

Developing Predictive Bioactivity Signatures from ToxCast's HTS Data

TestSmart DNT 2 Conference – Washington DC, November 12-14 2008

UNITED STATES ENVIRONMENTAL PROTECTION AGENCY

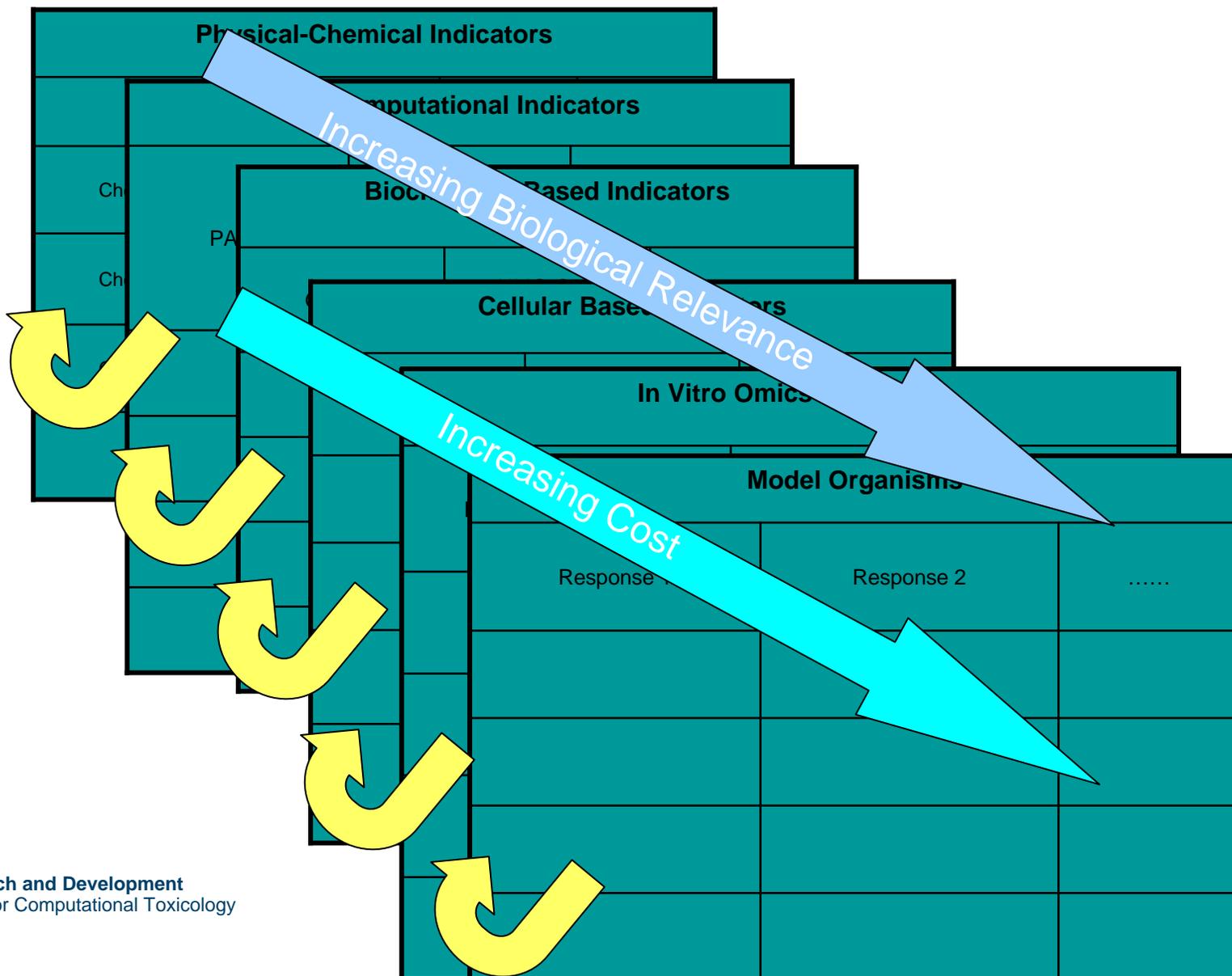


**COMPUTATIONAL
TOXICOLOGY**

Co-authors: David Dix, Keith Houck, Matt Martin, David Reif and Richard Judson

A Dream from TestSmart 1

Chemical Clusters
Bin 1
Bin 2
Bin 3
...
...
Bin ...



Future of Toxicity Testing

POLICYFORUM

TOXICOLOGY

Transforming Environmental Health Protection

Francis S. Collins,^{1*} George M. Gray,^{2*} John R. Bucher^{3*}

We propose a shift from primarily in vivo animal studies to in vitro assays, in vivo assays with lower organisms, and computational modeling for toxicity assessments.

In 2005, the U.S. Environmental Protection Agency (EPA), with support from the U.S. National Toxicology Program (NTP), funded a project at the National Research Council (NRC) to develop a long-range vision for toxicity testing and a strategic plan for implementing that vision. Both agencies wanted future toxicity testing and assessment paradigms to meet evolving regulatory needs. Challenges include the large numbers of substances that need to be tested and how to incorporate recent advances in molecular toxicology, computational sciences, and information technology, to rely increasingly on human as opposed to animal data; and to offer increased efficiency in design and costs (1–5). In response, the NRC Committee on Toxicity Testing and Assessment of Environmental Agents produced two reports that reviewed current toxicity testing, identified key issues, and developed a vision and implementation strategy to create a major shift in the assessment of chemical hazard and risk (6, 7). Although the NRC reports have laid out a solid theoretical rationale, comprehensive and rigorously gathered data (and comparisons with historical animal data) will determine whether the hypothesized improvements will be realized in practice. For this purpose, NTP, EPA, and the National Institutes of Health Chemical Genomics Center (NCGC) (organizations with expertise in experimental toxicology, computational toxicology, and high-throughput technologies, respectively) have established a collaborative research program.

EPA, NCGC, and NTP Joint Activities
In 2004, the NTP released its vision and roadmap for the 21st century (1), which established initiatives to integrate high-

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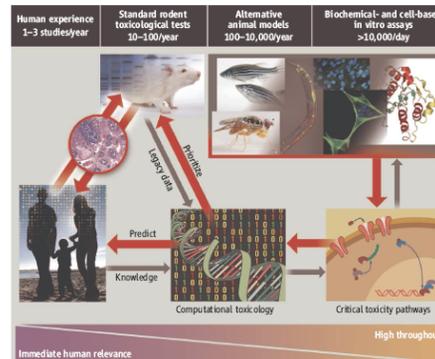
*The views expressed here are those of the individual authors and do not necessarily reflect the views and policies of their respective agencies.

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throughput screening (HTS) and other automated screening assays into its testing program. In 2005, the EPA established the National Center for Computational Toxicology (NCCT). Through these initiatives, NTP and EPA, with the NCGC, are promoting the evolution of toxicology from a predominantly observational science at the level of disease-specific models in vivo to a predominantly predictive science focused on broad inclusion of target-specific, mechanism-based, biological observations in vitro (1, 4) (see figure, below).

Toxicity pathways. In vitro and in vivo tools are being used to identify cellular responses after chemical exposure expected to result in adverse health effects (7). HTS methods are a primary means of discovery for drug development, and screening of >100,000 compounds per day is routine (8). However, drug-discovery HTS methods traditionally test compounds at one concentration,

usually between 2 and 10 μM, and tolerate high false-negative rates. In contrast, in the EPA, NCGC, and NTP combined effort, all compounds are tested at as many as 15 concentrations, generally ranging from ~5 nM to ~100 μM, to generate a concentration-response curve (9). This approach is highly reproducible, produces significantly lower false-positive and false-negative rates than the traditional HTS methods (9), and facilitates multitask comparisons. Finally, an informatics platform has been built to compare results among HTS screens; this is being expanded to allow comparisons with historical toxicologic NTP and EPA data (<http://ncg.nih.gov/pub/openhts>). HTS data collected by EPA and NTP, as well as by the NCGC and other Molecular Libraries Initiative centers (<http://mli.nih.gov/>), are being made publicly available through Web-based databases [e.g., PubChem (<http://pubchem.ncbi.nlm.nih.gov/>)]. In addition,



Transforming toxicology. The studies we propose will test whether high-throughput and computational toxicology approaches can yield data predictive of results from animal toxicity studies, will allow prioritization of chemicals for further testing, and can assist in prediction of risk to humans.

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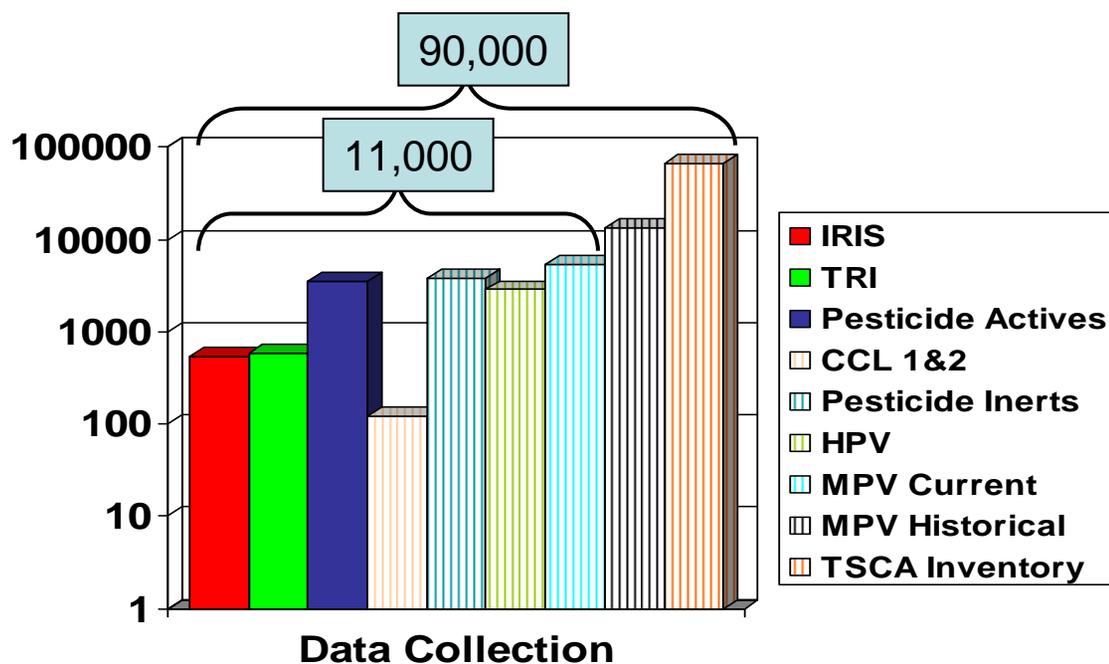
- ● Cancer
- ● ReproTox
- ● DevTox
- ● NeuroTox
- ● PulmonaryTox
- ● ImmunoTox



