

# An Update on ToxCast™

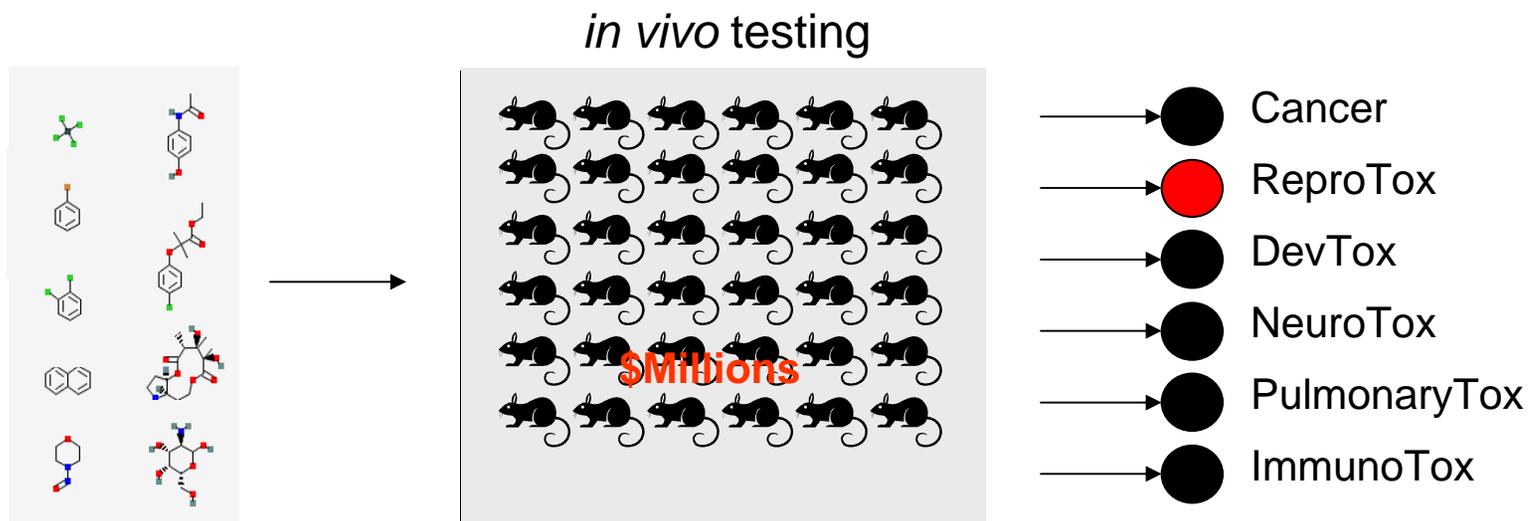
*OECD Molecular Screening Initiative  
Utrecht, The Netherlands*

