



Applicability Domain

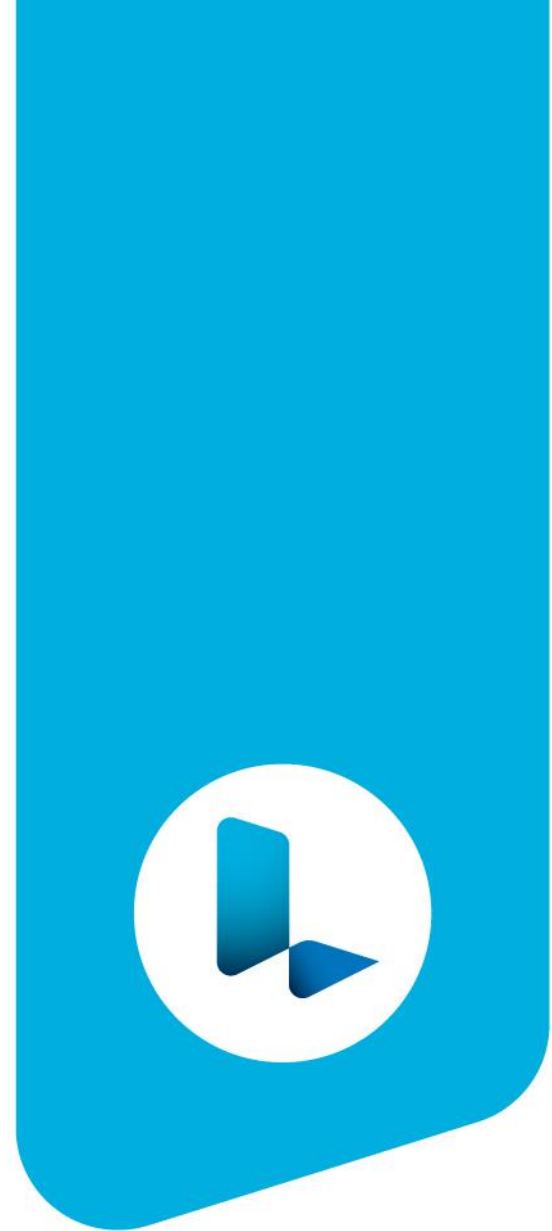
Towards a more formal definition

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Research Leader





shared **knowledge** • shared **progress**

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Meteor nexus

- Extensive Metabolism Knowledge**
Lhasa expert scientists implement Meteor biotransformations utilising public and proprietary data, ensuring vast coverage of biotransformations.
- An expert, knowledge-based system for the prediction of metabolic fate
- Reduces Risk in R&D**
Save time and money by identifying reactive, potentially adduct-forming intermediates and quickly assessing their toxicity through Derek, Sarah or Vitic Nexus.
- Accurate Metabolite Ranking**
The methodologies in Meteor rank predicted metabolites by taking into account experimental data and the chemical environment of the site of metabolism.
- Using structure activity relationships and a dictionary of biotransformations, Meteor provides transparent Phase 1 and 2 metabolism predictions.

Mirabilis TM

- Greater Efficiency**
Eliminates the need to set up costly and time consuming analytical methods to measure impurities that are unlikely to be present in the final drug product.
- Industry-standardised software tool for the calculation of purge factors of potentially mutagenic impurities in a synthetic route
- Expert ICH M7 Support**
The ICH M7 Guideline specifically allows for a control strategy that relies on an understanding of process controls in lieu of analytical testing.
- Transparent Predictions**
Provides expert commentary and detailed supporting information for calculated purge factors, enabling improved and justifiable decision making.
- Aiding Submission to Regulators**
Mirabilis provides a report which includes the purge calculation, scientific rationale and supporting evidence to aid in the submission to regulators.
- A semi-automated approach, built on expert knowledge, Mirabilis improves the efficiency of purge analysis.

Vitic nexus

- Expert ICH M7 Support**
Rapidly find relevant supporting examples for your impurities by structure, substructure or similarity searching.
- A toxicity database and information management system
- The Vitic database contains expert curated, high-quality and peer reviewed toxicity data from both published and unpublished sources.
- Current Toxicological Data**
Regular updates by Lhasa's dedicated expert data team ensures access to the latest toxicology data.
- Meeting Regulatory Requirements**
Analysis of actual and potential impurities by conducting database and literature searches across carcinogenicity and mutagenicity data.

Sarah nexus

- Transparent Predictions**
Predictions are clearly represented and supported by a measure of confidence. This high level of transparency facilitates expert review.
- Expert ICH M7 Support**
Sarah can be used as part of an ICH M7 workflow and predictions are accepted by regulators under the ICH M7 guideline.
- Confident Predictions**
The Sarah model is built upon a large, high-quality dataset that has been curated by Lhasa experts.
- A statistical-based system for the prediction of mutagenicity
- Using a unique machine-learning methodology, Sarah provides statistical-based predictions for mutagenicity.

