

2009 International Workshop on Virtual Tissues:

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UNITED STATES ENVIRONMENTAL PROTECTION AGENCY



Office of Research and Development
National Center for Computational Toxicology

April 21 2009



United States
Environmental Protection
Agency

EPA RTP

511 Acre Campus (shared with NIEHS)

2nd largest EPA facility (1.2m sq ft)

Longest solar power lighted road

100% Green Power

1800 employees working in 500 labs

Headquarters of two National

Laboratories, a National Center and
a Program Office

Top three ranking for postdocs

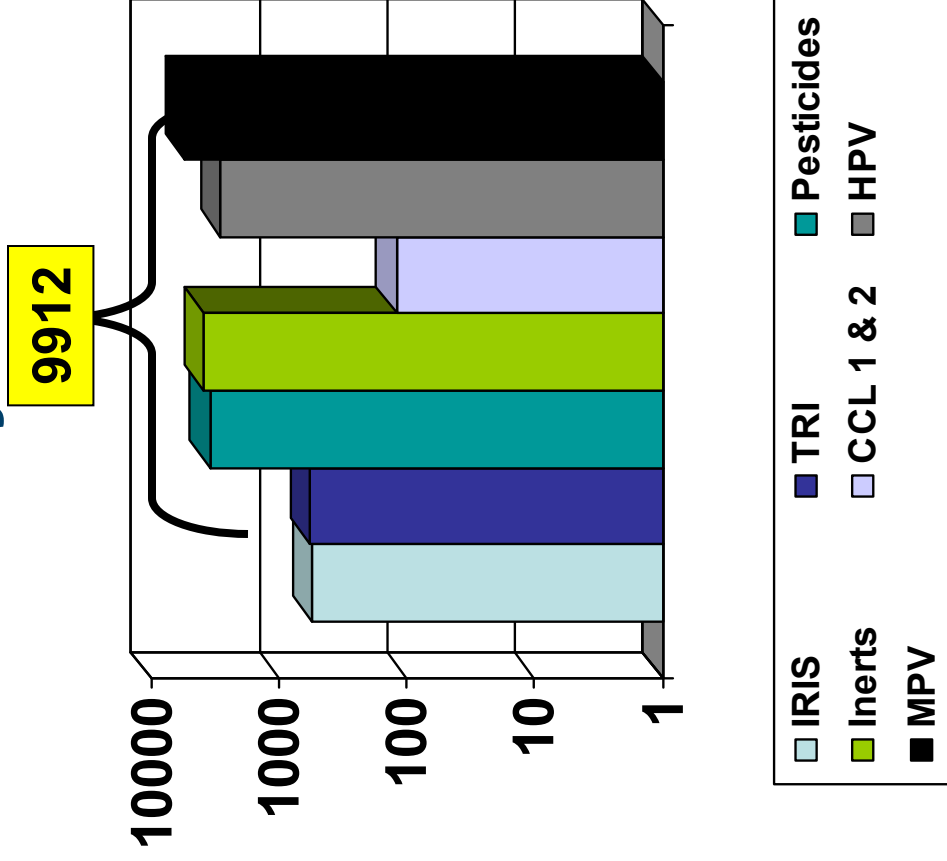


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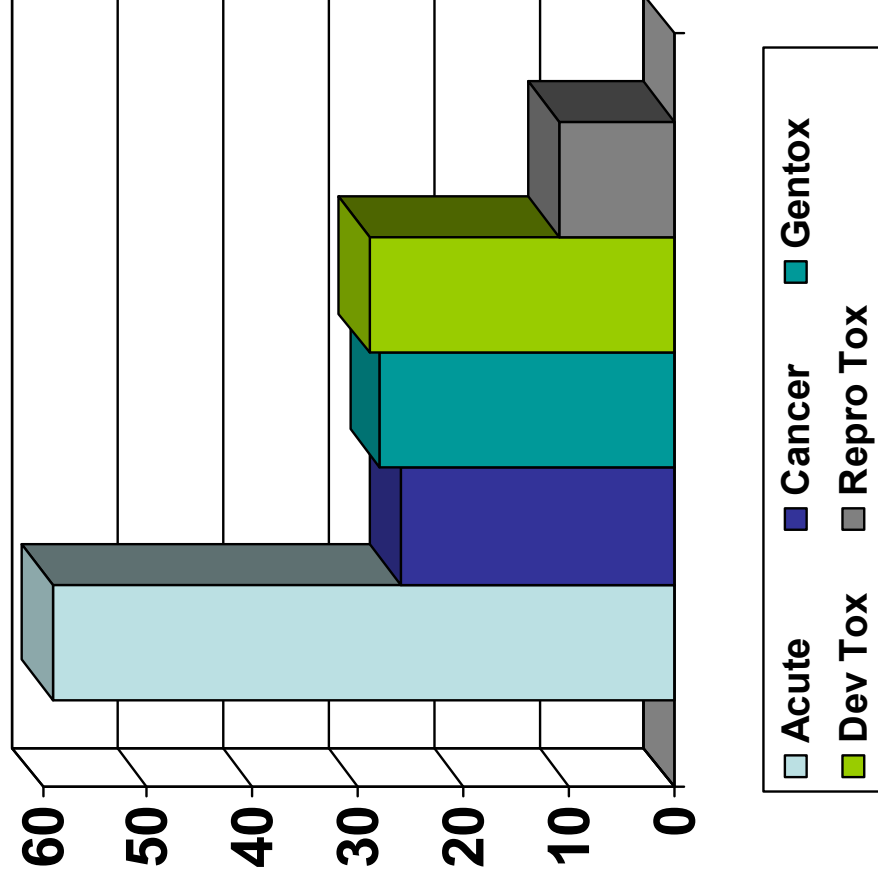
EPA's Need for Prioritization

Too Many Chemicals



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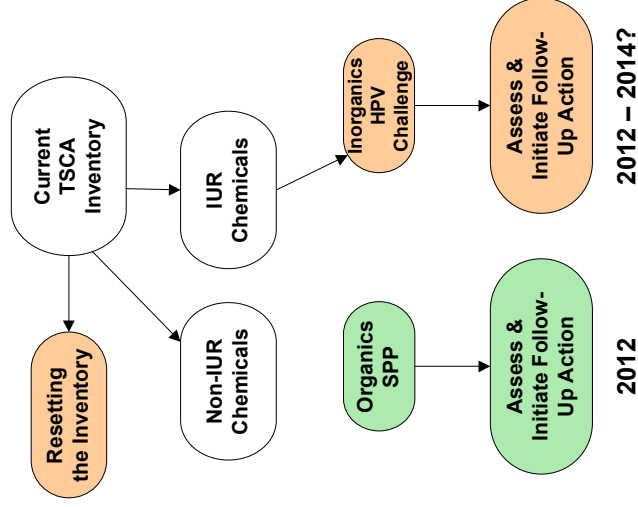
Too Little Data (%)



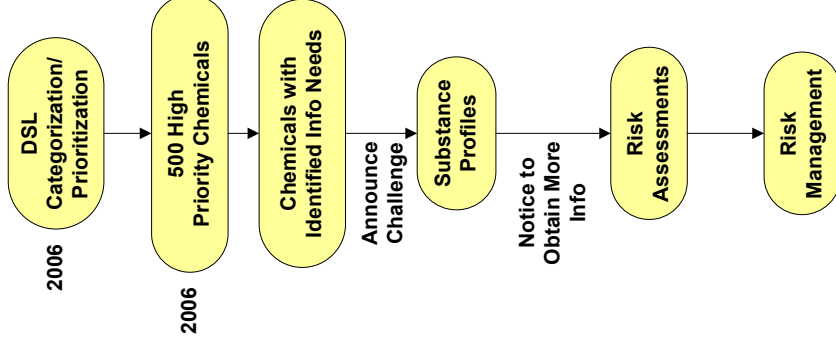
Judson, et al *EHP* (2009)