UNITED STATES ENVIRONMENTAL PROTECTION AGENCY



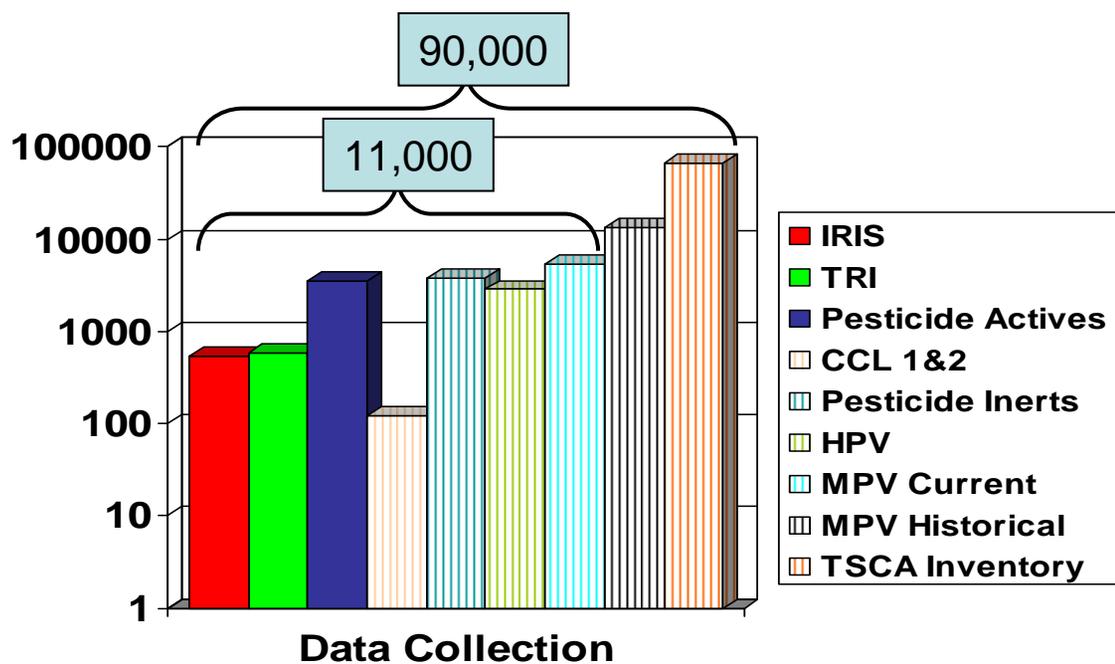
**COMPUTATIONAL  
TOXICOLOGY**

# Current Approach to Toxicity Testing

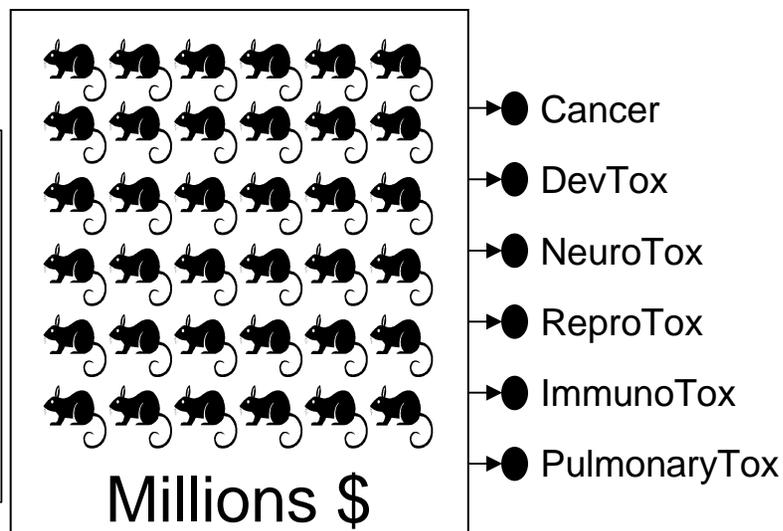


# The Problem

## Too Many Chemicals



## Too High a Cost



...and not enough data.

# Future of Toxicity Testing

## POLICYFORUM

### TOXICOLOGY

## Transforming Environmental Health Protection

Francis S. Collins,<sup>1\*</sup> George M. Gray,<sup>2\*</sup> John R. Bucher<sup>3\*</sup>

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.

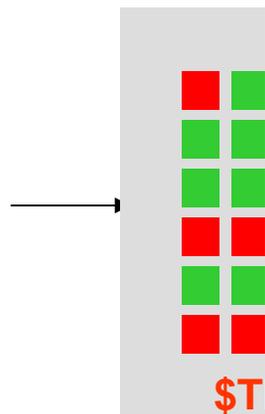
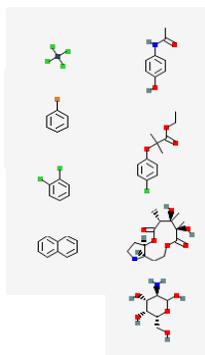
In 2005, the U.S. Environmental Protection Agency (EPA), with support from the U.S. National Toxicology Program (NTP), funded a project at the National Research Council (NRC) to develop a long-range vision for toxicity testing and a strategic plan for implementing that vision. Both agencies wanted future 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-

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 concentra-

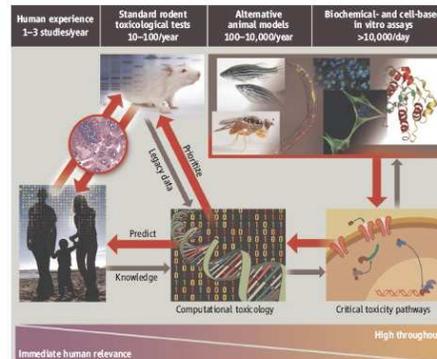
tion, usually between 2 and 10 μM, and tolerate high false-negative rates. In contrast, in the EPA, NCGC, and NTP combined effort, all compounds are tested at as many as 15 concentrations, generally ranging from ~5 nM to ~100 μM, to generate a concentration-response curve (9). This approach is highly 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://ncgc.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,



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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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**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.