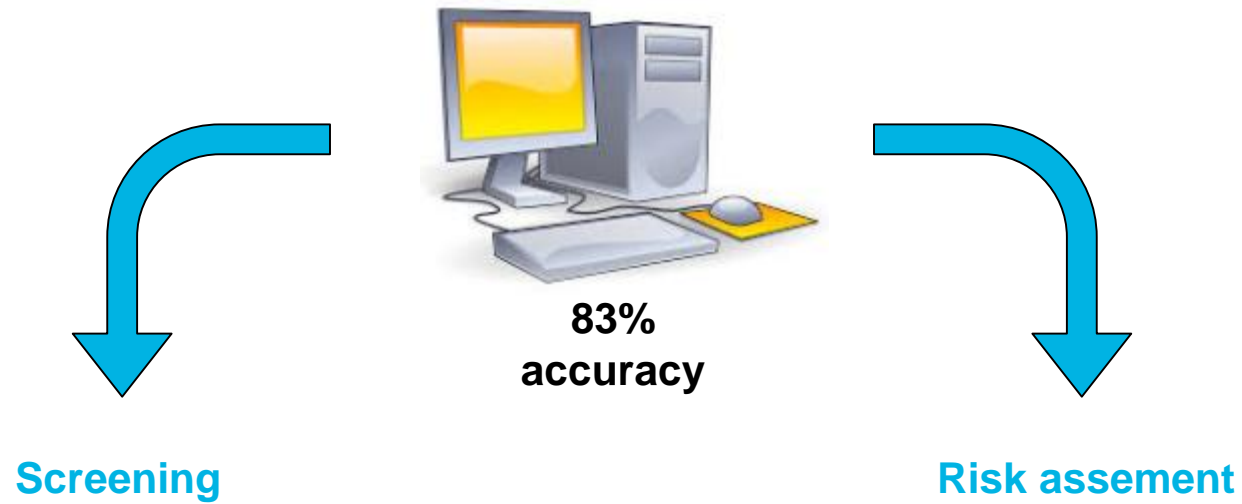
Derek nexus

- Skin Sensitisation Predictions**
Using a Nearest Neighbour approach, EC3 values are predicted for compounds that fire a skin sensitisation alert.
- Expert ICH M7 Support**
Derek can be used as part of an ICH M7 workflow and predictions are accepted by regulators under the ICH M7 guideline.
- Reducing Risk in R&D**
Save time and money by identifying potentially toxic chemicals early, thereby reducing risk in research and development.
- Variety of Endpoints Covered**
Derek predicts for various endpoints including carcinogenicity, mutagenicity, genotoxicity, teratogenicity, irritation and more.
- The preferred expert rule-based system for toxicity prediction. Winner of the Queen's Award for Enterprise: Innovation 2016
- Using structure activity relationships created by Lhasa's scientific experts, Derek provides scientifically robust and transparent toxicity predictions for query compounds.

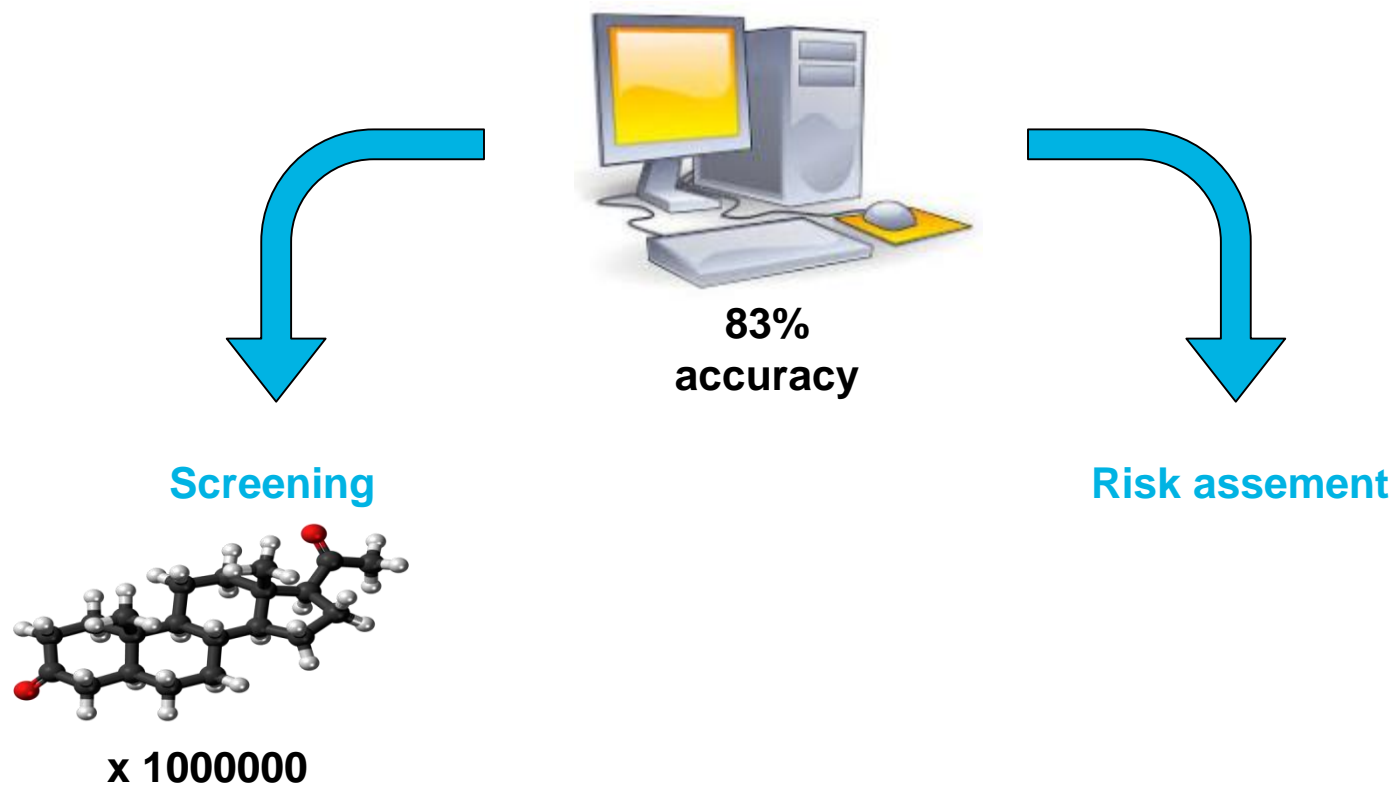
Zeneth TM

- Advanced Warning System**
Provides an instant understanding of what results might be expected when running stress tests.
- Transparent Predictions**
Predictions are clearly represented and contain detailed supporting evidence associated with the degradation pathway.
- Meeting Regulatory Requirements**
Numerous guidelines (including RDC 53) stipulate the need to understand the consequences of degradation, and focus on impurities and stability testing.
- An expert, knowledge-based system for the prediction of degradation pathways
- The only actively maintained system for the prediction of degradation. Zeneth uses structure activity relationships to provide forced degradation predictions for query compounds.

