

# Systematic Approaches to Biological/Chemical Read-Across for Hazard Identification



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- 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
- Ongoing research
- Summary

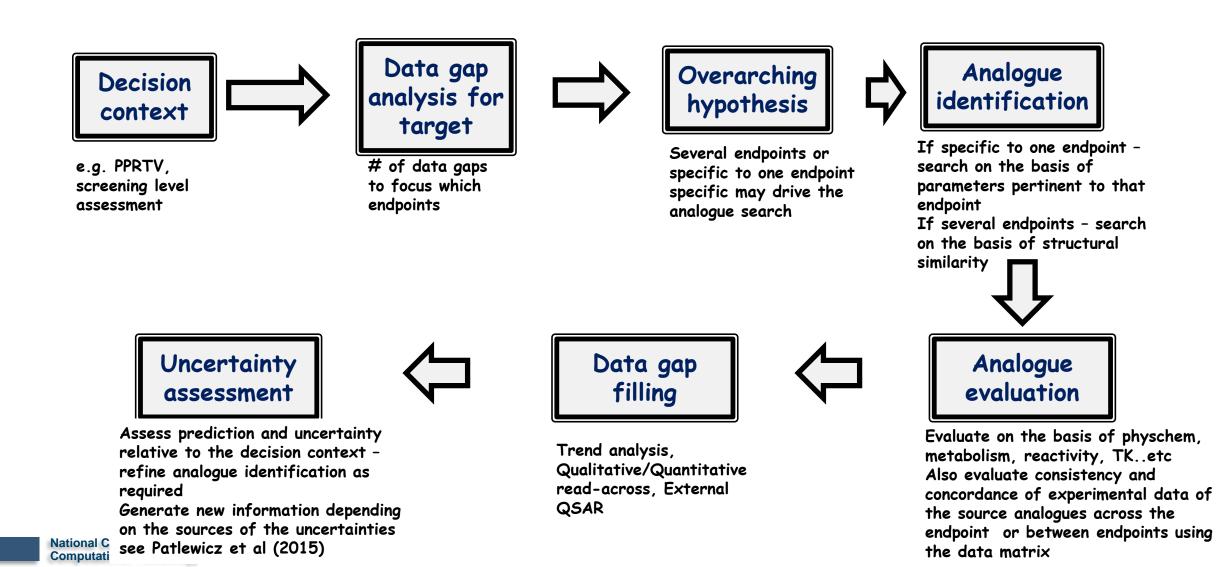




- •<u>Read-across</u> describes one of the <u>data gap filling techniques</u> used within <u>analogue</u> and <u>category</u> approaches
- "Analogue approach" refers to <u>grouping</u> based on a very limited number of chemicals (e.g. target substance + source substance)
- "<u>Category</u> approach" 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	To×Match	AMBIT	OECD Toolbox	CBRA	ToxRead
Analogue identification	×	×	×	×	×	×
Analogue Evaluation	NA	×	X by other tools available	×	×	X For Ames & BCF
Data gap analysis	NA	×	X Data matrix can be exported	X Data matrix viewable	NA	NA
Data gap filling	NA	×	User driven	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

CrossMark

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ABSTRACT

#### ARTICLE INFO

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Keywords: Category approach Analogue approach Data gap filling Read-across (Q)SAR Trend analysis Nearest neighbor 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 a 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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### Selected Read-Across Tools

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Data gap analysis	NA	×	X Data matrix can be exported	X Data matrix viewable	NA	NA
Data gap filling	NA	×	User driven	×	×	×
Uncertainty assessment	NA	NA	NA	X	NA	NA
Availability	Free	Free	Free	Free	Free	Free

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- 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
- Address the dissimilarities and whether these are significant from a toxicological standpoint
- Toxicokinetics





