

# Predictive Toxicology: Current Status and Future Outlook



**EBI-EMBL Industry Programme Workshop Predictive Toxicology** 

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# Regulatory Agencies Need to Make A Range of Decisions on Chemicals...

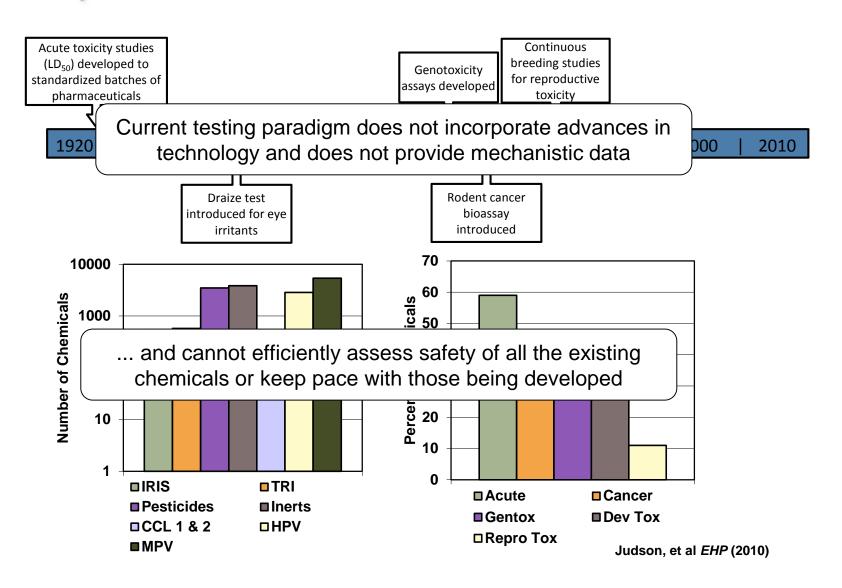
- Multiple drivers shape type of assessment
  - Regulatory scope
  - Economic considerations
  - Multiple applications
- Chemical assessments are "fit-forpurpose"
  - Prioritization (e.g., EDSP, PMN, SNUR)
  - Screening-level assessments (e.g., CCL, GreenChem)
  - Provisional assessments (e.g., PPRTVs)
  - Toxicity assessments (e.g., IRIS)
  - Risk assessments (e.g., MCLs, pesticides)

EPA	Workflow	Throughput	Data
ОРРТ	New chemicals: Premanufacture Notice (PMN) Existing chemicals: Significant New Use Rule (SNUR)	~1000/yr (90d/chem) ~84,000 total	III (II)
	Current Chemical Risk Assessments	~10 total	1
	DFE / Green Chemistry	~2500	I, II, III
OPP	Pesticide registration (PR)	~10 new/yr ~50 old/yr	1
	Pesticide re-registration	~1000/yr 24,576 total	1
OW	Chemical Contaminant List	6yr / ~6,000 total	1,11,111
	RegDet on CCL	Every 6yr / 90 total	1
	Unregulated Contaminant Monitoring	30/5yr	1
	Drinking Water Health Advisories		II, III
OLEM	Spills Brownfields Super Fund		

- Guideline animal testing data
- I Some *in vitro* bioactivity
- II Chemical structure data

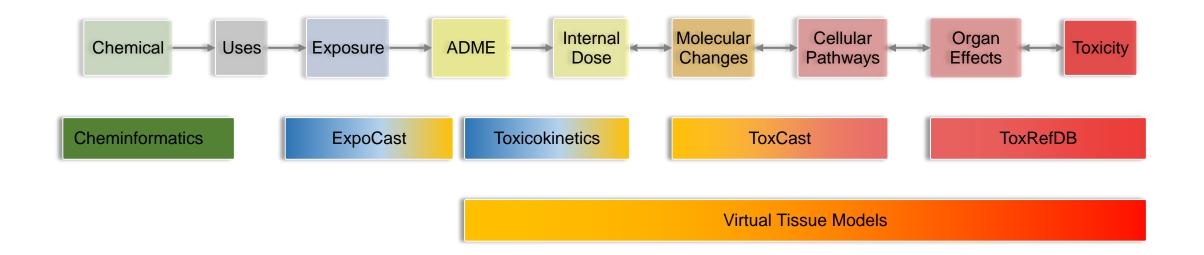


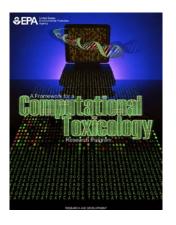
# Current System to Evaluate Chemicals is Antiquated and Inefficient





## Requires an Integrated and Multidisciplinary Solution





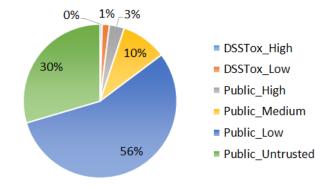
- National Center for Computational Toxicology established in 2005
- Currently staffed by ~60 employees
- Exists within the EPA's Office of Research and Development
- Home of the ToxCast and ExpoCast research efforts
- Key partner in U.S. Tox21 federal consortium



