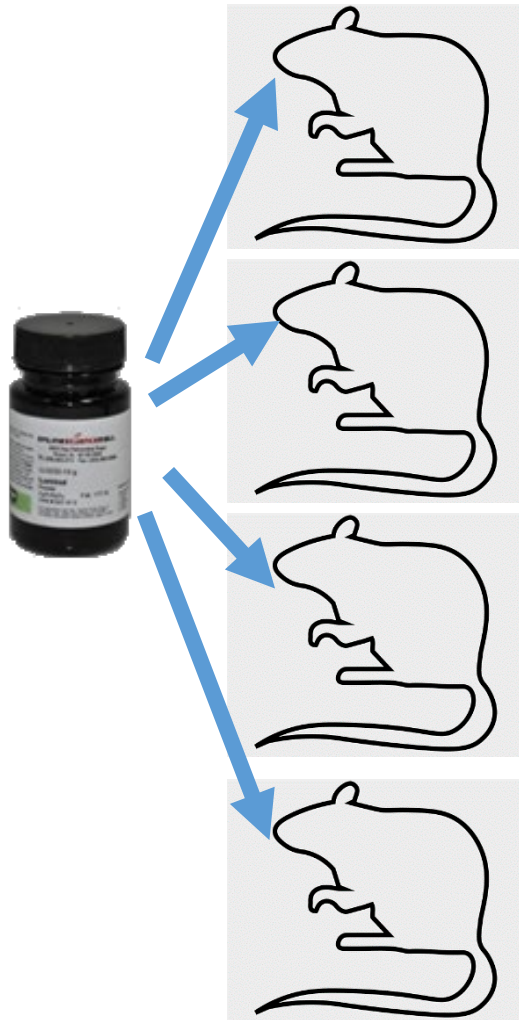


US EPA government photograph

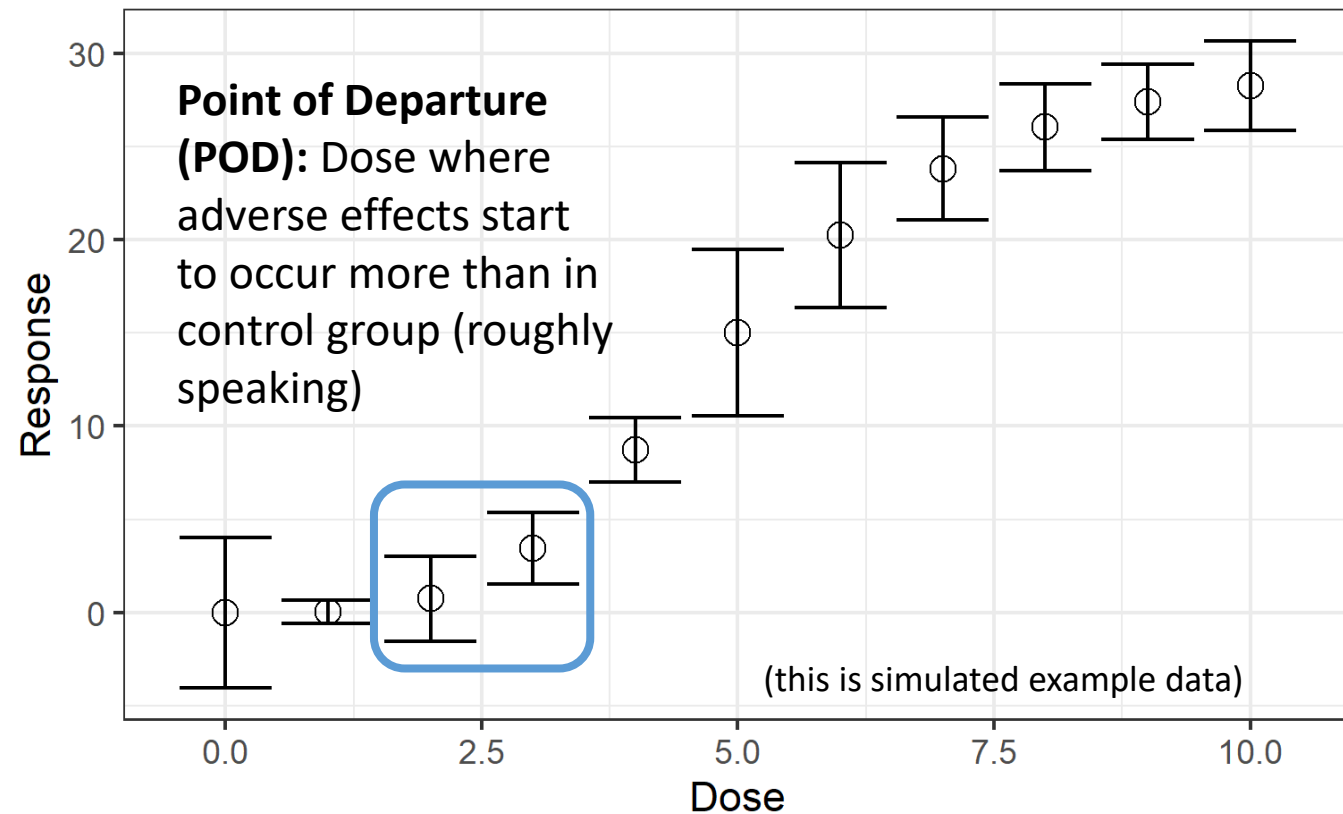
# Risk-based prioritization is a function of both hazard and exposure



# 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!

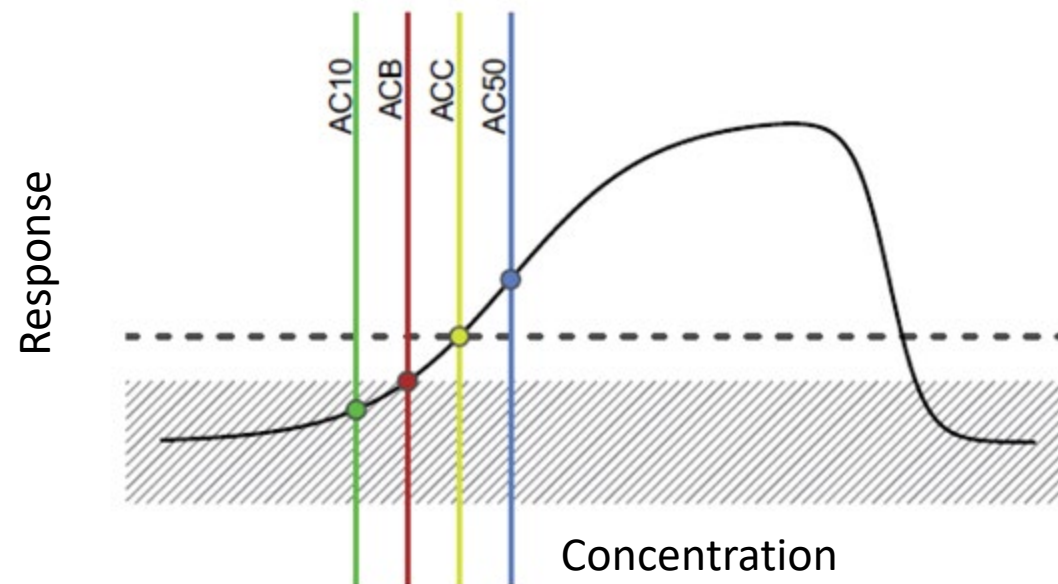
[Schmidt 2009; Dix et al. 2007; Kavlock et al. 2018; Filer *et al.*, 2016; Franzosa et al. 2021]

