

Filling in the Gaps: The role functional genomics can play in 21st century toxicology for environmental risk assessment



NAS Workshop: The Promise of Genome Editing Tools to Advance Environmental Health Research January 10-11, 2018

Keith Houck National Center for Computational Toxicology

The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA



All Understand Why We Need to Innovate In This Space...

Number of Chemicals/Combinations



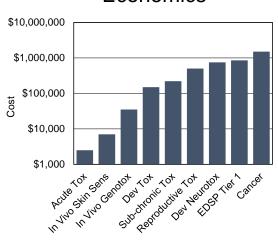
Lack of Data 70 60 50 50 40 40 10 0 Acute Cancer Gentox Dev Tox Repro Tox EDSP Tier 1

Modified from Judson et al., EHP 2010

Ethical Concerns

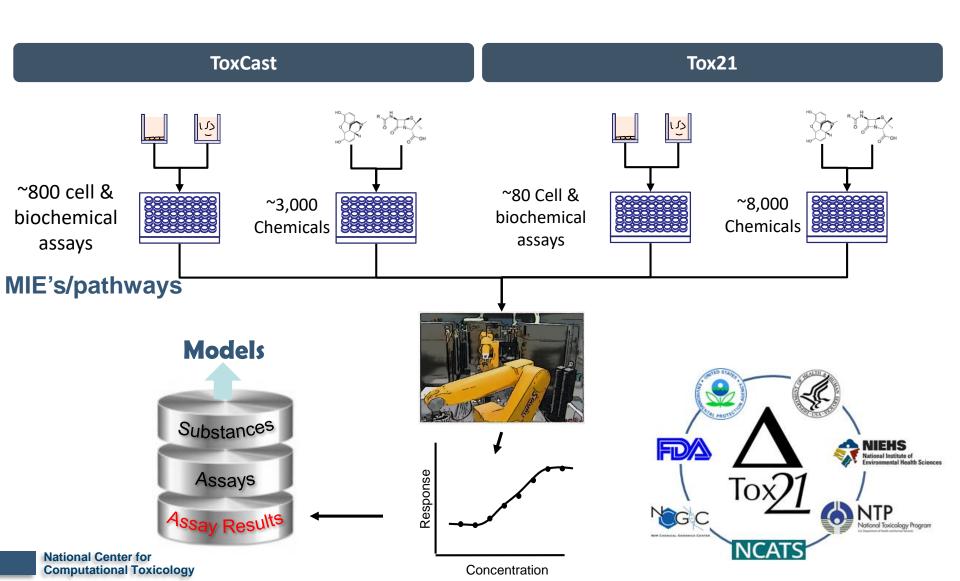


Economics





One Approach: High-Throughput Hazard Screening

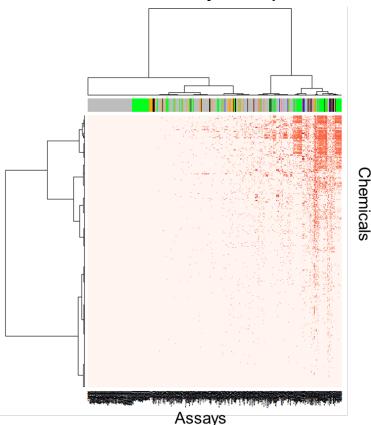


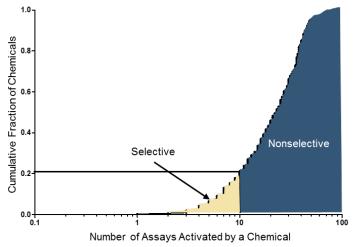
\$EPA

Environmental Protection

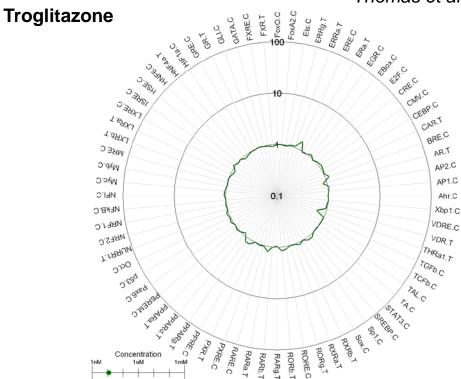
Promiscuous Chemical Response is the Rule

