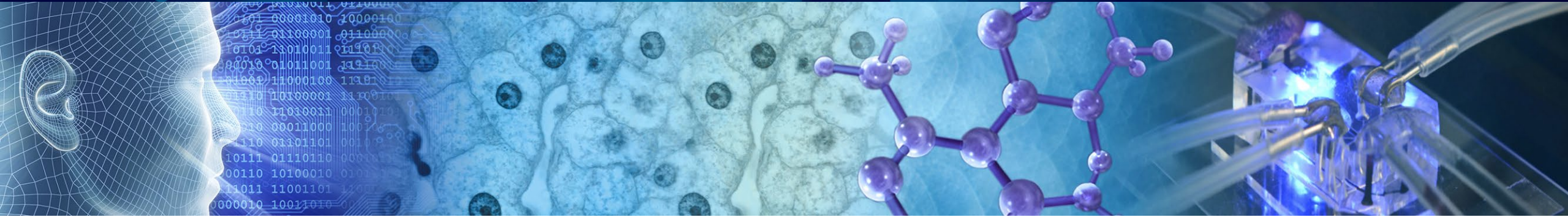




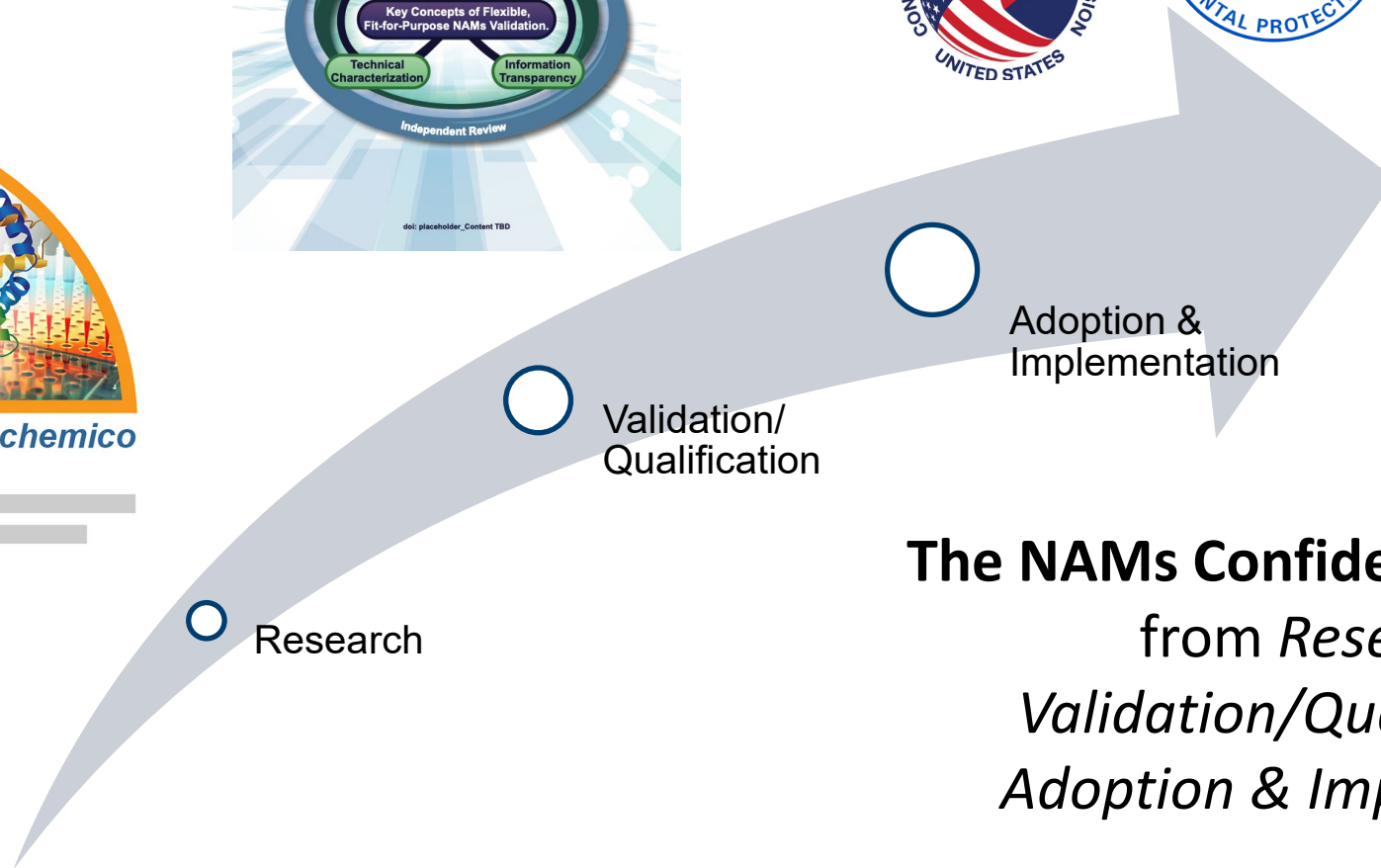
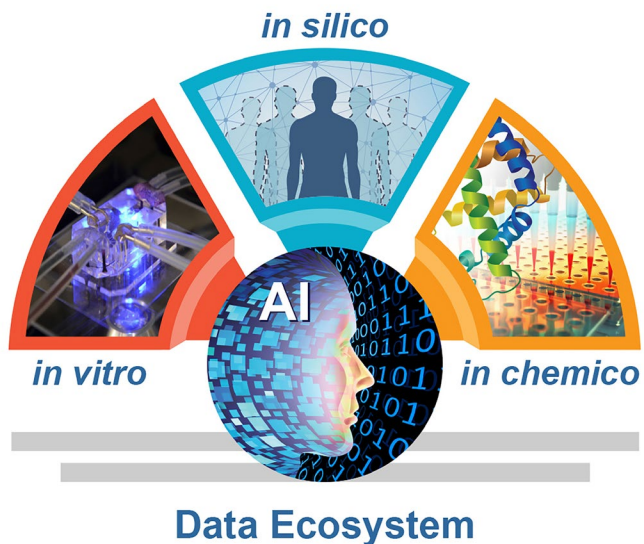
National Institute of  
Environmental Health Sciences  
*Division of Translational Toxicology*



# NICEATM Update: ICCVAM Public Forum 2025

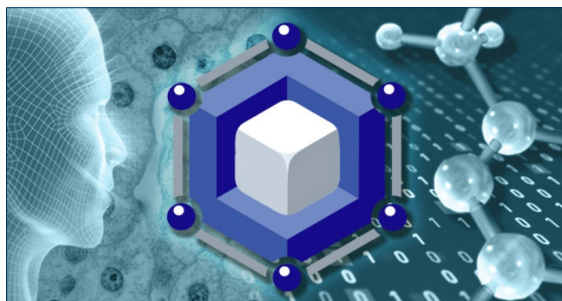
**Helena Hoegberg-Durdock**

**Acting Director, NTP Interagency Center for the Evaluation of  
Alternative Toxicological Methods**



**The NAMs Confidence Continuum:**  
from *Research* to  
*Validation/Qualification* to  
*Adoption & Implementation*





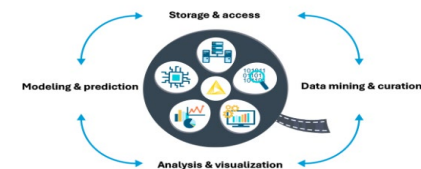
## Integrated Chemical Environment (ICE)

ICE is an open-access, user-friendly platform developed by NTP Interagency Center for the Evaluation of Alternative Toxicological Methods (NICEATM) that provides curated toxicologically relevant data and interactive computational tools to support development and evaluation of new testing approaches.



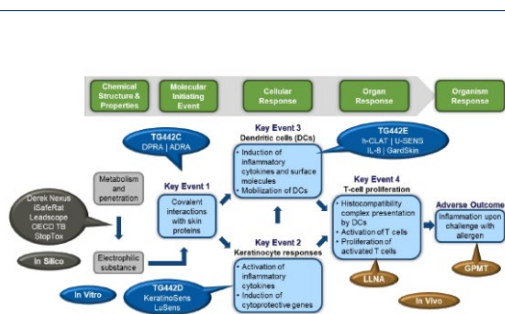
## Open (Quantitative) Structure-activity/property Relationship App (OPERA)

OPERA is a free and open-source/open-data suite of QSAR models providing predictions for physicochemical properties, environmental fate parameters, and toxicity endpoints to support non-animal approaches for predicting toxicity.



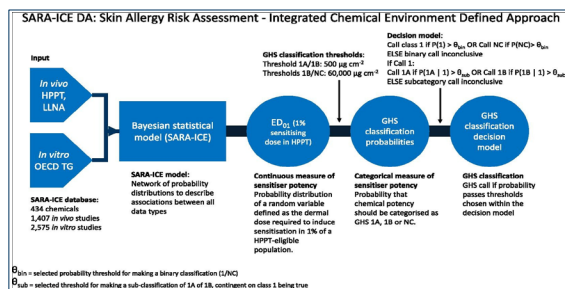
## Modeling and Visualization (MoVIZ) Pipeline

MoVIZ is a cheminformatics pipeline developed using the free and open-source KNIME analytics platform and aims to democratize computational methods through intuitive, well-documented, and user-friendly graphical interfaces. Among MoVIZ tools is a workflow facilitating chemical grouping based on supervised and unsupervised machine learning approaches.