An International Problem

U.S. ChAMP

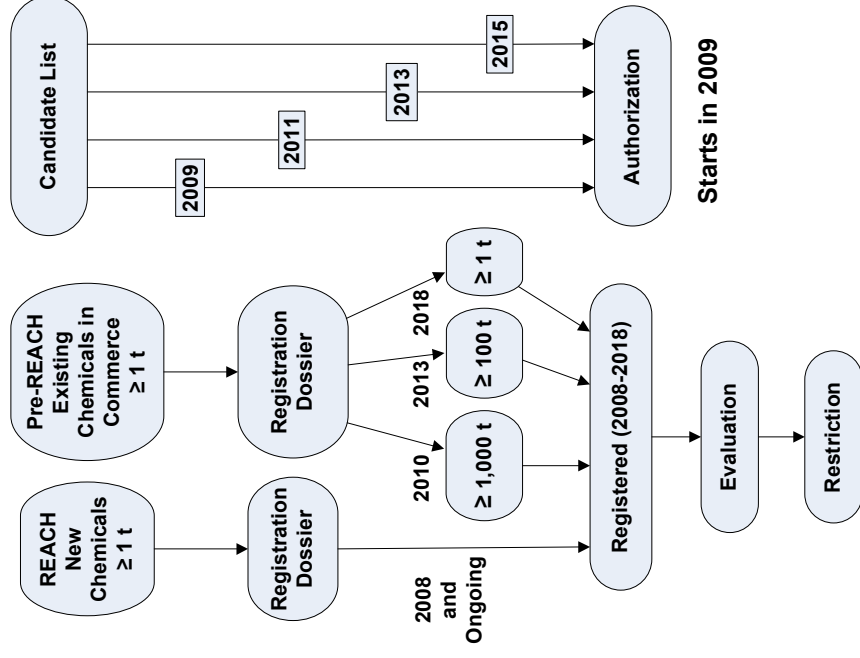


Canada Chemical Management Plan²



2011

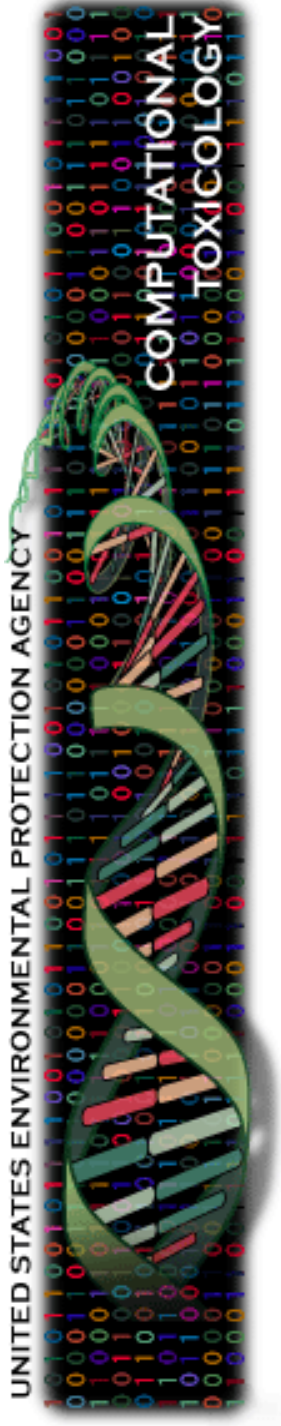
EU REACH Registration & Authorization Candidate List



¹ DSL = Canadian Environmental Protection Act Domestic Substances List

² Other aspects of the CMP are not shown on this figure.