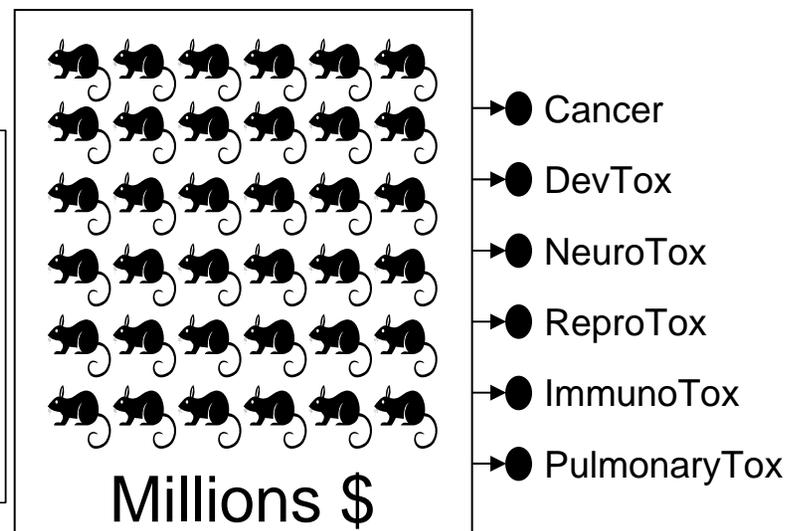
EPAs Contribution: The ToxCast Research Program

Change Needed Because

Too Many Chemicals



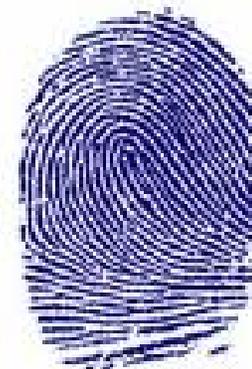
Too High a Cost



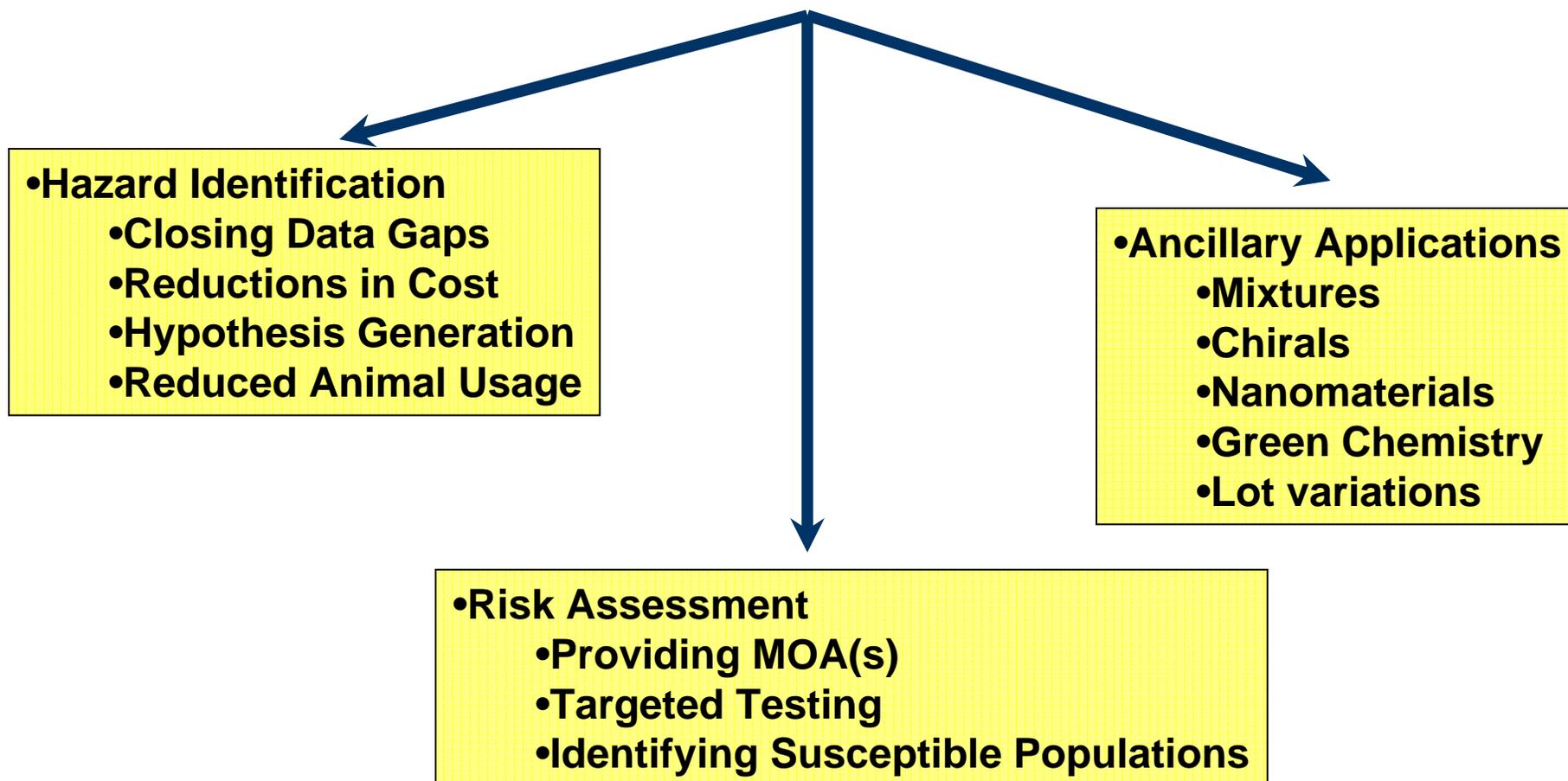
...and not enough data.

ToxCast™ Background

- 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 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://www.epa.gov/actor/>



Implications for Success



Key Challenges Of Pathway Profiling

- Find the Toxicity Pathways
 - Hepato vs developmental neurotoxicity
- Obtain HTS Assays for Them
 - Including metabolic capability
- Screen Chemical Libraries
 - Coverage of p-chem properties
- Link Results to in vivo Effects
 - Gold standard and dosimetry

Phased Development of ToxCast

Phase	Number of Chemicals	Chemical Criteria	Purpose	Number of Assays	Cost per Chemical	Target Date
Ia	320	Data Rich (pesticides)	Signature Development	552	\$20k	FY08 ¹
Ib	15	Nanomaterials	Pilot	166	\$10K	FY09
IIa	>300	Data Rich Chemicals	Validation	>400	~\$20-25k	FY09
IIb	>100	Known Human Toxicants	Extrapolation	>400	~\$20-25k	FY09
IIc	>300	Expanded Structure and Use Diversity	Extension	>400	~\$20-25k	FY10
IId	>12	Nanomaterials	PMN	>200	~\$15-20K	FY09-10
III	Thousands	Data poor	Reducing to Practice	>300	~\$15-20k	FY11-12

