


Addressing Toxicity Risk when Designing and Selecting Compounds in Early Drug Discovery

Lhasa Limited vICGM, 18th June 2014

- Scott McDonald – Lhasa Limited
- Matthew Segall – Optibrium



Overview

- Lhasa and Derek
 - Optibrium and StarDrop
 - Derek Nexus and StarDrop
- 


Who are Lhasa Limited

- Not-for-profit organisation
- Registered educational charity
- Controlled by our members
- Expertise in developing *in silico* prediction and database systems





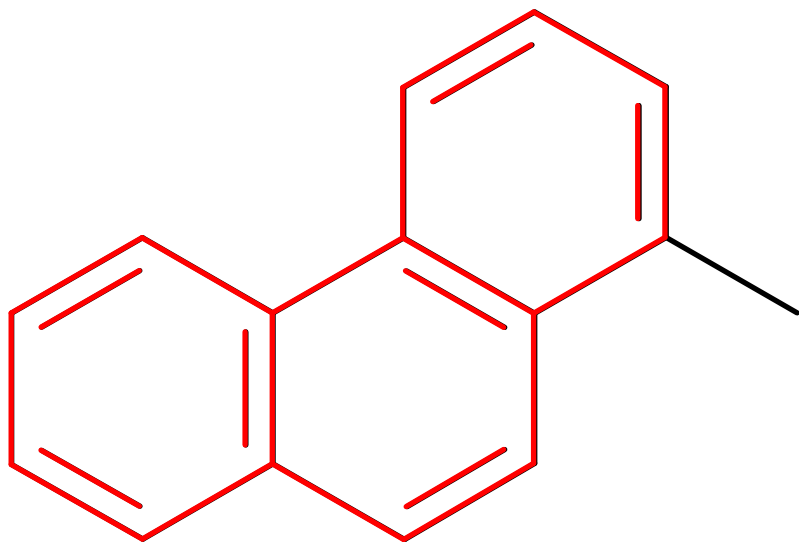
Derek Nexus

- Knowledge based expert system
 - Enables the evaluation of the potential toxicity of chemicals
 - Decision support tool
 - Accuracy
 - Transparency
 - Supporting data
 - Covers a broad range of toxicity endpoints
- 

Derek Nexus

Alert 754 – Mutagenicity in vitro


Bacterium - PROBABLE



- Knowledge base search for matching structural alerts
- Application of rules – level of likelihood
- Supporting information provided



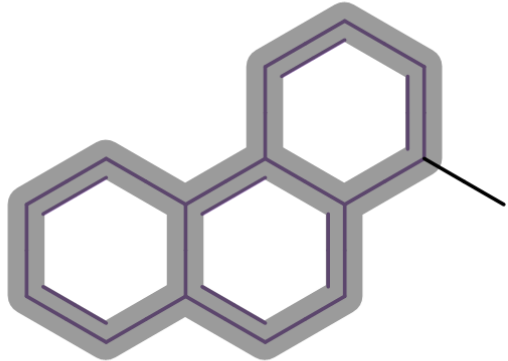
Levels of Likelihood

- Certain
 - There is proof that the proposition is true
 - Probable
 - At least one strong argument that the proposition is true and no arguments against it
 - Plausible
 - The weight of evidence supports the proposition
 - Equivocal
 - An equal weight of evidence for and against the proposition
 - Doubted, Improbable, Impossible, Open, Contradicted
- 

Nexus

File Window Prediction Reports Tools Help

Derek Prediction-2



Displaying 'Alert 754', click above to view the original structure

Prediction Navigator

Show predictions of at least: EQUIVOCAL

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

- Mutagenicity in vitro
 - bacterium - PROBABLE
 - Alert - 754: Phenanthrene derivative or hetero-analogue
 - Example - 1-methylphenanthrene
 - Skin sensitisation
 - mammal - PLAUSIBLE
 - Alert - 466: Bay-region polycyclic aromatic hydrocarbon

Alert Details Reasoning Explorer Prediction Constraints

754: Phenanthrene derivative or hetero-analogue

Alert Matches

Description Image

Comments

Mutagenicity alert: Ames test

This alert describes the mutagenicity of substituted phenanthrenes (I), methylene or carbonyl bridged phenanthrenes (II) and their hetero-analogues that are active in the Ames test. Examples include 1-methylphenanthrene [NTP 1989, LaVoie et al 1983], 3H-cyclopenta[c]phenanthrene [Marrocchi et al], 4,10-dimethylphenanthrene [LaVoie et al 1983], 9-fluorophenanthrene [LaVoie et al 1983] and 1-methyl-4H-cyclopenta[def]phenanthrene [Rice et al]. These examples have all been reported to be mutagenic in *Salmonella typhimurium* TA100 in the presence of rat S9 activation. It appears that molecules in these classes must contain activating features in order to be mutagenic; phenanthrene is negative [LaVoie et al 1983] and 4H-cyclopenta[def]phenanthrene is only positive with strong metabolic activation [NTP 1987].

The mutagenic activity of phenanthrenes and methyl bridged phenanthrene derivatives is likely to be mediated by electrophilic metabolites. It has been proposed that 7,8-dihydrodiol-5,6-epoxide is the ultimate mutagen. The corresponding proximate mutagen, 7,8-dihydrodiol, has been observed in vitro for 1,4- and 4,10-dimethylphenanthrene [LaVoie et al 1982] and 15,16-dihydro-1,11-methanocyclopenta[a]phenanthrene-17-one [Hadfield et al]. Furthermore, DNA adducts from the 7,8-dihydrodiol metabolite of 15,16-dihydro-11-methylcyclopenta[a]phenanthren-17-one have been observed in vitro [Coombs et al 1979]. Alternative routes of metabolic activation have been observed for some phenanthrenes, particularly those that are unsubstituted at the 4-position including formation of non-bay-region dihydrodiols. The proximate mutagenic metabolites of 1-methylphenanthrene and 9-methylphenanthrene have been identified as either the 3,4- or 5,6-dihydrodiols [LaVoie et al 1981]. DNA adducts from the former in HepG2 cells have been reported, albeit at a very low level compared with the more potent mutagens dibenzo[a]pyrene or dibenzo[a,h]anthracene [Staal et al]. Metabolism of the K-region and occasionally alkyl substituents can reduce the mutagenicity of molecules, e.g. 2-methylphenanthrene [LaVoie et al 1981].

The scope of this alert has been defined by the common structural features of the active compounds in this class and mechanistic considerations. Because the bay-region diol-epoxides are potential ultimate mutagens, the 7,8-dihydrodiol proximate mutagens have been included. Substituted benzoquinolines are also included because it is possible for them to be metabolised to bay-region diol-epoxides [Sasaki et al]. There are three features that

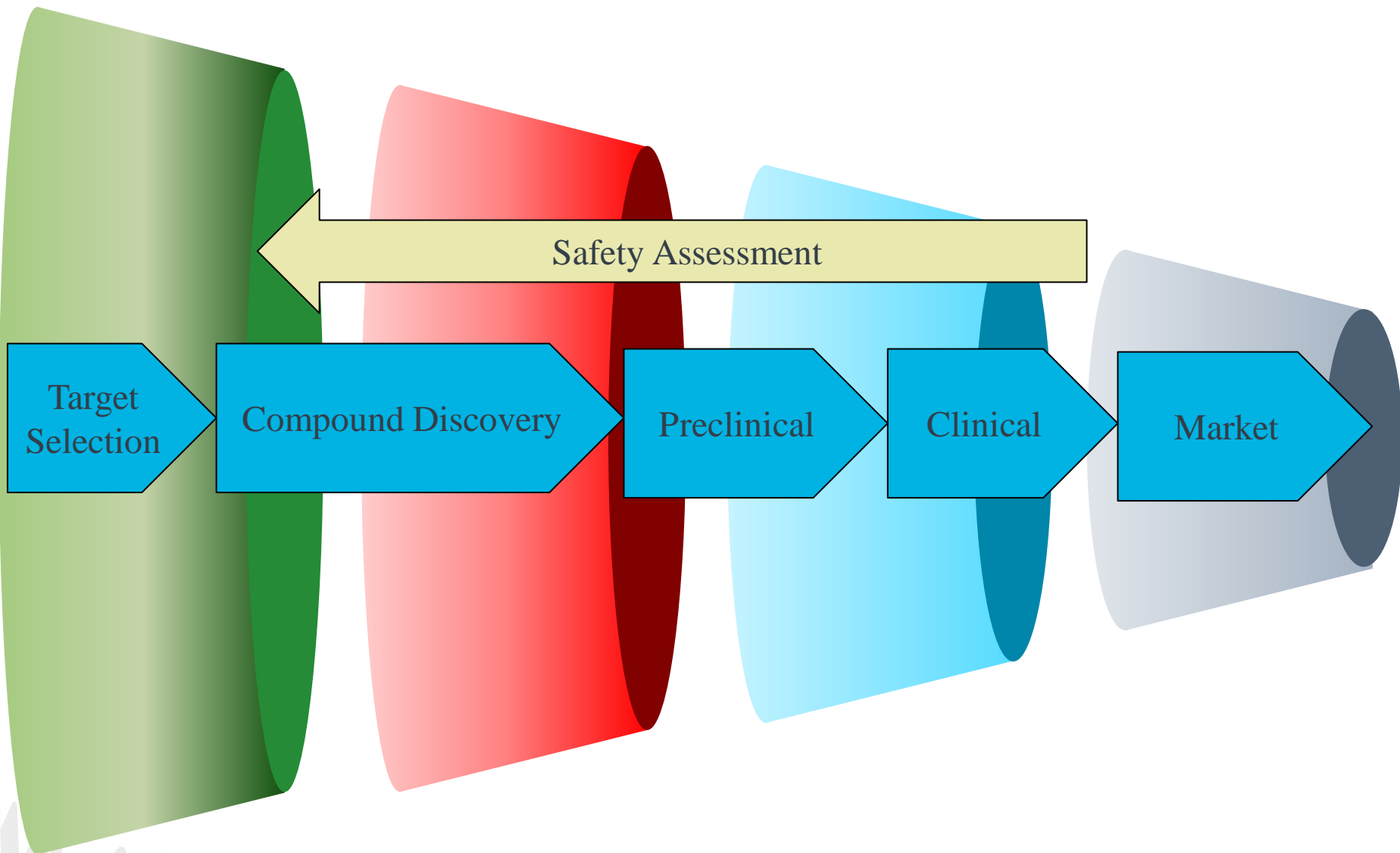
Validation Comments

Endpoints

References


ID	Title	Author	Source	Year	Supplemental
4632	Mutagenicity of s	LaVoie EJ, T	Mutation R	1983	DOI: 10.1016/0165-1218(83)90100-3, PI
7835	Identification of tl	Coombs M	Cancer Res	1979	PMID: 476652
7847	Mutagenicity, tun	LaVoie EJ, T	Cancer Res	1981	PMID: 7020927
784E	Tumor-initiating ;	LaVoie EJ, E	Cancer Res	1982	PMID: 7105001
7975	Metabolism of be	Amin S, Lin	Chemical R	2003	DOI: 10.1021/bx0200921, PMID: 125881
1401	Salmonella study .	National Tc	National Tc	1989	Note: available at "http://ntp-server.n
1401	Synthesis and mu	Marrocchi ,	Carcinoger	1996	DOI: 10.1093/carcin/17.9.2009, PMID: 8

Drug Discovery





Lhasa and Optibrium

- Collaboration commenced in 2013
 - Development of the Derek Nexus Module for StarDrop
 - Available as an optional module
 - Facilitates the design of safe drugs in hit-to-lead and lead optimisation
- 



optibrium

Addressing Toxicity Risk when Designing and Selecting Compounds in Early Drug Discovery