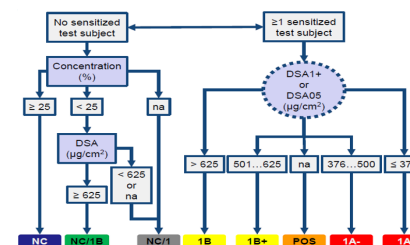
## The DASS App

The DASS App is an open-source web application to predict skin sensitization using defined approaches. It enables users to apply validated non-animal approaches to their own data.



## The Skin Sensitization Risk Assessment – Integrated Chemical Environment (SARA-ICE)

SARA-ICE is an open access web tool for quantitative prediction of a chemical's potential to cause skin sensitization in humans. It uses a Bayesian statistical model developed by NICEATM and Unilever to estimate human-relevant metric of skin sensitizer potency.



## The HPPT App (Coming Live Soon)

The HPPT App is an open-source web application that helps classify human predictive patch test (HPPT) data for skin sensitization potency using a WoE approach. It enables users to apply a GHS based approach for assigning skin sensitization potency subcategorizations to their own data.



ICE Data

- Curated in vivo, in vitro, and in silico toxicity data
- Measured and predicted chemical properties
- Predicted exposure
- Reported and predicted chemical use categories



Chemical  
Quick Lists

- Reference and Non-reference chemical lists
- Support the development and evaluation of new test methods

**Data and quick lists are accessible across ICE tools.**



Search



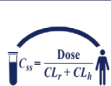
Chemical  
Quest



Curve  
Surfer



PBPK



IVIVE



Chemical  
Character-  
ization

- Explore ICE data through interactive visualizations
- Identify structurally similar chemicals
- Leverage computational models without coding
- Access data through APIs and bulk data downloads

**Tool Inter-connectivity: Send chemical and assay selections between tools.**

## News & Events

### ICE v4.2 Release

ICE updates include the following new resources and site improvements:

Updated cHTS  
PBPK and IVIVE tools allow user-provided physiochemical and ADME data  
Updated PCA plots in Chemical Characterization tool  
Updated Curve Surfer cards  
Updated Tox21 Chemical Quick List

ICE NEWS



## ICE Includes Property Predictions from OPERA

Learn more about QSAR property predictions generated by OPERA.

PAUSE



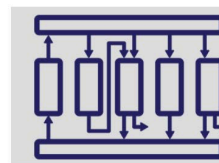
Search >



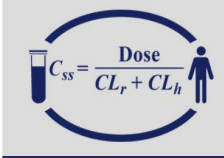
Chemical Quest >



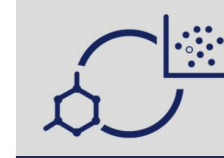
Curve Surfer >



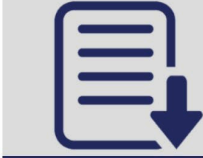
PBPK >



IVIVE >



Chemical Characterization >



Data >



Help Videos >



<https://ice.ntp.niehs.nih.gov/>

## ICE v4.1 August 2024

- Updated cHTS annotations from NCI Metathesaurus to OBO Foundry
- New PFAS chemical quick list and updated ROC chemical quick list
- ICE REST API updated to include Curve Surfer tool raw data
- Additional data visualizations in Search tool

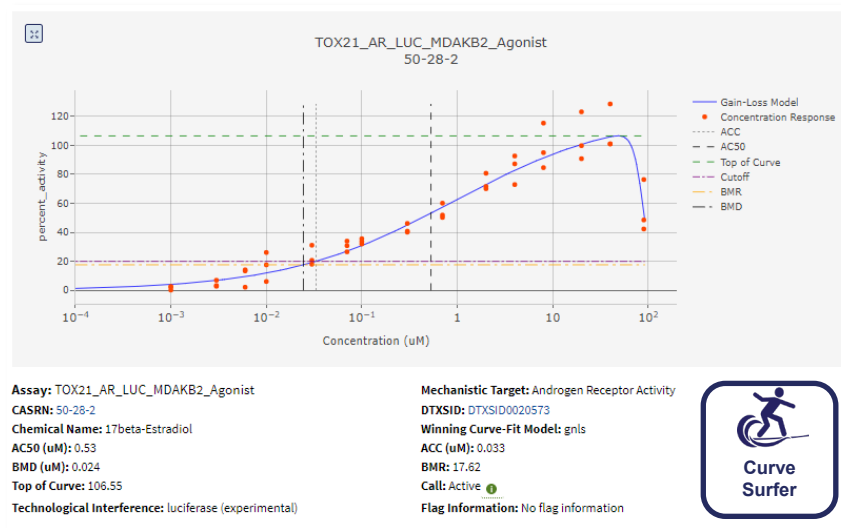
## ICE v4.1.1 February 2025

- New download file with ClassyFire (Djoumbou Feunang et al. 2016) chemical taxonomies for over one million chemicals
- Updated all ICE Tool Help videos

## ICE v4.2 July 2025

- cHTS updated to invitrodb v4.2
- Option to upload user's physicochemical and ADME data to run PBPK/IVIVE
- Updated PCAs in Chemical Characterization
- Updated Tox21 Chemical Quick List

### Curve Surfer cards show new points of departure, curation flags, and curve-fits



### Upload custom physicochemical parameters to run PBPK and IVIVE

**Upload Phys Chem Data**

Upload Drop file here

**Uploaded Files**

File Name

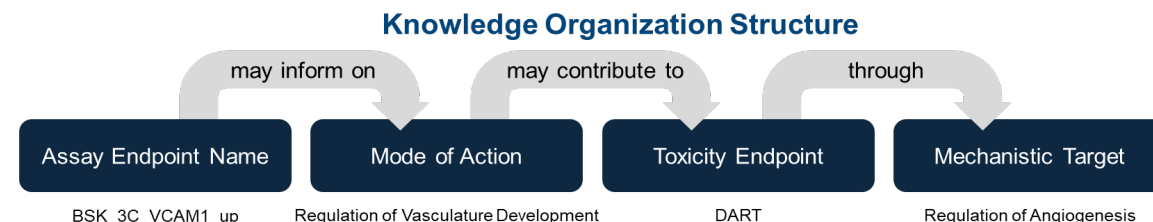
**Upload Custom PhysChem Data**

To load user assay data into your model, click "Upload." Supported file types: comma-delimited (csv), plain text (txt), and Excel (xlsx). To view example templates of data formats, click the links below.

Template PhysChem File for PBPK (Text)  
Template PhysChem File for PBPK (Excel)

Close

### Curated High Throughput Screening (cHTS) data are annotated to OBO Foundry ontologies to provide biological relevance





# OPERA as a Standalone Desktop Application

## OPERA standalone application:

- Free, opensource & open-data
- Single chemical and batch mode
- Multiple platforms (Windows and Linux)
- Embeddable libraries (java, C, C++, Python)
- **Command line & Graphical user interface**

## OPERA models:

- Physicochemical properties
- Environmental fate
- ADME properties
- Toxicity endpoints

## Input options:

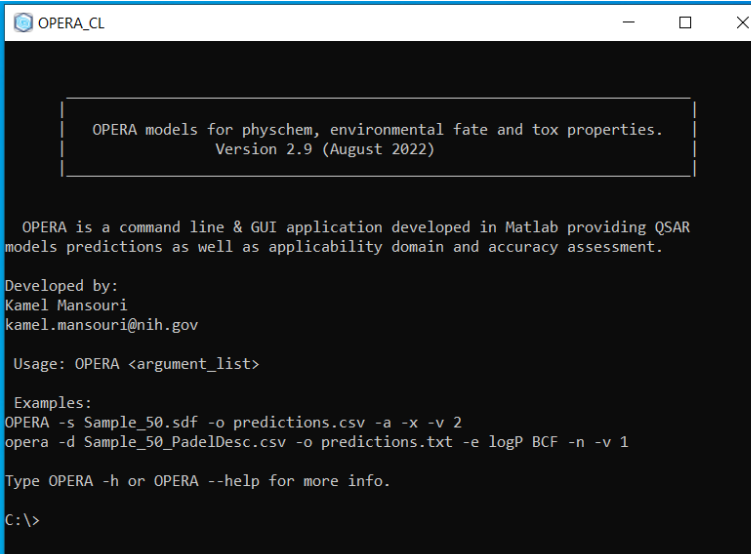
- Structure IDs (CAS, DTXSID, InChIKey)
- Structure files (SMILES, SDF, Mol)

