

Development and Validation of New Technologies in Drug Discovery and Toxicology

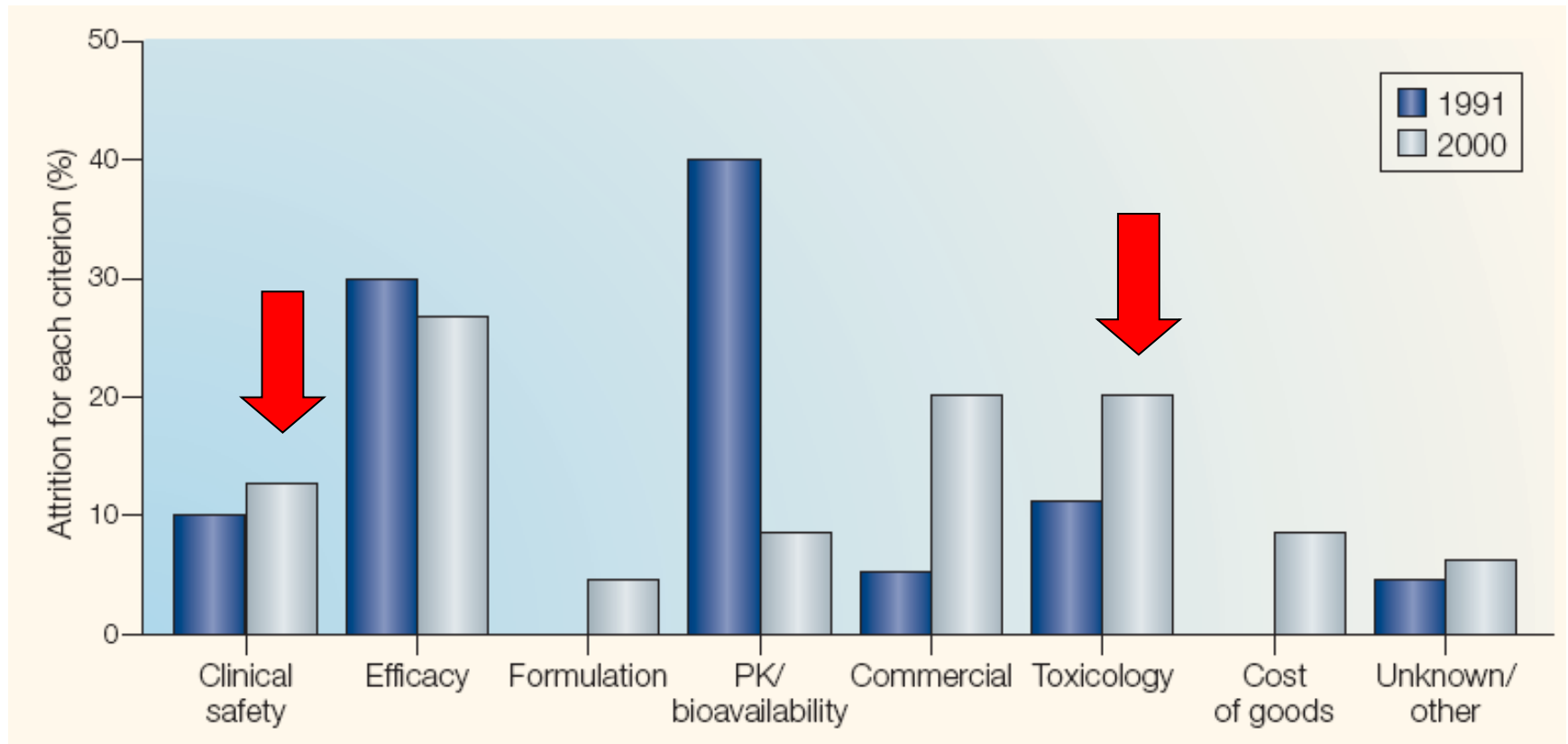


Christopher P. Austin, M.D.
Director, Division of Preclinical Innovation
National Center for Advancing Translational Sciences
National Institutes of Health

*AIMBE/NIH Summit on Validation and Qualification of New In Vitro Tools
March 19, 2012*

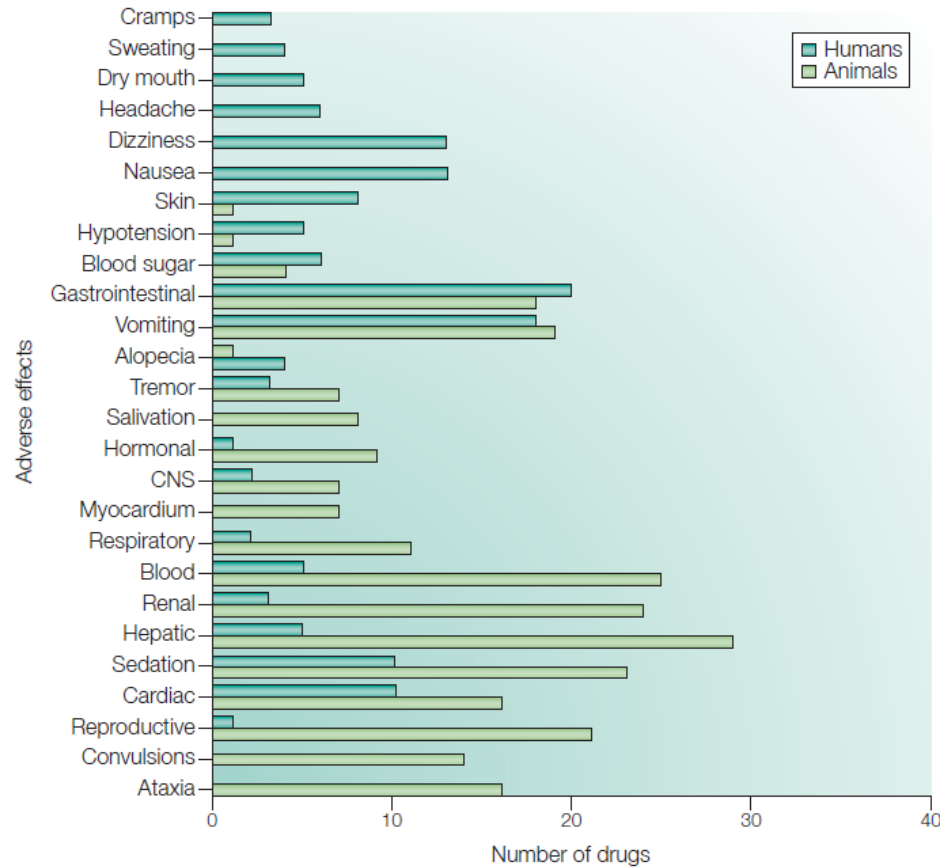
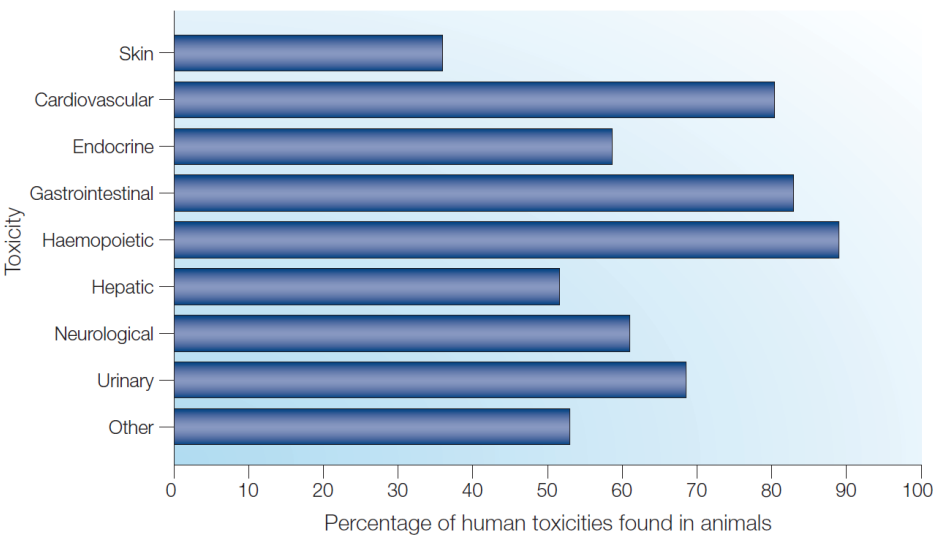


