

Challenges and opportunities for using AOP-based *in silico* models in regulatory contexts

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Overview

- The “AOP-based *in silico* model” concept
- AOP-based *in silico* model – Alttox’s Framework
- Examples: Balancing transparency, mechanistic interpretability *and* predictivity
- Challenges and opportunities for using these methods in regulatory contexts

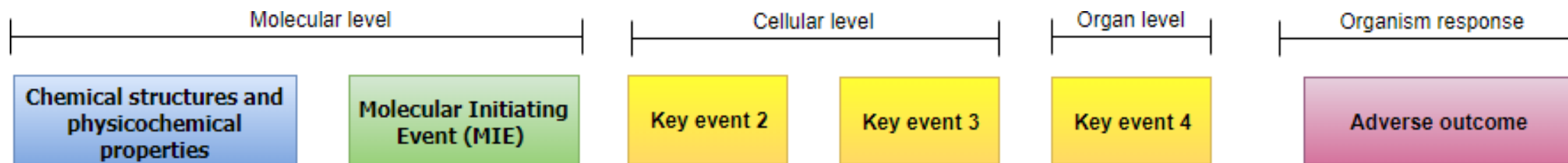
Introduction

Adverse outcome pathways (AOP) framework

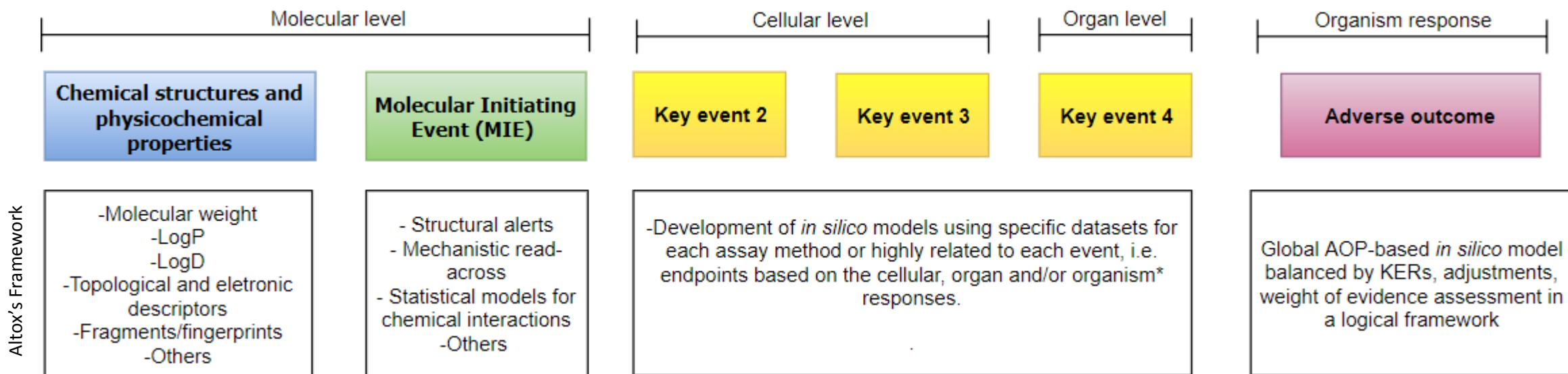
- An AOP is an analytical construct that describes a sequential chain of causally linked events at different levels of biological organisation that lead to an adverse health or ecotoxicological effect (OECD, 2012)
 - Regulatory challenge:** to define when there is sufficient confidence predictivity to use one or more alternative models for regulatory purposes

Proposed “AOP-based *in silico* model” Concept

- A framework composed by individual *in silico* models 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**.