¹Initiated April 2007

The ToxCast_320

309 Unique Structures

Replicates for QC

291 Pesticide Actives

9 Industrial Chemicals

8 Metabolites

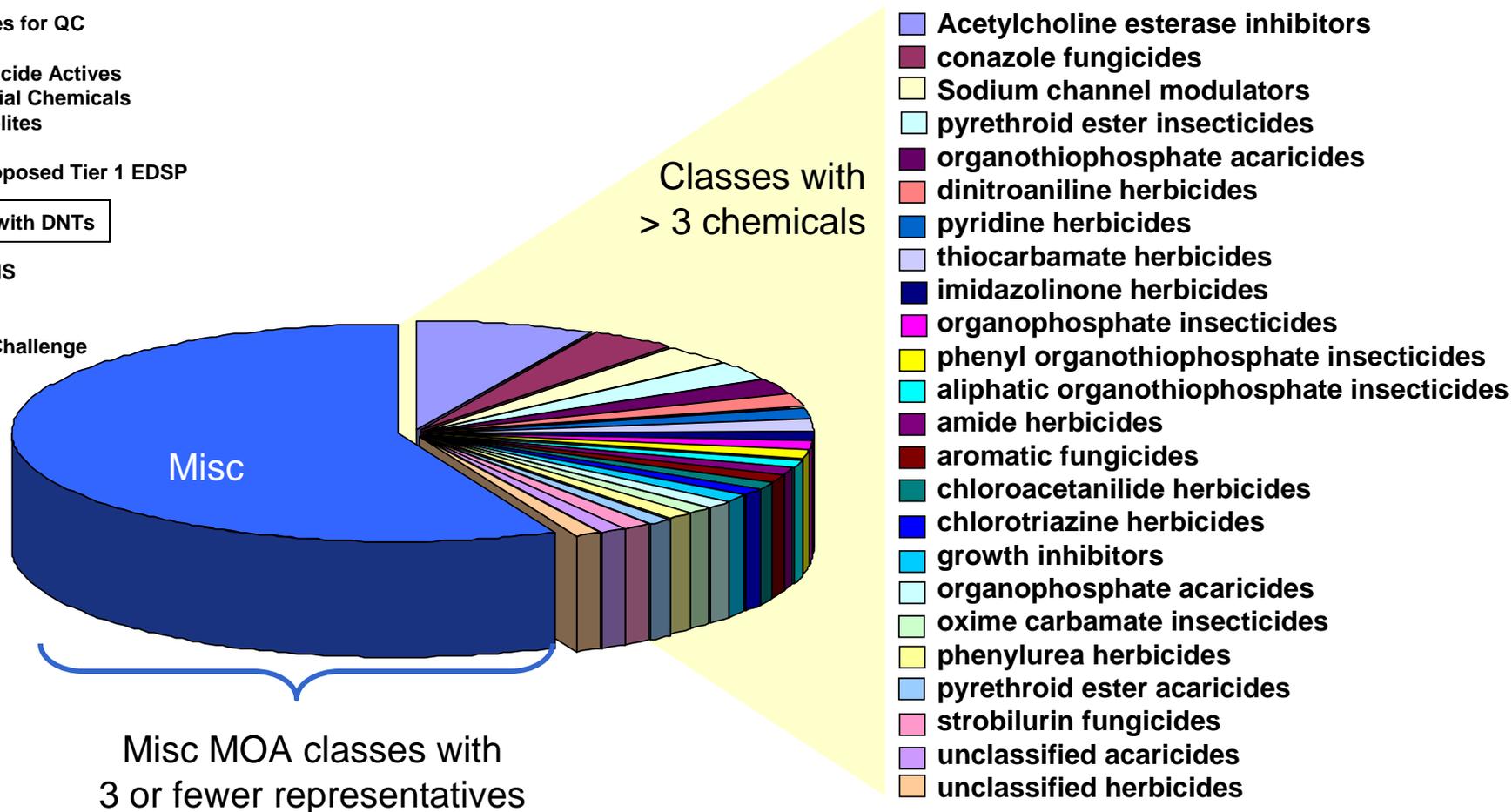
56/73 Proposed Tier 1 EDSP

53 of 80 with DNTs

122 in IRIS

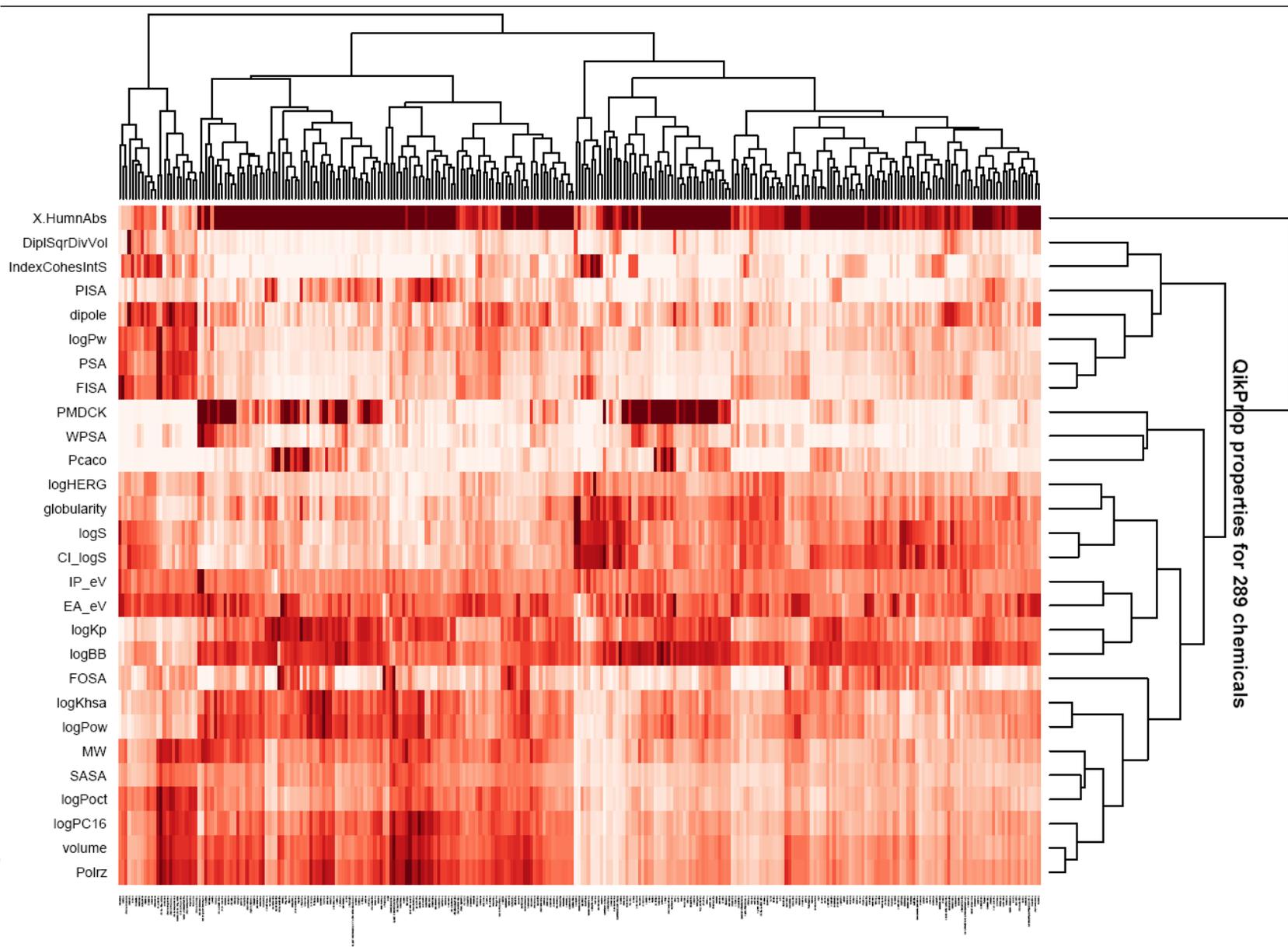
14 HPV

11 HPV Challenge

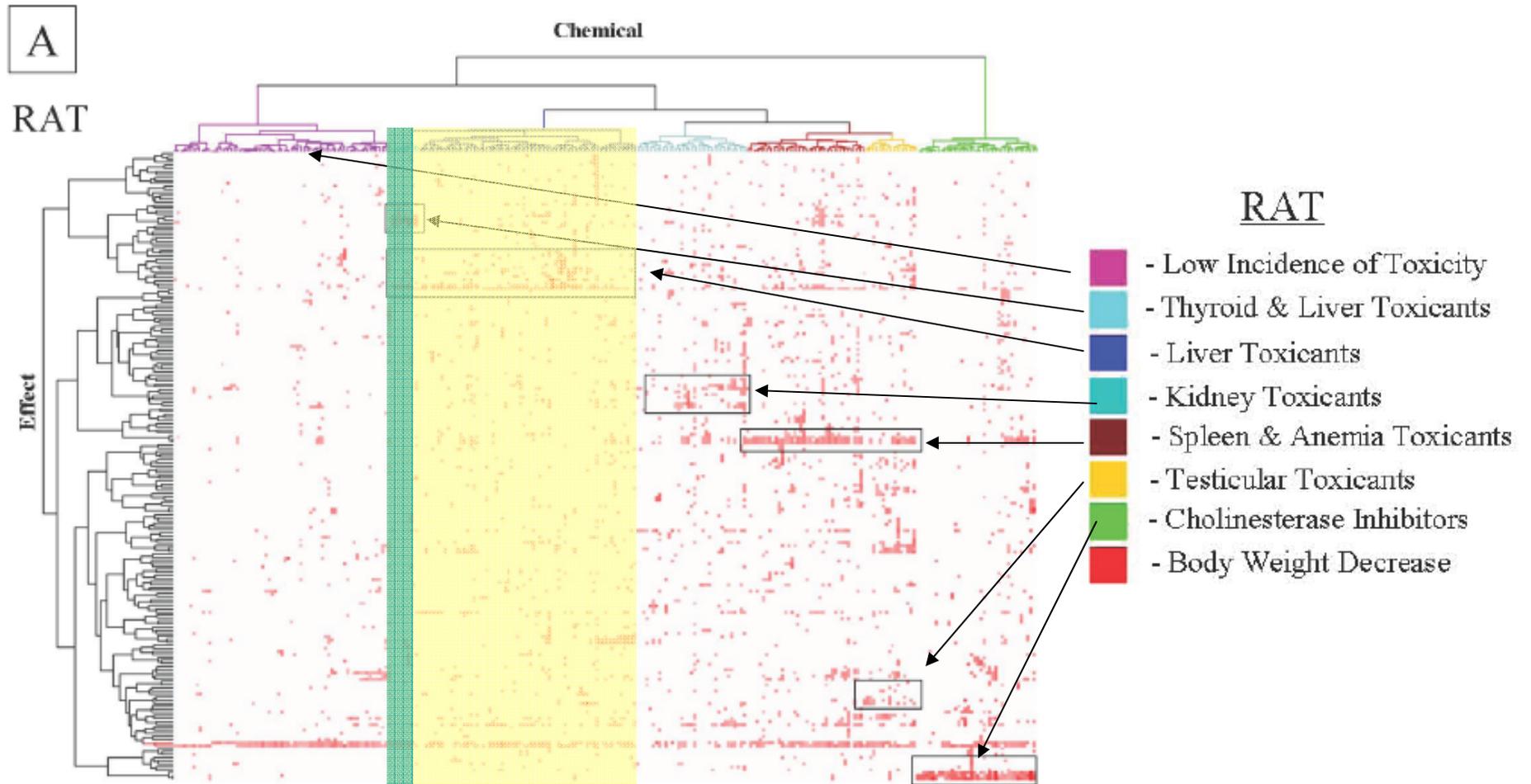


Classification based on OPPIN

Physical-Chemical Properties



>\$1B in Toxicology Now Stored in ToxRefDB





National Center for Computational Toxicology

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ToxRefDB Program Toxicology Reference Database

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.	Download (15.5 MB, ZIP)	Martin et al. (2008) " Profiling Chemicals Based on Chronic Toxicity Results from the U.S. EPA ToxRef Database " 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.	Download (2.7 MB, XLS)	Martin et al. (2008) " Profiling Chemicals Based on Chronic Toxicity Results from the U.S. EPA ToxRef Database " Environmental Health Perspectives doi:10.1289/ehp.0800074

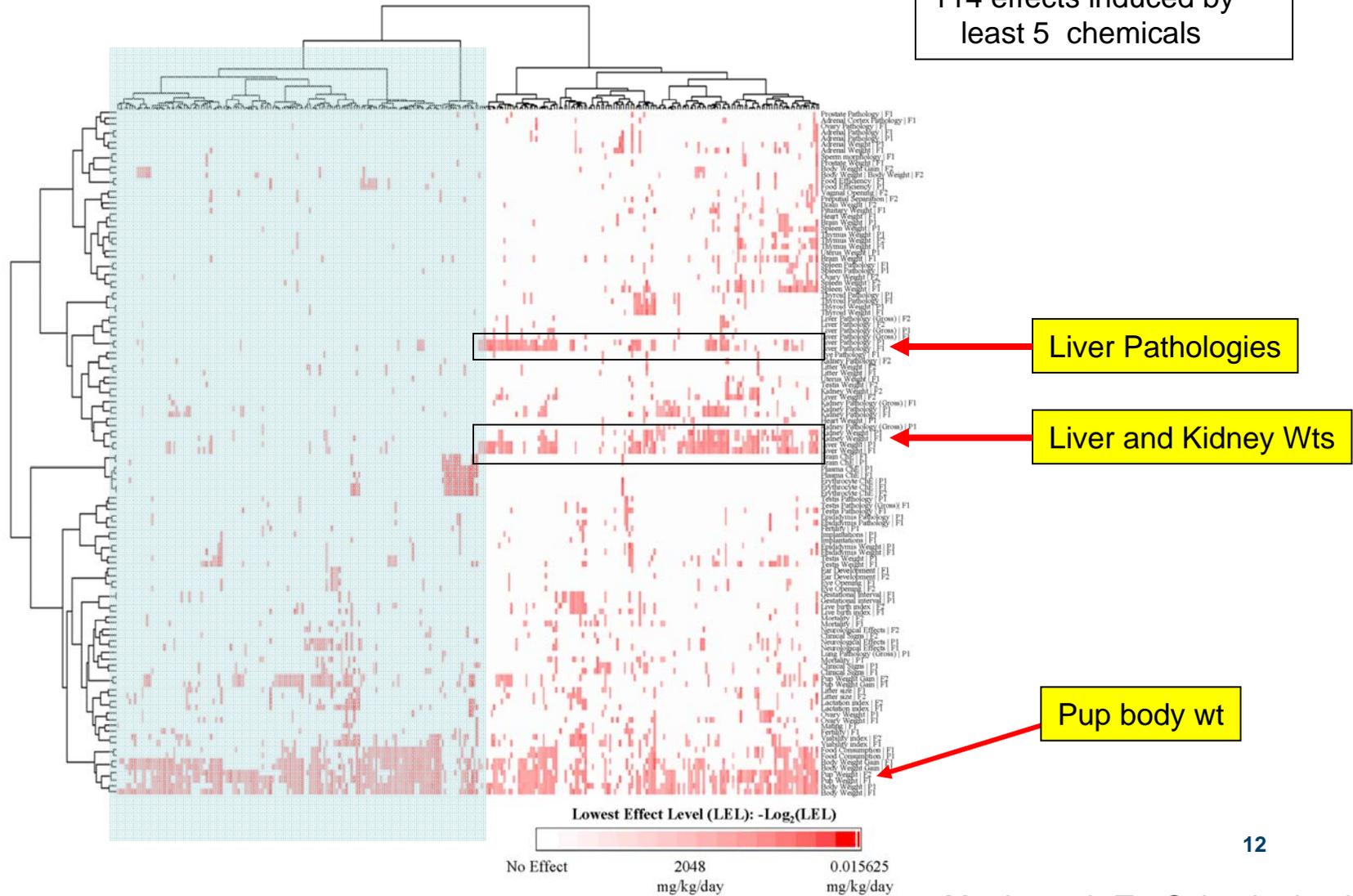
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Last updated on Tuesday, October 21st, 2008.
<http://www.epa.gov/ncct/toxrefdb/>
[Print As-Is](#)