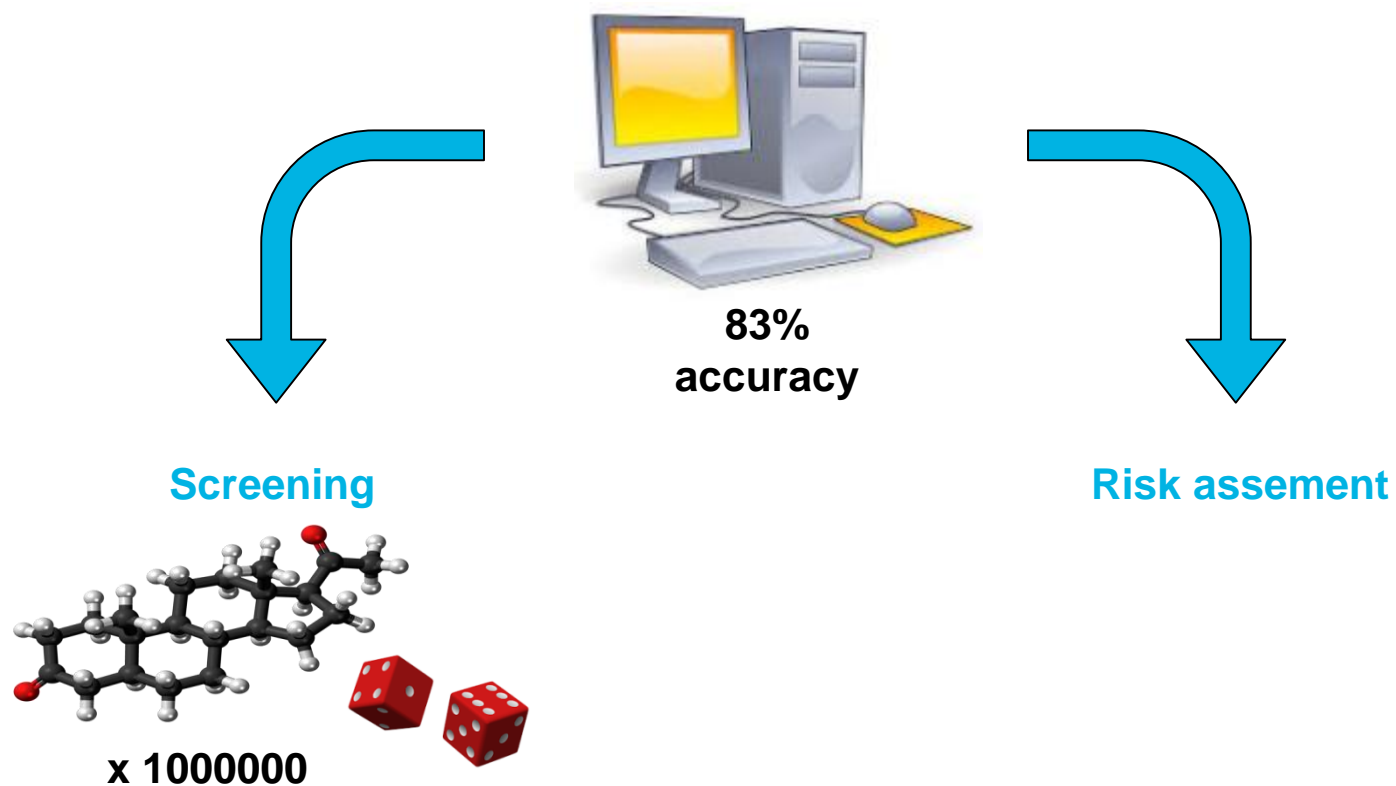
Can we trust a specific individual prediction?



