



國家衛生研究院
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 ChemDIS-ZF

Inferring Potential Effects of PFASs via a Novel Chemical-phenotype Inference System ZFinfer

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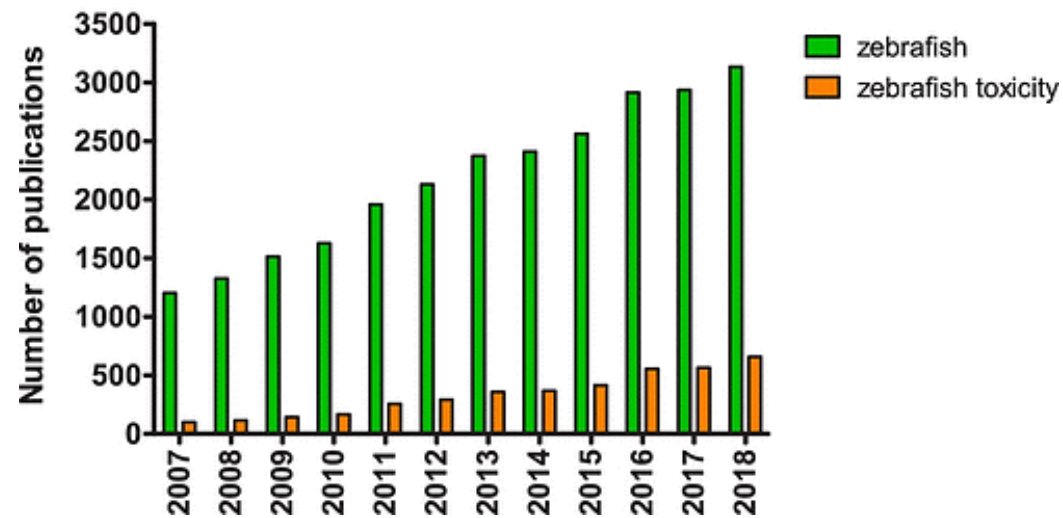
Zebrafish in Drug Discovery and Toxicology Studies

- Comparison to the human reference genome shows that approximately **70%** of human genes have at least one obvious zebrafish orthologue.
- Of the genes bearing morbidity descriptions, **82%** can be related to at least one zebrafish orthologue.

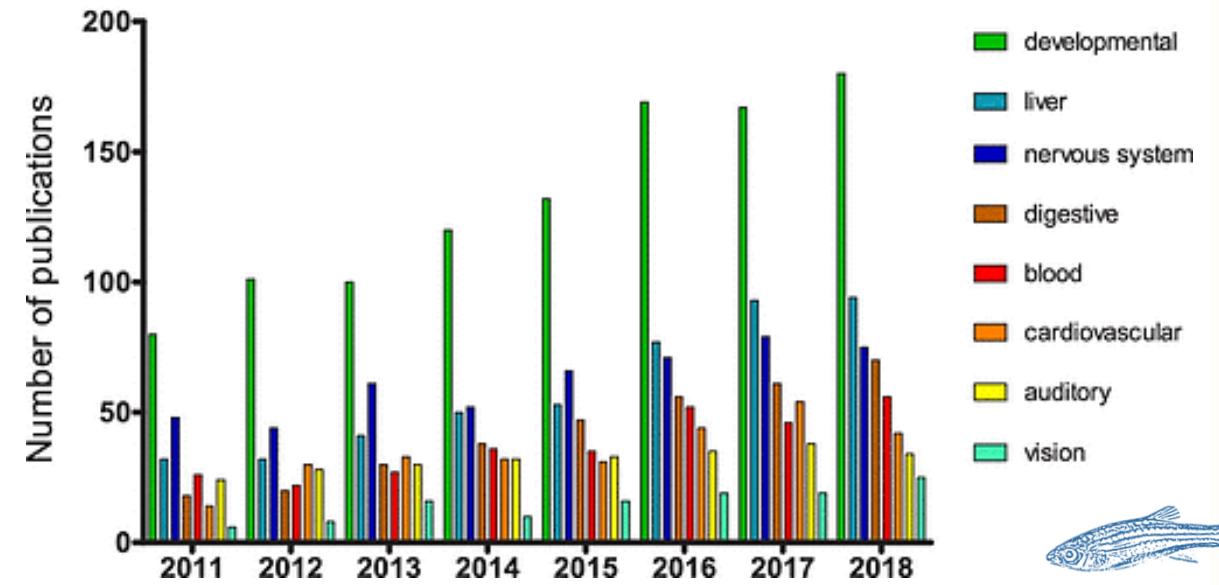
Howe, K. et al. *Nature*. 496, 7446 (2013)