Toxicity is the Most Common Reason for Drug Development Failure

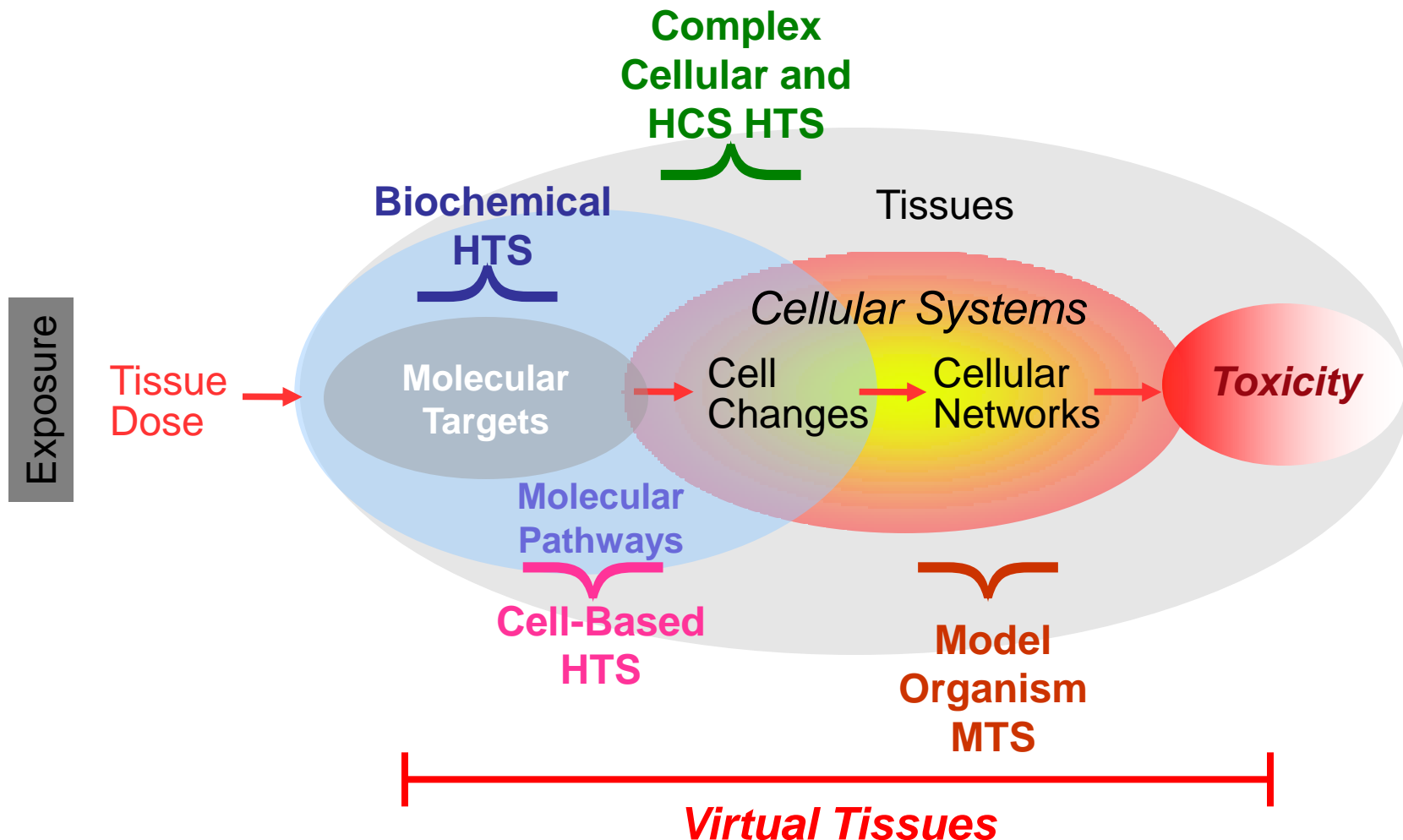


Preclinical (21%) + Clinical (12%) Tox = 33% of all failures

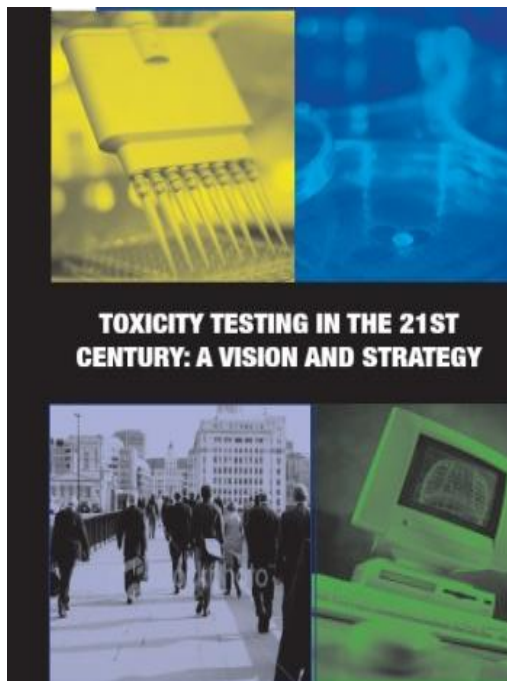
Poor concordance of human and animal drug toxicities



Grand Challenge: Predicting Toxicity



NAS "Toxicology in the 21st Century"



"This 2007 National Academy of Science report envisions a not-so-distant future in which virtually all routine toxicity testing would be conducted in vitro in human cells or cell lines by evaluating perturbations of cellular responses in a suite of toxicity pathway assays using high throughput robotic assisted methodologies."

POLICYFORUM

TOXICOLOGY

Transforming Environmental Health Protection

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

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.

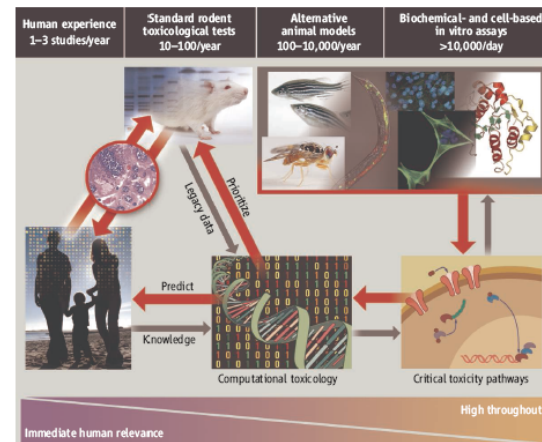
EPA, NCGC, and NTP Joint Activities
In 2004, the NTP released its vision and roadmap for the 21st century (1), which established initiatives to integrate high-

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 vivo (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-

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.

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 multiassay 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,



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.

¹Director, National Human Genome Research Institute (NHGRI), National Institutes of Health, Bethesda, MD 20892; ²Assistant Administrator for the Office of Research and Development, U.S. Environmental Protection Agency, Washington, DC 20460; ³Associate Director, U.S. National Toxicology Program, National Institute of Environmental Health Sciences (NIEHS), Research Triangle Park, NC 27709, USA.

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

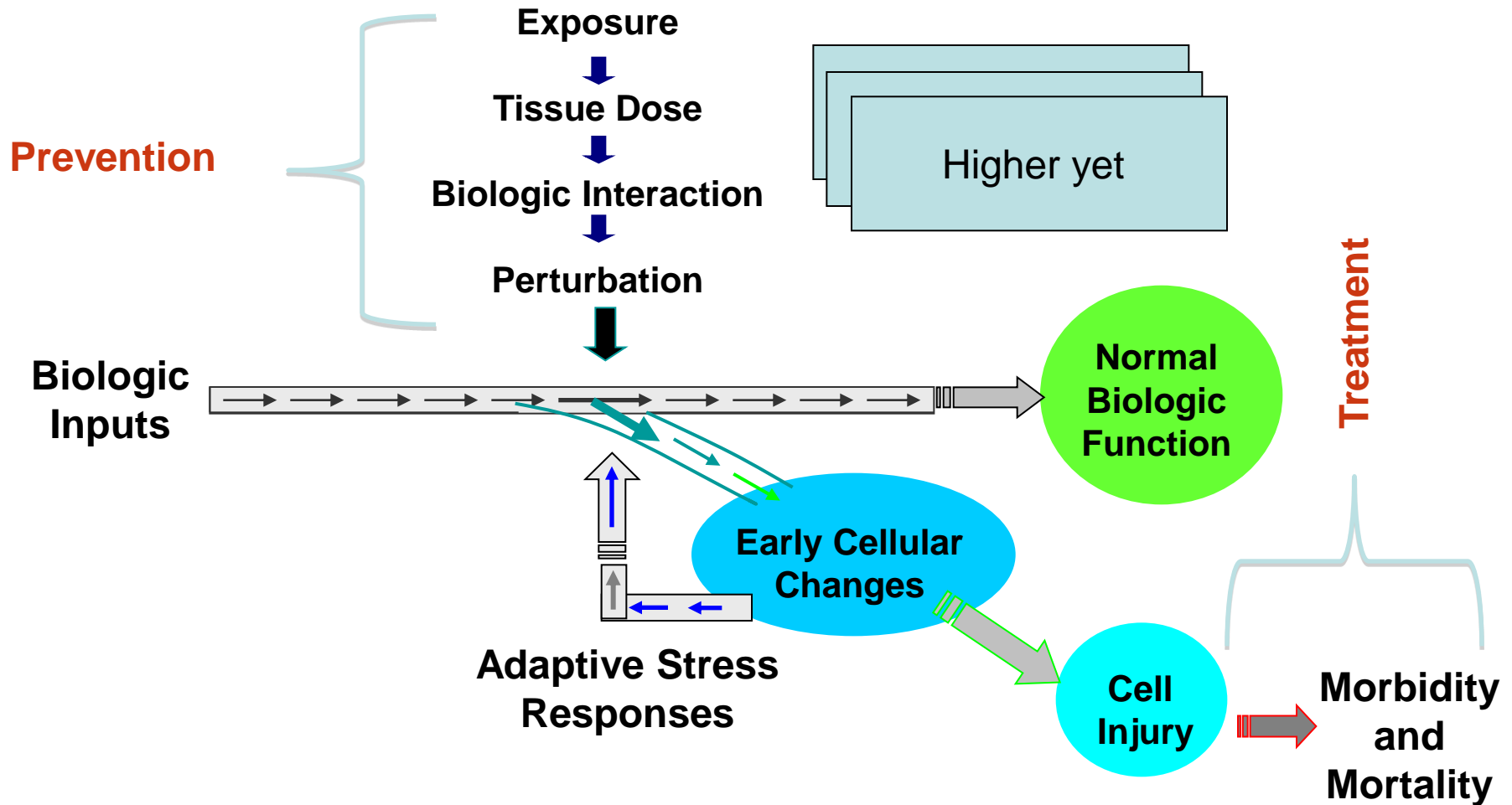
†Author for correspondence. E-mail: francis@mail.nih.gov

