

The Effectiveness of *In Silico* Methods in Predicting the Formation of Active Metabolites

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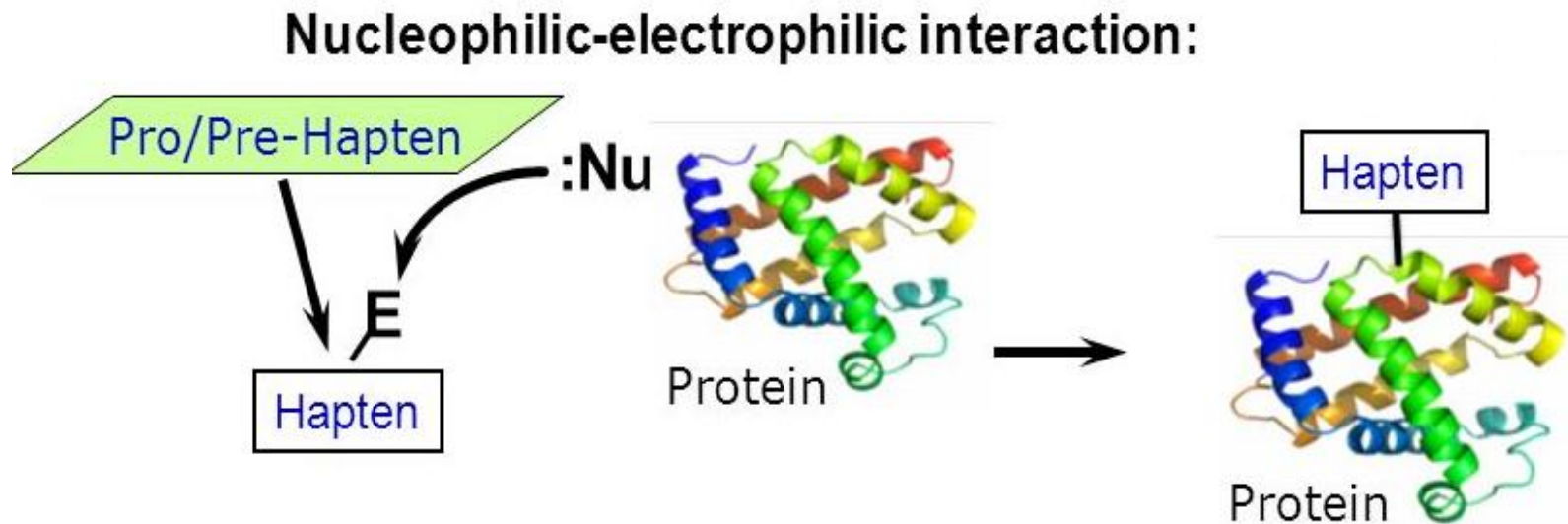
How good are *in silico* systems for predicting skin metabolism (pro-haptens)?

Why is this?

Do we need more experiments?

Skin Sensitisation: Electrophiles

- Pro-/pre-haptens are considered to make up about 25% of the chemical space related to skin sensitisation
- The chemistry of many pro-/pre-haptens is very closely related – indirect electrophiles is a more appropriate term



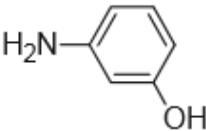
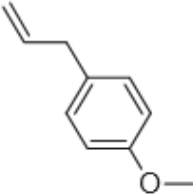
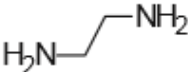
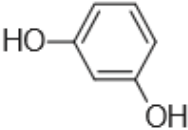
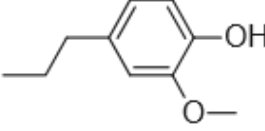
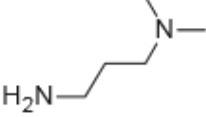
Skin Metabolism

- There are limited data around the exact expression levels of metabolising enzymes in the skin
- The most comprehensive review outlines (rat, mouse, pig, guinea pig and humans):
 - Low expression of the P450 system
 - Expression of oxidising enzymes (COX and FMO)
 - Expression of conjugating enzymes
- Essentially, the skin is capable of oxidation and conjugation

Skin Sensitisation: Pre-/Pro-Haptens

- Numerous chemicals can behave as both pre- and pro-haptens (for example, hydroquinones)
- However, some chemicals are definite pro-electrophiles
- These chemicals are typically not identified in the DPRA
- Interestingly, some of these were identified by other assay from the 'triple pack' (Keratinosens and HCLat)

Table 2
Pro-haptens in the dataset.

Name	Structure	In vivo		Cell free in chemico	Cell based <i>in vitro</i>		Reaction mechanistic domain
		LLNA	Human data	DPRA	KeratinoSens™	h-CLAT	
3-Aminophenol		1	NA	0	0	1	pro-MA
4-Allylanisole		1	NA	1	0	1	pro-MA
Ethylenediamine (free base)		1	1	0	1	1	pro-SB
Resorcinol		1	1	0	0	1	pro-MA
Dihydroeugenol (2-methoxy-4-propyl-phenol)		1	NA	0	1	1	pro-MA
3-Dimethylamino propylamine		1	1	0	1	1	pro-SB ^a

Notes: NA – not available, 1 = positive, 0 = negative, SB = Schiff base former, MA = Michael acceptor.

^a 3-Dimethylamino propylamine was originally assigned as a pre-SB in the original EC EURL-ECVAM report (Casati et al., 2016). It has been reconsidered as a pro-SB electrophile.

Pro-Hapten Identification for Skin Sensitisation: *In Silico* Tools



Plus a number of other commercial software packages – for a review see reference below

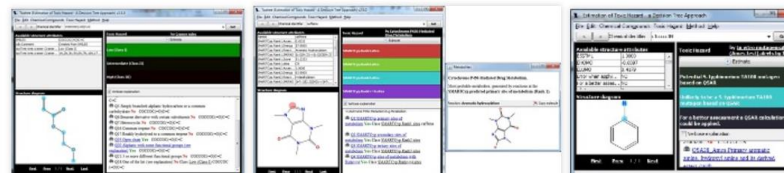
Toxtree - Toxic Hazard Estimation by decision tree approach

Toxtree is a full-featured and flexible user-friendly open source application, which is able to estimate toxic hazard by applying a decision tree approach.

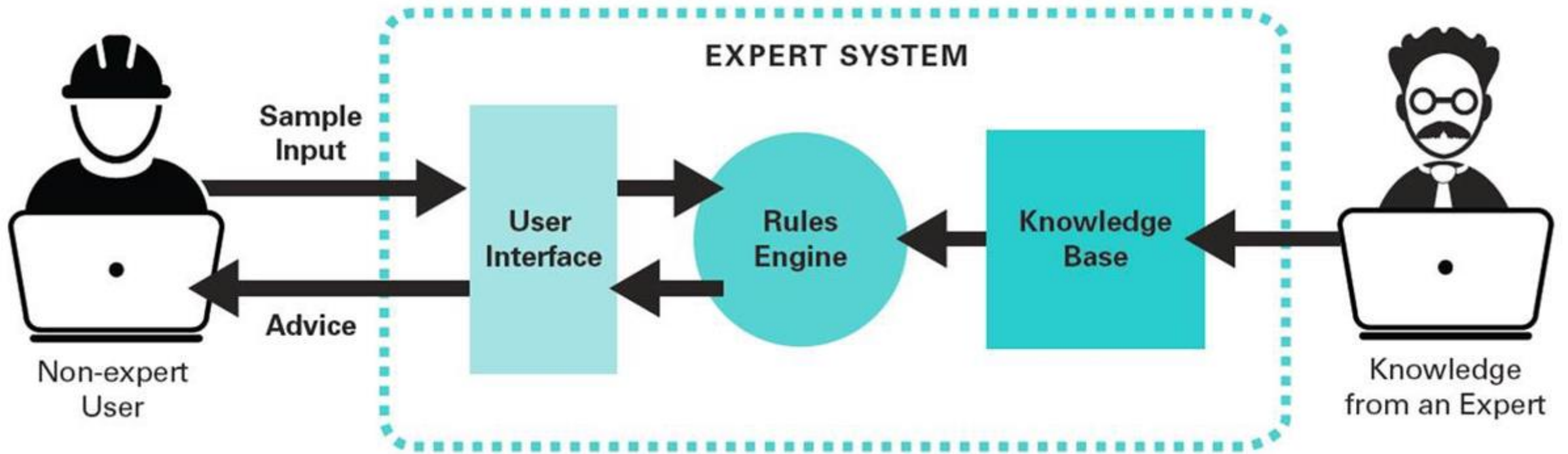
Toxtree could be applied to datasets from various compatible file types. User-defined molecular structures are also supported - they could be entered or by using the built-in 2D structure diagram editor.

