

OPERA: Open-Source QSAR Models for Regulatory Support

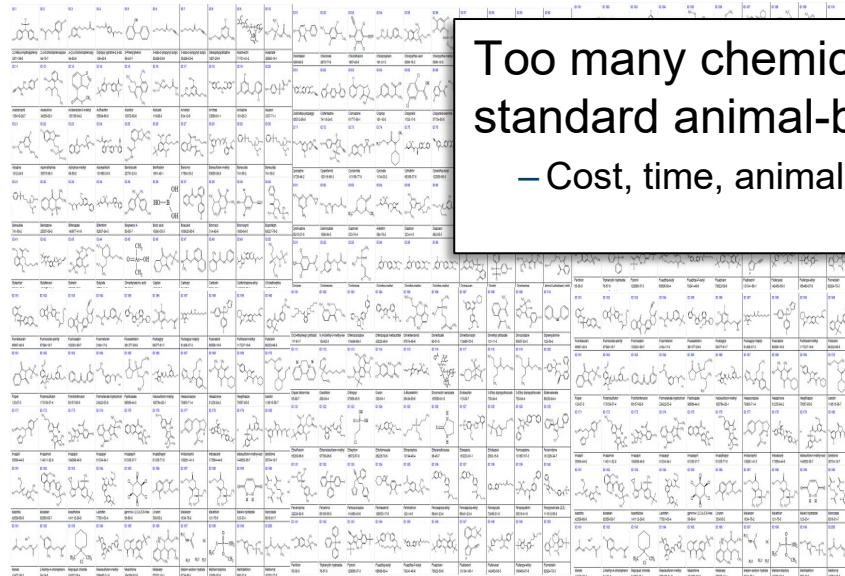
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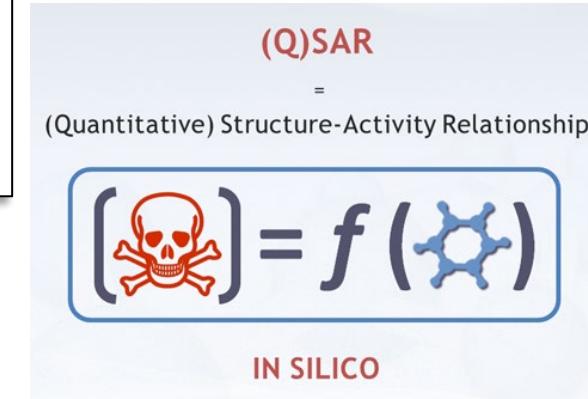
In silico screening



