

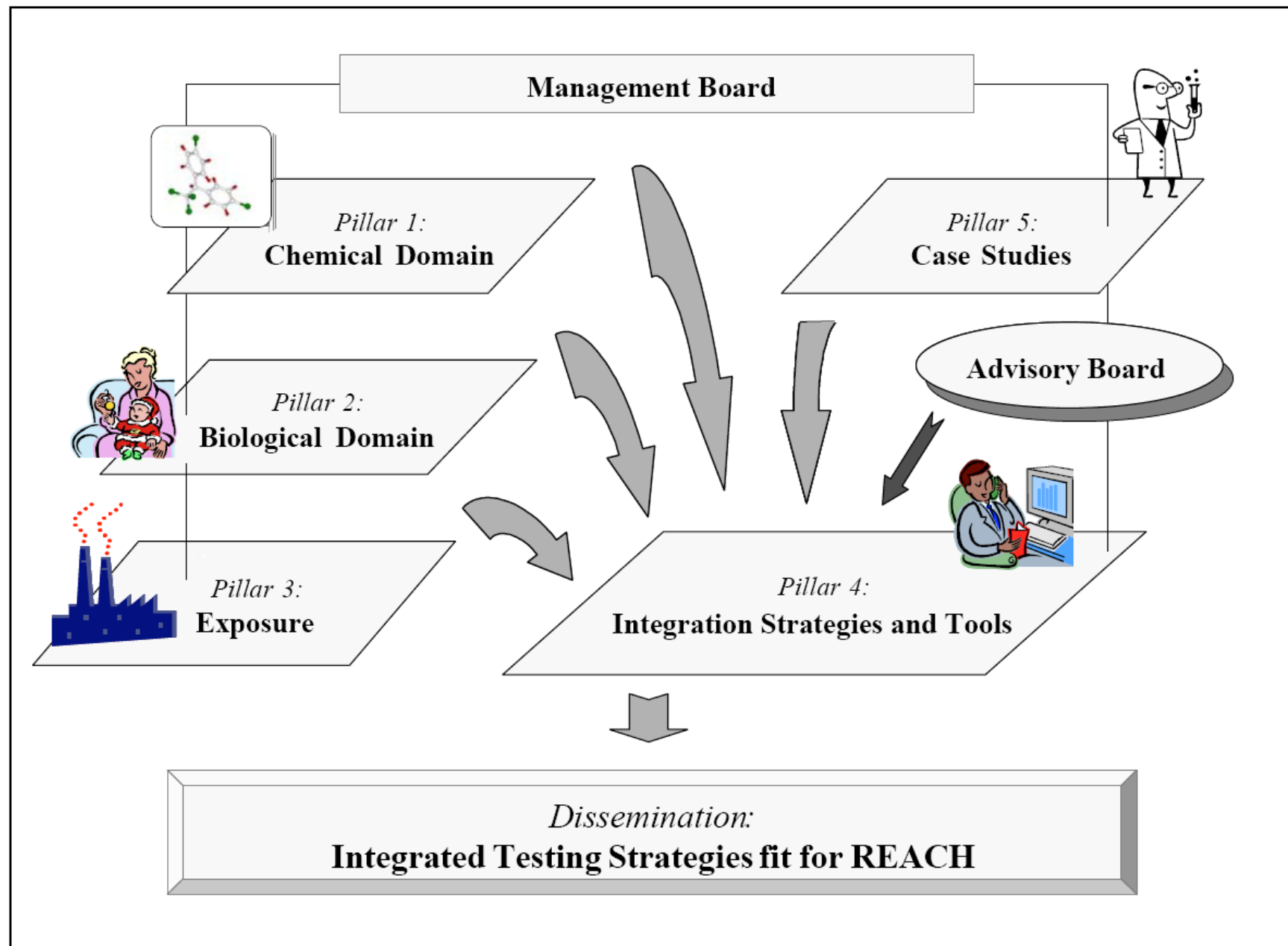
In Silico Models in the OSIRIS Web-Tool: Skin Sensitisation

