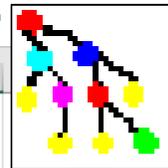


User requirements and concepts I

Falko Frenzel

Much more design properties and user functions!

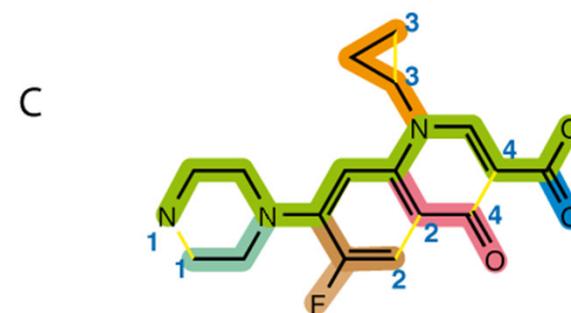
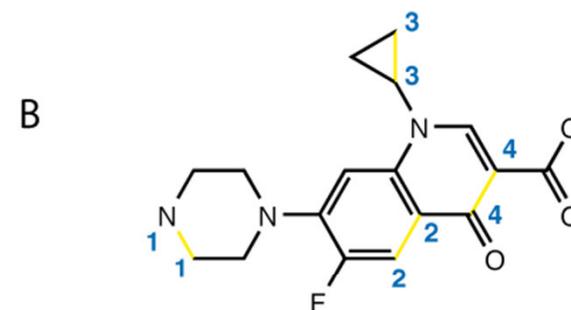
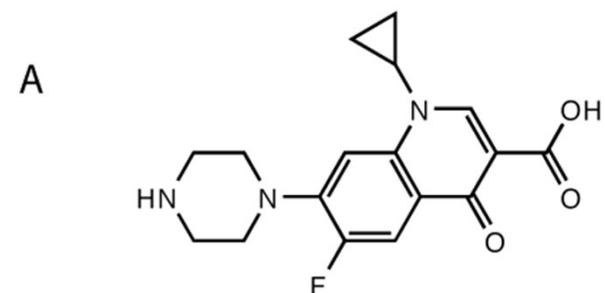



	A	B	E	F
1			A	B
6	Chemical structure notation	Downward compatible	0	1
7	Chemical structure notation	Support different notations	0	1
8	Chemical structure notation	Modul to draw structure should satisfy	0	1
9	Chemical structure notation	Markush/generic structures	0	?
10	Chemical structure notation	Search for structure similarities	1	1
11	Chemical structure notation	Search for similarities independent of the used chemical structure notation	?	1
12	Chemical structure notation	The similarity search filter could be combined with additional filter clauses	0	1
16	Assesment process	Usable for "non-guideline" experiments	0	1
17	Assesment process	Manage "tentative results"	0	1
18	Assesment process	Usable for "freestyle" studies	0	?
19	Assesment process	Manage of textual summaries of the interpretation of the results	1	1
20	Assesment process	Flexible reporting by flexible groups (Pivot tables)	0	1
21	Assesment process	Limitation of 7 columns per table was removed	0	1
22	Assesment process	Recalculations of values from one to another substance	0	1
23	Assesment process	Calculation of concentration factors in relation to other matrix	0	1
24	Assesment process	Grouping of metabolites according the OECD Guideline	0	1
25	Assesment process	Manage Q(SAR) responses in a user storable "List of similar substances"	0	1
26	Assesment process	Integrated start into Q(SAR) Tools with SMILES as the parameter	0	0
27	Assesment process	Manage response from the Q(SAR) tools according ECHA guide	0	0
28	Assesment process	Integrated start into predefined external substance databases	0	0
29	Assesment process	Prediction of metabolic pathways	0	?
30	Assesment process	Pooling of identical substances of different names across the studies	?	1
31	Assesment process	Mange substance groups by defined characteristics (e.g. according to functional groups, conjugates, ...)	0	0
52	Vizualisation	Metabolic pathway	1	1
53	Vizualisation	Overlay (merge) of different Metabolic pathways	1	1
54	Vizualisation	Compare Metabolic pathways	1	1
55	Vizualisation	Concentration time curves	0	1

Chemical Structure Notation

Currently used in Composer and MetaPath

- SMILES as a standard input for chemical structures
- depending on the chosen longest chain, different SMILES notations are possible
- therefore: SMILES are no good identifier



D

N1CCN(CC1)C(C(F)=C2)=CC(=C2C4=O)N(C3CC3)C=C4C(=O)O



image source: <https://upload.wikimedia.org/wikipedia/commons/0/00/SMILES.png>

Chemical Structure Notation

Future use of InChI?

