



# Toxicological Tipping Points & Cellular Stress

The 2<sup>nd</sup> Araucária Symposium on Cellular and Molecular Biology

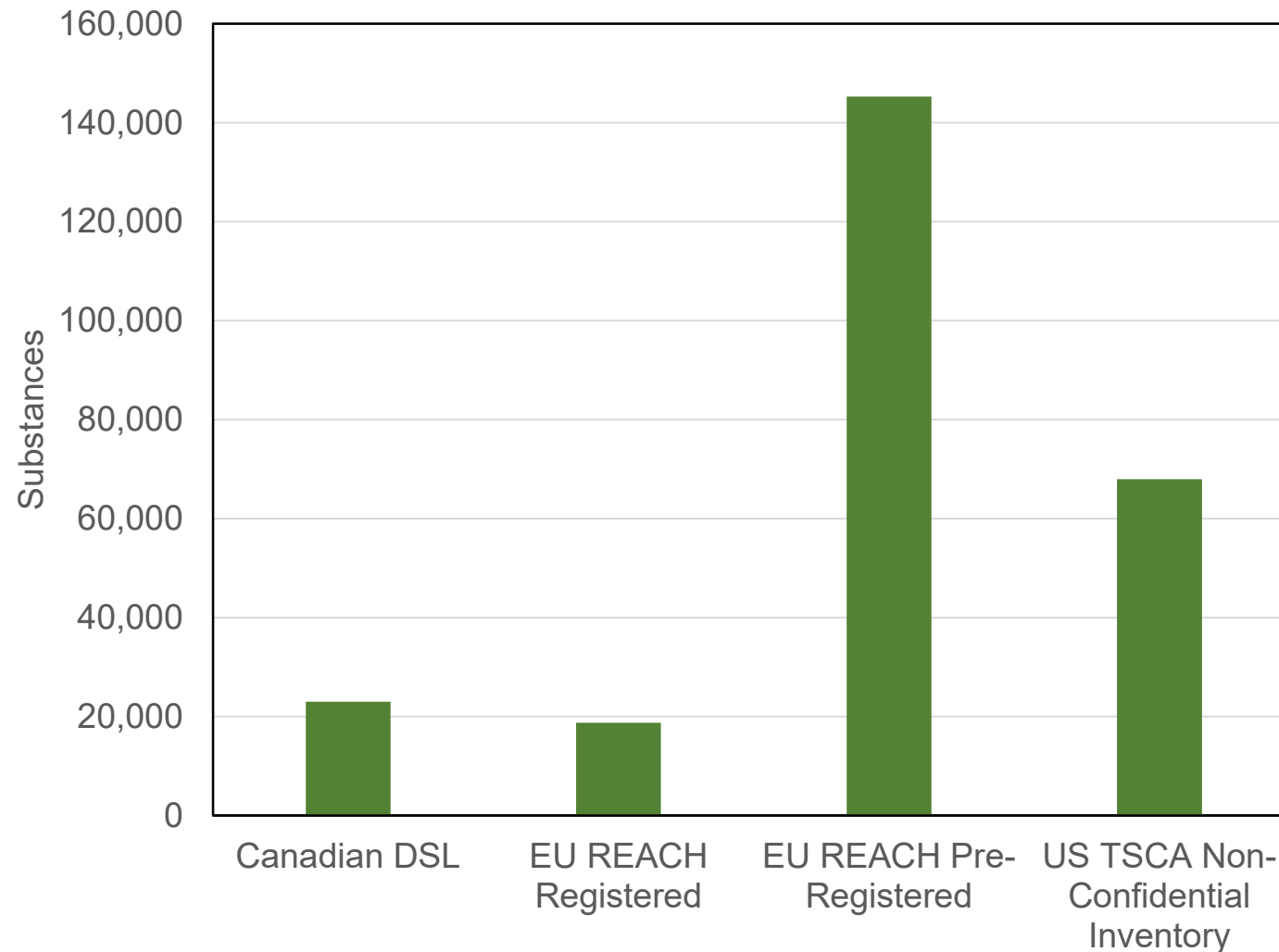
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Curitiba, Brazil

Imran Shah  
National Center for Computational Toxicology

*The views expressed in this presentation are those of the author[s] and do not necessarily reflect the views or policies of the U.S. Environmental Protection Agency.*



# Large number of chemicals in commerce





# Lack of Toxicity Data

## Toxicity Testing Strategies to Determine Needs and Priorities

Steering Committee on Identification of Toxic and Potentially Toxic  
Chemicals for Consideration by the National Toxicology Program

Board on Toxicology and Environmental Health Hazards

Commission on Life Sciences

National Research Council

- Major challenge is too many chemicals and not enough data
- Total # chemicals = 65,725
- Chemicals with no toxicity data of any kind = ~46,000

NATIONAL ACADEMY PRESS  
Washington, D. C. 1984

## The Toxicity Data Landscape for Environmental Chemicals

Richard Judson,<sup>1</sup> Ann Richard,<sup>1</sup> David J. Dix,<sup>1</sup> Keith Houck,<sup>1</sup> Matthew Martin,<sup>1</sup> Robert Kavlock,<sup>1</sup> Vicki Dellarco,<sup>2</sup> Tala Henry,<sup>2</sup> Todd Holderman,<sup>2</sup> Philip Sayre,<sup>2</sup> Shirlee Tan,<sup>4</sup> Thomas Carpenter,<sup>5</sup> and Edwin Smith<sup>6</sup>

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**OBJECTIVE:** Thousands of chemicals are in common use, but only a portion of them have undergone significant toxicologic evaluation, leading to the need to prioritize the remainder for targeted testing. To address this issue, the U.S. Environmental Protection Agency (EPA) and other organizations are developing chemical screening and prioritization programs. As part of these efforts, it is important to catalog, from widely dispersed sources, the toxicology information that is available. The main objective of this analysis is to define a list of environmental chemicals that are candidates for the U.S. EPA screening and prioritization process, and to catalog the available toxicology information.

**DATA SOURCES:** We are developing ACToR (Aggregated Computational Toxicology Resource), which combines information for hundreds of thousands of chemicals from > 200 public sources, including the U.S. EPA, National Institutes of Health, Food and Drug Administration, corresponding agencies in Canada, Europe, and Japan, and academic sources.

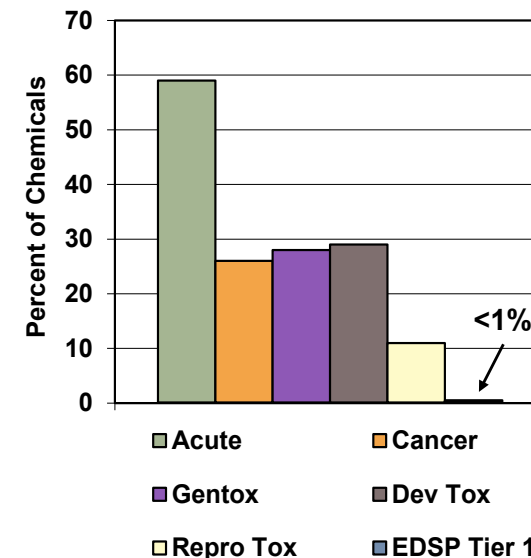
**DATA EXTRACTION:** ACToR contains chemical structure information; physical-chemical properties; *in vitro* assay data; tabular *in vivo* data; summary toxicology calls (e.g., a statement that a chemical is considered to be a human carcinogen); and links to online toxicology summaries. Here, we use data from ACToR to assess the toxicity data landscape for environmental chemicals.

**DATA SYNTHESIS:** We show results for analysis as part of the U.S. EPA ToxCast and medium-production-volume chemical water contaminants.

**CONCLUSIONS:** Approximately two-thirds of chemicals have some toxicity data available. About one-quarter of chemicals have evaluation data from sources such as the U.S. Risk Information System, and the National Toxicology Program. **KEY WORDS:** ACToR, carcinogenicity, reproductive, toxicity. *Environ Health Perspect* 115:103–110 (2007). Available via <http://ehpnet1.niehs.nih.gov/docs/2007/115-103-110/journal.html> [Online].