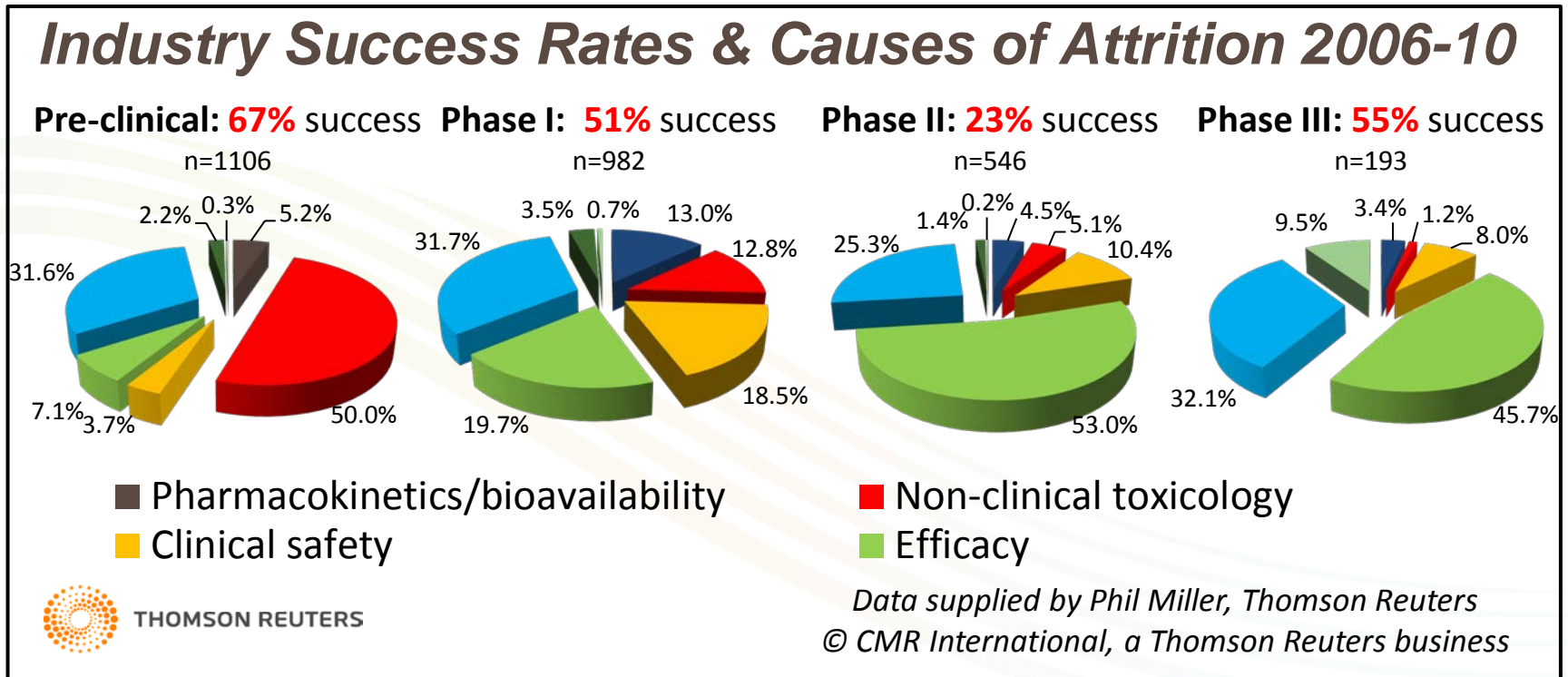
Lhasa Limited vICGM, 18th June 2014

Matthew Segall, Chris Barber

Overview

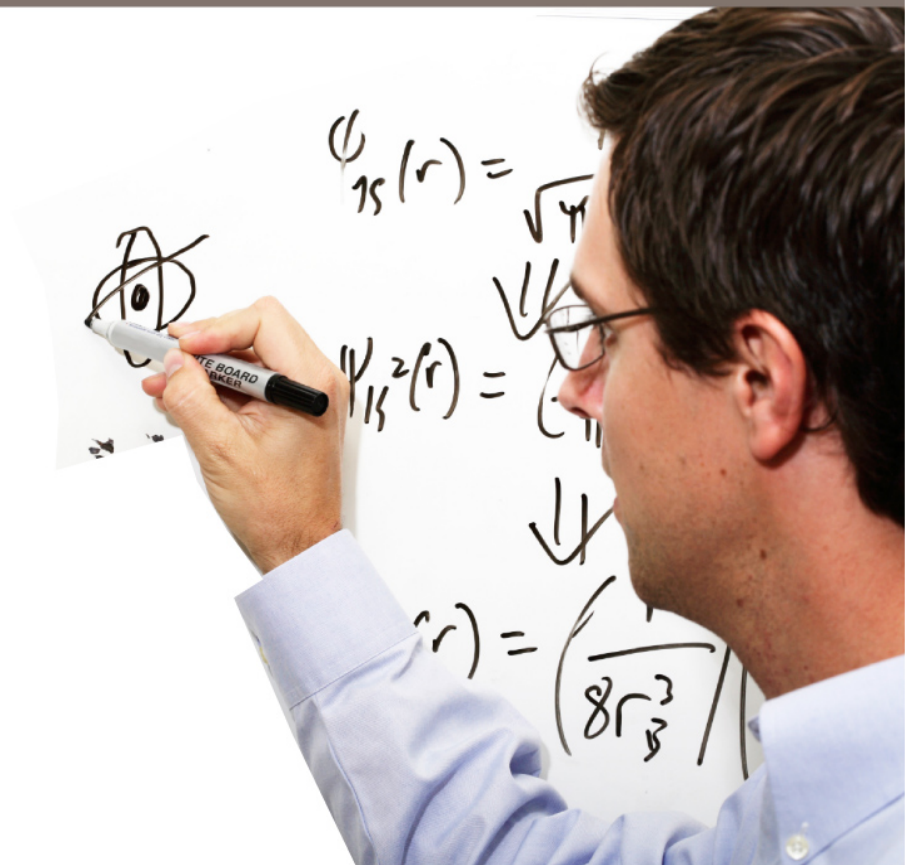
- Impact of toxicity in pharma. R&D
- Application of knowledge based prediction of toxicity
- Guiding compound selection and design
 - Multi-parameter optimisation
 - Glowing Molecule
- Example
 - Exploring a COX2 screening library
- Short overview of StarDrop and the Derek Nexus module
- Conclusions

Impact of Toxicity in Pharma R&D



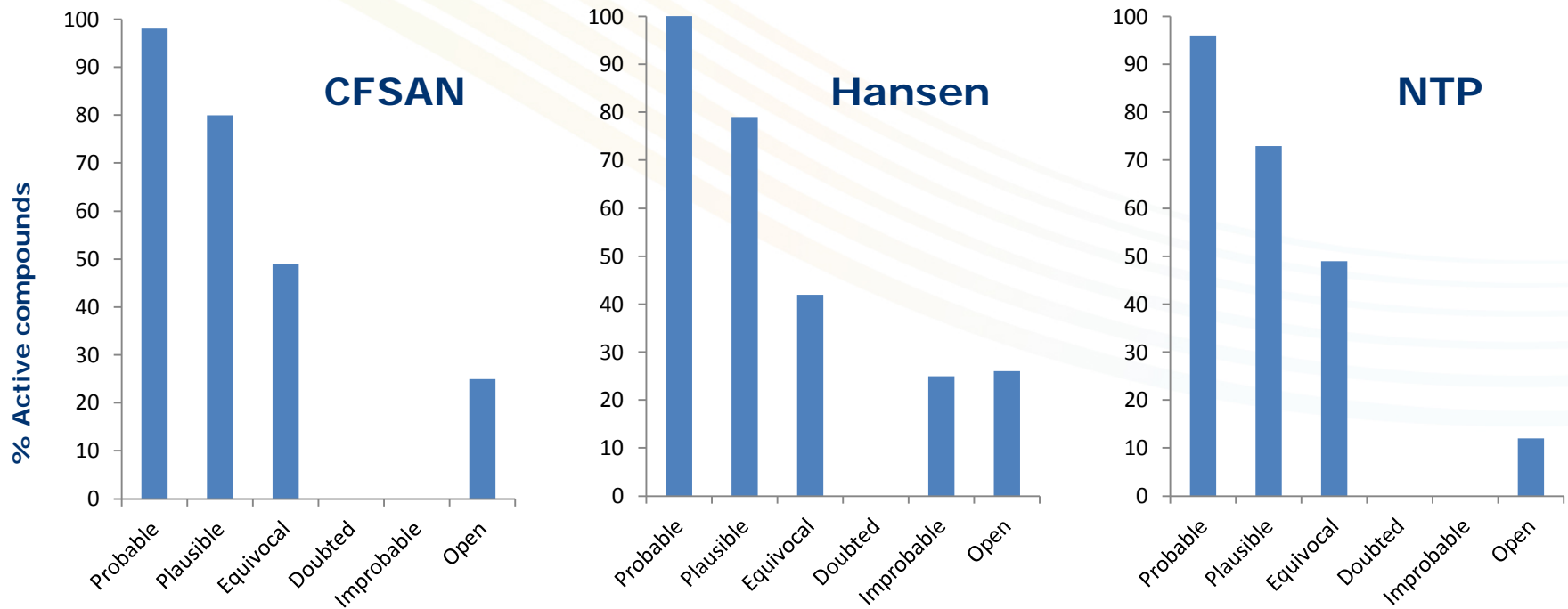
- 54% of pre-clinical failures due to tox/safety (18% of all candidates)
- 22% of all clinical candidates failed due to tox/safety
- 10.2% of approved drugs acquired black box warning, 2.9% withdrawn*

Application of Knowledge Based Prediction of Toxicity



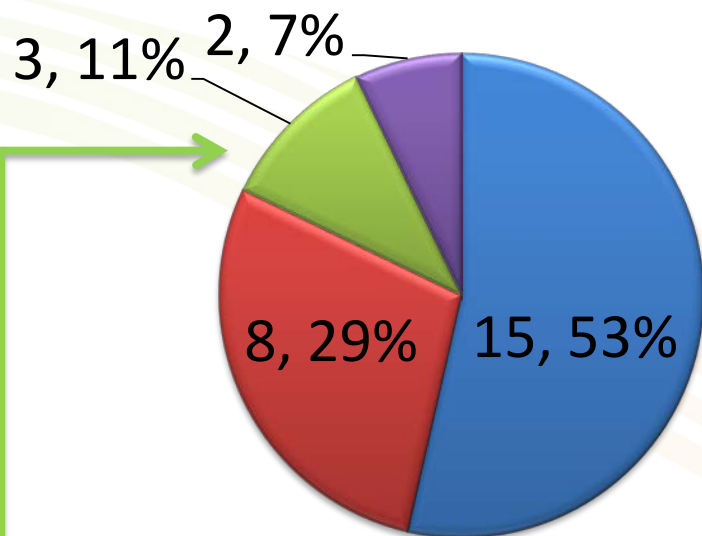
Relating confidence and accuracy

- Derek Nexus provides a level of confidence (likelihood) for each prediction
 - This correlates well with accuracy



How Well do Expert Systems Perform?

