

Evolution of the CompTox Program at EPA

From Primitive Beginnings to Sophisticated Application

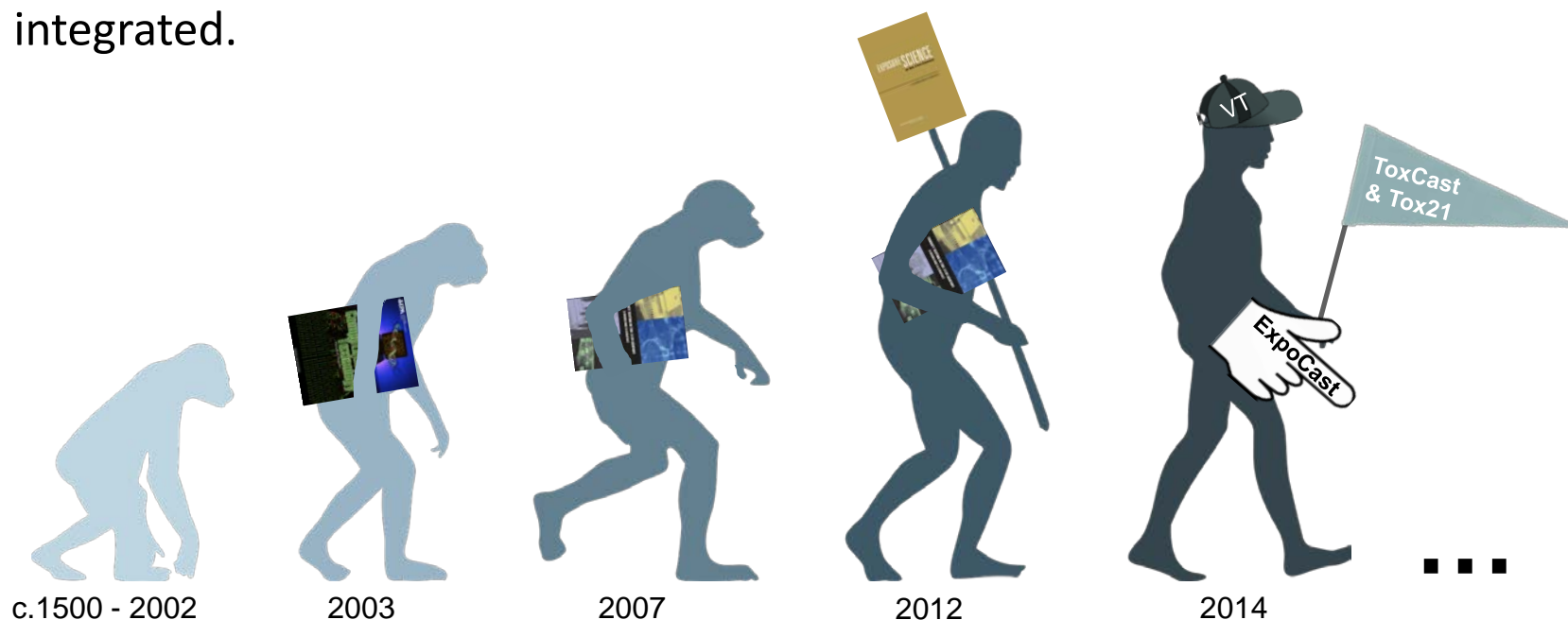


Global Chem
March 3, 2015

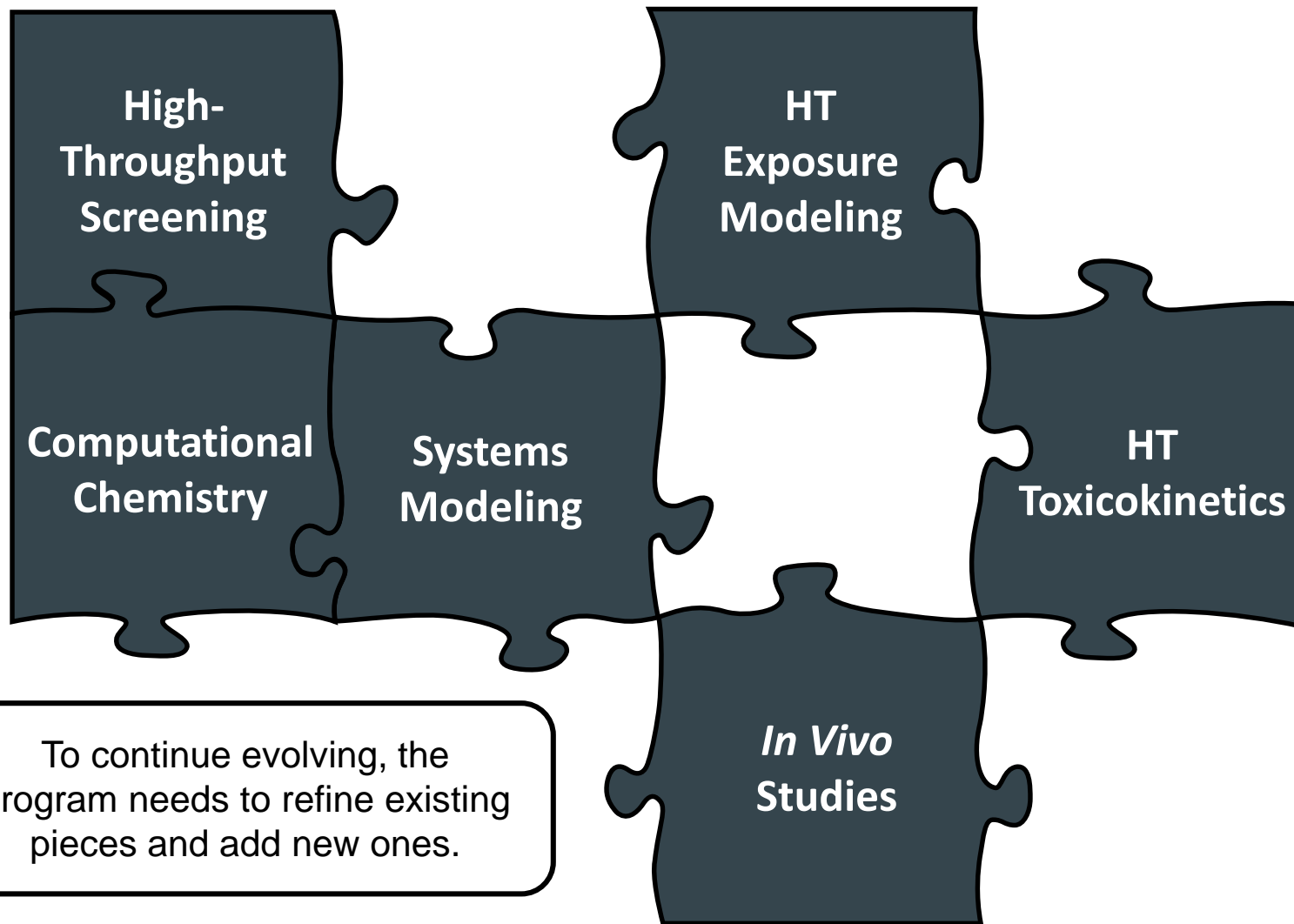
Rusty Thomas
Director
National Center for Computational Toxicology

Introduction

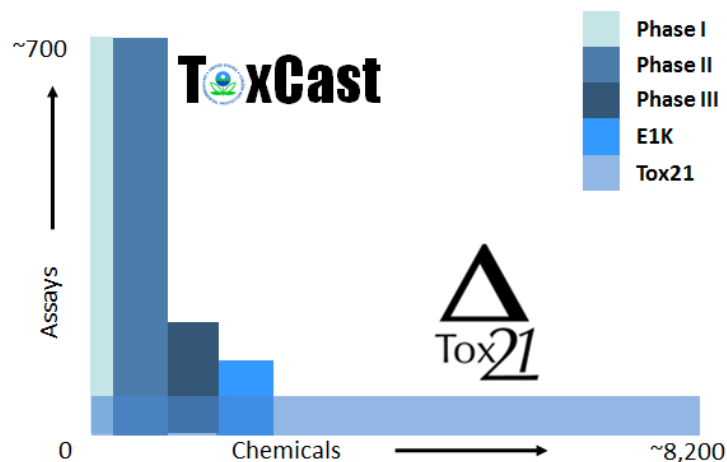
- The CompTox program at the Agency and the field in general have undergone punctuated and rapid evolution since its inception
- Diverse regulatory needs, complex scientific challenges, large influx of data, and guidance from expert committees have provided significant selection pressure
- The research will be increasingly broad in scope, multi-disciplinary, and highly integrated.



