

# Overview of the status of predictive computer models



Expert Meeting: identification of pre- and pro-haptens

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- Non-testing approaches
- Skin sensitisation biological understanding
- (Q)SAR models
- Read-across approaches
- Expert systems
  - -Knowledge based
  - -Statistical based
  - -Hybrid
- IATA
  - -IATA models

# Continuum of non-testing approaches



Agency

Properties of a chemical and how it will interact with a defined system are inherent in its molecular structure





- Multi-disciplinary approach of integrating chemistry and toxicology using statistical approaches..
- Requires data as inputs (calculated and/measured descriptors, toxicity data) and lots of it!







#### Physico-chemical Basis of Skin Agency Sensitisation

 Skin sensitisation potential is dependent on electrophilic reactivity of the skin sensitiser or a derivative (produced by metabolism or oxidation)



Reactive



Non-reactive

Michael acceptors

**Structural Features** 

- Schiff Base formers
- $\cdot$  S<sub>N</sub>2 electrophiles
- S<sub>N</sub>Ar electrophiles

no reactive groups

X = e.g. - CHO, COR, CN

- Acylating agents
- Non-reactive

X = e.g. F, CI, Br, I, -OC<sub>6</sub>H<sub>5</sub>



- By chemical inspection
- Protein Binding Profilers within the OECD Toolbox
- <u>http://www.oecd.org/chemicalsafety/risk-assessment/theoecdqsartoolbox.htm</u>



•Reactivity domains implemented within Toxtree <u>http://toxtree.sourceforge</u> .net/download.html







- By expert judgement
- Literature searching
- Simulating metabolites using tools notably the simulators within the OECD Toolbox and the additional capabilities that are available within the TIMES-SS platform (Refer to Saby's presentation)

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United States Environmental Protection Agency pre- and pro-haptens							General Mechanistic     DPRA Cysteine peptide depletion     DPRA Lysine peptide depletion     Protein binding by OECD     Protein binding potency     Endpoint Specific	
OSAR Toolbox 3.3.5.17 [Document 1]							Keratinocyte gene expression	
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# QSAR models - Quantitative Mechanistic Models (QMMs)



Michael Acceptors: Roberts & Natsch (2009) pEC3 = 0.24 log k + 2.11 n = 10, R<sup>2</sup> = 0.836, s = 0.11, and F = 40.8 k - reactivity (rate constant for reaction with a nucleophile)
Schiff Bases: Roberts et al (2006) pEC3 = 1.12(±0.07)Ss\* + 0.42(±0.04)log P - 0.62(±0.13) n = 16, R<sup>2</sup> = 0.952, s = 0.12, and F = 129.6

logP : partitioning (octanol-water partition coefficient)

- Ss\* : reactivity (sum of Taft s\* values)
- S<sub>N</sub>Ar: Roberts & Aptula (2014)

 $pEC3 = 2.81 (\pm 0.12)RP - 5.44(\pm 0.36)$ 

 $n = 10, R^2 = 0.987, s = 0.13, and F = 594$ 

 $\bullet$  RP – reactivity parameter, which is based on a combination of  $\sigma^{\star}$  and  $\sigma-$  substituent constants



- No models available yet for
  - $-S_N 2$  domain chemicals
  - -Acylating agents
- For non-reactive chemicals
  - -Possibility of using a TTC type value Dermal Sensitisation Threshold (DST) approach (Safford et al, 2011)
  - -NB Has since been extended to incorporate chemicals categorised as reactive and to exclude "High Potency" chemicals (see Safford et al, 2015 & Roberts et al, 2015)



# Read-across approaches

- Identify analogues and evaluate them on the basis of reaction domains, structural alerts to perform a "mechanistically based read-across"
- Tools such as the OECD Toolbox\*, Toxtree, Ochem's Toxalerts, Derek Nexus are helpful to evaluate "similarity" of potential analogues





- Knowledge based e.g. Derek Nexus
- Statistical based e.g. TOPKAT, MCASE
- Hybrid TIMES-SS



- Comprises alerts, reasoning rules and examples
- Alerts are substantiated by evidence and associated references.
- The reasoning rules describe the relationships between factors such as physicochemical properties, species etc that allows a confidence level to be associated with a given prediction.
- Calculator for LogKow to make skin permeability predictions (Kp) is used to refine the confidence levels for skin sensitization predictions.
- Some alerts do have information captured for metabolites (HQ - precursor to benzoquinone)



- Static model based on GPMT data
- Gives a yes/no outcome, a potency score (semi quantitative) and information on the applicability domain
- No accounting for metabolism



- Hybrid expert system
- Based on dataset of 875 chemicals tested by LLNA, GPMT and chemicals from the BfR list
- Structure-activity and structure metabolism rules
- Incorporates an autoxidation simulator
- Mechanistically transparent
- Predicts skin sensitisation effect in three classes: strong, weak and non-sensitisers
- Reliability of skin sensitisation prediction is assessed based on applicability domain, alert performance and mechanistic justification of the protein binding mechanism.

#### Moving towards IATA pipleline approaches



 Identify plausible MIEs
 Explore Linkages in Pathways to Downstream Effects

 Develop QSARs to predict MIEs from Structure or characterise other KEs as SARs

#### 20 AOP for skin sensitisation (SS) and assays mapped to KEs

Reisinger and Hoffmann et al./Toxicology in Vitro 29 (2015) 259-270





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AOP as implemented in the OECD Toolbox

Experimentary Skin Sensitization		
Full names		
<ul> <li>1 - Protein binding alerts</li> <li>2a - in chemico Peptide depletion assay DPRA (Cys)</li> <li>2b - in chemico Peptide depletion assay DPRA (Lys)</li> <li>2c - in chemico Glutathione depletion assay GSH (RC50)</li> <li>2d - in chemico Adduct formation assay LC-MS</li> <li>3 - in vitro Keratinocyte ARE (EC1.5, EC2, EC3)</li> <li>4a - in vitro Dendritic cell activity assay h-CLAT (expression 4b - in vitro Dendritic cell activity assay MUSST (expression 5 - in vivo Organ response (LLNA)</li> <li>6 - in vivo Organism response (GPMT)</li> </ul>	20 40 1 20 3 5 5 5 5 5 5 5 6 20 8 9 1 1 20 5 5 5 5 5 5 5 5 5 5 5 5 5	6
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### ITS-2 Integrated in AOP information using a Bayesian network





- A number of in silico approaches are available
  - profiling, read-across, QSAR, expert systems
    - Limited by the availability of measured reactivity data
- Shifting towards AOP informed IATA
  - Critical to understand the scope/technical limitations of the IATA elements e.g. analytical validity of individual assays
- Limited in our ability to systematically predict air oxidation products and metabolites by in silico tools
  - in silico approaches and even IATA developed to date do not typically account for pre-or pro-haptens