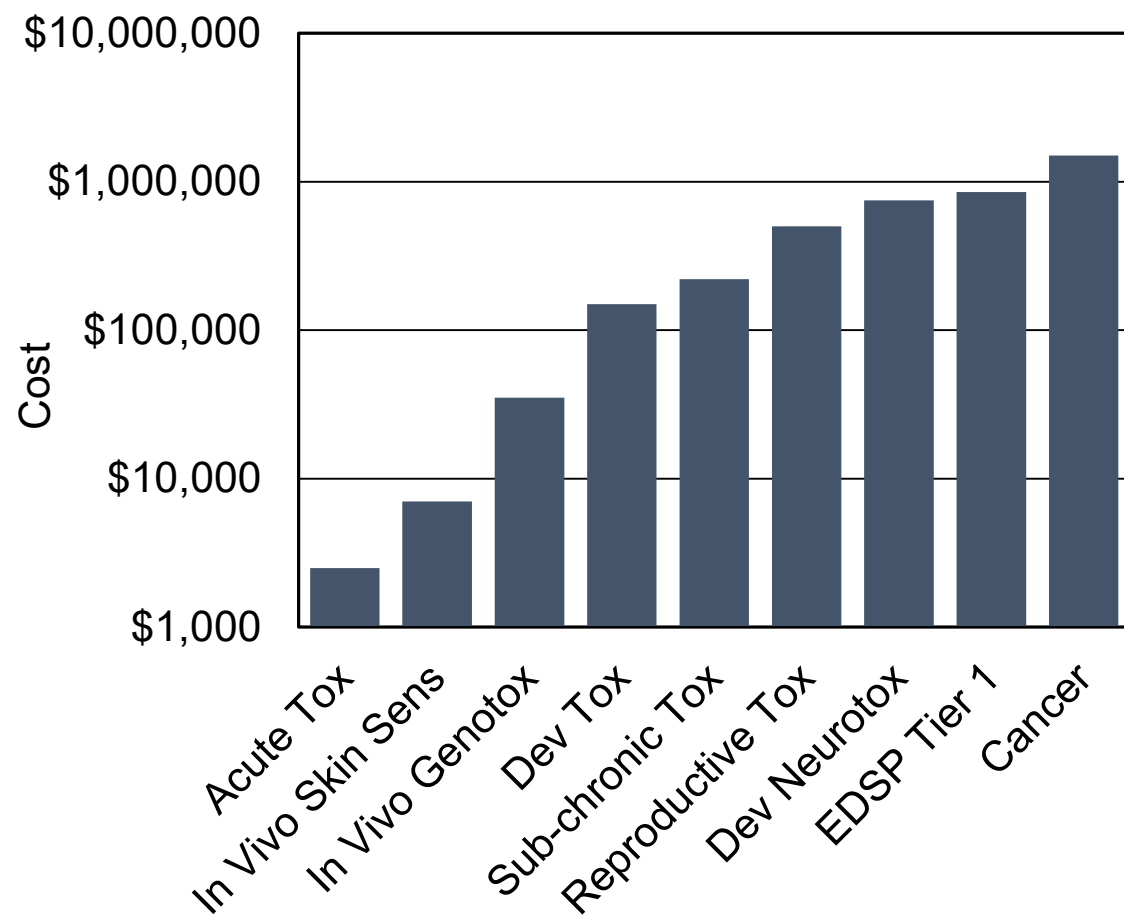
The U.S. Environmental Protection Agency (EPA) has a significant interest in developing more efficient and informative determination approaches in part of the large number of chemicals under its jurisdiction. Ultimately, it would be ideal to characterize the toxicologic profile of all chemicals in use in the United States. However, the size of this chemical inventory (in excess of 75,000 chemicals, with an estimated number in the Toxic Substances Control Act (TSCA 1976) inventory of 61,000 chemicals) makes this goal too large using current approaches to toxicity testing that rely on extensive animal testing, cost millions of dollars, and 2–3 years per chemical. The International Scientific Institute/Health and Environmental Sciences Institute (IHSI/HESI) released several reports describing focused, tier-based approaches for testing of agricultural chemicals, which ultimately lead to the use of fewer chemicals (Barton et al. 2006; Carmichael et al. 2006). The National Research Council

(Howard 2006). The European Union's Registration, Evaluation, and Authorization of Chemicals (REACH) program has recently released its first set of registered substances, which contains > 140,000 entries (REACH 2008). The exact number of chemicals in use is, in a sense, unknowable because it depends on where one sets the threshold of use and because use changes over time. The major point is that the number is relatively large and that only a relatively small subset of these chemicals have been sufficiently well characterized for their potential to cause human or ecologic toxicity to support regulatory action. This "data gap" is well documented (Allanson





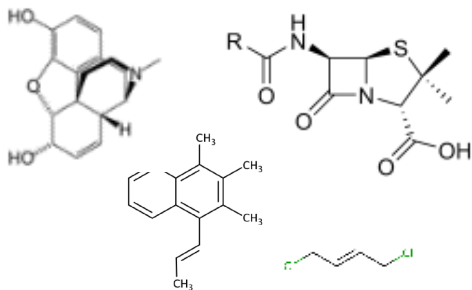
# Costs of Traditional Toxicity Testing





# How can high-throughput approaches help?

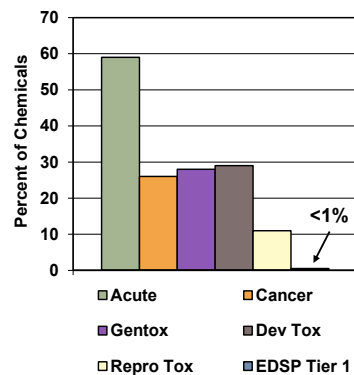
Number of Chemicals  
/Combinations



Comprehensive Tox  
Evaluation

DevTox  
Skinsens  
ImmunoTox  
MGR  
RepeatDoseTox  
AcuteTox  
Genotox  
2yrCarc

Limited Data

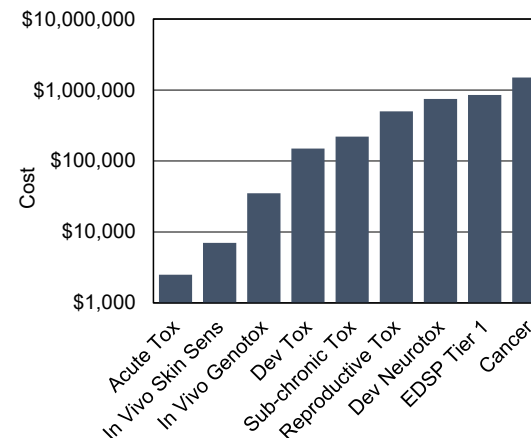


Modified from Judson *et al.*, EHP 2009



High-throughput  
Screening

Economics



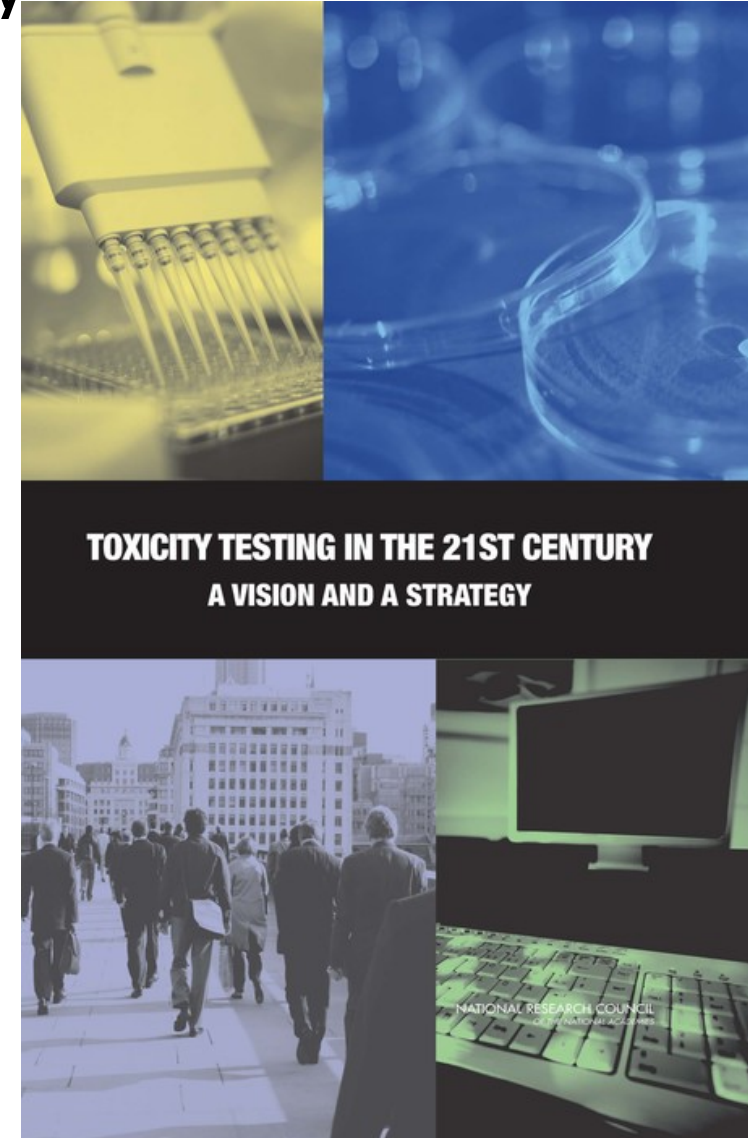


# Toxicity Testing in the 21<sup>st</sup> Century

*“Toxicity testing is approaching such a scientific pivot point. It is poised to take advantage of the revolutions in biology and biotechnology. Advances in toxicogenomics, bioinformatics, systems biology, epigenetics, and **computational toxicology** could transform toxicity testing from a system based on whole-animal testing to one founded primarily on in vitro methods ....”*

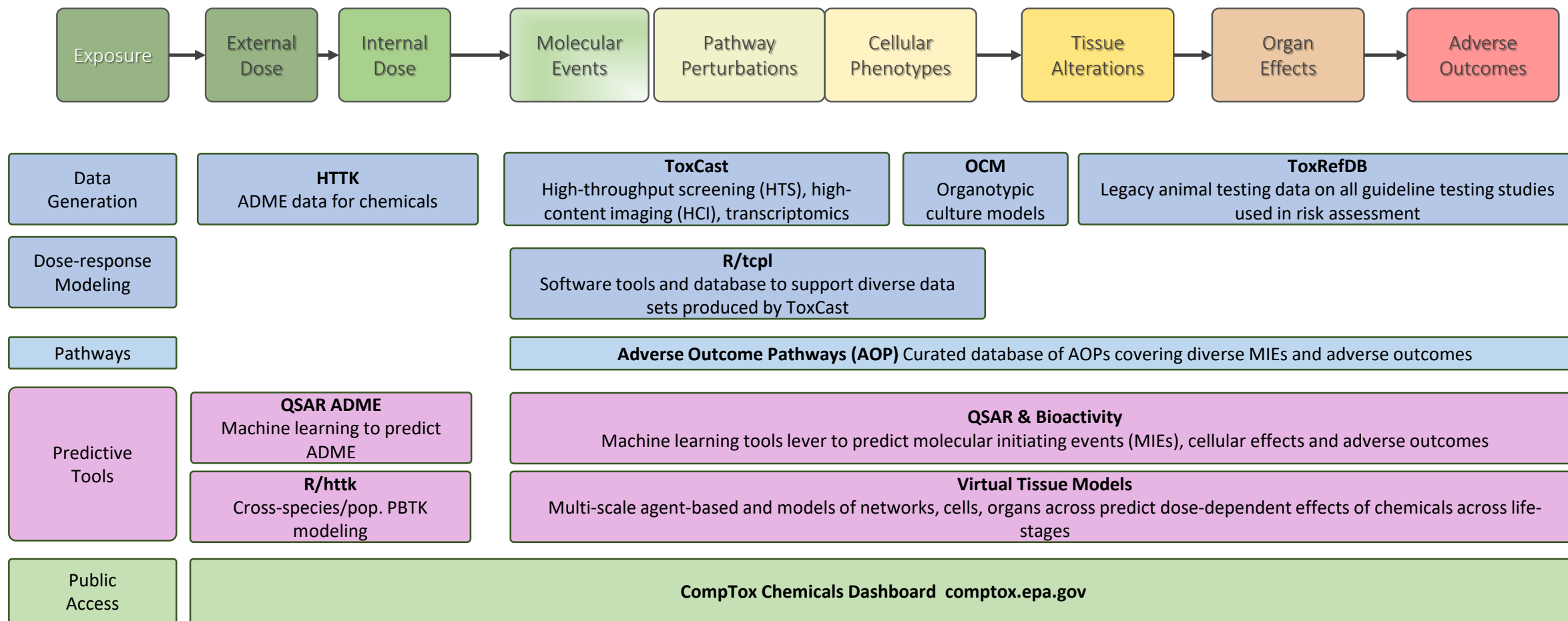
*NRC 2007*

- Three key components:
  - **High-throughput screening (HTS)** using *in vitro* assays to evaluate the molecular and cellular effects of thousands of chemicals
  - **Toxicity pathways**, when sufficiently perturbed or beyond adaptive capacity, lead to adverse health outcomes
  - **Dose-response modeling and *in vitro* to *in vivo* extrapolation** to estimate risk



