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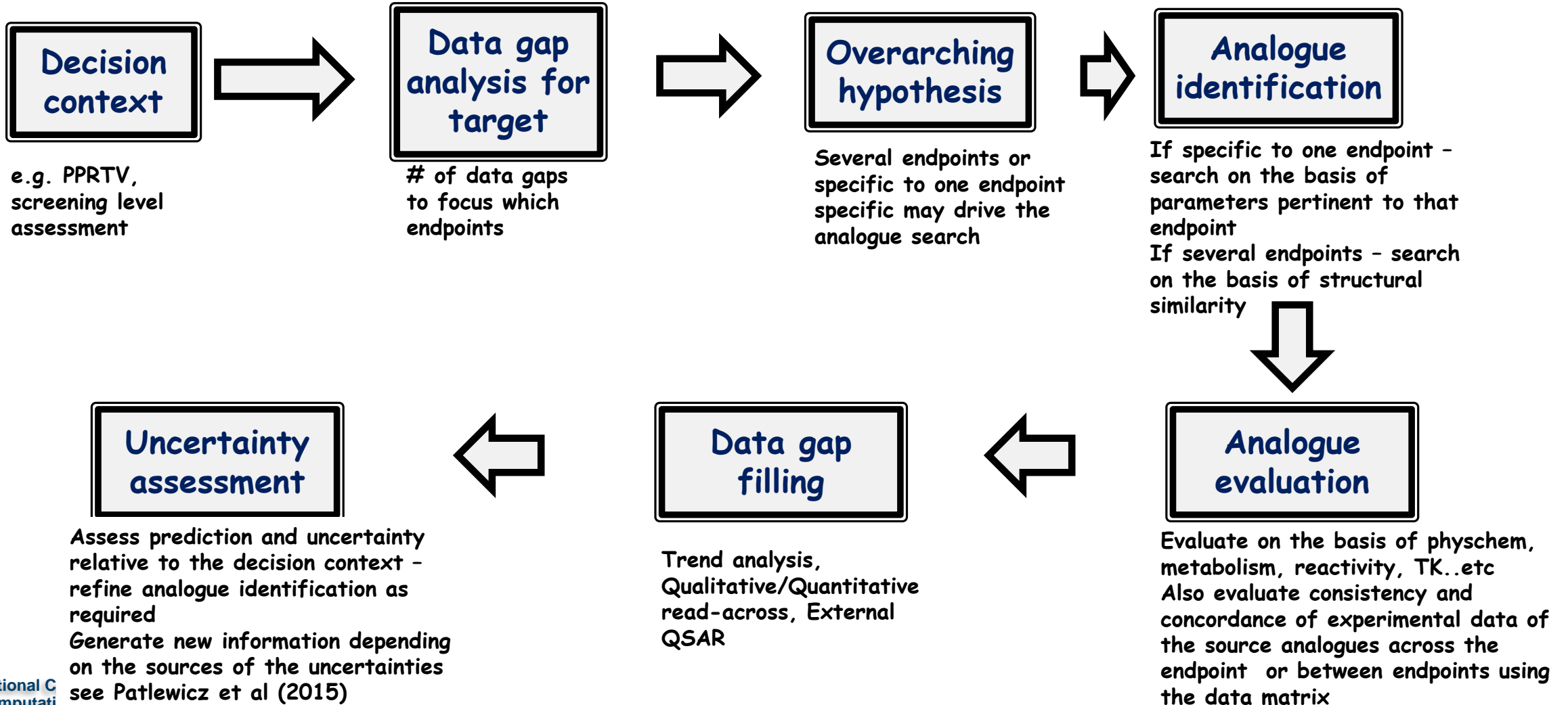
Outline

- Background, Concepts and Definitions
- Category workflow and selected tools for read-across
- Uncertainty assessment in read-across
- Quantifying Uncertainty & Assessing Performance of Read-Across
- From Research to Implementation
- Summary

Background & definitions

- Read-across describes one of the data gap filling techniques used within analogue and category approaches
- “Analogue approach” refers to grouping based on a very limited number of chemicals (e.g. target substance + source substance)
- “Category approach” is used when grouping is based on a more extensive range of analogues (e.g. 3 or more members)