Too many chemicals to test with
standard animal-based methods

- Cost, time, animal welfare

Alternative

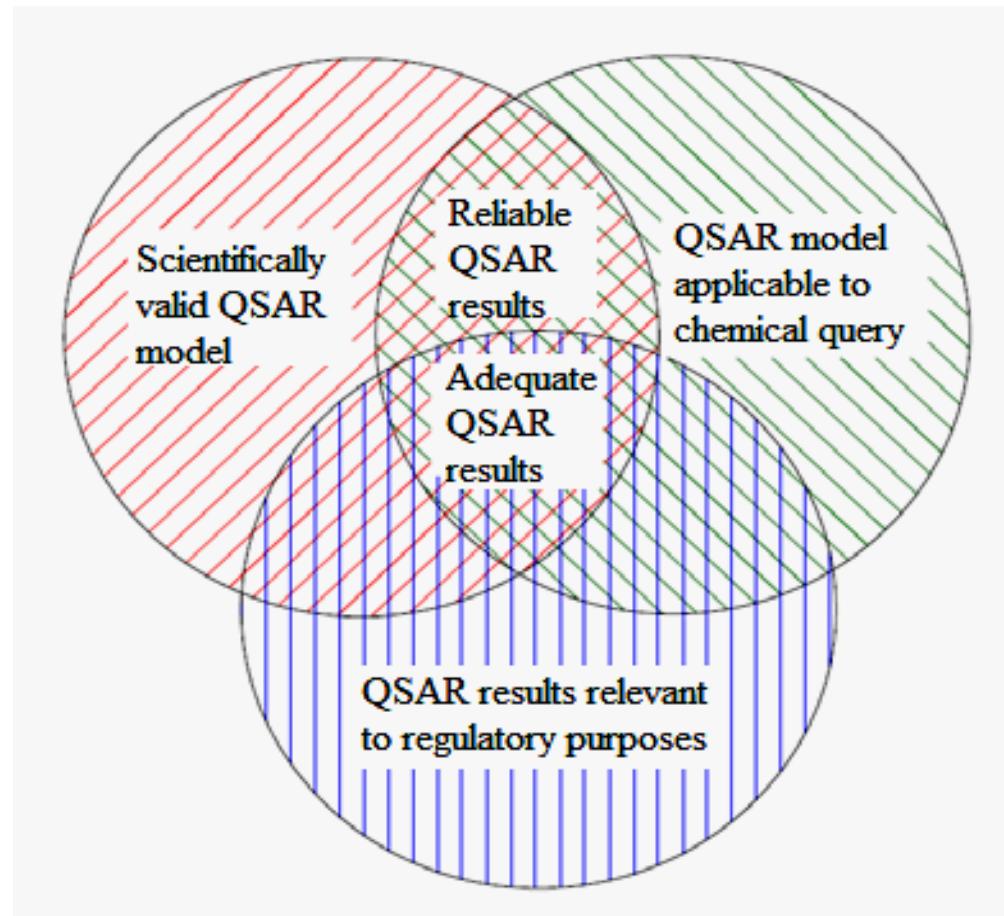


- Organic **pollutants** with exposure potential **accumulate** in body tissues
 - Cause **toxic effects** to wildlife and humans
- Existence of **gaps in the experimental data** for environmental endpoints
 - Need to fill the data gaps and bridge the **lack of knowledge**
- **Regulatory requirements:**
 - Reduce **animal testing, time and costs**
 - **Methodology:** use of **QSAR/QSPR** to predict the **endpoints** of interest.

QSARs for regulatory purposes

The 5 OECD Principles

- 1) A defined endpoint
- 2) An unambiguous algorithm
- 3) A defined domain of applicability
- 4) Appropriate measures of goodness-of-fit, robustness and predictivity
- 5) Mechanistic interpretation, if possible



[Journal of Cheminformatics](#)
December 2018, 10:10 | [Cite as](#)

OPERA models for predicting physicochemical properties
and environmental fate endpoints



OPERA approach

- Curated **open** access datasets (<https://doi.org/10.1186/s13321-018-0263-1>)
- **Open-source** code (github.com/NIEHS/OPERA)
- **Transparent** unambiguous algorithms (<https://qsardb.jrc.ec.europa.eu/qmrf/>)
- **Transparent** validated performances (<https://doi.org/10.1080/1062936X.2016.1253611>)
- **Defined** applicability domain and limitations of the models
- Predictions **available** through:
 - NICEATM's Integrated Chemical Environment (<https://ice.ntp.niehs.nih.gov/>)
 - The EPA's CompTox Dashboard (<https://comptox.epa.gov/dashboard>)
 - Free and open-source standalone application (github.com/NIEHS/OPERA)



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

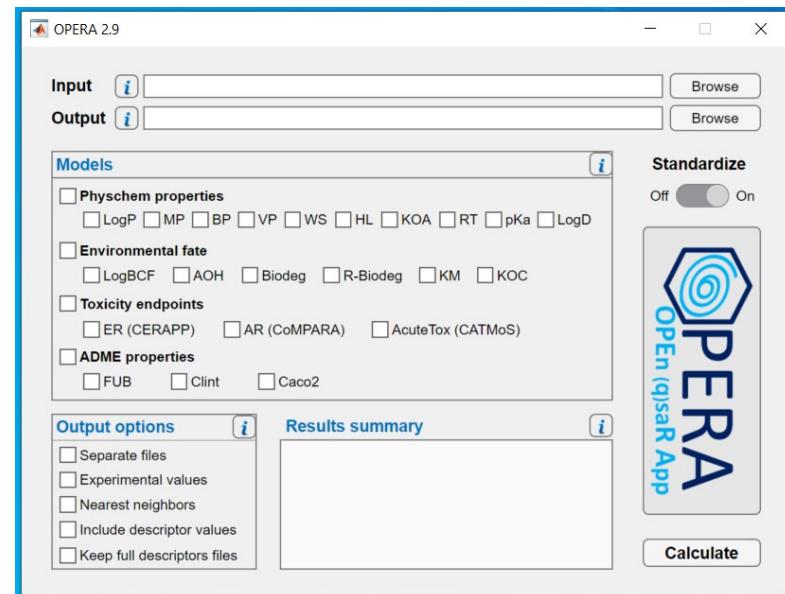
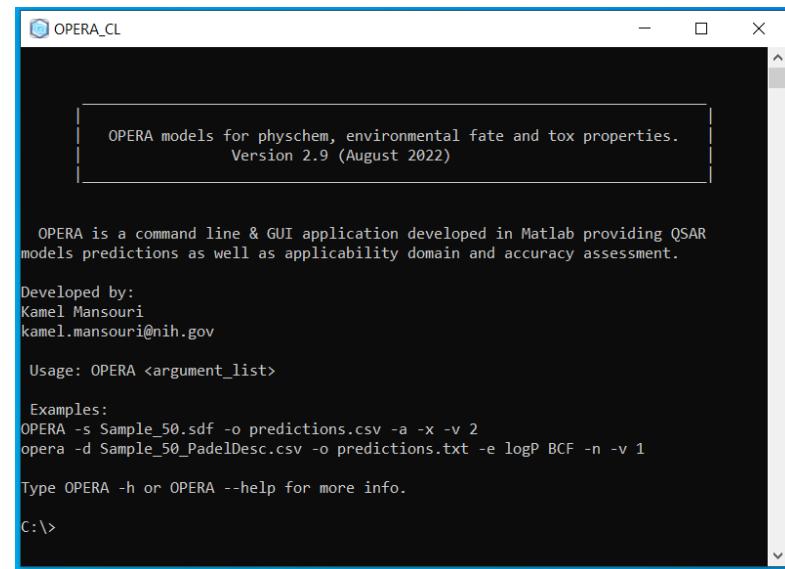
Input options:

- 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>



OPERA models (version 2.9)

Physchem properties		Chemicals	Version	Environmental fate		Chemicals	Version
BP	Boiling Point	7860	2.9	AOH	Atmospheric Hydroxylation Rate	692	2.6
HL	Henry's Law Constant	2233	2.9	BCF	Bioconcentration Factor	626	2.6
LogP	Octanol-water Partition Coefficient	18154	2.9	BioHL	Biodegradation Half-life	150	2.6
MP	Melting Point	22554	2.9	RB	Ready Biodegradability	1603	2.6
VP	Vapor Pressure	6764	2.9	KM	Fish Biotransformation Half-life	541	2.6
WS	Water Solubility	9943	2.9	KOC	Soil Adsorption Coefficient	728	2.6
pKa	Acid Dissociation Constant	6503	2.6				
KOA	Octanol/Air Partition Coefficient	270	2.6				

Toxicity endpoints		Chemicals	Version	ADME properties		Chemicals	Version
ER	Estrogen Receptor Activity	32464	2.6	FUB	Fraction unbound	3229	2.8
AR	Androgen Receptor Activity	47673	2.6	Clint	Intrinsic clearance	1346	2.8
AcuteTox	Acute Oral Systemic Toxicity	50660	2.6	CACO2	Caco-2 permeability	4601	2.8

New/updated since 2021

Version 3 coming soon!

PFAS in OPERA models

Models		PFAS/Total
BP	Boiling Point	346/7860
LogP	Octanol-water Partition Coef.	97/18154
MP	Melting Point	409/22554
VP	Vapor Pressure	178/6764
WS	Water Solubility	105/9943

Environmental Chemistry

VP Vapor Pressure					
Deviation of $\log P_p$ (Pa) estimates from experimental measurements at 25 °C					
	COSMOtherm	EPI Suite	NICEATM	ACD/Labs	TEST
MAE	1.22	1.08	1.46	1.53	1.31
RMSE	1.48	1.48	2.06	1.99	1.91

LogP Octanol-water Coef					
Deviation of octanol–water partition ratio ($\log K_{ow}$) estimates from experimental measurements at 25 °C					
	COSMOtherm	EPI Suite	NICEATM	ACD/Labs	OPERA
MAE	0.41	0.25	0.68	0.61	0.21
RMSE	0.50	0.29	1.13	0.70	0.28

WS Water solubility					
Deviation of water solubility ($\log S$; mg/L) estimates from experimental measurements at 25 °C					
	COSMOtherm	EPI Suite	NICEATM	TEST ^a	OPERA
MAE	0.35	1.82	2.38	0.95	0.23
RMSE	0.41	2.20	2.55	1.36	0.36

Property Estimation of Per- and Polyfluoroalkyl Substances: A Comparative Assessment of Estimation Methods

Alina Lampic and J. Mark Pamis*

Chemical Properties Research Group (Canadian Environmental Modelling Centre), Department of Chemistry, Trent University, Peterborough, Ontario, Canada

International collaborative projects

CERAPP

Collaborative Estrogen Receptor
Activity Prediction Project (2015/16)

Mansouri et al. (<https://doi.org/10.1289/ehp.1510267>)

CoMPARA

Collaborative Modeling Project for
Androgen Receptor Activity (2017/18)

Mansouri et al. (<https://doi.org/10.1289/EHP5580>)

CATMoS

Collaborative Acute Toxicity Modeling
Suite (2017/18)

Kleinstreuer et al. (<https://doi.org/10.1016/j.comtox.2018.08.002>)

Mansouri et al. (<https://doi.org/10.1289/EHP8495>)



