



Collaborative Acute Toxicity Modeling Suite (CATMoS)

Kamel Mansouri

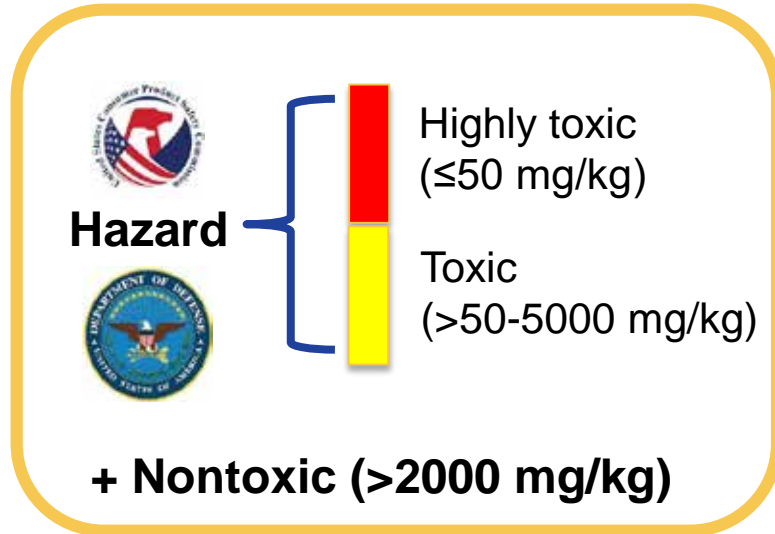
LEAD COMPUTATIONAL CHEMIST, ILS IN SUPPORT OF NICEATM

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(the author declares no conflict of interest)*



Agency-Based Modeling Endpoint Selection

Binary Models

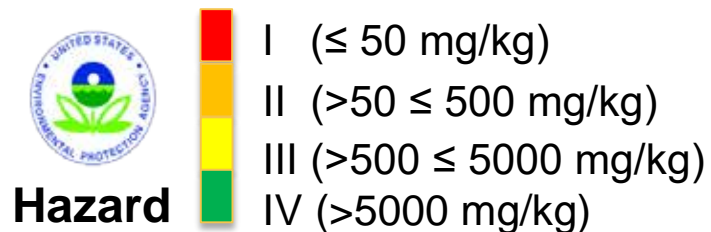


Continuous Model

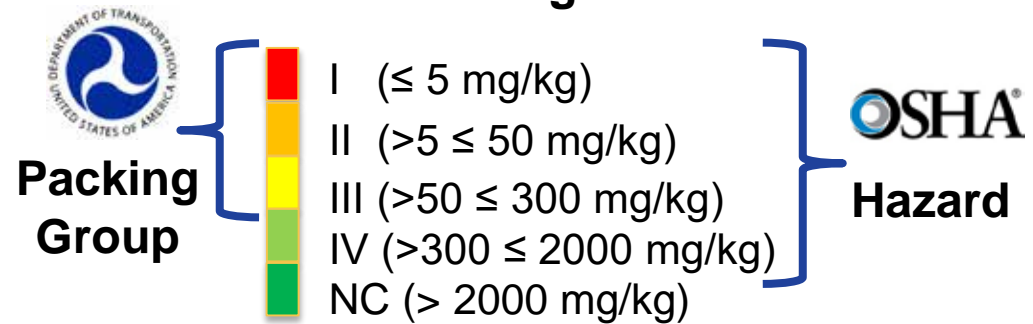


Categorical Models

EPA Categories



GHS Categories



Hazard



Available data for modeling

Rat oral LD50s:

16,297 chemicals total

34,508 LD50 values

15,688 chemicals total
21,200 LD50 values

QSAR-ready standardization

Desalted, stereochemistry stripped,
tautomers and nitro groups standardized,
valence corrected, structures neutralized

**11992 chemicals with
accurate structures**

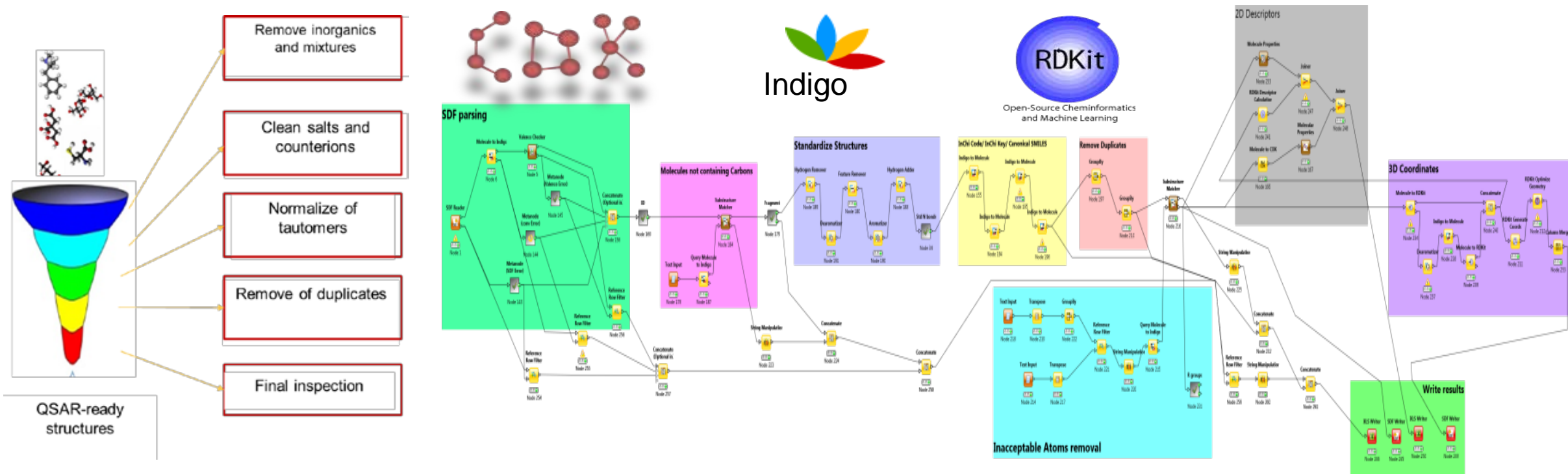
- Very toxic endpoint: 11886 entries (binary, 0/1)
- Non-toxic endpoint: 11871 entries (binary, 0/1)
- EPA endpoint: 11755 entries (categorical, 4 categories)
- GHS endpoint: 11845 entries (categorical, 5 categories)
- LD50 endpoint: 8908 entries (continuous values)



QSAR-ready KNIME workflow

Aim of the workflow:

- Combine different procedures and ideas
- Minimize the differences between the structures used for prediction
- Produce a flexible free and open source workflow to be shared



Fourches et al. J Chem Inf Model, 2010, 29, 476 – 488

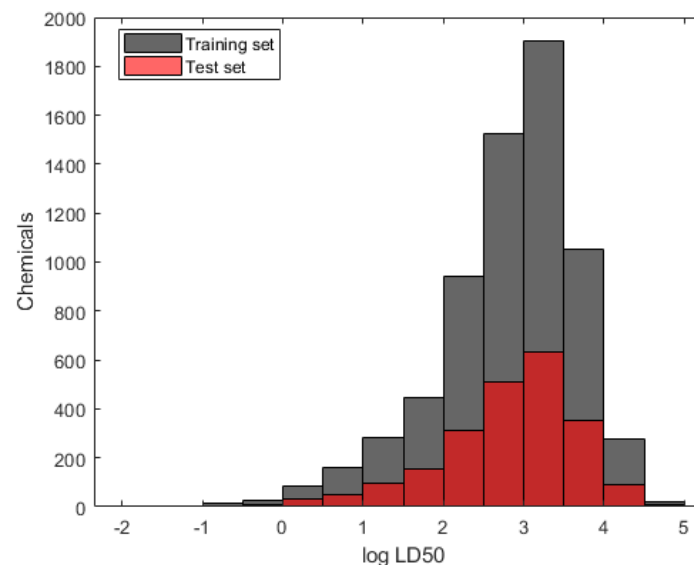
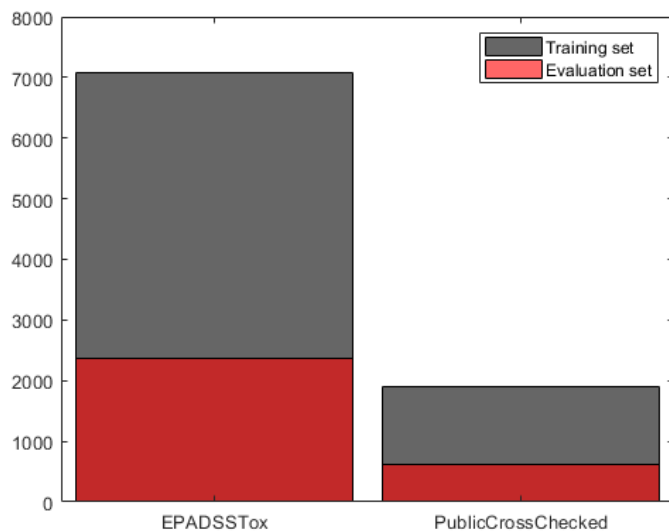
Wedebye et al. Danish EPA Environmental Project No. 1503, 2013

Mansouri et al. (<http://ehp.niehs.nih.gov/15-10267/>)



Establishing Modeling Dataset

- **Training and evaluation sets:**
 - 11,992 chemicals from the final inventory of chemicals with QSAR-ready structures having rat oral acute toxicity data were split into training and test sets:
 - 75% training set: 8,994 chemicals
 - 25% evaluation set: 2,998 chemicals
 - All endpoints training data included in same structure file
 - Similar distributions and variability for values and categories
 - Similar distribution of chemical structures sources





Establishing Modeling Dataset

- Prediction set:

Included lists of regulatory interest:

- ToxCast/Tox21
- EDSP
- TSCA
- Substances on the market (EPA Dashboard list)



After QSAR-ready standardization:

48137 structures to be predicted (including the evaluation set)



ChemMaps landscape of CATMoS chemicals

<http://www.chemmaps.com/chemmaps/DSSToxMap3D/>

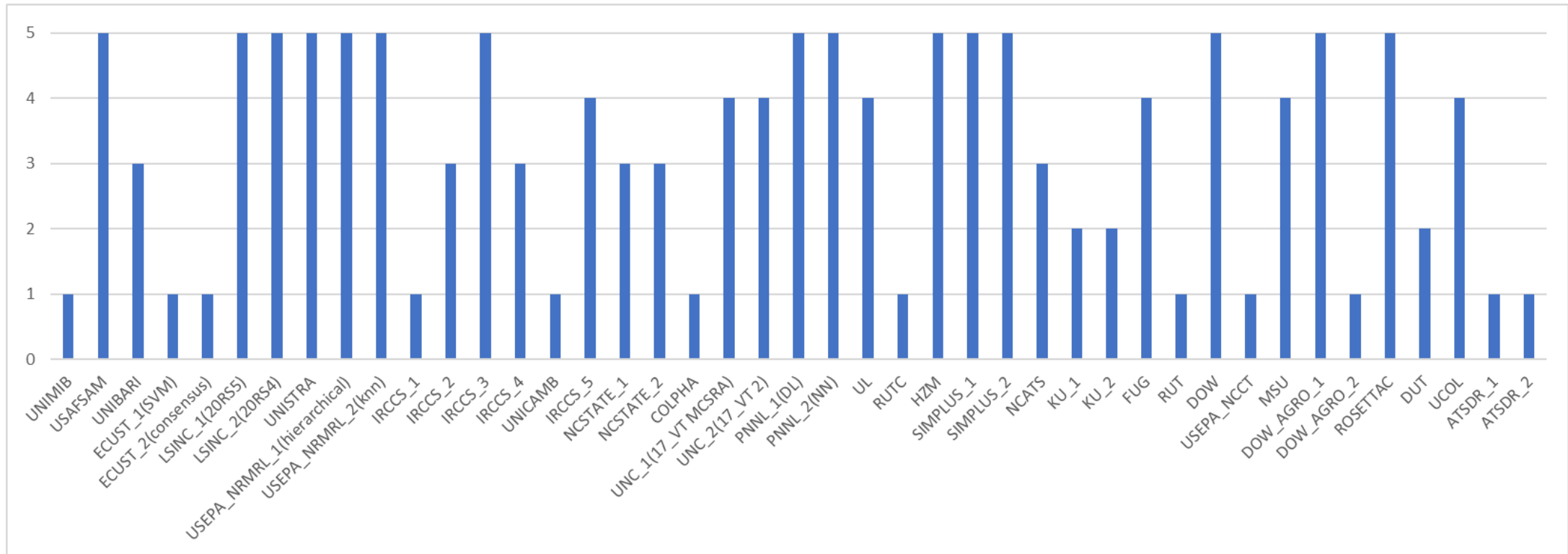




Submitted Models

- Non-toxic: 33 models
- Very Toxic: 32 models
- GHS categories: 23 models
- EPA categories: 26 models
- LD50: 25 models