Continuing to Evolve



Evolution of High-Throughput Screening

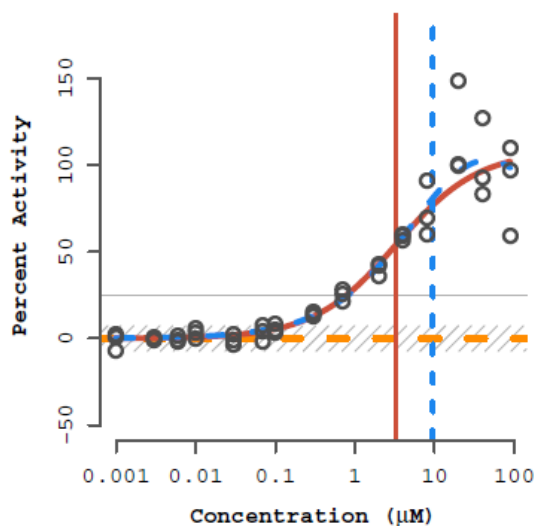


Present

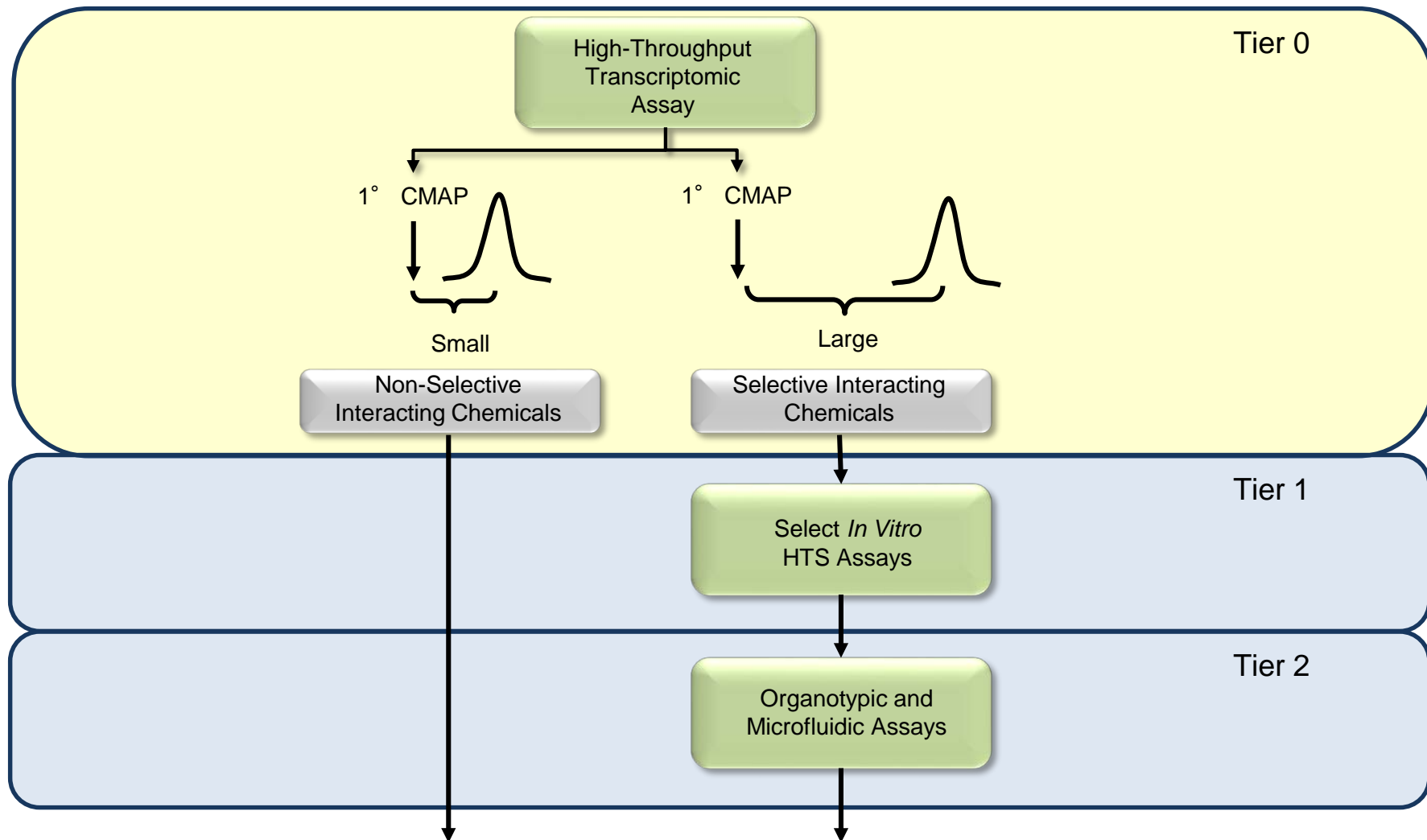
- ToxCast has screened ~2,000 chemicals across ~700 assay endpoints
- Tox21 has screened ~8,200 chemicals across ~50 endpoints
- ToxCast assay coverage represents over 327 genes and 293 pathways

