

# Mutagenicity Prediction Using *In Silico* Methods:

Gigabyte-Sized Petri Dishes.

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# Predictive Toxicology

- **What is it?**

- ◉ Collection of strategies employed to forecast the interaction between chemical compounds that lead to adverse effects in biological systems.

- **Why is it important?**

- ◉ ICH M7 guidelines for the assessment of mutagenic impurities in drugs explicitly include the use of *in silico* methods to fulfil regulatory requirements.

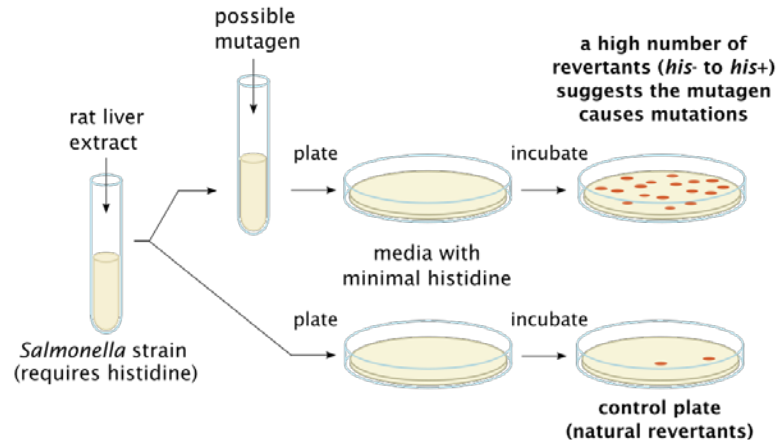
- ◉ Replaces animal testing (follows the 3R initiative: Replacement, Reduction and Refinement).

- ◉ Fast, cost effective, reliable and reproducible approach.

# Predictive Toxicology

- **Genotoxicity is the capacity of a chemical compound to cause damage to DNA.**
  - ⊙ It covers polyploidy, aneuploidy, mutagenicity, chromosome damage and non-inheritable DNA damage.
  - ⊙ Predicting genotoxicity is essential to ensure the safety of drugs, foods, etc...
  
- **How is genotoxicity predicted?**
  - ⊙ *In vitro* assays: **Replacement** of animal testing for cell/bacterial experiments.
  - ⊙ *In silico* prediction methods: **Replacement** of *in vitro* assays for algorithms in a computer.

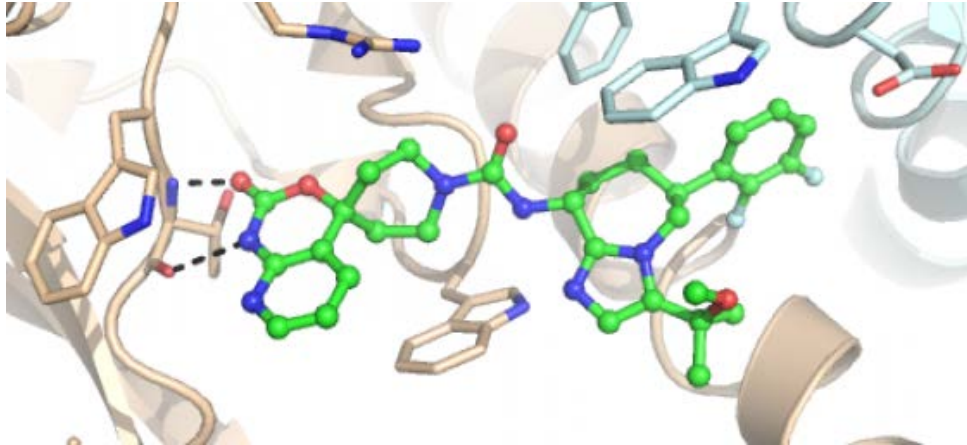
# Predictive Genotoxicity



- ***In vitro* tests**

- ⊙ Study the genotoxic effect of a specific chemical in bacterial, yeast or mammalian cells.
- ⊙ They are fast, easy to set up, cheaper than animal tests, simple to run, amenable for automation and provide quick results.
- ⊙ Main limitations come from the extrapolation of the *in vitro* data to *in vivo* systems and from the need to have the test substance isolated and purified in sufficient quantities to conduct the tests.

