New, Transparent, Statistical Approaches to Toxicity Prediction

Tokyo / Osaka
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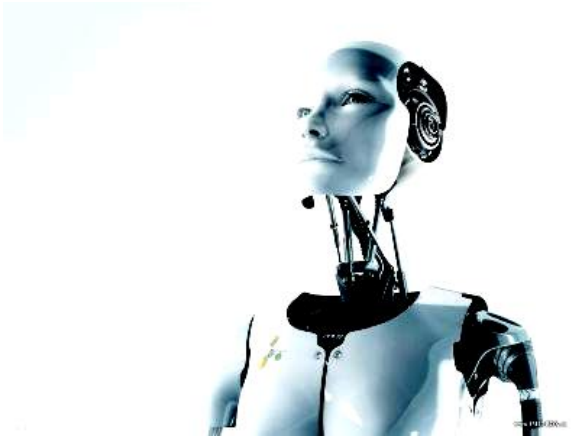
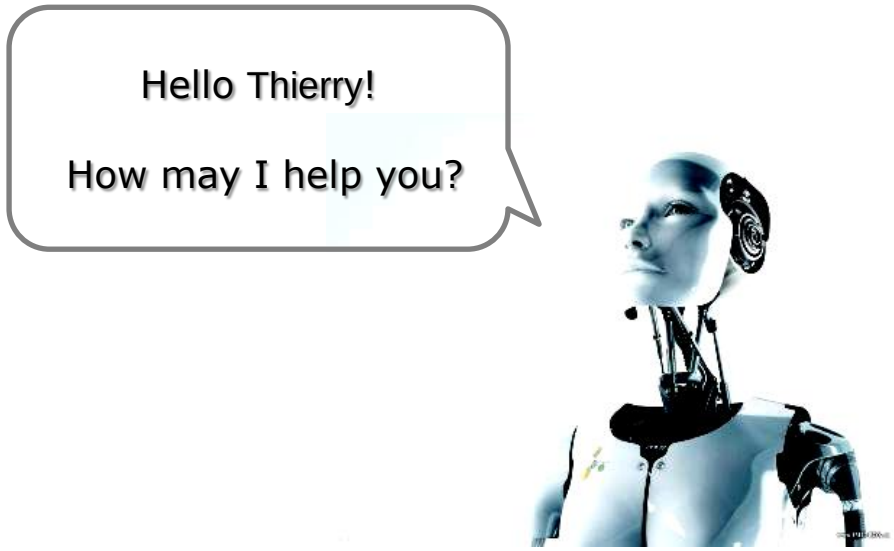
Toxicity prediction
The context of decision support

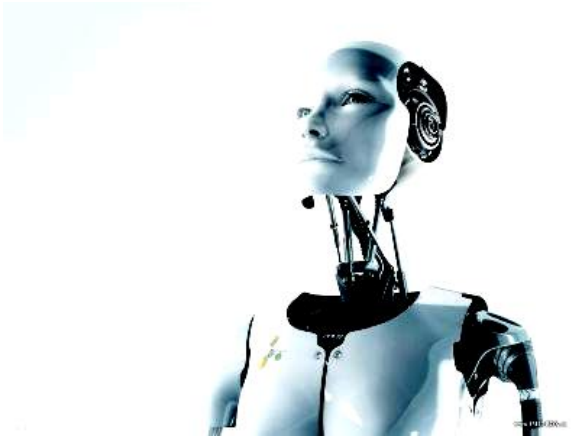
An ideal world

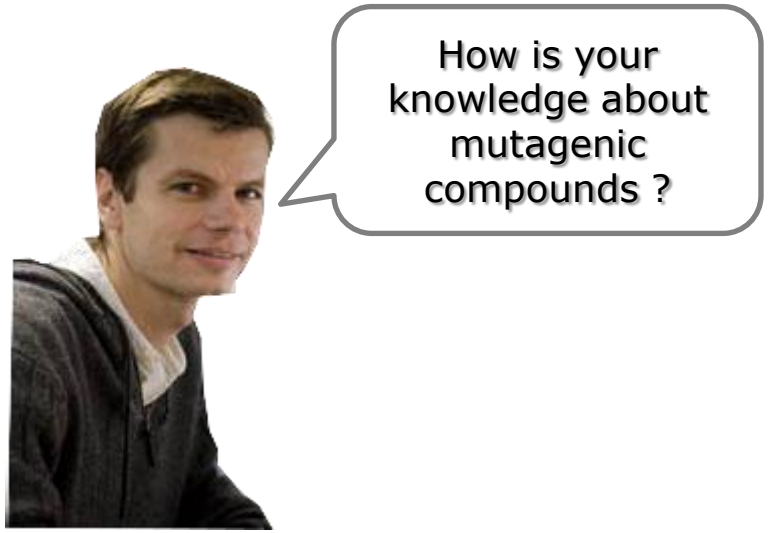


An ideal world



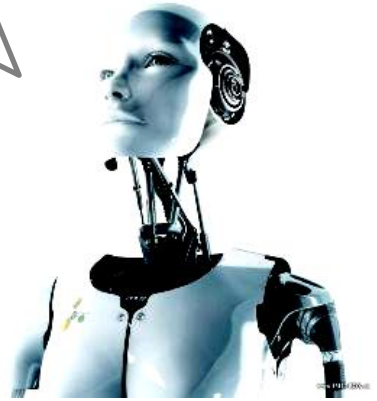
An ideal world



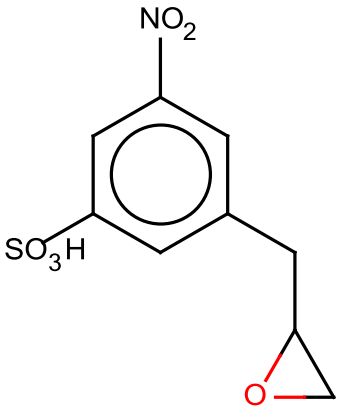


An ideal world

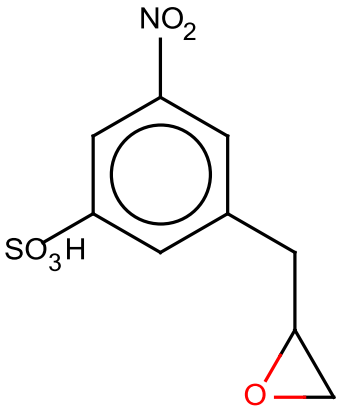
I have passed several tests with satisfactory results.



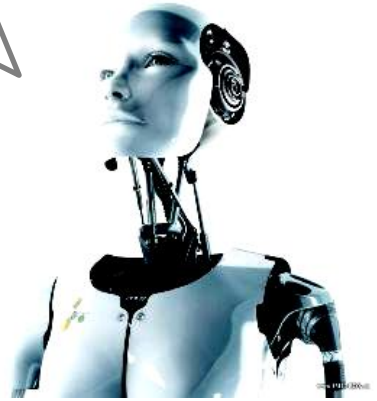
An ideal world



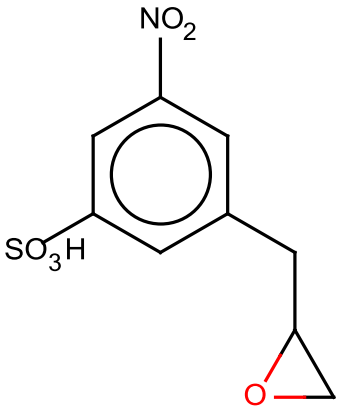
An ideal world



Yes, I have some understanding of this class of molecules.



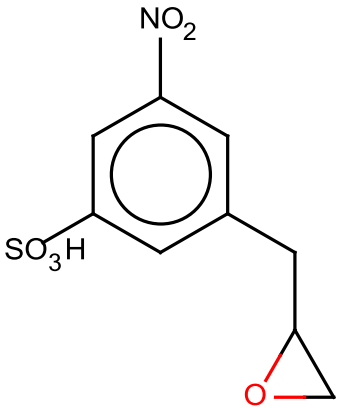
An ideal world



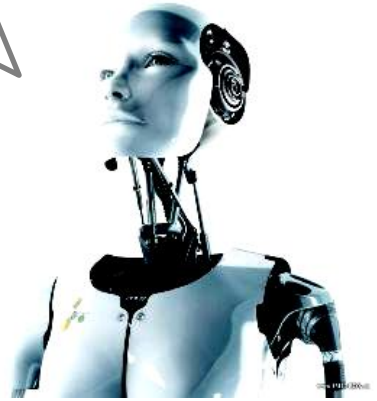
Great.

Could you please tell me if this compound is a mutagen?

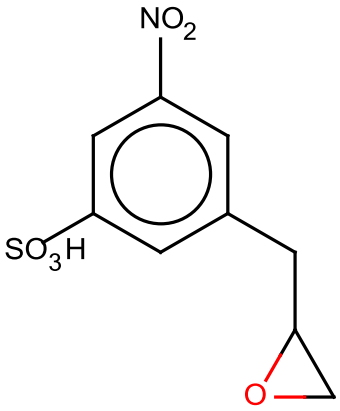
An ideal world

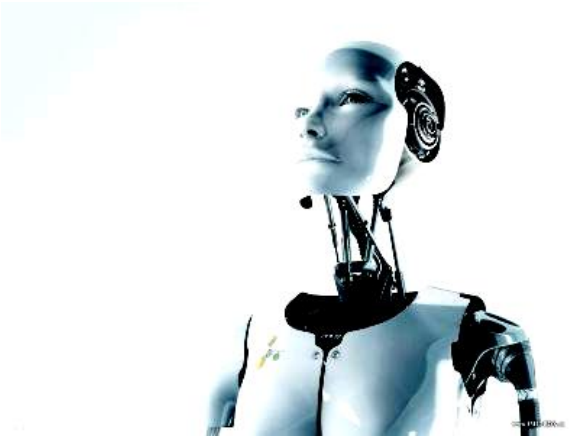


According to my knowledge it should be a mutagen; I am afraid so.

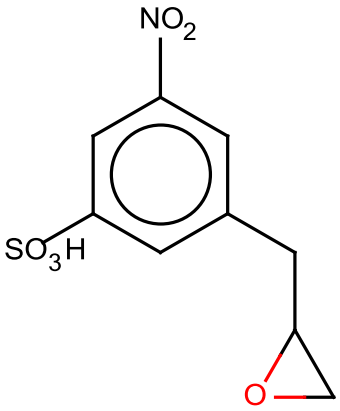


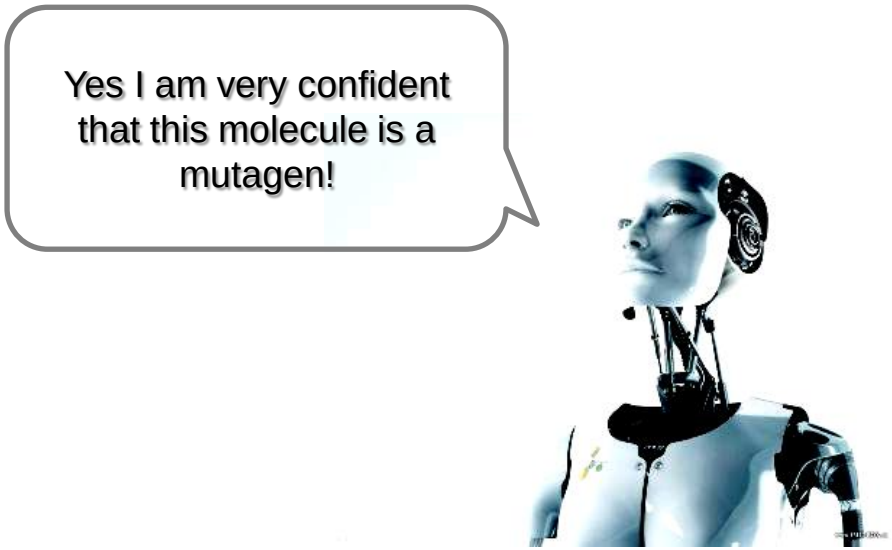
An ideal world





An ideal world





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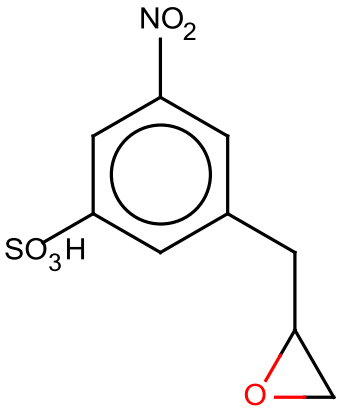
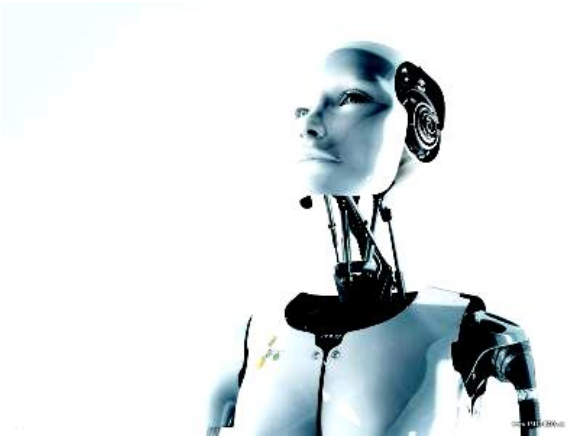
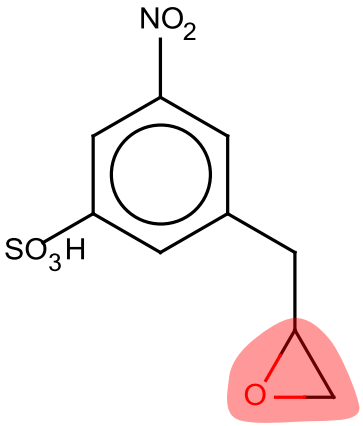
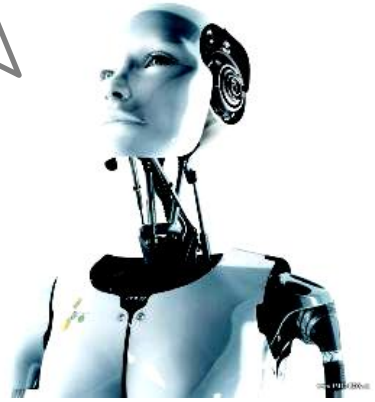




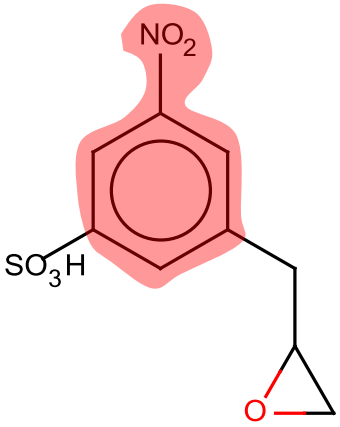
An ideal world



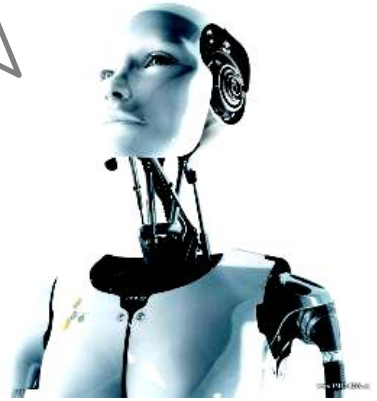
The main reason is the presence of an epoxide group



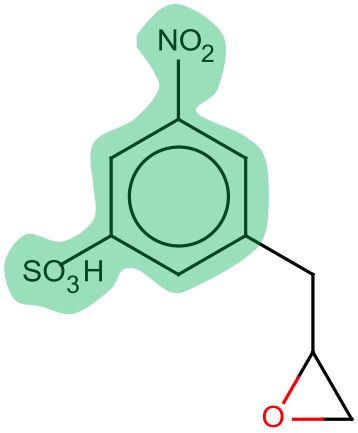
An ideal world



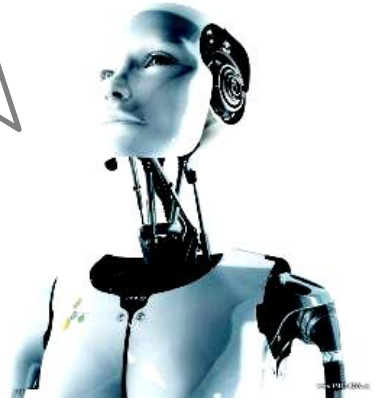
Additionally there is also a risk of mutagenicity due to the aromatic nitro group!