Category Workflow



Selected Read-Across Tools

Tool	AIM	ToxMatch	AMBIT	OECD Toolbox	CBRA	ToxRead
Analogue identification	X	X	X	X	X	X
Analogue Evaluation	NA	X	X by other tools available	X	X	X For Ames & BCF
Data gap analysis	NA	X	X Data matrix can be exported	X Data matrix viewable	NA	NA
Data gap filling	NA	X	User driven	X	X	X
Uncertainty assessment	NA	NA	NA	X	NA	NA
Availability	Free	Free	Free	Free	Free	Free

Selected Read-Across Tools – Review

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Navigating through the minefield of read-across tools: A review of in silico tools for grouping



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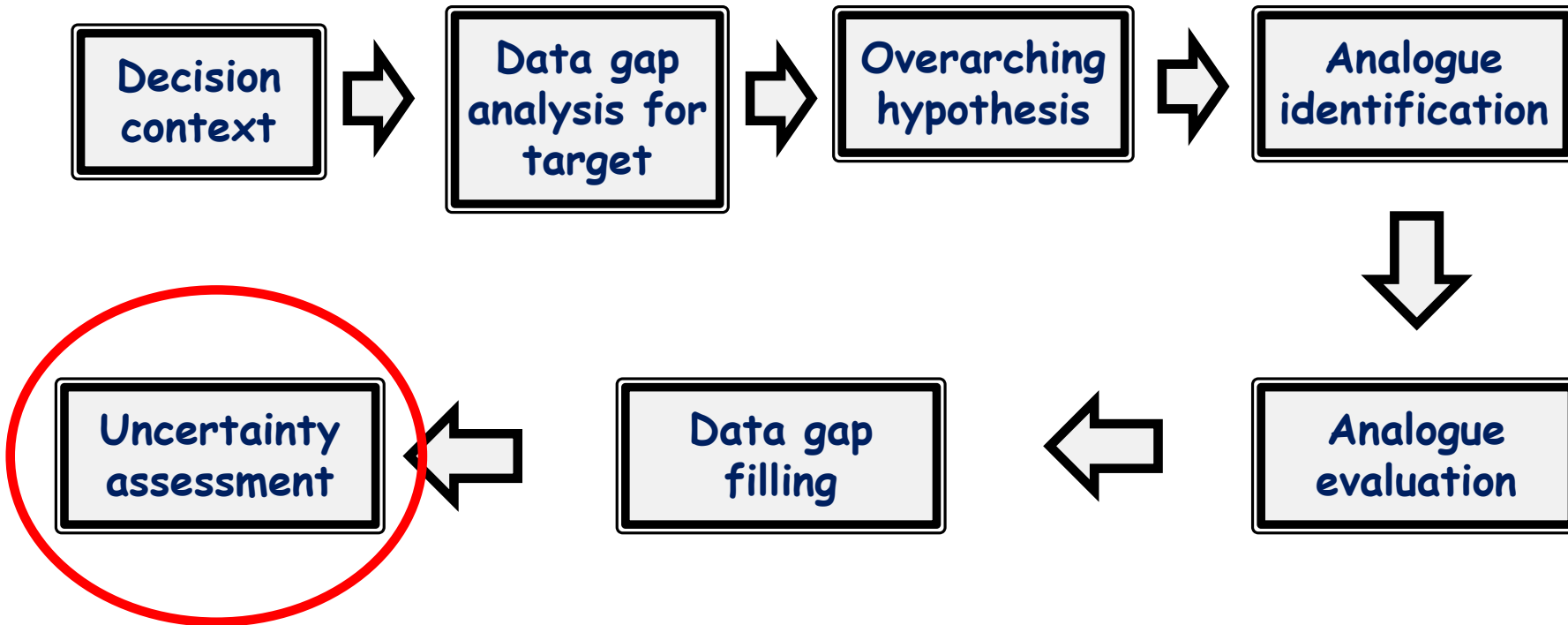
Nearest neighbor

ABSTRACT

Read-across is a popular data gap filling technique used within analogue and category approaches for regulatory purposes. In recent years there have been many efforts focused on the challenges involved in read-across development, its scientific justification and documentation. Tools have also been developed to facilitate read-across development and application. Here, we describe a number of publicly available read-across tools in the context of the category/analogue workflow and review their respective capabilities, strengths and weaknesses. No single tool addresses all aspects of the workflow. We highlight how the different tools complement each other and some of the opportunities for their further development to address the continued evolution of read-across.