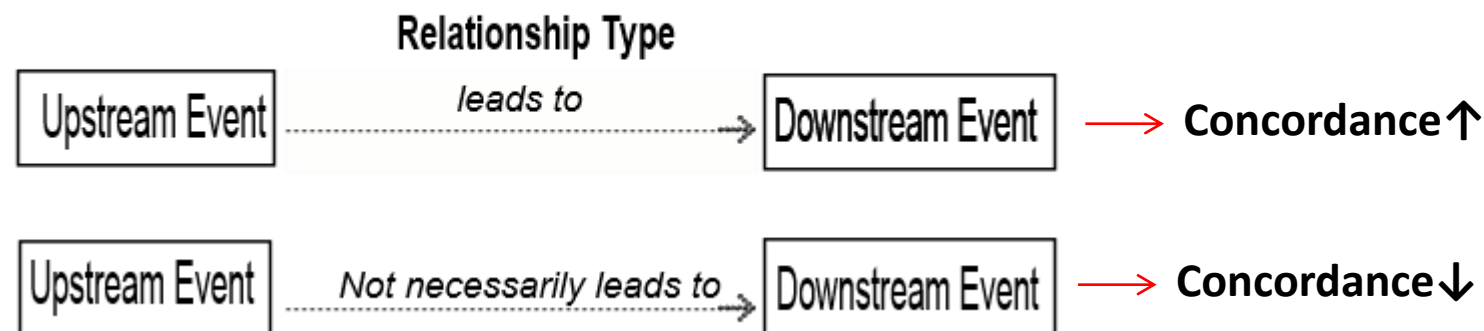


AOP-based *in silico* model – Alttox’s Framework

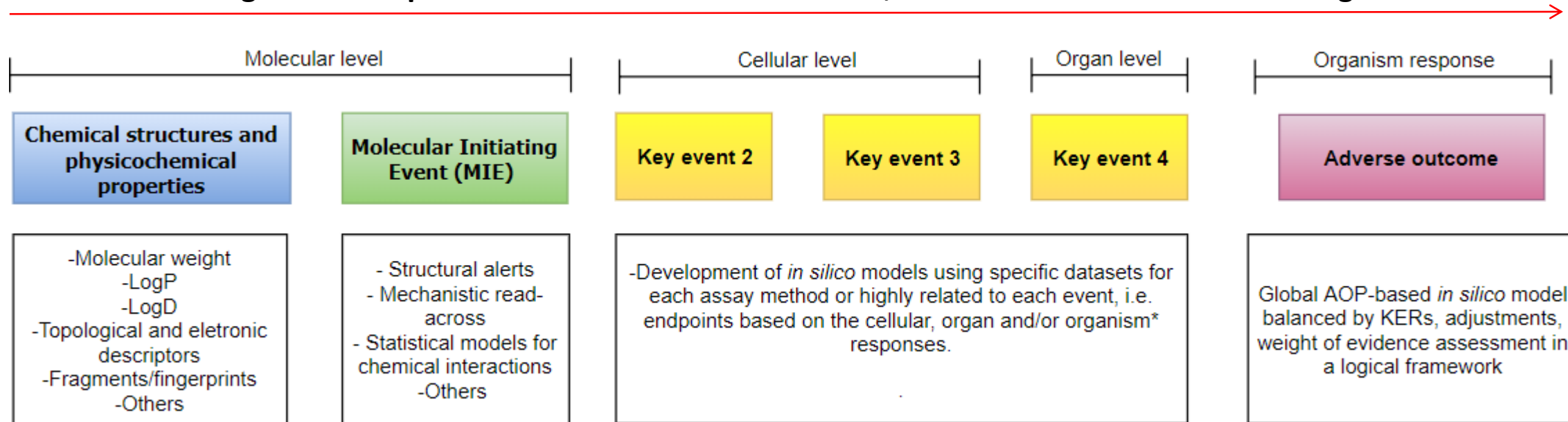


Key Event Relationships (KERs)

Key event relationships (KERs)

