

US EPA ARCHIVE DOCUMENT



# The ToxCast™ Program – predicting hazard, characterizing toxicity pathways and prioritizing the toxicity testing of environmental chemicals



Office of Research and Development  
National Center for Computational Toxicology

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# The Need For a New Approach

The image displays a collage of overlapping web browser windows from the EPA website. The windows show various pages including:

- EPA/OPP High Production Volume (HPV) Challenge Program: The HPV Voluntary Challenge Chemical List
- EPA/OSCP: Endocrine Disruptor Screening Program
- Drinking Water Contaminant Candidate List (CCL)
- EPA: Pesticides - Inert (other) Pesticide Ingredients in Pesticide Products
- EUROPA - European Commission - Enterprise & Industry - REACH

The REACH page is the most prominent, showing the title "THE NEW EU CHEMICALS LEGISLATION - REACH" and a detailed overview of the regulation. The EPA logo is visible in the top left of the browser windows.

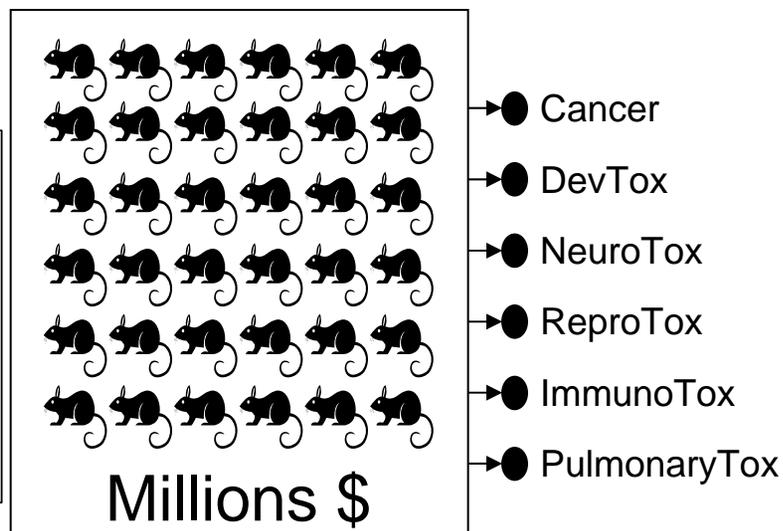
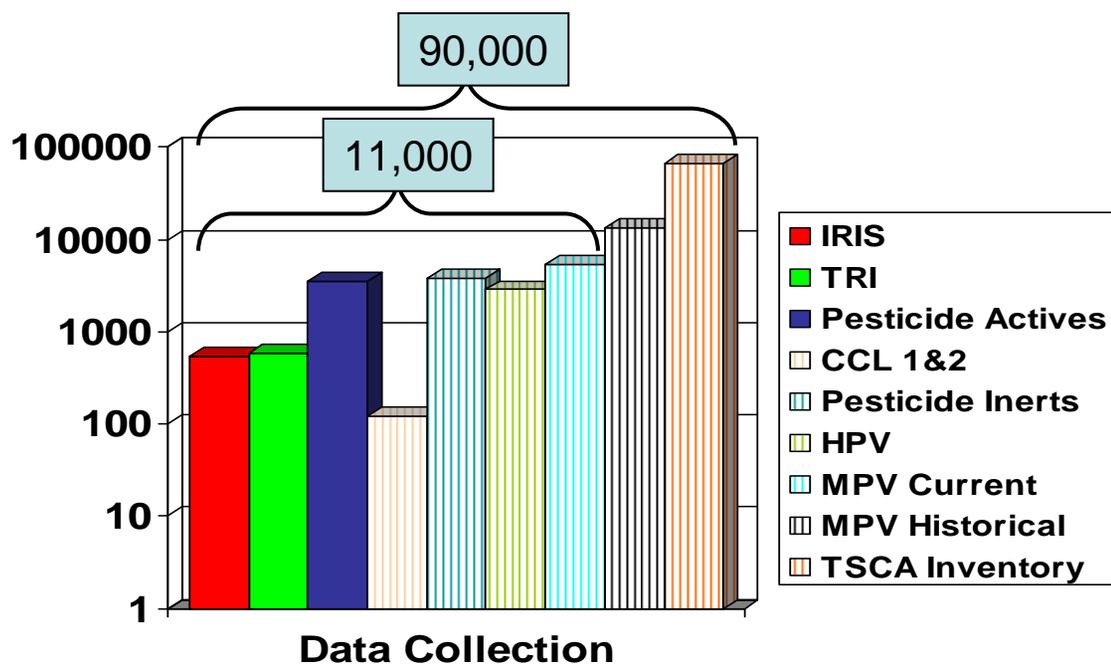
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# EPA's Need for Chemical Prioritization

## Too Many Chemicals

## Too High a Cost



...and not enough data.

## Ways to Prioritize:

- Animal studies
  - cost, time, ethical considerations
- QSAR
  - domain of applicability, availability of models
- Bioactivity Profiling
  - biologically relevant chemical characterization
  - HTS methods
  - ToxCast



# ToxCast™ : a computational toxicology approach based on high-throughput bioactivity profiling

- Research program of EPA's National Center for Computational Toxicology
- Addresses chemical screening and prioritization needs for pesticidal inerts, anti-microbials, CCLs, HPVs and MPVs
- Comprehensive use of HTS technologies to generate biological fingerprints and predictive signatures
- Coordinated with NIH: NTP and NHGRI/NCGC via Tox21
- Committed to stakeholder involvement and public release of data
  - Communities of Practice- Chemical Prioritization; Exposure
  - NCCT website- <http://www.epa.gov/ncct/toxcast>
  - ACToR- Aggregated Computational Toxicology Resource  
<http://actor.epa.gov/actor/>



# Phased Development of ToxCast

Phase	Number of Chemicals	Chemical Criteria	Purpose	Number of Assays	Cost per Chemical	Target Date
I	320	Data Rich (pesticides)	Signature Development	552	\$20k	FY08
Ib	15	Nanomaterials	Pilot	166	\$10K	FY09
Ila	>300	Data Rich Chemicals	Validation	>400	~\$20-25k	FY09
Ilb	>100	Known Human Toxicants	Extrapolation	>400	~\$20-25k	FY09
Ilc	>300	Expanded Structure and Use Diversity	Extension	>400	~\$20-25k	FY10
Ild	>12	Nanomaterials	PMN	>200	~\$15-20K	FY09-10
III	Thousands	Data poor	Prediction and Prioritization	>300	~\$15-20k	FY11-12



# ToxCast\_320 Phase I Chemicals

309 unique structures  
 3 triplicates, 5 duplicates for QC  
 8 metabolites

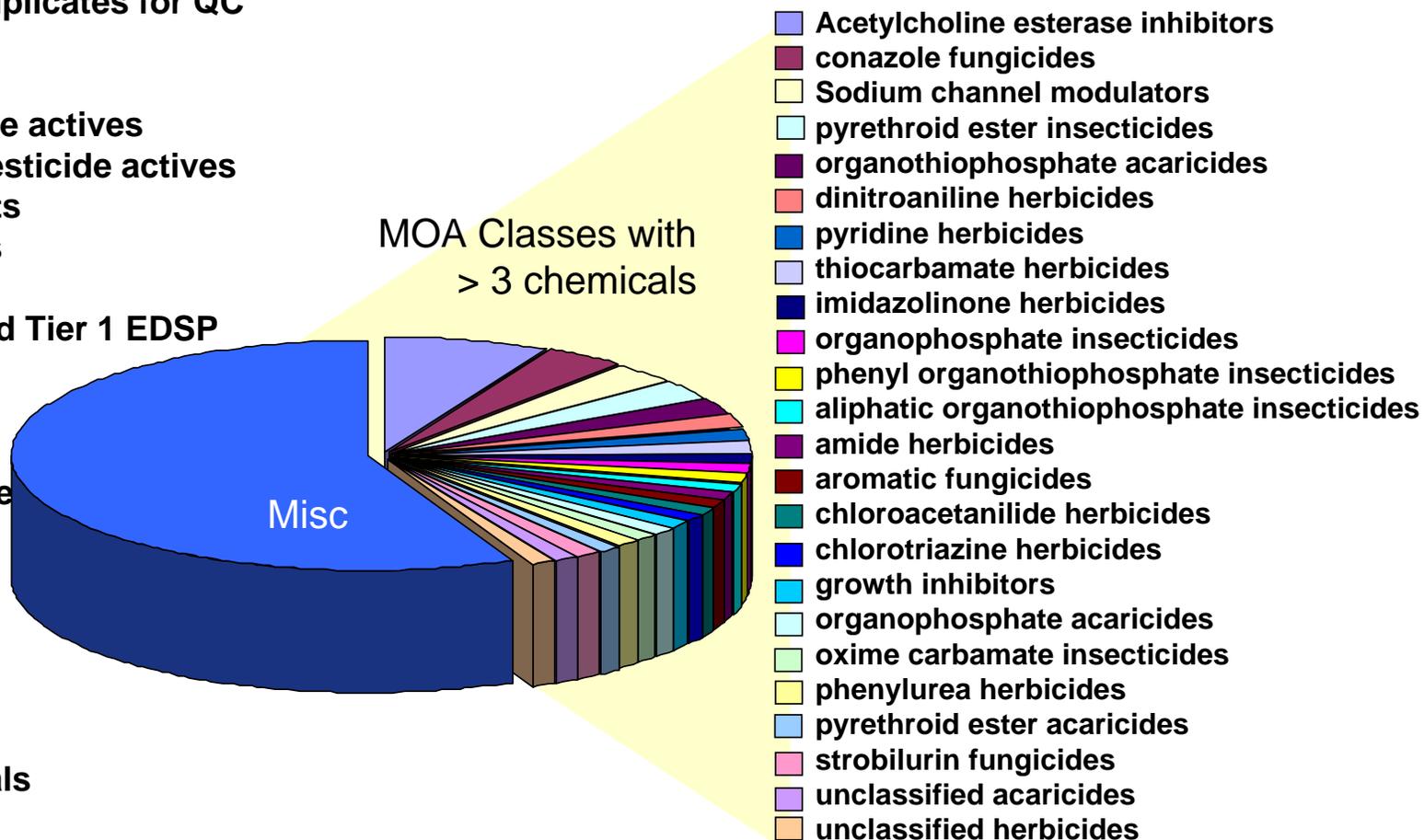
291 total pesticide actives  
 273 registered pesticide actives  
 22 pesticide inerts  
 33 antimicrobials

56 of 73 proposed Tier 1 EDSP

23 IUR  
 13 HPV  
 11 HPV Challenge

73 OW PCCL  
 11 CCL1  
 10 CCL2  
 25 CCL3

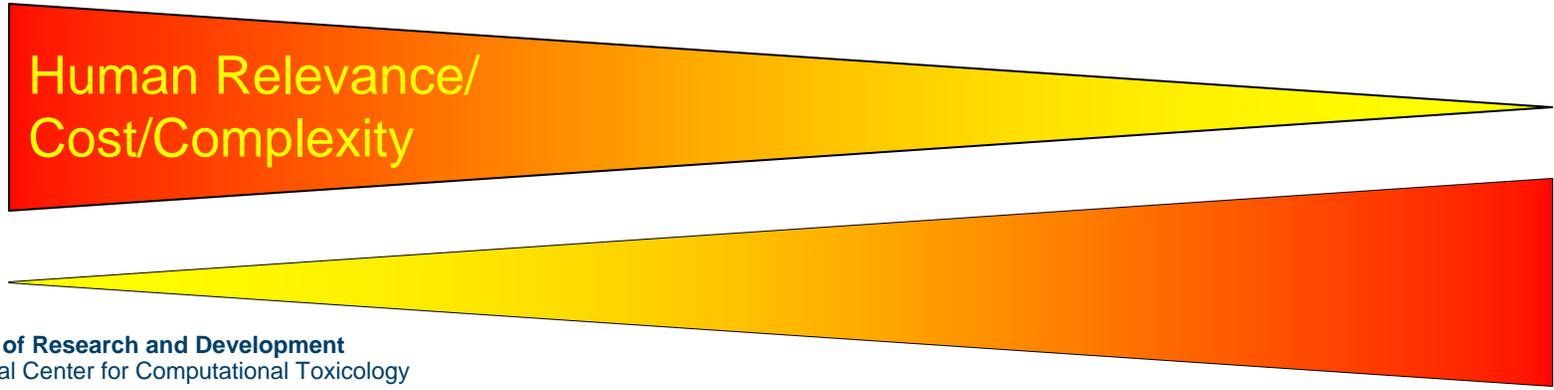
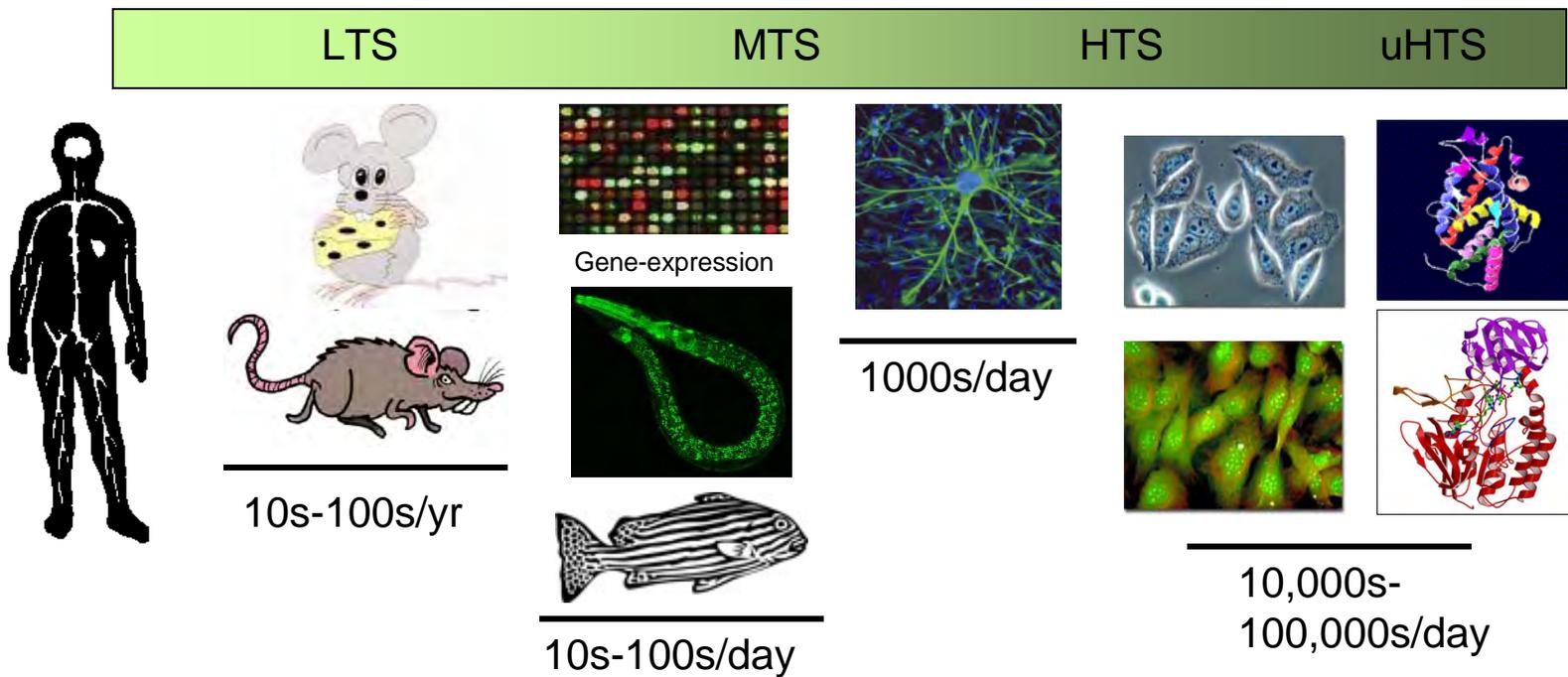
122 IRIS chemicals