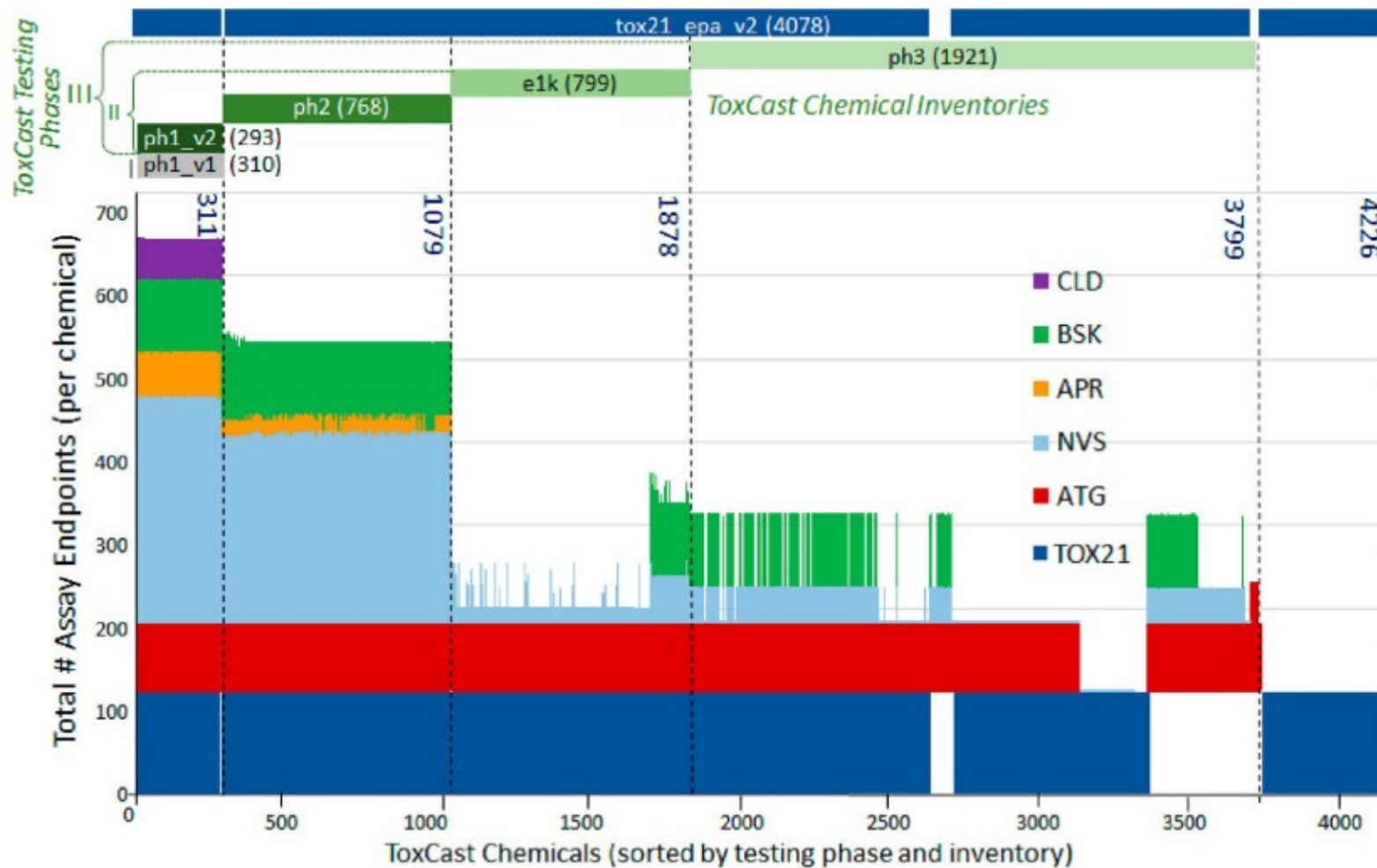


# Computational Toxicology





# ToxCast: 10 years later ...



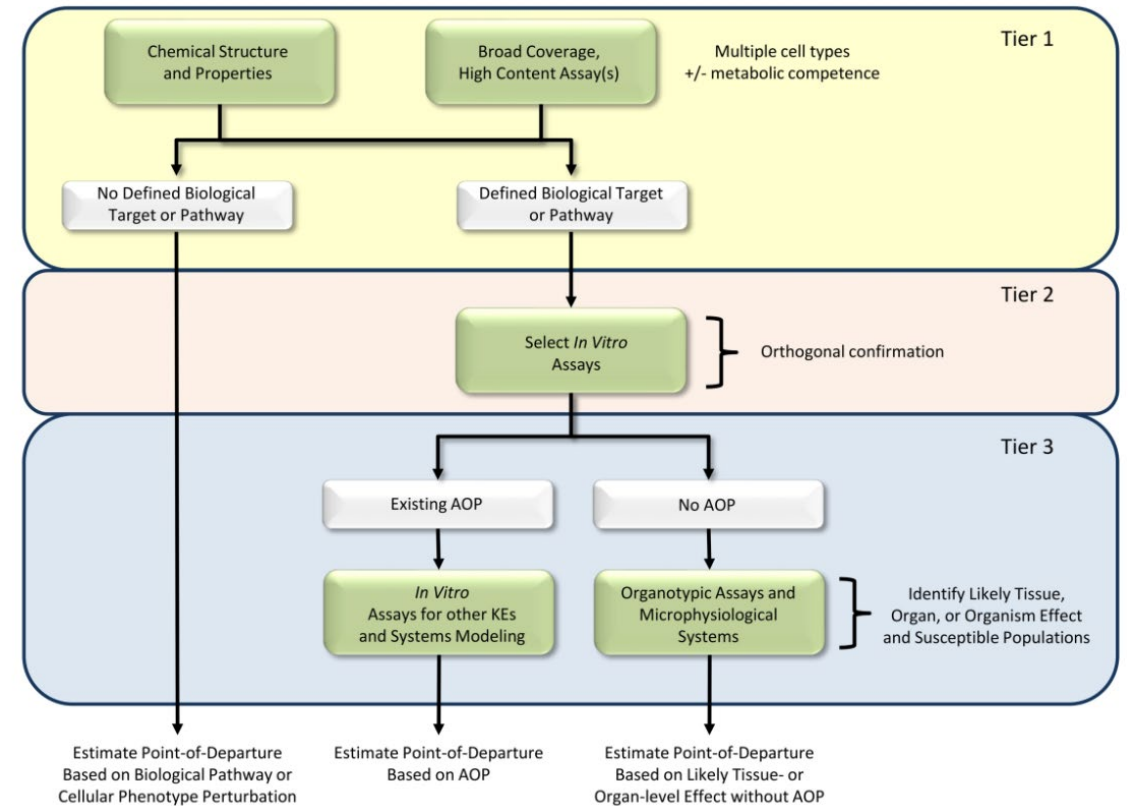
X chemicals  
Y assays  
Z genes  
A cells  
119 Publications



# Computational Toxicology: Future

- Expanded biological coverage to screen thousands of additional chemicals
  - High-content imaging (HCI)
  - High-throughput transcriptomics (HTTr)
- Tiered-testing to determine whether ...
  - Chemicals are potent activators of specific molecular initiating events (MIEs). Use adverse outcome pathway (AOP) framework identify adverse outcomes (AO) associated with MIE. Estimate point of departure (POD) using additional *in vitro* data and systems modeling
  - Chemicals are not potent but promiscuous. Estimate POD using pathway, cellular phenotype perturbations or other approach.
  - Chemicals are inactive.
- Chemical potency values derived from *in vitro* studies are quantitatively extrapolated to *in vivo* doses using toxicokinetic modeling

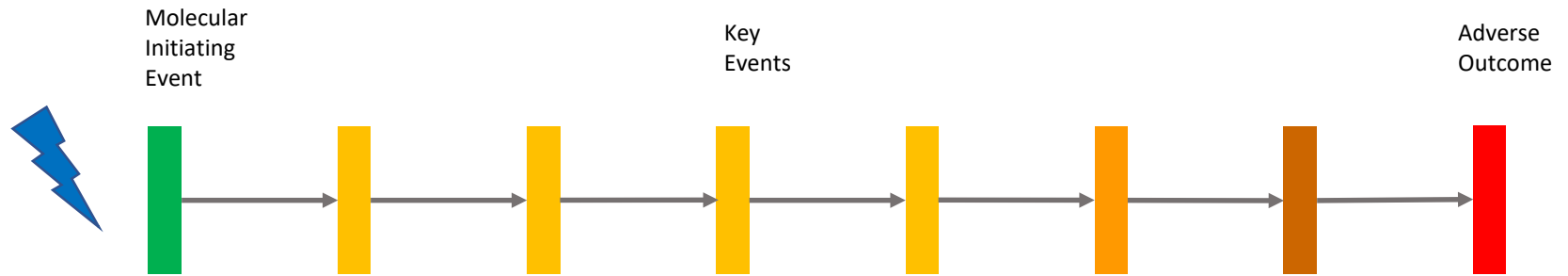
The next generation blueprint of computational toxicology at the U.S. Environmental Protection Agency