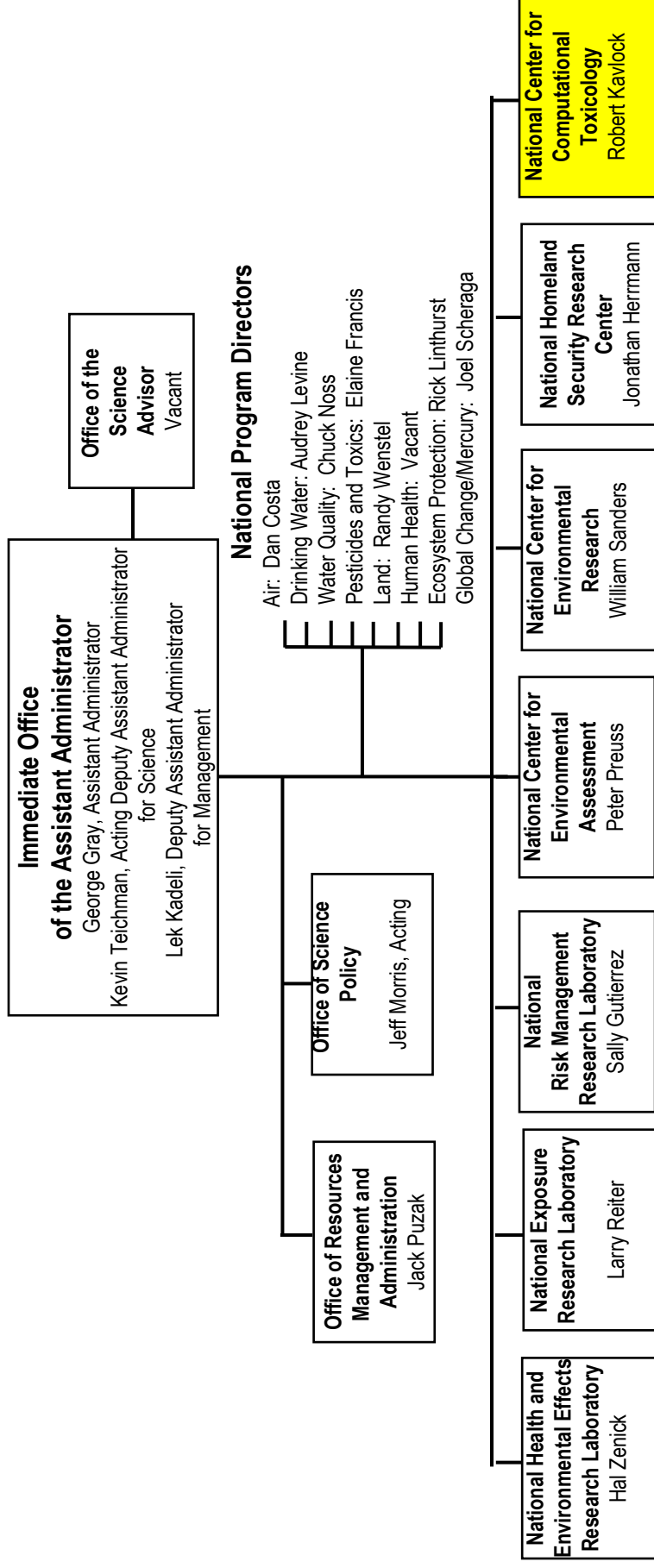




“ ...to integrate modern computing and information technology with molecular biology to improve Agency prioritization of data requirements and risk assessment of chemicals”

www.epa.gov/ncct

Office of Research and Development



Future of Toxicity Testing

POLICYFORUM

TOXICOLOGY

Transforming Environmental Health Protection

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

In 2005, the U.S. Environmental Protection Agency (EPA), with support from the U.S. National Toxicology Program (NTP), the National Institutes of Health (NIH), and the National Science Foundation (NSF), issued a joint vision statement for 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 new technologies 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 proposed a 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 2004, the NTP released its vision and roadmap for the 21st century (1), which established initiatives to integrate high-

EPA, NRC, and NTP Joint Activities
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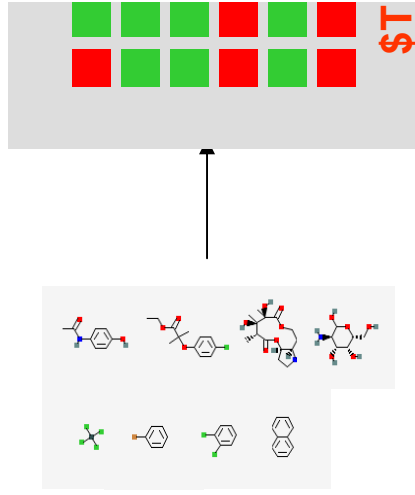
*The views expressed here are those of the individual authors and do not necessarily reflect the views and policies of their respective agencies.

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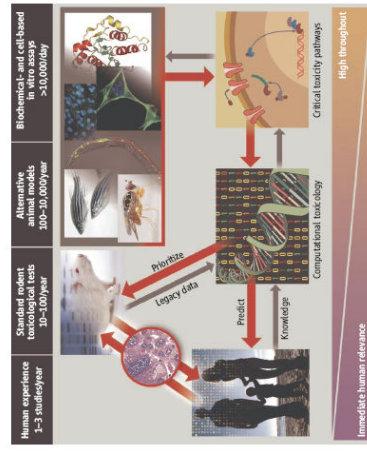
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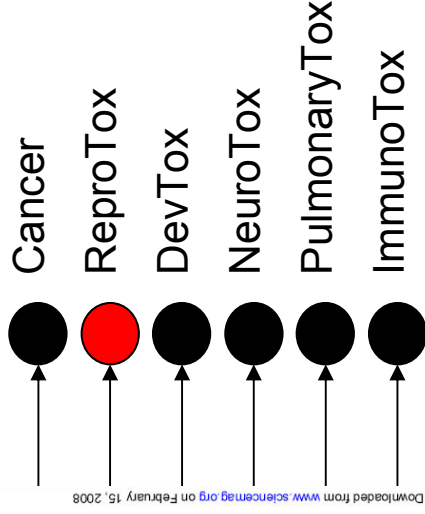
We propose a shift from primarily in vivo animal studies to in vitro assays, in vitro assays with lower organisms, and computational modeling for toxicity assessments.