# Predictive Genotoxicity



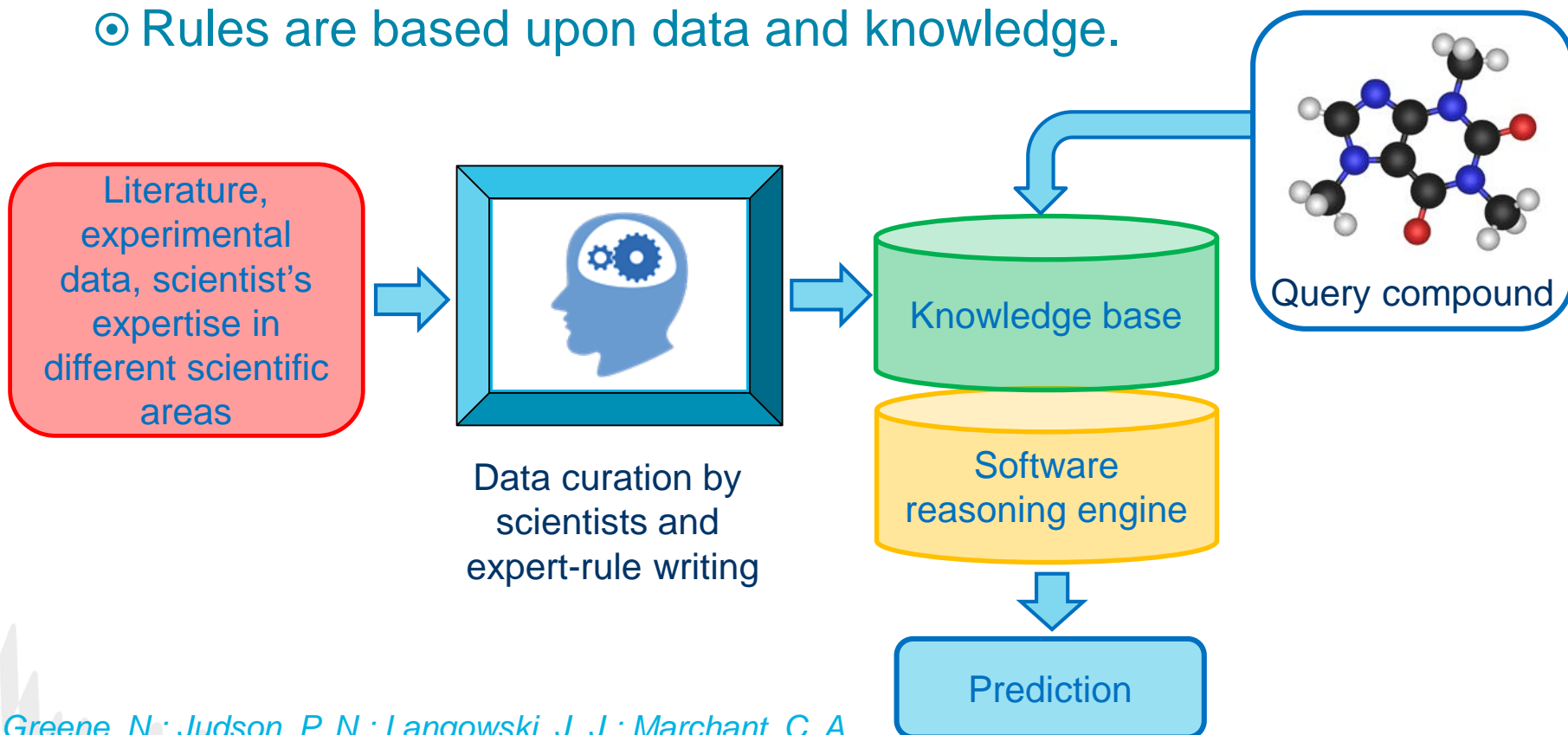
- ***In silico* tests**

- ⊙ Toxicity assessment that uses computational methods to model, simulate or predict toxicity of chemicals.
- ⊙ Higher throughput, faster, cheaper than *in vitro* methods and present a high reproducibility if the same model is used. Information about the mechanism of action can also be obtained.
- ⊙ Limitations root in the quality of the dataset, not accounting for ADME features and the complexity of certain endpoints (e.g. neurotoxicity).