### Uncertainty assessment

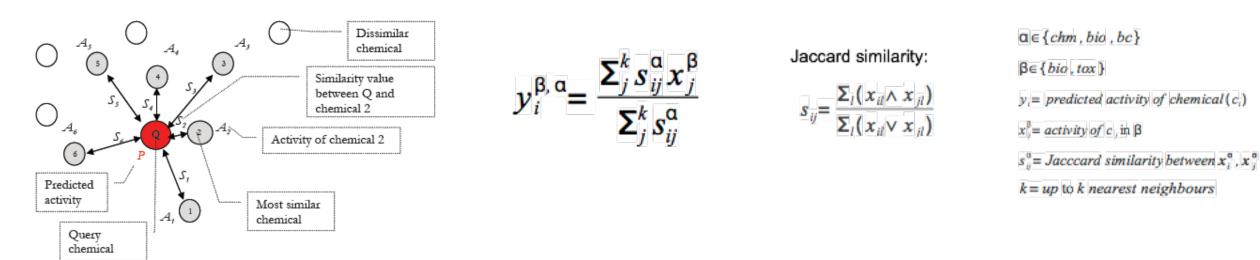
- There are several frameworks which aim to identify, document and address the uncertainties associated with read-across inferences/predictions
  - Blackburn & Stuard (2014)
  - Patlewicz et al (2015)
  - Schultz et al (2015)
  - ECHA RAAF (2015)
- However read-across acceptance relies on a subjective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

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#### PA 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 neighbours based on chemistry and/or bioactivity descriptors
- •Generalised version of Chemical-Biological Read-Across (CBRA) developed by Low et al (2013)
- •Goal: to systematically evaluate read-across performance and uncertainty using available data





## GenRA - Approach

#### I. Data III. GenRA II. Define Local neighbourhoods 1,778 Chemicals Use GenRA to predict toxicity Use K-means analysis to group 3,239 Structure descriptors (chm) effects in local neighbourhoods chemicals by similarity 820 Bioactivity hitcall (bio) ToxCast Evaluate impact of structural Use cluster stability analysis and/or bioactivity descriptors on ~ 100 local neighbourhoods 574 toxicity effects (tox) ToxRefDB prediction Quantify uncertainty CHR SUB MGR DEV Liver Kidney Adrenal Gland Thyroid Gland Testes Stomach Brain Heart Ovary Ev Uterus Bone Marrow Lymph Node Pituitary Gland Thymus Skin Pancreas Mammary Gland Urinary Bladder Epididymis Intestine Small Blood 🛑 Bone -Intestine Large Parathyroid Gland Skeletal Muscle Nerve Gallbladder Seminal Vesicle Salivary glands Harderian Gland Spinal cord Trachea Ear Blood vessel Parathyroid Vagina Esophagus Oral Mucosa

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100 200 300 400

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Penis Lacrimal Gland Mesentery Coordination Larynx Placenta Reflexes

Ureter



- The approach enabled a performance baseline for read-across predictions of toxicity effects within 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

	d States onmental Protection Home Advanced Search Batch Search cy	Lists		Search Chemistry Das	shboard	
Chemis	stry Dashboard			Submit Comment Copy -	Aa ▼ Aa	Aa
E V	Diethylene glycol 111-466   DTXSID8020462					
Е	Searched by CAS-RN: Found 1 result for '111-46-6'.     Q     Im     Im     L					
Τ.			Wikipedia			
0			Diethylene glycol (DEG) is an organic compound with the formula (HOCH2CH2)2O. It is a colorless, practically odorless, poisonous, and hygroscopic I 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 re poisoning since the early 20th century Read more			
			Intrinsic Properties			
	но он		Structural Identifiers			
			Related Compounds (Beta)			
			Presence in Lists			
E						
			Record Information			
N						
т	GenRA (Beta) Chimical Properties Synonyms External	inks Env. Fate/Tra	Insport Toxicity Values (Beta) Bioassays Exposure Literature Similar Molecules (Beta) Comments			
_	NN By: chm_mrgn ▼ K: 10 ▼ Sel by: tox_txrf ▼	Summary:	Grp: tox_txrf v By: tox_fp v Read-across			
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	2-Butoxyethanol	Diethylene glycol	Trietha Defna e glyceint Ebryle 2-Methou 2-Petrov			
	Z-(Hexylox/Jethanol	entaethylene glycol				
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## Current Category Workflow in GenRA



screening level assessment of hazard based on toxicity effects from ToxRef



Similarity context is structural characteristics using chemical fingerprints e.g. Morgan, torsion, chemotypes