Mark Cronin

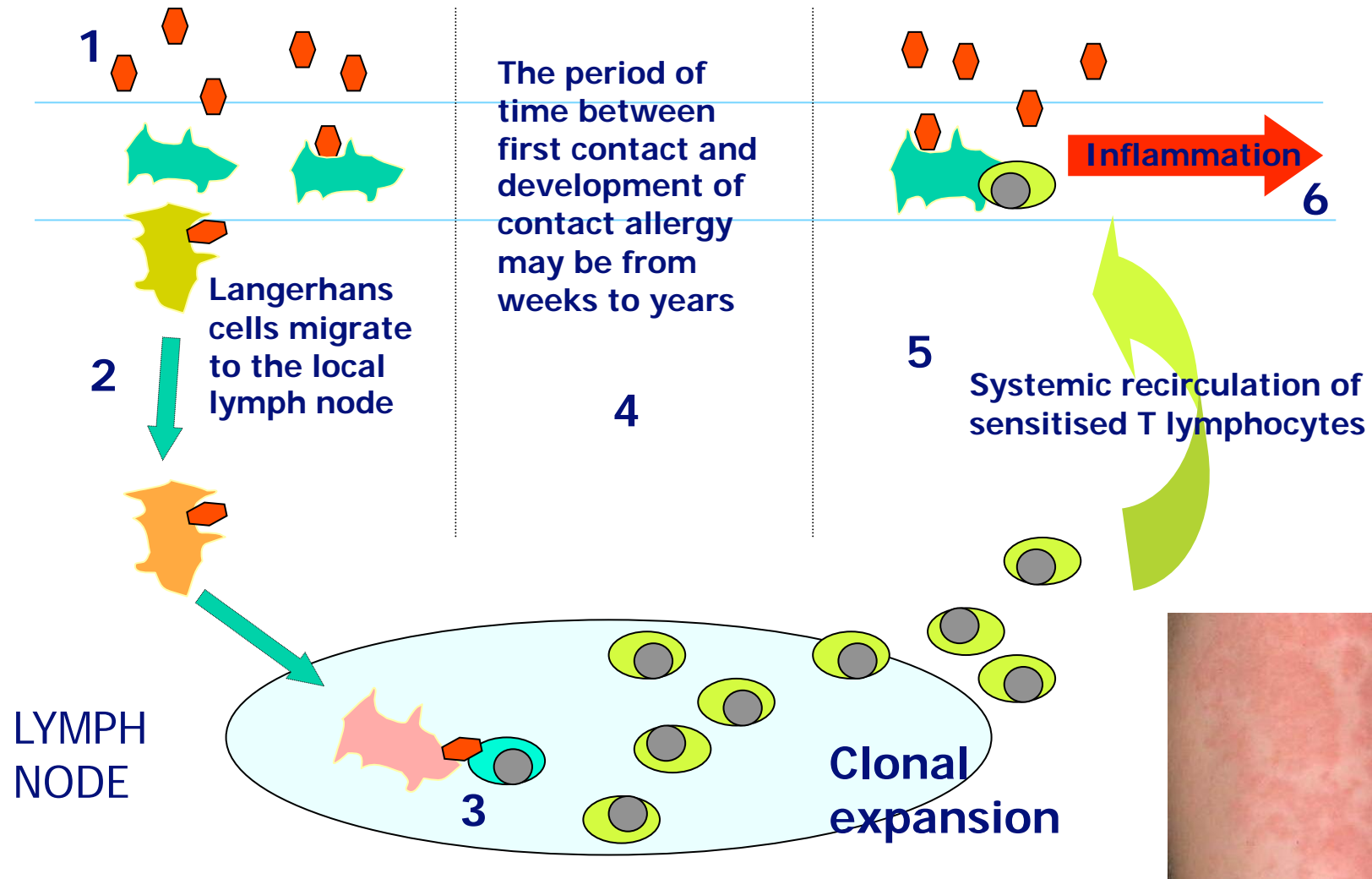
School of Pharmacy and Chemistry

Liverpool John Moores University

England



Skin Sensitisation: Current Knowledge



In Silico Approaches for Predicting Skin Sensitisation

- SMARTS Strings

- DEREK

- TIMES

- TOPKAT

- M-CASE

In the Web-Tool

- CAESAR

- OECD (Q)SAR Application Toolbox

In Silico Approaches for Predicting Skin Sensitisation

- SMARTS Strings – Fragments (organic chemistry)
 - DEREK - Structural alerts
 - TIMES - Hybrid – QSAR and metabolism
 - TOPKAT - QSAR
 - M-CASE - QSAR