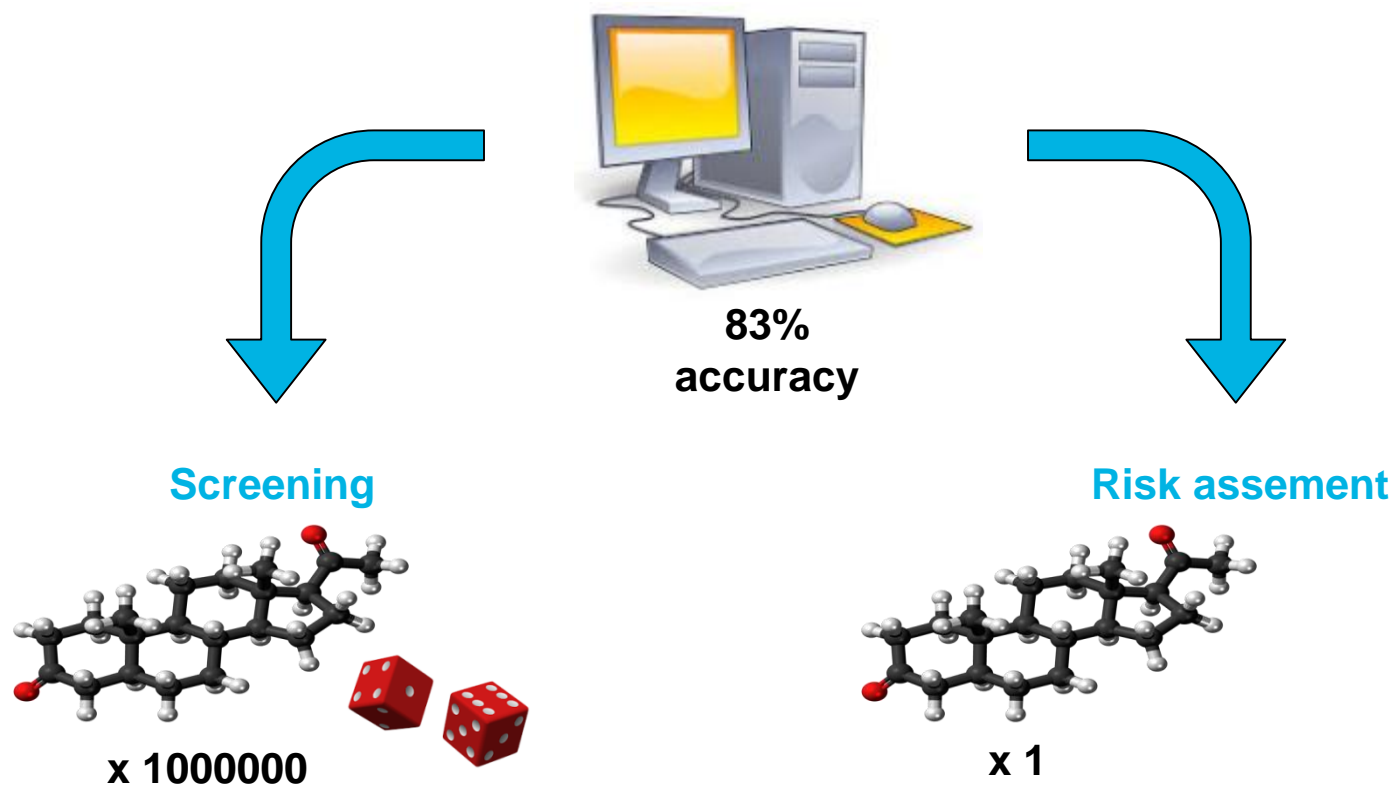
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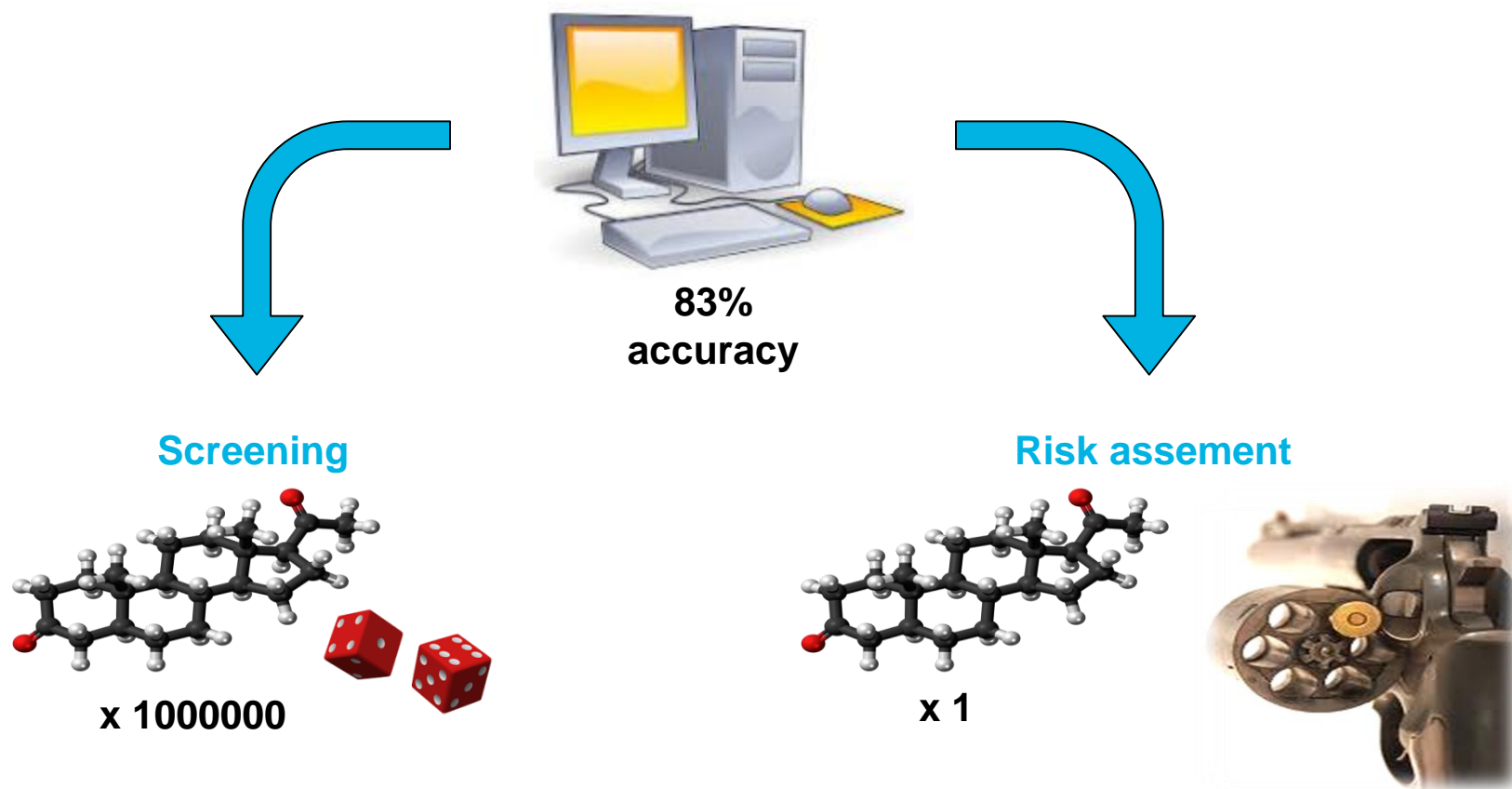
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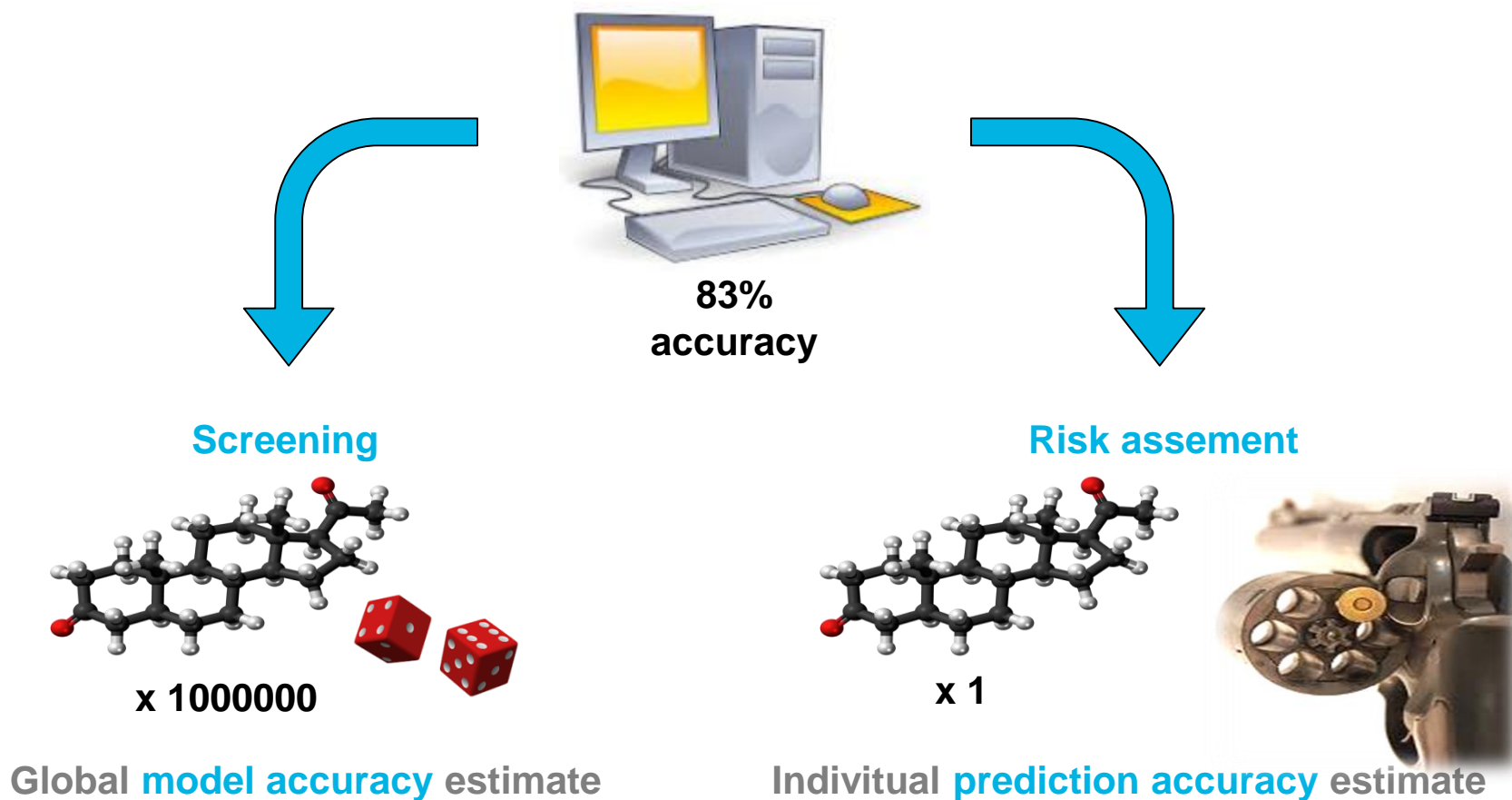
Can we trust a specific individual prediction?