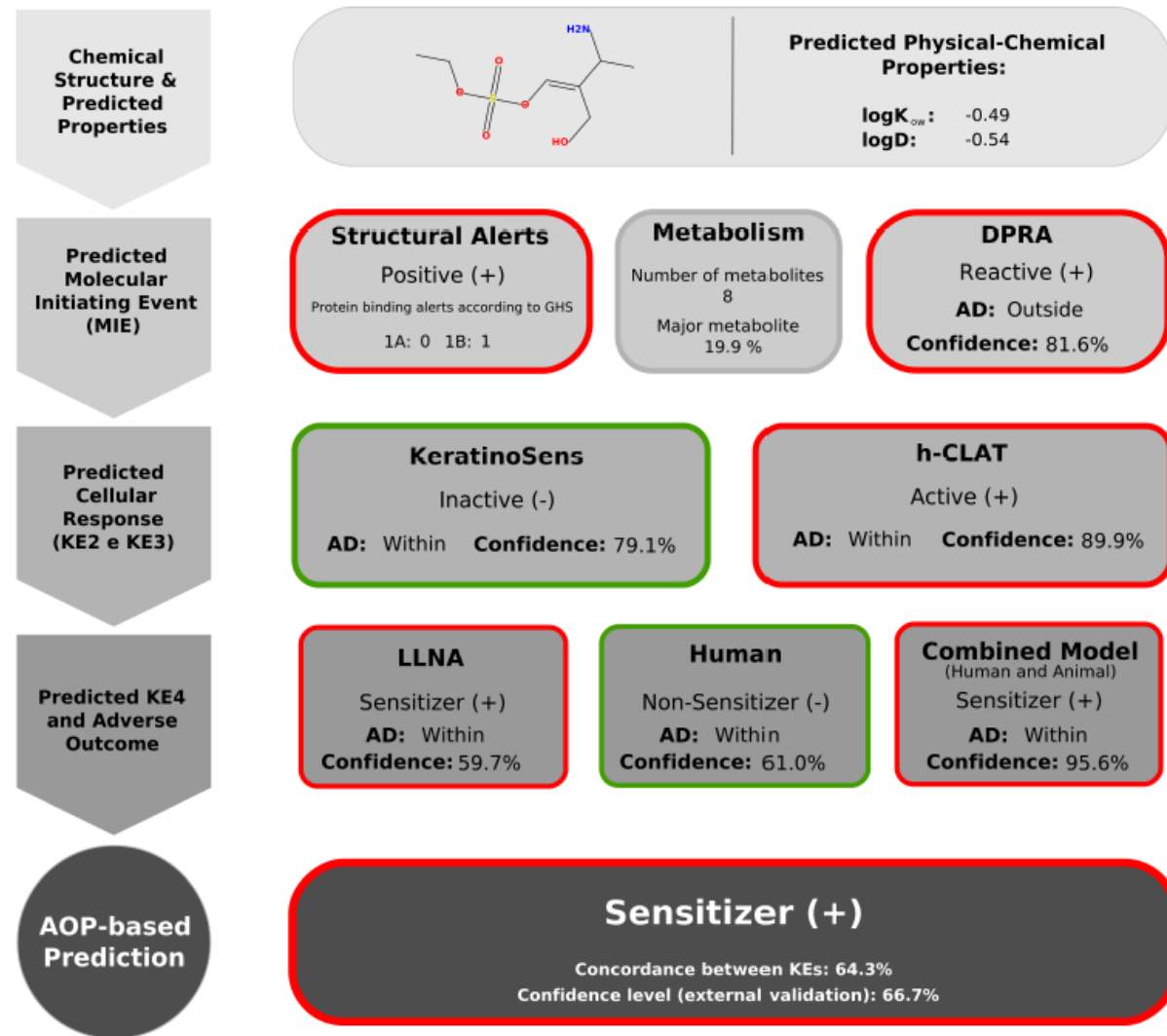


Different weights for the predictions based in the AOP level , KERs and confidence levels along the AOP



Example – AOP-Sens

- A logical framework balancing transparency, mechanistic interpretability and predictivity;
- Models to predict chemicals that **can activate** the AOP modular components (MIE/KEs);
- **A global model** integrating all multilevel predictions balancing predictivity, key events relationships and WoE adjustments, for predicting the outcome.



Structures and physicochemical properties

Molecular Query

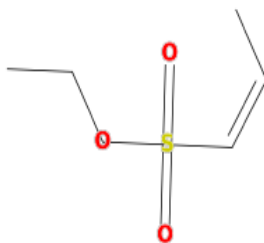
SMILES: C/C=C(S(=O)(=O)O)CC

Name: Molecule 2

logK_{ow}: 0.89

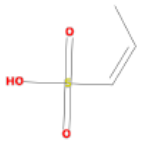
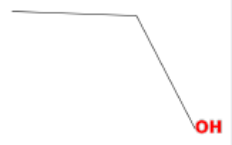
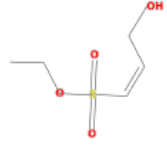
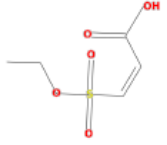
CAS: None

logD: 1.61



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).