Summary data coverage for target and source substances



Evaluate consistency and concordance of experimental data of the source analogues across the endpoint or between endpoints using the data matrix



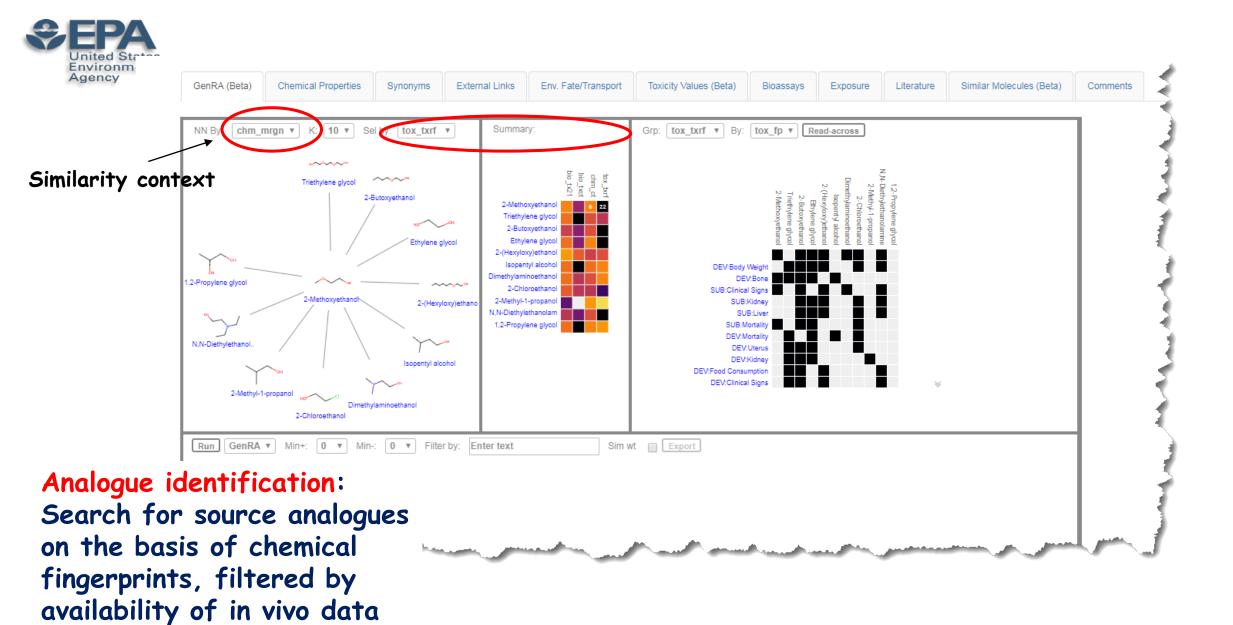
