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Science

Use of Computational Approaches for Genotoxicity Assessment of Pesticides

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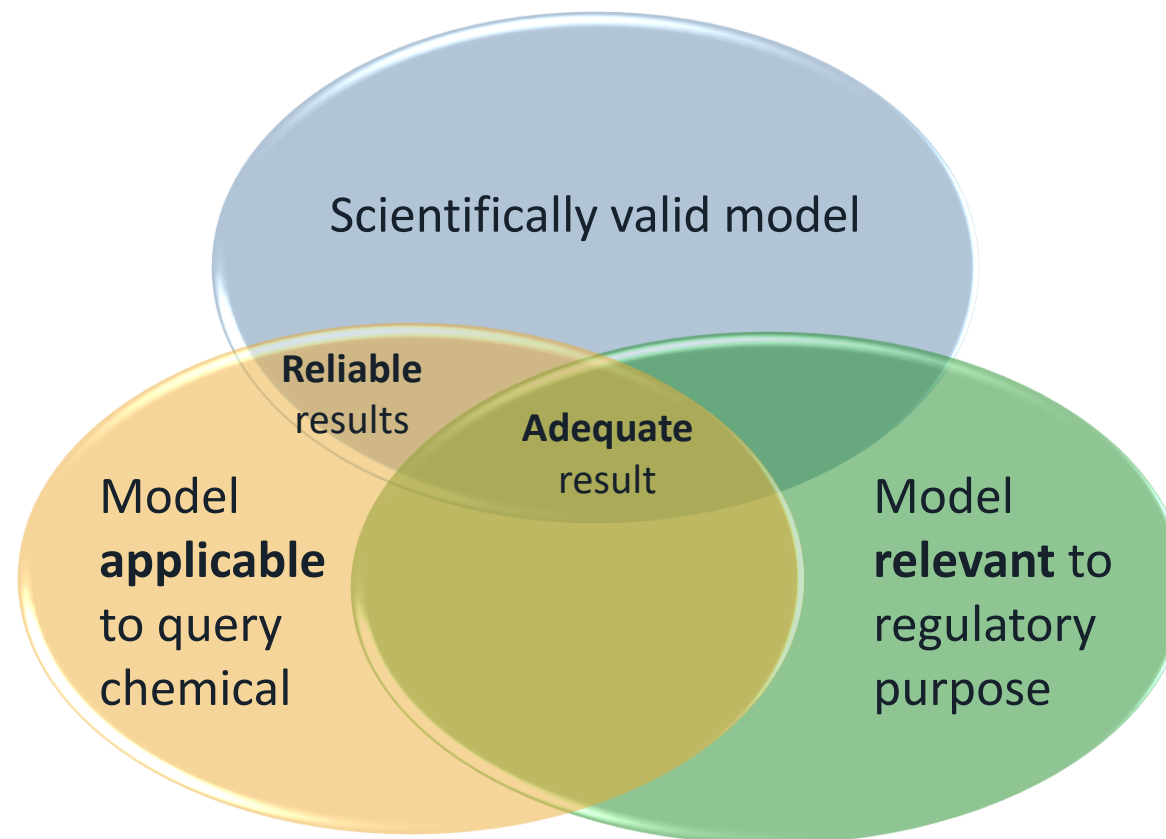
Review of Literature