- CDER approved drugs 2012 (n=27)



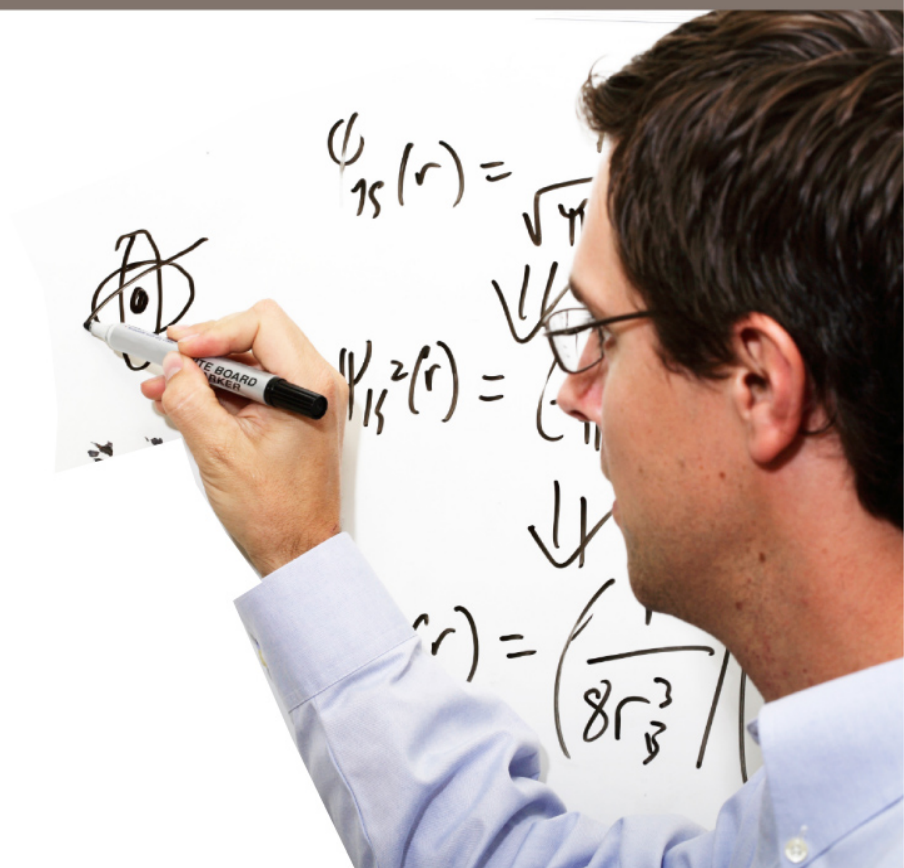
- true prediction of clean
- true prediction of dirty
- false prediction of dirty
- false prediction of clean

drug indication	prediction	dosing
Ingenol <i>actinic keratoses</i>	chromosomal damage	topical treatment (cytotoxic mechanism)
Acidinium <i>COPD</i>	hepatotoxic	inhaled (0.4mg dose)
Linaclotide <i>IBS</i>	hepatotoxic	metabolised in GI tract

Important Caveats/Questions

- Predict toxicity hazard
 - Risk \approx hazard + exposure
 - Risk also depends on dose, route of administration, therapeutic index...
- Knowledge-based prediction of toxicity widely used in preclinical development
 - Assessment of risk for regulatory submission
 - Design of experiments to support submissions
- Question: How can these predictions be applied effectively in early drug discovery?
 - We don't want to 'kill' potentially good compounds at an early stage due to uncertain predictions

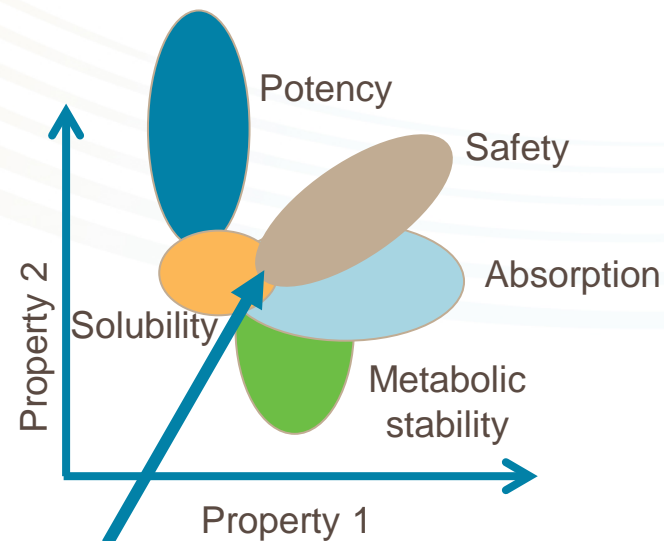
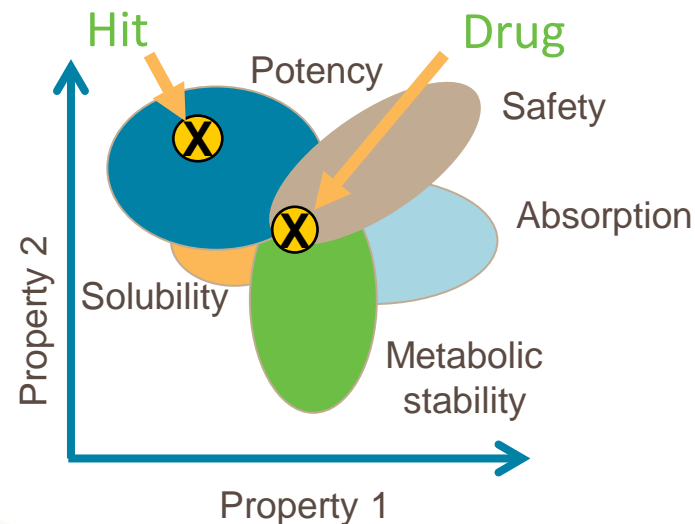
Guiding Compound Design and Selection



The Objectives

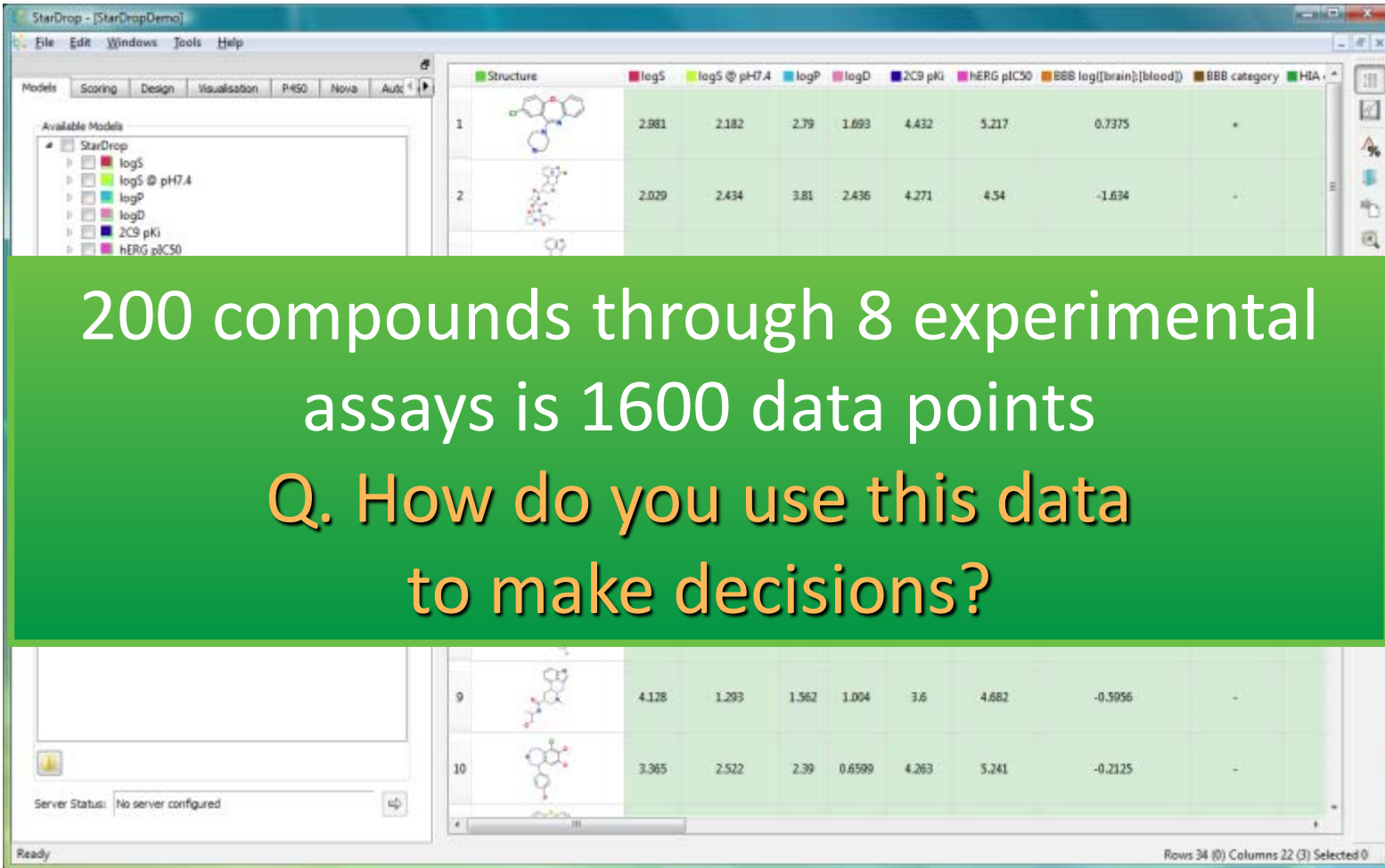
Multi-parameter optimisation

- Identify chemistries with an optimal **balance** of properties
- Quickly identify situations when such a balance is not possible
 - Fail fast, fail cheap
 - Only when **confident**



No good drug

The Challenge



The screenshot displays the StarDrop software interface. On the left, a tree view shows 'Available Models' including StarDrop, logS, logS @ pH7.4, logP, logD, 2C9 pKi, and hERG pIC50. The main window shows a table with 2 columns for chemical structures and 8 columns for assay results. The table contains 10 rows of data, with the first two rows highlighted in green. A green text box is overlaid on the table, containing the text: '200 compounds through 8 experimental assays is 1600 data points Q. How do you use this data to make decisions?'. At the bottom left, a 'Server Status' field shows 'No server configured'. At the bottom right, the status bar indicates 'Rows 34 (0) Columns 22 (3) Selected 0'.

200 compounds through 8 experimental assays is 1600 data points

Q. How do you use this data to make decisions?

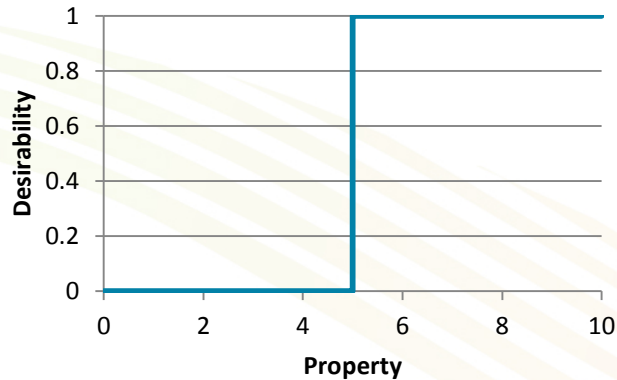
Approaches for MPO Filtering?



Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome

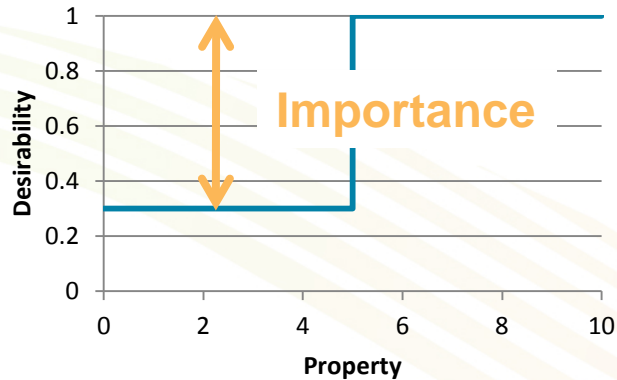


Simple filter: >5

Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome

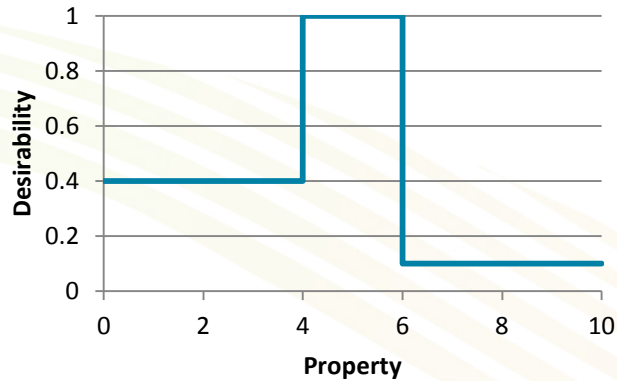


Desired value: >5

Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome

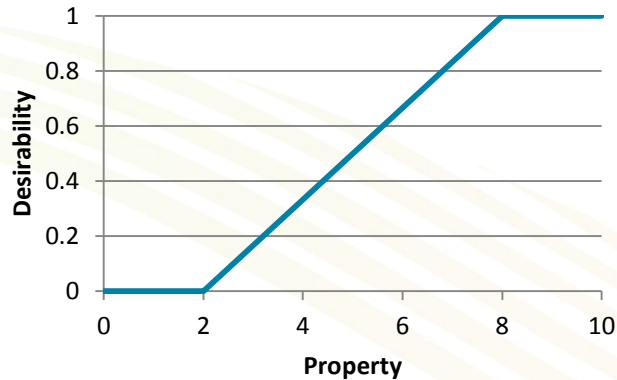


Rang: 4-6

Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome

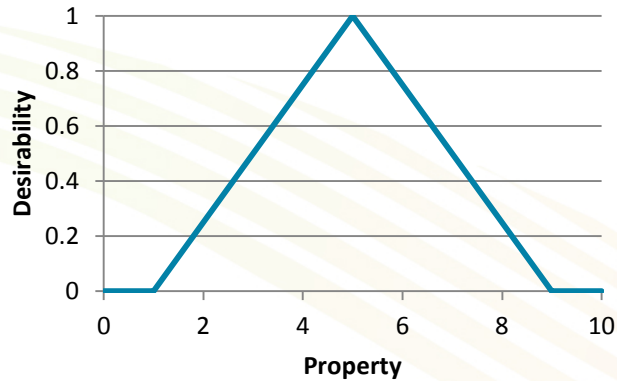


Trend: >8

Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome

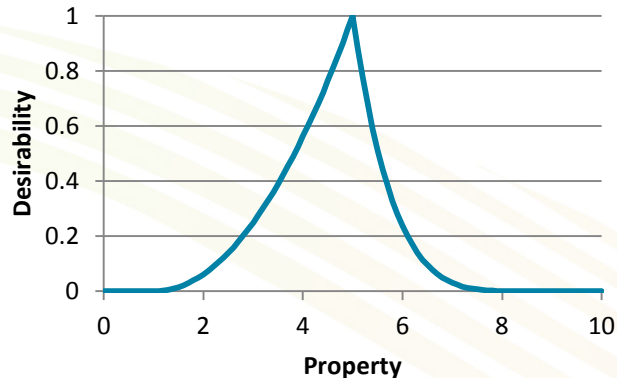


Ideal value: 5

Approaches for MPO

Desirability Functions*

- Relate property values to how 'desirable' the outcome



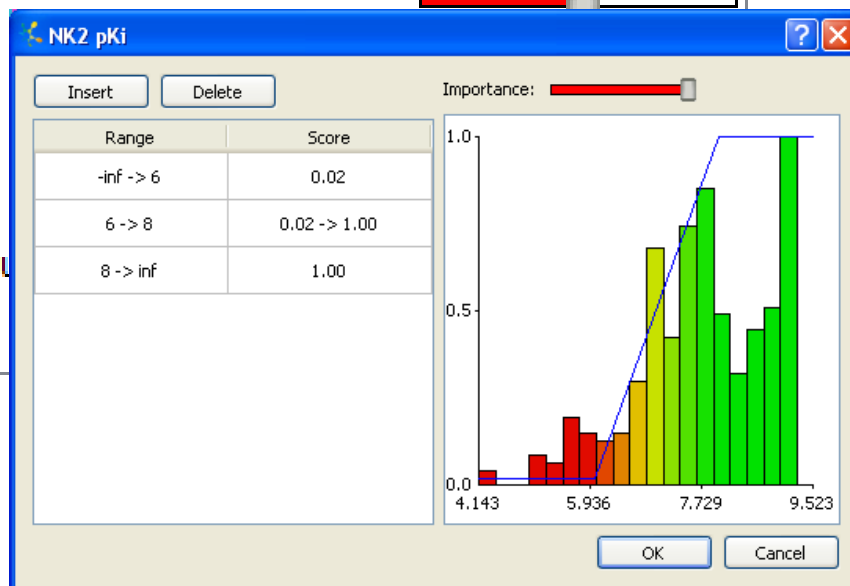
**Non-linear, ideal value: 5
(Derringer Function)**

- Combine multiple properties into 'desirability index'
 - Additive:
 - Multiplicative:
- Very flexible approach allowing parameters to be weighted
- But, does not explicitly consider uncertainty

Approaches for MPO

Probabilistic Scoring*

Profile	Desired Value	Importance
logS	> 1	
HIA category	+	
logP	0 -> 3.5	
BBB log([brain]:[blood])	-0.2 -> 1	
BBB category	+	
P-gp category	no	
hERG pIC50	≤ 5	
2C9 pKi	≤ 6	
2D6 affinity category	low medium	
PPB90 category	low	



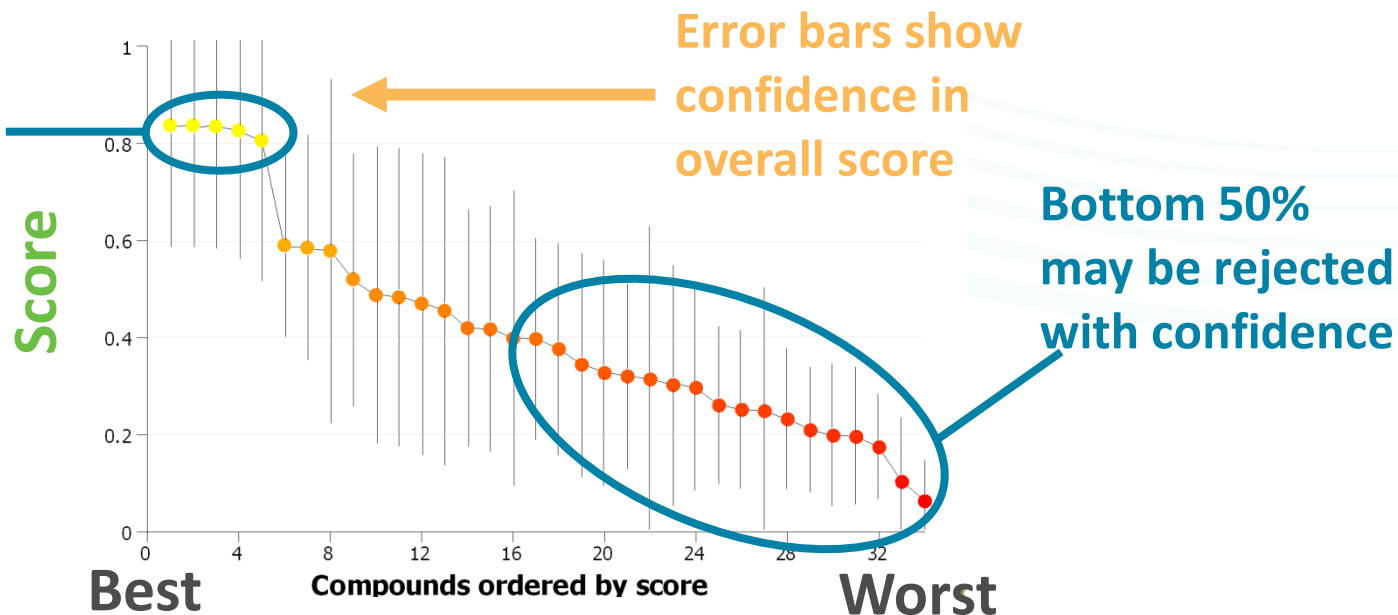
Approaches for MPO

Probabilistic Scoring*

- **Property data**
 - Experimental or predicted
- **Criteria for success**
 - Relative importance
- **Uncertainties in data**
 - Experimental or statistical

- **Score (Likelihood of Success)**
- **Confidence in score**

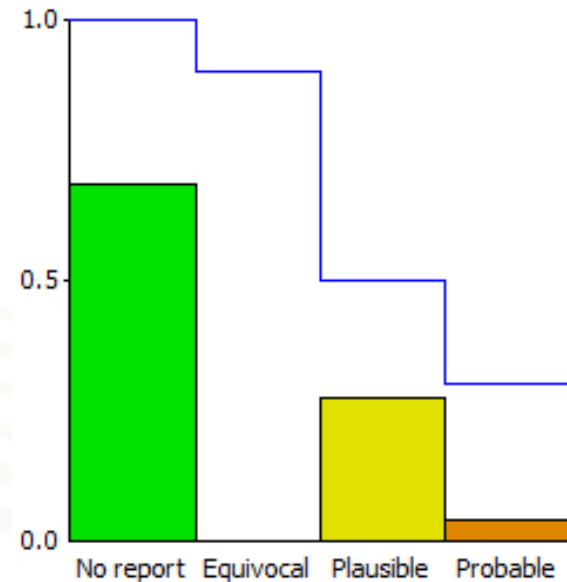
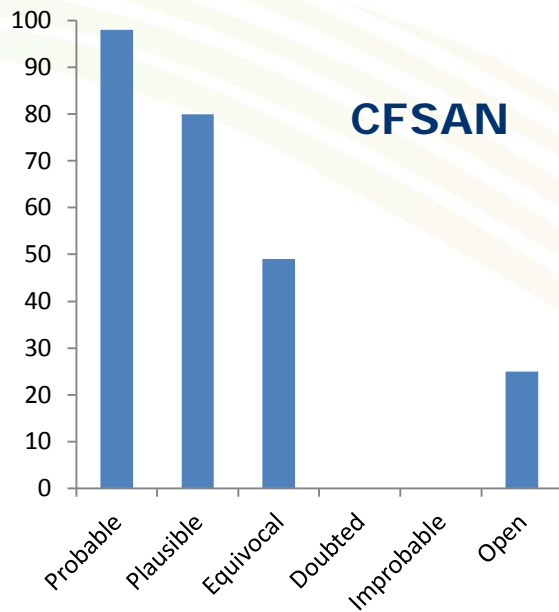
Data do not separate these as error bars overlap



