

Best practices to develop artificial intelligence models for predicting multilevel effects in Adverse Outcome Pathways (AOP)

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Outline

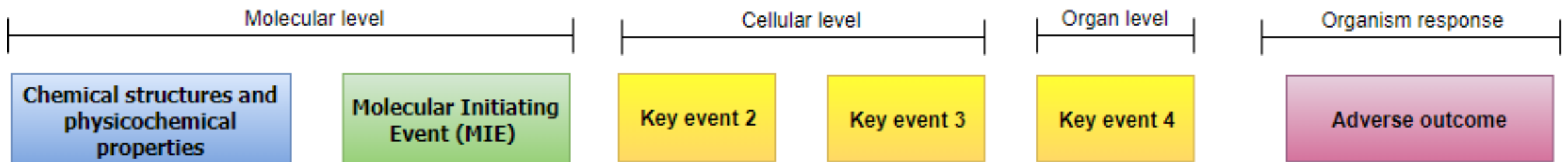
- The “AOP-based *in silico* model” overview
- Transparency vs mechanistic interpretability vs predictivity
- AOP-based *in silico* model – Framework
 - How to ensure accuracy and mechanistic interpretability?
 - Chemical Representation / Description
 - STR continuous x categorical
 - Prediction of MIE and KEs
 - Model validation and development
 - Chemical Space and Coverage
- Final remarks

Adverse outcome pathways (AOP) framework

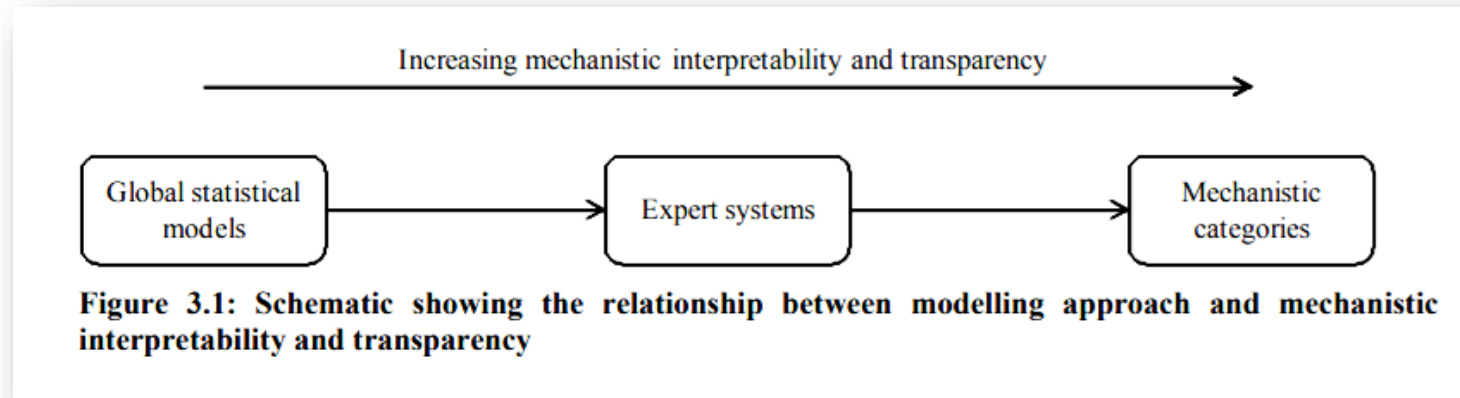
An AOP is an analytical construct that describes a sequential chain of causally linked events at different levels of biological organization that lead to an adverse health or ecotoxicological effect (OECD, 2012).

A proposed “AOP-based in silico model” Concept (Alttox)

An *in silico* framework used to identify chemicals that can **activate** the associated modular **AOP components** (MIE/KE) and based in these individual multilevel predictions, balanced by adjustments, relationships and weights, **to predict an adverse outcome**.



Transparency vs mechanistic interpretability vs predictivity



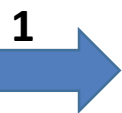
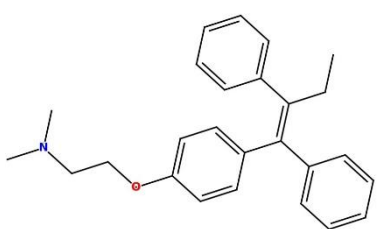
Frequently discussed assumptions:

- Global statistical models
- ↑ Accuracy
- ↓ Transparency
- ↓ Mechanistic interpretability
- Alert-based models
- ↓ Accuracy
- ↑ Transparency
- ↑ Mechanistic interpretability

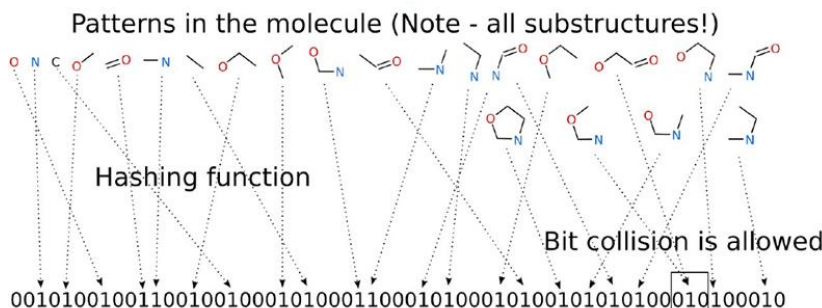
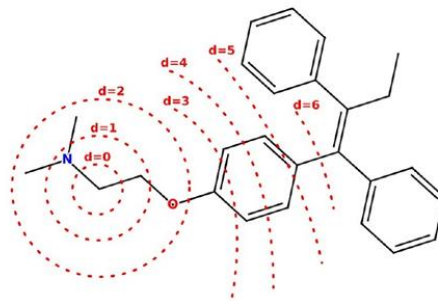
How to ensure accuracy and mechanistic interpretability?

chemical representation

Input Molecule



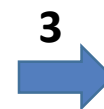
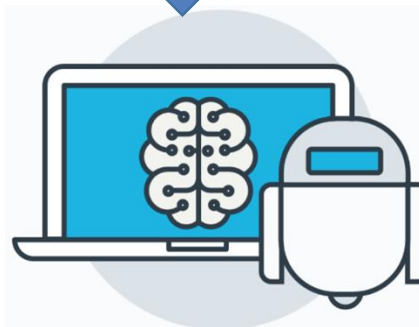
Molecular Pattern



