

Overview of toxicity prediction methods

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The challenges

Contaminants, drugs...

Increased use of chemical substances

Transformation products

Decline of species

One health

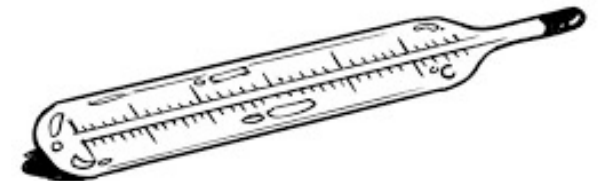
Lack of data

....



Are we adequate?
Are we effective?

- Which are our objectives?
- Do we have priorities?
- What is our timeline?
- Do we measure impact of our activities?





Not apostasy Not conversion

- Not to loose fundamental issues
- Need to focus our activities

Caravaggio, The conversion of S. Paul

Broad view

Priorities: hazard and exposure

One health

Single conceptual / in silico architecture



Safety (\neq lack of risk)

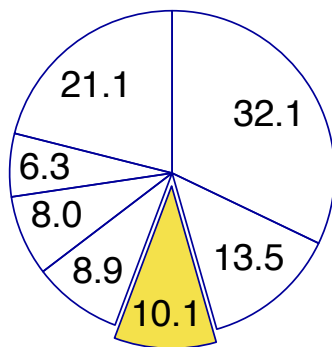
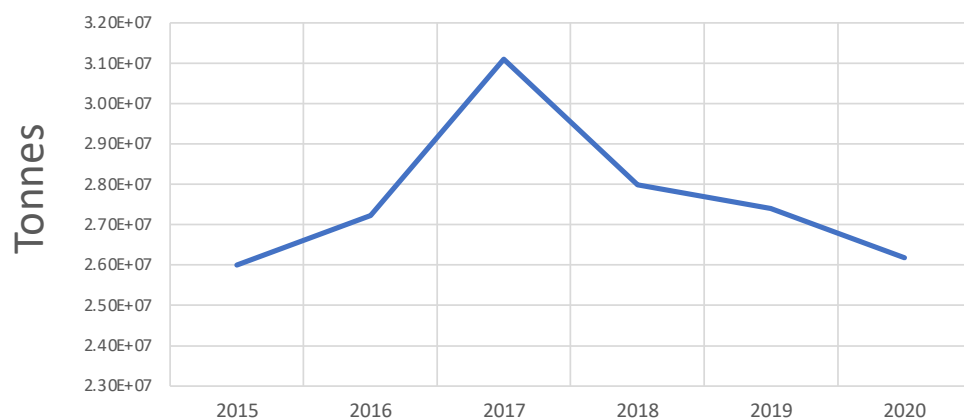
Beneficial aspects

Substitution

Green Deal

Risk. Global view. Are we exporting the risk?

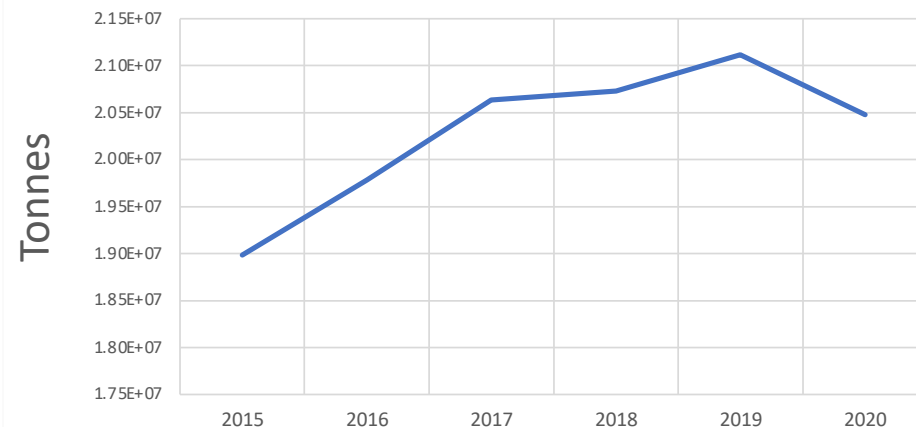
Production



Italy: 10% of the chemical market in EU (after Germany and France)

Trend Production / Import in Italy, 2015-2020

Import



Market of CMR and explosives

Previously we observed a similar shift in the period 2011-2015, in Italy, moving toward import, in particular for **CMR** (reduced diversity) **and explosive substances**

Marzo M, Leone C, Toma C, Roncaglioni A, Gianazzi S, Knauf R, Benfenati E. Impact of REACH legislation on the production and importation of CMR (carcinogen, mutagen and reproductive) and explosive chemicals in Italy from 2011 to 2015. Regul Toxicol Pharmacol 2019; 101 : 166-171 doi: 10.1016/j.yrtph.2018.11.013

Priorities and new methods

- Once we have the global view, we can cope with priorities
- Based on the global approach, global meter
- (1) In silico models for integrated view

Priorities: human (disease burden).

Ecotox (in The Netherlands 90% of the amphibians disappeared (for a disease), thus we should focus on amphibians, compared to fish)

(2) In silico models for priority endpoints

The new in silico tools for priorities

- Which in silico models are of higher relevance because addressing priority endpoints (hazard)?
- Which substances are of higher priorities (exposure)?

Mn Euro

EU28

Petrochemicals and Derivatives	143.568
Inorganic Industrial Chemicals	77.629
<i>Fertilizers</i>	24.065
<i>Industrial Gases</i>	11.644
<i>Other inorganics</i>	41.920
Specialty Chemicals	153.394
<i>Paints & Inks</i>	42.860
<i>Dyes & Pigments</i>	16.918
<i>Auxiliaries for Industry</i>	82.337
<i>Crop Protection</i>	11.279
Polymers	120.330
<i>Plastics & Synthetic rubber</i>	109.031
<i>Man-Made Fibres</i>	11.299

TOTAL	494.922
Pharmaceuticals	313.236
Personal Care Products	69.957

In silico platforms. Opportunities ... and challenges



Networks between different platforms

VEGA

OCHEM

Danish QSAR Database

AMBIT

Different models. Opportunities ... and challenges



increased confidence,
increased perspectives



different metrics, different
info (overlap ?)

