# **Omics** Applications

Case Study

Adam Biales

Great Lakes Toxicology and Ecology Division, US EPA, Duluth, MN

## Chemical Risk Assessment

#### Exposure:

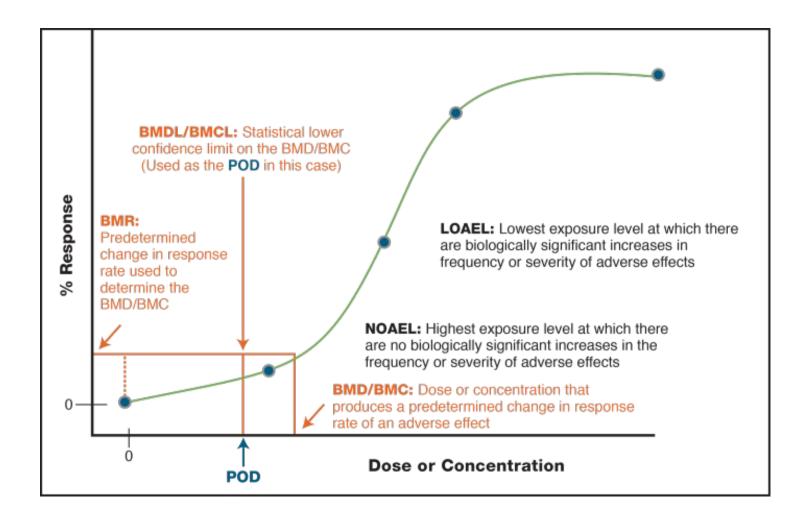
What concentrations occur in organisms or the environment?

### Hazard/Effect:

What concentrations cause adverse effects to exposed organisms?

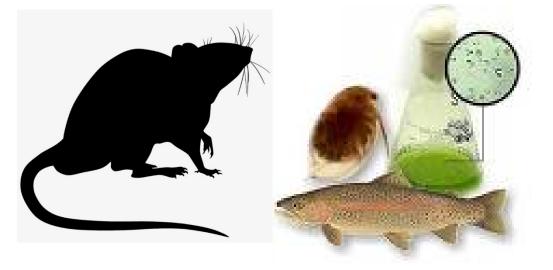
#### Safety:

At what concentration is there likely to be little or no hazard (adverse effects unlikely)?