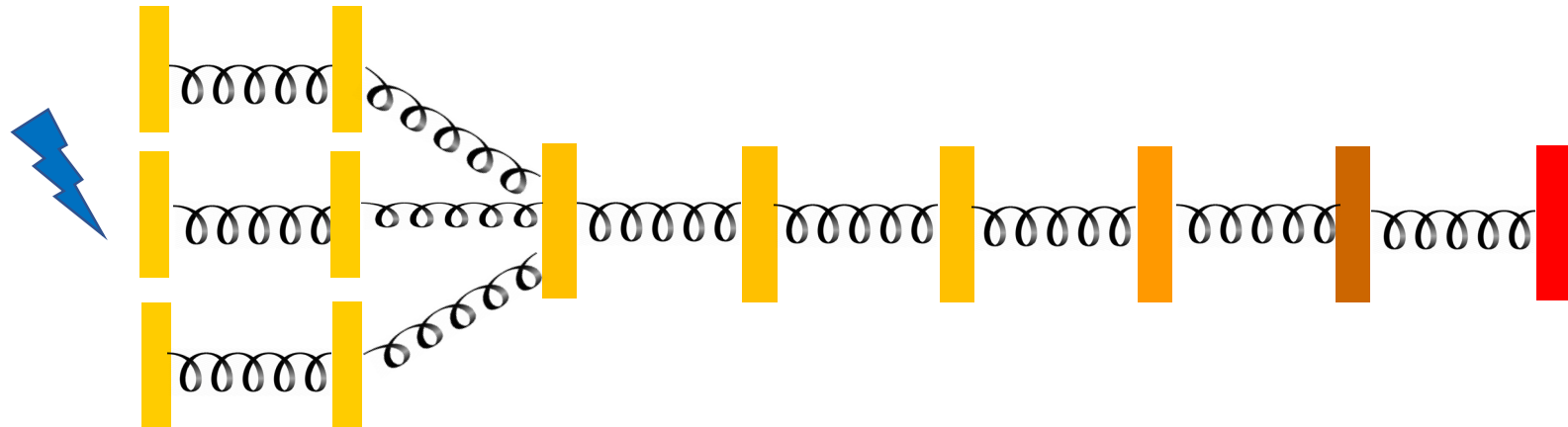


# How do chemical-induced perturbations propagate to adverse outcomes?

A. Activating MIE produces domino effect that results in adverse outcome (AOP)



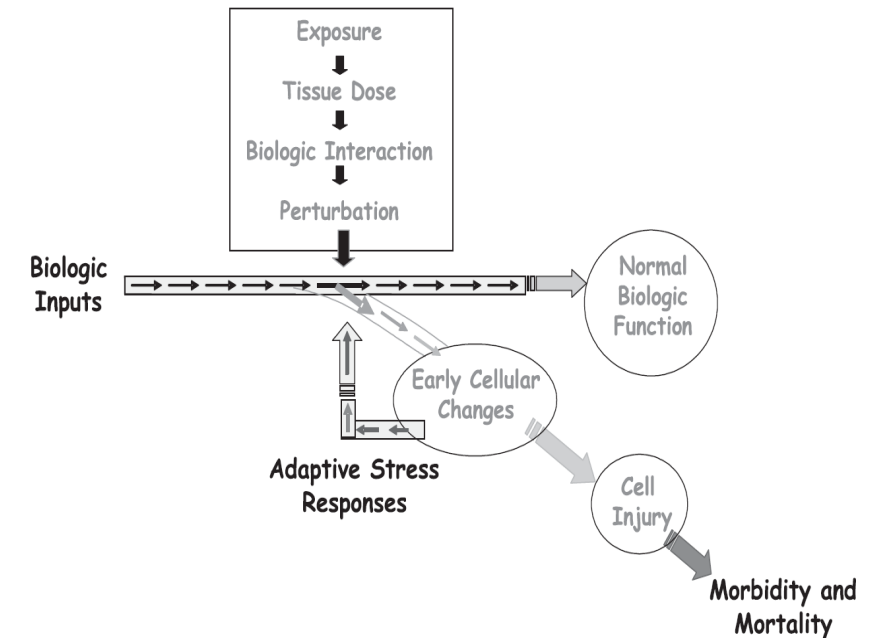
B. Multiple stress-response pathways are activated resulting in homeostatic adaptation. If perturbation exceeds "tipping point" then adverse outcome produced





# Toxicological Tipping Points

- Biological systems are resilient and adapt to environmental perturbations
- Tipping points are dose-dependent transitions in the system from normal to abnormal functions
- Key questions:-
  1. What type of *in vitro* data are suitable for identifying tipping points?
  2. How can we use these data to define dose-dependent transitions?
  3. Can we use tipping points as POD for risk assessment?



Krewski, Daniel, Daniel Acosta Jr, Melvin Andersen, Henry Anderson, John C Bailar 3rd, Kim Boekelheide, Robert Brent, et al. "Toxicity Testing in the 21st Century: a Vision and a Strategy." *Journal of Toxicology and Environmental Health. Part B, Critical Reviews* 13, no. 2-4 (February 2010): 51-138.

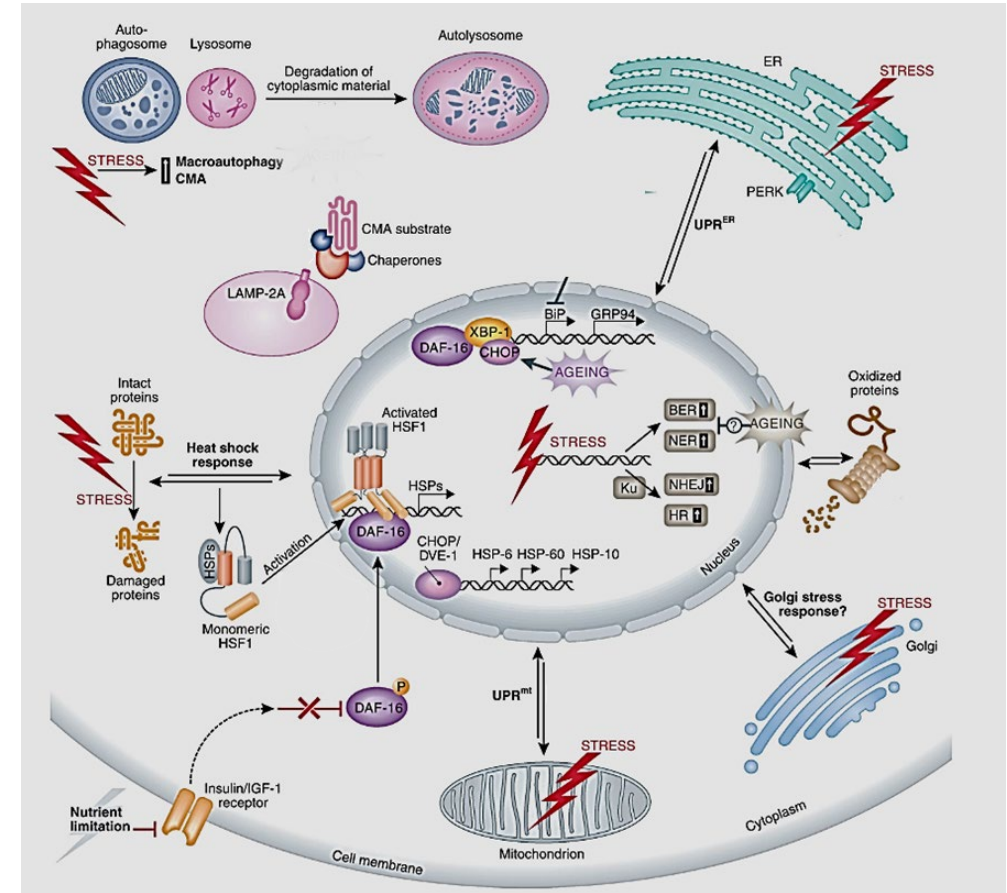


What type of data are suitable  
for analyzing tipping points?



# Cellular stress responses

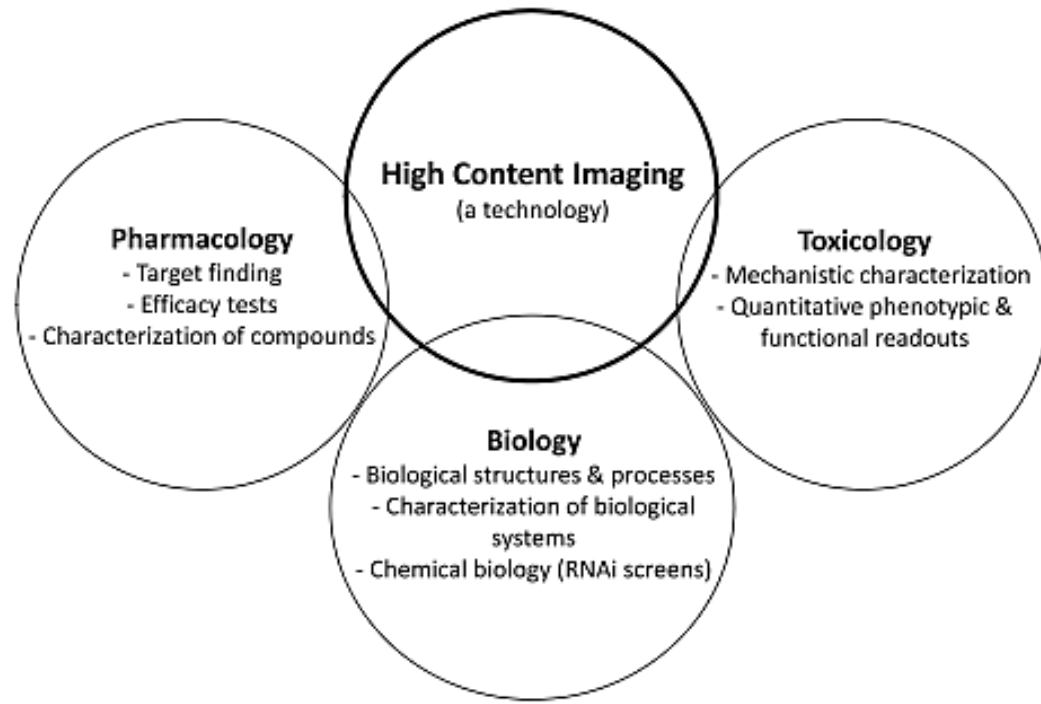
- Multiple stress-activated pathways are activated to counter chemical effects to maintain homeostasis:-
  - Heat-shock response
  - Ubiquitin-proteasome system
  - Endoplasmic reticulum (ER) stress
  - Mitochondrial stress
  - Lysosomal stress
  - DNA damage response
- There is cross-talk between stress-response pathways
- Overwhelming cellular adaptive capacity leads to cell death / autophagy