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

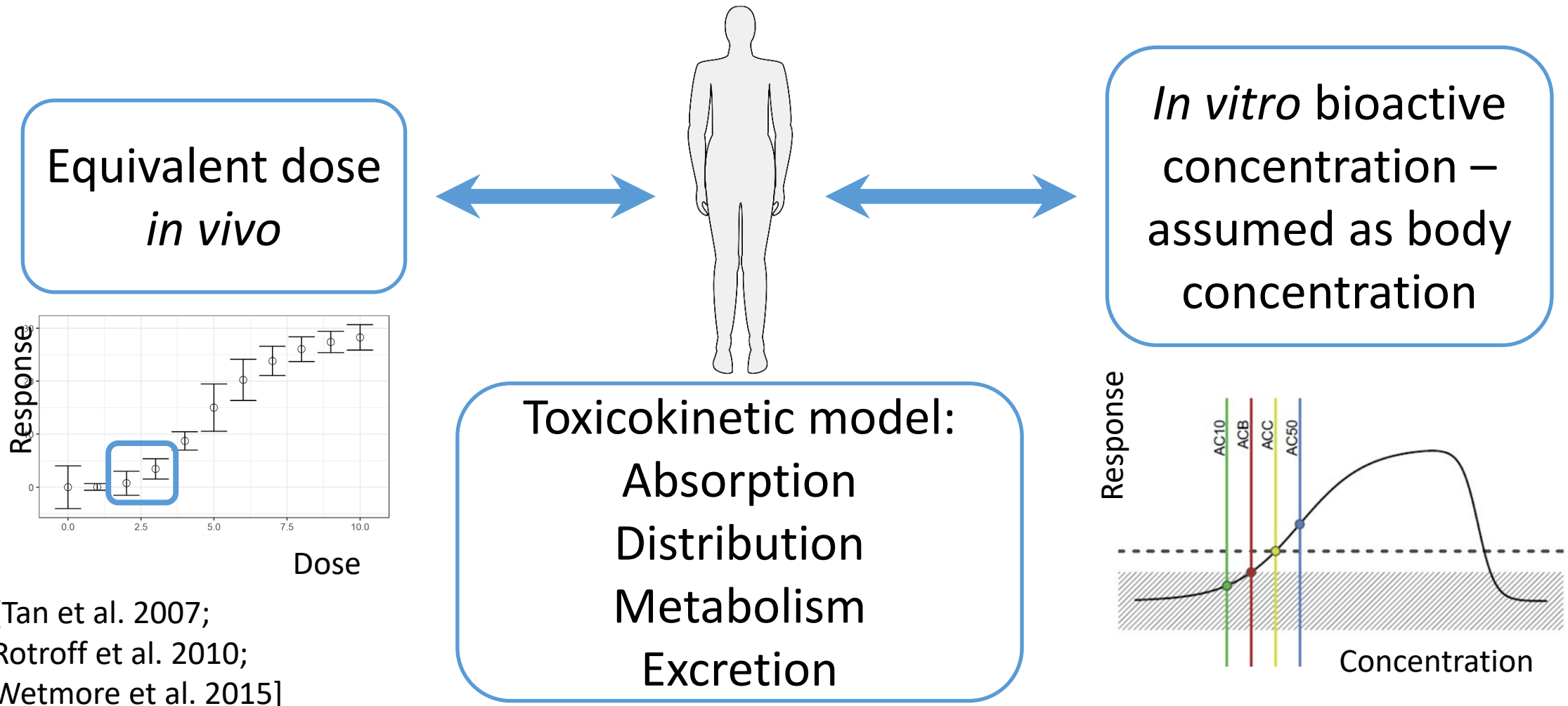
All data are public:

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

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



# Convert *in vitro* concentration to equivalent “*in vivo*” POD using toxicokinetic modeling



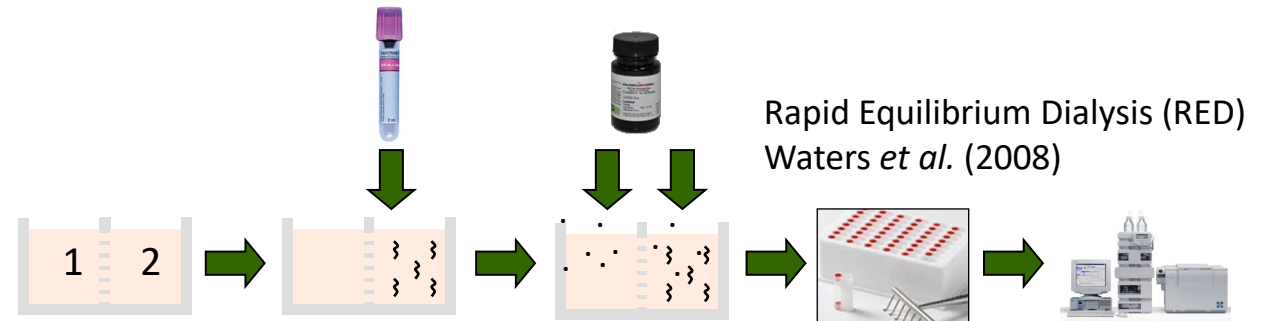
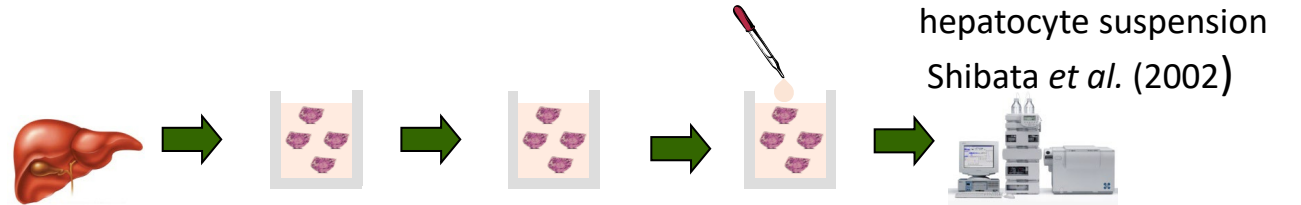
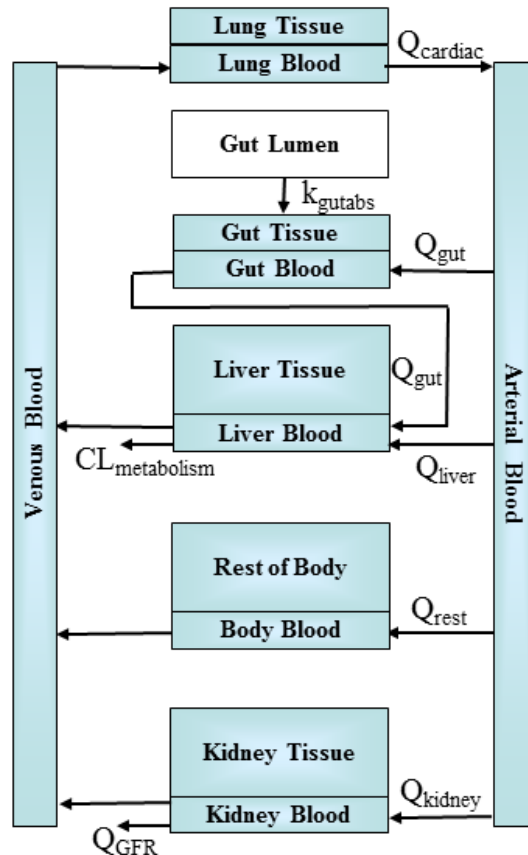
# New approach methodologies for toxicokinetics: *High-throughput toxicokinetics (HTTK)*