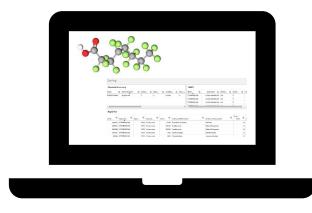
# Hazard/Safety Data

### Toxicity Testing



- Costly
- Time-consuming
- Animal intensive
- Lacking in mechanistic insight

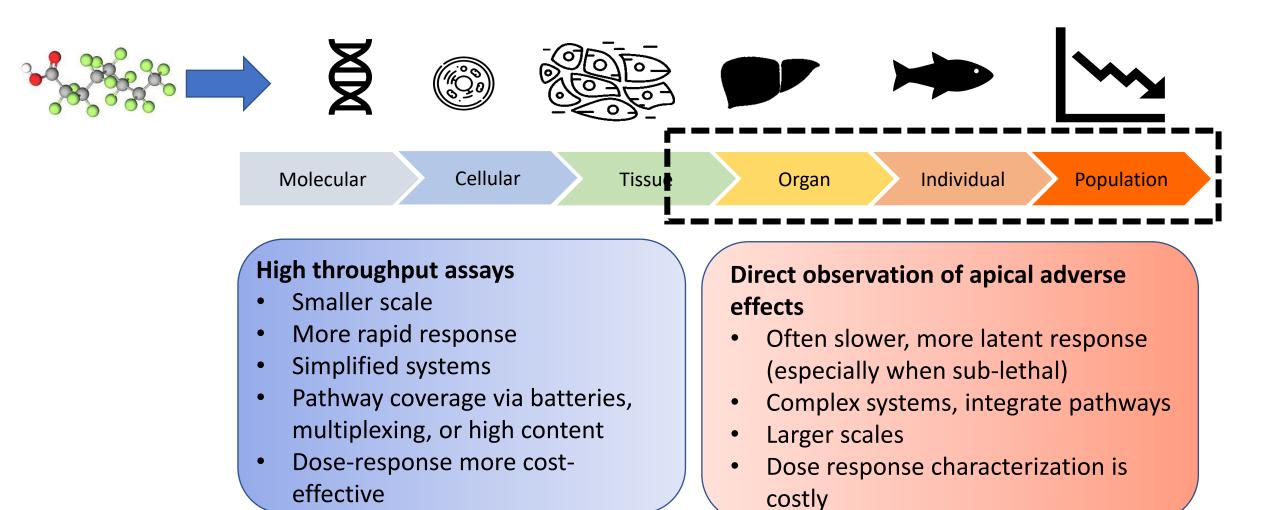
### **Structure-based Prediction**

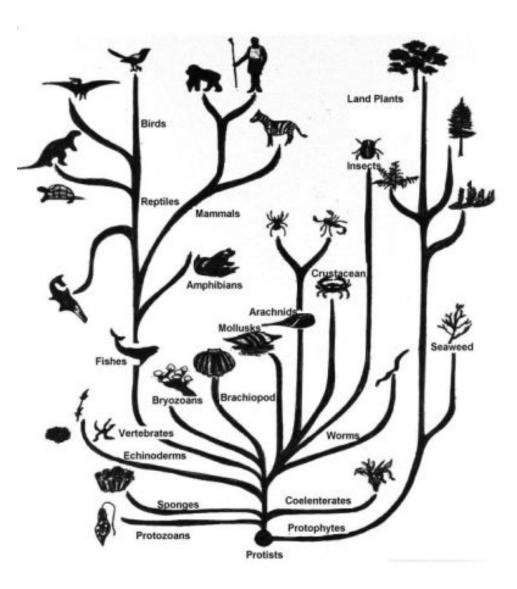


- Requires understanding about what chemical properties/structural features are associated with toxicity.
- Understanding of mechanism(s) of toxicity relevant to different structural groups.
- Traditional models don't work well for PFAS

# Approach – NAMs

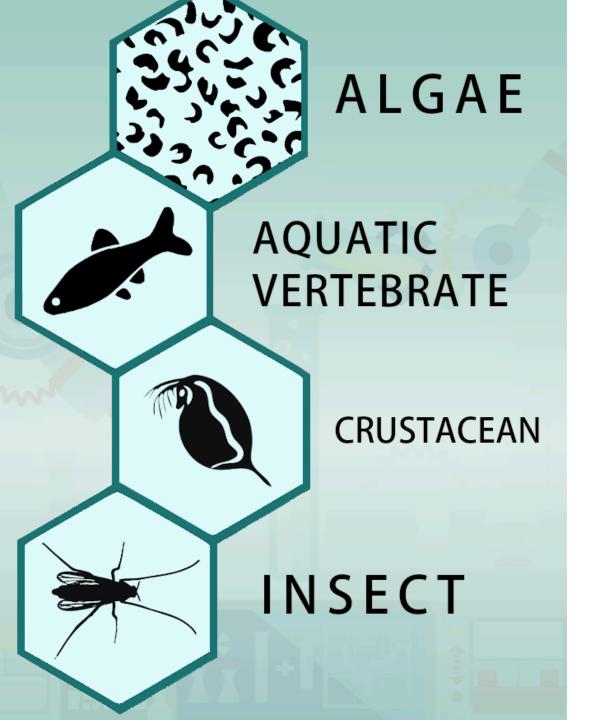
(New Approach Methodologies)





### **Ecotoxicology Perspective**

- Humans are just a tiny fraction of the biological diversity we are charged to protect.
- Many genes/pathways are conserved
- Unique physiology in other kingdoms, phyla, classes...
- How do we assure those pathways are covered?



High throughput assays for three major trophic levels of aquatic ecosystems

- Primary producers (e.g., algae)
- Primary consumers (e.g., zooplankton, aquatic inverts)
- Secondary consumers (e.g., fish)

Commonly used for GHS classification and labeling of chemicals for environmental hazard

Aquatic organisms highly vulnerable to exposure

#### **Eco HTP Assay Descriptions**





24 h exposure				
Control				
1   2   3   4   5   6   7   8   9   10   11   12     A   O <td></td>				

#### Phenotypic anchoring

- survival
- behavior
- Photo pigments

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Species	Guideline Test Method	Age at Start	Тетр
Daphnia magna	850.1010 Aquatic Invert Acute Toxicity	72-hour	20° C
Pimephales promelas	850.1075 Fish Acute Toxicity	24-hour	25° C
Chironomus dilutus	850.1790 Chironomid Sediment Toxicity	3 <sup>rd</sup> instar	20° C
Raphidocelis subcapitata	850.4500 Algal Toxicity	Log-phase	24° C

#### Exposures Design

- 1 ml deep 96-well plates
- 12 concentration 8 replicates per concentration
- 1 individual per well (algae ~5 x 10<sup>4</sup> cells/ml)
- 24-hour static exposures
- phenotypic endpoints assessed
  - animals: survival and behavior
  - algae: cell viability & division, photopigments
- then after homogenization, RNA extracted for transcriptomics