The Toxtree application is suitable for a standalone PC and can run on any platform, supported by Java 1.6 or higher.

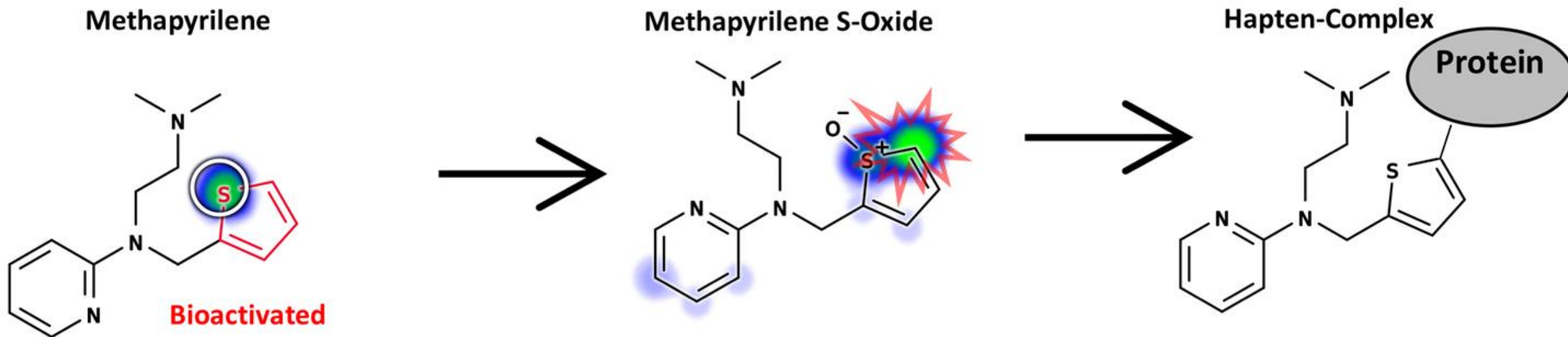
Toxtree has been designed with flexible capabilities for future extensions in mind (e.g. other classification schemes that could be developed at a future date). Custom decision trees with arbitrary rules can be built with the help of graphical user interface or by developing new plug-ins.



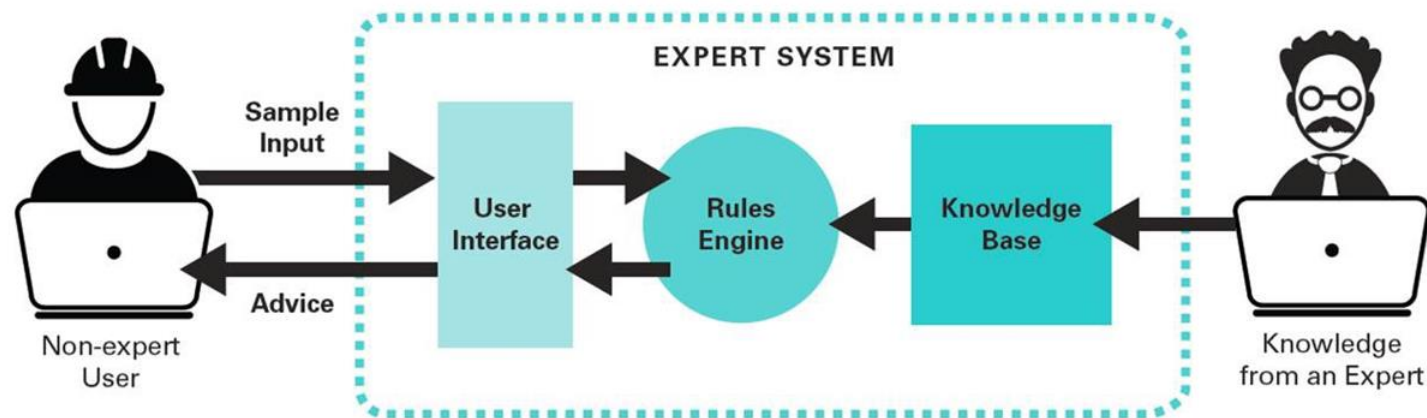
Expert Systems



Expert Systems – Knowledge Base



The knowledge base is constructed using structural alerts that relate chemistry to an effect (MIE, endpoint, metabolism etc.)



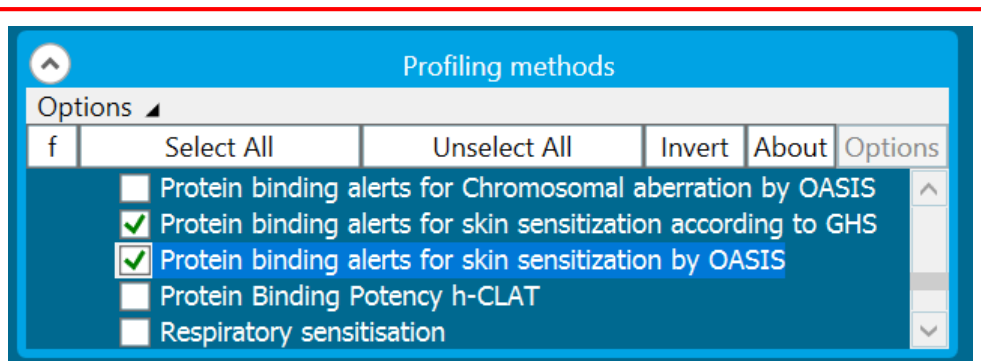
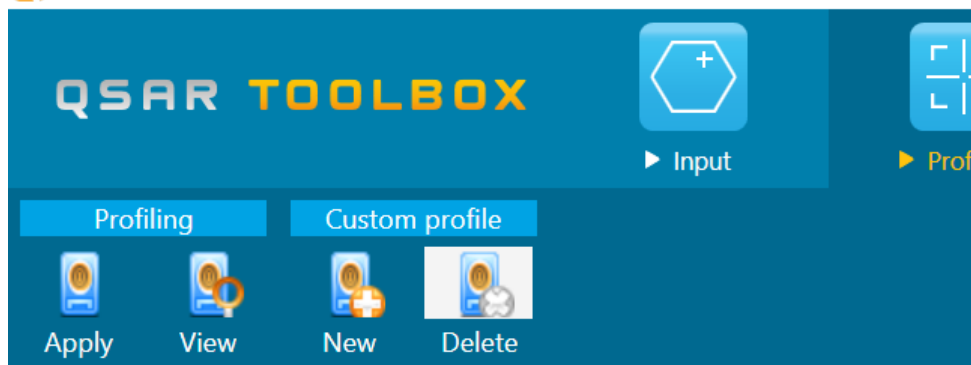
OECD QSAR Toolbox



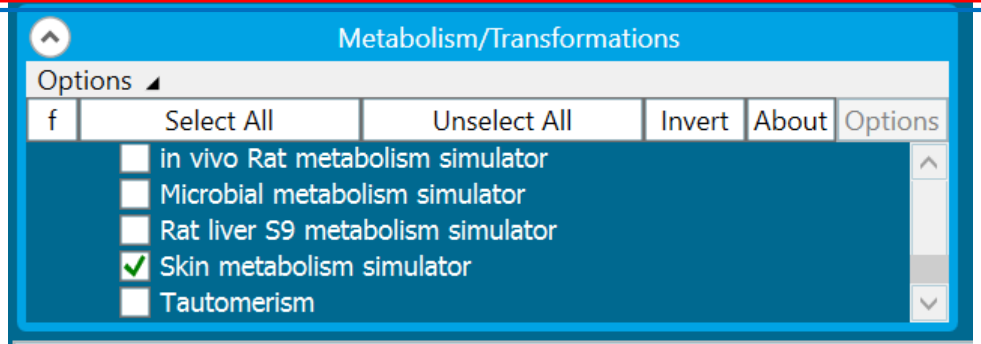
- The OECD QSAR Toolbox is an *in silico* tool designed to facilitate read-across predictions (it is not a true expert system)
- It is chemistry-based, featuring a range of structural alert-based profilers
- These profilers are organised on an MIE and/or endpoint basis
- Contains the knowledge of other freely available tools such as ToxTree (certainly in terms of skin sensitisation)

OECD QSAR Toolbox – Relevant Profilers

QSAR Toolbox 4.3.1 [Document 1]



Protein binding by OECD & OASIS
Protein binding alerts for skin sensitisation
according to GHS & OASIS



Skin metabolism simulator
Hydrolysis simulator
Dissociation simulator

OECD QSAR Toolbox – Skin Metabolism Simulator

- Rules initially derived from an understanding of liver metabolism and skin sensitisation data derived from the LLNA, GPMT and other sources
- Subsequent analysis has improved the skin metabolism rule base with the availability of biotransformations for 151 chemicals
- Full implementation is found in the commercial TIMES software, with a simplified version being available in the OECD QSAR Toolbox
- Full version adds probability and reliability scores for each predicted biotransformation

Derek Nexus (Lhasa Ltd.)