Uncertainty assessment

Assess prediction and uncertainty using AUC and p value metrics



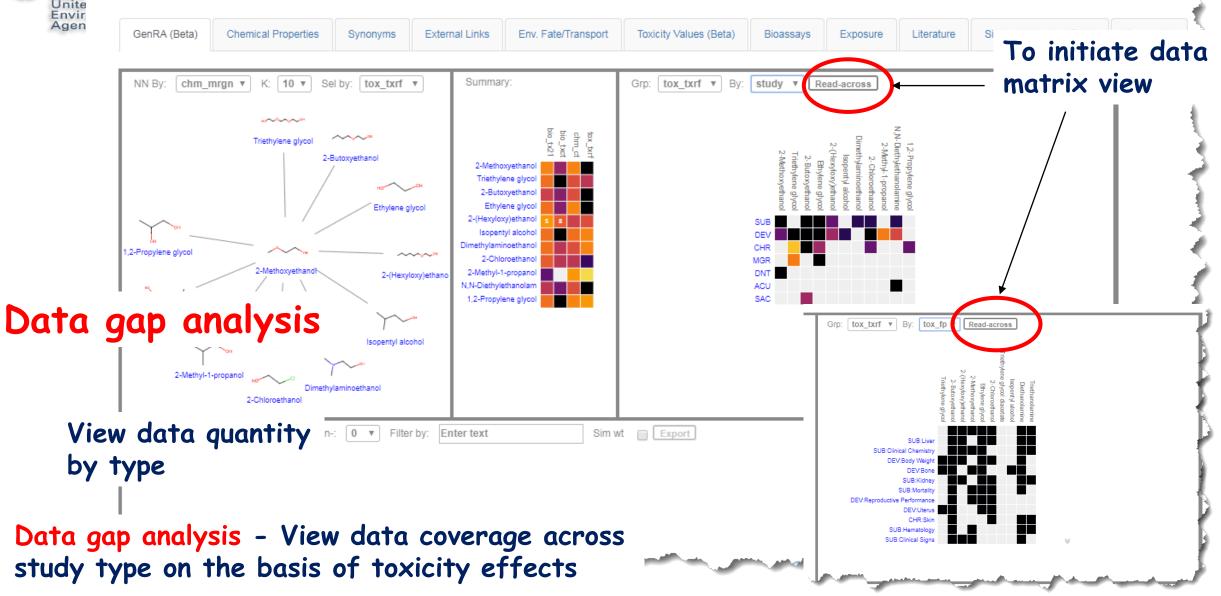


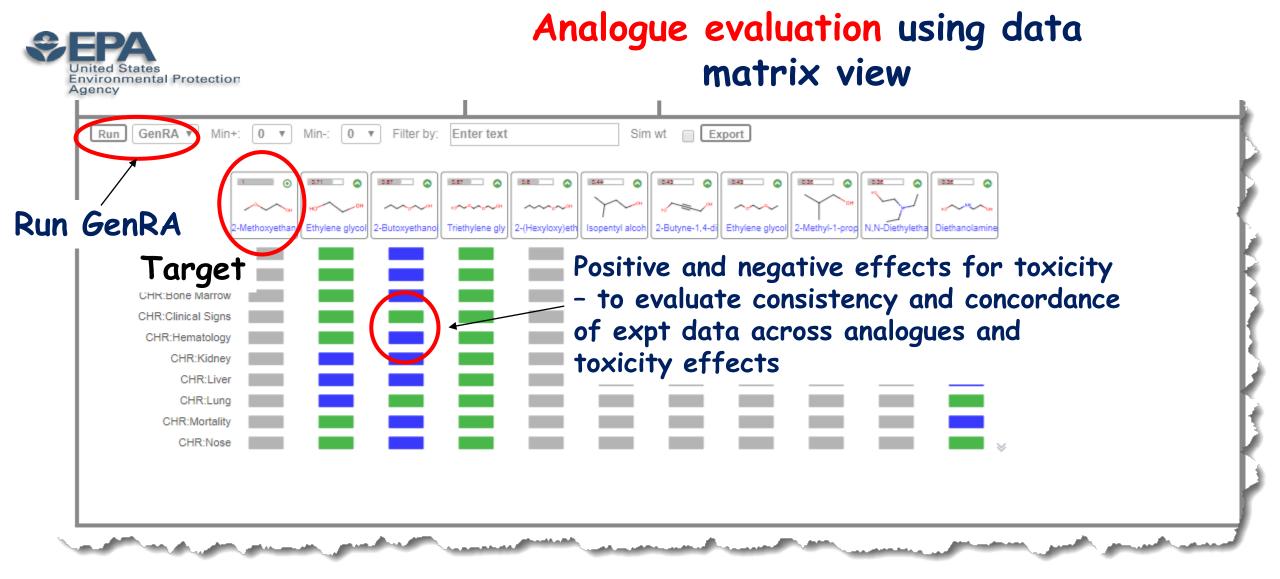
Similarity weighted average – many to one read-across