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

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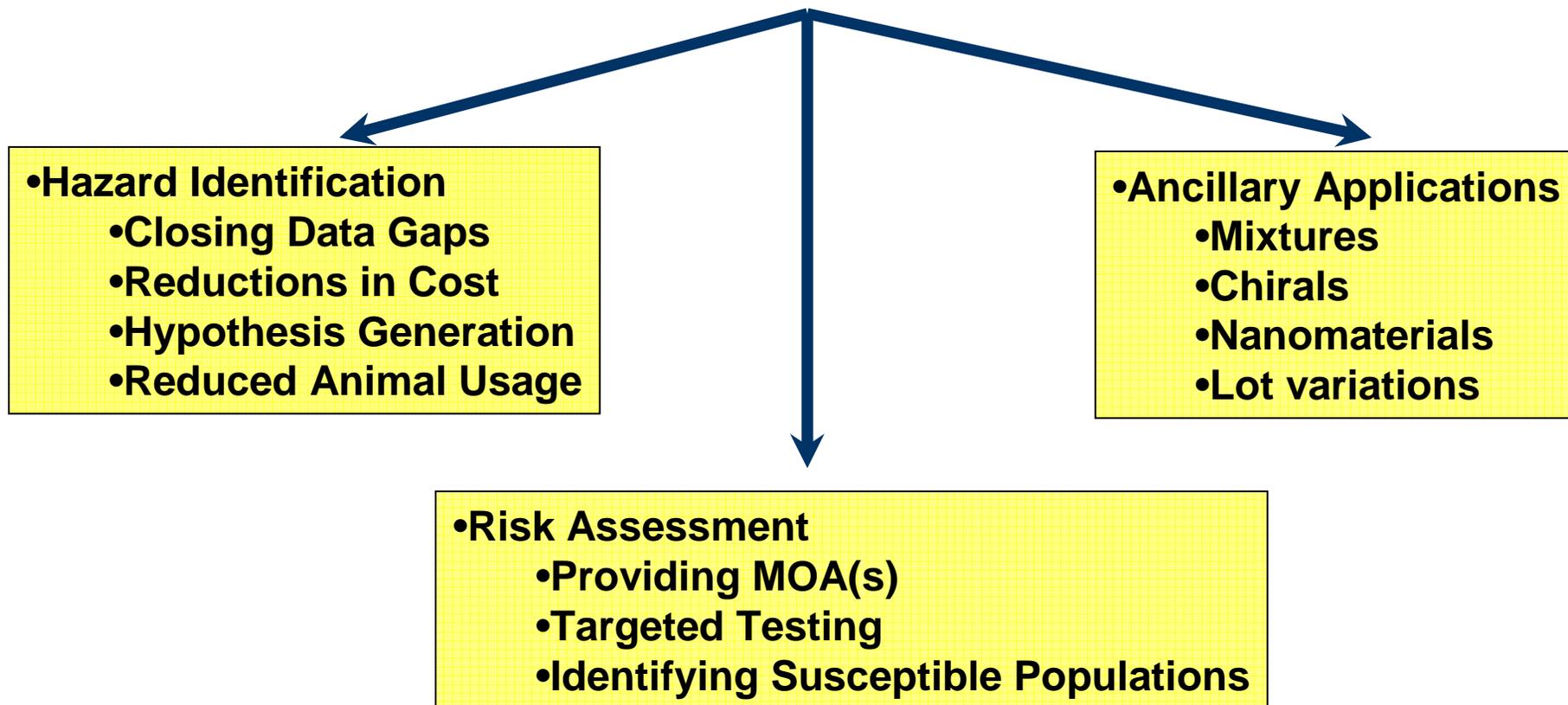


## EPAs Contribution: The ToxCast Research Program

# Key Challenges

- Find the Toxicity Pathways
  - Hepato vs developmental
- 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