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The Category Workflow



Sources of Uncertainty

- Analogue or category approach? (# analogues)
- Completeness of the data matrix - no. of data gaps
- Data quality for the underlying analogues for the target and source analogues
- Consistency of data across the data matrix - concordance of effects and potency across analogues

Sources of Uncertainty (cont'd)

- Overarching hypothesis/similarity rationale – how to identify similar analogues and justify their similarity for the endpoint of interest
- Address the dissimilarities and whether these are significant from a toxicological standpoint
- Presence vs. absence of toxicity
- Toxicokinetics

Uncertainty assessment

- There are several frameworks which aim to identify, document and address the uncertainties associated with read-across inferences/predictions
 - Blackburn & Stuard (2014)
 - Patlewicz et al (2015)
 - Schultz et al (2015)
 - ECHA RAAF (2015)
- However read-across acceptance relies on a subjective expert assessment
- There is no objective measure of read-across performance

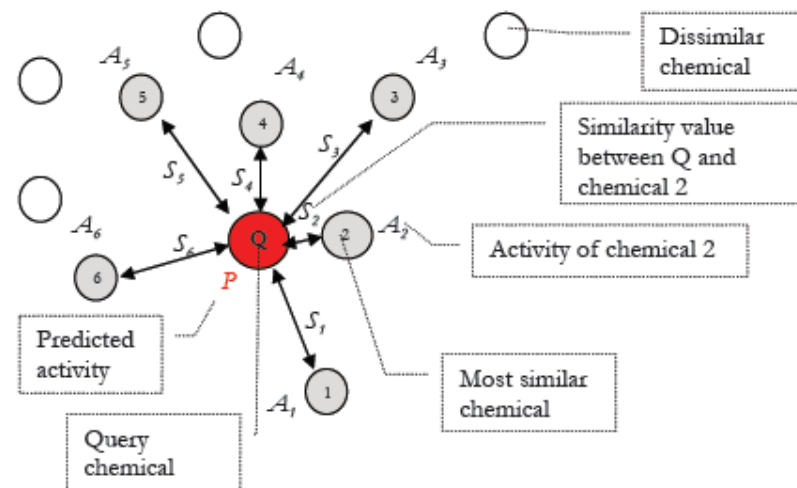
Quantifying Uncertainty & Assessing Performance of Read-Across

- GenRA (Generalised Read-Across) is a “local validity” approach
- Predicting toxicity as a similarity-weighted activity of nearest neighbors based on chemistry and bioactivity descriptors
- Generalised version of Chemical-Biological Read-Across (CBRA) developed by Low et al (2013)
- Systematically evaluates read-across performance and uncertainty using available data

$$y_i^{\beta, \alpha} = \frac{\sum_j^k s_{ij}^{\alpha} x_j^{\beta}}{\sum_j^k s_{ij}^{\alpha}}$$

Jaccard similarity:

$$s_{ij} = \frac{\sum_l (x_{il} \wedge x_{jl})}{\sum_l (x_{il} \vee x_{jl})}$$



GenRA - Approach

I. Data

1,778 Chemicals
3,239 Structure descriptors (chm)
820 Bioactivity assays (bio)
ToxCast
574 Apical outcomes (tox)
ToxRefDB

II. Define Local neighborhoods

Use K-means analysis to group chemicals by similarity
Use cluster stability analysis
~ 100 local neighborhoods

III. GenRA

Use GenRA to predict apical outcomes in local neighborhoods
Evaluate impact descriptors (chm, bio, bc) on prediction
Quantify uncertainty

GenRA - Performance in Each Cluster

- No preselection of descriptors was performed
- Tested and compared
 - Chemical descriptors
 - Bioactivity descriptors
 - Hybrid of chemical and bioactivity descriptors
- Use GenRA to predict the similarity weighted toxicity scores for each
 - Toxicity type (β)
 - Descriptor = {chm, bio, bc} (α)
 - No. of nearest neighbors (k)
 - Similarity score threshold (s_{ij}^{α})
- Calculate performance by comparing predicted y^{tox} and true x^{tox} for all chemicals using area under ROC curve (AUC)
- Bioactivity descriptors were often found to be more predictive of in vivo toxicity outcomes

GenRA - Insights and Next Steps

- The approach enabled a performance baseline for read-across predictions of specific study outcomes to be established but was still context dependent on the endpoint and the chemical
- Ongoing analysis:
- Consideration of other information to refine the analogue selection - e.g. **physicochemical similarity**, TK similarity, metabolic similarity, reactivity similarity...

