Omics Applications

Case Study

Chemical Risk Assessment

Exposure:

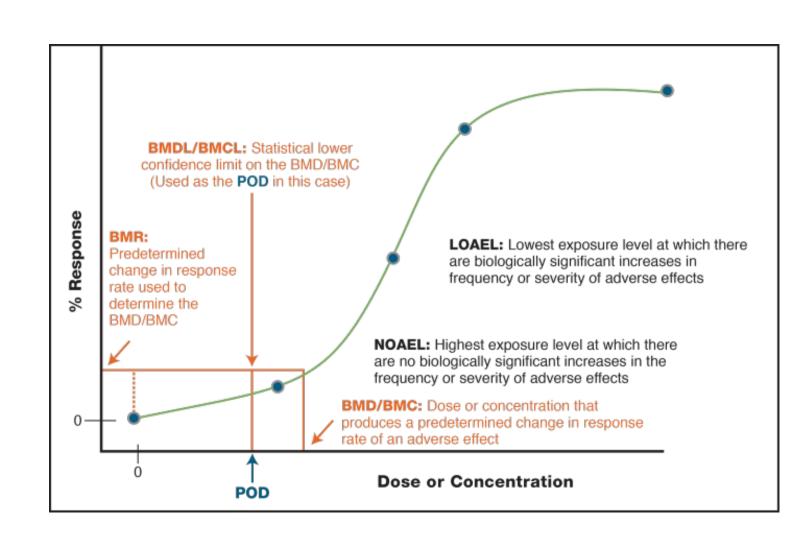
What concentrations occur in organisms or the environment?

Hazard/Effect:

What concentrations cause adverse effects to exposed organisms?

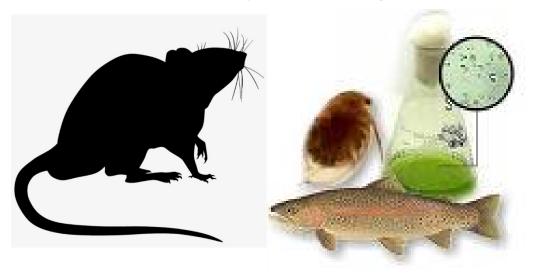
Safety:

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

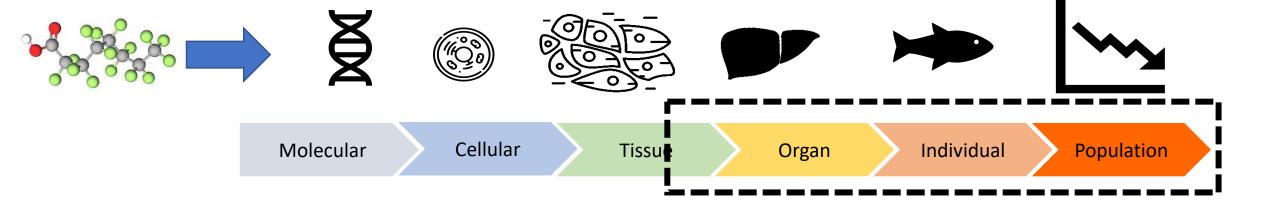
Structure-based Prediction



- 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)