Studies on zebrafish and toxicity studies with zebrafish



Toxicity studies using zebrafish




Cassar, S. et al. *Chemical Research in Toxicology*. 33, 95–118 (2020).

Zebrafish as an Alternative Model



Zebrafish is a useful model organism for toxicological research due to their **small size**, **fast reproduction**, and **genetic similarity** to humans.

However, as environmental pollutants increase, it becomes difficult to identify all hazards using zebrafish models alone. 

In silico models can assist in **identifying** chemical priorities for further experimental evaluation and provide insights into the underlying mechanisms.





Lin et al. *Journal of Cheminformatics* (2024) 16:91
<https://doi.org/10.1186/s13321-024-00891-4>

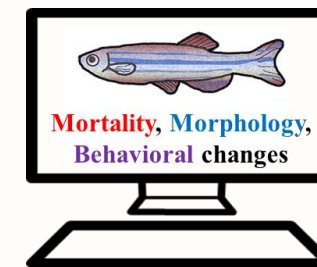
Journal of Cheminformatics

RESEARCH

Open Access

A novel multitask learning algorithm
 for tasks with distinct chemical space: zebrafish
 toxicity prediction as an example

Run-Hsin Lin^{1,2} , Pinpin Lin³ , Chia-Chi Wang⁴  and Chun-Wei Tung^{1,2*} 



| Zebrafish model

Bridging Data Gaps

The virtual zebrafish model speeds up **toxicity assessments** and **drug development**. It still has certain limitations, including the lack of:

- **Lack of Mechanistic Insights**

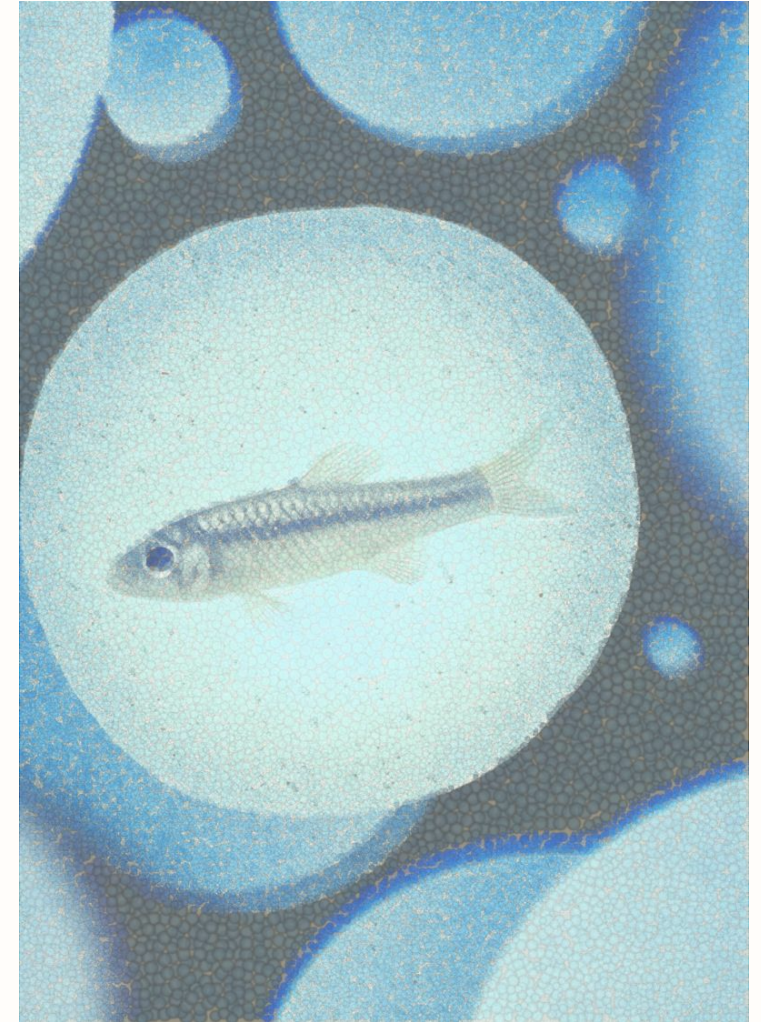
Mechanistic insights into how chemicals influence the overall effects on zebrafish.