Can we trust a specific individual prediction?



Can we trust a specific individual prediction?



Current understanding and definitions

OECD QSAR principles¹

- A defined endpoint
- An unambiguous algorithm
- **A defined domain of applicability**
- Appropriate measures of goodness-of-fit, robustness and predictivity
- A mechanistic interpretation, if possible

Common definition²

*“AD is the response and chemical structure **space** in which the model **makes** predictions with a **given reliability**”.*

 Guidance Document on the Validation of (Quantitative) Structure– Activity Relationship QSAR Models; OECD Series on Testing and Assessment No.69; OECD Environment Directorate, Environment, Health and Safety Division: Paris, 2007

 Setubal workshop report : Jaworska, J. S.; Comber, M.; Auer, C.; Van Leeuwen, C. Environ. Health Perspect. 2003, 111, 1358–1360

Current understanding and definitions



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Common definition²

"AD is the response and chemical structure space in which the model makes predictions with a given reliability".

Applicability

Boundaries

Likelihood ?

Reliability



A good fundation to build on

- Mathea M, Klingspohn W, Baumann K. **Chemoinformatic Classification Methods and their Applicability Domain**. Mol Inf. 2016 May 1;35(5):160–80.
- Gadaleta D, Mangiatordi GF, Catto M, Carotti A, Nicolotti O. **Applicability Domain for QSAR Models:: Where Theory Meets Reality**. International Journal of Quantitative Structure-Property Relationships. 2016 Jan;1(1):45–63.
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- Toccacheli P, Nourtdinov I, Gammerman A. **Conformal Predictors for Compound Activity Prediction**. arXiv:160304506 [cs] [Internet]. 2016 Mar 14 [cited 2016 May 11]; Available from: <http://arxiv.org/abs/1603.04506>
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- Sheridan RP. **The Relative Importance of Domain Applicability Metrics for Estimating Prediction Errors in QSAR Varies with Training Set Diversity**. J Chem Inf Model. 2015 Jun 22;55(6):1098–107.
- Cherkasov A, Muratov EN, Fourches D, Varnek A, Baskin II, Cronin M, et al. **QSAR Modeling: Where have you been? Where are you going to?** J Med Chem. 2014 Jun 26;57(12):4977–5010.
- Carrió P, Pinto M, Ecker G, Sanz F, Pastor M. **Applicability Domain Analysis (ADAN): A Robust Method for Assessing the Reliability of Drug Property Predictions**. J Chem Inf Model. 2014 May 27;54(5):1500–11.
- Toplak M, Močnik R, Polajnar M, Bosnić Z, Carlsson L, Hasselgren C, et al. **Assessment of Machine Learning Reliability Methods for Quantifying the Applicability Domain of QSAR Regression Models**. J Chem Inf Model. 2014 Feb 24;54(2):431–41.
- Dragos H, Gilles M, Alexandre V. **Predicting the Predictability: A Unified Approach to the Applicability Domain Problem of QSAR Models**. J Chem Inf Model. 2009 Jul 27;49(7):1762–76.

And many more...

A good fundation to build on

molecular informatics
models – molecules – systems

Full Paper

Structure Modification toward Applicability Domain of a QSAR/QSPR Model Considering Activity/Property

Shoki Ochi, Tomoyuki Miyao, Kimito Funatsu ✉

Volu

First published: 16 August 2017 | <https://doi.org/10.1002/minf.201700076>



ELSEVIER

Predicting skin sensitizers with confidence — Using conformal prediction to determine applicability domain of GARD

Andy Forreryd ^a ✉, Ulf Norinder ^{b, c} ✉, Tim Lindberg ^a ✉, Malin Lindstedt ^a ✉

JCIM
JOURNAL OF
CHEMICAL INFORMATION
AND MODELING

Conformal Regression for Quantitative Structure–Activity Relationship Modeling—Quantifying Prediction Uncertainty

Fredrik Svensson^{†‡}, Natalia Aniceto[†], Ulf Norinder[§], Isidro Cortes-Ciriano[†], Ola Spjuth[†], Lars Carlsson[¶], and Andreas Bender[†]

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A strategy on the definition of applicability domain of model based on population analysis

Yong-Huan Yun ^{a, b, c} ✉, Dong-Ming Wu ^{a, c}, Guang-Yi Li ^{a, c}, Qiao-Yan Zhang ^a ✉, Xia Yang ^a, Qin-Fen Li ^{a, c}, Dong-Sheng Cao ^d, Qing-Song Xu ^e

Current common methods

Molecule classes

- Organic-Organometallic-Inorganic
- Class of molecules (Arom. Amines)