Total: 139 models





Evaluation procedure

Qualitative evaluation:

- Documentation
- Defined endpoint
- Unambiguous algorithm
- Availability of code
- Applicability domain definition
- Availability of data used for modeling
- Mechanistic interpretation

Quantitative evaluation:

- Goodness of fit: training (Tr) statistics
- Predictivity: statistics on the evaluation set
- Robustness: balance between (Goodness of fit) & (Predictivity)

$$S = 0.3 * (\text{Goodness of fit}) + 0.45 * (\text{Predictivity}) + 0.25 * (\text{Robustness})$$

Categorical models (binary and multi-class):

$$\text{Goodness of fit} = 0.7 * (BA_{Tr}) + 0.3 * (1 - |Sn_{Tr} - Sp_{Tr}|)$$

$$\text{Predictivity} = 0.7 * (BA_{Eval}) + 0.3 * (1 - |Sn_{Eval} - Sp_{Eval}|)$$

$$\text{Robustness} = 1 - |BA_{Tr} - BA_{Eval}|$$

$$BA = \frac{(Sn + Sp)}{2}$$

$$Sn = \frac{TP}{TP + FN}$$

$$Sp = \frac{TN}{TN + FP}$$

Continuous models:

$$\text{Goodness of fit} = R_{Tr}^2$$

$$\text{Predictivity} = R_{Eval}^2$$

$$\text{Robustness} = 1 - |R_{Tr}^2 - R_{Eval}^2|$$

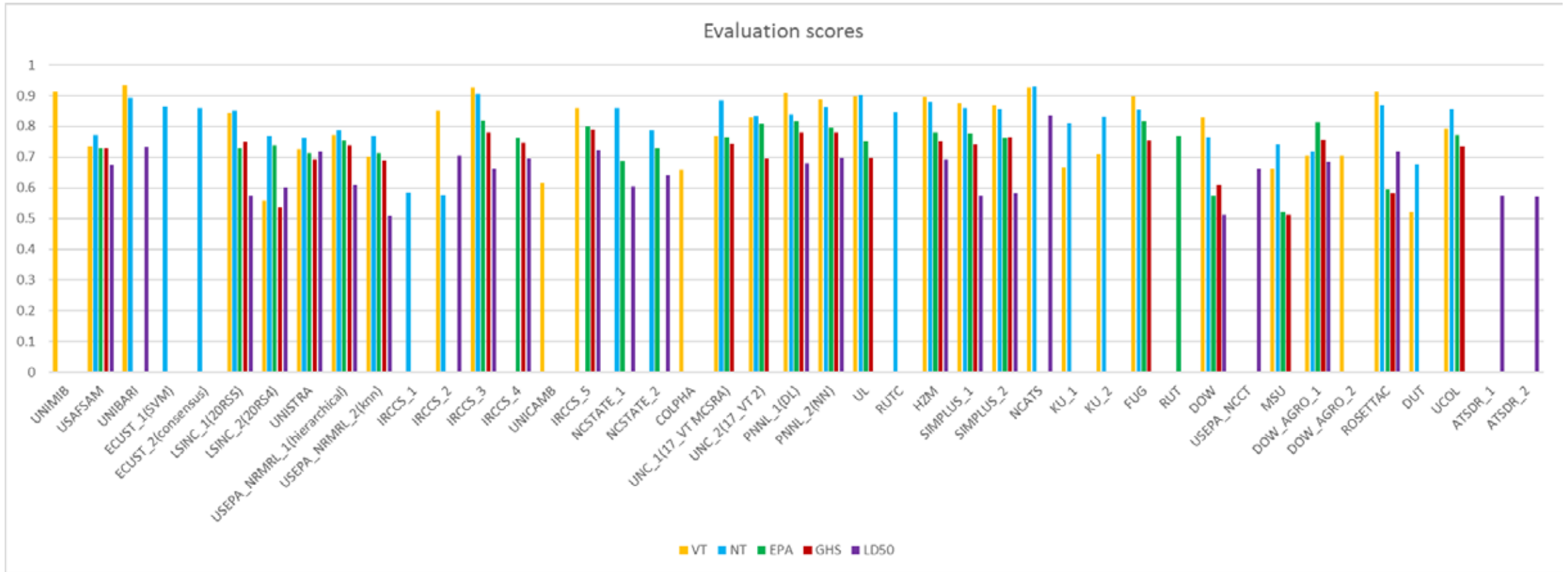
$$R^2 = 1 - \frac{\sum_{i=1}^{n_{TR}} (y_i - \hat{y}_i)^2}{\sum_{i=1}^{n_{TR}} (y_i - \bar{y})^2}$$