Species	Time to Load Plate	RNA Qty per Well
Daphnia magna	~45 minutes	~1000 ng
Pimephales promelas	~30 minutes	~1500 ng
Chironomus dilutus	~60 minutes	~900 ng
Raphidocelis subcapitata	~10 minutes	~300 ng

### Initial 10 Chemicals

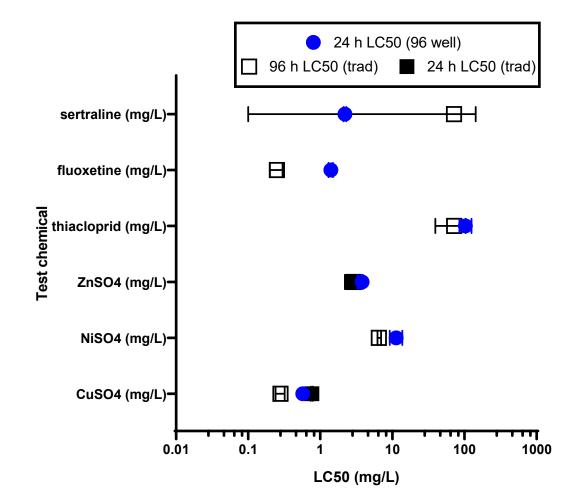


Metals CuSO4 NiSO4 ZnSO4 Selective Serotonin Reuptake Inhibitors (SSRI) Fluoxetine Paroxetine Sertraline

Neonicotinoids Clothianidin Imidacloprid Thiacloprid *Flupyradifurone* 

### Assay design

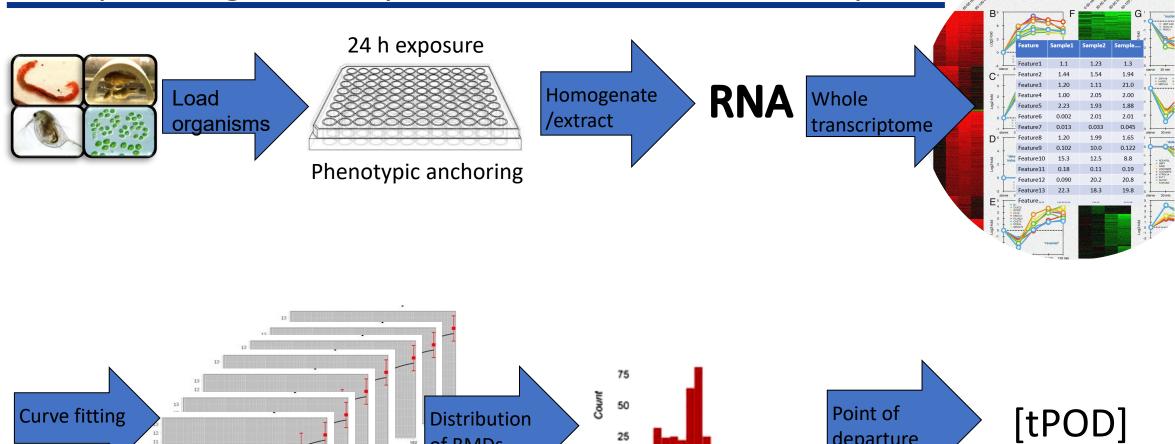
How does the 24 h, 96 well plate format compare with the traditional 24 or 96 h LC50 in a tank or beaker?



- Four of 10 chemicals were not toxic at the maximum concentration tested
- 24 h LC50s in 96 well plate format closely matched those in traditional tank/beaker format
- 24 h LC50s generally > 96 h LC50s as expected.
- 96 well format does not appear to be markedly altering overall sensitivity

#### Incorporating transcriptomics as assessment endpoint

of BMDs



25

0.0

**BMD** Express

10<sup>th</sup> centile BMD

12 24 38 48 69 12

BMD

departure



APCRA Case study: Transcriptomics-based PODs for Ecotoxicology

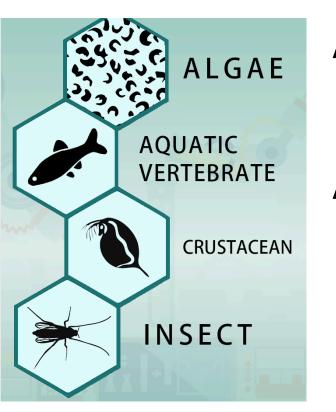
### 1. Generate transcriptomic PODs for $\approx$ 20 chemicals

- Initial focus on fathead minnow
- Parallel assays with additional taxa for future analyses

#### 2. Compare tPODs with available acute and chronic toxic toxicity data

### 3. Compare tPODs with in vitro-derived PODs

## Eco-HTTr Research at EPA



#### **Assay Optimization**

- How many replicate wells (animals)?
- How much genome coverage?
- Assay acceptance criteria?