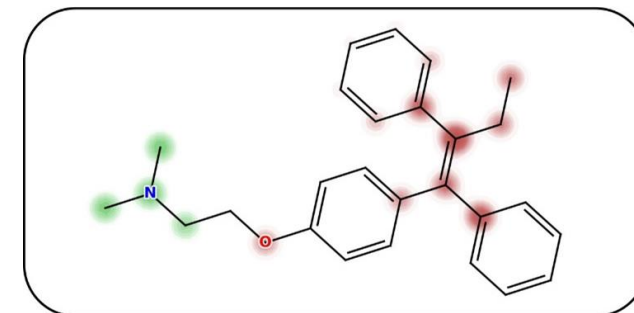
Hybrid Descriptor



Artificial intelligence model



STR Probability Mapping



PhysicalChemical Properties

Molecular weight (MW)
TPSA
(topological polar surface area)

Permeability

+

logK_{ow} (neutral cmpds)
logD (ionizable cmpds)


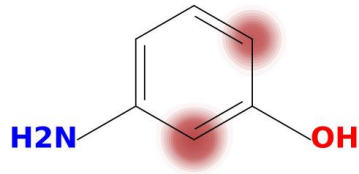
Lipophilicity

Transparency vs mechanistic interpretability vs predictivity


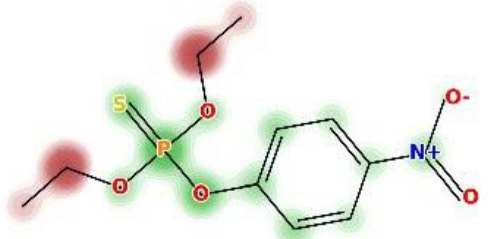
Structure-toxicity relationship (STR)

“the fragments more related to the absence/decrease of toxicity (green) or presence/increase (red)”

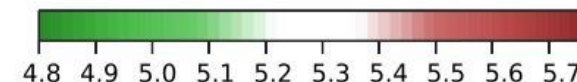
1 Categorical

Predicted Outcome/Assay	Predicted class (Confidence)	STR Contribution Mapping
 <p>Local Lymph Node Assay (LLNA, OECD 429)</p> <p>Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK_{ow}, logD)</p>	<p>Sensitizer (+) (78.4%)</p>	

2 Continuous

Predicted endpoint/Method	Predicted Value (Confidence)	STR Contribution Mapping
 <p>LC₅₀ (Fish, 96hrs)</p> <p>Deep Learning decision model implemented with hybrid descriptors</p>	<p>1.4 mg/L 4.7 μM (87.0%)</p>	 <p>parathion</p>

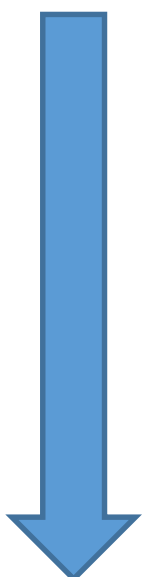
Overall Contribution = 5.33

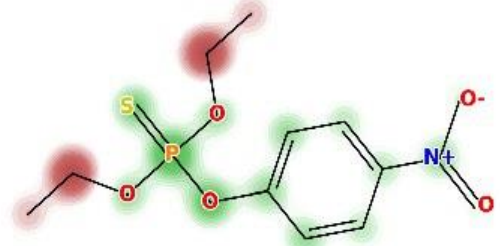
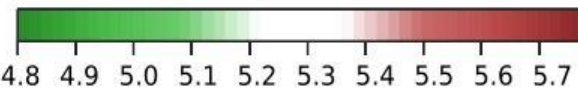
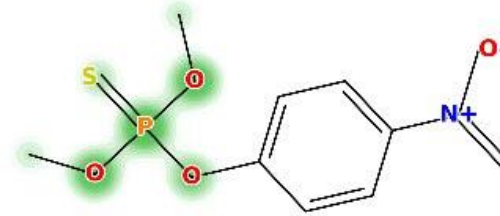
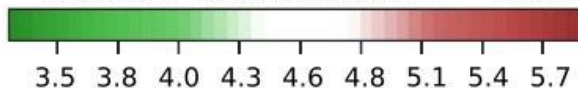


Structure-toxicity relationship (STR)



Reduce the Toxicity



Predicted endpoint/Method	Predicted Value (Confidence)	STR Contribution Mapping
<p>1</p> <p>LC₅₀ (Fish, 96hrs)</p> <p>Deep Learning decision model implemented with hybrid descriptors</p>	<p>1.4 mg/L 4.7 μM (87.0%)</p>	<p>parathion</p>  <p>Overall Contribution = 5.33</p> 
<p>2</p> <p>LC₅₀ (Fish, 96hrs)</p> <p>Deep Learning decision model implemented with hybrid descriptors</p>	<p>6.0 mg/L 22.6 μM (88.0%)</p>	<p>parathion methyl</p>  <p>Overall Contribution = 4.65</p> 

How to ensure accuracy and mechanistic interpretability?

Detecting mitigating factors (steric, electronic, and detoxifying) by statistical models with a visual probability mapping

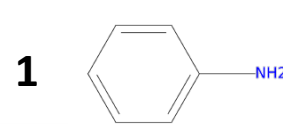
Theory

Mitigating factor	Example inactive chemical
Steric	
Electronic	
Detoxifying	

mitigating factors for aromatic amines (mitigating factor in red)

OECD - Organisation for Economic Co-operation and Development. Report of the expert consultation on scientific and regulatory evaluation of organic chemistry mechanism-based structural alerts for the Identification of DNA binding chemicals. Series on Testing and Assessment, No. 120 PART 1, 2010. Available in: <http://www.oecd.org/env/ehs/risk-assessment/45401393.pdf>

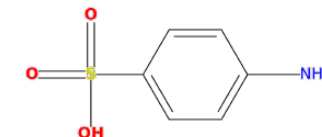
Alert model
Statistical models with probability map



Examples:

Genotox-iS™

2



Result: (+) Positive

Alerts were found in the molecule. The results are in the table below and a description is provided at the end of the report.

Category	Alert	Alert ID	References
in vitro mutagenicity (Ames alert) alerts by ISS		Primary aromatic amine, hydroxyl amine and its derived esters	Benigni, R., Giuliani, A., Franke, R., and Gruska, A. (2000). Quantitative structure-activity relationships of mutagenic and carcinogenic aromatic amines. <i>Chem.Revs.</i> 100, 3697-3714. Woo, Y. T. and Lai, D. Y. (2001). Aromatic amino and nitro-amino compounds and their halogenated derivatives. In 'Patty's Toxicology, Vol. 4.' (Eds E. Bingham, B. Cohnsen, and C. H. Powell.) pp. 969-1105. (John Wiley and Sons, Inc: New York.)

