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Exposure-based Chemical Priority Setting in the 21st Century

Course No: H02-014
Credit: 2 PDH

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This course was adapted from the U.S. Environmental Protection Agency (EPA), Publication No. 600H20234, “Exposure-based Chemical Priority Setting in the 21st Century”, which is in the public domain.

Exposure-based Chemical Priority Setting in the 21st Century

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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



US EPA Office of Research and Development

- The Office of Research and Development (ORD) is the scientific research arm of EPA
 - 562 peer-reviewed journal articles in 2018
- Research is conducted by ORD's four national centers, and three offices organized to address:
 - Public health and env. assessment; comp. tox. and exposure; env. measurement and modeling; and env. solutions and emergency response.
- 13 facilities across the United States
- Research conducted by a combination of Federal scientists (including uniformed members of the **Public Health Service**); contract researchers; and postdoctoral, graduate student, and post-baccalaureate trainees



Credit: the Research Triangle Foundation

ORD Facility in
Research Triangle Park, NC

Chemical Regulation in the United States

- Park *et al.* (2012): At least 3221 chemical signatures in pooled human blood samples, many appear to be exogenous
- A tapestry of laws covers the chemicals people are exposed to in the United States (Breyer, 2009)
- Chemical safety testing is primarily for food additives, pharmaceuticals, and pesticide active ingredients (NRC, 2007)
 - Different levels depending on category

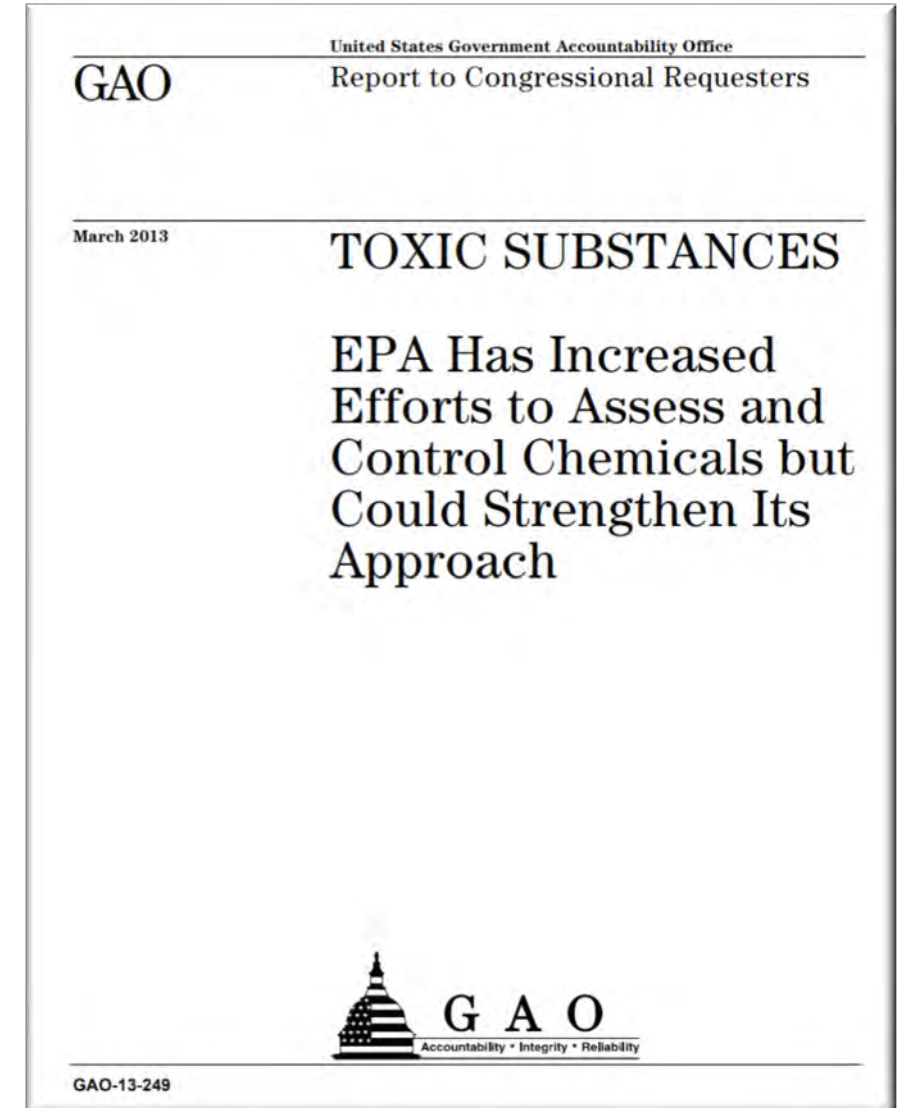


Chemical Regulation in the United States

- Most other chemicals, ranging from industrial waste to dyes to packing materials, are covered by the Toxic Substances Control Act (TSCA)
- Thousands of chemicals on the market were “grandfathered” in without assessment
Judson et al. (2009), Egeghy et al. (2012), Wetmore et al. (2015)

“Tens of thousands of chemicals are listed with the Environmental Protection Agency (EPA) for commercial use in the United States, with an average of 600 new chemicals listed each year.”

U.S. Government Accountability Office

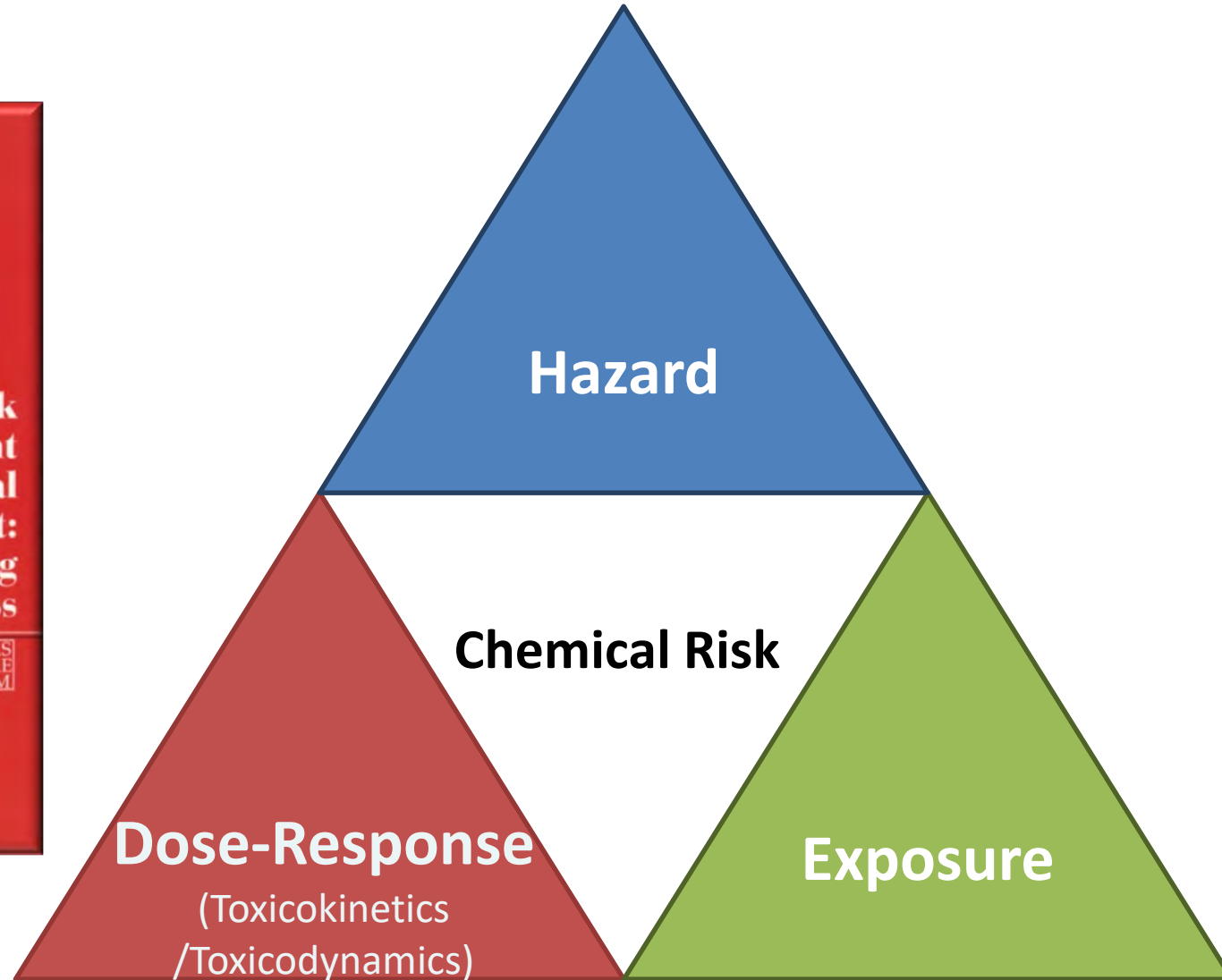


March, 2013

Three Components for Chemical Risk



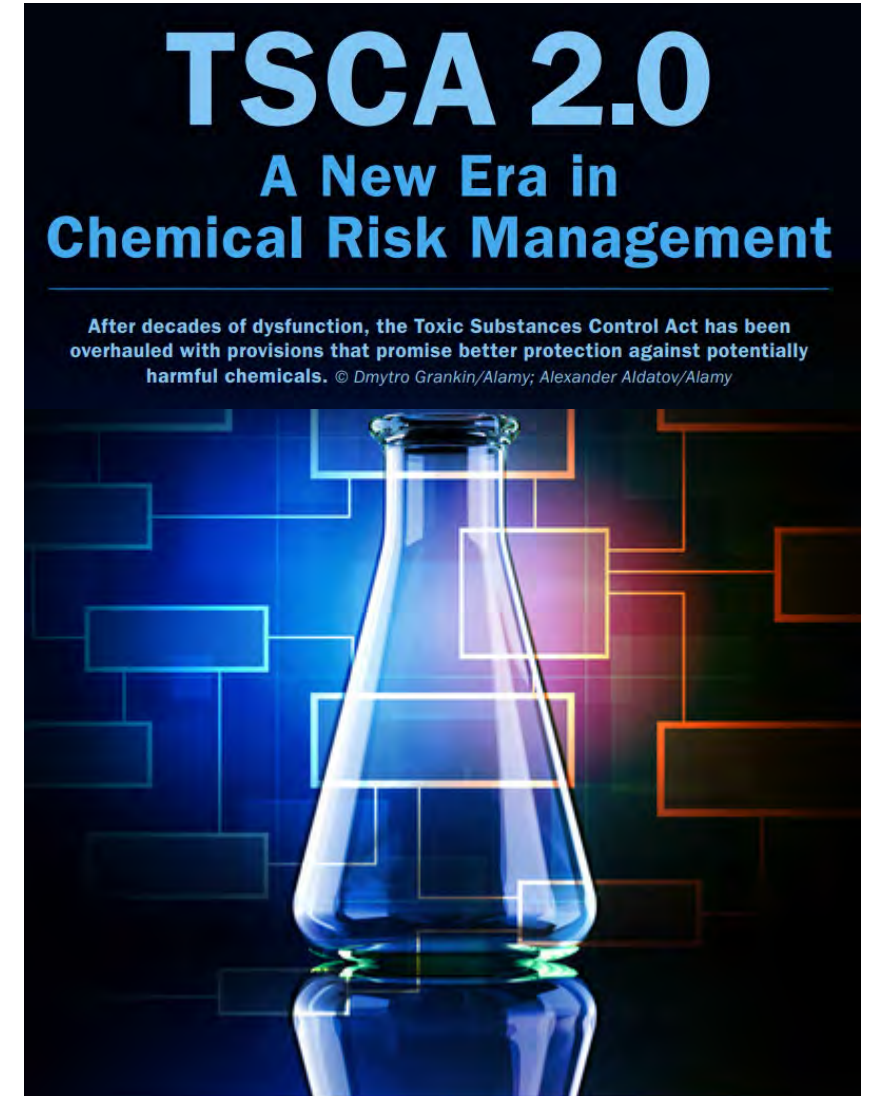
NRC (1983)



The National Academy of Sciences, Engineering and Medicine (1983) outlined three components for determining chemical risk.

Toxic Substances Control Act (TSCA)

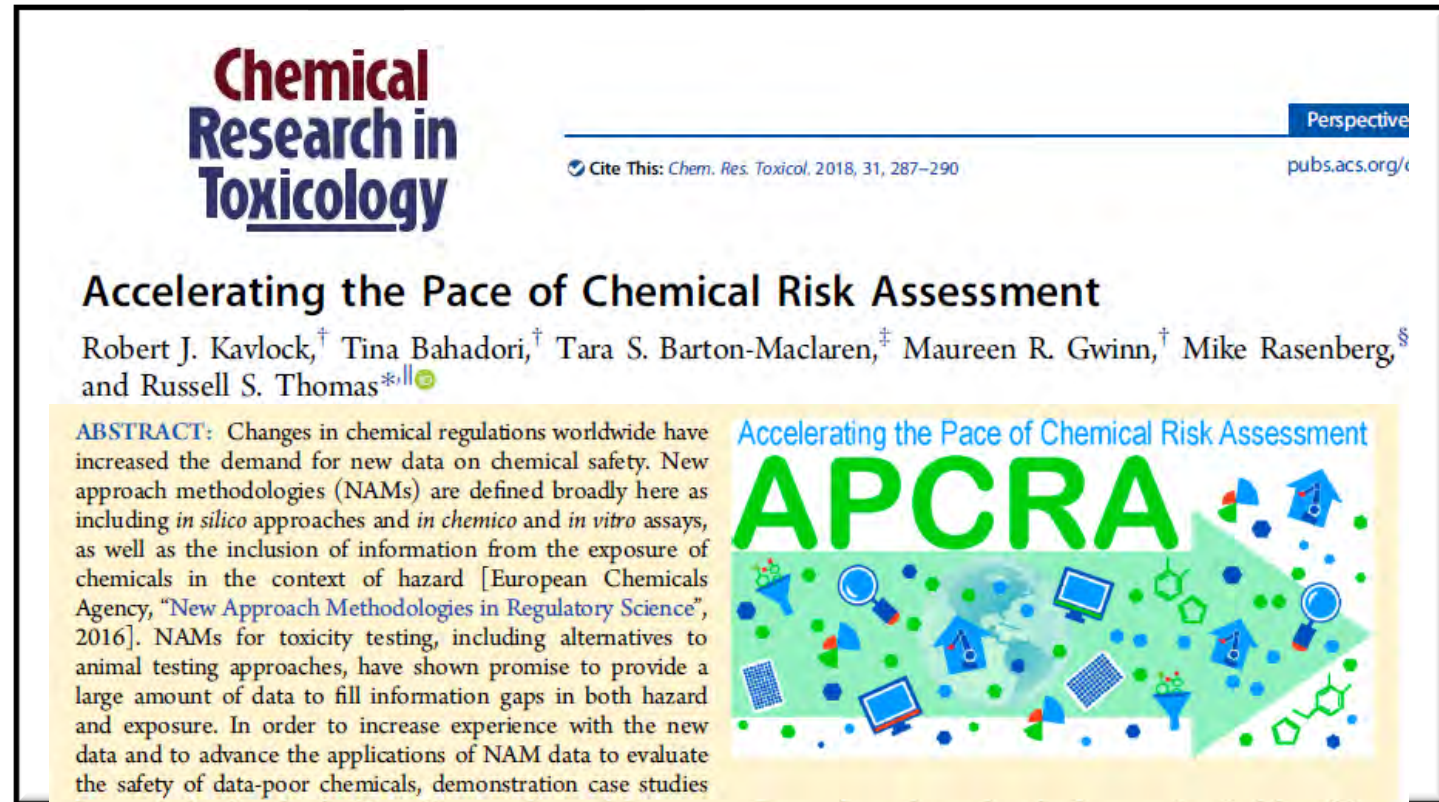
- TSCA was updated in June, 2016 to allow more rapid evaluation of chemicals (Frank R. Lautenberg Chemical Safety for the 21st Century Act)
- New approach methodologies (NAMs) are being considered to inform prioritization of chemicals for testing and evaluation (Kavlock et al., 2018)
- EPA has released a “A Working Approach for Identifying Potential Candidate Chemicals for Prioritization” (September, 2018)



Schmidt, C. W. (2016). TSCA 2.0: A new era in chemical risk management”, Environmental Health Perspectives, A182-A186.


New Approach Methodologies (NAMs)

- There are roughly 10,000 TSCA-relevant chemicals in commerce
 - Traditional methods are too resource-intensive to address all of these
- NAMs include:
 - High throughput screening (ToxCast)
 - High throughput exposure estimates (ExpoCast)
 - High throughput toxicokinetics (HTTK)
- TSCA Proof of concept: Examine ~200 chemicals with ToxCast, ExpoCast and HTTK
 - HTTK was rate limiter on number of chemicals
 - *“A Proof-of-Concept Case Study Integrating Publicly Available Information to Screen Candidates for Chemical Prioritization under TSCA”*



Replacing Animal Testing with NAMs

- Administrator of the EPA: “To aggressively pursue a reduction in animal testing, I am directing leadership and staff in the Office of Chemical Safety and Pollution Prevention and the Office of Research and Development [ORD] to prioritize ... the reduction of animal testing while ensuring protection of human health and the environment.”
- “These new approach methods (NAMs), include any technologies, methodologies, approaches or combinations thereof that can be used to provide information on chemical hazard and potential human exposure that can avoid or significantly reduce the use of testing on animals”
 - NAMs for filling information gaps for decision-making
 - integrating data streams into chemical risk assessment
 - making the information publicly available



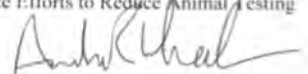
UNITED STATES ENVIRONMENTAL PROTECTION AGENCY
WASHINGTON, D.C. 20460

September 10, 2019

THE ADMINISTRATOR

MEMORANDUM

SUBJECT: Directive to Prioritize Efforts to Reduce Animal Testing

FROM: Andrew R. Wheeler 
Administrator

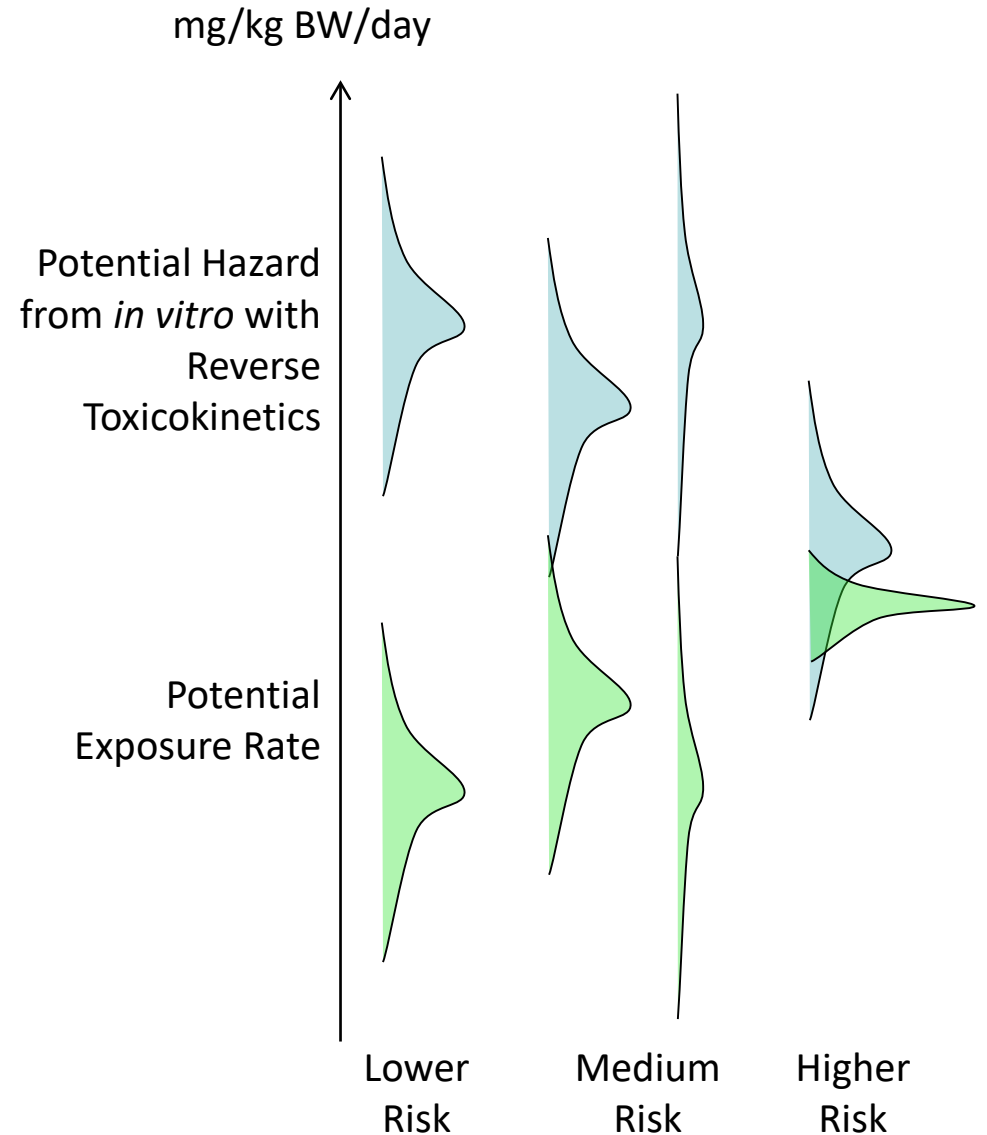
TO: Associate Deputy Administrator
General Counsel
Assistant Administrators
Inspector General
Chief Financial Officer
Chief of Staff
Associate Administrators
Regional Administrators

During my March 2019 all-hands address, I reiterated the U.S. Environmental Protection Agency's commitment to move away from animal testing. We are already making significant efforts to reduce, replace and refine our animal testing requirements under both statutory and strategic directives. For example, the *Toxic Substances Control Act*, amended June 22, 2016, by the Frank R. Lautenberg Chemical Safety for the 21st Century Act, requires the EPA to reduce reliance on animal testing. Also, Objective 3.3 of the *FY 2018-2022 U.S. EPA Strategic Plan* outlines a commitment to further reduce the reliance on animal testing within five years. More than 200,000 laboratory animals have been saved in recent years as a result of these collective efforts.

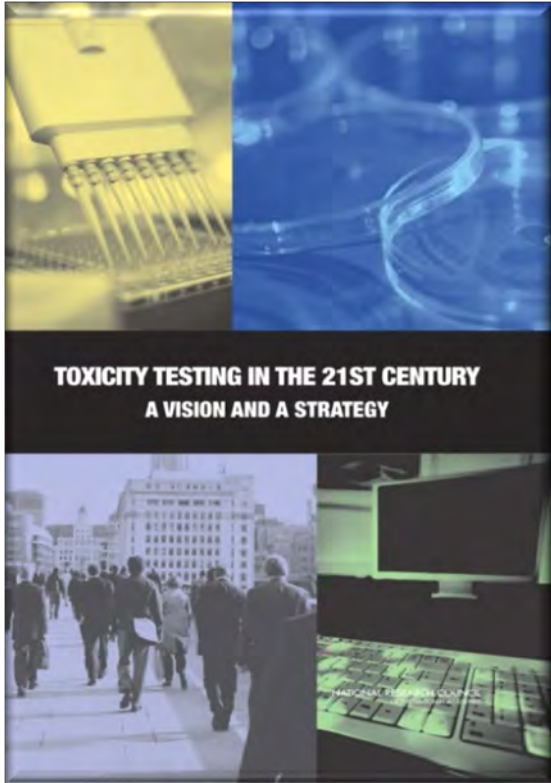
Scientific advancements exist today that allow us to better predict potential hazards for risk assessment purposes without the use of traditional methods that rely on animal testing. These new approach methods (NAMs), include any technologies, methodologies, approaches or combinations thereof that can be used to provide information on chemical hazard and potential human exposure that can avoid or significantly reduce the use of testing on animals. The benefits of NAMs are extensive, not only allowing us to decrease animals used while potentially evaluating more chemicals across a broader range of potential biological effects, but in a shorter timeframe with fewer resources while often achieving equal or greater biological predictivity than current animal models.