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

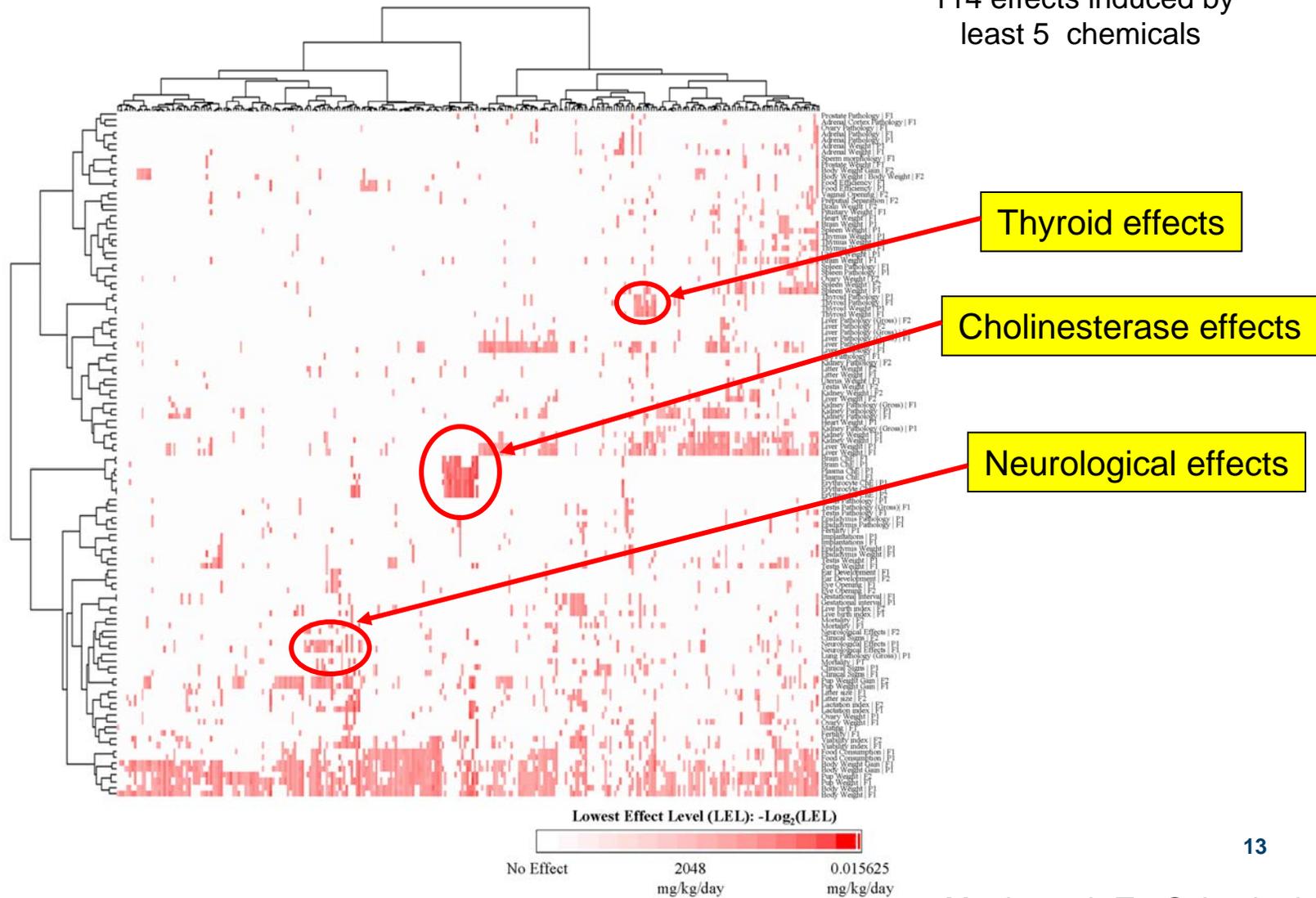
ToxRefDB Multigeneration Studies

114 effects induced by
least 5 chemicals



ToxRefDB Multigeneration Studies

114 effects induced by
least 5 chemicals

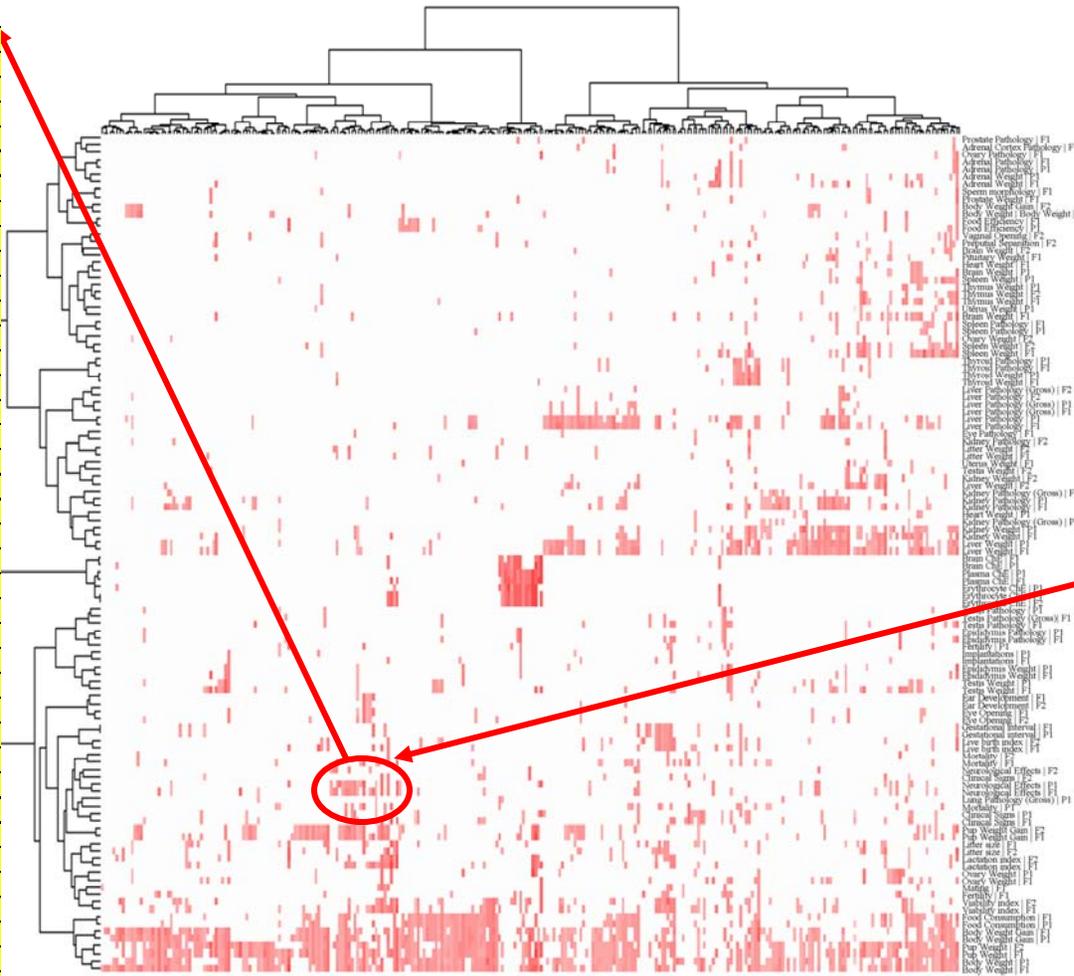


ToxRefDB Multigeneration Studies

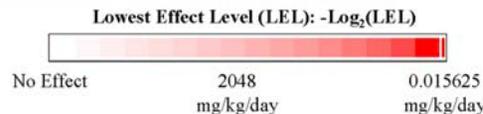
114 effects induced by
least 5 chemicals