In silico models. Future?

- Integrating multiple tools for the same endpoint
- Covering AOP, same overall toxicological category (e.g. muta+geno+carcino)
- Integrating hazard and exposure
- SSbD
- Integrating risk/benefit

In silico models. Predictions and ...

- **Reasoning** («*predicting*» mechanism, causality, ...)
- **Heuristics and expert systems** (supervised/unsupervised)
- Link with
 1. regulation,
 2. confidence,
 3. planning safer substances

Weight of evidence (WoE): EFSA Guidance

SCIENTIFIC OPINION



ADOPTED: 12 July 2017

doi: 10.2903/j.efsa.2017.4971

Guidance on the use of the weight of evidence approach in scientific assessments

EFSA Scientific Committee,
Anthony Hardy, Diane Benford, Thorhallur Halldorsson, Michael John Jeger,
Helle Katrine Knutsen, Simon More, Hanspeter Naegeli, Hubert Noteborn, Colin Ockleford,
Antonia Ricci, Guido Rychen, Josef R Schlatter, Vittorio Silano, Roland Solecki,
Dominique Turck, Emilio Benfenati, Qasim Mohammad Chaudhry, Peter Craig,
Geoff Frampton, Matthias Greiner, Andrew Hart, Christer Hogstrand, Claude Lambre,
Robert Luttik, David Makowski, Alfonso Siani, Helene Wahlstroem, Jaime Aguilera,
Jean-Lou Dorne, Antonio Fernandez Dumont, Michaela Hempen, Silvia Valtueña Martínez,
Laura Martino, Camilla Smeraldi, Andrea Terron, Nikolaos Georgiadis and Maged Younes

<https://www.efsa.europa.eu/en/efsajournal/pub/4971>

EFSA Guidance on WoE

Approach for WoE

1. Gather all info
2. Evaluate individual lines of evidence
3. Integrate the results

EFSA Guidance: integration

Criteria for integration

1. Relevance
2. Reliability
3. Agreement

In silico and read-across: integration

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Review article

Integrating *in silico* models and read-across methods for predicting toxicity of chemicals: A step-wise strategy



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^d Scientific Committee and Emerging Risks Unit, European Food Safety Authority, Via Carlo Magno 1A, Parma, Italy



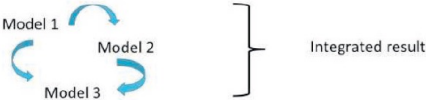
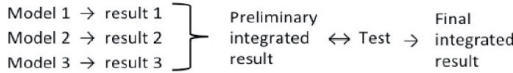
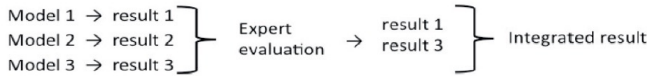
ARTICLE INFO

Handling Editor: Da Chen

ABSTRACT

In silico methods and models are increasingly used for predicting properties of chemicals for hazard identification and hazard characterisation in the absence of experimental toxicity data. Many *in silico* models are available and can be used individually or in an integrated fashion. Whilst such models offer major benefits to toxicologists, risk assessors and the global scientific community, the lack of a consistent framework for the integration of *in silico* results can lead to uncertainty and even contradictions across models and users, even for the same chemicals. In

Integration of in silico

Algebraic and voting methods	<p>Algebraic methods</p> <p>Model 1 → result 1 Model 2 → result 2 Model 3 → result 3</p> 
Weighing	<p>Weighing methods</p> <p>Model 1 → result 1 → transformed result 1 Model 2 → result 2 → transformed result 2 Model 3 → result 3 → transformed result 3</p> 
Hybrid	<p>Hybrid methods</p> 
Learning	<p>Learning methods</p> <p>Model 1 → result 1 Model 2 → result 2 Model 3 → result 3</p> 
Expert-based	<p>Expert-based integration</p> <p>Model 1 → result 1 Model 2 → result 2 Model 3 → result 3</p> 

Algebraic methods

Majority vote

Unanimity

Worst case

All models at the same level of reliability

Or you introduce thresholds (in / out: 2 levels or reliability)

Weighing methods

VEGA and mutagenicity is an example
Use of all models, in a quantitative way
(not in or out, binary, qualitative approach)