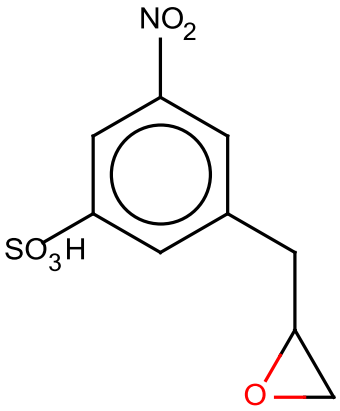
An ideal world

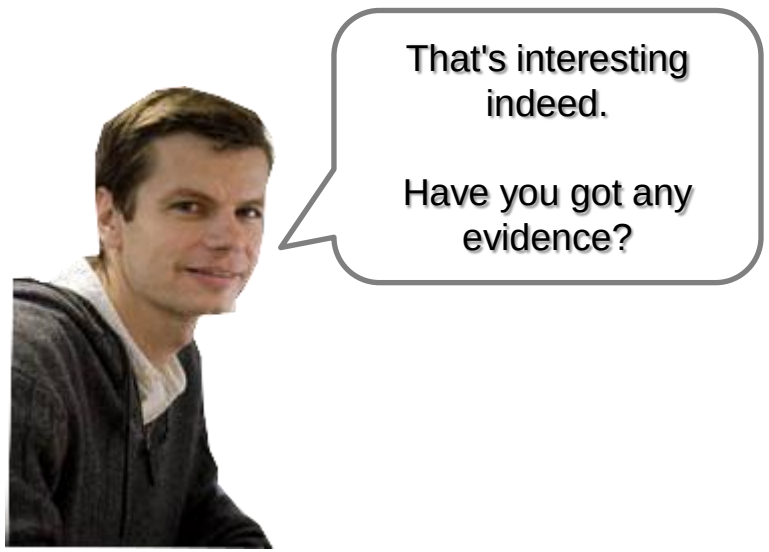


However, it seems that the sulfonic acid substituent deactivates the mutagenic effect of the nitro group



An ideal world





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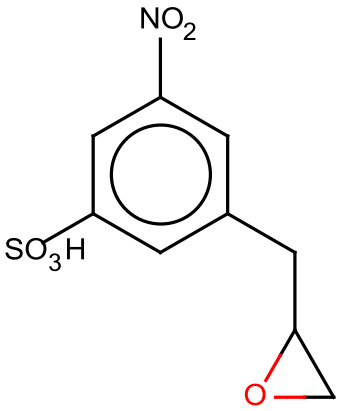
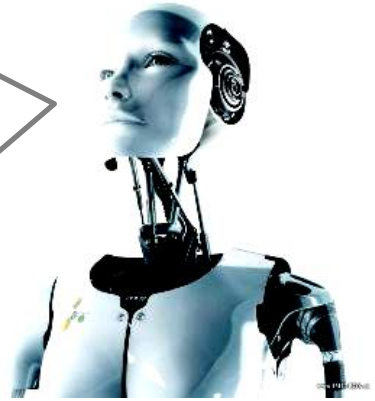
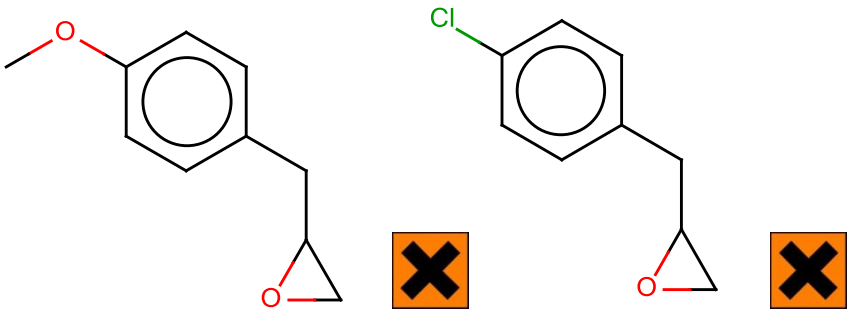


Yes.

Ames tests have shown two similar molecules containing epoxide groups as being mutagens





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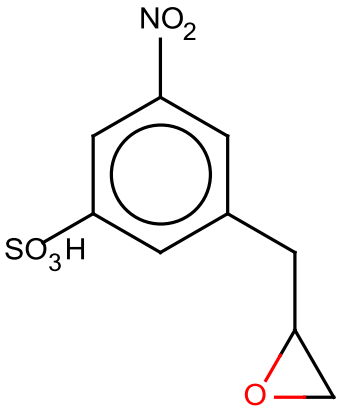
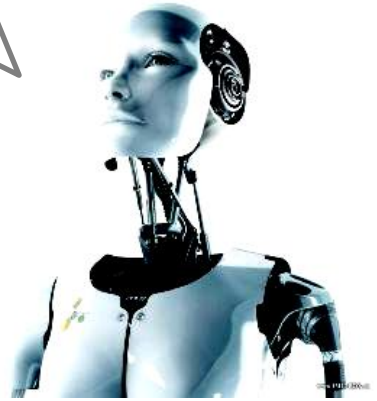
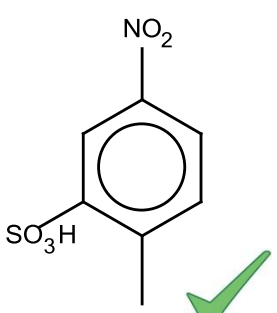
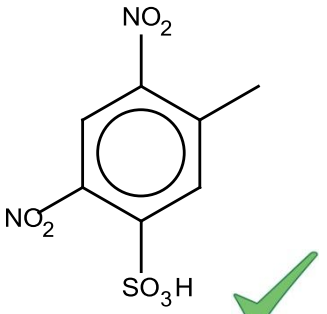


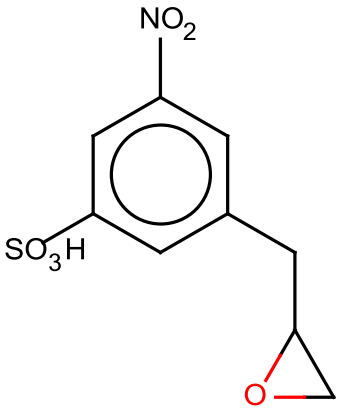
And these are examples of deactivated aromatic nitro groups







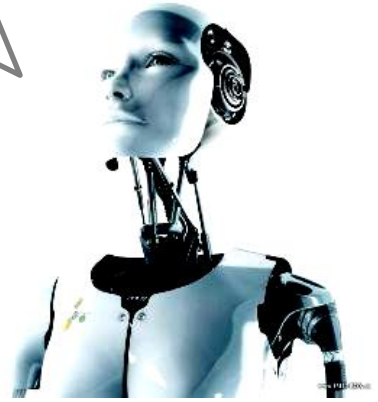
An ideal world



An ideal world

You are welcome.

Anything else I can help you with ?

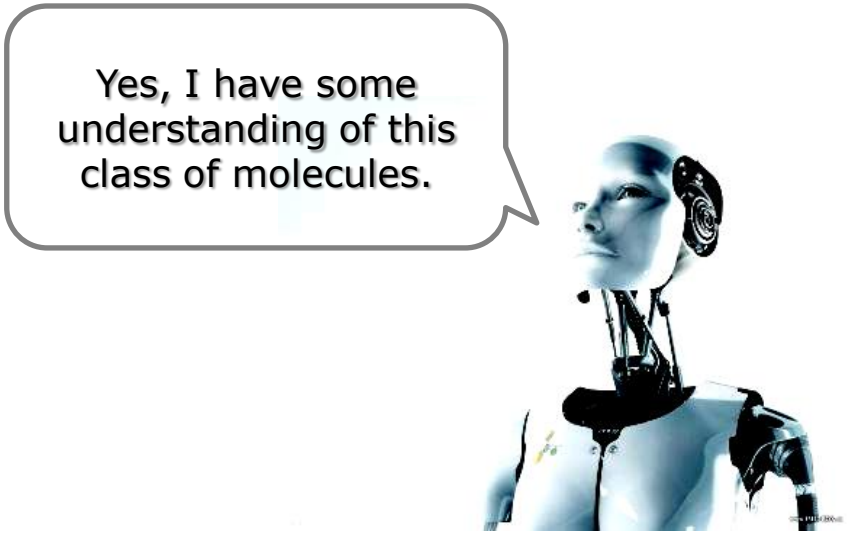


Model accuracy estimate

Model accuracy

Applicability domain

Model accuracy → Applicability Domain

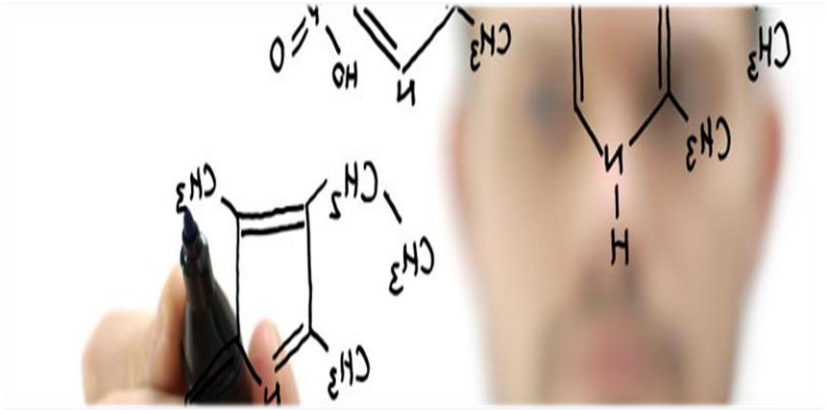


Individual prediction confidence

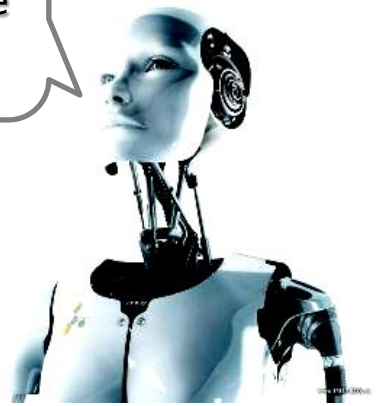
Model accuracy → Applicability Domain → Individual prediction confidence

Explanation

Model accuracy → Applicability Domain → Individual prediction confidence → **Interpretation**

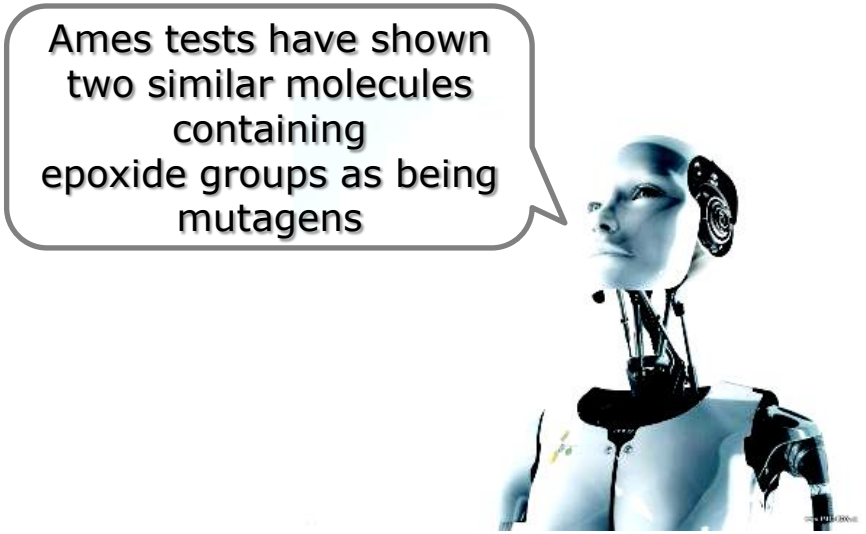


The main reason is probably the presence of an epoxide group



Supporting evidences

Model accuracy → Applicability Domain → Individual prediction confidence → Interpretation → Supporting Evidence



Interpretability accuracy trade-off

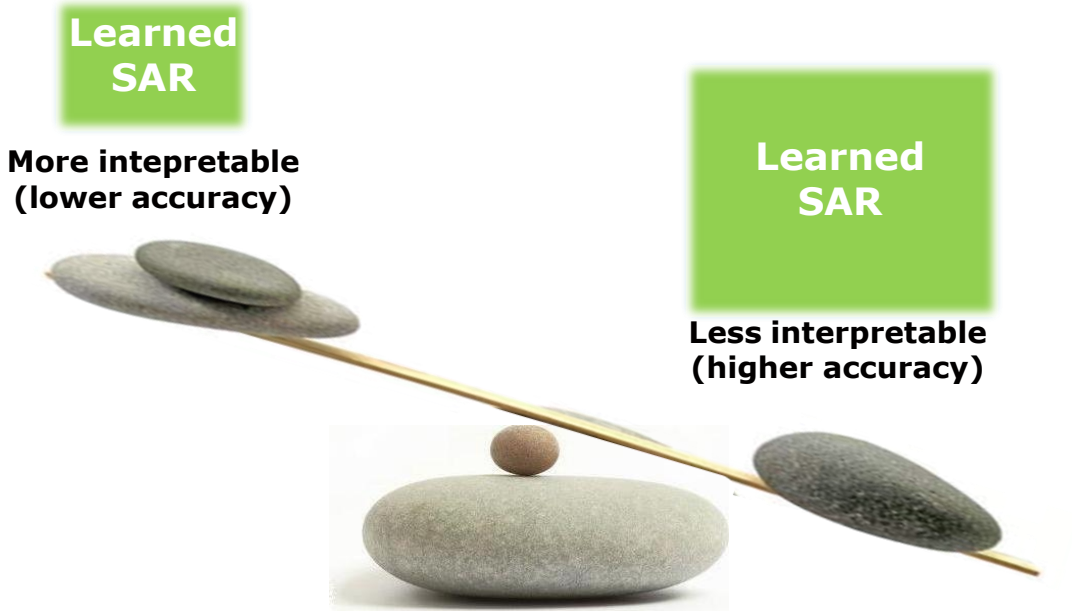
Learned SAR

More Accurate
(less interpretable)