# Need to Start with a High Quality Chemistry Foundation



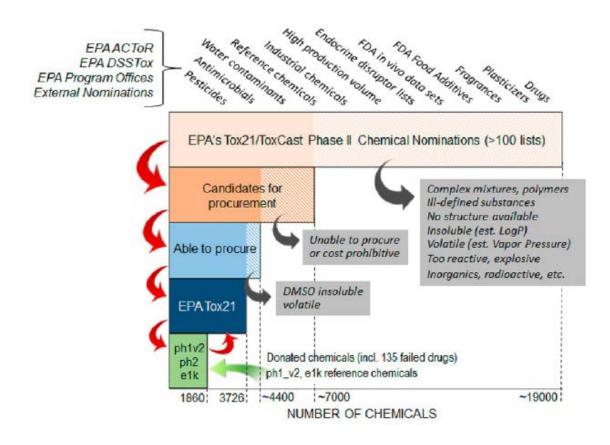
https://comptox.epa.gov



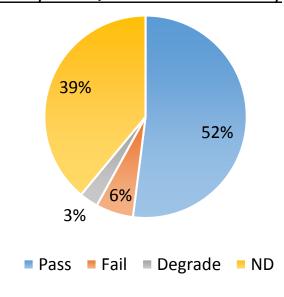
- Developing a centralized resource for curated chemical structure, identifier, and physical chemical properties of >700K unique substances with data quality flags
- Expand and curate training sets for QSAR models for phys-chem, environmental fate, and toxicological properties
- Use the centralized chemical resource as the foundation for an integrated hazard, bioactivity, pharmacokinetics, and exposure information



### Need to Start with a High Quality Chemistry Foundation



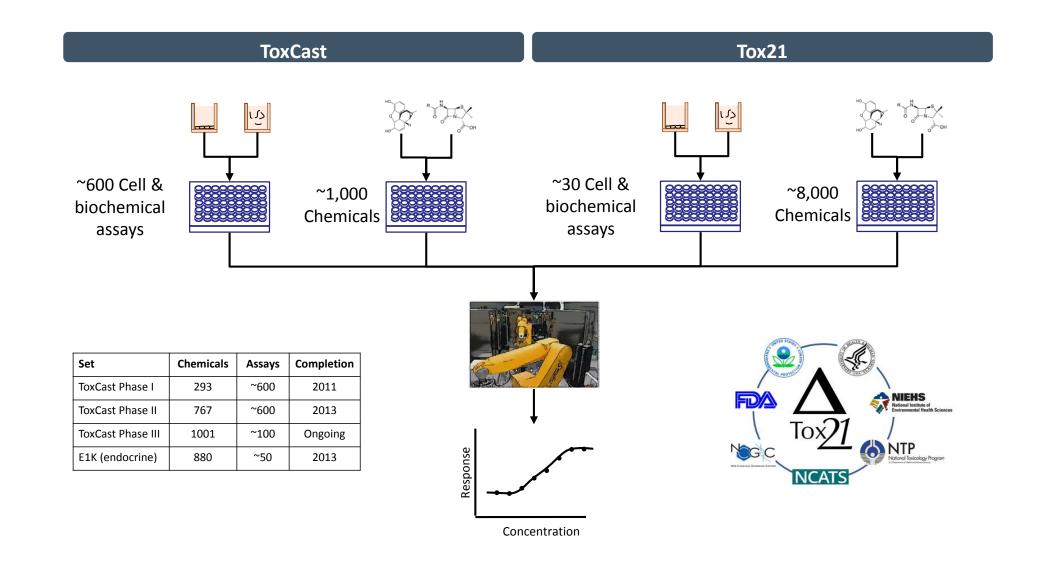
#### Analytical QC of Chemical Library



Pass = 
$$C$$
 (75%) or greater  
Fail =  $D$ ,  $F$ ,  $Ac$ ,  $Bc$ ,  $Cc$ 



### High-Throughput Bioactivity Screening





# ToxCast Incorporates a Diverse Array of High-Throughput *In Vitro* Assays

#### **Assay Provider**

ACEA
Apredica
Attagene
BioReliance
BioSeek
CeeTox
CellzDirect
Tox21/NCATS
NHEERL MESC
NHEERL Zebrafish
NovaScreen (Perkin Elmer)
Odyssey Thera
Vala Sciences

#### **Biological Response**

cell proliferation and death cell differentiation
Enzymatic activity
mitochondrial depolarization protein stabilization
oxidative phosphorylation reporter gene activation gene expression (qNPA) receptor binding receptor activity steroidogenesis

#### **Target Family**

response Element
transporter
cytokines
kinases
nuclear receptor
CYP450 / ADME
cholinesterase
phosphatases
proteases
XME metabolism
GPCRs
ion channels

#### **Assay Design**

viability reporter
morphology reporter
conformation reporter
enzyme reporter
membrane potential reporter
binding reporter
inducible reporter