Chemical Risk = Hazard x Exposure

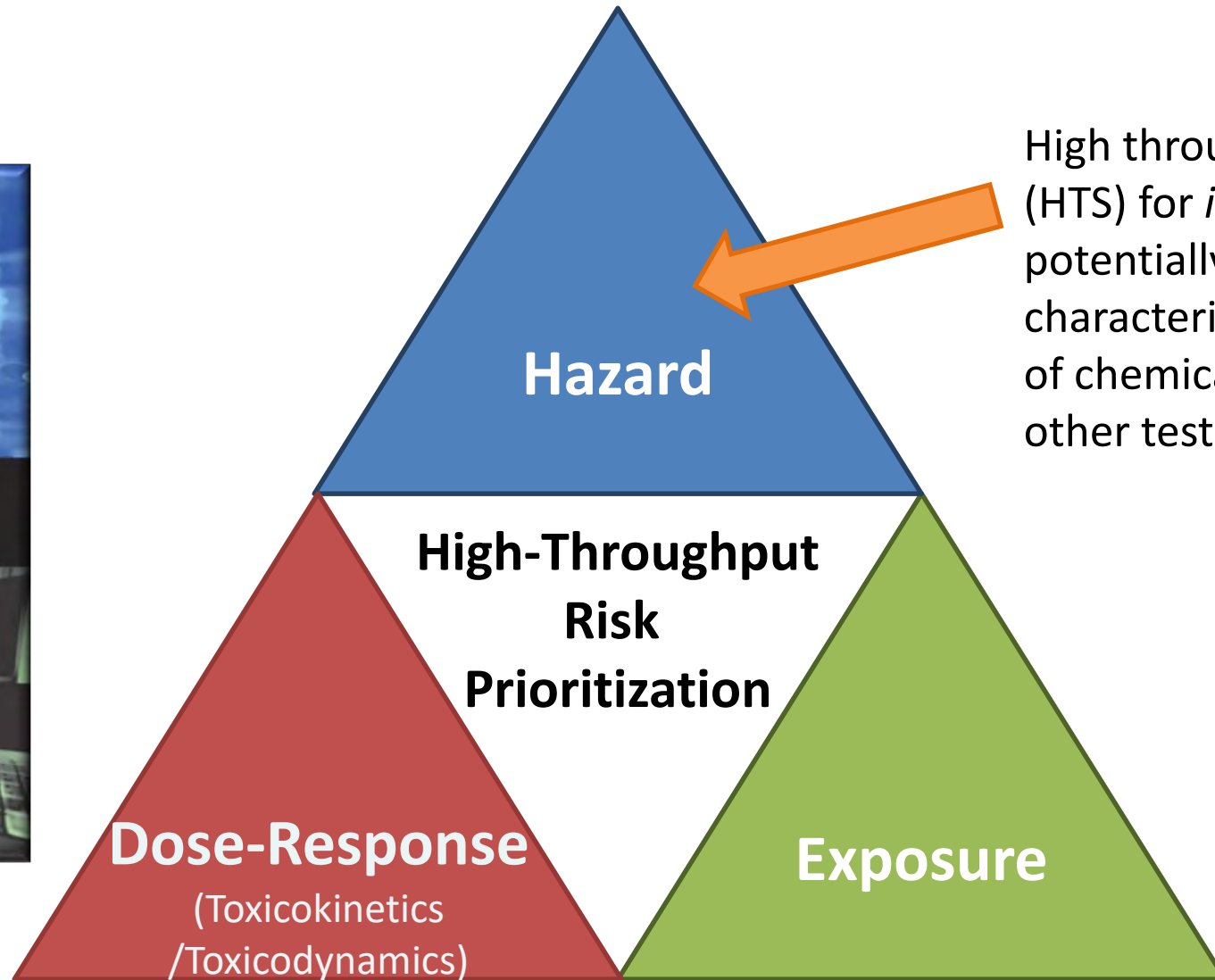
- The U.S. National Research Council (1983) identified chemical risk as a function of both inherent hazard and exposure
- Therefore, high throughput risk prioritization needs:
 1. High throughput hazard characterization (Dix et al., 2007, Collins et al., 2008)
 2. High throughput exposure forecasts (Wambaugh 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



NRC (2007)



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

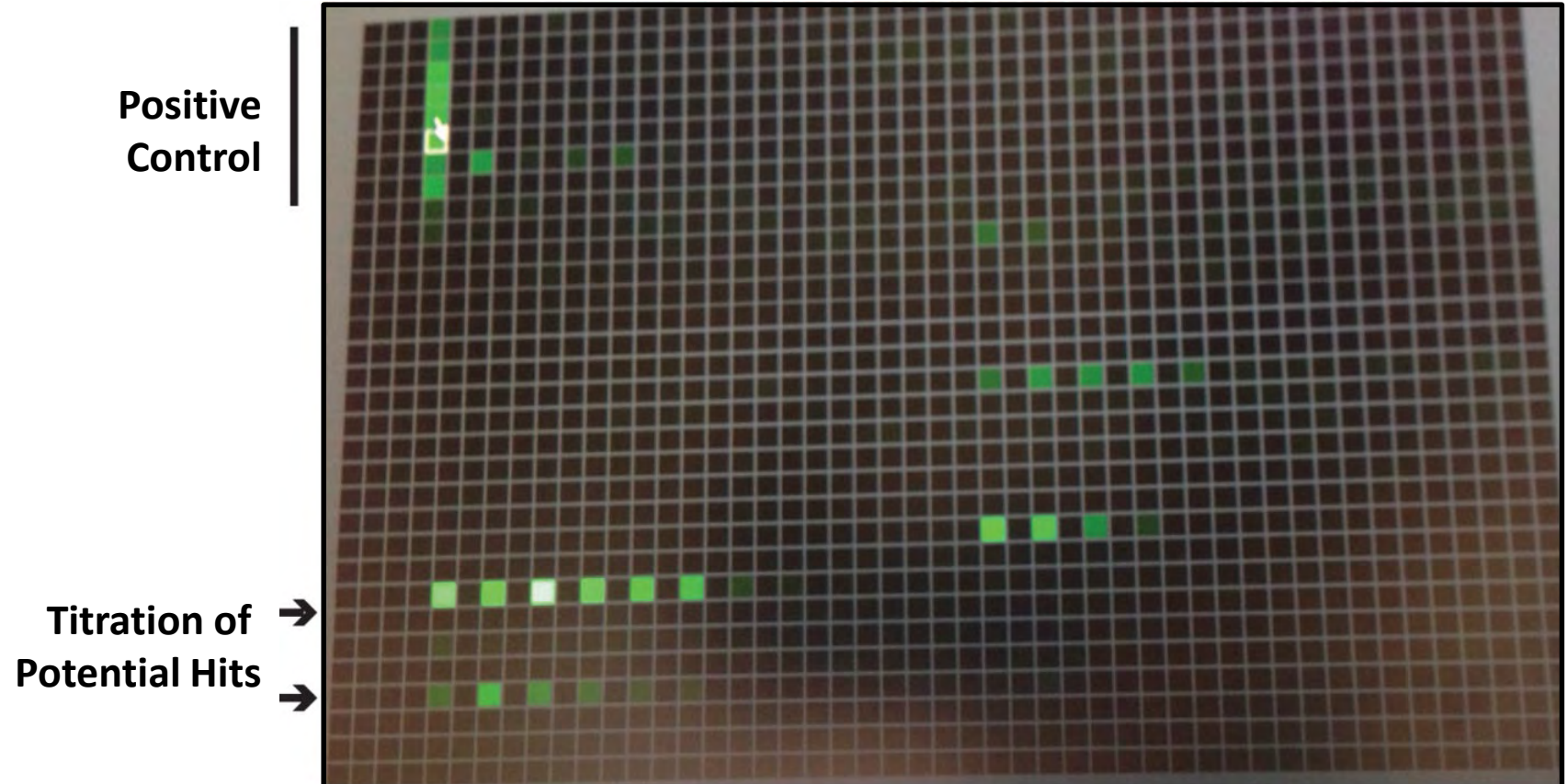
To perform high throughput risk prioritization, we need all three components

High-throughput Screening

Hertzberg and Pope (2000):

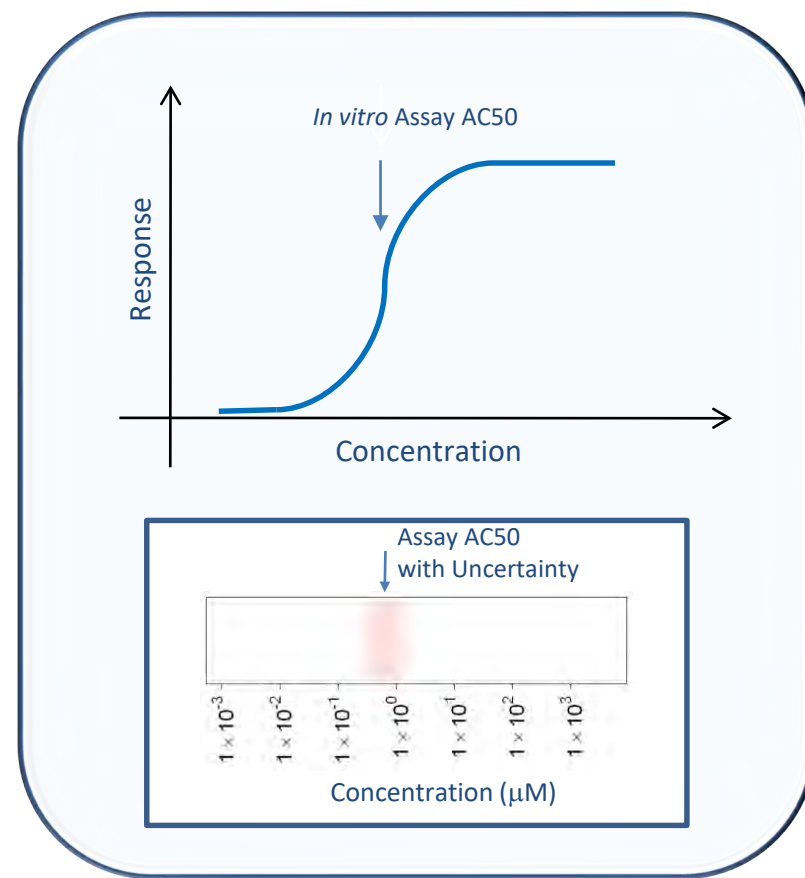
- “New technologies in high-throughput screening have significantly increased throughput and reduced assay volumes...”
- “...new fluorescence methods, detection platforms and liquid-handling technologies.”
- Typically assess many chemicals with a signal readout (e.g., green fluorescent protein).

Kaewkhaw et al. (2016)

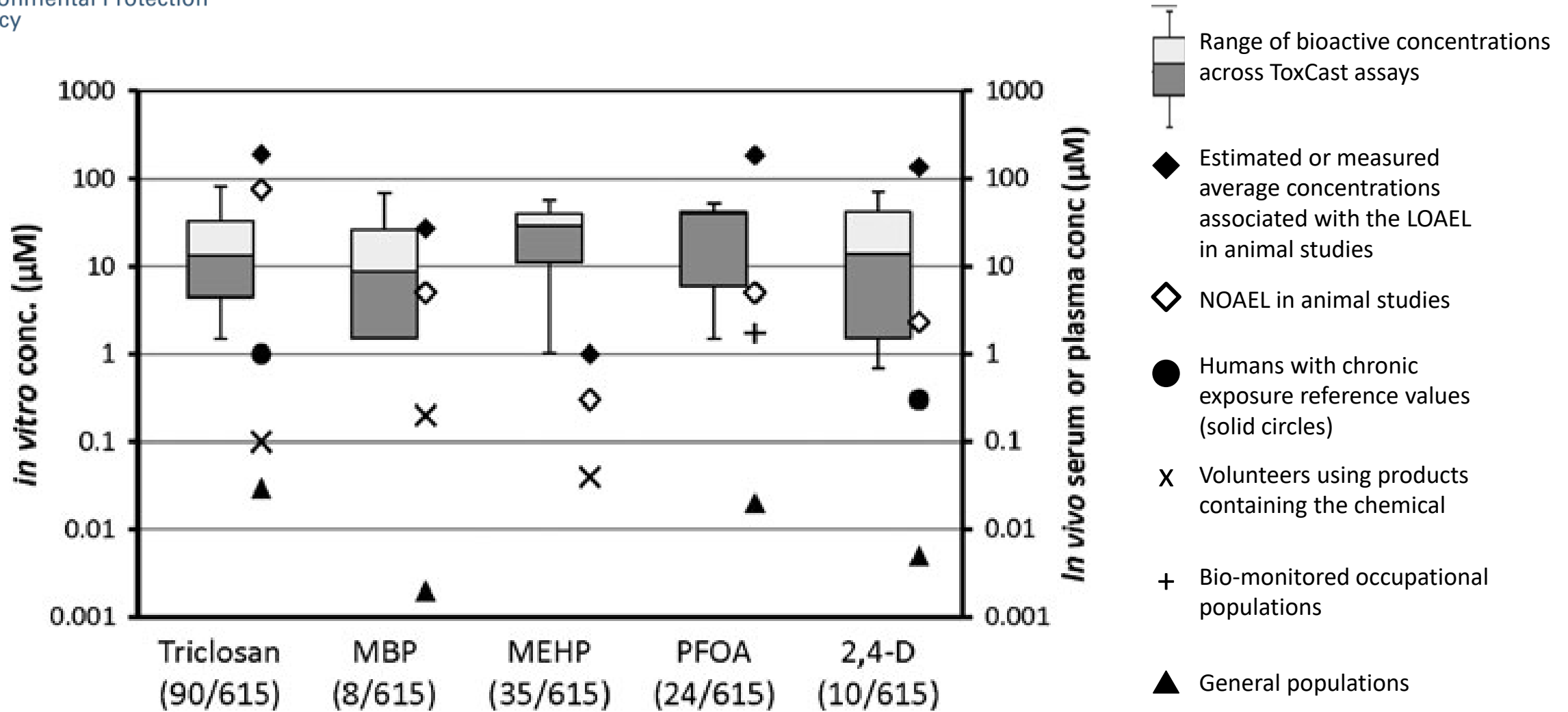


High-Throughput Bioactivity Screening Projects

- We attempt to estimate points of departure *in vitro* using high throughput screening (HTS)
- **Tox21:** Examining >8,000 chemicals using ~50 assays intended to identify interactions with biological pathways (Schmidt, 2009)
- **ToxCast:** For a subset (>2000) of Tox21 chemicals ran >1100 additional assays (Kavlock *et al.*, 2012)
- Most assays conducted in dose-response format (identify 50% activity concentration – AC_{50} – and efficacy if data described by a Hill function, Filer *et al.*, 2016)
- All data are public: <http://comptox.epa.gov/dashboard/>



The Margin Between Exposure and Hazard

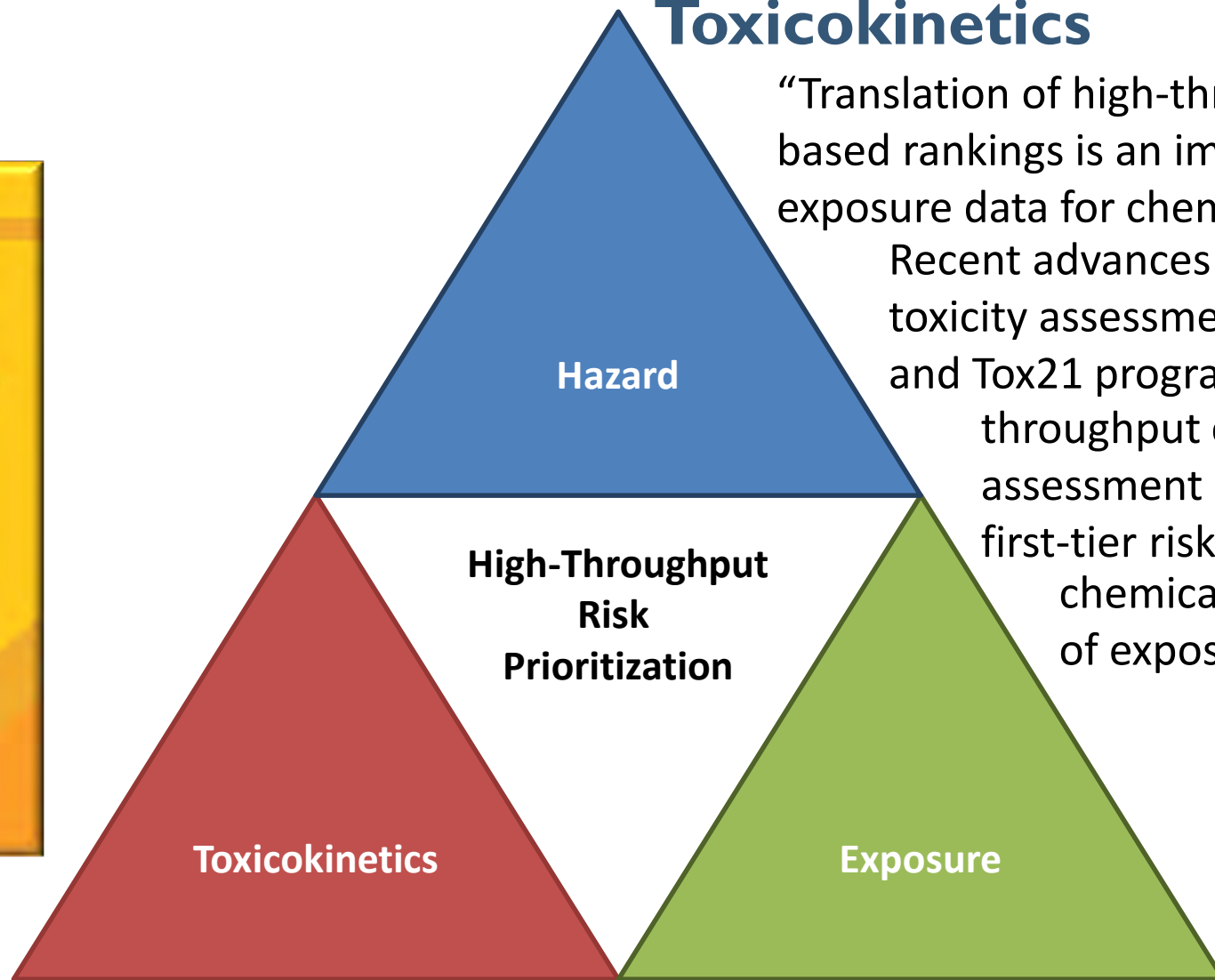


The five chemicals (as of 2011) with plasma biomonitoring AND ToxCast data... what do we do about the other 1000's?

Most Chemicals Lack Data on Exposure and Toxicokinetics



NASEM (2017)



“Translation of high-throughput data into risk-based rankings is an important application of exposure data for chemical priority-setting. Recent advances in high-throughput toxicity assessment, notably the ToxCast and Tox21 programs... and in high-throughput computational exposure assessment [ExpoCast] have enabled first-tier risk-based rankings of chemicals on the basis of margins of exposure” - National Academies of Sciences, Engineering, and Medicine (NASEM)

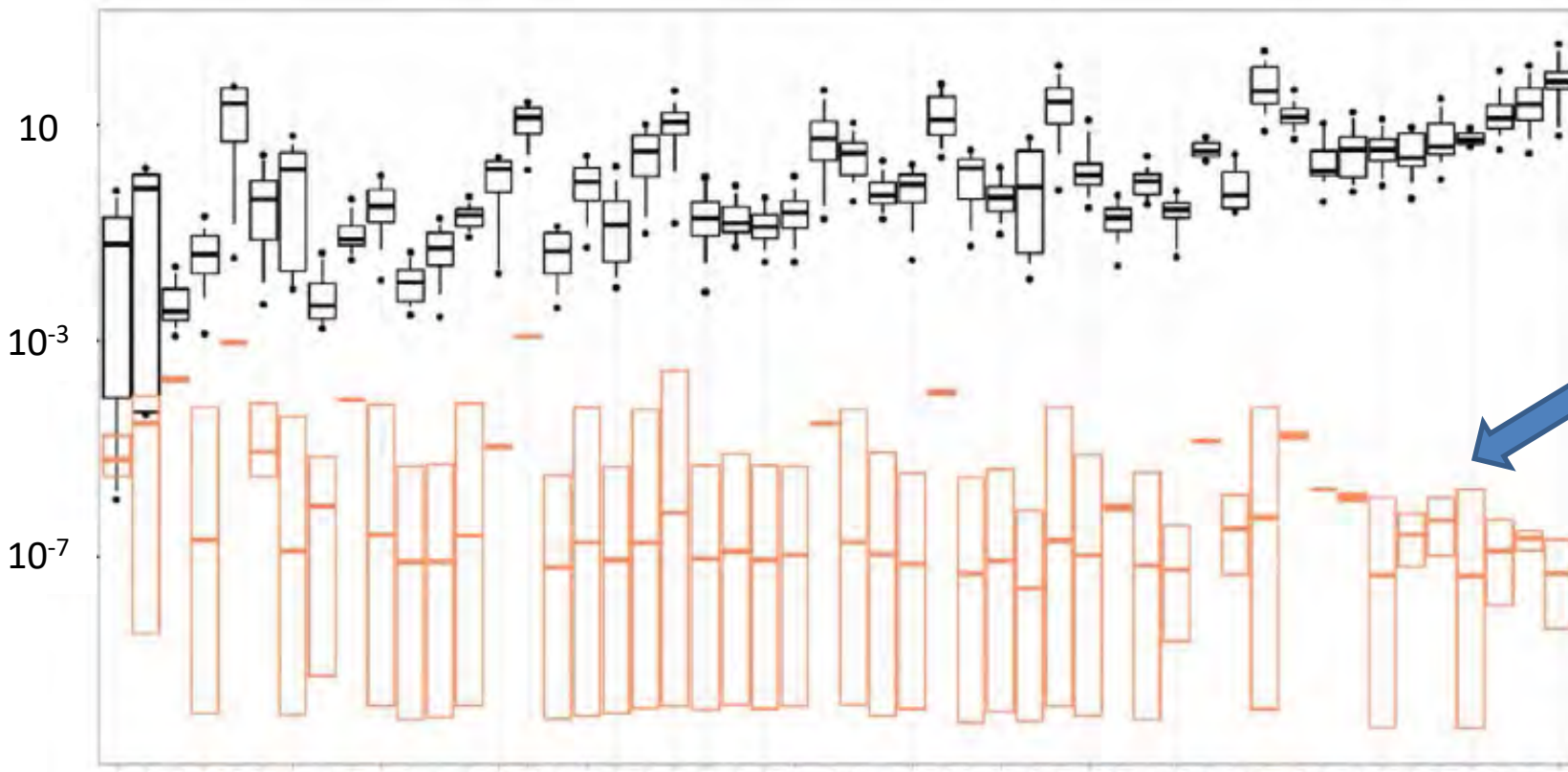
In order to perform risk-based ranking we need data on hazard, toxicokinetics, and exposure...

Chemical Prioritization NAMs

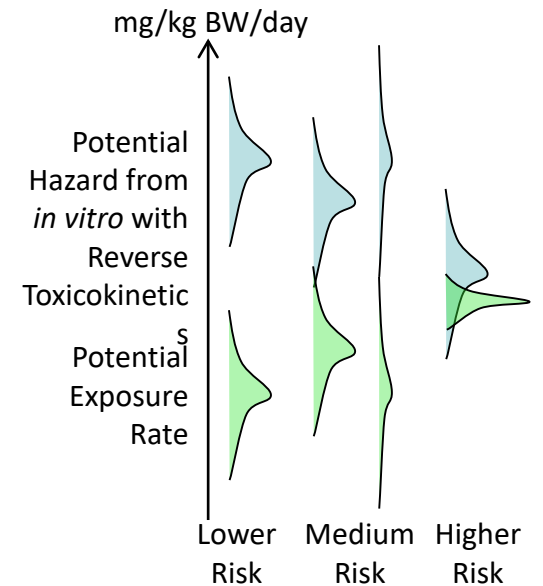
High throughput *in vitro* screening can estimate doses needed to cause bioactivity (e.g., Wetmore et al., 2015)

Exposure intake rates can be inferred from biomarkers (e.g., Ring et al., 2018)

Estimated Equivalent Dose or Predicted Exposure
(mg/kg BW/day)



Chemicals Monitored by CDC NHANES



Ring et al. (2017)

In Vitro - *In Vivo* Extrapolation (IVIVE)

IVIVE is the use of *in vitro* experimental data to predict phenomena *in vivo*

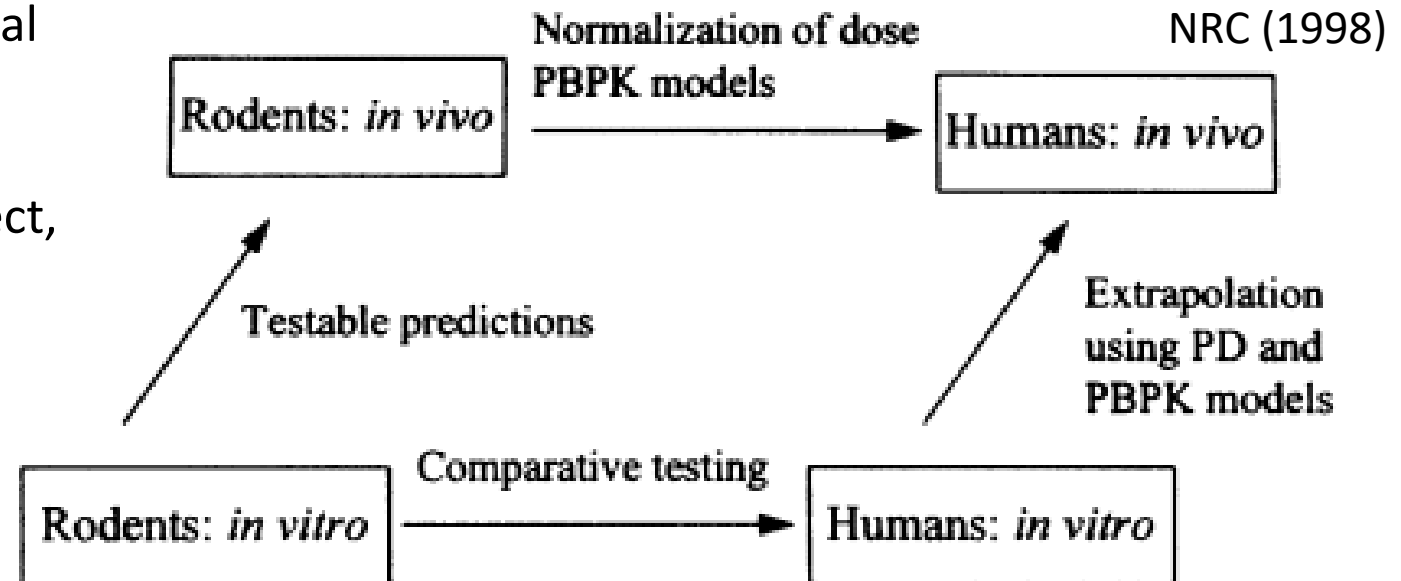
- IVIVE-PK/TK (**Pharmacokinetics/Toxicokinetics**):

- 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**):