- layer based coding system
- with defined standards – one unique structure encoding

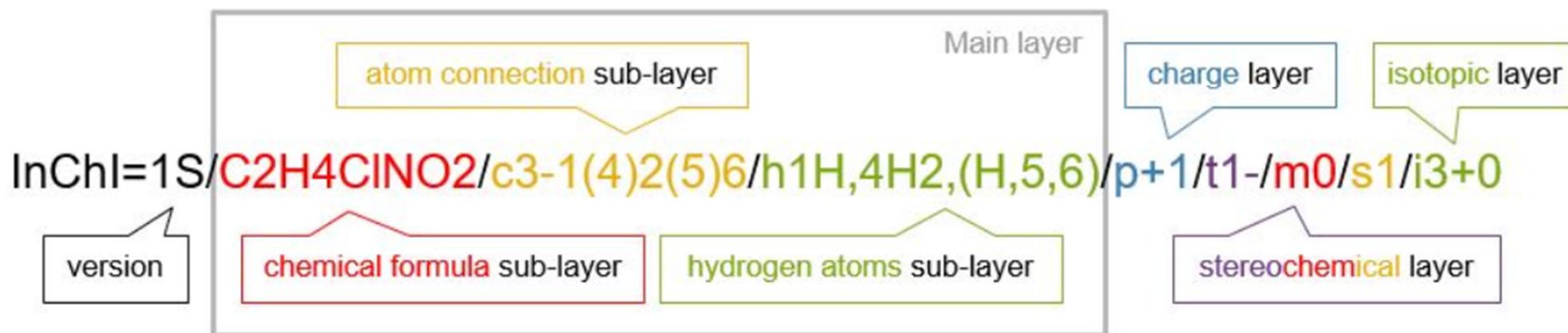
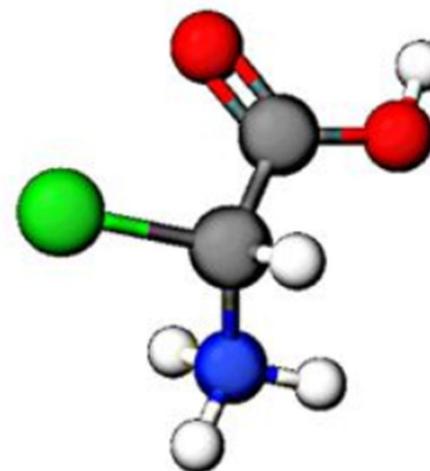
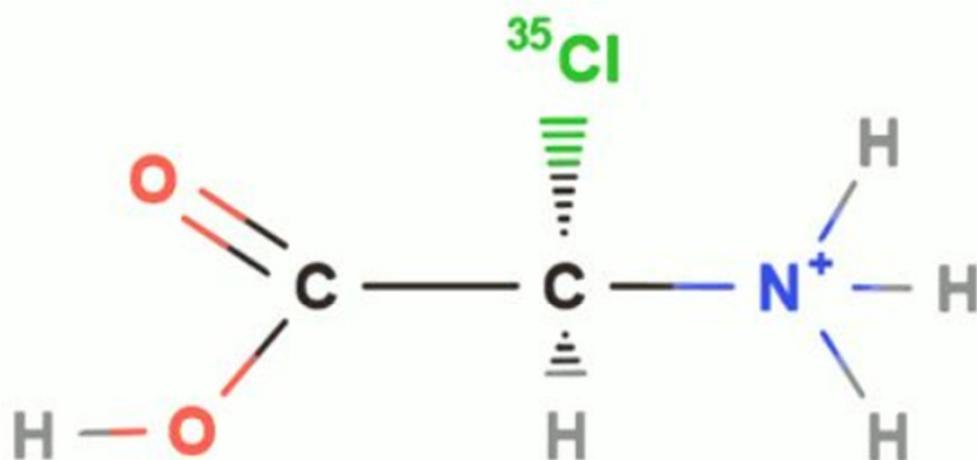