## Download and learn more:

<https://github.com/NIEHS/OPERA>

<https://ntp.niehs.nih.gov/go/opera>

**Over 10,000  
downloads**



```
OPERA_CL

OPERA models for physchem, environmental fate and tox properties.
Version 2.9 (August 2022)

OPERA is a command line & GUI application developed in Matlab providing QSAR
models predictions as well as applicability domain and accuracy assessment.

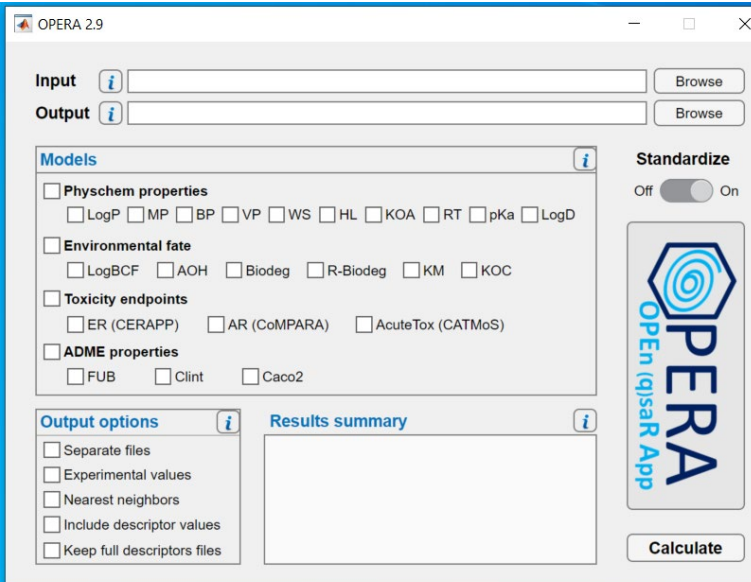
Developed by:
Kamel Mansouri
kamel.mansouri@nih.gov

Usage: OPERA <argument_list>


Examples:
OPERA -s Sample_50.sdf -o predictions.csv -a -x -v 2
opera -d Sample_50_PadelDesc.csv -o predictions.txt -e logP BCF -n -v 1


Type OPERA -h or OPERA --help for more info.


C:\>
```




OPERA 2.9

Input   Browse


Output   Browse

**Models** 


- ☐ **Physchem properties**  
☐ LogP ☐ MP ☐ BP ☐ VP ☐ WS ☐ HL ☐ KOA ☐ RT ☐ pKa ☐ LogD
- ☐ **Environmental fate**  
☐ LogBCF ☐ AOH ☐ Biodeg ☐ R-Biodeg ☐ KM ☐ KOC
- ☐ **Toxicity endpoints**  
☐ ER (CERAPP) ☐ AR (CoMPARA) ☐ AcuteTox (CATMoS)
- ☐ **ADME properties**  
☐ FUB ☐ Clint ☐ Caco2

**Output options** 

- ☐ Separate files
- ☐ Experimental values
- ☐ Nearest neighbors
- ☐ Include descriptor values
- ☐ Keep full descriptors files

**Results summary** 

☐ **Standardize**  
Off ☐ On

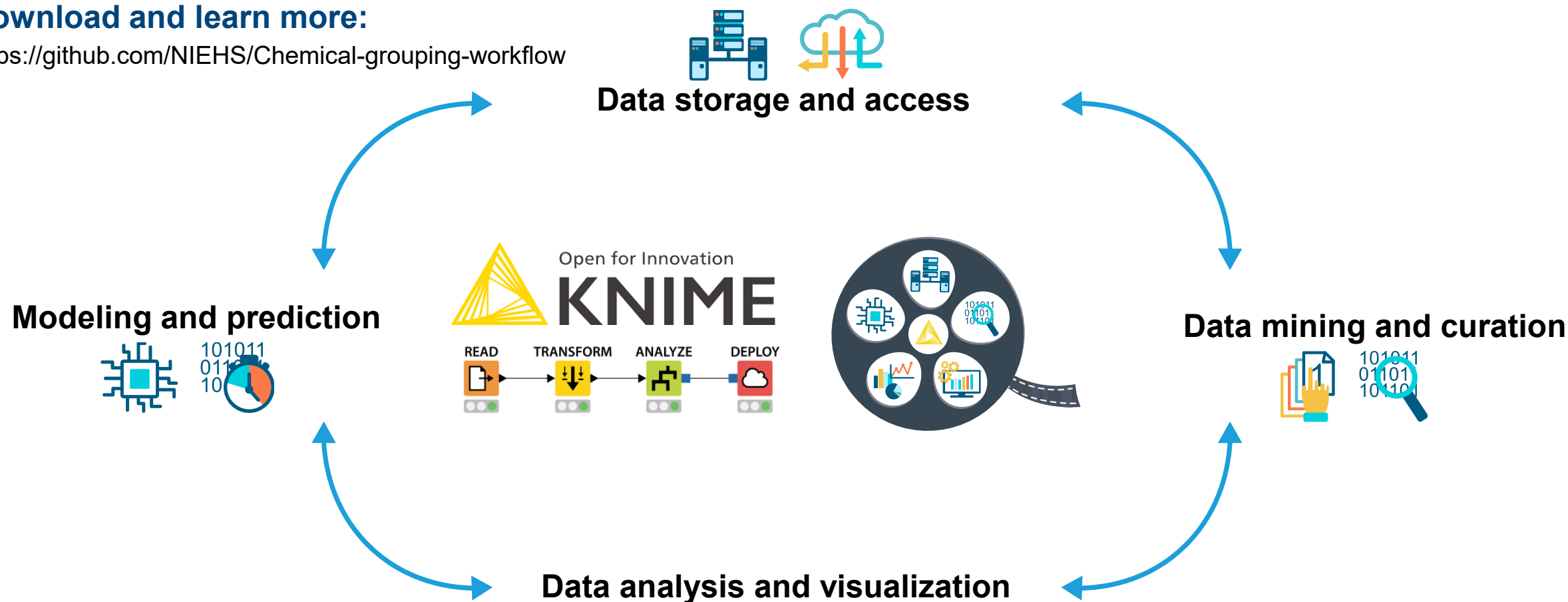
 **OPERA**  
Open (q)SAR App

Calculate

# Modeling and Visualization (MoVIZ) Pipeline

**Download and learn more:**

<https://github.com/NIEHS/Chemical-grouping-workflow>



Moreira-Filho et al.  
*Journal of Cheminformatics* (2024) 16:101  
<https://doi.org/10.1186/s13321-024-00894-1>

**SOFTWARE**

**Open Access**

Democratizing cheminformatics:  
interpretable chemical grouping using  
an automated KNIME workflow



*Journal of Cheminformatics*



Mansouri et al. *Journal of Cheminformatics* (2024) 16:19  
<https://doi.org/10.1186/s13321-024-00814-3>

**SOFTWARE**

**Open Access**

Free and open-source QSAR-ready workflow  
for automated standardization of chemical  
structures in support of QSAR modeling



*Journal of Cheminformatics*

Corpora  
Articles,  
Reports, etc.

Multi-layer Document Image/Text Object

- embed metadata for grounding
- markdown conversion: GROBID

LiteLLM Structured Data Extraction

- 100s of LLMs
- Configured Model Characteristics



Structured Data Output

- Extraction formatted data from LLM output



Human Centered Review and QC



Database ingestion



Fuzzy Performance Analysis

- Evaluate predicted matches
- Estimate performance
- Review Prompt Changes

Optimize Prompts and Model x Task

chemical_name	CASRN	lupac_name
1-(4-Chlorophenyl)-3-(3,4-dichlorophenyl)urea (Triclocarban)	101-20-2	not found
1-Bromo-4-chlorobutane	9840-78-9	not found
1-bromobutane	109-65-9	not found
1-Bromohexane	111-25-1	not found
1-Bromopentane	110-53-2	not found
1-Decanol	112-30-1	not found
1-Methyl-3-phenyl-1-piperazine	5271-27-2	not found
1,5-Hexadiene	502-42-7	not found
2-(Formylamino)-3-thiophenecarboxylic acid	43228-69-9	not found
2-Chloromethyl-3,5-dimethyl-4-methoxyphenyl HCl	86604-75-3	not found
2-Ethylhexyl-4-methylcinnamate (Octinoate)	3466-77-3	not found
2-Ethylhexyl-4-aminobenzoate	26218-04-2	not found
2-Hydroxy-4-methoxybenzophenone (Oxybenzone)	131-57-7	not found
2-Phenylethanol (Phenylethyl alcohol)	60-12-8	not found
2-Phenylhexanenitrile	3506-96-3	not found