\hat{y}_i and y_i are the estimated and observed responses



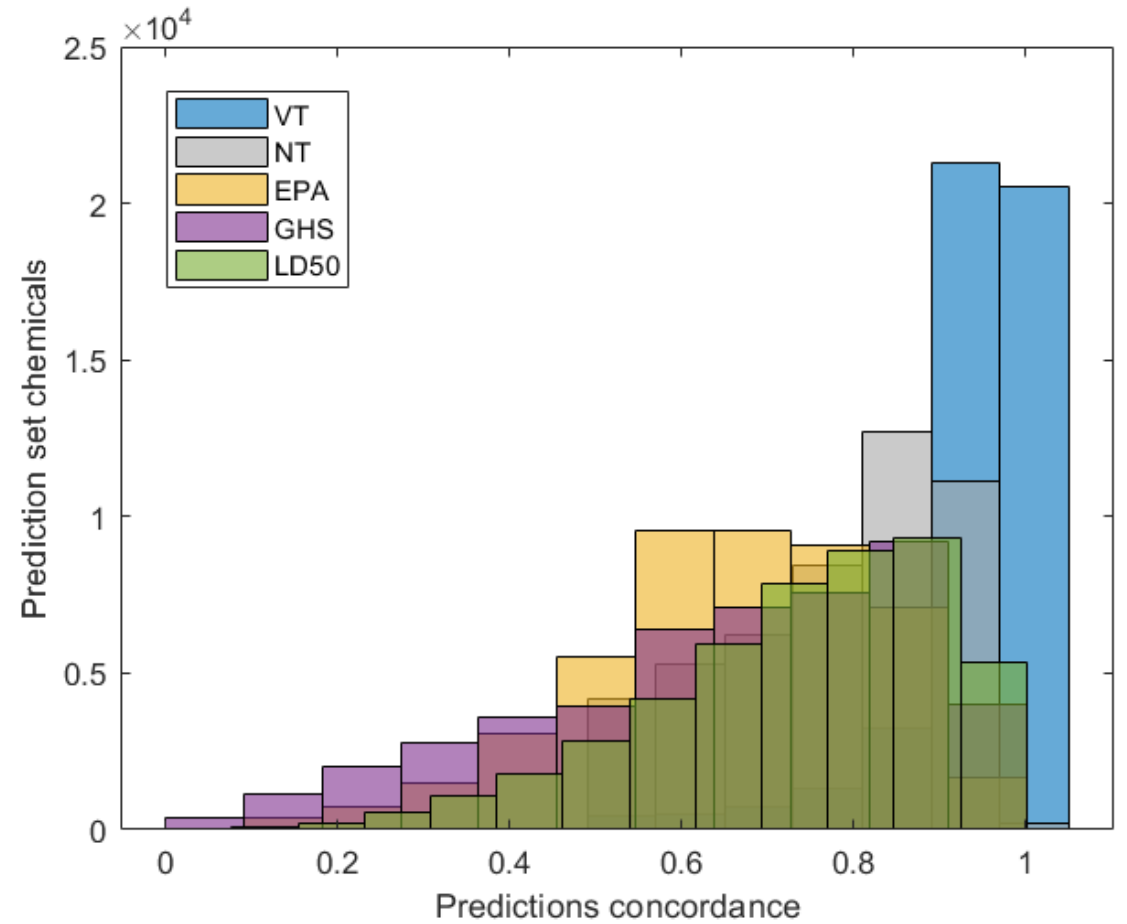
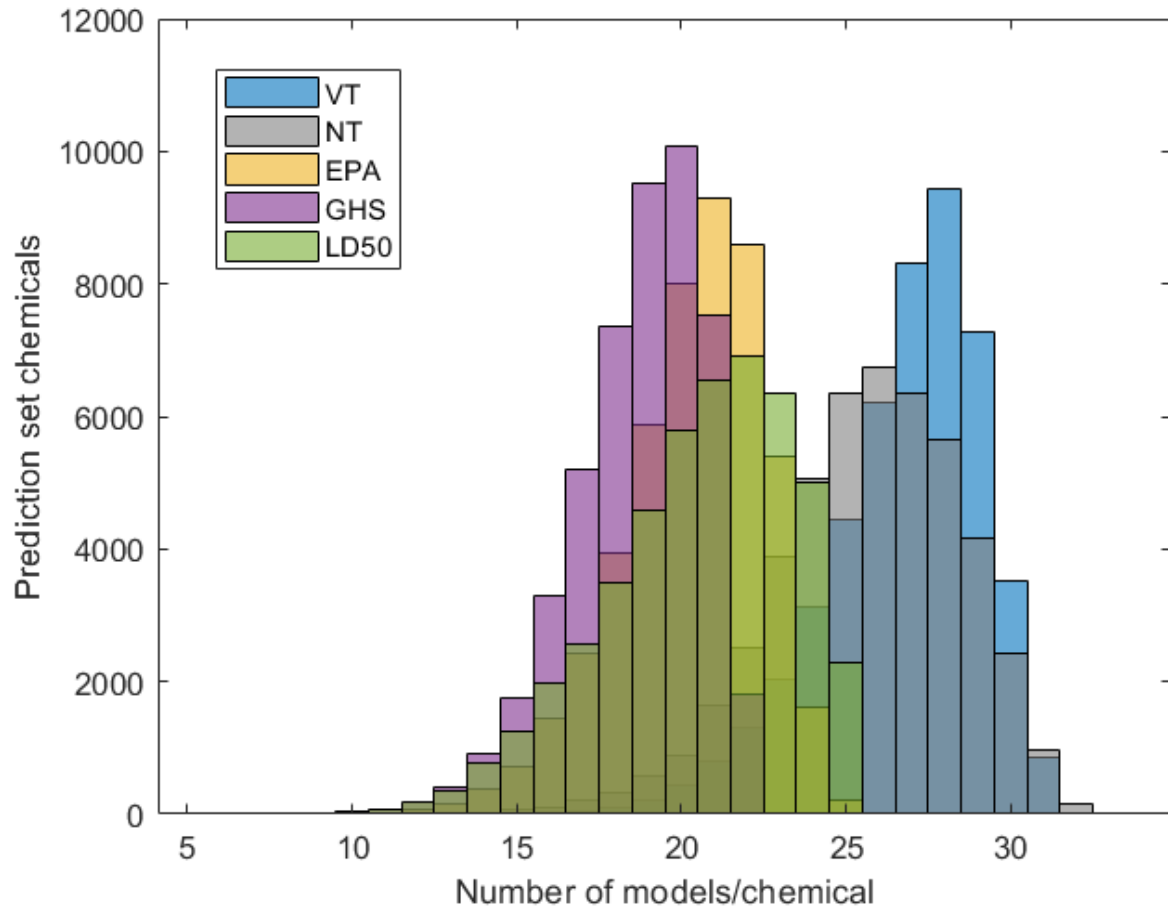
Evaluation results

Quantitative evaluation





Coverage and concordance of the models





CATMoS consensus modeling

Steps of combining the single models into consensus

Initial models & predictions

- VT (32 models)
- NT (33 models)
- GHS (23 models)
- EPA (26 models)
- LD50 (25 models)

Combining models

Step 1

Weighted average
/majority rule

Independent consensus models/predictions

- VT
- NT
- GHS
- EPA
- LD50

A consensus model
per endpoint
(~20-~30 models)

Weight of Evidence
approach (WoE)