Image source: [https://chem.libretexts.org/Courses/Fordham_University/Chem1102%3A_Drug_Discovery_-_From_the_Laboratory_to_the_Clinic/05%3A_Organic_Molecules/5.08%3A_Line_Notation_\(SMILES_and_InChI\)](https://chem.libretexts.org/Courses/Fordham_University/Chem1102%3A_Drug_Discovery_-_From_the_Laboratory_to_the_Clinic/05%3A_Organic_Molecules/5.08%3A_Line_Notation_(SMILES_and_InChI))

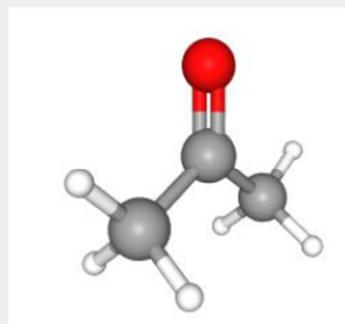
Chemical Structure Notation

SMILES

CC(=O)C

InChI

InChI=1S/C3H6O/c1-3(2)4/h1-2H3



Acetone

Chemical Structure Notation

Currently used:

SMILES (since 1980s)

Pro:

- *better* human readability
- broadly supported

Contra:

- mediocre stereochemical representation
- inconsistent across multiple parser
- ➔ this makes it hard for interoperability
- non unique data base entry

InChI (since 2006)

Pro:

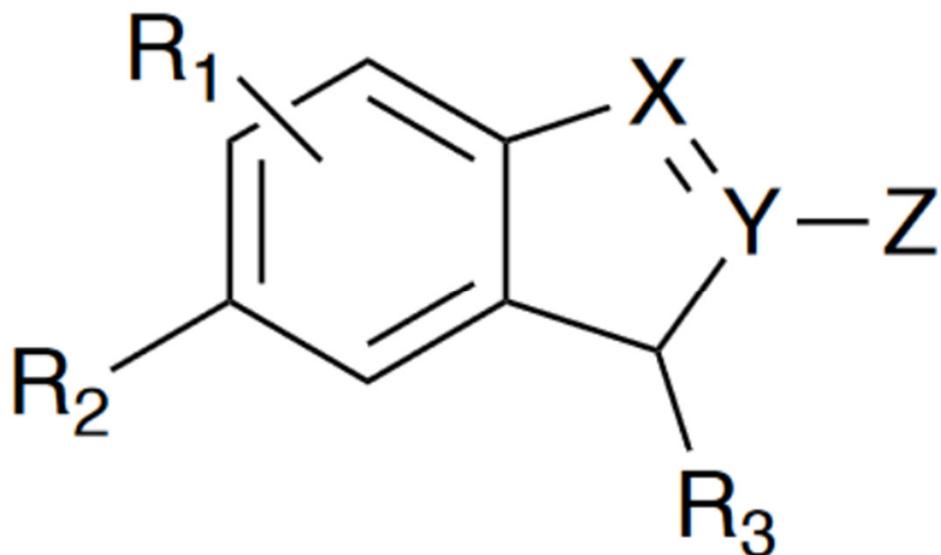
- unique string representation
- includes stereochemical properties
- standardised rules for interpretation
- broadly supported

Contra:

- hard to read by the naked eye (*minor*)
- is meant to be purely machine readable

Chemical Structure Notation

- Markush notations are more complex notations
- often used in patent descriptions



R₁ = H, Me
R₂ = Me, Et, Pr, Bu
R₃ = Ph, tolyl
X = N, CH
Y = C, N⁺
Z = Cl, CH₂Cl

image source: http://www-jmg.ch.cam.ac.uk/inchi/Variable_InChI.pdf

Switch to the voting system now regarding the relevance of Markush structures

Are there any questions? Please use the hand raise in the TEAMS environment.