# Implications for Success



# 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	>400	\$20k	FY07-08
IIa	>300	Data Rich Chemicals	Validation	>400	\$15-20k	FY09
IIb	>100	Known Human Toxicants	Extrapolation	>400	\$15-20k	FY09
IIc	>300	Expanded Structure and Use Diversity	Extension	>400	\$15-20k	FY10
III	Thousands	Data poor	Prediction and Prioritization	???	\$10-15k	FY11-12

- Affordable science-based system for categorizing chemicals
- Increasing confidence as database grows
- Identifies potential mechanisms of action
- Refines and reduces animal use for hazard ID and risk assessment

# Opportunities for Partnerships

- Phase I
  - Testing the ToxCast 320 in new assays
  - Targeted Testing of HTS results
- Phase II
  - Nominating chemicals in conjunction with providing standard toxicological testing results
  - Nominating biological pathways and assays
  - CRADA partnerships to accelerate depth and breadth
  - Targeted testing on HTS results
- Providing HTS assays that can be run at the NCGC
- Helping to create curated, public access databases of toxicity information



## Some Lessons Learned to Date

- Large amounts of quality HTS data can be economically obtained
- Large scale data sets will be required to understand potential for biological activity
- Value in having multiple assays with overlapping coverage of biological pathways and a variety of methodologies
- Concentration-response will be important for ultimate interpretation
- Data transparency will be important for acceptance
- Metabolic capabilities and coverage of developmental toxicity pathways will need additional attention
- Need to define the gold standard
- Partnerships are needed to bring critical mass and expertise

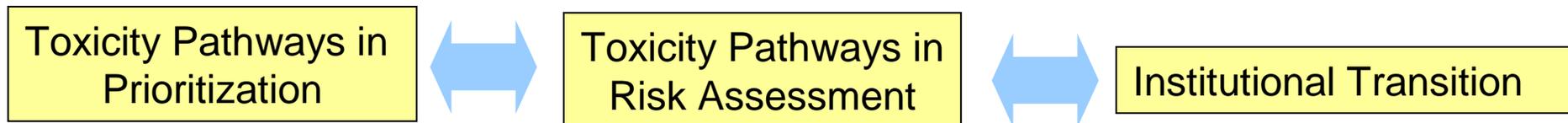
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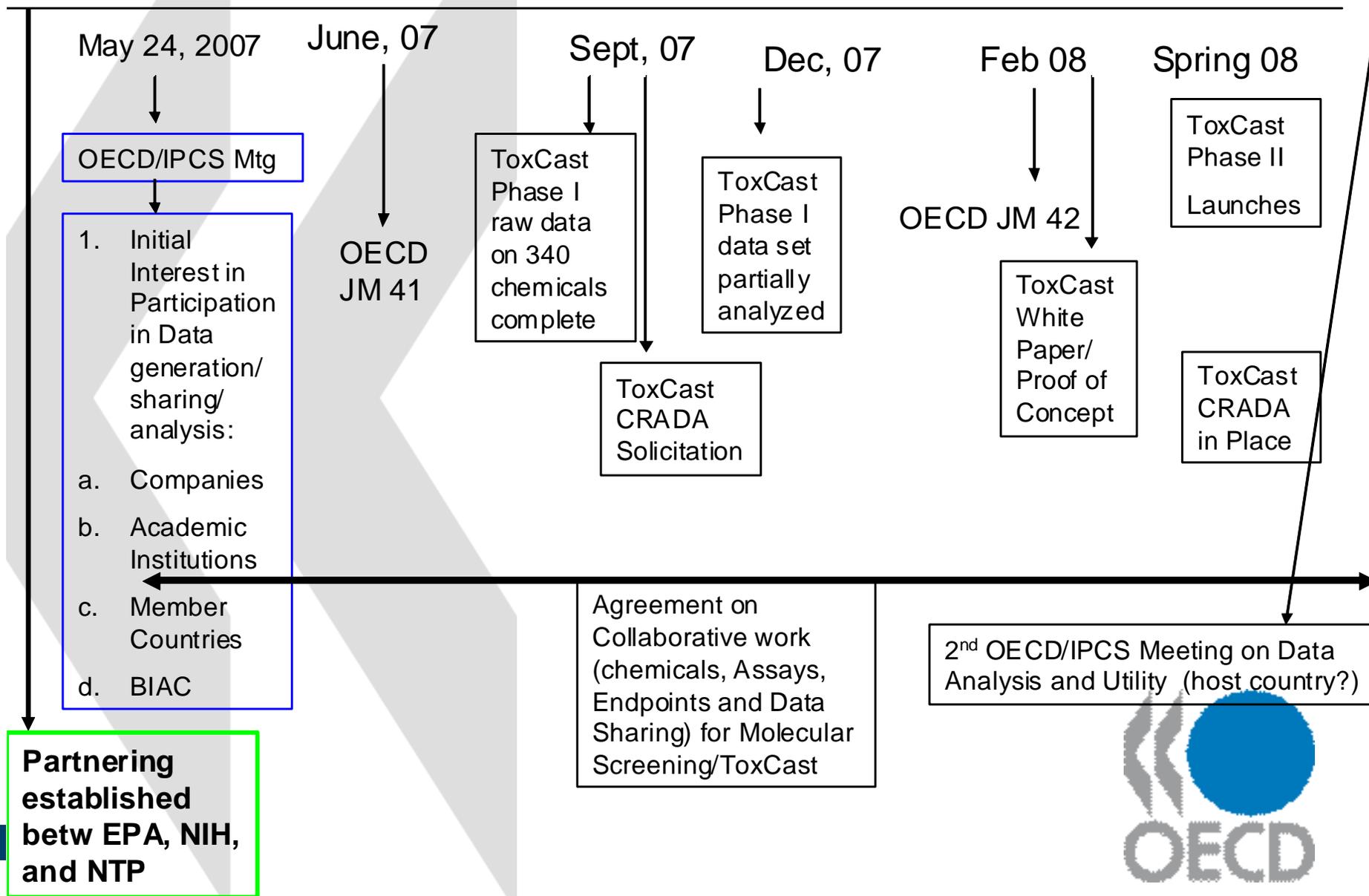