### **Assay Evaluation**

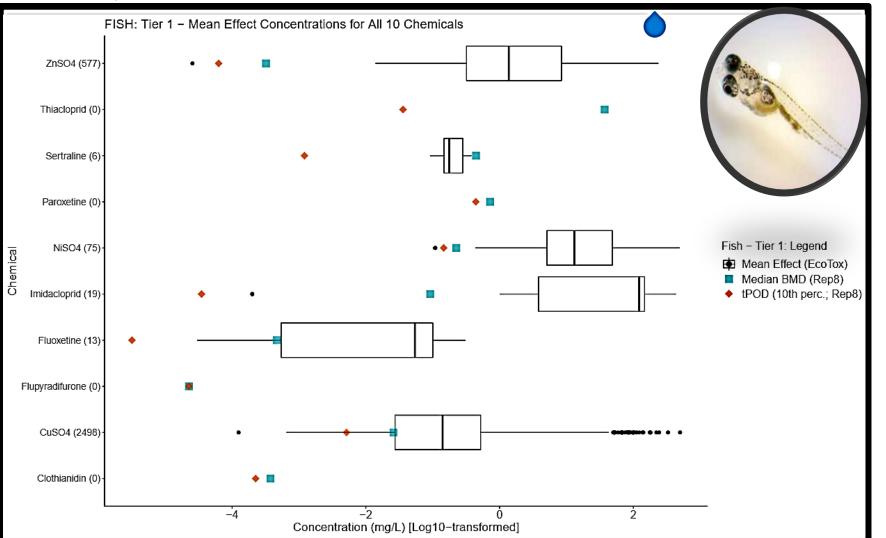
 Reliable point of departure
[tPOD] with defined uncertainty range

[tPOD] ≤ [Most sensitive chronic endpoint]
Effective provisional, protective value

[tPOD] <<< [Most sensitive chronic endpoint]
 Overly conservative

[tPOD] > [Most sensitive chronic endpoint]
 Not protective

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#### Comparison with In vivo, Adverse Effect Concentrations (Fish)

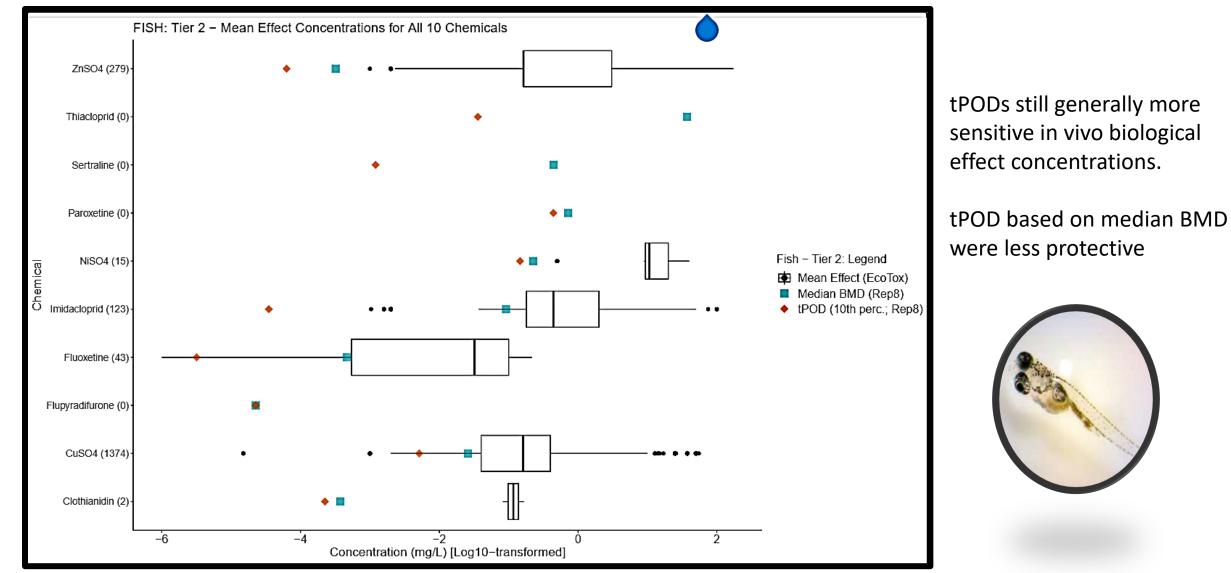
tPODs were generally more sensitive than apical adverse effect concentrations.