# *In silico* Predictive Genotoxicity

- **Expert Knowledge Software.**

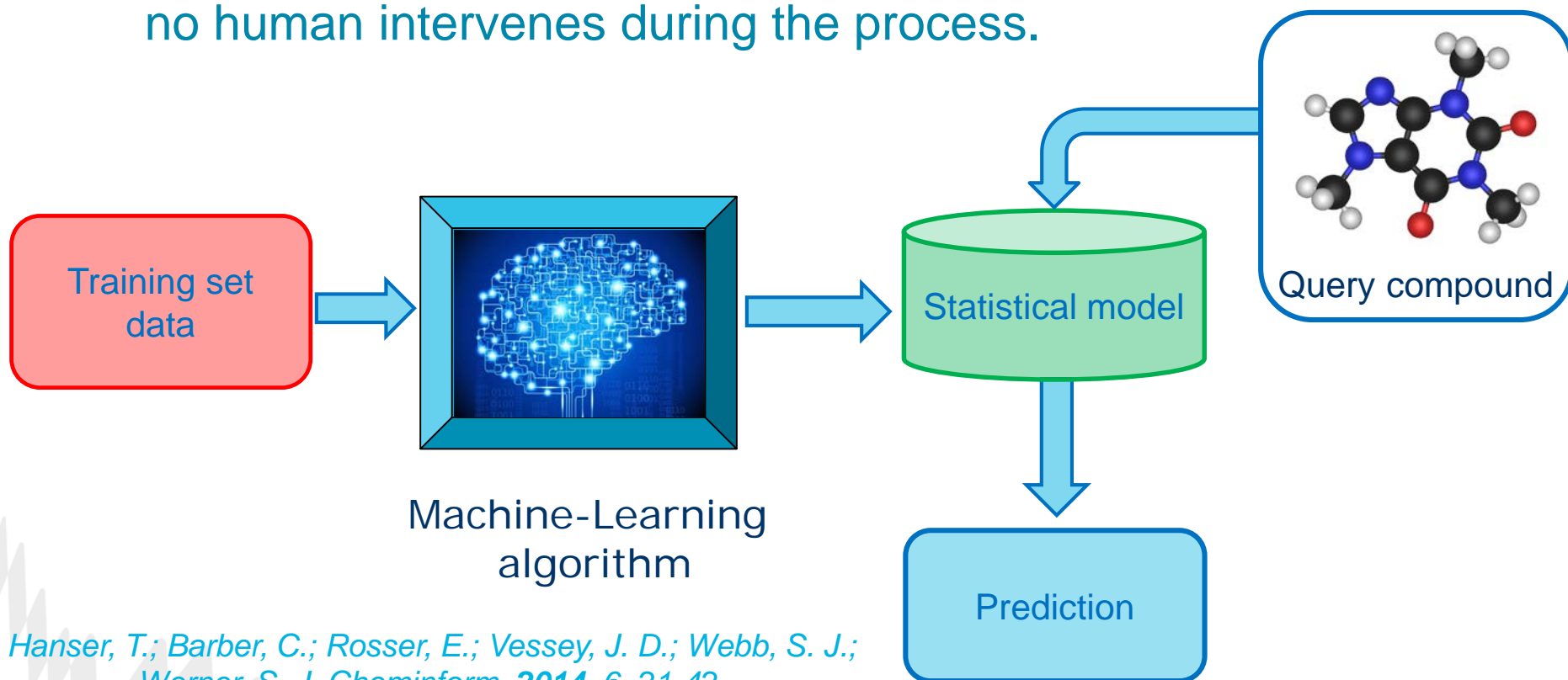
- ⊙ Scientists write expert rules that relate chemical structure to toxicity.
- ⊙ Rules are based upon data and knowledge.



# *In silico* Predictive Genotoxicity

- **Statistical Modelling Software.**

- ⊙ A training dataset is used to construct a statistical structure-activity model through machine learning algorithms.
- ⊙ This model is then used to predict the query compound toxicity; no human intervenes during the process.



# Importance of *in silico* methods: ICH M7

- “Assessment and Control of DNA Reactive (Mutagenic) Impurities in Pharmaceuticals to Limit Potential Carcinogenic Risk”
  - ⊙ ‘Global’ guidelines – America, Europe, Japan
  - ⊙ Finalised - June 2014
  - ⊙ In force since Jan 2016

identification

categorisation

qualification

Control of mutagenic impurities to limit potential carcinogenic risk

<http://www.ich.org/products/guidelines/multidisciplinary/article/multidisciplinary-guidelines.html>