## Generic physiologically-based TK (PBTK)

**model:** can be parameterized for many chemicals with minimal chemical-specific data requirements

+

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

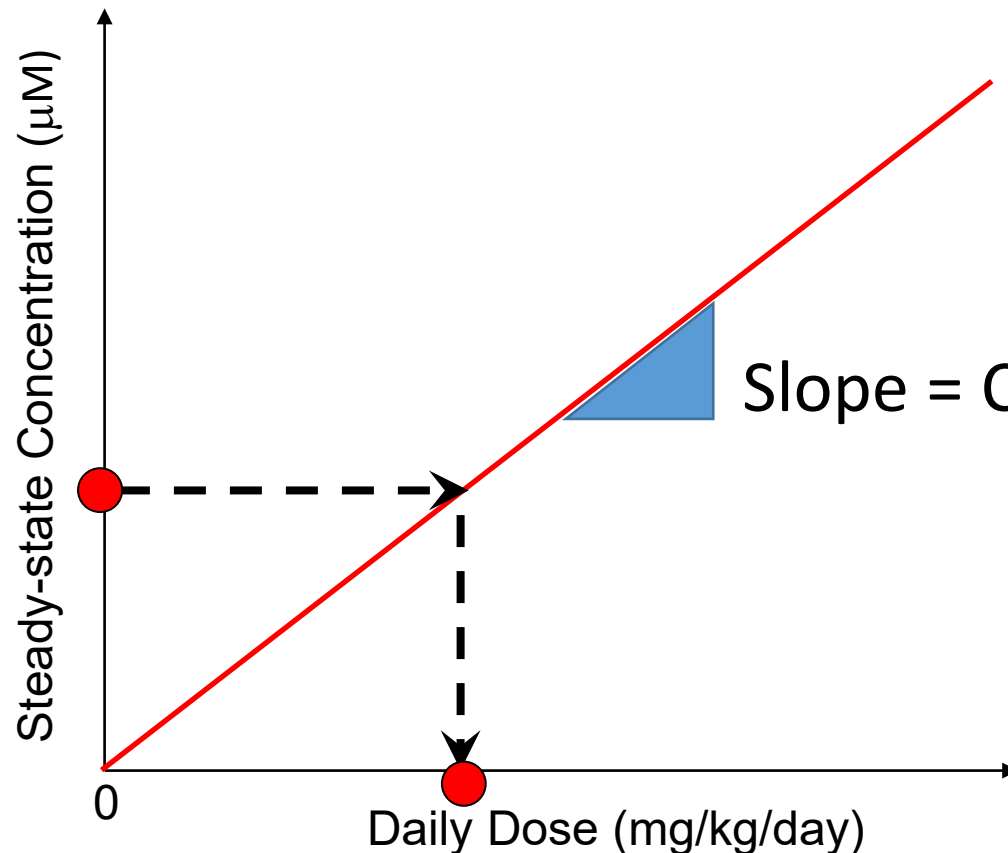


Rotroff *et al.* (2010)  
Wetmore *et al.* (2012)  
Wetmore *et al.* (2015)  
Wambaugh *et al.* (2019)

Wambaugh *et al.* (2015)  
Pearce *et al.* (2017a)  
Ring *et al.* (2017)  
Linakis *et al.* (2020)  
Breen *et al.* (2021)

For high-throughput applications: focus on steady-state plasma concentration ( $C_{ss}$ ) & simple TK models where  $C_{ss}$  is linear with dose

Concentration  
of interest:  
e.g. *in vitro*  
bioactive  
concentration



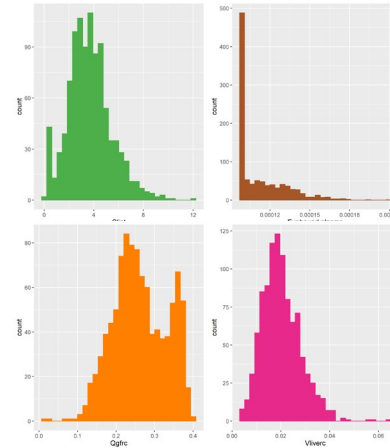
Slope =  $C_{ss}$  for 1 mg/kg/day

in `httk` R package:  
`calc_analytic_css()`

Equivalent dose

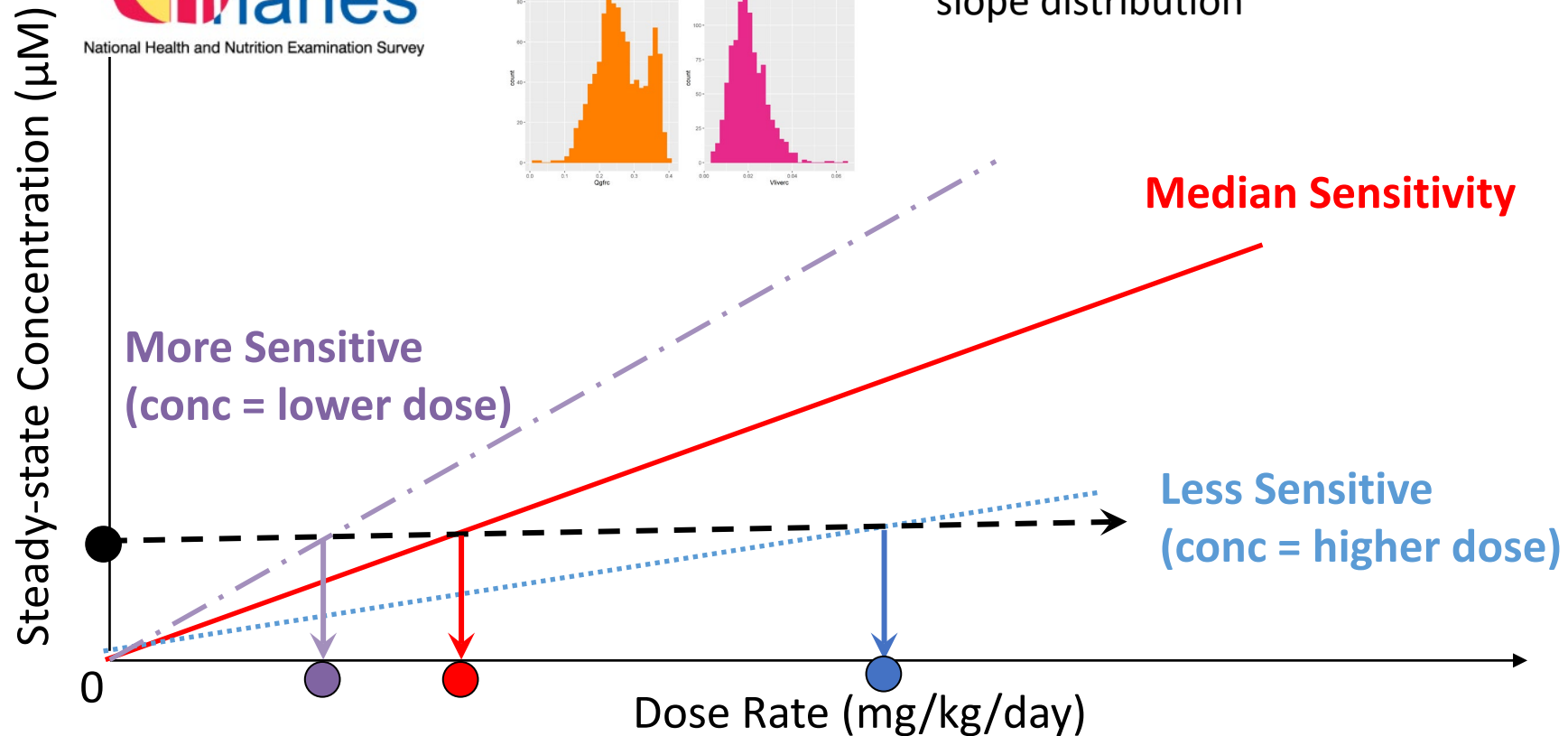
Wetmore *et al.* (2012)

# Inter-individual variability in TK is modeled using a correlated Monte Carlo approach based on CDC NHANES



Sample population TK parameters & calculate resulting population C<sub>ss</sub>-dose slope distribution

