Modelling Local and Systemic Toxicity: Incorporation of *In Silico* Predictions in the Development of Adverse Outcome Pathways

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In Silico Prediction

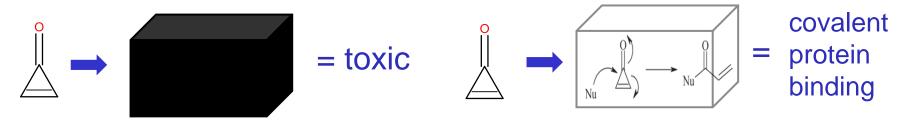
Activity (e.g. toxicity)of a chemical

 ∞

Molecular (structural) properties

Certain endpoints easier to predict than others

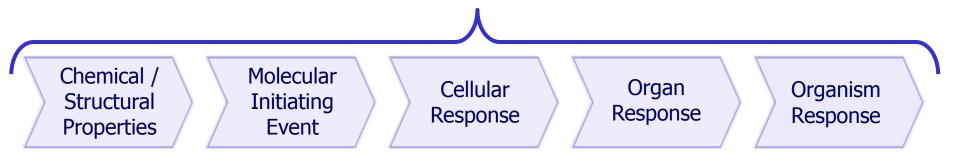
Methods need to be transparent



 Newer methods such as category formation (grouping) & readacross can be applied to complex endpoints & are transparent



Adverse Outcome Pathway for Skin Sensitisation



- The AOP provides a method to represent the Key Events involved
- Key Events are measurable and toxicologically relevant

- NB pathway may be non-linear; adaptive & regulatory responses may be involved
- Realistic exposure scenario need to be considered

Key Events in Skin Sensitisation AOP

Key Event 1 Molecular Initiating Event



Covalent binding interaction

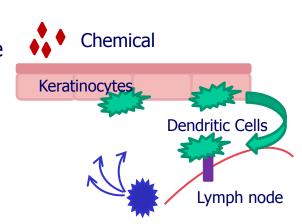
Electrophile

Nucleophile on skin protein (cysteine/lysine)

Key Event 2

Cellular response

Keratinocyte inflammatory response



Key Event 3

Cellular response

Dendritic cell activation

Key Event 4

Organ Response

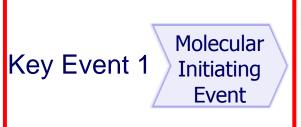
Lymph node - activation of T-cells proliferation of activated T-cells



Skin: 🖒

Inflammation on challenge with antigen

Key Events in Skin Sensitisation

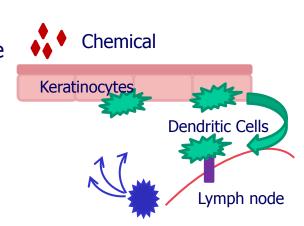




Covalent binding interaction

Nucleophile on skin protein (cysteine/lysine)

Keratinocyte inflammatory response



Key Event 3

Cellular response

Dendritic cell activation



Lymph node - activation of T-cells proliferation of activated T-cells



Skin: **C**Inflammation on challenge with antigen

Accumulating Information to Develop AOPs

- The Molecular Initiating Event (MIE) (initial interaction between chemical and biological system) is a Key Event
 - Hypothesis or evidence required for Key Events
 - Evidence can be accumulated from a range of sources