Learned SAR

Less Accurate
(More interpretable)

Interpretability accuracy trade-off



Interpretability accuracy trade-off

Learned SAR

Useful trade-off depending on the use-case

Model vs Knowledge vs Facts

Virtual screening

Model accuracy → Applicability Domain → Individual prediction confidence → Interpretation → Supporting Evidence

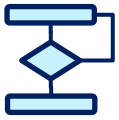
Lead Optimisation

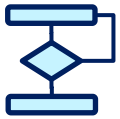
Model accuracy → Applicability Domain → Individual prediction confidence → Interpretation → Supporting Evidence

Regulatory decision process

Model accuracy → Applicability Domain → Individual prediction confidence → Interpretation → Supporting Evidence

Opaque Models	**Interpretable Models**	**Expert Knowledge**	**Facts**
SVM/RF/ANN/etc.	**DT, LR, MLR, PLS**	**Expert Systems**	**Databases**

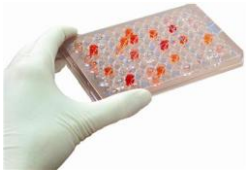


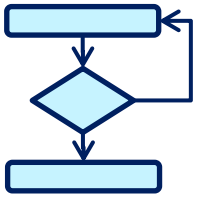


A priori knowledge unification
(**S**elf **O**rganizing **H**ypothesis **N**etworks)

Combining knowledge before prediction

Towards a unified framework





Facts, Expert Knowledge
Machine Learning

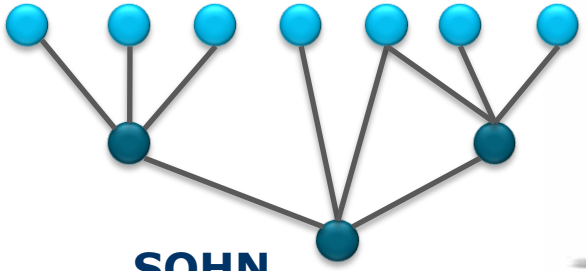
Knowledge Uniformisation

1

Simple interpretable hypotheses

2

Knowledge Organisation

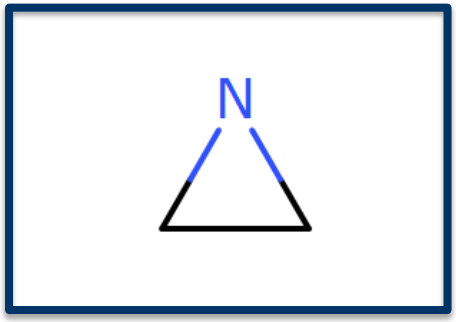


SOHN

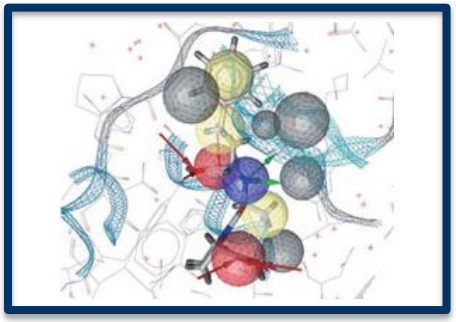
Use the SOHN structured knowledge

3

Hypothesis types



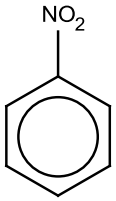
Structural Hypothesis



Pharmacophore Hypothesis

$2 < \log P < 3$

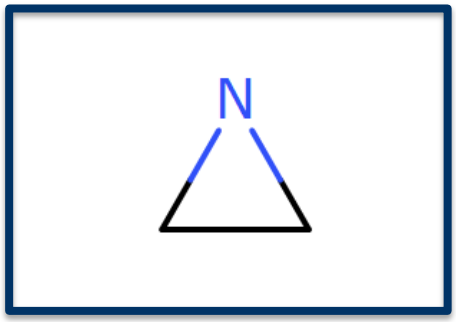
Physico-Chemical Hypothesis



$\tau > 0.8$

Similarity Hypothesis

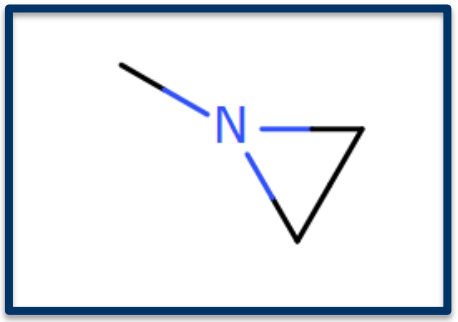
Hypotheses hierarchy



A

>

more general than



B

1 < logP < 4

C

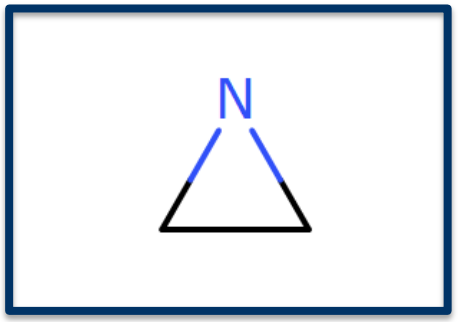
>

more general than

2 < logP < 3

D

Hypotheses hierarchy

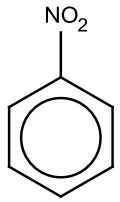


A ? more general than B

2 < logP < 3

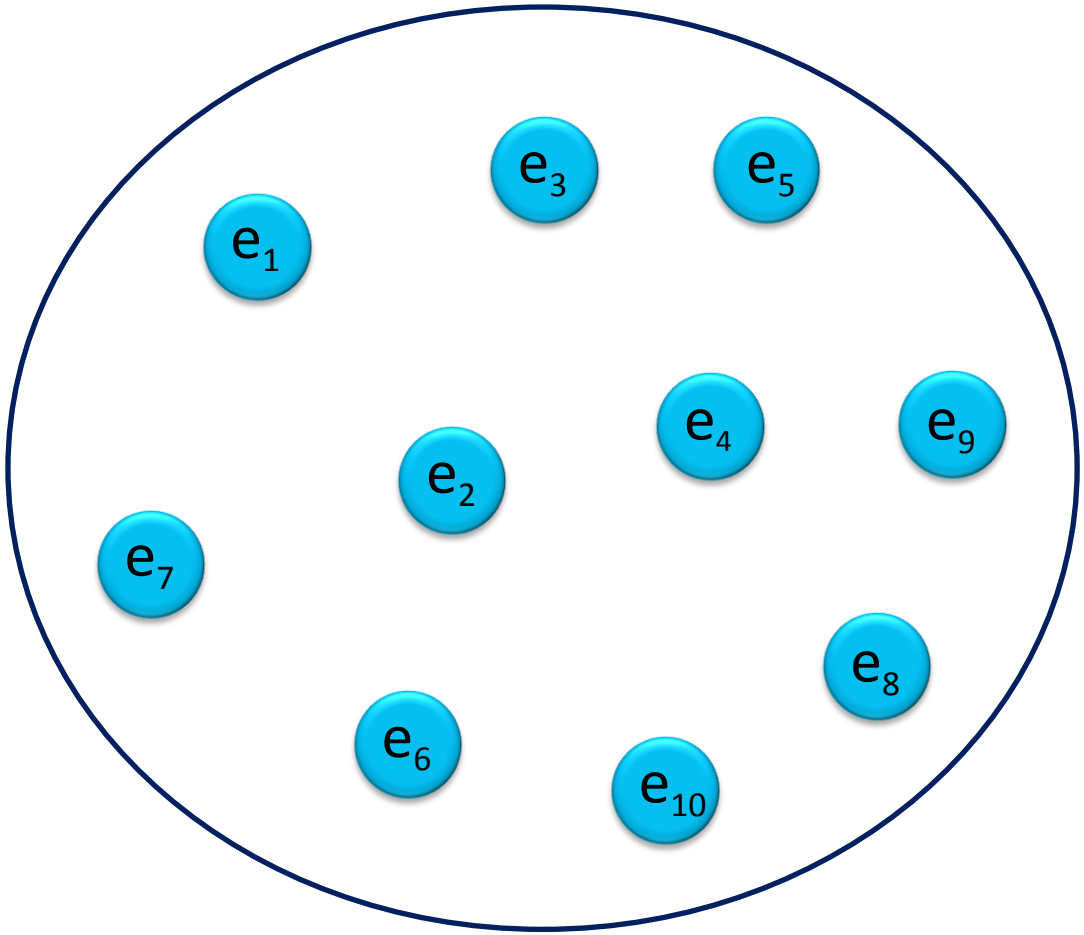
1 < logP < 4

C ? more general than D



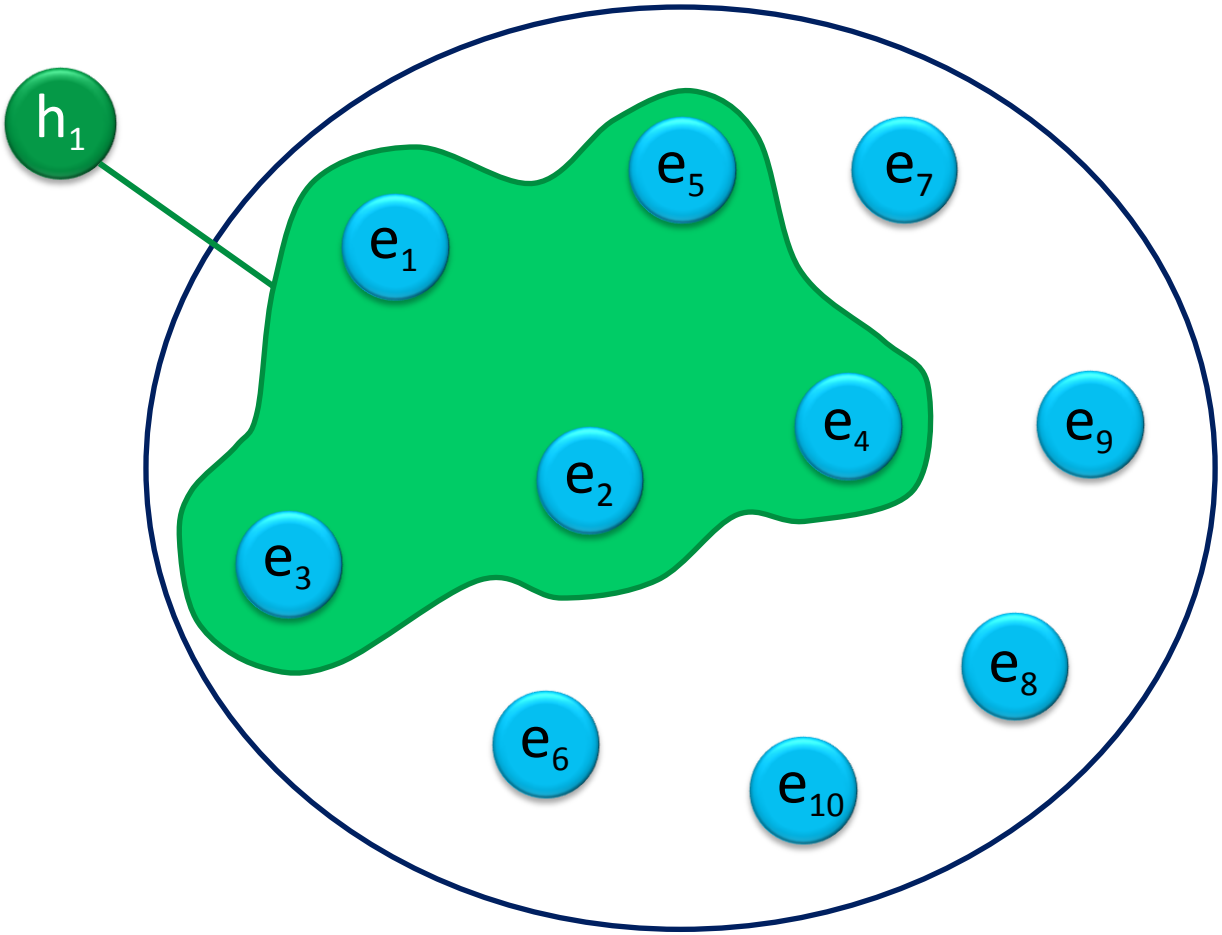
$\tau > 0.8$

Hypotheses hierarchy



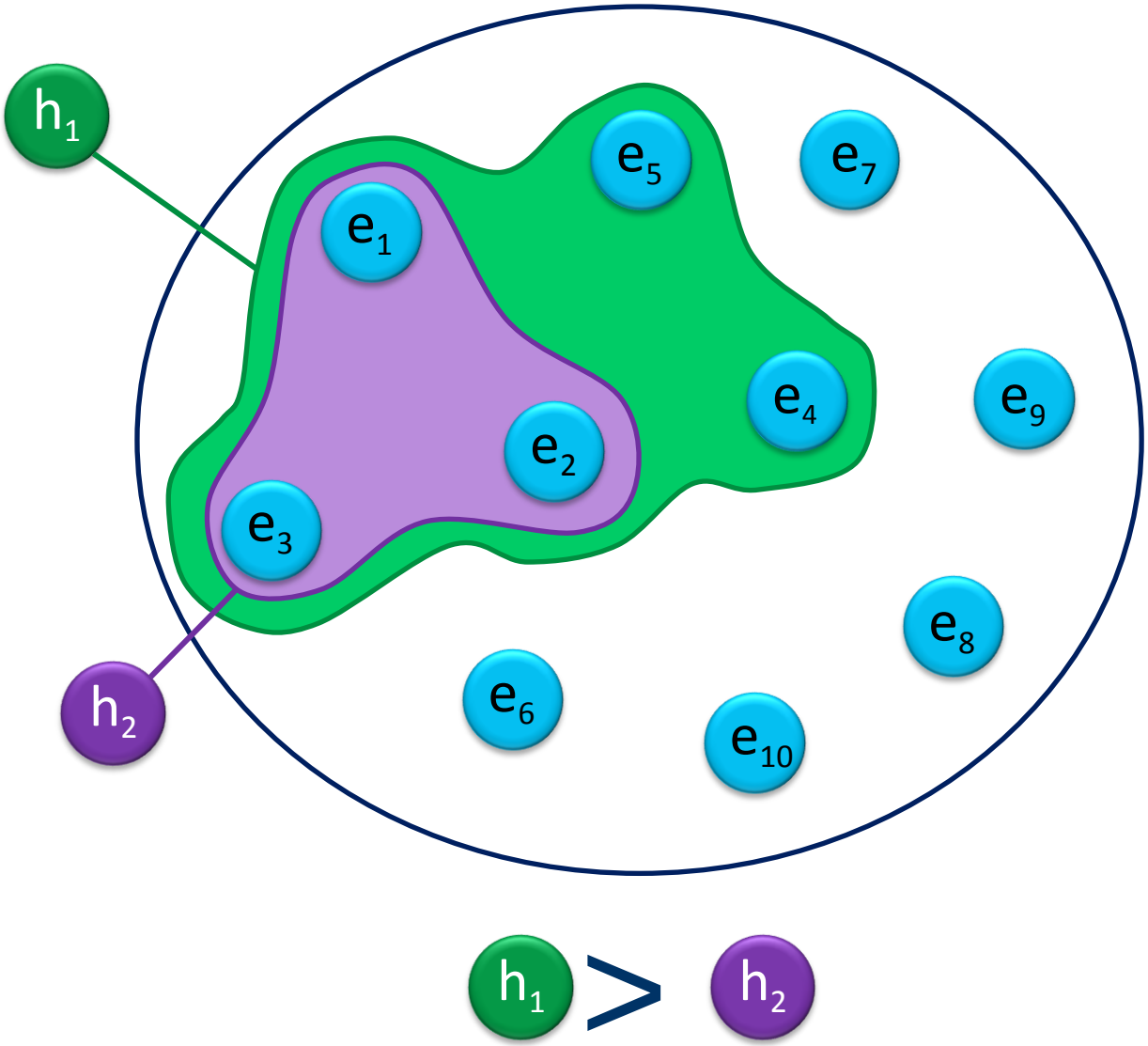
Reference dataset
(factual knowledge)