# High Content Imaging (HCI)

A powerful technology to interrogate the phenotypic state of cells by analyzing single cell data



## Examples

Cell source:

Primary cells, cell lines, co-cultures, stem cells, 3D cell models, lower organisms (e.g. *C. elegans*, zebrafish)

Readouts:

Reporters / dyes / biosensors

Cell/compartments:



Spatial imaging:

2D/3D Tracing Co-localization Segmentation

Imaging targets:

- DNA / RNA
- Transcription factors
- Enzymes and receptors
- Metabolites
- Potential differences, pH



# Study Design: Stress responses using HCl

Cell model: Human hepatoma HepG2

## Treatments:

Test chemicals: 978

Controls: (-) DMSO; (+) CCCP, Taxol

Conc: 0.39, 0.78, 1.56, 3.12, 6.24, 12.5, 25, 50, 100, 200  $\mu$ M

Duration: 1, 24 and 72 h.

Reps: 2 on plate

## Assay: High-content imaging (HCI)

Stress Kinase (SK): c-Jun

Oxidative Stress (OS): H2AX

Mitochondrial function (MMP): MitoTracker Red

Mitochondrial mass (MM): MitoTracker Red

Mitotic arrest (MA): PH3

Cytoskeletal stability (Mt): Tubulin

Cell cycle arrest (CCA): Hoechst 33342

Nuclear size (NS): Hoechst 33342

Cell number (CN): Hoechst 33342

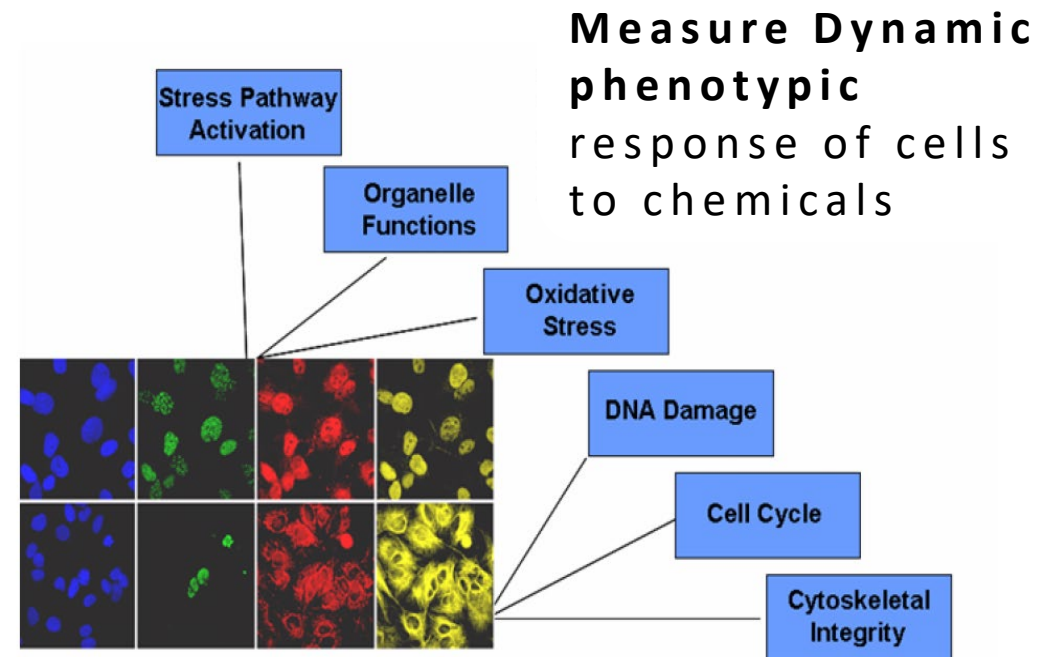
## Data:

~400 plates

~100,000 wells

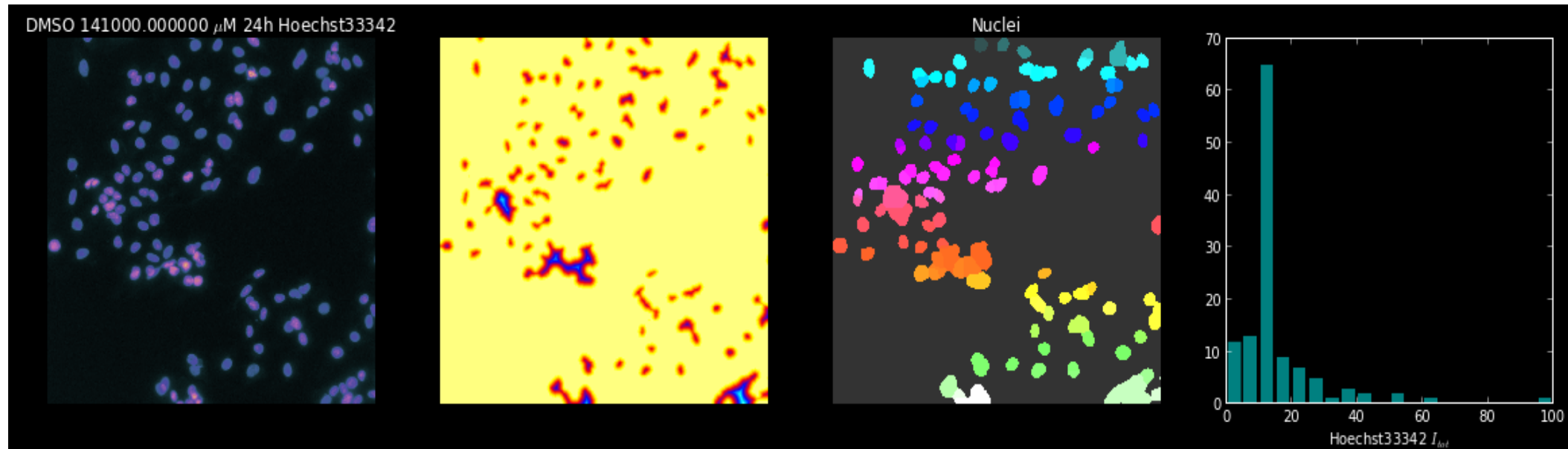
~2,400,000 images

~30,000 chemical-conc-time-response points





# Processing HCI data



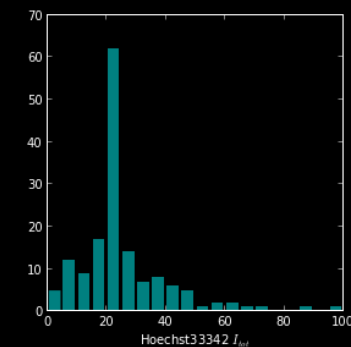
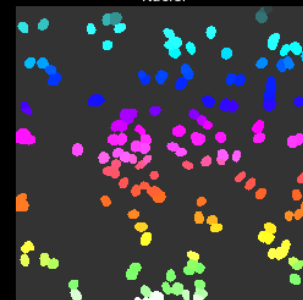
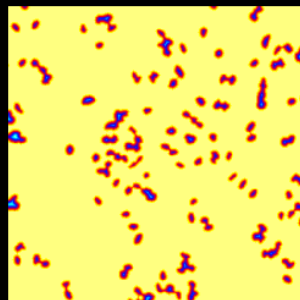
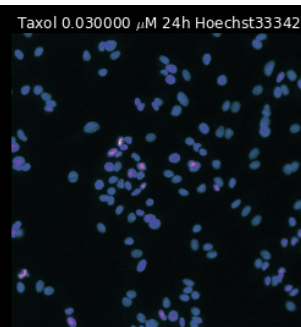
Raw Image (Hoechst)  $\longrightarrow$  Intensity Analysis  $\longrightarrow$  Object Identification  $\longrightarrow$  Nuclear intensity distribution

Image analysis and cell level feature feature extraction conducting by Cyprotex Inc. (proprietary software)



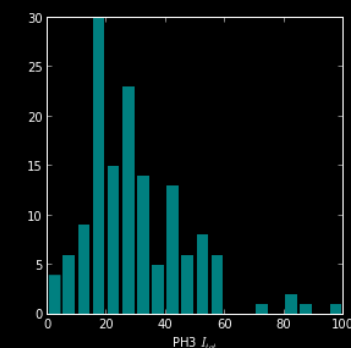
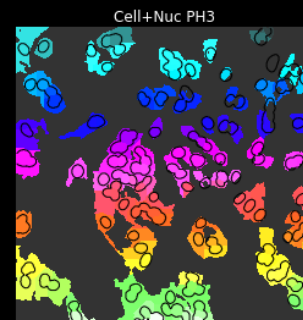
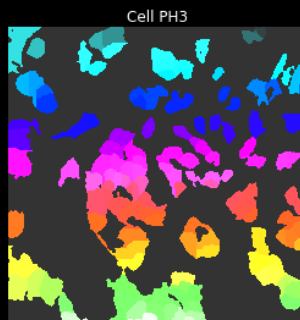
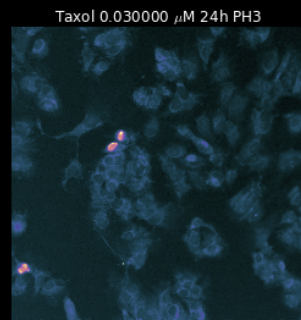
Taxol 0.03uM