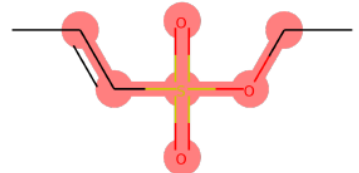
Metabolite (predicted structure)				
SMILES	<chem>CC=CS(=O)(=O)O</chem>	<chem>CCO</chem>	<chem>CCOS(=O)(=O)C=CCO</chem>	<chem>CCOS(=O)(=O)C=CC(=O)O</chem>
Reaction Rule	Hydrolysis	Hydrolysis	Aliphatic Hydroxylation	Carboxylation
Metabolite Score	17.4 %	17.4 %	4.9 %	1.6 %

Molecular Initiating Event

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
Skin Sensitization Category 1A (Protein binding Alerts, EC3 (LLNA) ≤ 2%; NOEL (HRIPT) ≤ 500 µg/cm ²)		Polarised Alkenes - sulfonates	Roberts, D., Api, 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.

STR Contribution Map

Reactive / Active (+)  Non-Reactive / Inactive (-)

Predicted endpoint/Method	Predicted class (Confidence)	STR Contribution Mapping
Direct Peptide Reactivity Assay (DPRA, OECD 442D)	Reactive (+) (82.8%)	
Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK _{ow} , logD)		

Cellular and organ levels

Key Events

STR Contribution Map

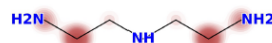


Predicted endpoint/Method	Predicted class (Confidence)	STR Contribution Mapping
KeratinoSens™ (OECD 442D) Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK _{ow} , logD)	Inactive (-) (89.7%)	
Human Cell Line Activation Test (h-CLAT, OECD 442E) Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK _{ow} , logD)	Inactive (-) (76.5%)	
Predicted Outcome/Assay	Predicted class (Confidence)	STR Contribution Mapping
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 (+) (76.4%)	

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 (+)
(76.4%)