Planned

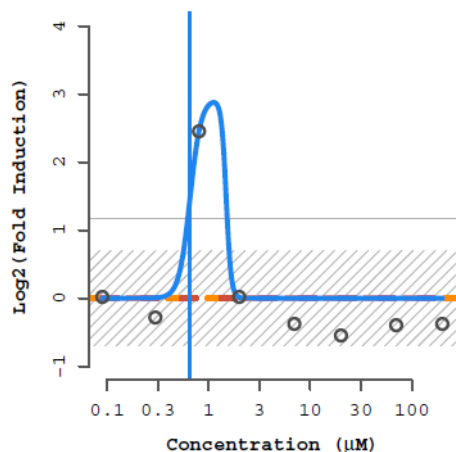
- Develop low cost, high-throughput global transcriptomic platform to screen across battery of cell types, concentration response, and time
- Capability to retrofit assays for metabolic competence
- Continue assay development for priority targets and complex organotypic cultures



Incorporating a Broad Biological Screening Platform



Efforts to Ensure HTS Data Quality and Increase Transparency



ASSAY: ARID117 (ATQ_Era_TRANS)

NAME: Thioglycolic acid
CHID: 26141 CASRN: 68-11-1
SPID(S): TX007664
L4ID: 420385

HILL MODEL (in red):
tp ga gw
val: 3.1e-11 -2.15 0.416
sd: NaN NaN NaN

GAIN-LOSS MODEL (in blue):
tp ga gw la lw
val: 2.93 -0.184 8 0.173 18
sd: 3.56 0.334 9.48 5.82 814

	CNST	HILL	GNLS
AIC:	20.14	26.14	17.79
PROB:	0.23	0.01	0.76
RMSE:	0.92	0.92	0.32

MAX_MEAN: 2.45 MAX_MED: 2.45 BMAD: 0.233

COFF: 1.17 HIT-CALL: 1 FITC: 50 ACTP: 0.77

FLAGS:

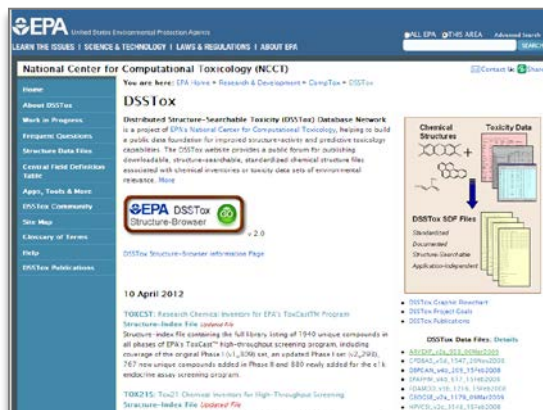
Only one conc above baseline, active
Borderline active

- Public release of Tox21 and ToxCast data on PubChem and EPA web site (raw and processed data)
- ToxCast data analysis pipeline has been completely revamped
 - More statistically rigorous and less prone to outliers
 - Data quality flags to indicate concerns with chemical purity and identity, noisy data, systematic assay errors, and activity in range of cytotoxicity
- Tox21 and ToxCast chemical libraries undergoing analytical QC and results will be publicly released
- Release of ToxCast “Owner’s Manual”
 - Chemical Procurement and QC
 - Data Analysis
 - Assay Characteristics and Performance
- External audit on ToxCast data and data analysis pipeline
- Continued offering of webinars and workshops to educate stakeholders on high-throughput screening data analysis and interpretation

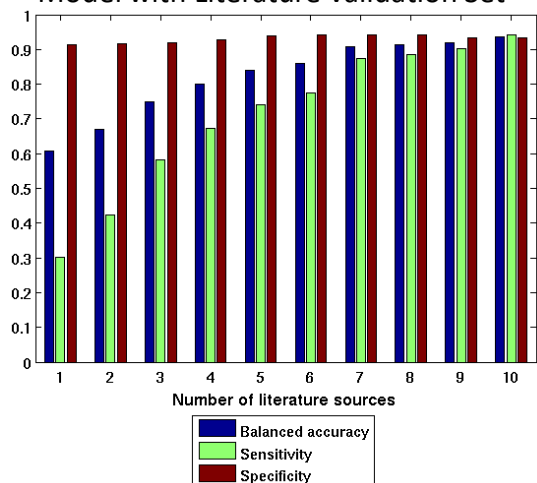
Evolution of Computational Chemistry

Present

- High quality, curated chemical structure database of 22,000 molecules in DSSTox
- Development of an open-source KNIME workflow for automated structure processing and QC
- Community consensus QSAR models for endocrine-related bioactivity (ER and AR) with 17 international participants