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U.S. NATIONAL INSTITUTE OF ENVIRONMENTAL HEALTH SCIENCES, NATIONAL INSTITUTES OF HEALTH



Activation of a Toxicity Pathway

NAS Report, 2007



The Tox21 Community



National Toxicology Program
Department of Health and Human Services



NIEHS
National Institute of
Environmental Health Sciences



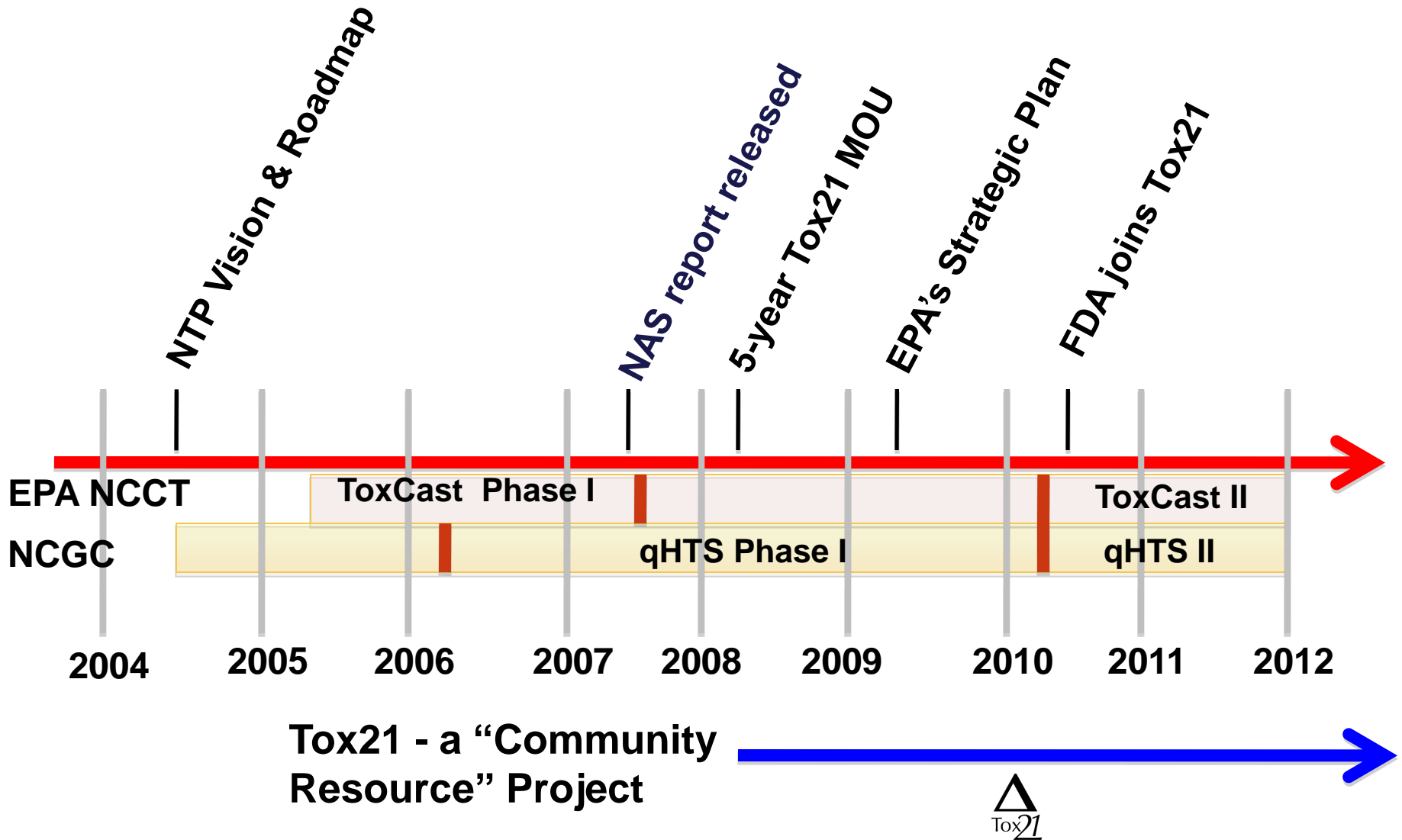
NIH CHEMICAL GENOMICS CENTER



National Institutes of Health
**NATIONAL CENTER FOR ADVANCING
TRANSLATIONAL SCIENCES**



Tox21 Timeline



Tox21 Partners have Complementary Expertises

Areas of Expertise	NIEHS/NTP	NCGC	EPA	FDA
Experimental Toxicology	✓		✓	✓
Human Toxicology				✓
qHTS		✓		
Epigenetics	✓	✓		
Low to Mid Throughput Assays	✓	✓	✓	✓
Lower Organism Systems	<i>C. elegans</i>		Zebrafish	Zebrafish/ <i>C. elegans</i>
<i>In Vitro</i> 3-D Model Systems	✓		✓	✓
Genetic Variability in Response	✓	✓		
Computational Toxicology	✓	✓	✓	✓
Human Exposure Assessment	✓		✓	
Validation Experience	✓		✓	✓

Operational Structure



Agency Points of Contact

Christopher Austin, M.D. (NCGC)
Tom Colatsky Ph.D. (FDA)
Robert Kavlock, Ph.D. (EPA)
Raymond Tice, Ph.D. (NTP)

Assays & Pathways Working Group

Co-Chairs

Kevin Gaido, Ph.D. (FDA)
Keith Houck, Ph.D. (EPA)
Kristine Witt, M.S. (NTP)
Menghang Xia, Ph.D. (NCGC)

- Identify toxicity pathways & corresponding assays
- Review nominated assays
- Prioritize assays for qHTS

Chemical Selection Working Group

Co-Chairs

William Leister, Ph.D. (NCGC)
Donna Mendrick, Ph.D. (FDA)
Ann Richard, Ph.D. (EPA)
Cynthia Smith, Ph.D. (NTP)

- Establish a 10K DMSO soluble compound library for qHTS
- Establish QC procedures
- Establish libraries of mixtures and aqueous soluble compounds for qHTS

Informatics Working Group

Co-Chairs

Ruili Huang, Ph.D. (NCGC)
Richard Judson, Ph.D. (EPA)
Jennifer Fostel, Ph.D. (NIEHS)
Weida Tong, Ph.D. (FDA)

- Characterize assay output and evaluate assay performance
- Develop prioritization schemes and prediction models
- Make all data publicly accessible via CEBS, PubChem, ACToR

Targeted Testing Working Group

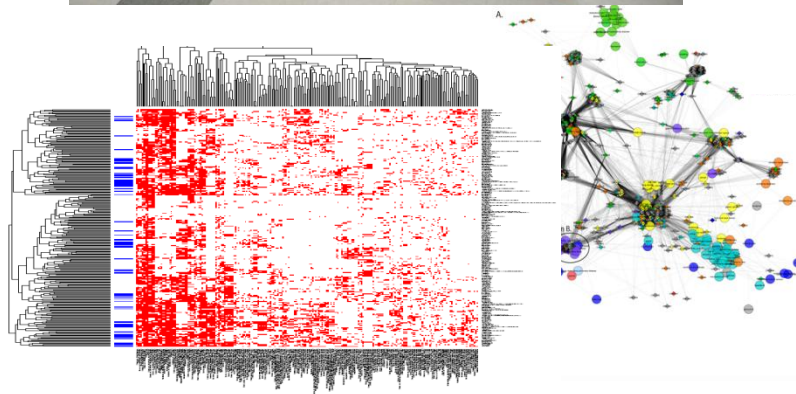
Co-Chairs

R. Daniel Benz, Ph.D. (FDA)
Kevin Crofton, Ph.D. (EPA)
Michael DeVito, Ph.D. (NTP)
David Gerhold, Ph.D. (NCGC)

- Evaluate the relevance of prioritization schemes and prediction models
- Prioritize substances for more complex testing
- Extrapolate *in vitro* conc to *in vivo* dose

Tox21 Goals

- Identify mechanisms of compound-induced biological activity in order to:
 - characterize toxicity/disease pathways
 - facilitate cross-species extrapolation
 - provide input to models for low-dose extrapolation
- Prioritize compounds for more extensive toxicological evaluation
- Develop predictive models for biological response in humans



Tox21 Implementation Strategy

- Develop
 - infrastructure to (1) support basic and applied research needed to develop the tests and pathway models, and (2) make all data/results available to scientific community
 - comprehensive suite of *in vitro* tests, preferably based on human cells, cell lines, or components
 - targeted animal tests to complement *in vitro* tests
 - computational models of toxicity pathways to support application of *in vitro* test results in hazard characterization and risk assessment
 - appropriate validation of tests and test strategies
 - evidence justifying that toxicity-pathway approach is adequately predictive of adverse health outcomes to use in decision-making