Step 2

Majority rule

Consistent consensus models/predictions

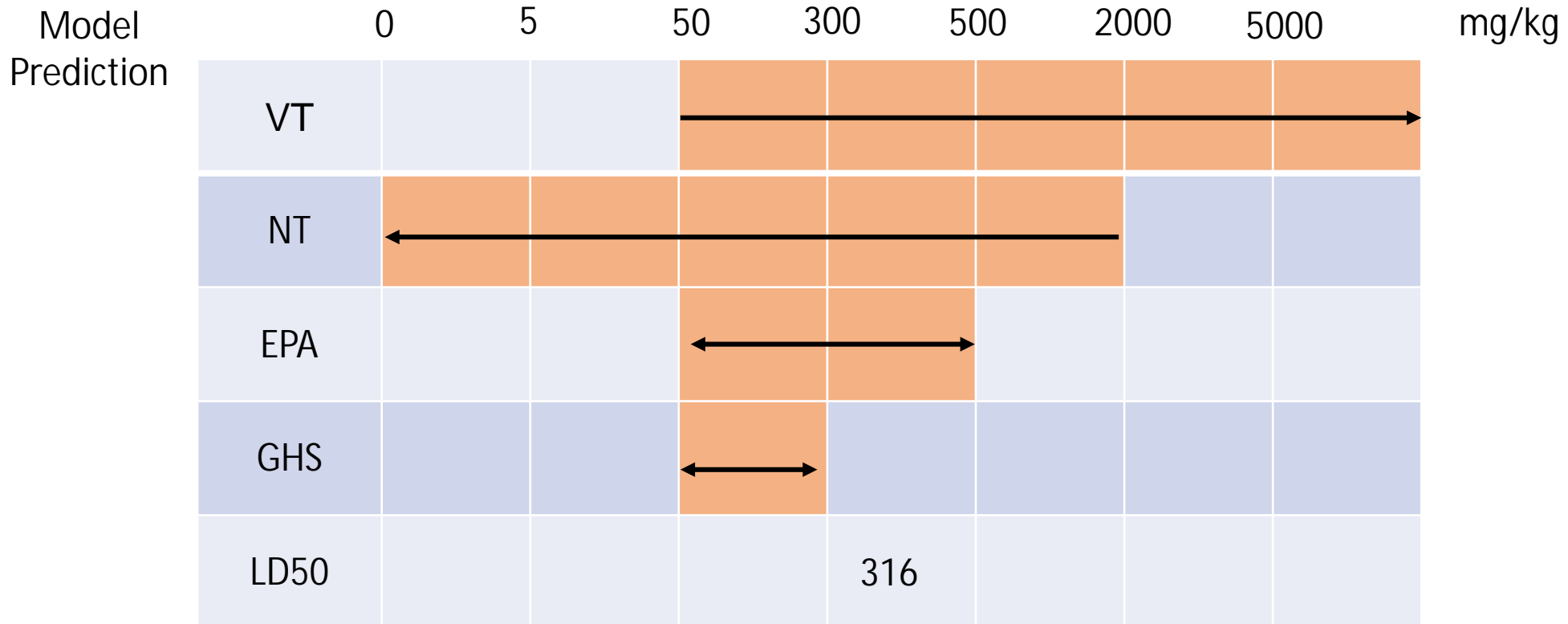
- VT
- NT
- GHS
- EPA
- LD50

Consensus
representing all
~140 models



WoE approach to combine the 5 endpoints

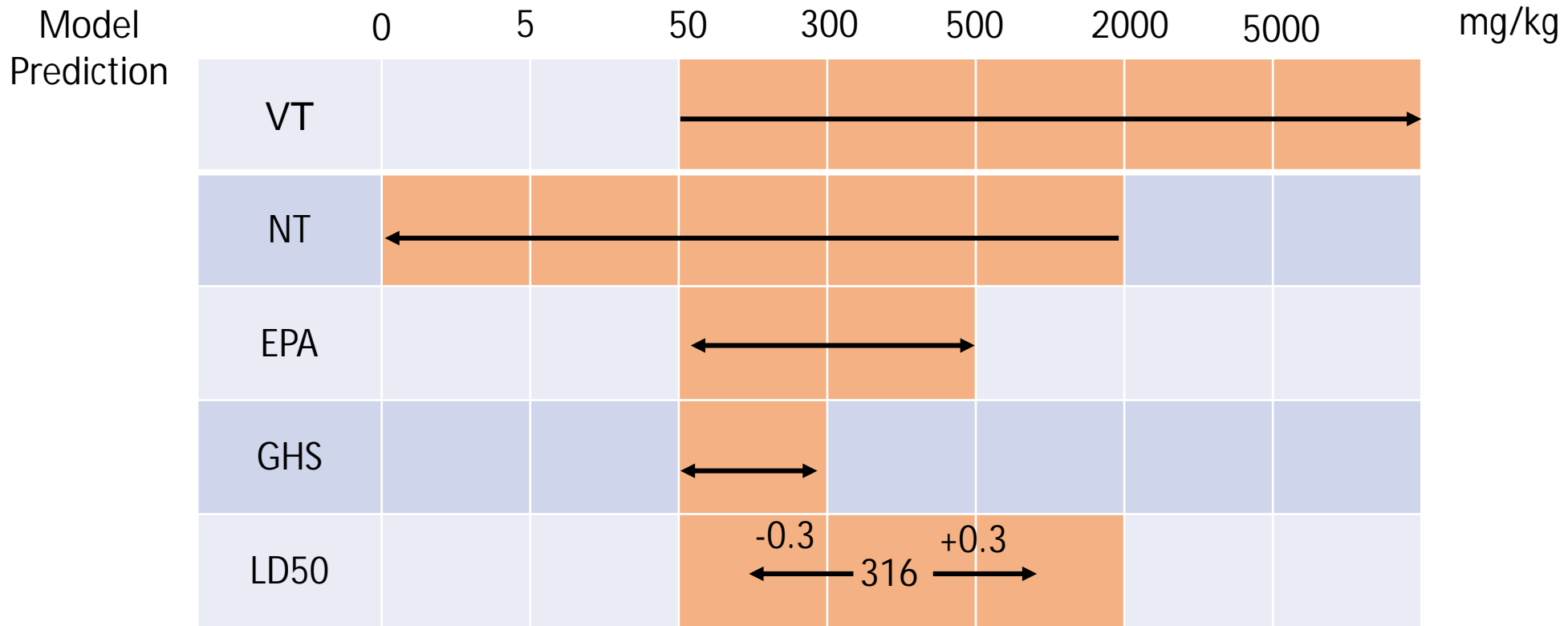
	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.5