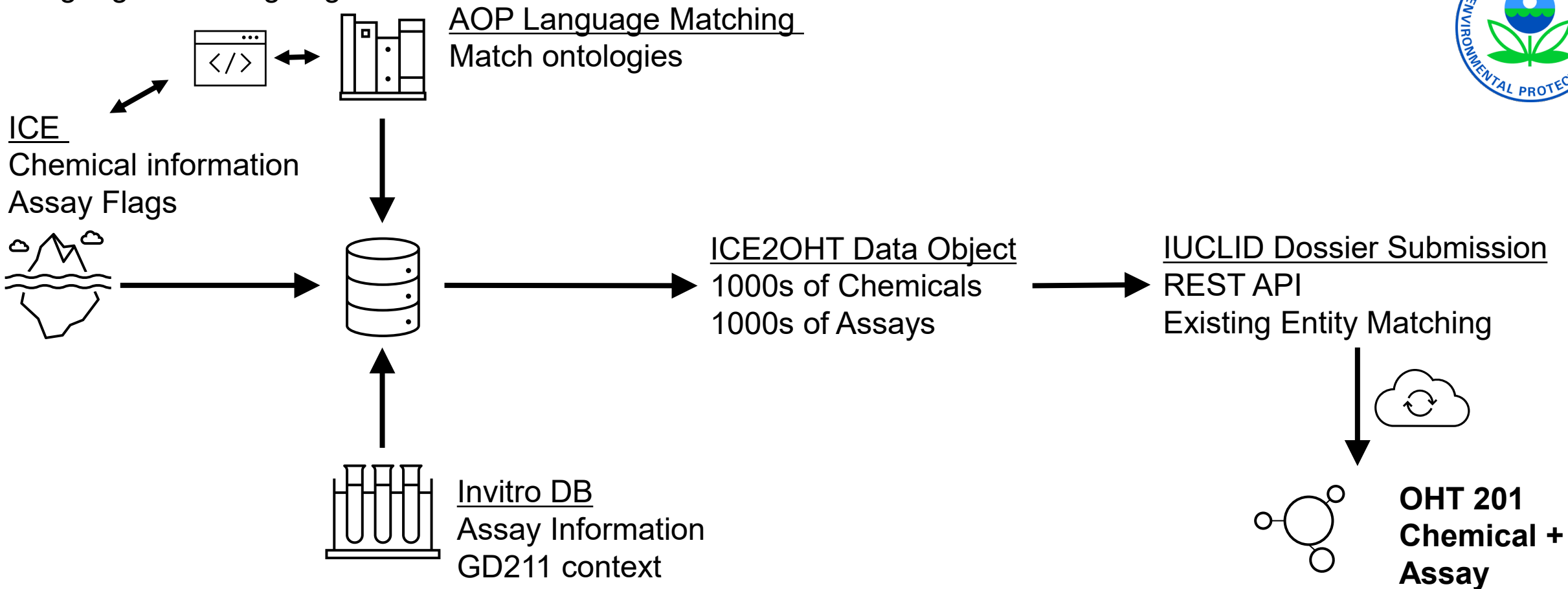
chemical_name	lab_id	experiment_id	test_id	physical_state	transdermal	between_lab_reproducibility	within_lab_reproducibility	predictive_capacity
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 1 (Bioscience Co.)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 2 (BIO)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 1	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 2	liquid	Yes	Yes	Yes	Yes
Decanol	Lab 3 (COSMOS)	not found	Bar 3	liquid	Yes	Yes	Yes	Yes

Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	100 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 3 (COSMOS)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source
Lab 2 (BIO)	Predictive	Human	UN GHS	Inherent	Specificity	96.7 %	30 not found	not found	not found	1000 VS vapt	30 of the source





Language matching engine



## DeTox: an *In-Silico* Alternative to Animal Testing for Predicting Developmental Toxicity Potential

**Authors:** Ricardo Scheufen Tieghi, Marielle Rath, José Teófilo Moreira-Filho, James Wellnitz, Holli-Joi Martin, Kathleen Gates, Helena T. Hogberg, Nicole Kleinstreuer, Alexander Tropsha, and Eugene N. Muratov | [AUTHORS INFO & AFFILIATIONS](#)

**Publication:** Environmental Health Perspectives • <https://doi.org/10.1289/EHP15307>

Model	# Compounds (Nontoxic/Toxic)	Accuracy	Sensitivity	Specificity	Positive Prediction Value	Negative Prediction Value
Overall	213 (63/150)	0.78	0.96	0.37	0.78	0.79
First Trimester	228 (67/161)	0.67	0.72	0.53	0.79	0.45
Second Trimester	256 (75/181)	0.74	0.82	0.53	0.81	0.56
Third Trimester	257 (76?181)	0.77	0.92	0.39	0.79	0.70

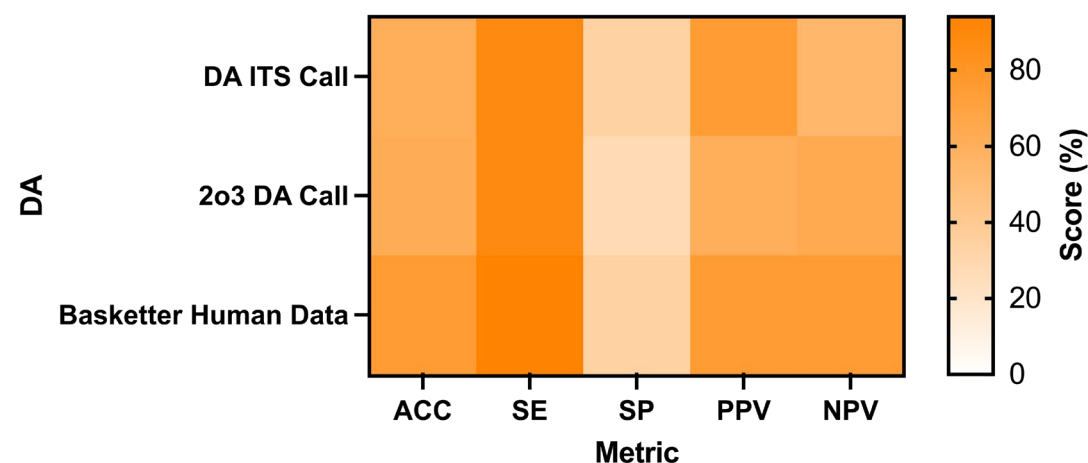
<https://detox.mml.unc.edu/>



Article

## A Novel Machine Learning Model and a Web Portal for Predicting the Human Skin Sensitization Effects of Chemical Agents

Ricardo Scheufen Tieghi <sup>1,2</sup>, José Teófilo Moreira-Filho <sup>1</sup>, Holli-Joi Martin <sup>2</sup>, James Wellnitz <sup>2</sup>, Miguel Canamary Otoch <sup>2</sup>, Marielle Rath <sup>2</sup>, Alexander Tropsha <sup>2,3,\*</sup>, Eugene N. Muratov <sup>2,\*</sup> and Nicole Kleinstreuer <sup>1,\*</sup>

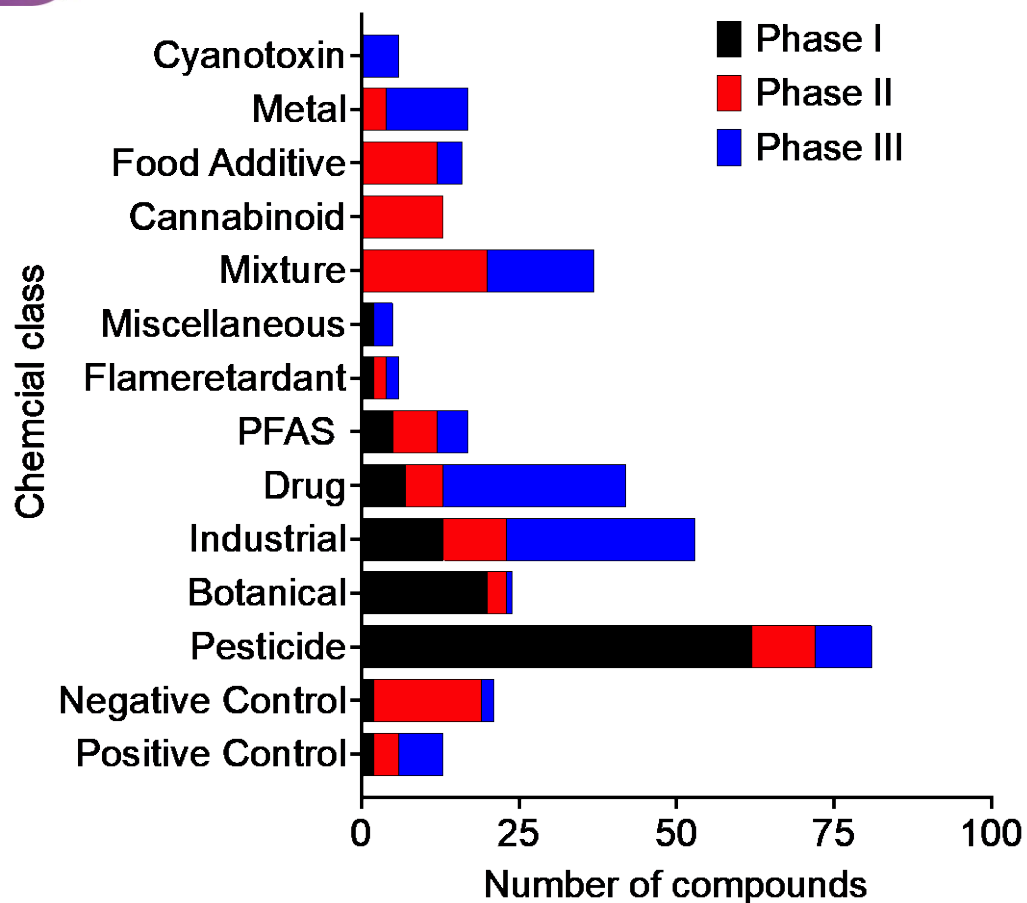


<https://husspred.mml.unc.edu/>



# DTTs Screening Efforts in a Battery of DNT In Vitro Assays

<https://www.niehs.nih.gov/research/atniehs/dtt/strategic-plan/health/developmental>