Probability Map

Non-Mutagen Mutagen

Method	Prediction (Confidence)	Probability Mapping (SAR)
Random Forest Machine learning decision model implemented with the 2D MACCS fingerprint	Non-Mutagen (69.3%)	
kNN k-nearest neighbors decision model implemented with the 2D Extended Connectivity Fingerprint	Mutagen (71.4%)	
Deep Learning 3D Deep Learning decision model implemented with the 3D conformer fingerprint like Extended Connectivity Fingerprint	Mutagen (86.8%)	

Result: (+) Positive

Alerts were found in the molecule. The results are in the table below and a description is provided at the end of the report.

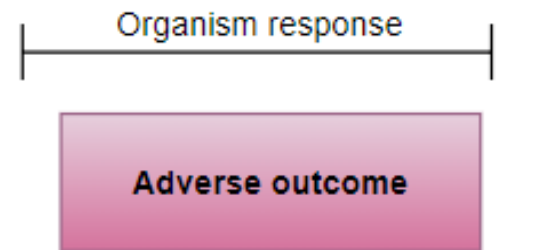
Category	Alert	Alert ID	References
in vitro mutagenicity (Ames alert) alerts by ISS		Primary aromatic amine, hydroxyl amine and its derived esters	Benigni, R., Giuliani, A., Franke, R., and Gruska, A. (2000). Quantitative structure-activity relationships of mutagenic and carcinogenic aromatic amines. <i>Chem.Revs.</i> 100, 3697-3714. Woo, Y. T. and Lai, D. Y. (2001). Aromatic amino and nitro-amino compounds and their halogenated derivatives. In 'Patty's Toxicology, Vol. 4.' (Eds E. Bingham, B. Cohnsen, and C. H. Powell.) pp. 969-1105. (John Wiley and Sons, Inc: New York.)

Probability Map

Non-Mutagen Mutagen

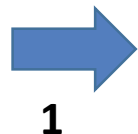
Method	Prediction (Confidence)	Probability Mapping (SAR)
Random Forest Machine learning decision model implemented with the 2D MACCS fingerprint	Non-Mutagen (98.2%)	
kNN k-nearest neighbors decision model implemented with the 2D Extended Connectivity Fingerprint	Non-Mutagen (85.7%)	
Deep Learning 3D Deep Learning decision model implemented with the 3D conformer fingerprint like Extended Connectivity Fingerprint	Non-Mutagen (95.0%)	

AOP-based *in silico* model



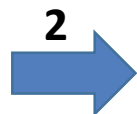
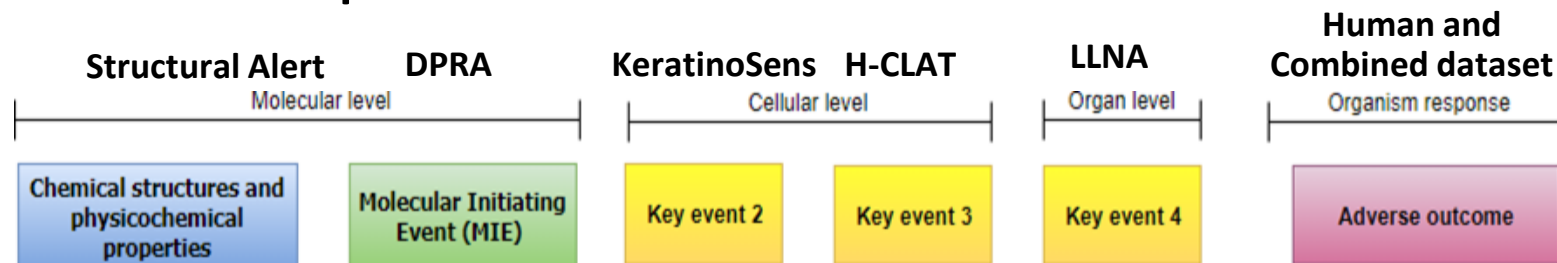
Global AOP-based *in silico* model balanced by KERs, adjustments, weight of evidence assessment in a logical framework

- To assess the results and predictivity measures in an **evidence assessment**.
- To integrate these results with *in chemico* and *in vitro* assessments, advancing in predictivity measures of an **IATA framework** for regulatory purposes.



Develop and Validate individual models

OECD Principles



key event relationships (KERs) integrating *in chemico*, *in vitro*, *ex vivo* and *in vivo* data in the *in silico* models

integrating all multilevel predictions balancing predictivity, key events relationships and WoE adjustments, and to predict an adverse outcome

Benchmark

Key event “Models” X
AOP X

Combined dataset (GMTP, LLNA and human data)
6971 Chemicals

$$\text{AOP-based algorithm} = \sum_{j=1}^n p_j c_j a_j w_j$$

Benchmark - Combined dataset (GMTP, LLNA and human data)

Model	Dataset	Specificity	Sensitivity	Accuracy	Coverage
Alert analysis	197 128: 1A 69: 1B	0.57	0.59	0.58	100%
DPRA	195	0.76	0.32	0.54	100%
DPRA AD		0.73	0.39	0.56	62%
KeratinoSens	190	0.74	0.27	0.51	100%
KeratinoSens AD		0.78	0.27	0.52	63%
h-CLAT	161	0.66	0.40	0.53	100%
h-CLAT AD		0.65	0.41	0.53	65%
LLNA	997	0.43	0.68	0.56	100%
LLNA AD		0.46	0.68	0.57	77%
Human Skin	389	0.83	0.33	0.57	100%
Human Skin AD		0.82	0.35	0.59	77%
Combined dataset (GMTP, LLNA and human data)*	6971	0.75	0.92	0.84	100%
Combined dataset (GMTP, LLNA and human data) AD*		0.75	0.92	0.84	93%
AOP-based prediction	-	0.76	0.71	0.74	100%