For statements you could use also the TEAMS chat. The chat will be recorded. **So no idea is lost.**

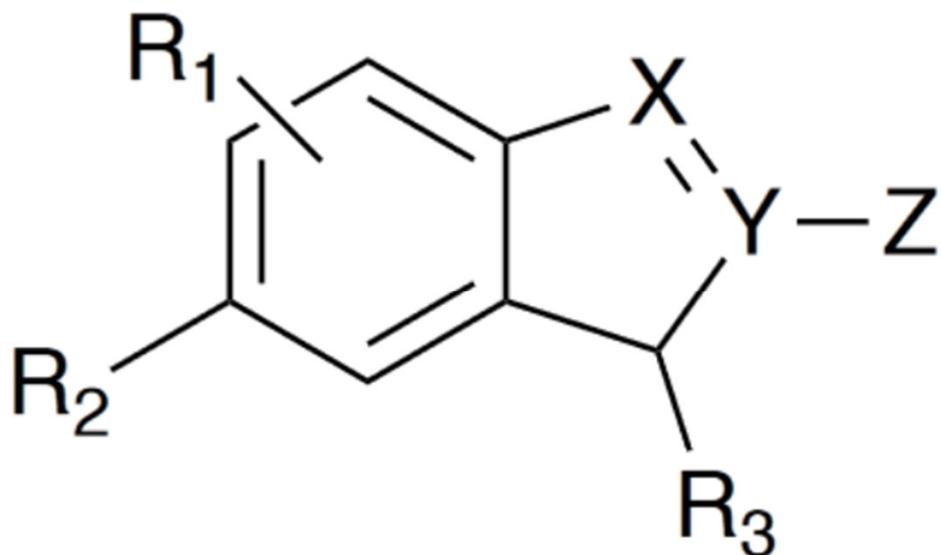
How often would you need a generic Markush Notation?

Would the "missing of Markush Notation" be a showstopper for the project?



Chemical Structure Notation

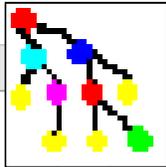
- Markush notations are more complex notations
- often used in patent descriptions
- hard to encode, but there are solutions on the way ([MarkInChI by the InChI Trust](#))
- currently no open standard available, due to financing issues ([InChI Trust Report 2016](#))



R₁ = H, Me
R₂ = Me, Et, Pr, Bu
R₃ = Ph, tolyl
X = N, CH
Y = C, N⁺
Z = Cl, CH₂Cl

image source: http://www-jmg.ch.cam.ac.uk/inchi/Variable_InChI.pdf

Much more design properties and user functions!

	A	B		E	F	
				A	B	C
						
1						
6	Chemical structure notation	Downward compatible	0	?	1	
7	Chemical structure notation	Support different notations	0	?	1	
8	Chemical structure notation	Modul to draw structure should satisfy	0	?	1	
9	Chemical structure notation	Markush/generic structures	0	?	?	
10	Chemical structure notation	Search for structure similarities	1	1	1	
11	Chemical structure notation	Search for similarities independent of the used chemical structure notation	?	?	1	
12	Chemical structure notation	The similarity search filter could be combined with additional filter clauses	0	0	1	
16	Assesment process	Usable for "non-guideline experiments "	0	0	1	
17	Assesment process	Manage "tentative results"	0	0	1	
18	Assesment process	Usable for "freestyle" studies	0	0	?	
19	Assesment process	Manage of textual summaries of the interpretation of the results	1	1	1	
20	Assesment process	Flexible reporting by flexible groups (Pivot tables)	0	0	1	
21	Assesment process	Limitation of 7 columns per table was removed	0	?	1	
22	Assesment process	Recalculations of values from one to another substance	0	0	1	
23	Assesment process	Calculation of concentration factors in relation to other matrix	0	0	1	
24	Assesment process	Grouping of metabolites according the OECD Guideline	0	0	1	
25	Assesment process	Manage Q(SAR) responses in a user storable "List of similar substances"	0	0	1	
26	Assesment process	Integrated start into Q(SAR) Tools with SMILES as the parameter	0	0	0	
27	Assesment process	Manage response from the Q(SAR) tools according ECHA guide	0	0	0	
28	Assesment process	Integrated start into predefined external substance databases	0	0	0	
29	Assesment process	Prediction of metabolic pathways	0	?	?	
30	Assesment process	Pooling of identical substances of different names across the studies	?	?	1	
31	Assesment process	Mange substance groups by defined characteristics (e.g. according to functional groups, conjugates, ...)	0	0	0	
52	Vizualisation	Metabolic pathway	1	1	1	
53	Vizualisation	Overlay (merge) of different Metabolic pathways	1	1	1	
54	Vizualisation	Compare Metabolic pathways	1	1	1	
55	Vizualisation	Concentration time curves	0	0	1	