- A true knowledge base system for predictive toxicology
- Contains structural alerts for a number of endpoints, including skin sensitisation
- The Derek Nexus knowledge base for skin sensitisation contains alerts for direct and indirect electrophiles
- Contains a reasoning engine that offers a five levels of confidence in the predictions (based on the extent of the underlying knowledge)
- Levels of confidence being: impossible, improbable, doubted, equivocal, plausible, probable, certain

Meteor Nexus (Lhasa Ltd.)



- No specific skin metabolism model
- Instead, there's a range of potential biotransformations
- As with Derek Nexus, these predictions have an associated level of confidence associated with them: improbable, doubted, equivocal, plausible, probable, certain
- As outlined earlier it is known that skin and liver metabolism differ, with differing expression levels of the key enzymes
- Careful considerations of the relevant enzymes is required in order to prevent Meteor being over predictive of potential metabolites

Profiling Examples

- The next slides outline how we can identify potential haptens for:
- Cinnamyl formate (CAS No 104-65-4)
- Isoeugenol acetate (93-29-8)
- Cinnamic aldehyde dimethyl acetate (4364-06-1)
- Ethylenediamine (107-15-3)

Cinnamyl Formate (104-65-4)

QSAR TOOLBOX

Input Profiling Data Category definition Data Gap Filling

Profiling Custom profile

Apply View New Delete

Documents

Document 1
CAS: 104654

Profiling methods

Options

Select All Unselect All Invert About Option

- Protein binding alerts for Chromosomal al
- Protein binding alerts for skin sensitization
- Protein binding alerts for skin sensitization
- Protein Binding Potency h-CLAT

Metabolism/Transformations

Options

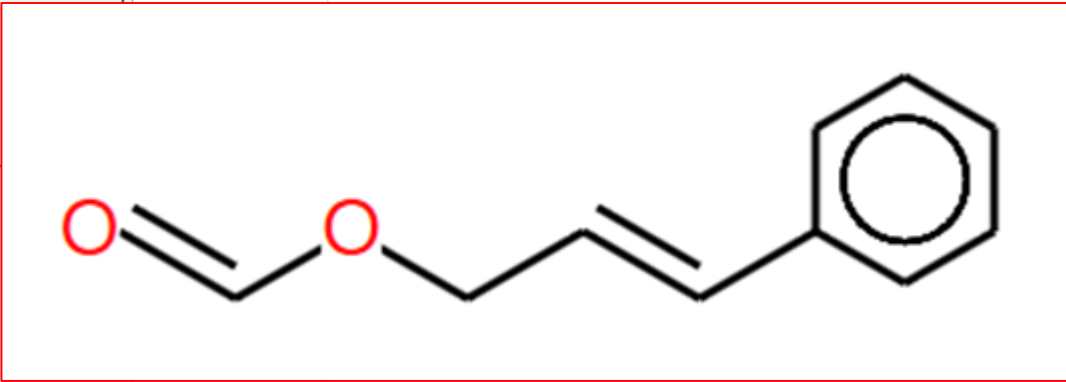
Select All Unselect All Invert About Option

- Microbial metabolism simulator
- Rat liver S9 metabolism simulator
- Skin metabolism simulator
- Tautomerism

Filter endpoint tree... 1 [target]

Structure

- Parameters
- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazards
- Profile
 - General Mechanistic
 - Protein binding by OASIS No alert found
 - Protein binding by OECD No alert found
 - Endpoint Specific
 - Protein binding alerts for skin sensitization according to GHS No alert found
 - Protein binding alerts for skin sensitization by OASIS No alert found
 - Metabolism/Transformations
 - Skin metabolism simulator 8 metabolite(s)



Cinnamyl Formate (104-65-4)

QSAR TOOLBOX

Input Profiling Data Category definition Data Gap Filling

Profiling Custom profile

Apply View New Delete

Documents

Document 1
CAS: 104654

Filter endpoint tree... 1 [target]

Structure

Profiling methods

Options

Select All Unselect All Invert

- Protein binding alerts for Chro
- Protein binding alerts for skin
- Protein binding alerts for skin
- Protein Binding Potency h-CL

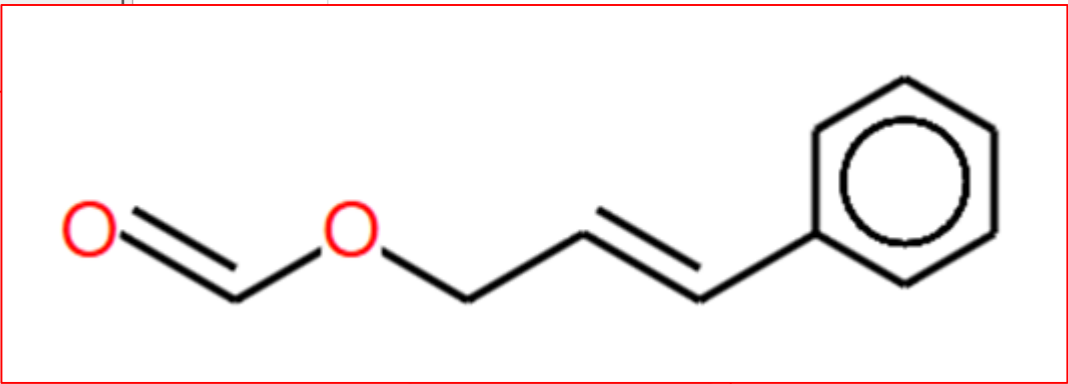
Metabolism/Transformation

Options

Select All Unselect All Invert

- Microbial metabolism simulat
- Rat liver S9 metabolism simul
- Skin metabolism simulator
- Tautomerism

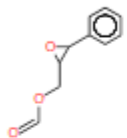
- Parameters**
- Physical Chemical Properties**
- Environmental Fate and Transport**
- Ecotoxicological Information**
- Human Health Hazards**
- Profile**
 - General Mechanistic**
 - Protein binding by OASIS
 - Protein binding by OECD
 - Endpoint Specific**
 - Protein binding alerts for skin sensitization according to GHS
 - Protein binding alerts for skin sensitization by OASIS
 - Metabolism/Transformations**
 - Skin metabolism simulator



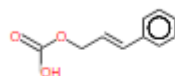
No alert found
No alert found
No alert found
No alert found
8 metabolite(s)

Cinnamyl Formate (104-65-4)

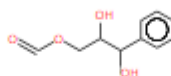
metabolite #1
No CAS number



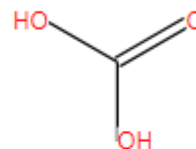
metabolite #2
No CAS number



metabolite #3
No CAS number



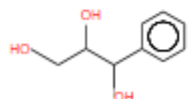
metabolite #4
No CAS number



metabolite #5
No CAS number



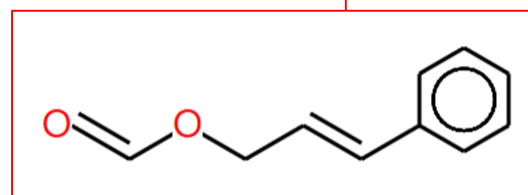
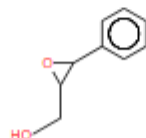
metabolite #6
No CAS number



metabolite #7
No CAS number



metabolite #8
No CAS number



Metabolism/Transformations
 Skin metabolism simulator

8 metabolite(s)

Cinnamyl Formate (104-65-4)