Application to Toxicity Alerts

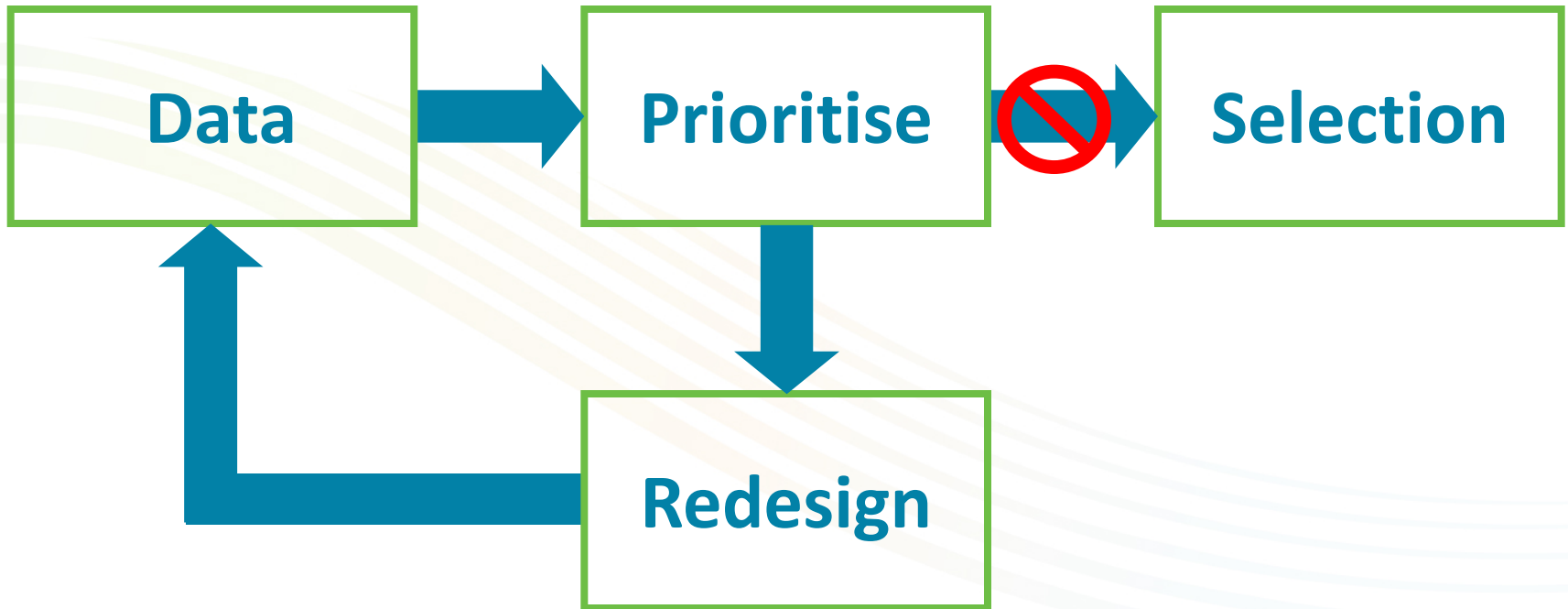
E.g. Mutagenicity

- Determine desirability function by reference to validation results:



- Also need to take into account:
 - Impact of toxicity on objective of project
 - Stage of the project, e.g. opportunity to redesign to reduce risk

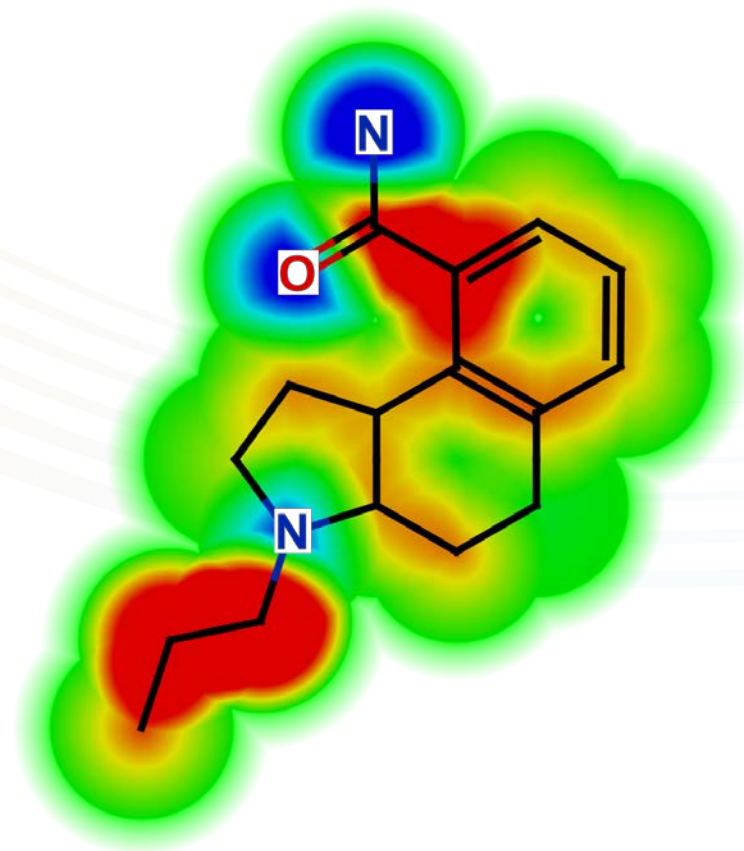
Guiding Interactive Redesign



Interpretation of a Model

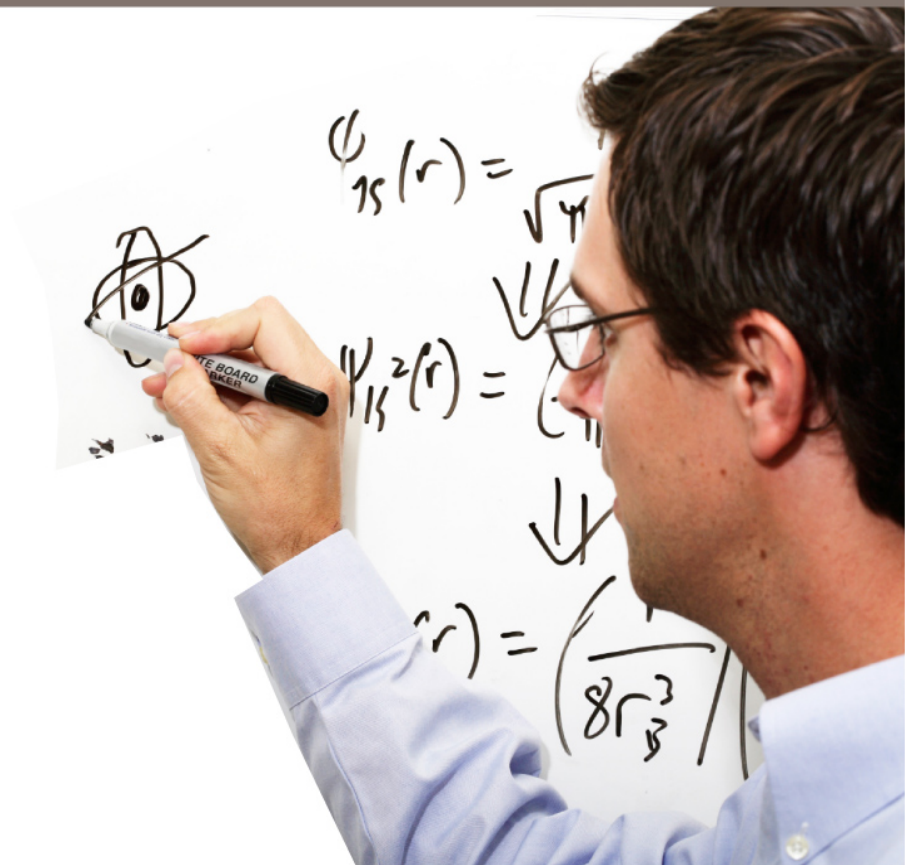
The 'Glowing Molecule'

- Provides visual interpretation of structural influences on predicted properties
 - “Why is a property value predicted?”
 - “Where can I change this property?”
 - Interpret SAR
 - Guide efficient redesign of molecules
- Avoid Black Boxes