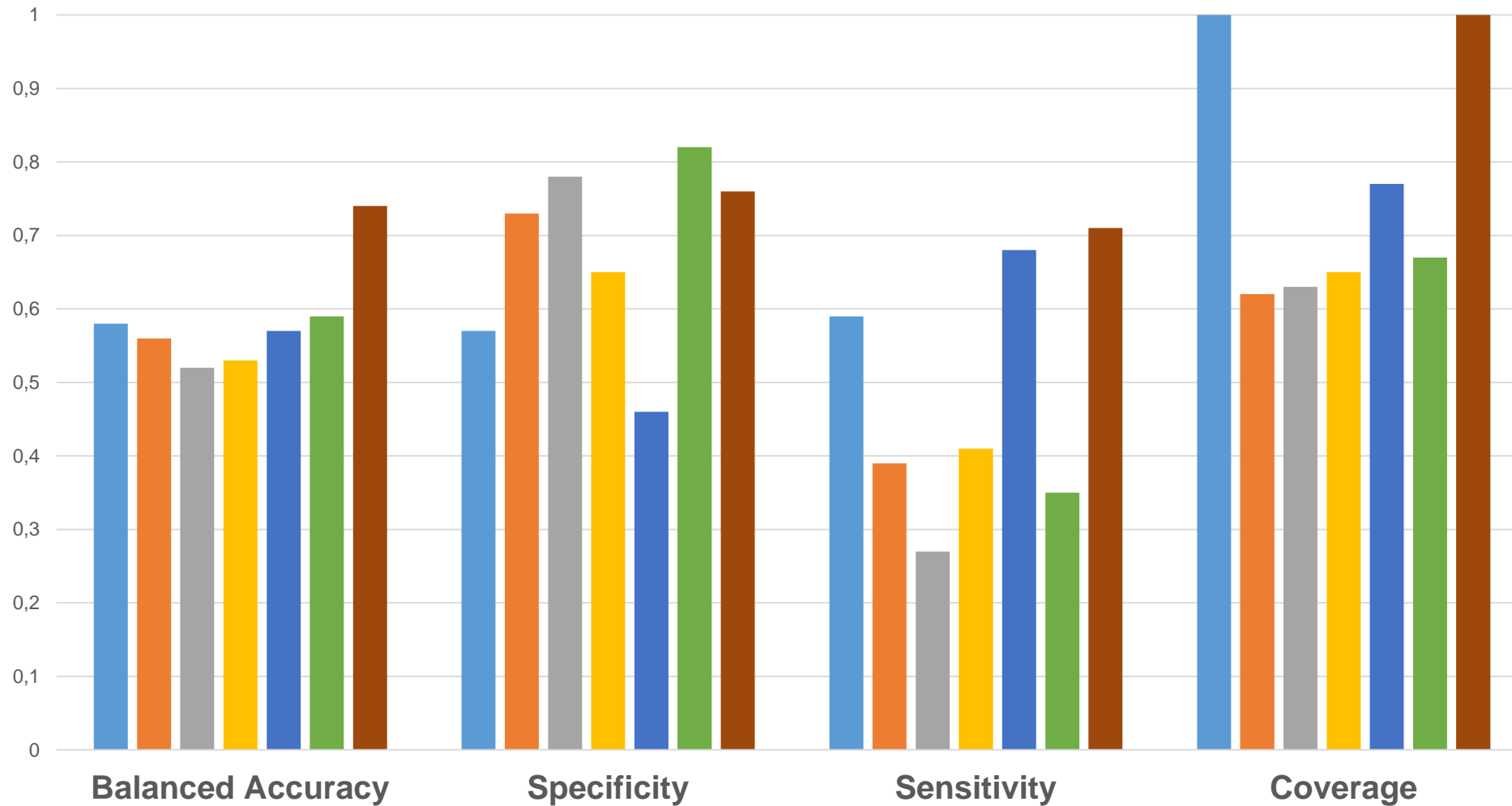
6971 Chemicals

Measures of goodness-of-fit, robustness and predictivity (External Validation)

Benchmark - Combined dataset (GMTP, LLNA and human data)

6971 Chemicals

Measures of goodness-of-fit, robustness and predictivity (External Validation)



Alert analysis

DPRA

KeratinoSens

h-CLAT

LLNA

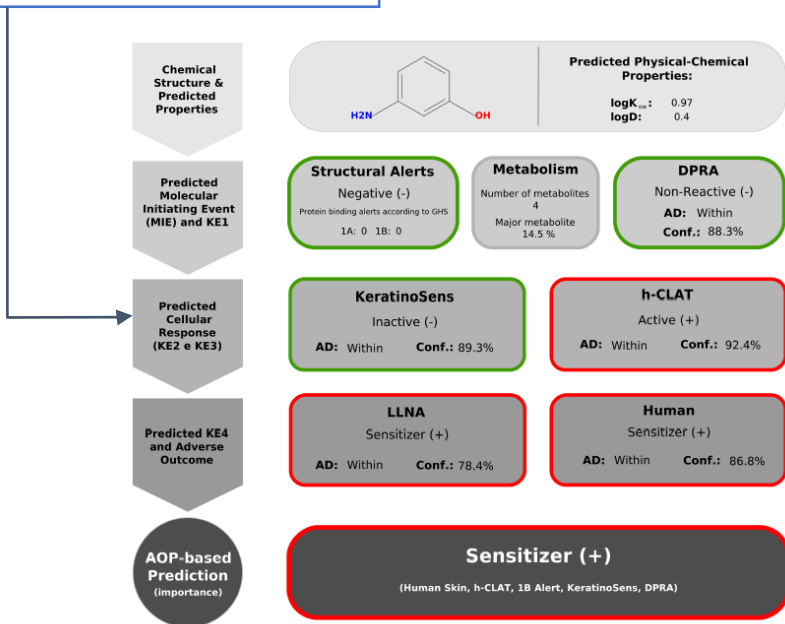
Human Skin

AOP-based prediction

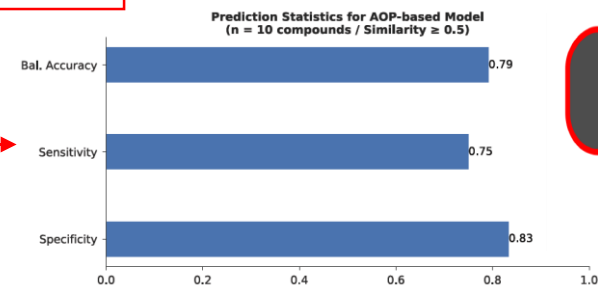
OECD Principles of (Q)SAR Validation for regulatory purposes

1. A defined endpoint;

2. An unambiguous algorithm;



4. Appropriate measures of goodness-of-fit, robustness and predictivity;

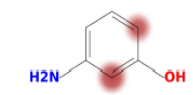


Predicted Outcome/Assay	Predicted class (Confidence)	STR Contribution Mapping
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Local Lymph Node Assay (LLNA, OECD 429)

Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK_{ow}, logD)

Sensitizer (+) (78.4%)

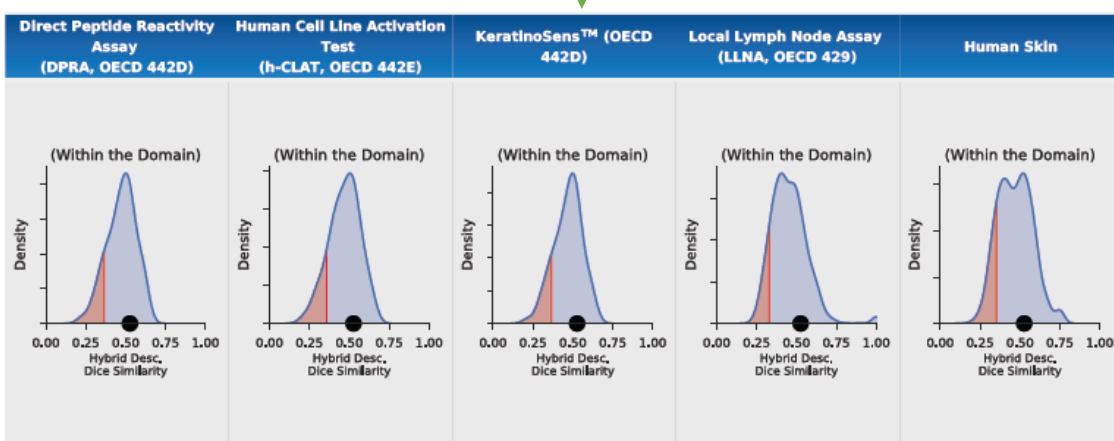


Sensitizer (+)

Concordance between KES: 65.3%
Confidence level (external validation): 100.0%



3. A defined domain of applicability;



5. A mechanistic interpretation, if possible.

Structural Alert Analysis

Result: (+) Positive

Alerts were found in the molecule. The results are in the table below and a description is provided at the end of the report.

Category	Alert	Alert ID	References
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Skin Sensitization Category 1A
(Protein binding Alerts, EC3 (LLNA) ≤ 2%; NOEL (IRPT) ≤ 500 µg/cm²)

Polarised Alkenes - sulfonates

Roberts, D., Apl, A.M., Safford, R., Lalko, J. Regulatory Toxicology and Pharmacology 72 (2015) 683-693.

Aptula A.O. et al., (2006) Chemical Research in Toxicology, 19, 1097.



Predicted endpoint/Method	Predicted class (Confidence)	STR Contribution Mapping
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Direct Peptide Reactivity Assay (DPRA, OECD 442D)

Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK_{ow}, logD)

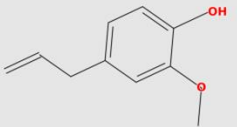
Reactive (+) (82.8%)



Prehaptens and prohaptens - Activation of weak or non-sensitizing substances into sensitizers

1

Chemical Structure & Predicted Properties



Predicted Physical-Chemical Properties:

logK_{ow}: 2.13
logD: 1.97

Structural Alerts

Negative (-)

Protein binding alerts according to GHS

1A: 0 1B: 0

Metabolism

Number of metabolites: 6

Major metabolite: 27.7 %

DPRA

Reactive (+)

AD: Within

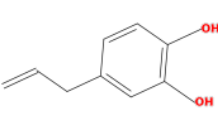
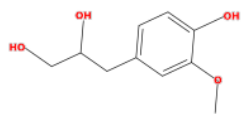
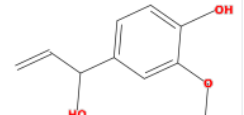
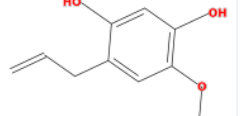
Confidence: 88.6%

Predicted Molecular Initiating Event (MIE)

2

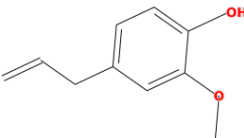
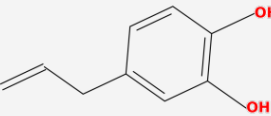
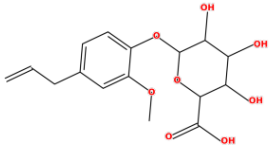
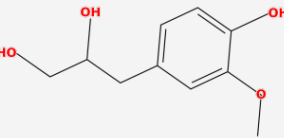
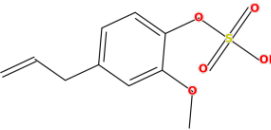
Metabolism prediction and potential for haptentation

To assess both direct and indirect haptens, this module predicts the potential for metabolic activation (pro-hapten formation) by known Phase I reactions, i.e., it can be used to identify potential skin sensitizers which require some type of metabolism to an active metabolite (pro-haptens) before initiation of the key event 1 (KE1) in a skin sensitization AOP (OECD Principle 5).

Metabolite (predicted structure)	SMILES	Reaction Rule	Metabolite Score	AOP-based Prediction
	<chem>C=CCc1ccc(O)c(O)c1</chem>	O-Demethylation	27.7 %	Sensitizer (+)
	<chem>COc1cc(CC(O)CO)ccc1O</chem>	Vinyl Oxidation	20.0 %	Sensitizer (+)
	<chem>C=CC(O)c1ccc(O)c(OC)c1</chem>	Benzylic Hydroxylation	7.3 %	Sensitizer (+)
	<chem>C=CCc1cc(O)c(O)c(O)c1</chem>	Aromatic Hydroxylation	5.6 %	Sensitizer (+)