Profiling results

- 8 metabolite(s)
 - 2 x Skin sensitization Category 1B
 - 2 x Epoxides, Aziridines and Sulfuranes_low activity
 - O=COCC1OC1c1ccccc1
 - OCC1OC1c1ccccc1
 - 6 x No alert found
 - OC(=O)OCC=Cc1ccccc1
 - OC(COC=O)C(O)c1ccccc1
 - OC(O)=O
 - OC=O
 - OCC(O)C(O)c1ccccc1
 - OCC=Cc1ccccc1

General Mechanism

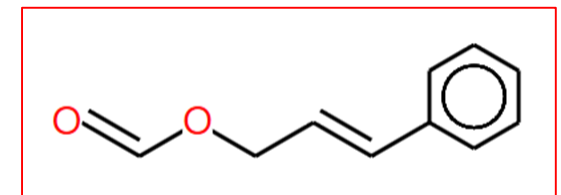
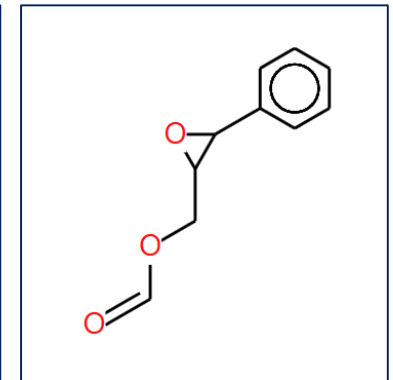
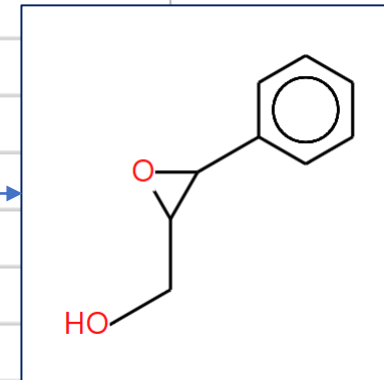
- Protein binding
- Protein binding

Endpoint Specific

- Protein binding
- Protein binding

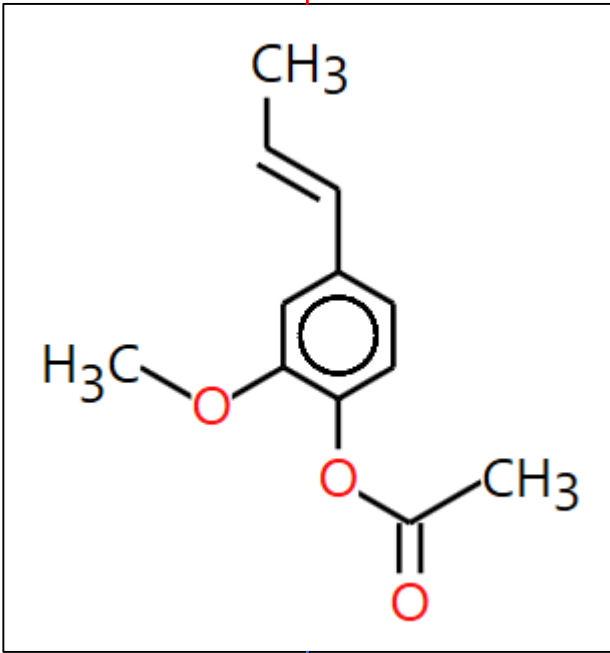
Metabolism/Transformation

- Skin metabolism
 - General Mechanism
 - Protein binding
 - Protein binding by OECD
 - Endpoint Specific
 - Protein binding alerts for skin sensitization according to GHS
 - Protein binding alerts for skin sensitization by OASIS



This is in contrast to the known liver metabolism in which cinnamic aldehyde is produced via the action of esterases followed by alcohol dehydrogenase. We might consider category 1B following metabolism to be a non-sensitizer.

Isoeugenol Acetate (93-29-8)



Metabolism/Transformations

Options

f	Select All	Unselect All	Invert
<input type="checkbox"/>			
<input type="checkbox"/>			
<input checked="" type="checkbox"/>			
<input type="checkbox"/>			

Microbial metabolism simulator
Rat liver S9 metabolism simulator
Skin metabolism simulator
Tautomerism

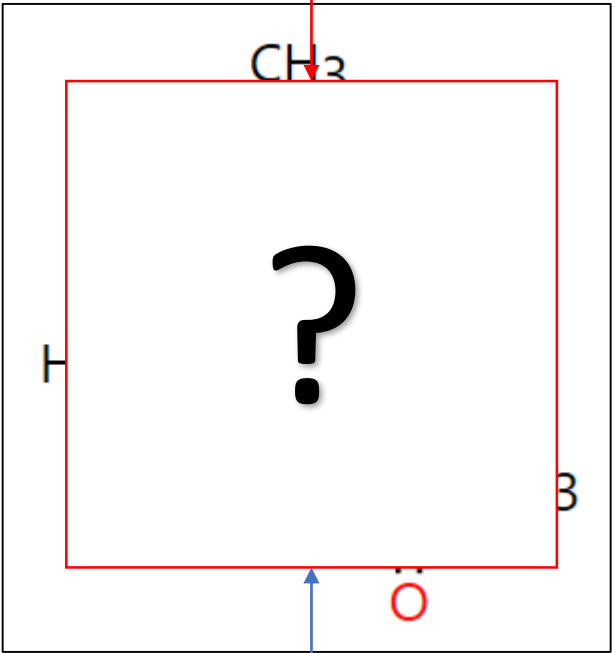
Metabolism versus hydrolysis

Metabolism/Transformations

Options

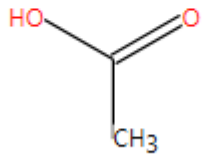
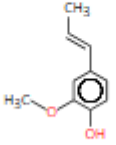
f	Select All	Unselect All	Invert
<input type="checkbox"/>			
<input type="checkbox"/>			
<input type="checkbox"/>			
<input checked="" type="checkbox"/>			

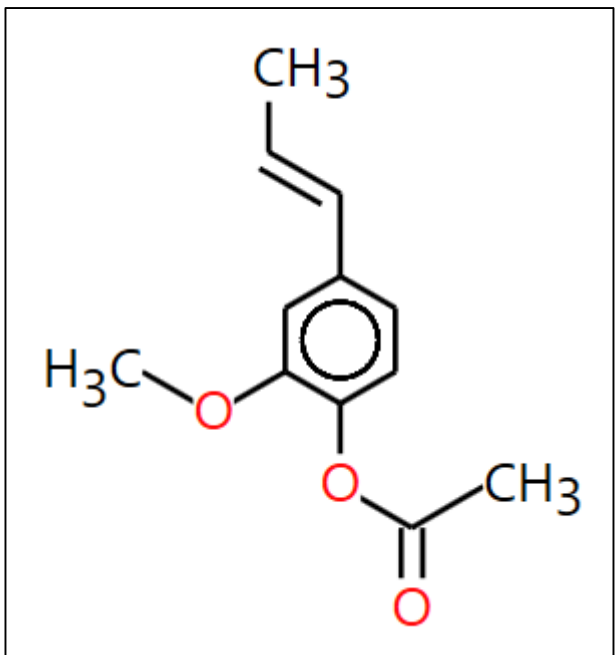
Dissociation simulator
Hydrolysis simulator (acidic)
Hydrolysis simulator (basic)
Hydrolysis simulator (neutral)

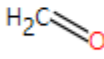
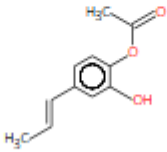



Isoeugenol Acetate (93-29-8)

Profile	
General Mechanistic	
Protein binding by OASIS	No alert found
Protein binding by OECD	Acylation
Endpoint Specific	
Protein binding alerts for skin sensitiz...	No alert found
Protein binding alerts for skin sensitiz...	No alert found
Metabolism/Transformations	
Hydrolysis simulator (neutral)	2 metabolite(s)
Skin metabolism simulator	3 metabolite(s)