Performance of Consensus ER QSAR Model with Literature Validation Set



Planned

- Increase curated chemical structure database to 150,000 molecules with addition of QC flags to indicate confidence in structural associations
- Chemical grouping and read-across effort that incorporates both chemical structure and bioactivity

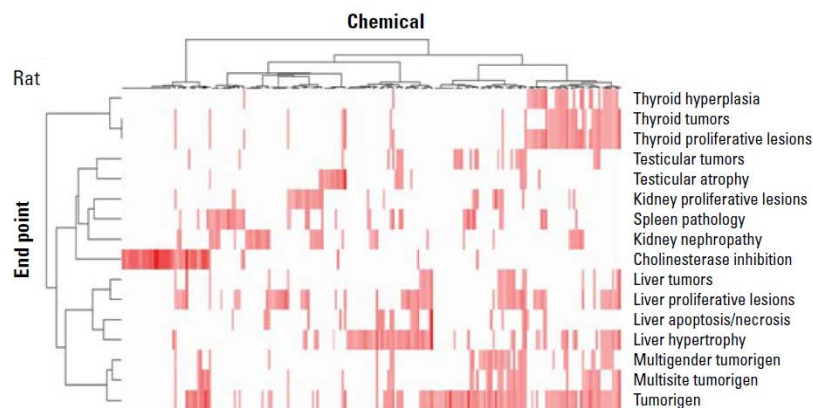
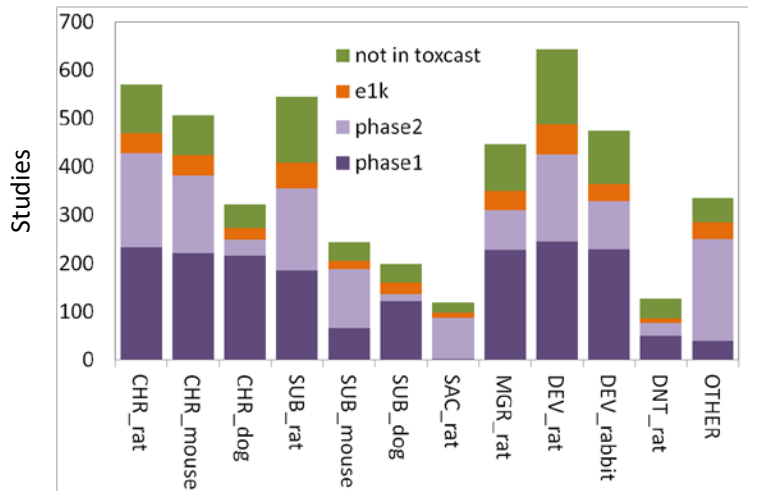
Evolution of ToxRefDB

Present

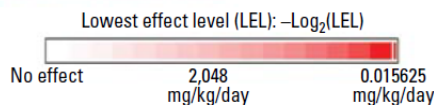
- *In vivo* endpoint data from 5,891 animal toxicology studies on ~1,110 unique chemicals
- A total of 654 chemicals (3311 studies) have LOAEL values, while 812 chemicals (2580 studies) have no adverse effect determination.
- Studies consist of chronic, subchronic, multigenerational, and developmental designs from DERs, NTP, open literature, and pharma

Planned

- Expand number of chemicals with critical effects and add quantitative dose-response data
- Add data from different sources (ECHA, E-Tox, Wignall et al.) and harmonize across databases
- Perform external QC review on ToxRefDB



