



A knowledge-based system for the prediction of forced degradation of organic compounds

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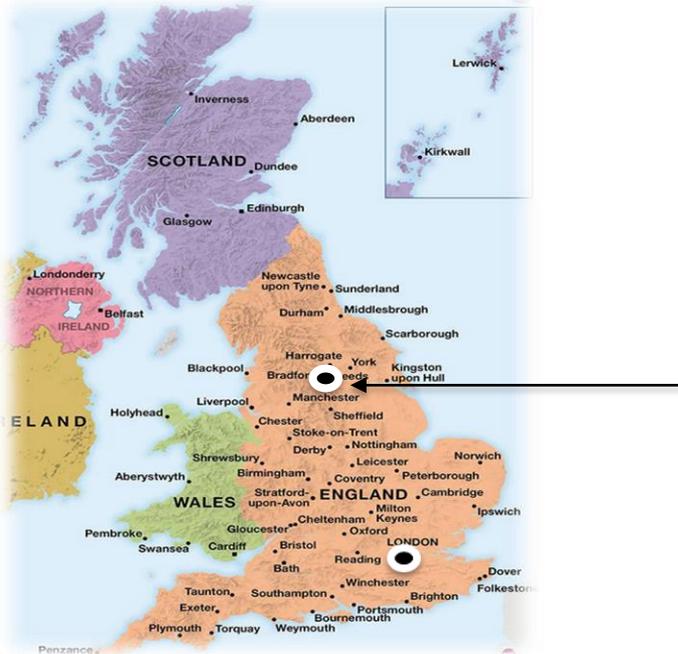


Contents

- Introduction to Lhasa Limited
- Regulatory requirements for forced degradation
- Degradation knowledge space
- Overview of Zeneth – what it is / how it works
- Future developments
- Summary

Introduction

- Established in 1983
- Located in central Leeds, UK
- Scientific software company driven by our [members](#)
- Produces [predictive](#) and [database systems](#) for the life sciences industry
- Not-for-profit organisation & educational charity



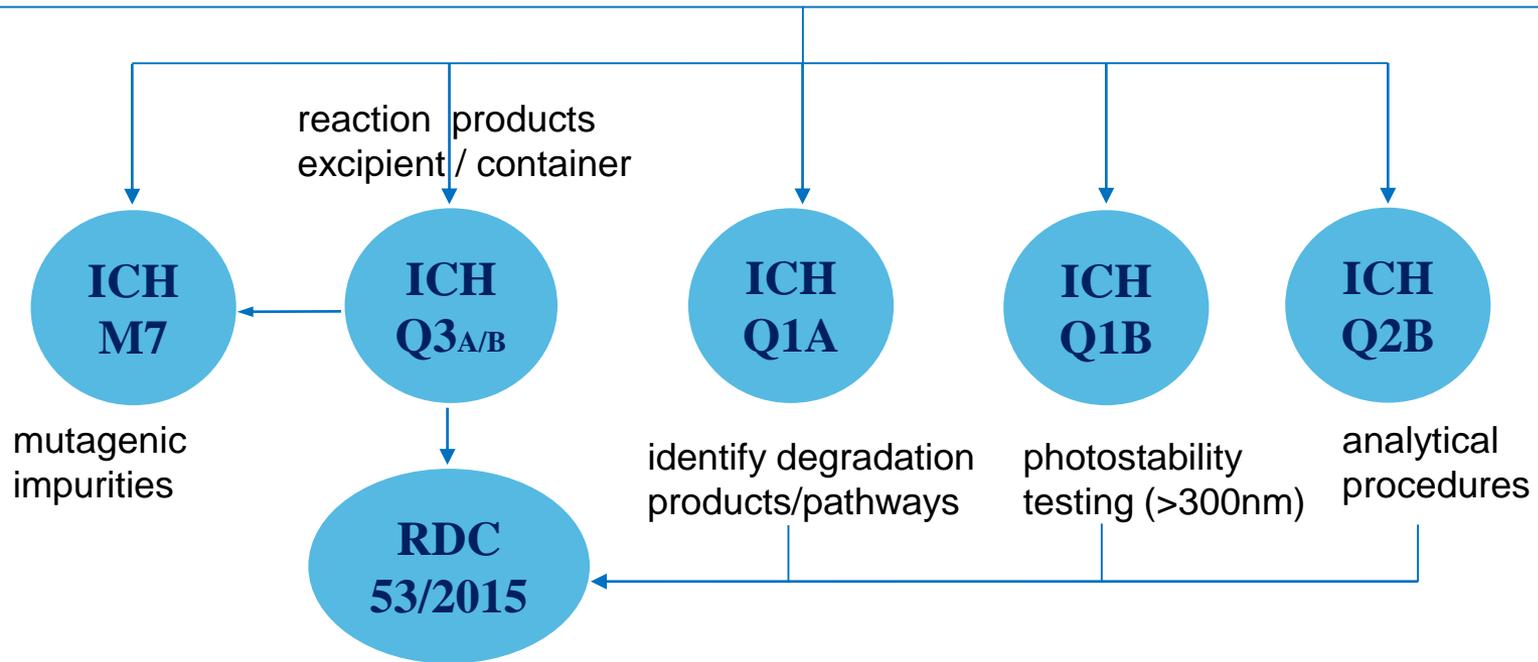
Membership worldwide



Pharmaceutical, Education, Regulatory bodies

Regulatory requirements

Degradation Product: A molecule resulting from a chemical change in the drug molecule brought about over time and/or by the action of light, temperature, pH, water, or by reaction with an excipient and/or the immediate container/closure system.