Consensus model (CNS-VEGA) : CAESAR + SARPY + TT-VEGA

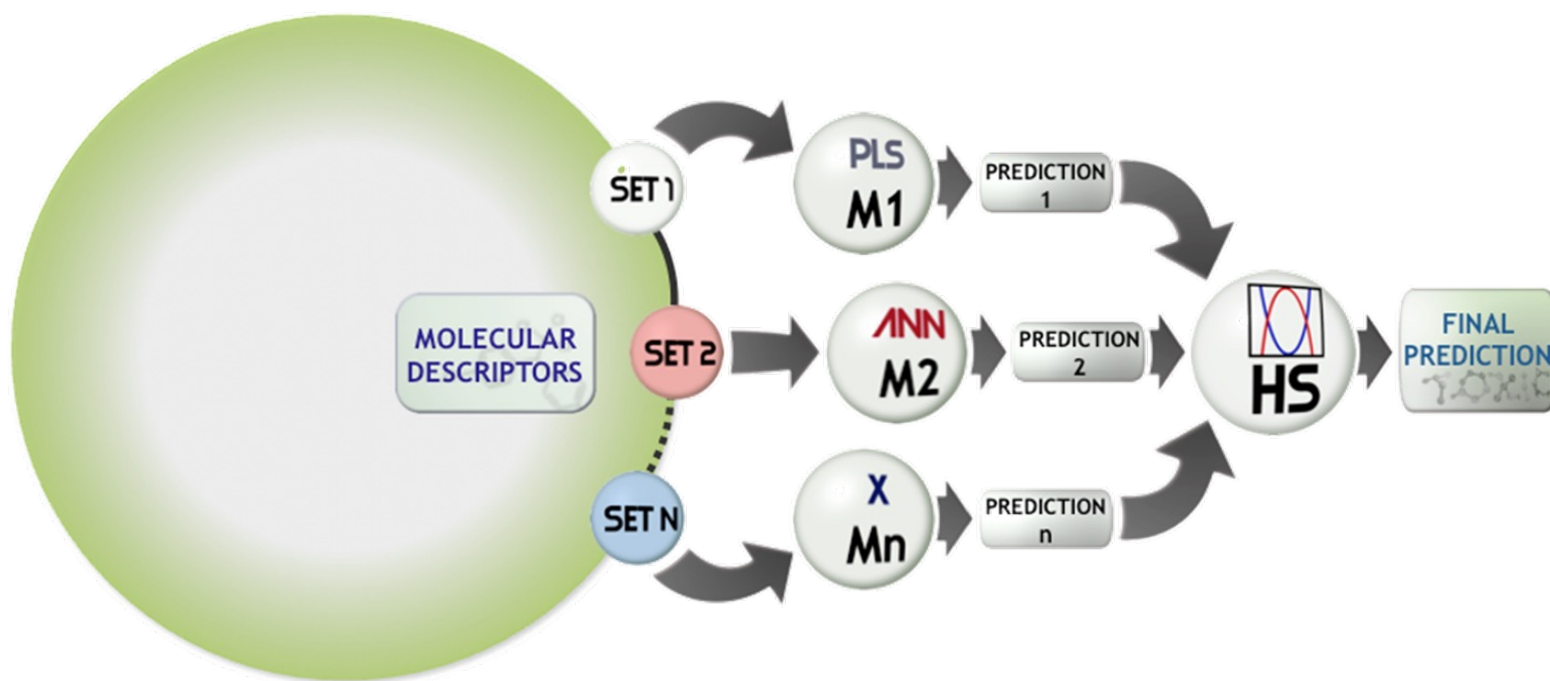
$$\text{CONSENSUS} = \frac{(\pm 1) * AD_{\text{CAESAR}} + (\pm 1) * AD_{\text{SARPY}} + (\pm 1) * AD_{\text{TTVEGA}}}{AD_{\text{CAESAR}} + AD_{\text{SARPY}} + AD_{\text{TTVEGA}}}$$

Algorithm extended now to 4 models



Hybrid models

The 5 CAESAR models in VEGA are hybrid models



Learning methods

Hybrid models are planned since their beginning to be within one single system

Learning methods takes pre-existing models, integrate them, and finds the best way to assemble them, ideally using a test set for this purpose.

The test set has to contain new substances, never used by any of the pre-existing models. This is often very difficult.

Expert-based methods

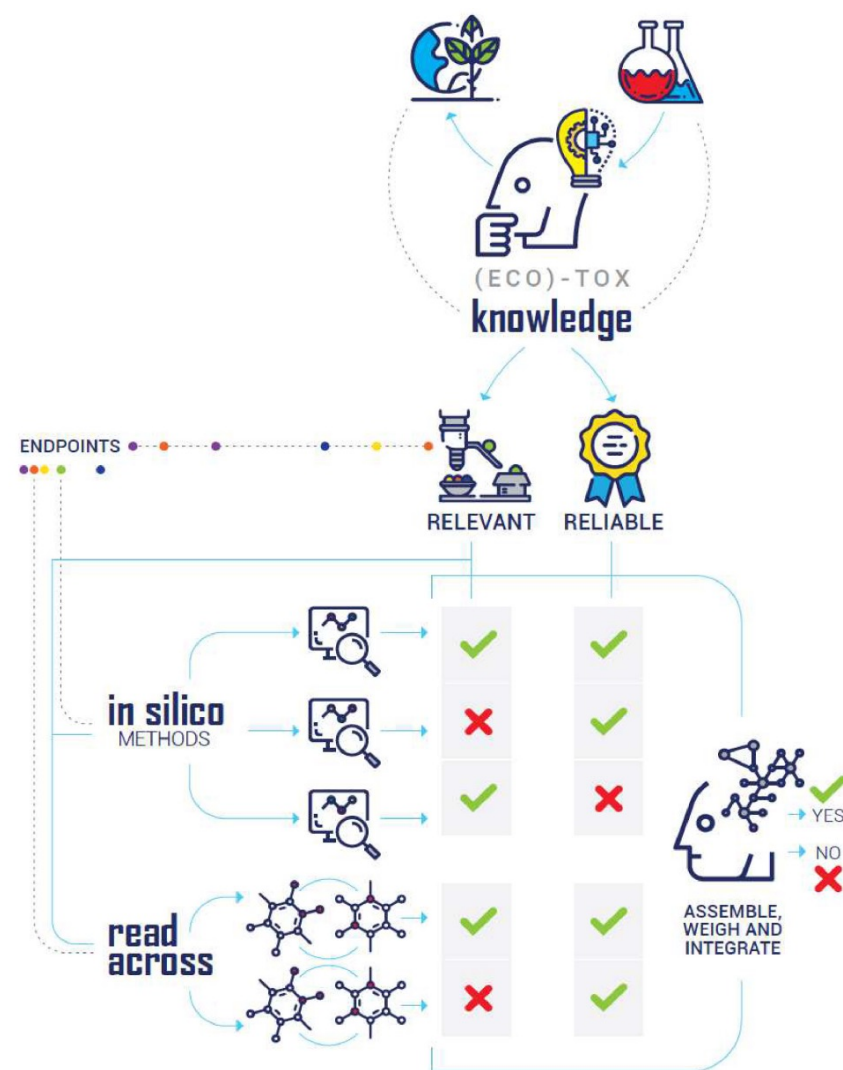
Experts may identify a preferred way to integrate results.