## Phase I: 115 Chemicals



## Phase II: 108 Chemicals



## Phase III: 128 Chemicals

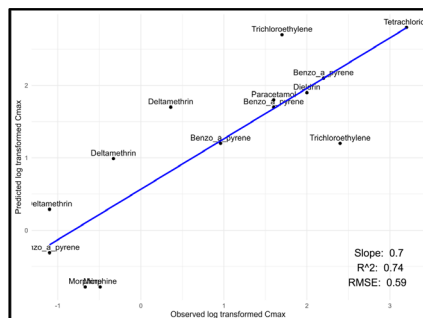
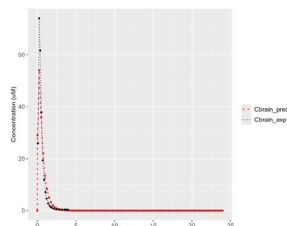




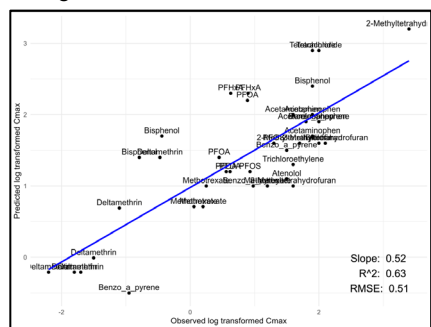
## Brain-Adipose Physiologically Based Kinetic (PBK) Model

- Built upon generic PBK model from EPA's httk R package (v2.2.2)
- Addition of brain and adipose compartments
- Diffusion-Limited brain compartment considering blood brain barrier permeability
- Validation using in vivo data for adipose, brain, and plasma concentrations

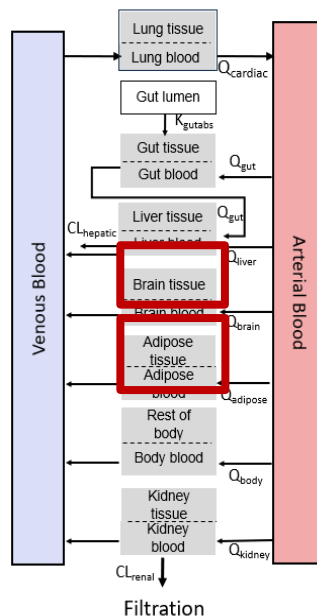
Acetaminophen  
Pred vs Obs  
Brain



Log10 Predicted vs Observed Brain

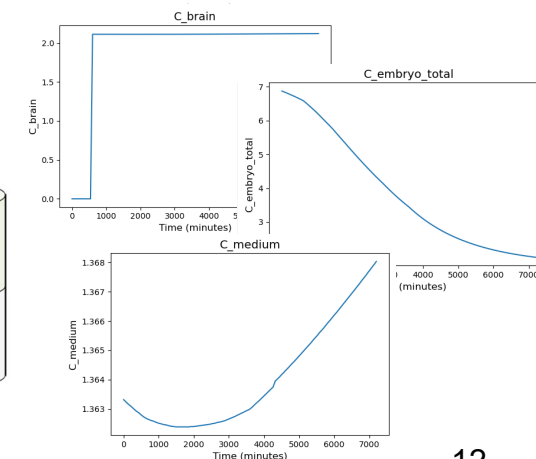
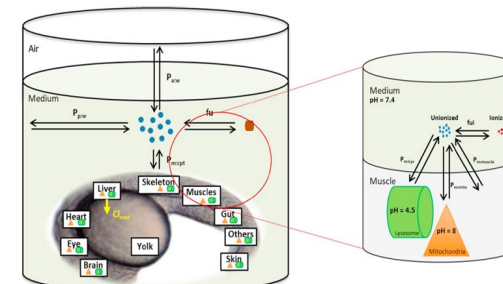


Log10 Predicted vs Observed Adipose



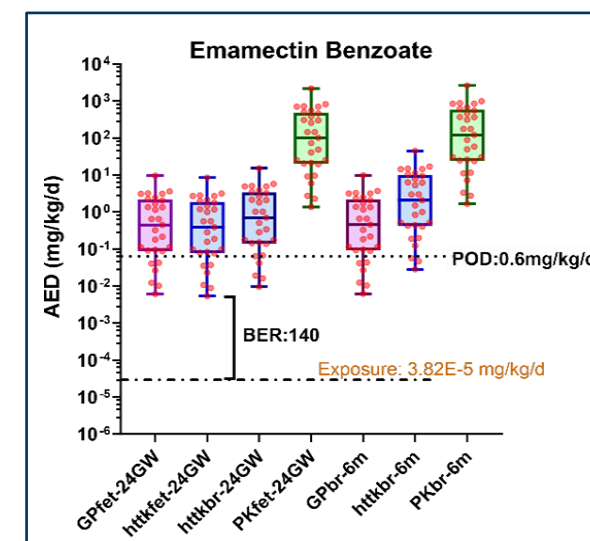
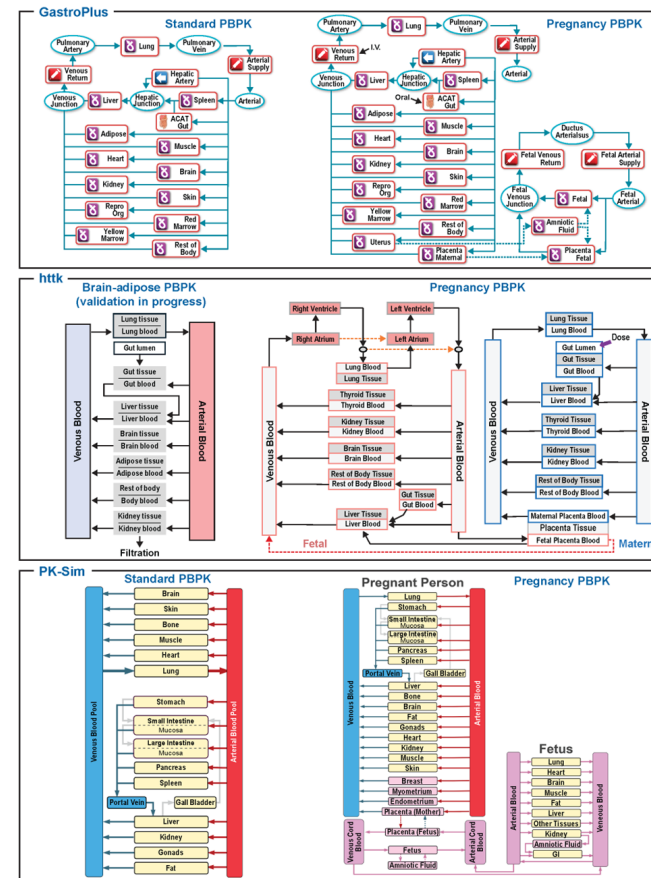
## Zebrafish PBK Model

- Model developed by Simeon et al., 2020
- Predictions for 10 organs and medium concentrations
- Incorporates developmental changes and metabolism
- Refining model with bioavailability data to expand applicability domain
- Applying to screening data from zebrafish behavioral assays conducted in extended DNT in vitro battery



## A Comparison of PBK Models

- DNT-IVIVE approach established to translate in vitro DNT activity doses into in vivo human doses that considers the site and period of brain development
- PBK models compared for DNT-IVIVE approach
- Chemicals bioactive in DNT NAMs from EPA with experimental toxicokinetic data
- Findings
  - In vivo data fall in the range of human administered equivalent dosages (AEDs), showing the concordance of DNT-IVIVE predictions with in vivo data
  - Approach relatively transferable across platforms, with differences explained by lipophilicity and partitioning algorithms
  - Uncertainties exist due to the lack of data available for validation



*Manuscript submitted to Tox Sci.*

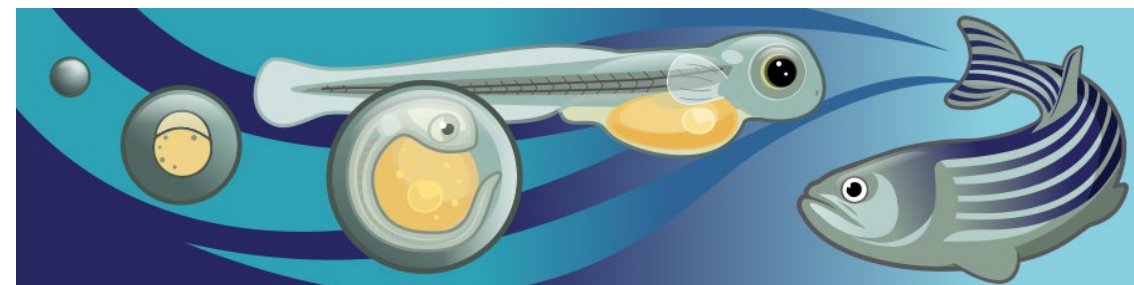
## **OECD DNT IATA Framework Template and Guidance of Support**