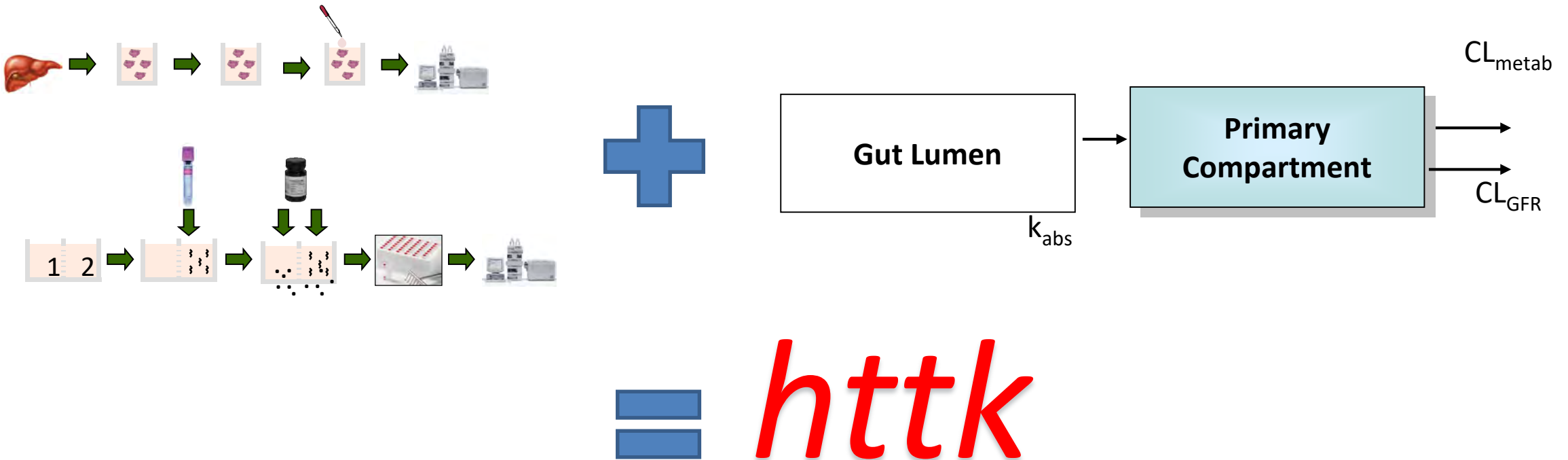
- Effect of molecules/chemicals at biological target *in vivo*
- Assay design/selection important
- Perturbation as adverse/therapeutic effect, reversible/ irreversible effects

- Both contribute to *in vivo* effect prediction



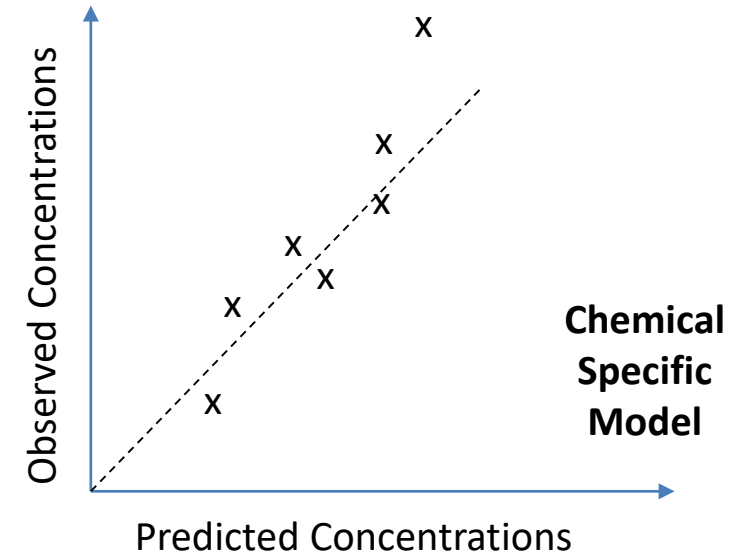
High Throughput Toxicokinetics (HTTK)

***In vitro* toxicokinetic data + generic toxicokinetic model
= high(er) throughput toxicokinetics**



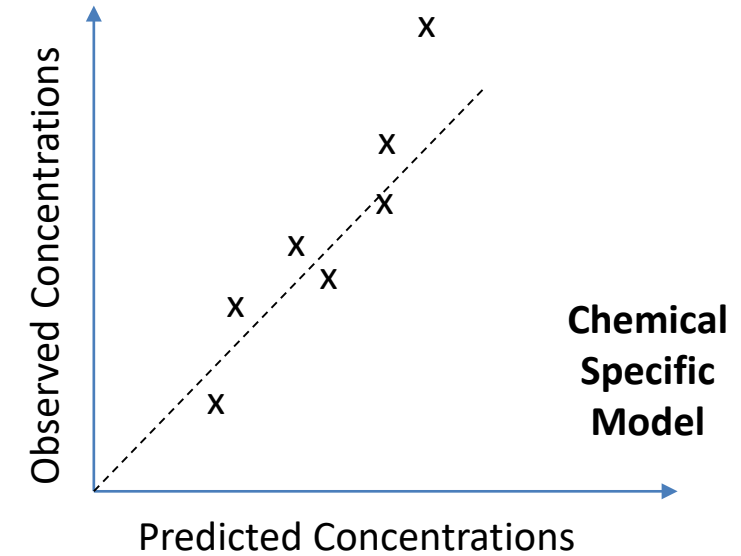
Building Confidence in TK Models

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



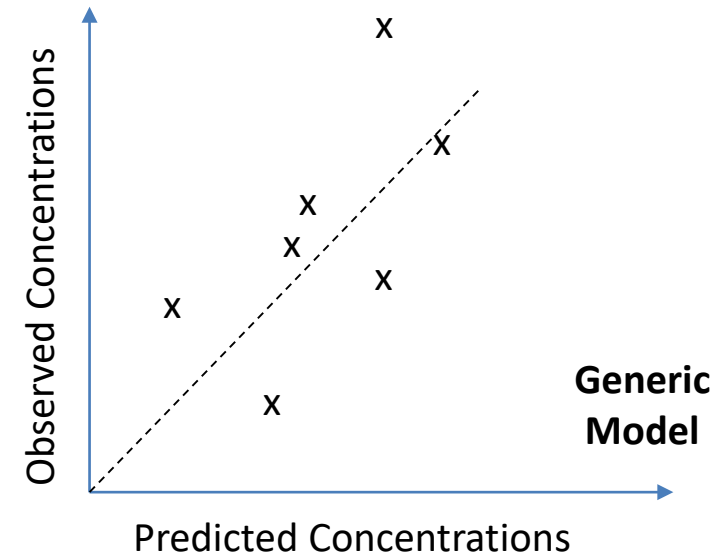
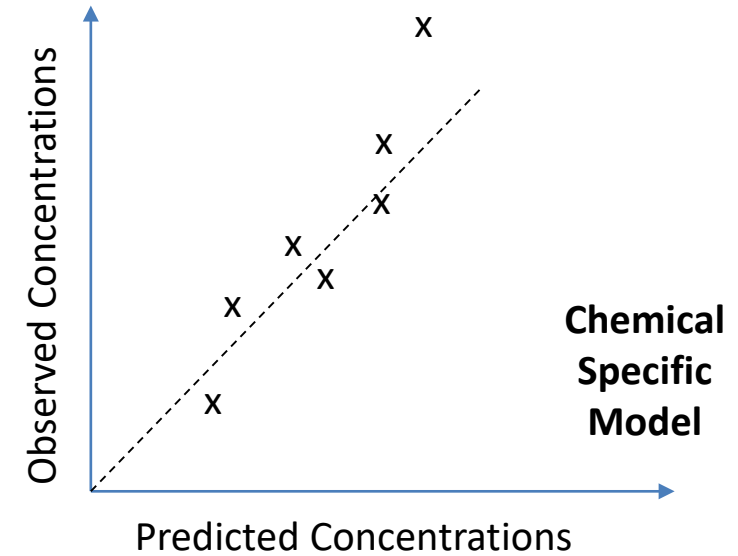
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- However, we do not typically have TK data



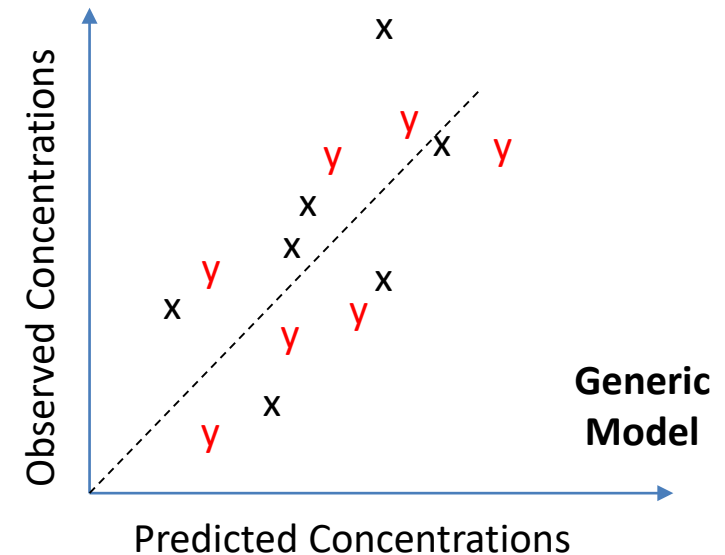
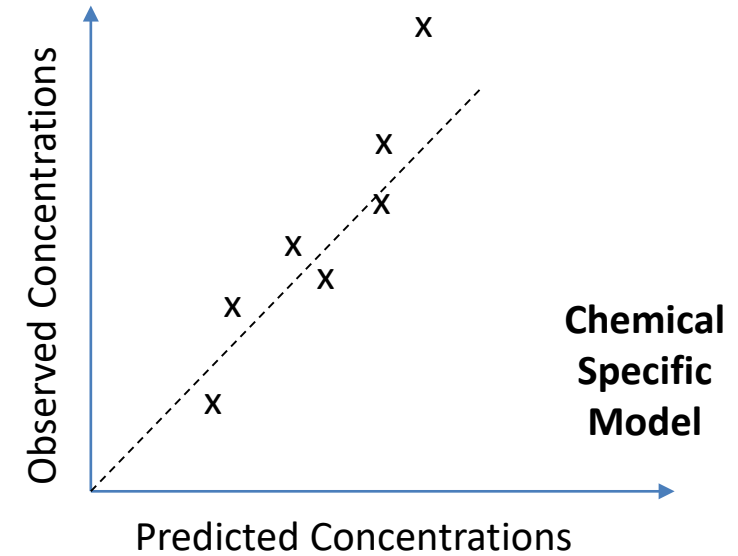
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- However, we do not typically have TK data
- We can parameterize a **generic TK model**, and evaluate that model for as many chemicals as we do have data
 - We do expect larger uncertainty, but also greater confidence in model implementation
 - Estimate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



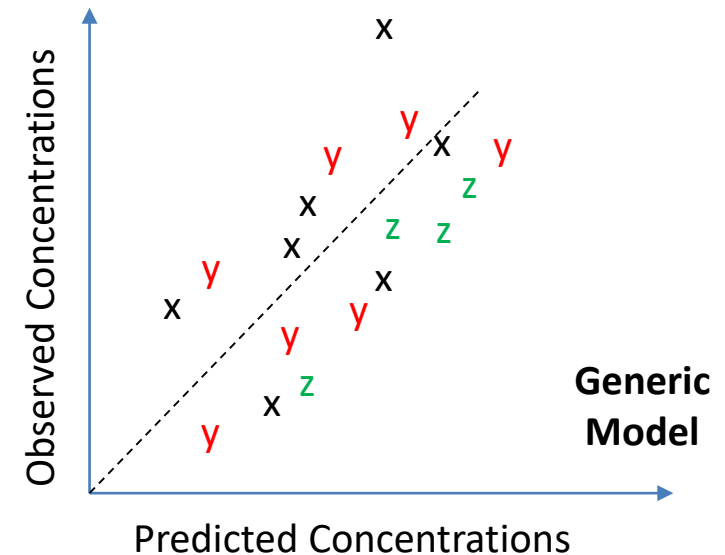
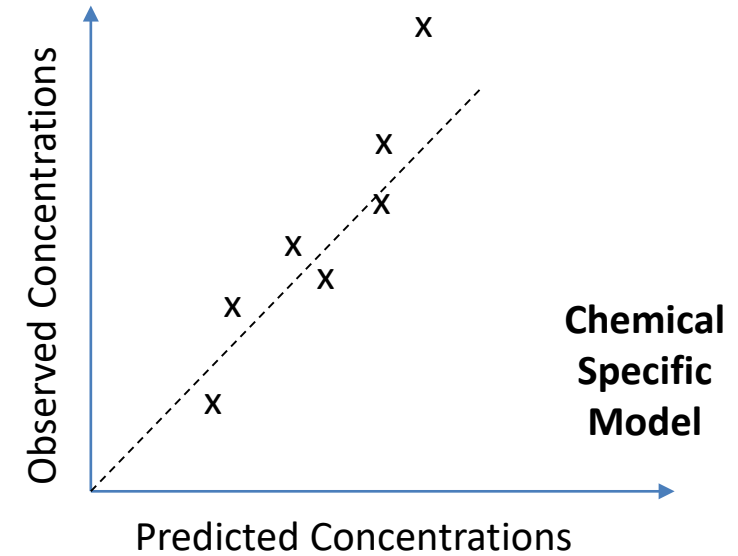
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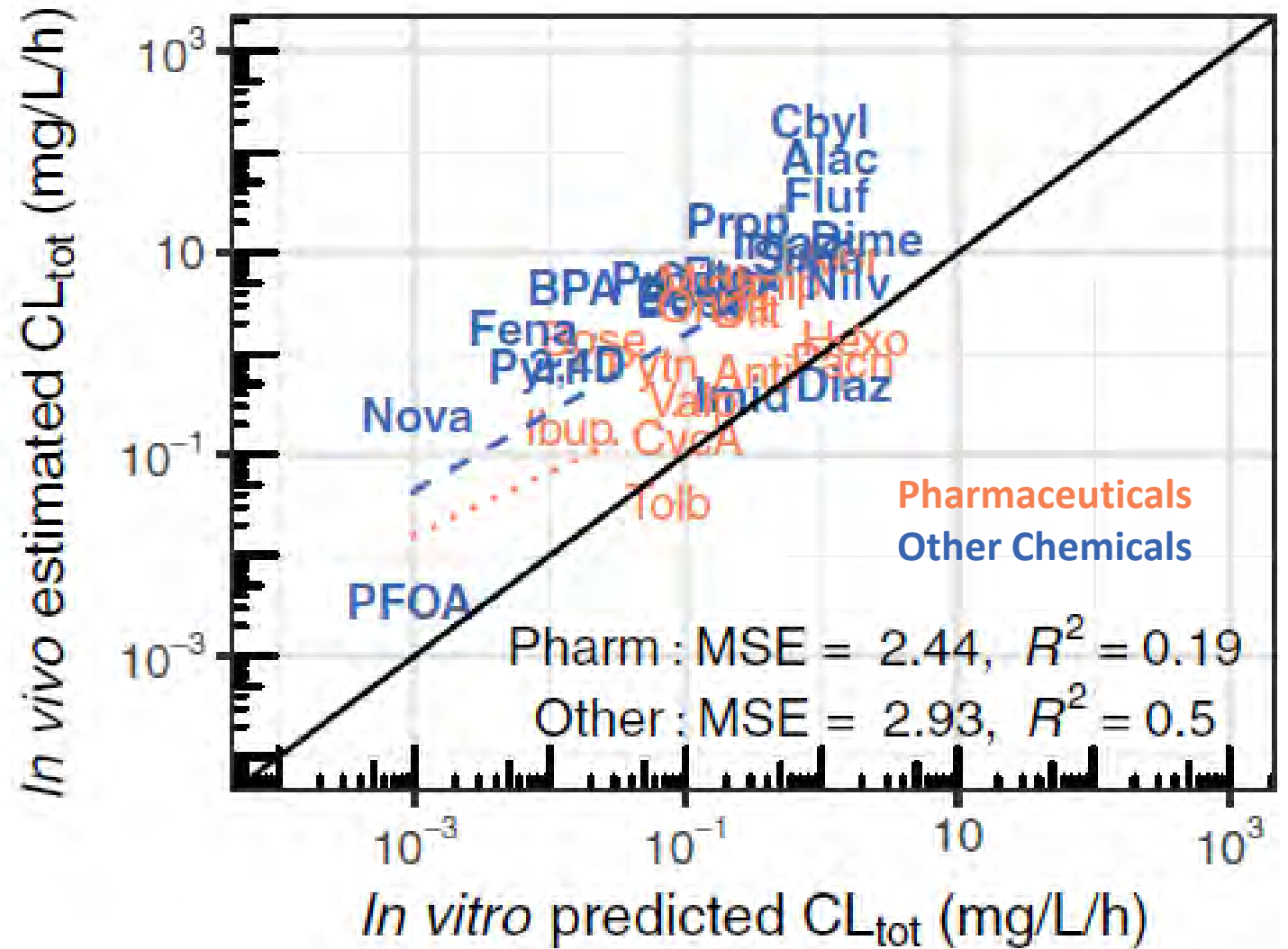
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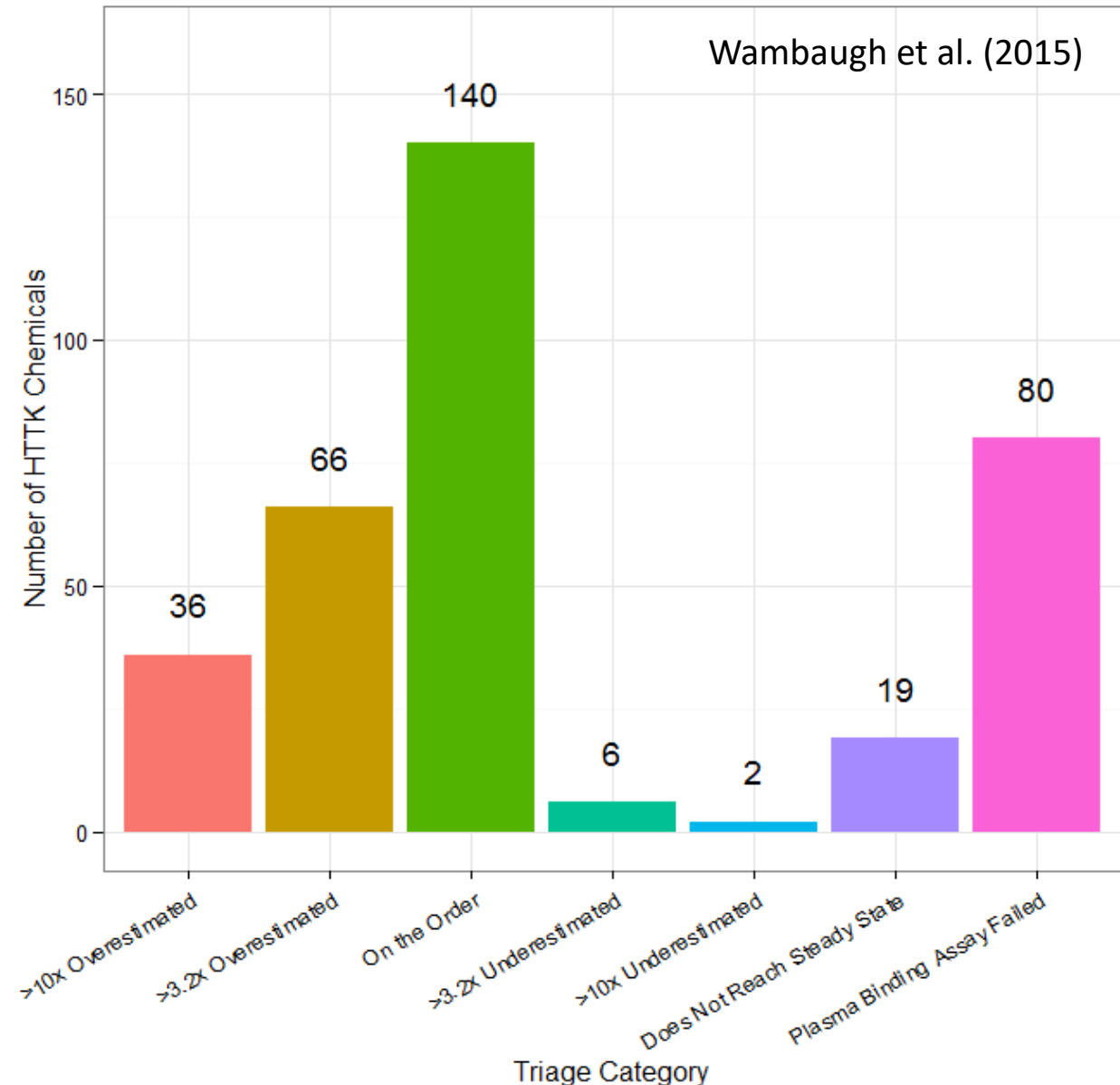
- The HHTK model estimates chemical clearance from the body by two processes:
 - hepatic metabolism (liver)
 - passive glomerular filtration (kidney)
- This appears to work better for pharmaceuticals than other chemicals:
 - ToxCast chemicals are overestimated
- Non-pharmaceuticals may be subject to extrahepatic metabolism and/or active transport

Evaluation Example



Toxicokinetic Triage: When Does TK IVIVE

- Through comparison to *in vivo* data, a cross-validated (random forest) predictor of success or failure of HTTK has been constructed
- All chemicals can be placed into one of seven confidence categories
 - Added categories for chemicals that do not reach steady-state or for which plasma binding assay fails
- Plurality of chemicals end up in the “on the order” bin (within a factor of 3.2x) which is consistent with Wang (2010)

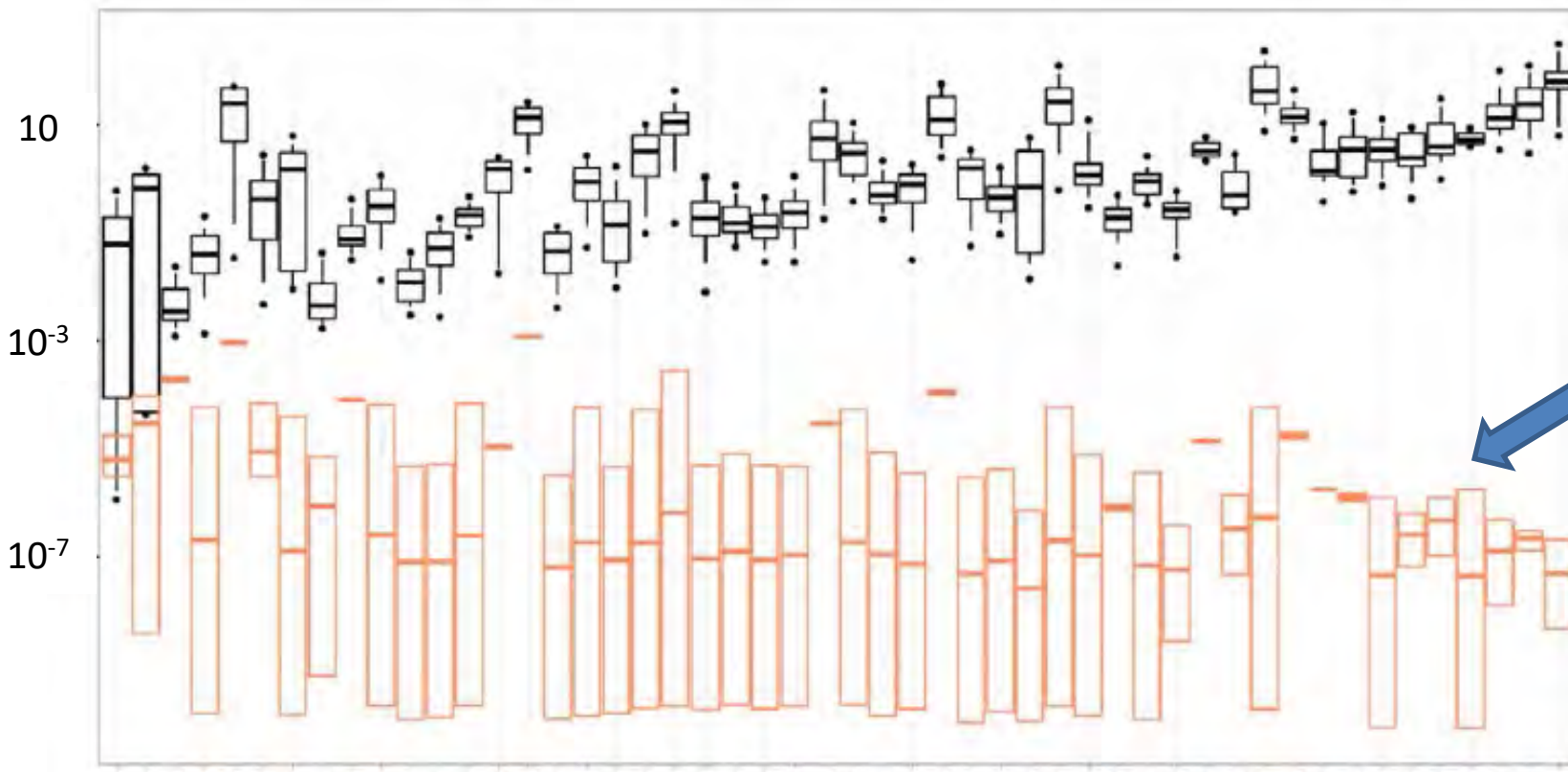


Chemical Prioritization NAMs

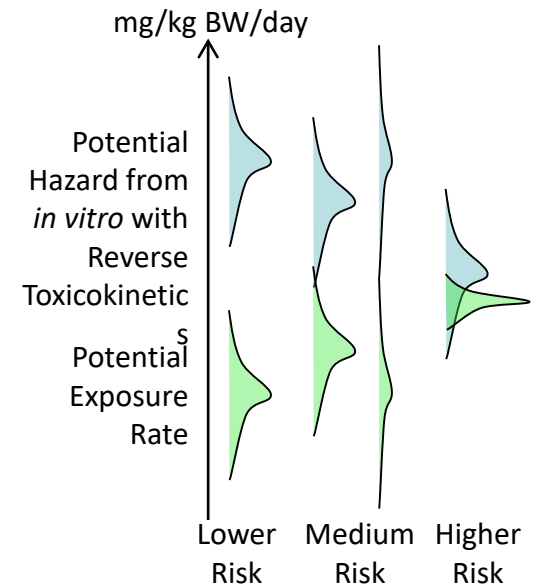
High throughput *in vitro* screening can estimate doses needed to cause bioactivity (e.g., Wetmore et al., 2015)