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### Data gap filling using GenRA within data matrix

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A	В	C C	D	E	F	G	Н	Ι	J	К
cls	target	analog	analog	analog	analog	analog	analog	analog	analog	
label	2-Methoxyethanol	Ethylene glycol	2-Butoxyethanol	Triethyle	2-(Hexylo	Isopentyl	2-Butyne-	Ethylene	2-Methyl-	1-propanol
dsstox_cid	DTXCID804182	DTXCID40597	DTXCID904097	DTXCID60	DTXCID60	DTXCID70	DTXCID90	DTXCID30	DTXCID60	1759
casrn	109-86-4	107-21-1	111-76-2	112-27-6	112-25-4	123-51-3	110-65-6	629-14-1	78-83-1	
jaccard	1	0.714285714	0.666666667	0.666667	0.6	0.444444	0.428571	0.428571	0.375	)
CHR:Bone Marrow	GenRA Neg Act=0 (0.326) AUC=0 p=0.685	no_effect	125.000 ppm	no_effect	t no_data	no_data	no_data	no_data	no_data	
DEV:Bone	GenRA TP Act=1 (1) AUC=0 p=1( 50.000 ppm)	750.000 mg/kg/day	100.000 ppm	5630.000	no_effect	0.500 p	no_effect	100.000	no_effect	:
MGR:Bone	GenRA Pos Act=1 (0.517) AUC=0 p=0.51	1333.330 mg/kg/day	no_data	no_effect	t no_data	no_data	no_data	no_data	no_data	
SUB:Bone	GenRA FN Act=0 (0.483) AUC=0 p=0.66( 546.000 mg/kg/day)	no_effect	500.000 ppm	no_data	no_effect	no_data	no_data	no_data	no_data	
SUB:Bone Marrow	GenRA FN Act=0 (0.483) AUC=0 p=0.65( 297.000 mg/kg/day)	no_effect	62.500 ppm	no_data	no_effect	no_data	no_data	no_data	no_data	
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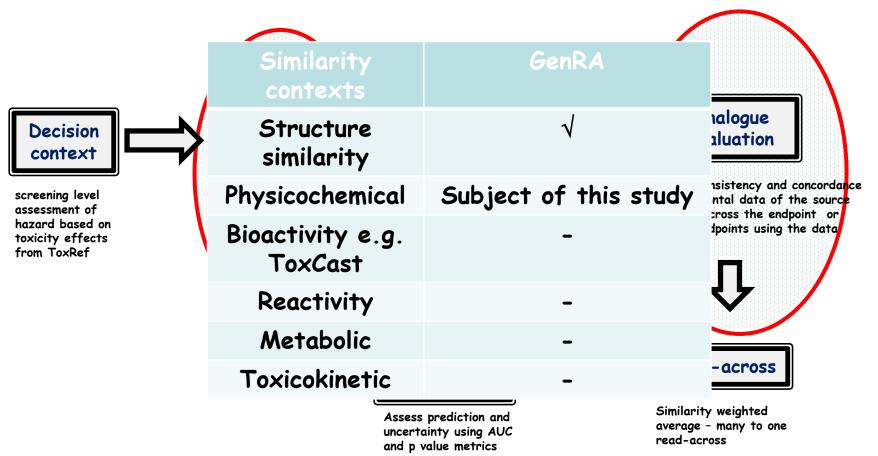