High throughput assays

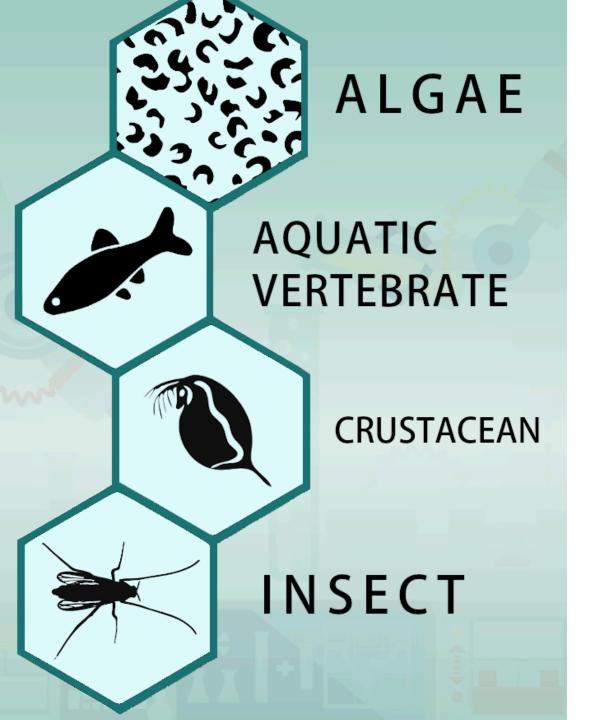
- Smaller scale
- More rapid response
- Simplified systems
- Pathway coverage via batteries, multiplexing, or high content
- Dose-response more costeffective

Direct observation of apical adverse effects

- Often slower, more latent response (especially when sub-lethal)
- Complex systems, integrate pathways
- Larger scales
- Dose response characterization is costly

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





Species	Guideline Test Method	Age at Start	Temp
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 rd 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 10⁴ 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

24 h exposure

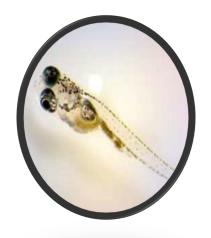
	Control
Replicates	A
	E

Phenotypic anchoring

- survival
- behavior
- Photo pigments

Office of Research and Development
Center for Computational Toxicology & Exposure

Initial 10 Chemicals



Metals

CuSO4

NiSO4

ZnSO4

Selective Serotonin

Reuptake Inhibitors (SSRI)