#### **Readout Type**

single multiplexed multiparametric

#### **Cell Format**

cell free
cell lines
primary cells
complex cultures
free embryos

#### **Species**

human
rat
mouse
zebrafish
sheep
boar
rabbit
cattle
guinea pig

#### **Tissue Source**

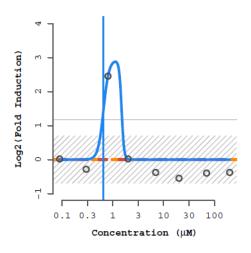
Lung Breast Liver Vascular Skin Kidney Cervix Testis Uterus Brain Intestinal Spleen Bladder Ovary Prostate **Pancreas** Inflammatory Bone

#### **Detection Technology**

qNPA and ELISA
Fluorescence & Luminescence
Alamar Blue Reduction
Arrayscan / Microscopy
Reporter gene activation
Spectrophotometry
Radioactivity
HPLC and HPEC
TR-FRET



## Efforts to Ensure HTS Data Quality and Increase Transparency



FLAGS:
Only one conc above baseline, active
Borderline active

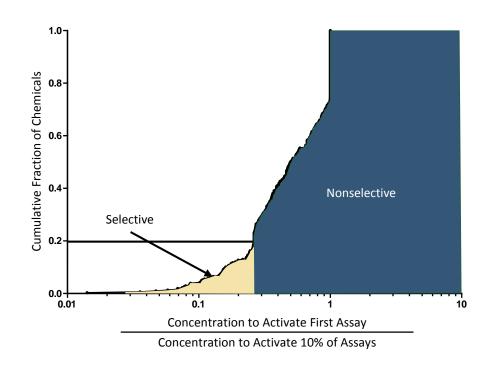
- Public release of Tox21 and ToxCast data on PubChem and EPA web site (raw and processed data)
- Transparent ToxCast data analysis pipeline
  - Data quality flags to indicate concerns with chemical purity and identity, noisy data, and systematic assay errors
  - Publicly available as an R package
- Tox21 and ToxCast chemical libraries have undergone analytical QC and results publicly available
- Public posting of ToxCast procedures
  - Chemical Procurement and QC
  - Data Analysis
  - Assay Characteristics and Performance
- External audit on ToxCast data and data analysis pipeline



## Most Environmental Chemicals are Nonselective for Biological Targets

### **ToxCast** ~600 Cell & ~1,000 biochemical Chemicals | assays Response

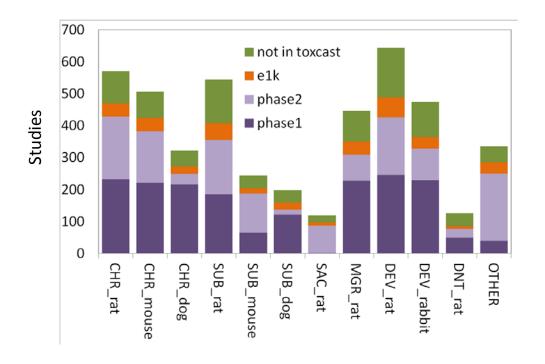
Concentration





### ToxRefDB: Digitizing Legacy in vivo Toxicology Data

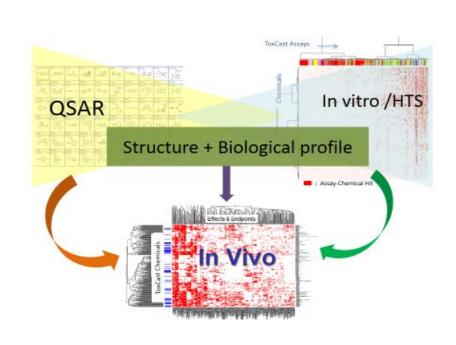
- ToxRefDB holds in vivo endpoint data from animal toxicology studies (DERs, NTP, open literature, pharma)
- Currently at 5567 studies on 1049 unique chemicals

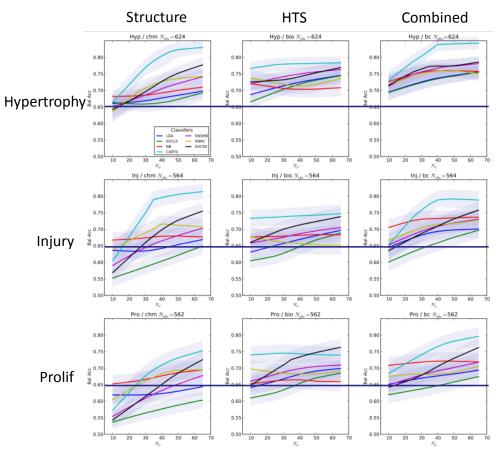


Data Source	Study Count	
EPA OPP DER	3279	
Open Literature	731	
National Toxicol Program	666	
Sanofi_Pharma	222	
Unpublished Submissions	50	
GSK Pharma	38	
Health Canada PMRA DER	23	