- EU (EFSA) and US (NICEATM) lead project, approved by WHPA in June 2024
- Develop an IATA framework template specific for DNT
  - Advance and provide guidance to address QIVIVE
  - Standardize uncertainty analyses for integration in WoE assessment
- Leverage on the existing and development of new DNT IATA case studies
- IATA framework template specific to the DNT regulatory endpoint
  - Several information sources
  - Multiple problem formulations
  - Consistent way to integrate data



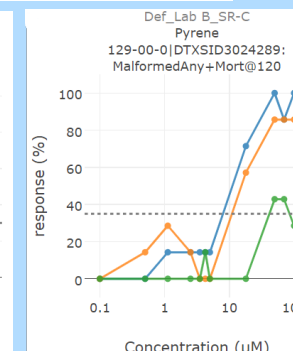
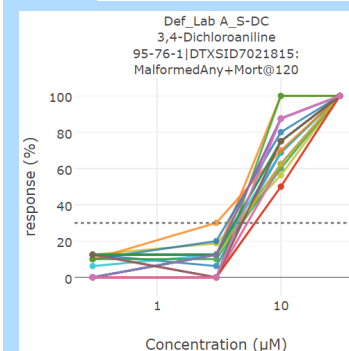
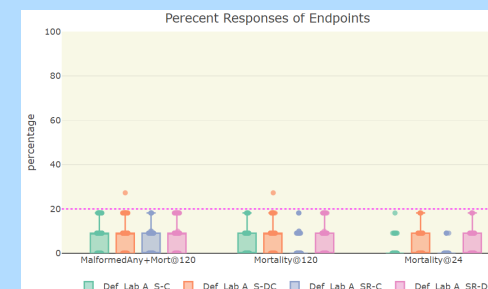


- Promotes broader adoption of zebrafish screening in toxicology.
- Interlaboratory studies were conducted to evaluate influence of experimental protocols on the assessment of developmental toxicity.
- Study design and data analysis approaches have been published.
- Results from the interlaboratory studies including a landscape analysis of phenotype responses are ongoing.



## Launch of SEAZIT-DIVER tool:

Explore and download protocol information, data, and visualizations



<https://seazit.dtt.niehs.nih.gov/seazit/>

Open Access Feature Paper Editor's Choice Article

## Interlaboratory Study on Zebrafish in Toxicology: Systematic Evaluation of the Application of Zebrafish in Toxicology's (SEAZIT's) Evaluation of Developmental Toxicity

by Jon T. Hamm<sup>1</sup>, Jui-Hua Hsieh<sup>2</sup> , Georgia K. Roberts<sup>2</sup>, Bradley Collins<sup>2</sup>, Jenni Gorospe<sup>3</sup>, Barney Sparrow<sup>3</sup>, Nigel J. Walker<sup>2</sup>, Lisa Truong<sup>4</sup> , Robyn L. Tanguay<sup>4</sup> , Sylvia Dyballa<sup>5</sup>, Rafael Miñana<sup>5,6</sup>, Valentina Schiavone<sup>5</sup> , Javier Terriente<sup>5</sup>, Andrea Weiner<sup>7</sup>, Arantza Muriana<sup>7</sup>, Celia Quevedo<sup>7</sup> and Kristen R. Ryan<sup>2,\*</sup>



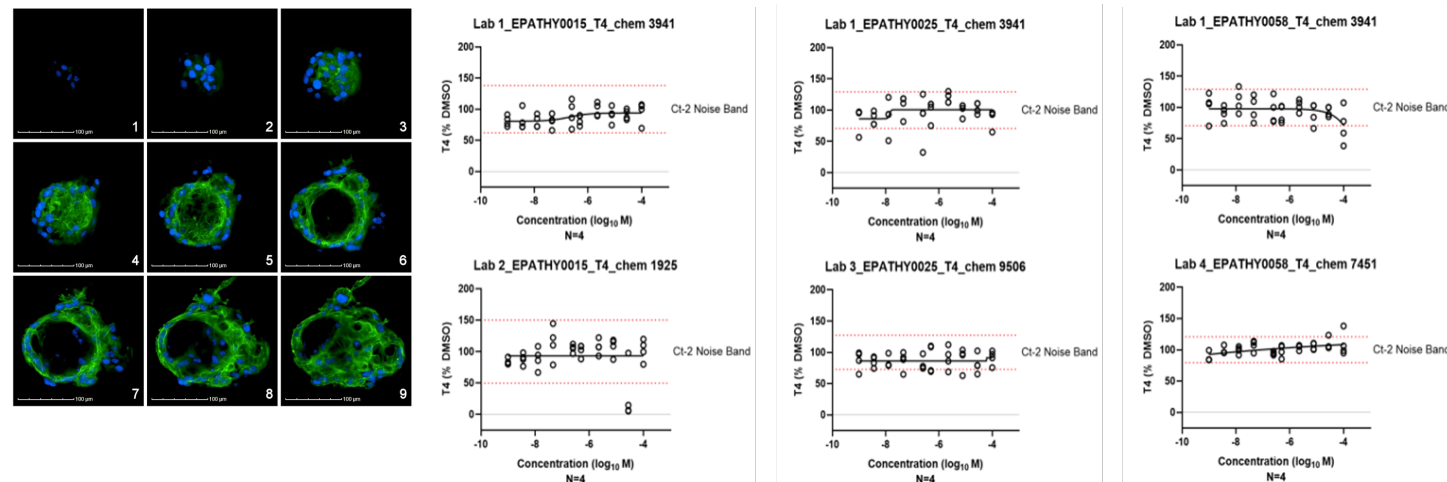
**SOT** Society of  
Toxicology  
academic.oup.com/toxsci

TOXICOLOGICAL SCIENCES, 2019, 1–16

doi: 10.1093/toxsci/afz238  
Advance Access Publication Date: December 6, 2019  
Research Article

## Development of an *In Vitro* Human Thyroid Microtissue Model for Chemical Screening

Chad Deisenroth <sup>\*,1</sup> Valerie Y. Soldatow,<sup>†</sup> Jermaine Ford,<sup>‡</sup> Wendy Stewart,<sup>\*</sup>  
Cassandra Brinkman,<sup>\*</sup> Edward L. LeCluyse,<sup>†</sup> Denise K. MacMillan,<sup>‡</sup> and  
Russell S. Thomas <sup>1\*</sup>



## Team Members

### Coordinator: NICEATM

Method Developer

Lab 1

Lab 4



Lab 2



**LifeNet Health**<sup>®</sup>  
Saving Lives. Restoring Health. Giving Hope.

Lab 3



**CORTEVA**<sup>™</sup>  
agriscience

### Status:

- Phase 1.2 complete (initial transfer phase, lab 2)
- Phase 1.3 complete (secondary transfer phase, labs 3 and 4)
- Phase 1.4 complete (transferability study)
- Pre-validation report and peer review publication are being drafted

## Identification of Skin Sensitizers

<https://ntp.niehs.nih.gov/go/ACDtest>

Substances with the potential to cause allergic contact dermatitis (ACD) are skin sensitizers; the process by which they cause ACD is skin sensitization. Non-animal approaches to identify potential skin sensitizers are becoming widely accepted. NICEATM, ICCVAM, and ICCVAM agencies have been at the forefront of advancing these non-animal approaches.



### GARD Models for Identifying Sensitizers

The GARD®skin assay uses a genomic profile to identify potential skin sensitizers. A NICEATM report evaluates the assay. [Read more »](#)



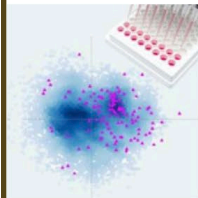
### Human Skin Sensitization Data

NICEATM and collaborators compiled human data as a resource to evaluate new testing approaches for skin sensitizers. [Read More »](#)



### Electrophilic Allergen Screening Assay (EASA)

The EASA is a chemical assay that measures a chemical's tendency to bind to proteins, the first step in skin sensitization. [Read More »](#)



### Defined Approaches to Identify Potential Skin Sensitizers

A defined approach uses input data and a data interpretation procedure to identify potential skin sensitizers. NICEATM and ICCVAM have developed and evaluated defined approaches. [Read More »](#)



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OECD » Publications » Guideline No. 497: Defined Approaches on Skin Sensitisation

## Guideline No. 497: Defined Approaches on Skin Sensitisation

Report

More info

OECD Guidelines for the Testing of Chemicals, Section 4 • 25 June 2025







## HPPT App (Rshiny tool) under development

**HPPT App** Home About Help

Upload HPPT Data File

Choose File No file selected.