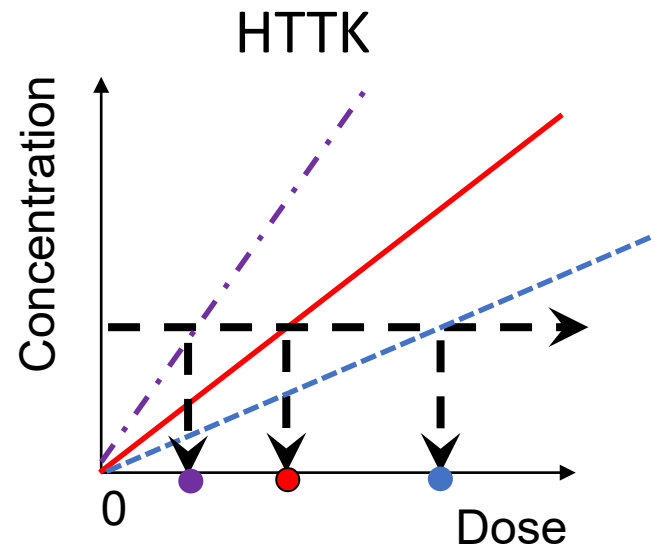
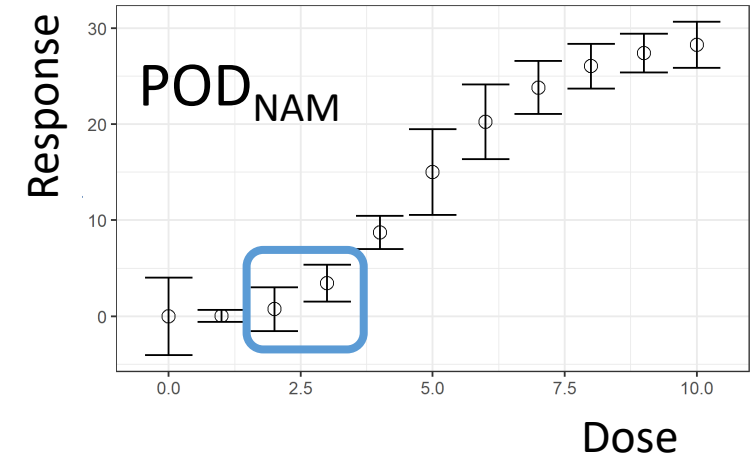
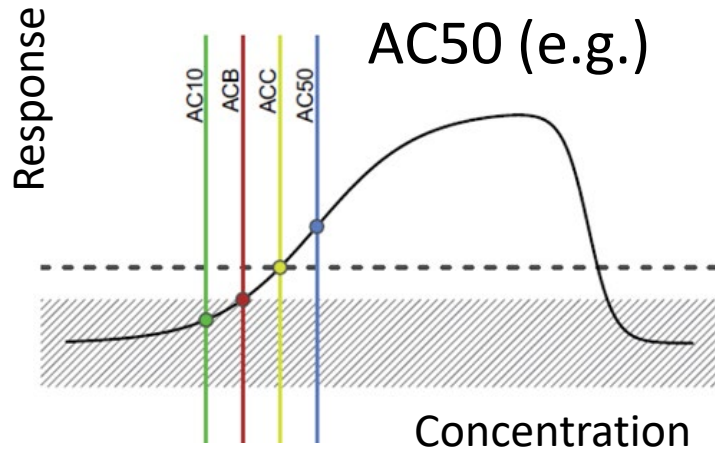
Ring et al.  
2017



[illegible]

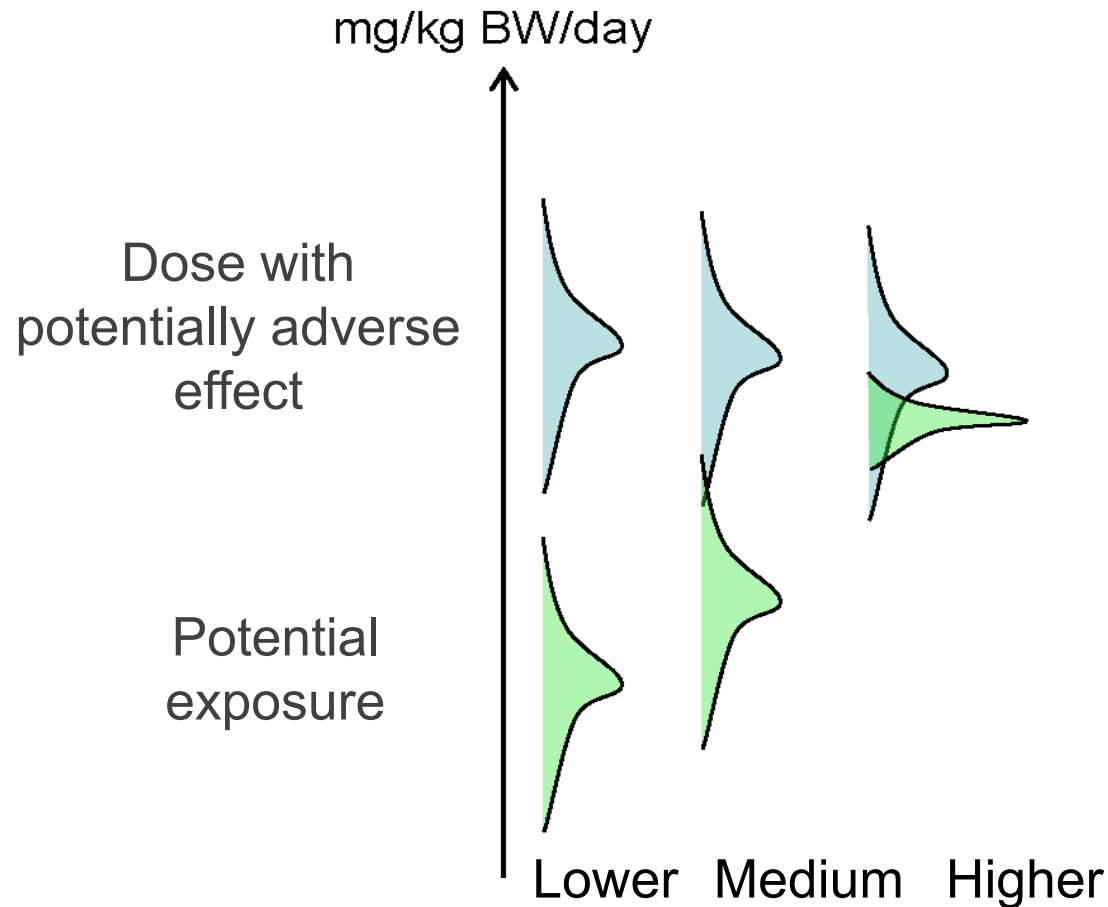
- Open source, transparent, and peer-reviewed tools and data for **high throughput toxicokinetics (HTTK)**
- Available publicly for free statistical software R
- Allows *in vitro-in vivo* extrapolation (IVIVE) and physiologically-based toxicokinetics (PBTK)
- Human-specific TK data for 987 chemicals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

So: We can map *in vitro* bioactive concentration to equivalent “*in vivo*”  $POD_{NAM}$  (dose) using HTTK



Note that this is like a data-driven  $UF_H$  (human TK variability)

Risk is a function of both hazard and *exposure* –  
new approach methodologies for exposure