In Silico

Creating
Structural Alerts
/ Profilers
Identifying MIEs

In Chemico

Direct Peptide
Reactivity
Assay
Glutathione
depletion

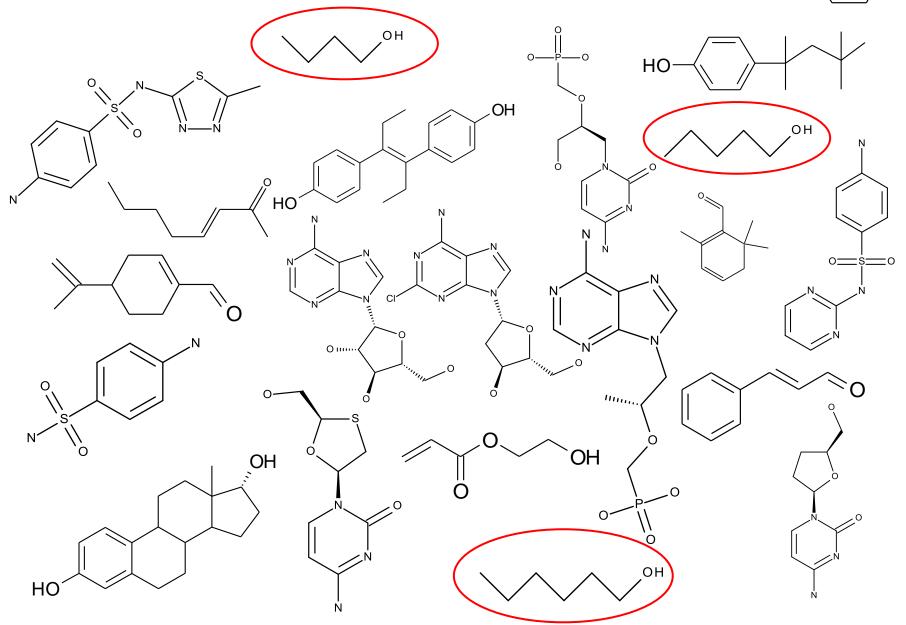
In Vitro

MUSST/h-CLAT
Ames Test
Mitochondrial
Damage
Oxidative Stress

In Vivo

GPMT LLNA Human Patch Test Toxicity Assays

 Capturing the chemistry underlying an MIE can be used to develop Structural Alerts / Profilers for grouping chemicals into categories