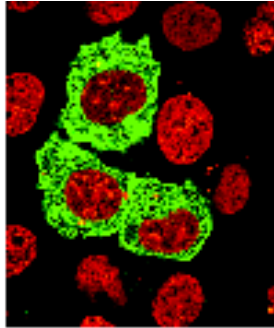
Tox21 Phase I: Proof of Principle

- NCGC screened 1408 compounds (1353 unique, 55 duplicates) from NTP and 1462 compounds (1384 unique, 78 duplicates) from EPA, with ~400 compound overlap in >100 qHTS assays.
- EPA via ToxCast™ screened 320 compounds (309 unique, 291 pesticide actives, 9 industrial, 56/73 proposed Tier 1 Endocrine Disruption Screening Program, 14 HPV, 11 HPV Challenge) in ~550 assays.
- Data released to the scientific community via:
 - EPA ACToR (Aggregated Computational Toxicology Resource; <http://epa.gov/actor>)
 - NLM PubChem (<http://pubchem.ncbi.nlm.nih.gov/>)
 - NTP CEBS (Chemical Effects in Biological Systems; <http://www.niehs.nih.gov/research/resources/databases/cebs/index.cfm>)

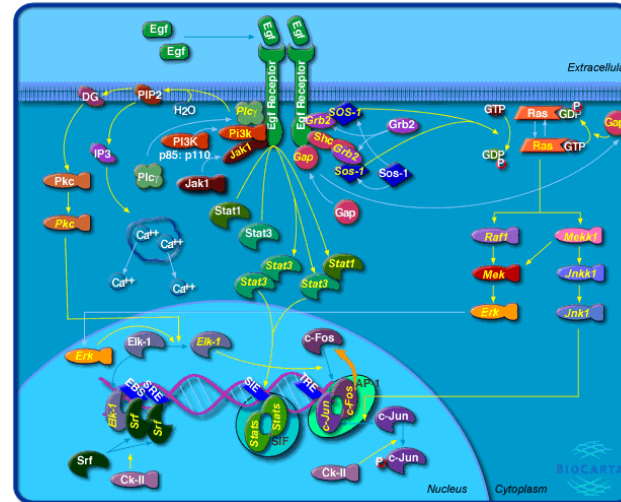
Range of screening assays performed

Extent of reductionism →

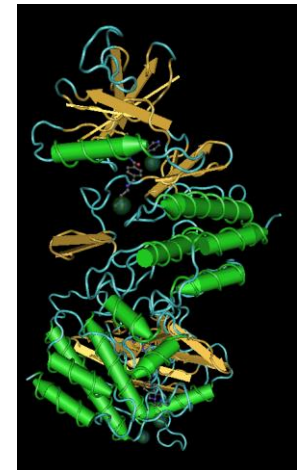
Phenotype
(Image-based HCS, GFP, etc)



Pathway
(Reporters, e.g., luciferase, β -lactamase)



Protein
(Enzyme readouts, interactions, etc)



Quantitative High-Throughput Screening (qHTS)

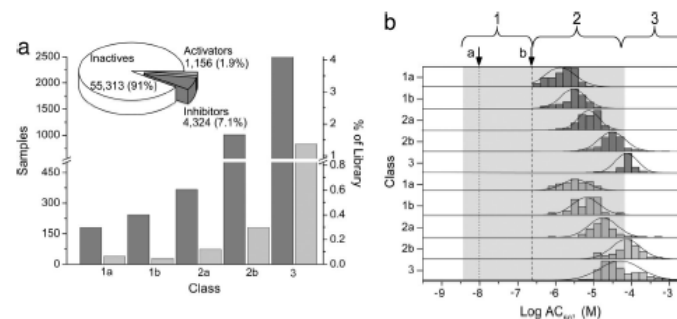
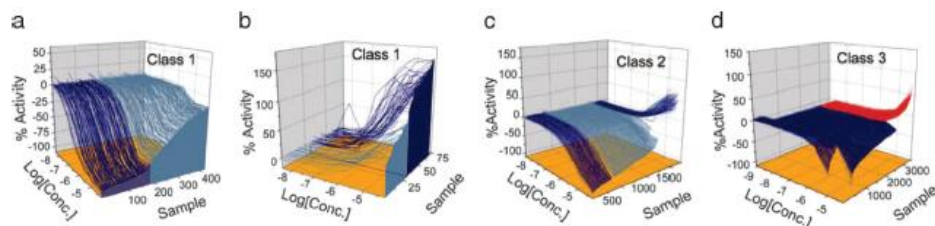
- Conventional HTS done at single concentration
 - typically 10 μM
- qHTS assays compounds at multiple concentrations
 - Tox21 assays all screened at 15 concentrations
 - Range = 2 nM – 100 μM
 - 1536-well plate format, assay volume $\sim 5 \mu\text{L}$, ~ 1000 cells/well
 - Concentration-response curve generated for each compound from primary screen
- Produces robust **activity profiles** of all compounds
 - Dramatically reduced FP and FN
- Informatics pipeline for data processing, curve fitting & classification, extraction of SAR

Quantitative high-throughput screening: A titration-based approach that efficiently identifies biological activities in large chemical libraries

James Inglesse*, Douglas S. Auld, Ajit Jadhav, Ronald L. Johnson, Anton Simeonov, Adam Yasgar, Wei Zheng, and Christopher P. Austin

NIH Chemical Genomics Center, National Human Genome Research Institute, National Institutes of Health, Bethesda, MD 20892-3370

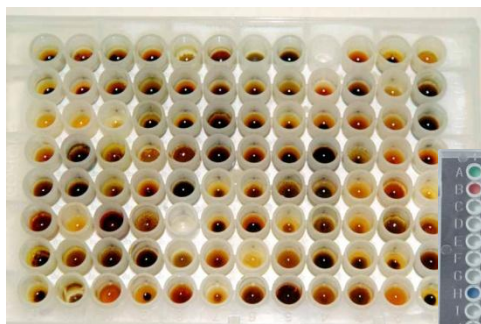
Communicated by Francis S. Collins, National Institutes of Health, Bethesda, MD, May 31, 2006 (received for review April 12, 2006)



PNAS | August 1, 2006 | vol. 103 | no. 31 | 11473-11478

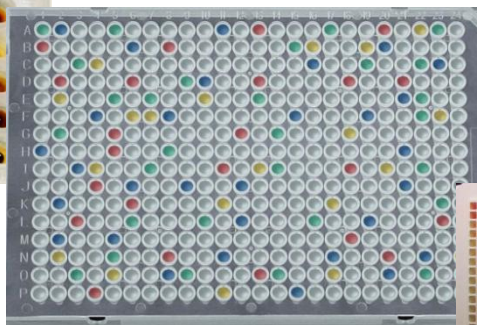
qHTS Screening Format

c



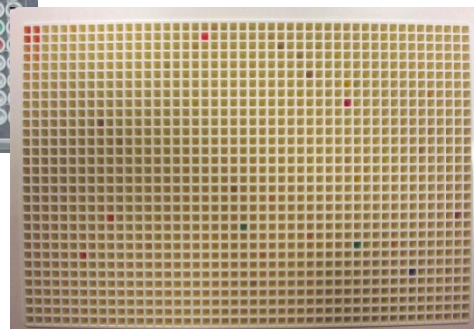
- 8 rows x 12 columns
- 88 test samples

96-well plate



- 16 rows x 32 columns
- 352 test samples

384-well plate
4 x 96-well plates



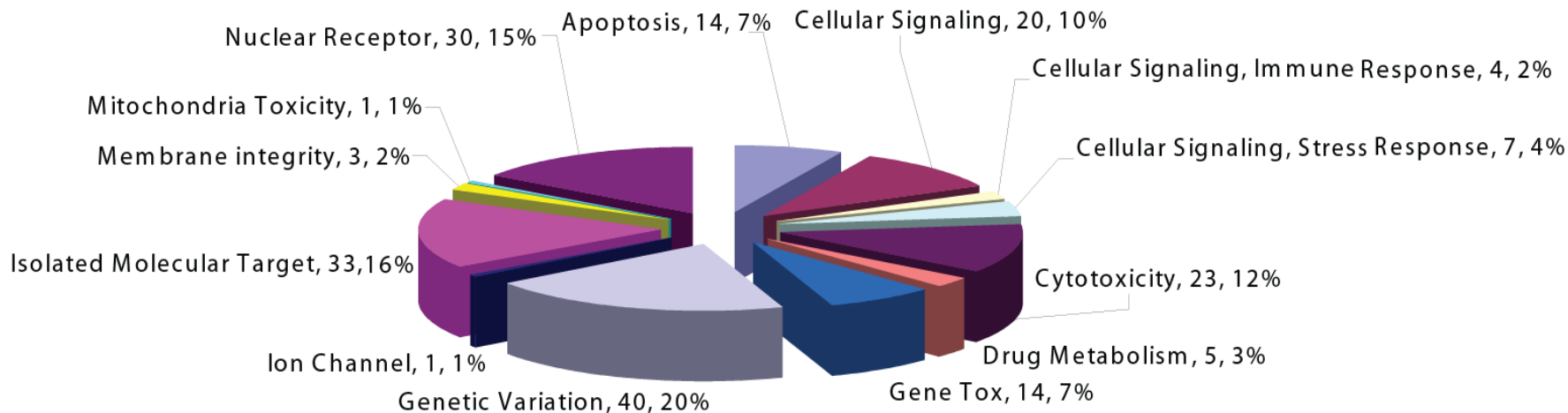
- 32 rows x 48 columns
- 1,408 test samples*