throughout screening (HTS) and other automated screening assays into its testing program. In 2005, the EPA established the National Center for Computational Toxicology (NCCT), with the NRC, 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 mechanisms and 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 (6). HTS methods are being used to screen additional test compounds at one concentration, usually between 2 and 10 μ M, and tolerance high false-negative rates. In contrast, in the EPA, NRC, and NTP combined effort, concentrations generally ranging from $\sim$ 5 nM to $\sim$ 100 μ M, 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 integrated approach to data analysis is being expanded to allow comparisons with historical toxicologic NTP and EPA data collected by EPA and NTP, as well as by the NRC and other Molecular Libraries Initiative centers (http://mli.nih.gov), are being made publicly available through the PubChem website (http://pubchem.ncbi.nlm.nih.gov). In addition,



Transforming toxicology. This 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.



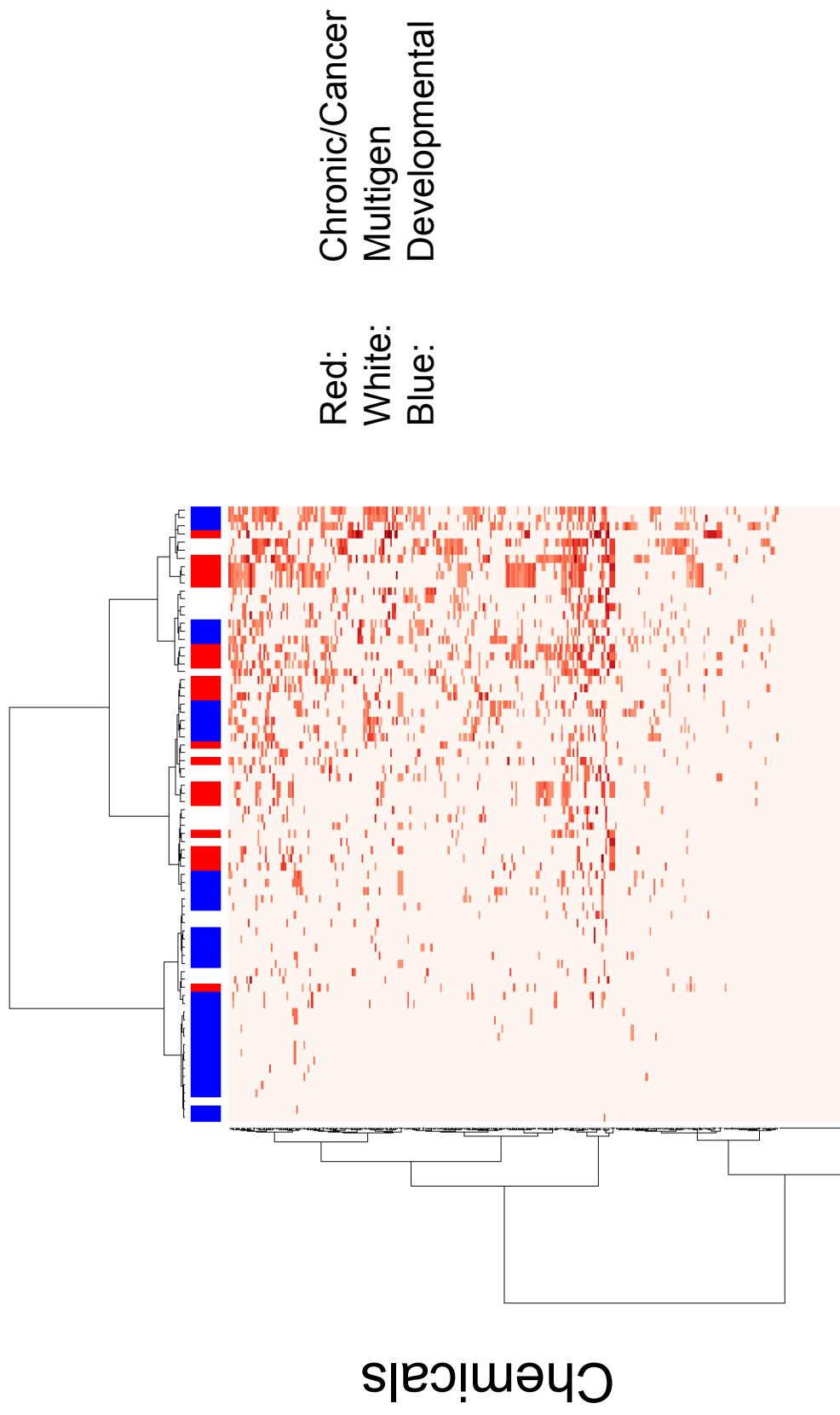
Downloaded from www.sciencemag.org on February 15, 2008