- **Insufficient Linkage to Toxicity Endpoints**

A clear linkage between the overall effects caused by chemicals and specific toxicity endpoints.

- **Potential Disease Outcomes**

Diseases that may result from toxic responses caused by chemicals.



ChemDIS-ZF: Chemical–Disease Inference System for Zebrafish

• Extensive Database

 MongoDB.



Chemical-Protein Interaction data

STITCH

419,328 Chemicals
23,180 Zebrafish Proteins

Gene-Phenotype/Disease Interaction data



38,116 GO + Zebrafish anatomy
and development ontology (ZFA)
65,800 Zebrafish Phenotype Ontology (ZP)
5,227 Disease Ontology (DO)

Gene-Pathway Interaction data



12,511 KEGG Pathways
161,192 Reactome Pathways

Gene Ontology (GO): Gene functions, including cellular component, molecular function, and biological processes.

Zebrafish Anatomy Ontology (ZFA): Zebrafish Genes to Anatomical System.

Zebrafish Phenotype Ontology (ZP): Consists of structural or process terms affected by specific genes and quality terms from the Phenotype Quality Ontology.

• Enrichment Analysis System

Phenotype enrichment analysis



GO analysis



Pathway analysis



Disease inference



All the enrichment analyses were based on **hypergeometric tests** with multiple testing corrections using the **Benjamini–Hochberg method**.

System Validation (ex: Phenotype enrichment analysis - ZFinfer)



777

ToxCast Chemicals

Chemicals from the ToxCast used for validation.

Multidimensional in vivo hazard assessment using zebrafish

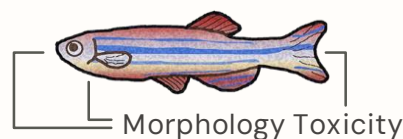


Lowest Effect Level in Zebrafish



Chemical-induced Toxicity References for validation.

17 Morphology Toxicity by Lowest Effect Level



17 Morphology Toxicity Endpoints related phenotypes

Endpoint	Description
YSE	Yolk Sac Edema
AXIS	Axis
EYE	Eye
SNOU	Snout
JAW	Jaw
OTIC	Otic
HEART	Heart
BRAIN	Brain
SOMI	Somite
PFIN	Pectoral Fin
CFIN	Caudal Fin
PIG	Pigmentation
CIRC	Circulation
TRUN	Trunk Length
SWIM	Swim Bladder
NC	Notochord Distortion
TR	Touch Response

189 Phenotypes (GO + ZFA)
3,542 Phenotypes (ZP)

Performance Measurement

- Sensitivity
- Specificity
- Accuracy
- Balanced Accuracy
- Positive Predictive Value
- Negative Predictive Value

System Performances

0.37

Average Sensitivity

Demonstrated sensitivity in 17 morphology endpoints.

0.72

Demonstrated average sensitivity in **Heart, Trunk length, Eyes, and Brain.**

- Sensitivity in critical morphological endpoints

- Sensitivity in different toxicity levels

0.49

Demonstrated average sensitivity in **Strong Toxicity** (<1 μM).

- System Validation by ECOTOX Knowledgebase Data



51

Priority Pollutants

Pollutant chemicals from the USEPA priority pollutant list used for validation.