[Download Input File Template \(.xlsx\)](#)

Select Potency Measure

DSA1+

Run

Download All Results

Chemical Identifier	Conc (%)	DSA (ug/cm2)	No. Test Subjects	No. Positive	Call	DSA1+	DSA5%	Ex.C	WES_indiv
All	All	All	All	All	All	All	All	All	All
5870-93-9	12	8100	24	5	Active	1620	1944	1B	1
626-82-4	4	2700	30	2	Active	1350	2025	1B	1
Ammoniated mercury	2	1296	100	14	Active	93	463	1A	2
DTXSID8039241	4	2700	25	1	Active	2700	3375	1B	1
10032-02-7	6	3888	25	0	Inactive	N/A	N/A	NC/1B	0.5

**Overall Weight of Evidence GHS Classification**

Skin sensitization classifications for each input chemical using the modified GHS classification approach. Results are based on consolidating classifications from three individual methods (Table 2). Detailed information is available in the App's Userguide on the Help page.

CSV Excel

Show 10 entries

Search:

Chemical Identifier	Wof_bin	Wof_sub	Wof_border	Total Tests
All	All	All	All	All
10032-02-7	N/A	N/A	NC/1B	1
5870-93-9	1	1B	1B	1
626-82-4	1	1B	1B	1
Ammoniated mercury	1	1A	1A	1
DTXSID8039241	1	1B	1B	1

Showing 1 to 5 of 5 entries

Previous 1 Next

## Database and proposed HPPT GHS classification approach

Archives of Toxicology (2023) 97:2825–2837  
<https://doi.org/10.1007/s00204-023-03530-3>

REVIEW ARTICLE



### A database of human predictive patch test data for skin sensitization

Judy Strickland<sup>1</sup> · Jaleh Abedini<sup>1</sup> · David G. Allen<sup>1</sup> · John Gordon<sup>2</sup> · Victoria Hull<sup>1</sup> · Nicole C. Kleinstreuer<sup>3</sup> · Hon-Sum Ko<sup>4</sup> · Joanna Matheson<sup>2</sup> · Hermann-Josef Thierse<sup>5</sup> · James Truax<sup>1</sup> · Jens T. Vanselow<sup>5</sup> · Matthias Herzler<sup>5</sup>

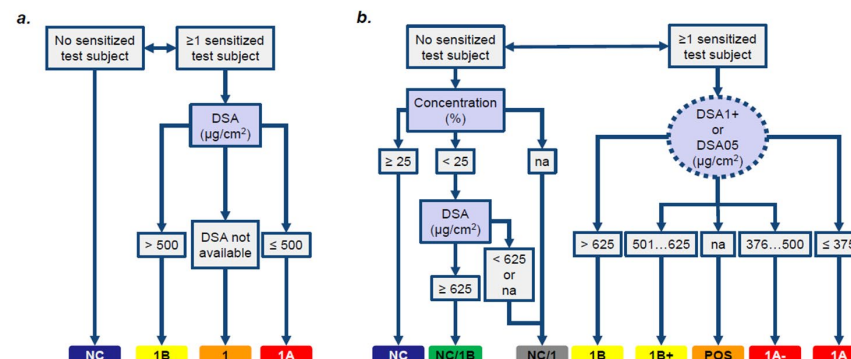
Archives of Toxicology (2024) 98:1253–1269  
<https://doi.org/10.1007/s00204-023-03656-4>

REVIEW ARTICLE



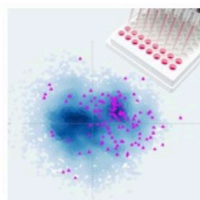
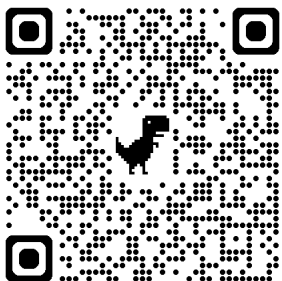
### Use of human predictive patch test (HPPT) data for the classification of skin sensitization hazard and potency

Matthias Herzler<sup>1</sup> · Jaleh Abedini<sup>2</sup> · David G. Allen<sup>2</sup> · Dori Germolec<sup>3</sup> · John Gordon<sup>4</sup> · Hon-Sum Ko<sup>5</sup> · Joanna Matheson<sup>4</sup> · Emily Reinke<sup>2</sup> · Judy Strickland<sup>2</sup> · Hermann-Josef Thierse<sup>1</sup> · Kim To<sup>2</sup> · James Truax<sup>2</sup> · Jens T. Vanselow<sup>1</sup> · Nicole Kleinstreuer<sup>6</sup>



(a) Standard GHS Classification Approach (b) Modified Classification Approach 18

# The Skin Allergy Risk Assessment (SARA) – ICE Model



## Defined Approaches to Identify Potential Skin Sensitizers

A defined approach uses input data and a data interpretation procedure to identify potential skin sensitizers. NICEATM and ICCVAM have developed and evaluated defined approaches.

[Read More »](#)



Contents lists available at [ScienceDirect](#)

## Current Research in Toxicology

journal homepage: [www.journals.elsevier.com/current-research-in-toxicology](http://www.journals.elsevier.com/current-research-in-toxicology)



### Research Paper

The skin allergy risk assessment-integrated chemical environment (SARA-ICE) defined approach to derive points of departure for skin sensitization

Emily N. Reinke<sup>a,\*</sup>, Joe Reynolds<sup>b</sup>, Nicola Gilmour<sup>b</sup>, Georgia Reynolds<sup>b</sup>, Judy Strickland<sup>a,1</sup>, Dori Germolec<sup>c</sup>, David G. Allen<sup>a,2</sup>, Gavin Maxwell<sup>b</sup>, Nicole C. Kleinstreuer<sup>c</sup>



OECD Publications > Guideline No. 497: Defined Approaches on Skin Sensitisation

## Guideline No. 497: Defined Approaches on Skin Sensitisation

Report  
More info



OECD Guidelines for the Testing of Chemicals, Section 4 • 25 June 2025

A SARA-ICE is a Bayesian statistical model which infers a human-relevant metric of sensitiser potency (termed ED<sub>01</sub>), the dose with a 1% chance of human skin sensitisation.

- Accounts for variability of the input data and explicitly quantifies uncertainty
- Utilises any combination of human repeat insult patch test (HRIPT), LLNA, direct peptide reactivity assay (DPRA), KeratinoSens™, h-CLAT, U-SENS™ data
- Derive a Point-of-departure for use in risk assessment (geometric mean)

## Steps for Using the SARA-ICE Tool

Steps 1-4 outline the process for conducting an analysis using the SARA-ICE tool. Refer to the User Guide for the purpose of the tool, the types of data that can be analyzed, how to use the tool and underlying models, and additional details on each step:

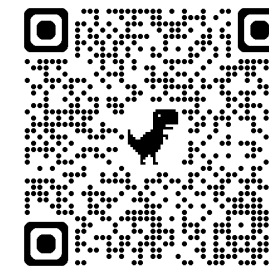
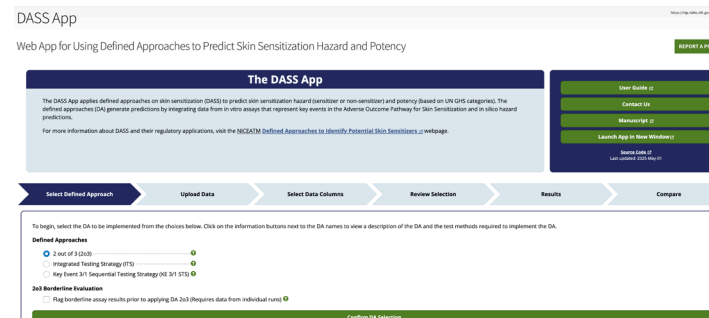
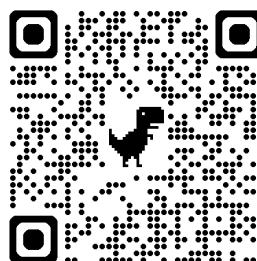
[Download User Guide](#)

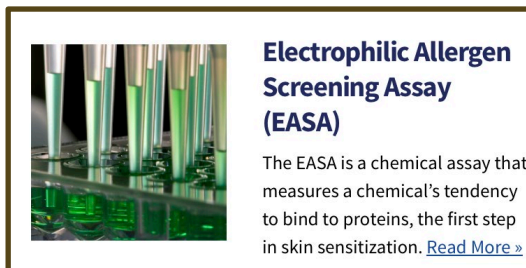
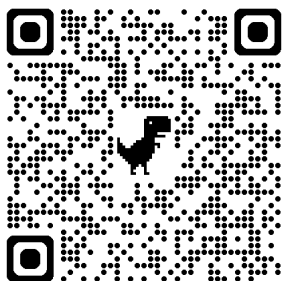
1. Create and [Select Input File](#)
2. [Select Model](#) either the 'SARA-ICE OECD TG 497 Defined Approach (version 1.0)' or the 'SARA-ICE Extended Model (version 1.0)'
3. After selecting a model, the analysis will automatically start running. You can [View Analysis](#) during and after processing
4. After analysis is finished, you can [Download Analysis](#)