Feature representation

- Unseen features

Agreement based

- RF consensus
- kNN

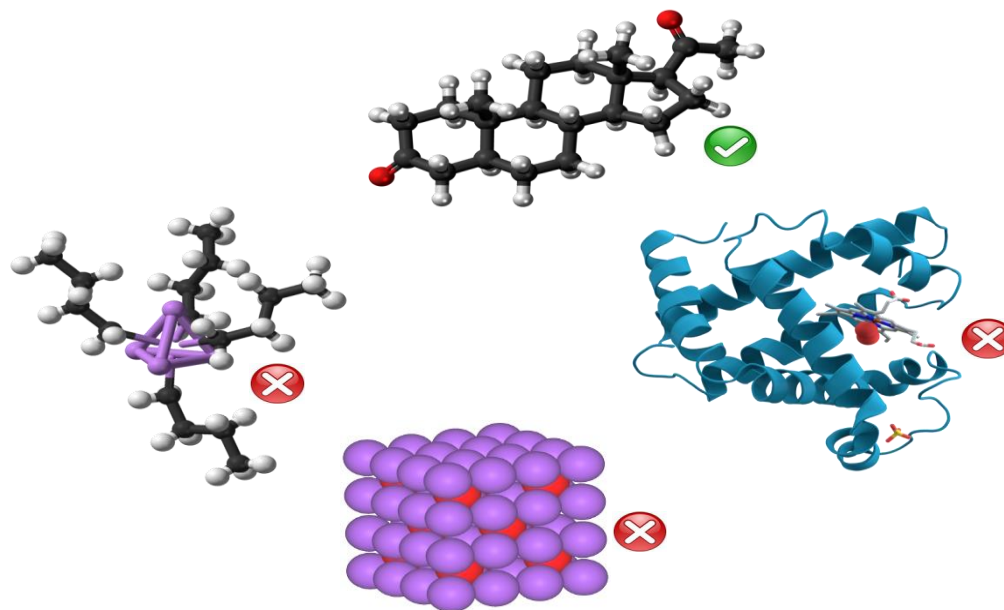
Descriptor ranges

- Box
- Convex hull

Distance based methods

- Distance to data points
- Density

Response domain



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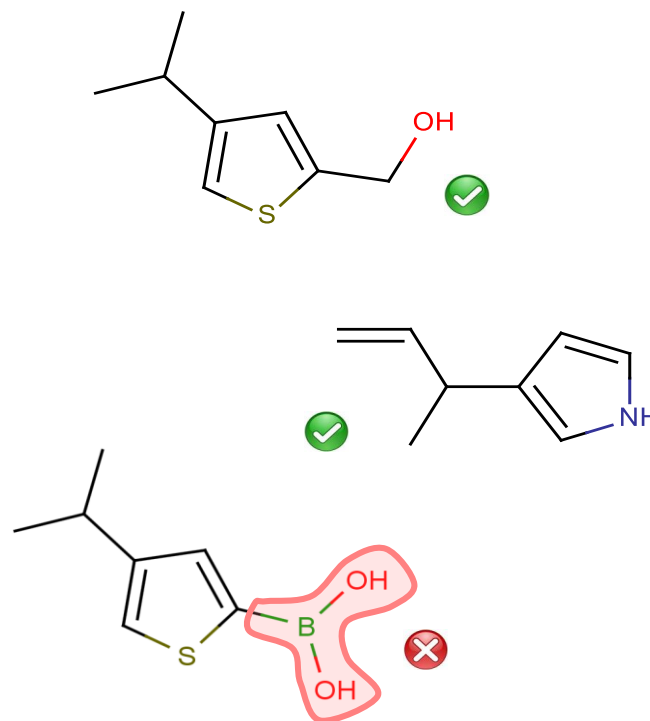
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No boronic acids in the training set

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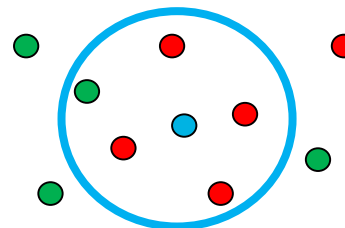
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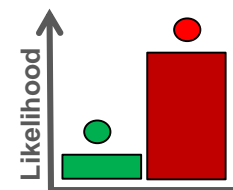
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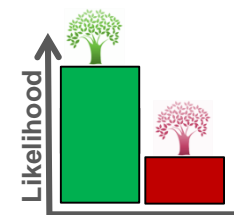
Response domain



Nearest Neighbours



Random forest



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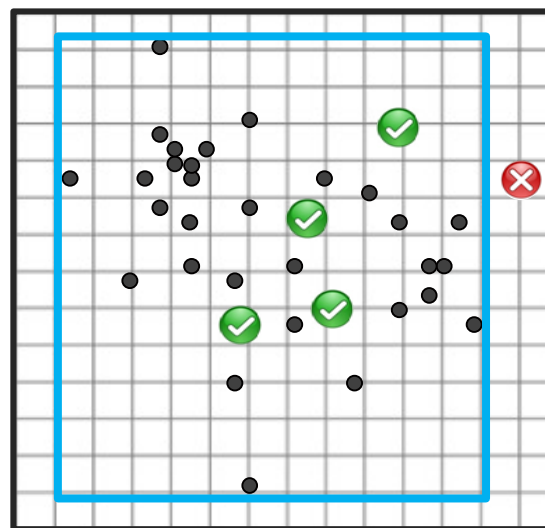
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Response domain



Box ✓ ✗

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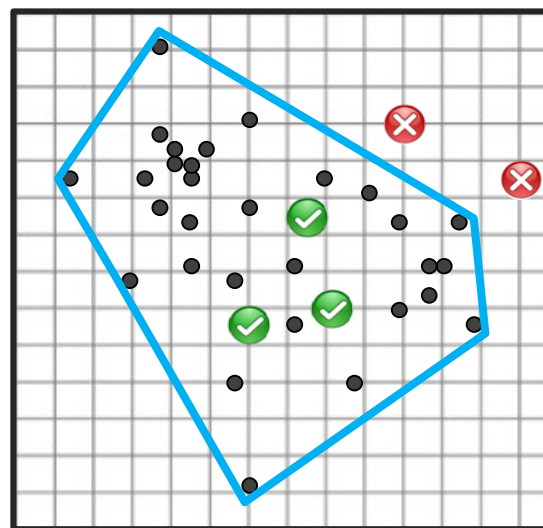
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Response domain



Convex hull ✓ ✗

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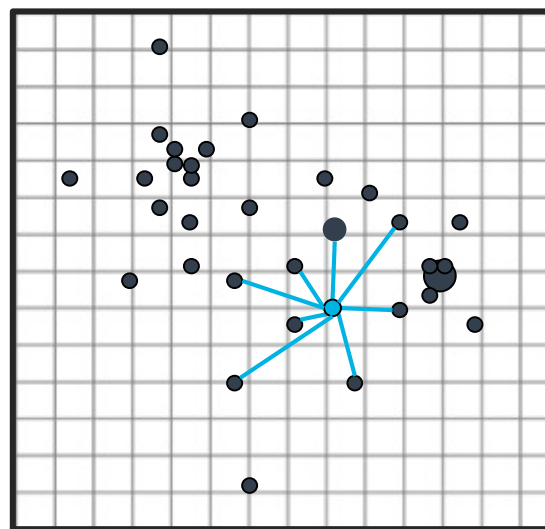
- Box
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Distance based methods

- Distance to data points
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Response domain

Distance to data



Distance to data points



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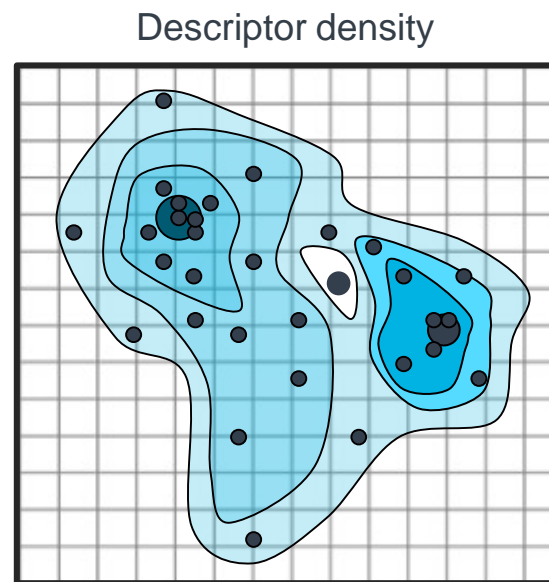
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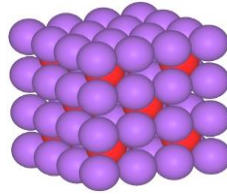
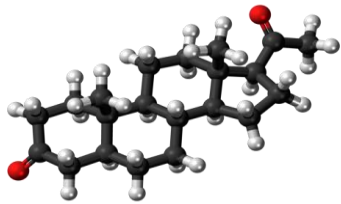
- Distance to data points
- Density

Response domain



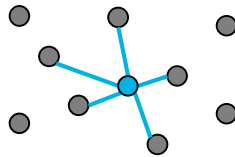
Density of data 

Mixture of different concepts



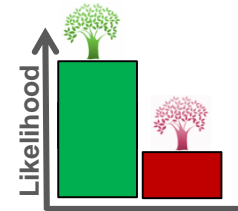
Applicability

(can I use this model to make a prediction ?)