1536-well plate
16 x 96-well plates

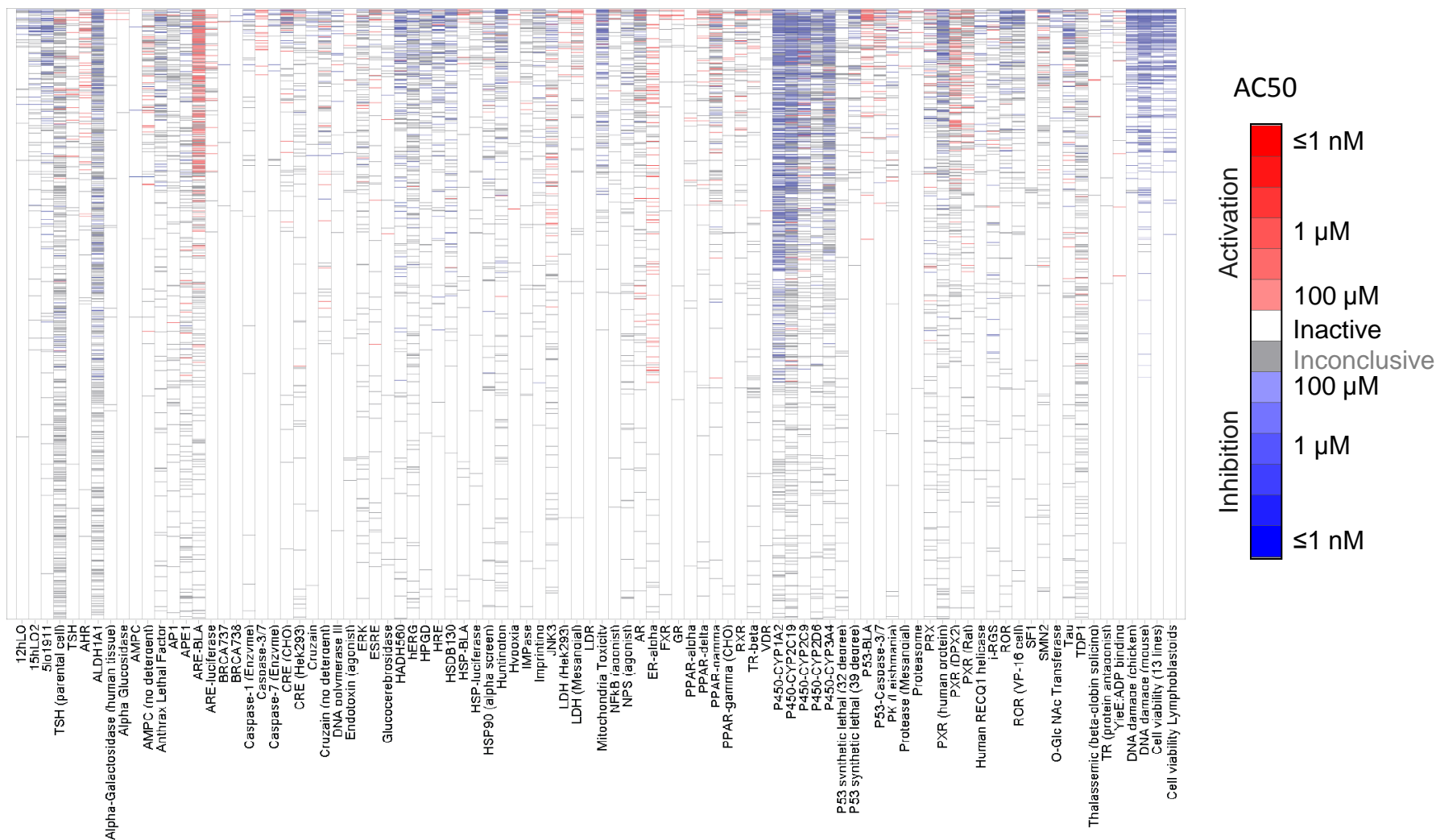
* wells remaining after subtraction of control wells; NCGC uses left 4 columns of 1536-well plate for controls

Phase I: NCGC qHTS Assays Screened

- Phenotypic readouts
 - Cytotoxicity
 - Apoptosis: caspase 3/7, 8, 9)
 - Membrane integrity: LDH, protease release
 - Mitochondrial toxicity (membrane potential)
 - Gene tox: p53, ELG1, DNA damage gene deficient lines (DT40 lines and mouse)
- Cell Signaling
 - Stress response: ARE, ESRE, HSP, Hypoxia, AP-1
 - Immune response: IL-8, TNF α , TTP
 - Other: AP-1, CRE, ERK, HRE, JNK3, NFkB, LDR
- Drug metabolism
 - CYP1A2, CYP2C19, CYP2C9, CYP2D6, CYP3A4
- Target specific assays
 - Nuclear receptors: AR, AhR, ER α , FXR, GR, LXR, PPAR α , PPAR δ , PPAR γ , PXR, RXR, TR β , VDR, ROR α , ROR γ
 - hERG channel
 - Isolated molecular targets: 12hLO, 15hLO1, 15hLO2, ALDH1A1, HADH560, HPGD, HSD17b4, α -Glucosidase, α -Galactosidase, Glucocerebrosidase, APE1, TDP1, DNA polymerase III, RECQ1 helicase, RGS4, BRCA, IMPase, O-Glc NAc Transferase, Caspase-1/7, CBF β -RUNX1, PK, Tau, Cruzain, β -Lactamase, PRX, YjeE, NPS, Proteasome, SF1, SMN2, beta-globin splicing, Anthrax Lethal Factor, TSHR
- Genetic variation: 87 HapMap lines



Phase I qHTS Compound Activity Profile

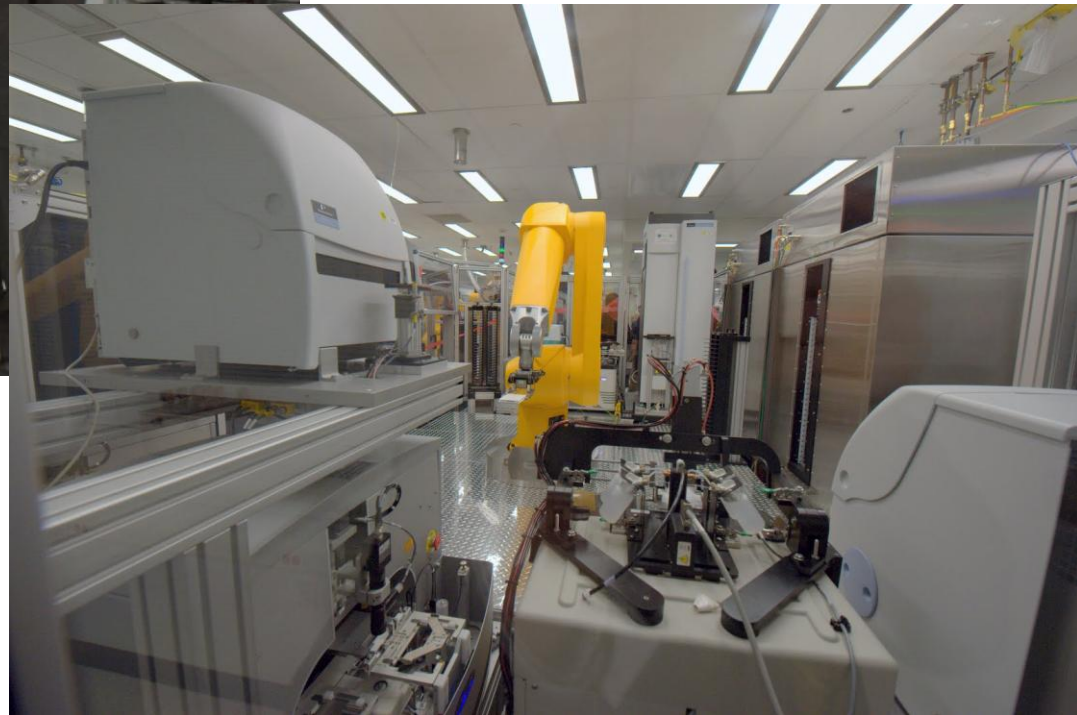


Compounds screened: NTP-1408 and EPA-1462

Tox21 Robot Ribbon-Cutting March 10, 2011



(L to R): Eric Green (Director, NHGRI/NIH),
Linda Birnbaum (Director, NIEHS/NIH),
Janet Woodcock (Director, CDER/FDA),
Lek Kadelli (Asst Administrator, ORD/EPA)



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Coverage of Tox21
engineering enthusiastic
but not always
scientifically accurate...



Natural Environment
Secures Peace of
Mind for the Mankind

Robots could reduce animal tests

U.S. scientists are taking the first step towards testing potentially hazardous chemicals on cells grown in a laboratory, without using live animals.

Two government agencies are looking into the merits of using high-speed automated robots to carry out tests.