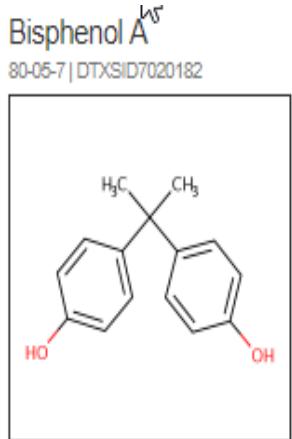
Endocrine Disruptor Screening Program



Acute Toxicity Workgroup: alternative methods

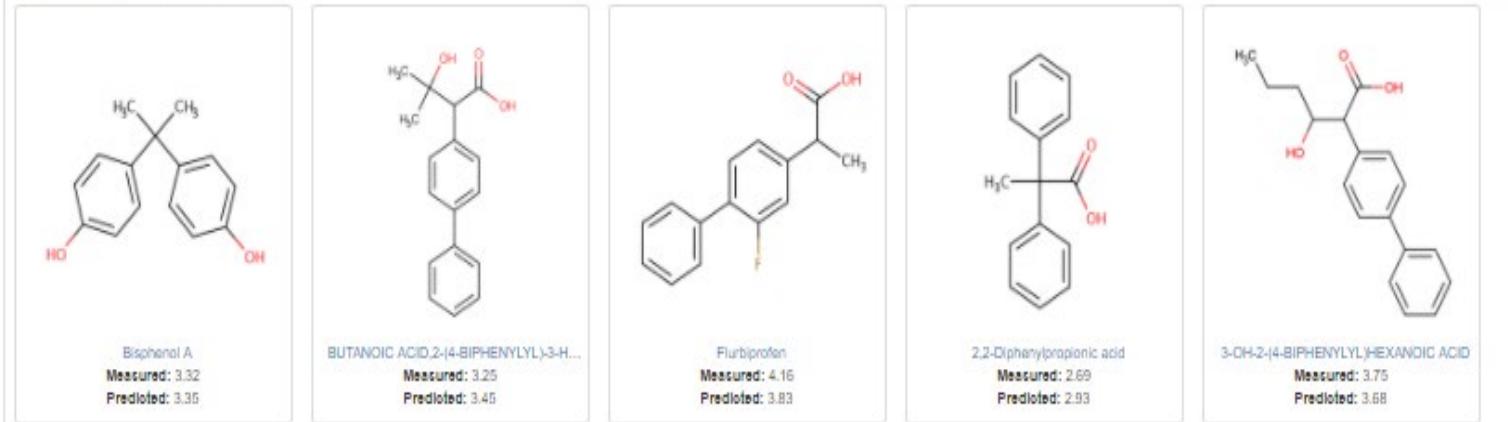
ICCVAM: Interagency Coordinating Committee on the Validation of Alternative Methods

OPERA output



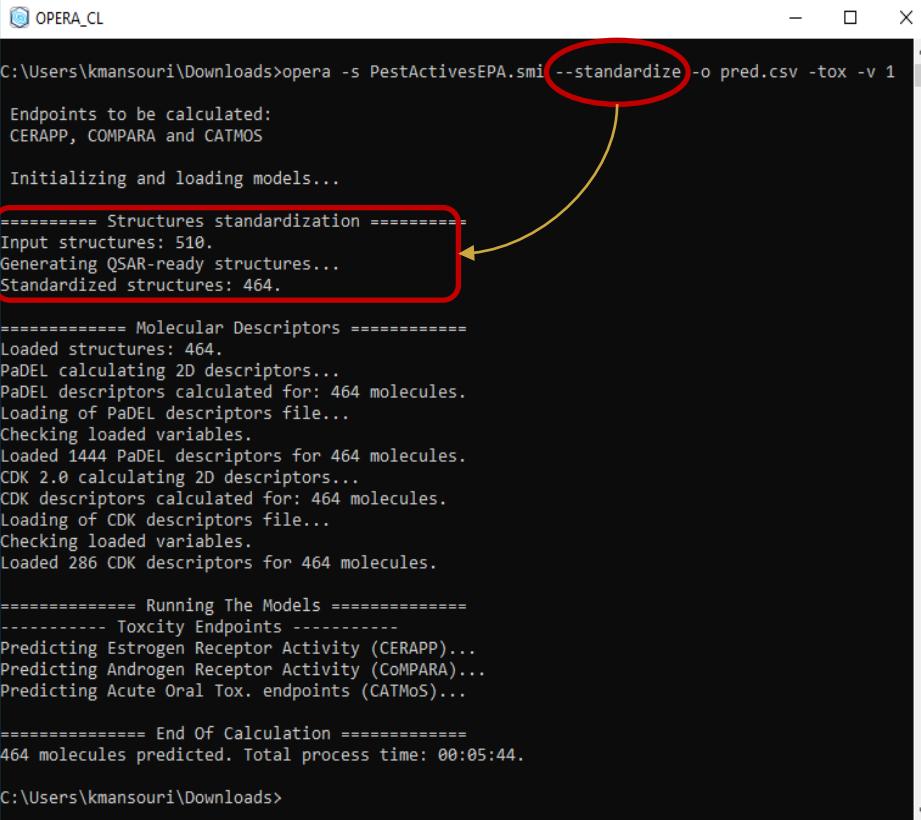
- Predicted value
- Experimental value, if available
- Global applicability domain
- Local applicability domain index [0:1]
- Accuracy index [0:1]
- Confidence interval [min:max]
- 5 nearest neighbors:
 - Identifiers (CAS, DTXSID, InChiKey)
 - Experimental values
 - Predicted values
- Molecular descriptors values