Pragmatic approach.

Often combining some criteria for reasoning, and introducing thresholds, and conservative assumptions.

Thus, the criteria are not only statistical. They should be declared.

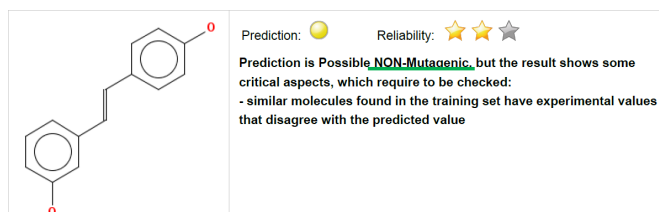
Integrating in silico and read-across








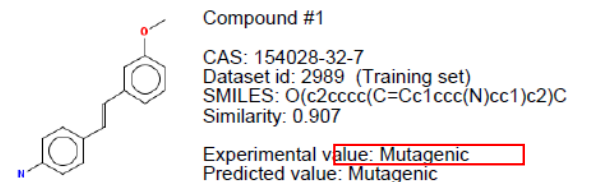
Use all lines of evidence

1. VEGA in silico models
2. Read-across
3. Reasoning
 - Check agreement

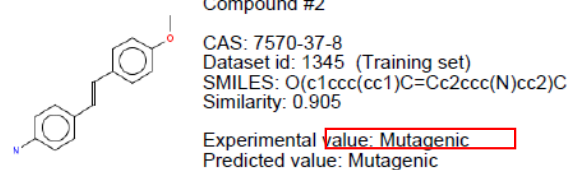
ADI concordance



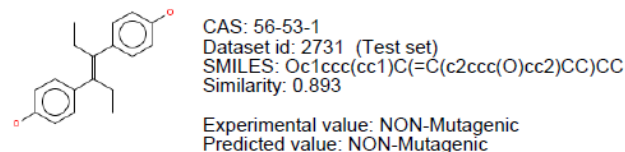
	Global AD Index AD index = 0.719 Explanation: the predicted compound could be out of the Applicability Domain of the model.
	Similar molecules with known experimental value Similarity index = 0.901 Explanation: strongly similar compounds with known experimental value in the training set have been found.
	Accuracy of prediction for similar molecules Accuracy index = 1 Explanation: accuracy of prediction for similar molecules found in the training set is good.
	Concordance for similar molecules Concordance index = 0.33 Explanation: similar molecules found in the training set have experimental values that disagree with the predicted value.
	Atom Centered Fragments similarity check ACF index = 1 Explanation: all atom centered fragment of the compound have been found in the compounds of the training set.



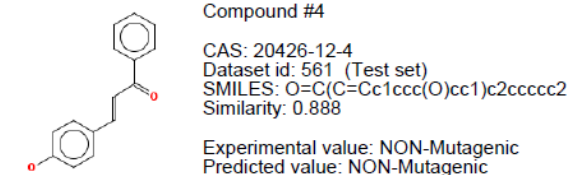
Alerts (not found in the target): SM44; SM104



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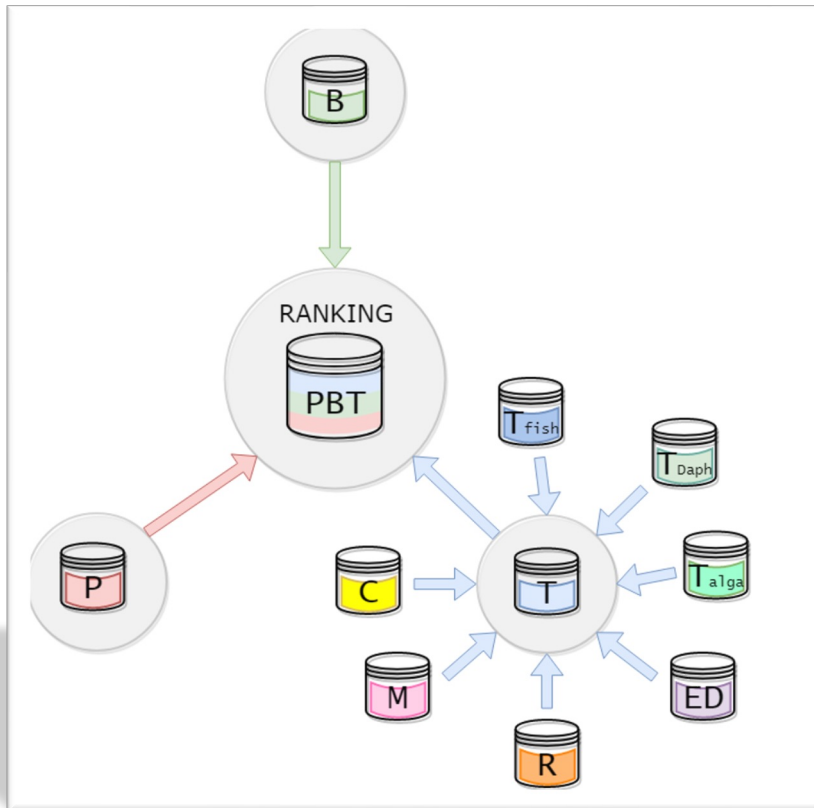