### Selected Read-Across Tools

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



### Ongoing research





### Physchem Similarity Context

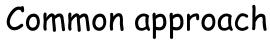
- Important context of similarity in read-across
- Models "bioavailability"
- Properties selected: Lipinski Rule of 5 (LogP, MW, # HB donors/acceptors)
- Two approaches investigated as a means to identify source analogs and evaluate their predictive performance relative to GenRA:

```
Approach 1: "Filter"
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Subcategorise from a set of analogues identified based on structural similarity

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Approach 2: "Search
Expansion"
```

"Frontload" both structure and physchem into analogue identification



Novel approach



### Approaches considered

Approach 1: Filter

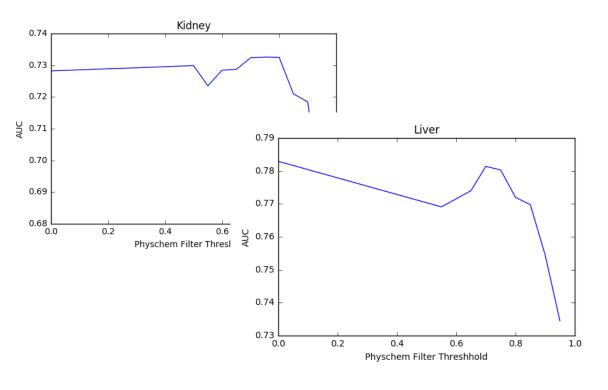
- Similarity search using Jaccard distance of Morgan chemical fingerprints to find source analogues. (Default of 10 nearest neighbours (k=10))
- Calculate physchem similarity between target and source analogues using a generalised Jaccard similarity metric
- Reduce neighbourhood based on the physchem similarity threshold



### Approaches considered

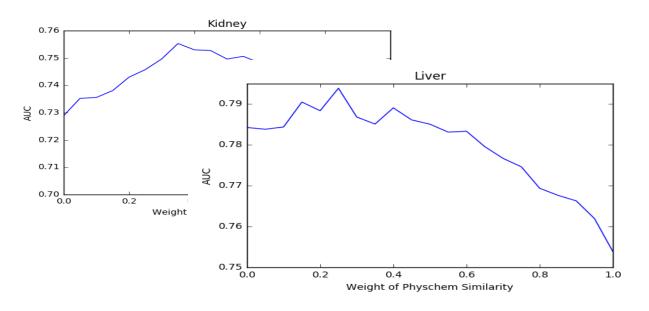
Approach 1: Filter

• This approach did not perform as well as GenRA for the entire dataset, nor did it significantly improve any target organ predictions.



#### Approach 2: Search Expansion

- This approach shows a small improvement over baseline for entire dataset, but large improvement in certain organs.
- Target organ predictions that were significantly improved: Intestine Large, Intestine Small, Mammary Gland, Pancreas, Ureter, Urinary Bladder

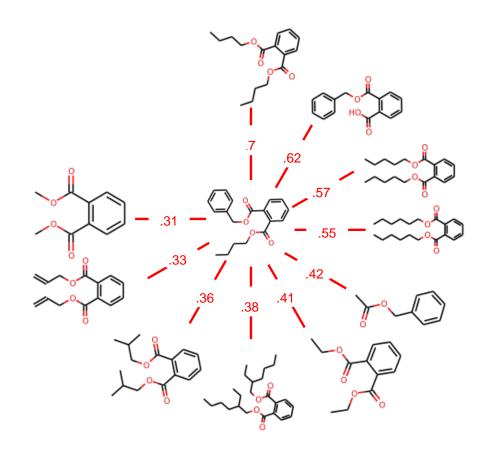


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#### Target organ toxicity predictions aggregated over study type to organ level