# Moving Forward with ToxCast

- Completion of Data Acquisition and Data Mining for Phase I
- Publication and Public Release of all Data
- Data Summit, Fall/Winter 2008
- Tox21 MOU partnership with NTP/NIEHS and NCCG/NIH/CDI
  - Four Working Groups
  - Total of ~7000 chemicals for screening
  - Subset to feed Phase II of ToxCast
- Communities of Practice – Prioritization (Dix), Exposure (Hubal)
- Launch Phase II in FY09
- EPA Research Strategy



## Action Line from May 2007



# Evolution of Phase I

- **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 (March, 2008)**
  - Organ culture: liver, kidney, lung (Hamner Institutes)
  - HTS Genotoxicity (Gentronix)
  - Toxicity and signaling pathways (Invitrogen)
  - NR Activation and translocation (CellzDirect)
  - 3D Cellular microarray with metabolism (Solidus)
  - C. elegans (NIEHS)
  - Functional markers from microscale cultured hepatocytes (MIT)

**9 Assay Sources  
& 412 Endpoints**

**+3 Assay Sources  
& 16 Endpoints**

**+7 Assay Sources  
& 123 Endpoints**

**19 Assay Sources, 551 Endpoints**





# Comparing Activities by Chemical Class

Conazole  
Fungicides vs.  
NovaScreen  
Assays

NAME	CYP2C19	CYP2C9	CYP3A1	Dopamine Transporter (Human)	CYP2D2	Androgen Receptor	Dopamine Transporter (Rat)	CYP2B6	CYP2D1	CYP3A4	Progesterone Receptor	Benzodiazepine Receptor
Cyproconazole	1	1	1	1	1	0	1	0	0	1	0	0
Difenoconazole	1	1	1	1	1	0	0	1	1	0	0	0
Diniconazole	1	1	1	0	1	0	0	0	1	1	1	0
Fenbuconazole	1	1	0	0	0	0	0	0	0	1	0	0
Flusilazole	1	1	1	0	1	1	0	1	1	NA	1	1
Hexaconazole	1	1	1	1	1	0	1	1	1	NA	1	0
Imazalil	1	1	1	1	1	1	1	1	1	1	1	1
Myclobutanil	1	1	1	1	0	0	0	0	0	NA	0	0
Paclobutrazol	1	0	1	1	0	1	1	0	1	1	0	0
Prochloraz	1	1	1	1	1	1	1	1	1	NA	1	1
Propiconazole	1	1	1	0	0	0	0	1	0	NA	0	1
Tetraconazole	1	1	1	0	1	1	0	1	0	1	1	0
Triadimefon	1	1	0	1	1	1	1	0	0	1	0	1
Triadimenol	1	0	0	1	0	1	1	0	0	0	0	0
Triflumizole	1	1	1	1	1	1	0	1	1	1	1	1
Triticonazole	1	1	1	1	0	1	1	0	0	NA	0	0
Totals	16	14	13	11	10	9	8	8	8	8	7	6