The Tox21 Phase II 10K Compound Library

NCGC

- Pharmaceutical Collection

EPA

- ToxCast I and II compounds
- Antimicrobial Registration Program
- Endocrine Disruptor Screening Program
- OECD Molecular Screening Working Group List
- FDA Drug Induced Liver Injury Project
- Failed Drugs from Pharma

NTP

- NTP-studied compounds
- NTP nominations and related compounds
- ICCVAM/NICEATM validation and reference compounds
- External collaborators (e.g., Silent Spring Institute, U.S. Army Public Health Command)
- Defined mixtures

The NCGC Pharmaceutical Collection: A Comprehensive Resource of Clinically Approved Drugs Enabling Repurposing and Chemical Genomics

**Ruili Huang,* Noel Southall,* Yuhong Wang, Adam Yasgar, Paul Shinn,
Ajit Jadhav, Dac-Trung Nguyen, Christopher P. Austin†**

Small-molecule compounds approved for use as drugs may be “repurposed” for new indications and studied to determine the mechanisms of their beneficial and adverse effects. A comprehensive collection of all small-molecule drugs approved for human use would be invaluable for systematic repurposing across human diseases, particularly for rare and neglected diseases, for which the cost and time required for development of a new chemical entity are often prohibitive. Previous efforts to build such a comprehensive collection have been limited by the complexities, redundancies, and semantic inconsistencies of drug naming within and among regulatory agencies worldwide; a lack of clear conceptualization of what constitutes a drug; and a lack of access to physical samples. We report here the creation of a definitive, complete, and nonredundant list of all approved molecular entities as a freely available electronic resource and a physical collection of small molecules amenable to high-throughput screening.

The NCGC Pharmaceutical Collection - Windows Internet Explorer

http://tripod.nih.gov/npc/

File Edit View Favorites Tools Help

The NCGC Pharmaceutical Collection

The NCGC Pharmaceutical Collection

current version 1.1.0

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- [Has the NPC screening library been characterized analytically?](#)
- [How do I cite the NPC resource?](#)
- [Contact](#)
- [Acknowledgements](#)

What is the NCGC Pharmaceutical Collection (NPC)?

The NCGC Pharmaceutical Collection (NPC) is a comprehensive, publicly-accessible collection of approved and investigational drugs for high-throughput screening that provides a valuable resource for both validating new models of disease and better understanding the molecular basis of disease pathology and intervention. The NPC has already generated several useful probes for studying a diverse cross section of biology, including novel targets and pathways. NCGC provides access to its set of approved drugs and bioactives through the [Therapeutics for Rare and Neglected Diseases \(TRND\)](#) program and as part of the compound collection for the [Tox21 initiative](#), a collaborative effort for toxicity screening among several government agencies including the US Environmental Protection Agency (EPA), the National Toxicology Program (NTP), the US Food and Drugs Administration (FDA), and the NCGC. Of the nearly 2750 small molecular entities (MEs) that have been approved for clinical use by US (FDA), EU (EMA), Japanese (NHI), and Canadian (HC) authorities and that are amenable to HTS screening, we currently possess 2400 as part of our screening collection.

How do I get access to the NPC?

The NPC resource currently consists of (i) the physical collection suitable for high throughput screening (HTS) and (ii) the informatics browser and database. Putting together the physical collection has been surprisingly challenging in terms of the time and effort required in the informatics, compound management and synthetic chemistry related activities required for this endeavor. We provide access to the NPC screening library through collaboration. Please contact our Scientific Director Dr. [Chris Austin](#) for additional information.

The other half of the NPC resource is the NPC browser. This is a self-contained software that is actively developed and maintained by the informatics group to provide electronic access to the NPC content. The latest version of the NPC browser for various platforms can be downloaded [below](#). Please let us know if your platform is not listed. Note that a fairly modern hardware (preferably with at least 2Gb of memory) is required to run the browser effectively.

How do I download the NPC browser?

File	Platform	Size	MD5 checksum
npc-browser_windows_1_1_0.exe	Windows	94.3 MB	9329911fba2e6ff44f2645f38337cc3b
npc-browser_macos_1_1_0.dmg	Mac OS X	78.7 MB	b9e58f73ba1f13a74f6525a2b88c7807
npc-browser_unix_1_1_0.sh	Linux	95.3 MB	7afedcc3741bd0ab8c96938a9ab9c852

Links

- [NCGC](#)
- [MLI](#)
- [NHGRI](#)
- [PubChem](#)
- [ClinicalTrials.gov](#)
- [DailyMed](#)
- [Drug Portal](#)

Presentations

- [ACS Spring 2010](#)
- [Chemaxon UGM 2008](#)

Tools

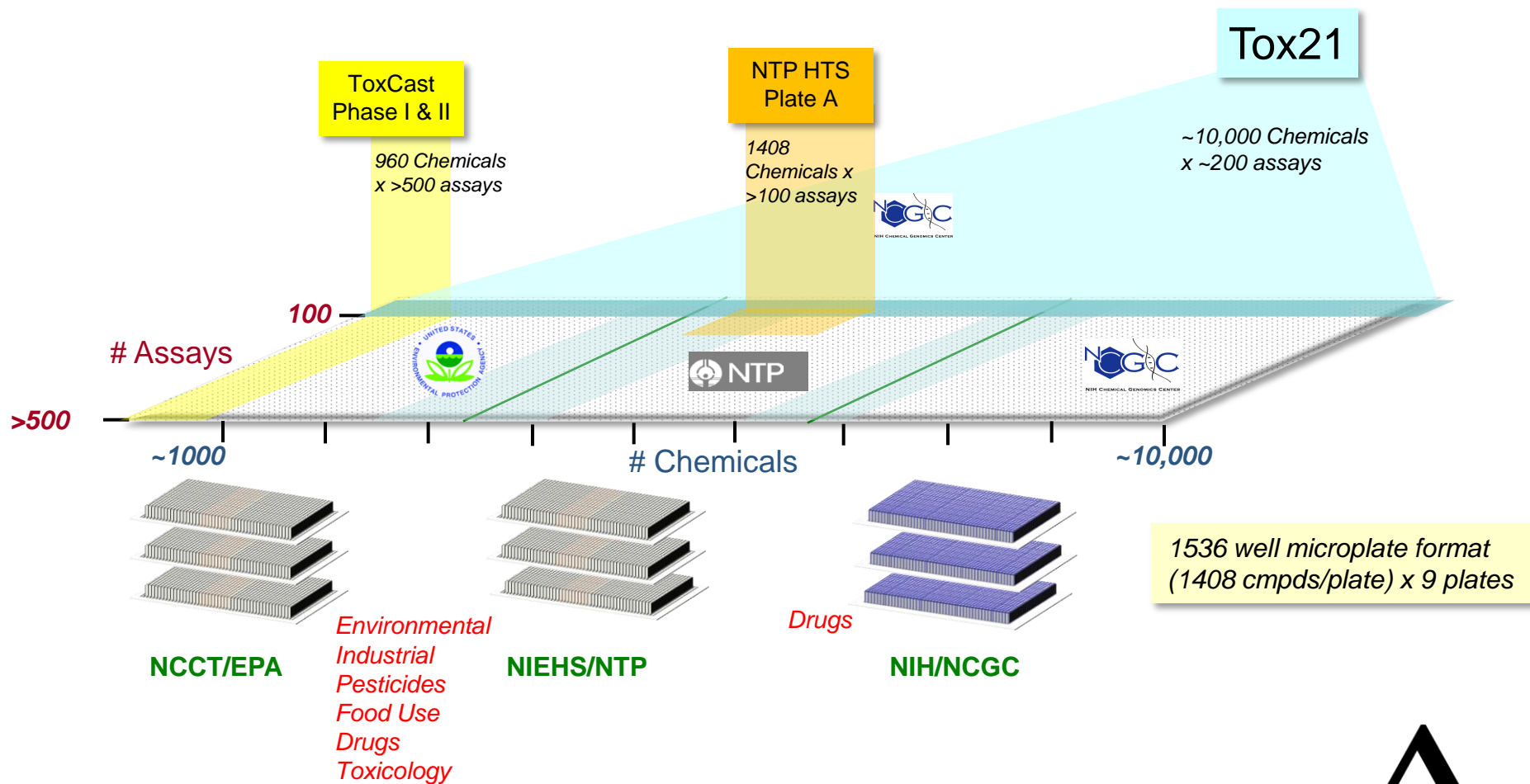
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Local intranet 100%

Tox21 Chemicals x Assays Landscape

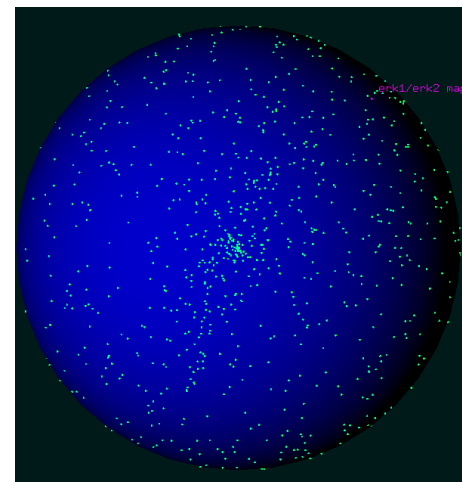


Phase II qHTS Strategic Screening Strategy

- Assay selection based on
 - Phase I experience
 - Information from *in vivo* toxicological investigations
 - Advice of basic researchers and nominated assays
 - Maps of disease-associated cellular pathways
- Stage I
 - nuclear receptor activation or inhibition (AR, AhR, ER, FXR, GR, LXR, PPAR,, PXR, RXR, TR, VDR, ROR)
 - induction of stress response pathways (e.g., DNA damage, heat shock, hypoxia, inflammation, oxidative)
- Stage II
 - other disease-associated pathways (e.g., obesity/diabetes, autism) and move to HTS gene array assays applicable to all cell types

Facilitating Choice of Assays for Tox21: The NCGC BioPlanet of Pathways

- Hosts the universe of pathways
 - All pathway annotations from manually curated, public sources
 - Integrates ~1,100 unique human pathways from different data sources
 - Annotates pathways by source, species, biological function/process, disease/toxicity relevance, assay availability
 - Easy visualization, browsing, analysis of pathways
- Facilitates pathway assay selection/prioritization for Tox21 Phase 2
 - Disease, Toxicity pathways
 - Assay availability
 - Tox21/ToxCast/NCGC/PubChem
 - Commercial assays
 - Develop new assays for pathways with no coverage
- Will be publicly available: <http://spotlite.nih.gov/tox21/>
- Future developments
 - Link compound activity data
 - Incorporate other data forms: sequence, gene/protein expression data, etc.
 - Other species: rat, mouse, etc.
 - Organize assays according to pathways/diseases/toxicity endpoints



The NCGC Universe of Human Pathways

Detailed view of a pathway

Pathways

The screenshot displays the NCGC BioPlanet interface. On the left, a large blue sphere represents the 'Pathways' universe, with a red dot labeled 'Homologous recombination'. The top left panel contains search filters for Category, Gene/Pathway, and various databases like PubChem, Tox21, and NCGC. The top right panel shows a detailed network diagram of a pathway with nodes like FLJ10195, RPA70, and SOSS-B1. The bottom right panel shows a 'Find Gene' window for 'thioredoxin' with a table of related genes.