# High-Throughput Screening Assays

*batch testing of chemicals for pharmacological/toxicological endpoints using automated liquid handling, detectors, and data acquisition*



# ToxCast Phase I Datasets

20 Assay sources  
554 Endpoints

- **ToxCast 1.0 (April, 2007)**
  - Enzyme inhibition/receptor binding HTS (Novascreen)
  - NR/transcription factors (Attagene, NCGC)
  - Cellular impedance (ACEA)
  - Complex cell interactions (BioSeek)
  - Hepatocellular HCS (Cellumen)
  - Hepatic, renal and airway cytotoxicity (IVAL)
  - In vitro hepatogenomics (IVAL, Expression Analysis)
  - Zebrafish developmental toxicity (Phylonix)
- **ToxCast 1.1 (January, 2008)**
  - Neurite outgrowth HCS (NHEERL)
  - Cell proliferation (NHEERL)
  - Zebrafish developmental toxicity (NHEERL)
- **ToxCast 1.2 (June, 2008)**
  - XME Gene Regulation (CellzDirect)
  - HTS Genotoxicity (Gentronix)
  - Organ toxicity; dosimetry (Hamner Institutes)
  - Toxicity and signaling pathways (Invitrogen)
  - C. elegans WormTox (NIEHS)
  - Gene markers from microscale cultured hepatocytes (MIT)
  - 3D Cellular Zebrafish vascular/cardiotoxicity (Zygogen)
  - microarray with metabolism (Solidus)
  - HTS stress response (NHEERL+NCGC)

# ToxCast Assays

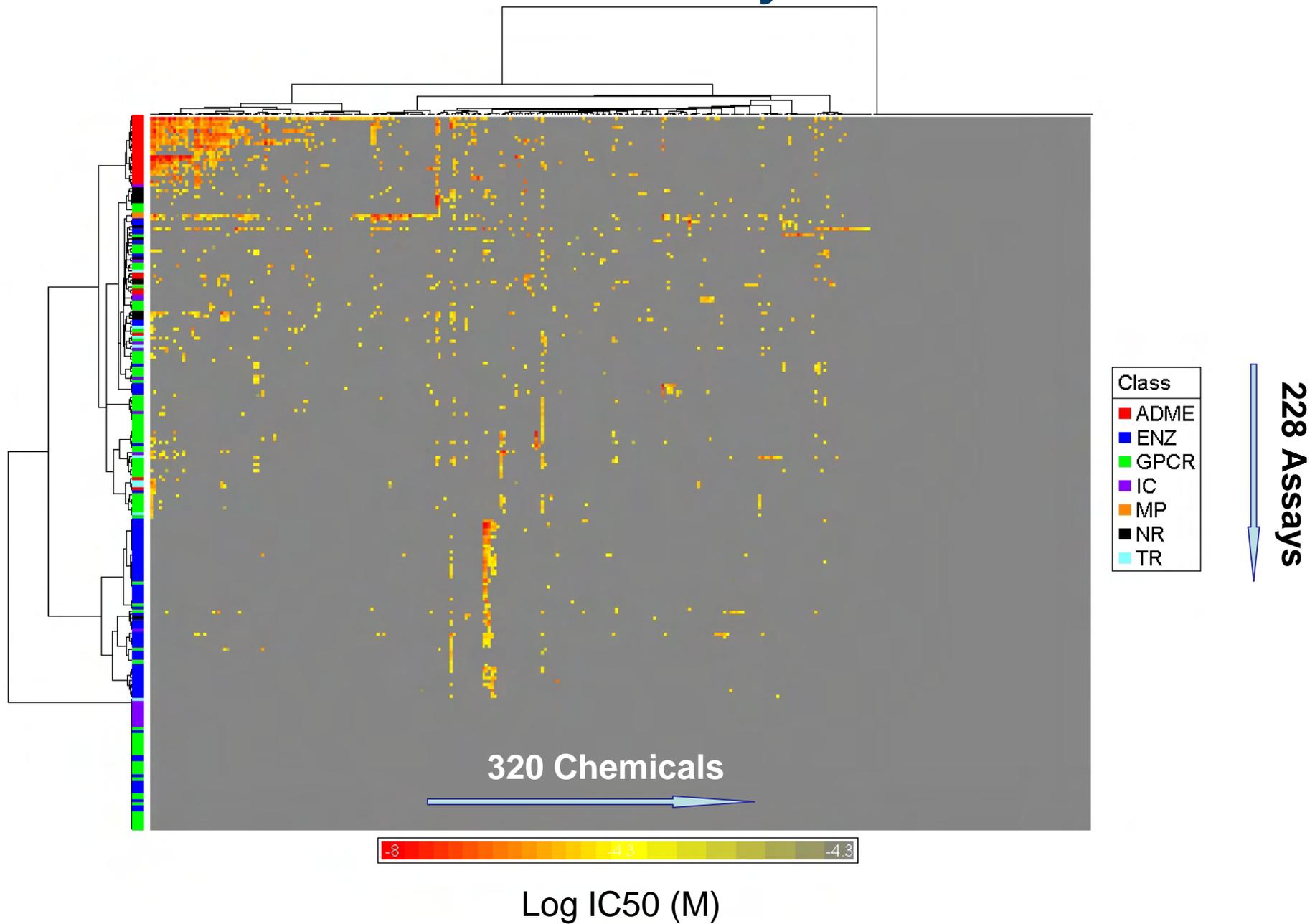
## Biochemical Assays

- Protein families
  - GPCR
  - NR
  - Kinase
  - Phosphatase
  - Protease
  - Other enzyme
  - Ion channel
  - Transporter
- Assay formats
  - Radioligand binding
  - Enzyme activity
  - Co-activator recruitment

## Cellular Assays

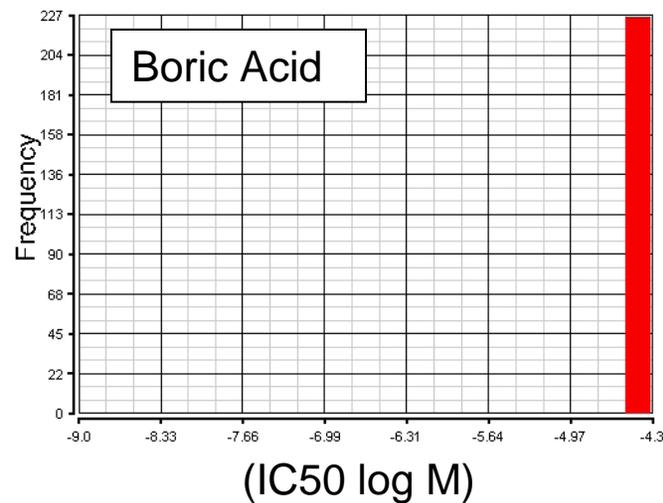
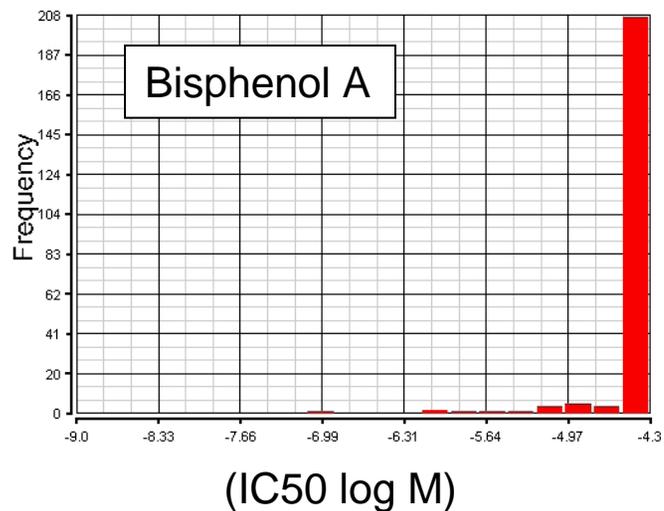
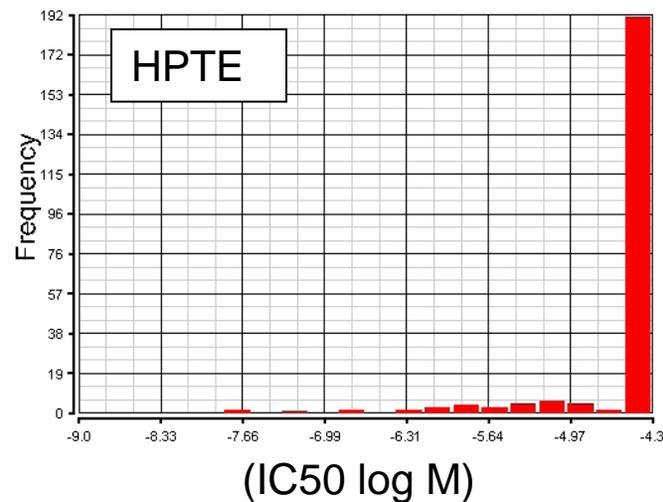
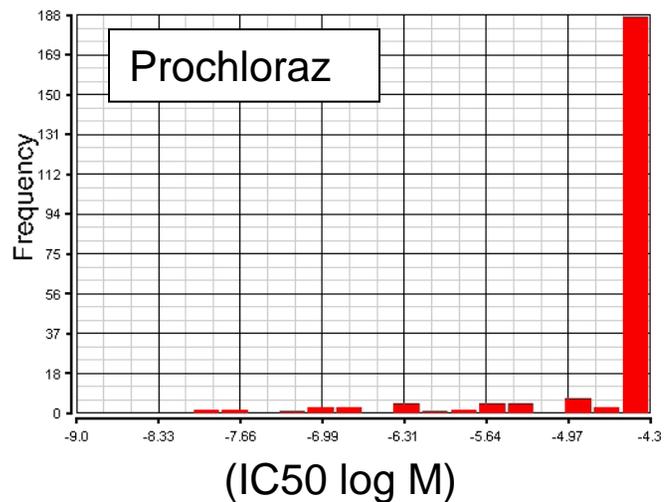
- Cell lines
  - HepG2 human hepatoblastoma
  - A549 human lung carcinoma
  - HEK 293 human embryonic kidney
- Primary cells
  - Human endothelial cells
  - Human monocytes
  - Human keratinocytes
  - Human fibroblasts
  - Human proximal tubule kidney cells
  - Human small airway epithelial cells
- Biotransformation competent cells
  - Primary rat hepatocytes
  - Primary human hepatocytes
- Assay formats
  - Cytotoxicity
  - Reporter gene
  - Gene expression
  - Biomarker production
  - High-content imaging for cellular phenotype