EPAs Contribution: The ToxCast Research Program

Office of Research and Development
National Center for Computational Toxicology

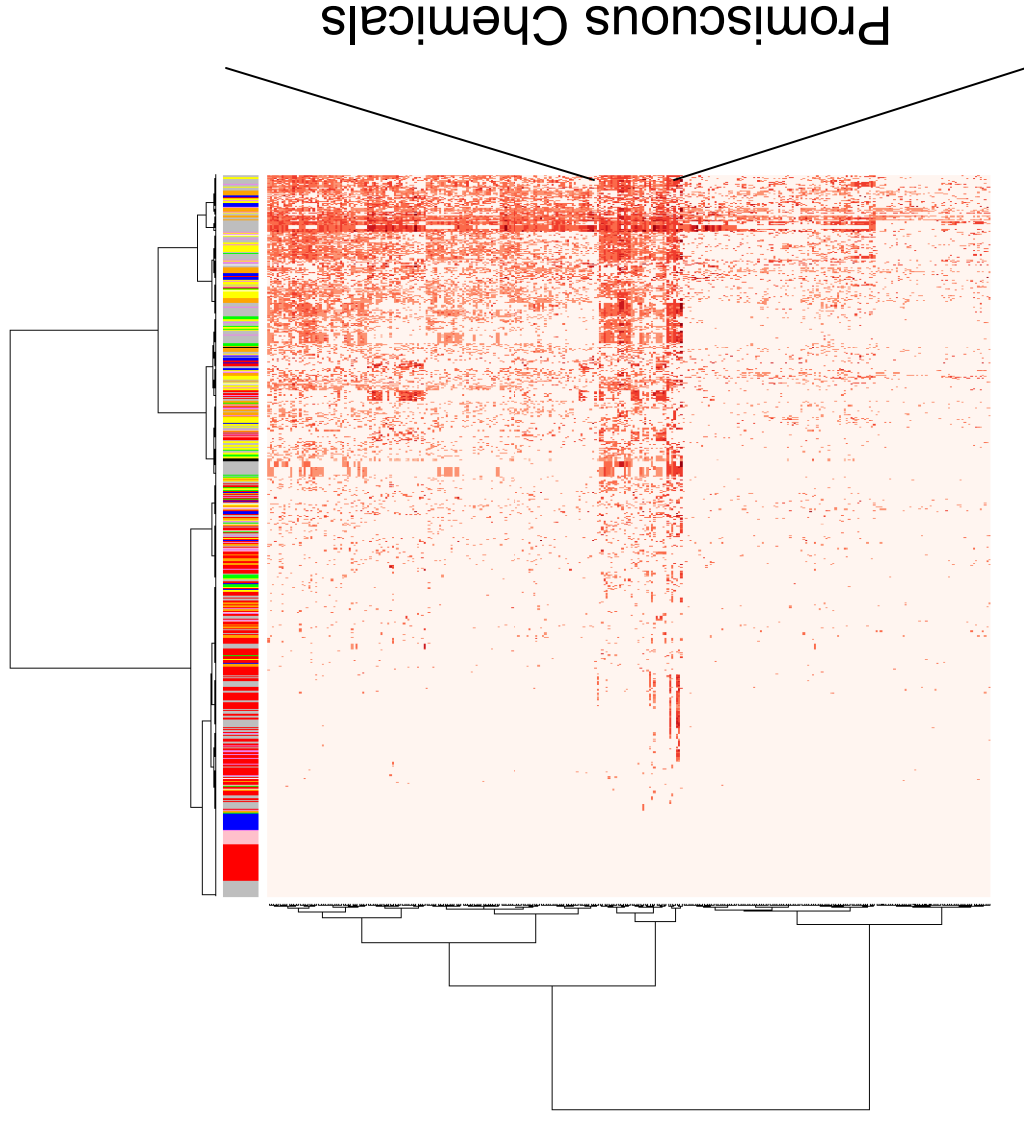
www.epa.gov/ncct/toxcast