Fluoxetine

Paroxetine

Sertraline

Neonicotinoids

Clothianidin

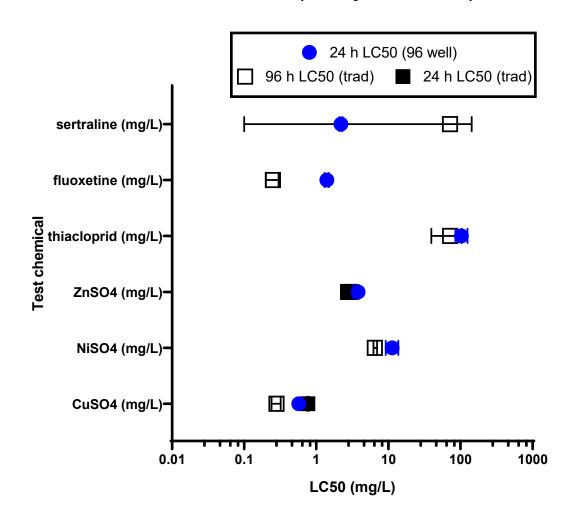
Imidacloprid

Thiacloprid

Flupyradifurone

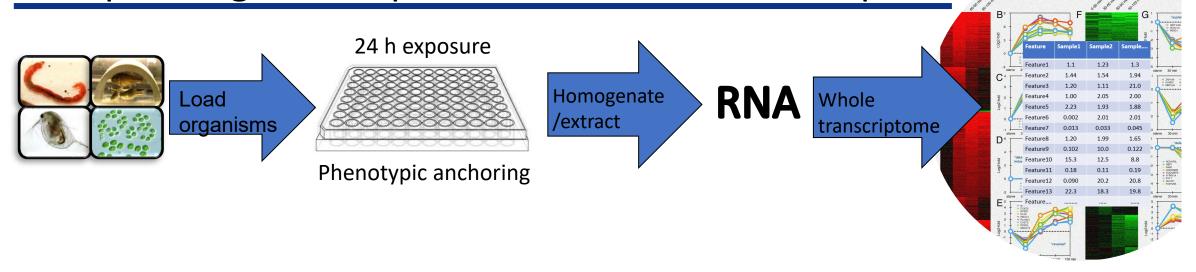
Assay design

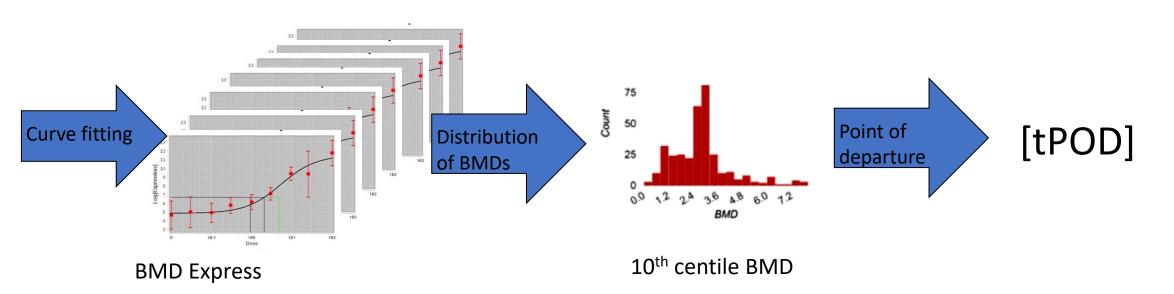
How does the 24 h, 96 well plate format compare with the traditional 24 or 96 h LC50 in a tank or beaker?



- 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







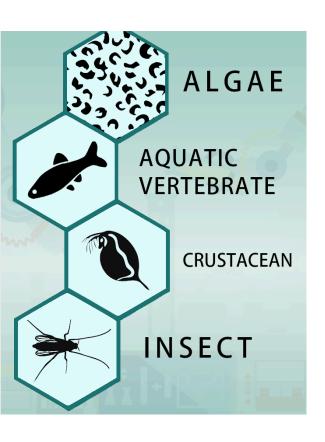
APCRA Case study: Transcriptomics-based PODs for Ecotoxicology

- 1. Generate transcriptomic PODs for ≈ 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

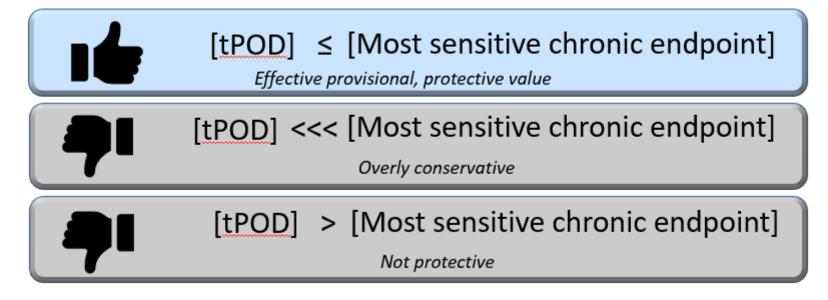


Assay Optimization

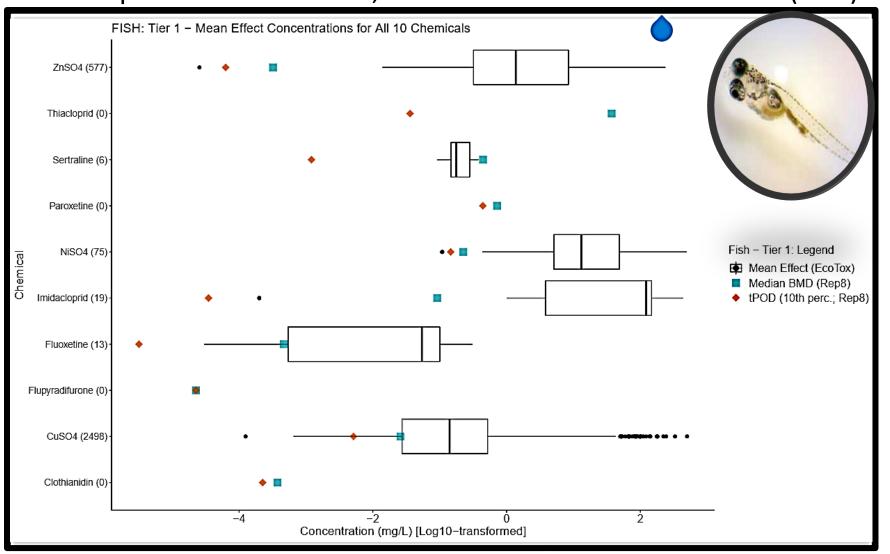
- How many replicate wells (animals)?
- How much genome coverage?
- Assay acceptance criteria?