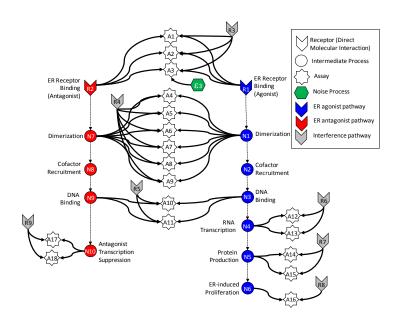
## Predicting Target Organ Toxicities by Machine Learning







## Developing a Pathway Model to Predict Endocrine Activity



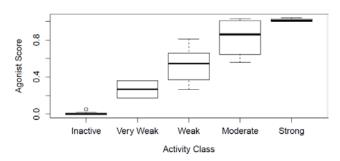
Performance for *In Vitro* Reference Chemicals

# True Pos	28
# True Neg	12
# False Pos	0
# False Neg	4
PPV	1.0
NPV	0.75
ВА	0.94
Sensitivity	0.88
Specificity	1.0

Performance for *In Vivo* Uterotrophic Studies

# True Pos	28
# True Neg	12
# False Pos	1
# False Neg	1
PPV	0.97
NPV	0.92
ВА	0.95
Sensitivity	0.97
Specificity	0.92

ER Pathway Model Integrating 18 In Vitro Assays



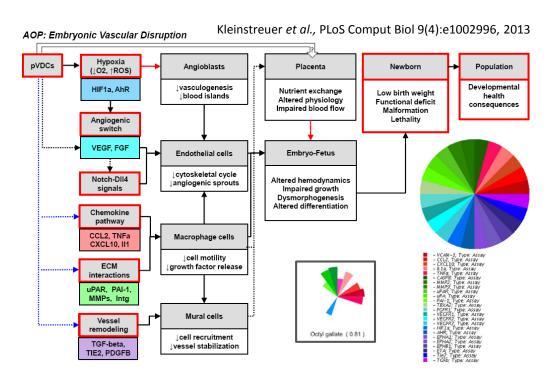
Judson et al., ToxSci (in press)

#### **ER Active Hit Rate**

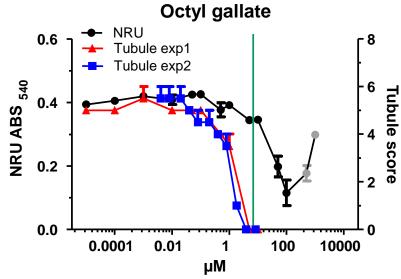
1431 EDSP chemicals run *in vitro*71 (5%) have a significant ER score



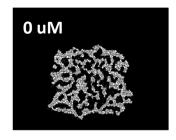
## Systems Biology Models To Scale Targets to Pathways and Networks – Virtual Tissues

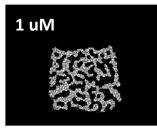


### Human Tubulogenesis Assay (FICAM: T Heinonin)



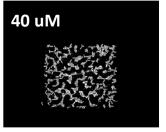
#### **Cell ABM of Octyl Gallate (NCCT: G Nagaraj)**