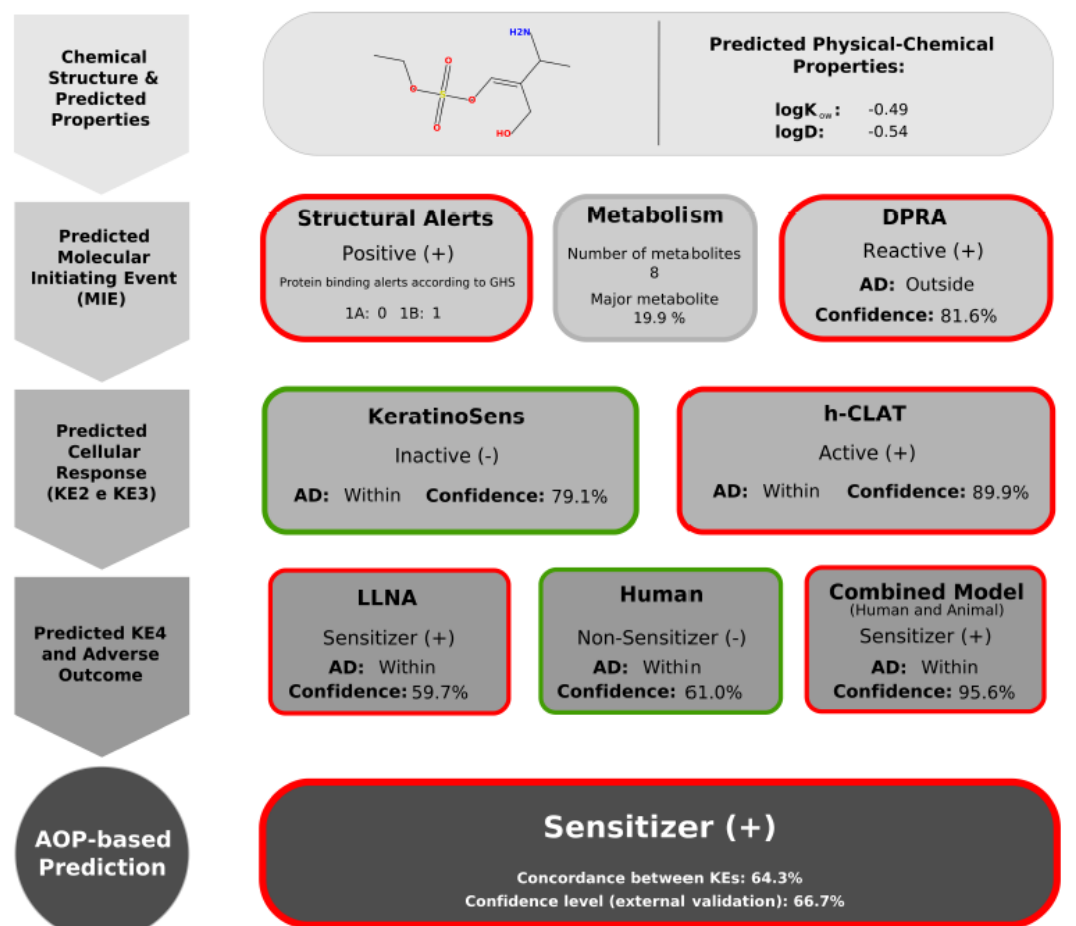


Organism response

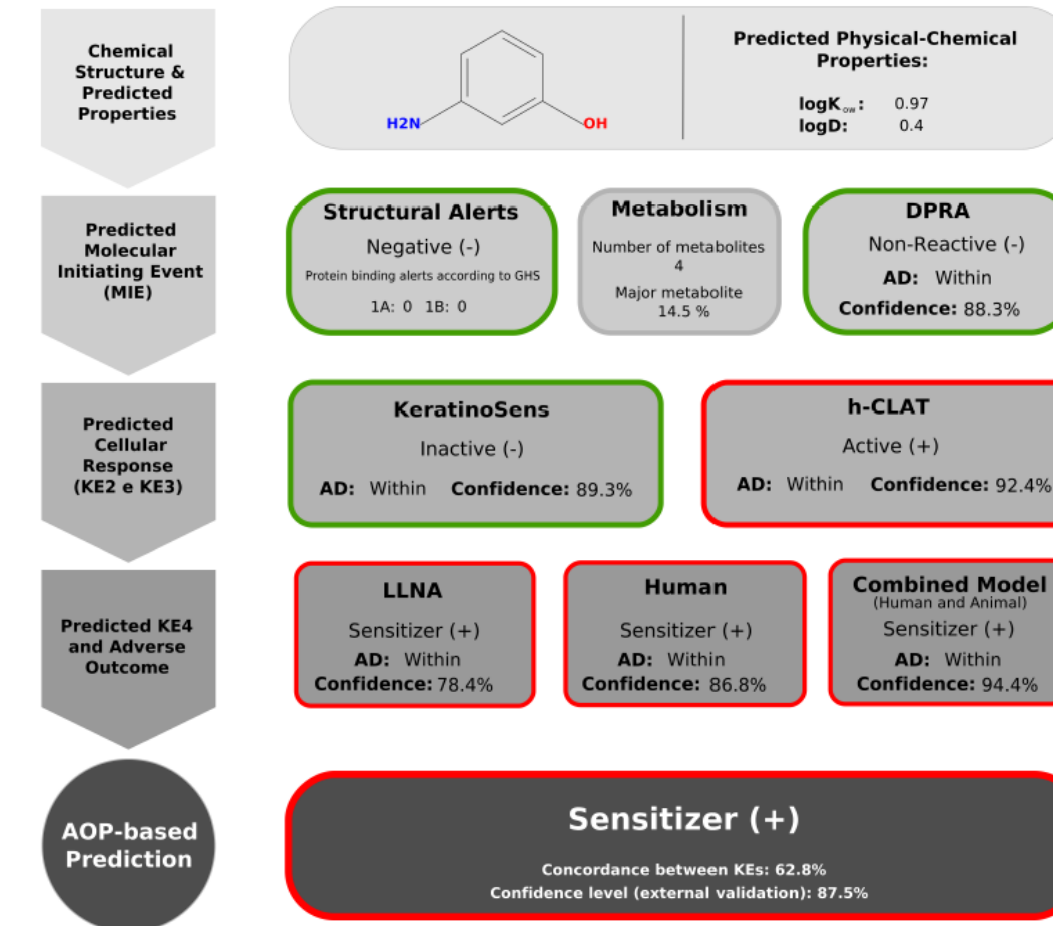
$$\text{AOP-based algorithm} = \sum_{j=1}^n p_j c_j a_j w_j$$

Where p is a prediction result, c is the confidence level, a is the applicability domain and w is a weight (adjusted by KERs)