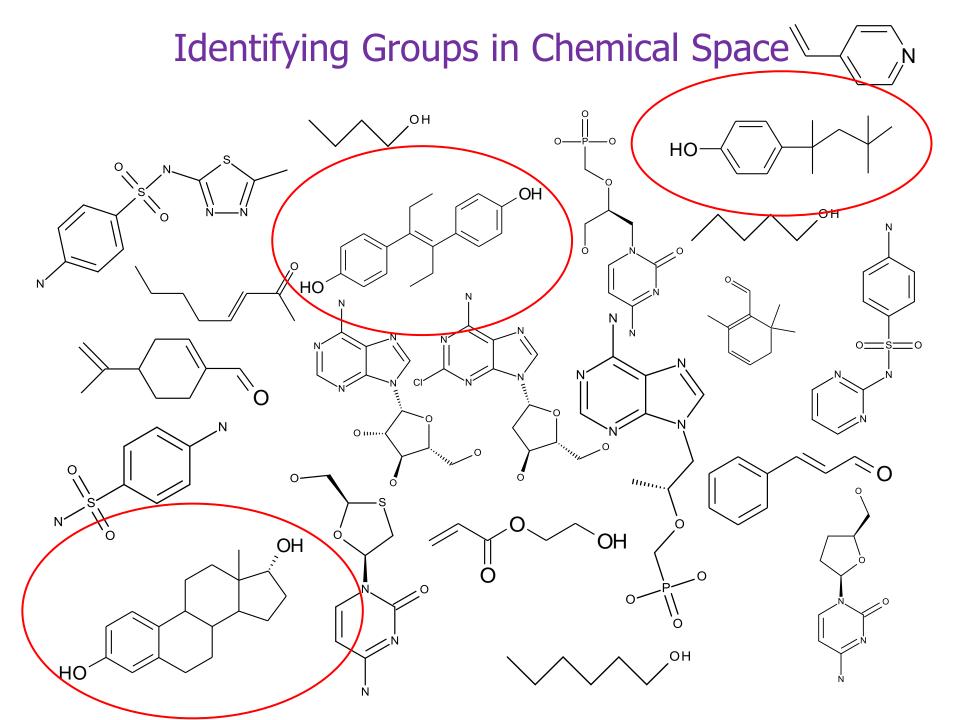


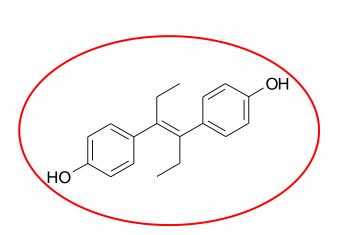


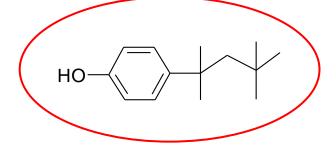


Structural Analogues

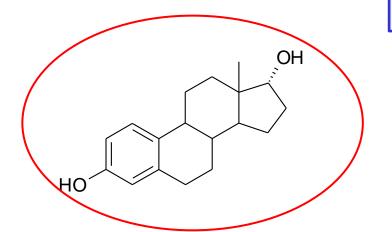


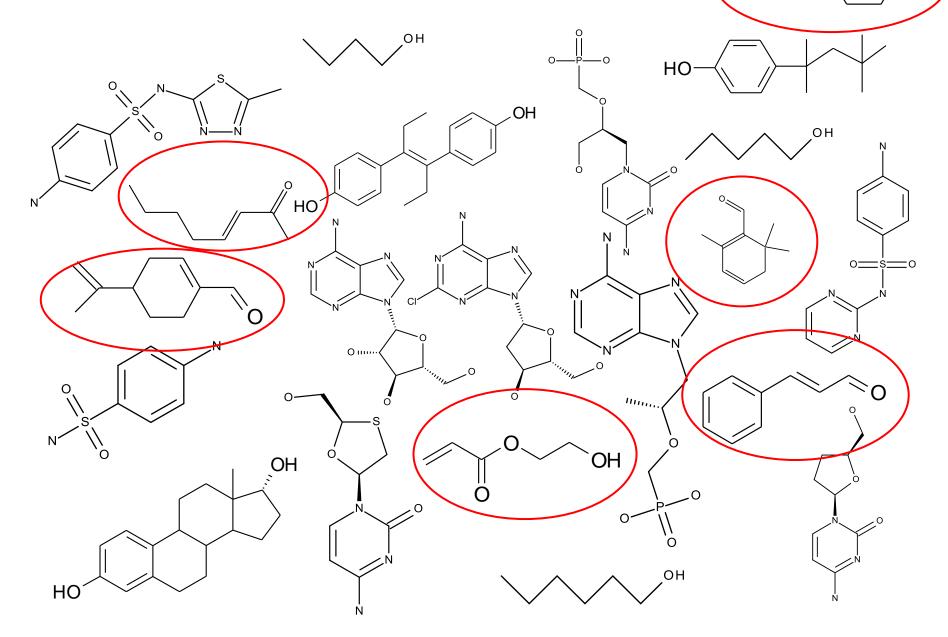


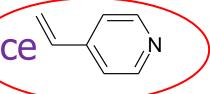


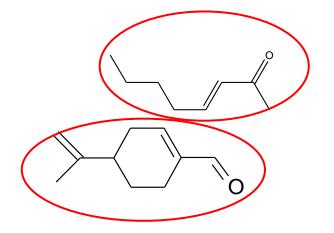


Mode of Action Analogues (Binding to oestrogen receptor)

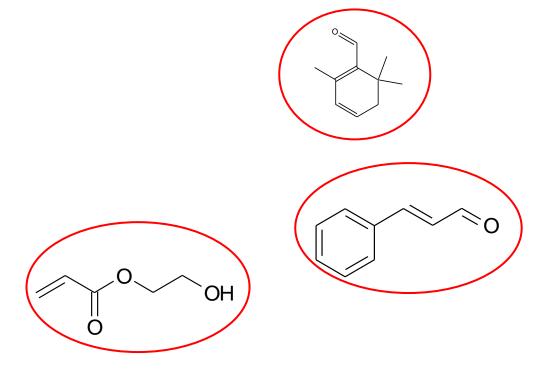








Mechanistic analogues (Michael addition reaction)



Mechanistic Structural Alerts and Profilers Developed:

Protein binding relating to skin & respiratory sensitisation

Use knowledge of mechanistic chemistry domain e.g.