Hoechst33342



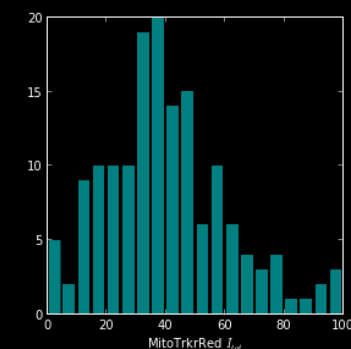
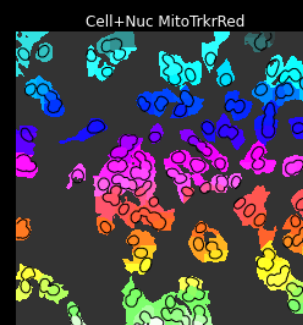
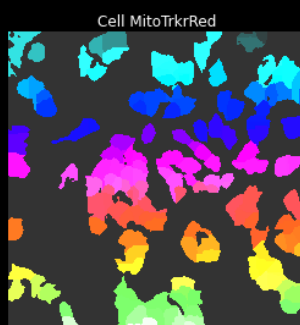
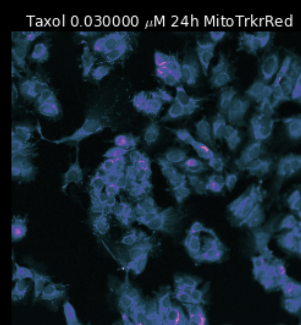
nuclear size (**NS**)  
cell cycle arrest (**CCA**)  
cell number (**CN**)

Phospho-Histone3



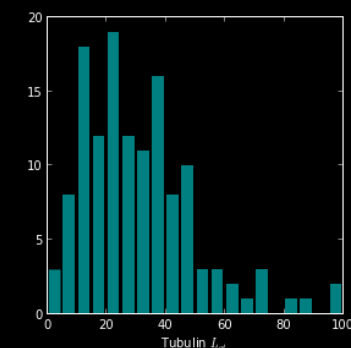
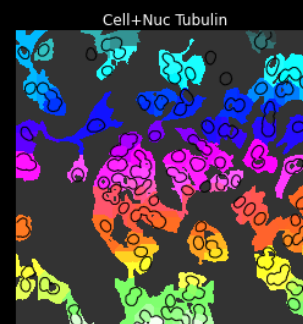
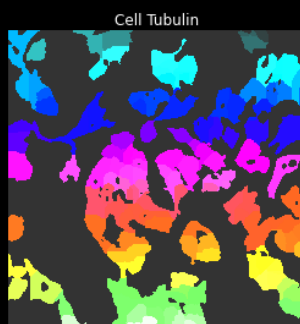
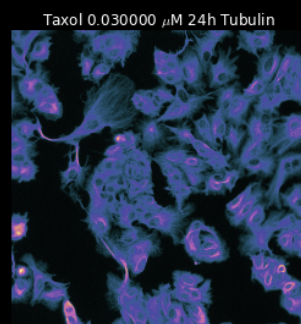
mitotic arrest (**MA**)

MitoTracker Red



mitochondrial mass (**MM**)  
mitochondrial membrane potential (**MMP**)

Phospho-Tubulin



microtubules (**Mt**)

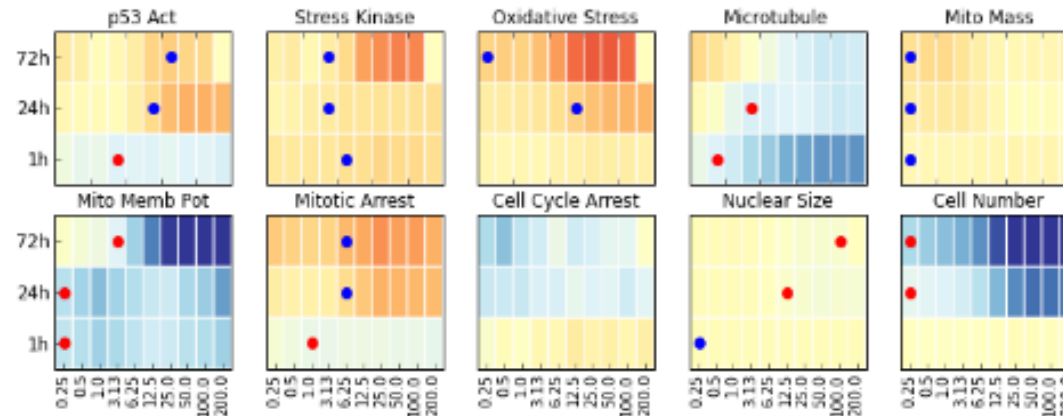
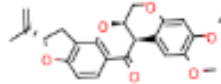
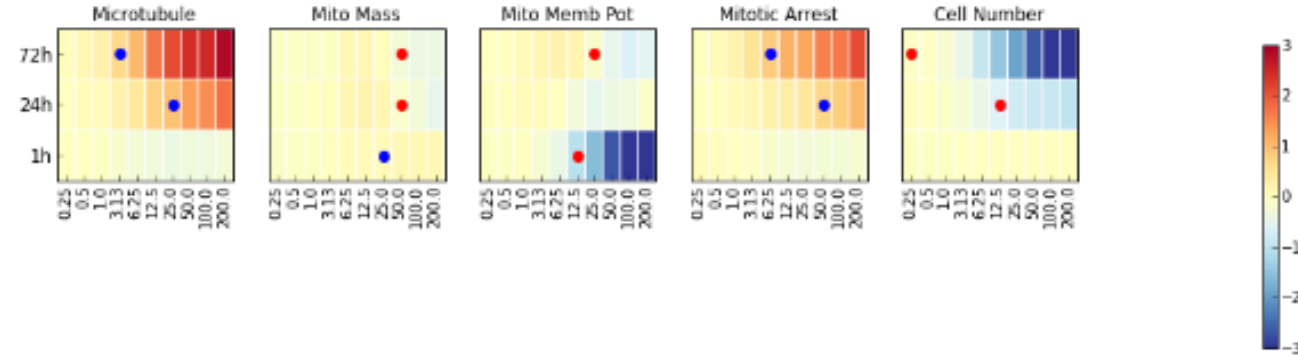
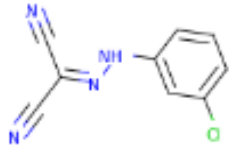


# HCI to bioactivity profiles

- Plate-level median-normalization – fold-change from background
- Report  $\log_2(\text{fold change})$ : decrease (BLUE), increase (RED) or no effect (YELLOW)
- Multi-dimensional bioactivity profile – “deviation from normal state” of HepG2 cells



# Mitochondrial disruptors

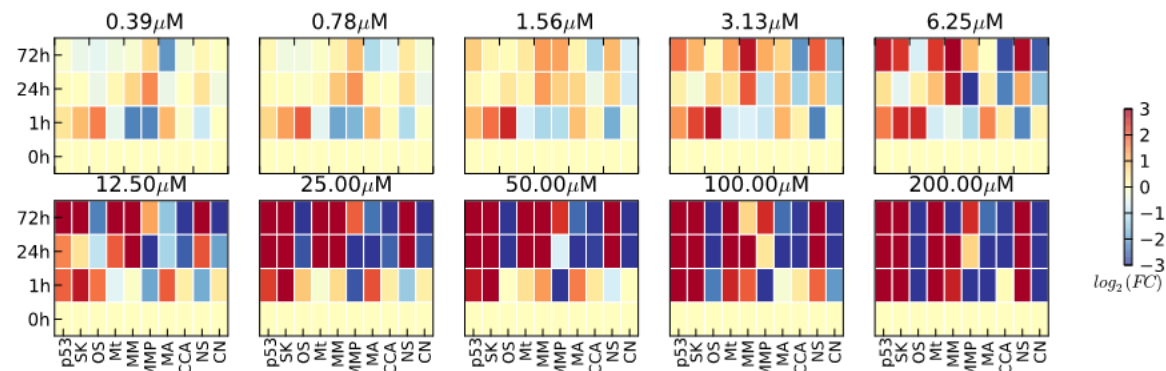


- log2(fold change): decrease (BLUE), increase (RED) or no effect (YELLOW)
- Bioactivity profile: “deviation from normal state” of HepG2 cells

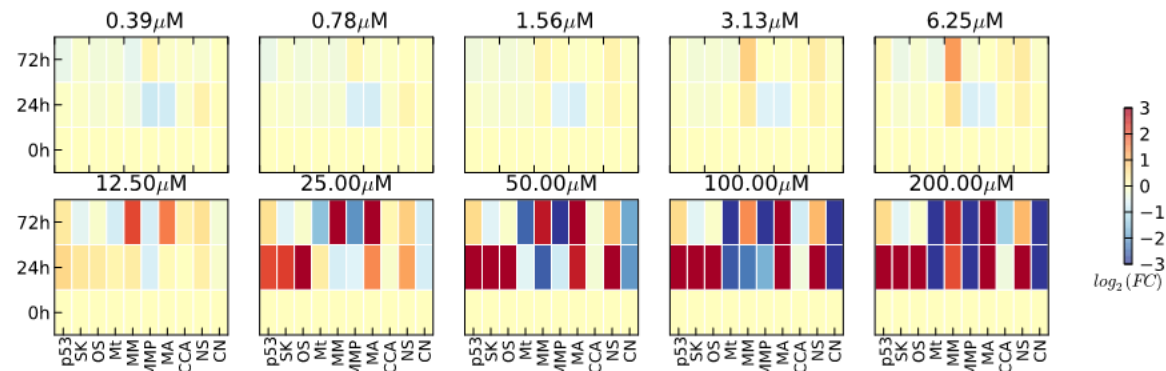
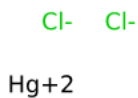