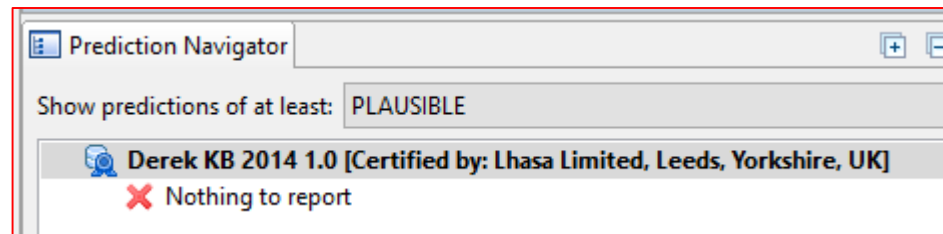
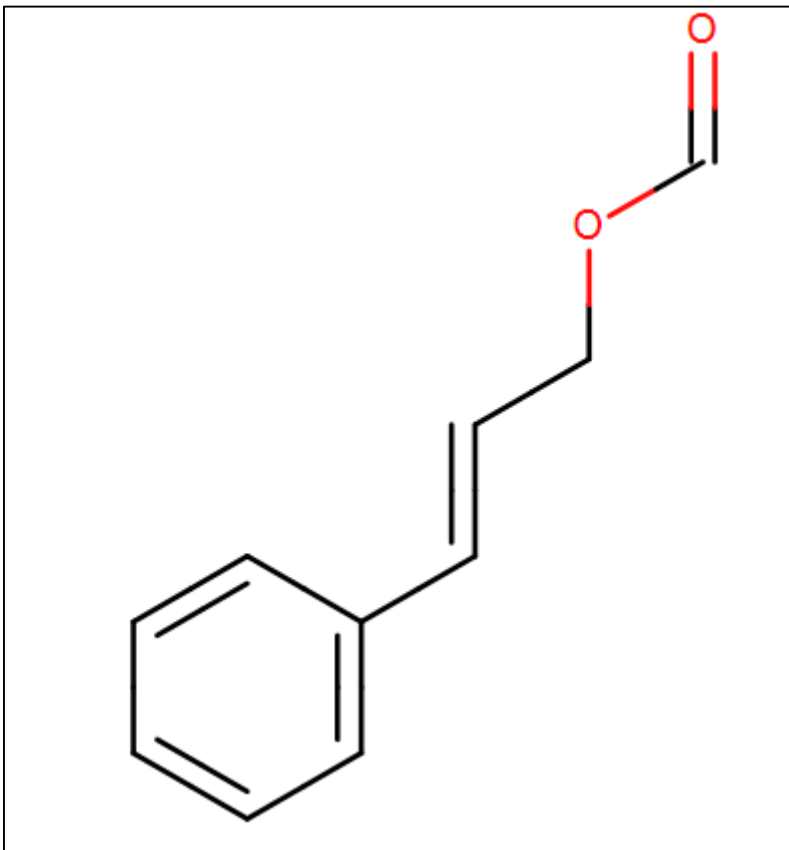
metabolite #1 No CAS number	metabolite #2 No CAS number
	



metabolite #1 No CAS number	metabolite #2 No CAS number	metabolite #3 No CAS number
		

Summary: OECD QSAR Toolbox

- The examples we have looked at showed that we can identify potential electrophiles due to skin metabolism and hydrolysis
- Profilers are available for the prediction of skin metabolism, hydrolysis and dissociation
- However, we cannot offer any indication of the rate of these reactions or the ratios in which the metabolites exist (probability/reliability scores are available in TIMES)
- Knowledge bases are only as good as the data on which they are built



Derek Nexus returns 'nothing to report' if there is no information about the chemical of interest within the knowledge base

The next step is to investigate metabolism

Biotransformations

Filter biotransformations...

- Phase I
 - Redox
 - Non-Redox
- Phase II
 - Glucuronidation
 - Glutathione Conjugation
 - Sulphonation
 - Acetylation
 - Methylation
 - Conjugation with Amino Acids
 - Conjugation with Other Acids
 - Glucosidation

Select All Deselect All

Enzyme -- Select an enzyme --

- Phase I
 - Redox
 - Oxidation
 - Reduction
 - Non-Redox
 - Hydrolytic Reactions
 - Dehydration
 - Hydration
 - Hydrolytic Fragmentation or Ring-Opening
 - Non Hydrolytic Fragmentation or Ring-Opening
 - Ring-Closure Reactions
 - Other Elimination Reactions
 - Decarboxylation
 - Miscellaneous

- Phase II
 - Glucuronidation
 - O-Glucuronidation
 - N-Glucuronidation
 - S-Glucuronidation
 - Glutathione Conjugation
 - Alkyl Halides
 - Halogenated Alkenes
 - Alkynes
 - Esters
 - Epoxides and Episulphides
 - Arene Oxides
 - Oxygen-Containing Functional Groups
 - Nitrogen-Containing Functional Groups
 - alpha,beta-Unsaturated Compounds
 - Aromatic Compounds


- Sulphonation
 - O-Sulphonation
 - N-Sulphonation
- Acetylation
 - N-Acetylation
- Methylation
 - O-Methylation
 - N-Methylation
 - S-Methylation
- Conjugation with Amino Acids
 - Glycine
 - Glutamine
 - Taurine
 - Lysine
 - Cystine
- Conjugation with Other Acids
 - Pyruvic Acid
 - alpha-Ketoglutaric Acid

- Glucosidation
 - [439] Glucosidation of Alcohols
 - [440] Glucosidation of Carboxylic Acids
 - [441] Glucosidation of Primary Aromatic Amines
 - [442] Glucosidation of Barbiturates
 - [443] Glucosidation at Aromatic Nitrogen

The benefit of this approach is that you can design your own profiling scheme for enzymes/biotransformations of interest


In this example we have selected oxidation and reduction phase 1 transformations only

Cinnamyl Formate (104-65-4)

 **Species**

Filter species...

- Mammals
 - Dog
 - Primates
 - Rodents

 **Processing Options**

Processing direction: Breadth first Depth first User defined


Phase constraints: Grow from phase II products

Max. Depth:

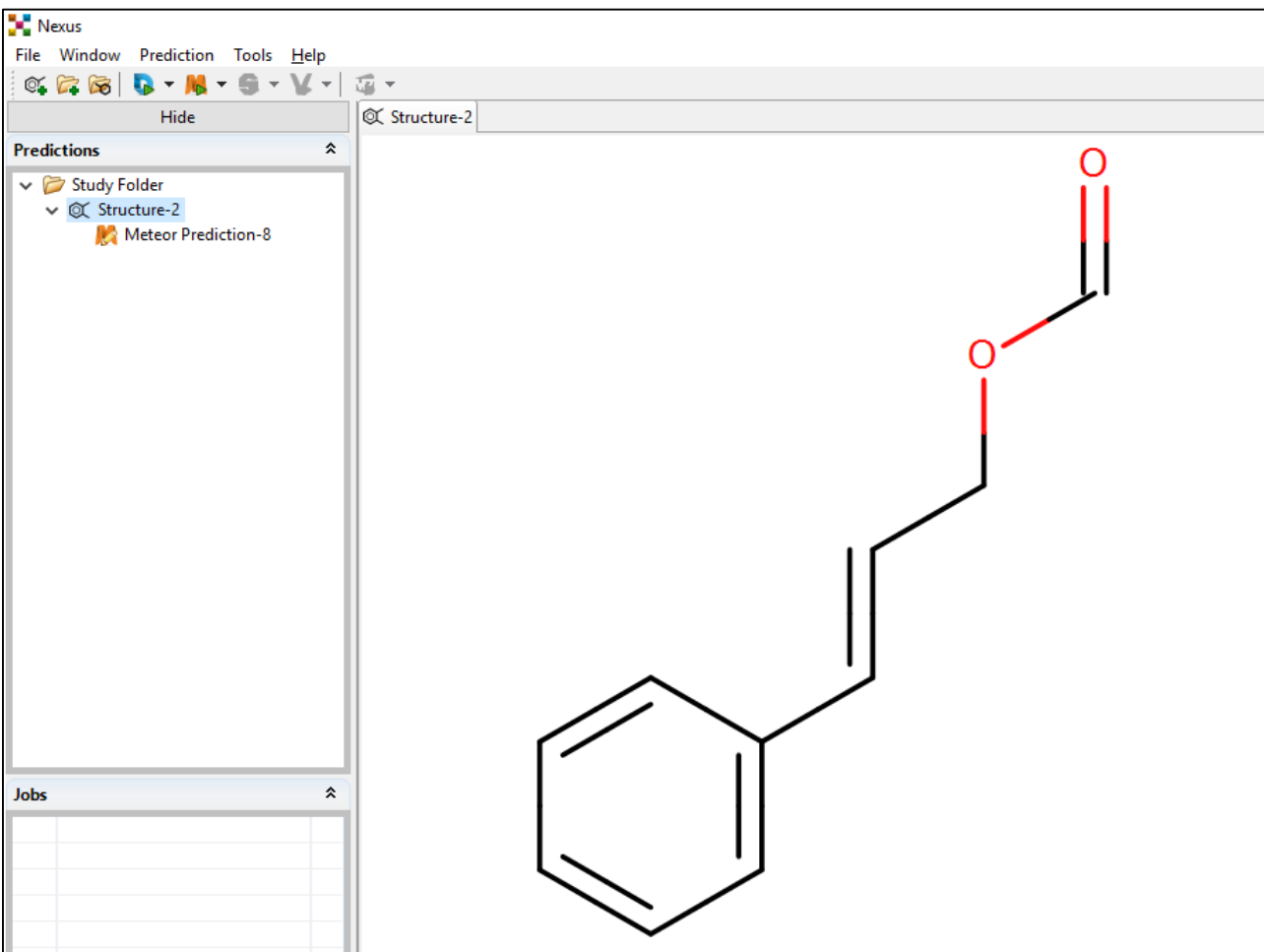
Max. Metabolites:

Min. Likelihood:

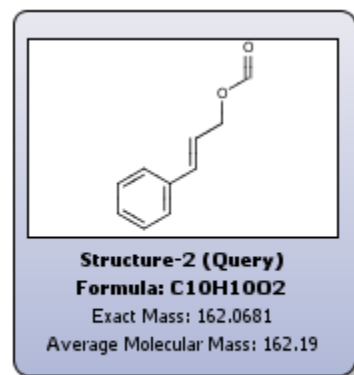
Apply Relative Reactions

 **Biotransformation**

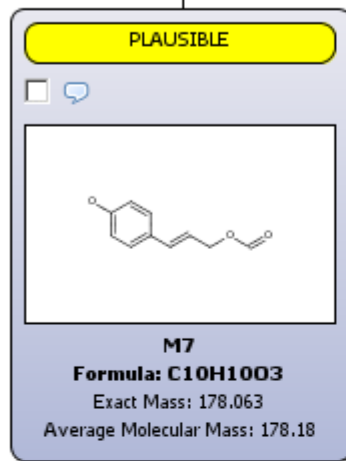
- PLAUSIBLE
- PROBABLE
- PLAUSIBLE
- EQUIVOCAL
- DOUBTED
- IMPROBABLE



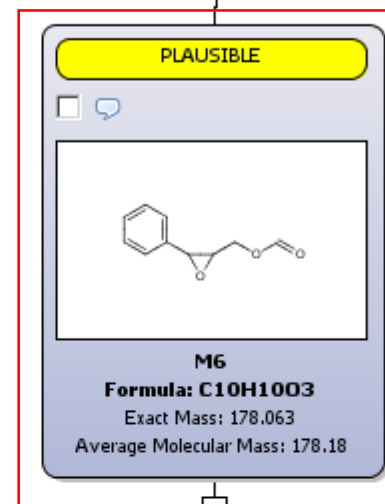
Cinnamyl Formate (104-65-4)



078 Para Hydroxylation of Monosubstituted Benzene Compounds

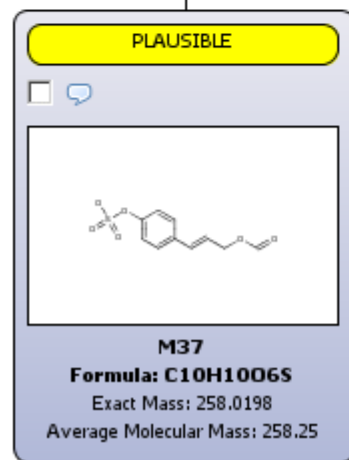


365 Epoxidation of Z-1,2-Disubstituted Alkenes

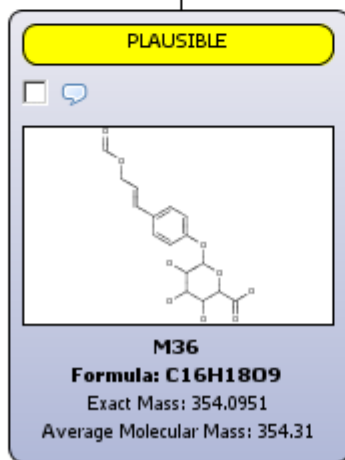


Phase I metabolism

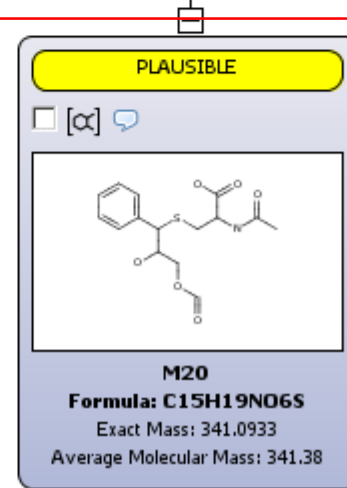
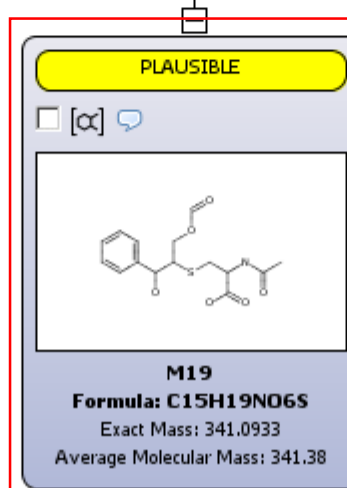
020 O-Sulphation of Aromatic Alcohols



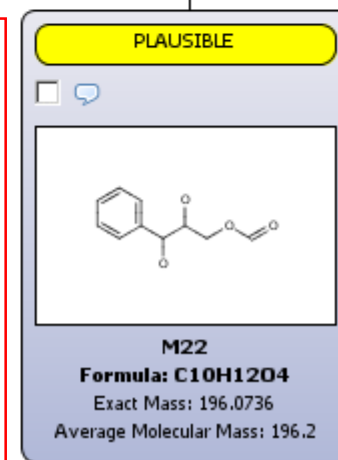
027 Glucuronidation of Aromatic Alcohols



058 Conjugation of Epoxides with Glutathione

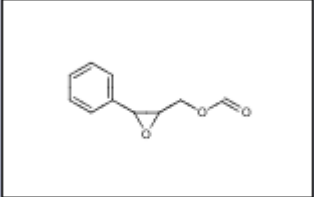


089 Vicinal Diols from Epoxides

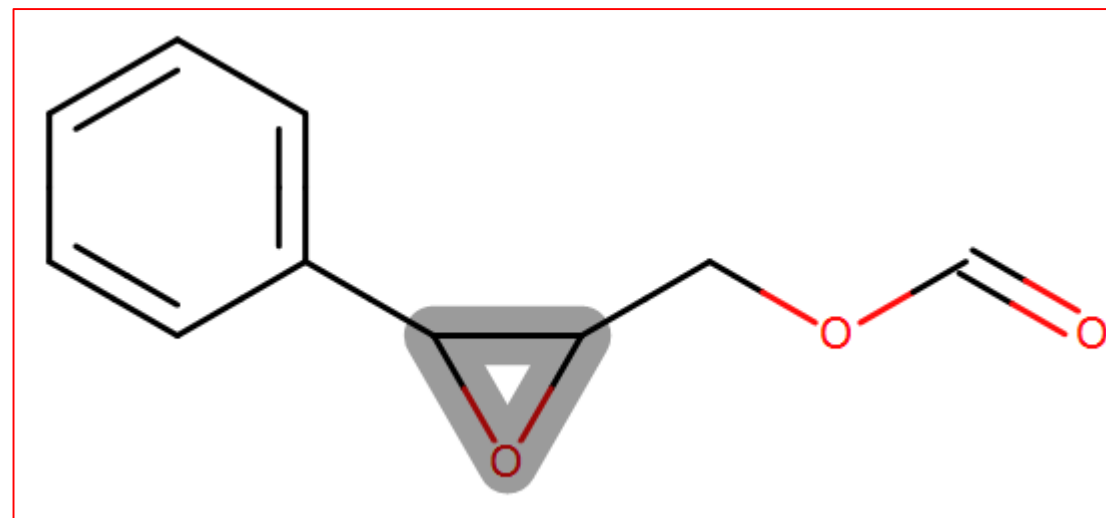


Phase II metabolism

PLAUSIBLE



M6
Formula: C10H10O3
Exact Mass: 178.063
Average Molecular Mass: 178.18



Endpoints

Derek KB 2014 1.0

Filter endpoints...