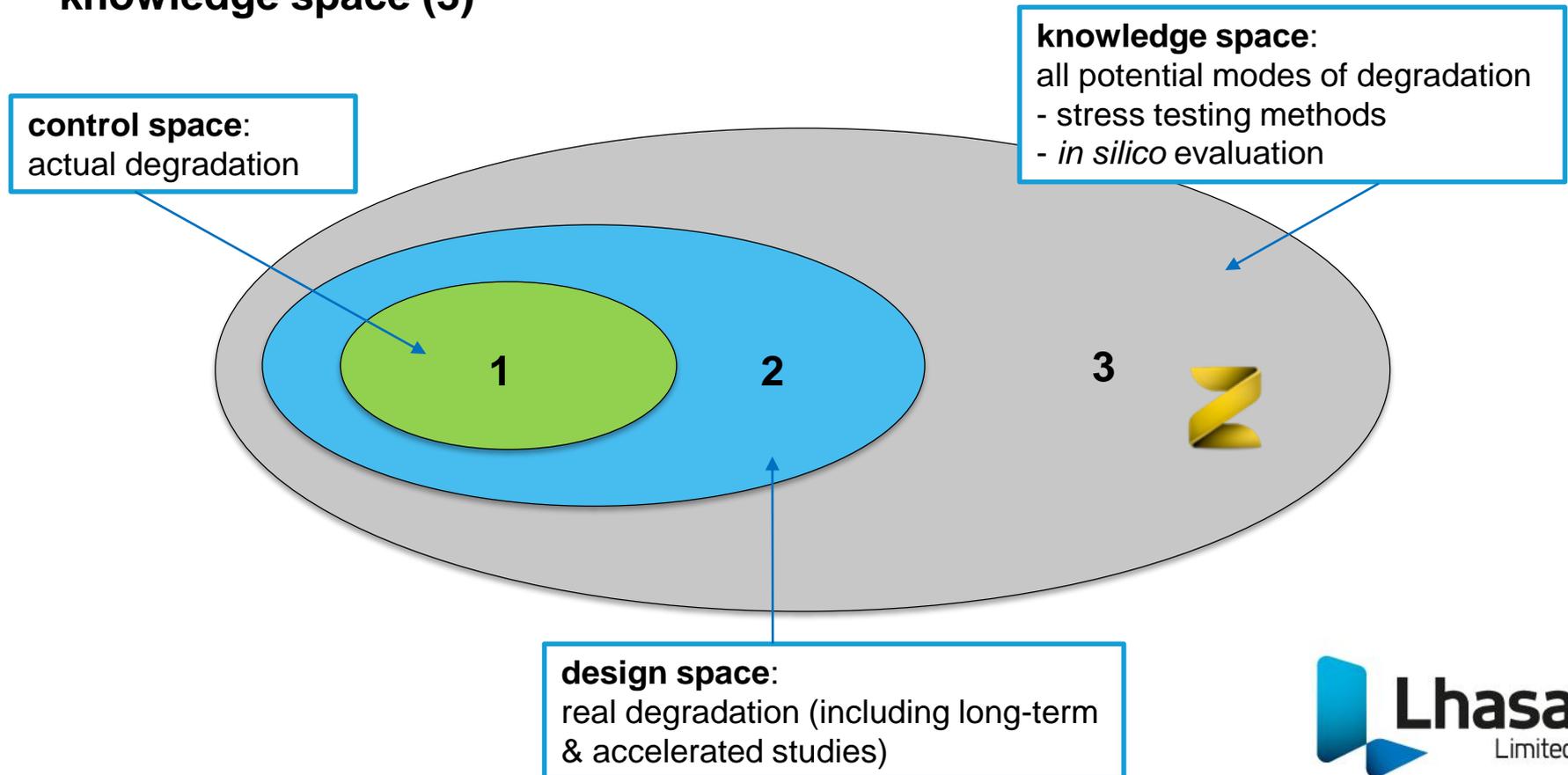
Forced degradation studies are performed to:

- Identify all potential degradation products & pathways
- Assess the intrinsic stability of the drug substance (API) & product
- Validate the stability-indicating methods/procedures

Degradation knowledge space

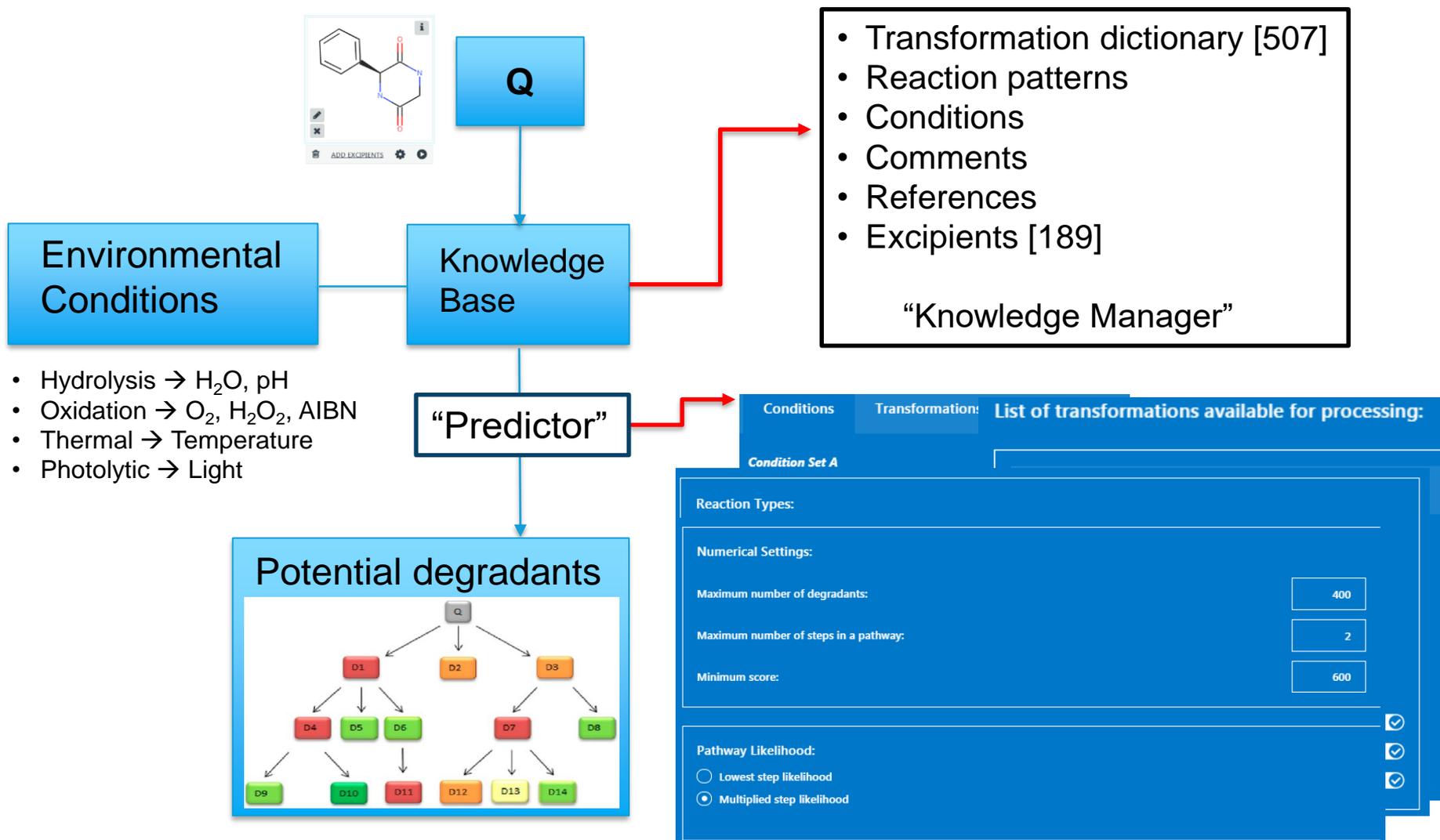
Aim of forced degradation studies on the drug product/API:

- Ensure all relevant degradation products are “known” by applying stress conditions
- Increase the confidence of the **control space (1)** by having a well designed **knowledge space (3)**

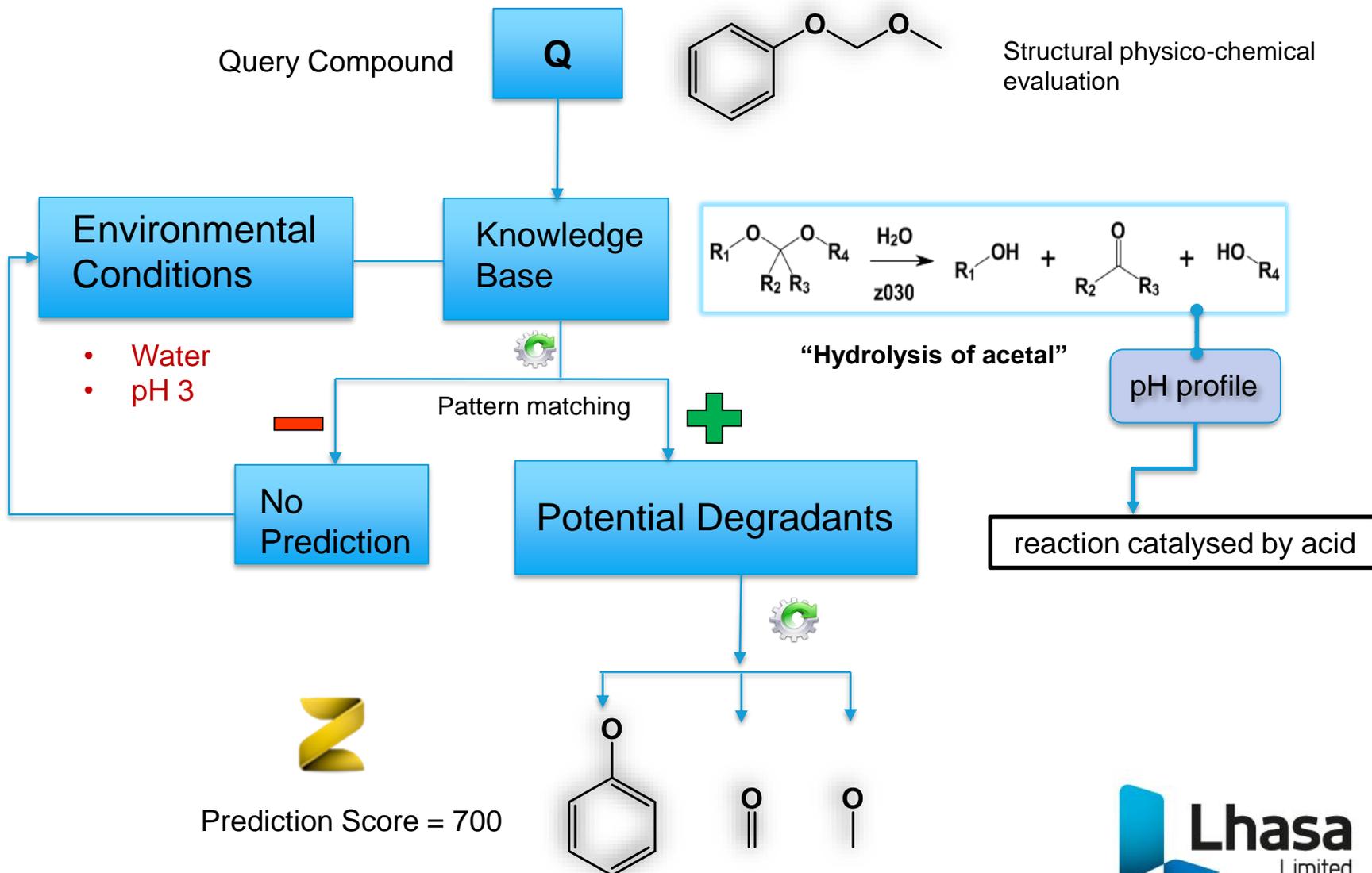


What is Zeneth?