Hypotheses hierarchy



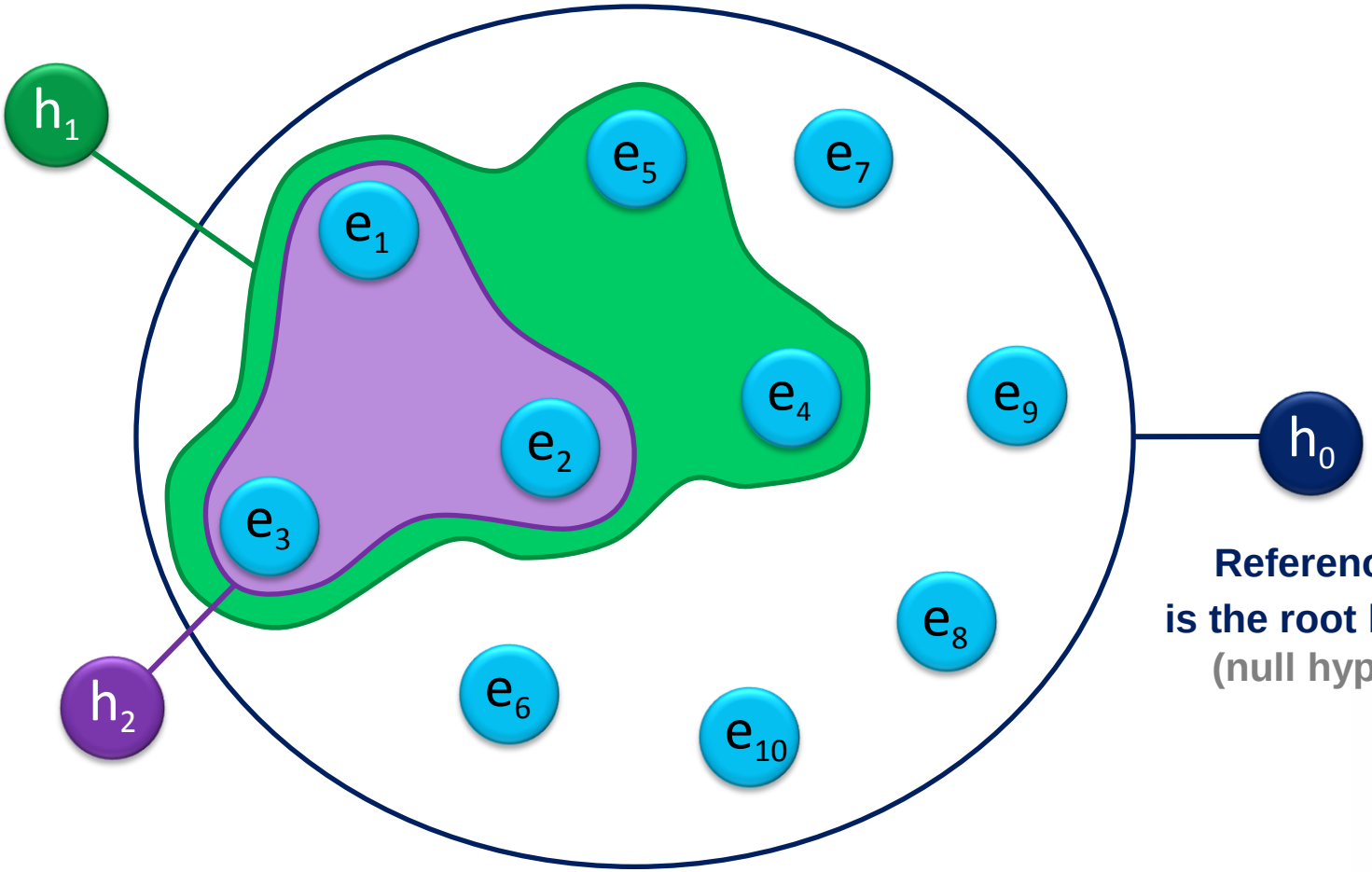
Reference dataset

Hypotheses hierarchy



Reference dataset

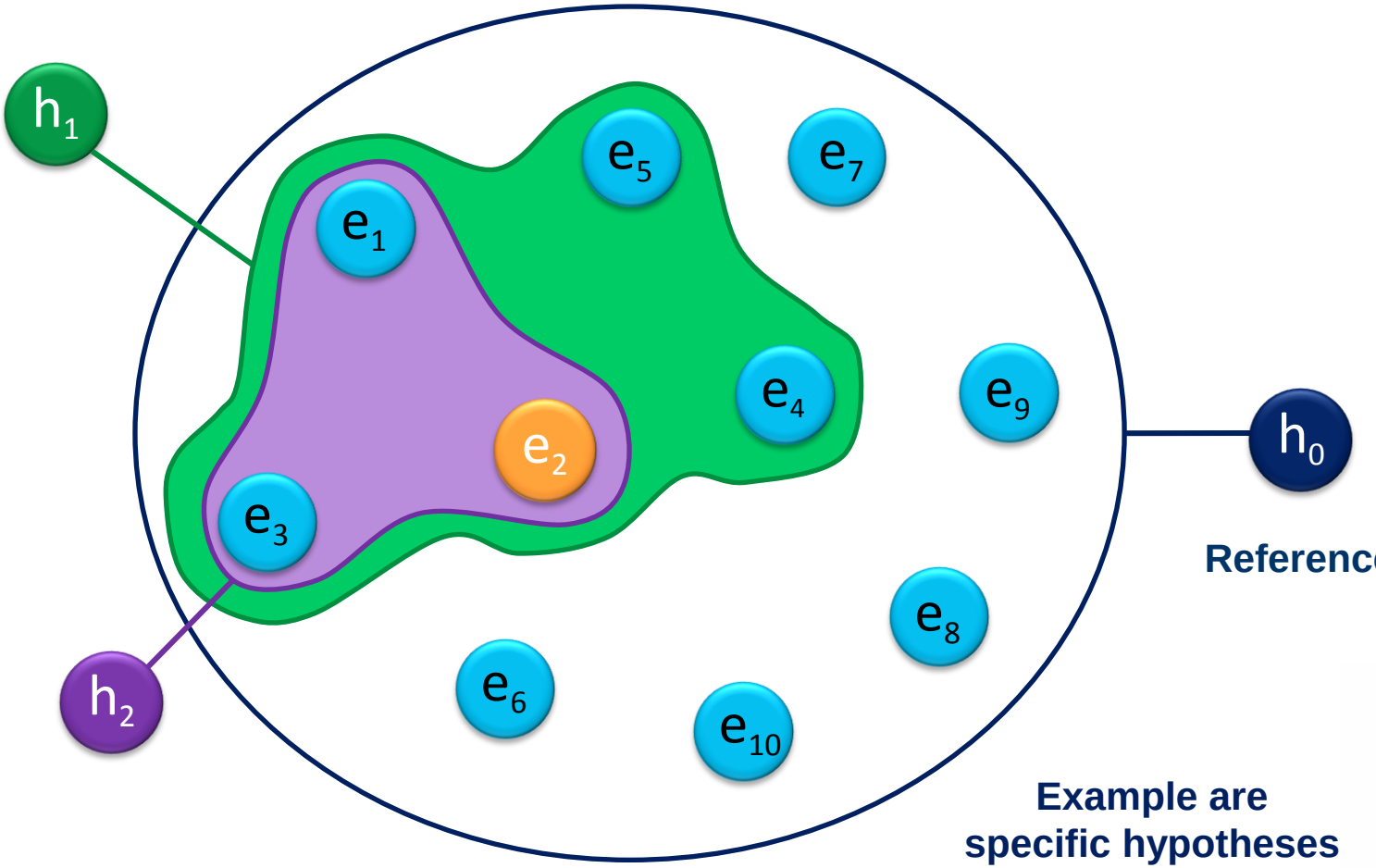
Hypotheses hierarchy



Reference dataset
is the root hypothesis
(null hypothesis)