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Estimated Equivalent Dose or Predicted Exposure
(mg/kg BW/day)



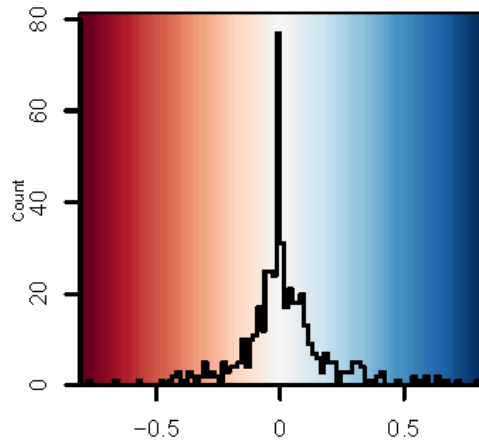
Chemicals Monitored by CDC NHANES



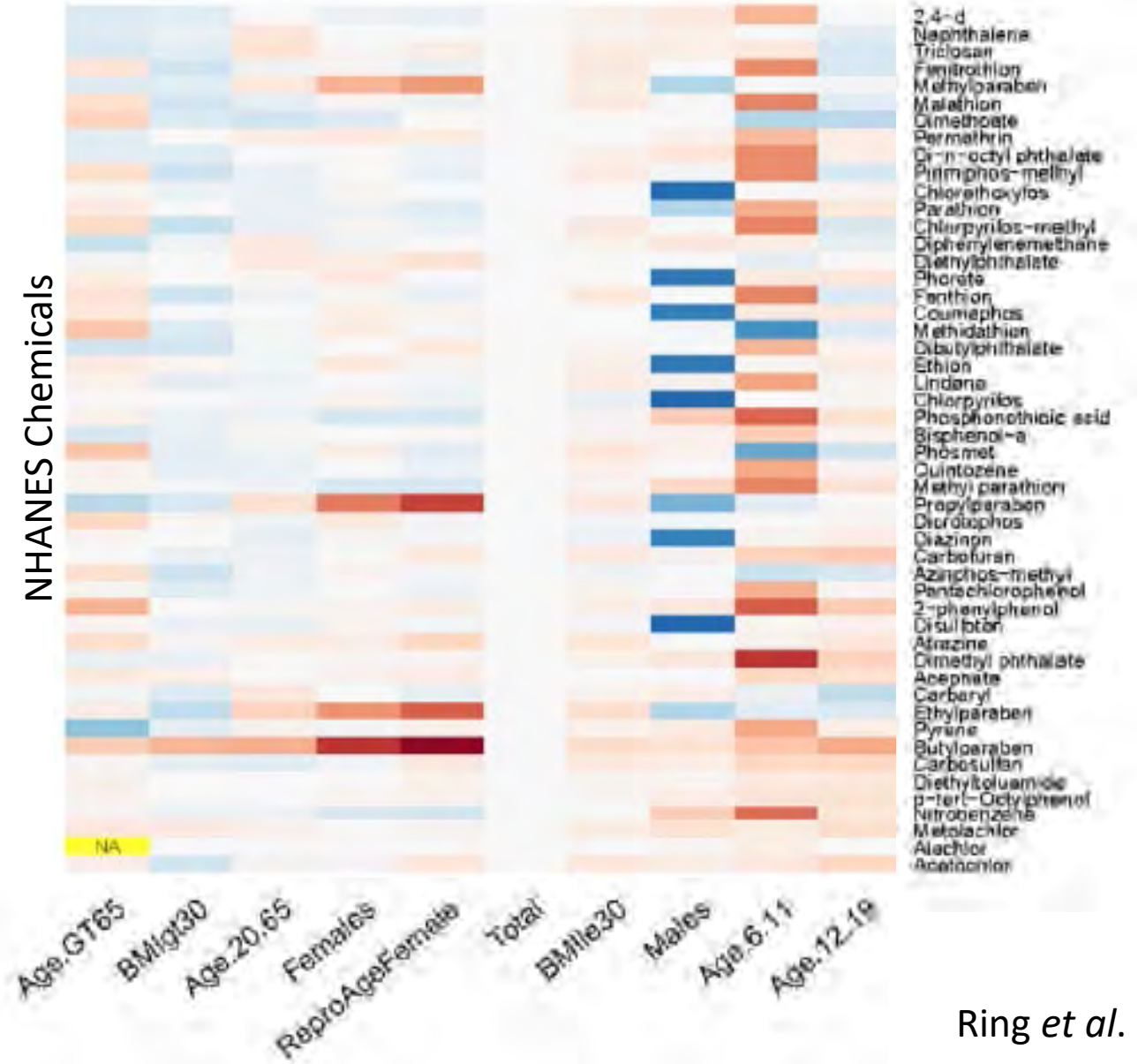
Ring et al. (2017)

Life-stage and Demographic Variation in Exposure

- Wambaugh *et al.* (2014) made steady-state inferences of exposure rate (mg/kg/day) from NHANES data for various demographic groups

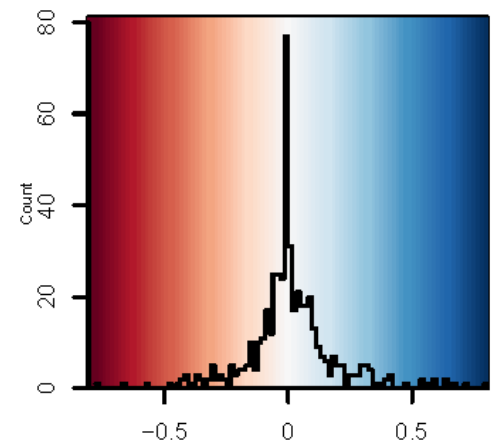


Change in Exposure
Relative to Total Population

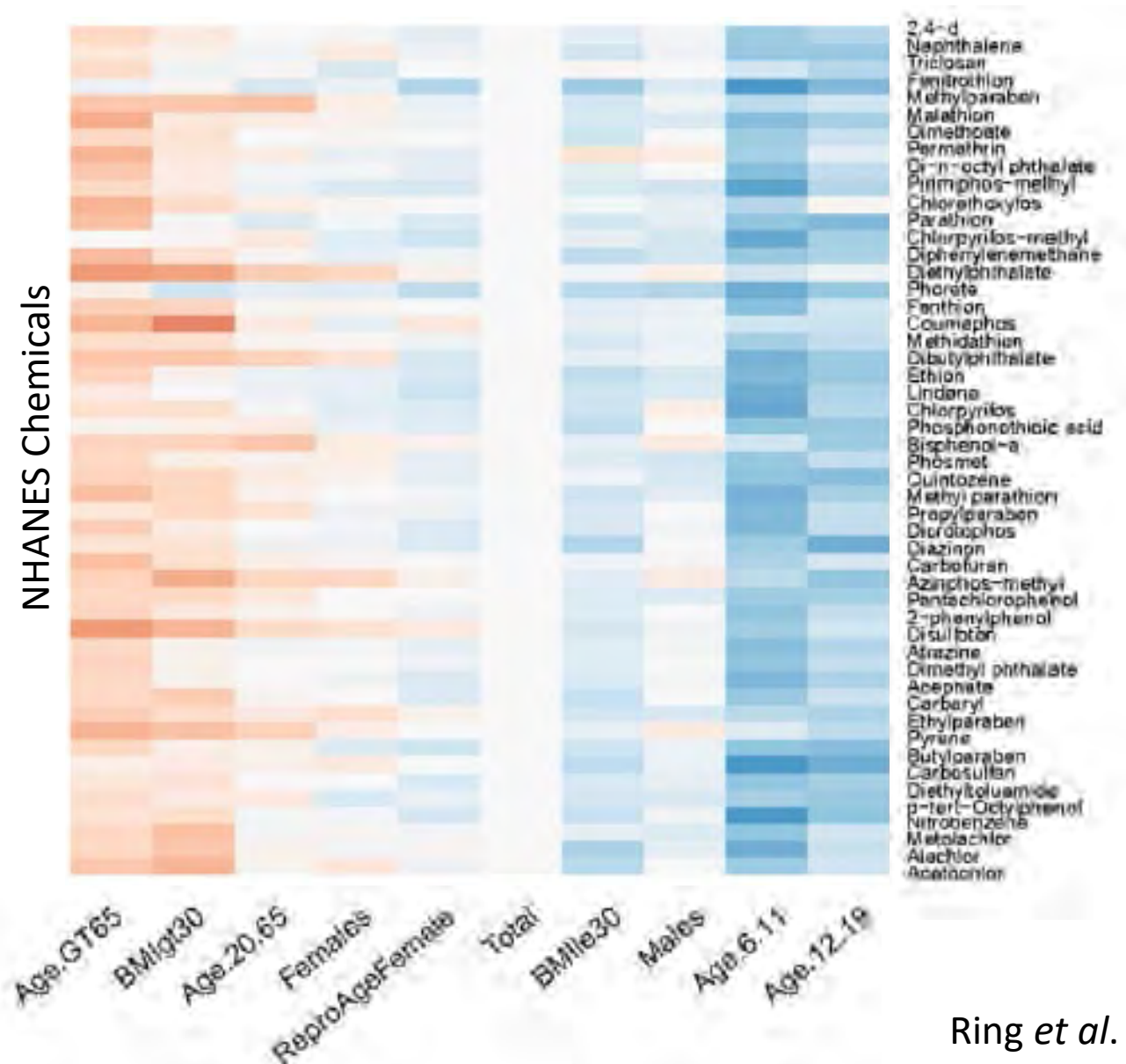


Life-stage and Demographic Variation in TK

- Ring *et al.* (2017) made demographic-specific predictions of change in plasma concentrations for a 1 mg/kg bw/day exposure

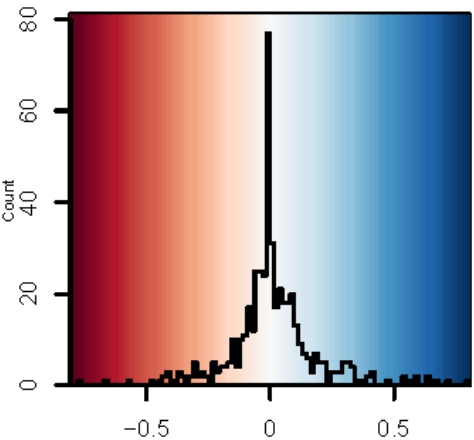
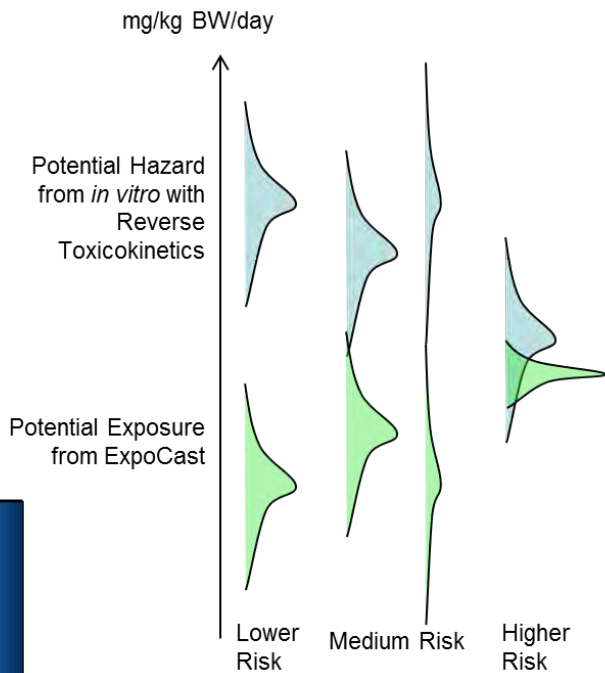


Change in Toxicokinetics
Relative to Total Population

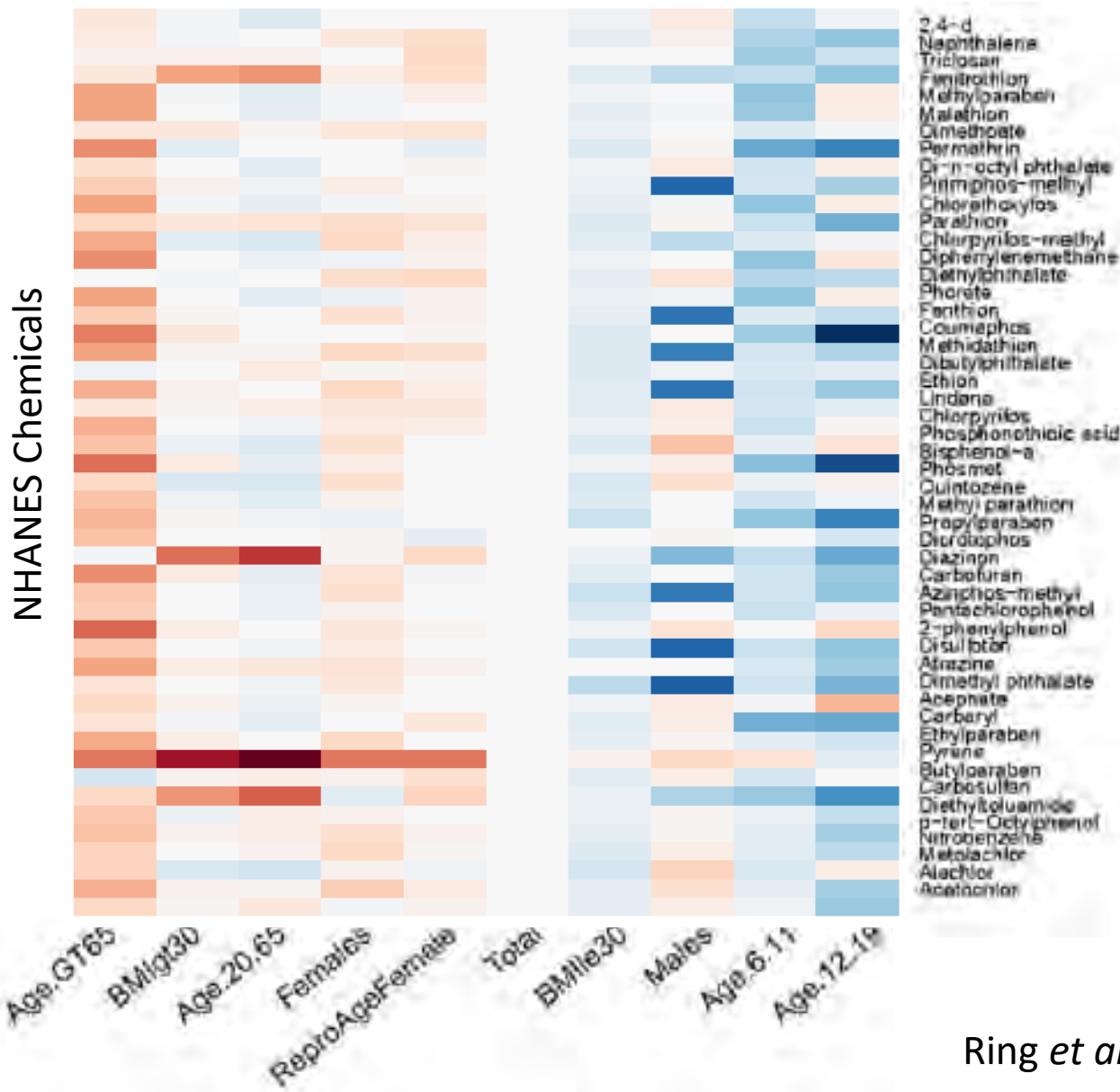


Life-stage and Demographic Variation in Risk Priority

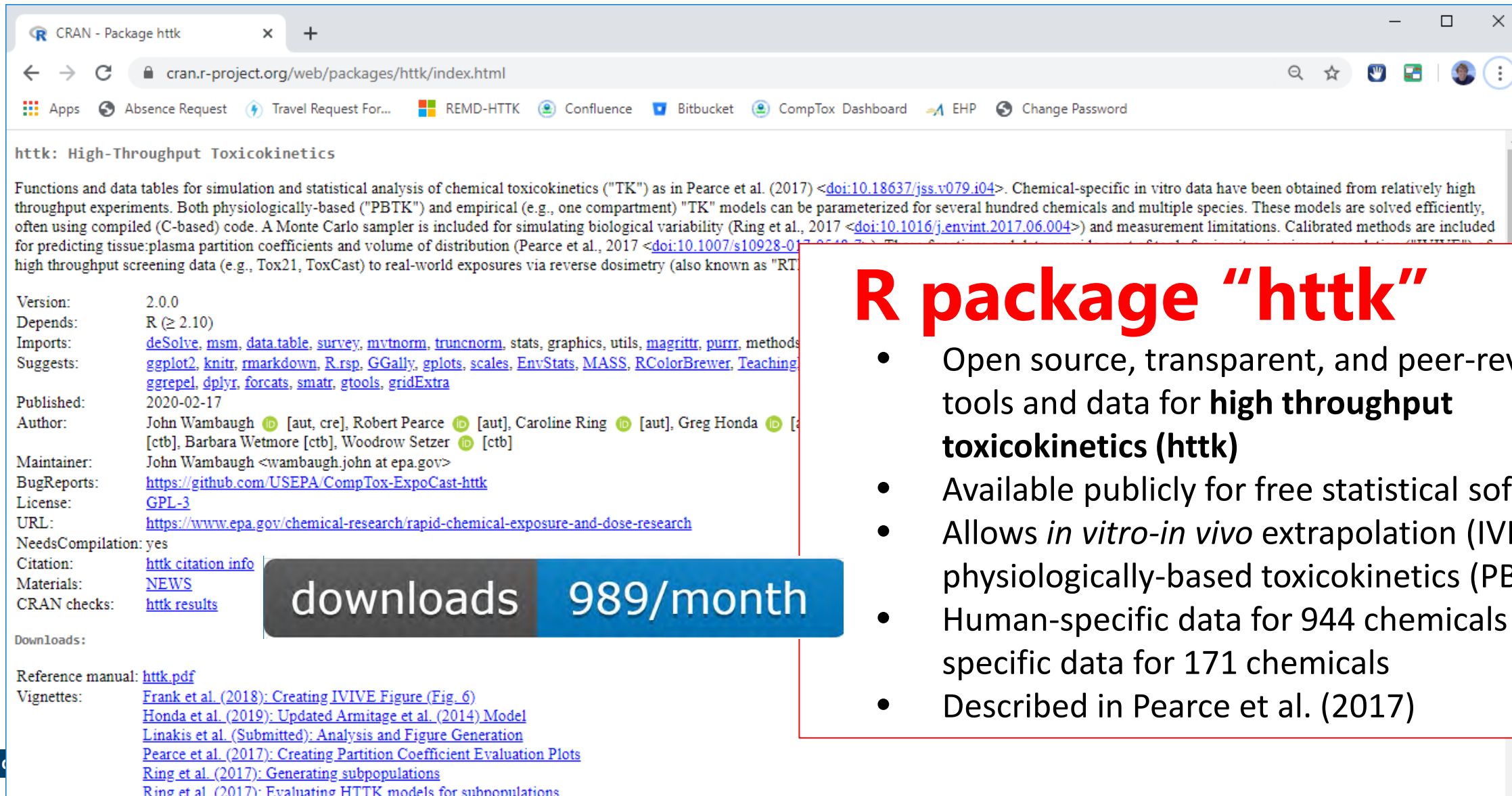
- Can calculate margin between bioactivity and exposure for specific populations



Change in Risk Relative to
Total Population



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



The screenshot shows the CRAN package page for 'httk'. The browser address bar shows 'cran.r-project.org/web/packages/httk/index.html'. The page title is 'httk: High-Throughput Toxicokinetics'. The description states: 'Functions and data tables for simulation and statistical analysis of chemical toxicokinetics ("TK") as in Pearce et al. (2017) <doi:10.18637/jss.v079.i04>. Chemical-specific in vitro data have been obtained from relatively high throughput experiments. Both physiologically-based ("PBTk") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability (Ring et al., 2017 <doi:10.1016/j.envint.2017.06.004>) and measurement limitations. Calibrated methods are included for predicting tissue:plasma partition coefficients and volume of distribution (Pearce et al., 2017 <doi:10.1007/s10928-017-0648-7>). The package also includes methods for simulating high throughput screening data (e.g., Tox21, ToxCast) to real-world exposures via reverse dosimetry (also known as "RT...').

Version: 2.0.0
Depends: R (≥ 2.10)
Imports: deSolve, msm, data.table, survey, mvtnorm, truncnorm, stats, graphics, utils, magrittr, purrr, methods
Suggests: ggplot2, knitr, rmarkdown, R.rsp, GGally, gplots, scales, EnvStats, MASS, RColorBrewer, TeachingDemos, ggrepel, dplyr, forcats, smatr, gtools, gridExtra
Published: 2020-02-17
Author: John Wambaugh [aut, cre], Robert Pearce [aut], Caroline Ring [aut], Greg Honda [aut], Barbara Wetmore [ctb], Woodrow Setzer [ctb]
Maintainer: John Wambaugh <wambaugh.john@epa.gov>
BugReports: <https://github.com/USEPA/CompTox-ExpoCast-httk>
License: GPL-3
URL: <https://www.epa.gov/chemical-research/rapid-chemical-exposure-and-dose-research>
NeedsCompilation: yes
Citation: [httk citation info](#)
Materials: [NEWS](#)
CRAN checks: [httk results](#)

Downloads: 989/month

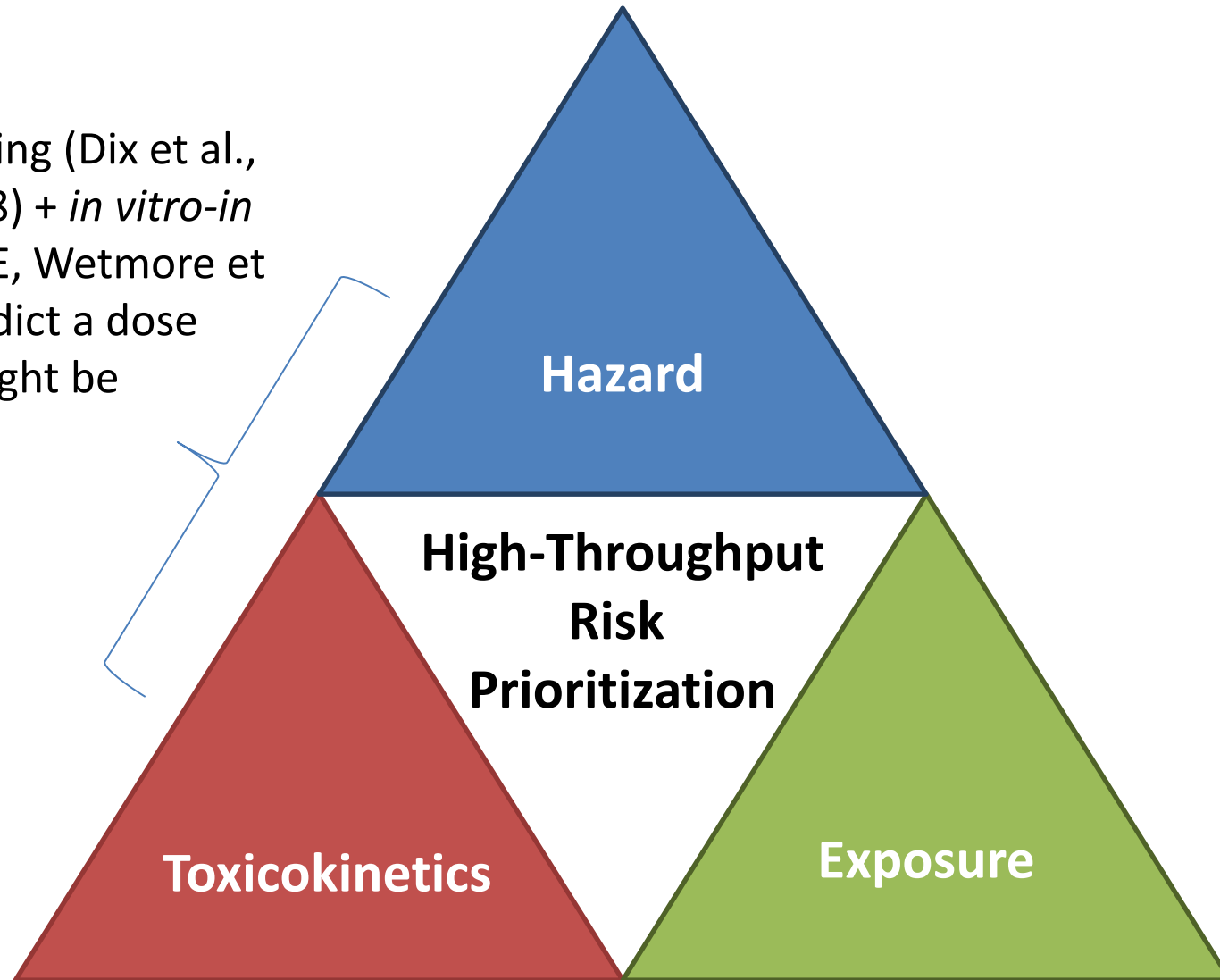
Reference manual: [httk.pdf](#)
Vignettes: Frank et al. (2018): [Creating IVIVE Figure \(Fig. 6\)](#)
Honda et al. (2019): [Updated Armitage et al. \(2014\) Model](#)
Linakis et al. (Submitted): [Analysis and Figure Generation](#)
Pearce et al. (2017): [Creating Partition Coefficient Evaluation Plots](#)
Ring et al. (2017): [Generating subpopulations](#)
Ring et al. (2017): [Evaluating HHTK models for subpopulations](#)