.....Expert **knowledge-based** system for the **prediction** of forced degradation

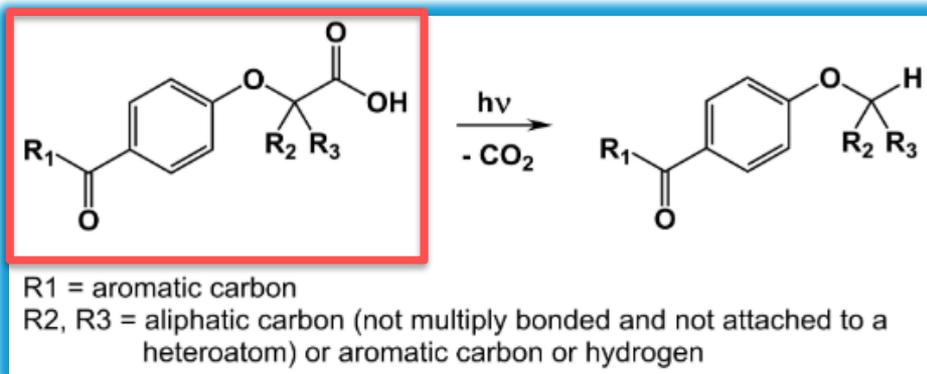


What is Zeneth?



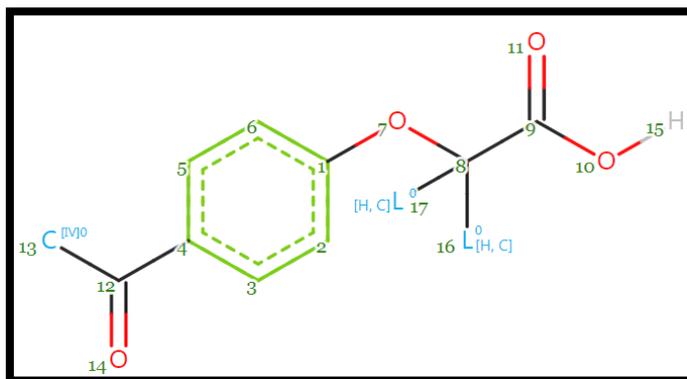
Chemical patterns

z457:
Photodecarboxylation
of aryloxyacetic acid



Explora language

1



Atom 13: `requiredBondType(AROMATIC)`

Atoms 16, 17: `(heteroCount == 0 and allowedBondType(SINGLE))`
or `requiredBondType(AROMATIC)`

Script
[LTL]

2

```
breakBond(10, 15);  
hideStructure(15);  
breakBond(8, 9);  
setBondType(9, 10, DOUBLE);  
hideStructure(9);  
setCharge(8, -1);  
captureIntermediate("");  
setIntermediateTypeMechanistic();  
setIntermediateAsPresumed();  
unsetCharge(8);  
  
setScore(0.7);
```

Prediction



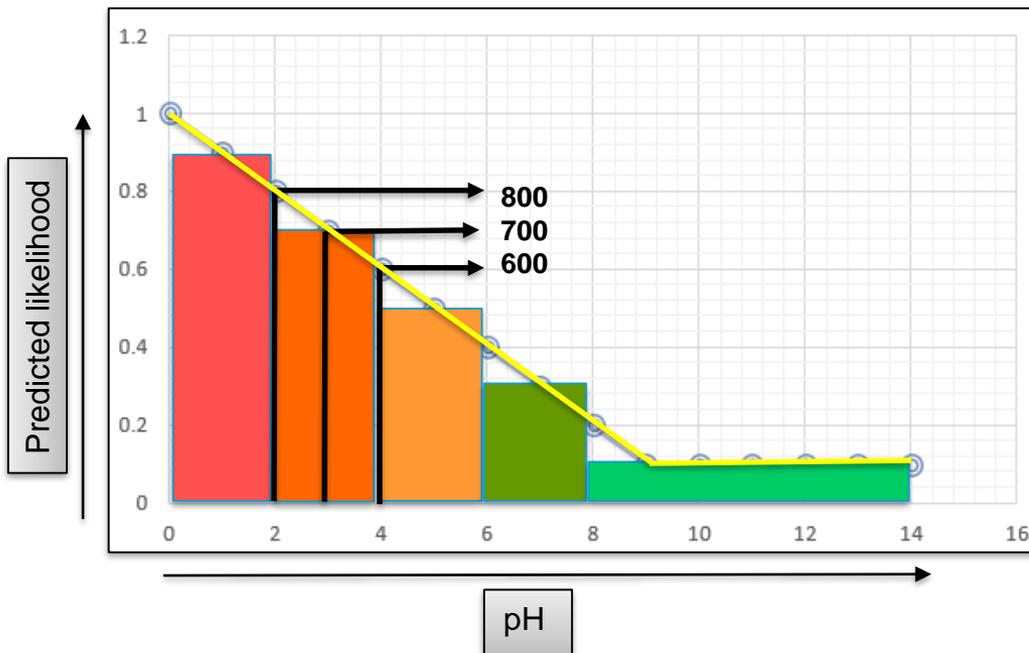
Degradant likelihood/scoring

Most transformations have a pH profile:

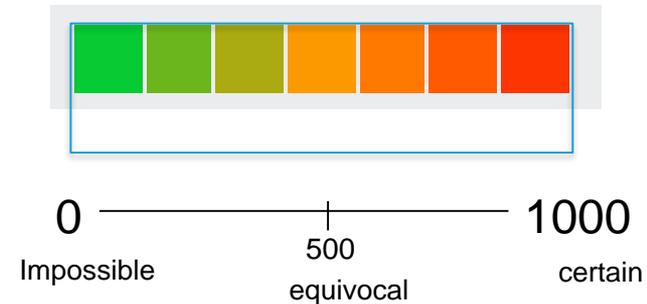
- Rule 34: If [transformation 030] is [certain] then [transformation 030 products] is [hydrolysis with pH profile 5]
- Rule 121: If [conditions: water] is [certain] then [hydrolysis with pH profile 5] is [likelihood with pH profile 5]
- Rule 80: If [pH >= 8] is [certain] then [likelihood with pH profile 5] is [very unlikely]
- Rule 81: If [6 <= pH < 8] is [certain] then [likelihood with pH profile 5] is [unlikely]
- Rule 82: If [4 <= pH < 6] is [certain] then [likelihood with pH profile 5] is [equivocal]
- **Rule 83: If [2 <= pH < 4] is [certain] then [likelihood with pH profile 5] is [likely]**
- Rule 84: If [pH < 2] is [certain] then [likelihood with pH profile 5] is [very likely]

pH 2 = likely
pH 3 = likely
pH 3.9 = likely

Line Formula = $\text{MIN}(\text{MAX}(-\text{pH}/10+1, 0.1)) \times 1000 \rightarrow \text{score}$