$h_0 > h_1 > h_2 >$

Hypotheses hierarchy

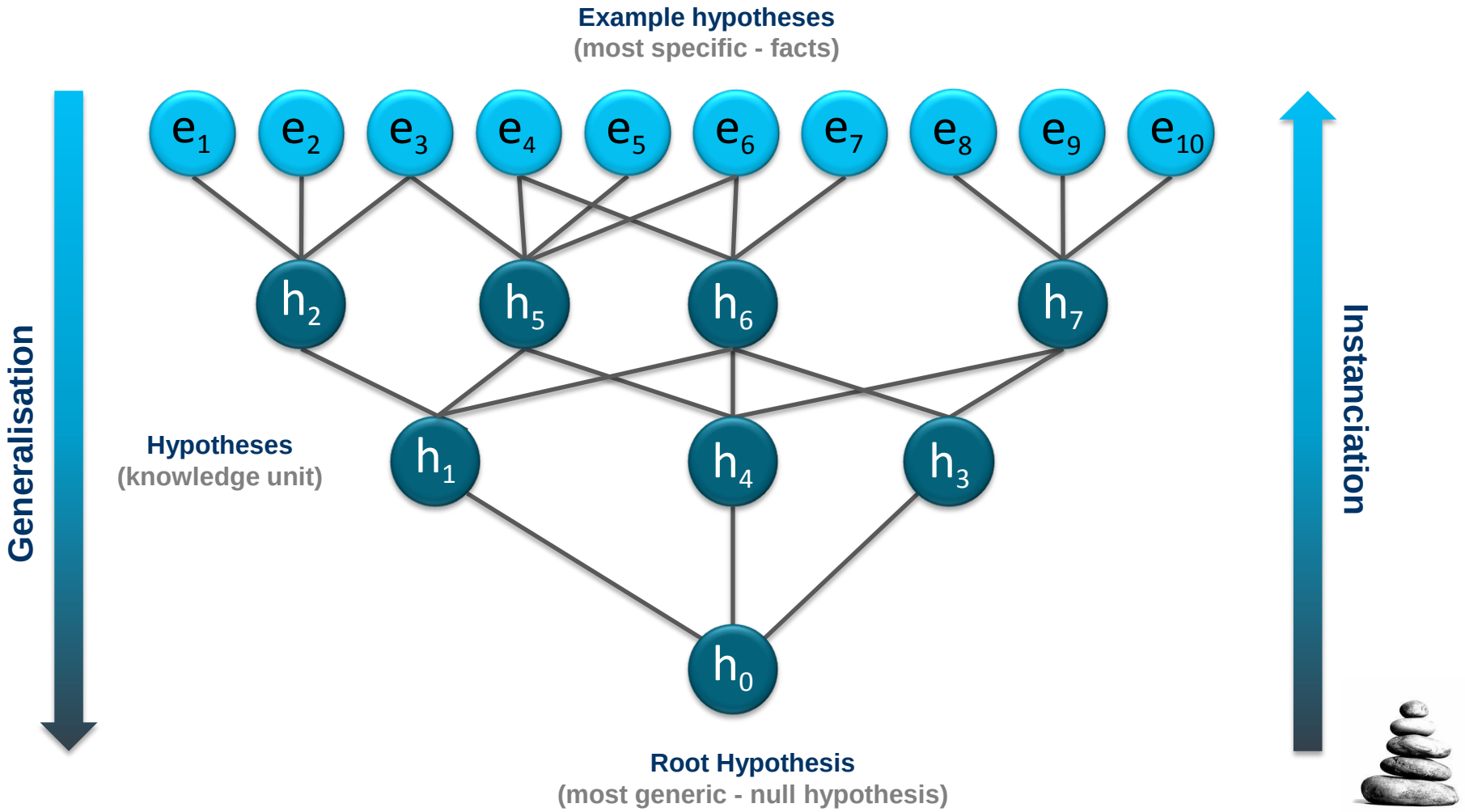


Reference dataset

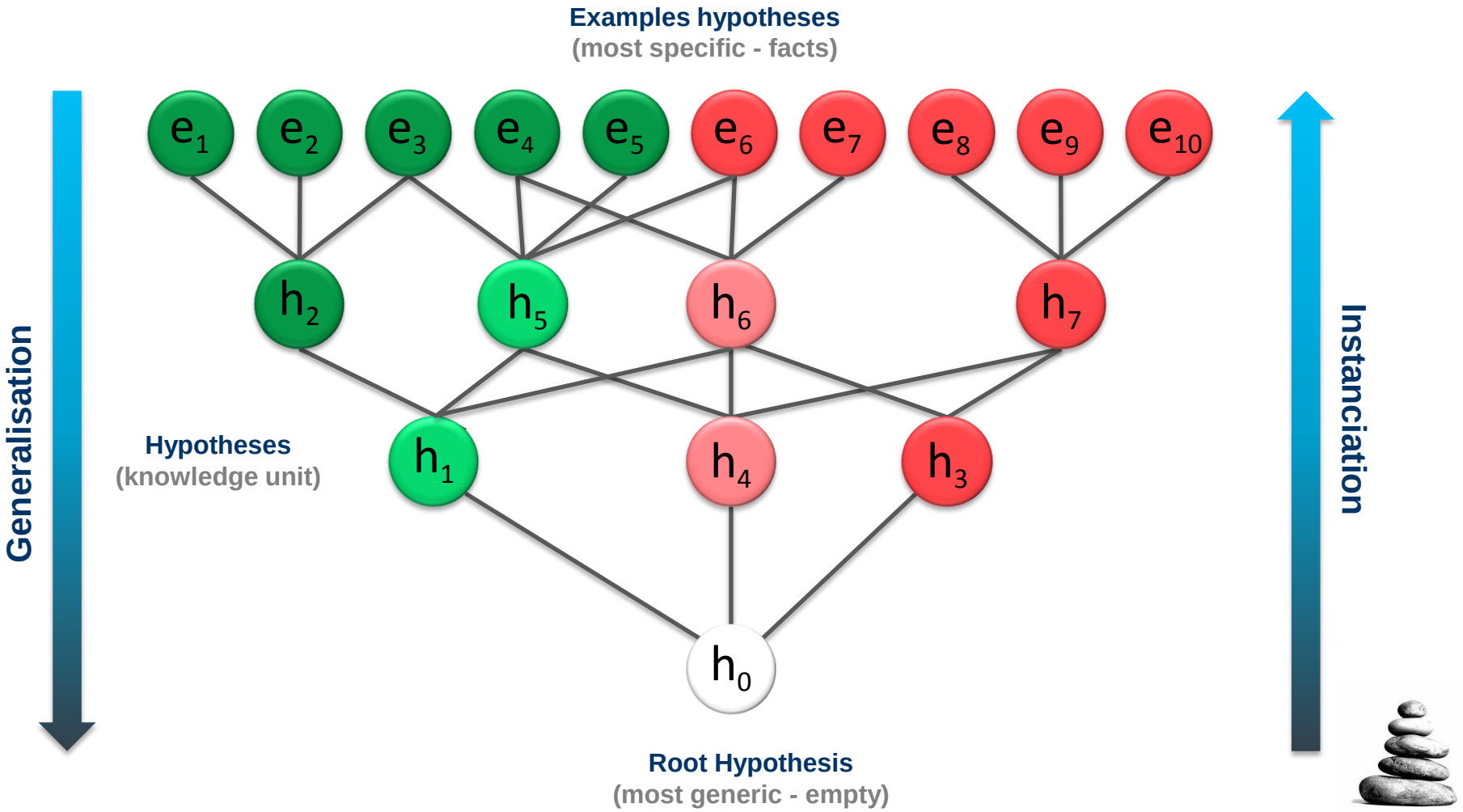
Example are specific hypotheses

$h_0 > h_1 > h_2 > e_2$

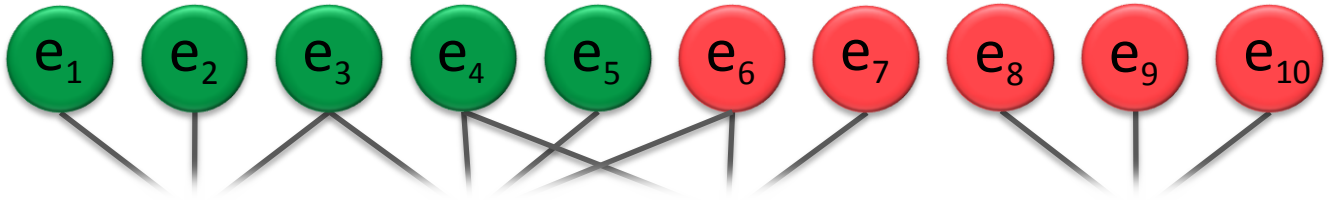
Hypotheses hierarchy



Hypotheses hierarchy



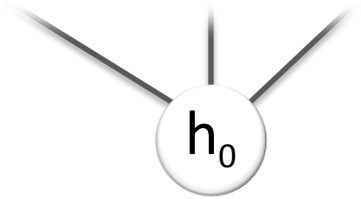
Hypotheses hierarchy



Good hypotheses (knowledge) combine strong signal and high coverage

We expect the hypothesis sources to provide good hypotheses

(can be analysed using information theory e.g. Shanon Entropy)

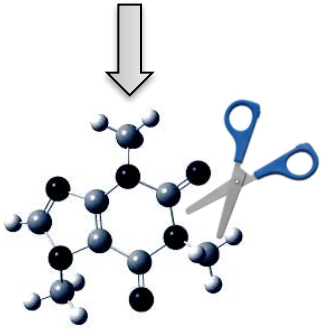


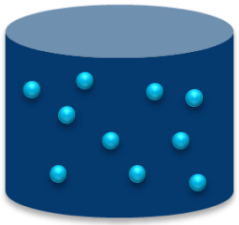
Generalisation

Instanciation

Example

Experimental data

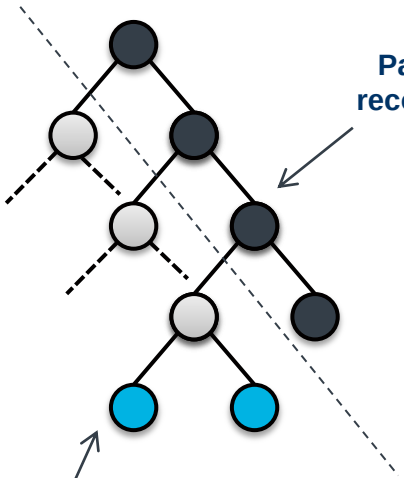




Fragment dictionary

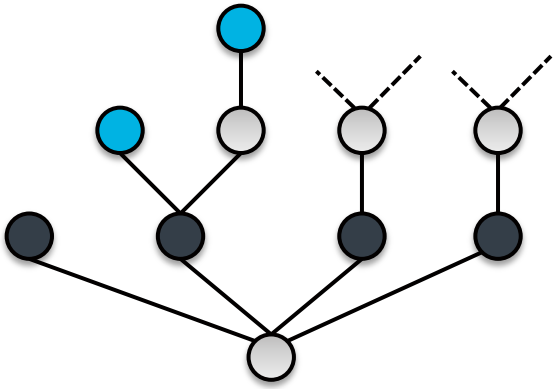
Mine hypotheses

Decision Tree (Fragments)



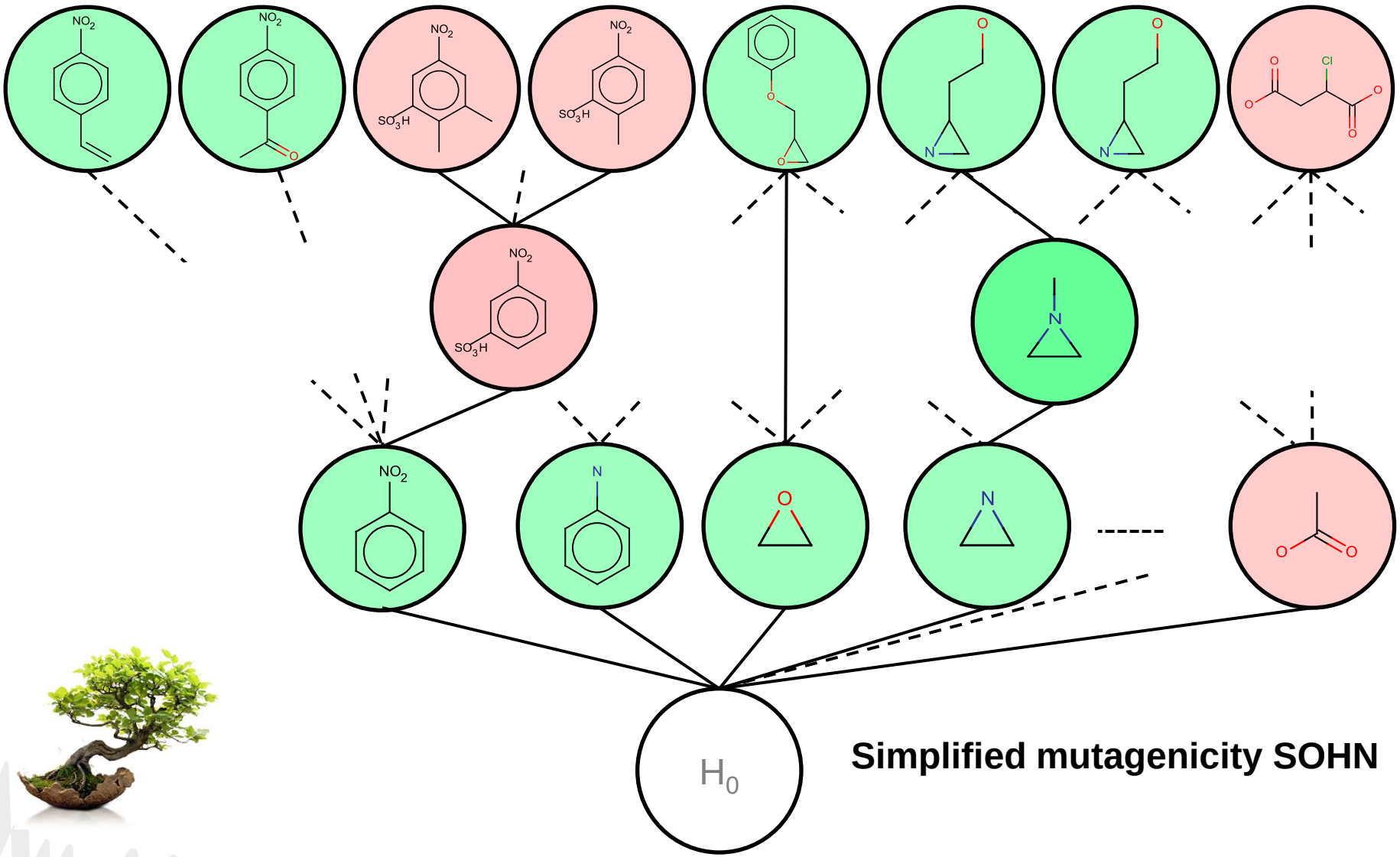
Patterns recognition

Patterns refinement



SOHN (Hypotheses)