# Biochemical Assay Results

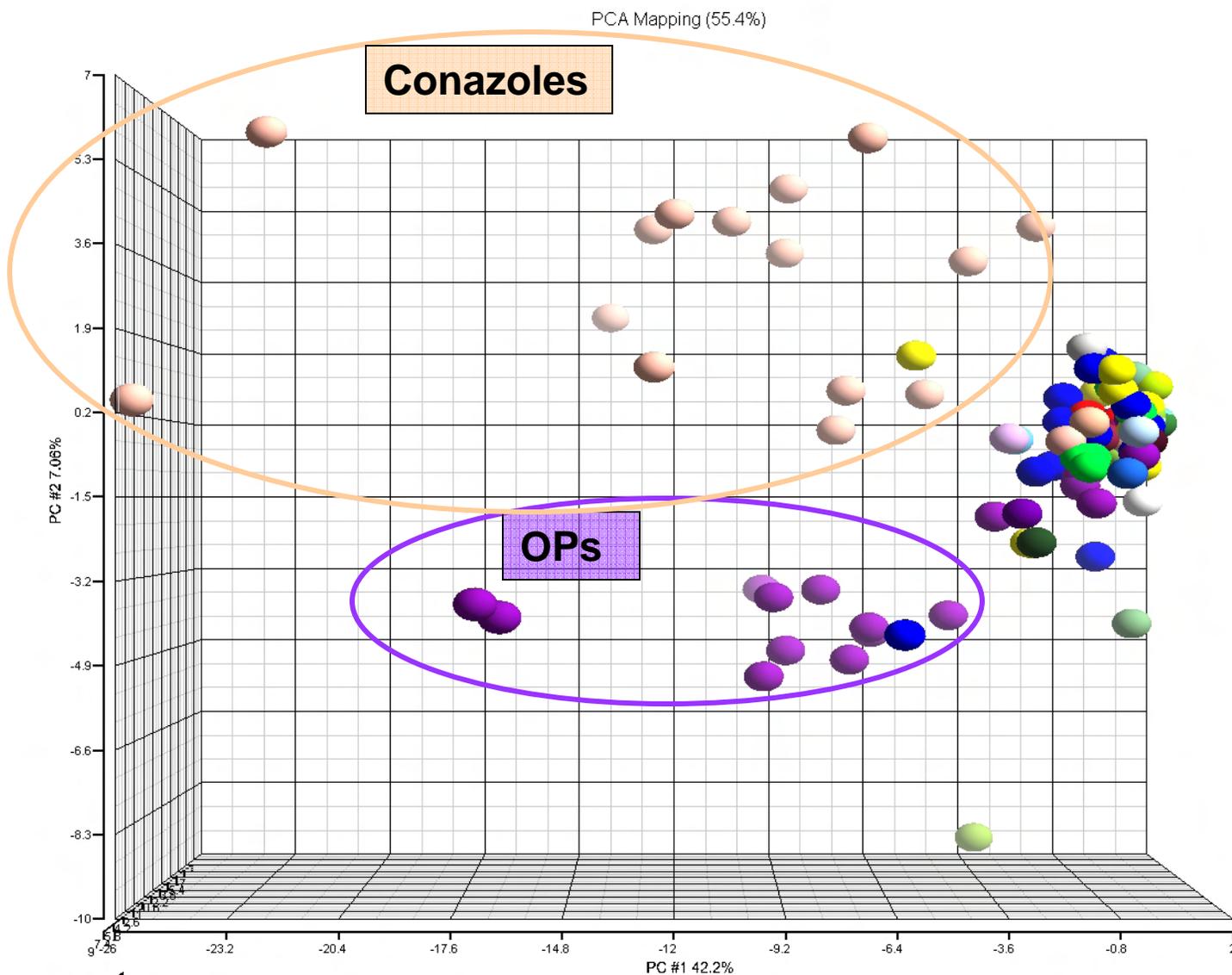




# Examples of Chemical Activity in Biochemical Assays

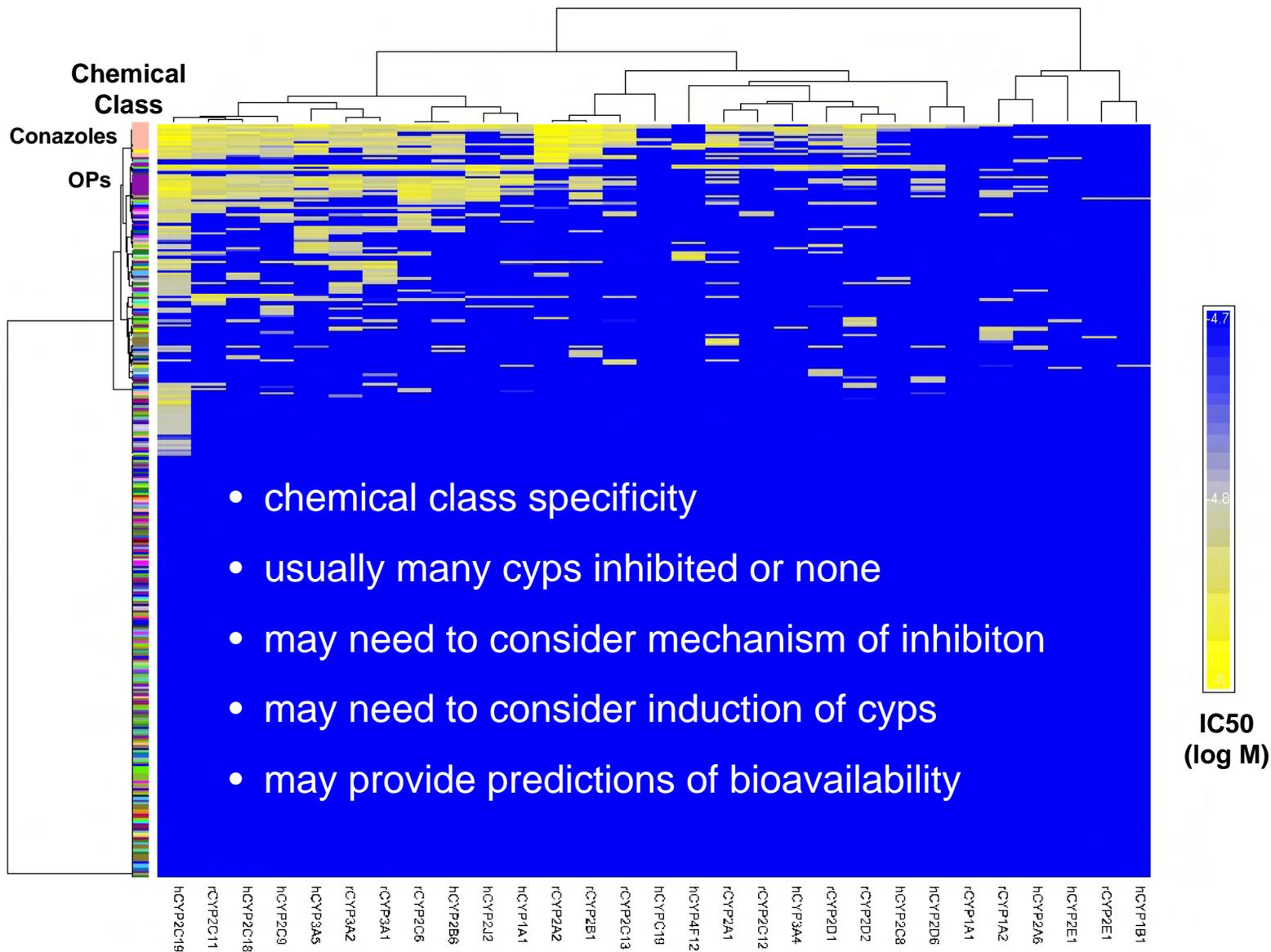


# PCA Mapping of CYP Inhibition



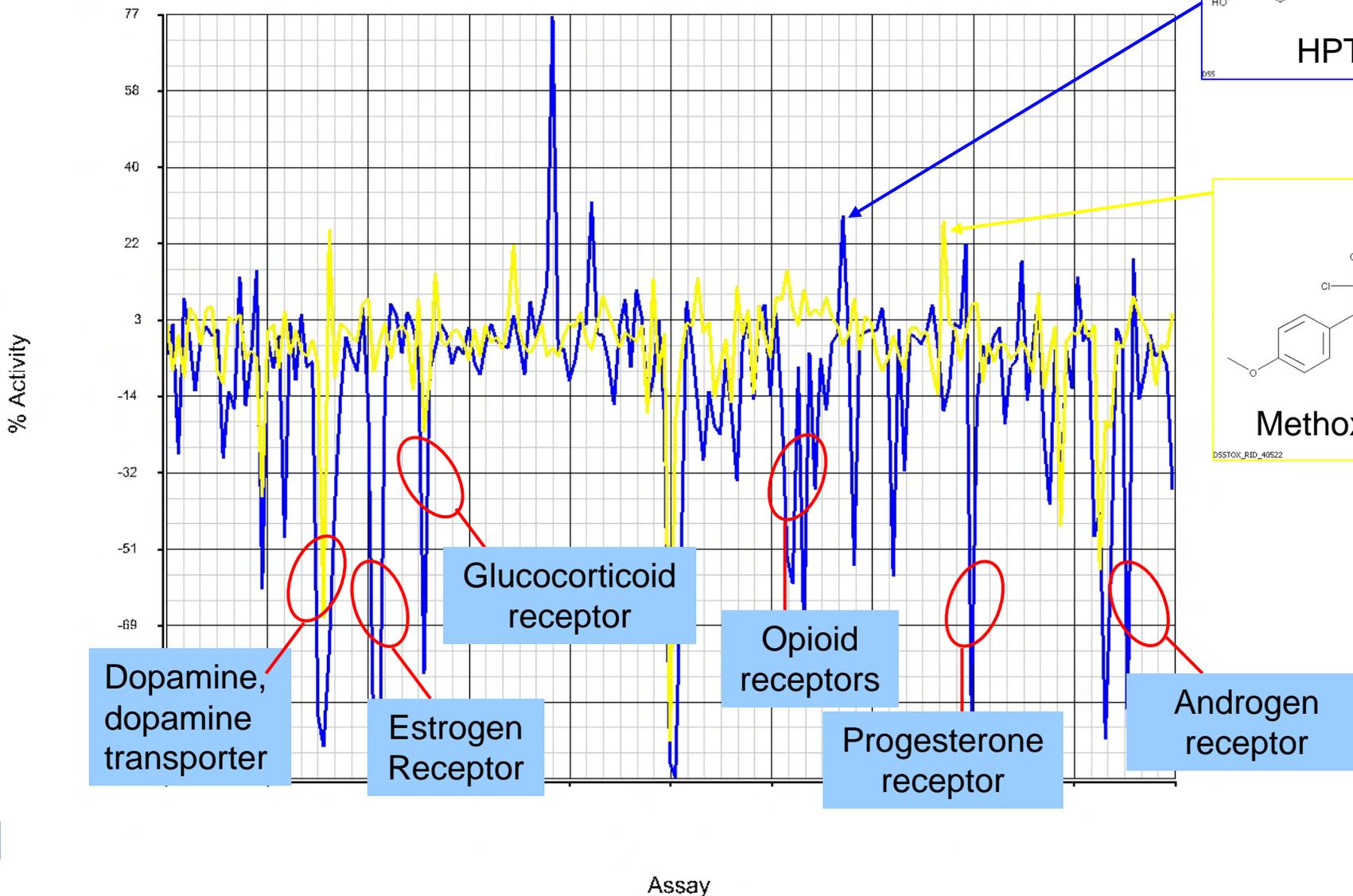
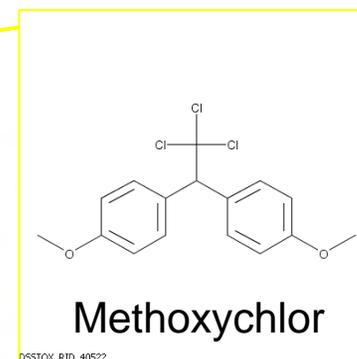
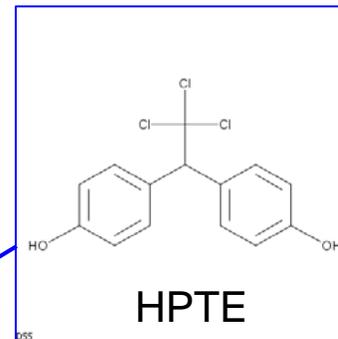


# CYP Inhibition





# ToxCast and Biotransformation



# Cellular Assays

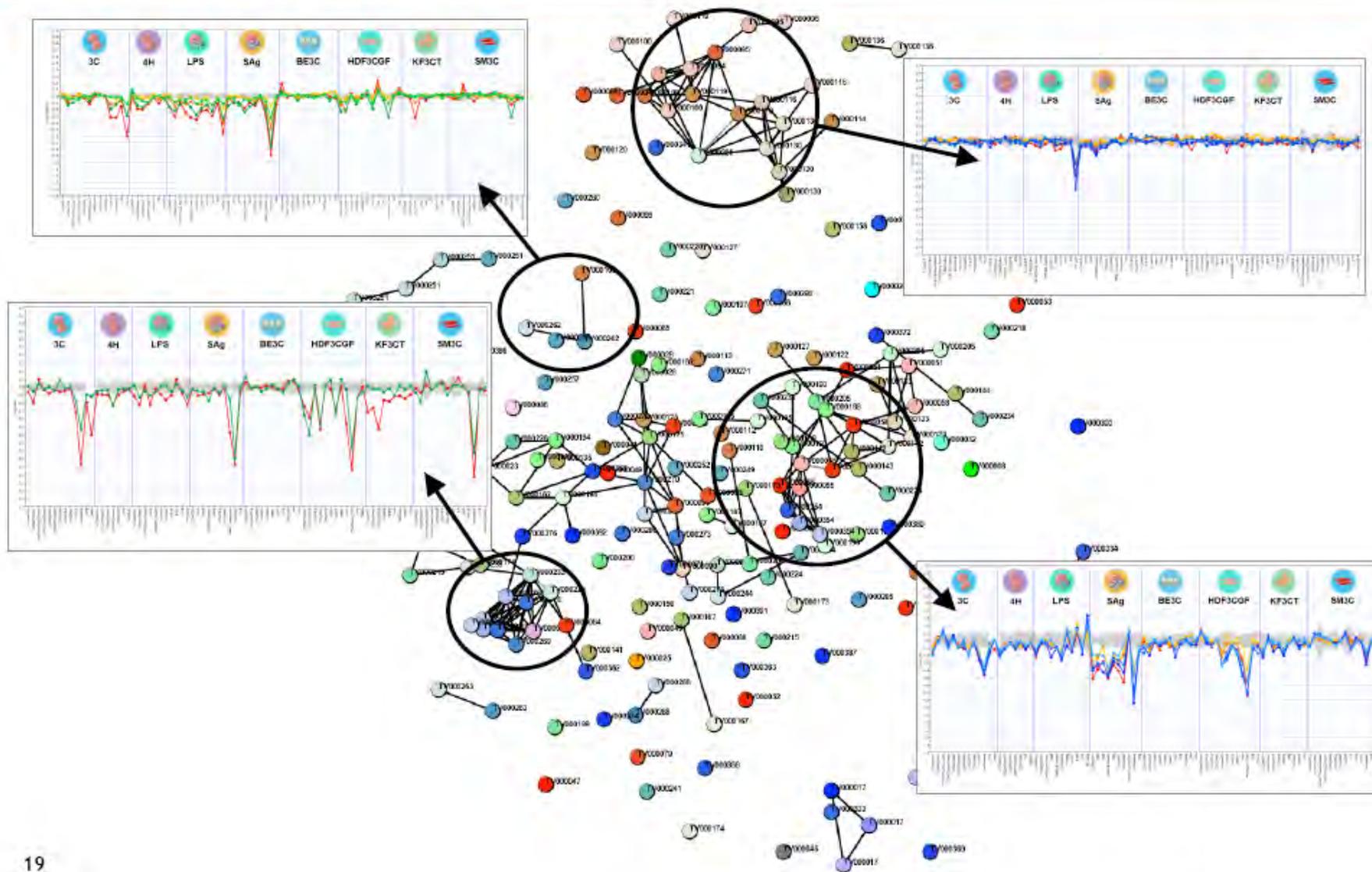
- Types of Assays
  - Known toxicity pathways and targets
    - biomarker measurements
    - reporter gene assays
  - General cytotoxicity
  - Toxicity cellular phenotypes
- Cell lines and primary cells
- Generally screened at up to 100  $\mu\text{M}$  or used maximally tolerated concentration defined by general cytotoxicity determination
- Concentration-response format used and  $\text{EC}_{50}$  generated