From research to implementation: *GenRA* prototype

- Intent is to integrate objective read-across functionality as part of ongoing dashboard efforts see <https://comptox.epa.gov/dashboard>
- A limited release of *GenRA* is currently available on EPA's development server

Integration via a GenRA tab

Chemistry Dashboard

Diethylene glycol
111-46-6 | DTXSID8020462

© Searched by CAS-RN: Found 1 result for '111-46-6'.

Wikipedia

Diethylene glycol (DEG) is an organic compound with the formula (HOCH2CH2)2O. It is a colorless, practically odorless, poisonous, and hygroscopic liquid with a sweetish taste. It is miscible in water, alcohol, ether, acetone, and ethylene glycol. DEG is a widely used solvent. It can be a contaminant in consumer products; this has resulted in numerous epidemics of poisoning since the early 20th century.... [Read more](#)

Intrinsic Properties

Structural Identifiers

Related Compounds (Beta)

Presence in Lists

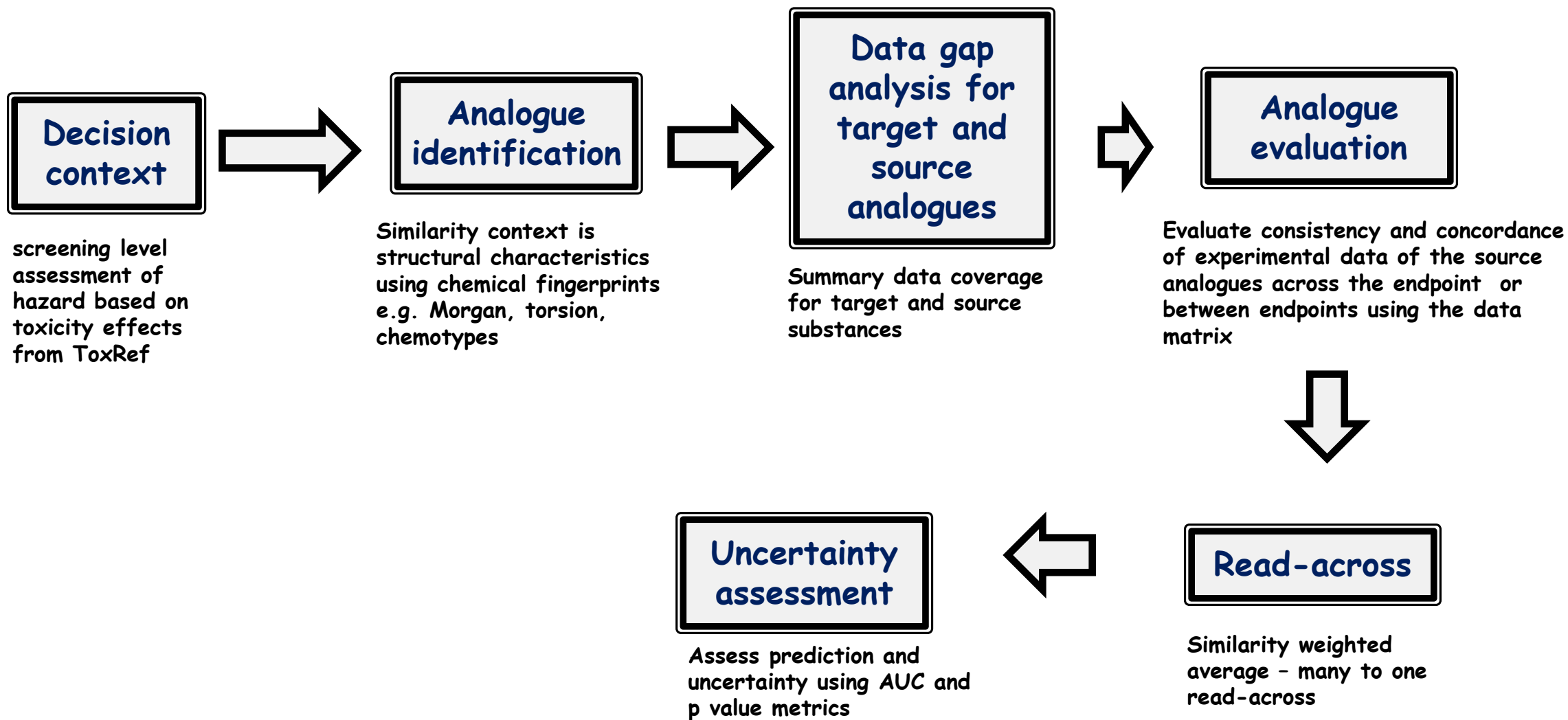
Record Information

GenRA (Beta) | Chemical Properties | Synonyms | External Links | Env. Fate/Transport | Toxicity Values (Beta) | Bioassays | Exposure | Literature | Similar Molecules (Beta) | Comments