Results



Results



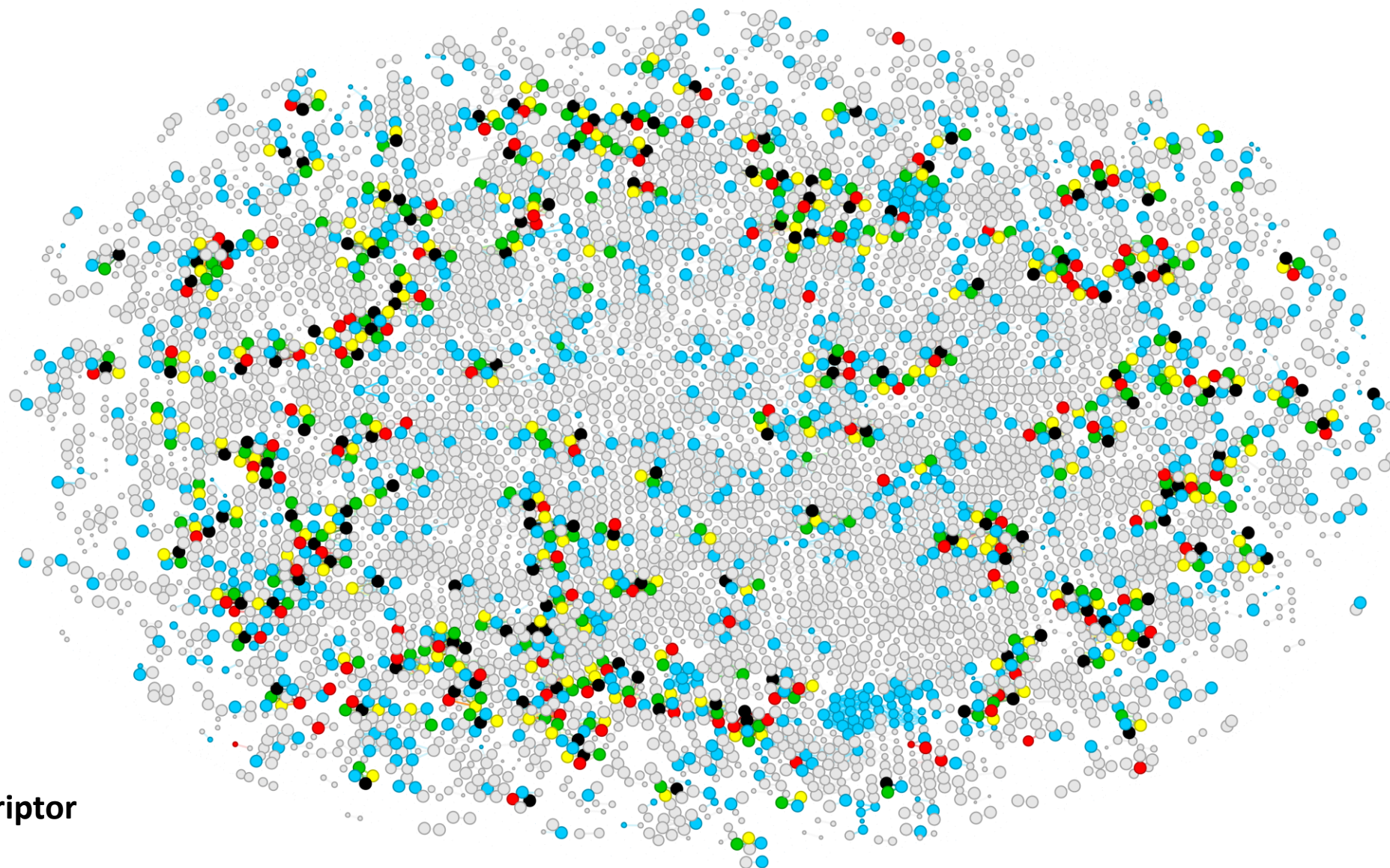
Benchmark - Combined dataset (GMTP, LLNA and human data)

(Q)SAR Models	Model Dataset	Specificity	Sensitivity	Accuracy	<i>N (external dataset)</i>
Alert analysis	197 128: 1A 69: 1B	0.57	0.59	0.58	6422
DPPRA	195	0.76	0.32	0.54	6422
KeratinoSens	190	0.78	0.27	0.52	4050
H-CLAT	161	0.65	0.41	0.53	4178
LLNA	997	0.46	0.68	0.57	4932
Human Skin	389	0.82	0.35	0.59	4177
Combined dataset (GMTP, LLNA and human data)**	6971	0.75	0.92	0.84	1284
AOP-based prediction	-	0.76	0.71	0.74	6422