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)
- Human-specific data for 944 chemicals and rat-specific data for 171 chemicals
- Described in Pearce et al. (2017)

$$\text{Risk} = \text{Hazard} \times \text{Exposure}$$

High throughput screening (Dix et al., 2006, Collins et al., 2008) + *in vitro-in vivo* extrapolation (IVIVE, Wetmore et al., 2012, 2015) can predict a dose (mg/kg bw/day) that might be adverse

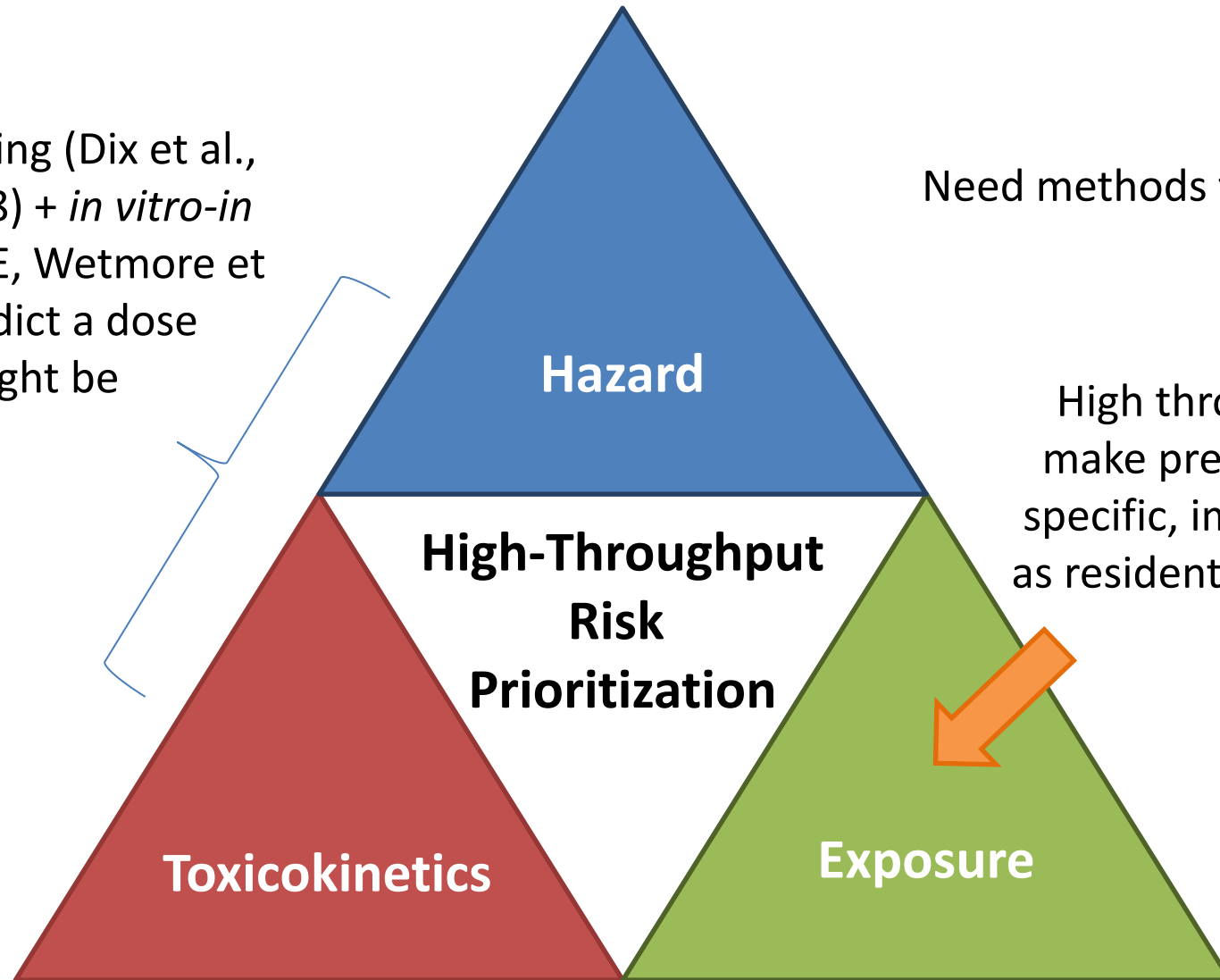


Risk = Hazard x Exposure

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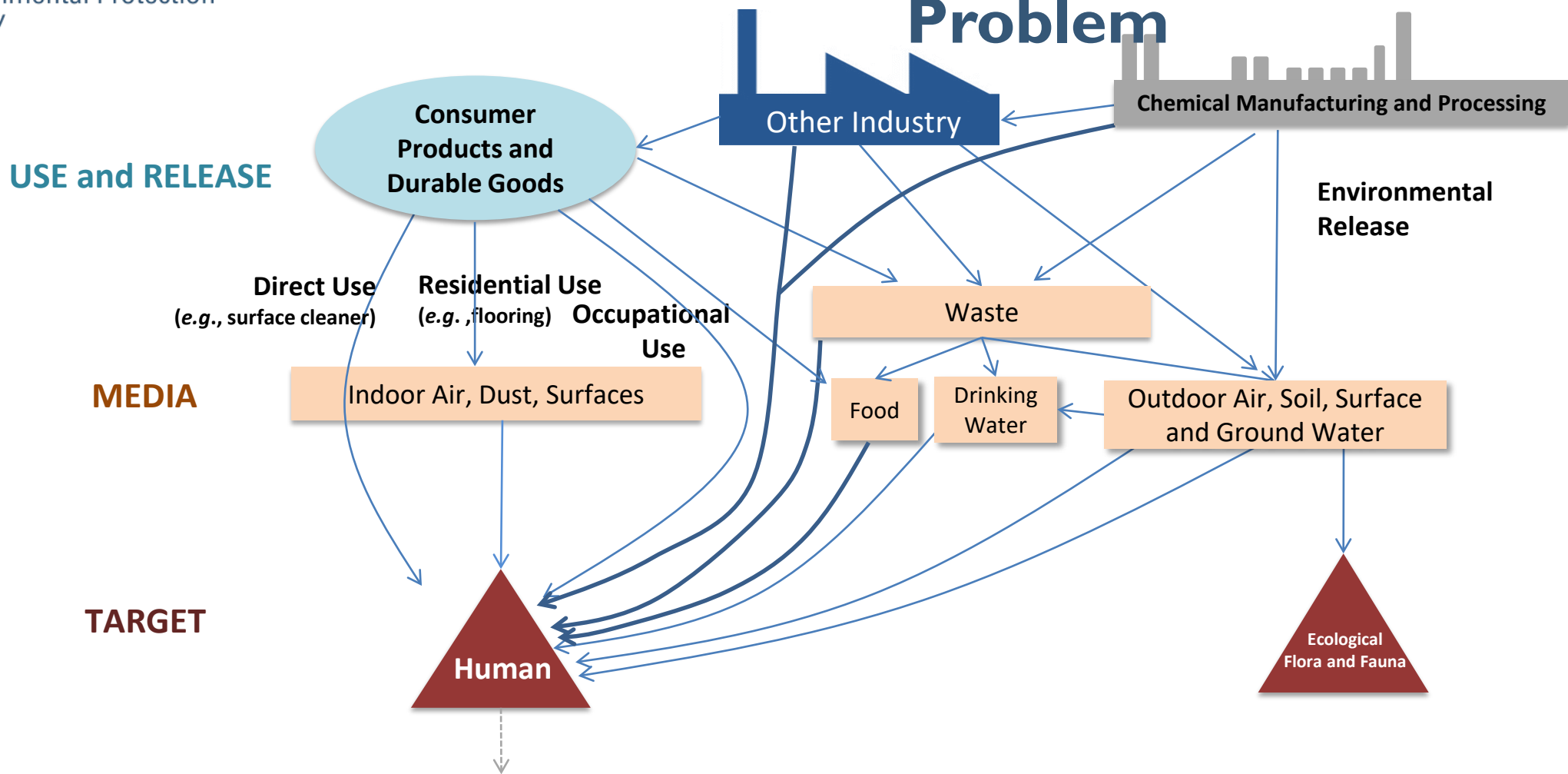
Need methods to forecast exposure for thousands of chemicals (Wetmore et al., 2015)

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

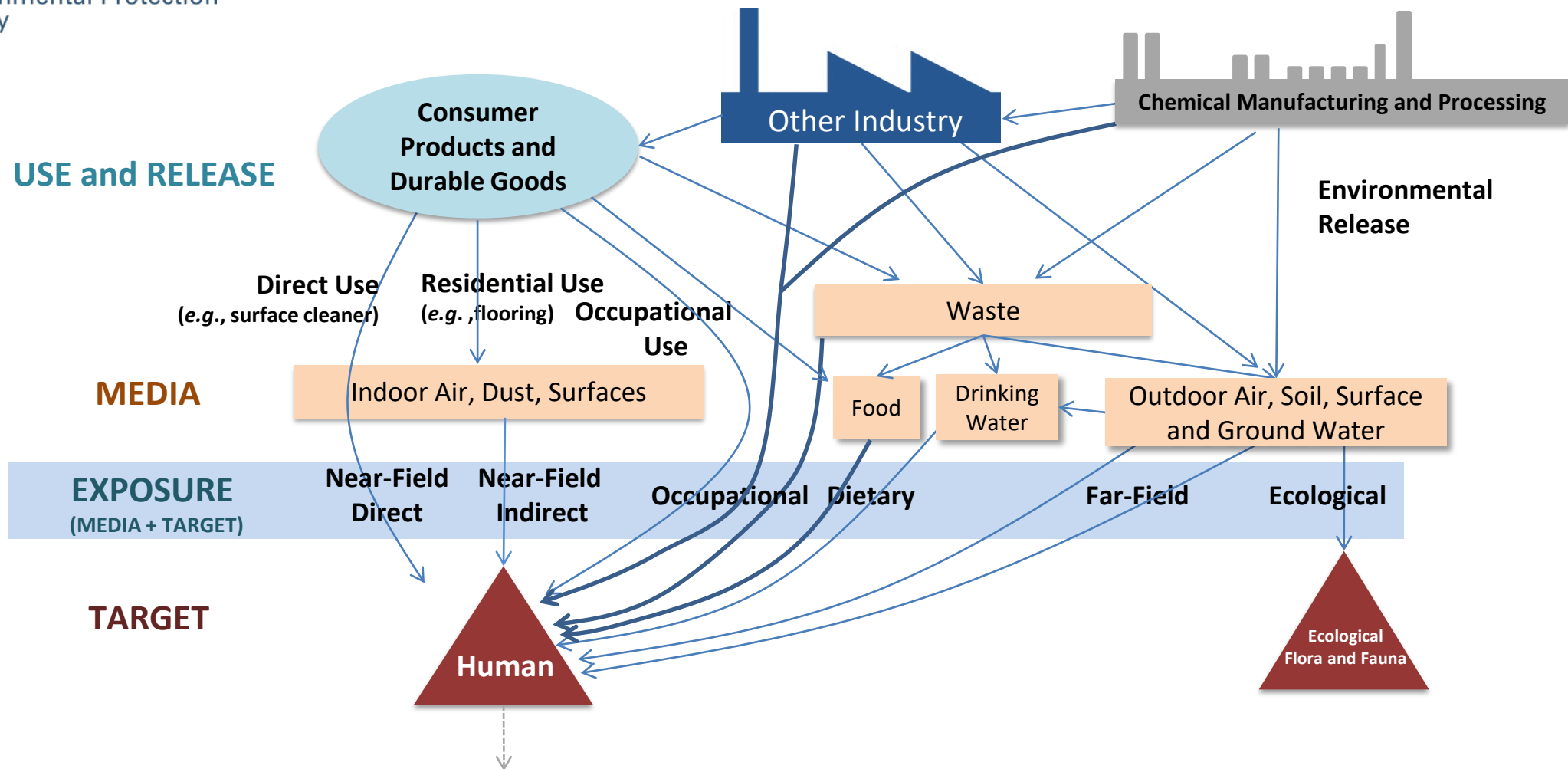


NRC (1983)

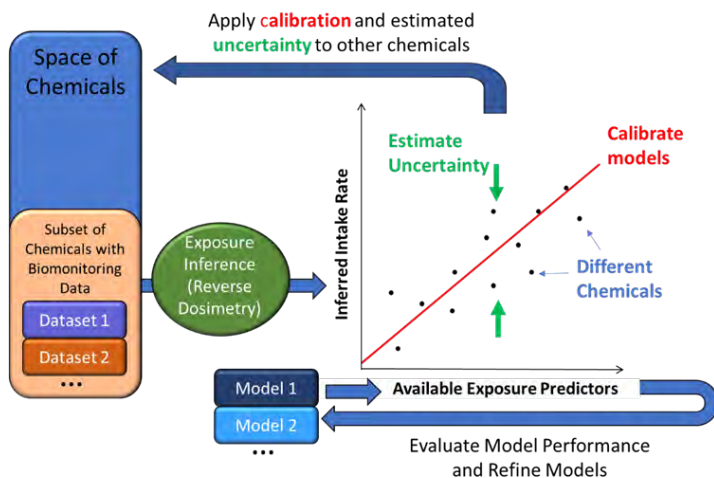
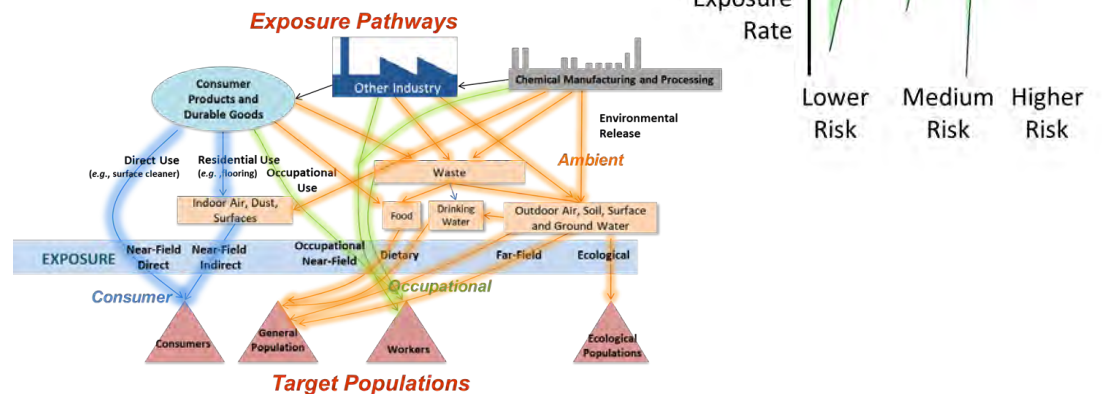
Understanding Exposure is a Systems Problem



Exposure event is often unobservable



- Can try to predict exposure by characterizing pathway
- Some pathways have much higher average exposures: In home “Near field” sources significant (Wallace, *et al.*, 1987)



Author's Personal Copy
Available online at www.sciencedirect.com

ScienceDirect

Current Opinion in
Toxicology

New approach methodologies for exposure science

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Abstract

Chemical risk assessment relies on knowledge of hazard, the dose–response relationship, and exposure to characterize potential risks to public health and the environment. A chemical with minimal toxicity might pose a risk if exposures are extensive, repeated, and/or occurring during critical windows across the human life span. Exposure assessment involves understanding human activity, and this activity is confounded by interindividual variability that is both biological and behavioral. Exposures further vary between the general population and susceptible or occupationally exposed populations. Recent computational exposure efforts have tackled these problems through the creation of new tools and predictive models. These tools include machine learning to draw inferences from existing data and computer-enhanced screening analyses to generate new data. Mathematical models provide frameworks describing

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New Approach Methodologies for Exposure Science

Exposure NAM Class	Description	Traditional Approach	Makes Use of					
			Measurement	Toxicokinetics	Models	Descriptors	Evaluation	Machine Learning
Measurements	New techniques including screening analyses capable of detecting hundreds of chemicals present in a sample	Targeted (chemical-specific) analyses	-	●	●	●		●
Toxicokinetics	High throughput methods using in vitro data to generate chemical-specific models	Analyses based on in vivo animal studies	●	-		●		●
HTE Models	Models capable of making predictions for thousands of chemicals	Models requiring detailed, chemical- and scenario-specific information	●	●	-	●		
Chemical Descriptors	Informatic approaches for organizing chemical information in a machine-readable format	Tools targeted at single chemical analyses by humans				-		●
Evaluation	Statistical approaches that use the data from many chemicals to estimate the uncertainty in a prediction for a new chemical	Comparison of model predictions to data on a per chemical basis	●	●	●	●	-	●
Machine Learning	Computer algorithms to identify patterns	Manual Inspection of the Data	●	●		●		-
Prioritization	Integration of exposure and other NAMs to identify chemicals for follow-up study	Expert decision making	●	●	●	●	●	●

What Do We Know About Exposure?

Biomonitoring Data

- Centers for Disease Control and Prevention (CDC) National Health and Nutrition Examination Survey (NHANES) provides an important tool for monitoring public health
- Large, ongoing CDC survey of US population: demographic, body measures, medical exam, biomonitoring (health and exposure), ...
- Designed to be representative of US population according to census data
- Data sets publicly available (<http://www.cdc.gov/nchs/nhanes.htm>)
- Includes measurements of:
 - Body weight
 - Height
 - **Chemical analysis of blood and urine**



What Do We Know About Exposure?

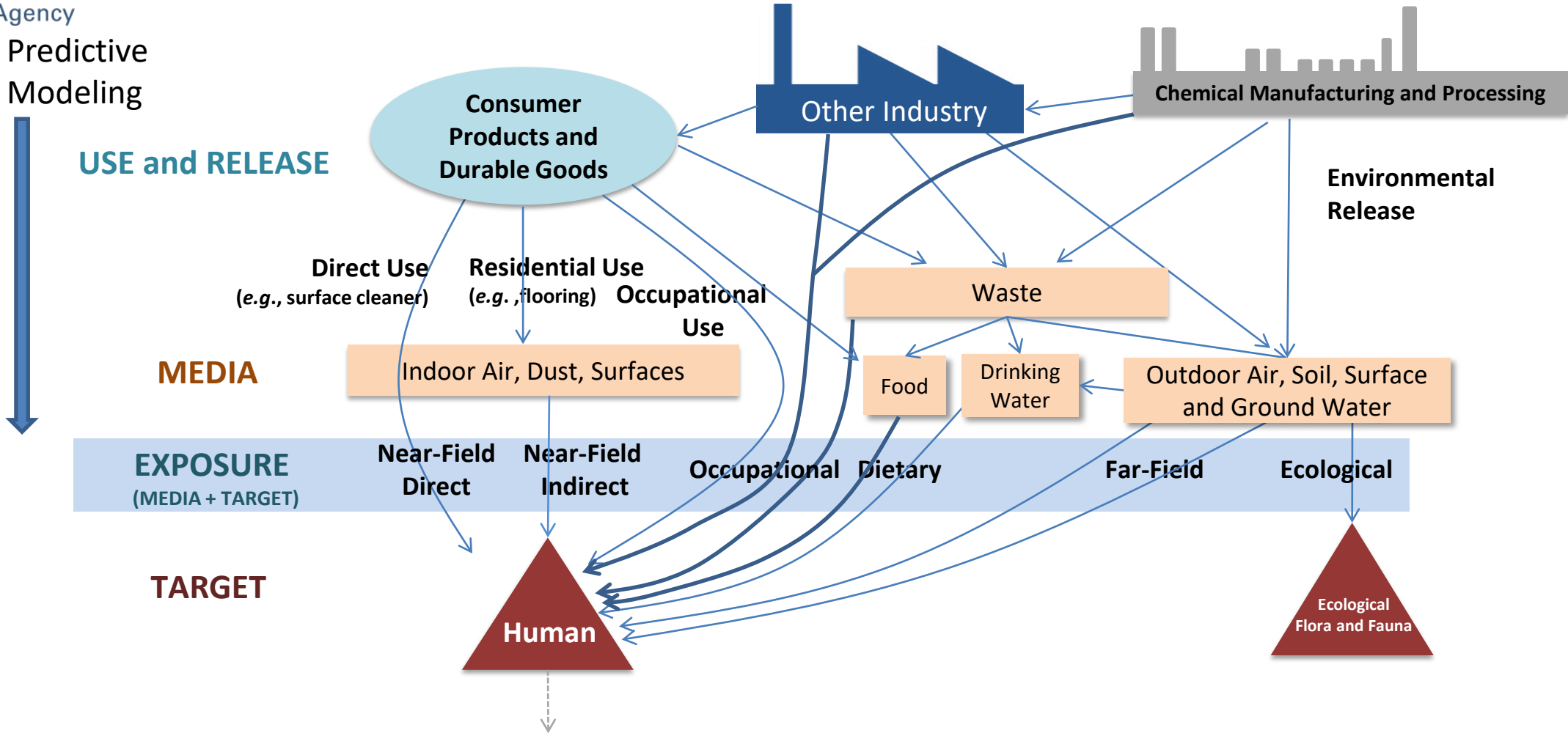
Exposure Models

- Human chemical exposures can be coarsely grouped into “**near field**” sources that are close to the exposed individual (consumer or occupational exposures) ‘**far-field**’ scenarios wherein individuals are exposed to chemicals that were released or used far away (ambient exposure) (Arnot *et al.*, 2006).
- A model captures knowledge and a hypothesis of how the world works (MacLeod *et al.*, 2010)
- EPA’s EXPOsure toolBOX (EPA ExpoBox) is a toolbox created to assist individuals from within government, industry, academia, and the general public with assessing exposure
 - Includes many, many models

<https://www.epa.gov/expobox>

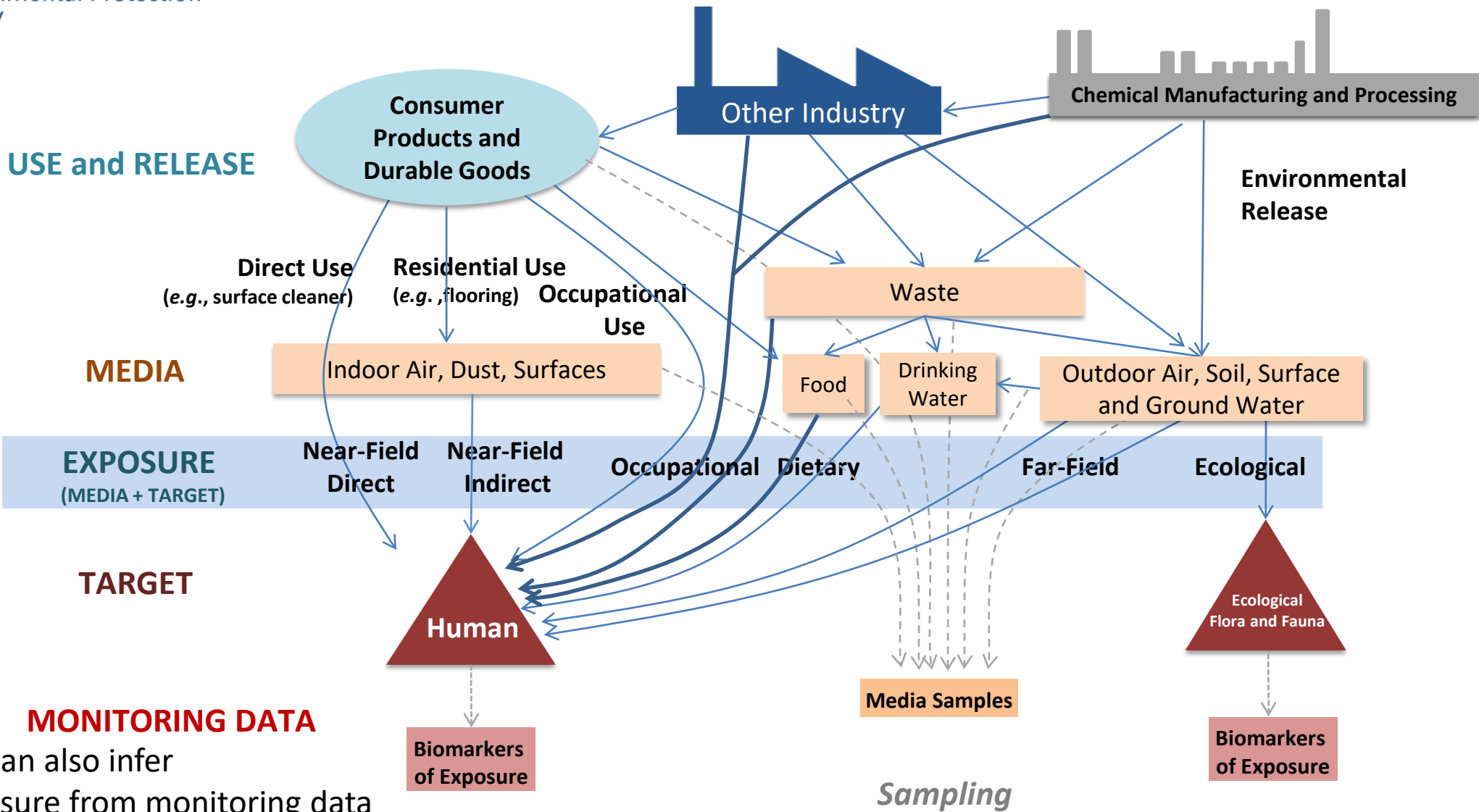
“Now it would be very remarkable if any system existing in the real world could be exactly represented by any simple model. However, cunningly chosen parsimonious models often do provide remarkably useful approximations... The only question of interest is ‘Is the model illuminating and useful?’” George Box