Acylation

Schiff Base Formation

104 structural alerts developed for protein binding (46 associated with skin sensitisation)²; 52 structural alerts for respiratory

52 structural alerts for respiratory sensitisation

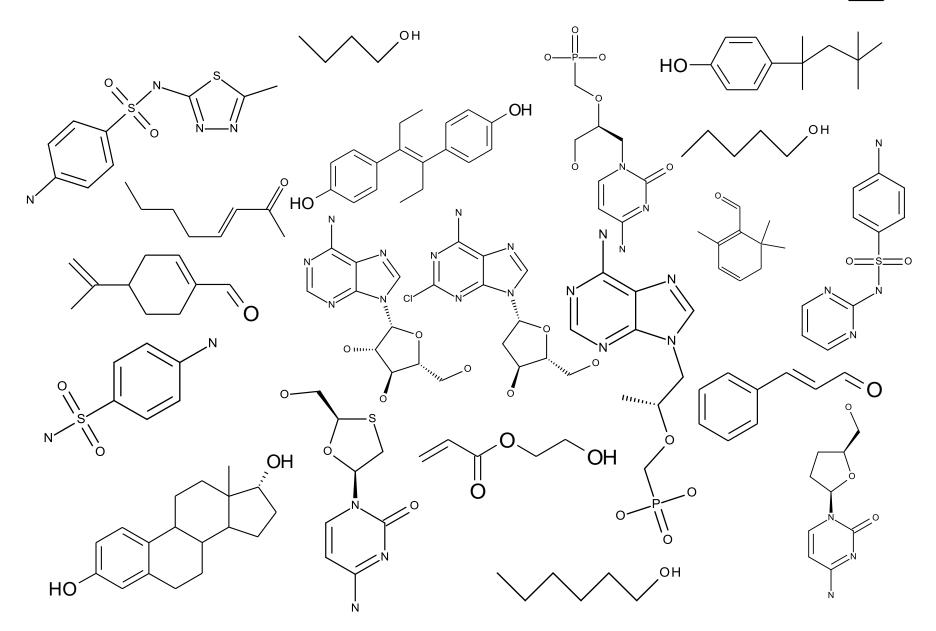
Also 57 structural alerts developed for DNA binding²

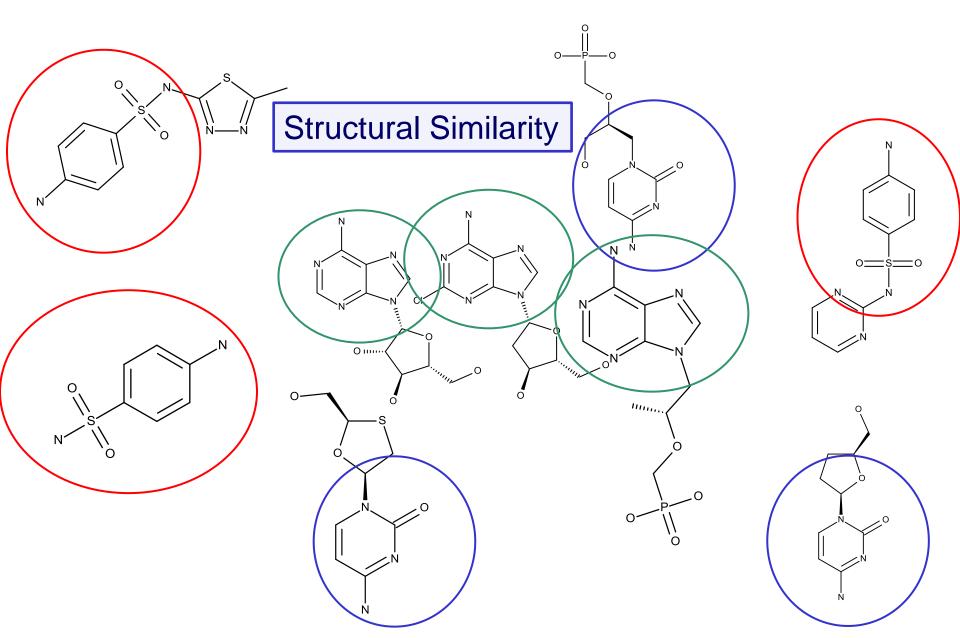


Derive associated SMARTS¹ patterns (e.g.CC=OC;