Alerts (not found in the target): SM158



Alerts (not found in the target): SM158; SM172

Integrating multiple endpoints/pathways. The JANUS example

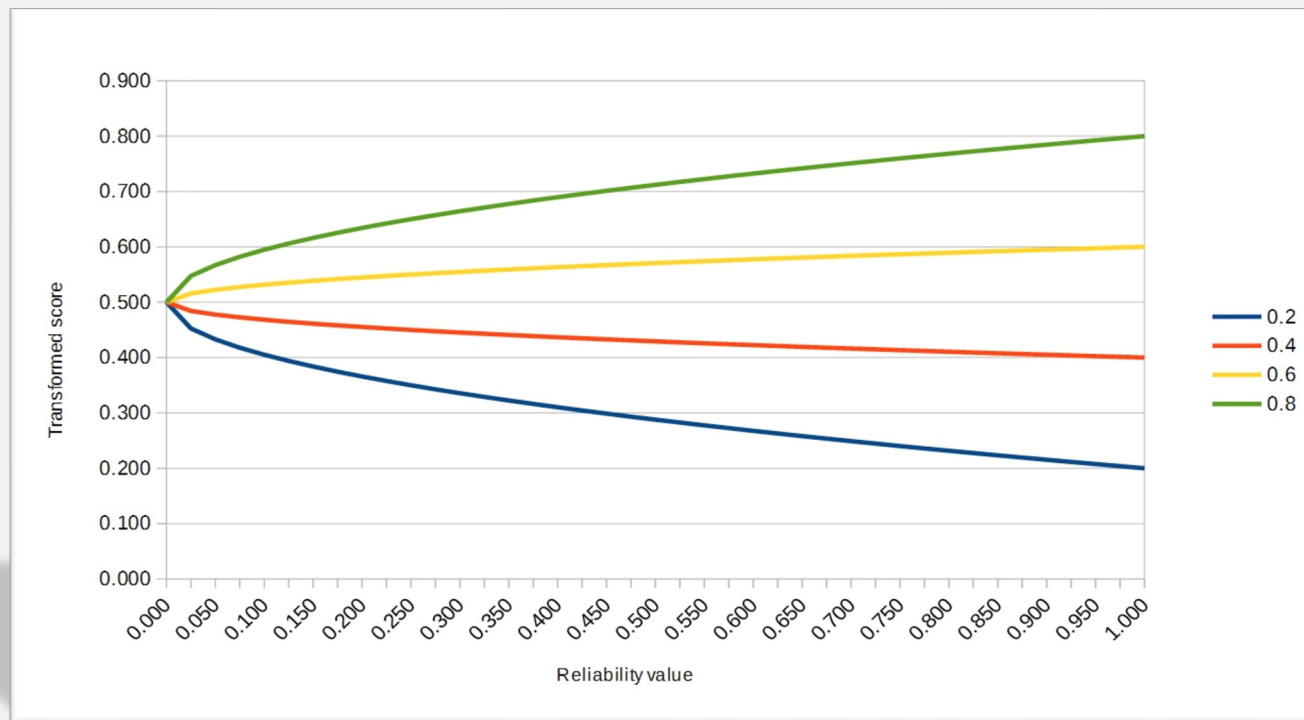


One single platform covering multiple endpoints, using 48 separate in silico models for CMR, PBT and ED (parental and degradation products).
Done for German UBA

www.vegahub.eu

Uncertainty and effect

Example: how the score (on the Y axis) changes depending on the reliability value (X axis) for four example property's value (0.2, 0.4, 0.6 and 0.8)



Ranking

Molecule identification			LogP		PERSISTENCE		BCF		TOXICITY		Ranking		
d/No.	CAS	SMILES	Remarks	Prediction	Reliability	Prediction	Reliability	Prediction	Reliability	Prediction	Reliability	Des(PBT)	Des(PB)
1	371-067-1	CC1=CC=C(C=C1)C2=CC=CC=C2	benzene	0.03	0.9	0.03	0.9	0.000353	0.9	0.310	0.9	0.310	0.311
2	128-03-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
3	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
4	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
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43	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
44	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
45	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
46	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
47	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
48	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
49	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
50	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
51	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
52	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
53	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
54	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
55	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
56	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
57	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
58	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
59	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
60	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
61	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
62	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
63	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
64	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
65	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
66	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
67	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
68	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
69	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
70	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
71	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
72	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
73	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
74	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
75	100-101-0	CC1=CC=C(C=C1)C2=CC=CC=C2C3=CC=CC=C3C4=CC=CC=C4	benzene	0.10	0.9	0.10	0.9	0.001050	0.9	0.319	0.9	0.319	0.312
76	100-101-0</												

