



# New QSAR system to predict acute inhalation toxicity

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Acute inhalation toxicity is the adverse effect caused by a substance after a single uninterrupted exposure by inhalation over a period of 24 hours or less. Environmental pollution and workplace/household exposure require special attention to the inhalation toxicity of chemicals that can cause damage to the respiratory organs and organism as a whole. While regulatory bodies, e.g. European Union, require inhalation toxicity evaluation to be performed for high volume production chemicals, the use of animals is not permitted in such evaluations.

In place of animal testing, it is often recommended to use search and retrieval of existing data, identification of structural alerts to judge toxicity, grouping of chemicals for read-across analysis, and the development and use of quantitative structure-activity models. However, the volume of experimental data and number of reliable QSAR models, concerning the acute inhalation toxicity is limited which hinders all in-silico approaches.

In this study, we developed a set of QSAR models built using publicly available acute inhalation toxicity data in rat. The set consists of 4 categorical models, designed to predict one of the four GHS categories for inhalation toxicity, and a model to predict LC50 values directly.

All models demonstrated leave group out cross-validation performance in the range of 52-71% sensitivity, 71-74 % specificity, 64-73 % positive predictivity, and 61-68 % negative predictivity. The inhalation toxicity set has assign the correct GHS category to 75% of the external validation compounds. Coverage was maintained in the range of 71-84 %. Identified alerts are discussed along with the advantages and disadvantages of using the categorical versus the direct model. Overall, this approach provides a reliable QSAR method for predicting acute inhalation toxicity of chemical substances, only using their structures.

# Regulatory Context

- Acute inhalation toxicity data are used to satisfy hazard classification and labeling requirements, to estimate the toxicity of mixtures, and to assess human health and environmental risk.
- The derivation of either a point estimate of the LC50 value (using TG 403) or a range estimate of the LC50 (using TG 436) generally meets the acute inhalation toxicity regulatory requirements for classification and labeling of industrial chemicals, consumer products, and many pesticide applications.
- (Q)SAR models for Inhalation toxicity endpoints may be used as early screening tools during drug development or to support regulatory safety decisions when experimental data are limited or unavailable.
- Predictions may be used to contribute to the weight-of-evidence supporting a regulatory safety decision.

# Objectives

Build optimized InhTOX (Q)SAR models for

- GHS categorical prediction (range estimate of LC50)
- Point estimate of LC50 predictions