Grouping and Calculation

- raw data will collect all necessary information
- grouping and calculation are “pivot” tables

Column Group name	Column group	Column group	...	Optional Mean	Optional SD
Row group name					
Row group 1					
Row group 2					
Row group 3					
Sum of all rows					

Dose group / and or ID for animal	DG1	DG2	...
Excretion product 1	TRR per excretion product or organ		
Excretion product 2			
Excretion product ...			
Organ 1			
Organ 2			
Organ ...			
Sum of all TRR			

Grouping and Calculation

Grouping

- grouping based on substances (e.g. similarity based), dose groups, sample groups, matrices and studies should be available by default

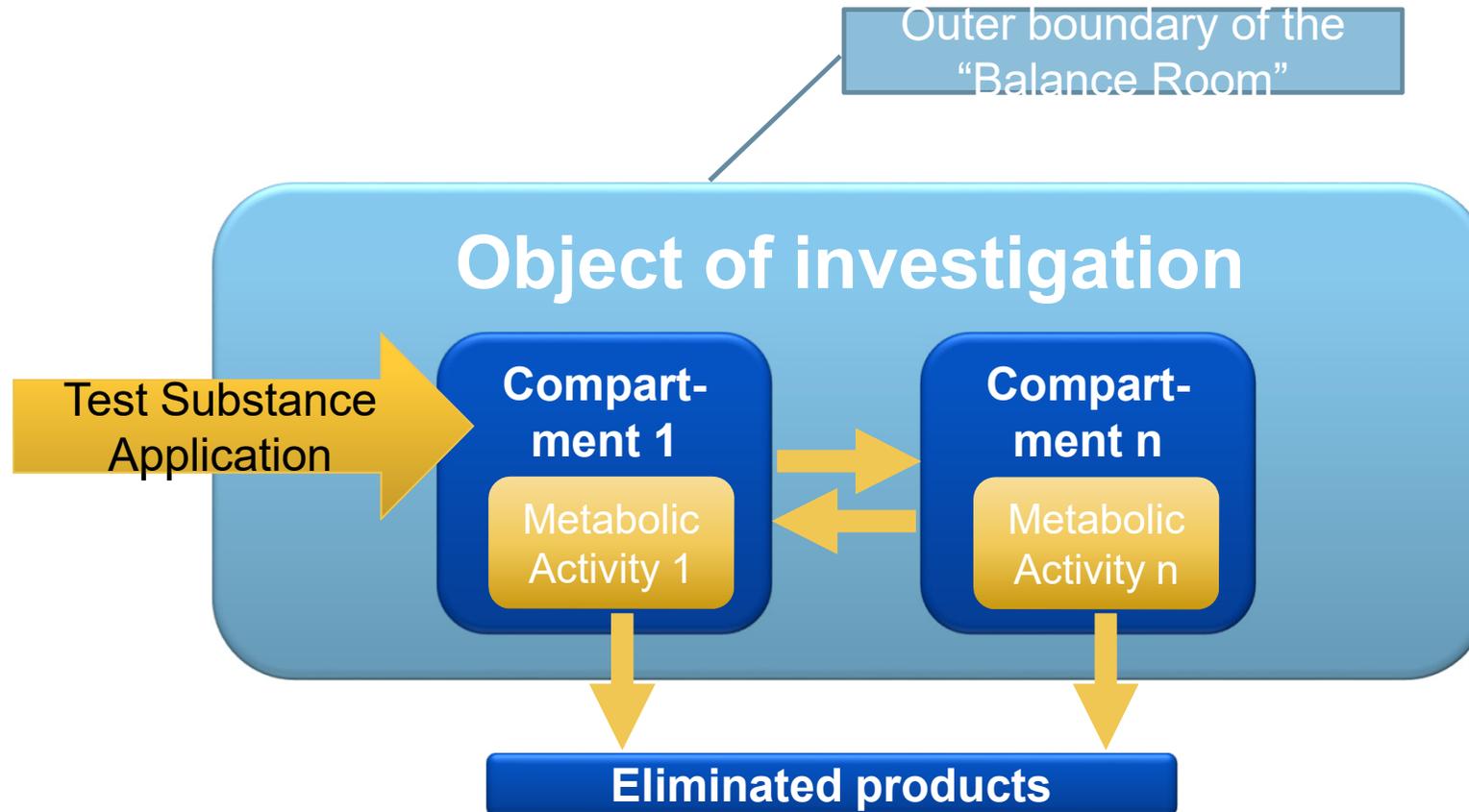
Crop	Soybean			
Study reference	6.2.1/03, Desai 2016; 2013MET-IFP0730, ASB2019-7510			
No and Rate	3 x 125 g as/ha			
N rate	3N rate			