WoE approach to combine the 5 endpoints

	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.5

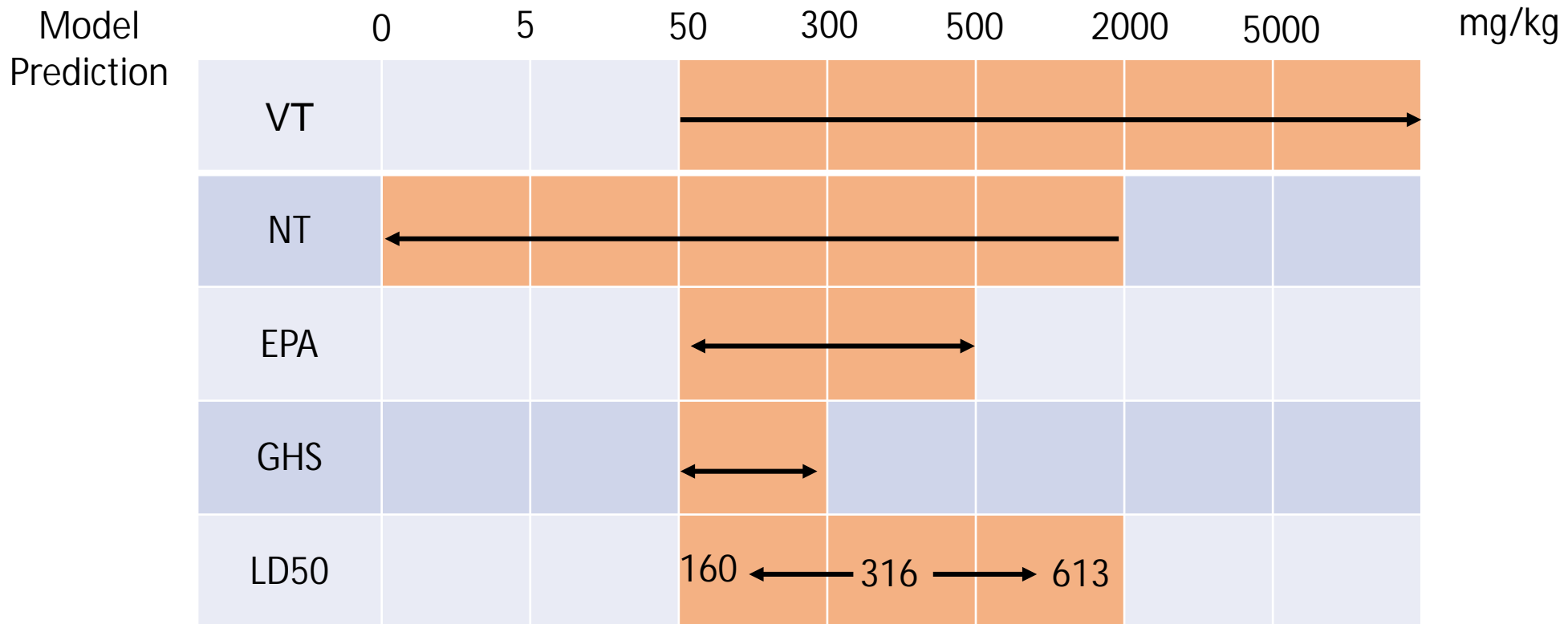


Variability range (log units) for LD50



WoE approach to combine the 5 endpoints

	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.5



Variability range (log units) for LD50



WoE approach to combine the 5 endpoints

	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.5

Model Prediction	0	5	50	300	500	2000	5000	mg/kg
VT	0	0	1	1	1	1	1	
NT	1	1	1	1	1	0	0	
EPA	0	0	1	1		0	0	
GHS	0	0	1	0	0	0	0	
LD50	0	0	160 1	1	613 1			
WoE	1	1	5	4	3	1	1	



WoE approach to combine the 5 endpoints

Original: independent calls

	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.5



WoE: consistent calls

	VT	NT	EPA	GHS	LD50
molX	0	0	2	3	2.36

Model Prediction	Winning bin							mg/kg
	0	5	50	300	500	2000	5000	
VT	0	0	1	1	1	1	1	
NT	1	1	1	1	1	0	0	
EPA	0	0	1	1		0	0	
GHS	0	0	1	0	0	0	0	
<u>LD50?</u>	0	0	1	1	1			
WoE	1	1	5	4	3	1	1	

How to adjust quantitative LD50?
Avg of Lower CI and upper bin threshold



$$(160+300)/2 = 230\text{mg/kg}$$





Performance Assessment

Consensus Model Statistics

	Very Toxic		Non-Toxic		EPA		GHS	
	Train	Eval	Train	Eval	Train	Eval	Train	Eval
Sensitivity	0.87	0.70	0.88	0.67	0.81	0.62	0.80	0.58
Specificity	0.99	0.97	0.97	0.90	0.92	0.86	0.95	0.90
Balanced Accuracy	0.93	0.84	0.92	0.78	0.87	0.74	0.88	0.74
<i>In vivo</i> Balanced Accuracy	0.81		0.89		0.82		0.79	

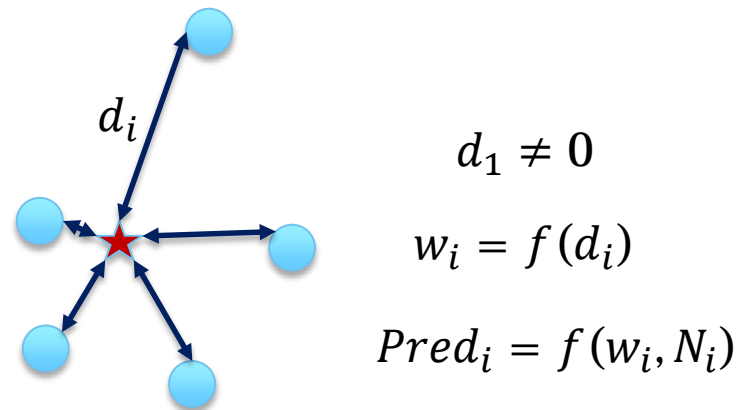
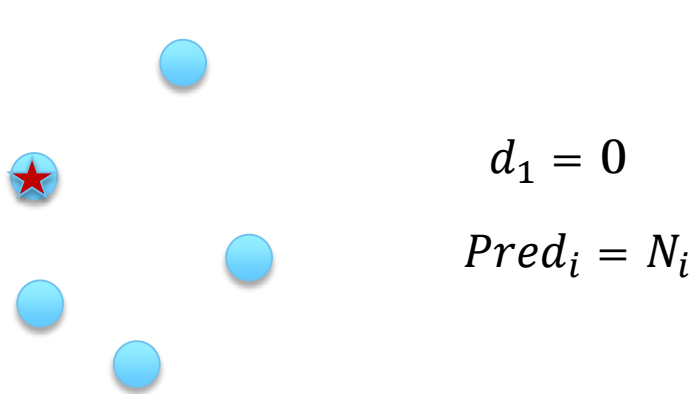
	LD50 values		LD50 values
	Train	Eval	<i>In Vivo</i>
R2	0.85	0.65	0.80
RMSE	0.30	0.49	0.42

The consensus predictions perform just as well as replicate *in vivo* data do at predicting oral acute toxicity outcome