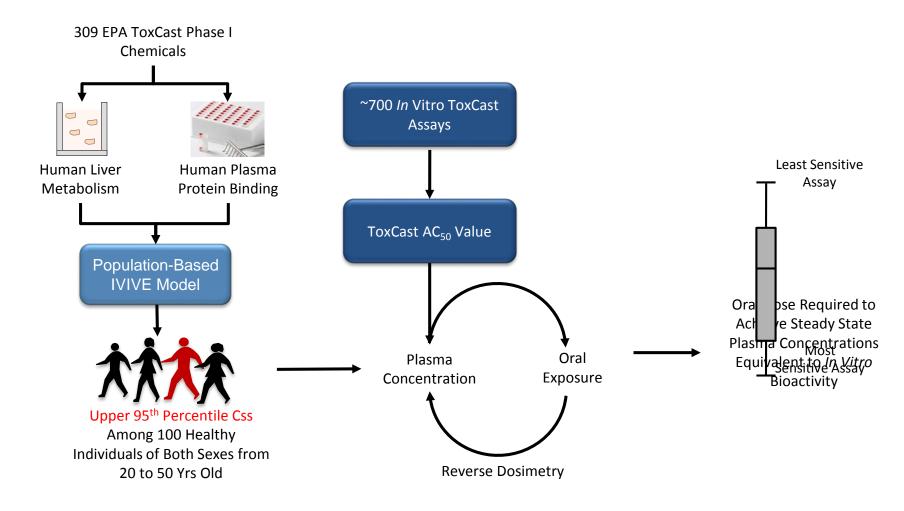






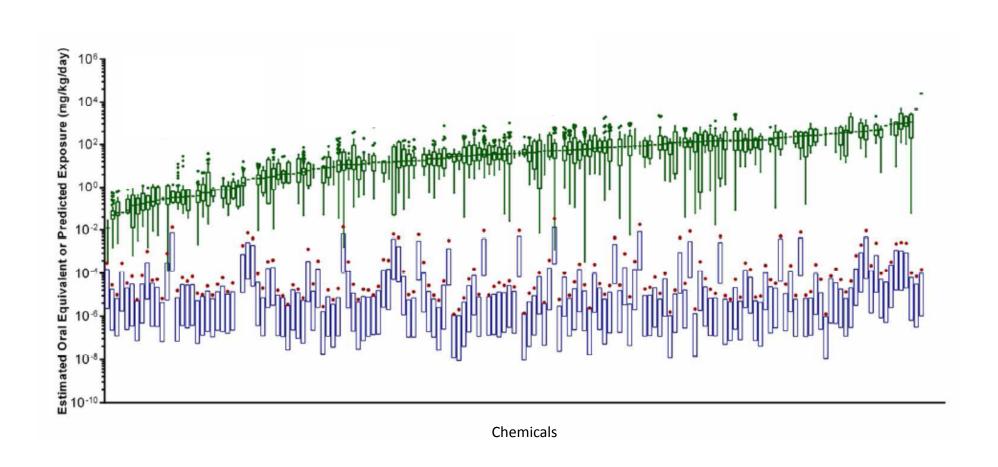


## Providing Context by Incorporating Toxicokinetics and Exposure



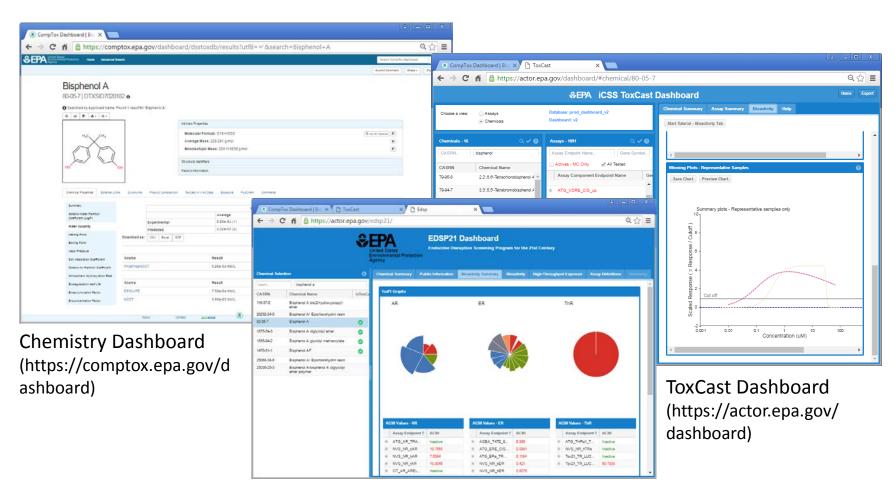


## Comparing Bioactivity with Exposure Predictions for Risk Context





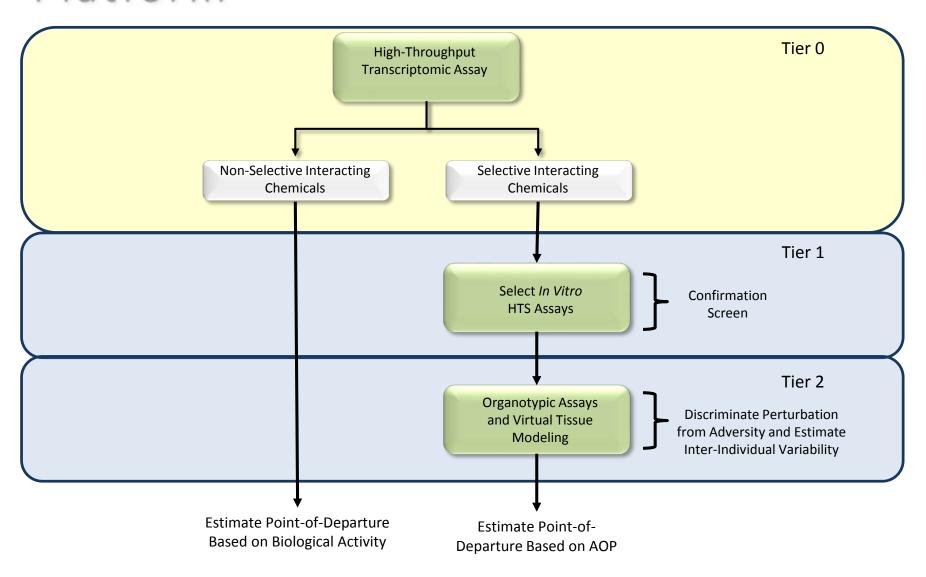
## Delivering Data to Stakeholders and Scientific Community



EDSP21 Dashboard (https://actor.epa.gov/edsp1)



## Developing a Broad Hazard Screening Platform

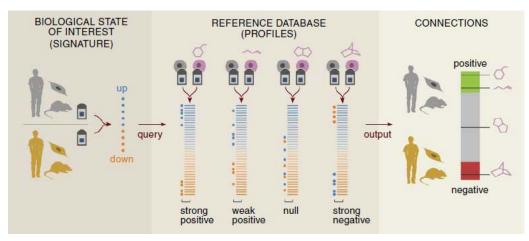




# How Would a HT Transcriptomic Platform be Deployed?

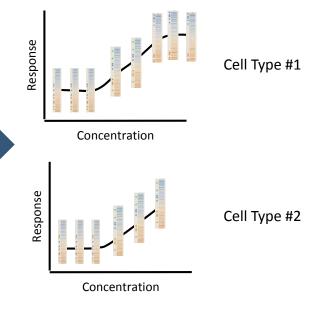
High-Throughput Transcriptomic Assay Tier 0