 - CAESAR – QSAR
 - OECD (Q)SAR Application Toolbox - Categories
-

In Silico Approaches for Predicting Skin Sensitisation

- SMARTS Strings – Mechanistic (chemistry)
 - DEREK - Mechanistic (chemistry / biology)
 - TIMES - Mechanistic (chemistry / biology)
 - TOPKAT – Non-mechanistic
 - M-CASE - Non-mechanistic

 - CAESAR – Non-mechanistic
 - OECD (Q)SAR Toolbox – Mechanistic (chemistry)
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In Silico Approaches for Predicting Skin Sensitisation

- SMARTS Strings – Freely available (via e-mail)
 - DEREK - Commercial (not for profit)
 - TIMES - Commercial
 - TOPKAT - Commercial
 - M-CASE - Commercial

 - CAESAR – Freely available via internet server
 - OECD (Q)SAR Toolbox – Freely downloadable
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SMARTS Strings Described Reaction Chemistry Mechanistic Applicability Domains

Michael acceptor

S_NAr

S_N2

Schiff base

Acyl transfer

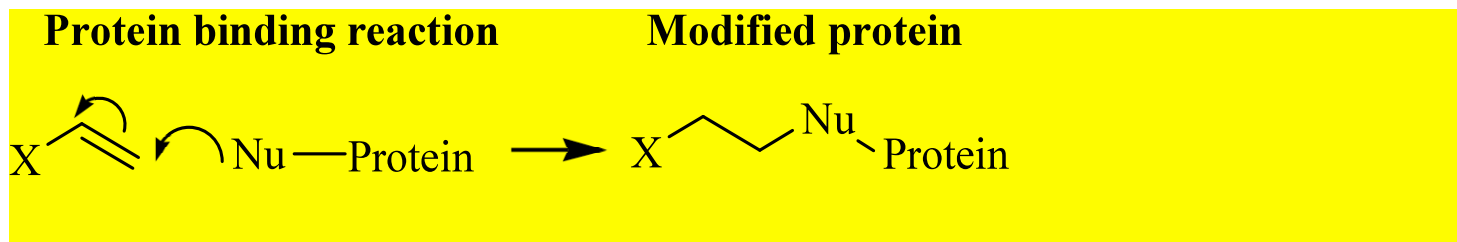
S_N1

Non-reactive, non-prereactive

Enoch SJ *et al* (2008) *Chem. Res. Toxicol.* 21:513-520

Rules for Mechanistic Domain Assignment

Michael Acceptor Domain



Identification characteristics:

Double or triple bond with substituent X
-CHO, -COR, -CO₂R, -CN, -SO₂R, -NO₂, also
2- or 4- pyridino
p-, o-quinones

SMARTS Strings – Identify Chemistry

<i>Mechanism</i>	<i>SMARTS pattern</i>
S_NAr	<pre>c1([F, Cl, Br, I, \$(N(=O) ~ O)])c([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))]cc[F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))]ccl c1([F, Cl, Br, I, \$(N(=O) ~ O)])c([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))]cccc1([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))] c1([F, Cl, Br, I, \$(N(=O) ~ O)])ncc([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))]ccl c1([F, Cl, Br, I, \$(N(=O) ~ O)])ncccc1([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O))] c1([F, Cl, Br, I, \$(N(=O) ~ O)])ncccn1 c1([F, Cl, Br, I, \$(N(=O) ~ O)])nncnc1 c1([F, Cl, Br, I, \$(N(=O) ~ O)])ncc([F, Cl, Br, I, \$(N(=O) ~ O), \$(C#N), \$(C=O), \$(C(F)(F)F), \$(S=O)])nc1 c1nc([F, Cl, Br, I, \$(N(=O) ~ O)])ncn1</pre>

A simple tool is freely available to use the SMARTS strings

Enoch SJ *et al* (2008) *SAR QSAR Environ. Res.* 19:555-578

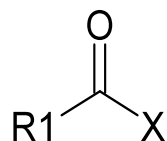
Derek for Windows / Derek Nexus

- Knowledge based expert system
 - www.lhasalimited.org
 - Searches molecule for structural alerts (fragments) of a molecule associated with skin sensitisation
 - Fragments developed by expert toxicologists
 - If no fragments are found, then DfW states "*nothing to report*" – not necessarily a negative prediction
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Derek for Windows / Derek Nexus