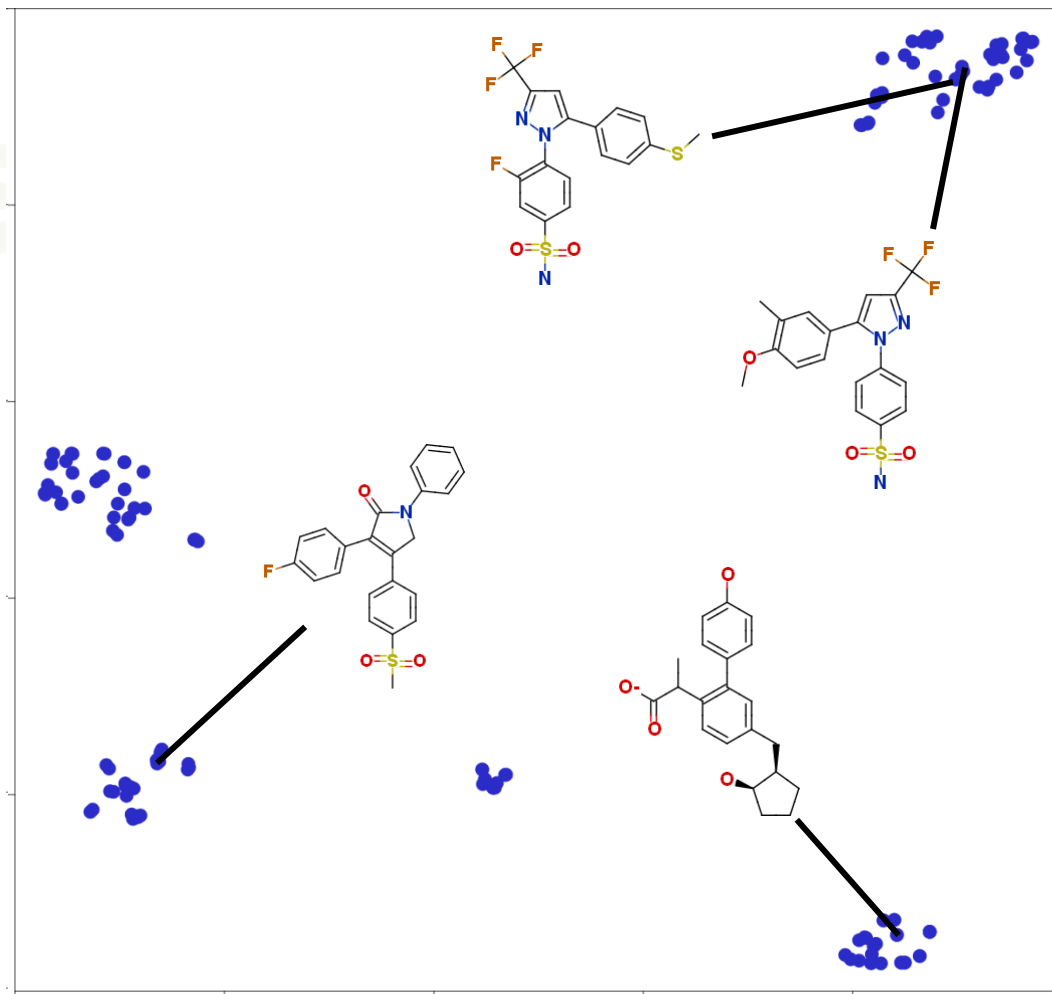
Example Application

Exploring a COX2 screening library



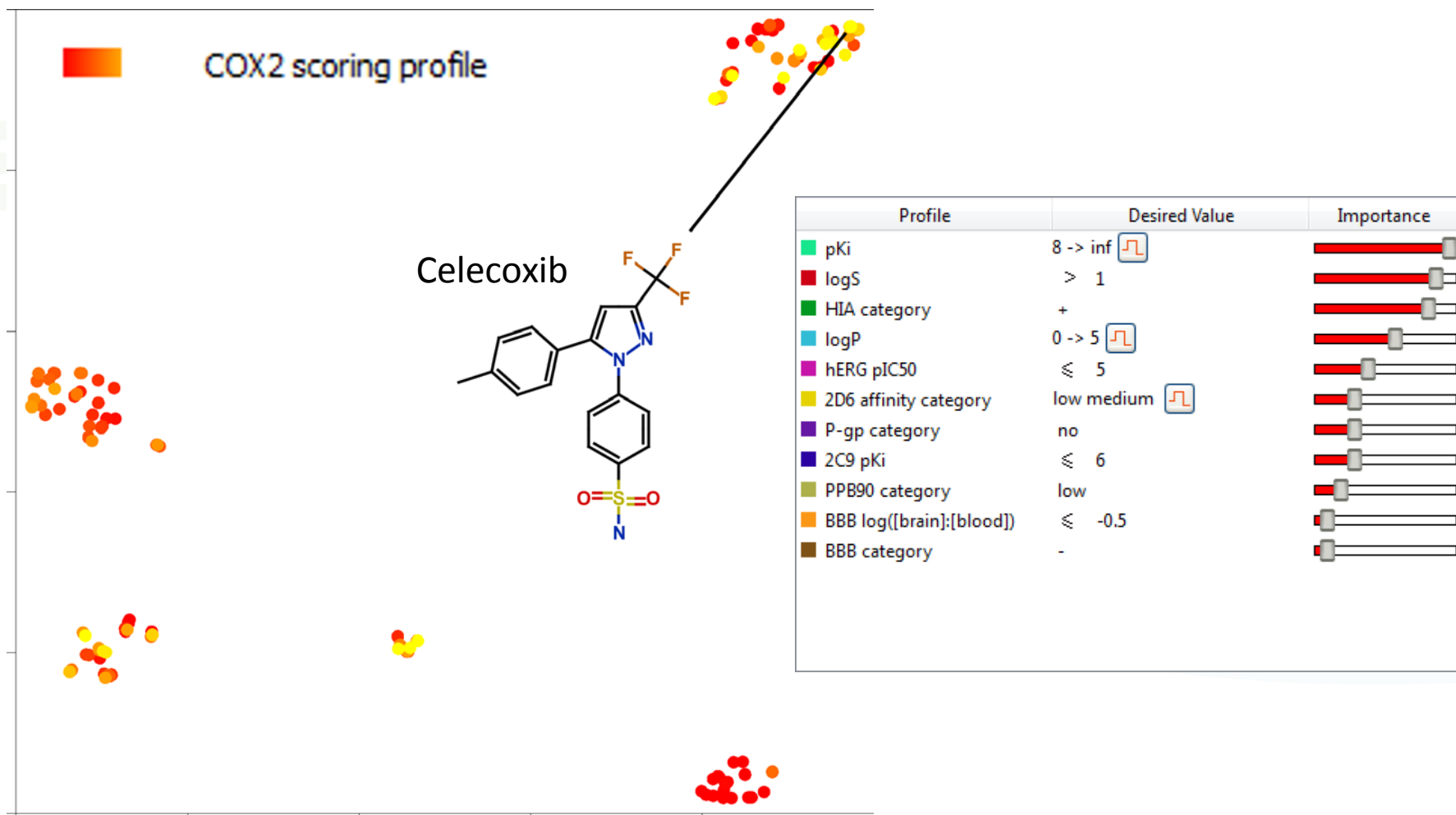
COX2 library

Chemical space



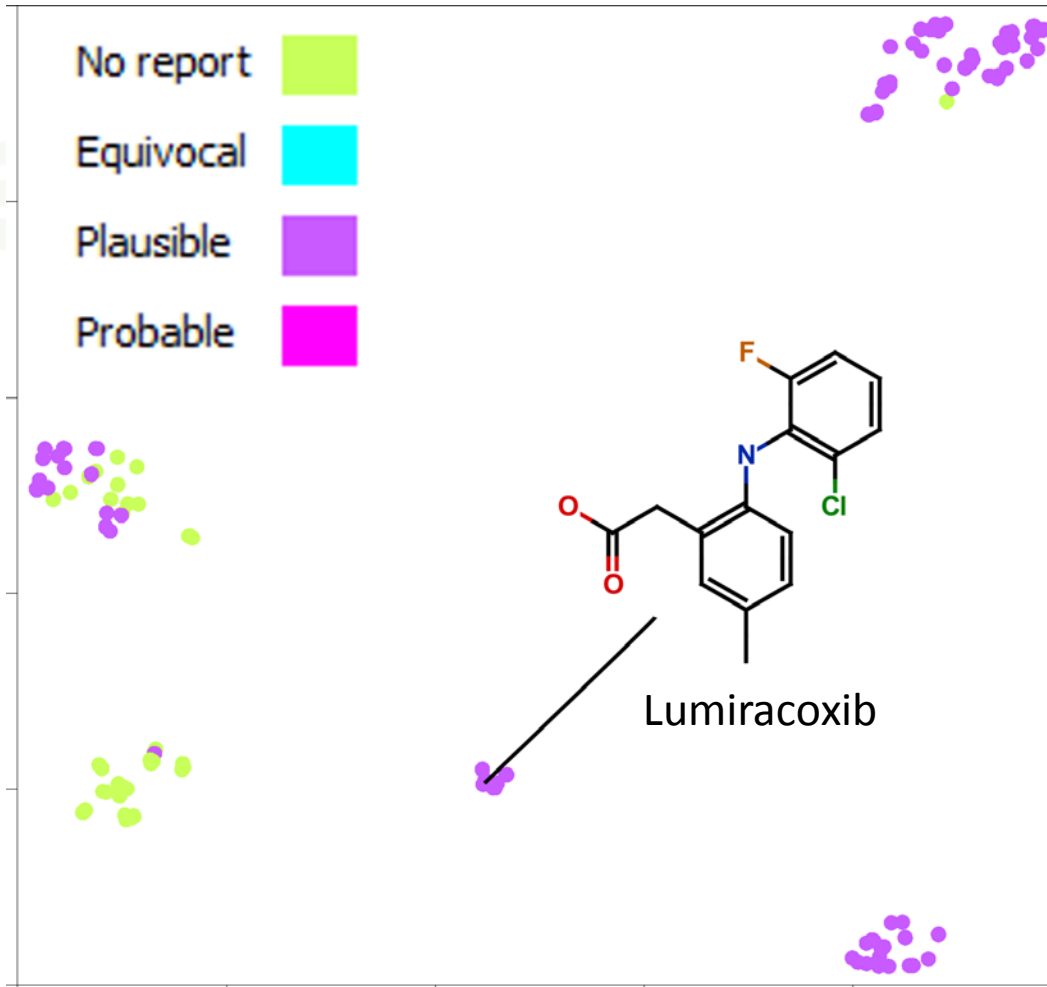
COX2 library

Scored excluding toxicity endpoints



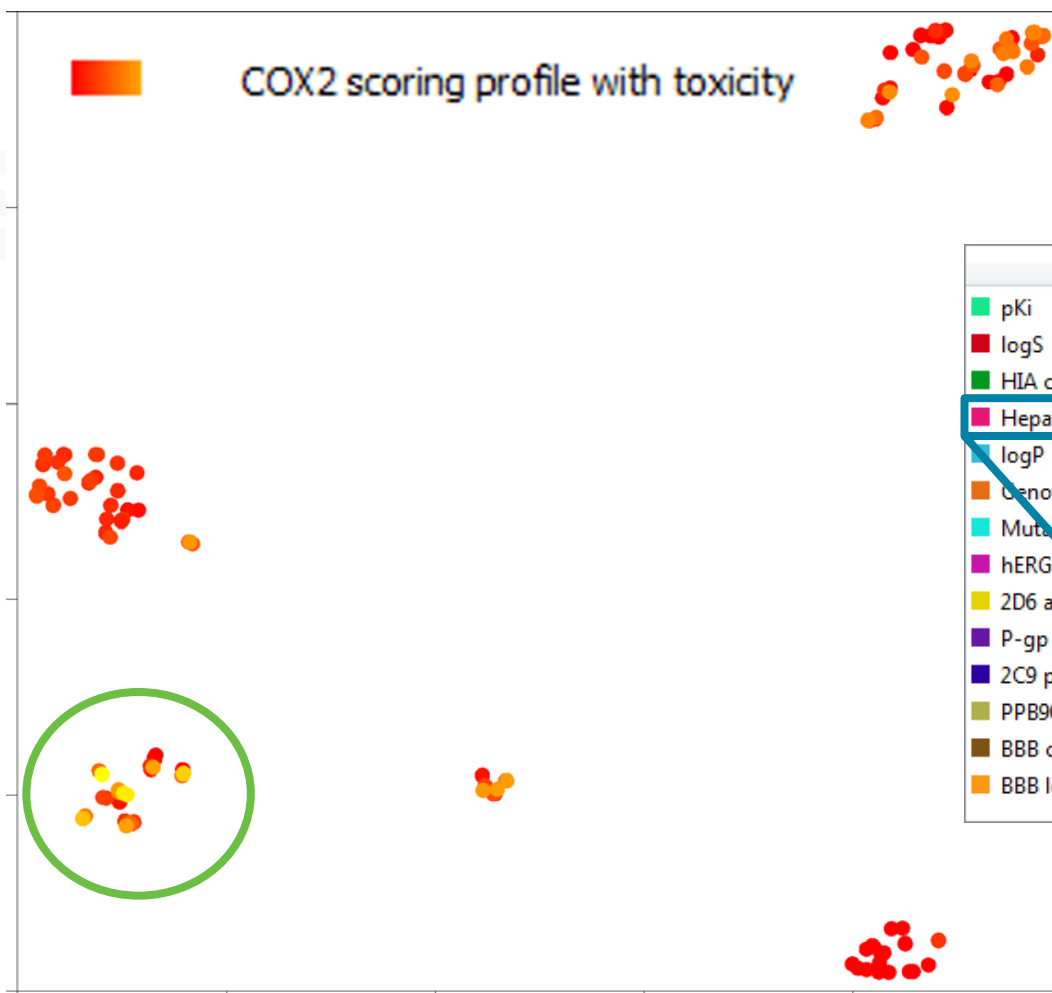
COX2 library

Predicted hepatotoxicity

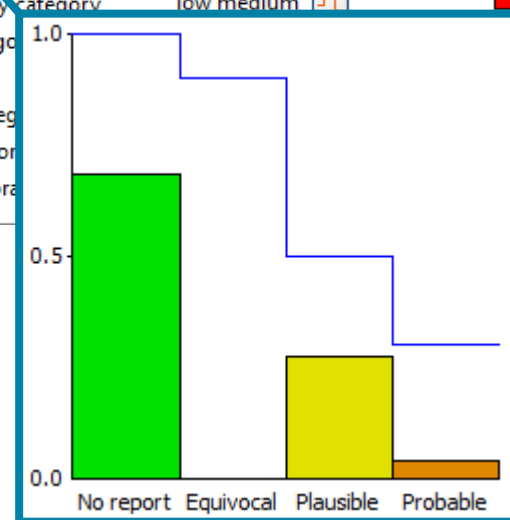


COX2 library

Scored including toxicity endpoints

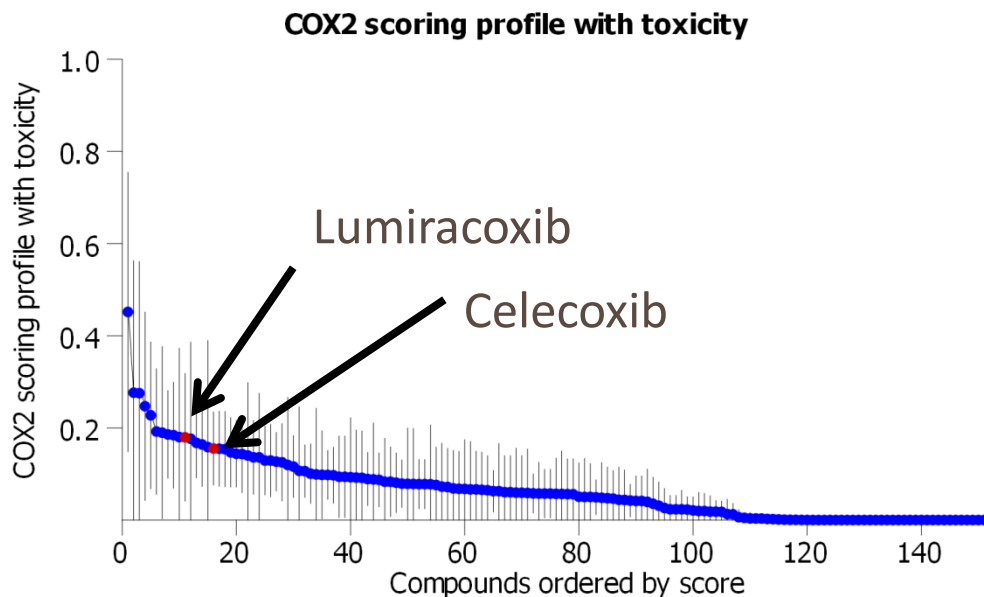


Profile	Desired Value	Importance
pKi	8 -> inf	High
logS	> 1	High
HIA category	+	High
Hepatotoxicity	No report	High
logP	0 -> 5	High
Cytotoxicity in vitro	No report	High
Mutagenicity in vitro	No report	High
hERG pIC50	≤ 5	High
2D6 affinity category	low medium	High
P-gp category		Medium
2C9 pKi		Medium
PPB90 category		Medium
BBB category		Medium
BBB log([bra		Medium



But wait...