- Identify predominant mechanisms as a function of concentration
- Group chemicals by similar mechanism/bioactivity
- Identify a concentration that results in no transcriptional effects



Lamb et al. Science (2006)

Broad CMAPdb: 7,000 profiles; 1,309 compounds NIH LINCs CMAPdb: 9,000 shRNAs, 3,000 over expression ORFs, and 4,000 compounds in 20 cell types/lines (cell lines and primary cells)



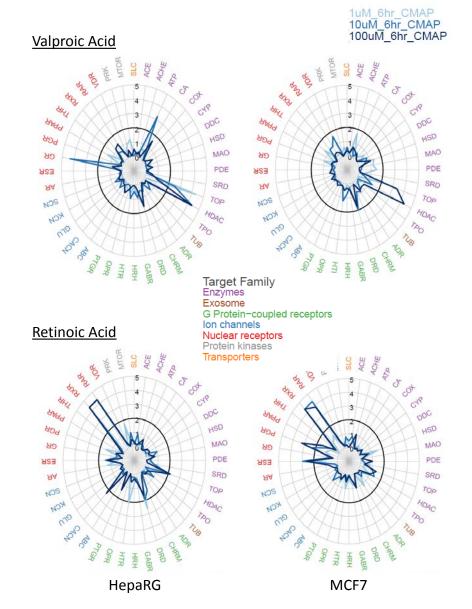
Cell Type #3



### Using HT Transcriptomics to Identify Mode-of-Action

Target Family	Total Profiles	Target Genes	Chemicals	Cell Lines
Cytokine receptors	3	1	1	3
Enzymes	336	40	112	5
Exosome	14	1	4	4
G protein-coupled receptors	585	16	192	4
Ion channels	194	8	65	3
Nuclear receptors	227	10	71	5
Protein kinases	19	8	6	4
Transporters	102	2	35	3

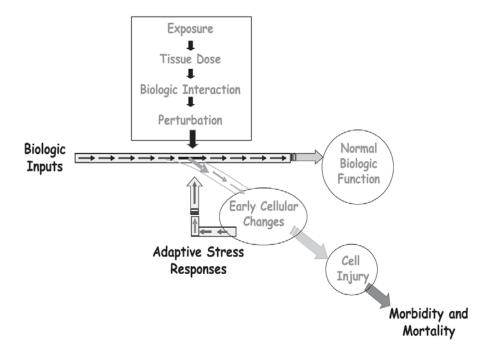
- Developed local database of Broad's CMAP data (~3,000 profiles)
- Annotated targets using KEGG (1,571 profiles)
- Significant genes identified using a z-score cutoff of 2
- Incorporated "JG" scoring method (Jiang and Gentleman 2007)
- Determine significance using a permuted rank approach across target family





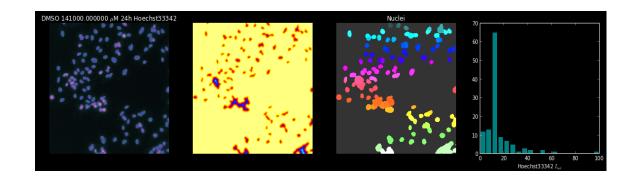
### Distinguishing Adaptation from Adversity

☐ **Tipping Point**: Threshold between adaptation and adversity



☐ Can we use **Tipping Point** to define a point of departure (PoD) for risk assessment?

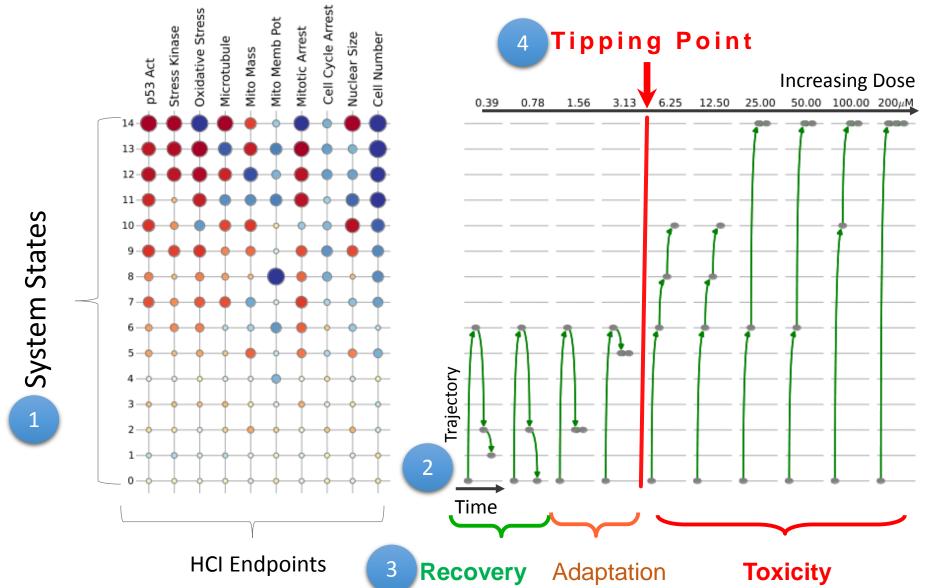
☐ Use ToxCast High Content Imaging (HCI) data to identify Tipping Points