# Importance of *in silico* methods: ICH M7

| Class | Definition   | Proposed action for control   |
|-------|--|---|
| 1     | Known mutagenic carcinogens  | Control at or below compound-specific acceptable limit  |
| 2     | Known mutagens with unknown mutagenic potential (bacterial mutagenicity positive, no rodent carcinogenicity data)  | Control at or below acceptable limits (TTC)   |
| 3     | Alerting structure, unrelated to the structure of the drug substance; no mutagenicity data   | Control at or below acceptable limits (acceptable TTC) or conduct bacterial mutagenicity assay;<br>If non-mutagenic=Class 5<br>If mutagenic=Class 2 |
| 4     | Alerting structure, same alert in drug substance or compounds related to the drug substance (e.g., process intermediates) which have been tested and are non-mutagenic | Treat as non-mutagenic impurity.  |
| 5     | No structural alerts, or alerting structure with sufficient data to demonstrate lack of mutagenicity or carcinogenicity  | Treat as non-mutagenic impurity.  |

<http://www.ich.org/products/guidelines/multidisciplinary/article/multidisciplinary-guidelines.html>

# Focussing on the *in silico* predictions...

- Q(SAR) Requirements:

- ⊙ must predict the outcome of a bacterial mutagenicity assay
- ⊙ one expert rule-based; the second statistical-based
- ⊙ both should follow the OECD principles

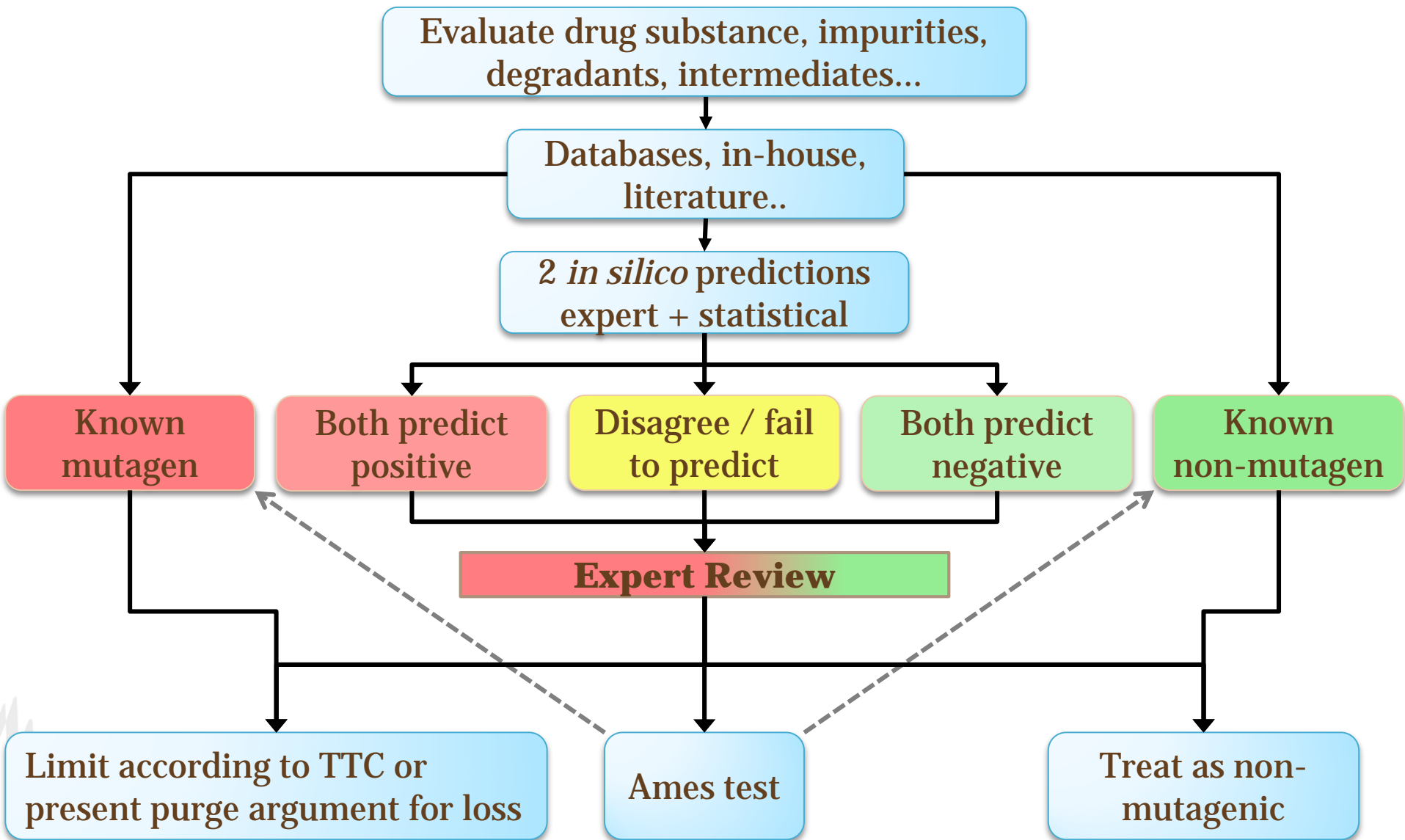
1. *a defined endpoint;*
2. *an unambiguous algorithm;*
3. *a defined domain of applicability;*
4. *appropriate measures of goodness-of-fit, robustness and predictivity;*
5. *a mechanistic interpretation, if possible.*