Reliable point of departure [tPOD] with defined uncertainty range

Assay Evaluation



Comparison with In vivo, Adverse Effect Concentrations (Fish)



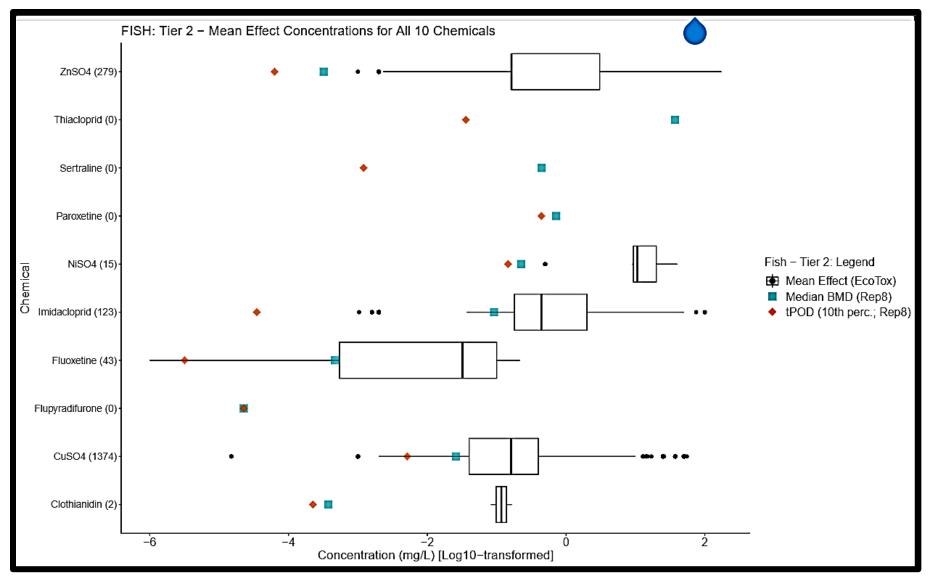
tPODs were generally more sensitive than apical adverse effect concentrations.

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)