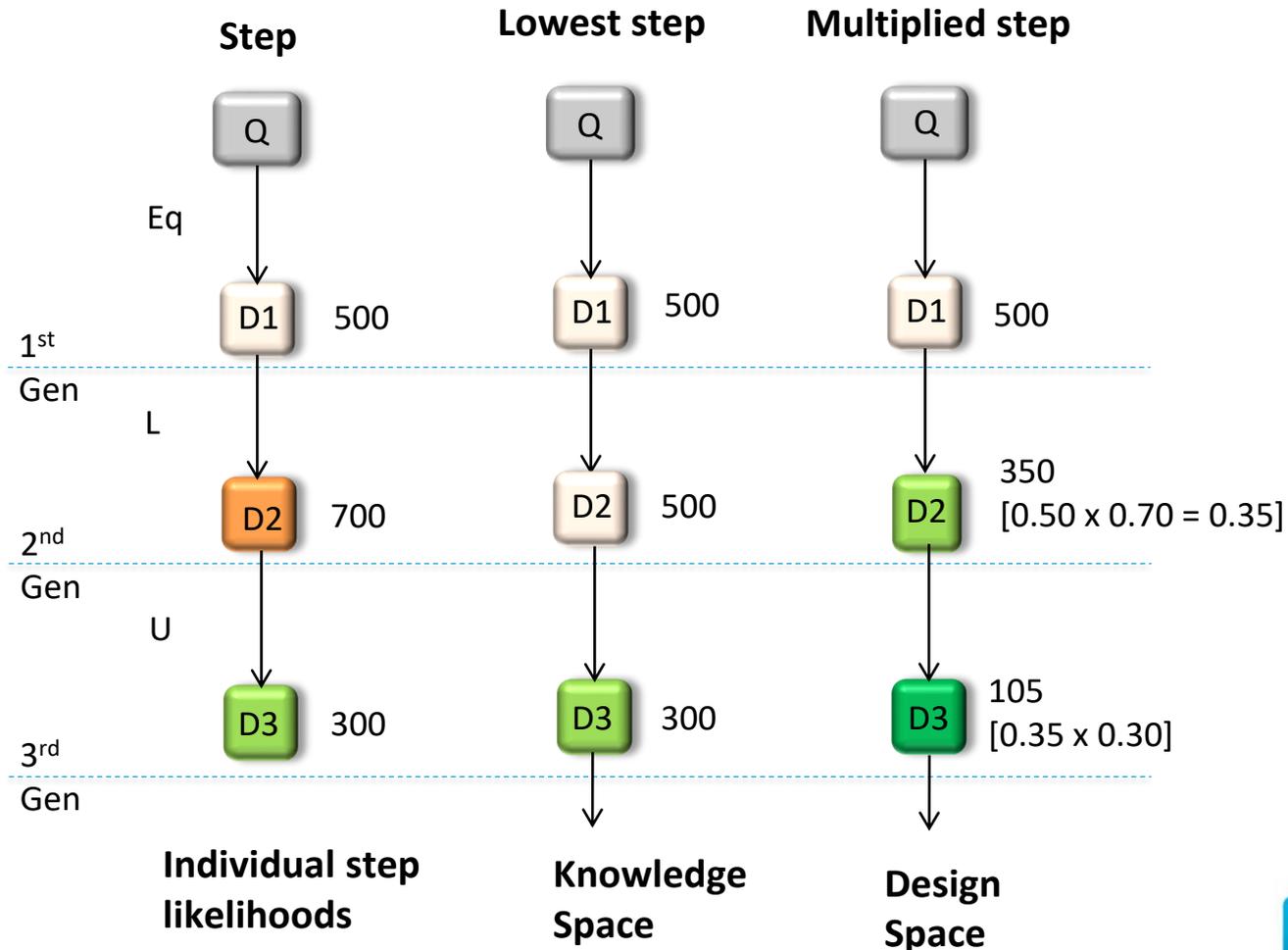


Score:

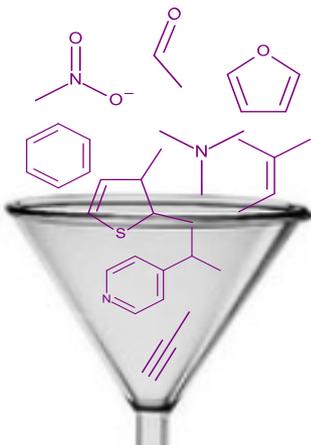


Scoring: multiplied step likelihood

Child degradant cannot have a higher score than the parent



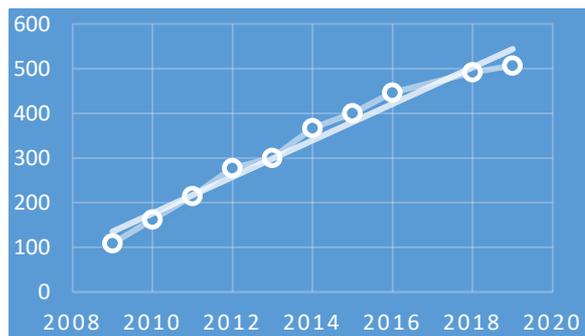
Building the knowledge base



Knowledge base

Oxidation [148]
Hydrolysis [118]
Condensation [109]
Elimination [60]
Isomerisation [28]
Photolysis [44]

- External requests (members)
 - provide data for a reaction
 - proprietary data via data sharing initiatives
- Group discussion with members
 - discuss proposals for new reactions
 - provide feedback on new & existing chemistry
- Internal requests, updates to existing knowledge
 - general maintenance
 - investigation of primary literature
 - specialised degradation chemistry books
- Strategic decision (e.g. cover a new reaction type)