Application method	Foliar spray, ~BBCH 15/16+60+79			
Label	Phenyl (ph) and pyrazole (py) label			
DALA	21 DAA 1	7 DAA 2	7 DAA 2	7 DAA 2
Sample	Forage	Forage	Hay	Hay
TRR mg/kg (combustion)	0,301	0,514	1,794	1,598
	% TRR (ph)	% TRR (py)	% TRR (ph)	% TRR (py)
TRR (extraction+comb. of PES)	103,0	100,7	101,9	98,0
M351 Fluindapyr (IR9792/F9990)	14,80	11,40	6,64	10,2
M367/3 3-OH-fluindapyr	4,04	2,40	4,44	4,40
M353 3-OH-methyl-N-desmethyl fluindapyr				
M381 1-COOH fluindapyr				
M162 N-desmethyl pyrazole COOH				
M176 Pyrazole-COOH		3,27		4,17
M175 Pyrazole-carboxamide		3,49		4,37
M337 N-desmethyl fluindapyr (free and N-conj.)	16,53	19,36	18,17	16,09
M337 N-desmethyl-fluindapyr	[4.60]	[4.16]	[0.97]	[1.59]
M499 N-desmethyl-fluindapyr-N-glu	[11.1]	[15.2]	[17.2]	[14.5]
M565 N-desmethyl-fluindapyr-N-glu-mal	[0.63]	[0]	[0]	[0]
M424 N-desmethyl-fluindapyr-N-serine				