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



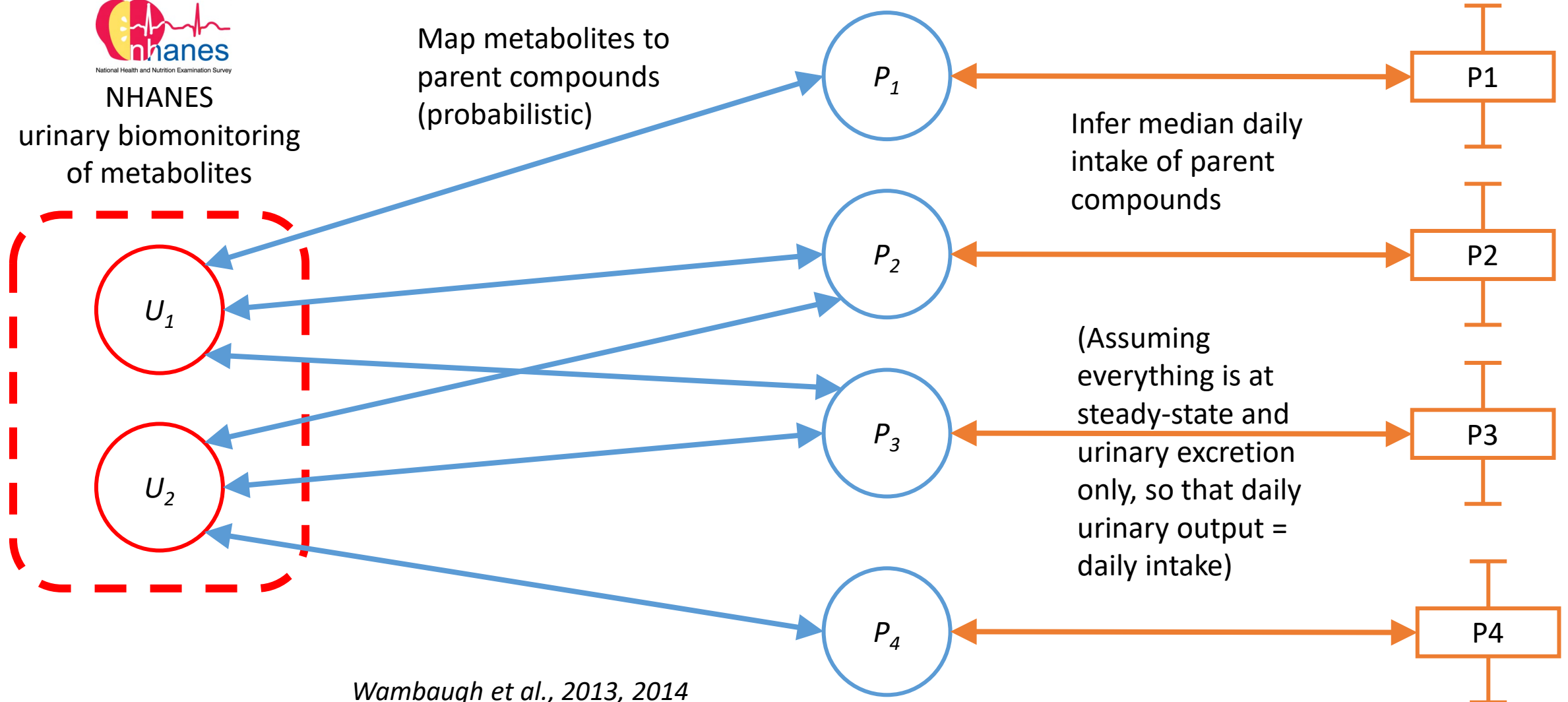
NHANES  
urinary biomonitoring  
of metabolites

Map metabolites to  
parent compounds  
(probabilistic)

Infer median daily  
intake of parent  
compounds

(Assuming  
everything is at  
steady-state and  
urinary excretion  
only, so that daily  
urinary output =  
daily intake)

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

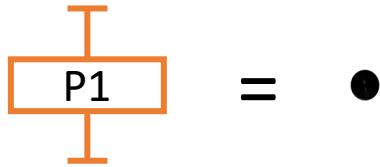


# SEEM3: A consensus model for aggregate exposure

SEEM3 = Systematic Empirical Evaluation of Models, version 3

Ring et al. (2019)

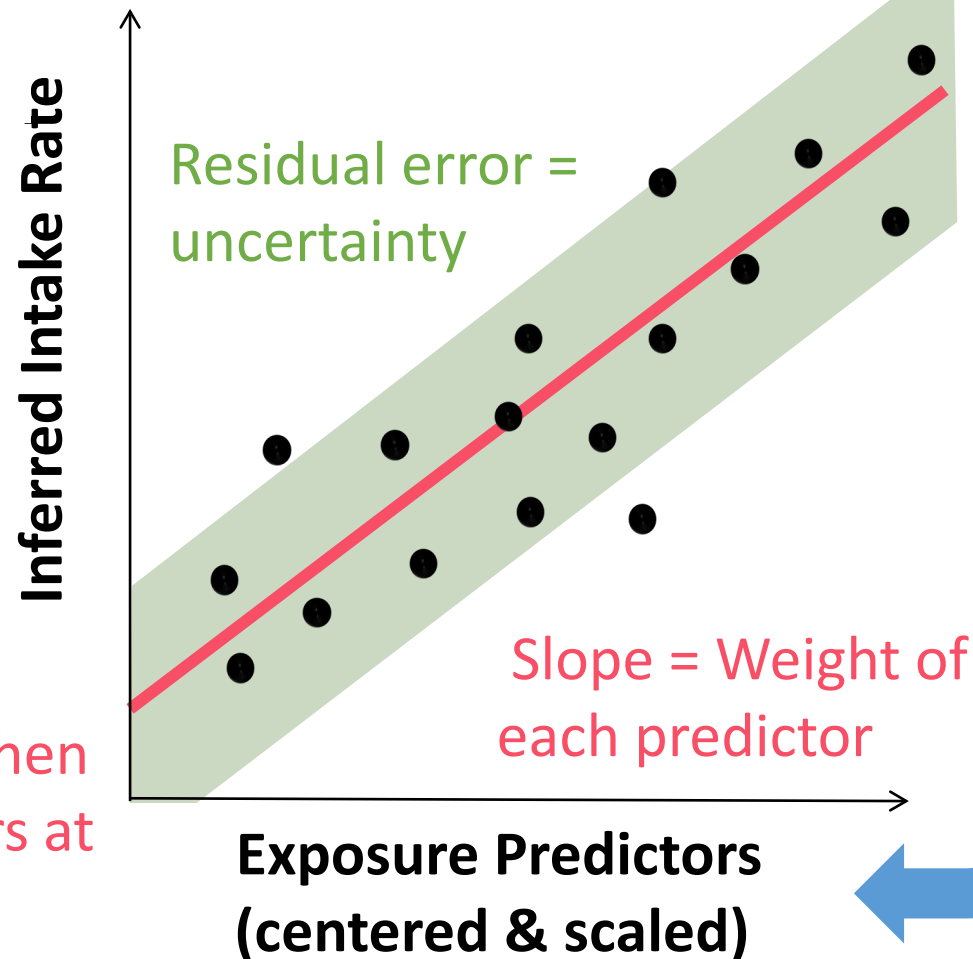
Train model on inferred  
exposures from NHANES  
biomonitoring data