F1 Actives

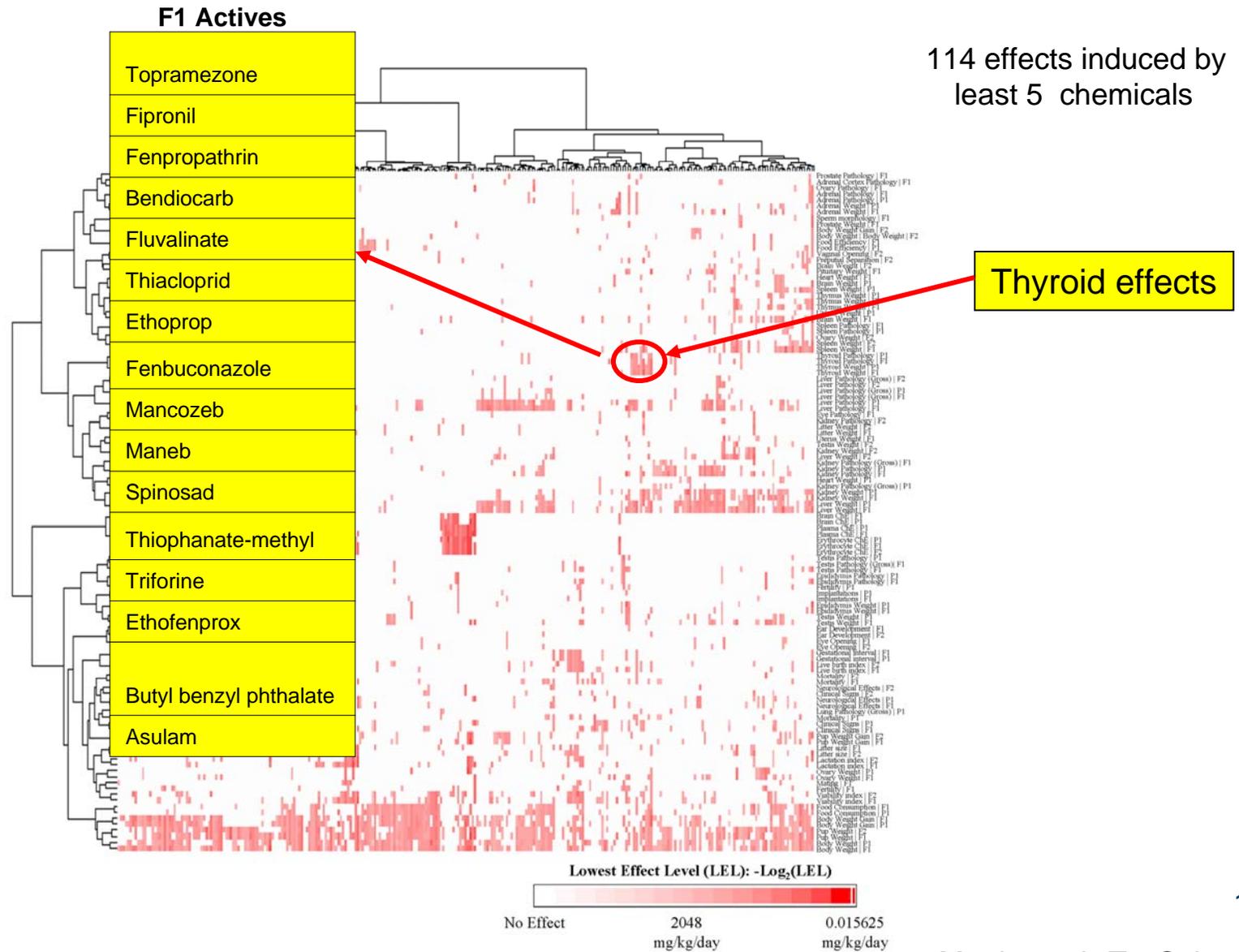
Phorate
Chlorethoxyfos
Sodium pyriithione
Potassium perfluorooct
Emamectin benzoate
Methidathion
Bifenthrin
Propetamphos
Cyfluthrin
Fluvalinate
Fenpropathrin
Oxamyl
Diquat dibromide
Paclobutrazol
Oryzalin
Ethalfuralin
Deltamethrin
Cypermethrin
Tri-allate
Tetraconazole
Oryzalin
Permethrin
S-Bioallethrin
Iprodione
Flufenpyr-ethyl
Clomazone
Fludioxonil
Ethofenprox
Vinclozolin
Diphenylamine
Dicamba
2,4-Dichlorophenol
Mepiquat chloride
Quintozene
Dinotefuran
Forchlorfenuron
Flumiclorac-pentyl
Clofencet
Pyriithiobac-sodium