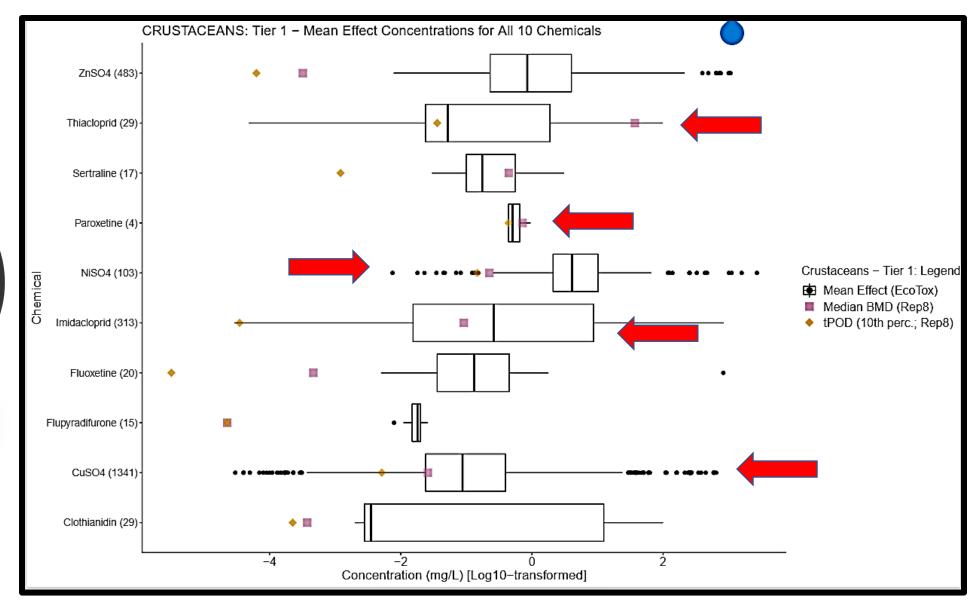


tPODs still generally more sensitive in vivo biological effect concentrations.

tPOD based on median BMD were less protective



Fish-based tPODs are not protective of all aquatic organisms



Preliminary Evaluation

 Preliminary data suggest tPOD is promising as a lower bound estimate of toxicity to fish.

• Appears more conservative than $\frac{1}{2} \log - \text{different regulatory}$ 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

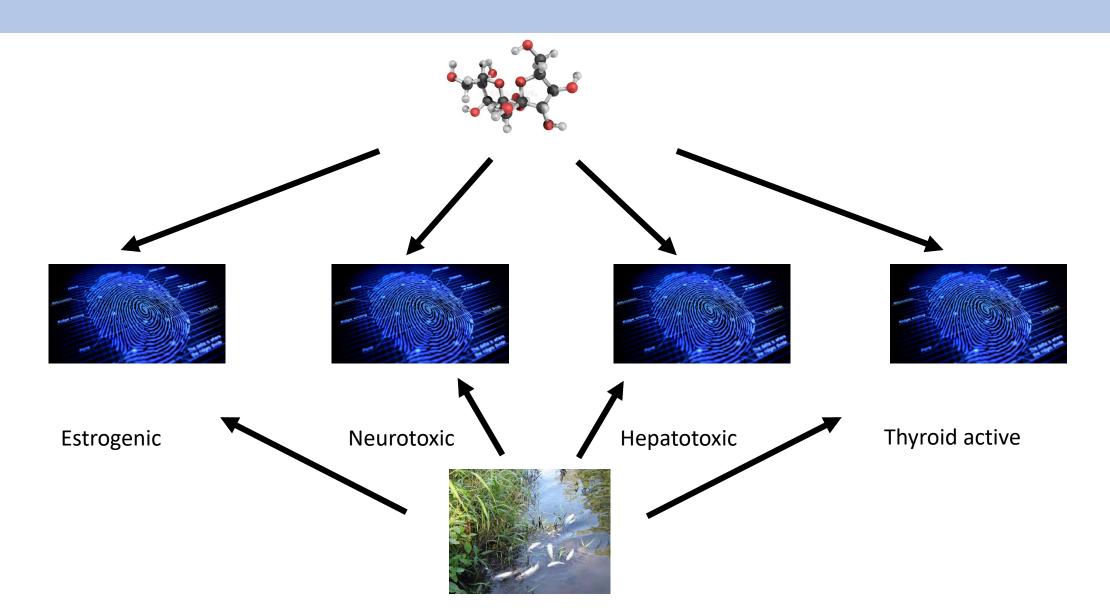


Current Limitations

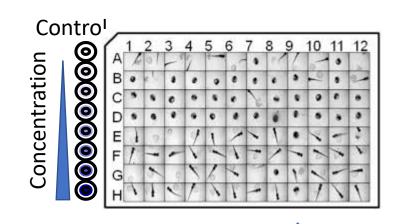
- Cumulative Risk
 - Chemical in water or tissue
 - Interactions Mixtures
 - Nonchemical stressors e.g. DO
- Lamp post
 - Look for what you can look for
- Apical endpoints
 - Uninformative
 - Read-across
 - Prediction of mixture effects