Models to Predict Exposure



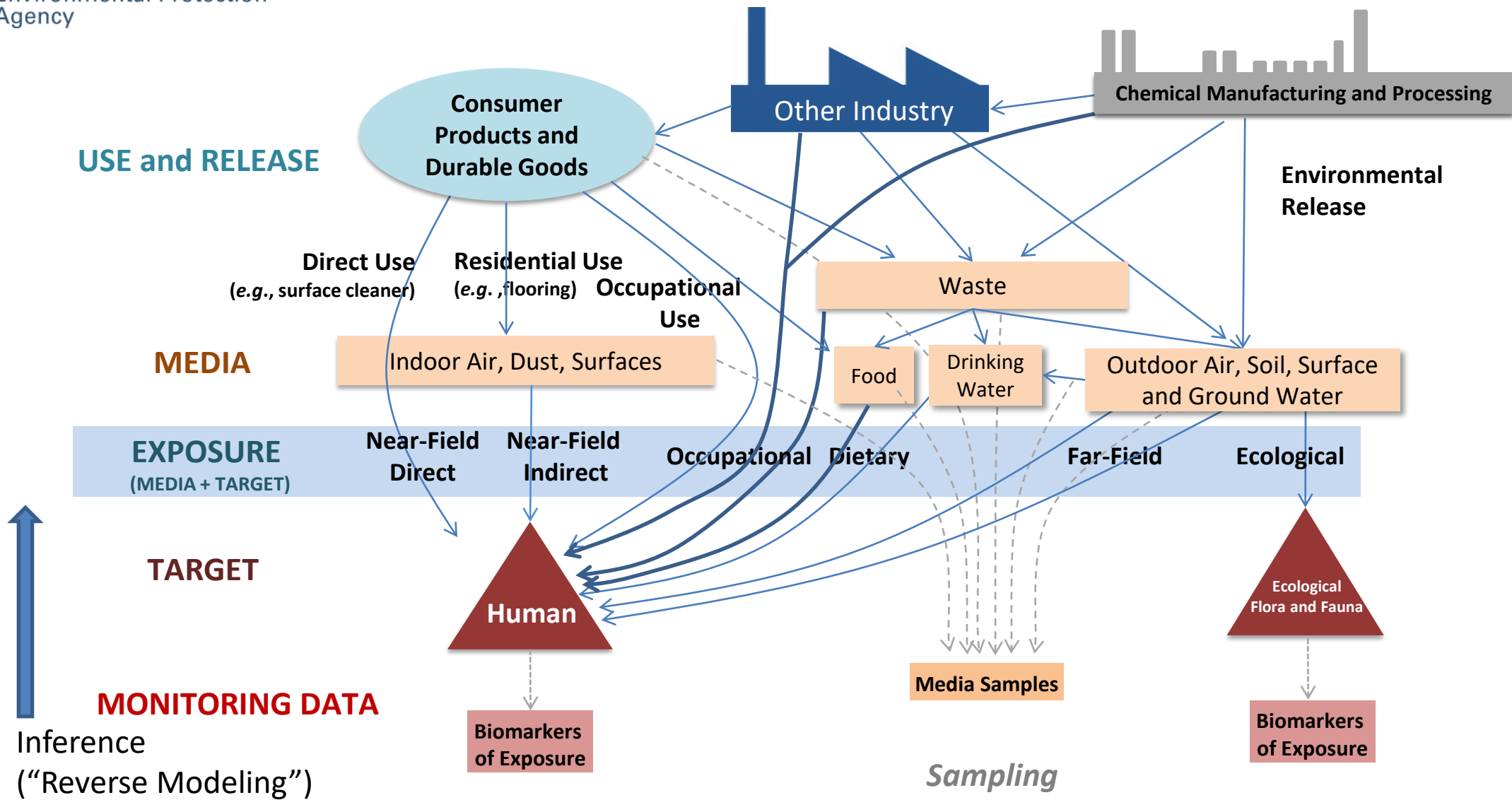
We can try to predict exposure by describing the process leading to exposure

Monitoring Data



We can also infer
exposure from monitoring data

Models to Infer Exposure



Evaluating Models with Monitoring Data

Predictive
Modeling

USE and RELEASE

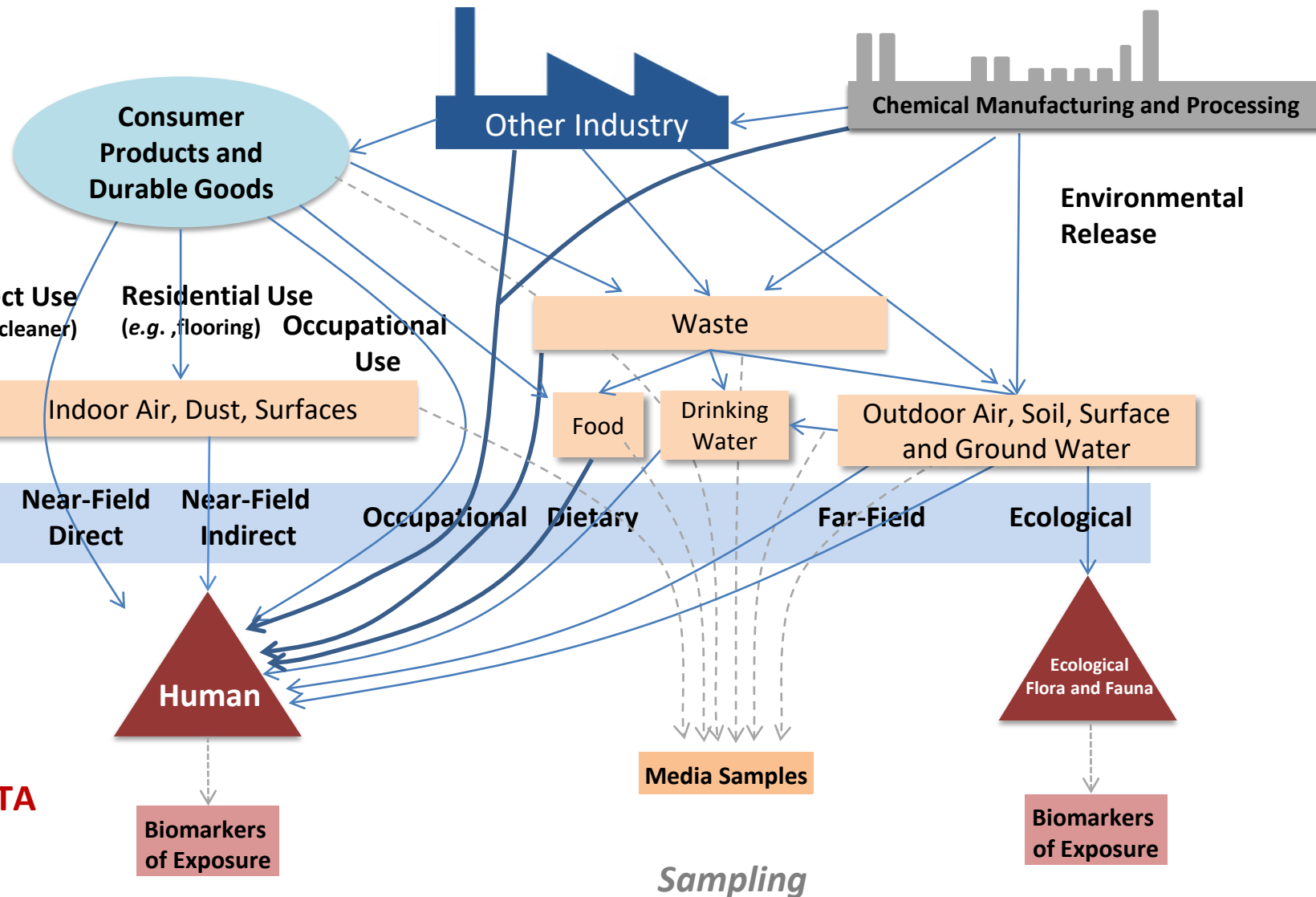
MEDIA

EXPOSURE
(MEDIA + TARGET)

TARGET

MONITORING DATA

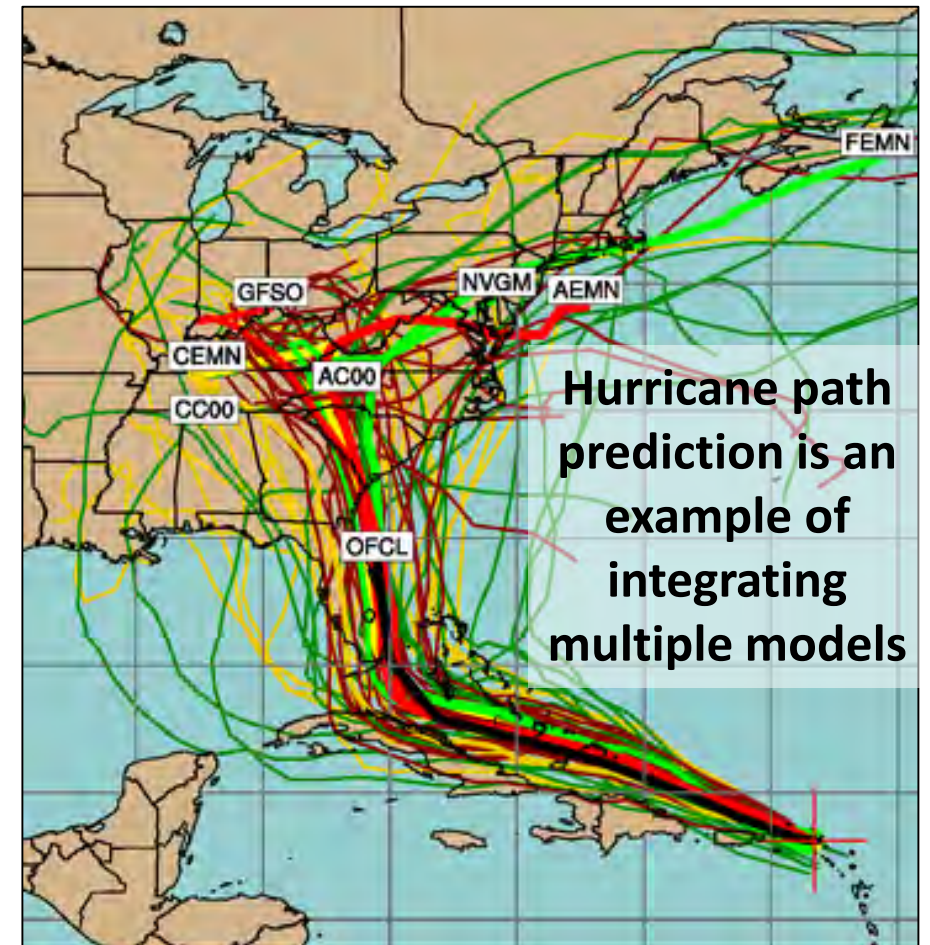
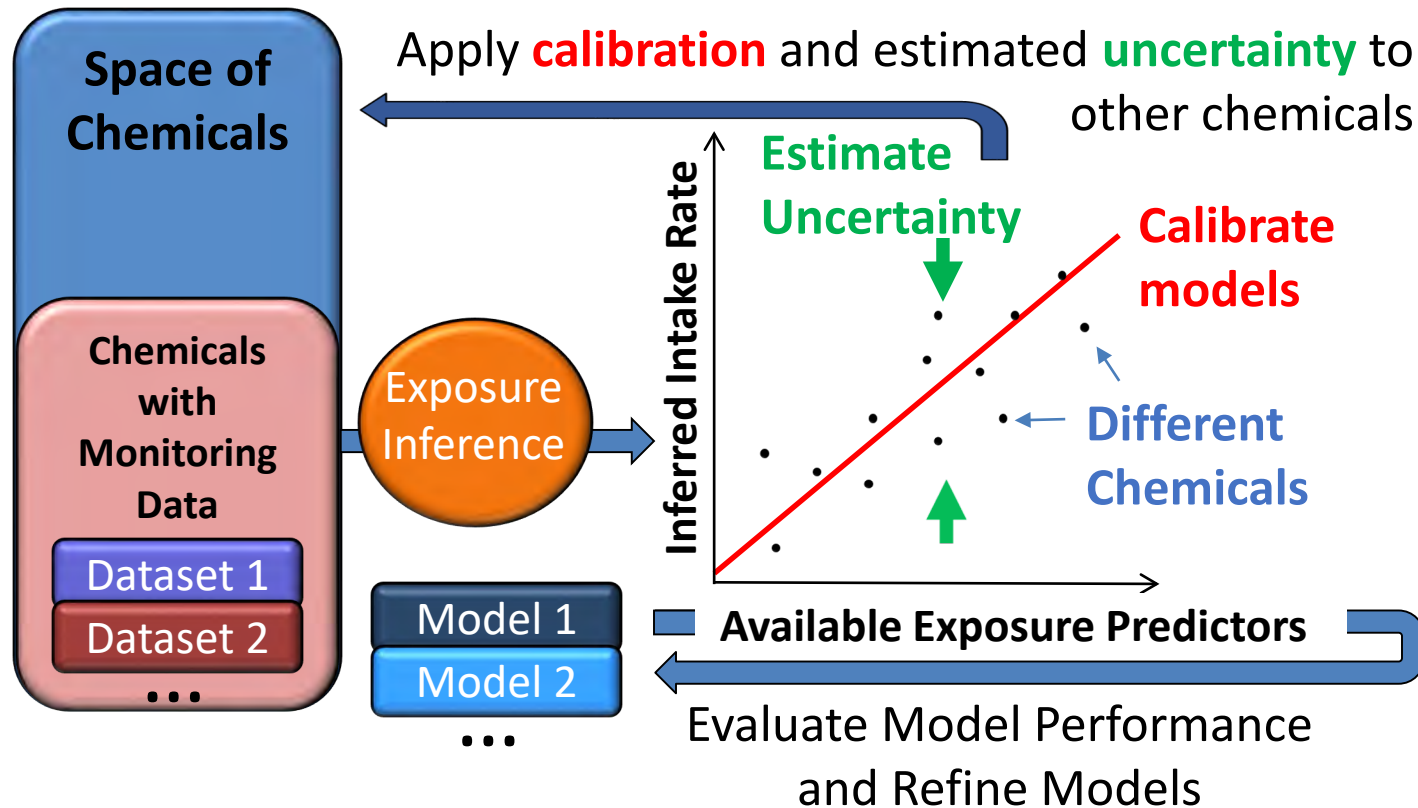
Inference
("Reverse Modeling")



Sampling

Evaluation NAMs: The SEEM Framework

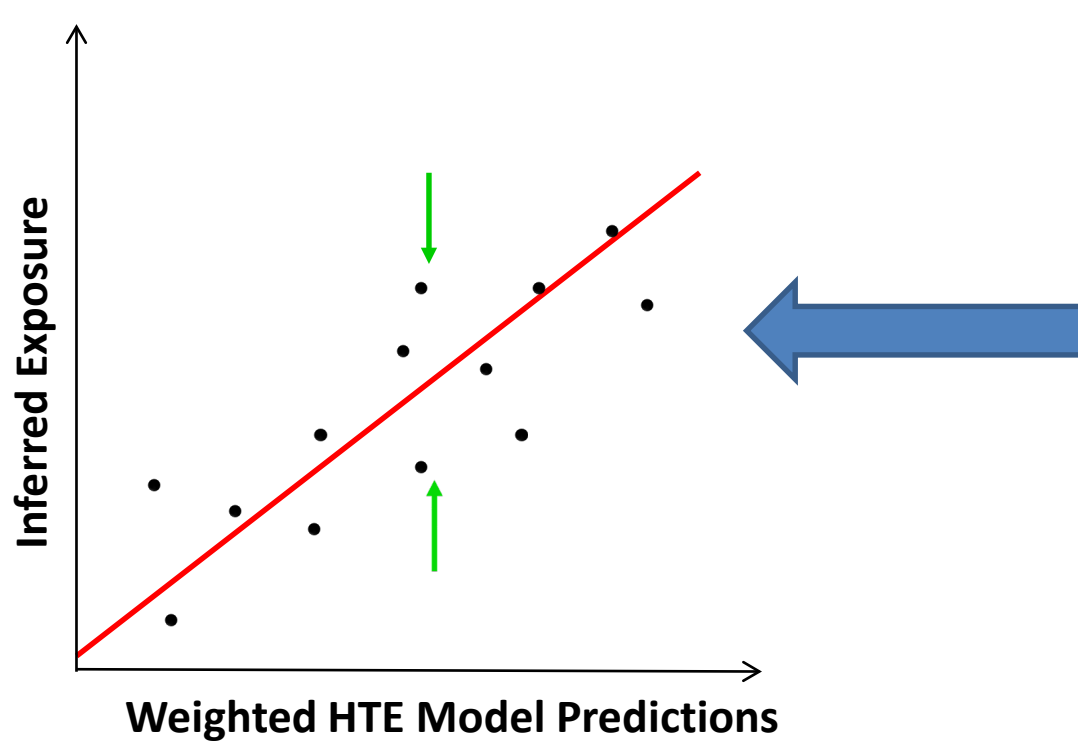
- We use Bayesian methods to incorporate multiple models into consensus predictions for 1000s of chemicals within the **Systematic Empirical Evaluation of Models (SEEM)** (Wambaugh et al., 2013, 2014; Ring et al., 2018)



SEEM is a Linear Regression

Multiple regression models:

$$\text{Log(Parent Exposure)} = a + m * \log(\text{Model Prediction}) + b * \text{Near Field} + \varepsilon$$

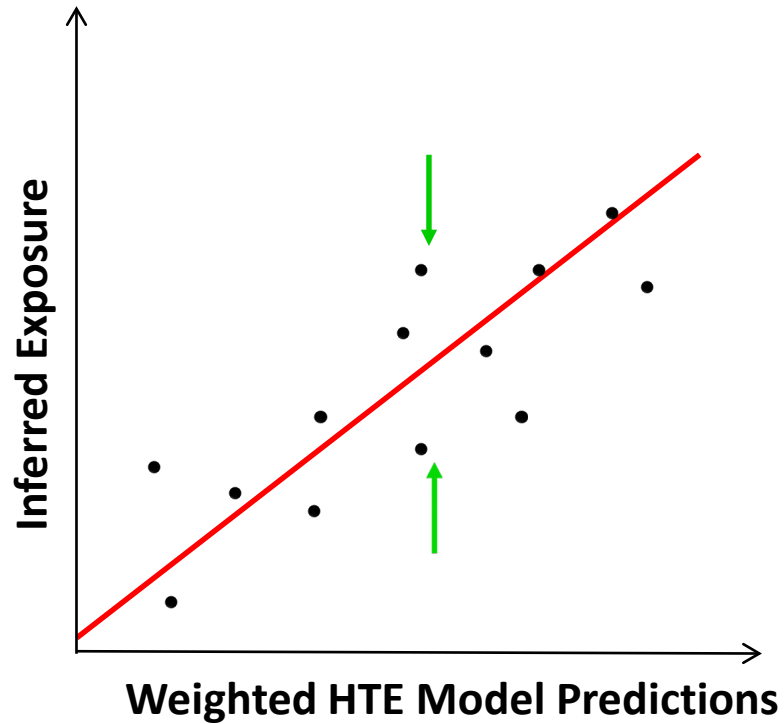


$\varepsilon \sim N(0, \sigma^2)$
Residual error,
unexplained by
the regression
model

SEEM is a Linear Regression

Multiple regression models:

$$\text{Log(Parent Exposure)} = a + m * \log(\text{Model Prediction}) + b * \text{Near Field} + \varepsilon$$



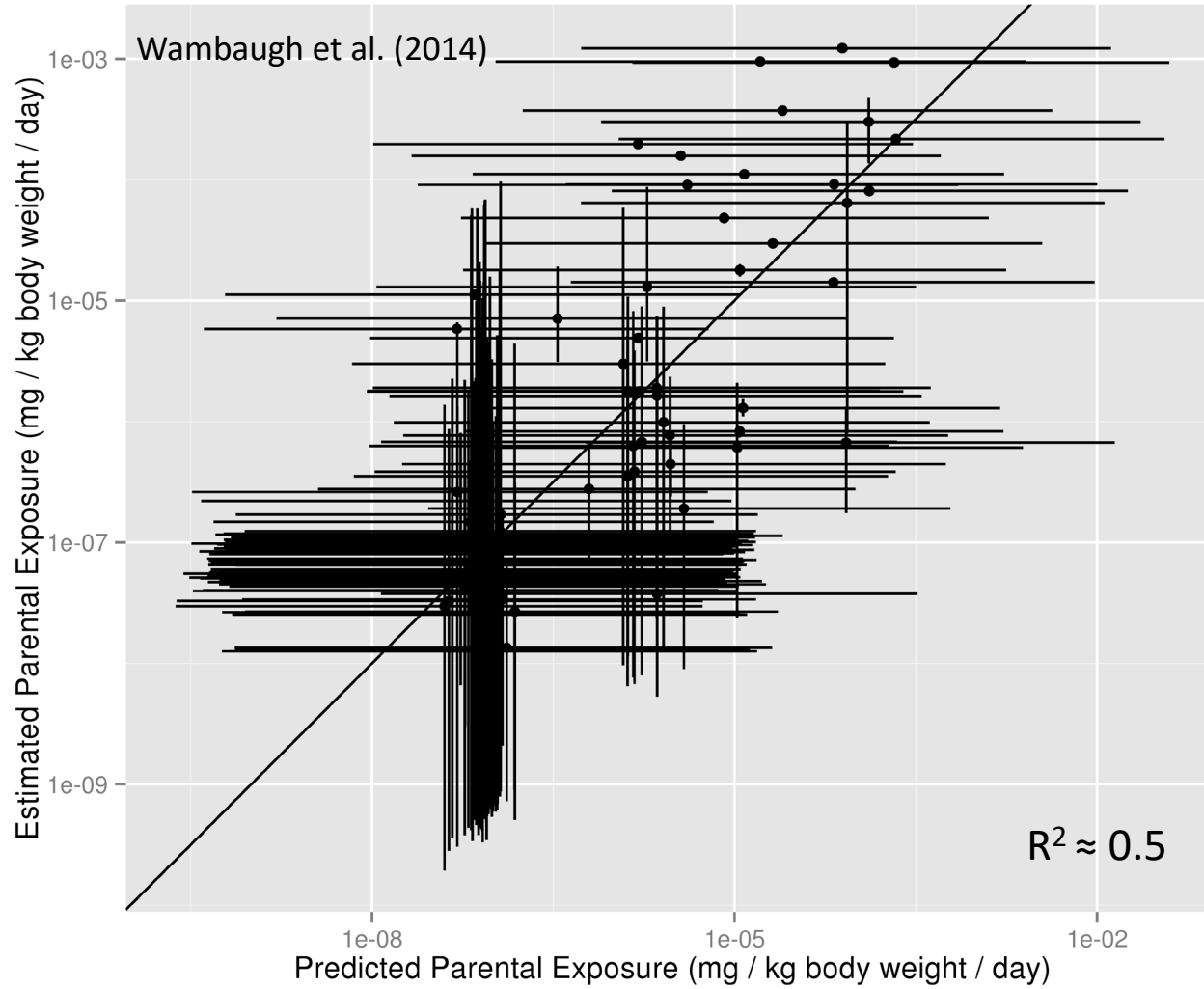
Not all models have predictions for all chemicals

- We can run SHEDS-HT (Isaacs et al., 2014) for ~2500 chemicals

What do we do for the rest?

- Assign the average value?
- Zero?