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

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

**SEEM3 is a multiple linear regression!**

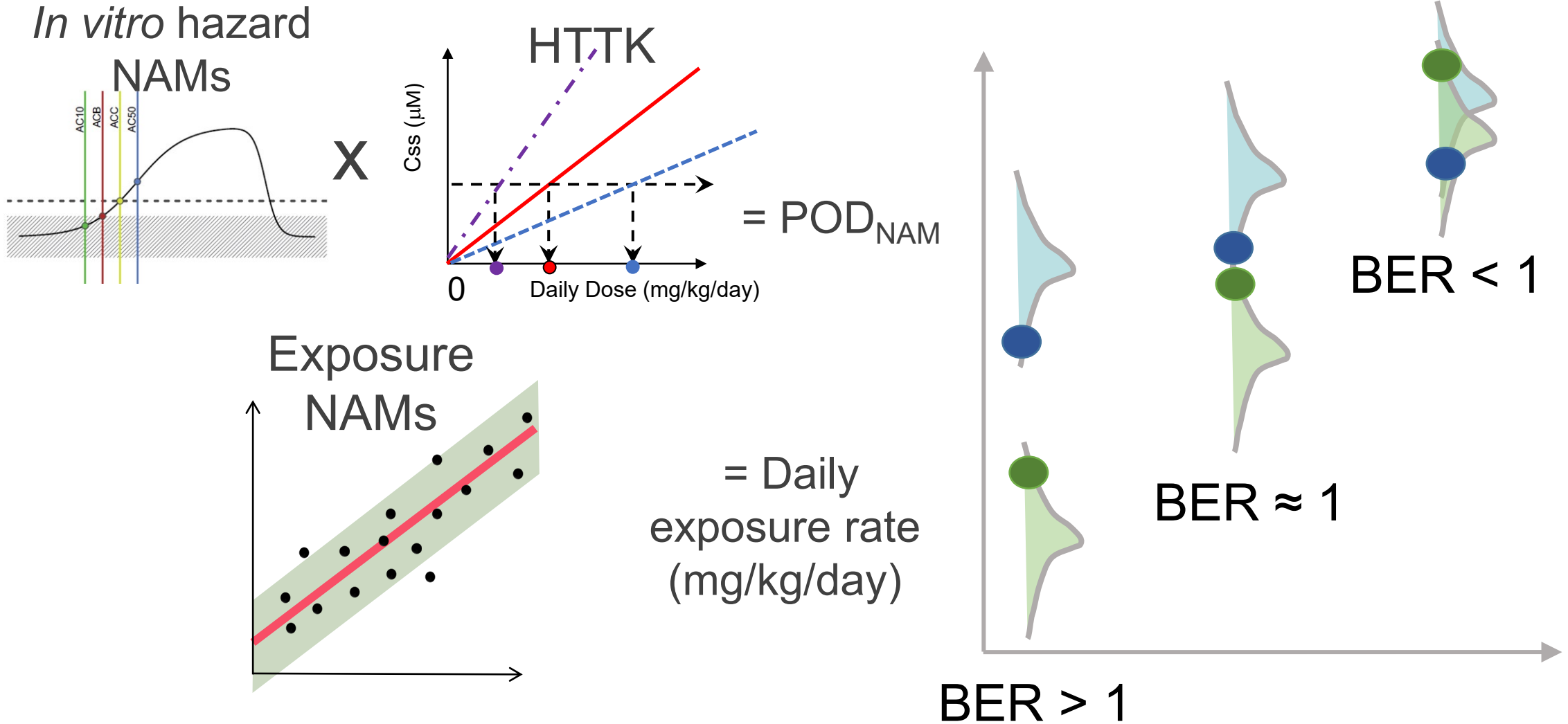


## **Exposure Predictors:**

- 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

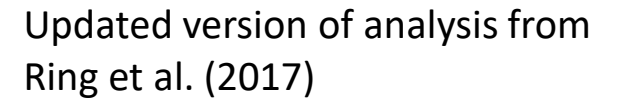
Missing predictor data:  
Impute mean

# Bioactivity-exposure ratio (BER):

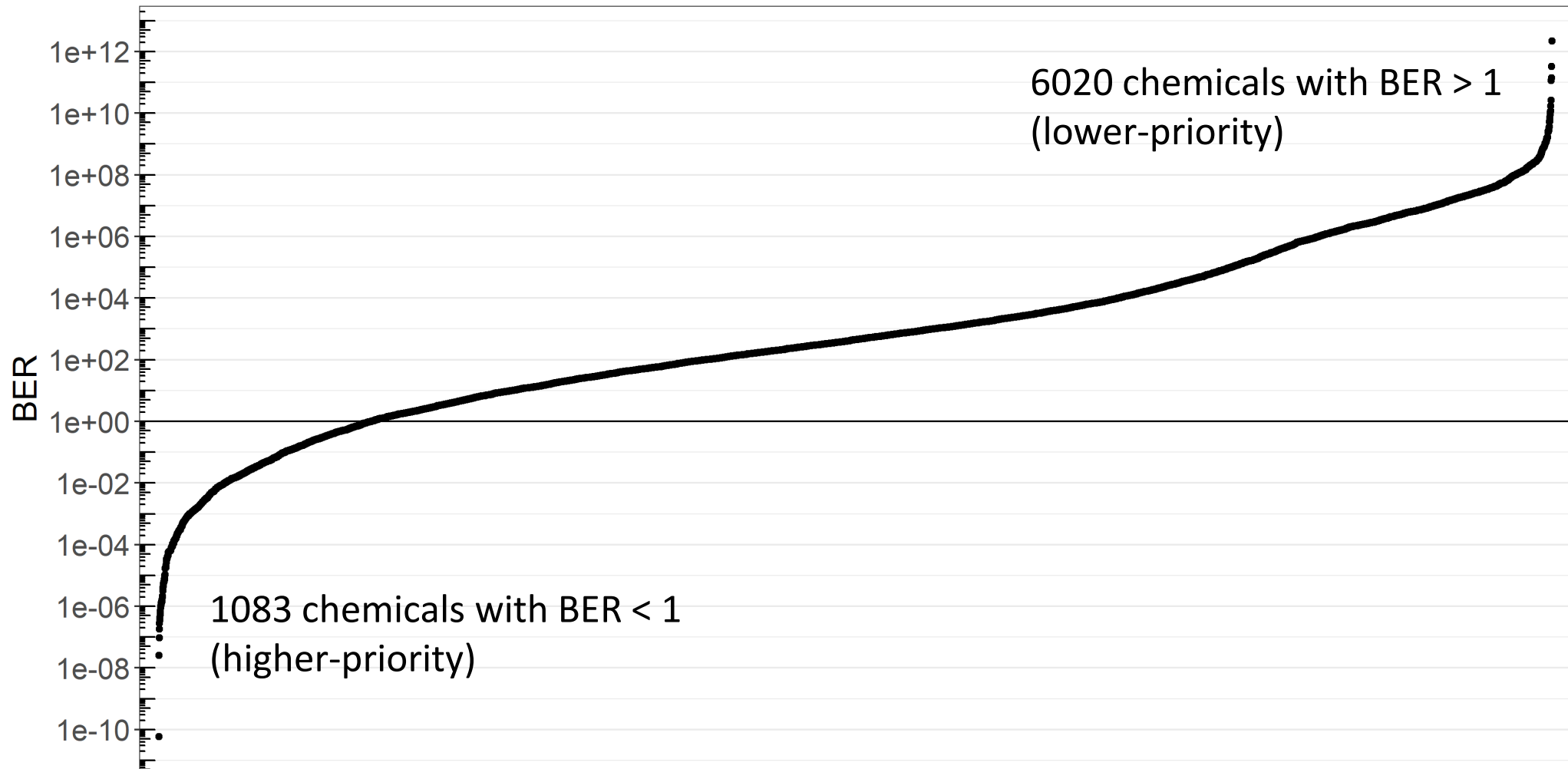
$$\text{POD}_{\text{NAM}} \div \text{NAM-predicted exposure}$$


Wetmore et al. (2015)

Paul Friedman et al. (2017)

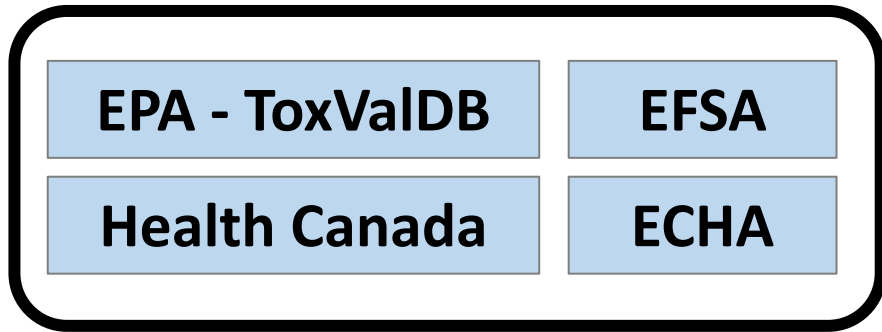


An even-more high-throughput application: BER prioritization of 7104 chemicals based on HTTK IVIVE of ToxCast AC50s and HT exposure predictions from SEEM3 model