# Primary Human Cell Systems (BioSeek, Inc.)

System	Cell Types	Environment	Readouts
3C	 Endothelial cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	MCP-1, VCAM-1, ICAM-1, Thrombomodulin, Tissue Factor, E-selectin, uPAR, IL-8, MIG, HLA-DR, Prolif., Vis., SRB (13)
4H	 Endothelial cells	IL-4+histamine	VEGFR2, P-selectin, VCAM-1, uPAR, Eotaxin-3, MCP-1, SRB (7)
LPS	 Peripheral Blood Mononuclear Cells + Endothelial cells	TLR4	CD40, VCAM-1, Tissue Factor, MCP-1, E-selectin, IL-1a, IL-8, M-CSF, TNF-a, PGE2, SRB (11)
SAg	 Peripheral Blood Mononuclear Cells + Endothelial cells	TCR	MCP-1, CD38, CD40, CD69, E-selectin, IL-8, MIG, PBMC Cytotox, SRB, Proliferation (10)
BE3C	 Bronchial epithelial cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	uPAR, IP-10, MIG, HLA-DR, IL-1a, MMP-1, PAI-1, SRB, TGF-b1, tPA, uPA(11)
HDF3CGF	 Fibroblasts	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$ +bFGF+EGF+PDGF-BB	VCAM-1, IP-10, IL-8, MIG, Collagen III, M-CSF, MMP-1, PAI-1, Proliferation, TIMP-1, EGFR, SRB (12)
KF3CT	 Keratinocytes + Fibroblasts	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$ +TGF- $\beta$	MCP-1, ICAM-1, IP-10, IL-1a, MMP-9, TGF-b1, TIMP-2, uPA, SRB (9)
SM3C	 Vascular smooth muscle cells	IL-1 $\beta$ +TNF- $\alpha$ +IFN- $\gamma$	MCP-1, VCAM-1, Thrombomodulin, Tissue Factor, IL-6, LDLR, SAA, uPAR, IL-8, MIG, HLA-DR, M-CSF, Prolif., SRB (14)

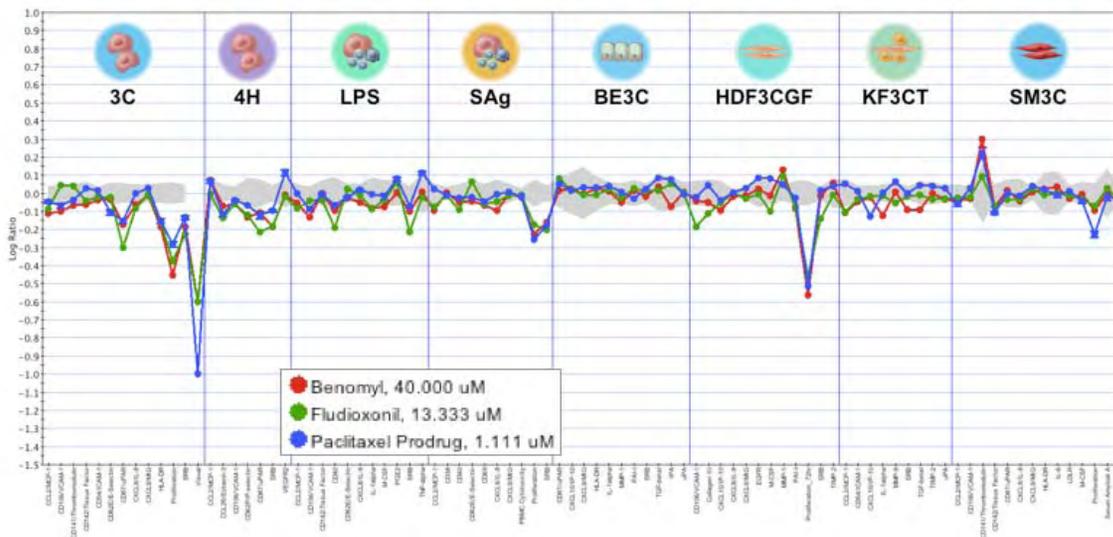
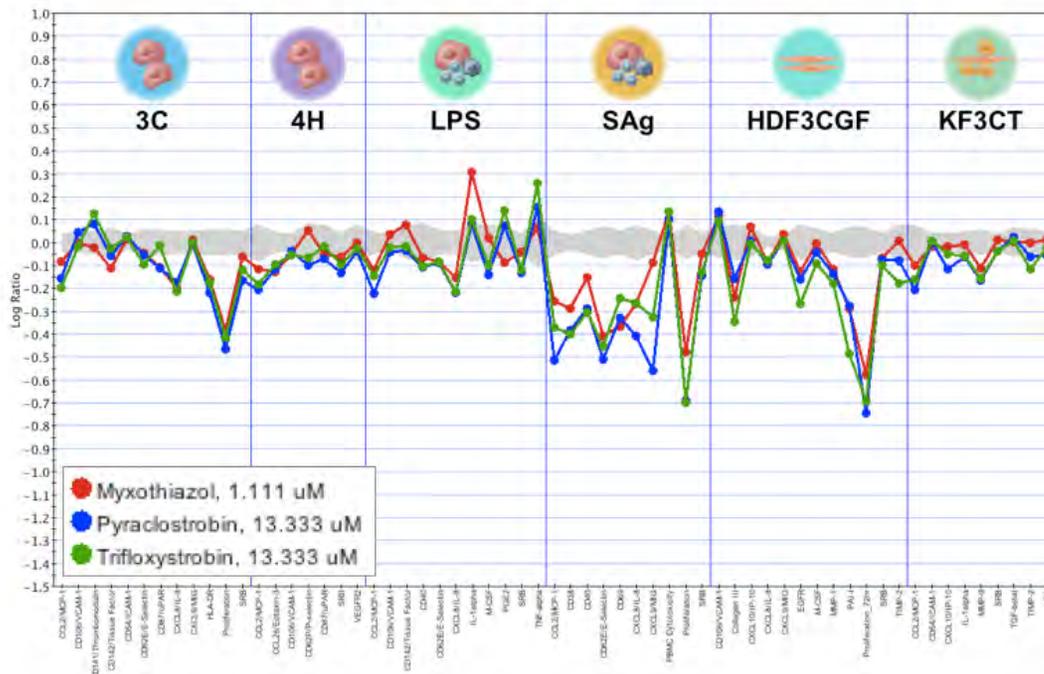
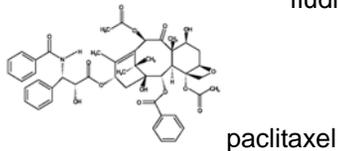
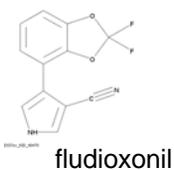
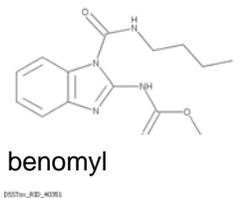
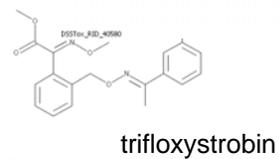
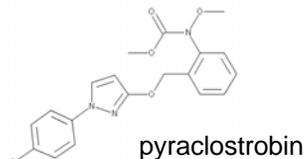
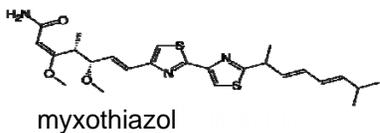
- 8 Assay systems
- 87 endpoints
- 4 concentrations

# Functional Similarity Map of ToxCast Library





# Mitochondrial Dysfunction and Endoplasmic Reticulum Stress Classes



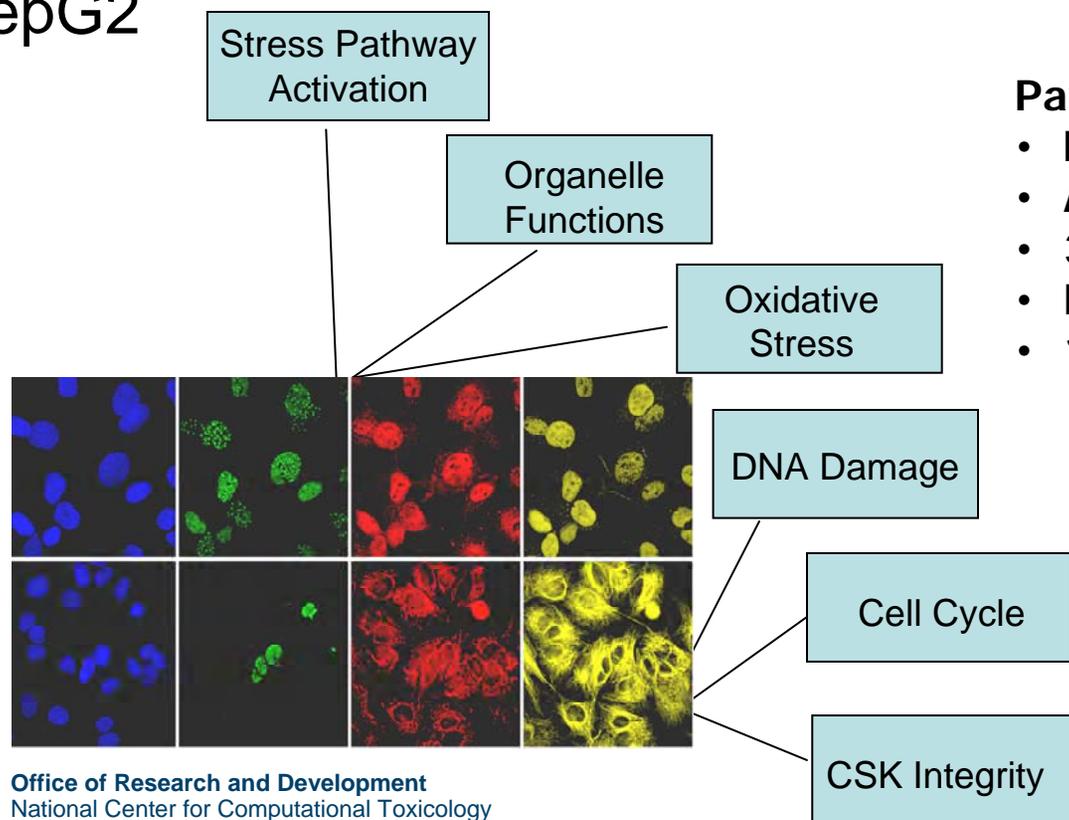
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## Use of BioSeek Data in ToxCast

- Individual assay endpoints become part of larger ToxCast data set for developing predictive models
- BioMAP signatures used to provide mechanistic understanding of potential mechanism/mode of action
- May be able to validate signatures with other phenotypic assays

# High-Content Screening of Cellular Phenotypic Toxicity Parameters (Cellumen, Inc.)

- Technology: automated fluorescent microscopy
- Objective: Determine effects of chemicals on toxicity biomarkers in a cell culture of human liver hepatoma HepG2



## Panel 1 design\*:

- Multiple mechanisms of toxicity
- Acute, early & chronic exposure
- 384-well capacity
- HepG2
- 1<sup>o</sup> rat hepatocytes

# CellCiphr™ Cytotoxicity Panel

- 10-point conc-response (200  $\mu$ M-39 nM)
- Three time points (1 hr, 24 hr, 72 hr)
- 11 endpoints per assay

Biomarker	Positive Control	Z'
Stress Pathway	Anisomycin	.63
Oxidative Stress	Camptothecin	.7
Mitochondrial Function	CCCP	.55
Mitochondrial Mass	CCCP	.35
Cell Loss	Camptothecin	.56
Cell Cycle	Paclitaxel	.54
DNA Degradation	Paclitaxel	.6
Nuclear Size	Paclitaxel	.63
DNA Damage	Camptothecin	.43
Mitotic Arrest	Paclitaxel	.63
Cytoskeletal Integrity	Paclitaxel	.3



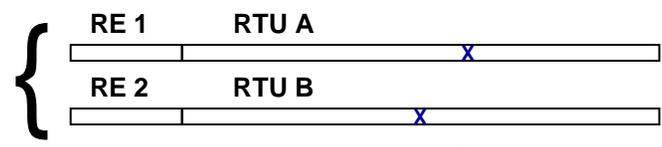
# Multiplexed Reporter Gene Assay (Attogene, Inc.)

- Measures activation/inhibition of transcription factors (TF)
- TF integrate signals arising from changing cellular environments and coordinate cellular response to such change
- Similar to genomics but many fewer TF than genes
- Compounds with similar mechanism of toxicity should bear similar patterns
- Patterns should reflect the changes that precede or accompany the compounds' toxicity
- Use signatures for prediction of toxicological outcomes of compounds

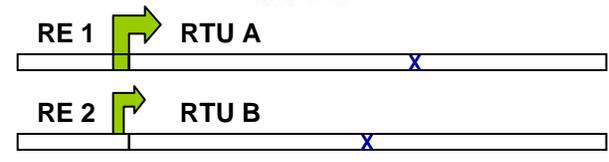


# Multiplexed Reporter Gene Assay

Library of RTUs

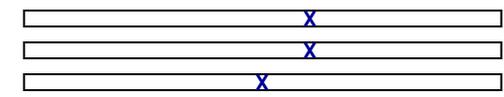


Cell Transfection

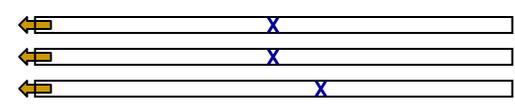


Transcription

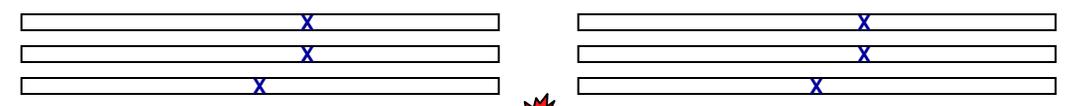
RNA Isolation



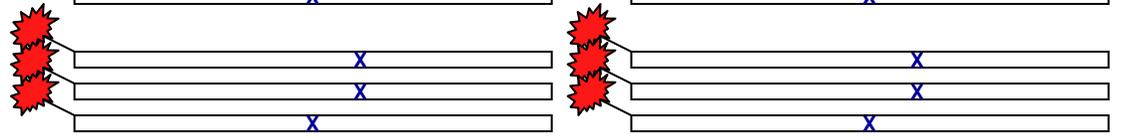
Reverse transcription



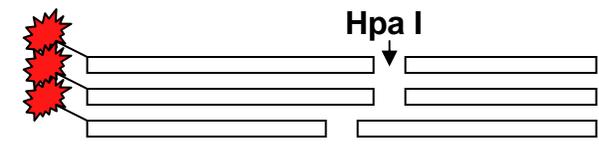
PCR amplification



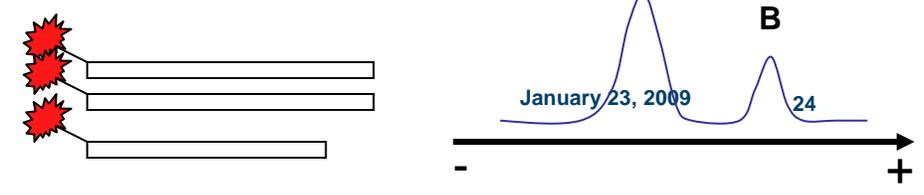
Labeling



Processing (Hpa I)