Nearest Neighbors from the Training Set



MoleculeID	DTXSID7020182
LogP_exp	3.32
LogP_pred	3.32
LogP_predRange	[3.27:3.37]
AD_LogP	1
AD_index_LogP	1
Conf_index_LogP	0.948
LogP_CAS_neighbor_1	80-05-7
LogP_CAS_neighbor_2	93371-55-2
LogP_CAS_neighbor_3	85-27-8
LogP_CAS_neighbor_4	5104-49-4
LogP_CAS_neighbor_5	5558-66-7
LogP_InChiKey_neighbor_1	IISBACLAFKSPIT-I
LogP_InChiKey_neighbor_2	ZNPAJPOVURPJF
LogP_InChiKey_neighbor_3	PQSXNIMHIHYFE
LogP_InChiKey_neighbor_4	SYTBZMRGLBWN
LogP_InChiKey_neighbor_5	ODELFXJUOVNEF
LogP_DTXSID_neighbor_1	DTXSID7020182
LogP_DTXSID_neighbor_2	DTXSID5087493C
LogP_DTXSID_neighbor_3	DTXSID50100531
LogP_DTXSID_neighbor_4	DTXSID0037231
LogP_DTXSID_neighbor_5	DTXSID4020412E
LogP_Exp_neighbor_1	3.32
LogP_Exp_neighbor_2	3.25
LogP_Exp_neighbor_3	2.11
LogP_Exp_neighbor_4	4.16
LogP_Exp_neighbor_5	2.69
LogP_pred_neighbor_1	3.18
LogP_pred_neighbor_2	3.66
LogP_pred_neighbor_3	3.5
LogP_pred_neighbor_4	3.05
LogP_pred_neighbor_5	3.44

QSAR-ready standardization tool



OPERA_CL

```
C:\Users\kmansouri\Downloads>opera -s PestActivesEPA.smi --standardize -o pred.csv -tox -v 1
Endpoints to be calculated:
CERAPP, COMPARA and CATMOS
Initializing and loading models...
===== Structures standardization =====
Input structures: 510.
Generating QSAR-ready structures...
Standardized structures: 464.

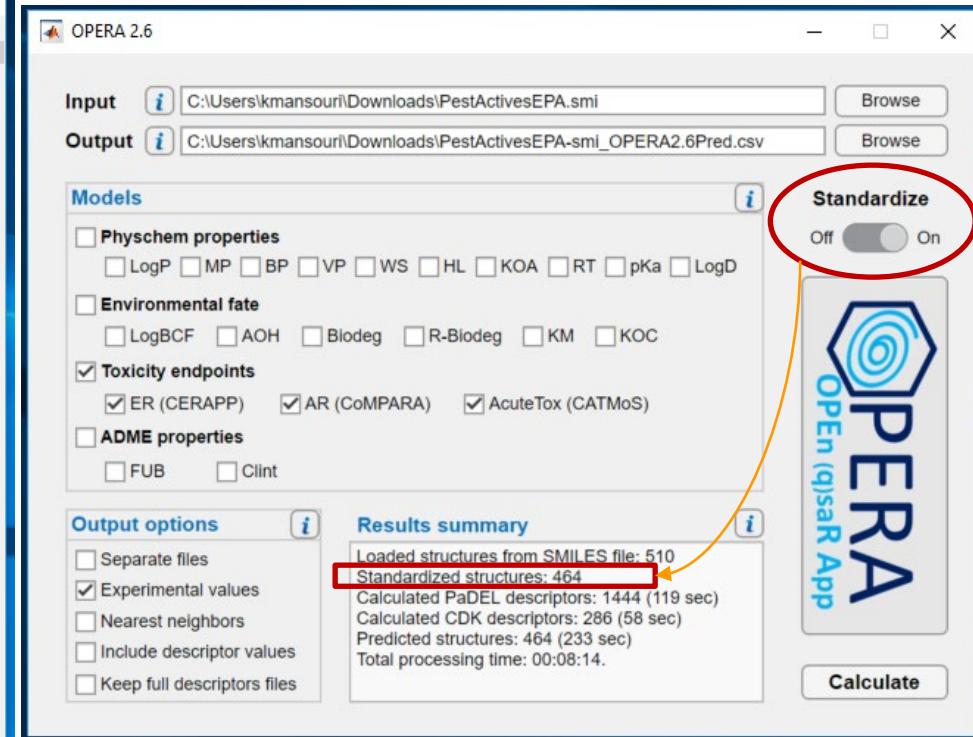
===== Molecular Descriptors =====
Loaded structures: 464.
PaDEL calculating 2D descriptors...
PaDEL descriptors calculated for: 464 molecules.
Loading of PaDEL descriptors file...
Checking loaded variables.
Loaded 1444 PaDEL descriptors for 464 molecules.
CDK 2.0 calculating 2D descriptors...
CDK descriptors calculated for: 464 molecules.
Loading of CDK descriptors file...
Checking loaded variables.
Loaded 286 CDK descriptors for 464 molecules.

===== Running The Models =====
----- Toxicity Endpoints -----
Predicting Estrogen Receptor Activity (CERAPP)...
Predicting Androgen Receptor Activity (CoMPARA)...
Predicting Acute Oral Tox. endpoints (CATMoS)...

===== End Of Calculation =====
464 molecules predicted. Total process time: 00:05:44.

C:\Users\kmansouri\Downloads>
```