Example for user defined substance groups where results should be aggregated.

Grouping and Calculation

Calculation

- the raw data are delivered with the study
- data might be delivered in concentration per compartment



- distribution coefficient across the test system might be of interest
- BfR has experience in designing and implementing such calculations (e.g. RUEDIS)

Switch to the voting system now regarding the grouping/calculation functionalities

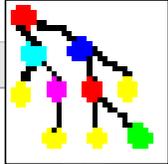
Are there any questions? Please use the hand raise in the TEAMS environment.

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Is there a real need for the evaluators to create flexible reports for substance grouping to calculate concentration factors etc.?

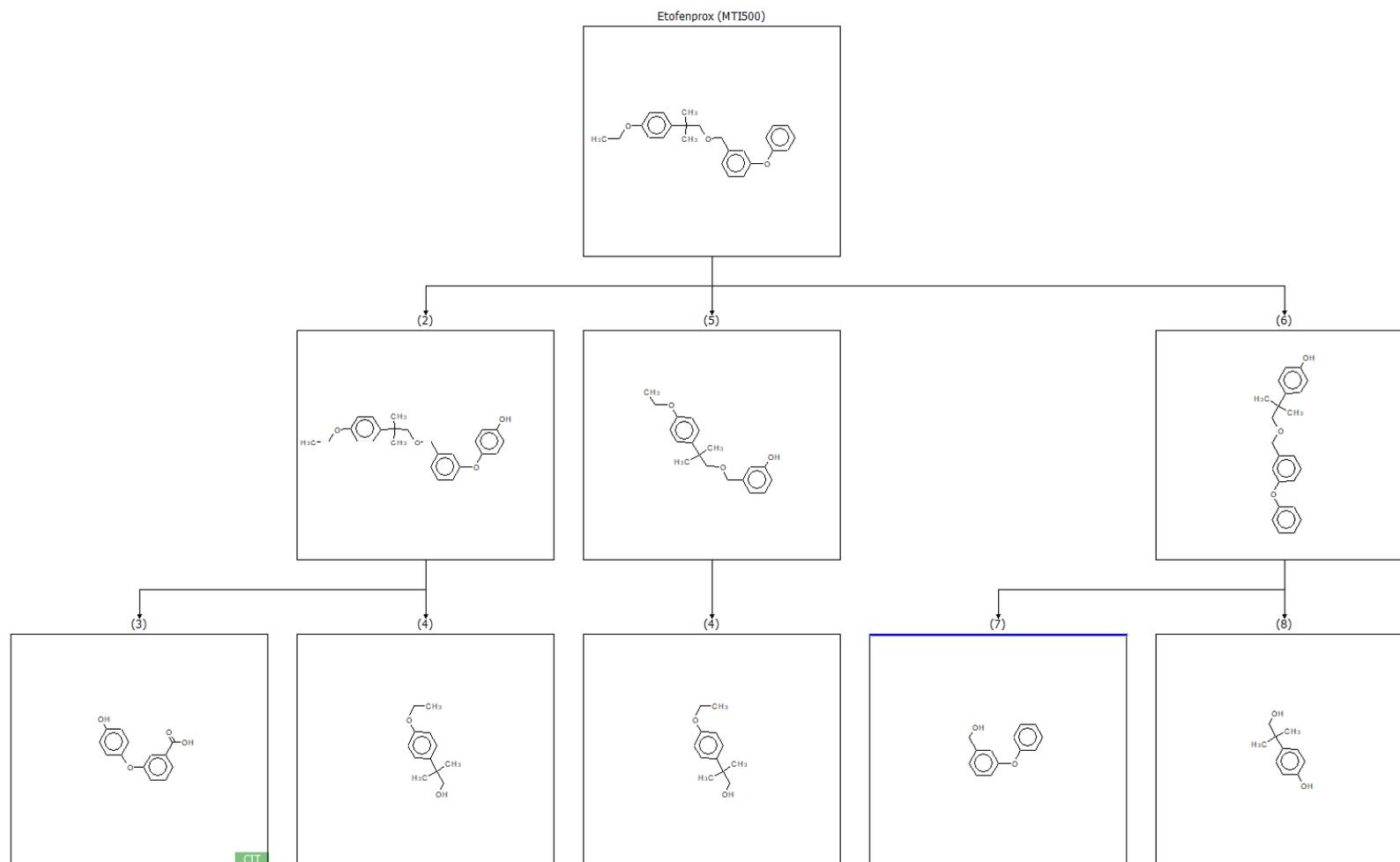


Much more design properties and user functions!

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				A	B	C
						
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10	Chemical structure notation	Search for structure similarities	1	1	1	
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16	Assesment process	Usable for "non-guideline experiments "	0	0	1	
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28	Assesment process	Integrated start into predefined external substance databases	0	0	0	
29	Assesment process	Prediction of metabolic pathways	0	?	?	
30	Assesment process	Pooling of identical substances of different names across the studies	?	?	1	
31	Assesment process	Mange substance groups by defined characteristics (e.g. according to functional groups, conjugates, \	0	0	0	
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53	Vizualisation	Overlay (merge) of different Metabolic pathways	1	1	1	
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55	Vizualisation	Concentration time curves	0	0	1	

Visualisation

- implementation of [PathVisio](#) (OpenSource) which is used in [WikiPathways](#)
- in general the currently used visualisation tools in MetaPath are quite good



Visualisation

- implementation of [PathVisio](#) (OpenSource) which is used in [WikiPathways](#)
- in general the currently used visualisation tools in MetaPath are quite good
- the visualisation is capable of showing the „degradation“ process of a test substance and can merge multiple networks
- network data are always structured similarly:
 - **source**
 - **target**
 - **connection (type)**
 - additional data / meta data
- given this information in text format any modern visualisation tool can create networks/pathways (even webbased ones such as [D3.js](#))

Visualisation

- implementation of [PathVisio](#) (OpenSource) which is used in [WikiPathways](#)
- in general the currently used visualisation tools in MetaPath are quite good