Reliability

(is the prediction reliable?)



Decidability

(can I make a clear decision)

Mixture of different concepts



Applicability

(can I use this model to make a prediction?)



Reliability

(is the prediction reliable?)



Decidability

(can I make a clear decision)

Mixture of different concepts



Applicability Domain

Towards an extended and more formal framework

Confidence in the prediction if ...



My model can be applied for this query compound

Applicability
domain



The prediction is reliable enough for my use case

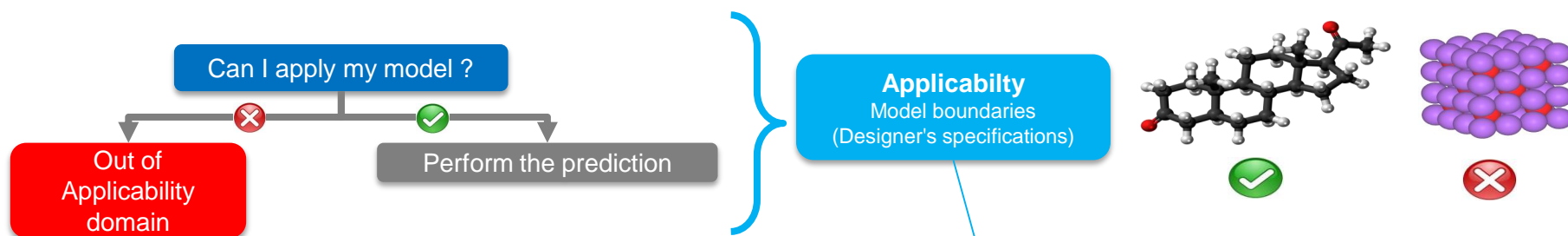
Reliability
domain



I can make a clear decision

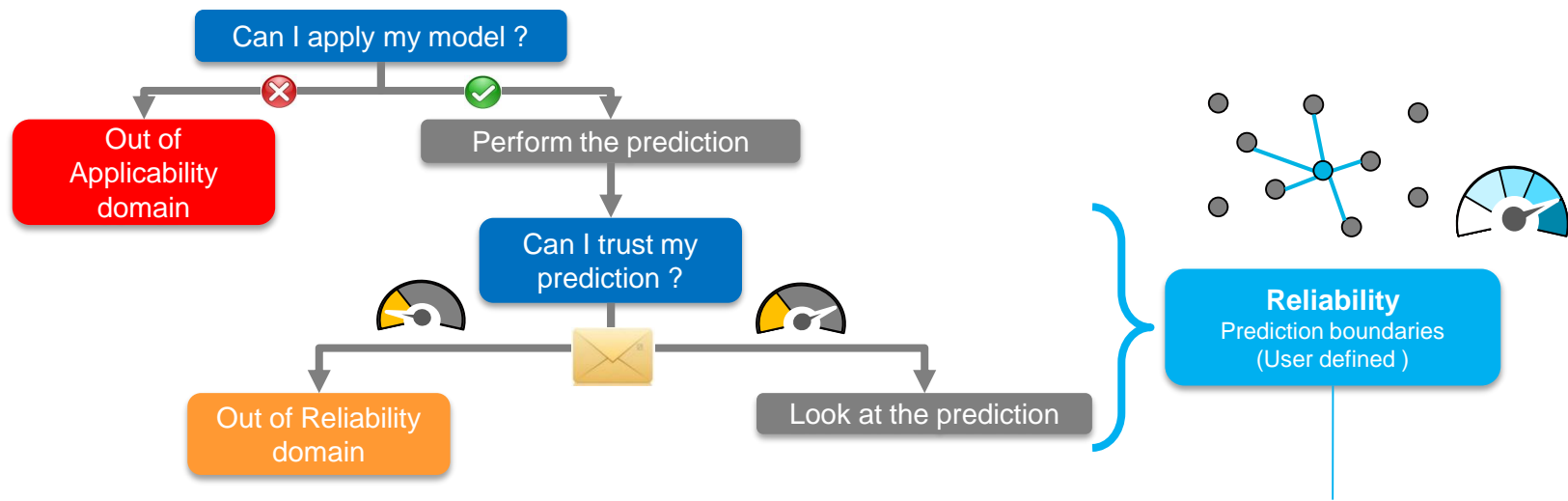
Decidability
domain

Applicability (of the model)