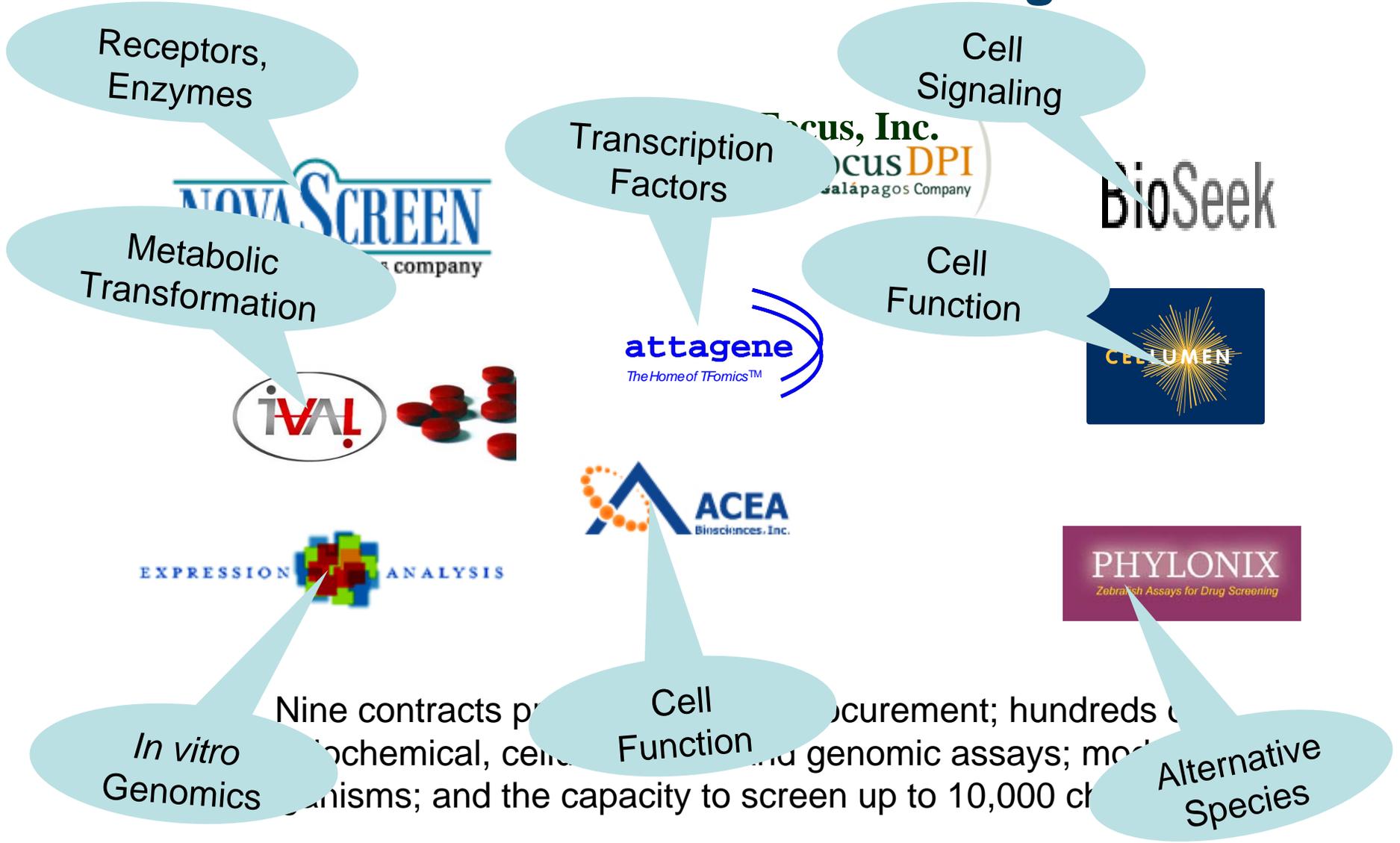
Neurological effects



ToxRefDB Multigeneration Studies



ToxCast Contracts for Generating HTS Data

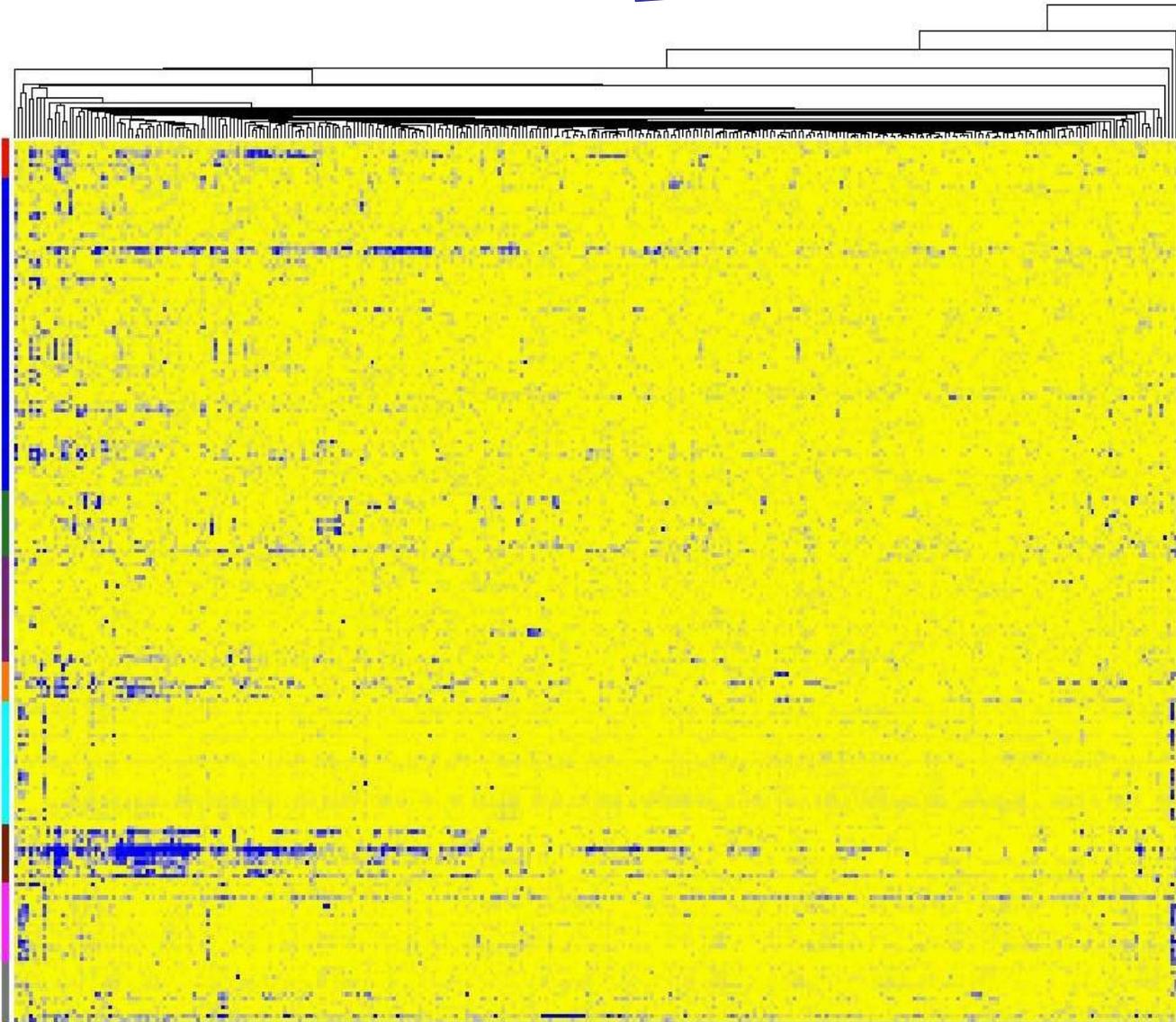




320 Chemicals



Transporter
GPCR
Enzyme, other
Ion channel
NR
Kinase
CYP450
Phosphatase
Protease

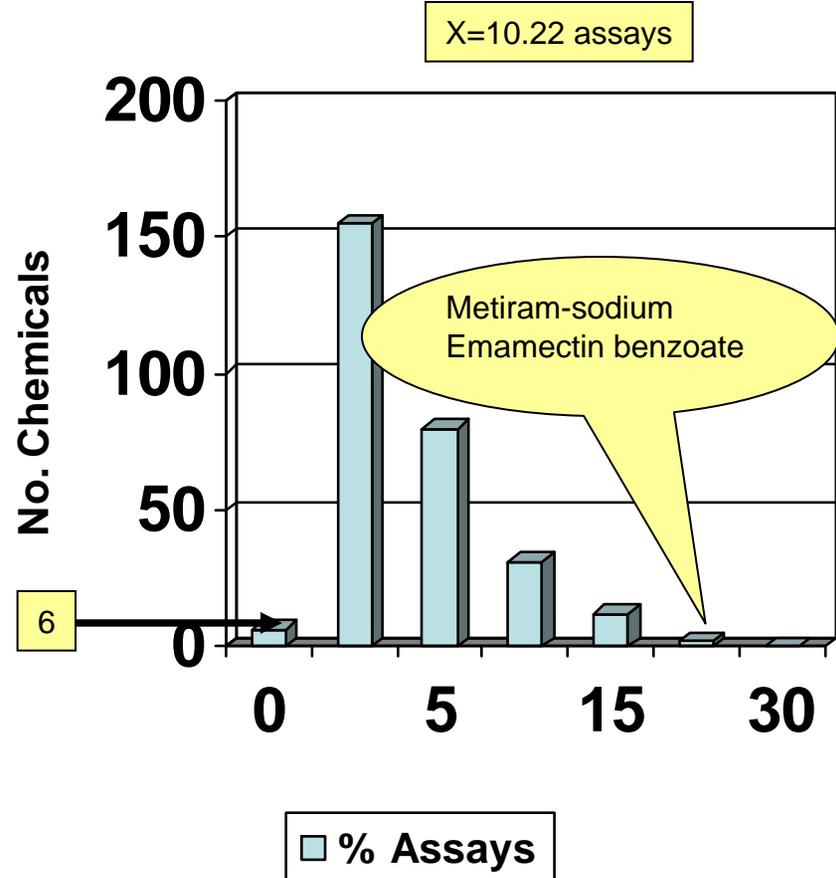
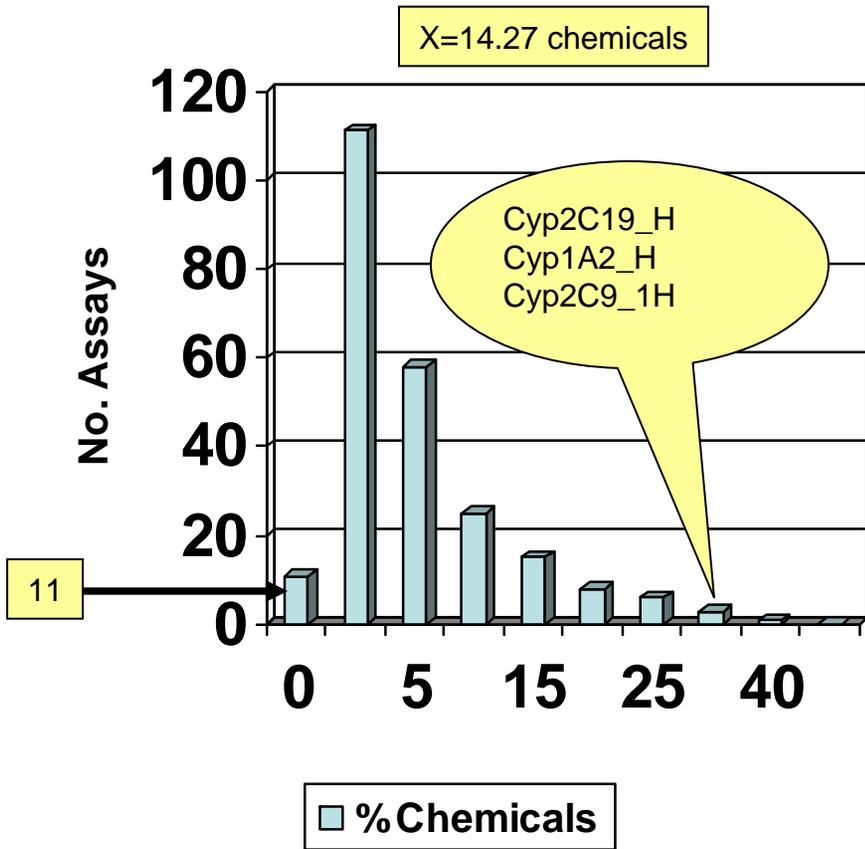


201 Assays

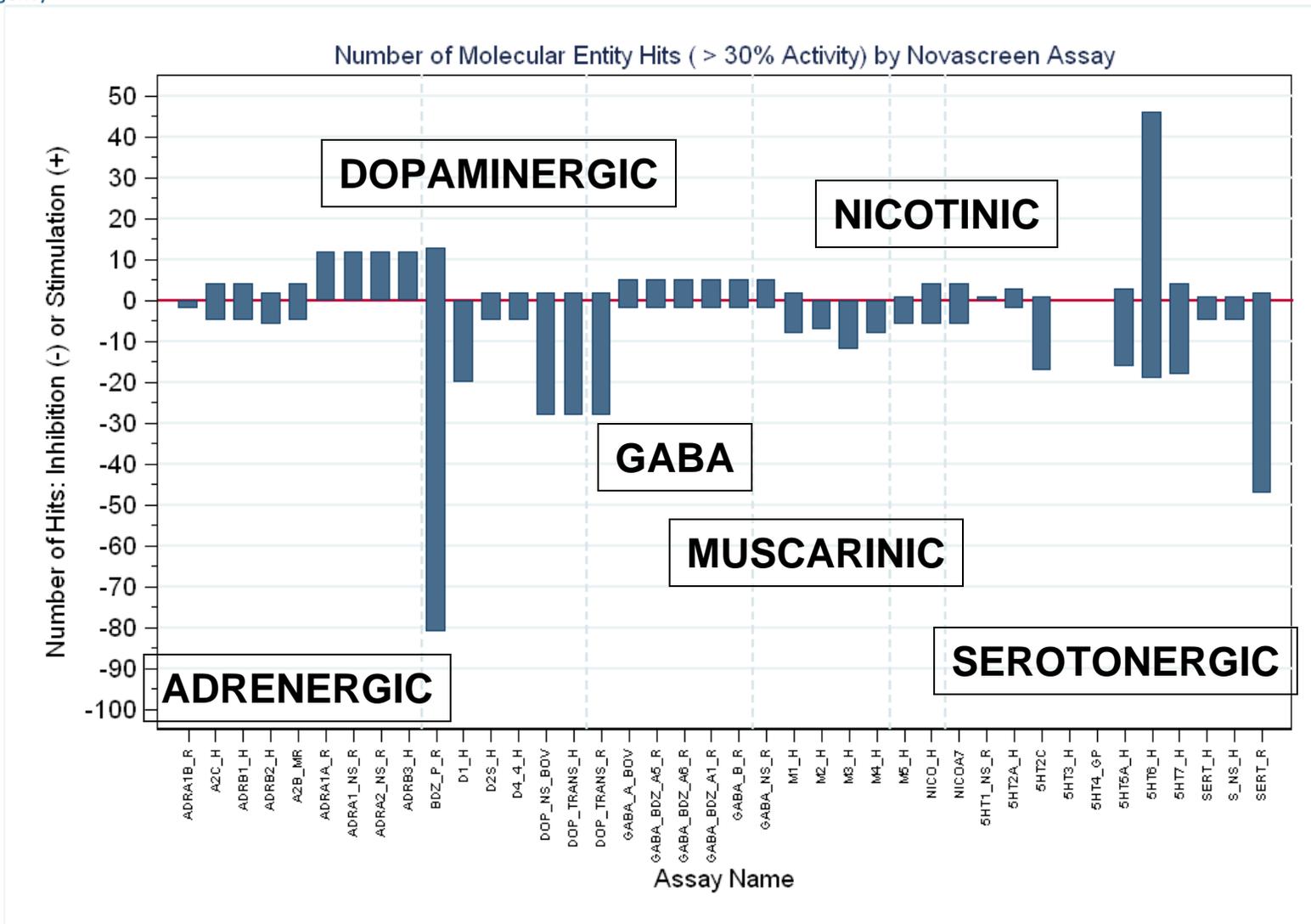


Activity (% of Control)

NovaScreen Descriptive Statistics (30% Cutoff)

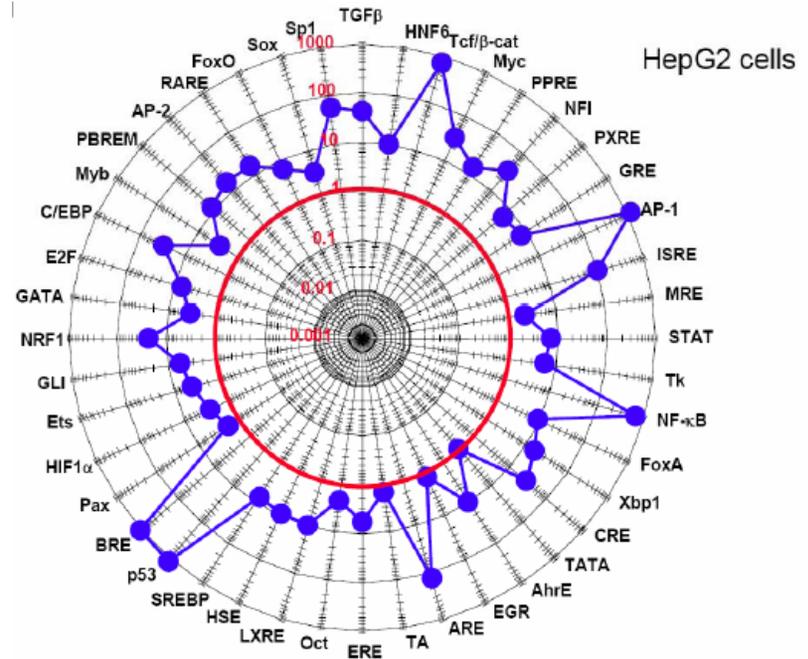
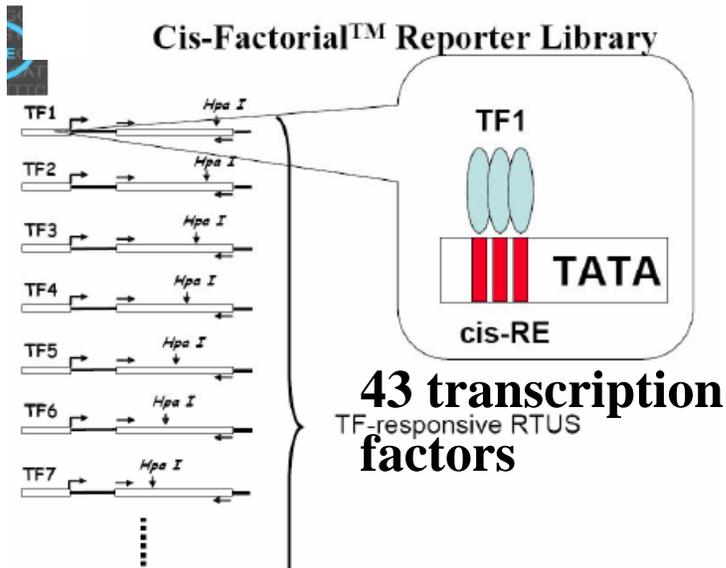