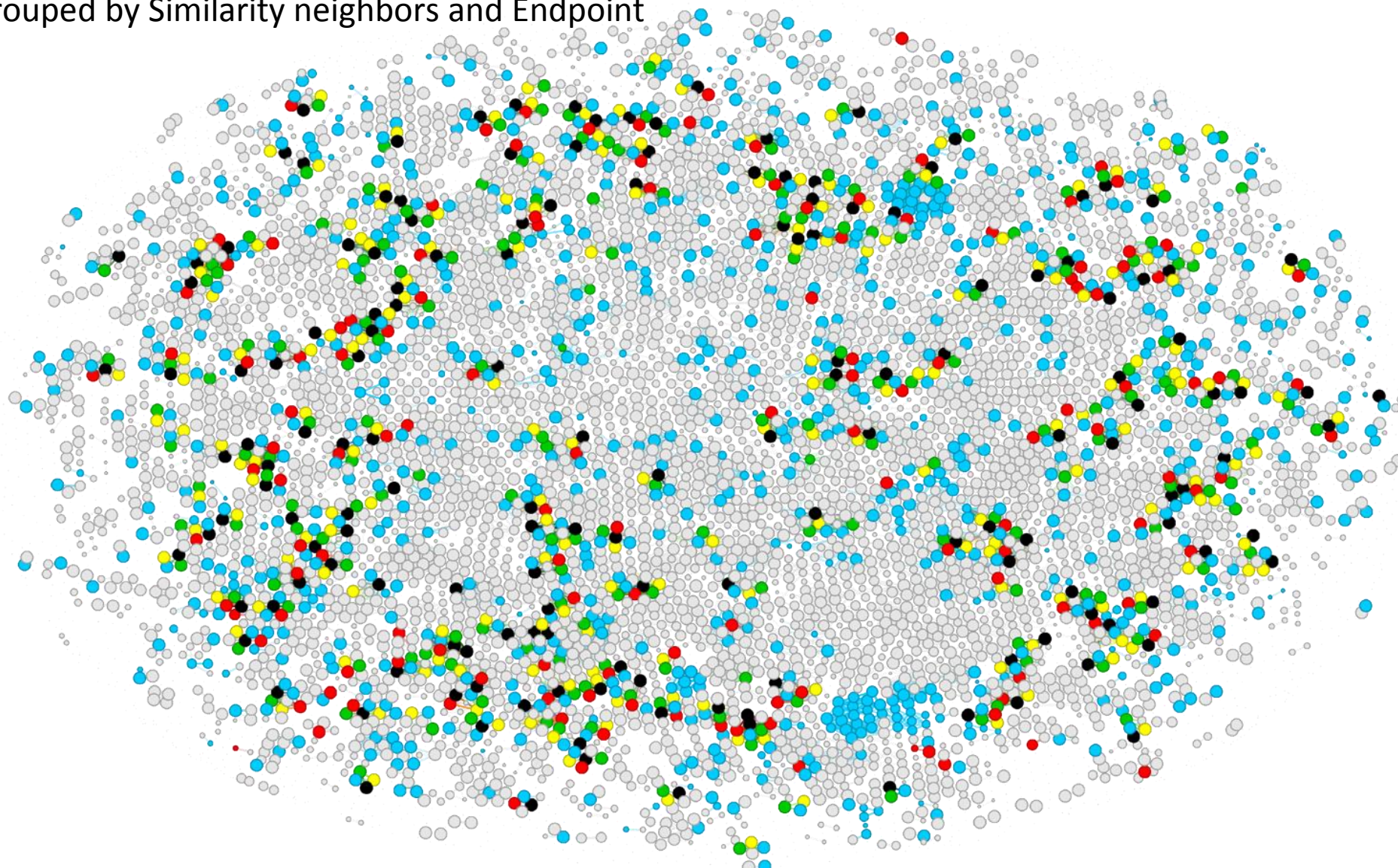
3

Pred-CYP2D™

Metabolite (Chemical Name)	SMILES (Reaction Rule)	Metabolite Score	Metabolite Database	Bioassay Database
Parent Molecule (Query)				
 (Eugenol)	<chem>C=CCc1ccc(O)c(OC)c1</chem>	-	Human endogenous metabolite	Approved Drugs, Compounds that have been investigated (clinical trials)
 (4-(prop-2-en-1-yl)benzene-1,2-diol)	<chem>C=CCc1ccc(O)c(O)c1</chem> (O-Demethylation)	27.7 %	Human endogenous metabolite	<i>in vitro</i> active at 10 uM
	<chem>C=CCc1ccc(OC2OC(C(=O)O)C(O)C(O)C2O)c(OC)c1</chem> (O-Glucuronidation)	25.0 %	-	-
 (3-(4-Hydroxy-3-methoxyphenyl)-1,2-propanediol)	<chem>COc1cc(CC(O)CO)ccc1O</chem> (Vinyl Oxidation)	20.0 %	Human endogenous metabolite	Tested in man but not approved or in trials
 ([2-methoxy-4-(prop-2-en-1-yl)phenyl]oxidanesulfonic acid)	<chem>C=CCc1ccc(OS(=O)(=O)O)c(OC)c1</chem> (Sulfation)	11.9 %	Human endogenous metabolite	<i>in vitro</i> active at 10 uM

Chemical and Toxicological Space for Skin Sensitization

Grouped by Similarity neighbors and Endpoint



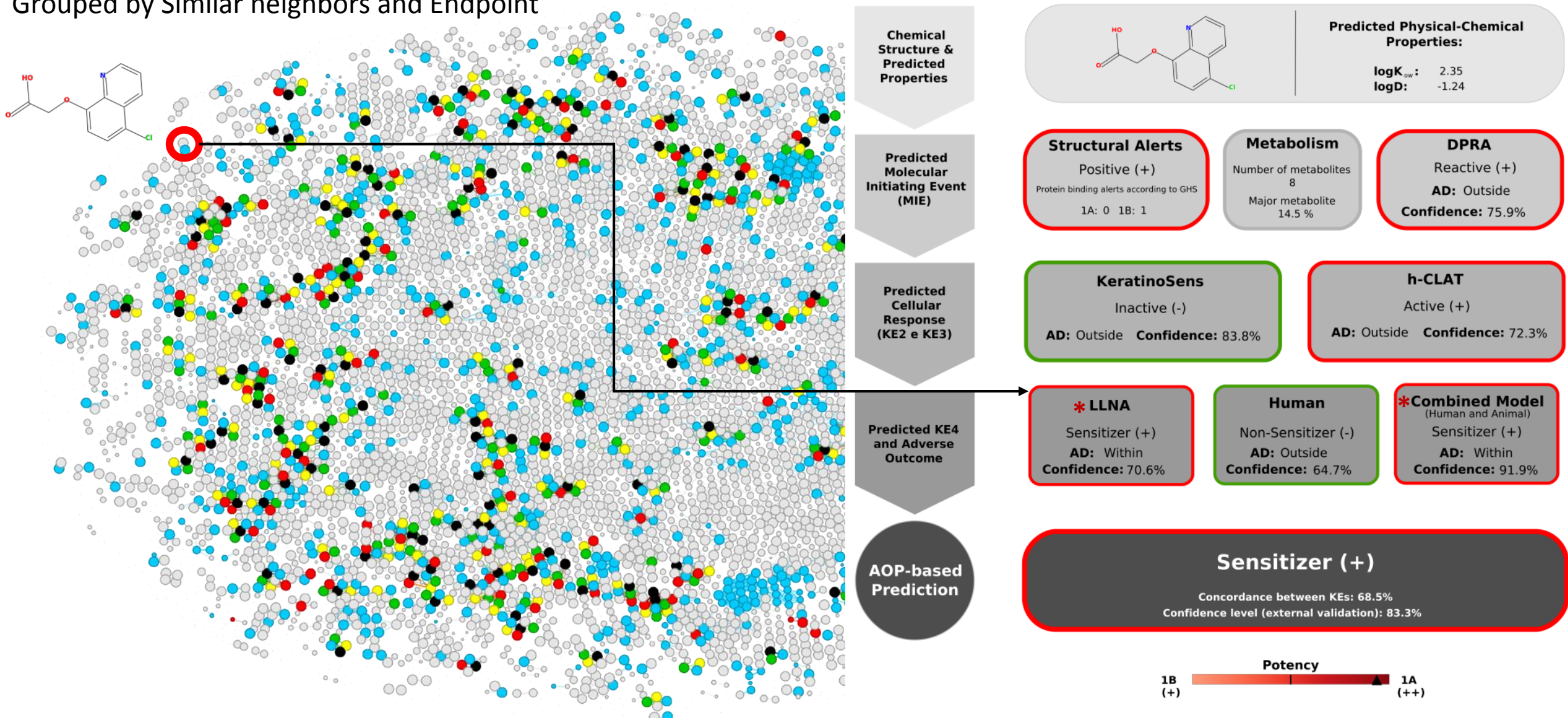
Diversity?