Martin *et al.*, EHP 2009



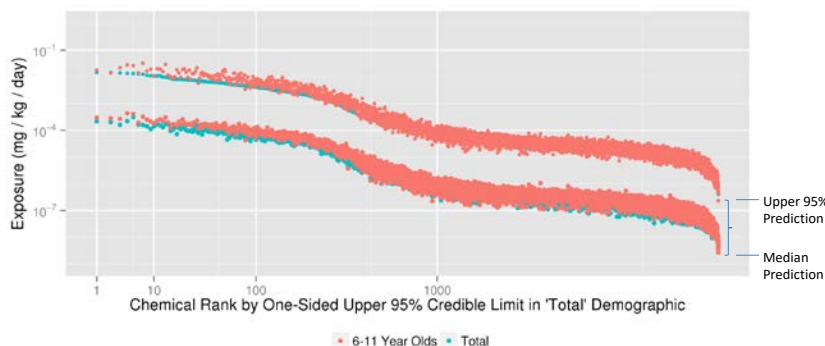
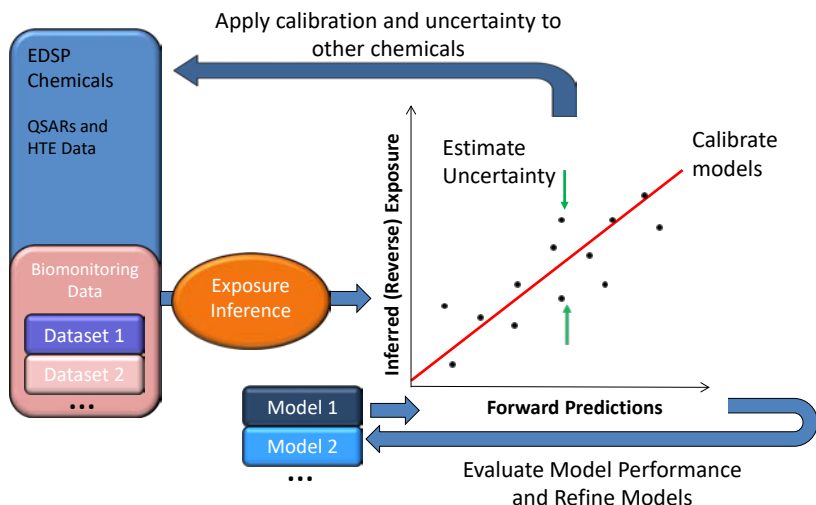
Evolution of High-Throughput Exposure Modeling

Present

- Computational exposure framework (SEEM) developed that provides exposure estimates for over 7,000 chemicals based on production volume and five chemical use categories
- A database of chemical-product categories (CPCat) that maps over 45,000 chemicals to ~8,000 product uses or functions

Planned

- Obtain additional biomonitoring data to expand domain of applicability
- Survey a range of consumer products and home goods for chemical ingredients and emissivity
- Incorporate SHEDS-HT into SEEM framework



Wambaugh *et al.*, EST 2014

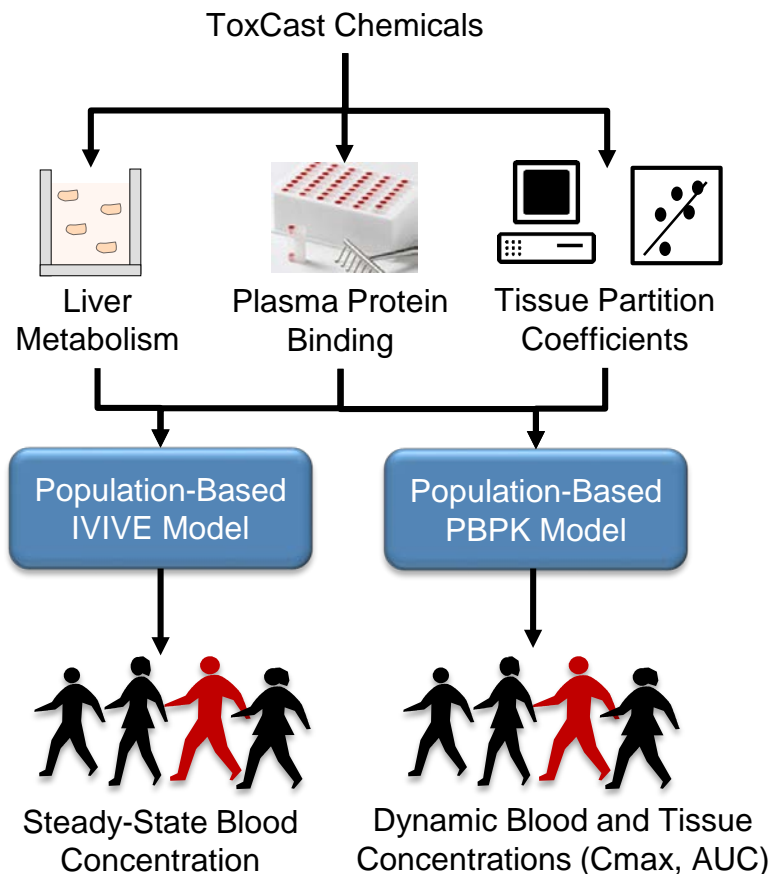
Evolution of High-Throughput Toxicokinetics

Present

- Steady-state IVIVE models for hundreds of chemicals based on limited high-throughput *in vitro* assays
- Implemented structure-based methods to estimate tissue partitioning
- Developed HT-Physiologically-Based Pharmacokinetic (HT-PBPK) models for hundreds of chemicals