- The absence of structural alerts from both is sufficient to conclude that the impurity is of no mutagenic concern

- ⊙ Expert review can provide

- ⊙ additional supportive evidence
- ⊙ reason to dismiss an *in silico* prediction
- ⊙ rationale to support the final conclusion

# *In silico* workflow under ICH M7



# Lhasa Limited Solutions



**Derek Nexus** – Expert knowledge-based toxicity prediction software



**Meteor Nexus** – Expert decision support software for predicting the metabolic fate of chemicals in mammals



**Vitic Nexus** - Chemical database and information management system, offering researchers and scientists rapid access to searchable toxicological information



**Zeneth** - Expert decision support software for predicting the forced degradation pathways of organic compounds



**Sarah Nexus** - Statistical-based software for the prediction of mutagenicity

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# Of expert and statistical systems...

## Expert system

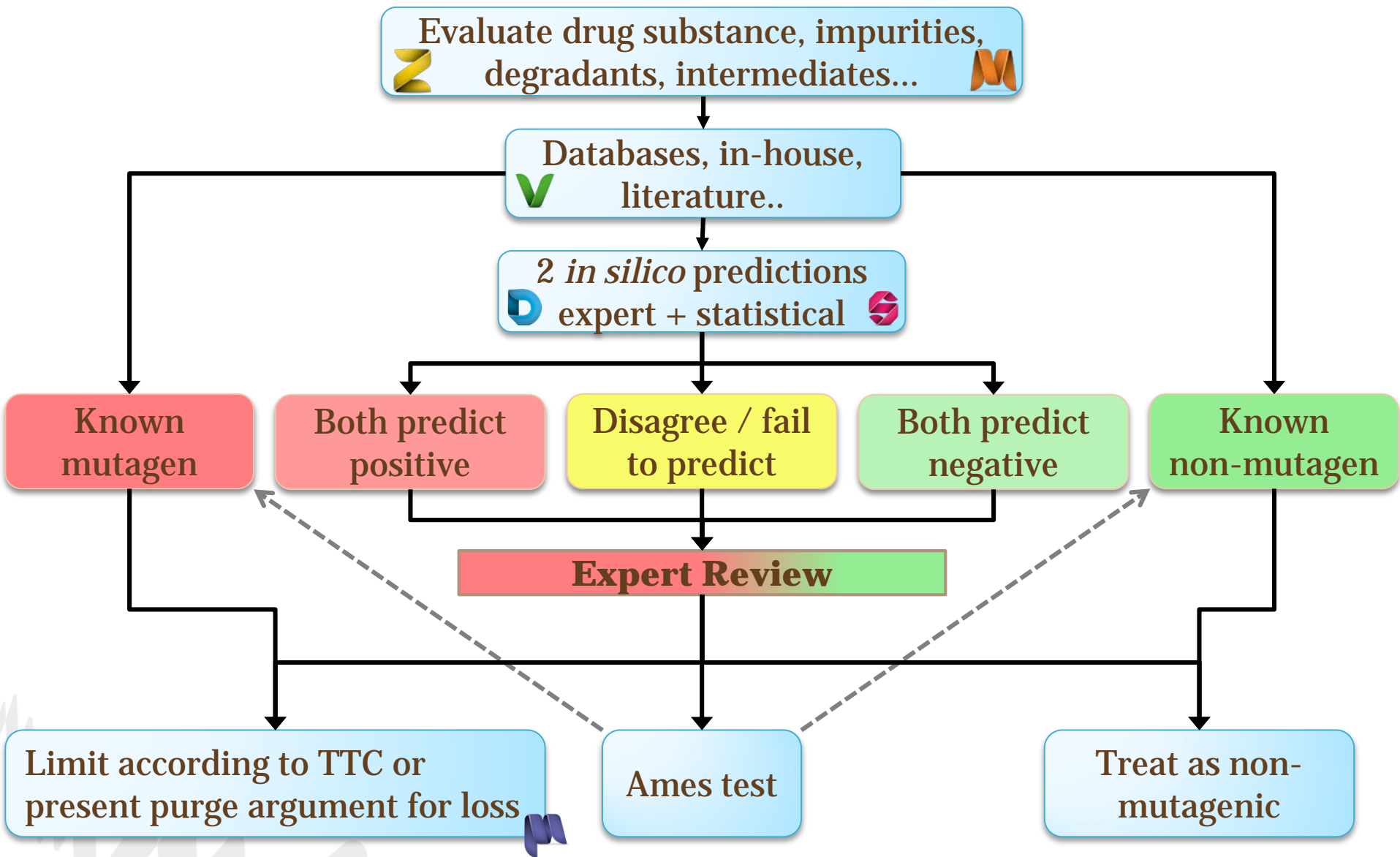


## Statistical system



|                  |   |   |
|------------------|---|---|
| Data             | <ul style="list-style-type: none"><li>• Public <i>and confidential data</i></li><li>• Wider knowledge - mechanism, reactivity, related assays...</li></ul>  | <ul style="list-style-type: none"><li>• Transparent systems can only show non-confidential data</li></ul>   |
| Methodology      | <ul style="list-style-type: none"><li>• Human-written rules based upon data &amp; knowledge</li></ul>   | <ul style="list-style-type: none"><li>• <i>Machine-learnt fragments</i></li><li>• Machine-learnt significance of each fragment</li></ul>  |
| Scope of alert   | <ul style="list-style-type: none"><li>• Human defined (Markush)</li></ul>   | <ul style="list-style-type: none"><li>• Fragments learnt by model</li></ul>   |