SOME NEUROTRANSMITTER TARGETS

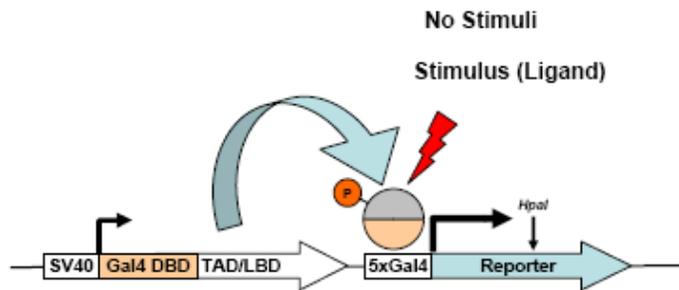


Transcription Factor Activity Profiling

Cis-Factorial™ Biosensors



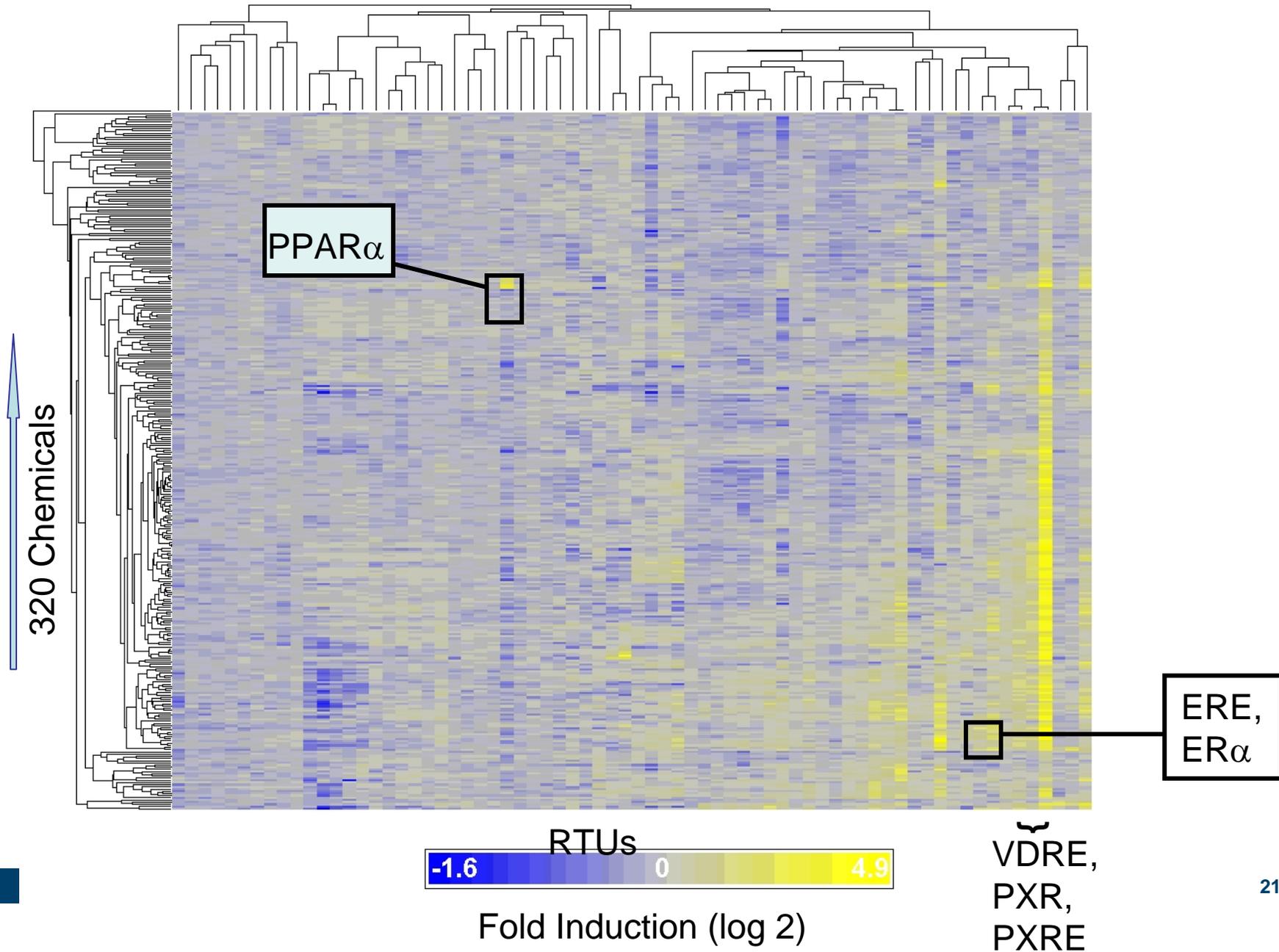
Trans-Factorial™ Biosensors



24 nuclear receptors



Hierarchical Cluster Attagene Results



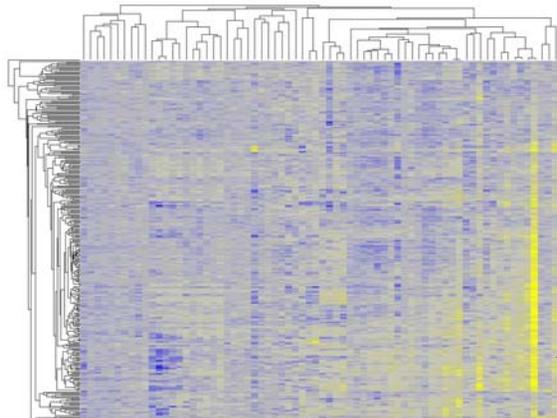
ToxCast Phase I Assays/Datasets/Publications

- **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)**
 - ★ – NR Activation and translocation (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 microarray with metabolism (Solidus)
 - Zebrafish vascular/cardiotoxicity (Zygogen)
 - HTS stress response (NHEERL+NCGC)