ECOTOX Knowledgebase

Chemical Toxicity Records

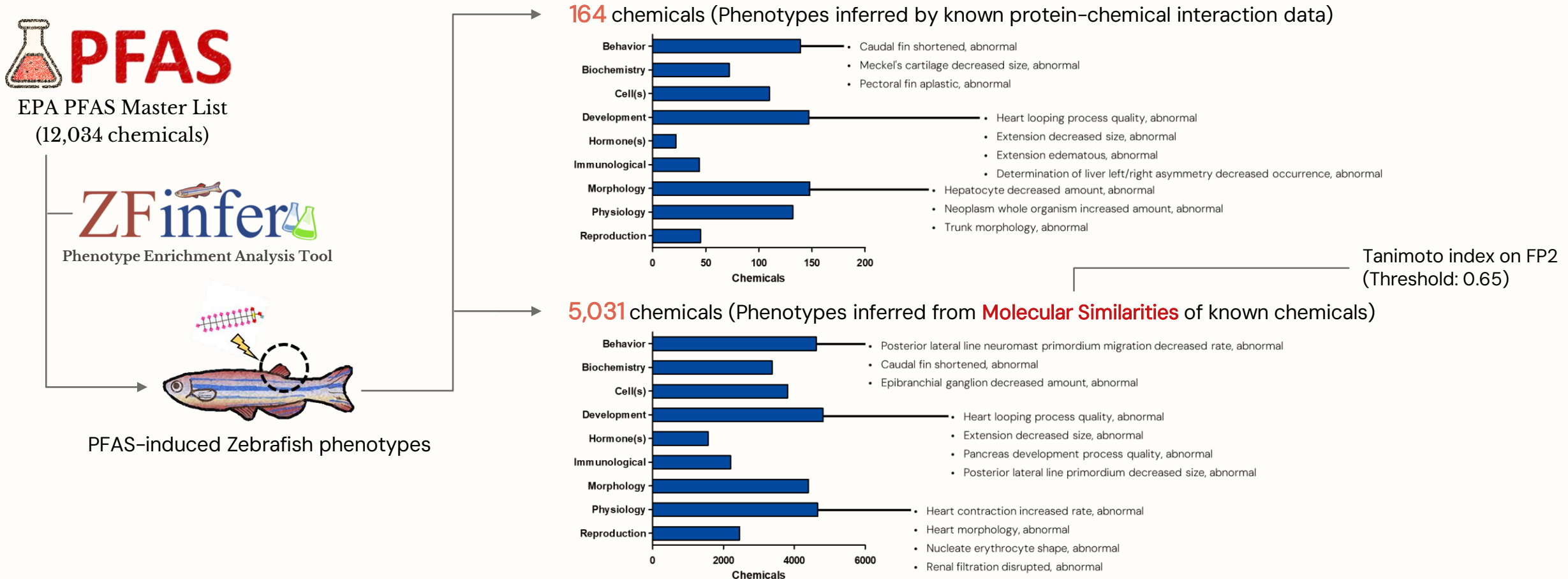
93%

Rediscovery Rate

Rediscovery rate in inferring relevant phenotypic effect groups from ECOTOX Knowledgebase.

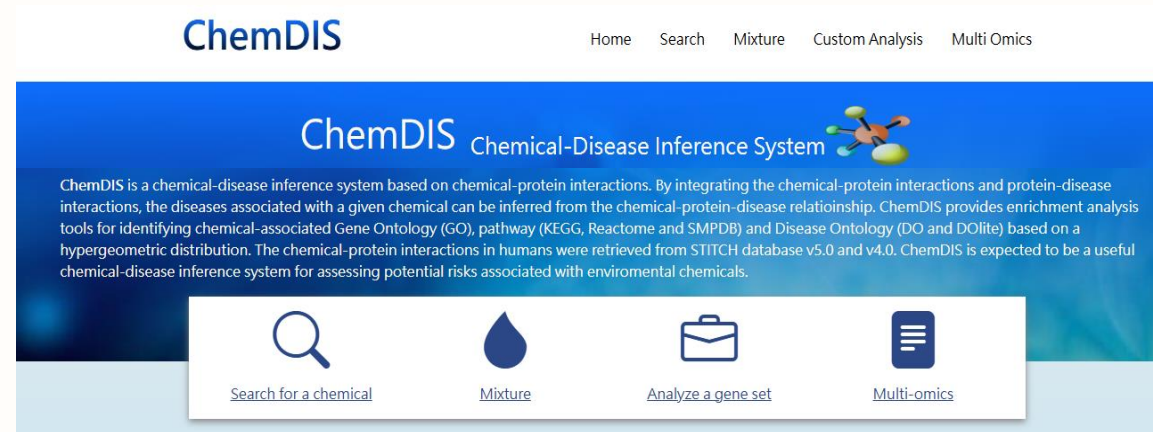
System Applications (ex: Phenotypes Inference of Environmental Pollutants)

Per- and poly-fluoroalkyl substances (PFAS) are widely used in various industries, and studies show that PFOS and PFOA can be toxic, but the effects of other PFAS are not well understood.