- 967 chemicals (ToxCast)
- HepG2 cells culture
- 10 concentrations
- 3 Time points
- 10 HCI Assays
- 400 plates
- . 100,000 wells
- . 2,400,000 images



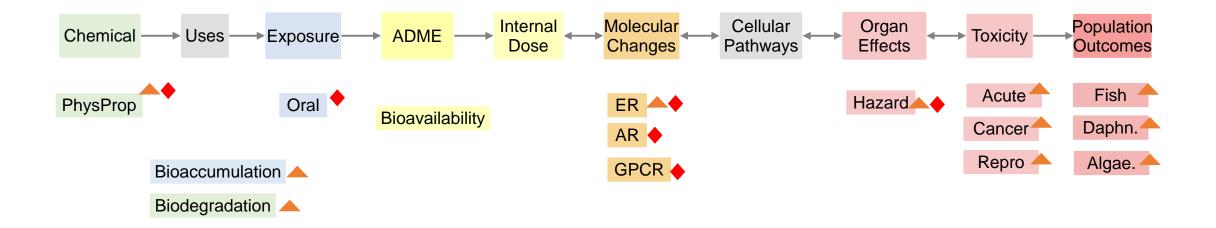
### Tipping Point Analysis





#### Thousands of chemicals have limited data!

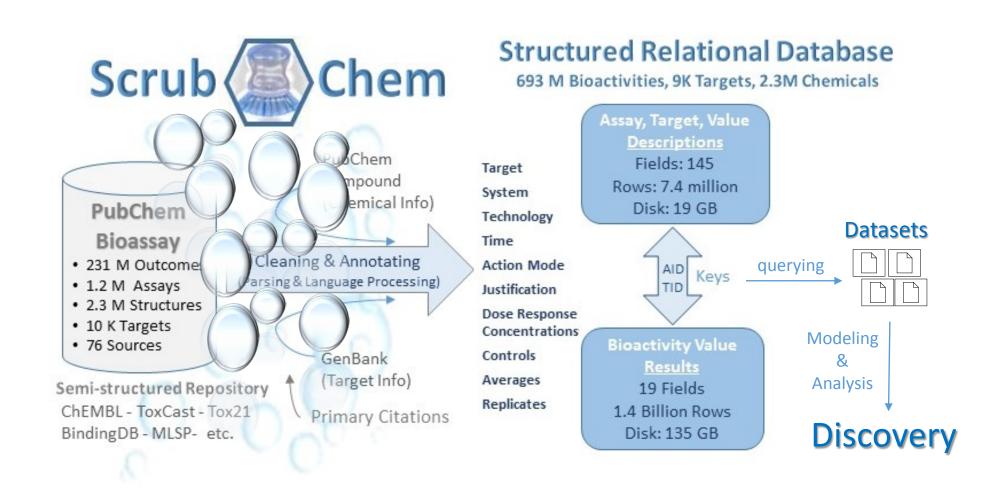
- Tens of thousands of environmental chemicals have very limited exposure/biological data
  - → Need more effective tools to describe chemical properties, effects and linkages
  - → Need predictive models to fill data gaps



- existing/legacy tools: TIMES, LeadScope, ECOSAR, EPIWIN
- developed internally: Read-across/GenRA, Machine Learning: classification and regression



### Structuring PubChem Data for Analysis



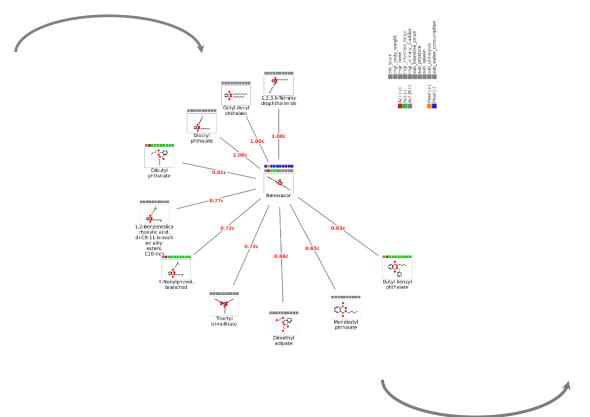


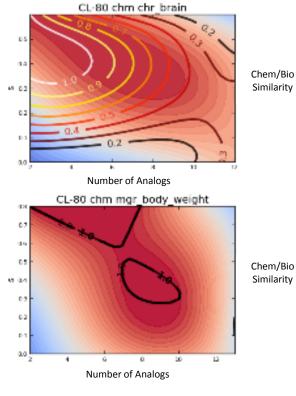
#### Generalised Read-across (GenRA)