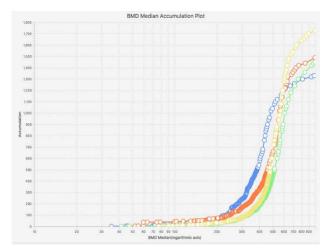


Biomarker Library



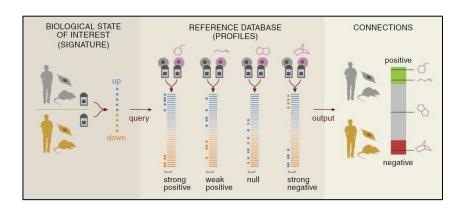
Omics-based Biomarkers





Points of Departure





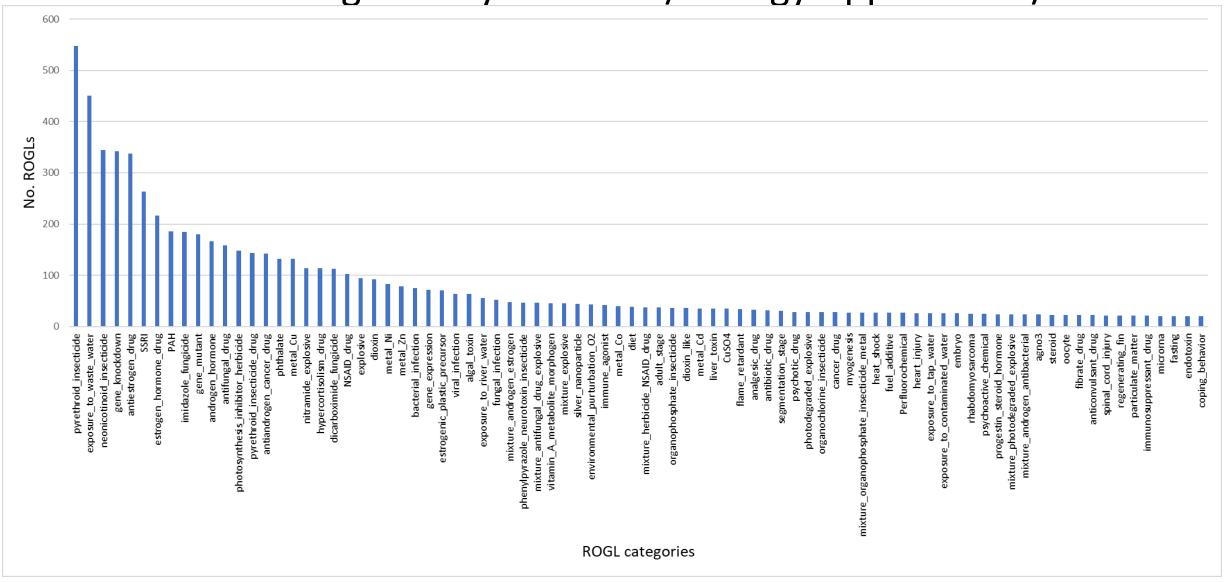
Connectivity Mapping (CMAP)

Scaled up ROGL Library

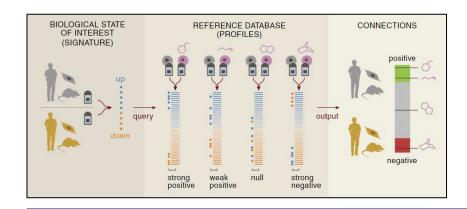
Metrics	Current Effort 2021
No. studies/datasets (GSE)	450 = 70 (FHM) + 153 (ZF Affy) + 227 (ZF Agilent)
No. samples	11639 = 4222 (FHM) + 2147 (ZF Affy) + 4440 (ZF Agilent) + 830 (RNAseq)
No. microarray platforms	42 = 9 (FHM Agilent) + 5 (ZF Affy) + 28 (ZF Agilent)
Profiling technology	array & RNAseq
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