tPOD based on median BMD were less protective

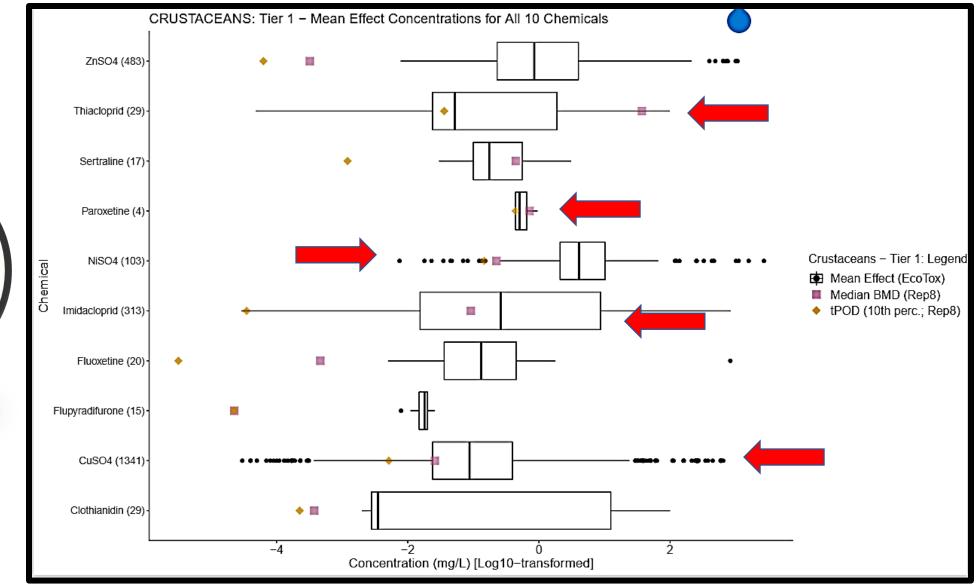
In some cases 2 orders of magnitude more protective

Still in the process of more detailed QA of the ECOTOX records with lower effect conc.

#### Comparison with In vivo, Biological Effect Concentrations (Fish)



#### Fish-based tPODs are not protective of all aquatic organisms





- Preliminary data suggest tPOD is promising as a lower bound estimate of toxicity to fish.
- Appears more conservative than ½ log different regulatory programs will need to weigh in on whether too conservative (need to test more chemicals).
- There does appear to be a need for taxa-specific tPOD determinations

# Molecular Biomarkers

Connectivity Mapping

US EPA, Center for Computational Toxicology and Exposure, Great Lakes Toxicology and Ecology Division