Command line



Graphical user interface



Predictions on NICEATM's ICE

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

The screenshot displays the NICEATM's ICE interface. At the top left is the National Toxicology Program logo. The top right features a search bar and navigation links: Calendar & Events, News & Media, Get Involved, and Support. A large banner in the center promotes the OPERA OPEn (q)saR App, which includes a play button and three sub-tools: Search, Chemical Quest, and Curve Surfer.

The main workspace is divided into sections:

- Input:** Includes tabs for Results, Run, Reset, and Union or Intersection (set to Union). Below these are sections for Chemical Input (Select Chemicals, Quick List CASRN, User CASRNs) and Assay Input (Select Assays, listing various assays like CERAPP, ER Binding, CoMPARA, AR Binding, etc.).
- Chemical Input:** Shows a table of substances with their properties and assay results. The table includes columns for Substance Name, CASRN, DTXSID, and various assay calls (e.g., CATMoS, Rat Acute Oral Toxicity LD50, CoMPARA, AR Agonist Call, CERAPP, ER Antagonist Call).

Substance Name	CASRN	DTXSID	CATMoS, Rat Acute Oral Toxicity LD50	CoMPARA, AR Agonist Call	CoMPARA, AR Antagonist Call	CoMPARA, AR Binding Call	CERAPP, ER Agonist Call	CERAPP, ER Antagonist Call	CERAPP, ER Binding Call
Acetaminophen	103-90-2	DTXSID2020006	1625	Inactive	Inactive	Inactive	Inactive	Inactive	Inactive
Endosulfan	115-29-7	DTXSID1020560	2.26	Inactive	Inactive	Inactive	Inactive	Active	Active
3-Phenylprop-2-enal	104-55-2	DTXSID1024835	2568	Inactive	Inactive	Inactive	Inactive	Active	Active
Acrolein	107-02-8	DTXSID5020023	40	Inactive	Inactive	Inactive	Inactive	Inactive	Inactive
Linalool	78-70-6	DTXSID7025502	2097	Inactive	Inactive	Inactive	Inactive	Inactive	Inactive



Predictions on the EPA Dashboard

Single chemical

EPA United States Environmental Protection Agency

Endosulfan
115-29-7 | DTXSID1020560
Searched by DSSTox Substance Id.

ToxCast: Models
ToxCast Model Predictions

[Download ToxCast Model Predictions](#)

Model	Receptor	Agonist	Antagonist	Binding
ToxCast Pathway Model (AUC)	Androgen	0.00	7.34e-2	-
ToxCast Pathway Model (AUC)	Estrogen	2.95e-2	0.00	-
COMPARA (Consensus)	Androgen	Inactive	Inactive	Inactive
CERAPP Potency Level (From Literature)	Estrogen	-	Inactive (NaN)	Active (Weak)
CERAPP Potency Level (Consensus)	Estrogen	Inactive (Inactive)	Active (VeryWeak)	Active (VeryWeak)

DETAILS
EXECUTIVE SUMMARY
PROPERTIES
ENV. FATE/TRANSPORT
HAZARD
ADME
EXPOSURE
BIOACTIVITY
 TOXCAST: SUMMARY
 EDSP21
 TOXCAST/TOX21
 PUBCHEM
TOXCAST: MODELS
SIMILAR COMPOUNDS

<https://comptox.epa.gov/dashboard/dsstoxdb/results?search=DTXSID1020560#bioactivity-toxcast-models>

Batch search

EPA United States Environmental Protection Agency

Batch Search

Step 1 Step 2 Step 3 Step Five: Click "Download"

Please enter one identifier per line

Select Input Type(s)

Identifiers

Chemical Name ?

CASRN ?

InChIKey ?

DSSTox Substance ID ?

DSSTox Compound ID ?

InChIKey Skeleton ?

MS-Ready Formula(e) ?

Exact Formula(e) ?

Monoisotopic Mass ?

[Display All Chemicals](#) [Download Chemical Data](#)

Enter Identifiers to Search

104-55-2
78-70-6
103-90-2
107-02-8
115-29-7

Select Output Format:

Customize Results

Select All

Select All in Lists

Chemical Identifiers

DTXSID ?

Chemical Name ?

DTXCID ?

CAS-RN ?

InChIKey ?

IUPAC Name ?

Structures

Mol File ?

SMILES ?

InChI String ?

MS-Ready SMILES ?

QSAR-Ready SMILES ?

Intrinsic And Predicted Properties

Molecular Formula ?