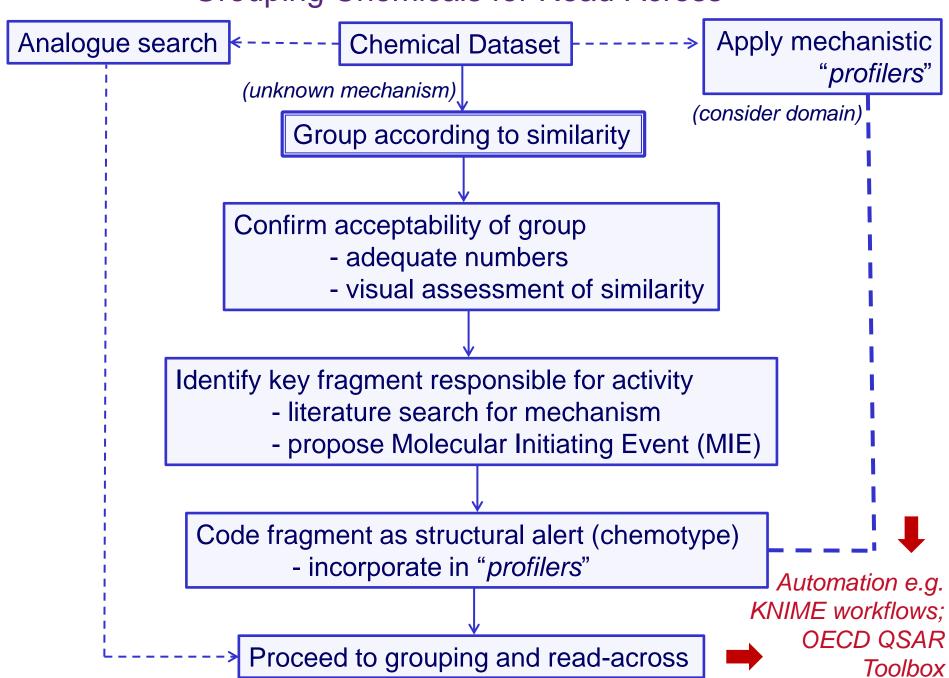


Encode into profilers





Grouping Chemicals for Read-Across



Structural Alerts for Hepatotoxicity

951 compound dataset* (650 +ve)

Grouped by structural similarity
- using Toxmatch software

Example SMARTS Patterns

- 1. C=CC=CC=CC=O
 - 2. CICCNCCCI
 - 3. O=C1CCC2C3CCC4CCCC4C3CCC2=C1

Group (category) acceptable if:

contained ≥ 6 compounds

and similarity index ≥ 0.6

and

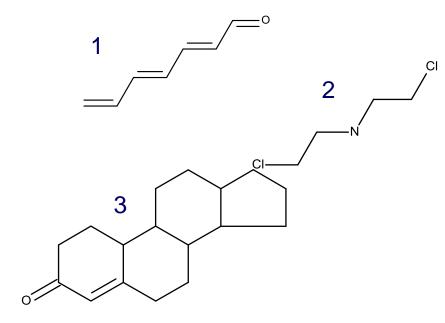
visually appeared similar

16 structural alerts*
(key fragments) identified

Alerts used to re-screen dataset no. category members

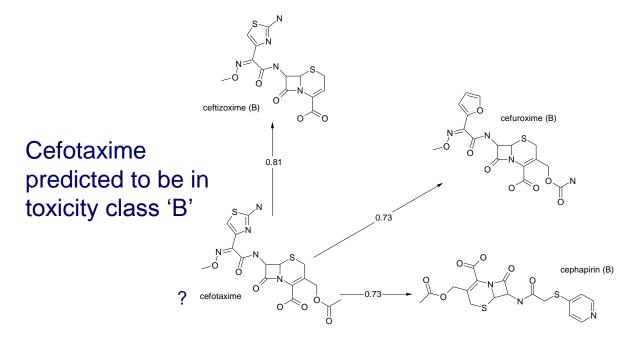
Literature search for putative mechanisms of toxicity