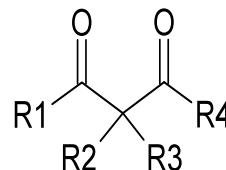
- Derek (ver 11) contains 65 alerts for skin sensitisation

Alert 401 - Carboxylic acid halide



R1 = Any
X = F, Cl, Br

Alert 420 - 1,3-Diketone



R1, R4 = C
R2, R3 = Any

- It does not have a “skin sensitisation database”
 - Alerts are supported by data
 - Some approaches to applicability domain exist
-

TIMES

- A hybrid system
 - QSARs for reactive chemistry
 - Prediction of skin metabolism
 - www.oasis-lmc.org
 - Identifies reactive chemistry associated with protein binding (47 alerting groups)
 - Skin metabolism simulator predicts plausible metabolites (from 236 transformations)
 - Approximately 800 compounds (LLNA, GPMT, BfR)
 - Assessment of applicability domain
-

TOPKAT

- A (statistical) QSAR system based on QSARs developed from discriminant analysis of 2-D structural descriptors
 - Molecular connectivities, e-state indices
 - Approximately 300 compounds (GPMT)
 - Predicts non/ weak / moderate / strong sensitisers
 - Optimum prediction space (OPS) for applicability domain
 - <http://accelrys.com/products/discovery-studio/predictive-toxicology.html>
-

M-CASE

- A (statistical) QSAR system based on molecular fragments
 - Molecules are divided up into fragments from two to n atoms
 - Fragments related statistically to any biological activity
 - Fragments are related to sensitisation (biophores) or non-sensitisation (biophobes)
 - Fragments could be used as structural alerts or more commonly in a regression approach
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M-CASE

- It is possible to put a mechanistic interpretation on fragments
 - Over 1000 compounds (Human data)
 - Identifies fragments in a molecule not found in the training set i.e. a form of applicability domain
 - www.multicase.com
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OECD Toolbox – Not in Webtool

OECD Toolbox 1.1.01

OECD
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Options Tracks

Chemical input Profiling Endpoints Category definition Filling data gap Report

Apply

Profiling methods

- Benigni/Bossa rulebase
- BFR rulebase for eye irritation
- BFR rulebase for skin irritation
- BioWin MITI fragments
- Cramer rules
- DNA Binding
- EcoSAR Classification
- ER-binding
- OASIS Acute Toxicity MOA
- Organic functional groups
- Protein Binding
- Superfragment profiling
- Verhaar scheme

Empiric

- Chemical elements
- Groups of elements
- Lipinski Rule

Metabolism

Documented

- Observed Liver metabolism
- Observed Microbial metabolism

Simulated

- GI tract simulator
- Hydrolysis
- Liver metabolism simulator

Show Category Boundaries

Create a new profiler

Delete profiler

Structure

1 (Target)

O=[N+]([O-])c1ccc(Cl)c(Cl)c1

Substance Information

Profile

- Database Affiliation
 - CANADA Bioaccu...
 - Danish EPA
 - EPISUITE_OBS_D...
 - OASIS Aquatic
 - OASIS Genotox
 - US-EPA ECOTOX
- Inventory Affiliation
 - Danish EPA
 - EU EINECS
 - US EPA TSCA
- OECD categorization
 - (N/A)
- Substance type
 - Discrete chemical
- US EPA Categorization
 - Neutral Organics
- Protein Binding
 - Nucleophilic substit...
- Chemical elements
 - Group 14 - Carbon C
 - Group 15 - Nitrogen N
 - Group 16 - Oxygen O
 - Group 17 - Halogen...
- Groups of elements
 - Halogens
 - Non-Metals
 - Molecule satisfies t...
- Lipinski Rule

Single chemical

- A tool to Profile for reactive chemistry

- Allows for category formation

CAESAR Model – Not in Webtool

- QSAR approach based on nearest neighbours as defined by 2-D descriptors
- 211 compounds (LLNA)
- www.caesar-project.eu



Conclusions

- Webtool contain five QSAR approaches
 - QSAR approaches will overlap, but differ significantly
 - Mechanistic vs non-mechanistic
 - Modelling approaches
 - QSAR vs protein binding
 - Number of compounds
 - LLNA vs GPMT vs Human
 - This is not an exercise to find the “best” QSAR
 - Different approaches may provide complementary information
 - We need a method to integrate the predictions
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