Wouldn't we miss Celecoxib?!

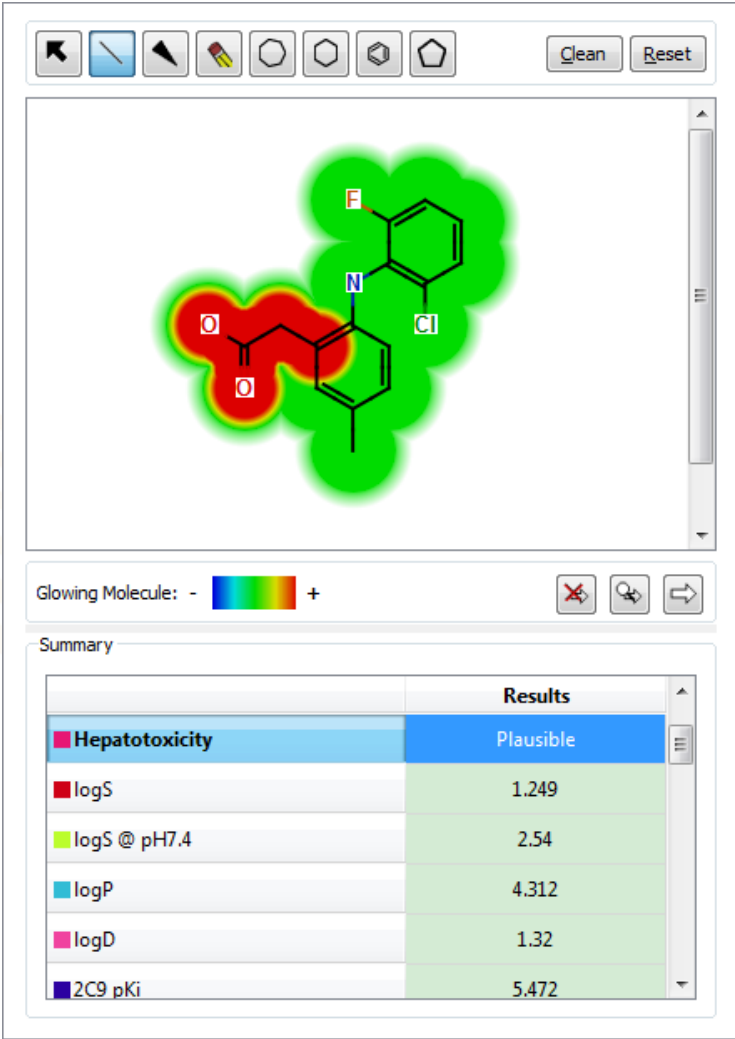


- Celecoxib and Lumiracoxib would not be rejected outright
 - Highlighted hazard, confirm experimentally and consider context
- Celecoxib does exhibit signs of hepatotoxicity, but is 'saved' by its low dose and high therapeutic index*

Consider Redesign Strategies

Lumiracoxib

- Glowing molecule highlights 2-Arylacetic acid alert
- Interactively explore strategies for reducing risk
 - Monitor changes in multiple properties simultaneously



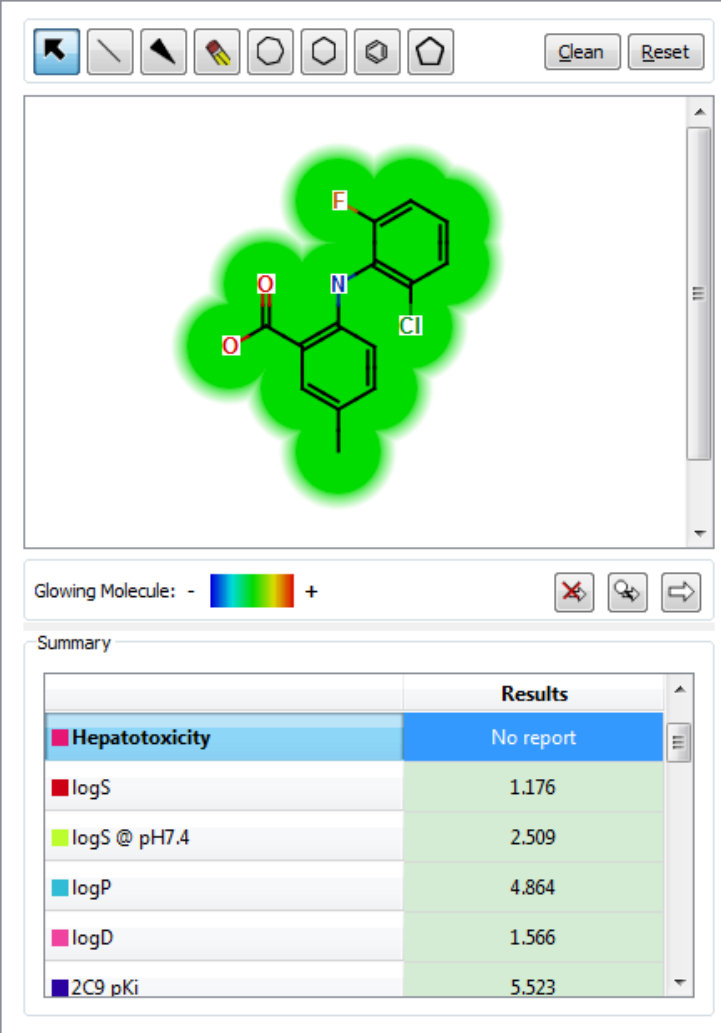
The screenshot displays a molecular design software interface. At the top, there is a toolbar with various icons for navigation and editing, along with 'Clean' and 'Reset' buttons. The central area shows a 3D molecular model of Lumiracoxib, which is highlighted with a glowing effect. The molecule is colored in a gradient from red to green, indicating different levels of risk or alert. Below the molecule, there is a 'Glowing Molecule' control with a color bar and a '+' sign. To the right of the color bar are three icons: a red 'X', a magnifying glass, and a right-pointing arrow. Below this is a 'Summary' section containing a table with the following data:

	Results
■ Hepatotoxicity	Plausible
■ logS	1.249
■ logS @ pH7.4	2.54
■ logP	4.312
■ logD	1.32
■ 2C9 pKi	5.472

Consider Redesign Strategies

Lumiracoxib

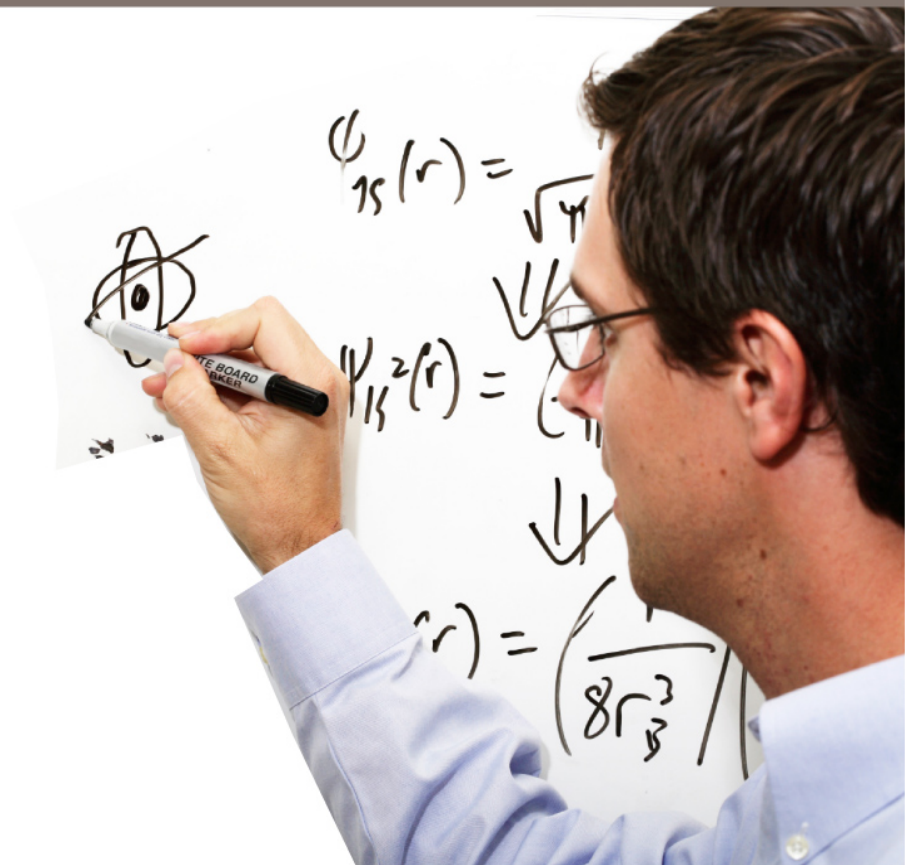
- Glowing molecule highlights 2-Arylacetic acid alert
- Interactively explore strategies for reducing risk
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Summary

	Results
■ Hepatotoxicity	No report
■ logS	1.176
■ logS @ pH7.4	2.509
■ logP	4.864
■ logD	1.566
■ 2C9 pKi	5.523