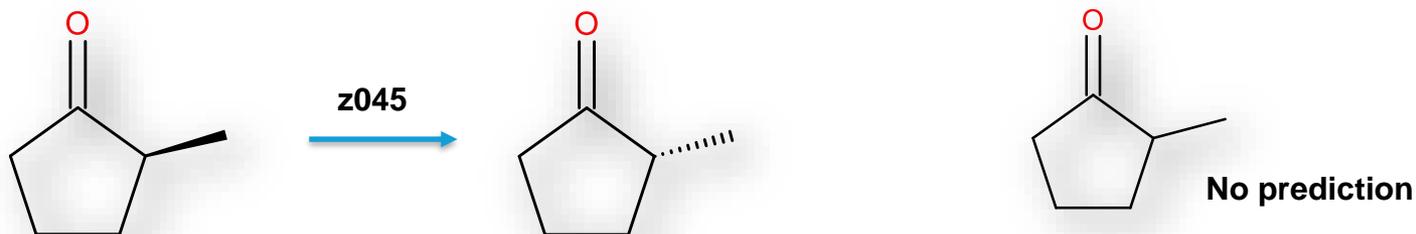


New knowledge added & existing knowledge continuously maintained

Stereochemistry

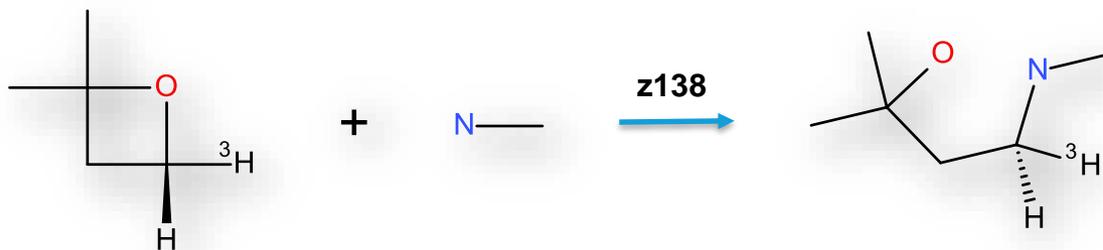
- Epimerisations

This transformation operates only on carbon centres that have a defined stereocentre:



- Nucleophilic substitutions

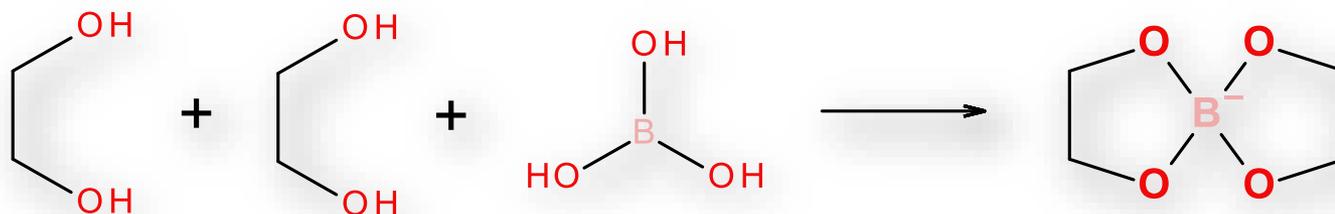
S_N2 -type reactions exhibit inversion of configuration at a stereocentre:



Three-component reactions

The program has the capacity to express three-component reactions:

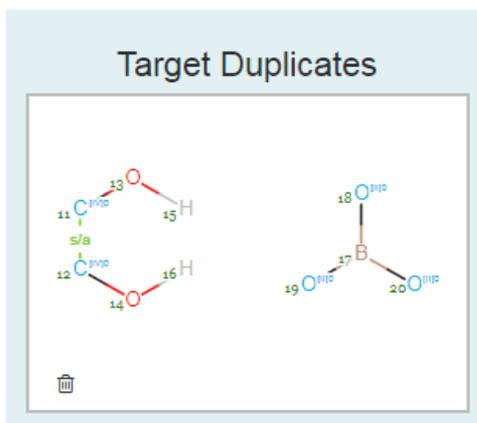
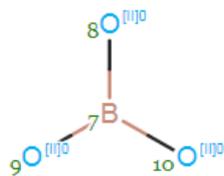
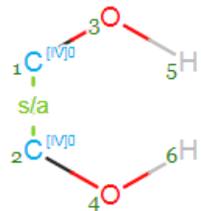
New transformation z505



2 oxygen nucleophiles

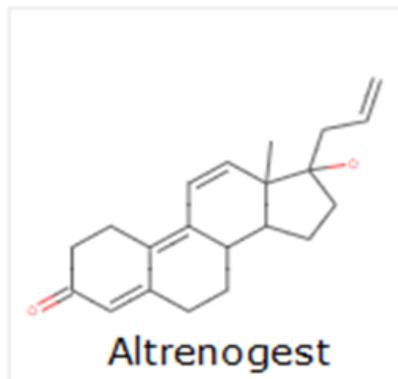
boric acid

spiroborate anion



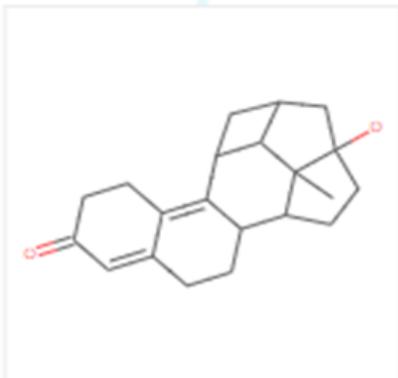
```
1 breakBond(3, 5);
2 hideStructure(5);
3 breakBond(7, 8);
4 hideStructure(8);
5 makeBond(3, 7, SINGLE);
6 captureIntermediate("");
7 setIntermediateTypeObservable();
8 breakBond(4, 6);
9 hideStructure(6);
10 breakBond(7, 9);
11 hideStructure(9);
12 makeBond(4, 7, SINGLE);
13 captureIntermediate("");
14 setIntermediateTypeObservable();
15 breakBond(13, 15);
16 hideStructure(15);
```