## Help & Support

Have questions or need to report an issue? Please email [ICE-support@niehs.nih.gov](mailto:ICE-support@niehs.nih.gov)

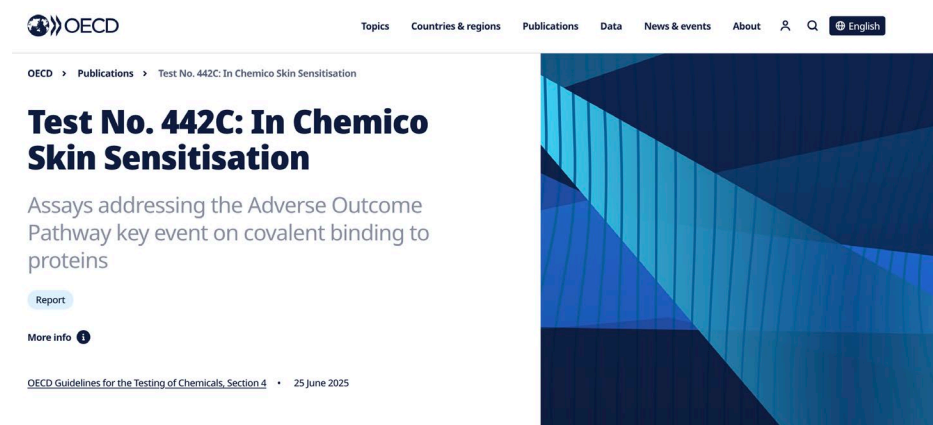
Version 0.2.5-beta





- Addresses KE1 in the Skin Sensitization AOP
- In chemico plate-based assay
  - Measures protein reactivity of a chemical via fluorescent or colorimetric probes
- Multi-lab validation study
  - Participating labs: U.S. FDA, DoD, CPSC/NIST, BRT, Inc.
  - Utilize 2019 OECD Performance Standards for KE1-based assays for validation study
  - Peer Review of validation study nearly finished
- Accepted on to 2024 OECD workplan for inclusion in TG 442C – meetings with expert group summer 2025

Lab #	Balanced Accuracy	Sensitivity	Specificity	Within Lab Reproducibility	Between Lab Reproducibility
1	76%	85%	67%	94%	96%
2	82%	92%	71%	100%	
3	84%	85%	83%	97%	
4	84%	85%	83%	94%	
Mean	82%	87%	76%	96%	





# Defined Approaches for Predicting Eye Irritation Classifications

NICEATM, PETA Science Consortium International, and EPA Office of Pesticide Programs collaborated to test eye irritation/corrosion potential of 29 agrochemical formulations in a common set of in vitro methods

- Developed four defined approaches (DAs) applicable to GHS and EPA classification systems:
  - BCOP with histopathology (DA-BCOP+)
  - EO + BCOP with histopathology (DA-EO+)
  - TTL+ BCOP with histopathology (DA-TTL+)
  - EyeIRR-IS + BCOP with histopathology (DA-EyeIRR-IS+)
- Instead of evaluating direct concordance of the four individual DAs with historical in vivo data, for each formulation, we assessed orthogonal concordance of GHS and EPA classifications predicted across all five approaches
- Also evaluated associated hazard labeling (GHS) and PPE labeling (EPA) predicted across all five approaches
- All DAs performed **as well as or better** than the in vivo test and generally resulted in hazard/PPE labeling that is **more protective of human health**
- Publications:
  - NTP report (2021; addended 2025) <https://doi.org/10.22427/NTP-NICEATM-1>
  - van der Zalm et al. (2024) <https://doi.org/10.1080/15569527.2023.2275029>
  - Daniel et al. (2025) <https://doi.org/10.1080/15569527.2025.2499552>

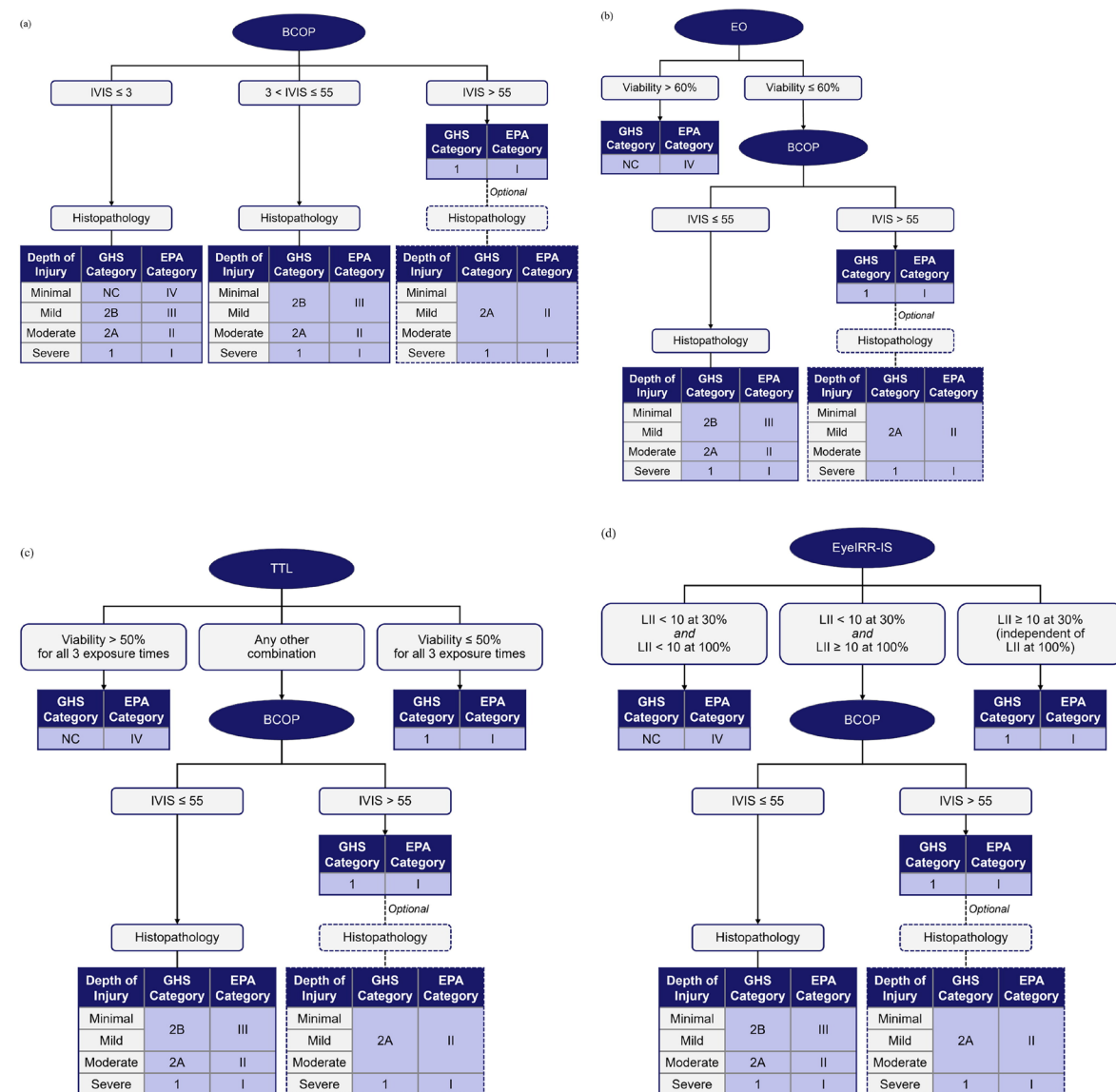


Figure 7 Daniel et al., 2025



# Advancing Quantitative Analysis in Human Health Assessments through Probabilistic Methods

## Workshop Monday, October 7-8, 2024

Convened international experts to facilitate discussion of probabilistic methods in human health risk assessment. The workshop provided examples of application of probabilistic methods in chemical risk assessments, highlighted ongoing research, and discussed the needs and challenges for the regular use of these methods.

### Broken into four subtopics, the agenda featured sessions on:

- Probabilistic exposure
- Toxicokinetics
- Benchmark dose modeling
- Toxicity value determination

### Accomplishments:

- Greater understanding of probabilistic methods within the context of human health assessments
- Guided discussion for how these methods may be implemented



<https://ntp.niehs.nih.gov/whatwestudy/niceatm/3rs-meetings/past-meetings/probabilistic-methods-wksp>





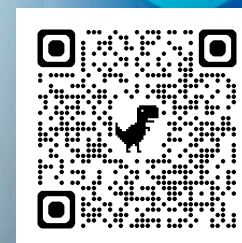
National Institute of  
Environmental Health Sciences  
*Division of Translational Toxicology*

# Acknowledgments

## The NICEATM Group



## NIEHS/DTT Contributors



<https://ntp.niehs.nih.gov/iccvmreport/2023>



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