Gene ID	Gene Symbol	Gene Description
5889	RAD51C RAD51L2 MGC104277	RAD51 homolog C (S. cerevisiae)
5425	POLD2	polymerase (DNA directed), delta 2, regulatory subunit 50kDa
6119	REPA3 RPA3	replication protein A3, 14kDa
6742	SOSS-B1 SSBP1 SSBP	single-stranded DNA binding protein 1
25788	FSBP RDH54 RAD54B	RAD54 homolog B (S. cerevisiae)
29935	MGC120334 HSU24186 MGC120333 RPA4	replication protein A4, 34kDa
7516	XRCC2 DKFZp781P0919	X-ray repair complementing defective repair in Chinese hamster cells 2
5888	HsT16930 HRAD51 RAD51A HsRad51 BRCC5 RECA RAD51	RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)
5424	CDC2 POLD1 POLD	polymerase (DNA directed), delta 1, catalytic subunit 125kDa
4361	HNGS1 MRE11B ATLD MRE11A MRE11	MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
675	GLM3 BRCC2 BROVCA2 FANCD1 FANCD FAD1 BRCA2 FAD FANCB FADC	breast cancer 2, early onset
4683	FLJ10155 NBN MGC87362 NBS1 NBS ATV AT-V2 P95 AT-V1	nibrin
7156	TOP3 TOP3A	topoisomerase (DNA) III alpha
7517	XRCC3	X-ray repair complementing defective repair in Chinese hamster cells 3
146956	FLJ31364 EME1 MMS4L	essential meiotic endonuclease 1 homolog 1 (S. pombe)
10111	hRad50 NBSLD RAD50-2 RAD50	RAD50 homolog (S. cerevisiae)
641	RECQL3 MGC131618 BS BLM RECQ2 MGC131620 MGC126616 RECQL2	Bloom syndrome, RecQ helicase-like
10714	POLD3 P66 P68 MGC119643 KIAA0039 MGC119642	polymerase (DNA-directed), delta 3, accessory subunit
5893	RAD52	RAD52 homolog (S. cerevisiae)
6118	RPA2 RPA32 REPA2	replication protein A2, 32kDa
8940	FLJ39376 TOP3B	topoisomerase (DNA) III beta
8438	hRAD54 RAD54A HR54 hHR54 RAD54L	RAD54-like (S. cerevisiae)
80198	FLJ21012 MUS81 FLJ44872	MUS81 endonuclease homolog (S. cerevisiae)

Gene information

~1100 human pathways mapped to the pathway globe

"Disease" pathways (per OMIM)

The NCGC BioPlanet

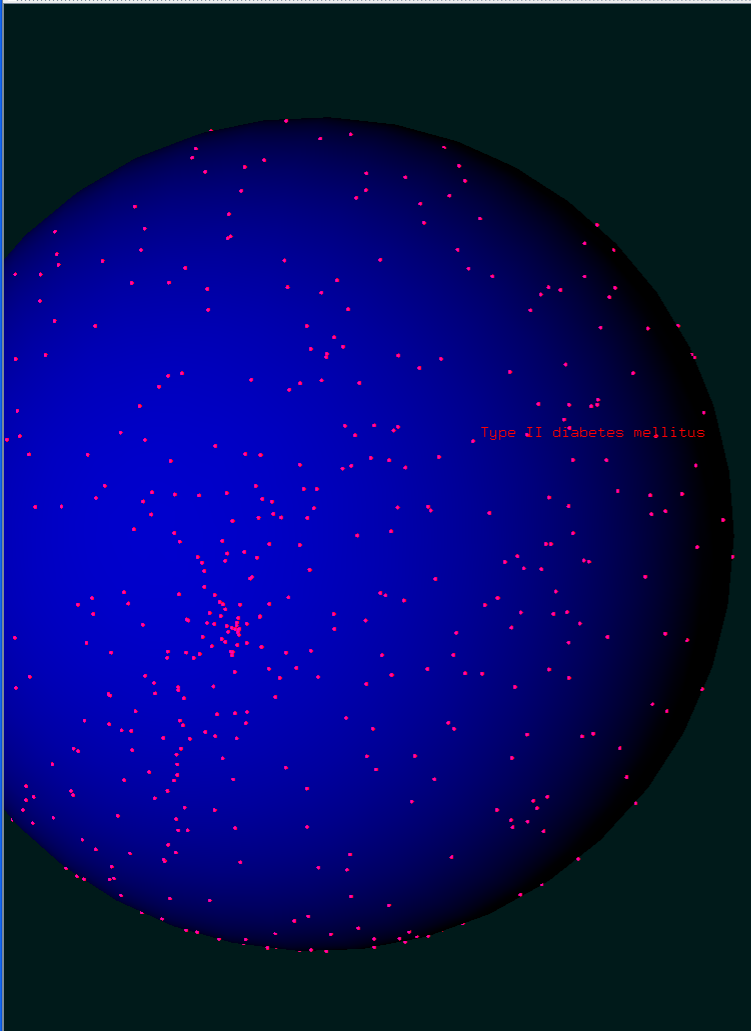
Category: Toxicity Disease PubChem

Gene/Keyword:

Find

And Or 0 25 50 75 100

4 pathways found.



Type II diabetes mellitus

Find Gene: IRDN

Information

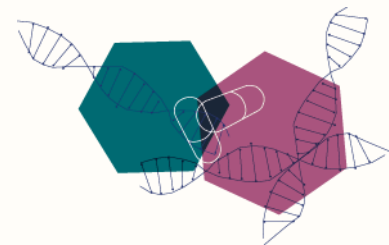
hsa04930: Type II diabetes mellitus

Gene ID	Gene Symbol	Gene Description
5602	PRKM10 MGC50974 JNK3A FLJ33785 MAPK10 FLJ12099 p54bSAPK JNK3 p493F12	mitogen-activated protein kinase 10
5291	PIK3CB PI3KBETA DKFZp779K1237 PIK3C1 PI3K P110BETA MGC133043	phosphoinositide-3-kinase, catalytic, beta polypeptide
6517	SLC2A4 GLUT4	solute carrier family 2 (facilitated glucose transporter), member 4
6833	TNDM2 ABC36 SUR HI MRP8 HRINS SUR1 HHF1 PHH1 ABCC8	ATP-binding cassette, sub-family C (CFTR/MRP), member 8
5296	P85B p85 p85-BETA PIK3R2	phosphoinositide-3-kinase, regulatory subunit 2 (beta)
5581	nPKC-epsilon PRKCE MGC125656 PKCE MGC125657	protein kinase C, epsilon
5293	P110DELTA PIK3CD p110D PI3K	phosphoinositide-3-kinase, catalytic, delta polypeptide
5594	MAPK2 MAPK1 p41mapk ERK2 p38 ERK P42MAPK p40 p41 PRKM2 PRKM1 ERT1	mitogen-activated protein kinase 1
776	Cav1.3 CACN4 CACH3 CACNL1A2 CCHL1A2 CACNA1D	calcium channel, voltage-dependent, L type, alpha 1D subunit
8471	IRS4 IRS-4 PY160	insulin receptor substrate 4
3098	HK1 HK1 HXK1 HK1-1b HK1-1a HK1-1c	hexokinase 1
2475	FLJ44809 FRAP2 FRAP1 RAPT1 FRAP MTOR RAFT1	mechanistic target of rapamycin (serine/threonine kinase)
5599	JNK PRKM8 JNK1A2 JNK1 JNK2B1/2 MAPK8 SAPK1	mitogen-activated protein kinase 8
7124	TNFA TNF TNFSF2 DIF TNF-alpha	tumor necrosis factor (TNF superfamily, member 2)
5295	p85-ALPHA GRB1 p85 PIK3R1	phosphoinositide-3-kinase, regulatory subunit 1 (alpha)
3101	HXK3 HK3 HKIII	hexokinase 3 (white cell)
777	CACNL1A6 CACNA1E CACH6 Cav2.3 BII	calcium channel, voltage-dependent, R type, alpha 1E subunit
3630	IRDN INS ILPR	insulin
8835	SOCS-2 Cish2 CIS2 SOCS2 SSI2 STATI2 SSI-2	suppressor of cytokine signaling 2
23533	FOAP-2 p101 PIK3R5 F730038115Rik P101-PI3K	phosphoinositide-3-kinase, regulatory subunit 5
389692	hMafA RIPE3b1 MAFA	v-maf musculoaponeurotic fibrosarcoma oncogene homolog A (avian)

NCATS: Pursuing Opportunities for Disruptive Innovation



To catalyze the generation of innovative methods and technologies that will enhance the development, testing, and implementation of diagnostics and therapeutics across a wide range of human diseases and conditions.