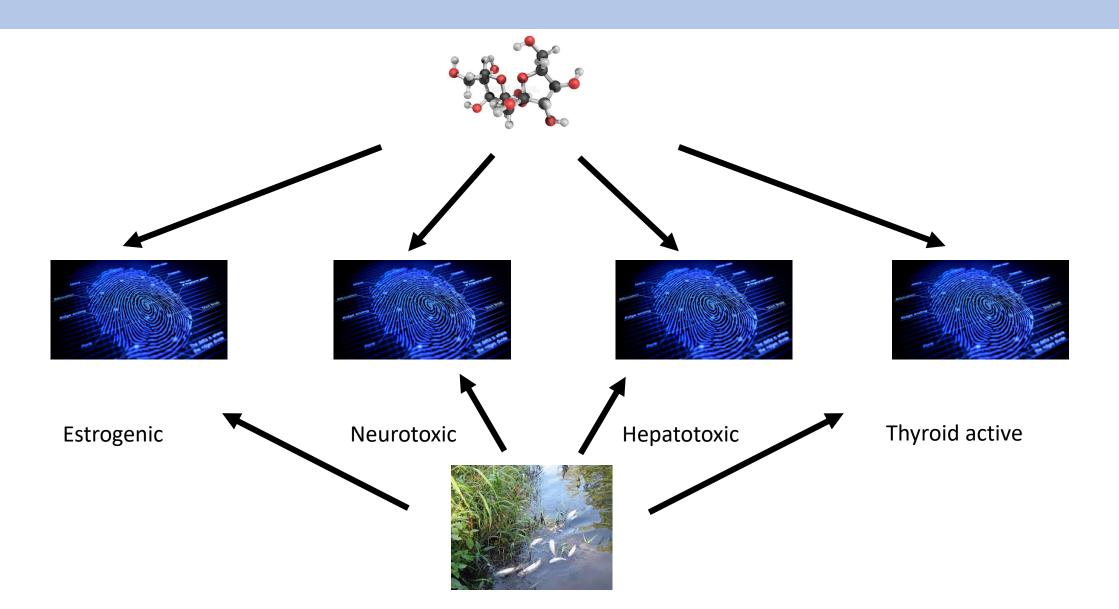


# **Current Limitations**

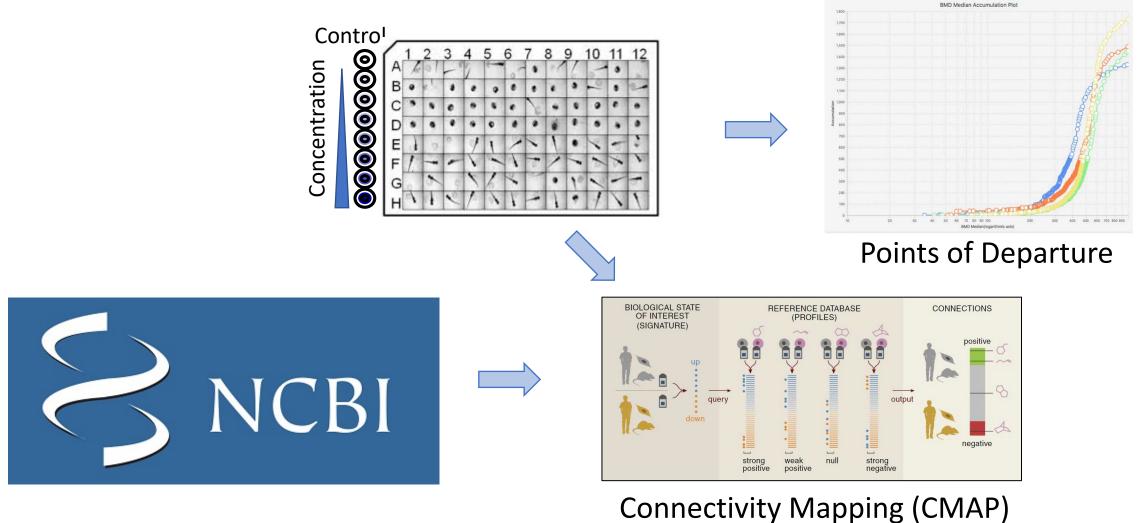
- Cumulative Risk
  - Chemical in water or tissue
  - Interactions Mixtures
  - Nonchemical stressors e.g. DO
- Lamp post
  - Look for what you can look for
- Apical endpoints
  - Uninformative
  - Read-across
  - Prediction of mixture effects



## **Biomarker Library**



## **Omics-based Biomarkers**

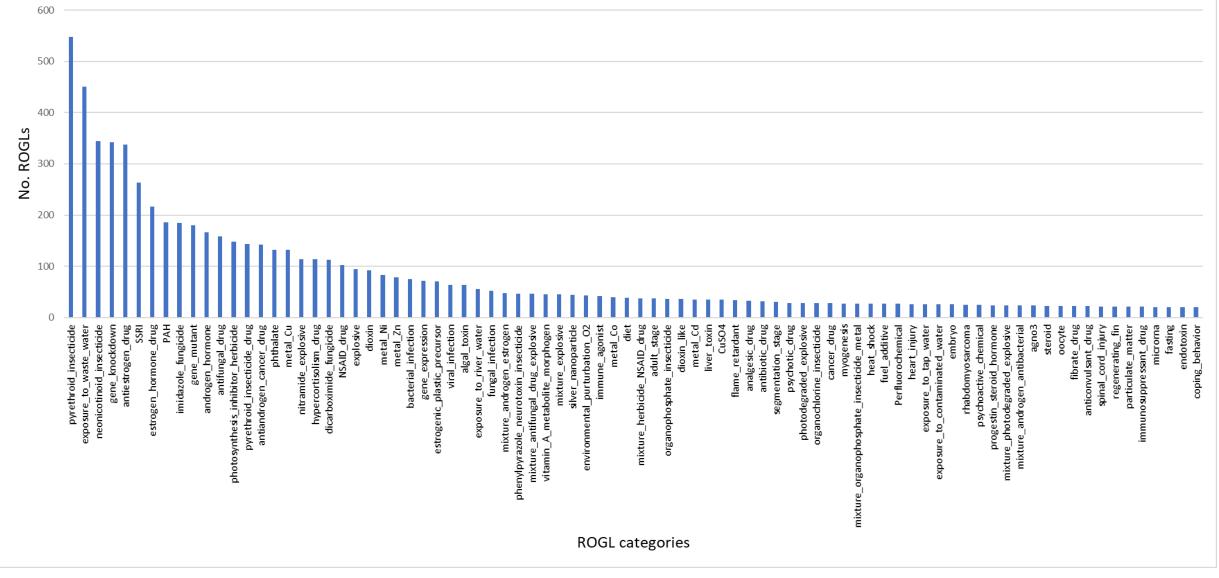


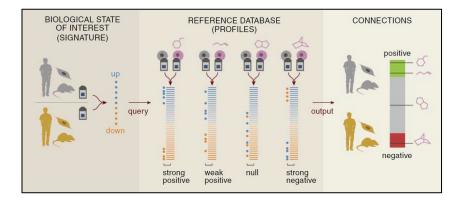
Justin Lamb et al. Science 2006;313:1929-1935