Average Mass ?

Monoisotopic Mass ?

TEST Model Predictions ?

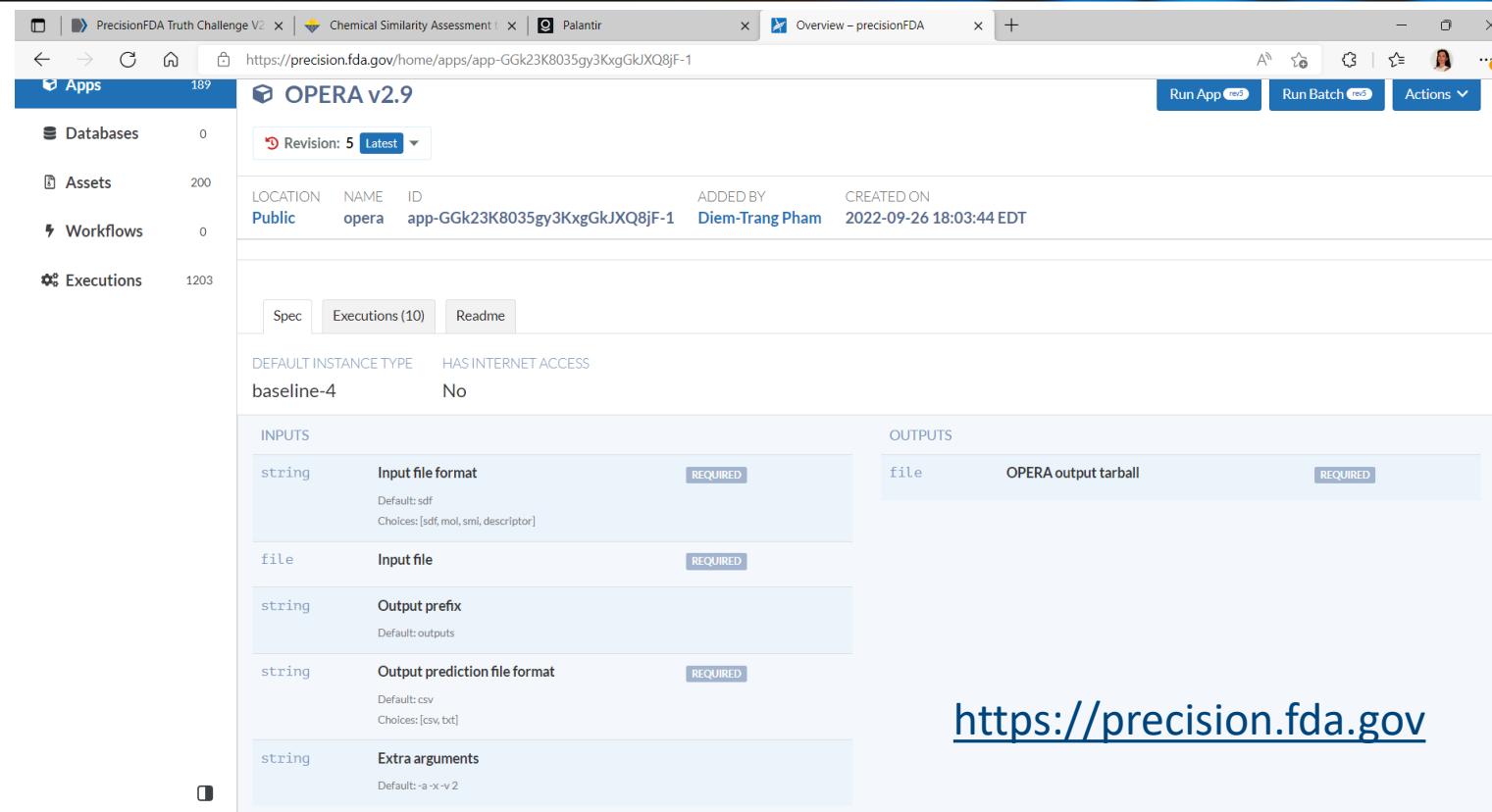
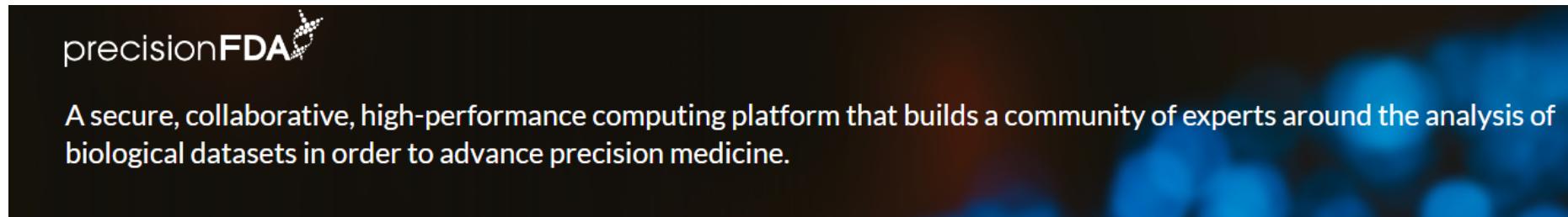
OPERA Model Predictions ?