## ACToR: Aggregated Computational Toxicology Resource

**ACToR: Aggregated Computational Toxicology Resource**

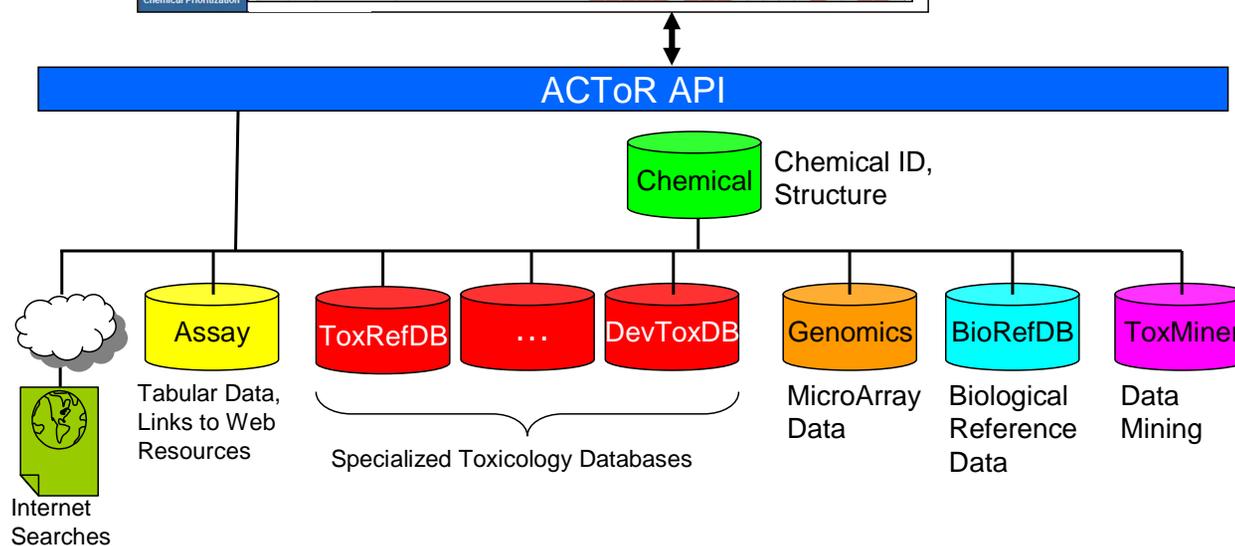
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You are here: [Home](#) > [ACToR](#) > Data Collection