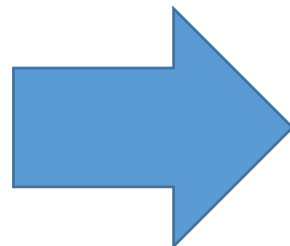
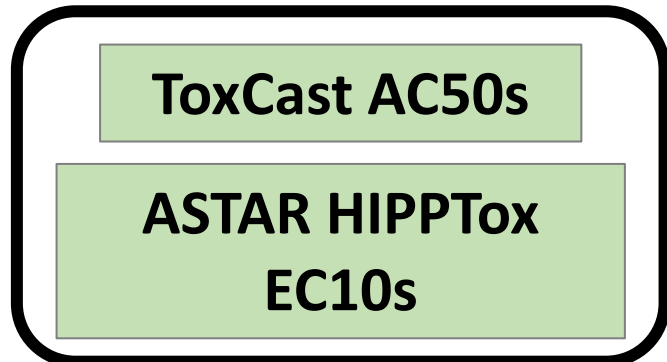


# Case study: Is $POD_{NAM}$ health-protective vs. traditional *in vivo* POD? (Paul Friedman et al. 2019)

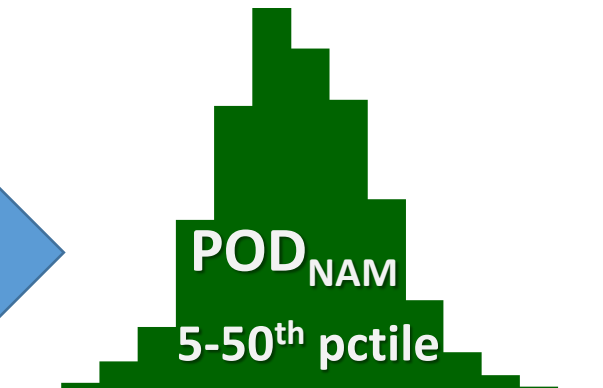
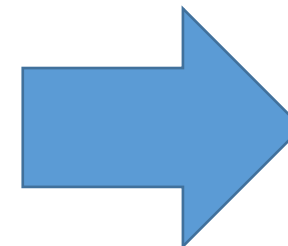
*In vivo* PODs (mg/kg/day)



NAM-based *in vitro* bioactive conc. ( $\mu$ M)



HTTK



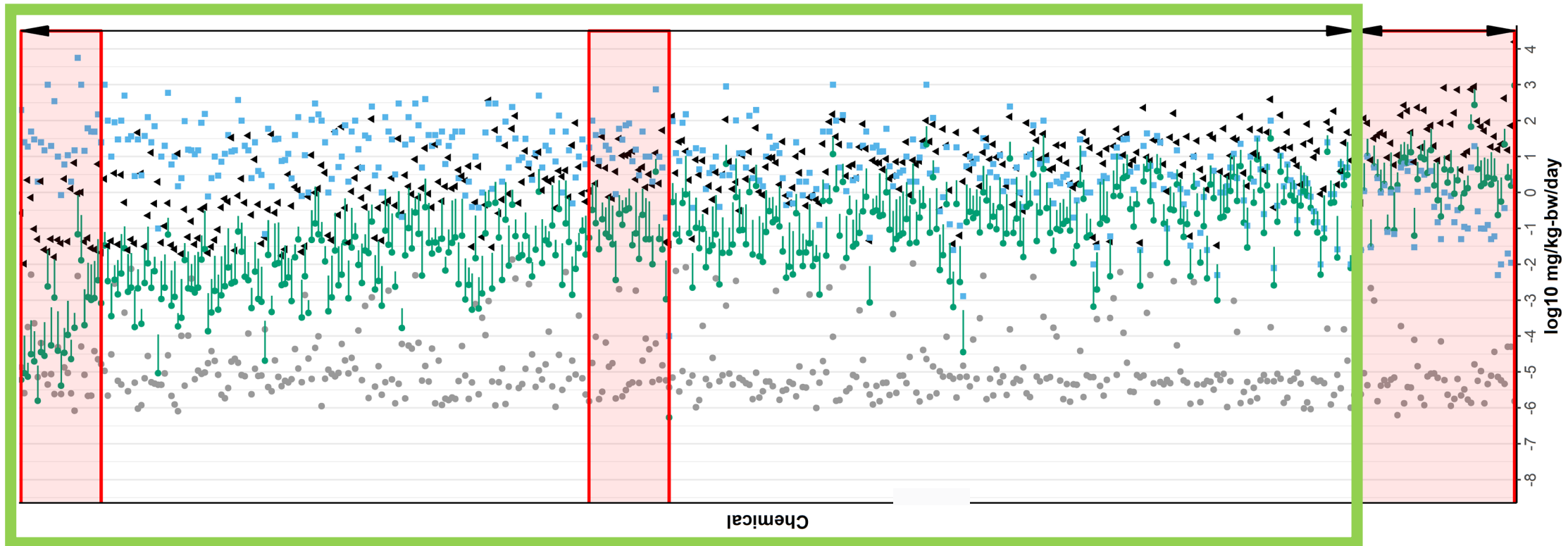
$POD_{NAM} < POD_{traditional}$   
(most of the time)

Adapted from Figure 3, Paul Friedman et al. 2019

$POD_{traditional}$

$POD_{NAM}$  (median –  
most-sensitive 5%)

NAM-pred exposure



400/448 chemicals = 89% of the time,  
 $POD_{NAM} < POD_{trad}$

When  $POD_{NAM} > POD_{traditional}$ ,  
specific chemical features are  
more likely

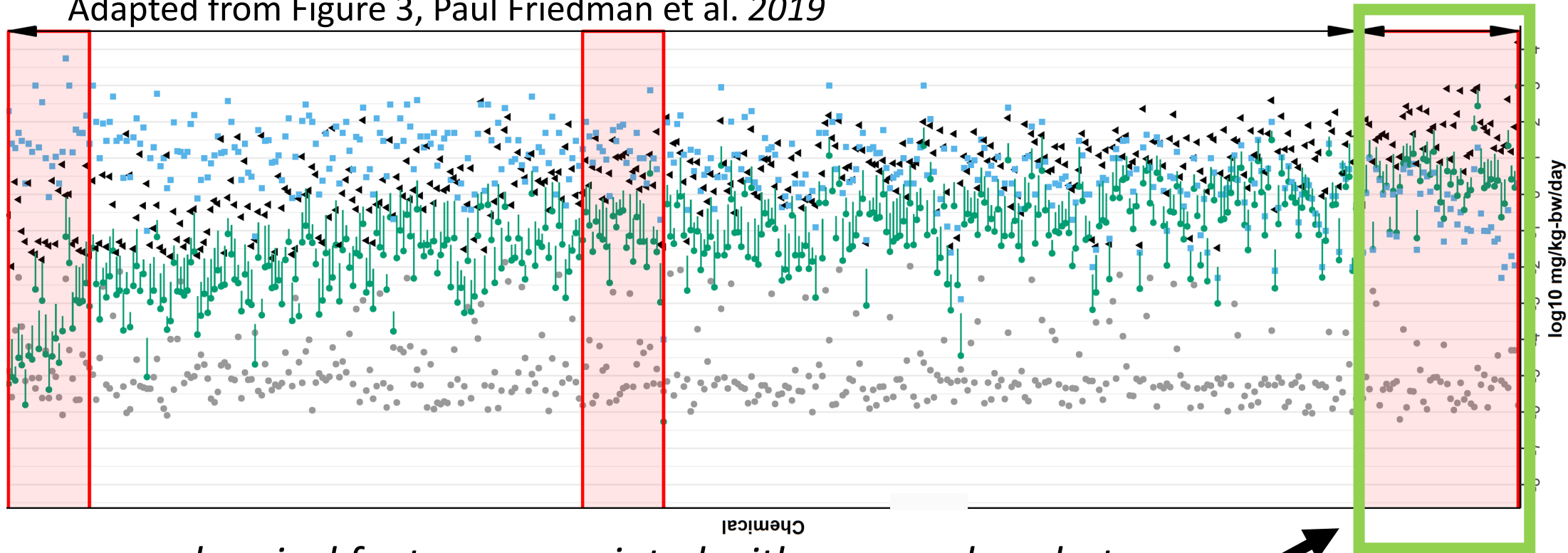
$POD_{traditional}$

$POD_{NAM}$  (median –  
most-sensitive 5%)