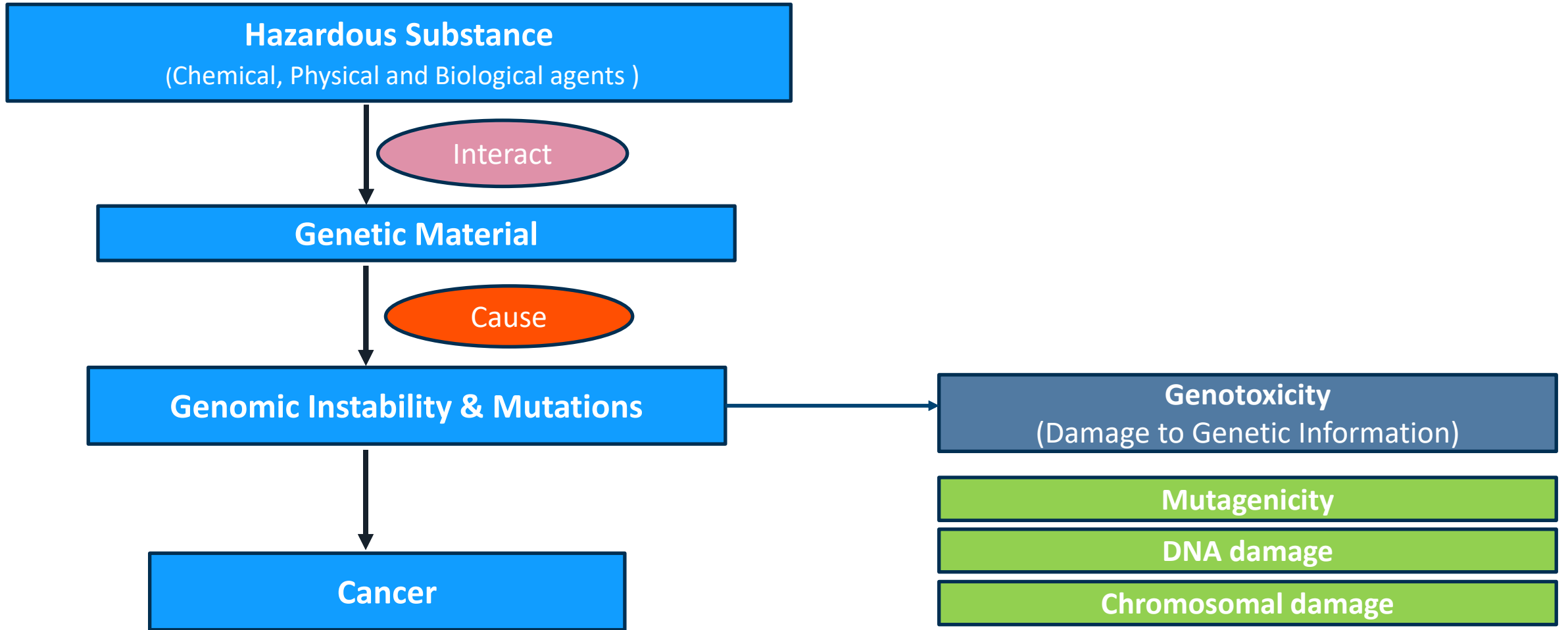
Demo of various *in silico* tools

Our work

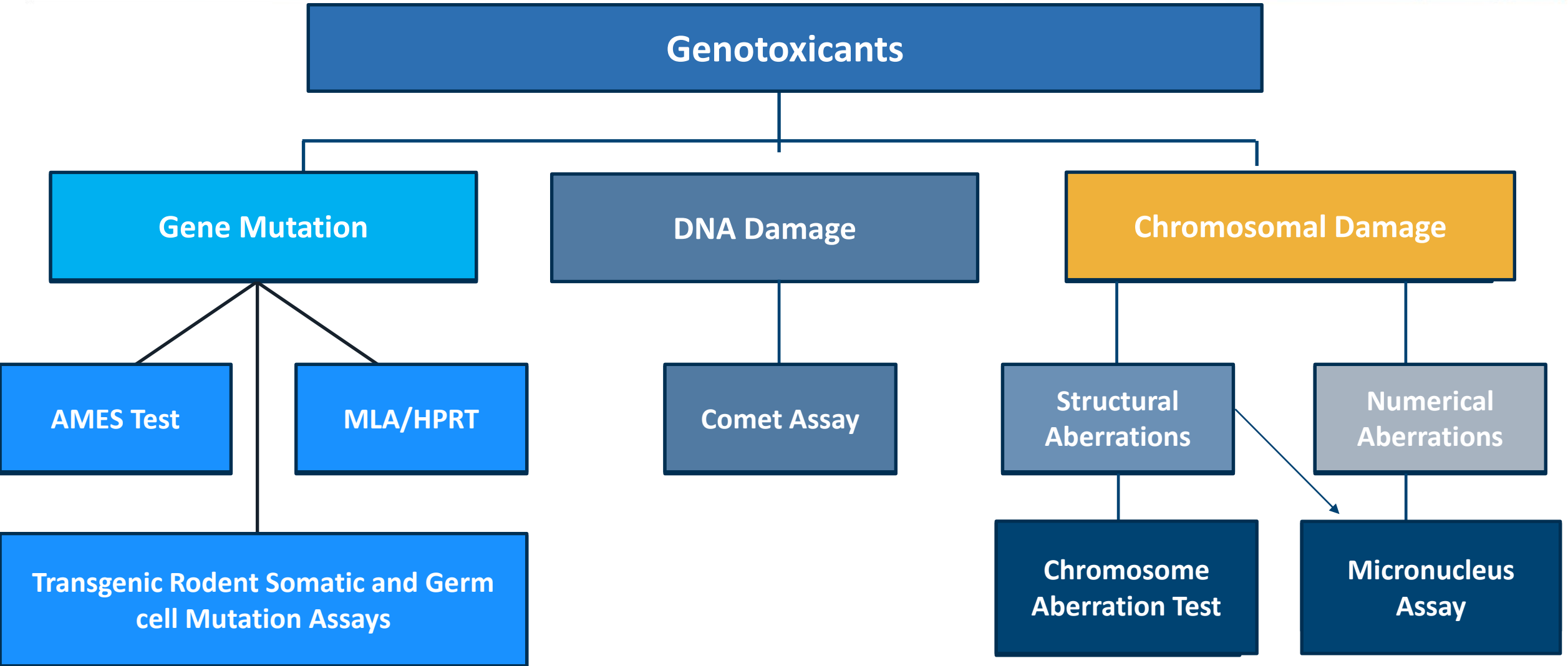
Opportunities for Collaboration

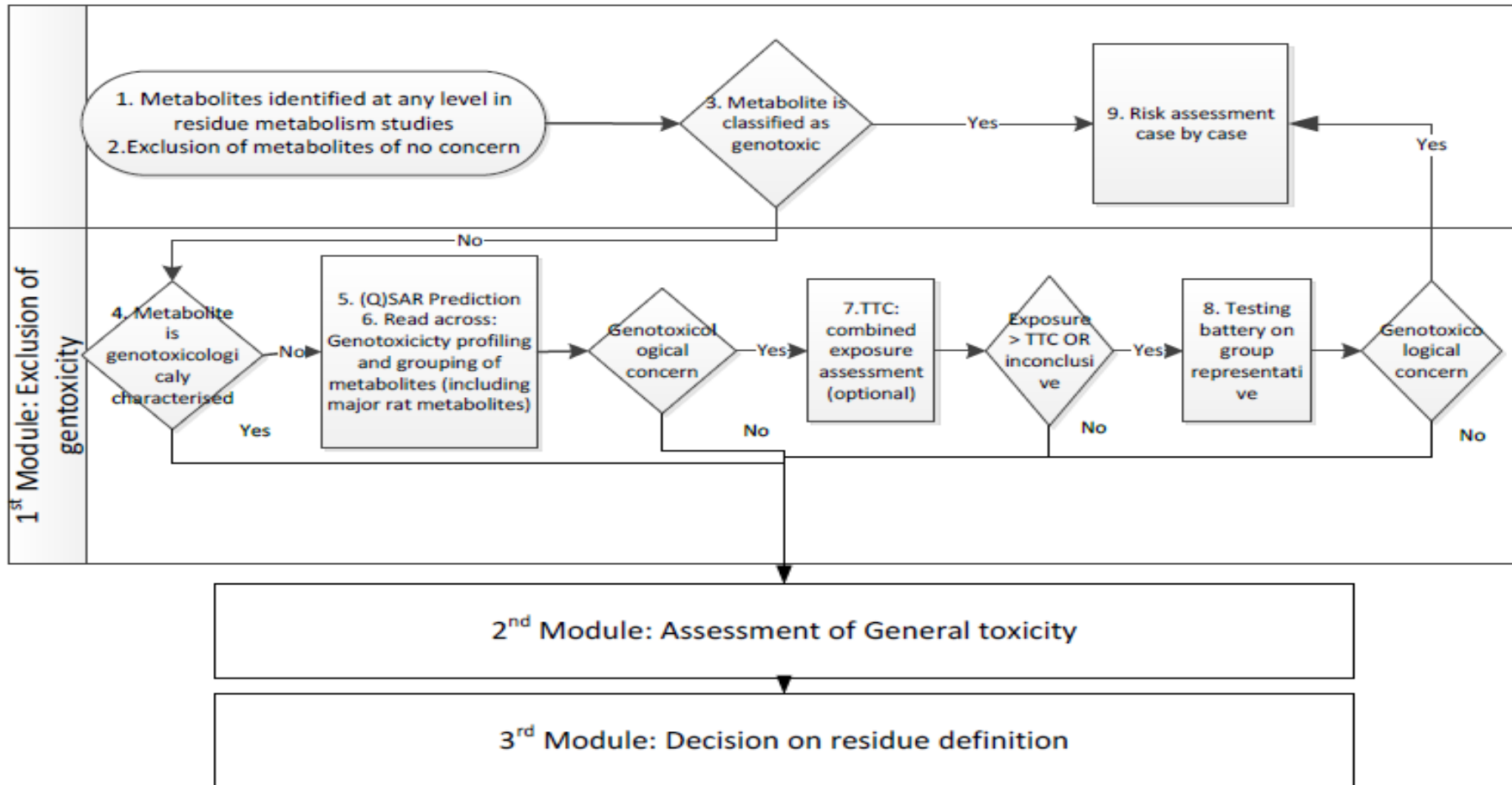
Most comprehensive guidance currently available for applying QSAR analysis is provided in the REACH guidance on Information Requirements and Chemical Safety Assessment (ECHA, 2008).