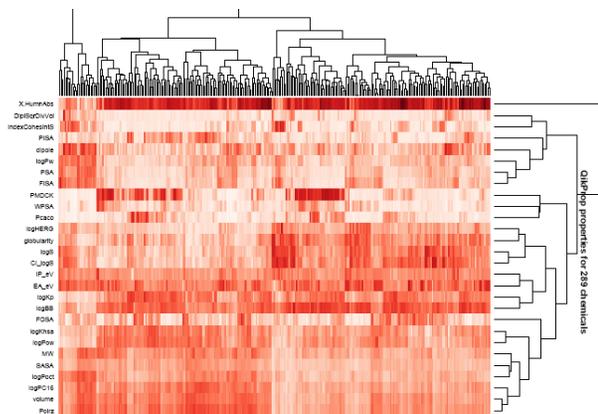
21 Assay Sources
>550 Endpoints

★ Nearing publication

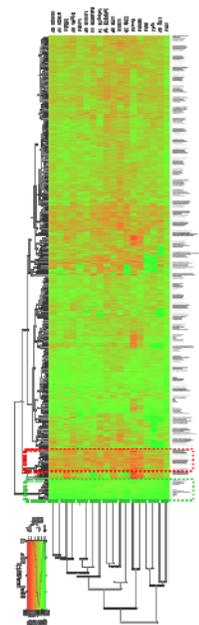
Correlating HTS to Toxicity



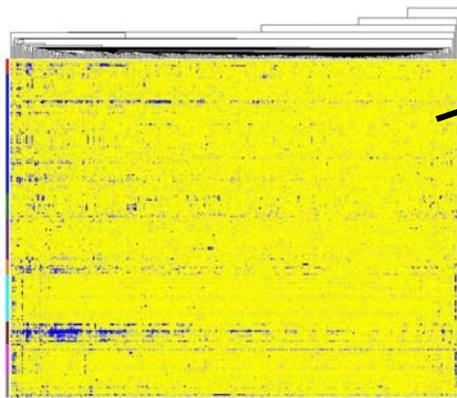
Cellular Assays



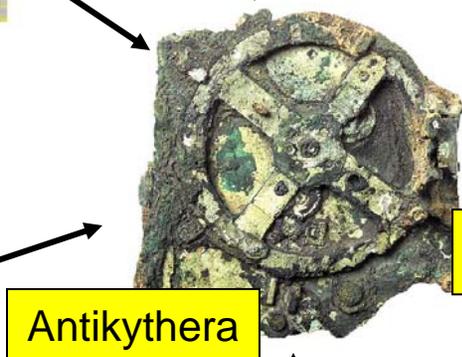
Physical chemical properties



In silico Predictions

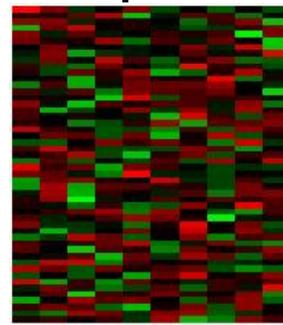


Biochemical Assays

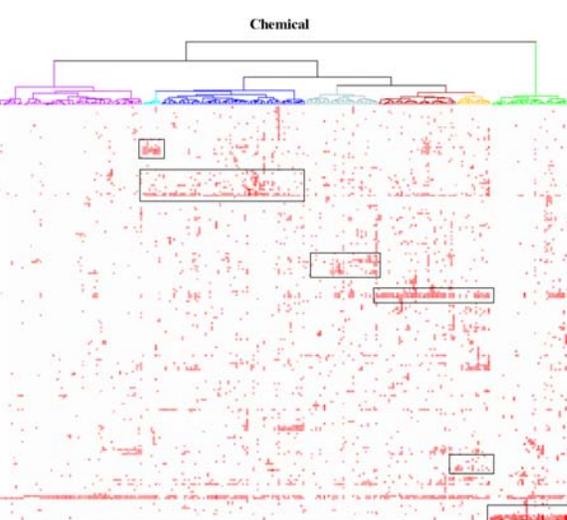


Antikythera

ToxMiner



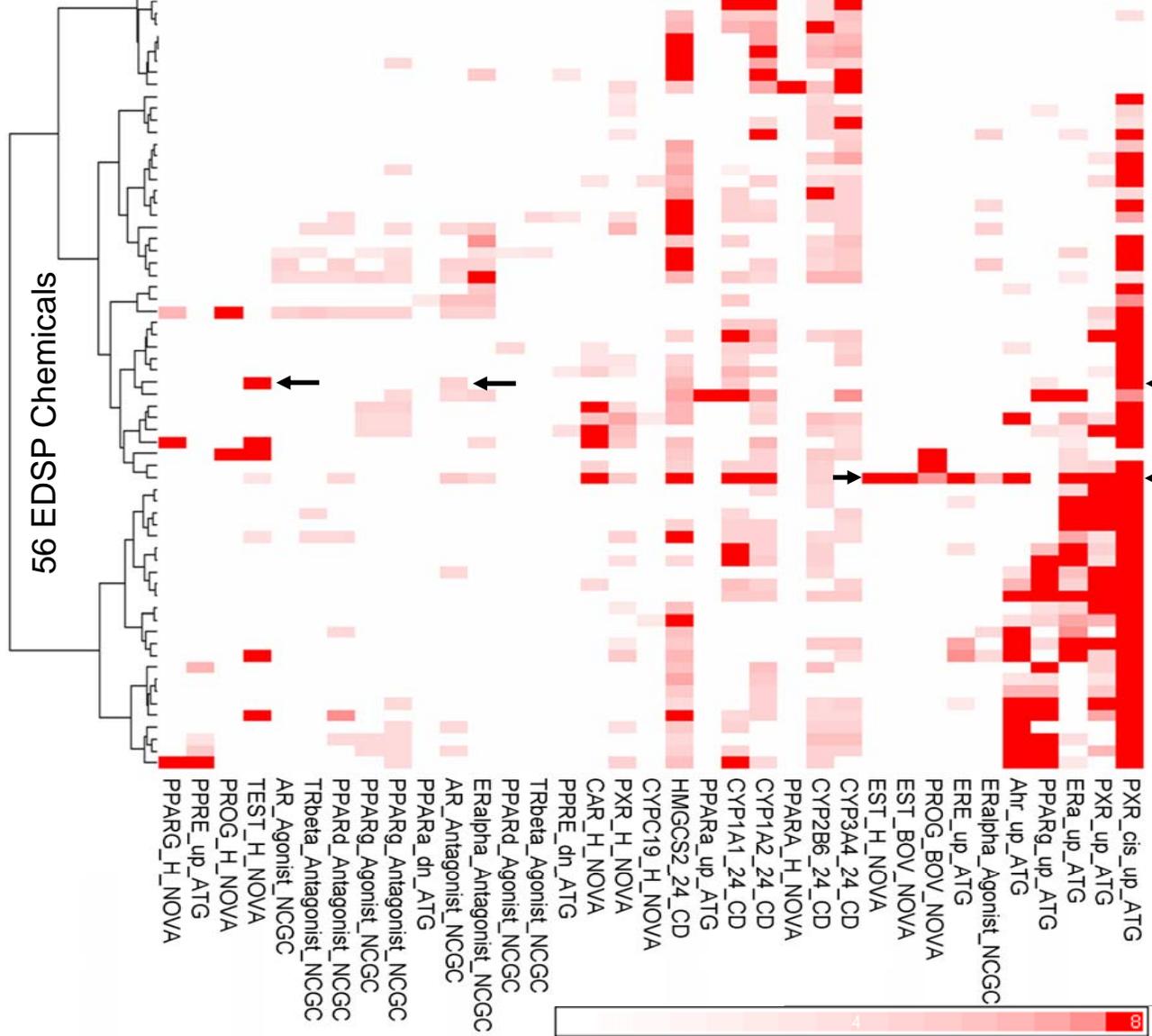
Genomic Signatures



Toxicology Endpoints

ToxCast MoA Profiling – Endocrine Targets

35 ToxCast Assays



ToxCast Predictive Modeling for Toxicity Endpoints

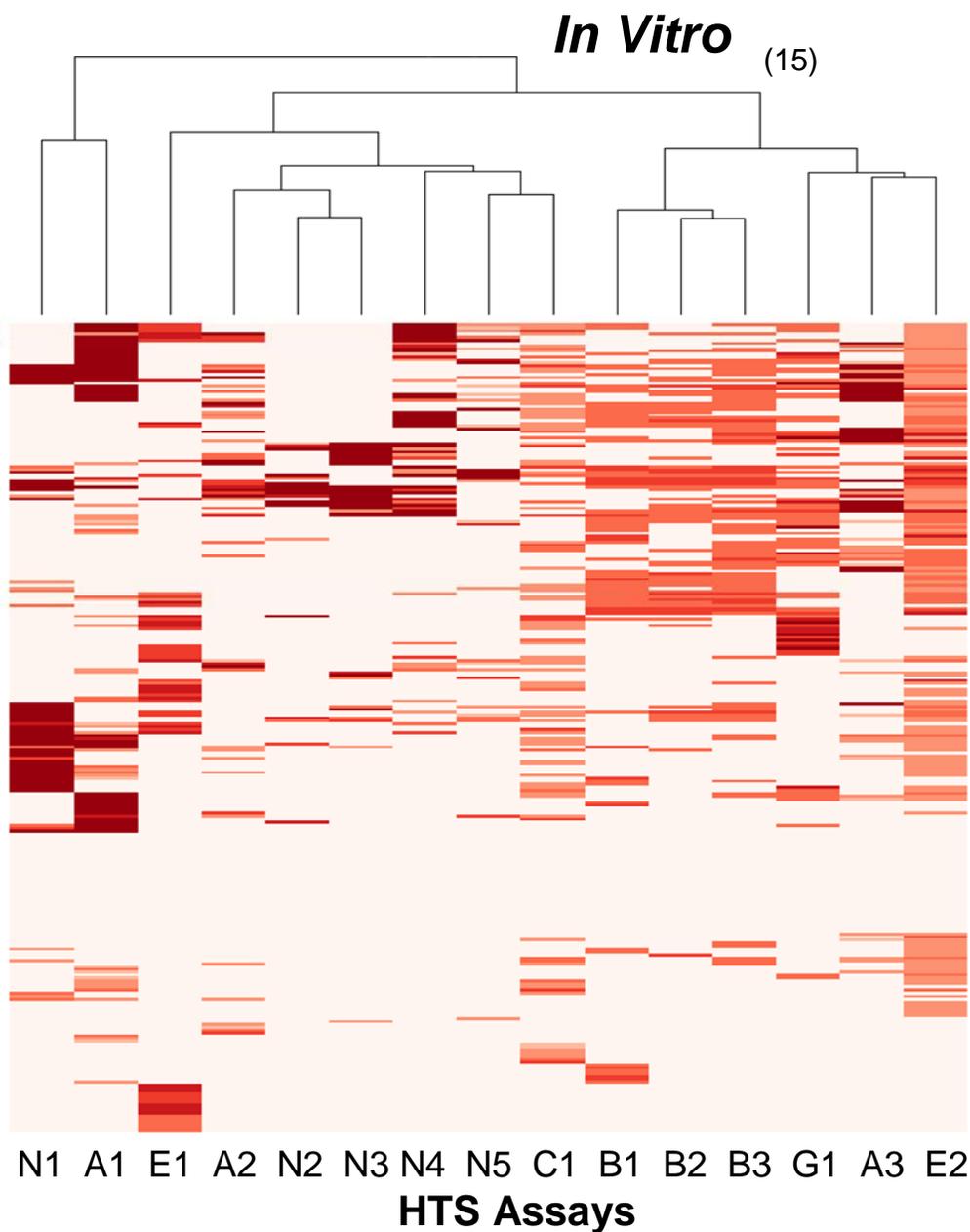
Chronic Rat Liver Apoptosis/Necrosis

In Vivo (23)

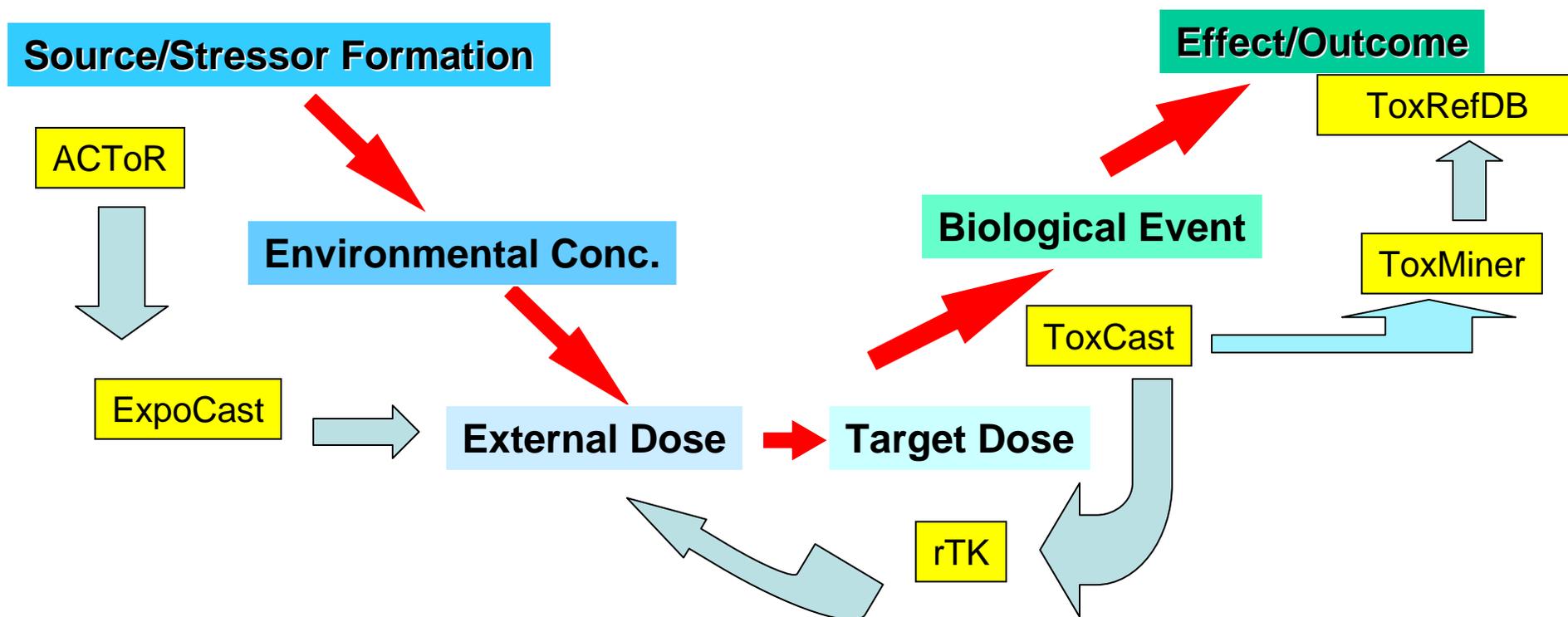
Positive cluster

Negative cluster

Methods 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



A High Throughput Vision for the Source to Outcome Continuum



Summary

- The international community needs better predictive tools for assessing the hazards and risks of chemicals
- It is technically feasible to collect bioactivity data on virtually all chemicals of potential concern
- ToxCast is providing a proof of concept for obtaining predictive, broad-based spectra of bioactivity
- A critical need remains the elucidation of the majority of key biological processes involved in toxic responses
- The time is right to rapidly move this field along



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.

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