- Carcinogenicity (ALL)
- Genotoxicity (ALL)
- Irritation (ALL)
- Miscellaneous endpoints (ALL)
- Neurotoxicity (ALL)
- Organ toxicity (ALL)
- Reproductive toxicity (ALL)
- Respiratory sensitisation (ALL)
- Skin sensitisation (ALL)
 - Photoallergenicity
 - Skin sensitisation

Species

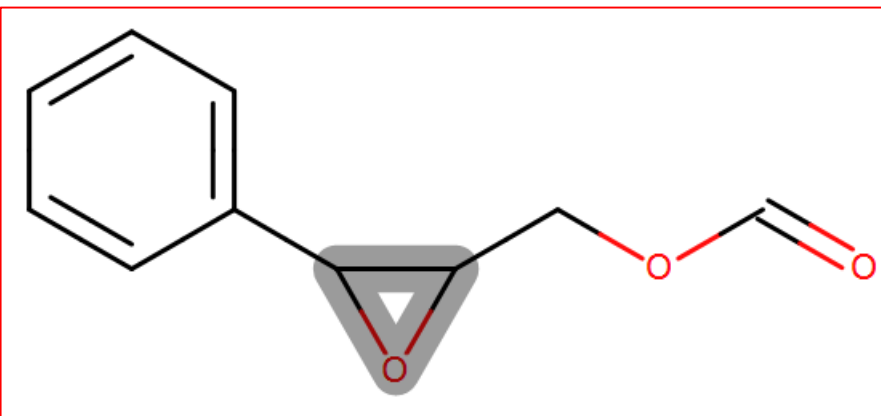
Filter species...

- bacterium
 - Escherichia coli
 - Salmonella typhimurium
- chicken
- mammal
 - dog
 - primate
 - human
 - monkey
 - rabbit
- rodent
 - guinea pig
 - hamster
 - mouse
 - rat
- unspecified
- various

Prediction Navigator

Show predictions of at least:

- Derek KB 2014 1.0 [Certified by: Lhasa Limited, Leeds, Yorkshire, UK]
 - Skin sensitisation
 - mammal - PLAUSIBLE
 - Alert - 433: Epoxide



Comments

The activity of glycidyl ethers, amines, esters and amides is described in a separate alert.

The presence of a skin sensitisation structural alert within a molecule indicates the molecule has the potential to cause skin sensitisation. Whether or not the molecule will be a skin

Validation Comments

Skin sensitisation: guinea pig maximisation test, local lymph node assay

The alert has demonstrated the following predictive performance:

1) Cronin and Basketter data set: 2 compounds activate this alert of which 2 are reported positive (positive predictivity = 100%)

Endpoints

ID	Name	Parent
4189	Skin sensitisation	Skin sensitisation (ALL)

References

ID	Title	Author	Source	Year	Supplier
4853	Multivariate QSAR analysis of a skin sensitization databas	Cronin MTD and Basketter DA.	SAR and QSAR in Environmental Rese	1994	Note: d
4914	The use of nuclear magnetic resonance spectrometry (1H	Betso JE, Carreon RE and Miner VM.	Toxicology and Applied Pharmacolog	1991	

Examples

3-phenyl-oxirane-2-carboxylic acid

-methyl-3-phenyl-oxiranecarboxyli

Prediction Navigator

Show predictions of at least: **PLAUSIBLE**

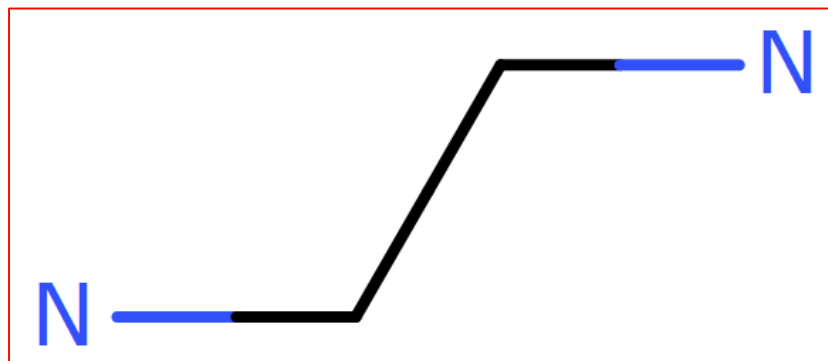
- Derek KB 2014 1.0** [Certified by: Lhasa Limited, Leeds, Yorkshire, UK]
 - Skin sensitisation**
 - mammal - PLAUSIBLE**
 - Alert - 433: Epoxide**

Summary: Meteor and Derek Nexus

- The example we have looked at showed that we can identify potential electrophiles due to skin metabolism
- The knowledge base in Meteor Nexus identifies potential metabolites, these can be screened for skin sensitisation potential using Derek Nexus
- Derek Nexus also contains structural alerts for some pro-electrophiles
- Predictions in both systems are given a level of confidence; however, potential biotransformation rates and ratios are not available
- Knowledge bases are only as good as the data on which they are built

OECD QSAR Toolbox & Nexus

- Occasionally these two leading tools will give differing profiling results
- This is usually around whether an alert has been encoded as a part of the 'toxicity' or 'metabolism' knowledge
- For example, consider profiling ethylenediamine (a pro-electrophile from the Patlewicz et al study)



Derek Nexus contains direct and indirect alerts for skin sensitisation in its knowledge base

Ethylenediamine
(107-15-3)

Prediction Navigator

Show predictions of at least:

Derek KB 2014 1.0 [Certified by: Lhasa Limited, Leeds, Yorkshire, UK]

- Skin sensitisation
 - mammal - PROBABLE
 - Alert - 435: Diamine
 - Example - 1,2-diaminoethane

Comments

The presence of a skin sensitisation structural alert within a molecule indicates the molecule has the potential to cause skin sensitisation. The presence of a skin sensitisation structural alert within a molecule indicates the molecule has the potential to cause skin sensitisation. The presence of a skin sensitisation structural alert within a molecule indicates the molecule has the potential to cause skin sensitisation. Generally, small lipophilic molecules are more readily absorbed into the skin.

Validation Comments

Skin sensitisation: guinea pig maximisation test, local lymph node assay

The alert has demonstrated the following predictive performance:

1) Cronin and Basketter data set: 1 compound activates this alert of which 1 is reported positive (positive predictivity = 100%)

References

ID	Title	Author
4847	Chemikalien und Kontaktallergie: Eine Bewertende Zusammenfassung	Kayser D and Schlede E (editors).
4853	Multivariate QSAR analysis of a skin sensitization database	Cronin MTD and Basketter DA.
4868	A comparison of three guinea-pig sensitization procedures	Goodwin BFJ, Crevel RWR and Johnstone IB.
4910	Contact allergy to impurities in surfactants: amount, chemical structure and skin sensitization	Angelini G, Riquano L, Foti C, Vena GA.

Ethylenediamine (107-15-3)

QSAR TOOLBOX

Documents

Document 1

[Search chemical](#)

Profiling methods

Options

f	Select All	Unselect All	Invert
<input type="checkbox"/>	Oncologic Primary Classification		
<input type="checkbox"/>	Protein binding alerts for Chromosomal		
<input checked="" type="checkbox"/>	Protein binding alerts for skin sensitizat		
<input checked="" type="checkbox"/>	Protein binding alerts for skin sensitizat		

Metabolism/Transformations

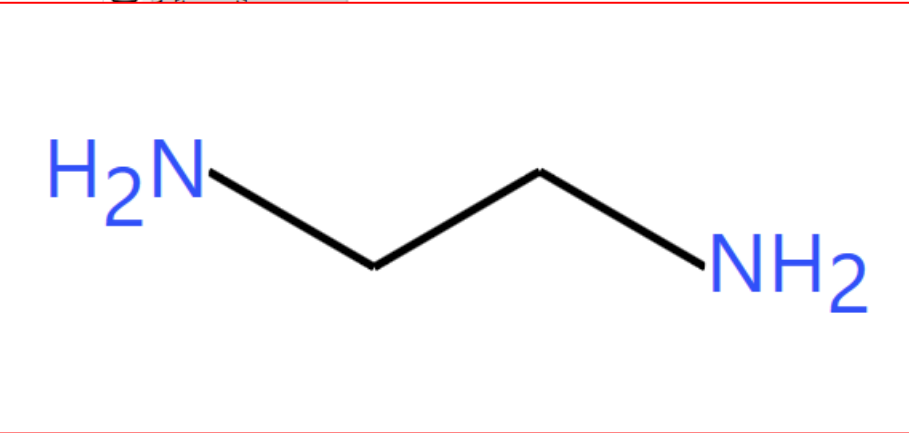
Options

f	Select All	Unselect All	Invert
<input type="checkbox"/>	Microbial metabolism simulator		
<input type="checkbox"/>	Rat liver S9 metabolism simulator		
<input checked="" type="checkbox"/>	Skin metabolism simulator		
<input type="checkbox"/>	Tautomerism		

Filter endpoint tree...