ToxCast In Vivo Data from ToxRefDB



ToxCast In vitro data

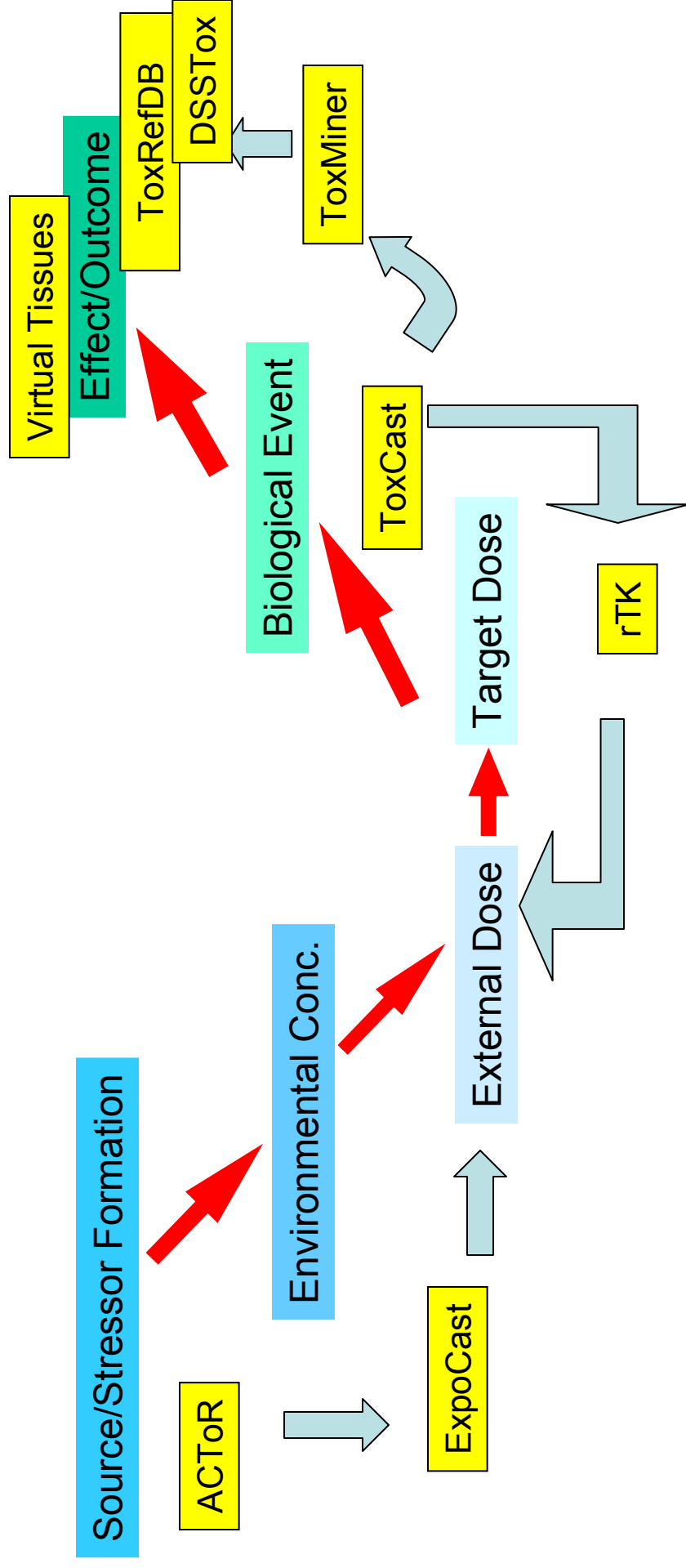
- Novascreen
- Attagene
- BioSeek
- Cellumen
- CellzDirect
- NCGC
- Gene assays
- Pathway assays