1000 chemicals/ 800 assay endpoints





Thomas et al., 2013

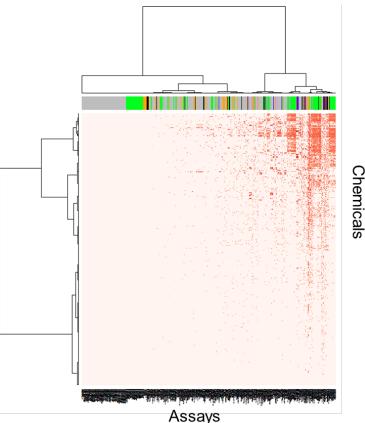


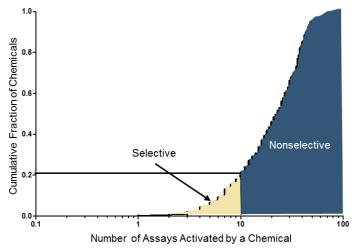
\$EPA

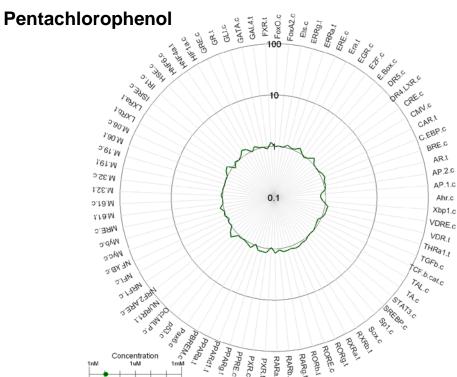
Promiscuous Chemical Response is the Rule

United States Environmental Protection Agency

1000 chemicals/ 800 assay endpoints



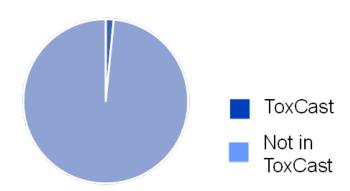




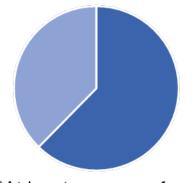


Beginning to Address Concerns for Increased Biological Coverage

Gene Coverage



Pathway Coverage*



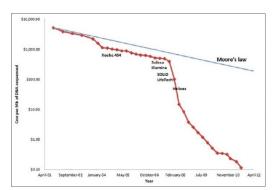
*At least one gene from

pathway represented

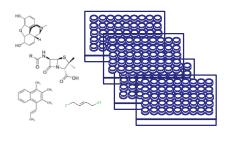
National Center for Computational Toxicology

High-throughput Genomics (HTTr)

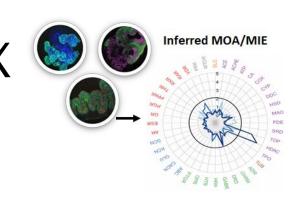




Thousands of chemicals



Multiple Cell Types

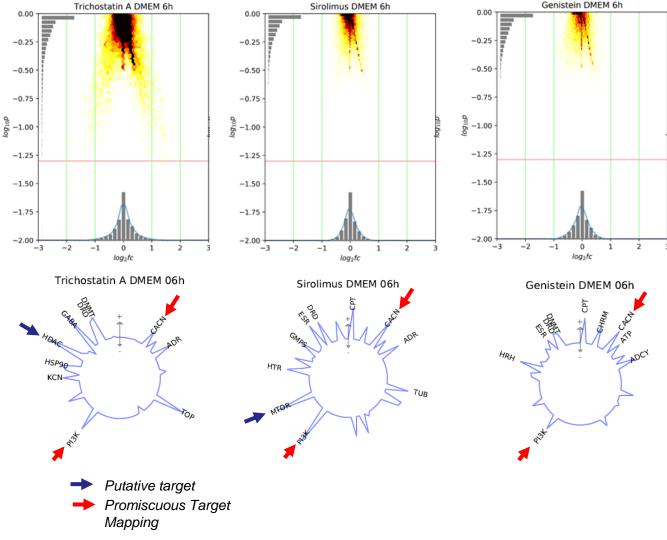


Requirements:

- Low cost
- Whole genome
- 384 well
- Automatable

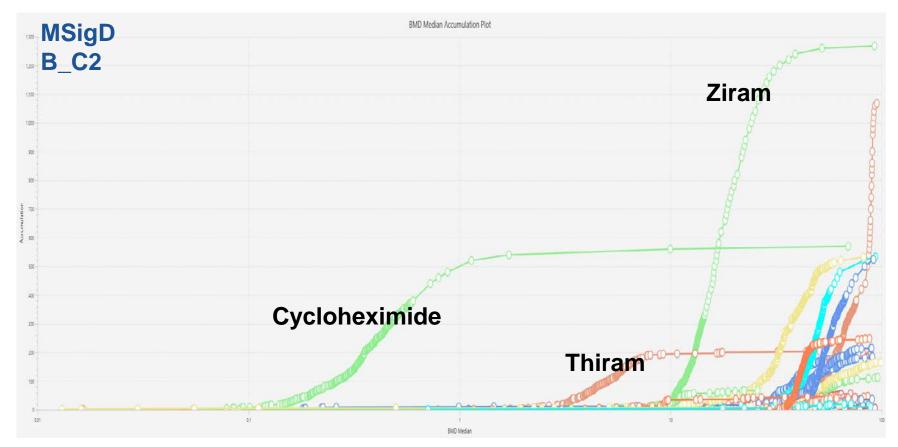


Connectivity Mapping Demonstrates Multiple Pathway Matches



- Differential gene expression observed with reference chemicals
- Putative targets identified using Connectivity Mapping
- Large degree of promiscuity of predicted targets observed
- Currently evaluating additional methods for MIE prediction

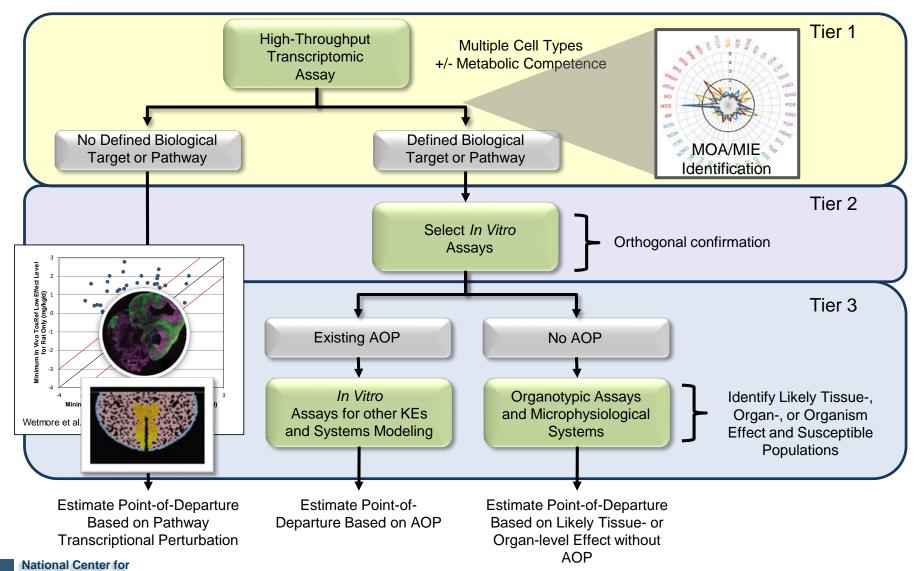
Pathway Potencies by BMD Analysis Environmental Protection



 Broad range of pathway level potency estimates and number of pathways affected across chemicals.



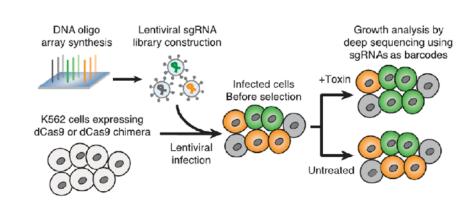
Framework for Integrating Hazard Components...





