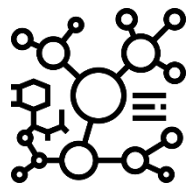


VERA: VIRTUAL EXTENSIVE READ-ACROSS

A new tool for automated read-across

CASE STUDY:
*using the new VERA tool, for automated read-across
assessment of carcinogenicity*

VERA: the new concept of similarity



**STRUCTURAL
SIMILARITY**

VEGA

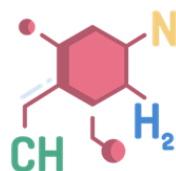
**Endpoint
Specific**

SAs

**TOXICOLOGICAL
SIMILARITY**



Based on literature searching and
implemented as SMARTS



**GROUPING
SIMILARITY**

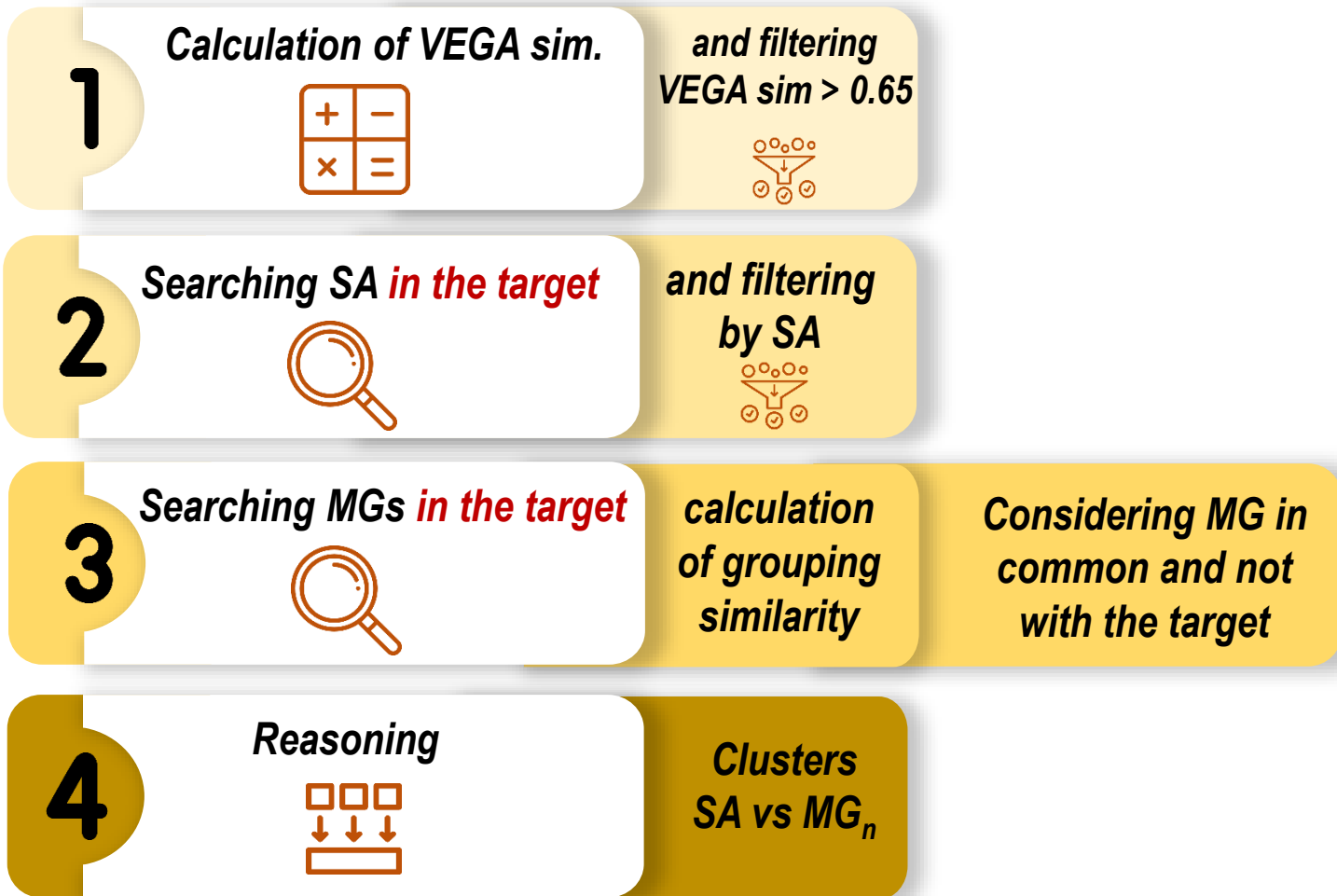
MGs

From RdKit and other manually
implemented as SMARTS

General Workflow of VERA

VERA

VIRTUAL EXTENSIVE READ-ACROSS



Target Molecule

VEGA similarity > 0.65

Similarity	SMILES	Experimental value
8	0.938	CC(=O)OCC1=CC=CC=C1
1	0.915	CC(=O)OCC1=CC=C(C=C1)N
2	0.914	CC(=O)OCC1=CC=C(C=C1)C
3	0.910	CC(=O)OCC1=CC=C(C=C1)O
4	0.908	CC(=O)OCC1=CC=C(C=C1)C(=O)O
...
483	0.891	CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=CC=C1
484	0.890	CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C
485	0.890	CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)O
486	0.890	CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C(=O)O
487	0.890	CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=CC=C1

Searching SA in target