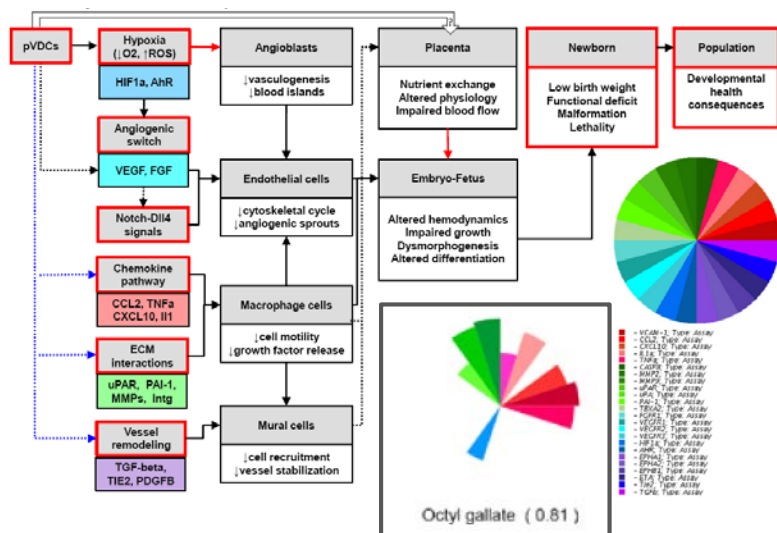
Planned

- Develop computational modeling framework to compare IVIVE with *in vivo* data and allow explicit estimates of uncertainty
- Separate chemical classes where we do either a good or poor job of predicting pharmacokinetics



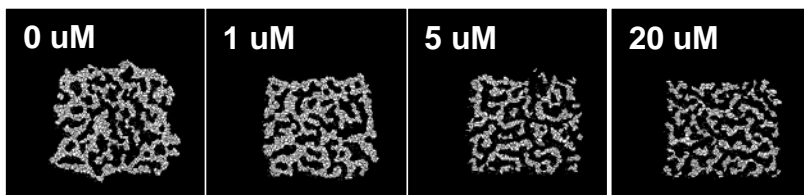
Evolution of Systems Modeling and Virtual Tissues