Summary:

Grid interface where windows are dynamically updated in subsequent windows

Current Category Workflow in GenRA



GenRA (Beta) Chemical Properties Synonyms External Links Env. Fate/Transport Toxicity Values (Beta) Bioassays Exposure Literature Similar Molecules (Beta) Comments

NN By: **chm_mrgn** K: 10 Sel: **tox_txf** Summary: Grp: **tox_txf** By: **tox_fp** Read-across

Similarity context

Chemical structures shown in the similarity context include: Triethylene glycol, 2-Butoxyethanol, Ethylene glycol, 1,2-Propylene glycol, 2-Methoxyethanol, 2-(Hexyloxy)ethanol, N,N-Diethylethanolamine, Isopentyl alcohol, 2-Methyl-1-propanol, 2-Chloroethanol, and Dimethylaminoethanol.

Heatmap data (approximate values):

Chemical	tox_txf	chm_mrgn	bio_txf	bio_mrgn
2-Methoxyethanol	22	6	6	6
Triethylene glycol	6	6	6	6
2-Butoxyethanol	6	6	6	6
Ethylene glycol	6	6	6	6
2-(Hexyloxy)ethanol	6	6	6	6
Isopentyl alcohol	6	6	6	6
Dimethylaminoethanol	6	6	6	6
2-Chloroethanol	6	6	6	6
2-Methyl-1-propanol	6	6	6	6
N,N-Diethylethanolamine	6	6	6	6
1,2-Propylene glycol	6	6	6	6

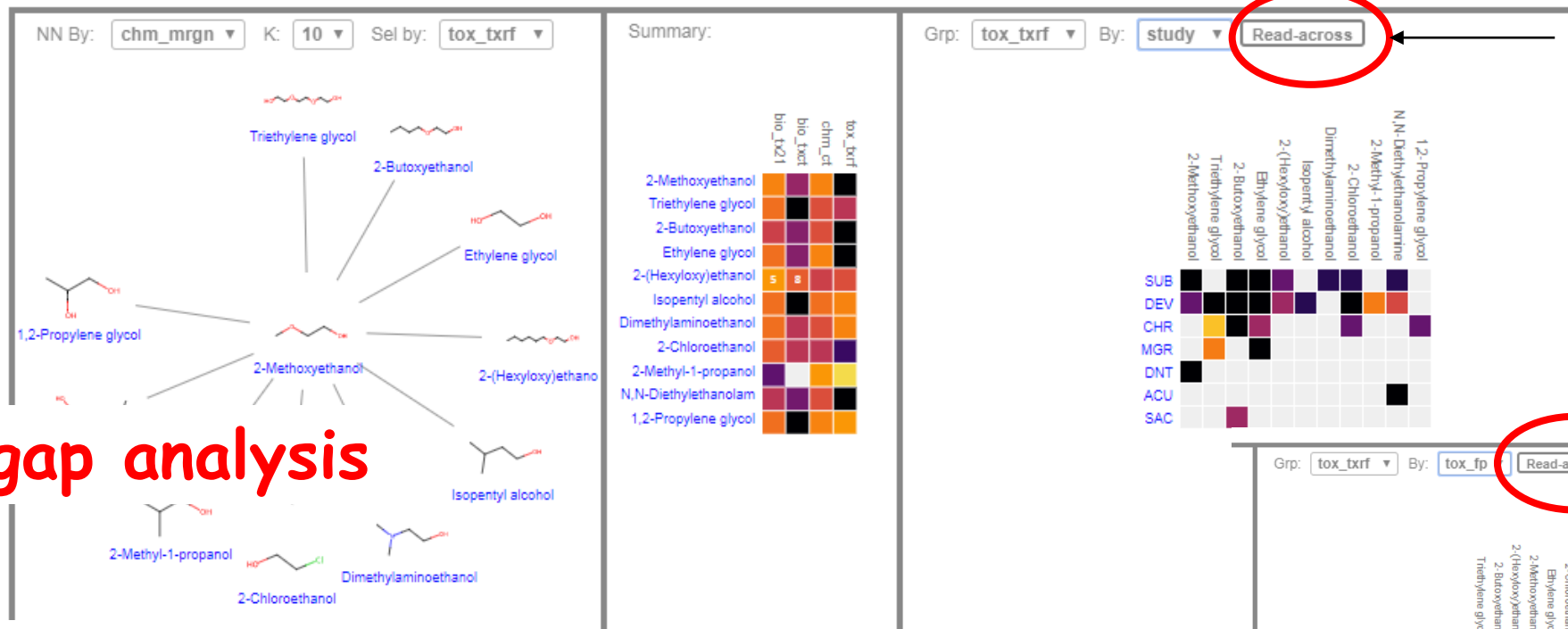
Heatmap data (approximate values):