# Complex perturbations

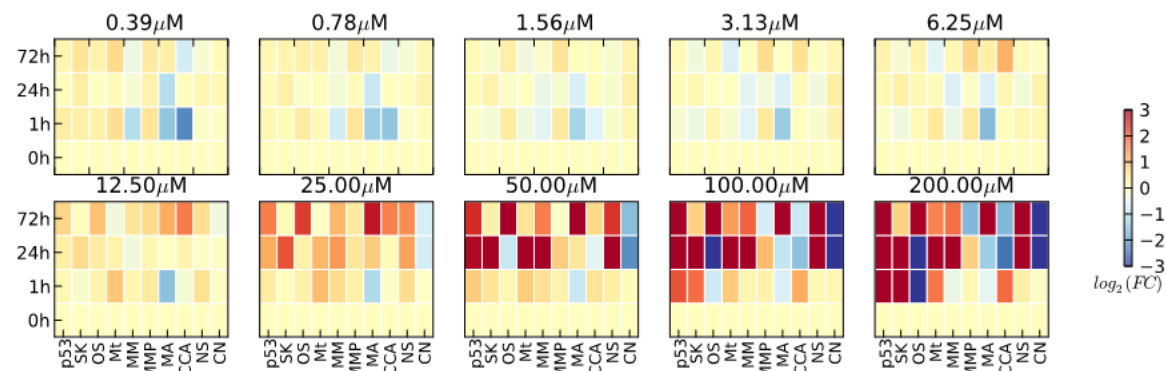
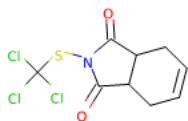
Fluazinam  
79622-59-6  
TX000777



Mercuric chloride  
7487-94-7  
TX007182



Captan  
133-06-2  
TX001434





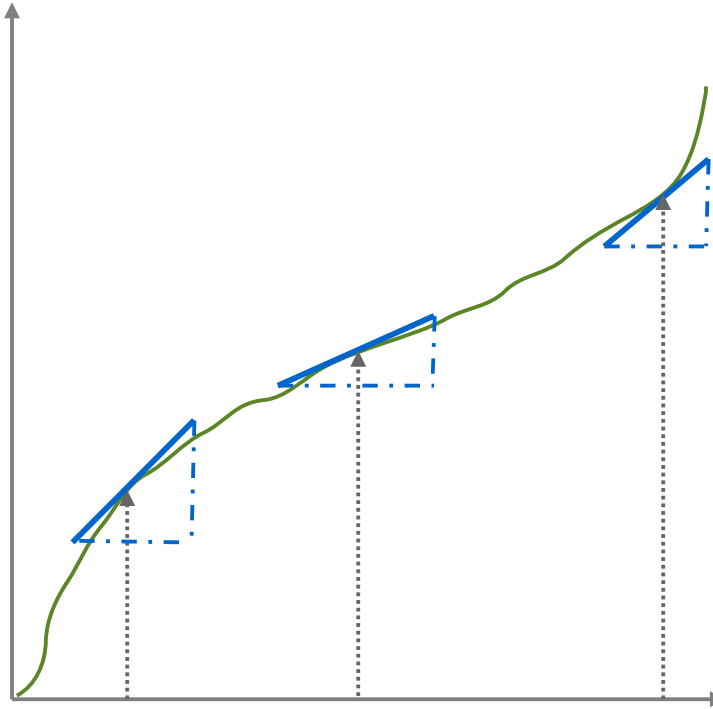
How can we use HCI data to  
identify tipping points?



Trajectories: system perturbation,  $|X|$



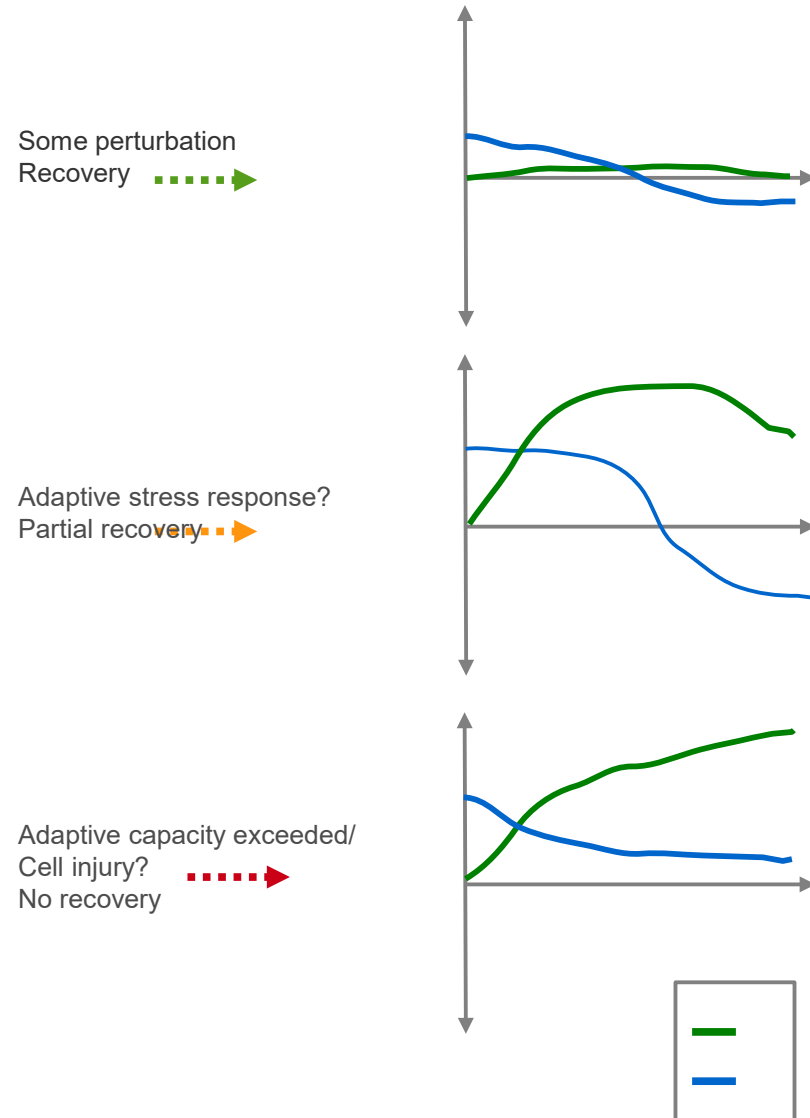
# Trajectories and pert. “velocity”