### Case Study: Butyl Benzyl Phthalate GenRA: Baseline



Endpoint	Baseline Prediction
Body Weight	.78
Clinical Chemistry	.27
Food Consumption	0
Hematology	0
Kidney	.27
Liver	1
Mortality	.27
Pancreas	.27
Prostate	0
Skin	.27
Spleen	0
Tissue NOS	0
Urinary Bladder	0

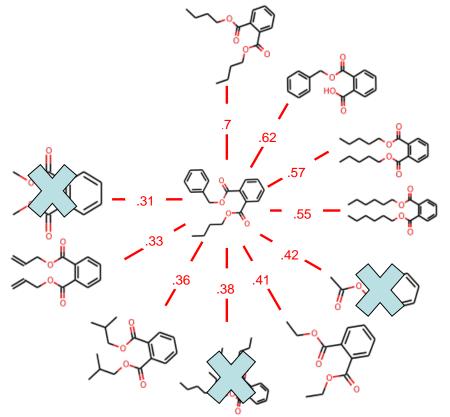
• Chronic studies

- All true positive effects
- Predictions between 0 and 1
- Higher prediction indicates more and stronger positive neighbours



### Case Study: Butyl Benzyl Phthalate Approach 1: Filter

## Filter out chemicals with physchem similarity <0.8



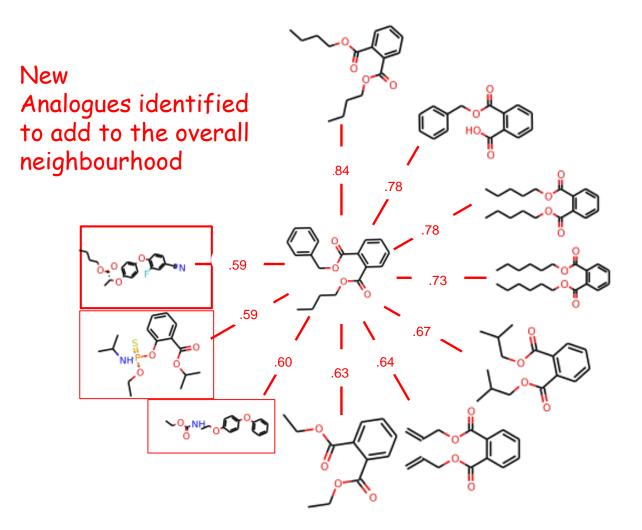
#### Endpoint

**Body Weight** Clinical Chemistry **Food Consumption** Hematology Kidney Liver Mortality Pancreas Prostate Skin Spleen Tissue NOS Urinary Bladder

- Filtering overturns incorrect predictions for 4 endpoints.
- BUT if filtering is too stringent, significant analogues are excluded resulting in a worse performance c.f original GenRA baseline



### Case Study: Butyl Benzyl Phthalate Approach 2: Search Expansion



Endpoint		Baseline Prediction		Structure + Pchem Prediction						
Body Weight	-	78		.79						
Clinical Chemistry		27		.60						
Food Consumption	• Ac	<ul> <li>Adding phys-chem to</li> </ul>								
Hematology										
Kidney		similarity search								
Liver	OV	overturns incorrect predictions for 2								
Mortality	br									
Pancreas	•	•								
Prostate	en	endpoints								
Skin	• In	• Improves many								
Spleen		others								
Tissue NOS	01	11613								
Urinary Bladder	C	)		0						





- Many challenges still remain in read-across
- Quantifying the uncertainty of read-across prediction is a critical issue
- Work in NCCT is focused on systematic and objective approaches to readacross development, evaluation and application
- Established a "baseline" GenRA approach and highlighted progress to implement this approach into a practical tool
- Illustrated the impact on performance that physicochemical similarity can have on analogue identification & evaluation as part of ongoing research





- Rusty Thomas
- Imran Shah
- George Helman
- Tony Williams
- Prachi Pradeep
- Richard Judson
- Jeff Edwards
- Chris Grulke