Extended CATMoS predictions

Weighted read-across



★ New chemical to be predicted

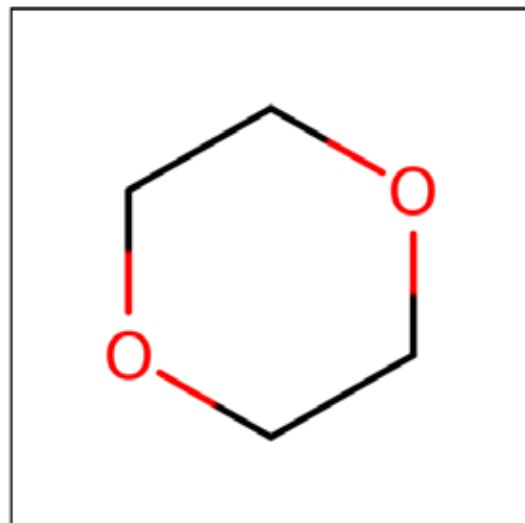
● Nearest neighbors (N_i)

d_i : Euclidean distance based on the selected descriptors for each endpoint

➔ Automated, similarity-endpoint dependent read-across: weighted kNN



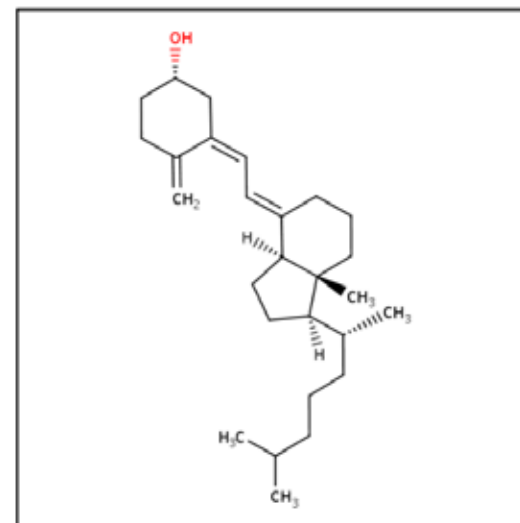
CATMoS prediction examples



1,4-Dioxane
123-91-1 | DTXSID4020533
Molecular Formula: C₄H₈O₂
Average Mass: 88.106 g/mol

LD50: 4200 mg/kg
log₁₀ LD50= 3.62

<https://comptox.epa.gov/dashboard/dsstoxdb/results?search=DTXSID4020533>



Vitamin D3
67-97-0 | DTXSID6026294
Molecular Formula: C₂₇H₄₄O
Average Mass: 384.648 g/mol

LD50: 42 mg/kg
log₁₀ LD50= 1.62

<https://comptox.epa.gov/dashboard/dsstoxdb/results?search=DTXSID6026294>

CATMoS predictions:

MoleculeID	CATMoS_VT_pred	CATMoS_NT_pred	CATMoS_EPA_pred	CATMoS_GHS_pred	CATMoS_LD50_pred	AD_CATMoS	AD_index_CATMoS	Conf_index_CATMoS
'123-91-1'	0	1	3	5	3.4053	1	1	0.9500
'67-97-0'	1	0	1	2	1.2845	1	1	0.8684



Collaboration with ATWG partners and ICCVAM agencies

Agency	No. Substances	Agency	No. Substances
Air Force	421	EPA OPP	36
Army Public Health Command	18	EPA OPPT	8
Army Edgewood Chemical Biological Center	42	EPA NCCT	4815
CPSC	110	FDA CFSAN	22
DOT	3671		