*Velocity – rate of change of aggregate system perturbation*

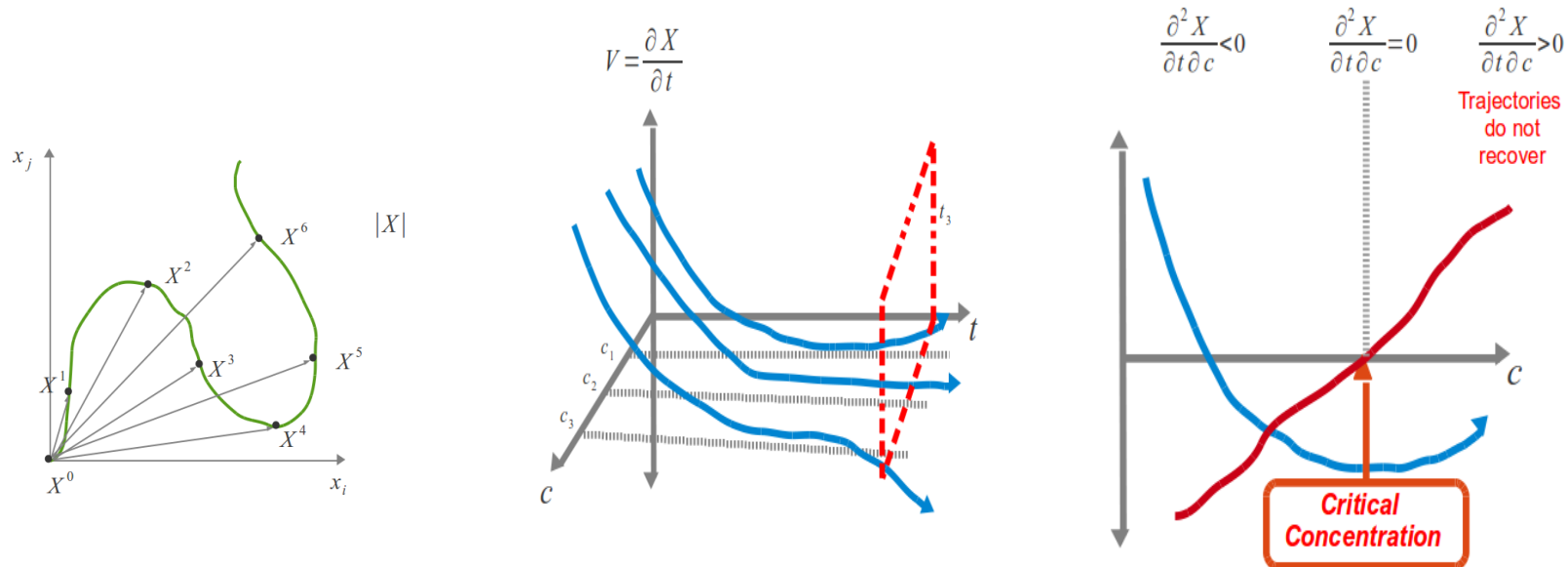


# Trajectories: perturbation & velocity

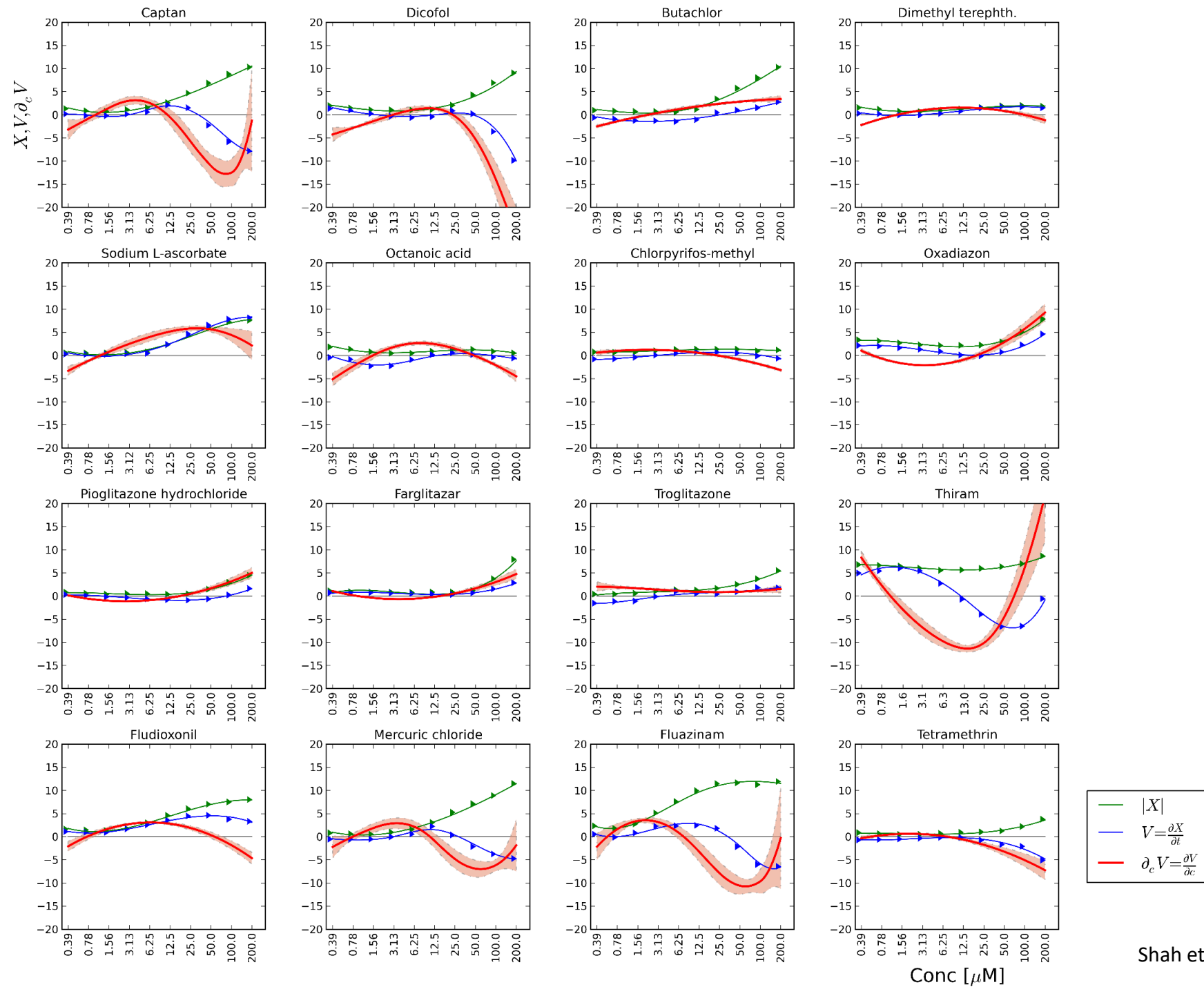




# Tipping point for system recovery

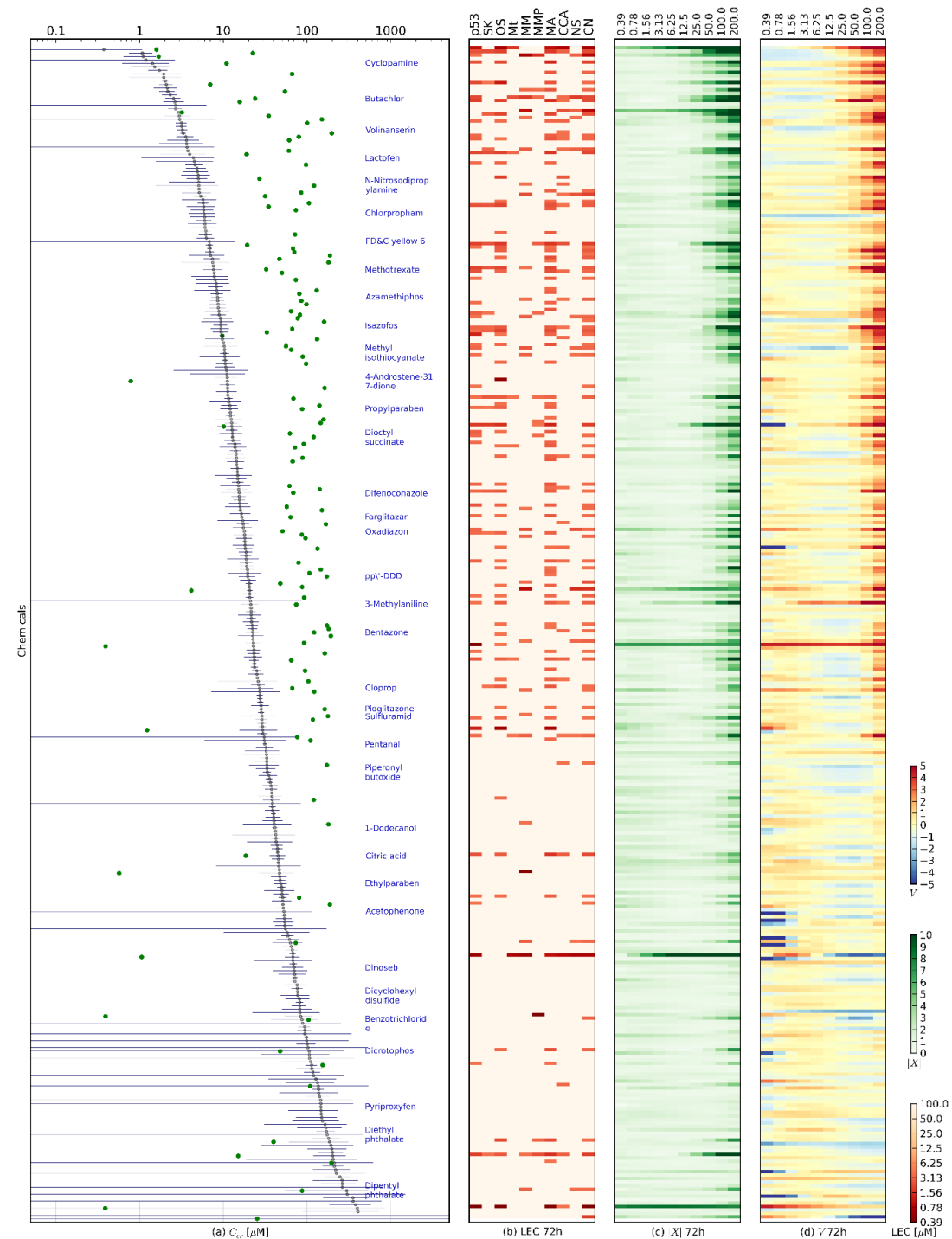








- ~340/967 chemicals had critical concentration
- ~170/967 chemicals always produced recovery
- Critical concentration ● more sensitive than cell loss





## Using ToxCast™ Data to Reconstruct Dynamic Cell State Trajectories and Estimate Toxicological Points of Departure

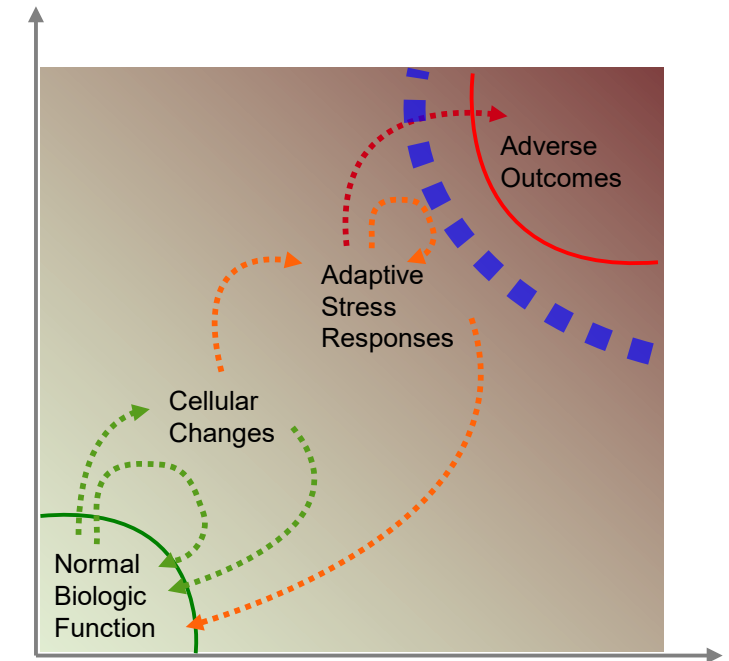
*Imran Shah,<sup>1</sup> R. Woodrow Setzer,<sup>1</sup> John Jack,<sup>2</sup> Keith A. Houck,<sup>1</sup> Richard S. Judson,<sup>1</sup> Thomas B. Knudsen,<sup>1</sup> Jie Liu,<sup>3</sup> Matthew T. Martin,<sup>1</sup> David M. Reif,<sup>4</sup> Ann M. Richard,<sup>1</sup> Russell S. Thomas,<sup>1</sup> Kevin M. Crofton,<sup>1</sup> David J. Dix,<sup>1</sup> and Robert J. Kavlock<sup>1</sup>*

<sup>1</sup>National Center for Computational Toxicology, Office of Research and Development, U.S. Environmental Protection Agency, Research Triangle Park, North Carolina, USA; <sup>2</sup>Department of Statistics, North Carolina State University, Raleigh, North Carolina, USA; <sup>3</sup>Oak Ridge Institute for Science Education (ORISE), U.S. Department of Energy, Oak Ridge, Tennessee, USA; <sup>4</sup>Department of Biological Sciences, North Carolina State University, Raleigh, North Carolina, USA



# Summary

- Biological systems are resilient
- The boundary between adaptation and adversity is complex !
- System trajectory analysis can reveal the tipping point between adaptation and adversity
- Tipping points can be calculated from HTS data to estimate critical concentrations of chemicals
- Additional work underway to evaluate utility of critical concentrations for risk assessment



Biologic Perturbations:

System Trajectories:

- Some perturbation/  
Recovery
- Adaptive stress response/  
Recovery
- Adaptive capacity exceeded/  
Cell injury/  
No recovery



# Acknowledgements and Questions

Todor Antonijevic  
Woody Setzer  
Thomas Knudsen  
Rusty Thomas