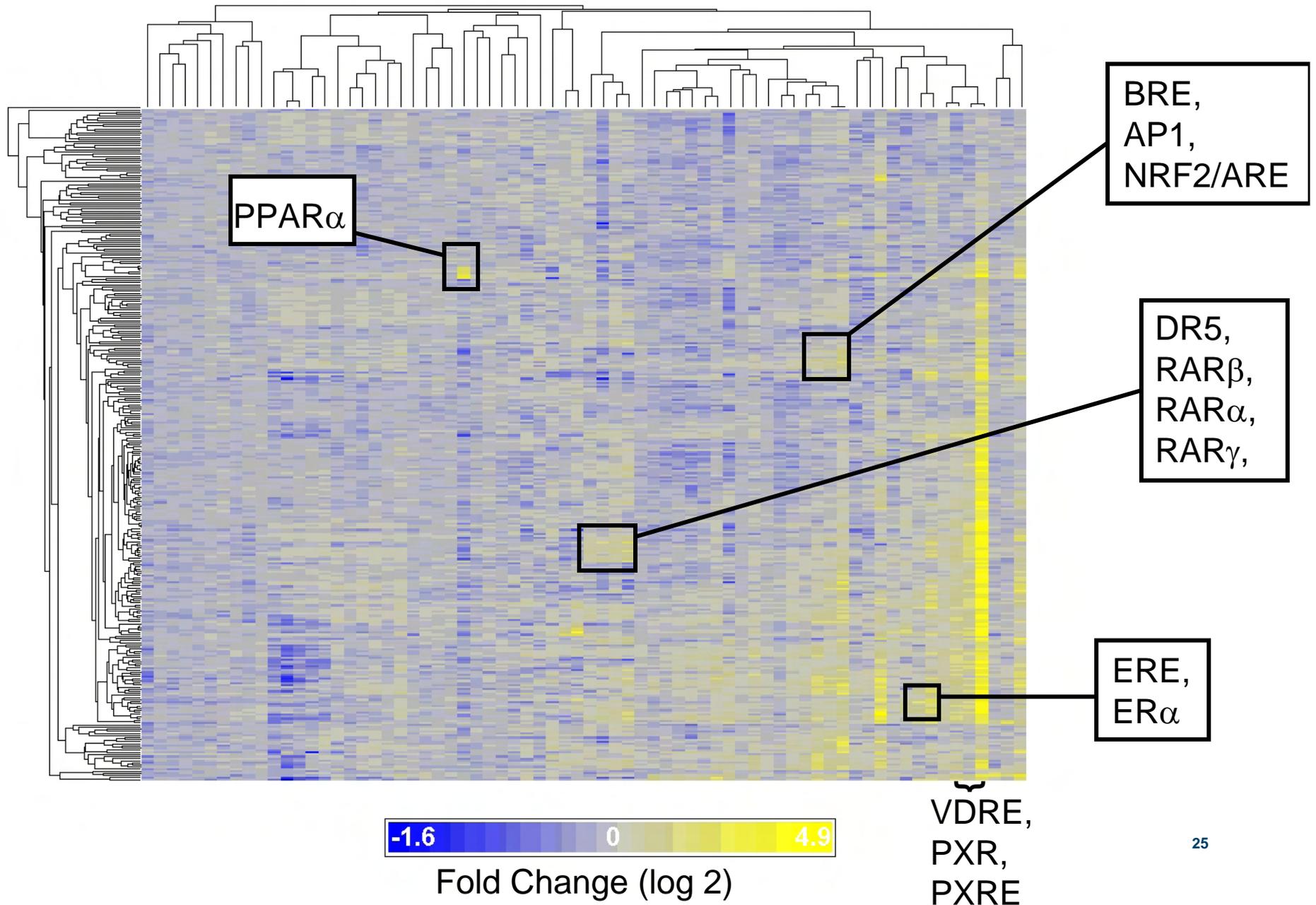


Separation and detection (capillary electrophoresis)



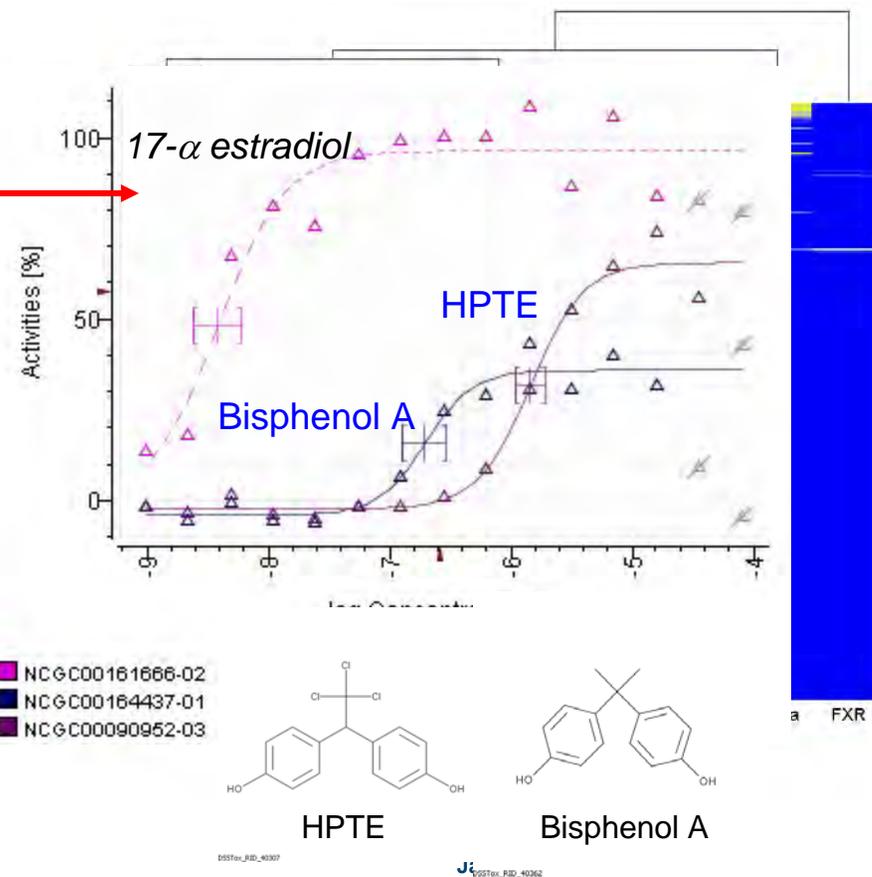
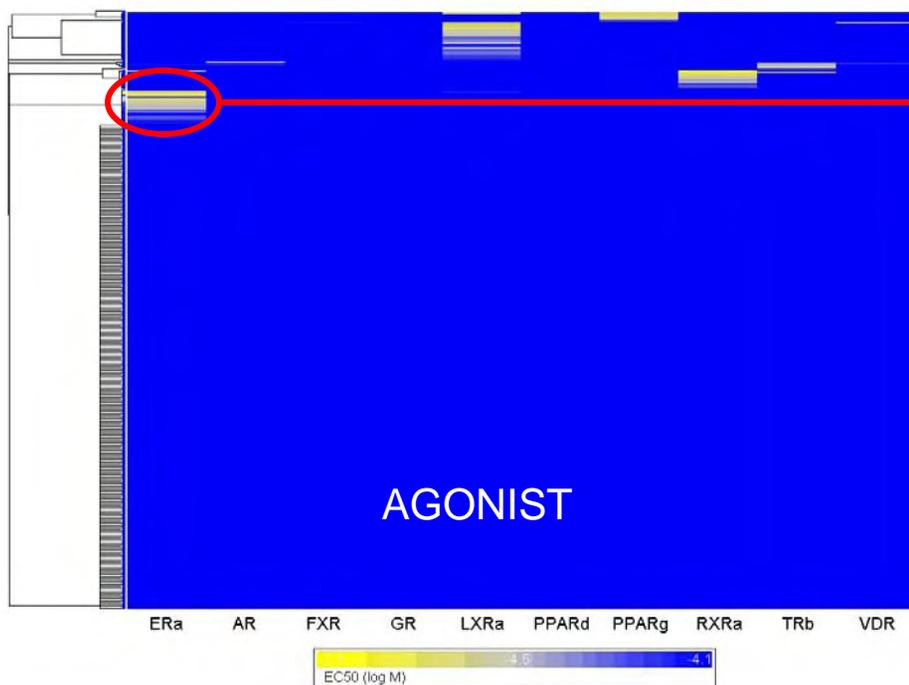


# Hierarchical Cluster Attagene Results

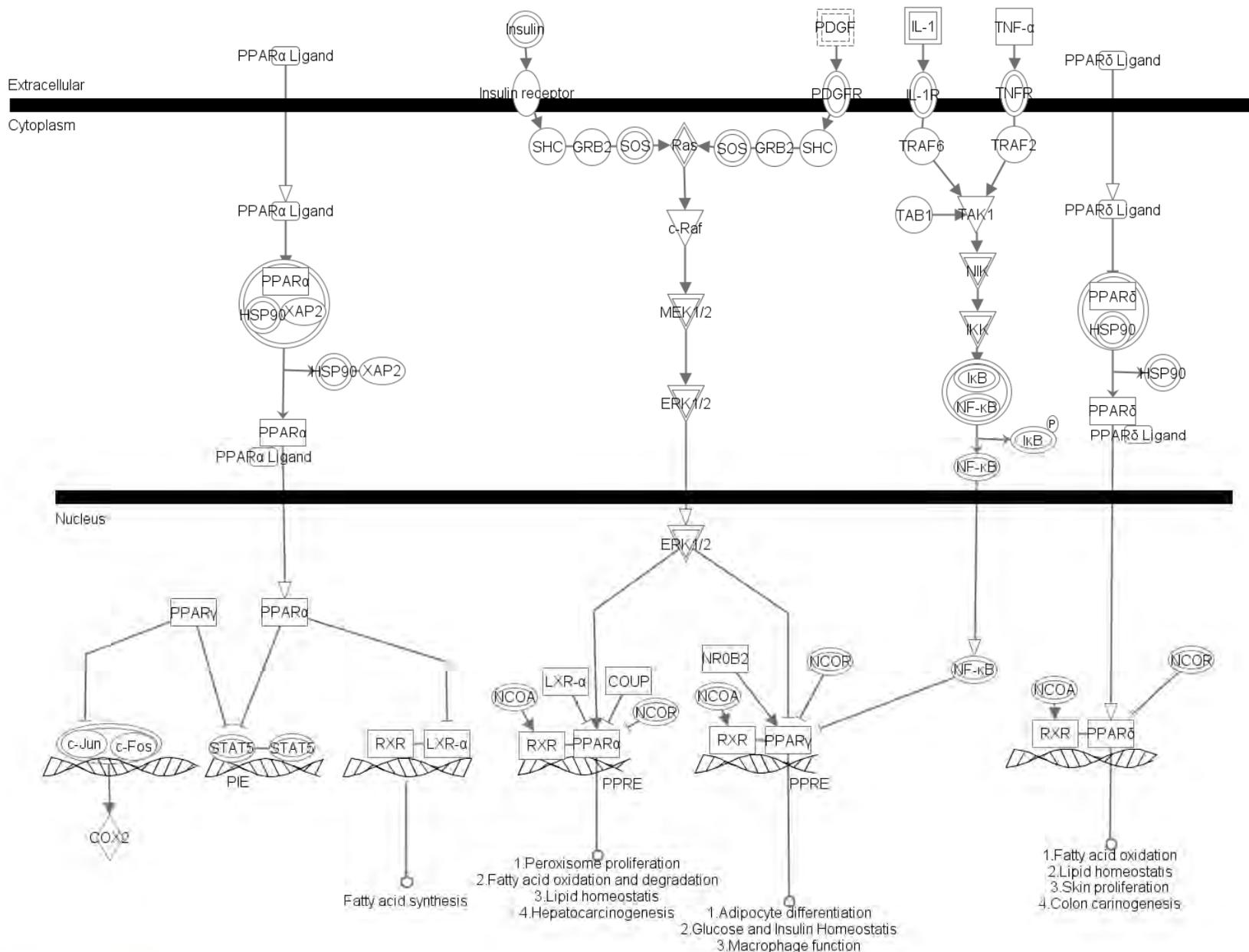


# Nuclear Receptor Screening (NCGC)

- 10 Nuclear Receptors (more in queue)
- Cellular Reporter Assays
- Agonist and Antagonist modes
- Concentration-Response Format (15 conc)



PPAR Signaling



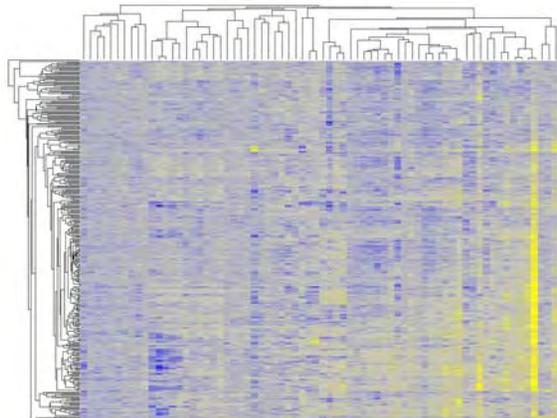
# ToxCast Covers a Wide Swath of Biological Space

## Molecular Pathways Identified by Analyses of ToxCast Assays

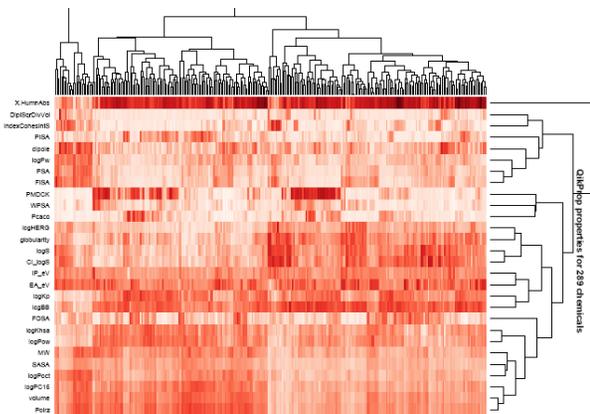
	<b>GeneGO</b>	<b>Ingenuity</b>	<b>David-KEGG</b>	<b>Total GeneID</b>
<b>Human</b>	<b>81</b>	<b>60</b>	<b>42</b>	<b>317</b>
<b>Rat</b>	<b>34</b>	<b>39</b>	<b>9</b>	<b>51</b>