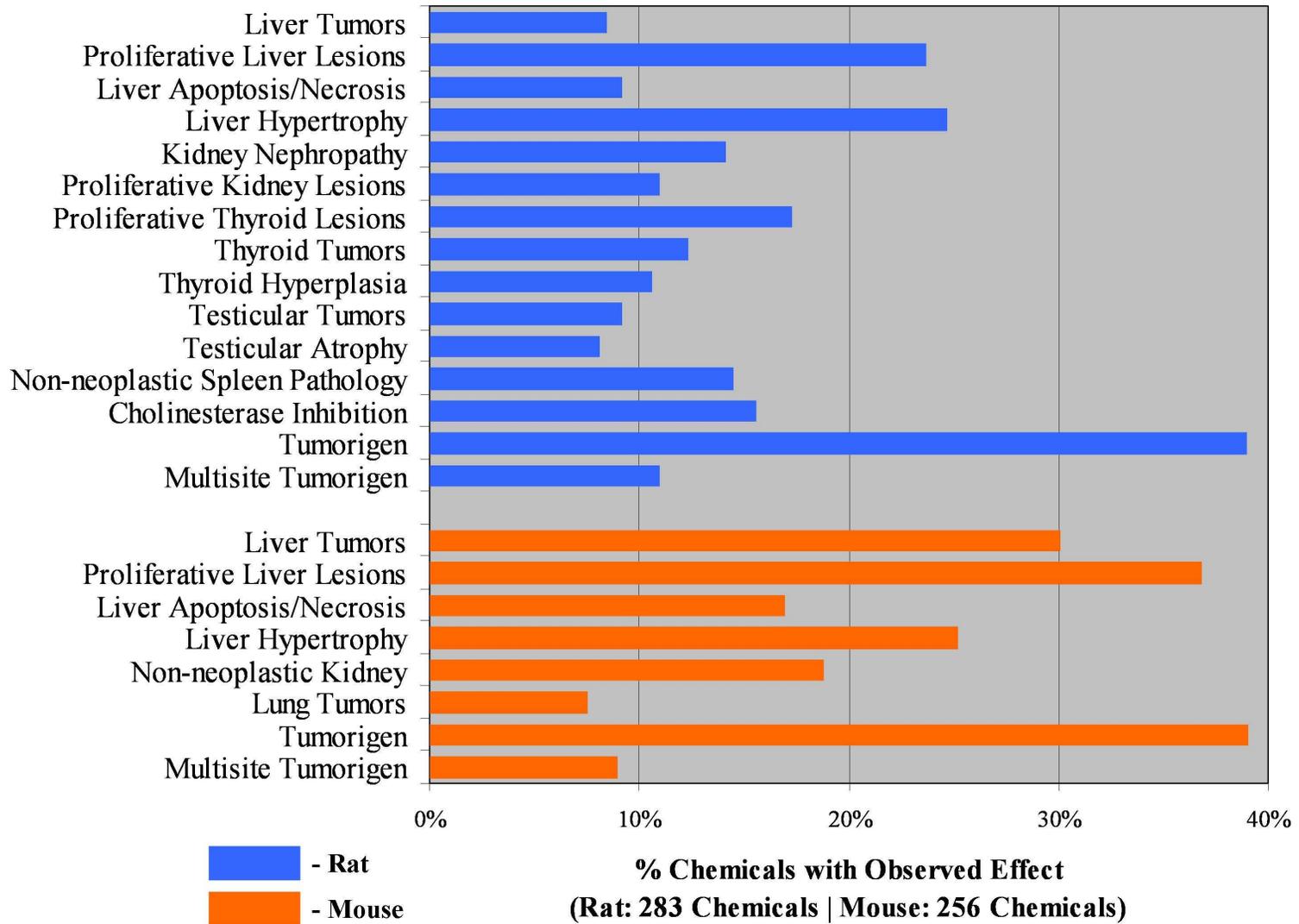
**Data Collection: ToxCast\_320**

SCID	GCID	CASRN	Name	Hazard	SubchronicTox	ChronicTox	CardiogenTox	DevTox	ReproTox	NeuroTox	ImmunTox	DermatTox	RespiratoryTox	HepatTox	Endocrine	CardioTox	FoodSafe	ToxOther
<a href="#">12622</a>	<a href="#">447</a>	94-75-7	2,4-D	11	6	1	7	16	23	8	4	3	2			1	1	7
<a href="#">12623</a>	<a href="#">6424</a>	94-82-6	2,4-DB	5	4	1	4	8	7	6	5	2			1			2
<a href="#">12624</a>	<a href="#">7712</a>	136-45-8	2,5-Pyridinedicarboxylic acid, dipropyl ester	3	1	1	1	5	2	1								1
<a href="#">12625</a>	<a href="#">1174</a>	90-43-7	2-Phenylphenol	6	2	1	2	10	1	3	2	1				1	1	2
<a href="#">12626</a>	<a href="#">4555</a>	55406-53-6	3-Iodo-2-propynylbutylcarbamate	6	2	1	2	3	3	2	2							1
<a href="#">12627</a>	<a href="#">4555</a>	55406-53-6	3-Iodo-2-propynylbutylcarbamate	6	2	1	2	3	3	2	2							1