Intramolecular cycloadditions



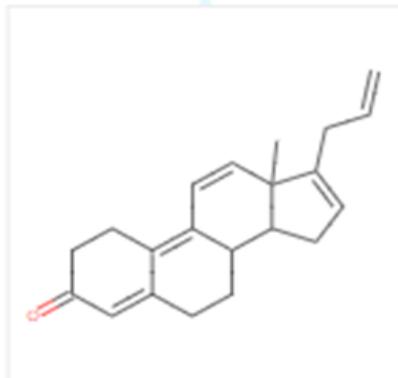
267 - Intramolecular photochemical cycloaddition of two alkenes

700

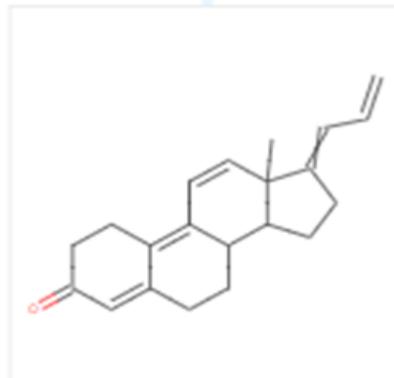


064 - Dehydration of alcohol

300

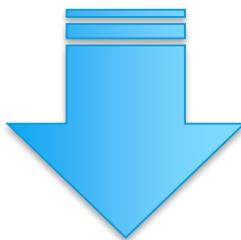


300



Chemical descriptors

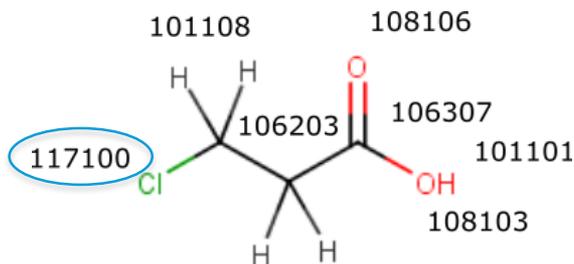
- pK_a
- Bond Dissociation Energy (BDE)
- Electrophilicity/Nucleophilicity
- Photochemical Reactivity
- Aromaticity
- Steric Hindrance



Reasoning Engine

pK_a Calculator

- pK_a of a molecule determines its protonation state at a given pH
- pK_a tends to be governed by electronic effects that decrease with the distance from the centre of protonation
- Generate a distance spectrum for each atom type from the pK_a centre

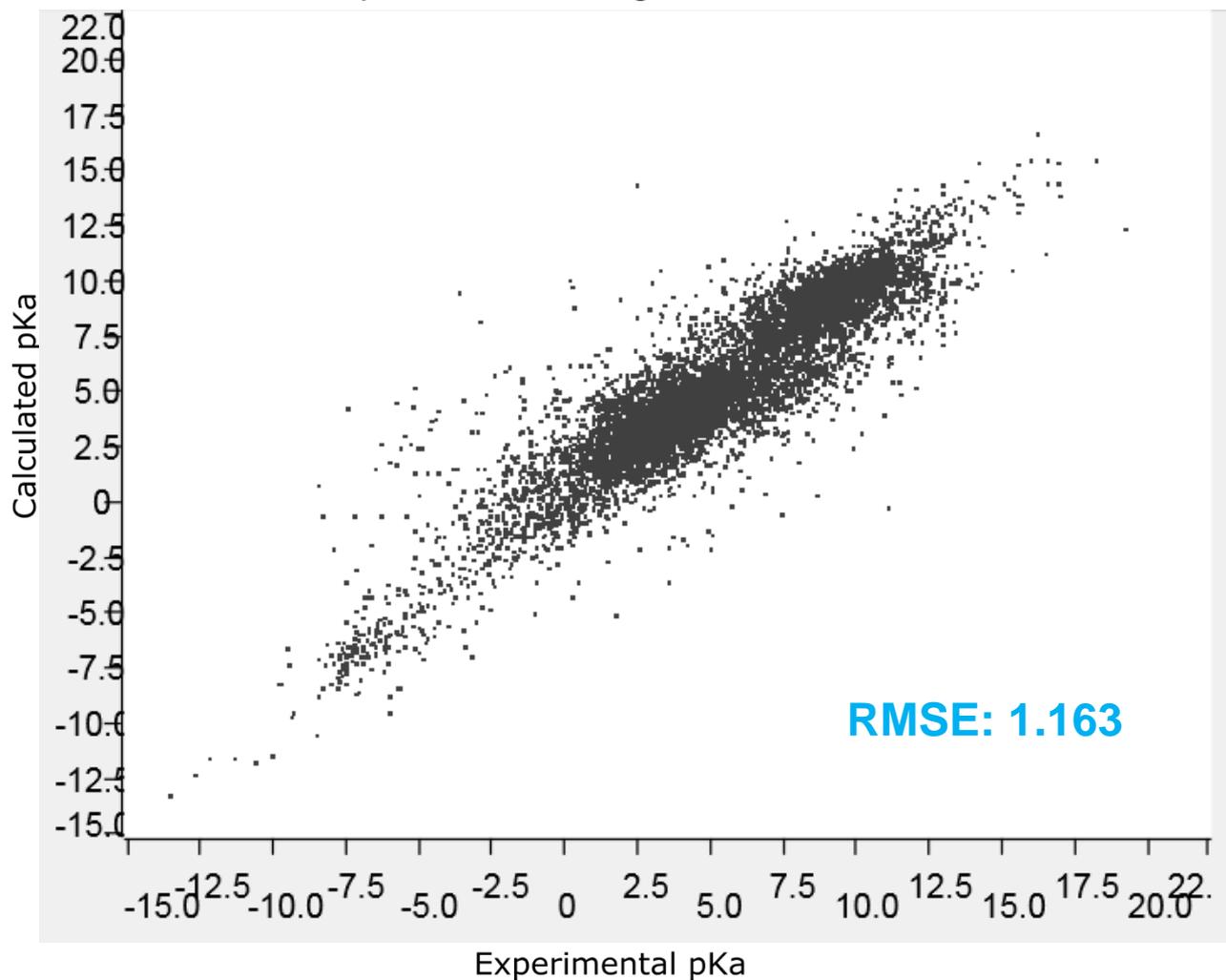


Dist	101101	101108	106307	106203	108103	108106	117100
1	1		1				
2				0.25		0.25	
3		0.22		0.11			
4		0.12					0.06
Sum	1	0.34	1	0.36	0	0.25	0.06