Example



Simplified mutagenicity SOHN

Knowledge Mining

Paths contain valuable information

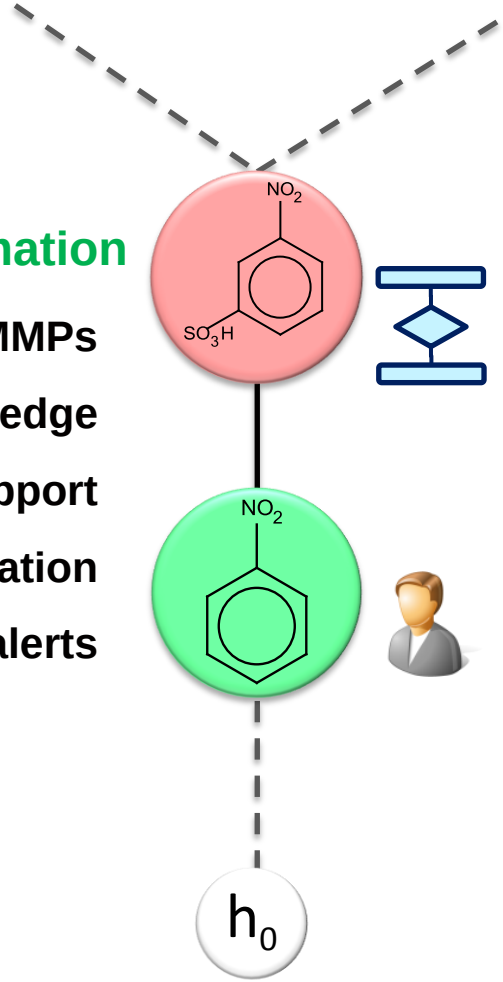
Identify Activity Cliffs / MMPs

Refine expert knowledge

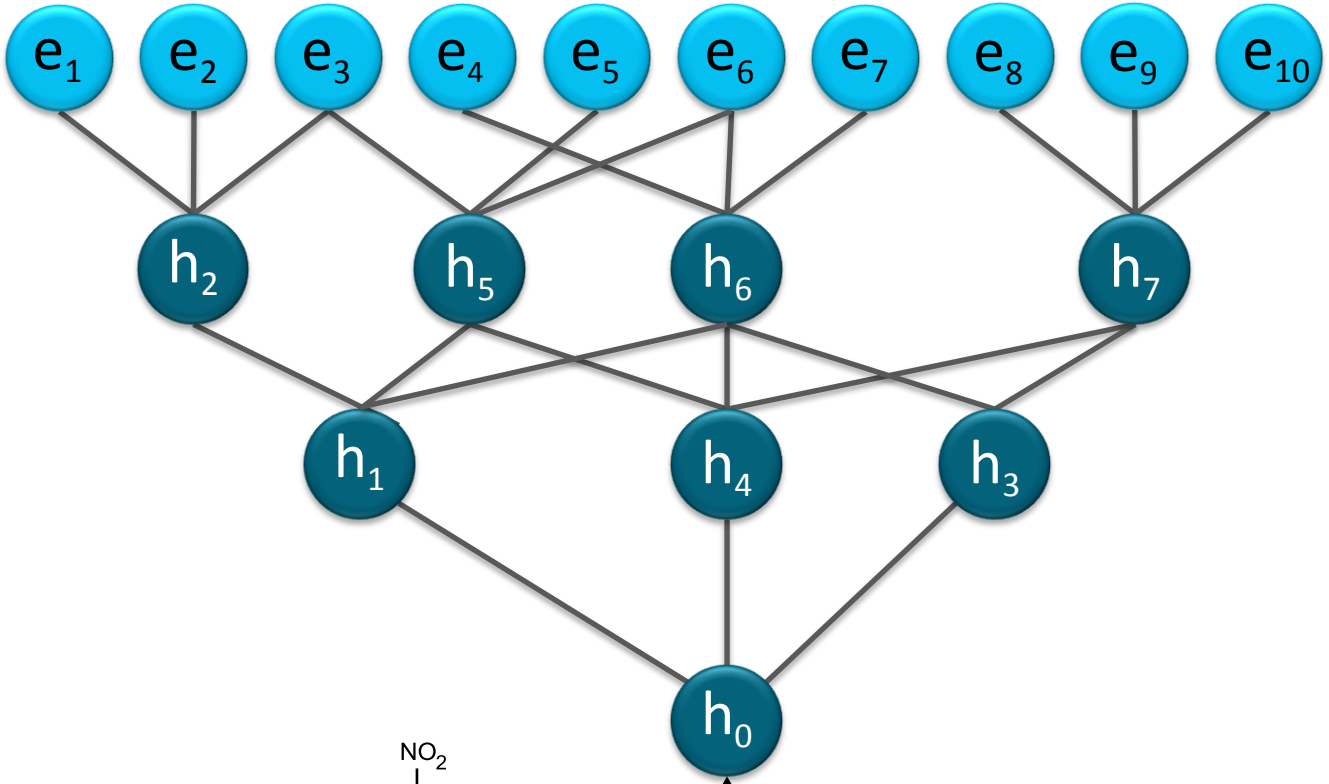
Lead optimisation support

Extended intepretation

New alerts

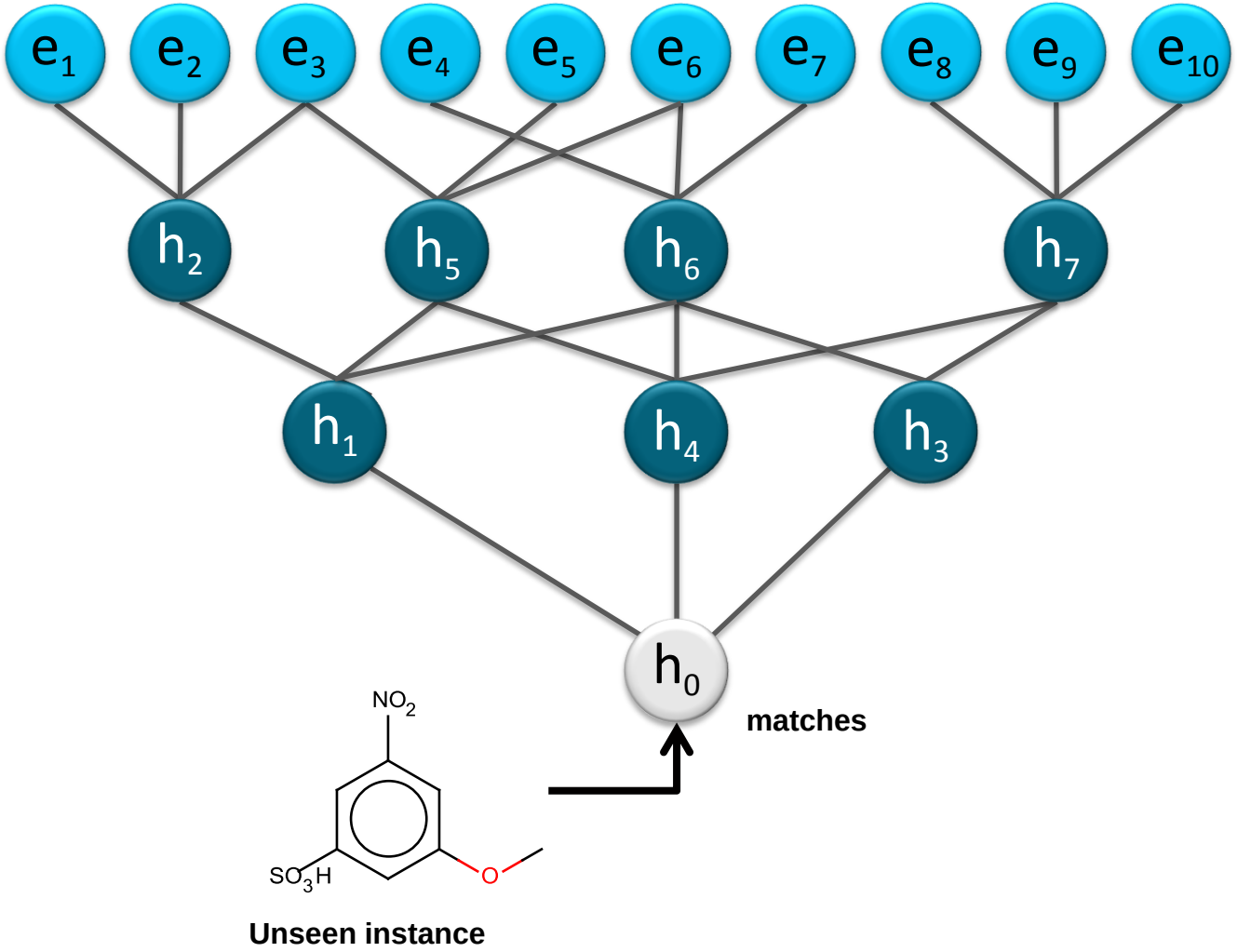


Prediction

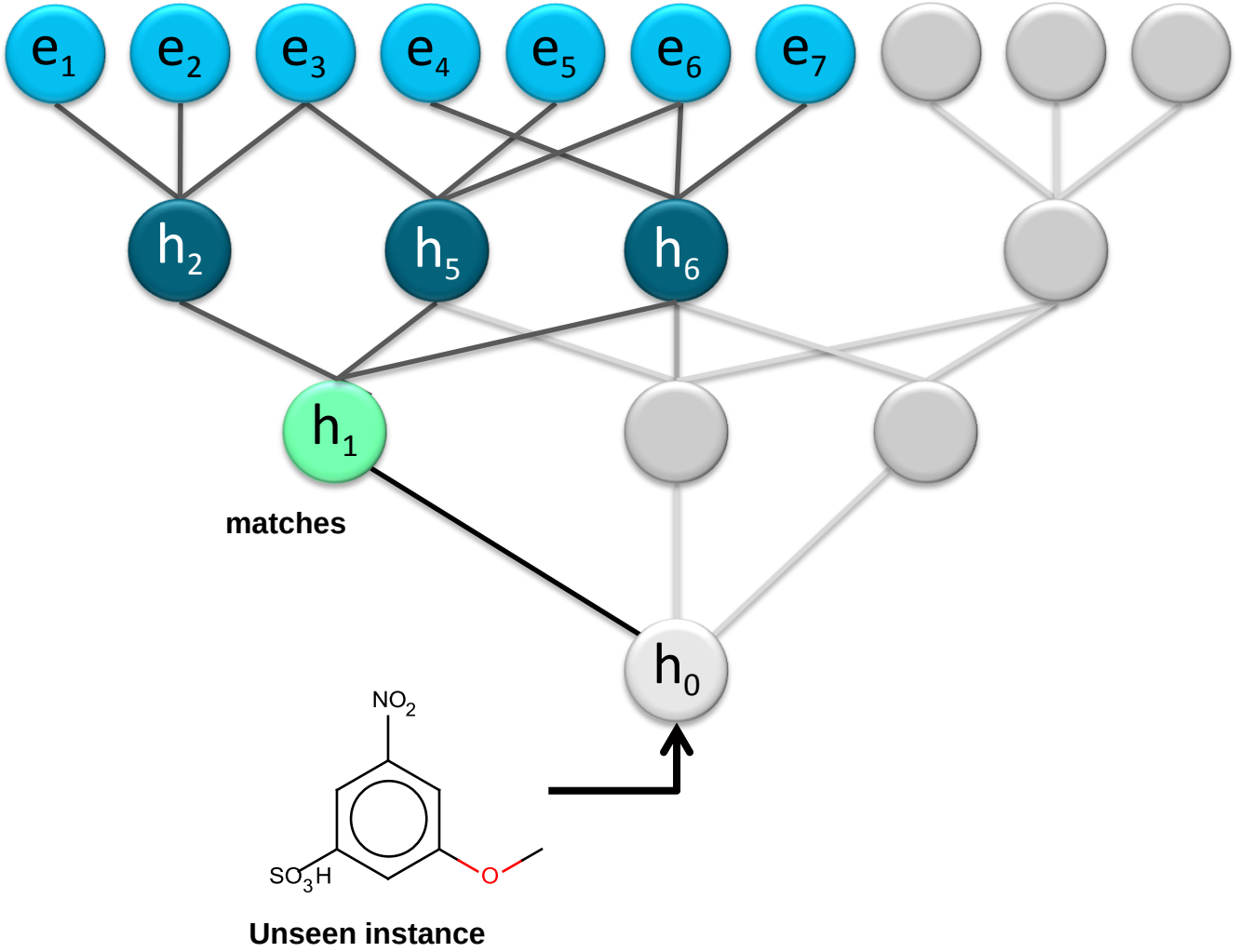


Unseen instance

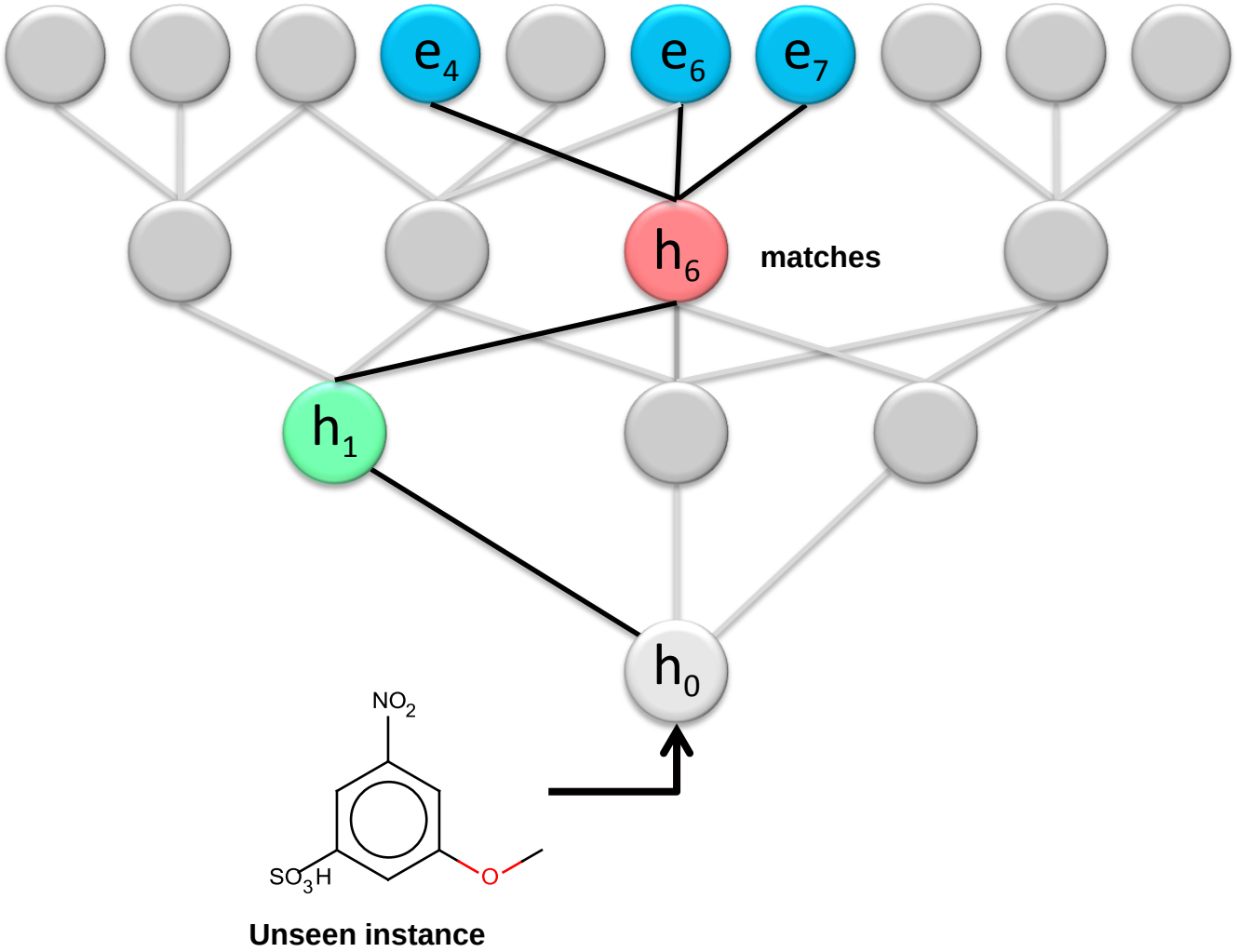
Prediction


e1
e2
e3
e4
e5
e6
e7
e8
e9
e10
h2
h5
h6
h7
h1
h4
h3
h0
matches
NO2
SO3H
O
Unseen instance