780 RNA-seq datasets

ROGL Categories by Chemical/Biology Applications/MOAs



^{*}among the 259 categories, only those >= 20 are displayed



CMAP: Effects-based linkages

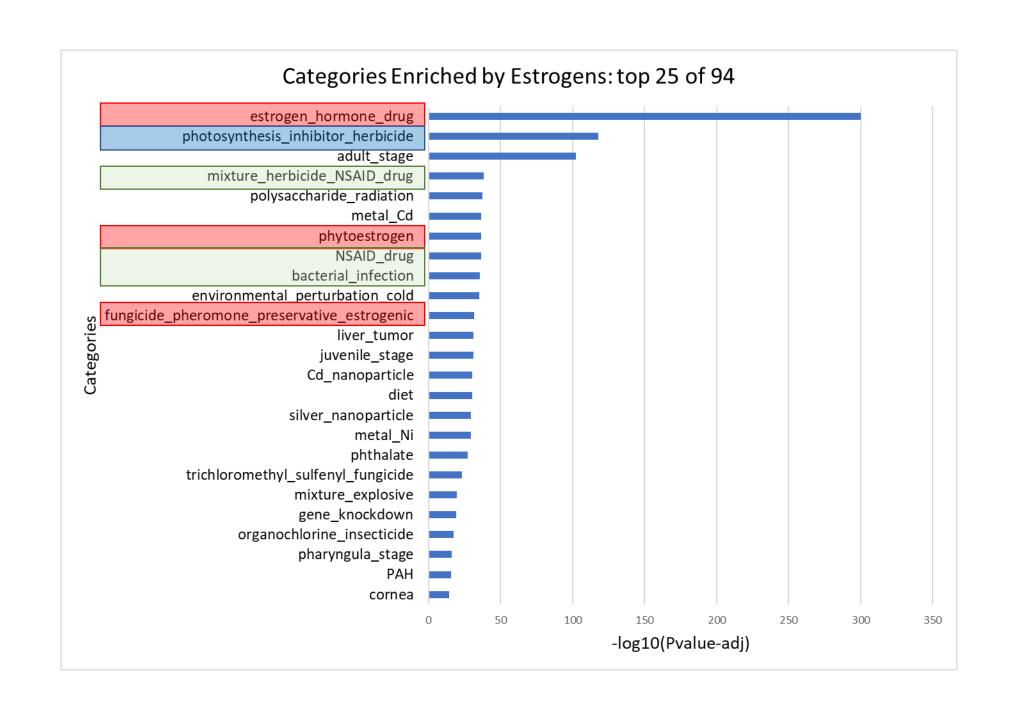
NCIT -National Cancer Institute Thesaurus

▼ 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 ← Gene Product Manufactured Object Molecular Abnormality ► Corganism ► - Property or Attribute ► - Retired Concept

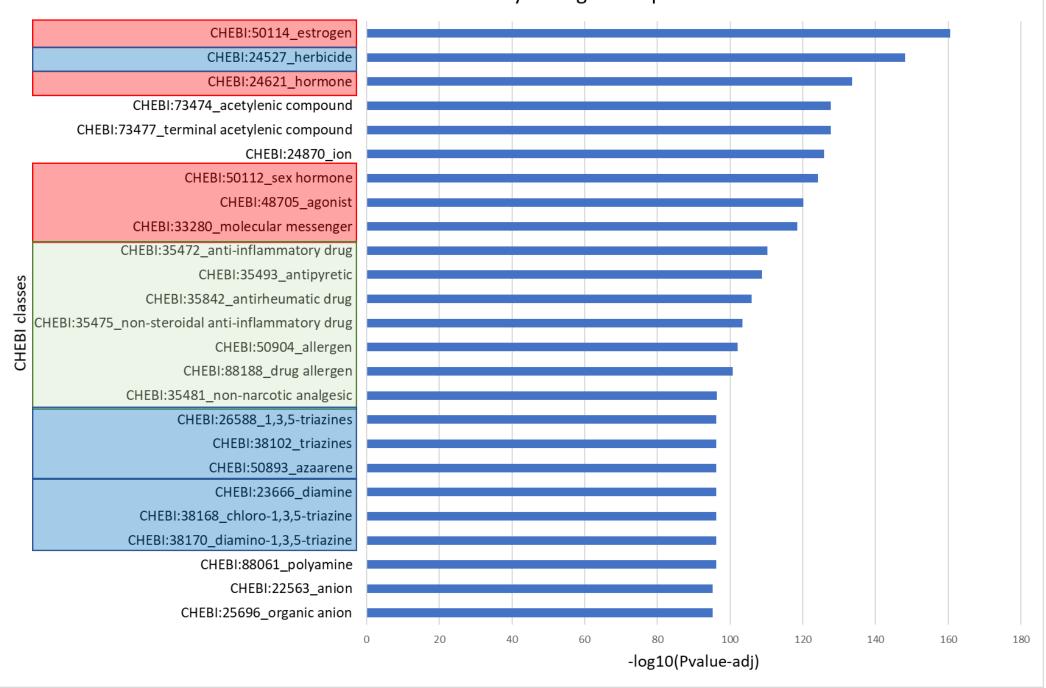
CHEBI – Chemical Entities of Biological Interest