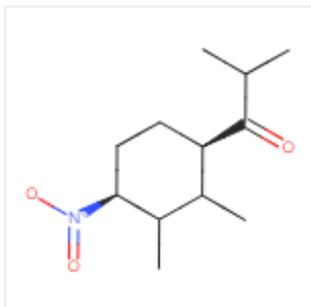
= distance spectrum

pK_a Performance

Preliminary results using 5-fold cross-validation



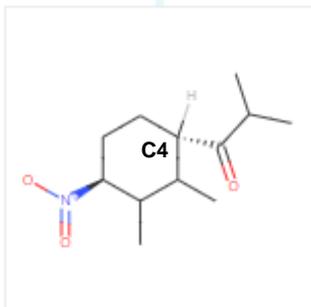
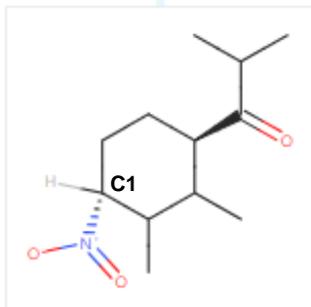
pK_a Calculator implemented



045 - Epimerisation alpha to electron-withdrawing group

900

777



pH = 12

z045:

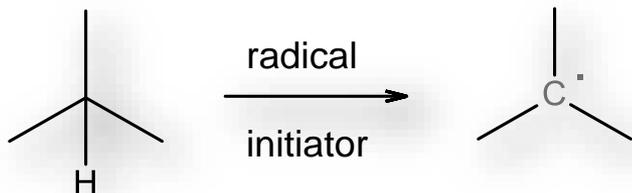
Epimerisation alpha to electron-withdrawing group

```
if (!isDefinedStereocentre(1)) { abort(); }  
if (isBridgehead(1)) { abort(); }  
  
substitutionWithInversion(4, 1, 3);  
hideStructure(3);  
  
real pH = getpH();  
real pKa = getpKa(1);  
real raw_score = (max(pH, 14 - pH) + 6 - pKa) / 10;  
if (raw_score < 0.1) { abort(); }  
  
setScore(min(raw_score, 0.9));
```

Epimerisation is more likely at C1 than C4

Other descriptors: BDE

General pathway of a transformation initiated by hydrogen abstraction:



- The radical initiator considered in forced degradation is peroxy ($R-O-O\bullet$)
- Peroxy is quite selective and attacks mainly “labile” C-H bonds e.g. allylic, benzylic, α to a heteroatom etc.
- The incorporation of a BDE descriptor would aid the assessment of likelihood of these reactions

BUT ...

To implement it within the Zeneth reasoning framework, both speed and performance is critical

Summary

- Zeneth is an expert knowledge-based system for the prediction of forced degradation
- Helps understand mechanistic pathways as well as aid degradant identification and structure elucidation
- Zeneth generates predictions from its knowledge base which currently contains 507 transformations
- Our members drive the development of the Zeneth program and directly influence the scientific content
- Zeneth has recently undergone a complete re-design with the result of massively improved usability as well as improvements to the quality and speed of predictions.

Acknowledgements

Zeneth Team, Lhasa Limited

Scientists: Martin Ott, Rachel Hemingway and Tony Long

Development Team:

Andrzej Paetz, Alan Hayward, Christos Wild, Daragh Connolly, Diane Bureau, Lucy Wilcock, Mohit Garg, Nandini-Arjun Mandya, Paul Cunningham, Pawel Maslej, Piotr Gorny, Roderic Green, Tony Clay and Jeff Plante

Account Manager: Maggie Coombs

Business Analysts: Lisa Leivers, Faye Stringer

Meet the team at Booth 4!



Multiplied step likelihood

Likelihood values:						
VL/L/Eq/U/VU:		0.900	0.700	0.500	0.300	0.100
Bin boundaries:		0.999	0.800	0.600	0.400	0.200

Two-step pathways:

*	bin		VL	L	Eq	U	VU	
VL	0.900	0.800	1.000	0.810	0.630	0.450	0.270	0.090
L	0.700	0.600	0.800	0.630	0.490	0.350	0.210	0.070
Eq	0.500	0.400	0.600	0.450	0.350	0.250	0.150	0.050
U	0.300	0.200	0.400	0.270	0.210	0.150	0.090	0.030
VU	0.100	0.000	0.200	0.090	0.070	0.050	0.030	0.010

Three-step pathways:

VL	0.900	bin		VL	L	Eq	U	VU
VL	0.900	0.800	1.000	0.729	0.567	0.405	0.243	0.081
L	0.700	0.600	0.800	0.567	0.441	0.315	0.189	0.063
Eq	0.500	0.400	0.600	0.405	0.315	0.225	0.135	0.045
U	0.300	0.200	0.400	0.243	0.189	0.135	0.081	0.027
VU	0.100	0.000	0.200	0.081	0.063	0.045	0.027	0.009

L	0.700	bin		VL	L	Eq	U	VU
VL	0.900	0.800	1.000	0.567	0.441	0.315	0.189	0.063
L	0.700	0.600	0.800	0.441	0.343	0.245	0.147	0.049
Eq	0.500	0.400	0.600	0.315	0.245	0.175	0.105	0.035
U	0.300	0.200	0.400	0.189	0.147	0.105	0.063	0.021
VU	0.100	0.000	0.200	0.063	0.049	0.035	0.021	0.007