** Internal validation

Chemical and Toxicological Space for Skin Sensitization

Grouped by Similarity neighbors and Endpoint



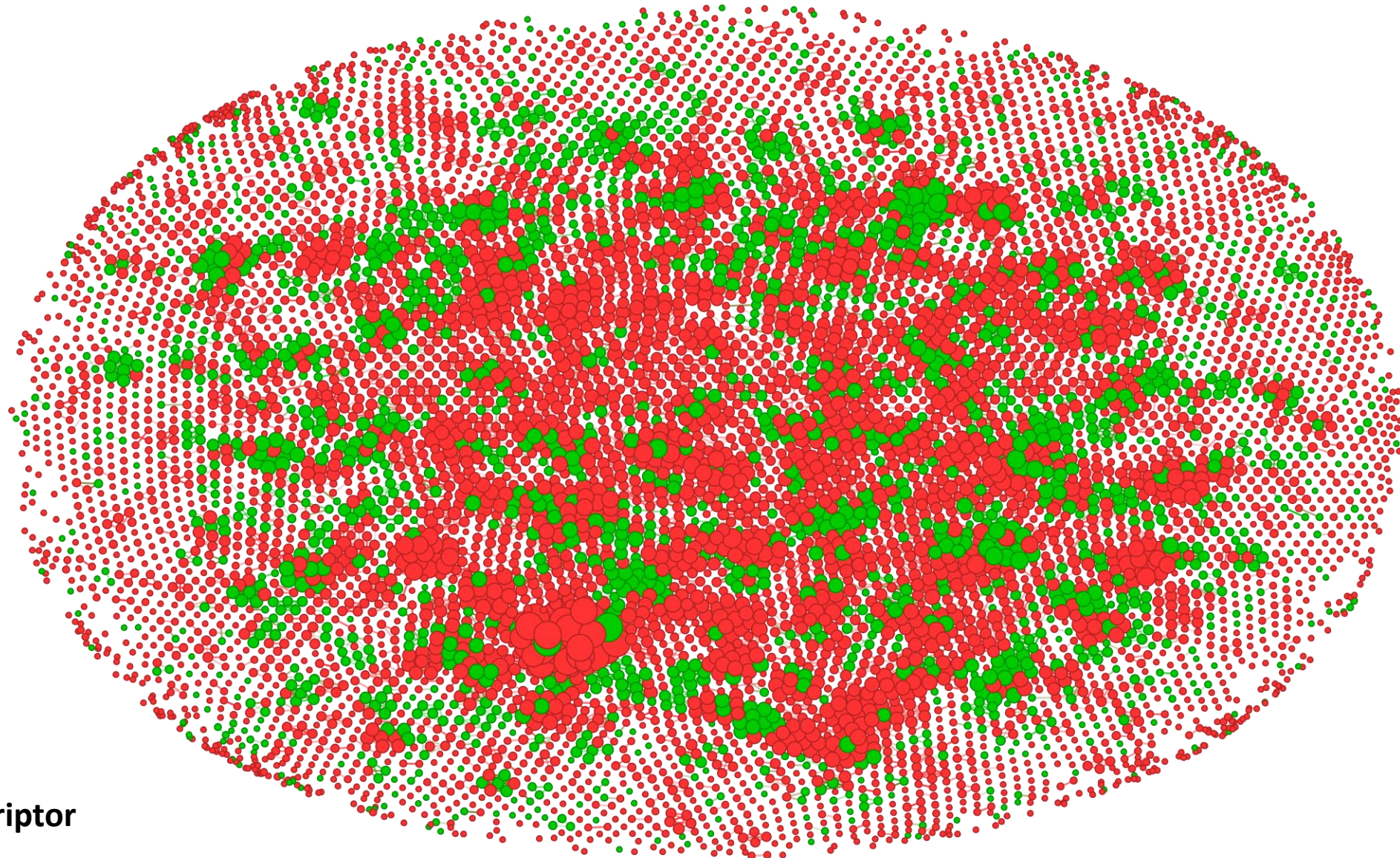
Hybrid descriptor

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

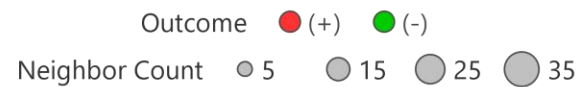
Combined dataset = (GMTP, LLNA and human data)

Chemical and Toxicological Space for Skin Sensitization

Grouped by Similarity neighbors and all Endpoints Outcome: Positive (+) or Negative (-)