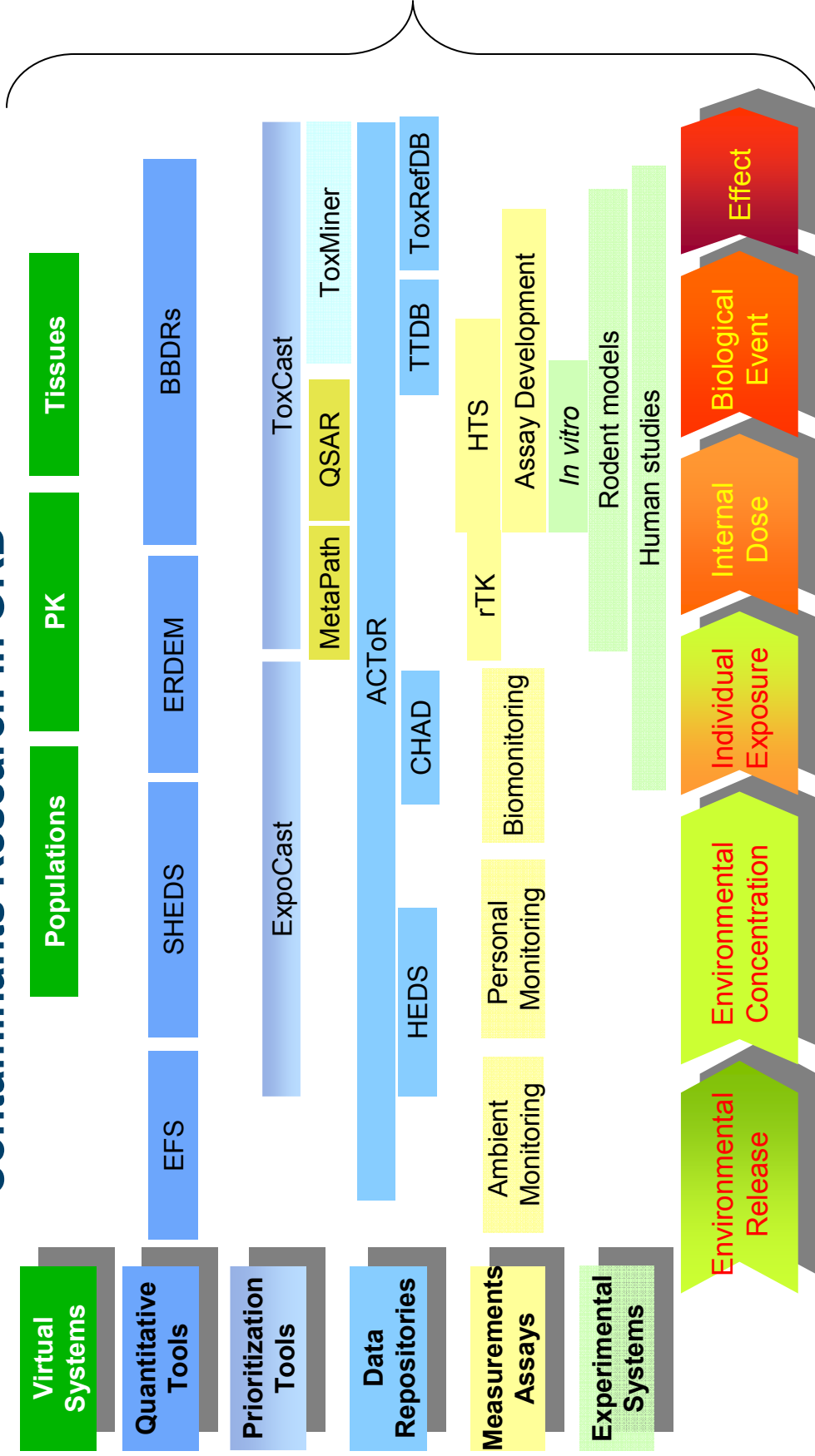
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Applying Computational Toxicology Along the Source to Outcome Continuum



Contaminants Research in ORD

Improved Risk Assessment & Risk Management



v-Tissues: EPA Vision

- Gain deeper understanding of molecular and cellular pathways to efficiently analyze thousands of environmental contaminants
- Predict dose-dependent human adverse outcomes (development/cancer/immune etc)
 - Use *in vitro* data
 - Accurately model low-dose response
 - Evaluate role of genomic variation in response
- Reduce dependence on animal testing

v-Tissues 2009: Vision

- How do we link molecular activity to phenotypes?
- What is the state-of-the-art in modeling disease phenotypes *in silico*?
- What role do tissues play in understanding/simulating clinical outcomes?
- How far can we get with “Virtual Tissues” using *in vitro* data (reducing dependence on animals)?
- How do we leverage transatlantic collaboration to get there?

v-Tissues 2009: Scope

- Which translational research gaps can be bridged by Virtual Tissues?
- What are the short-term products and long-term applications?
- What key computational and experimental hurdles must be overcome?
- How can EU and US international collaborations help achieve these goals?