NCATS: Programs & Initiatives

Clinical and Translational Science Activities

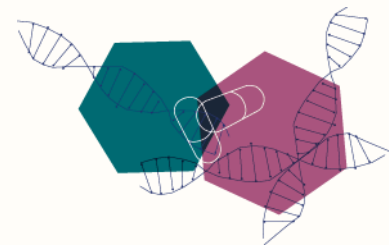
- Clinical and Translational Science Awards

Rare Diseases Research and Therapeutics

- Therapeutics for Rare and Neglected Diseases
- Office of Rare Diseases Research

Re-engineering Translational Sciences

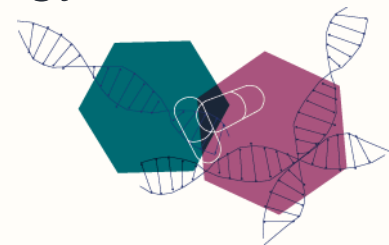
- NIH Chemical Genomics Center
- Bridging Interventional Development Gaps
- Toxicology in the 21st Century



NCATS: Fostering Collaboration

NIH–FDA–DARPA Collaboration for Tissue Chip

- Aims to develop a tissue chip that mimics human physiology to screen for safe, effective drugs
 - Liver, heart, lung, other cell types
 - Designed for multiple types of readouts
- NIH and Defense Advanced Research Projects Agency (DARPA) contribute \$70M over 5 years; FDA provides guidance
- Received first proposals in late January 2012
 - Seeking best ideas in engineering, biology, toxicology





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Program Snapshot

The NIH is partnering with the U.S. Food and Drug Administration (FDA) and the Defense Advanced Research Projects Agency (DARPA) to advance the field of regulatory science, a specialized research area that aims to improve assessment of experimental therapies, preventives, and diagnostics. The Common Fund's **Regulatory Science** program is fostering the development, evaluation and availability of new or improved tools, methods, standards, and applied science that support a better understanding and improved evaluation of product safety, quality, effectiveness, and manufacturing throughout the product life cycle.

During the initial phase of the program, launched in fiscal year 2010, four new research awards in high priority areas of regulatory science were supported. Expansion of the program in FY 2012 focuses on developing new cell-based technologies, called microsystems, to predict more accurately drug safety and efficacy in humans.

[Read more...](#)

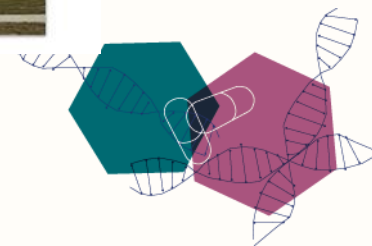
Program Highlights

In 2010, in partnership with the U.S. Food and Drug Administration (FDA), the Common Fund awarded four new grants in the Regulatory Science program. The awards address four distinct,

New collaboration and funding announcement to accelerate therapeutics development

The NIH announces a new collaboration and funding opportunity through the Common Fund's Regulatory Science program, and involving NIH, DARPA, and the FDA, to advance the development of new technologies aimed at streamlining the drug development pipeline. The initiative will support the development of human microsystems, or organ "chips," that can be used to screen for safe and effective drugs far more swiftly and efficiently than current methods, and before they are tested in humans. These microsystems will use specific cell types that reflect the biology of several different organs and tissues, and will be integrated together to model the connection between different organ systems in the human body. This integration will allow researchers to assess how drugs metabolized by one organ affect other organs or systems. It is hoped that the development of such microsystems will allow faster and more accurate measures of drug toxicology and efficacy, thereby reducing the time and cost associated with new therapeutics development.

[View the funding opportunity](#)
[Read Frequently Asked Questions \(FAQs\) for the funding opportunity](#)
[Read the press release from NIH Director announcing NIH-DARPA-FDA collaboration](#)





For Immediate Release
Friday, September 16, 2011

Contact:
[NIH Communications](#)
301-496-5787

NIH, DARPA and FDA collaborate to develop cutting-edge technologies to predict drug safety

President Obama announced today that the National Institutes of Health will collaborate with the Defense Advanced Research Projects Agency (DARPA), and the U.S. Food and Drug Administration to develop a chip to screen for safe and effective drugs far more swiftly and efficiently than current methods, and before they are tested in humans. The chip will be loaded with specific cell types that reflect human biology. It will be designed to allow multiple different

readouts that can indicate whether a particular compound will be safe and effective. The agencies will run separate and independent programs, but they will also run collaborative programs. This fall, the two agencies, in coordination with the Food and Drug Administration, will convene academic institutions, and other research organizations to discuss advances in engineering, biology, and toxicology

"Drug toxicity is one of the most common reasons drugs fail in clinical trials," said NIH Director Francis S. Collins. "We need to know which ones are safe and effective before we test them in humans. This is an unprecedented opportunity to speed development of new drugs."

Over the next five years, the NIH plans to commit \$1.5 billion to this effort. This groundbreaking effort is an example of the National Center for Advancing Translational Science's mission to provide science-based solutions to reduce costs and help determine how this new technology can be used in clinical studies.

"We know the development pipeline has bottlenecked at the point of testing drugs in humans. What we need are entirely novel approaches to test drugs before they are tested in humans. Biomedical discoveries that have been made in recent years have shown that this is possible."

As proposed, NCATS will study the steps in the process that create bottlenecks, and experiment with innovative methods to streamline the process. By focusing on developing innovative new tools and methods for therapeutics development, as opposed to developing therapeutics themselves, NCATS will enable others to bring safer and more effective medical products to market in less time. In this way, NCATS will complement, and not compete with, the work of the private sector and other NIH translational science efforts.

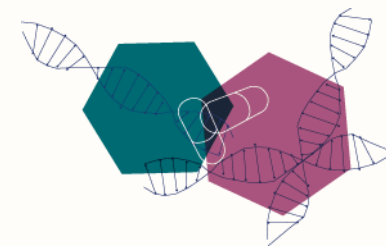
About the National Institutes of Health (NIH): NIH, the nation's medical research agency, includes 27 Institutes and Centers and is a component of the U.S. Department of Health and Human Services. NIH is the primary federal agency conducting and supporting basic, clinical, and translational medical research, and is investigating the causes, treatments, and cures for both common and rare diseases. For more information about NIH and its programs, visit www.nih.gov.

NIH...Turning Discovery Into Health



Current Funding Opportunities

Title	NIH Guide	RFA Number	Common Fund Contact	Application Receipt Date
Pre-Application Teleconference for RFA-RM-11-022 - Integrated Microphysiological Systems for Drug Efficacy and Toxicity Testing in Human Health and Disease (UH2/UH3) and RFA-RM-12-001 Stem/Progenitor Cell-Derived Human Micro-organs and -tissues (U18) <ul style="list-style-type: none"> Read Frequently Asked Questions (FAQs) for the funding opportunity View slides from the December 16, 2011 Pre-Application Teleconference 	12/5/11	NOT-RM-12-007	Margaret Sutherland sutherlandm@ninds.nih.gov 301 496-5680	N/A
Integrated Microphysiological Systems for Drug Efficacy and Toxicity Testing in Human Health and Disease (UH2/UH3)	11/22/11	RFA-RM-11-022	Danilo A. Tagle tagled@ninds.nih.gov 301 496-5745	1/26/12
Stem/Progenitor Cell-Derived Human Micro-organs and -tissues (U18)	11/23/11	RFA-RM-12-001	Margaret Sutherland sutherlandm@ninds.nih.gov 301 496-5680	1/26/12



Related/Coordinated Initiatives

- ICCVAM, NICETAM, ECVAM
- REACH: European Community Regulation on chemicals and their safe use
 - **R**egistration, **E**valuation, **A**uthorisation and Restriction of **C**hemical substances
- IMI eTox: drug safety database from the pharmaceutical industry legacy toxicology reports and public toxicology data
- OECD Molecular Screening project: internationally collaborative efforts to define screening applications in a regulatory context

Take-home points

- Opportunity and imperative now exists to transform preclinical toxicology from an empirical animal-based exercise to a predictive mechanism-based science
- Tox21 and the DARPA-NIH toxicology initiative are two of the initiatives driving this vision forward
- All data and results from both initiatives are being made public for all researchers to use/compute on
- U.S. efforts are being coordinated with related international initiatives including REACH, eTox, OECD
- For the most part these are research initiatives so far, not broadly applied in regulatory contexts

Further Information

NCATS.nih.gov

austinc@mail.nih.gov



Tox21: <http://www.epa.gov/ncct/Tox21/>

ICCVAM/NICETAM: <http://iccvam.niehs.nih.gov/>

REACH: http://ec.europa.eu/environment/chemicals/reach/reach_intro.htm

eTox: <http://www.etoxproject.eu/>

OECD Molecular Screening project:

http://www.oecd.org/document/14/0,3746,en_2649_34377_48906062_1_1_1_1,00.html