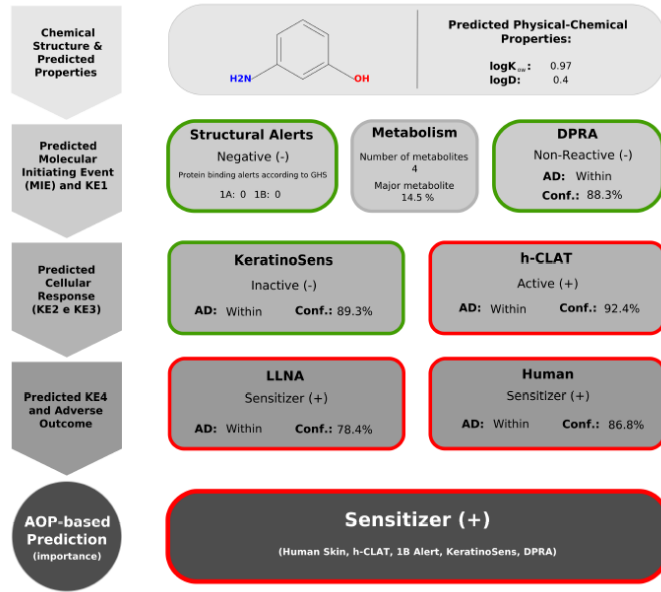


Hybrid descriptor



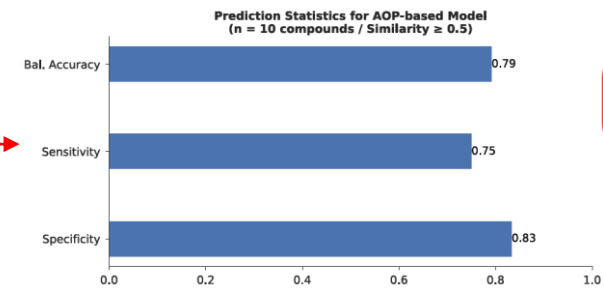
OECD Principles of (Q)SAR Validation for regulatory purposes

1. A defined endpoint;



2. An unambiguous algorithm;

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



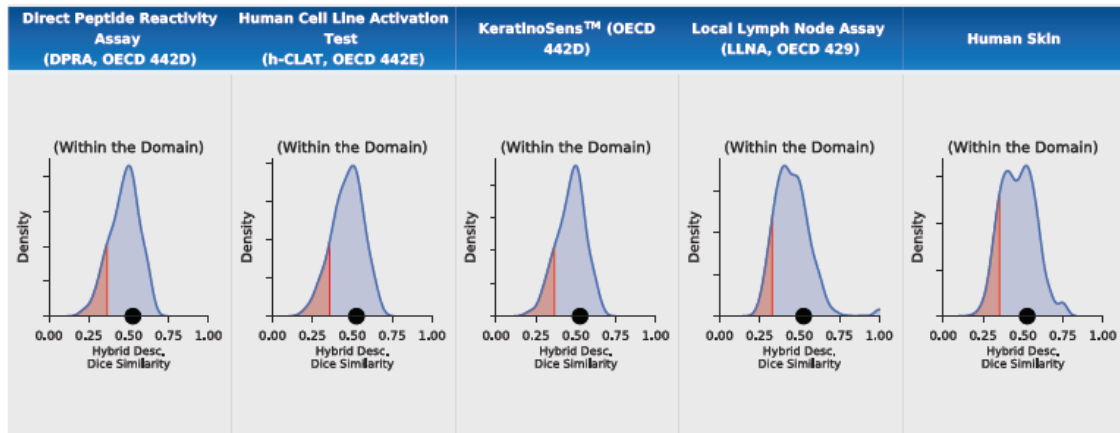
Predicted Outcome/Assay	Predicted class (Confidence)	STR Contribution Mapping
Local Lymph Node Assay (LLNA, OECD 429) Deep Learning categorical model implemented with hybrid descriptors (ECFP6 fingerprint and physicochemical properties: MW, TPSA, logK _{ow} , logD)	Sensitizer (+) (78.4%)	

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

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Predicted endpoint/Method	Predicted class (Confidence)	STR Contribution Mapping
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%)	

Challenges and opportunities

- Balancing predictivity and mechanistic interpretation of the models at different levels (adverse outcome pathways - AOP)
- Looking for patterns with predictivity level similar or higher than biological assays
- Filling gaps in safety assessments with *in silico* predictions
- Making new regulatory criteria based in predictive integrated approaches containing *in silico* models

We are seeking partnerships and collaborations with our tools!



Genotox-iSTM

Artificial intelligence and advanced machine learning algorithms for mutagenicity assessment



Thank you for your attention!

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