Genotoxicity Testing





Module 1 exclusion of genotoxicity

SCIENTIFIC OPINION

Guidance on the establishment of the residue definition for dietary risk assessment¹EFSA Panel on Plant Protection Products and their Residues (PPR)^{2,3}

European Food Safety Authority (EFSA), Parma, Italy

- Genotoxicity assessment should be assisted by application of (Q)SAR and read across of metabolites.
- Use of computational models for predictions of genotoxicity should not be based on the use of any single model alone, but on a “weight of evidence” approach including all available information provided by the models (e.g. applicability domain, proposed mechanistic information, prediction for the similar substance).
- To maximize the sensitivity and specificity of the prediction, at least two independent (Q)SAR models, where possible, should be applied for each genotoxicity endpoint, including both knowledge based and statistical based models.

SCIENTIFIC OPINION

Guidance on the establishment of the residue definition for dietary risk assessment¹EFSA Panel on Plant Protection Products and their Residues (PPR)^{2,3}

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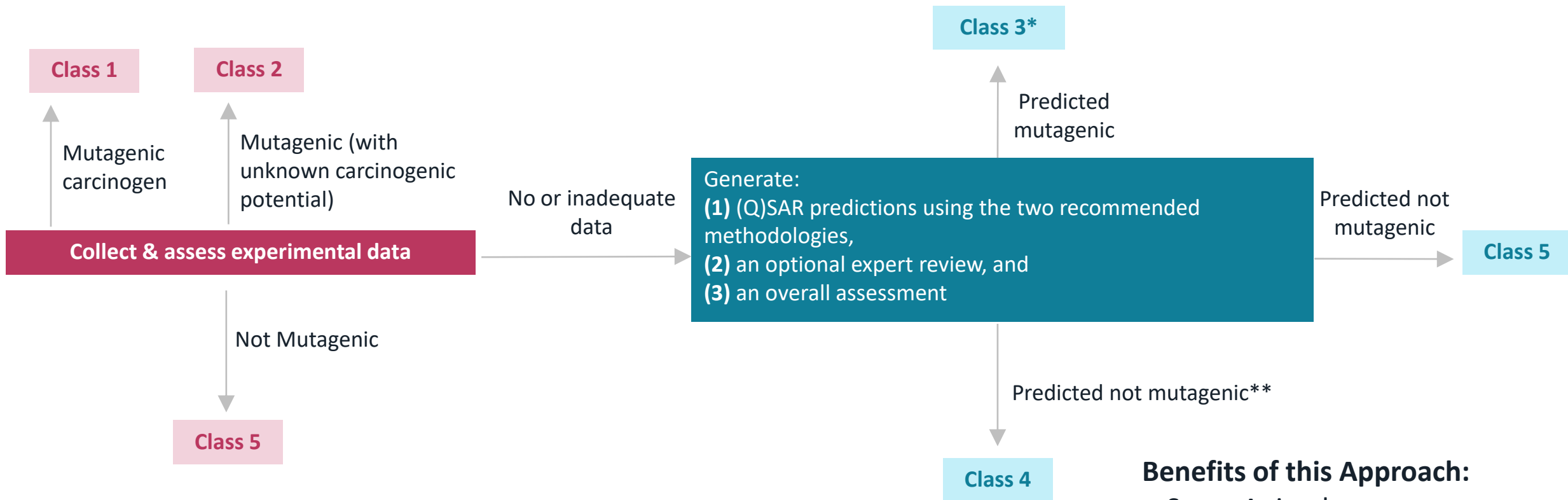
Case Study: Study of genotoxicity potential of Isoproturon and 12 metabolites using QSAR & Read-across

a) In silico:

- In order to predict the genotoxic potential (gene mutation and chromosomal aberrations) of the minor rat and plant specific metabolites, four models have been applied: VEGA software, DEREK Nexus, and Toxtree.

b) Read-across:

OECD Toolbox used



**Or perform an Ames test*

***Based on a shared alert with a known negative*

Benefits of this Approach:

- Saves Animals
- Saves Time
- Saves Money

Biological evaluation of medical devices —

Part 17:

Toxicological risk assessment of medical device constituents

- Use of *in silico* analysis to predict nature of harms is an example when a chemical specific POD is not available.
- When nature of harm is not understood, a computer-based model (also known as *in silico* analysis) can be used to predict the nature of the harm to health for the identified constituent.

The use of a combination of rule-based and statistical-based (Q)SAR software is one of the preferred options proposed in the EFSA guidance document.

There are several (Q)SAR software tools available, which allow to predict the DNA reactivity of a substance.

Potential applicable alerts are:

- in vitro mutagenicity (Ames test) alerts by ISS (ToxTree)
- mutagenicity in vitro (Sarah Nexus)
- bacterial mutagenicity OECD 471 (CaseUltra)
- bacterial mutation alerts (Leadscope)

In addition, Annex III of REACH, available on the ECHA website, consists of a compilation of (Q)SAR predicted toxicities for some 33.000 substances, including genotoxicity or carcinogenicity alerts, where applicable



CompTox Chemicals Dashboard-EPA: <https://comptox.epa.gov/dashboard/>



Test compound: Endosulfan (SMILES: C1C2C(COS(=O)O1)C3(C(=C(C2(C3(Cl)Cl)Cl)Cl)Cl)Cl)



- *In silico* evaluation of several agrochemicals including inorganic metals, carbamates, chlorinated hydrocarbons, pyrethroids, organophosphate insecticides, fungicides, herbicides, fumigants, nematicides and solvents.
- Three softwares viz. QSAR Toolbox by OECD, Toxtree and TEST by US EPA were used.
- Selection of approximately 60 agrochemicals for *in silico* analysis were based on published data for various endpoints viz. (i) Ames, (ii) *in vitro* mammalian cell gene mutation, (iii) *in vitro* micronucleus, (iv) *in vitro* chromosomal aberration, (v) *in vivo* micronucleus, (vi) *in vivo* chromosomal aberration, (vii) rodent carcinogenicity and (viii) skin sensitization.
- QSAR Toolbox, Toxtree and TEST (for Ames only) had accuracy of 80%, 66% and 77%, respectively. Additionally, QSAR Toolbox and Toxtree had an accuracy of 90% and 69% for carcinogenicity endpoints, respectively.



Predicting genotoxicity and carcinogenicity of drugs and chemicals using OECD QSAR toolbox, Derek Nexus and TEST



EVALUATION OF PHOTOTOXIC POTENTIAL OF SEVEN SOLVENTS USING *IN SILICO* DEREK AND *IN VITRO* 3T3 NRU METHOD.

M. Shameer P¹, V. Ahuja¹, M. Krishnappa¹, D. Venkataraman¹



Toxicology Letters

Volume 368, Supplement, 1 September 2022, Page S286



Late breaking abstracts

LP-07 *In silico* analysis using Derek and Sarah along with ICH M7 expert review accurately predicts Mutagenicity and Carcinogenicity of Nitrosamines

V. Ahuja, M. Krishnappa

<https://www.sciencedirect.com/science/article/abs/pii/S0378427422016770>



Prediction of Mutagenicity, Genotoxicity, and Carcinogenicity of Drugs and Chemicals Using Derek Nexus

The Toxicologist
Supplement to *Toxicological Sciences*



Predicting skin sensitization of agrochemicals using OECD QSAR toolbox, Derek Nexus, ToxTree, PredSkin and SkinSensPred

Gowrav Adiga.P, Deepa Venkataramulu, Mohan Krishnappa and Varun Ahuja



Toxicology Letters

Volume 350, Supplement, September 2021, Page S250



In silico toxicity prediction using Derek Nexus® for skin sensitization, phototoxicity, hepatotoxicity and *in vitro* hERG inhibition

V. Ahuja¹, M. Krishnappa¹, H. Kandarova²

<https://www.sciencedirect.com/science/article/abs/pii/S0378427421008171>



Questions