Filter similar compounds with SA

SMILES	VEGA_sim	Experimental value	grp_sim
CC(=O)OCC1=CC=CC=C1	0.875	1	0.887
CC(=O)OCC1=CC=C(C=C1)N	0.890	1	0.799
CC(=O)OCC1=CC=C(C=C1)C	0.878	1	0.795
CC(=O)OCC1=CC=C(C=C1)O	0.887	0	0.791
CC(=O)OCC1=CC=C(C=C1)C(=O)O	0.938	1	0.744
CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=CC=C1	0.914	1	0.744
CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C	0.894	1	0.744
CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)O	0.742	1	0.744
CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C(=O)O	0.848	1	0.723
CC(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=C(C=C1)C(=O)OCC1=CC=CC=C1	0.765	1	0.718

Searching MGs in similar compounds...
- calculation of grouping similarity

Searching MGs in target

10 most similar compounds

TOX TOX TOX NO TOX TOX
TOX TOX TOX TOX TOX

$$GSI = \frac{1+nMGs \in common}{2 \cdot ntargetMGs} - \frac{nMGs NOT \in common}{8 \cdot nsimilarMGs}$$

Endpoints

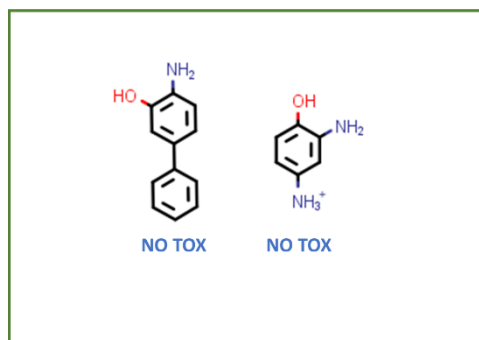


CARCINOGENICITY Classification



FISH ACUTE TOXICITY Continuous

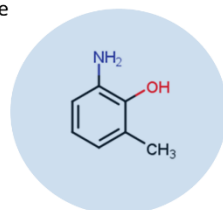
Grouping: SA + MG1



Cluster: INACTIVE

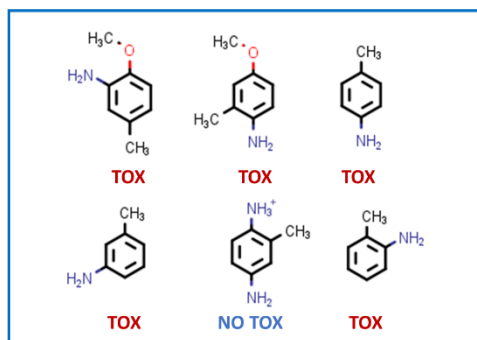
MG1: possible exception rule

VERA ASSESSMENT:



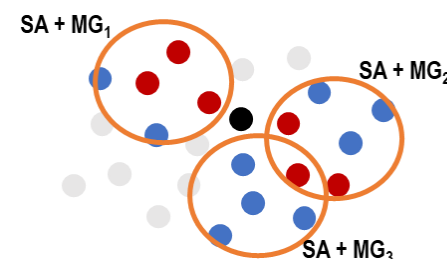
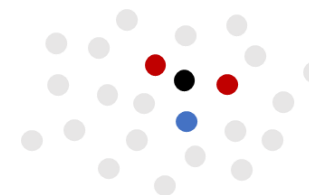
Not Toxic due to exception rules

Grouping: SA + MG2



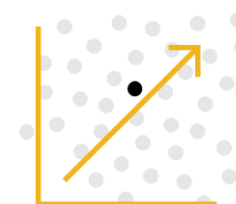
Cluster: ACTIVE

1 kNN model
If VEGA sim > 0.85



Grouping 2
If VEGA sim > 0.75

3 Local linear models














Publication

Open Access

Article

Virtual Extensive Read-Across: A New Open-Access Software for Chemical Read-Across and Its Application to the Carcinogenicity Assessment of Botanicals

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Published: 5 October 2022

Availability

VERA
VIRTUAL EXTENSIVE READ-ACROSS

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VERA

[Read more](#)

Description VERA, a new open-access software for chemical read-across

Within the EC-funded LIFE project CONCERT REACH, the new open-access software for chemical read-across VERA has been developed.


Last Release March 9, 2023



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VERA

VERA Virtual-Extensive-Read-Across is a new open-access software for chemical read-across.





Within the EC-funded LIFE project [CONCERT REACH](#), the new open-access software for chemical read-across VERA has been developed. Read-across applies the principle of similarity to identify the most similar substances to represent a given target substance in data-poor situations. However, differences between the target and the source substances exist. VERA (Virtual Extensive Read-Across) provides a means to assess the similarity between chemicals using structural alerts specific to the property, pre-defined molecular groups and structural similarity. The software finds the most similar compounds with a certain feature e.g. structural alerts and molecular groups and provides clusters of similar substances while comparing these similar substances within different clusters.

[Introduction](#)

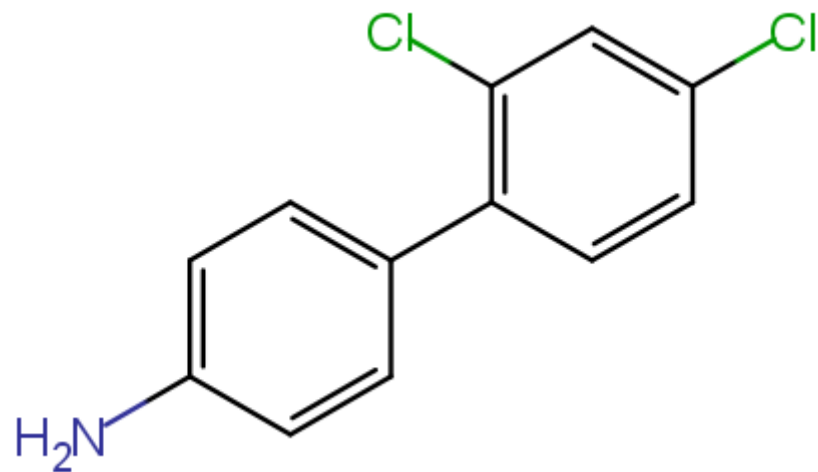
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[Screenshots](#)



CASE STUDY

CARCINOGENICITY



4-(2,4-dichlorophenyl)aniline

- Pharmaceutical precursor
- Pesticide precursor
- Starting material for synthesizing dyes
- Potentially building block for polymers