Functional genomics: Defining Relevancy

- Most chemicals have apparent polypharmacology—what is the critical/relevant MOA?
 - -Could use potency to define but this may not be linked to adversity
 - Transcriptomics is high content but function is generally inferred
- Functional genomics allows for bridging between genotype and phenotype
- Previously mostly used in prokaryotic systems such as S. cerevisiae
- Advent of CRISPR-Cas9
 opens door for higher
 throughput applications in
 mammalian cells

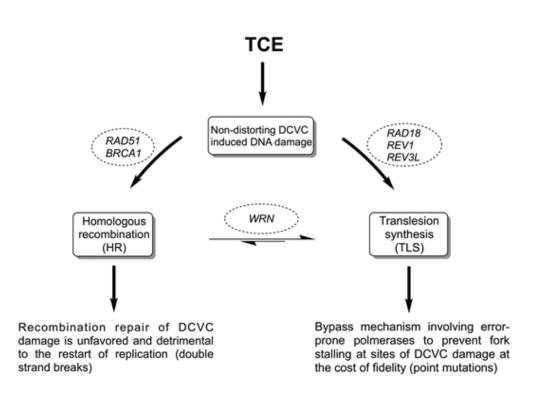


Gilbert et al., Cell, 2014



Environmental Application of Functional Genomics

- Trichloroethylene (TCE) metabolite study in yeast mutants (n=4607)
- Genome-wide profiling of yeast mutants (n=4607) identified the error-prone translesion synthesis (TLS) pathway conferring sensitivity to TCE metabolite DCVC
- Results were confirmed in a eukaryotic system using DT40 avian kidney cells



De la Rosa et al., Toxicological Sciences, 2017



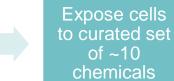
Pilot Project

- Collaboration between University of Florida (Chris Vulpe) and USEPA (NCCT, Keith Houck)
- Funded by USEPA SMARTi award to Keith Houck and Audrey Bone
- Goal of the project is to test the feasibility of using CRISPR-Cas9 genome editing in human cells for screening environmental chemicals in a functional genomics toxicology format



Experimental Design

Use CRISPR-Cas9 genome
editing
technology
with a
targeted, short
guide RNA
library (n=3675
genes) in X
cells





Evaluate results in context of putative mechanisms of cytotoxicity for each chemical



Chemical Selection

CION

Triclosan

Criteria

- –Mix of uses (pharmaceutical, pesticide, consumer, industrial)
- –Well-characterized mechanisms of cytotoxicity
 - Mitochondrial toxicity
 - DNA damage
 - Oxidative stress
 - Microtubule disruption
 - Proteosome inhibition
- -Known cytotoxic in Tox21/ToxCast assays without metabolic activation

- 11 chemicals
 - -Colchicine
 - -Triphenyltin chloride
 - -Triglycidyl isocyanurate
 - -Cytembena
 - -Propargite
 - -Octhilinone
 - -Triclosan
 - -Tralopyril
 - Dibutyltin dichloride
 - -Malachite green
 - Bisphenol A glycidyl methacrylate



Weaknesses

- Current approach limited to identifying genes that lead to reduction in cell viability
- Problems inherent to CRISPR tech
 - Unintended mutations
- "Pathway" analysis

- Larger perspectivesame problems as other HTT approaches
 - Metabolic capacity
 - -Toxicokinetic integration
- BUT functional aspects increases confidence in relevance of specific genes to adverse effects



Future Needs for Pathway to Integration of Functional Genomics into Environmental Risk Assessment

- More proof-of-concept and pilot studies with wellcharacterized chemicals
- Development of assay technology that facilitates expansion of functional endpoints beyond cell viability
 - -in vivo assays in organisms suited to HTT such as C. elegans or Danio rerio
 - -cell line engineering of pathway reporter lethality assays
- Dependent on continued integration of HTT into risk assessment paradigms

SEPA Other Uses of Genome Editing Tools

- Validation of HTTr signatures
- Validation of AOPs

Environmental Protection

- –What is the effect of knockdown/activation of MIE's/KE's to the adverse endpoint?
- -Can this help with the development of quantitative AOP's?
- Rapid generation of specific animal model MOAs
 - Sensitivity/resistance to specific MOA's
 - Modulation of critical ADME parameters
- Generation of sensitive population models in vivo and in vitro
 - Single genetic mutation disease models
 - Engineered ADME genetic variability
 - SNP recapitulation of sensitive populations from GWAS-type studies (complex)
 - -Validation of sensitivities identified through GWAS-type analysis



Thank You for Your Attention!

Acknowledgements

EPA Colleagues:

NCCT

Audrey Bone (*Bayer*)

Joshua Harrill

Imran Shah

Rusty Thomas

NCEA

Tina Bahadori

Jeff Dean

Jason Lambert

NIEHS

Allison Harrill

Univ. of Florida
Chris Vulpe
Abderrahmane Tagmount



EPA's National Center for Computational Toxicology