JANUS. Details for mutagenicity

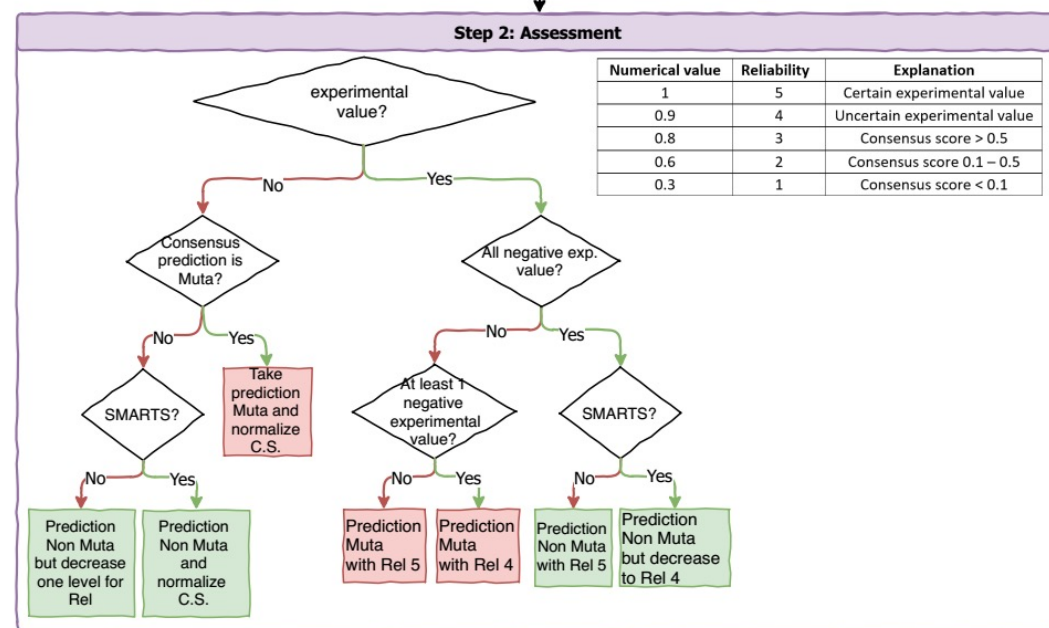
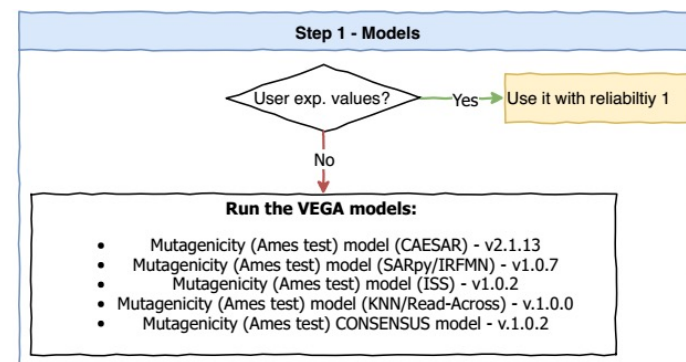
✓ Only in classification

- Based on **4 qualitative models** + a **consensus** model

✓ Metabolism SMARTS (5 SMARTS associated to mutagenicity)

- Used **only with Non-Muta** values (if matched → reliability reduce)
- Reliability based on the output** and the **consensus score**

Numeric value	Reliability	Explanation
1	5	Certain exp. values
0.9	4	Uncertain exp. values
0.8	3	Consensus score > 0.5
0.6	2	Consensus score 0.1 – 0.5
0.3	1	Consensus score < 0.1



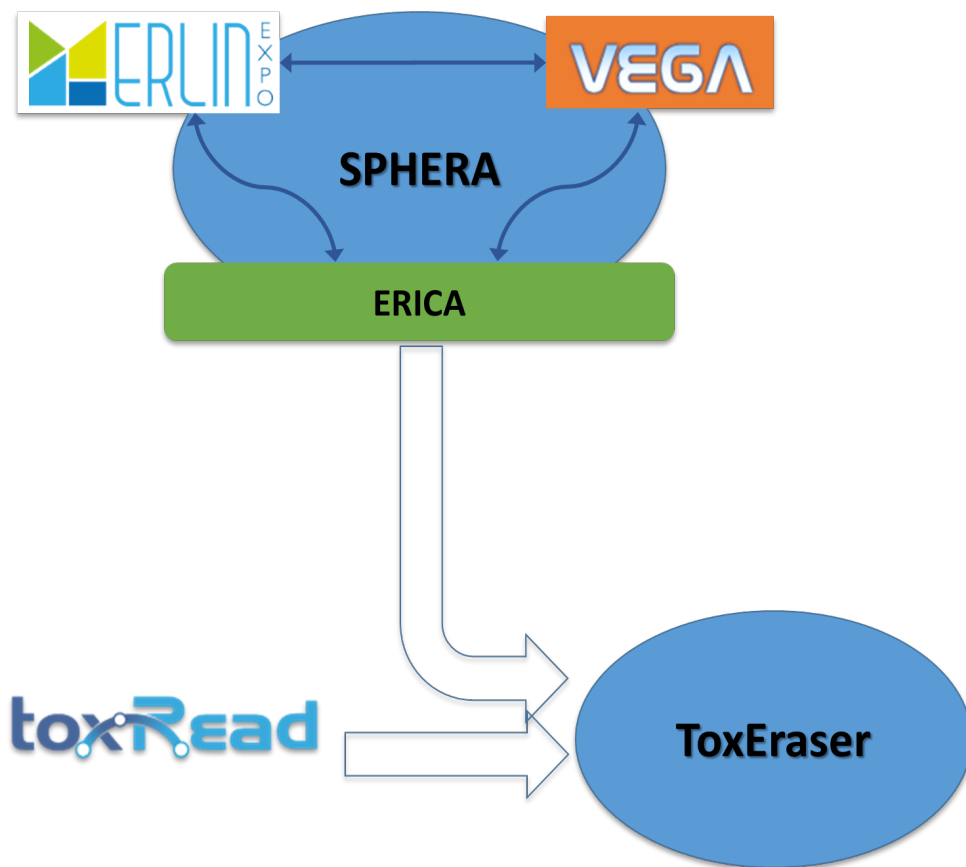
JANUS. One score

Janus Result

NUMBER OF COMPOUNDS 3

☒ PBT ☐ CMR ☐ ED ☒ PARTIAL SCORES ☒ FINAL SCORES

	No. ▼	Metabolite	Id	SMILES	Label	P	rel.	score	B [log(L/kg)]	rel.	score	T [mg/l]	rel.	score	Score(vPvB)	Score(SVHC)	Score(PBT)
Q	1		Molecule 1	OC(OC)C	PBT-CMRE	nP	0.45	0.211	0.16	0.9	0.078	2.18	0.65	0.264	0.129	0.853	0.242
Q	2	Metabolite	Molecule 1 [M-01]	O=CC	PBT-CMRE	nP	0.5	0.243	0.31	0.5	0.191	1.08	0.7	0.301	0.215	0.523	0.291
Q	3	Metabolite	Molecule 1 [M-02]	OC	PBT-CMRE	nP	0.8	0.174	0.15	0.75	0.114	0.78	0.7	0.325	0.141	0.875	0.26