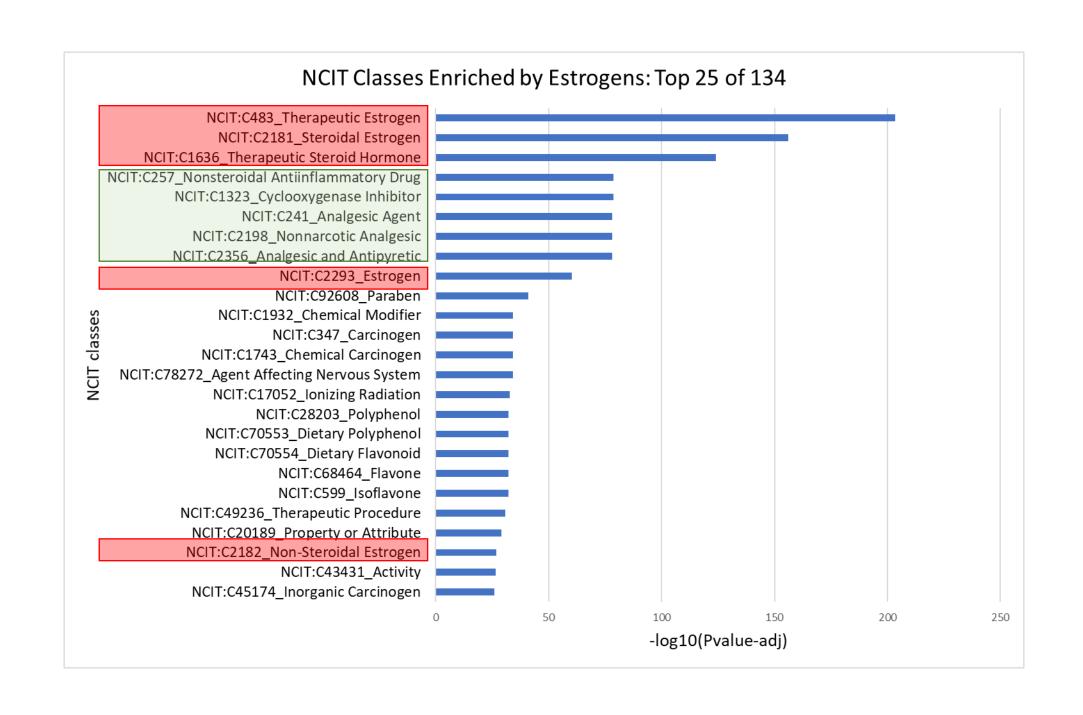
- Structure
- Role
 - Chemical context (ligand)
 - Biological context (hormone)
 - Application (pesticide)
- Subatomic Particle

- 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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CMAP HTTr-POD

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