SEEM Analysis of NHANES Data

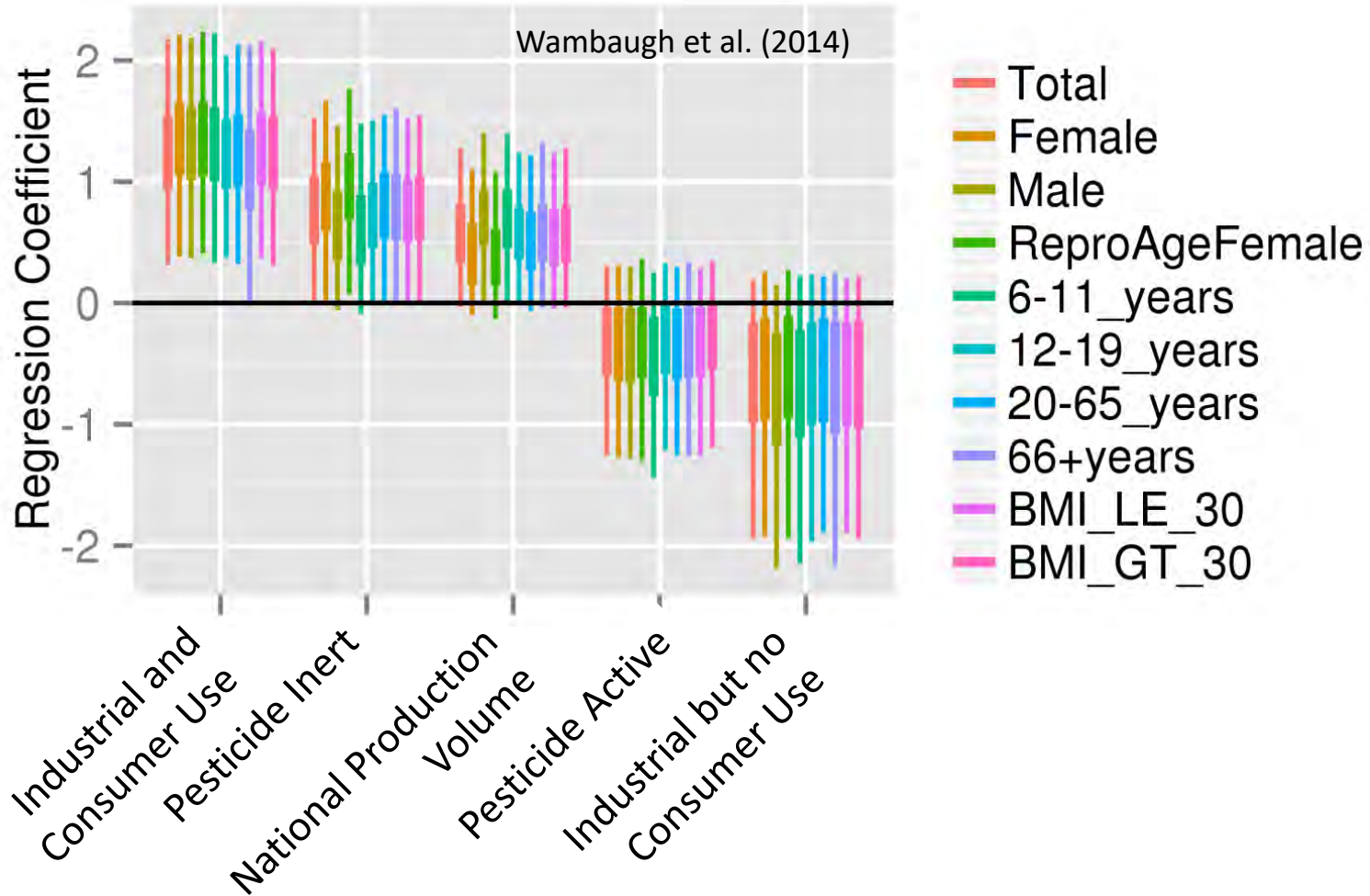


$R^2 \approx 0.5$ indicates that we can predict 50% of the chemical to chemical variability in median NHANES exposure rates

Same five predictors work for all NHANES demographic groups analyzed – stratified by age, sex, and body-mass index:

- Industrial and Consumer use
- Pesticide Inert
- Pesticide Active
- Industrial but no Consumer use
- Production Volume

Heuristics of Exposure



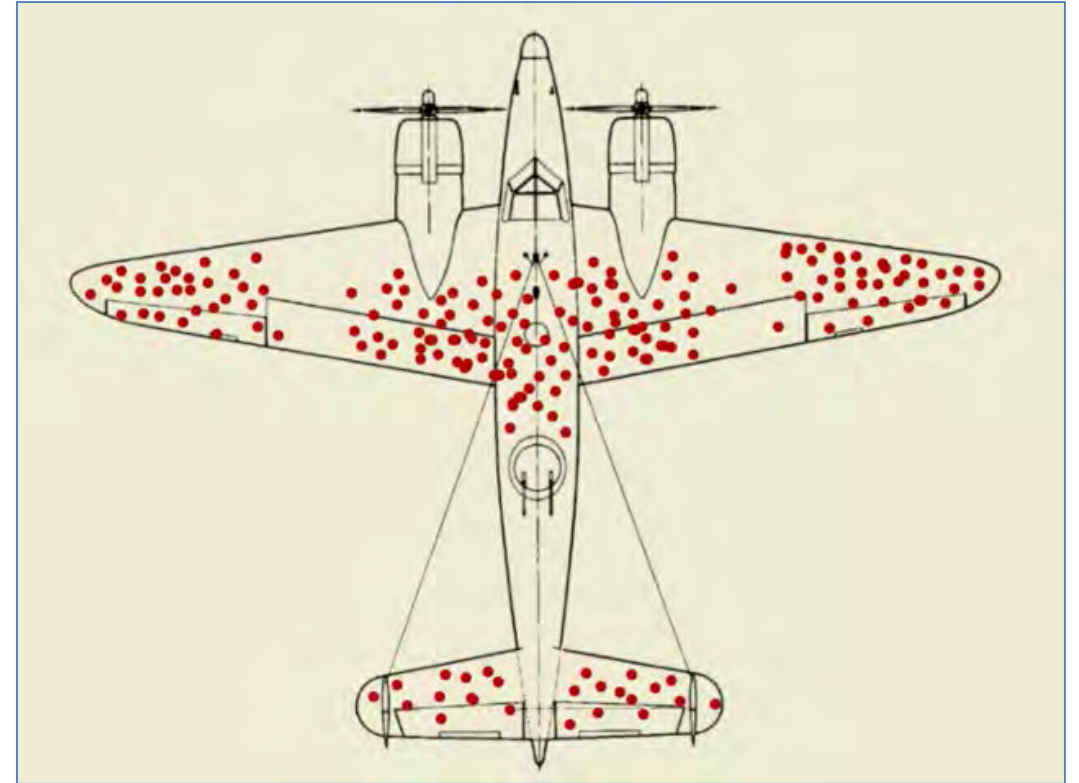
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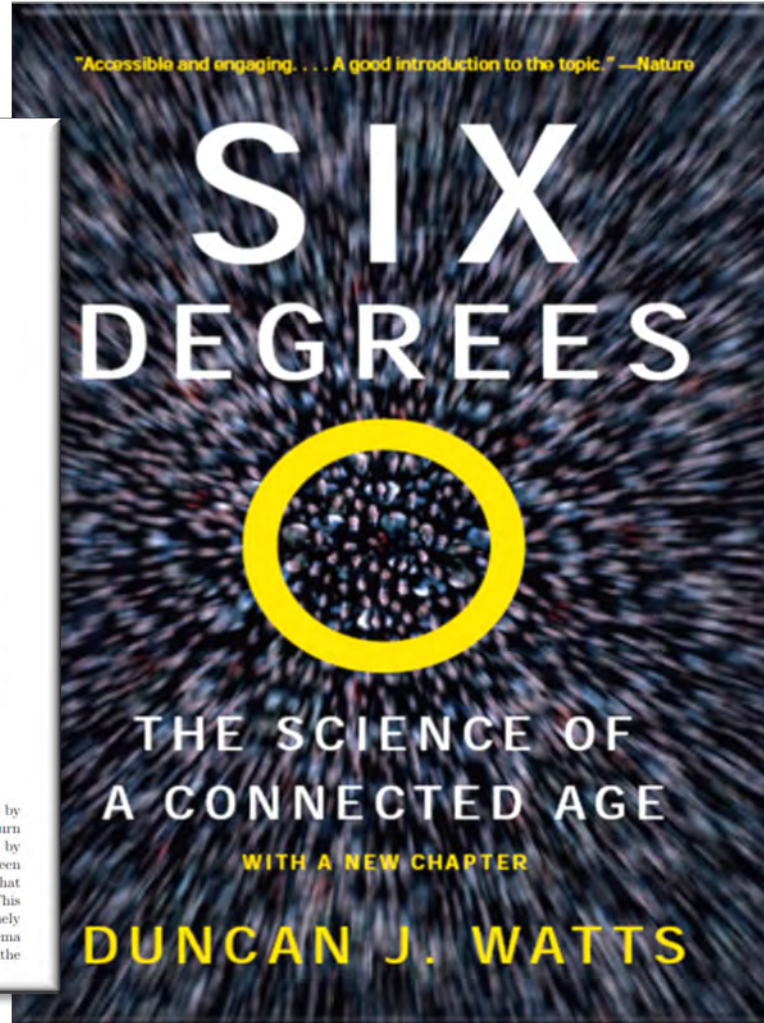
- Industrial and Consumer use
- Pesticide Inert
- Pesticide Active
- Industrial but no Consumer use
- Production Volume

Correlation is Not Causation

- Wambaugh et al. (2014) found that “pesticide inerts” had higher than average levels in biomonitoring data, while “pesticide actives” had lower than average
- In World War II, the Royal Air Force (UK) wanted to armor planes against anti-aircraft fire
 - Initial proposal was to place armor wherever bullet holes were most common
 - Mathematician Abraham Wald pointed out that they were looking at the planes that had returned
 - *See Drum, Kevin (2010) “The Counterintuitive World”*
- Pesticide inerts have many other uses, but there are more stringent reporting requirements for pesticides
 - **Exposure is occurring by other pathways**

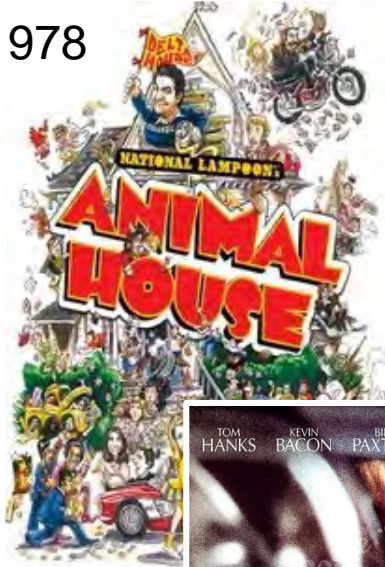


The Six Degrees of Kevin Bacon



Kevin Bacon

1978



1984



1992



1995



Kevin Bacon

1990



Michael B. Jordan



Connectedness to Michael B. Jordan

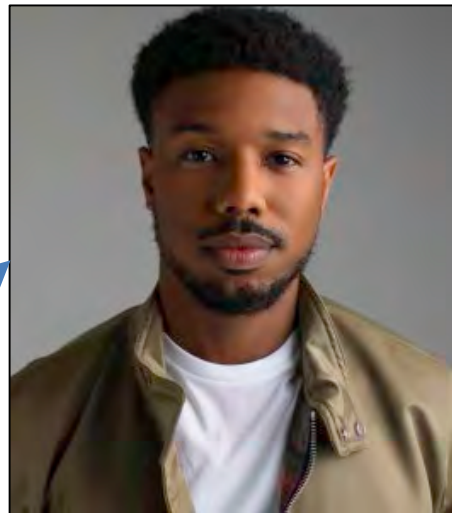


Frances McDormand
Best Actress Winner 2018

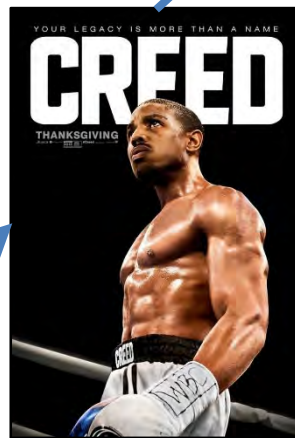
Hail Caesar
McDormand &
Channing Tatum



GI Joe: Retaliation
Tatum & Bruce Willis

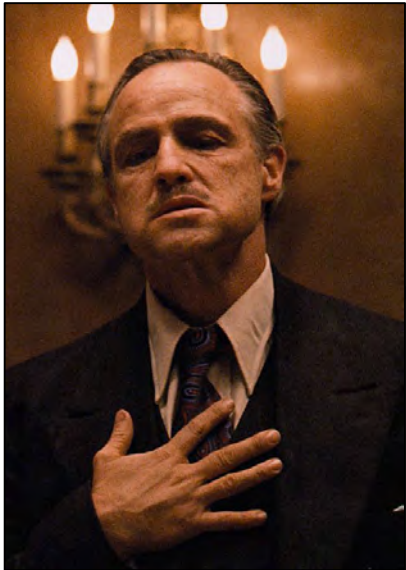


Expendables
Willis &
Sylvester Stallone



Creed
Stallone & Jordan

Connectedness to Michael B. Jordan

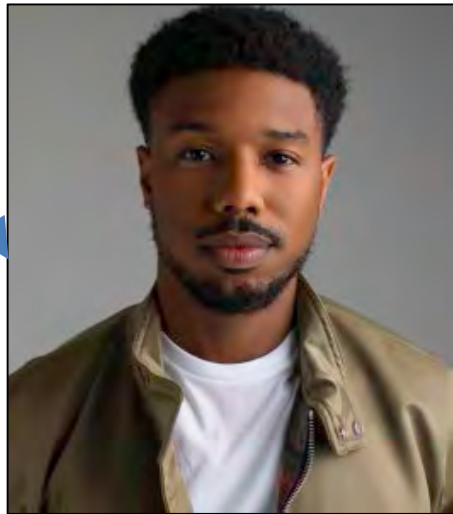


Marlon Brando
Best Actor 1954 and 1972
Died 2004

**Avengers:
Infinity War**
Paltrow &
Chadwick
Boseman



Black Panther
Boseman & Jordan



Superman
with Gene Hackman



The Royal Tenenbaums
Hackman & Gwyneth Paltrow

letters to nature

typically slower than $\sim 1 \text{ km s}^{-1}$) might differ significantly from what is assumed by current modelling efforts²⁷. The expected equation-of-state differences among small bodies (ice versus rock, for instance) presents another dimension of study; having recently adapted our code for massively parallel architectures (K. M. Olson and E.A. manuscript in preparation), we are now ready to perform a more comprehensive analysis.

The exploratory simulations presented here suggest that when a young, non-porous asteroid (if such exist) suffers extensive impact damage, the resulting fracture pattern largely defines the asteroid's response to future impacts. The stochastic nature of collisions implies that small asteroid interiors may be as diverse as their shapes and spin states. Detailed numerical simulations of impacts, using accurate shape models and rheologies, could shed light on how asteroid collisional response depends on internal configuration and shape, and hence on how planetesimals evolve. Detailed simulations are also required before one can predict the quantitative effects of nuclear explosions on Earth-crossing comets and asteroids, either for hazard mitigation²⁸ through disruption and deflection, or for resource exploitation²⁹. Such predictions would require detailed reconnaissance concerning the composition and internal structure of the targeted object. □

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Watts and Strogatz (1998)

Collective dynamics of 'small-world' networks

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Networks of coupled dynamical systems have been used to model biological oscillators¹, Josephson junction arrays², excitable media³, neural networks^{4–7}, spatial games⁸, genetic control networks⁹ and many other self-organizing systems. Ordinarily, the connection topology is assumed to be either completely regular or completely random. But many biological, technological and social networks lie somewhere between these two extremes. Here we explore simple models of networks that can be tuned through this middle ground: regular networks 'rewired' to introduce increasing amounts of disorder. We find that these systems can be highly clustered, like regular lattices, yet have small characteristic path lengths, like random graphs. We call them 'small-world' networks, by analogy with the small-world phenomenon^{10,11} (popularly known as six degrees of separation¹²). The neural network of the worm *Caenorhabditis elegans*, the power grid of the western United States, and the collaboration graph of film actors are shown to be small-world networks. Models of dynamical systems with small-world coupling display enhanced signal-propagation speed, computational power, and synchronizability. In particular, infectious diseases spread more easily in small-world networks than in regular lattices.

To interpolate between regular and random networks, we consider the following random rewiring procedure (Fig. 1). Starting from a ring lattice with n vertices and k edges per vertex, we rewire each edge at random with probability p . This construction allows us to 'tune' the graph between regularity ($p = 0$) and disorder ($p = 1$), and thereby to probe the intermediate region $0 < p < 1$, about which little is known.

We quantify the structural properties of these graphs by their characteristic path length $L(p)$ and clustering coefficient $C(p)$, as defined in Fig. 2 legend. Here $L(p)$ measures the typical separation between two vertices in the graph (a global property), whereas $C(p)$ measures the cliquishness of a typical neighbourhood (a local property). The networks of interest to us have many vertices with sparse connections, but not so sparse that the graph is in danger of becoming disconnected. Specifically, we require $n \gg k \gg \ln(n) \gg 1$, where $k \gg \ln(n)$ guarantees that a random graph will be connected¹³. In this regime, we find that $L \sim n/2k \gg 1$ and $C \sim 3/4$ as $p \rightarrow 0$, while $L \sim L_{\text{rand}} = -\ln(n)/\ln(k)$ and $C \sim C_{\text{rand}} = -\ln(k) \ll 1$ as $p \rightarrow 1$. Thus the regular lattice at $p = 0$ is a highly clustered, large world where L grows linearly with n , whereas the random network at $p = 1$ is a poorly clustered, small world where L grows only logarithmically with n . These limiting cases might lead one to suspect that large C is always associated with large L , and small C with small L .

On the contrary, Fig. 2 reveals that there is a broad interval of p over which $L(p)$ is almost as small as L_{rand} yet $C(p) \gg C_{\text{rand}}$. These small-world networks result from the immediate drop in $L(p)$ caused by the introduction of a few long-range edges. Such 'short cuts' connect vertices that would otherwise be much farther apart than L_{rand} . For small p , each short cut has a highly nonlinear effect on L , contracting the distance not just between the pair of vertices that it connects, but between their immediate neighbourhoods, neighbourhoods of neighbourhoods and so on. By contrast, an edge

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Small World Networks

Travers and Milgram (1977):

296 arbitrary individuals in Nebraska and Boston were asked to give a letter to an acquaintance most likely to help it reach a target person in Massachusetts. 64 reached the target person, average number of intermediaries was 5.2

Collins and Chow (1998)

It's a small world

James J. Collins and Carson C. Chow

The concept of Six Degrees of Separation has been formalized in so-called 'small-world networks'. The principles involved could be of use in settings as diverse as improving networks of cellular phones and understanding the spread of infections.

A few years ago, on American campuses, it was popular to play Six Degrees of Kevin Bacon. In this game, participants attempt to link the actor Kevin Bacon to any other actor through as few common films and co-stars as possible. Links are formed directly between Bacon and another actor if they appeared in the same film or indirectly through a chain of co-stars in different films (Fig. 1).

In the world of mathematics, a similar amusement involves assessing one's Erdős number, which measures the number of links needed to connect one to the prolific mathematician Paul Erdős through jointly authored papers. For example, individuals have an Erdős number of 1 if they co-authored a paper with Erdős. If one of their co-authors wrote a paper with Erdős, then they have an Erdős number of 2, and so forth. It has been pointed out¹ that Dan Kleiman has a combined Erdős/Bacon number of 3 because he wrote a paper with Erdős and appeared in *Good Will Hunting* with Minnie Driver, who appeared with Bacon in *Sleepers*.

These games are related to the popular concept of Six Degrees of Separation², which is based on the notion that everyone in the world is connected to everyone else through a chain of at most six mutual acquaintances. If two people have one mutual acquaintance, then they have one degree of separation. The estimate of six degrees of separation, which is related to the small-world phenomenon³, arises from pioneering empirical work by Milgram⁴ and can be understood heuristically from a somewhat unrealistic assumption of random connectivity. That is, if each person knows about one hundred individuals, and given that there are about a billion people on the Earth, then seven connections or six degrees of separation are enough to link everyone together.

On page 440 of this issue⁵, Watts and Strogatz formalize this idea in what they call small-world networks. They demonstrate through numerical simulations that a network need not be very random to get this small-world effect. They consider a connected network with nodes and links. In the friendship analogy, each node represents a person and each link represents a single connection to an acquaintance. They then define



Figure 1 Three degrees. Because Kevin Bacon has appeared in many films, most actors have low Bacon numbers and the game Six Degrees of Separation has declined in popularity. It is possible to centre the game around a newer star such as Leonardo DiCaprio. These film stills, running clockwise, show that in this case there are at most three degrees of separation between DiCaprio and Helena Bonham-Carter, through Kate Winslet (*Titanic*, Columbia TriStar; *Sense and Sensibility*, Columbia TriStar), Emma Thompson (*Sense and Sensibility*, Much Ado About Nothing, Entertainment Films) and Kenneth Branagh (*Much Ado About Nothing*, *Frankenstein*, Columbia TriStar). Short cuts between cliques could be created in this game through some of DiCaprio's well-connected co-stars such as Sharon Stone (*The Quick and the Dead*, TriStar; not shown).

news and views

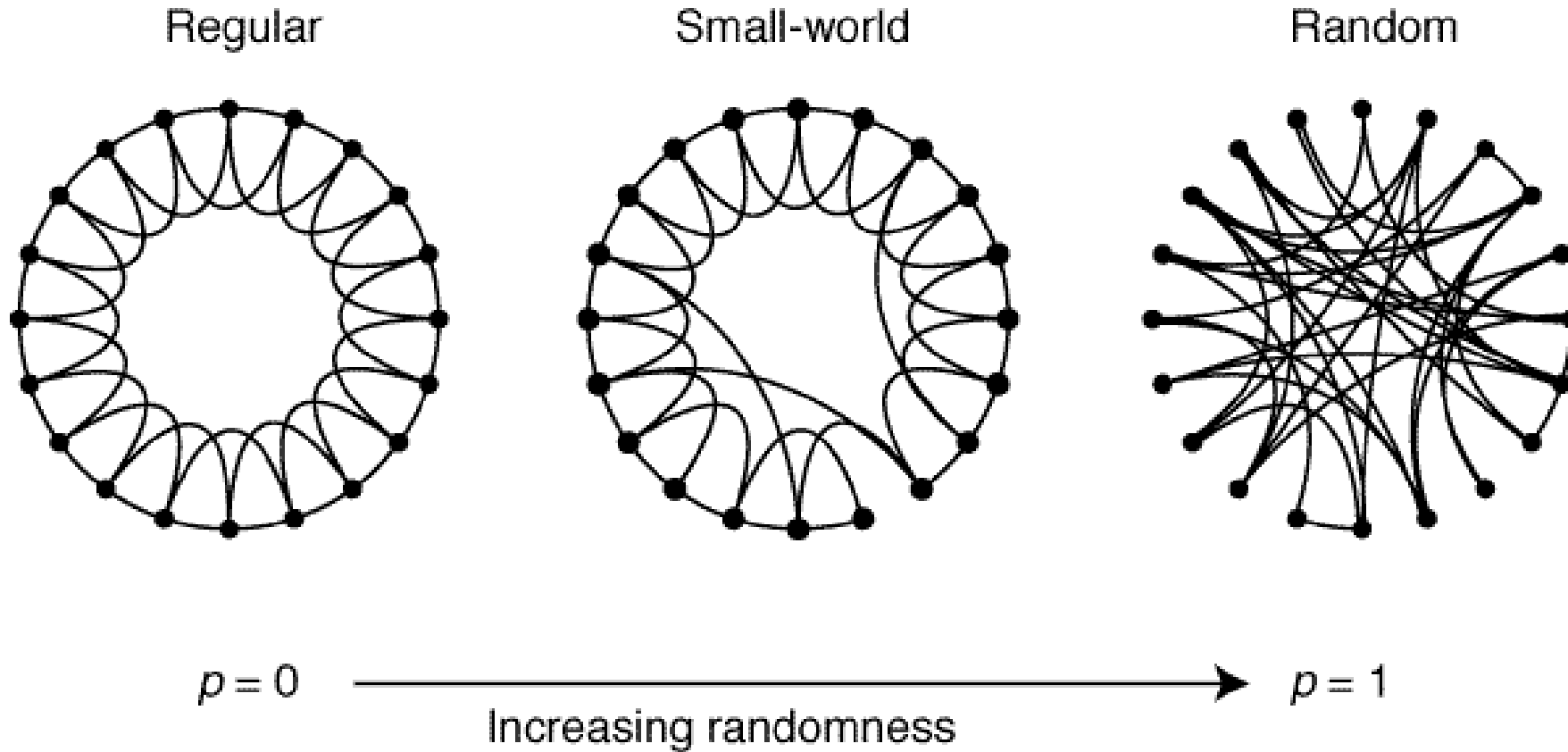
length is short, scaling logarithmically with the size of the network.

What Watts and Strogatz⁵ do is to shift gradually from a regular network to a random network by increasing the probability of making random connections from 0 to 1 (see Fig. 1, page 441). They then measure the characteristic path length and the amount of clustering of the network as a function of the amount of randomness. They find that path length and clustering depend differently on the amount of randomness in the network. The characteristic path length drops quickly, whereas the amount of clustering drops rather slowly. This leads to a small-world network in which the amount of clustering is high and the characteristic path length is short. So a small world can exist even when the cliquishness is imperceptibly different from that of a large world.

The explanation for this effect is that it only takes a few short cuts between cliques to turn a large world into a small world. In the friendship analogy, it only takes a small number of well-connected people to make a world small. The interesting and surprising thing is that it is impossible to determine whether or not you live in a small world or a large world from local information alone. The average person (node) is not directly associated with the key people (the clique-linkers).

Small-world connectivity has consequences that could be good or bad,

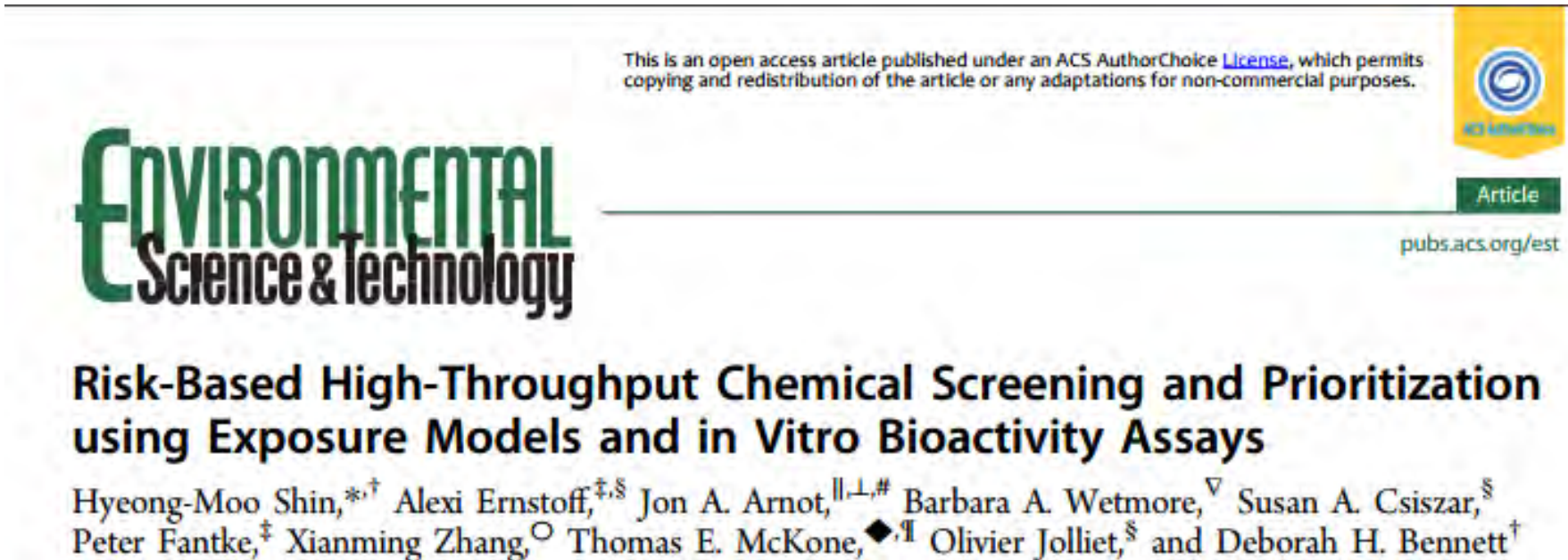
Complex is Not the Same as Random



Watts and Strogatz (1998)

Knowledge of Exposure Pathways Limits High Throughput Exposure Models

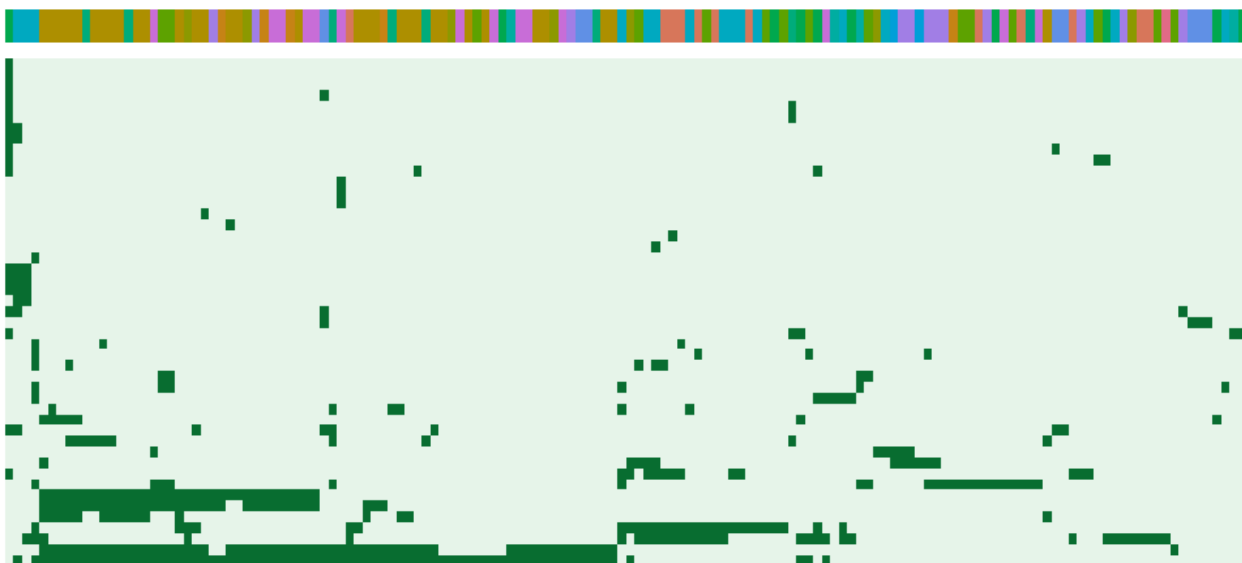
“In particular, the assumption that 100% of [quantity emitted, applied, or ingested] is being applied to each individual use scenario is a very conservative assumption for many compound / use scenario pairs.”