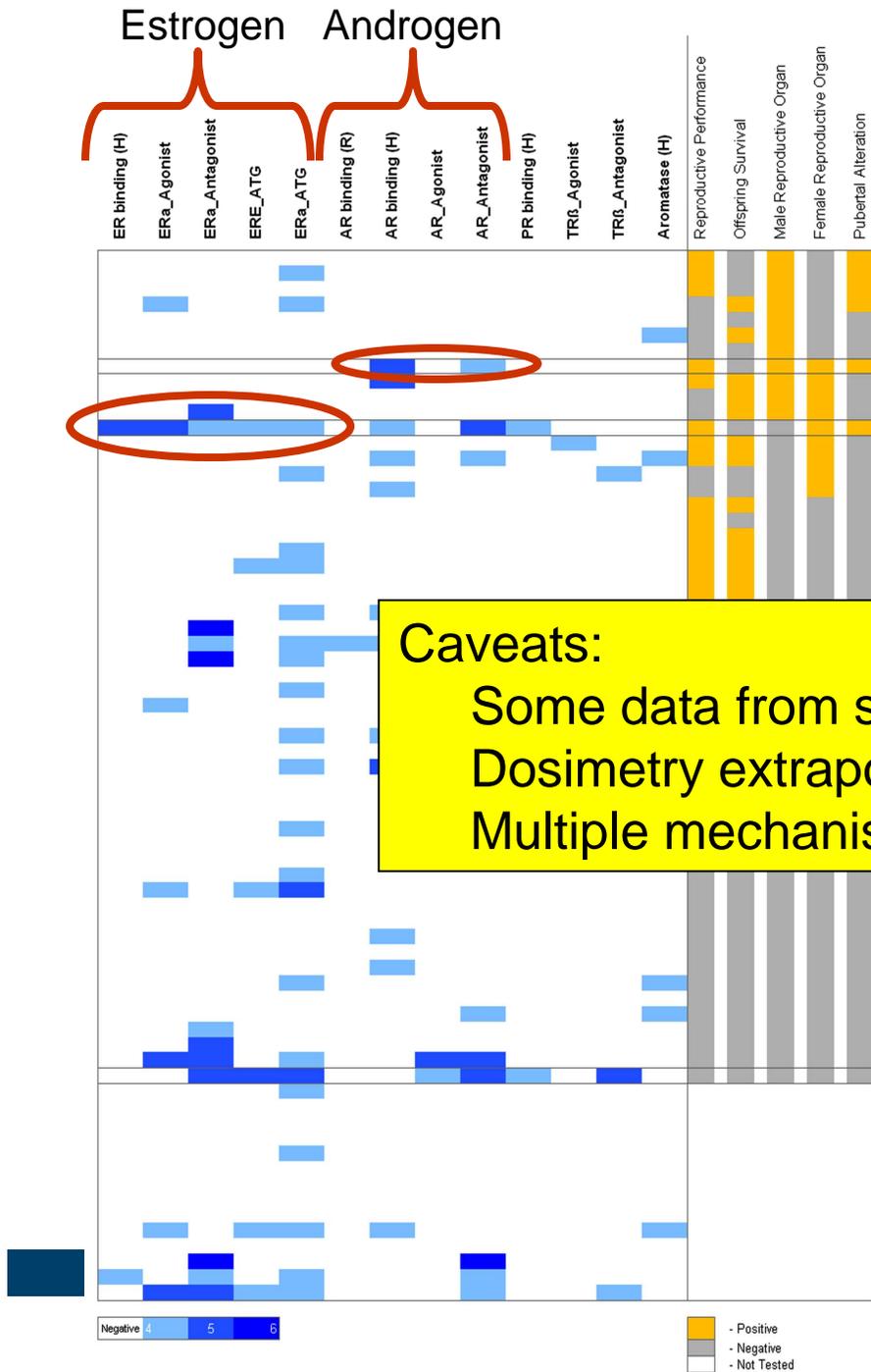
ACToR Web  
Browser



# Common Phenotypes in Chronic Rodent Studies



# ToxCast Endocrine Profiling



HTS Data from receptor binding (Novascreen), single gene reporter (NCGC)

Priority Chemicals contained in the ToxCast 320

## The ToxCast Team





## National Center for Computational Toxicology

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The EPA Web site will be unavailable on Sunday, March 2, 2008 from 8:00 pm until 10:00 pm ET.

## 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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- [ToxCast™ Assays](#)
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# Assay Read Across for EDSP Chemicals (beta version)