Structure

- Structure info
- Parameters
- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazards
- Profile
 - General Mechanistic
 - Protein binding by OASIS
 - Protein binding by OECD
 - Endpoint Specific
 - Protein binding alerts for skin sensitization according to GHS
 - Protein binding alerts for skin sensitization by OASIS
 - Metabolism/Transformations
 - Skin metabolism simulator



No alert found
No alert found
No alert found
No alert found
5 metabolite(s)

Ethylenediamine
(107-15-3)

QSAR TOOLBOX

Documents

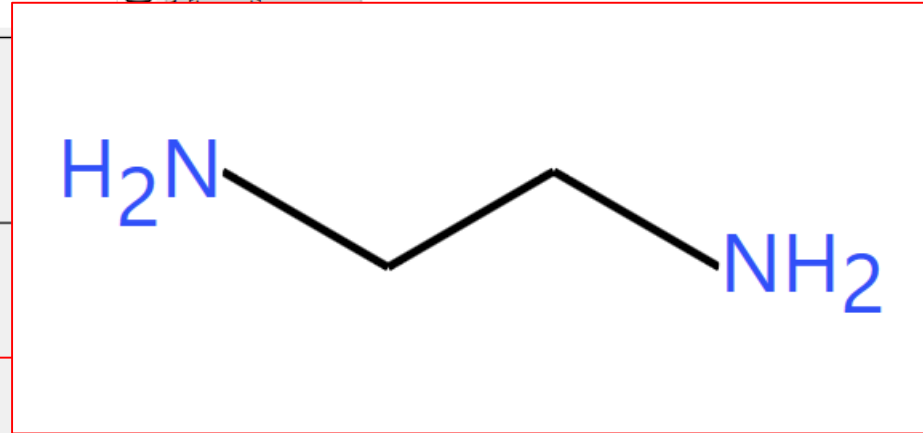
Document 1

[Search chemical](#)

Filter endpoint tree...

Structure

- Structure info
- Parameters
- Physical Chemical Properties



Profiling methods

Options

- [-] **Metabolism/Transformations**
 - [-] **Skin metabolism simulator**
 - [-] **General Mechanistic**
 - Protein binding by OASIS
 - Protein binding by OECD
 - [-] **Endpoint Specific**
 - Protein binding alerts for skin sensitization according to GHS
 - Protein binding alerts for skin sensitization by OASIS

5 metabolite(s)
1 x Schiff base formatio...
1 x Schiff Base Formers...
1 x Skin sensitization Ca...
1 x Schiff base formatio...

Profiling results

▲ 5 metabolite(s)

▲ 2 x No alert found

N

NCC(O)=O

▲ 2 x Skin sensitization Category 1B

▲ 2 x Aldehydes

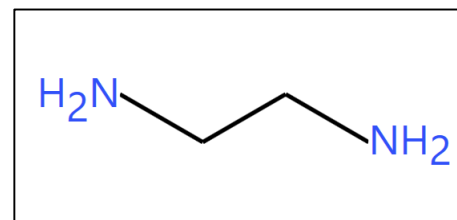
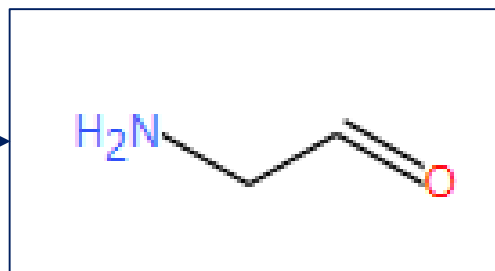
NCC=O

OC(=O)C=O

▲ 1 x Skin sensitization Category 1A

▲ 1 x Bis Aldehydes

O=CC=O



Ethylenediamine
(107-15-3)

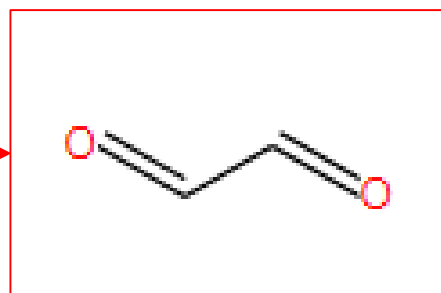
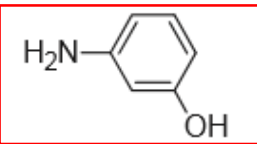
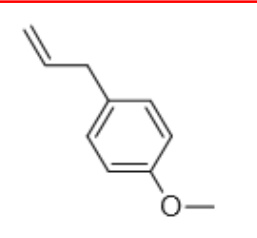
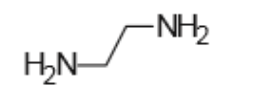
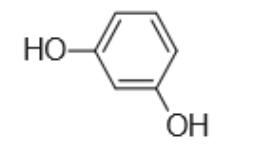
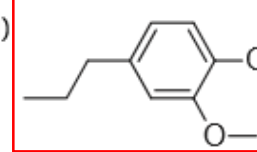
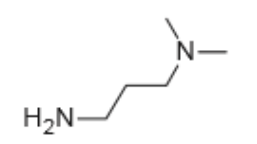
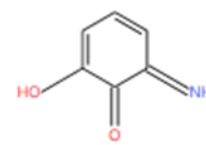
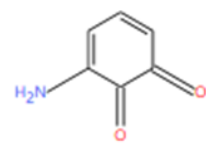


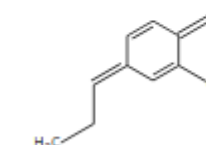
Table 2
Pro-haptens in the dataset.

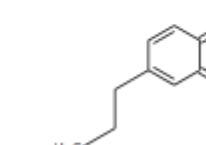
Name	Structure	In vivo		Cell free in chemico		Schiff base former		Michael acceptor	
		LLNA	Human data	DPP4	DPP4	SB	MA	SB	MA
3-Aminophenol		1	NA	0	0	1	0	1	0
4-Allylanisole		1	NA	1	0	1	0	1	0
Ethylenediamine (free base)		1	1	0	0	1	0	1	0
Resorcinol		1	1	0	0	0	0	1	0
Dihydroeugenol (2-methoxy-4-propyl-phenol)		1	NA	0	0	1	0	1	0
3-Dimethylamino propylamine		1	1	0	0	1	0	1	0

metabolite #1
No CAS number


metabolite #2
No CAS number


No metabolites predicted

metabolite #2
No CAS number


metabolite #3
No CAS number


Notes: NA – not available, 1 = positive, 0 = negative, SB = Schiff base former, MA = Michael acceptor.

^a 3-Dimethylamino propylamine was originally assigned as a pre-SB in the original EC EURL-ECVAM report (Casati et al., 2016). It has been reconsidered as a pro-SB electrophile.

How good are *in silico* systems for predicting skin metabolism (pro-haptens)?

Why is this?

Do we need more experiments?

Conclusions

- There are number of 'expert' systems available that can predict the potential of a chemical to react covalently with proteins
- The Nexus programmes are true expert systems, whilst we have used the knowledge within the OECD QSAR Toolbox as it were
- The encoded knowledge in both systems is very good at identifying potential direct and indirect electrophiles capable of causing skin sensitisation
- However, neither system can predict the rates of metabolite formation or the ratios of the various metabolites
- Lhasa products are commercial, whilst the OECD QSAR Toolbox is free