Presence in Lists:

DRUGSNORMAN: Pharm
 40CFR355: Extremely Hazardous
 AERGLS: Acute Exposure
 Amphibole Minerals ?
 ANDROGEN: Androgen
 ARTICLE: Bench-Mark Discrepancy
 ARTICLE: Collaborative Efficacy
 ARTICLE: Collaborative Efficacy
 ATSDR: Minimal Risk Level
 ATSDR: Toxic Substances
 CAA/EPAs Office of Environ
 Canadian Domestic Substances
 Chemicals in human blood
 DRUGS: Antibiotics ?
 DRUGS: DrugBank database
 DRUGS: Opioids and related
 DRUGS: Statins drugs ?
 DRUGSNORMAN: ITNA
 DRUGSNORMAN: Pharmacokinetic
 DRUGSNORMAN: Target
 DRUGS/WIKILIST: Veterinary

https://comptox.epa.gov/dashboard/dsstoxdb/batch_search

Predictions on FDA's precision platform



The screenshot displays the precisionFDA platform interface. At the top, there is a navigation bar with tabs for "PrecisionFDA Truth Challenge V2", "Chemical Similarity Assessment", "Palantir", and "Overview – precisionFDA". Below the navigation bar, a sidebar on the left shows statistics for "Apps": 189 Databases, 200 Assets, 0 Workflows, and 1203 Executions. The main content area is titled "OPERA v2.9" and shows the following details:

- Revision: 5 (Latest)
- LOCATION: Public
- NAME: opera
- ID: app-GGk23K8035gy3KxgGkJXQ8jf-1
- ADDED BY: Diem-Trang Pham
- CREATED ON: 2022-09-26 18:03:44 EDT

Below this, there are tabs for "Spec", "Executions (10)", and "Readme". Further down, it shows "DEFAULT INSTANCE TYPE: baseline-4" and "HAS INTERNET ACCESS: No". The "INPUTS" section contains fields for "Input file format" (string, REQUIRED), "Input file" (file, REQUIRED), "Output prefix" (string), "Output prediction file format" (string, REQUIRED), and "Extra arguments" (string). The "OUTPUTS" section contains a field for "OPERA output tarball" (file, REQUIRED).

<https://precision.fda.gov>

OPERA in the OECD Toolbox

QSAR TOOLBOX

Gap Filling Workflow Editor

Trend analysis Read across (Q)SAR Automated Standardized New Import Export Delete

Documents Document 1 # [C: 1;Md: 0;P: 0] CAS: 58082

Structure info Parameters Physical Chemical Properties

- Autoflammability / Self-ignition temperature
- Boiling point
- Chemical reactivity
- Density
- Dissociation Constant (pKa)
- Explosive properties
- Flammability
- Flash point
- Melting / freezing point
- Oxidation reduction potential
- Oxidising properties
- Particle size
- Partition Coefficient:

 - Solubility in organic solvents / fat solubility
 - Stability in organic solvents and identity
 - Surface tension
 - Vapour pressure
 - Viscosity
 - Water solubility

- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazards

Data Gap Filling Settings

Only endpoint relevant

At this position:

QSARs	2
Automated workflows	0
Standardized workflows	0

Input Profiling Data Category definition Data Gap Filling Report

Filter endpoint tree... 1 [target]

Structure

Boiling Point Adapted Stein and Brown Method (EPISUITE) (1.0)

Opera BP (2.6)

Details for 2 (Q)SAR models

QSAR name	#	Predicted	Domain	OMRF	Test set	Training set
Boiling Point Adapted Stein and Brown Method (EPISUITE) (1.0)	1					
Opera BP (2.6)	1					

Toolbox Repository

Tools / QSARs / OPERA models

OPERA models



<https://repository.qsartoolbox.org/>

OPERA is a free and open-source/open-data suite of physicochemical properties, environmental fate, ADME and toxicological information including applicability domain and accuracy data and standardized QSAR-ready chemical structures. Operates on a user-friendly graphical interface for Windows and Linux operating systems.

Current version: 1.0

Supported Toolbox versions: 4.5

Developer: NIEHS

Category: QSARs

Downloads: 57

Rating:  0 

Description: OPERA is a free and open-source/open-data suite of physicochemical properties, environmental fate, ADME and toxicological information including applicability domain and accuracy data and standardized QSAR-ready chemical structures. Operates on a user-friendly graphical interface for Windows and Linux operating systems.

Acknowledgments

The NICEATM Group



US EPA
All international collaborators



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



Subscribe to NICEATM News email list
<https://ntp.niehs.nih.gov/go/niceatm>

Thank you for your attention!



Question



OR

Comment