Integrating hazard + exposure

Integrating **VEGA**, **ToxRead**, **MERLIN-Expo**, and **ERICA** in a **platform for risk assessment and substitution** of risky substances

1. Identification of the risky substances
 2. Identification of possible substitutes
- Application to 6 case studies*

LIFE VERMEER project - Case studies



Food Contact Materials



Biocides



Oil fractions



Solvents



Dispersants



Cosmetics



VERMEER-Cosmolife - input



- The user is asked to provide the information regarding the ingredient, its concentration and the product type
- The software allows to add single or multiple ingredients
- Ingredients can be entered using **INCI**, **CAS** or **SMILES**

The screenshot shows the main interface of the Sphera Cosmolife software. At the top, there's a title bar "Sphera Cosmolife - v. 0.20". Below it, a text field for "Directory for output:" is set to "C:\Users\GSelvestrel\Documents" with a "Select directory" button. The main area is titled "List of ingredients" and contains a table with columns "Id" and "Concentration". Below the table are icons for adding (+), editing (pencil), deleting (-), saving (floppy disk), printing (printer), and exiting (X). A "Product Type" dropdown menu is open, showing a list of product types: Shower gel, Shampoo, Hair styling products (leave-on), Hair styling products (rinse-off), Body lotion, Face cream, Face cream (applied on neck), and Face cream (applied on back of neck). A "Run calculation" button is visible. At the bottom left, an "Application log" shows the status: "* Initializing core..." and "* Ready.".

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The "New ingredient" dialog box is shown, allowing users to add a new ingredient. It has a title bar "New ingredient" with a close button (X). The form contains several input fields: "Ingredient Id" (empty), "SMILES" (containing OCCOc1ccccc1), "SMILES (Neutral form)" (containing OCCOc1ccccc1), "CAS" (containing 122-99-6), "INCI" (containing PHENOXYETHANOL), and "Concentration (%)" (containing 0.5). There are magnifying glass icons next to the CAS and INCI fields. At the bottom, there are "Cancel" and "Add ingredient" buttons.

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VERMEER-Cosmolife - summary



The software provides a preliminary output table with a summary of the hazard and exposure features of the ingredients

Processed product

Product type: **Body lotion**

Ingredients:

	Ingredient Id	CAS	INCI	Conc. %	Annex	Mutagenicity	Skin Sensitization	Dermal abs.	MoS	TTC
Details	DISODIUM EDTA	139-33-3	DISODIUM EDTA	0.05	-	NON-Mutagen (EXPERIMENTAL value)	Sensitizer (low reliability)	10%	162337.66	0.0015 mg/kg bw/day
Details	GLYCERIN	56-81-5	GLYCERIN	5.0	-	NON-Mutagen (EXPERIMENTAL value)	NON-Sensitizer (EXPERIMENTAL value)	80%	52.37	0.03 mg/kg bw/day
Details	POTASSIUM BENZOATE	582-25-2	POTASSIUM BENZOATE	2.0	V	NON-Mutagen (EXPERIMENTAL value)	NON-Sensitizer (EXPERIMENTAL value)	80%	14.03	0.03 mg/kg bw/day
Details	EUGENOL	97-53-0	EUGENOL	0.001	III	NON-Mutagen (EXPERIMENTAL value)	Sensitizer (EXPERIMENTAL value)	80%	233360.39	0.03 mg/kg bw/day
Details	GERANIOL	106-24-1	GERANIOL	0.001	III	NON-Mutagen (EXPERIMENTAL value)	Sensitizer (EXPERIMENTAL value)	40%	158441.56	0.03 mg/kg bw/day

VERMEER-Cosmolife – internal exposure

The software calculates the SED following 3 different approaches:

Absorption=100%
(Oral/inhalation)

Absorption=50%
(Dermal; defined
by SCCS)

Kroes Approach
(Refined
approach)



1. Implementation in VEGA of two models that allow predicting **Kp (Skin Permeability Coefficient)**

Example: **Potts and Guy: $\log Kp = 0,71 \log(K_{o/w}) - 0.0061 MW - 2.7$ (cm/h)**

2. Once predicted Kp, we obtain **Jmax (Maximum flux)**

$$Jmax = Kp * C_{water, sat} \text{ (mg/cm}^2\text{/h)}$$

3. Based on Jmax, Kroes proposed three default exposure values:

$$\%A = \begin{cases} 10\% & \text{if } Jmax \leq 0.1 \mu\text{g/cm}^2\text{/h} \\ 40\% & \text{if } 0.1 < Jmax \leq 10 \mu\text{g/cm}^2\text{/h} \\ 80\% & \text{if } Jmax > 10 \mu\text{g/cm}^2\text{/h} \end{cases}$$

VERMEER-Cosmolife - hazard

The software includes the HAZARD IDENTIFICATION, considering different toxicological endpoints, including NOAEL

4. Hazard identification

Mutagenicity - Ames test (Janus workflow prediction)	NON-Mutagen (EXPERIMENTAL value)
In vitro micronucleus assay (IRFMN model prediction)	Inactive (EXPERIMENTAL value)
Chromosomal aberration test (Coral model prediction)	Inactive (good reliability)
Skin sensitization (Consensus of Caesar and JRC models)	NON-Sensitizer (EXPERIMENTAL value)
Skin sensitization (Caesar model prediction)	NON-Sensitizer (good reliability)
Skin sensitization (IRFMN/JRC model prediction)	NON-Sensitizer (EXPERIMENTAL value)

Output Example

Other endpoints, such as Carcinogenicity, Reproductive/Developmental toxicity, Endocrine Disruptors, Skin Irritation and Eye Irritation and Acute Toxicity will be included soon

VERMEER-Cosmolife - MoS

The software provides the Risk Characterization, considering the process proposed within the SCCS Notes of Guidance.