Chemical	DEV:Body Weight	DEV:Bone	SUB:Clinical Signs	SUB:Kidney	SUB:Liver	SUB:Mortality	DEV:Mortality	DEV:Uterus	DEV:Kidney	DEV:Food Consumption	DEV:Clinical Signs
1,2-Propylene glycol	1	1	1	1	1	1	1	1	1	1	1
N,N-Diethylethanolamine	1	1	1	1	1	1	1	1	1	1	1
2-Methyl-1-propanol	1	1	1	1	1	1	1	1	1	1	1
2-Chloroethanol	1	1	1	1	1	1	1	1	1	1	1
2-Methoxyethanol	1	1	1	1	1	1	1	1	1	1	1
Triethylene glycol	1	1	1	1	1	1	1	1	1	1	1
2-Butoxyethanol	1	1	1	1	1	1	1	1	1	1	1
Ethylene glycol	1	1	1	1	1	1	1	1	1	1	1
2-(Hexyloxy)ethanol	1	1	1	1	1	1	1	1	1	1	1
Isopentyl alcohol	1	1	1	1	1	1	1	1	1	1	1
Dimethylaminoethanol	1	1	1	1	1	1	1	1	1	1	1

Run GenRA Min+: 0 Min-: 0 Filter by: Enter text Sim wt Export

Analogue identification:
Search for source analogues
on the basis of chemical
fingerprints, filtered by
availability of in vivo data