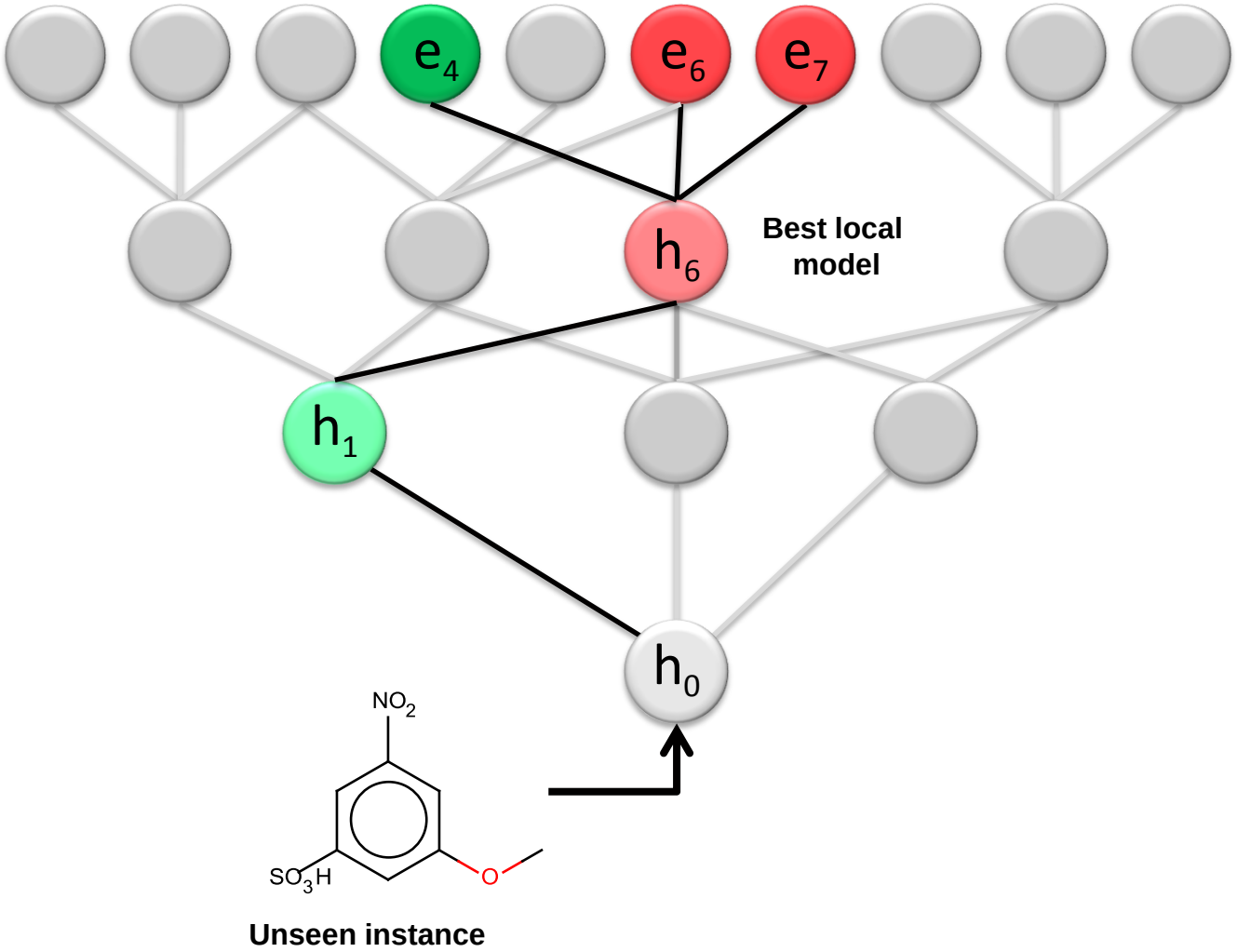
Prediction


e1
e2
e3
e4
e5
e6
e7
h2
h5
h6
h1
matches
h0
NO2
SO3H
O
Unseen instance

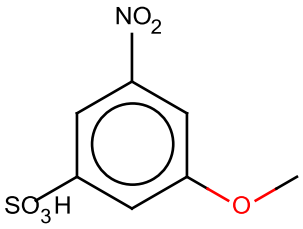
Prediction



Prediction



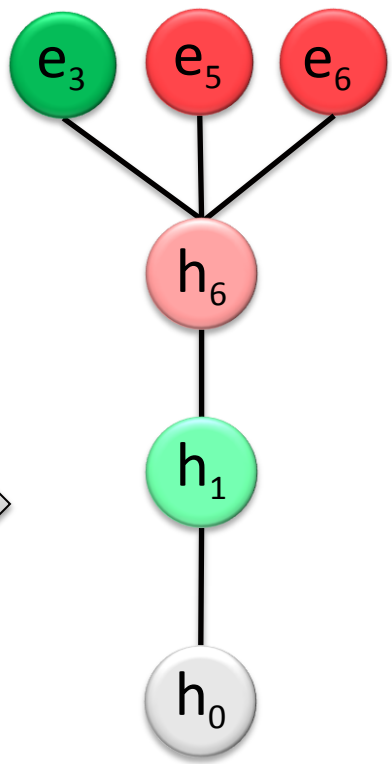
Prediction



Unseen instance

Explore SOHN

Supporting examples

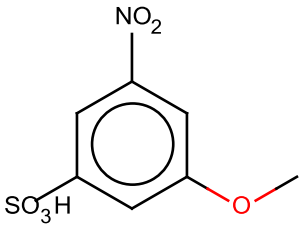


Most relevant part of the knowledge

Prediction (instance based)

Interpretation (induction based)

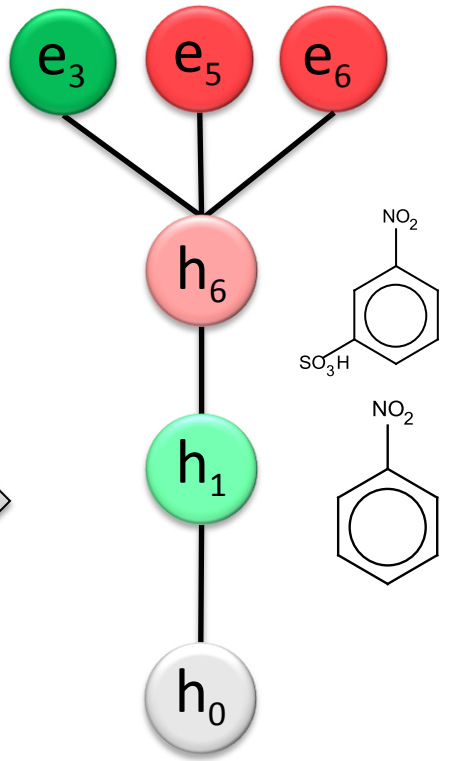
Prediction



Unseen instance

Explore SOHN

Supporting examples



Most relevant part of the knowledge

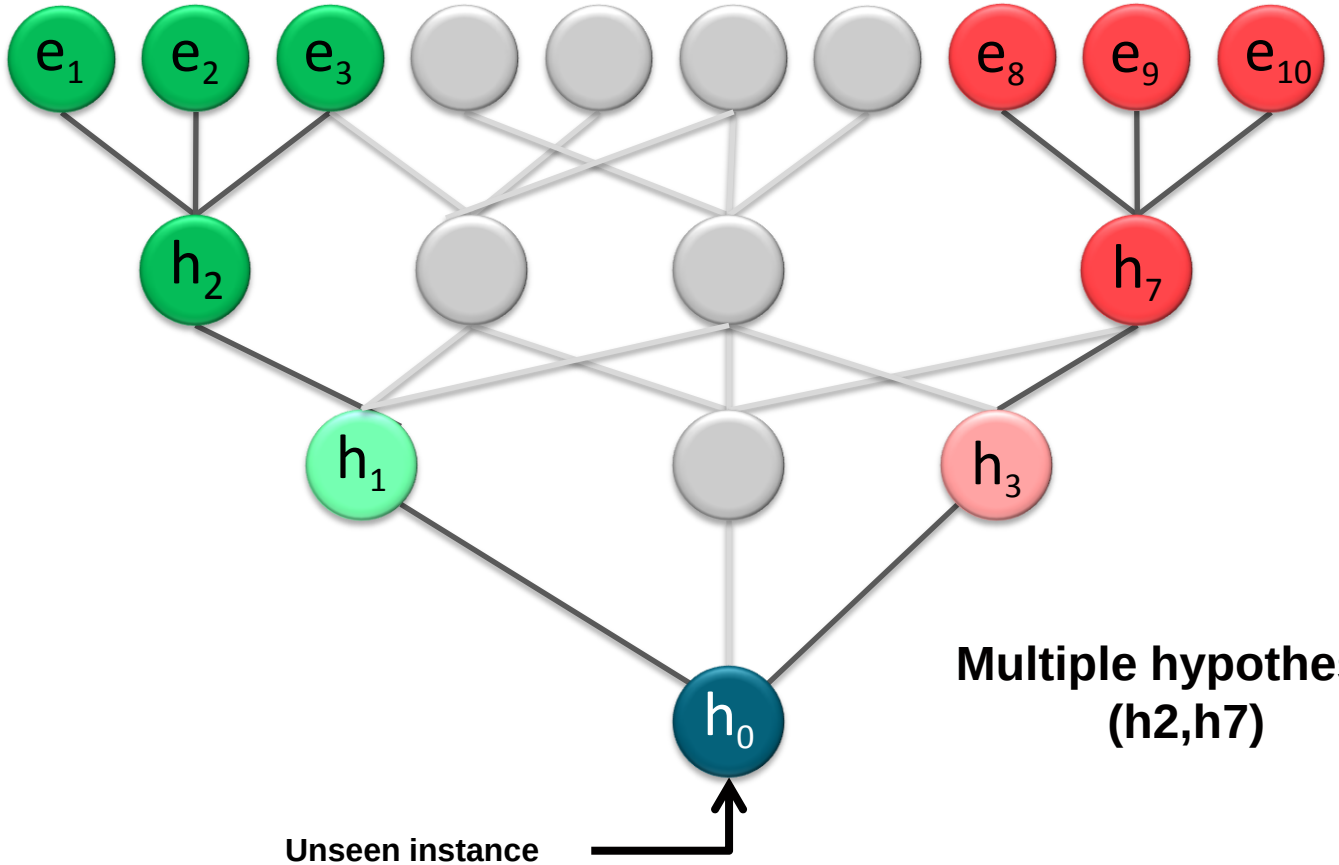
Local KNN model
CLASS

Example variance & similarity
CONFIDENCE

Path : INTERPRETATION

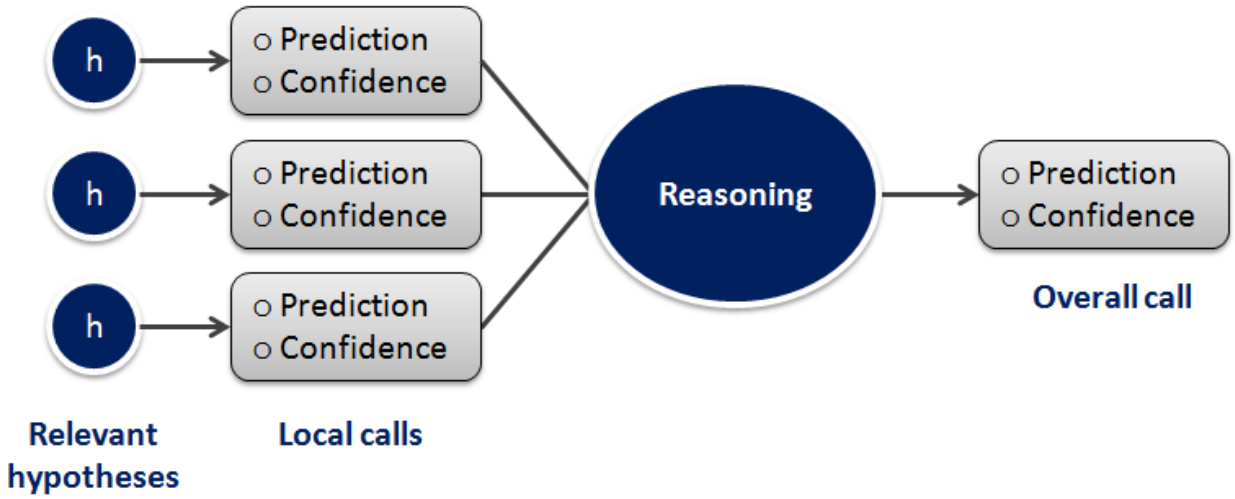
Aromatic nitro deactivated by the sulfonatic acid group in meta position

Prediction



Multiple hypotheses (h2,h7)

Prediction



Reasoning

- **Weighted / Confidence**
- **Most confident**
- **Conservative** (1+ve enough)
- **Average**

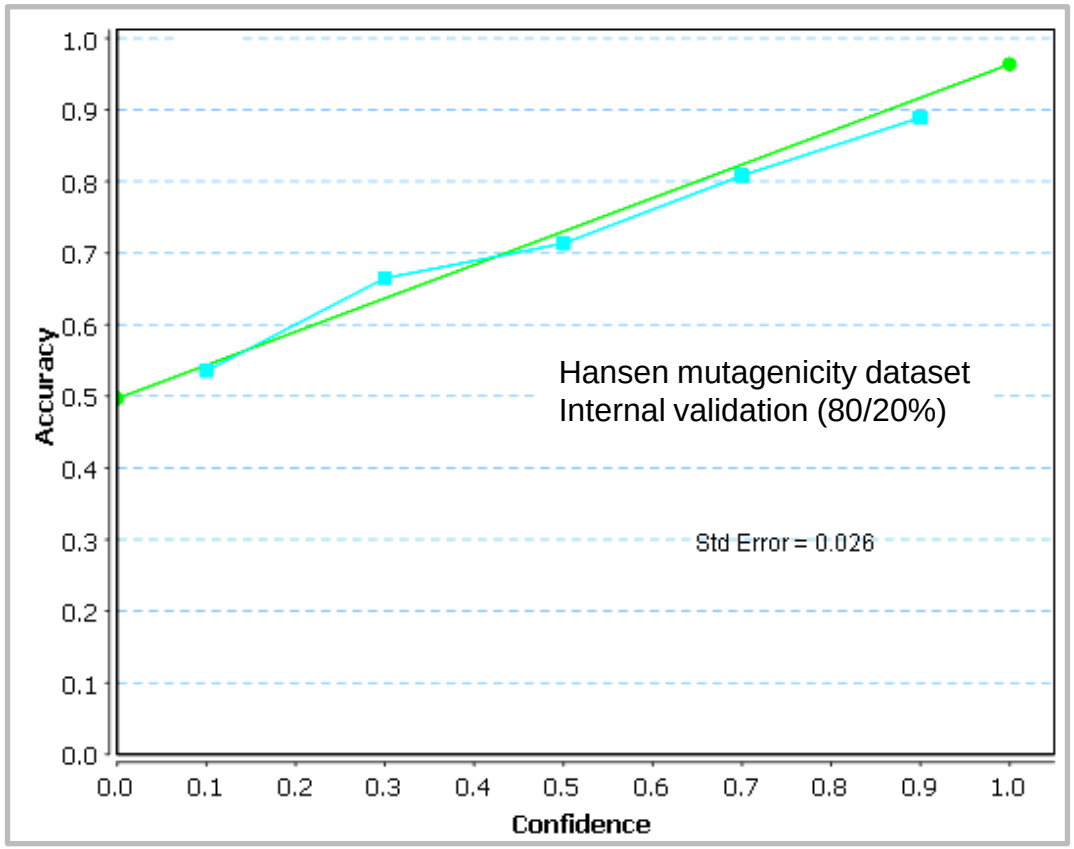
Flexibility / Use case

Weighted / Confidence

$$s_x = \frac{\sum_{h=1}^{m} s_{h,x} \times confidence_{h,x}}{\sum_{h=1}^{m} confidence_{h,x}}$$

$$confidence_x = |s_x|$$

Individual prediction confidence



Confidence / Accuracy correlation

Example
Mutagenicity prediction

Results: Mutagenicity

SOHN details

Dataset	Training	Test	Sensitivity	Specificity	Balanced Accuracy
A (internal 20%)	6560 (80%)	1640 (20%)	78.5	81.4	80
B (external)	8200 (100%)	800 (100%)	53	82	67

Comparison with other ML methods

Balanced Accuracy	SVM	RF	KNN	DT	SOHN
A (internal 20%)	81	80	76	77	80
B (external)	64	63	61	60	67

Public dataset A : 8200 public structures (balanced : 50% +ve)
Proprietary dataset B : 800 proprietary structures (biased : 29% +ve)

SVM : best results using PubChem fingerprints (optimised parameters)
RF : best results using MACCS keys / 100 trees

Application

sarah nexus

Integration into Lhasa Limited Nexus Suite

Nexus

File Window Prediction Reports Tools Help

Hide

Predictions

- Ames study
  - Query compound
    - Sarah prediction

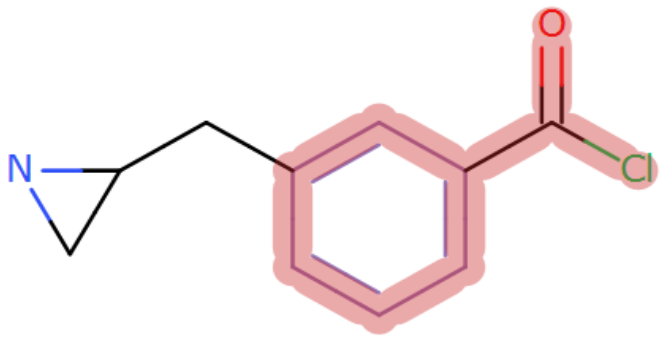
Jobs

Sarah prediction

For the '**Mutagenicity**' endpoint the prediction is:

POSITIVE

with **71%** confidence



Displaying 'H-739 matches', click above to view the original structure

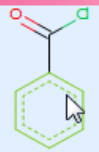
Prediction Constraints

Model:	Lhasa
Endpoint:	Mutagenicity
Reasoning type:	Weighted
Equivocal:	0%
Sensitivity:	0%
Certified model:	Yes
Prediction date:	12 March 2014 15:22

Prediction options

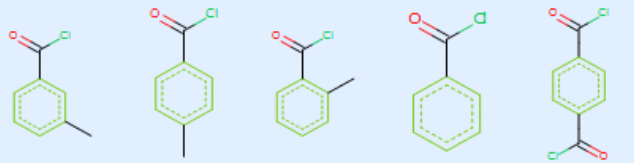
Results

The compound is predicted to be positive with 71% confidence for the 'Mutagenicity' endpoint in the 'Lhasa' model. Supporting hypotheses containing similar examples from the training set have been found.