For cosmetics, the focus is on systemic effects and a MoS (Margin of Safety) is calculated, according to the formula:

$$MoS = \frac{POD \text{ (Point of Departure)}}{SED \text{ (Systemic Exposure Dose)}}$$

MoS > 100



6. Risk characterization

Output Example

MoS - Margin of Safety with 100% absorption	101.46
MoS - Margin of Safety with 50% absorption	202.92
MoS - Margin of Safety with 10% absorption (from Kroes thresholds)	1014.61

ToxEraser – planning substitution



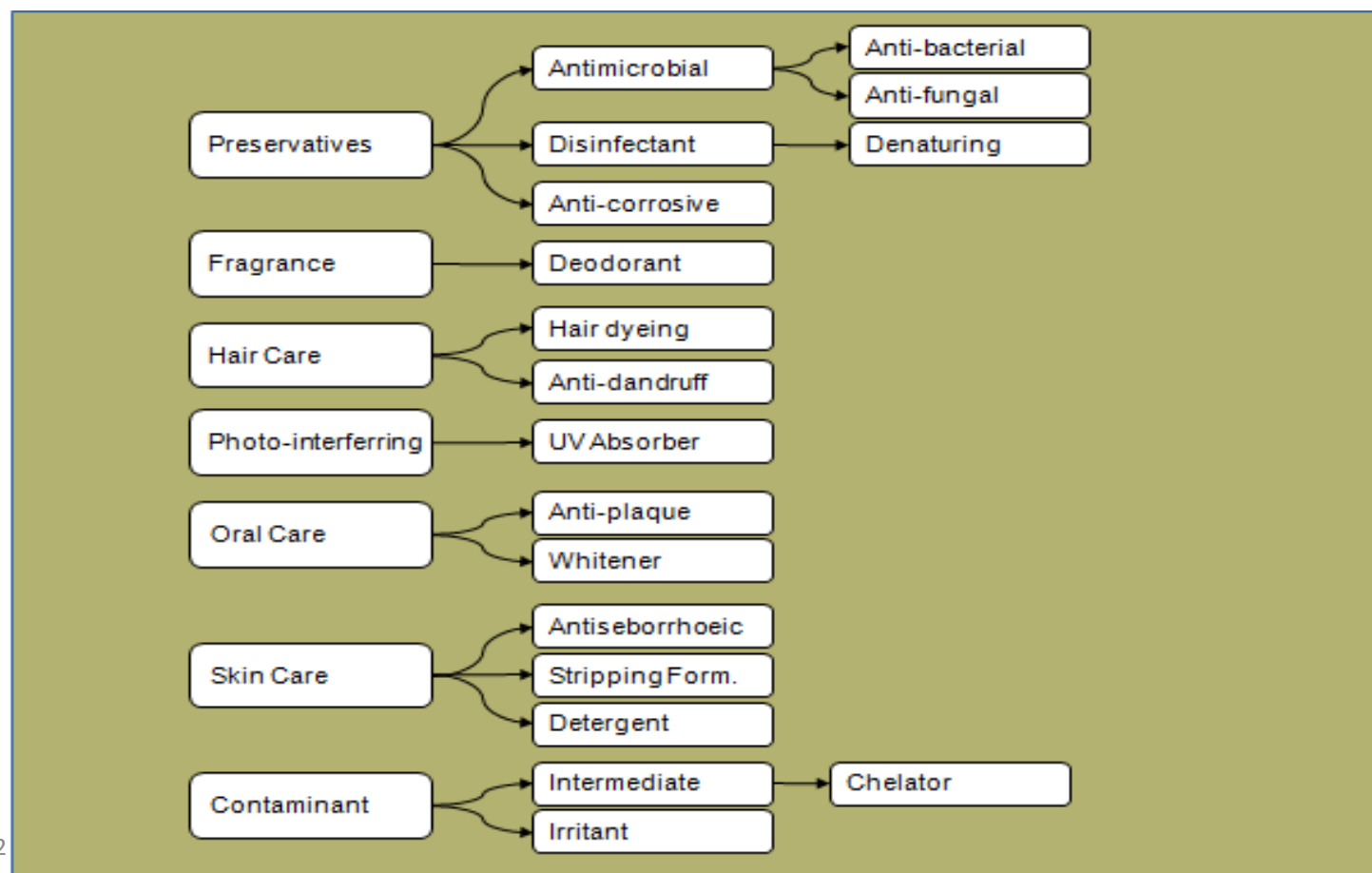
Systematic retrieval of information, concerning:

- safety
- functional uses
- similarity (read-across)

ToxEraser – functional use



Cosmetics according to their functional uses: a hierarchical ontology



www.vegahub.eu

Welcome to the VEGA HUB

Offering a family of tools to evaluate chemical hazard: VEGA, ToxRead, ToxWeight, ToxDelta, and JANUS.

VEGA is the QSAR software with tens of models for individual properties.

ToxDelta

ToxWeight



Do you need assistance for a property prediction ?

CONTACT US

ToxRead

VEGA

Life-Sphera

ToxEraser


JANUS



Unique sw environment

Chemical features and tools associated to different aspects?

Need of a single conceptual scheme:

- *Functional properties*
 - Phys-chem properties
 - Toxicological properties
 - Environmental properties
- 
- Industry
- Same for authorities
and industries

Conclusions

- Global challenges, broad view
 - Multitask methodologies
 - Set up priorities
 - Players: politics, economy, science
 - Joining efforts to get higher targets
-
- In silico models for
 1. *new paradigm: holistic*
 2. *prioritization*
 3. *models for the most relevant endpoints*
 4. *safer substances*

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