Web Tool

- ChemDIS Website



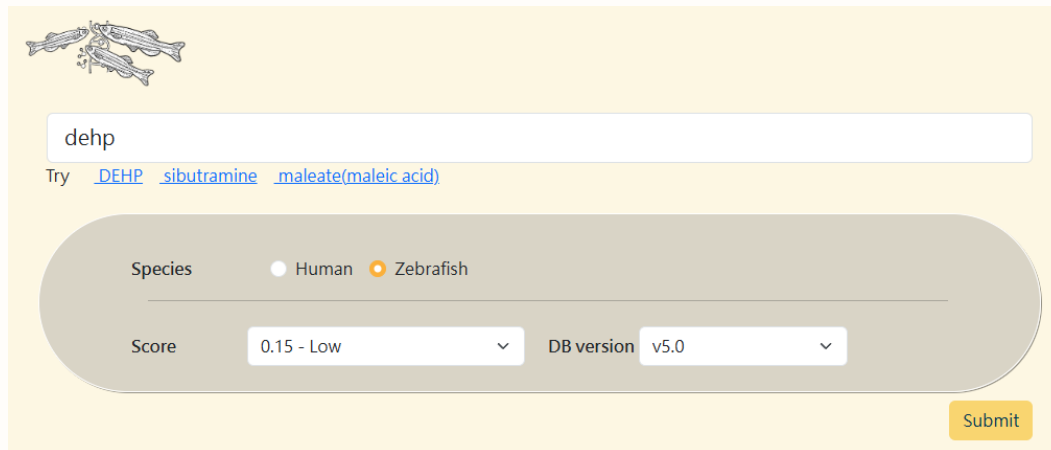
ChemDIS Home Search Mixture Custom Analysis Multi Omics


ChemDIS Chemical-Disease Inference System

ChemDIS is a chemical-disease inference system based on chemical-protein interactions. By integrating the chemical-protein interactions and protein-disease interactions, the diseases associated with a given chemical can be inferred from the chemical-protein-disease relationship. ChemDIS provides enrichment analysis tools for identifying chemical-associated Gene Ontology (GO), pathway (KEGG, Reactome and SMPDB) and Disease Ontology (DO and DOLite) based on a hypergeometric distribution. The chemical-protein interactions in humans were retrieved from STITCH database v5.0 and v4.0. ChemDIS is expected to be a useful chemical-disease inference system for assessing potential risks associated with environmental chemicals.

Search for a chemical Mixture Analyze a gene set Multi-omics

- Chemical Search






dehp

Try [_DEHP](#) [_sibutramine](#) [_maleate\(maleic acid\)](#)

Species Human Zebrafish

Score DB version

Phenotype 

Show: entries/page < < page 1/9 > >

Type	ID	Description	Gene Ratio	Bg Ratio	P	Adj. P	Genes
PhenotypeZP Phenotype	ZFA-0000303	female organism	10/67	68/4131	8.75e-8	1.09e-6	[+] cyp19a1a cyp17a1 nr3c1 esr2a fdx1b...
Phenotype	ZFA-0000242	male organism	10/67	67/4131	7.56e-8	1.09e-6	[+] esr2a esr2b esr1 fdx1b cyp11a1.2...
Phenotype	GO:0007530	sex determination	4/67	9/4131	7.50e-6	0.00006	esr2a fshr esr1 esr2b
Phenotype	ZFA-0000403	ovary	5/67	25/4131	0.00004	0.00025	cyp19a1a fshr esr1 cyp17a1 lhcg
Phenotype	ZFA-0009016	germ line cell	3/67	7/4131	0.00014	0.00068	cyp19a1a nr0b1 fshr
Phenotype	ZFA-0000598	testis	5/67	34/4131	0.00019	0.00078	esr2b fdx1b cyp11a1.2 fshr cyp17a1
Phenotype	ZFA-0001264	ovarian follicle stage IV	3/67	10/4131	0.00045	0.00161	fshr esr1 lhcg
Phenotype	ZFA-0005272	immature gonad	2/67	5/4131	0.00251	0.00785	cyp19a1a nr0b1
Phenotype	ZFA-0001345	interrenal gland	2/67	7/4131	0.00516	0.01290	cyp21a2 fdx1b
Phenotype	GO:0030238	male sex determination	3/67	22/4131	0.00504	0.01290	cyp19a1a cyp17a1 nr0b1