AOP for Developmental Vascular Disruption



Kleinstreuer *et al.*, PLoS Comp Bio, 2013

Model Simulations of Dev Vascular Disruption



Knudsen *et al.*, unpublished

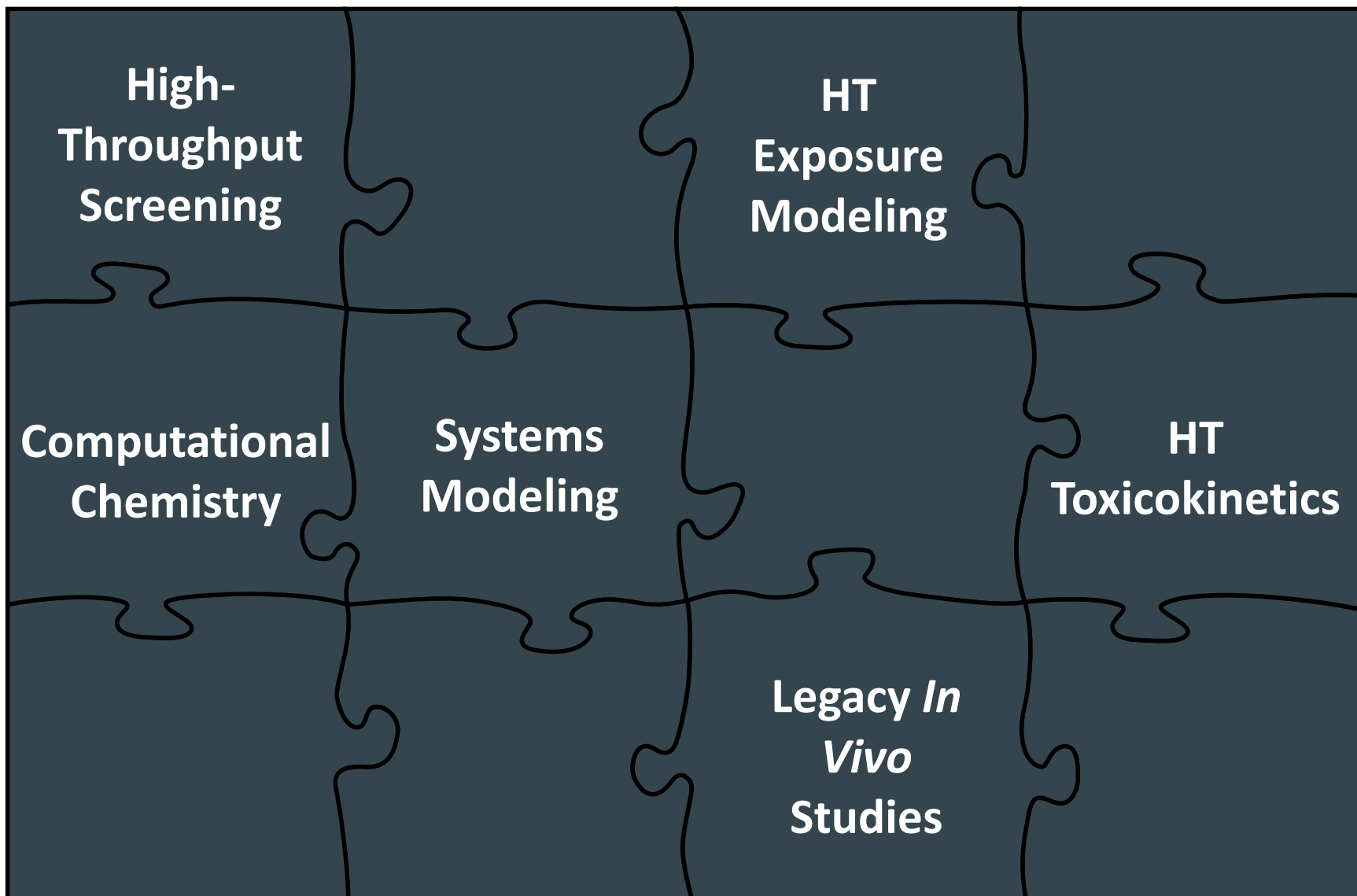
Present

- AOPs for embryonic vascular disruption, cleft palate, hypospadias, and limb (digit) defects
- Developed computational models for each AOP and used ToxCast data to parameterize models
- Validated model results with orthogonal organotypic assays and reference teratogens
- Developed first-generation methods for identifying cellular ‘tipping points’

Planned

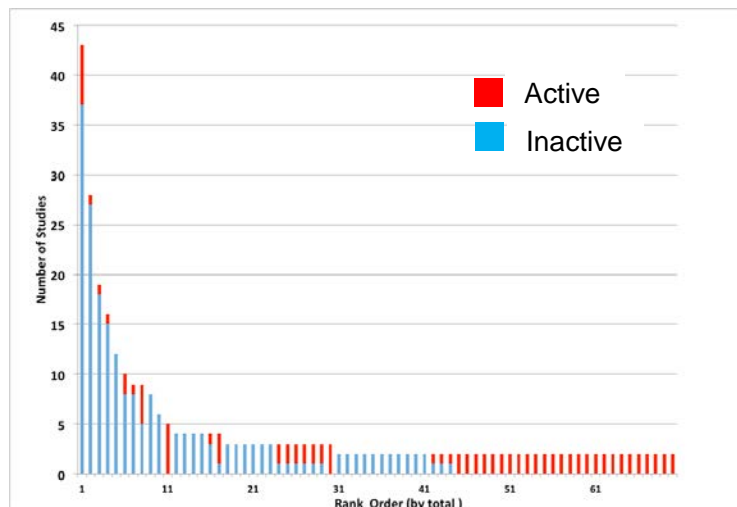
- Refinement and validation of methods identifying concentration-dependent ‘tipping points’
- Evaluate impact of model assumptions on shape of the dose response curve and model potential sensitive populations

Fitting the Pieces Together for the Next Evolutionary Chapter

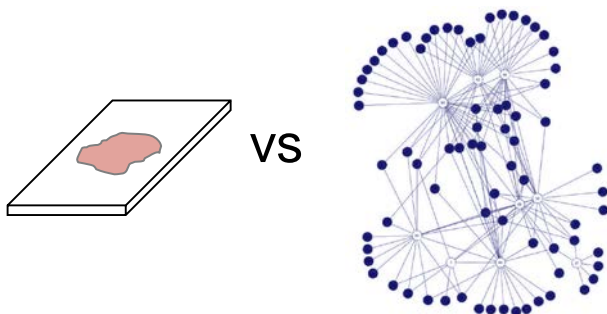


Challenges

Concordance of *In Vivo* Uterotrophic Studies



- Dealing with the “V” word
 - Defining a fit-for-purpose framework(s) that is time and resource efficient
 - Role of *in vivo* rodent studies
 - Incorporating the inherent uncertainty of the *in vivo* studies
- Moving from an apical to a molecular paradigm and defining adversity
- Predicting human safety vs. toxicity
- Integrating multiple data streams from the new approaches in a risk-based, weight of evidence assessment
- Ensuring a comprehensive screening and testing paradigm
- Quantifying uncertainty and variability
- Application to cumulative risk/mixtures



Questions?