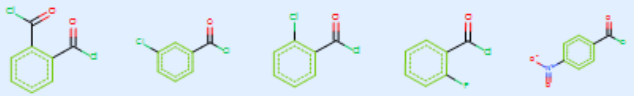
Structure	ID	Hypothesis Result	Confidence
Hypothesis			
	H-739	Positive	77%

Training Set Examples

1 of 16 (+Ve) 2 of 16 (+Ve) 3 of 16 (+Ve) 4 of 16 (+Ve) 5 of 16 (+Ve)



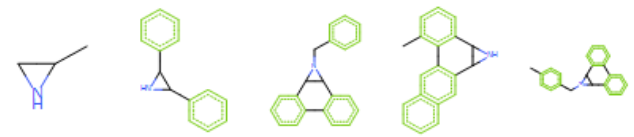
6 of 16 (+Ve) 7 of 16 (+Ve) 8 of 16 (+Ve) 9 of 16 (+Ve) 10 of 16



Structure	ID	Hypothesis Result	Confidence
Hypothesis			
	H-689	Positive	64%

Training Set Examples

1 of 56 (+Ve) 2 of 56 (+Ve) 3 of 56 (+Ve) 4 of 56 (+Ve) 5 of 56 (+Ve)



6 of 56 (+Ve) 7 of 56 (+Ve) 8 of 56 (+Ve) 9 of 56 (+Ve) 10 of 56

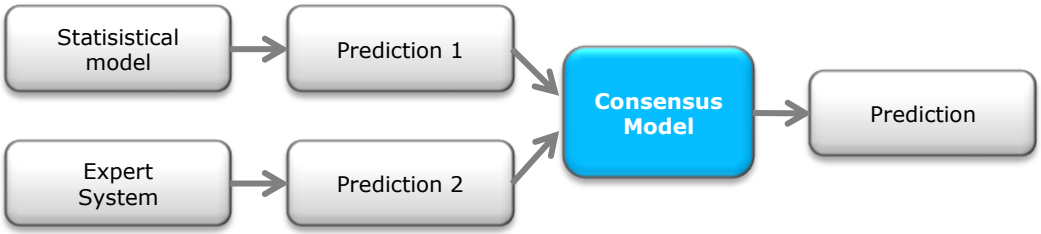
Hypotheses: 2

Combining Statistical Models with Expert Systems

(ICHM7)

Combining Statistical and Expert system

Consensus model approach

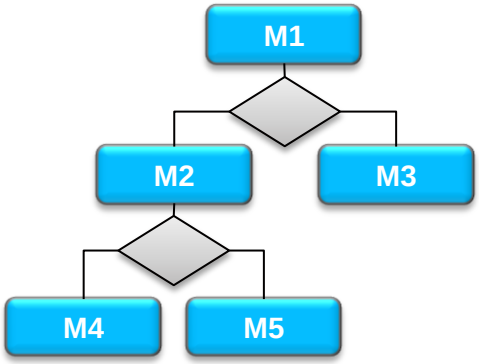


Importance of individual prediction accuracy estimate (confidence)

Model selection

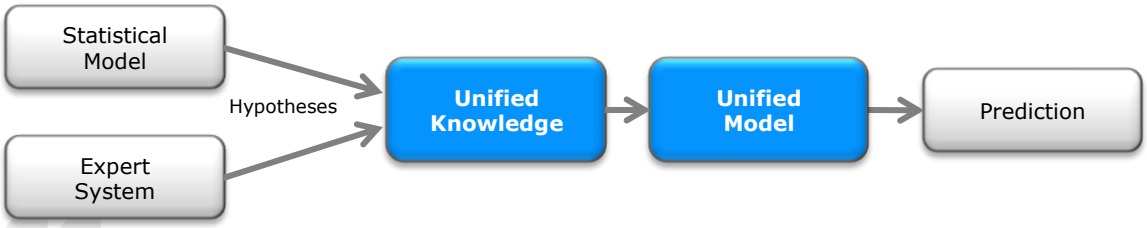
M1	M2	M3	M1	M4
M1	M1	M3	M1	M2
M3	M1	M1	M4	M1
M1	M2	M5	M1	M5
M4	M1	M1	M3	M1

Chemical space performance partition

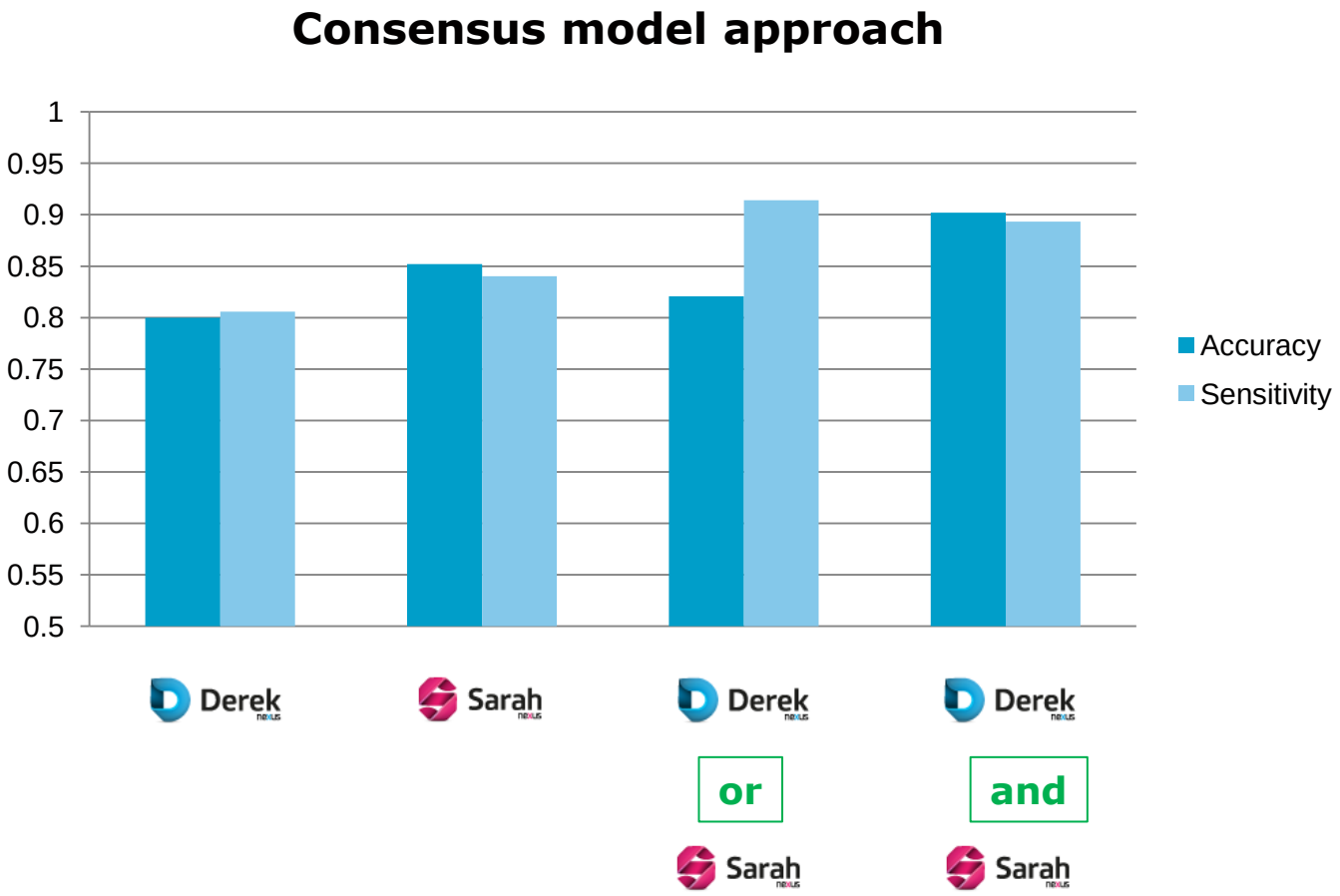


Decision trees (Meta Model)

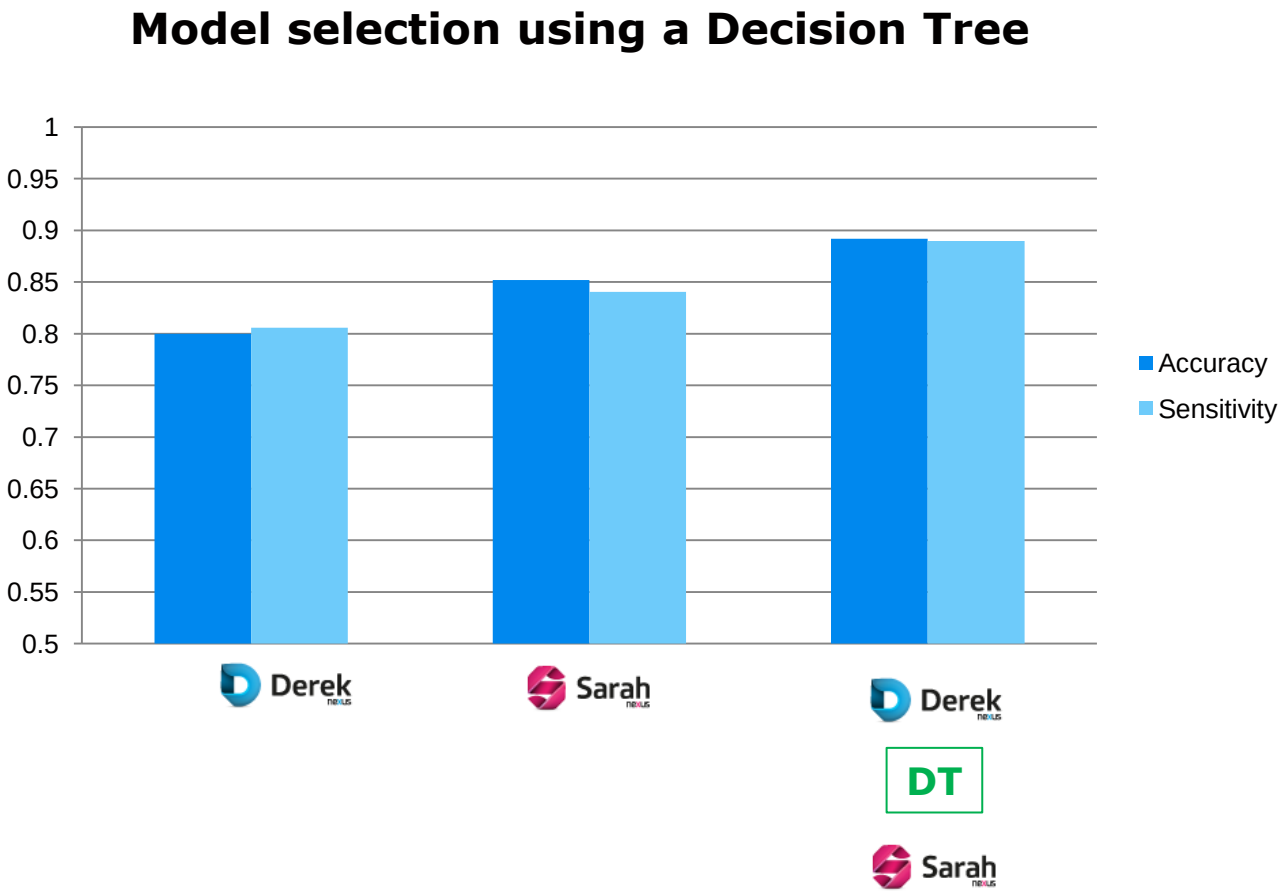
Unified Knowledge approach



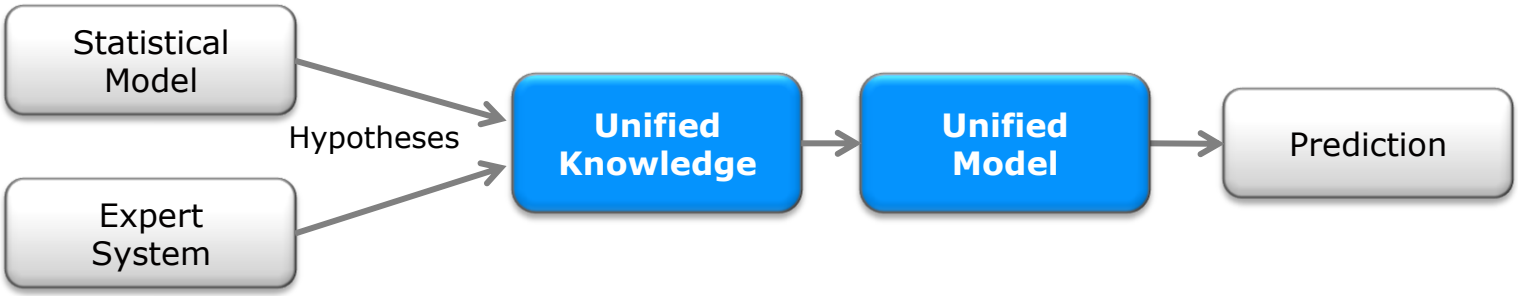
Combining Statistical and Expert system



Combining Statistical and Expert system



Unified Knowledge approach



Advantages

- Combining different source of knowledge (Machine learning, Expert Knowledge)
- Automatic knowledge organisation within a local model hiearachy
- Optimised knowledge selection
- Single prediction algorithm
- Transparent predictions with indication of the origin of the knowledge used
- Harmonised confidence level for individual prediction

Conclusion

• The expert plays the key role in accessing the toxicity of a compound and needs transparent and accurate tools to help him in this task (OECD guidelines)

• Finding the right trade-off between transparency and accuracy is challenging

• One approach is to combine the knowledge from different sources including expert systems and statistical models (ICHM7)

• These different sources can be integrated into a single framework to provide transparent and accurate predictions (SOHN approach)

Arigatou gozaimasu
有難う 御座います

Thank you

Lhasa Limited

shared **knowledge** • shared **progress**

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