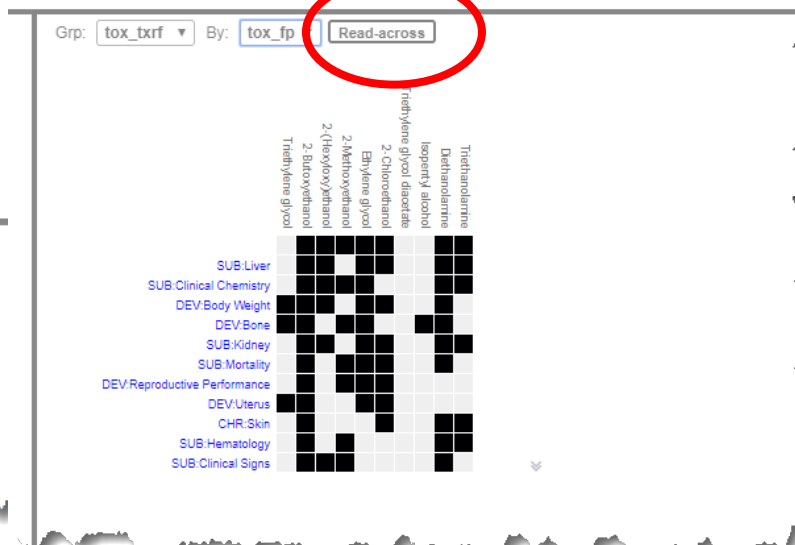
To initiate data matrix view



Data gap analysis

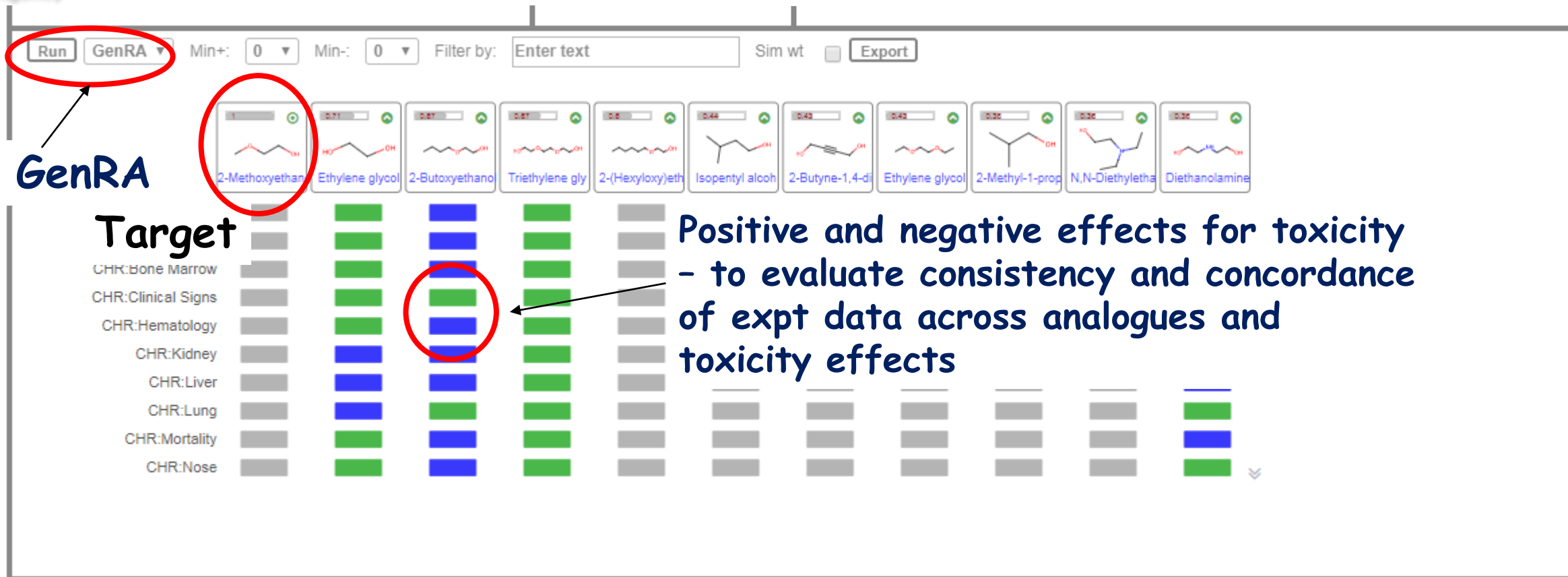
View data quantity by type

Data gap analysis - View data coverage across study type on the basis of toxicity effects



Analogue evaluation using data matrix view

Run GenRA



Export to a csv file

Selected Read-Across Tools

Tool	AIM	ToxMatch	AMBIT	OECD Toolbox	CBRA	ToxRead	GenRA
Analogue identification	X	X	X	X	X	X	X
Analogue Evaluation	NA	X	X by other tools available	X	X	X For Ames & BCF	NA
Data gap analysis	NA	X	X Data matrix can be exported	X Data matrix viewable	NA	NA	X Data matrix can be exported
Data gap filling	NA	X	User driven	X	X	X	X
Uncertainty assessment	NA	NA	NA	X	NA	NA	X
Availability	Free	Free	Free	Free	Free	Free	Beta for Internal testing

Summary

- Still many challenges remain in read-across
- Quantifying the uncertainty of read-across prediction is a critical issue
- Have illustrated the research directions being taken within NCCT and work to implement these into practical tools

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