New Search
Basic
Protein
GO
Pathway
DO
Phenotype

New Approaches: Mixtures



The **mixture** of chemicals raises concerns about **health** and **environmental effects**, as neglecting these interactions could result in underestimating risks.

Drakvik, E. et al. *Environment international*. 134 (2020)

Current **regulations** on unintentionally produced mixtures and emerging contaminants are **unclear** due to the random and unknown nature of these mixtures. To effectively address these challenges, we need to:

1. Improve our **understanding** of potential exposures to chemical mixtures in the environment.
2. Improve the **analysis** of toxicological **mechanisms** and **interaction** patterns of single and mixtures.
3. Identify specific mixtures that are required to be addressed with **priority**.

European Commission and Joint Research Centre (2018)

System Updates (In progress)



Toxicogenomic-based Mixture Effect Analysis System for Zebrafish

- Mixture chemicals Search

Search results: rifampin, efavirenz

Species: Human Zebrafish

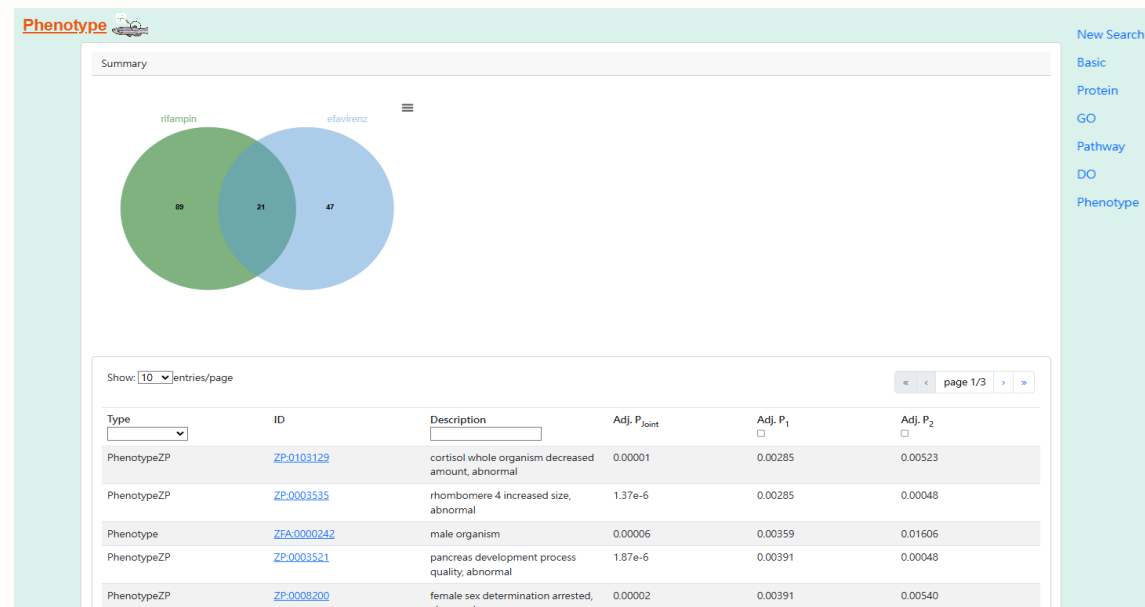
Score: 0.15 - Low

DB version: v5.0

Example1 Example2 Example3 Submit

- Mixture Effect Analysis

Mixture effect enrichment analysis will be evaluated by using the overall and overlapped genes that interact with chemicals in mixtures.



Thanks for your attention

BioCompLab.

<https://cwtung.nhri.edu.tw/>

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