Chemical Use Identifies Relevant Pathways

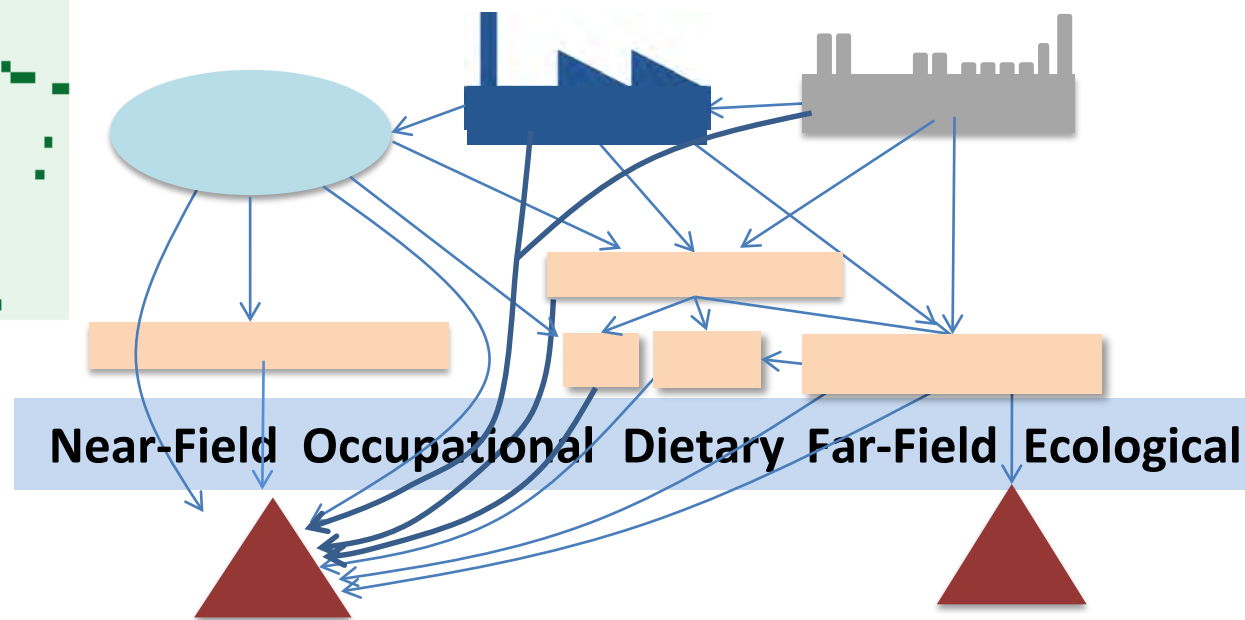
>2000 chemicals with Material Safety Data Sheets
(MSDS) in CPCPdb (Goldsmith *et al.*, 2014)

106 NHANES Chemicals



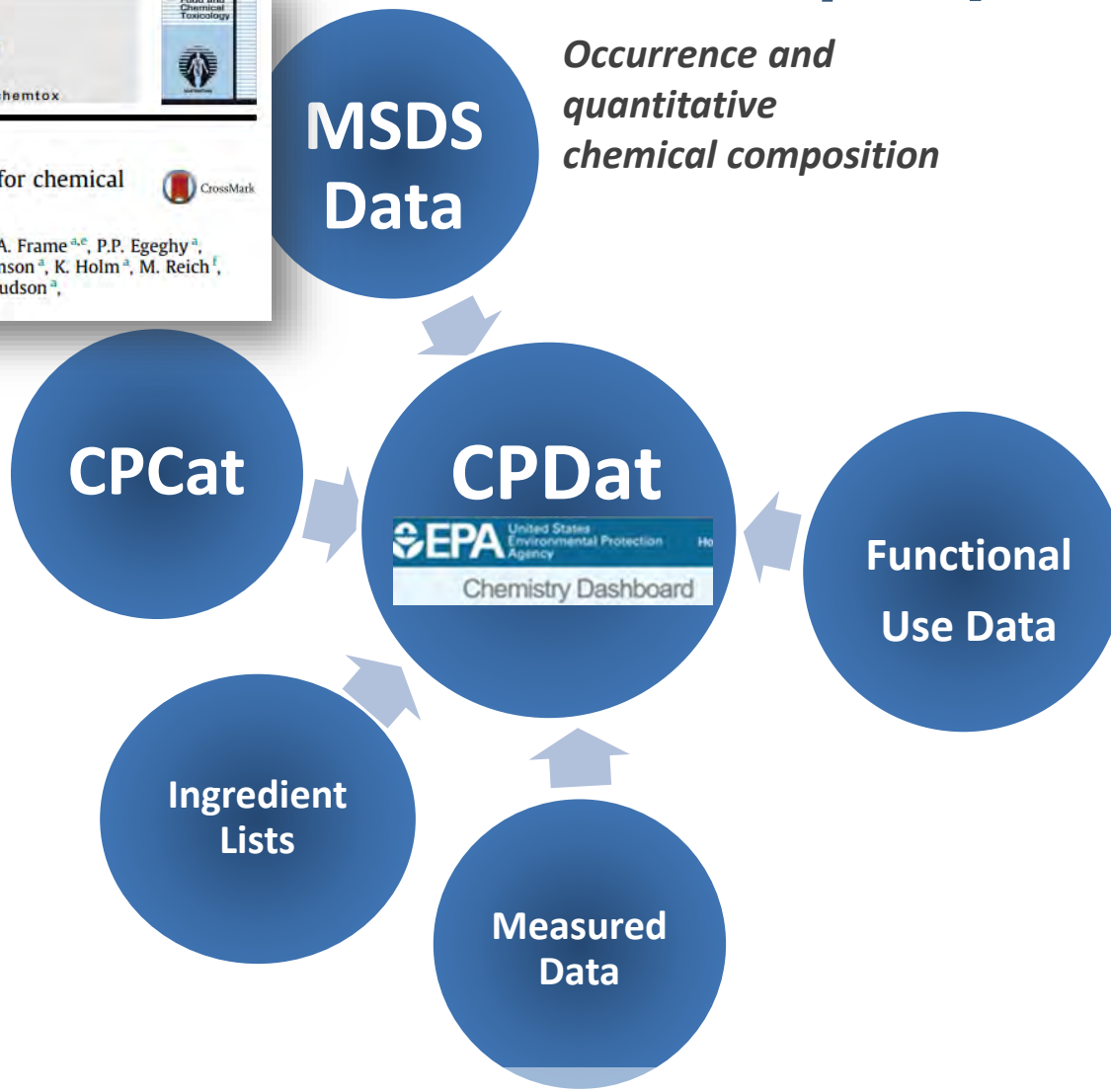
- | | |
|-----------------|---------------------|
| Apparel | Health |
| Auto and Tires | Home |
| Baby | Home Improvement |
| Beauty | Patio and Garden |
| Craft and Party | Pets |
| Electronics | Sports and Outdoors |
| Grocery | Toys |

Some pathways have
much higher average
exposures!



Near field sources have been known to be important at least since 1987 –
see Wallace, *et al.*

How Can we Know Chemical Use? Chemical Property NAMs



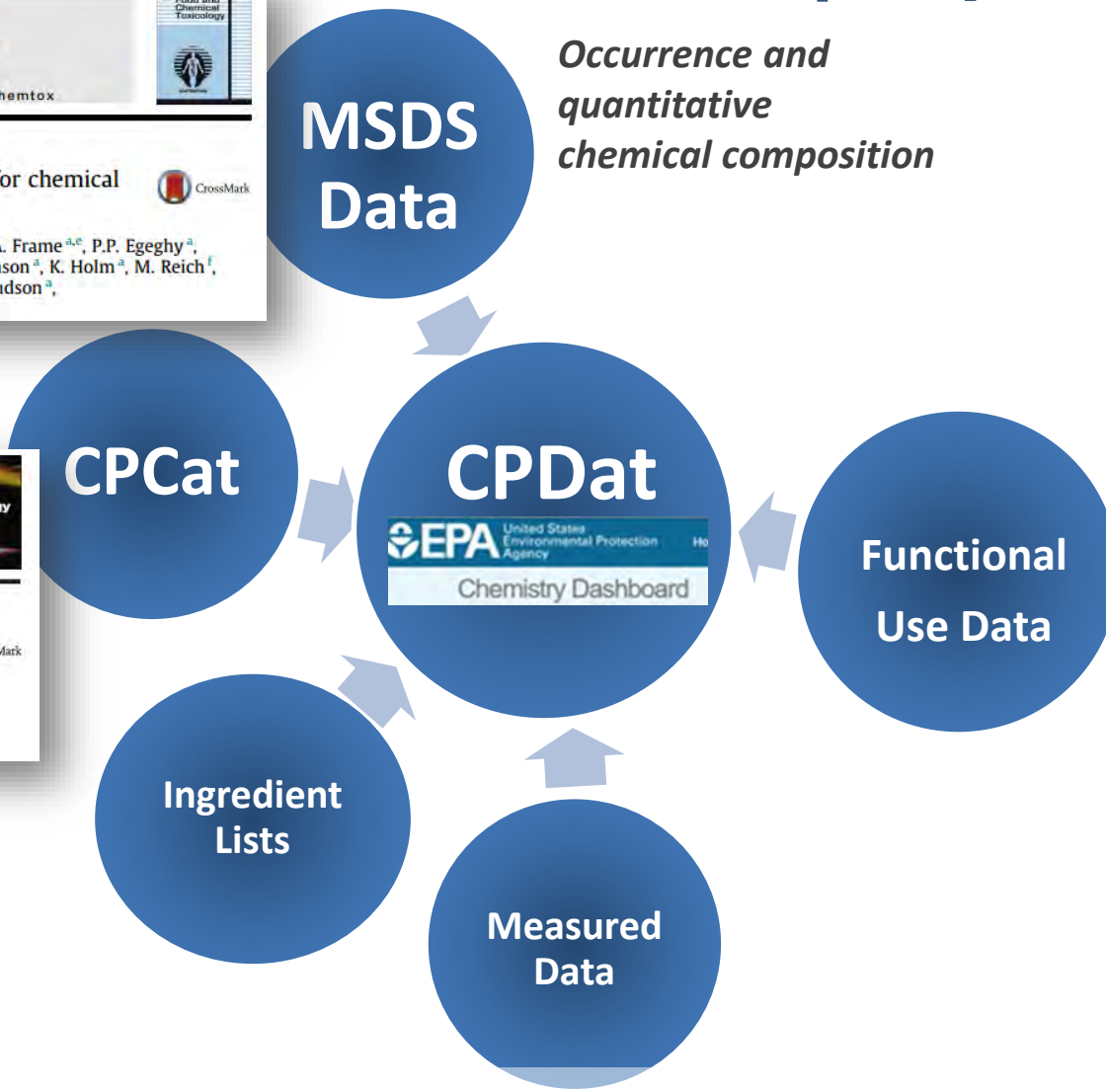
Product-specific
uses determined
using web spider
to click through
categories (e.g.,
home goods, bath
soaps, baby) to
find each product

**Material Safety
Data Sheet**
COM-35604

How Can we Know Chemical Use? Chemical Property NAMs



Broad "index" of chemical uses



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

How Can we Know Chemical Use? Chemical Property NAMs



Broad "index" of chemical uses



MSDS Data

Occurrence and quantitative chemical composition

CPCat

CPDat



Functional Use Data

Ingredient Lists

Occurrence data

Measured Data

How Can we Know Chemical Use?

Chemical Property NAMs



Broad "index" of chemical uses



MSDS Data

Occurrence and quantitative chemical composition

CPCat

CPDat

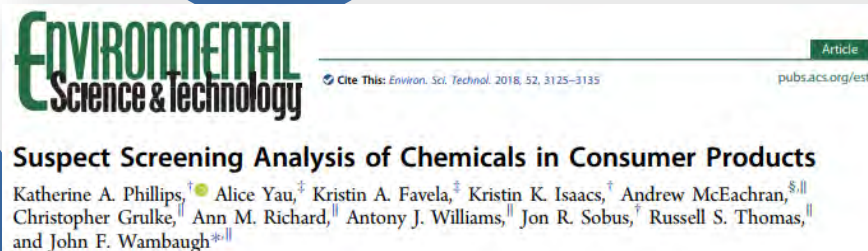


Functional Use Data

Ingredient Lists

Occurrence data

Measured Data



Measurement of chemicals in consumer products

How Can we Know Chemical Use?

Chemical Property NAMs



Broad "index" of chemical uses



MSDS Data

Occurrence and quantitative chemical composition

CPCat

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Green Chemistry

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Katherine A. Phillips,^{a,c} John F. Wambaugh,^b Christopher M. Grulke,^b Kathie L. Dionisio^c and Kristin K. Isaacs^c

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Environmental Science & Technology

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Suspect Screening Analysis of Chemicals in Consumer Products

Katherine A. Phillips,[†] Alice Yau,[‡] Kristin A. Favela,[‡] Kristin K. Isaacs,[‡] Andrew McEachran,^{§,||} Christopher Grulke,^{||} Ann M. Richard,^{||} Antony J. Williams,^{||} Jon R. Sobus,[†] Russell S. Thomas,^{||} and John F. Wambaugh^{*,||}

Measurement of chemicals in consumer products

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

How Can we Know Chemical Properties?

SCIENTIFIC DATA

Contents lists available at ScienceDirect

Food and Chemical Toxicology

ELSEVIER

journal homepage: www.elsevier.com/locate/foodchemtox

Development of a consumer product ingredient database for chemical exposure screening and prioritization

M.-R. Goldsmith^{a,*}, C.M. Grulke^a, R.D. Brooks^b, T.R. Transue^c, Y.M. Tan^a, A. Frame^{a,c}, P.P. Egeghy^a, R. Edwards^d, D.T. Chang^a, R. Tornero-Velez^a, K. Isaacs^a, A. Wang^{a,c}, J. Johnson^a, K. Holm^a, M. Reich^f, J. Mitchell^g, D.A. Vallero^a, L. Phillips^a, M. Phillips^a, J.F. Wambaugh^a, R.S. Judson^a, T.J. Buckley^a, C.C. Dary^a

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Broad "index" of chemical uses

Contents lists available at ScienceDirect

Toxicology Reports

ELSEVIER

journal homepage: www.elsevier.com/locate/toxrep

Exploring consumer exposure pathways and patterns of use for chemicals in the environment

Kathie L. Dionisio^a, Alicia M. Frame^{b,1}, Michael-Rock Goldsmith^{a,2}, John F. Wambaugh^b, Alan Liddell^{c,3}, Tommy Cathey^d, Doris Smith^b, James Vail^b, Alexi S. Ernstoff^e, Peter Fantke^e, Olivier Jolliet^f

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www.nature.com/jes

ORIGINAL ARTICLE

Consumer product chemical weight fractions from ingredient lists

Kristin K. Isaacs¹, Katherine A. Phillips¹, Derya Biryol^{1,2}, Kathie L. Dionisio¹ and Paul S. Price¹

MSDS Data

Occurrence and quantitative chemical composition

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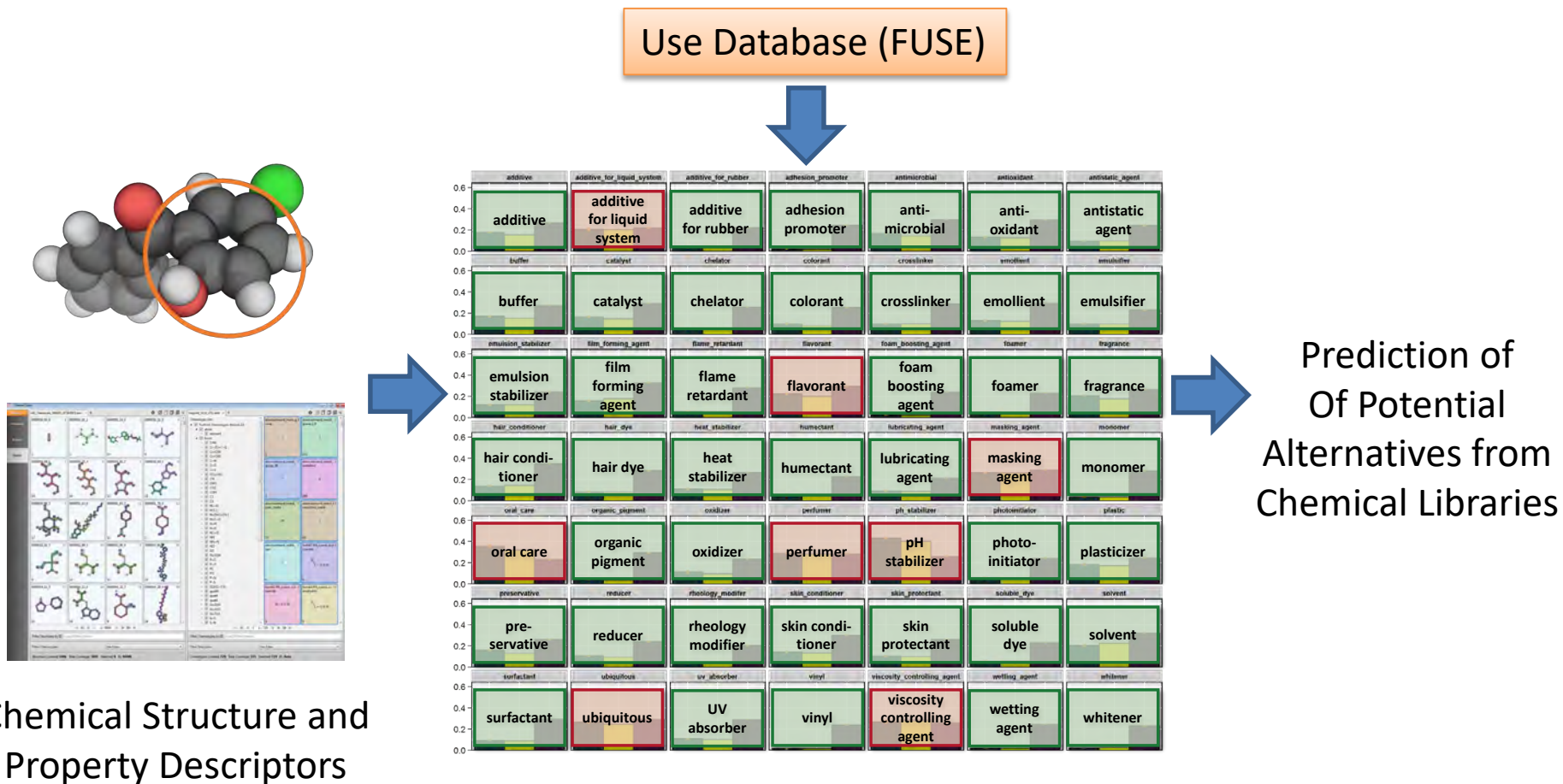
Katherine A. Phillips,[†] Alice Yau,[‡] Kristin A. Favela,[‡] Kristin K. Isaacs,[‡] Andrew McEachran,^{§,||} Christopher Grulke,^{||} Ann M. Richard,^{||} Antony J. Williams,^{||} Jon R. Sobus,[†] Russell S. Thomas,^{||} and John F. Wambaugh^{*,||}

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Exposure NAM: Machine Learning to Fill Data Gaps

EXAMPLE: Predicting Function Based on Structure



Machine Learning Based Classification Models
(Random Forest, Breiman, 2001)

Phillips et al. (2017)

What is “High Throughput”?

- Tox21: Testing one assay across 10,000 chemicals takes 1-2 days, but only 50 assays have been developed so far that can run that fast
- ToxCast: ~1100 off-the-shelf (pharma) assay-endpoints tested for up to 4,000 chemicals over the past decade, now developing new assays as well

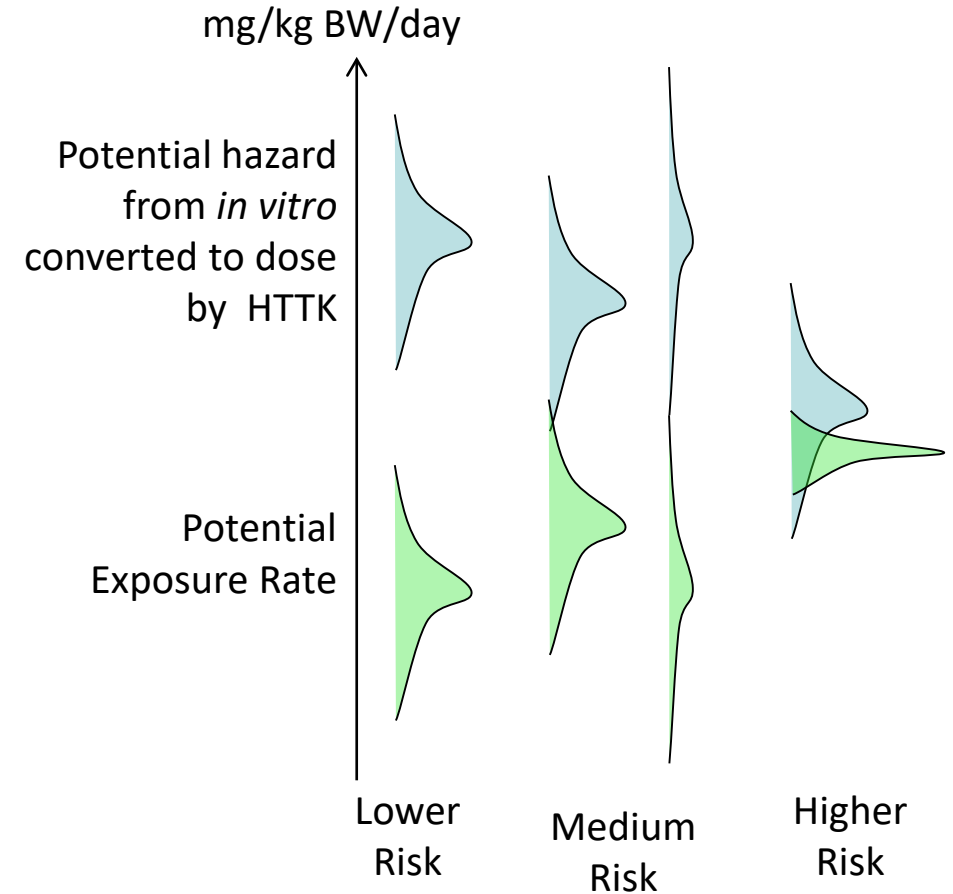
HTS tox assays often use single readout, such as fluorescence, across many chemicals, measuring concentration for toxicokinetics or exposure requires chemical-specific methods...

- ExpoCast: Ring et al. made *in silico* predictions for ~480,000 chemicals from structure, but based on NHANES monitoring for ~120 chemicals
 - Quantitative non-targeted analysis (NTA) may eventually provide greater evaluation data to reduce uncertainty
- HTTK: *In vitro* data on 944 chemicals collected for humans, starting with Rotroff et al. (2010)
 - Work continues to develop *in silico* tools, e.g. Sipes et al. (2016)

Our work is not done...

Summary

- A tapestry of laws covers the chemicals people are exposed to in the United States (Breyer, 2009)
- Many chemicals, ranging from industrial waste to dyes to packing materials, are covered by the recently updated Toxic Substances Control Act (TSCA) and administered by the EPA
- New approach methodologies (NAMs) are being developed to prioritize these existing and new chemicals for testing
- All data are being made public:
 - The CompTox Chemicals Dashboard (A search engine for chemicals) <http://comptox.epa.gov/>
 - R package “httk”: <https://CRAN.R-project.org/package=httk>



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



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