Data	
DPRA	195
KeratinoSens	190
h-CLAT	161
LLNA	997
Human Skin	389
Combined dataset (GMTP, LLNA and human data)*	6971

Source ○ Combined dataset ● DPRA ● Human Skin ● KeratinoSens ● LLNA ● h-CLAT

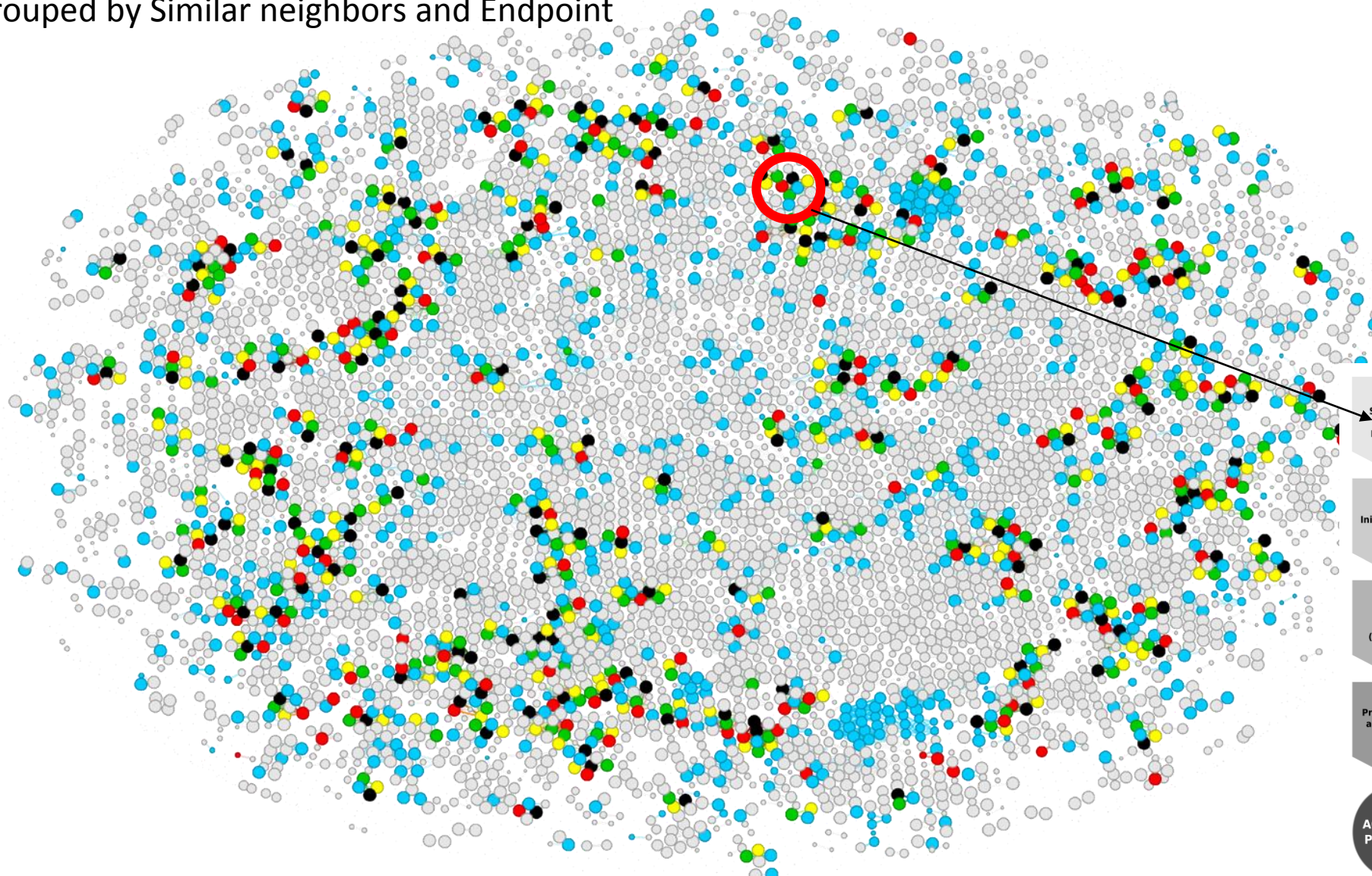
Chemical and Toxicological Space for Skin Sensitization

Grouped by Similar neighbors and Endpoint



Chemical and Toxicological Space for Skin Sensitization

Grouped by Similar neighbors and Endpoint



Chemical Structure & Predicted Properties

Predicted Physical-Chemical Properties:

logK_{ow}: 0.97
logD: 0.4

Predicted Molecular Initiating Event (MIE)

Structural Alerts

Negative (-)