### Scaled up ROGL Library

Metrics	Current Effort 2021	
No. studies/datasets (GSE)	<b>450</b> = 70 (FHM) + 153 (ZF Affy) + 227 (ZF Agilent)	780 RNA-seq datasets
No. samples	<b>11639</b> = 4222 (FHM) + 2147 (ZF Affy) + 4440 (ZF Agilent) + 830 (RNAseq)	
No. microarray platforms	<b>42</b> = 9 (FHM Agilent) + 5 (ZF Affy) + 28 (ZF Agilent)	
Profiling technology	array & RNAseq	
NO. ROGLs	8021 = 7191 (array) + 830 (RNAseq); 4491 sets (combo of platform/chemical/dose/duration/ tissue/lifestage)	
NO. sets of query signatures	1188	
Signature cross-mapping	Ensembl ZF gene orthologs; EPA FHM genome	
Performance across platforms/species	Much better	21

### ROGL Categories by Chemical/Biology Applications/MOAs





### **CMAP: Effects-based linkages**

#### NCIT -National Cancer Institute Thesaurus

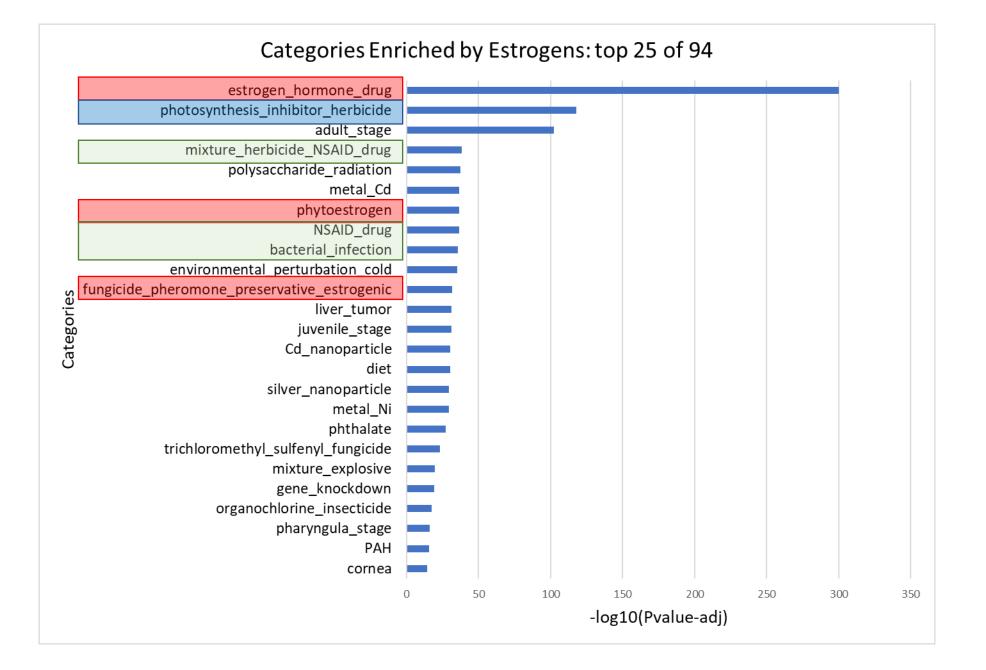
- owl:Thing
  - 🕨 🗝 Abnormal Cell
  - ► -- Activity
  - ► ←● Anatomic Structure, System, or Substance
  - ► ←● Biochemical Pathway
  - 🕨 🗝 Biological Process
  - Chemotherapy Regimen or Agent Combination
  - ► ←● Conceptual Entity
  - Diagnostic or Prognostic Factor
  - ► -- Disease, Disorder or Finding
  - 🛏 🗝 Drug, Food, Chemical or Biomedical Material
  - Experimental Organism Anatomical Concept
  - 🛏 🗝 Experimental Organism Diagnosis
  - ⊫ ←● Gene
  - ► ← e Gene Product
  - ► ←● Manufactured Object
  - 🕨 🗝 Molecular Abnormality
  - ► ←● Organism
  - ---- Property or Attribute
  - Retired Concept