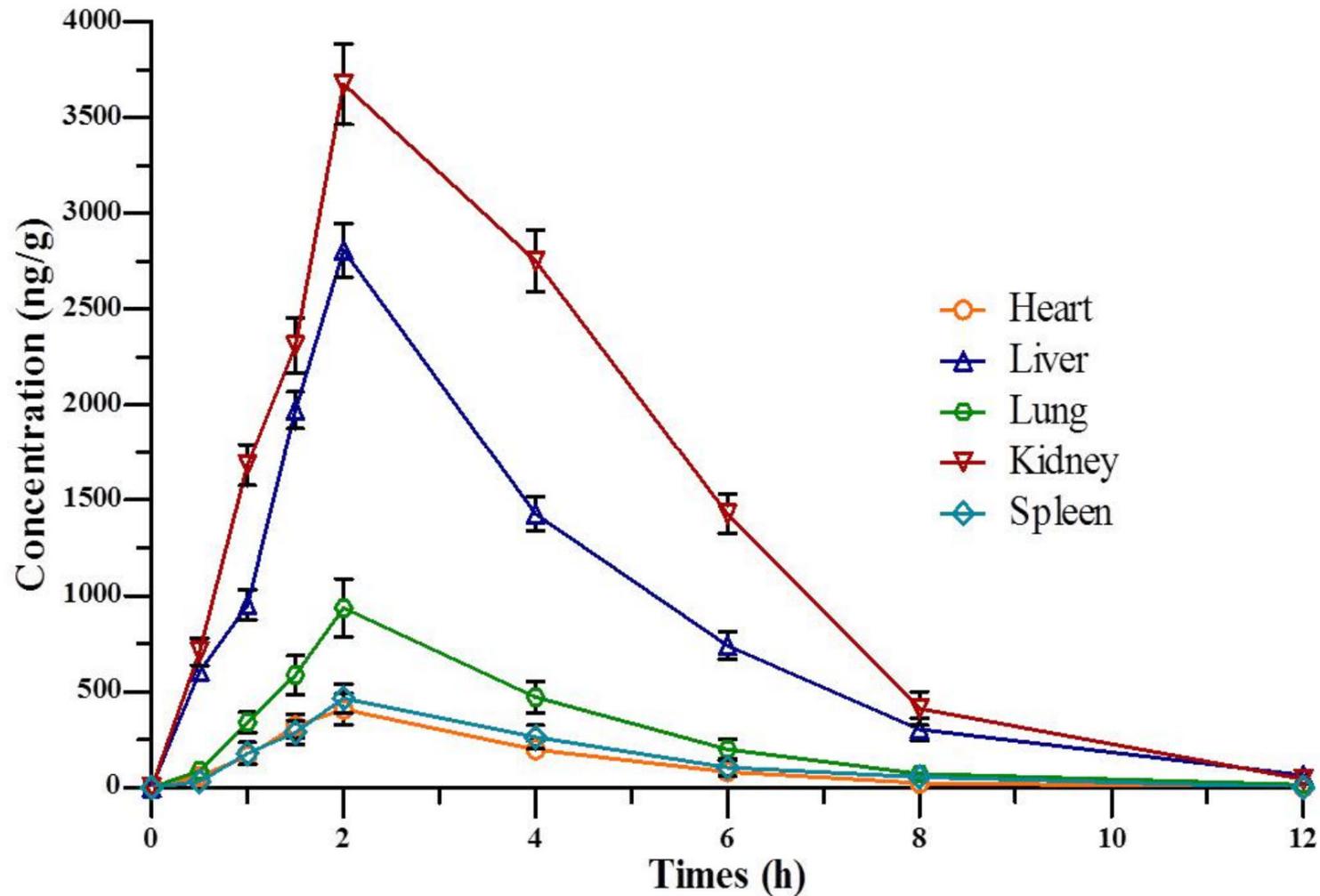


image source: Ambhore et al., 2020 ([10.5530/jyp.2020.12.11](https://doi.org/10.5530/jyp.2020.12.11))

Switch to the voting system now regarding the visualization functionalities

Are there any questions? Please use the hand raise in the TEAMS environment.

For statements you could use also the TEAMS chat. The chat will be recorded. **So no idea is lost.**

Do you have ideas for additional visualisations which you would recommend to include in the project?



Thank you for your attention

Falko Frenzel

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