Protein binding alerts according to GHS

1A: 0 1B: 0

Metabolism

Number of metabolites: 4

Major metabolite: 14.5%

DPRA

Non-Reactive (-)

AD: Within

Confidence: 88.3%

Predicted Cellular Response (KE2 e KE3)

KeratinoSens

Inactive (-)

AD: Within Confidence: 89.3%

h-CLAT

Active (+)

AD: Within Confidence: 92.4%

Predicted KE4 and Adverse Outcome

LLNA

Sensitizer (+)

AD: Within Confidence: 78.4%

Human

Sensitizer (+)

AD: Within Confidence: 86.8%

Combined Model
(Human and Animal)

Sensitizer (+)

AD: Within Confidence: 94.4%

AOP-based Prediction

Sensitizer (+)

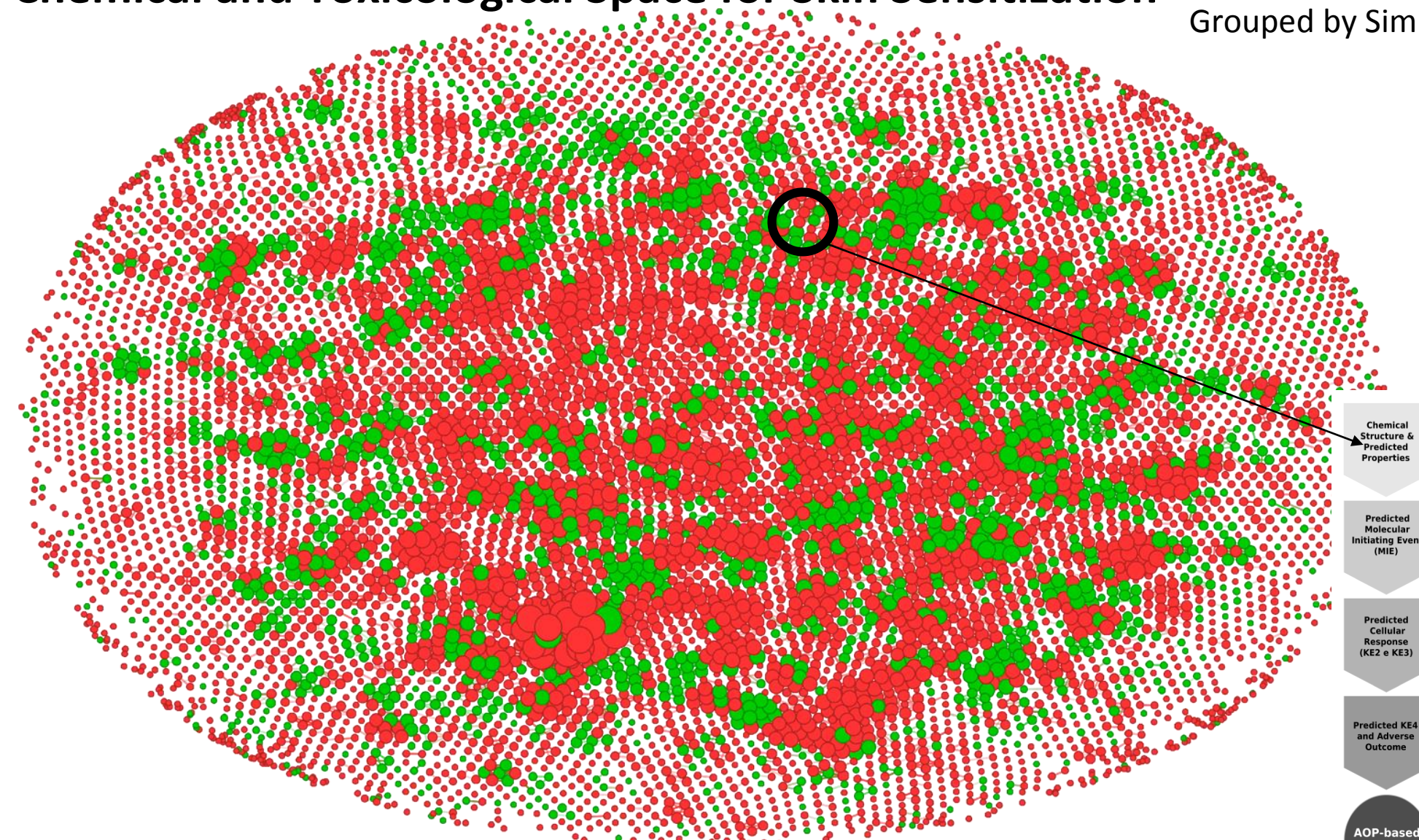
Concordance between KEs: 62.8%
Confidence level (external validation): 87.5%

Source ○ Combined dataset ● DPRA ● Human Skin ● KeratinoSens ● LLNA ● h-CLAT



Chemical and Toxicological Space for Skin Sensitization

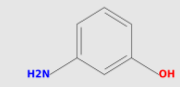
Grouped by Similar neighbors and Endpoint



Outcome ● (+) ● (-)

Neighbor Count ○ 5 ○ 15 ○ 25 ○ 35

Chemical Structure & Predicted Properties



Predicted Physical-Chemical Properties:

logK_{ow}: 0.97
logD: 0.4

Structural Alerts

Negative (-)

Protein binding alerts according to GHS

1A: 0 1B: 0

Metabolism

Number of metabolites: 4

Major metabolite: 14.5%

DPRA

Non-Reactive (-)

AD: Within

Confidence: 88.3%

KeratiNoSens

Inactive (-)

AD: Within Confidence: 89.3%

h-CLAT

Active (+)

AD: Within Confidence: 92.4%

LLNA

Sensitizer (+)

AD: Within Confidence: 78.4%

Human

Sensitizer (+)

AD: Within Confidence: 86.8%

Combined Model
(Human and Animal)

Sensitizer (+)

AD: Within Confidence: 94.4%

Sensitizer (+)

Concordance between KEs: 62.8%
Confidence level (external validation): 87.5%

Potency

1B (+)

1A (++)

AOP-based Prediction

Final Remarks

- A logical framework balancing transparency, mechanistic interpretability and predictivity in a sequential chain of causally linked events at different levels;
- To assess key event relationships (KERs) integrating *in chemico*, *in vitro*, *ex vivo* and *in vivo* data in the *in silico* models;
- To design new regulatory decision trees based in predictive Integrated Approaches to Testing and Assessment (IATA) containing *in silico* models ;



Thank you for your attention!

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