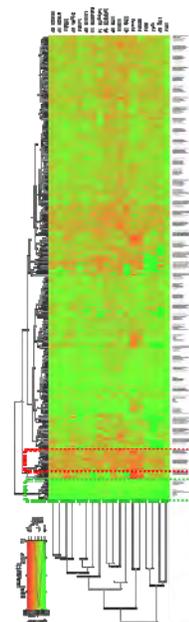
# ToxCast Data Analysis



-1.6 0 4.5

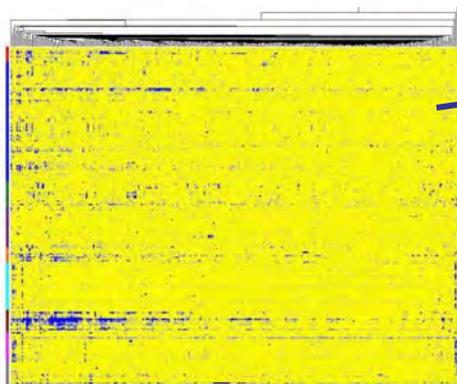


Physical chemical properties

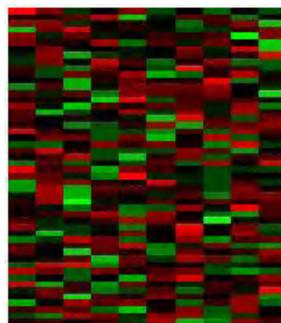


In silico Predictions

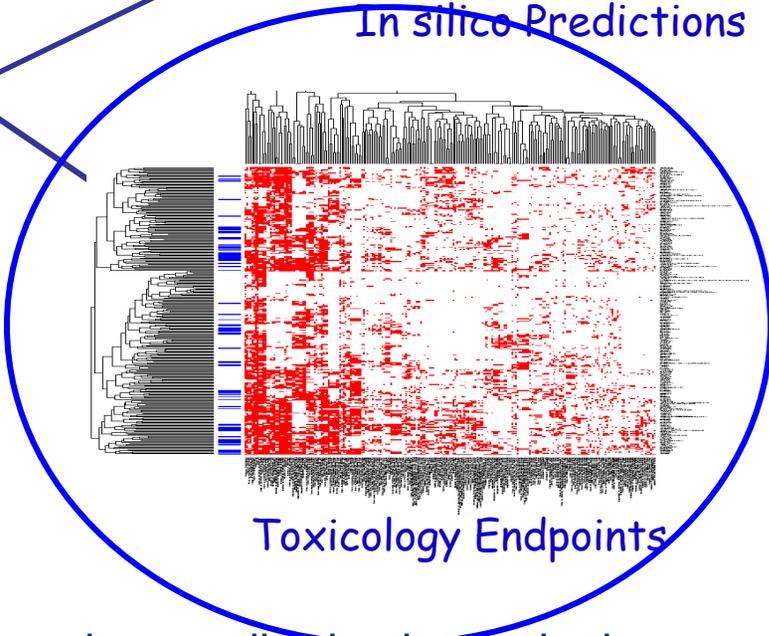
Profile Matching



Biochemical Assays



Genomic Signatures



Toxicology Endpoints

Find "Signatures" from *in vitro* & *in silico* assays that predict *In vivo* endpoints.

US EPA National Center for Computational Toxicology | US EPA - Windows Internet Explorer

US EPA http://www.epa.gov/ncct/toxrefdb/

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You are here: [EPA Home](#) » [National Center for Computational Toxicology](#) » [Toxicology Reference Database \(ToxRefDB\)](#)

**ToxRefDB Program**  
**Toxicology Reference Database**

Home  
 Basic Information  
 Organization  
 Post Doc Profiles  
 Framework  
 Databases and Models  
 Research Activities  
 ACToR  
 DSSTox  
 ToxCast™  
 ToxRefDB  
 Virtual Liver  
 v-Embryo™  
 Conferences and Seminars  
 Publications  
 BOSC Information  
 EPA Community of Practice  
 Jobs and Opportunities  
 Related Information

ToxRefDB was developed by the National Center for Computational Toxicology (NCCT) in partnership with EPA's Office of Pesticide Programs (OPP), to store data from in vivo animal toxicity studies. The initial focus was populating ToxRefDB with pesticide registration toxicity data that has been historically stored as hard-copy and scanned documents by OPP. A significant portion of these data have now been processed into ToxRefDB in a standardized and structured format. ToxRefDB currently includes chronic, cancer, sub-chronic, developmental, and reproductive studies on hundreds of chemicals, many of which are pesticide active ingredients. These data are now accessible and computable within ToxRefDB, and are serving as reference toxicity data for ORD research and OPP retrospective analyses. The primary research application of ToxRefDB is to provide toxicity endpoints for the development of ToxCast™ predictive signatures.

Data Set	Description	Download	Publication
Data Entry Tool & Controlled Vocabulary	The Data Entry Tool provided the user interface for all initial data input into ToxRefDB. The controlled vocabulary standardized the capturing of regulatory animal toxicity studies performed across various study types.	<a href="#">Download</a> (15.5 MB, ZIP)	Martin et al. (2008) " <a href="#">Profiling Chemicals Based on Chronic Toxicity Results from the U.S. EPA ToxRef Database</a> " Environmental Health Perspectives doi:10.1289/ehp.0800074
Chronic & Cancer Endpoints	Based on incidence, severity and potency, 26 primarily tissue-specific pathology endpoints were selected to uniformly classify 310 chemicals included in the manuscript's analysis. The 310 chemicals in this analysis largely overlap with the 320 ToxCast Phase I chemicals.	<a href="#">Download</a> (2.7 MB, XLS)	Martin et al. (2008) " <a href="#">Profiling Chemicals Based on Chronic Toxicity Results from the U.S. EPA ToxRef Database</a> " Environmental Health Perspectives doi:10.1289/ehp.0800074

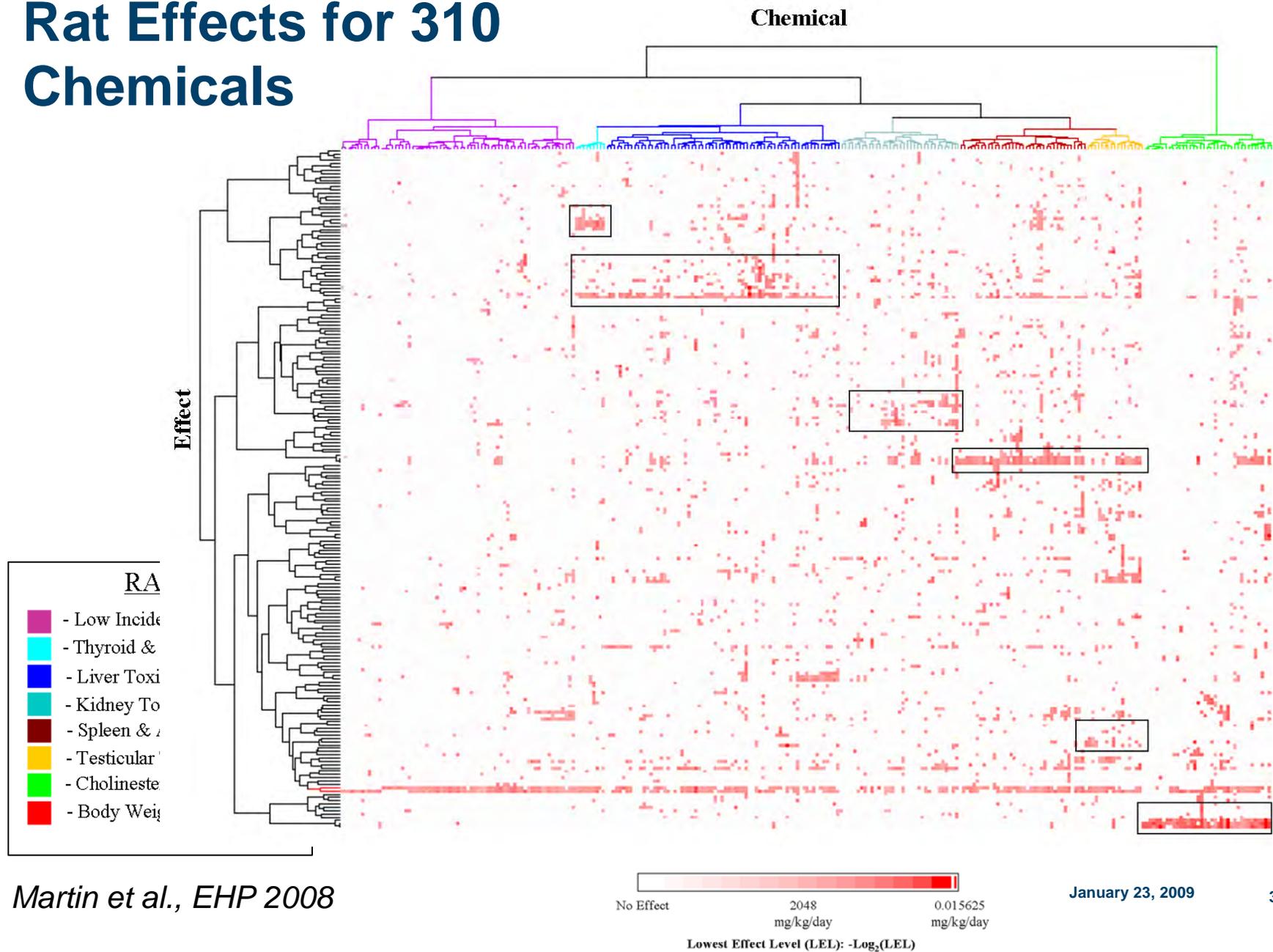
[EPA Home](#) | [Privacy and Security Notice](#) | [Contact Us](#)

Last updated on Tuesday, October 21st, 2008.  
<http://www.epa.gov/ncct/toxrefdb/>  
[Print As-Is](#)

