

ToxCast: Developing Predictive Signatures of Chemically Induced Toxicity

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UNITED STATES ENVIRONMENTAL PROTECTION AGENCY

Office of Research and Development National Center for Computational Toxicology

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ized 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-

POLICYFORUM

Health Protection

funded a project at the National Research

implementing that vision. Both agencies

wanted future toxicity testing and assessment

paradigms to meet evolving regulatory needs.

stances that need to be tested and how to incor-

ogy, computational sciences, and information

technology; to rely increasingly on human as

opposed to animal data; and to offer increased

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 assess-

ment of chemical hazard and risk (6, 7). Although the NRC reports have laid out a solid

theoretical rationale, comprehensive and rig-

orously gathered data (and comparisons with historical animal data) will determine whether the hypothesized improvements will be real

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Transforming Environmental

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n 2005, the U.S. Environmental Protection throughput screening (HTS) and other auto-Agency (EPA), with support from the U.S. mated screening assays into its testing at high false-negative rates. In contrast, in National Toxicology Program (NTP), program. In 2005, the EPA established the the EPA, NCGC, and NTP combined effort, National Center for Computational Toxi-Council (NRC) to develop a long-range vision cology (NCCT). Through these initiatives, for toxicity testing and a strategic plan for 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 Challenges include the large numbers of sub- predominantly predictive science focused the traditional HTS methods (9), and facilion broad inclusion of target-specific, mechtates multiassay comparisons. Finally, an porate recent advances in molecular toxicol- anism-based, biological observations in informatics platform has been built to comvitro (1, 4) (see figure, below). pare results among HTS screens; this is Toxicity pathways. In vitro and in vivo tools are being used to identify cellular

being expanded to allow comparisons with historical toxicologic NTP and EPA data efficiency in design and costs (I-5). In response after chemical exposure expected to result in adverse health effects (7). HTS (http://ncgc.nih.gov/pub/openhts). HTS data collected by EPA and NTP, as well as by the NCGC and other Molecular Libraries methods are a primary means of discovery for drug development, and screening of >100,000 compounds per day is routine (8). being made publicly available through Web-However, drug-discovery HTS methods trabased databases [e.g., PubChem (http:// ditionally test compounds at one concentra- pubchem.ncbi.nlm.nih.gov)]. In addition,

Future of Toxicity Testing

for toxicity assessments

We propose a shift from primarily in vivo animal

studies to in vitro assays, in vivo assays with ower organisms, and computational modeling



icology 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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tion, usually between 2 and 10 µM, and tolerall 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





EPAs Contribution: The ToxCast Research Program

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www.epa.gov/ncct/toxcast

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Change Needed Because

Too Many Chemicals

Too High a Cost



...and not enough data.

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Judson, et al EHP submitted



High-Throughput Screening Assays

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



National Center for Computational Toxicology



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://134.67.216.45:22722/servlet/ActorPrototype2008Q1?page=0 http://www.epa.gov/actor/





Implications for Success

•Hazard Identification •Closing Data Gaps •Reductions in Cost •Hypothesis Generation •Reduced Animal Usage

•Ancillary Applications •Mixtures •Chirals •Nanomaterials •Green Chemistry •Lot variations

•Risk Assessment •Providing MOA(s) •Targeted Testing •Identifying Susceptible Populations





•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



Phased Development of ToxCast

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

¹Initiated April 2007 7



The ToxCast_320





Physical-Chemical Properties





\$1B in Toxicology Now Stored in ToxRefDB



Martin, et al *EHP*, *submitted* 10



ToxRefDB Endpoints from Chronic Rodent Studies for Training ToxCast Predictions



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Transcription Factor Activity Profiling

Cis-FactorialTM Biosensors







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24 nuclear receptors



ToxCast Phase I Assays/Datasets/Publications

- ToxCast 1.0 (April, 2007)
 - Enzyme inhibition/receptor binding HTS (Novascreen)
- MR/transcription factors (Attagene, NCGC)
- 🛨 Cellular impedance (ACEA)
- \star Complex cell interactions (BioSeek)
- ★ Hepatocelluar 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)
- \star 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)
- \star 3D Cellular microarray with metabolism (Solidus)
 - Zebrafish vascular/cardiotoxicity (Zygogen)
 - HTS stress response (NHEERL+NCGC)

21 Assay Sources >550 Endpoints







Descriptive Profiling of Endocrine Activity

A number of ToxCast assays examine the ability of a chemical to interact with hormone systems, including binding to hormone receptors and activating, or inhibiting, hormonally sensitive genes. Here is the profile for 56 of the 73 proposed EDSP priority chemicals in a number of relevant assays.

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Predictive (Meta) Analysis of HTS Results Against a Liver Phenotype

Rat Liver Apoptosis Necrosis

ACEA_LOCinc ATG_RARa_up

CYP3A4_H CYP2A2_R

CYP1A2_H

ATG_PPARg_up

DOP_TRANS_H SCRTSE_B_H

MitoFunctionl_1 BioSeek26 BioSeek20 BioSeek63 AR_Antagonist ATG_BRE_up

ACEA_LOCdec

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- 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



Activity (% of Control)

201 Assays



NovaScreen Descriptive Statistics (30% Cutoff)





Real-time Cell Electronic Sensing (RT-CES) Assay

Judson et al., in preparation



Figure 2 – Cell growth curve for Diquat Dibromide.

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Figure 7 – Comparison of average growth rate values for replicates chemicals across all concentrations.

Sepan ToxCast Website: www.epa.gov/ncct/toxcast

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 > ToxCast™ Program

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[™].

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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.