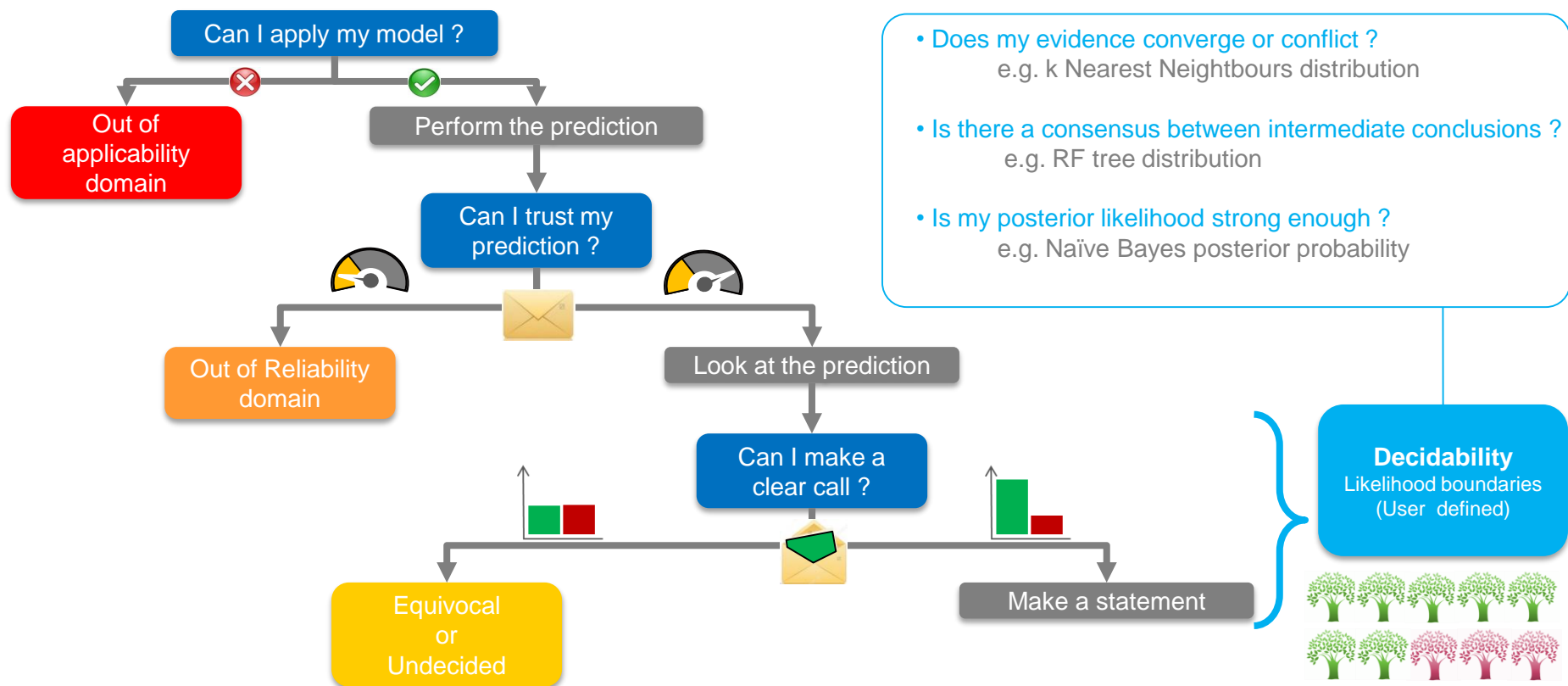
- Is the class of my query compound supported by the model ?
e.g. exclude polymers, proteins, inorganic molecules, etc.
- Is my query compound in the range of the descriptor of the training set ?
e.g. inside convex hull, minimum information density
- Did my model see all the structural features present in the query compound ?
e.g. not in domain, contains unseen boronic acid functional group

Reliability (of the prediction)

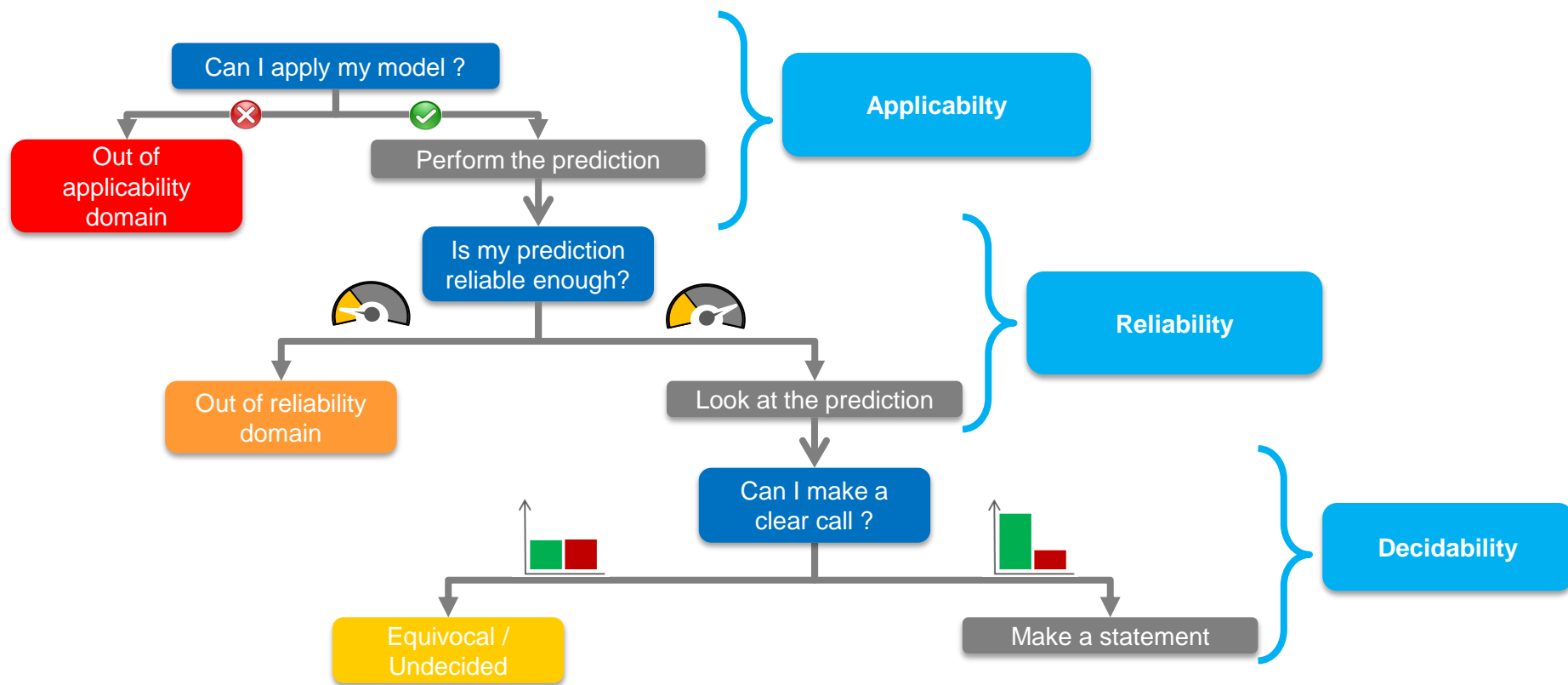


- How close are the nearest neighbours ?
- How reliable are these nearest data points ?
e.g. GLP compliance
- How well did my model predict these data points ?
e.g. performance during CV

Decidability (of the outcome)



Intuitive, non ambiguous and formal decision framework



Articulation of the method

- Applicability domain is **not a monolithic concept**, there are 3 key layers
- Separation of concern can **help clarify and formalise** the notion of AD
- Purpose: Initiate a constructive discussion among our QSAR community to build a **common understanding** together
- **Harmonize the way we define and present AD** to the end users across models and applications
- **Remove confusion** for the end user and improve the value of our AD model

Collaborators

- **Stéphane Werner**
- **Jean-François Marchaland**
- **Sébastien Guenes**
- **Lilia Fisk**
- **Chris Barber**



Thank you
for your
kind attention

shared **knowledge** • shared **progress**

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