ToxRefDB website: <http://www.epa.gov/ncct/toxrefdb/>

Local intranet 100%

# ToxRefDB Chronic Rat Effects for 310 Chemicals

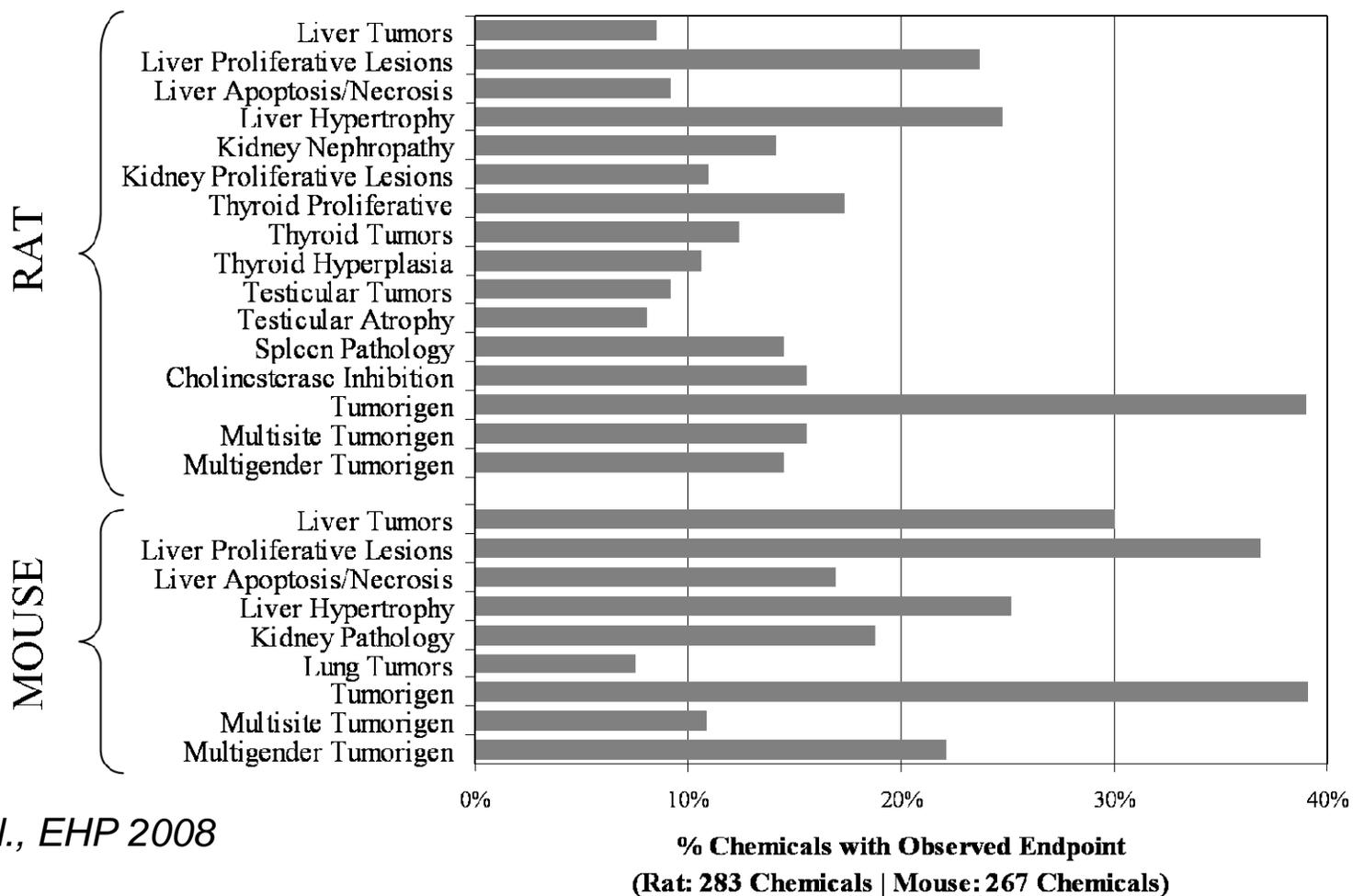


Martin et al., EHP 2008

January 23, 2009



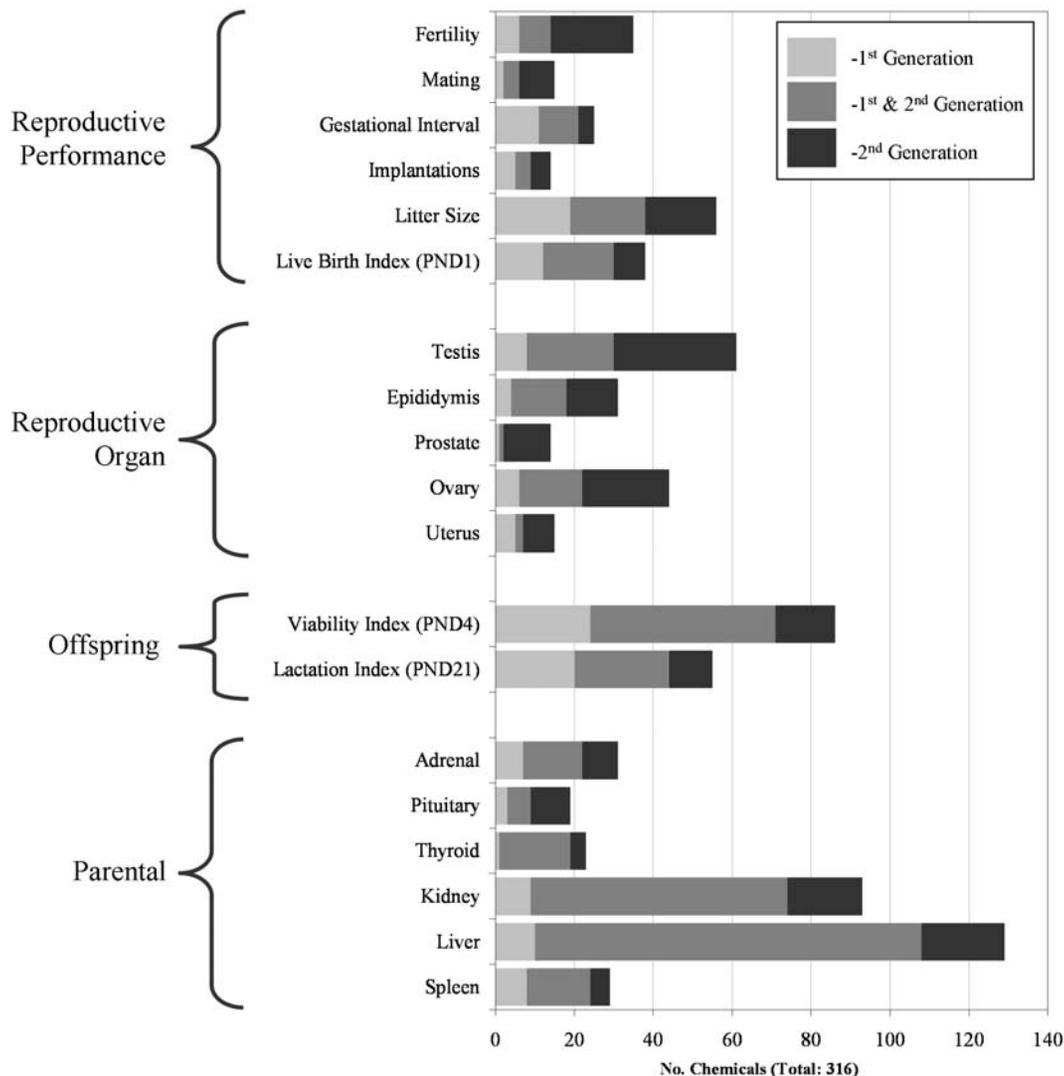
# Selected Chronic Rat & Mouse Endpoints for ToxCast Predictive Modeling



Martin et al., EHP 2008



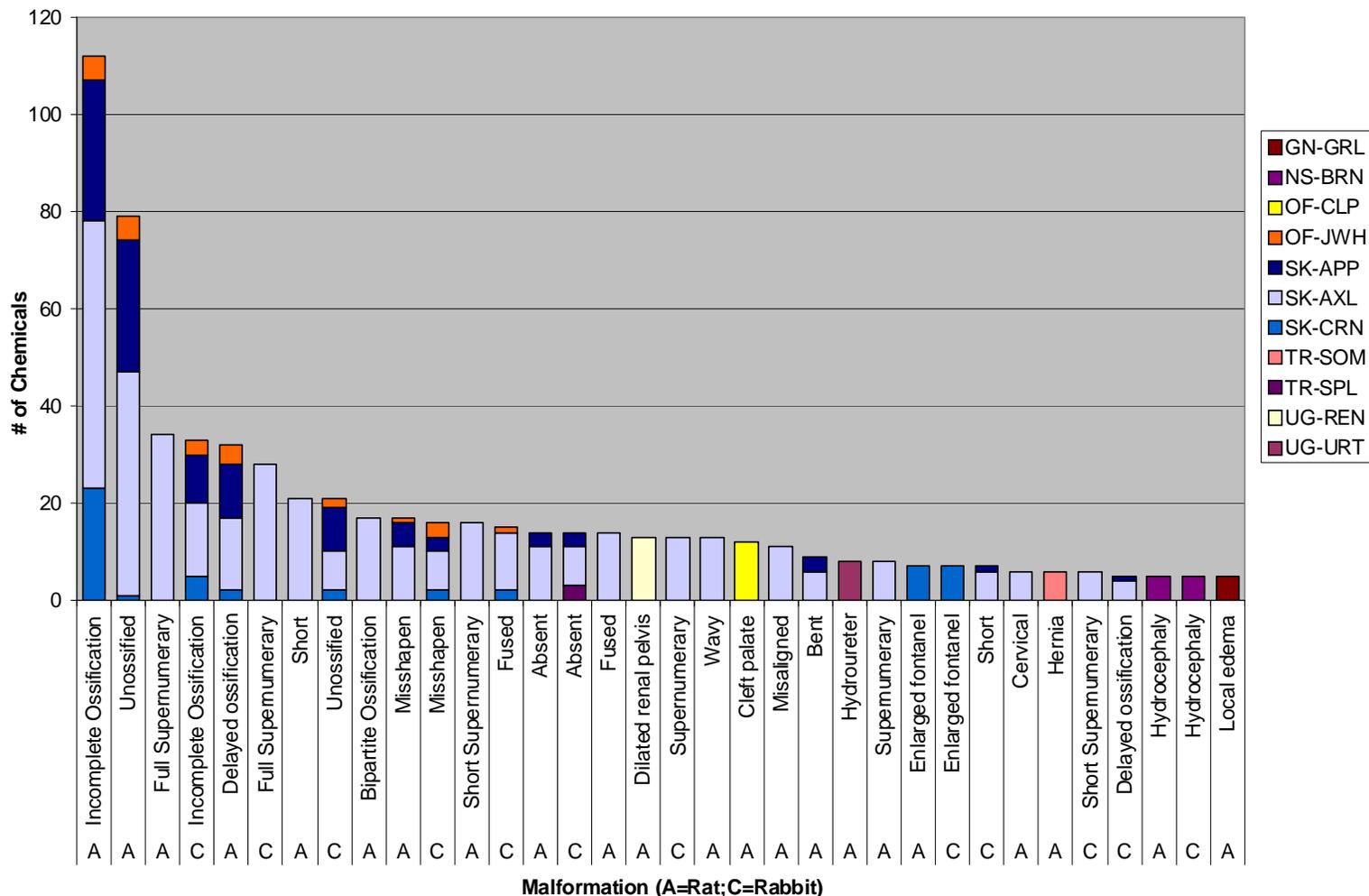
# Selected Rat Reproductive Endpoints for ToxCast Predictive Modeling



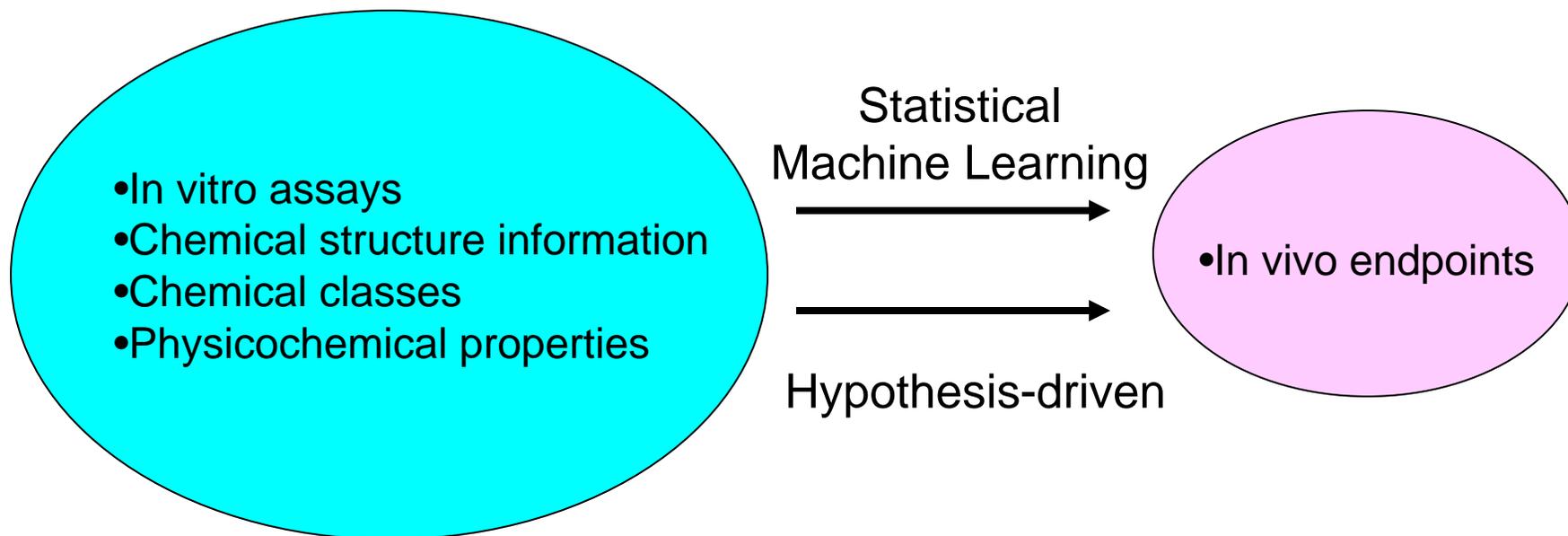
Martin et al., sub



# Selected Developmental Rat & Rabbit Endpoints for ToxCast Predictive Modeling

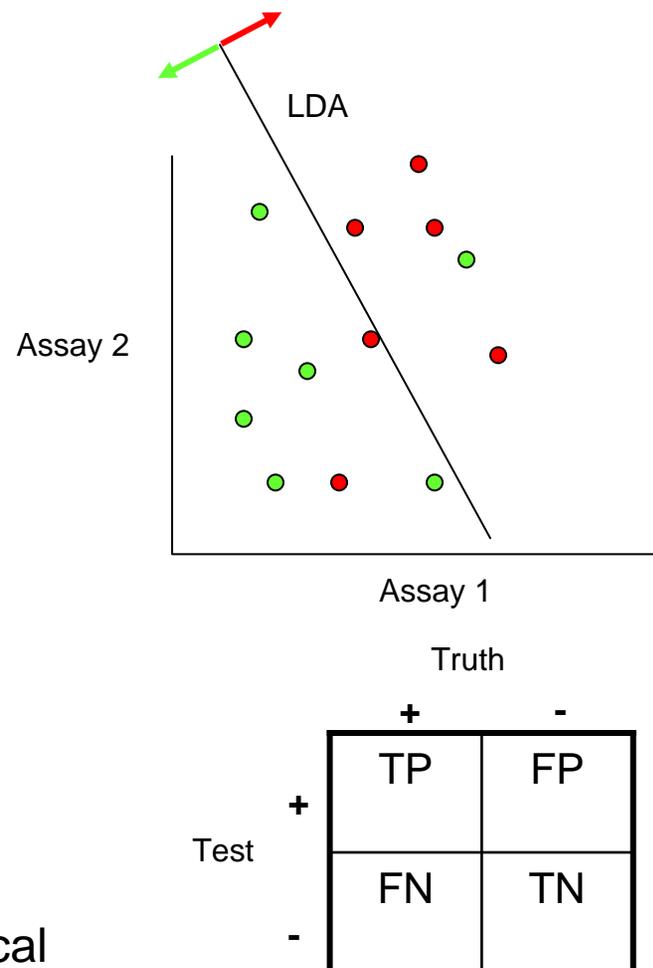


# ToxCast Analysis Approaches



# Association Analysis / Signatures

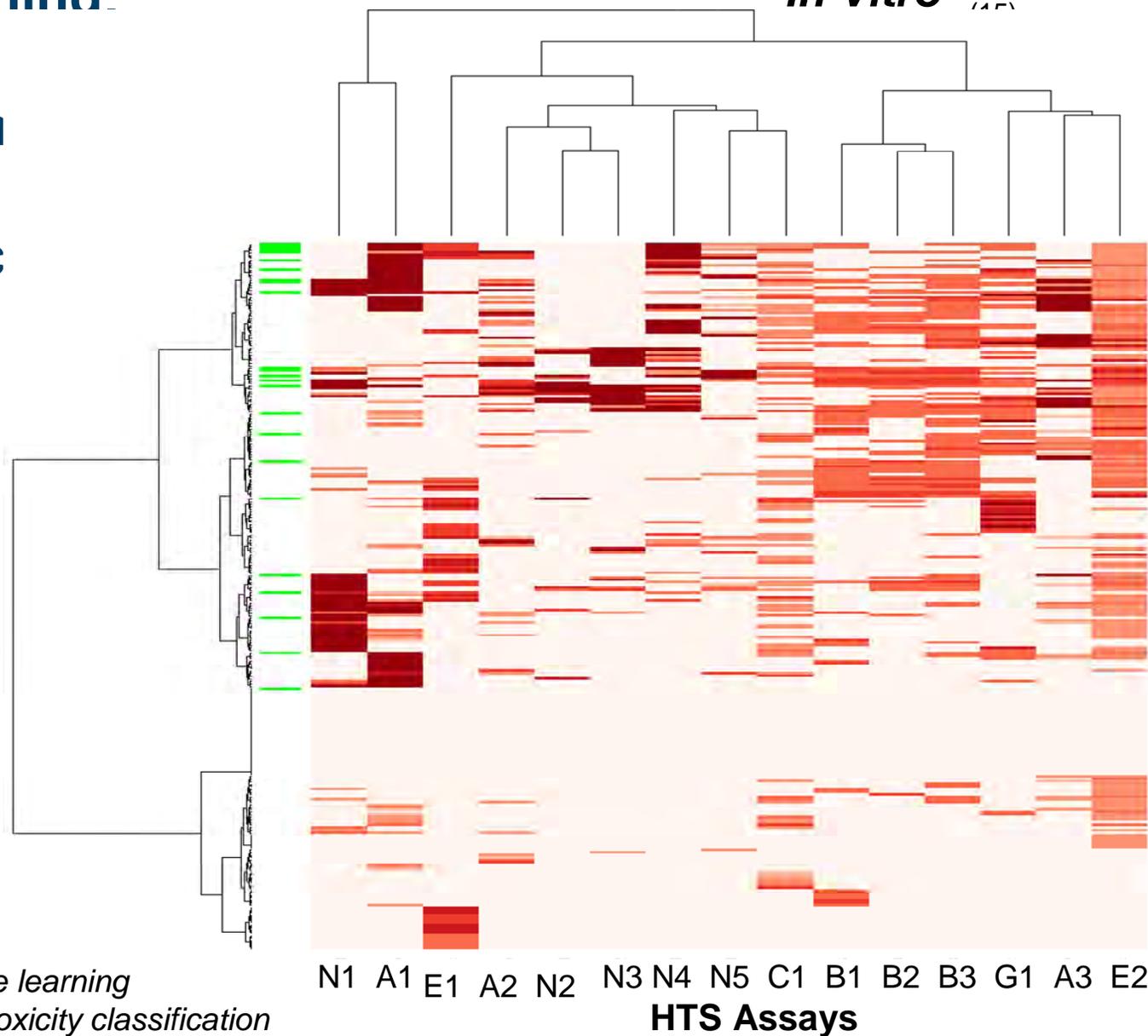
- Use Machine Learning methods
  - SLR: Stepwise Logistic Regression
  - LDA: Linear Discriminant Analysis
  - SVM: Support Vector Machines
  - Many others
- For each binary endpoint, build models of form
  - $Predictor = F(\text{assay values})$
  - If
    - $Predictor$  for a chemical meets criteria
  - Then
    - Predict endpoint to be positive for the chemical