NAM-pred exposure

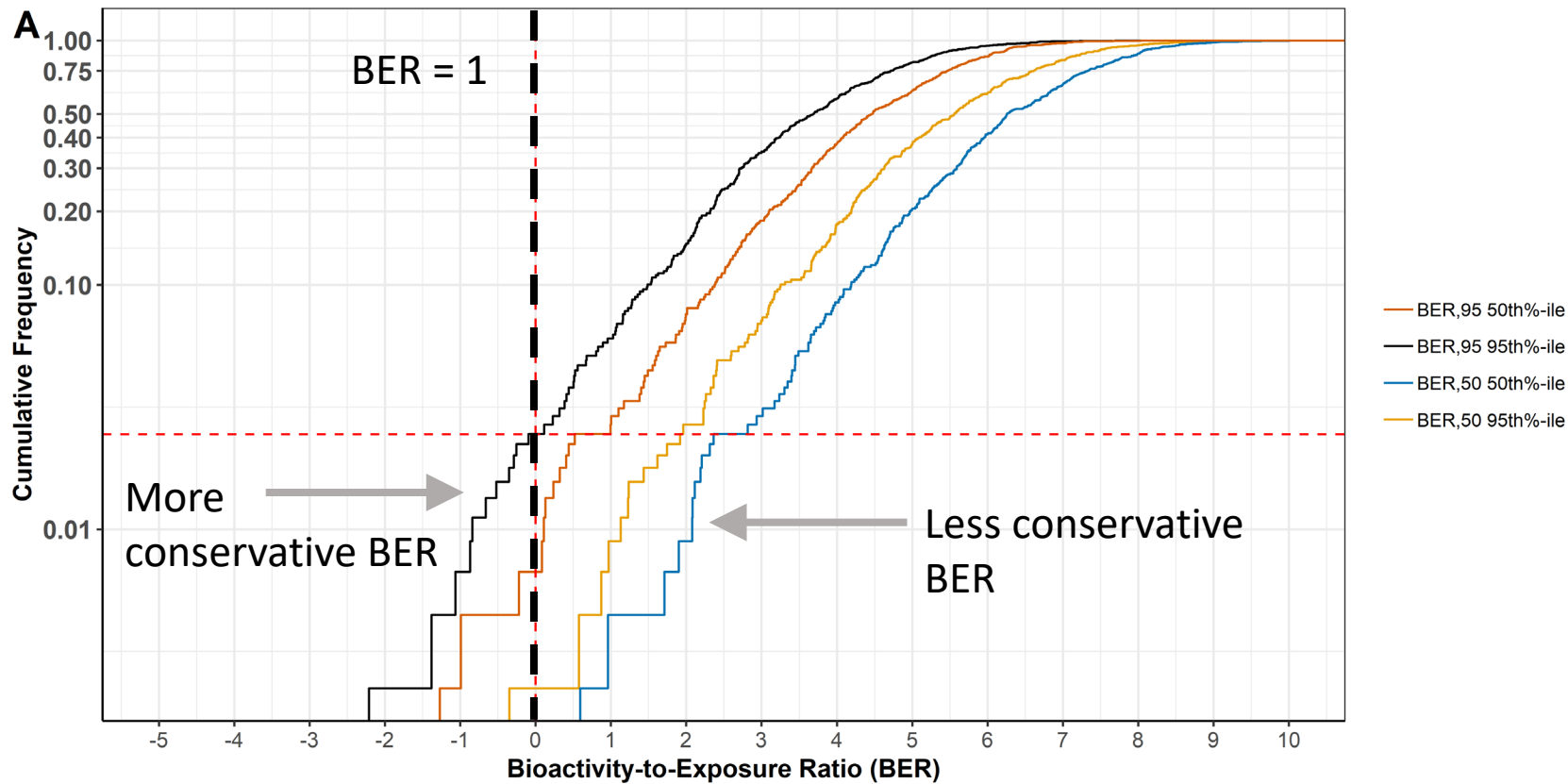


Adapted from Figure 3, Paul Friedman et al. 2019

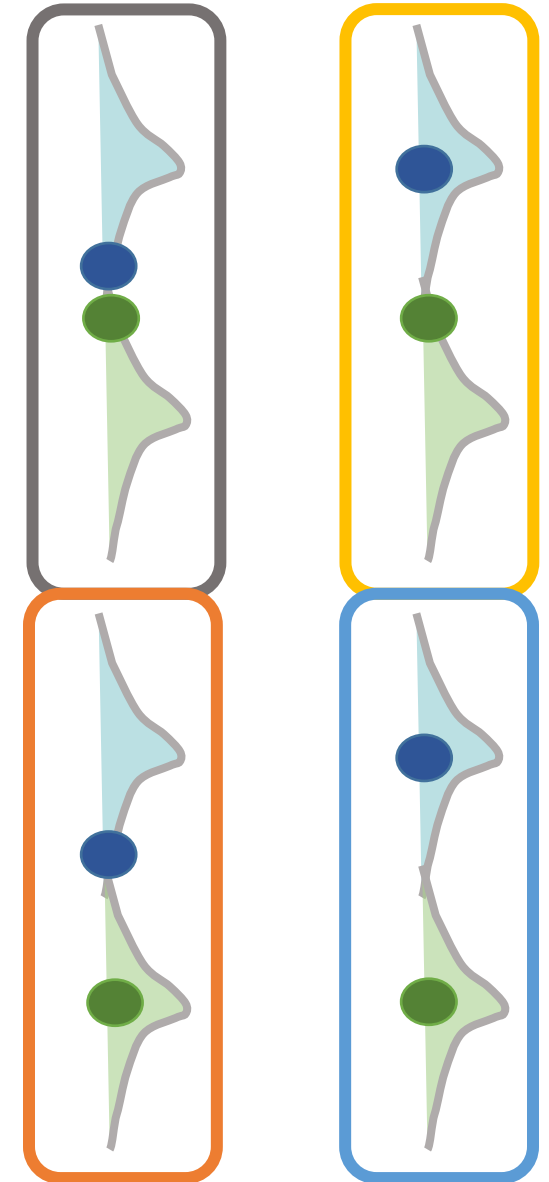


*chemical features associated with organophosphate  
pesticides and carbamates more likely*

BERs can be calculated to reflect different levels of uncertainty/conservatism based on use case



Adapted from Figure 4, Paul Friedman et al. 2019



# Conclusions

- Risk-based prioritization involves both hazard and exposure
- NAMs can help fill data gaps in both the hazard and the exposure components of risk
- *In vitro* bioactive concentrations can be converted to equivalent “in vivo”  $POD_{NAM}$  doses, using high-throughput toxicokinetics (HTTK)
- $POD_{NAM}$  incorporates data-driven inter-individual TK variability via HTTK
- Exposure NAMs can rapidly predict median population aggregate exposures
- $POD_{NAM}$  vs. exposure quantified using *bioactivity-exposure ratio* (BER)
- BER-based prioritization gives a useful starting point
- $POD_{NAM}$  is typically more conservative than traditional *in vivo* PODs
- BER-based approach allows flexible consideration of uncertainty

***NAMs for hazard, exposure, and toxicokinetics provide a useful way to rapidly prioritize chemicals***

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