| Interpretability | <ul style="list-style-type: none"><li>• Expert commentary</li><li>• Mechanistic explanation</li><li>• Supporting examples</li><li>• References</li><li>• Confidence</li><li>• Highlights areas of uncertainty</li></ul> | <ul style="list-style-type: none"><li>• Transparent methodology</li><li>• No uninterpretable descriptors or definitions of applicability</li><li>• Learning summarised</li><li>• Direct link to training set</li><li>• Confidence in prediction</li></ul> |

# How Lhasa fits into ICH M7 workflow



# Conclusions

- ⊙ **Predictive Toxicology** tries to forecast toxicity by studying the perturbation of biological pathways on a molecular basis.
- ⊙ It moves away from the direct observation of adverse effects in animals to *in vitro* and *in silico* tests.
- ⊙ *In vitro* assays provide experimental data in a fast way.
- ⊙ *In silico* methods are faster, more cost effective and have a higher throughput than *in vitro* assays.
- ⊙ Reliability of *in silico* methods results has motivated the possibility of being employed as a substitute for *in vitro* assays, as stated by ICH M7 guidelines for mutagenicity.
- ⊙ Improvements in computational power and new algorithms will cause a sheer increase in the predictive capabilities of *in silico* methodologies.



# Thank you for your attention!

## Not-for-profit Educational charity

Teaching  
lectures

Interns

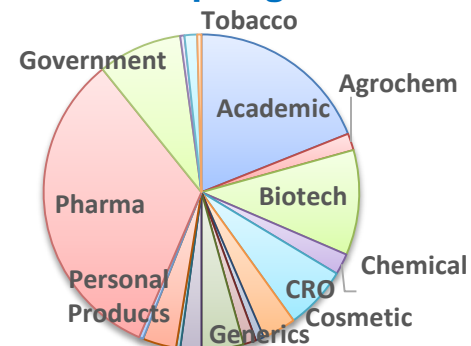
Sponsor  
PhDs

Undergrad  
projects

## Located in Leeds, UK



## A membership organisation



## Predictive software (expert & machine learnt)



Purge



Degradation



Toxicity



Metabolism



## Regulators (members & collaborators)



## Data & knowledge sharing Honest broker



MIP-DILI



Proprietary data mining