Evaluate and optimize CATMoS predictions based on lists of interest



Running CATMoS Consensus models

OPERA Standalone application

```
OPERA_CL
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OPERA models for physchem, environmental fate and tox properties.
Version 2.3 (June 2019)

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

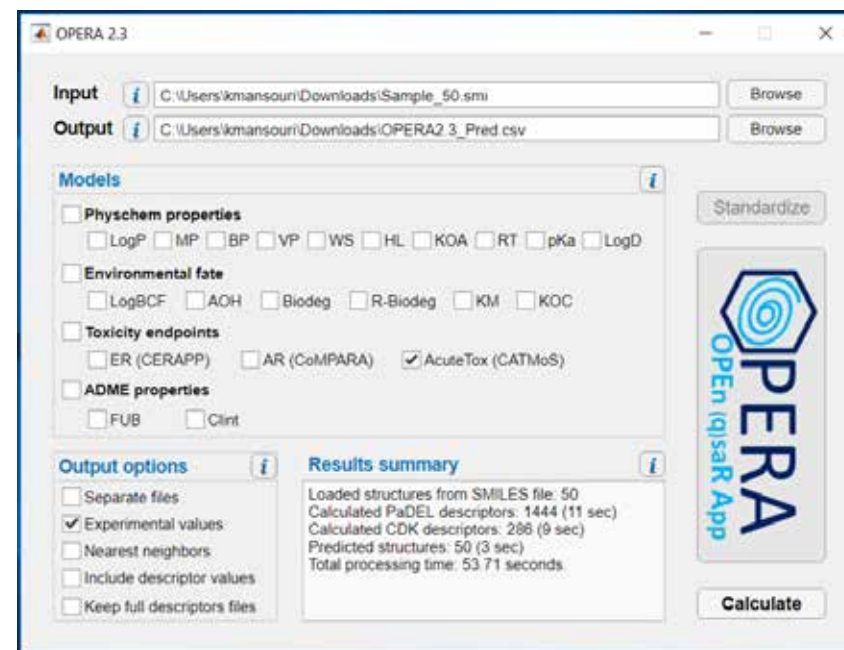
Developed by:
Kamel Mansouri
mansourikamel@gmail.com

Usage: OPERA <argument_list>

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

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

Command line



Graphical user interface

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

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

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



Since OPERA v1.5

Physchem & Environmental fate:

Model	Property
AOH	Atmospheric Hydroxylation Rate
BCF	Bioconcentration Factor
BioHL	Biodegradation Half-life
RB	Ready Biodegradability
BP	Boiling Point
HL	Henry's Law Constant
KM	Fish Biotransformation Half-life
KOA	Octanol/Air Partition Coefficient
LogP	Octanol-water Partition Coefficient
MP	Melting Point
KOC	Soil Adsorption Coefficient
VP	Vapor Pressure
WS	Water solubility
RT	HPLC retention time

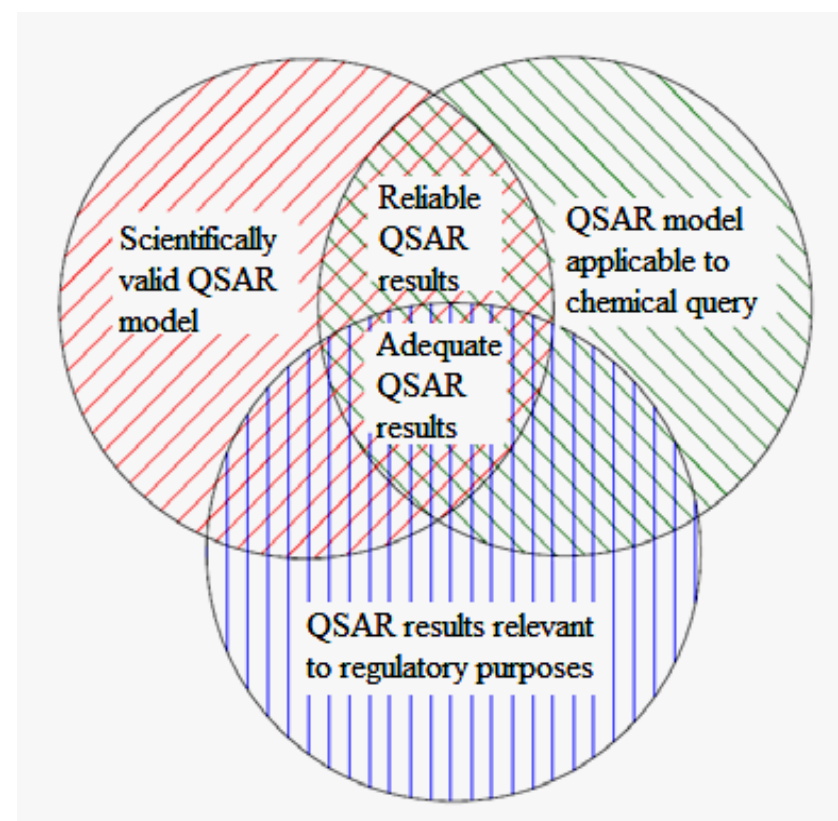
New since OPERA 2.0

- Physchem properties:
 - General structural properties
 - pKa
 - Log D
- ADME properties
 - Plasma fraction unbound (FuB)
 - Intrinsic clearance (Clint)
- Toxicity endpoints
 - ER activity (CERAPP)
<https://ehp.niehs.nih.gov/15-10267/>
 - AR activity (CoMPARA)
<https://doi.org/10.13140/RG.2.2.19612.80009>
 - **Acute toxicity (CATMoS)**
<https://doi.org/10.1016/j.comtox.2018.08.002>



Inform Regulatory Decisions

Principle	Description
1) A defined endpoint	Any physicochemical, biological or environmental effect that can be measured and therefore modelled.
2) An unambiguous algorithm	Ensure transparency in the description of the model algorithm.
3) A defined domain of applicability	Define limitations in terms of the types of chemical structures , physicochemical properties and mechanisms of action for which the models can generate reliable predictions .
4) Appropriate measures of goodness-of-fit, robustness and predictivity	a) The internal fitting performance of a model b) the predictivity of a model, determined by using an appropriate external test set .
5) Mechanistic interpretation, if possible	Mechanistic associations between the descriptors used in a model and the endpoint being predicted .





Soon on NTP/ICE and EPA CompTox dashboard

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

National Toxicology Program
U.S. Department of Health and Human Services

Integrated Chemical Environment

Chemicals

Input

Results

Assay	Description	Assay Type
NHK NRU	Acute Oral Toxicity	in vitro
3T3 NRU	Acute Oral Toxicity	in vitro

Search EAD by chemical: Select assay data to display:

EAD 95th Acute Oral Toxicity Log Axis

EAD 95th Box and Whisker

Search $C_{50} = \frac{Dose}{CL_r + CL_b}$ Machine Learning Chemical Characterization

<https://comptox.epa.gov/dashboard>

EPA United States Environmental Protection Agency

Chemistry Dashboard

OPERA Models: LogP: Octanol-Water

Bisphenol A
80-05-7 | DTXSID07020182

Prediction

Model Results

Predicted value: 3.33

Global applicability domain:

Local applicability domain:

Confidence level: 3.75

Accuracy & AD assessment

Performance

Weighted KNN model

QMRFS

Fold CV (75%)		Training (75%)		Test (25%)	
Q2	RMSE	R2	RMSE	R2	RMSE
0.00	0.00	0.00	0.07	0.00	0.70

Nearest Neighbors

Chemical	Measured	Predicted
Bisphenol A	3.33	3.33
BUTYRIC ACID 2-(4-METHYLBENZOYL)-	4.25	3.45
Phthalic acid	4.15	3.83
2,2-Dimethylpropanoic acid	3.69	2.93
3-(4-HYDROXYBENZYL)PENTANOIC ACID	3.75	3.18

Discover: About/Disclaimer, Accessibility

Connect: ACToR, USTox

Ask: Contact, Help

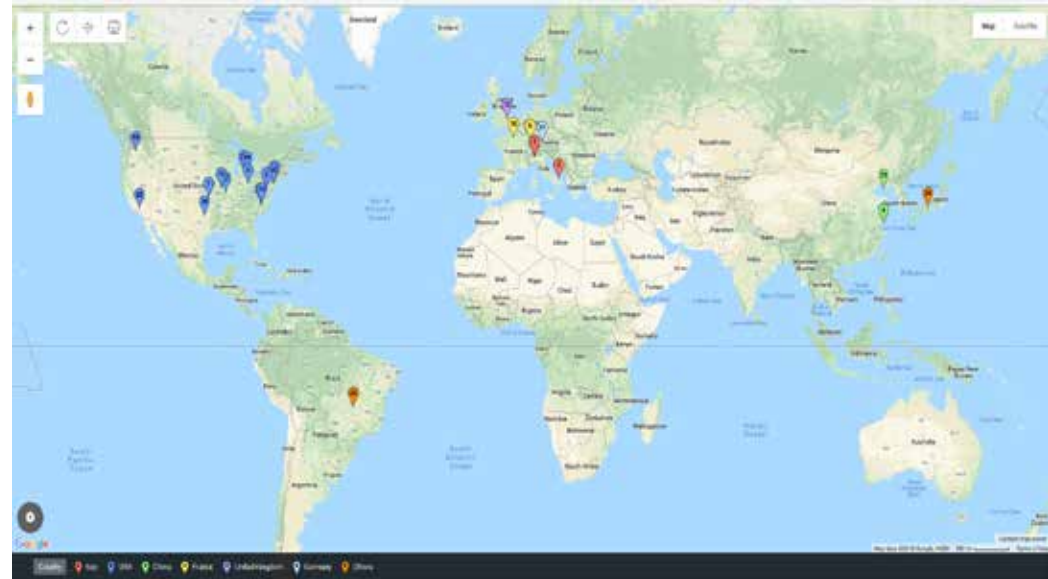


Acknowledgements

THANK YOU!

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 - Nicole Kleinstreuer
 - Warren Casey

All CATMoS international collaborators



Feedback welcome: [Kamel Mansouri \(kmansouri@ils-inc.com\)](mailto:kmansouri@ils-inc.com)

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