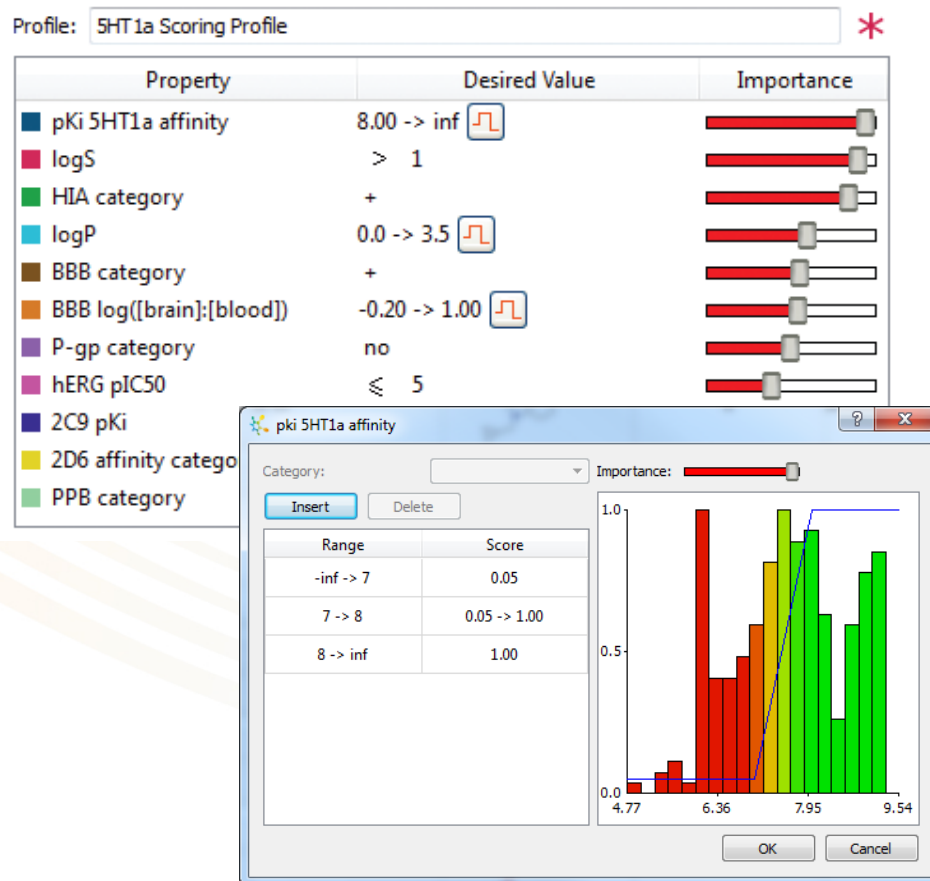
Introduction to StarDrop and the Derek Nexus module



StarDrop Helps to Guide Decisions

From selection... to design

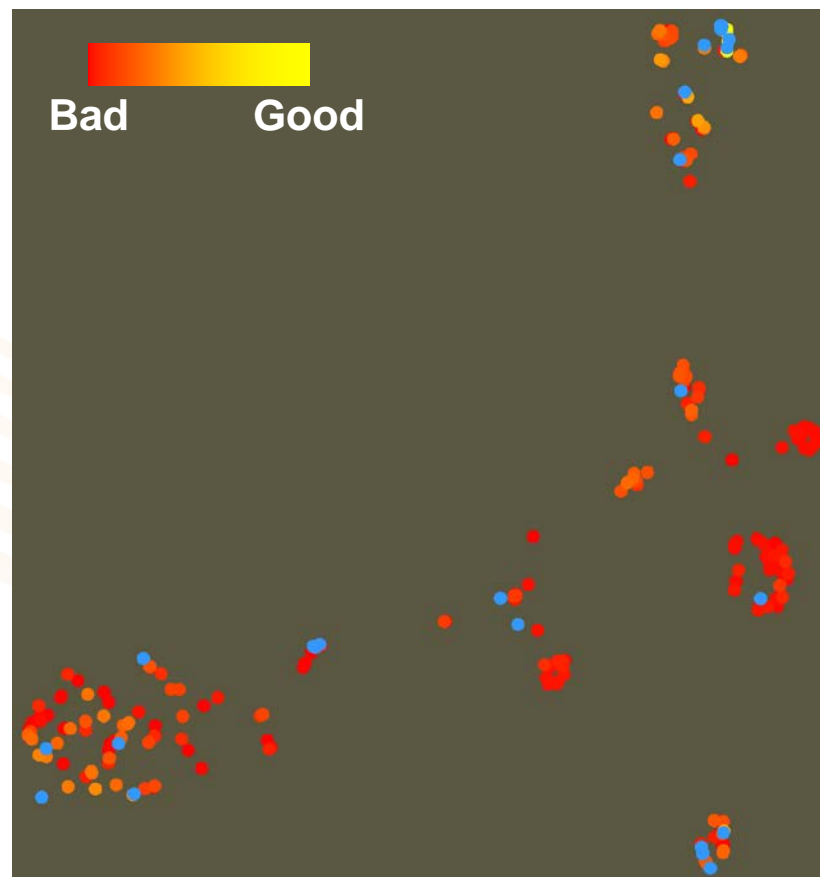
- Probabilistic Scoring*
 - User-defined profile and weights
 - Use data from any source
 - Allow for uncertainty
 - Score for likelihood of success
- Chemical Space and Selection
 - View property distributions across chemical diversity
 - Balance quality and diversity
- Glowing Molecule
 - Interactively explore new ideas
 - Link compounds structure with properties
- Interactive Visualisation
 - R-group analysis
 - Matched Molecular Pair analysis



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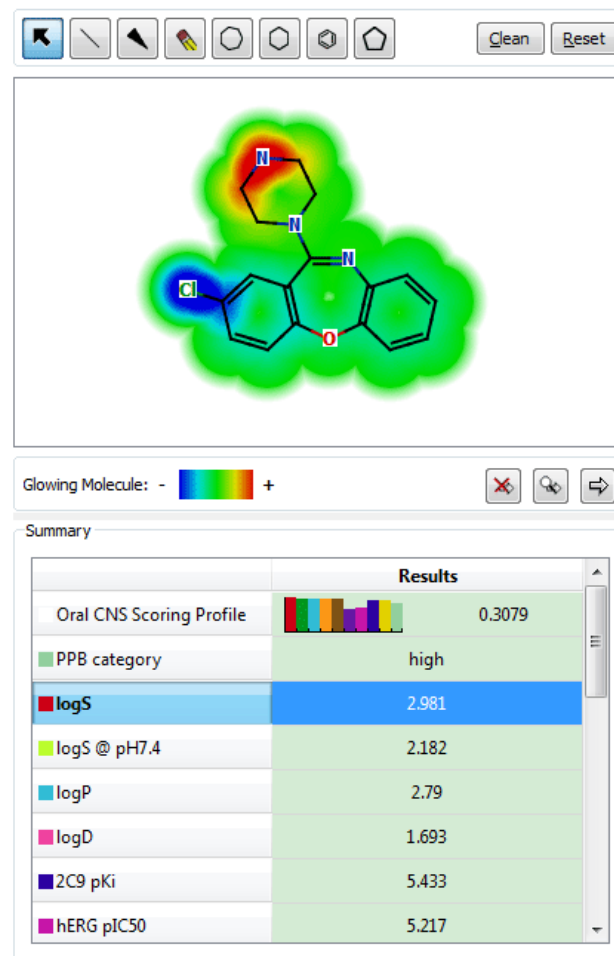
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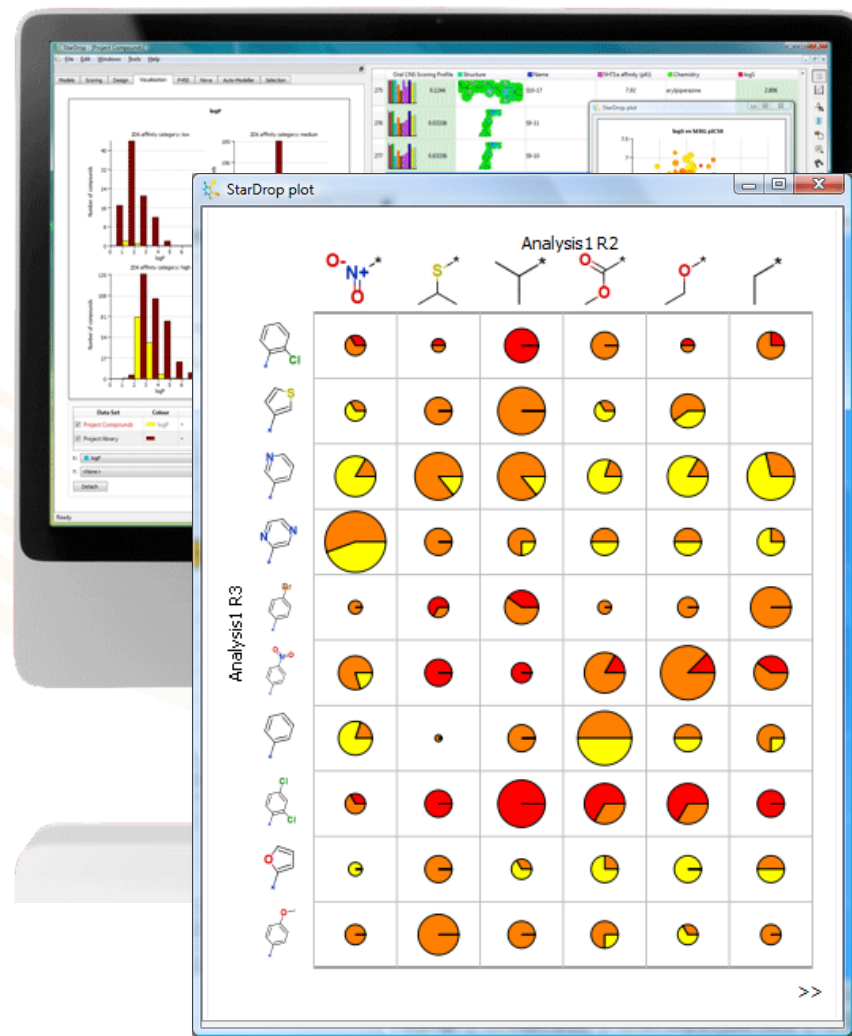
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StarDrop Helps to Guide Decisions

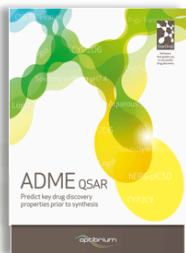
From selection... to design

- Probabilistic Scoring
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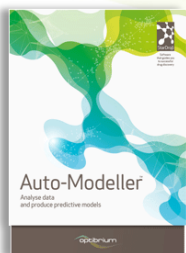
StarDrop Plug-in Modules and Integration

Extend Core Capabilities



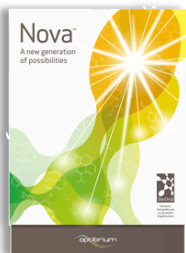
ADME QSAR

High quality predictive models of key ADME properties



Auto-Modeller

Build and validate robust models tailored to your chemistry



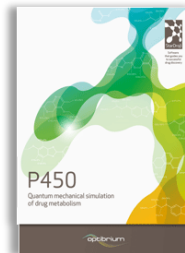
Nova

Generate and prioritise new, relevant compound ideas



BIOSTER™

Explore >20k precedented transformations with the Nova module



P450

QM simulations identify sites of metabolism and lability for major P450s



torch3D™

Understand and apply 3D SAR to identify and optimise novel actives



Derek Nexus™

Knowledge-based prediction of >40 toxicity endpoints



MPO Explorer™

Develop multi-parameter optimisation strategies

Derek Nexus™ Module for StarDrop

Differences with full Derek Nexus

- Derek Nexus for StarDrop provides unique features for medicinal chemists and drug discovery projects, e.g.
 - Visualisation to explore toxicity risk of different chemistries
 - Probabilistic scoring to balance toxicity risk against other factors
 - Interactive design with Glowing Molecule to guide redesign and reduce risk of toxicity
- The full Derek Nexus platform from Lhasa Limited provides access to full Derek knowledge base for expert toxicologists
 - Information on mechanism of action, biological data and references
 - Detailed annotation of structural alerts
 - Helps to design toxicology experiments
- Reporting feature in StarDrop helps collaboration between drug discovery projects and preclinical toxicology



Conclusion

- Addressing toxicity early in the drug discovery process is key to improving success rate and productivity
- Knowledge based predictions provide a reliable way to identify toxicity hazards (potential risk)
- Results need to be used in context of other requirements of a successful drug
- Need to take confidence into account
 - Avoid rejecting good compounds due to uncertain data
- Reference:
 - Segall and Barber, Drug Discov. Today **19** (2014), pp. 688-693
 - Download (p)reprint from www.optibrium.com/community/publications