# Machine Learning: ToxCast Prediction Modeling of Chemical Rat Liver Apoptosis/Necrosis

*In Vivo*  
(23)

*In Vitro* (45)



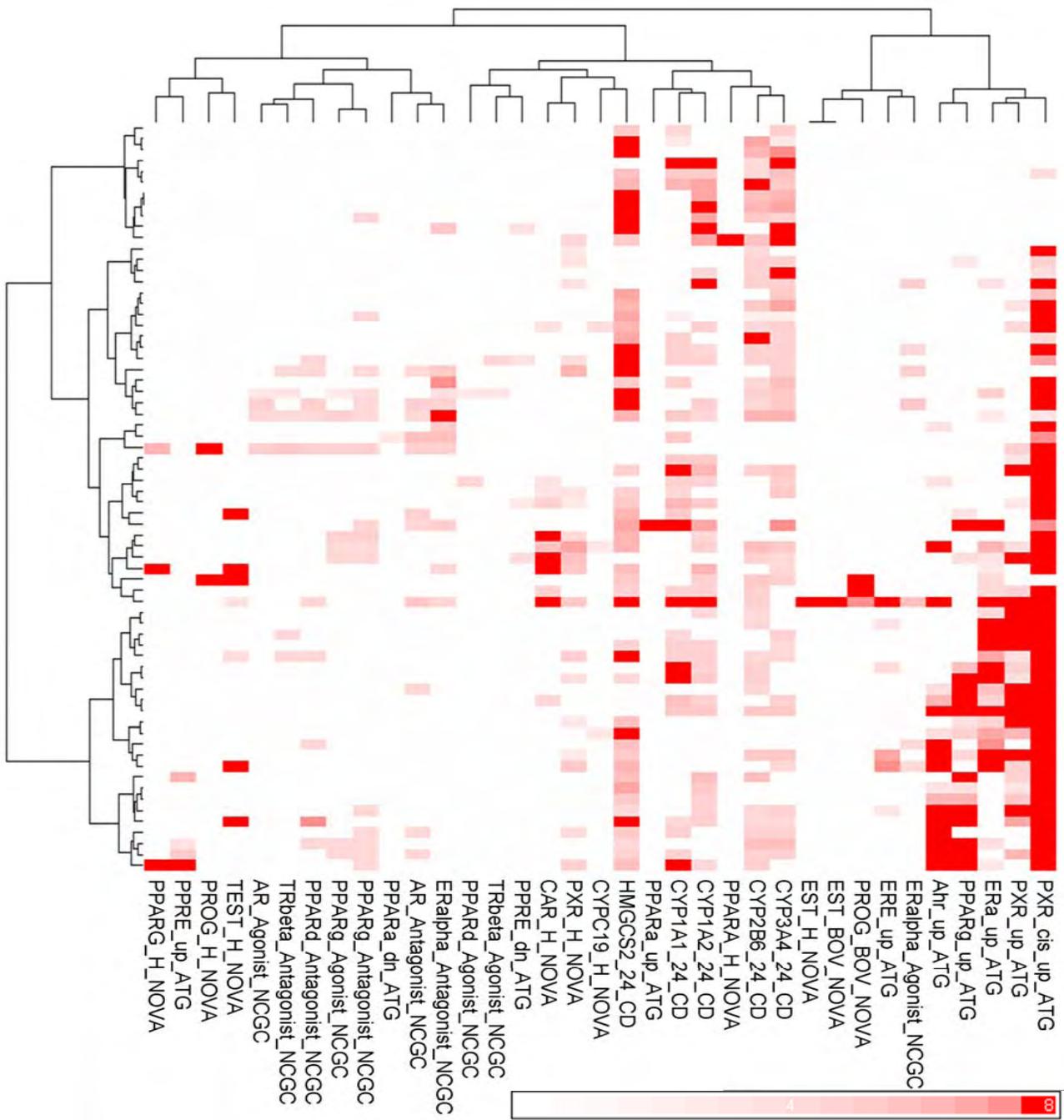
Methods described in:

*Judson et al 2008*

*A comparison of machine learning algorithms for chemical toxicity classification using a simulated multi-scale data model.*

*BMC Bioinformatics 9:241*

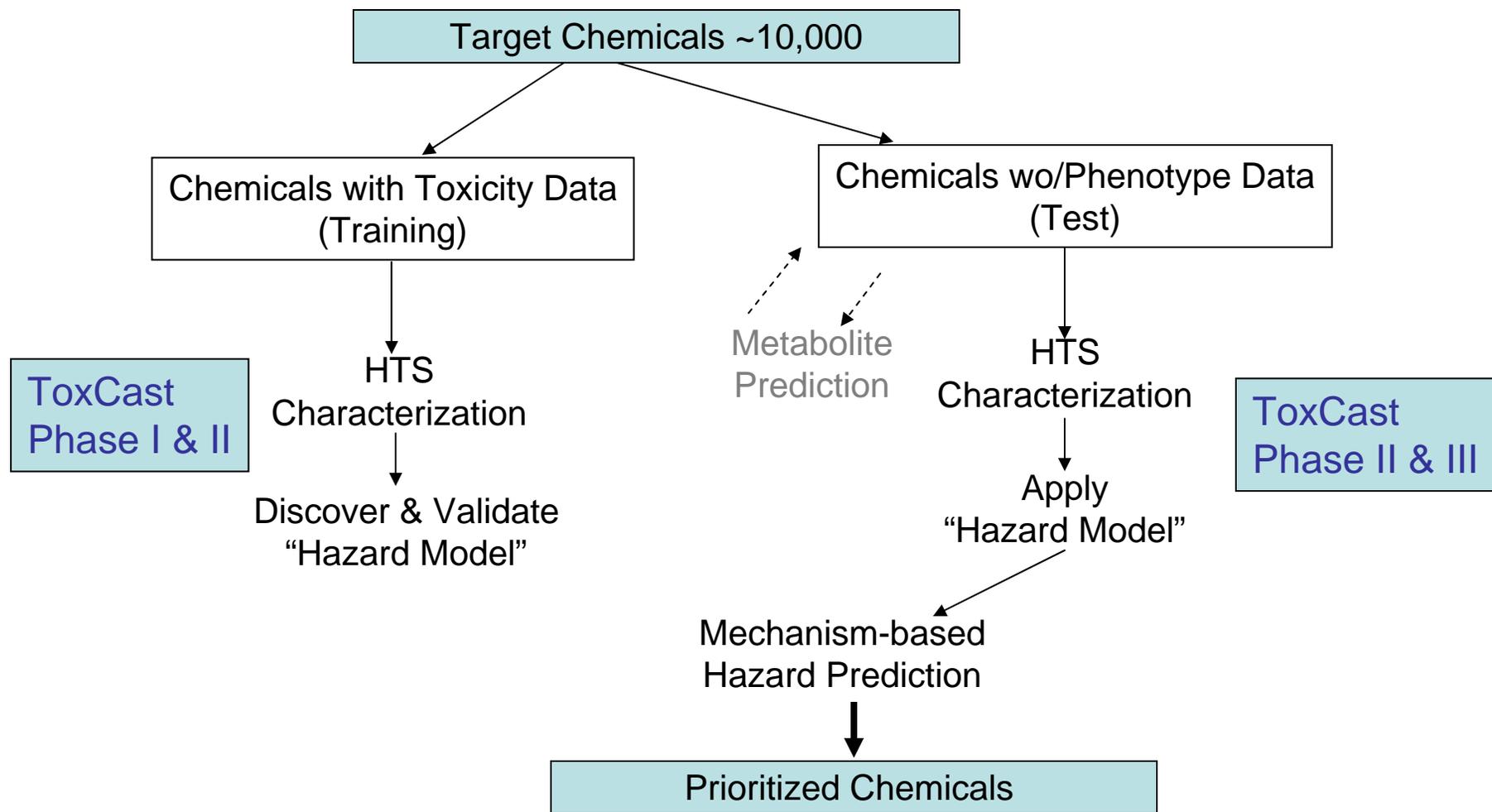
# Hypothesis Driven: ToxCast Endocrine Profiling



← Vinclozolin  
← Bisphenol A



# ToxCast Screening and Prioritization

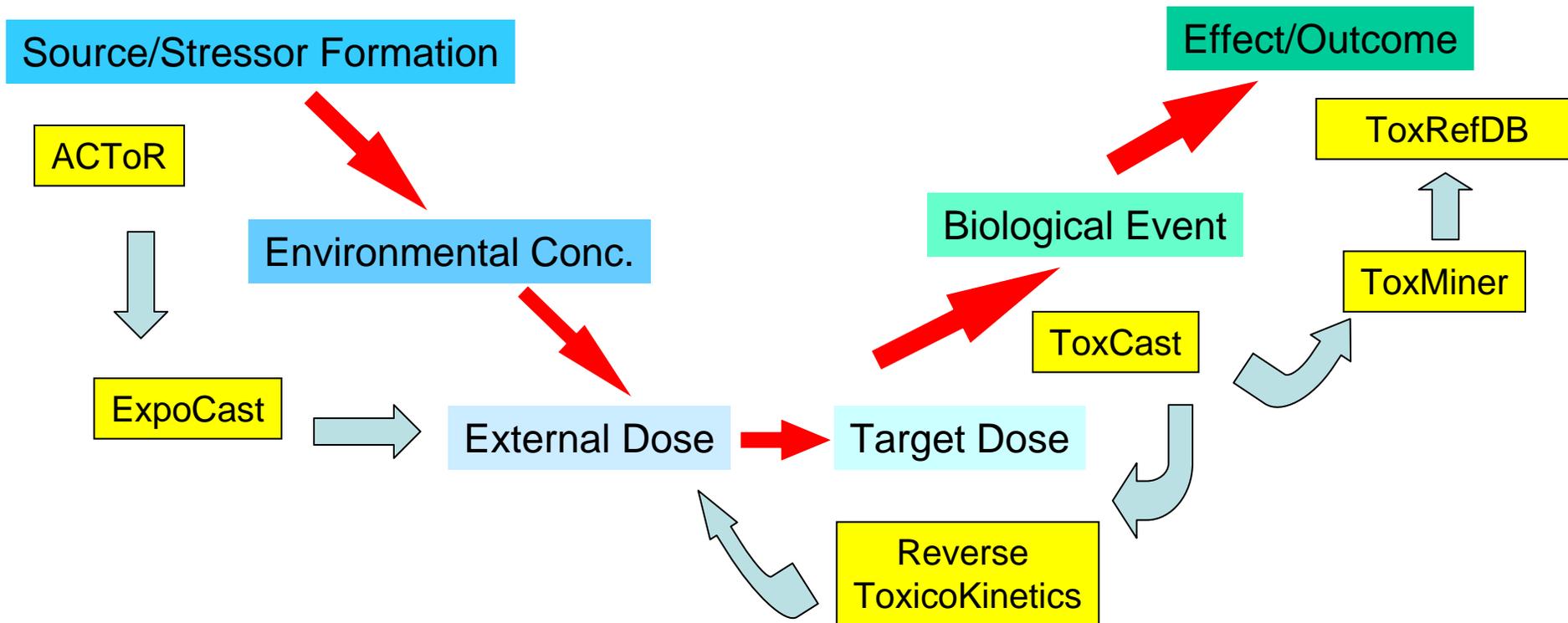


## Moving Forward with ToxCast

- First predictive toxicity signatures based on ToxCast data submitted for publication April 2009
- ToxCast data available to collaborators now, publicly available May 2009 at 1<sup>st</sup> ToxCast Data Analysis Summit
- EPA & partners examining methods for analyzing ToxCast data, identifying predictive signatures from Phase I for validation in Phase II
- Phase II testing will commence June 2009 on upwards of 700 additional chemicals.



# Applying Computational Toxicology Along the Source to Outcome Continuum



## National Center for Computational Toxicology

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## ToxCast™ Program

### Predicting Hazard, Characterizing Toxicity Pathways, and Prioritizing the Toxicity Testing of Environmental Chemicals

#### Introduction

In 2007, EPA launched ToxCast™ in order to develop a cost-effective approach for prioritizing the toxicity testing of large numbers of chemicals in a short period of time. Using data from state-of-the-art high throughput screening (HTS) bioassays developed in the pharmaceutical industry, ToxCast™ is building computational models to forecast the potential human toxicity of chemicals. These hazard predictions will provide EPA regulatory programs with science-based information helpful in prioritizing chemicals for more detailed toxicological evaluations, and lead to more efficient use of animal testing.

In its first phase, ToxCast™ is profiling over 300 well-characterized chemicals (primarily pesticides) in over 400 HTS endpoints. These endpoints include biochemical assays of protein function, cell-based transcriptional reporter assays, multi-cell interaction assays, transcriptomics on primary cell cultures, and developmental assays in zebrafish embryos. Almost all of the compounds being examined in Phase 1 of ToxCast™ have been tested in traditional toxicology tests, including developmental toxicity, multi-generation studies, and sub-chronic and chronic rodent bioassays. ToxRefDB, a relational database being created to house this information, will contain nearly \$1B worth of toxicity studies in animals when completed. ToxRefDB is integrated into a more comprehensive data management system developed by NCCT called ACToR (Aggregated Computational Toxicology Resource), that manages the large-scale datasets of ToxCast™. ACToR is comprised of several independent data repositories linked to a common database of chemical structures and properties, and to tools for development of predictive HTS and genomic bioactivity signatures that strongly correlate with specific toxicity endpoints from ToxRefDB. These ToxCast™ signatures will be defined and evaluated by their ability to predict outcomes from existing mammalian toxicity testing, and identify toxicity pathways that are relevant to human health effects.

The second phase of ToxCast™ will screen additional compounds representing broader chemical structure and use classes, in order to evaluate the predictive bioactivity signatures developed in Phase I. Following successful conclusion of Phases I and II, ToxCast™ will provide EPA regulatory programs an efficient tool for rapidly and efficiently screening compounds and prioritizing further toxicity testing.

#### ToxCast™ Navigation

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