#### CHEBI – Chemical Entities of Biological Interest

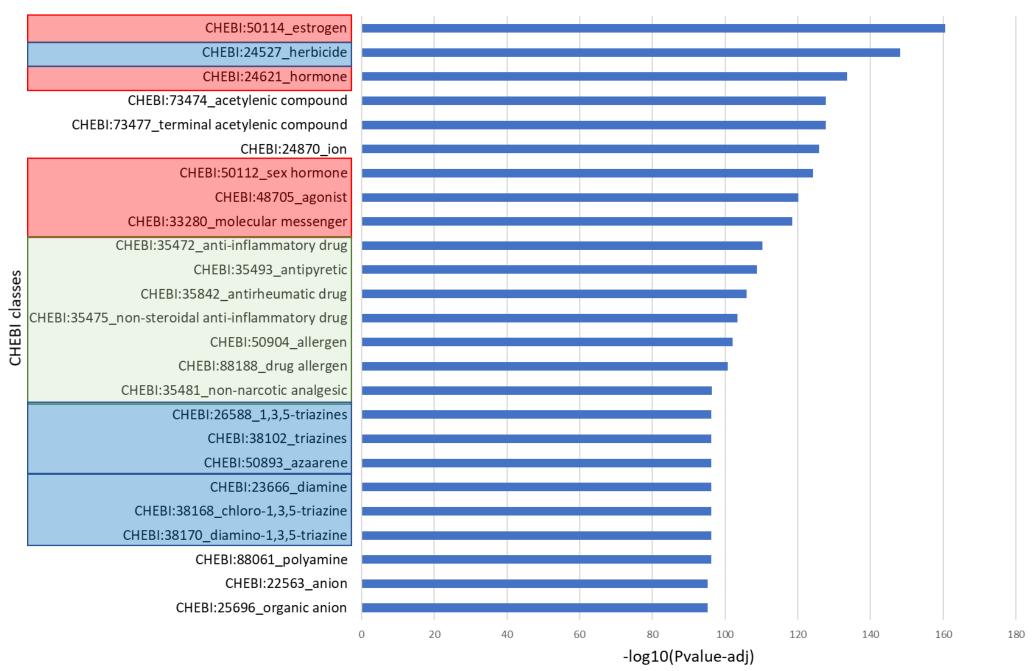
- Structure
- Role

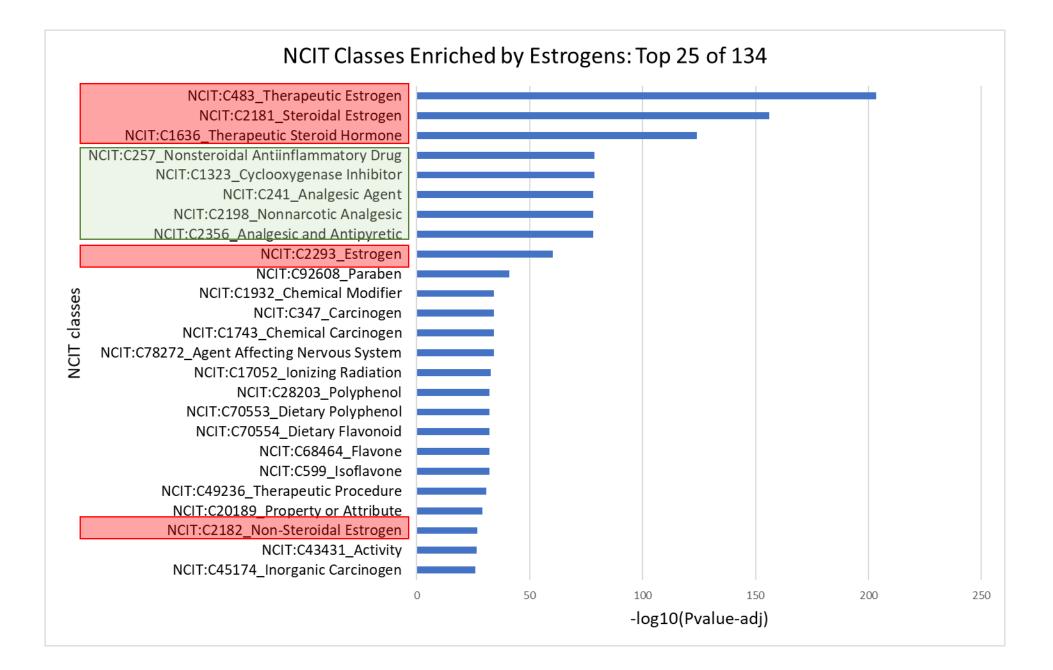
- Chemical context (ligand)
- Biological context (hormone)
- Application (pesticide)
- Subatomic Particle

- Ontologies
  - Controlled vocabulary
  - Maintained by experts in the field
  - Evaluated and edited
- Enrichment
  - Discovery
    - Structural moieties
    - Biological connections
      - Roles
  - Weight of Evidence
    - Do the effects-based linkages reflect enrichment



#### CHEBI Classes Enriched by Estrogens: Top 25 of 402







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- Logan Everett
- Leah Wehmas
- Russ Hockett
- Teresa Norberg-King
- Kathy Jensen
- Jenna Cavallin
- David Murphy
- Brett Blackwell

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