Mechanisms proposed for ~25% of hepatotoxicants in dataset



Using the Information to Predict Toxicity

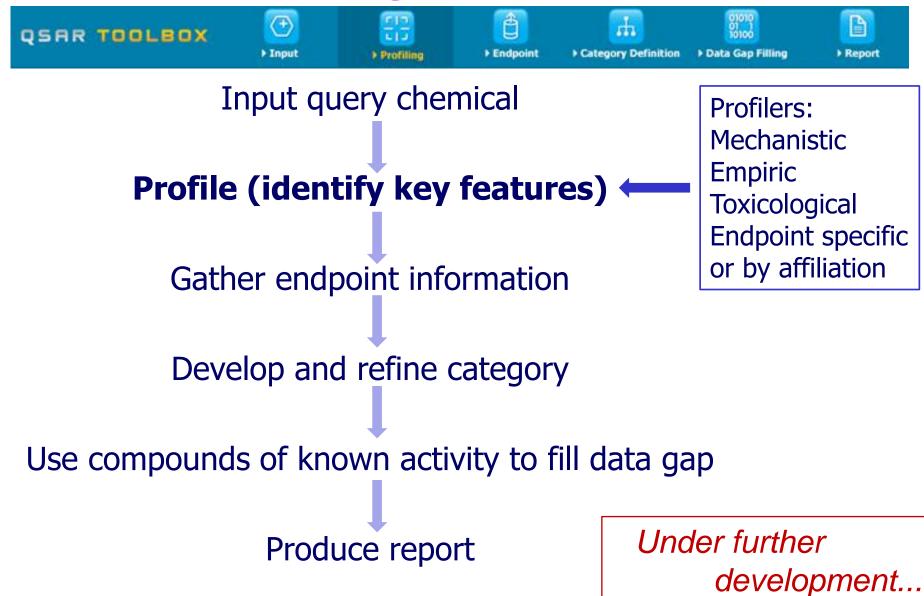
- Profilers can be used to form groups of compounds (categories)
- Structural or mechanistic knowledge of category members can be used to infer information concerning an (unknown) compound of interest i.e. a read-across prediction



Enables
 transparent,
 justifiable
 predictions to be
 made

Tools for Category Formation & Read-Across:

OECD QSAR Toolbox



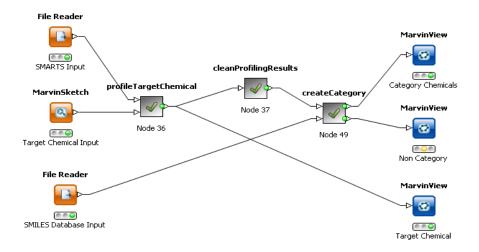
The COSMOS Project



Integrated *in silico* models for the prediction of repeated dose toxicity of **COSM**etics to **O**ptimise **S**afety

- Developing tools for predicting repeat dose toxicity
 - database of relevant toxicity & ADME data
 - building (Q)SARs
 - identifying structural alerts creating profilers for category formation
- Freely available as KNIME workflows





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Conclusions and Outlook

- AOPs provide a framework for organising information
 - The MIE is a key event () in an AOP
- Understanding mechanistic chemistry enables structural alerts / profilers to be built associated with an MIE
- Categories (groups) can be based on structural similarity & potential mechanisms investigated
- Literature or experimental evidence provides support for AOPs
 - Link in silico investigation to directed in vitro analysis
 - Confirmation of effects; defining chemical space of alert
- Improvements needed in tools to capture and use information
 - Toolbox; Effectopedia
 - Mitigating factors need to be considered
 - More quantitative predictions in future

Category formation (grouping) & read-across provide more transparent, acceptable methods of predicting toxicity

Acknowledgements

Staff Mark Cronin Steve Enoch Andrea-Nicole Richarz Katarzyna Przybylak Mark Hewitt Mark Nelms Fabian Steinmetz Przymyslaw Piechota



The funding from the European Community's Seventh Framework Program (FP7/2007-2013) COSMOS Project under grant agreement n°266835 and from Cosmetics Europe is gratefully acknowledged. Funding from the eTOX project, grant agreement no. 115002 under the Innovative Medicines Initiative Joint Undertaking (IMI-JU) is also gratefully acknowledged.

Thank you for your attention!