Chemicals are clustered on the basis of chemical descriptors to identify local neighbourhoods

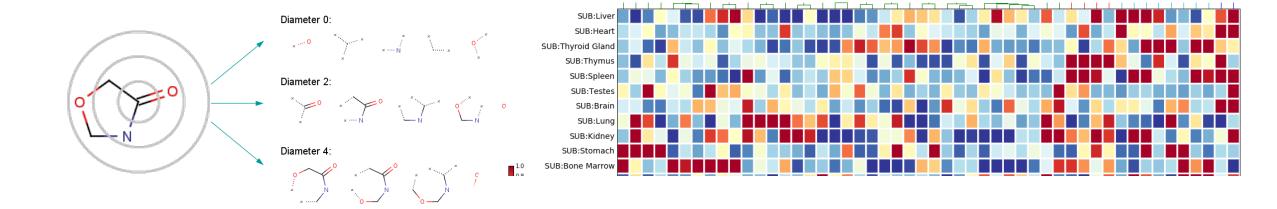
The Read-across toxicity prediction is a similarity-weighted activity of nearest neighbours based on chemistry and bioactivity descriptors

Uncertainties can be evaluated across the local neighbourhoods



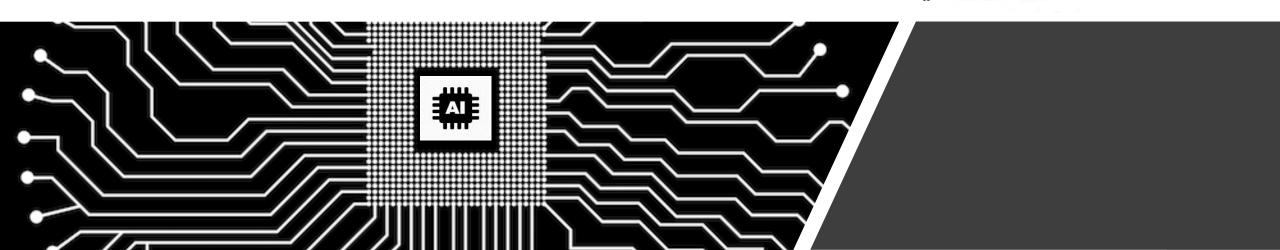


Shah, et al. RTP 2016



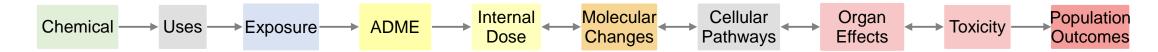
# Machine Learning to Predict Chemical Effects



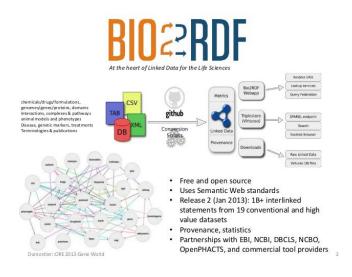


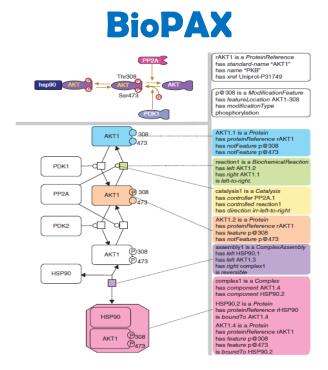


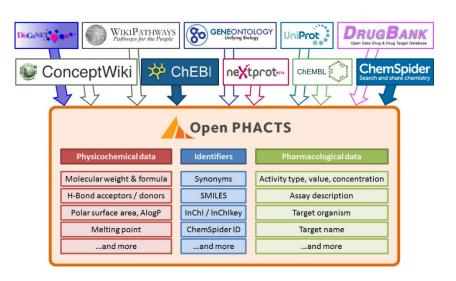
### Use Semantic Tools to Link Disparate Data



Can we use semantic tools (OWL/RDF, existing ontologies to meaningfully integrated disparate resources?









#### Data Challenges

- Transparently sharing complex data streams—adequately capturing chemical (treatment dose and time), biological (experimental modal, assay, etc.) context to ease re-use
- Systematically integrating disparate data streams—representing linkages across molecular, cellular, tissue, organs. This is vital for relating early molecular changes to adverse (e.g. histopathological) outcomes
- Effectively extracting evidence from unstructured textual data— the literature is one of the largest resource for information about apical outcomes
- Using linked data to better discriminate between adaptation vs adversity predicting which molecular markers lead to apical outcomes
- Quantifying and incorporating uncertainty and variability in predictions
- Legal defensibility of new methods and assessment products



### Acknowledgements and Questions

Tox21 Colleagues:

NTP Crew FDA Collaborators NCATS Collaborators

EPA Colleagues:

NERL NHEERL NCEA



**EPA's National Center for Computational Toxicology**