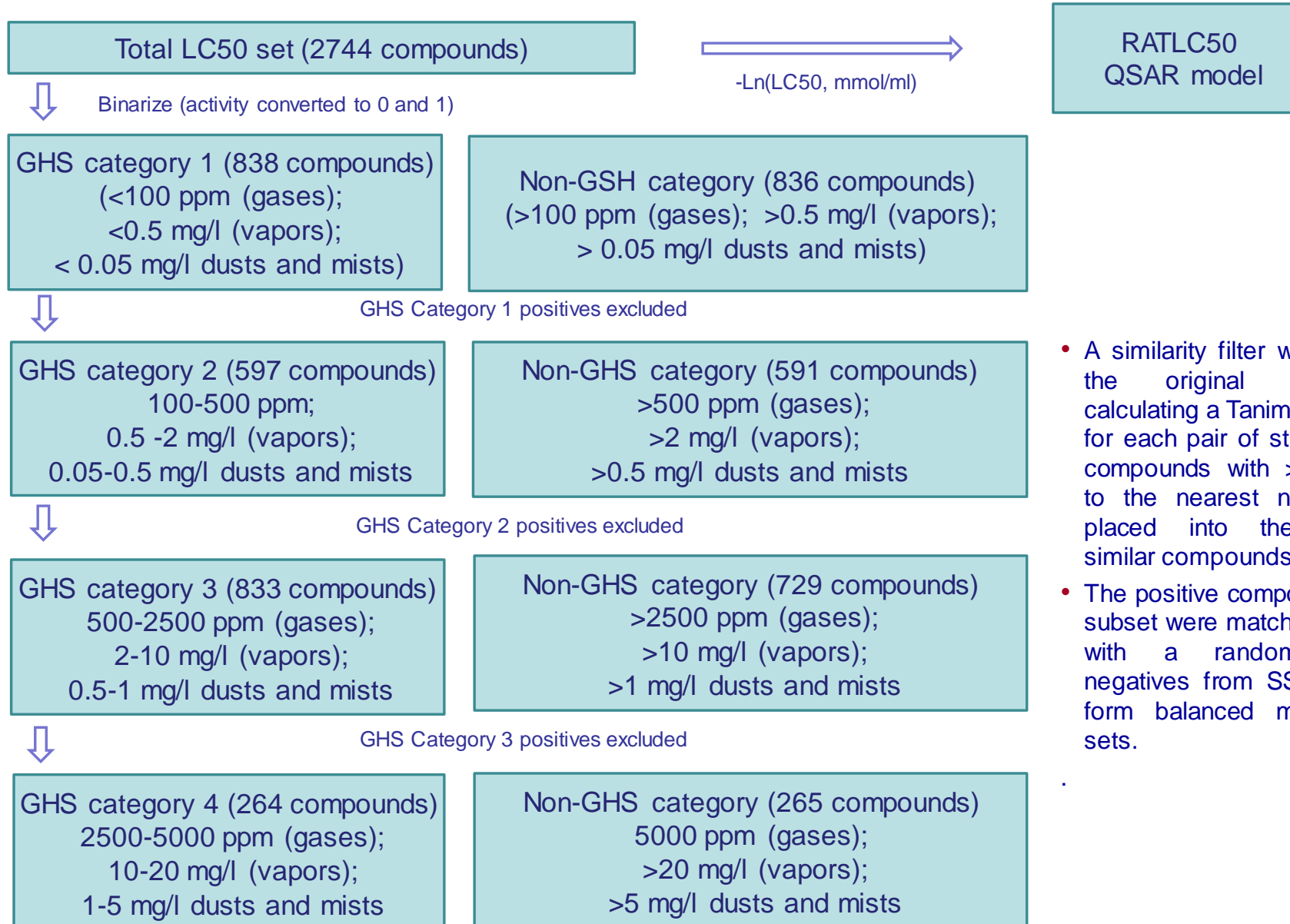
# Software and Training Sets for (Q)SAR Models

- All training sets used to construct (Q)SAR models were generated from an existing non-proprietary database of Inhalation Toxicity assays sourced from publicly available databases and the published literature[1-2].
  - [1] RTECS, Registry Of Toxic Effects of Chemical Substances. [Atlanta, Ga.] :U.S. Dept. of Health and Human Services, Public Health Service, Centers for Disease Control and Prevention, National Institute for Occupational Safety and Health, 1997
  - [2] O. A. Raevsky, E. A. Modina, O. E. Raevskaya SAR models of the inhalation toxicity of organic compounds Pharmaceutical Chemistry Journal 45(3):165-169, 2011
- CASE Ultra (CU) v1.6.2.3 (MultiCASE, Inc.) software was used to build statistical QSAR models.
- A binary scoring system was used to classify assay results as negative (0) or positive (1) for GHS model bundle.
- The endpoints of acute rat inhalation toxicity (4 hours exposure) was chosen for this project. The training sets consisted of 2744 findings, where 30% were Category 1, 21% were Category 2, 30% were Category 3 and 10% were Category 4.

# Data preprocessing

- A **binary scoring system** was used to classify assay results as negative (0) or positive (1) for GHS model bundle.
- The endpoints of **acute rat inhalation toxicity (4 hours exposure)** was chosen for this project. The training sets consisted of 2744 findings, where 30% were Category 1, 21% were Category 2, 30% were Category 3 and 10% were Category 4.
- After **downsampling**, newly constructed model training sets were created containing an equal number of positive and negative findings with the total size of models ranging from 529 to 1674 records. The remaining compounds were used to create supporting datasets to decrease “out of domain” calls produced by the final model.

## Similarity-Based Downsampling



- A similarity filter was applied to the original dataset by calculating a Tanimoto coefficient for each pair of structures. Only compounds with  $>0.3$  similarity to the nearest neighbor were placed into the structurally similar compounds.
- The positive compounds of GHS subset were matched in 1:1 ratio with a randomly selected negatives from SSC subset, to form balanced model training sets.

# Composition of InhTOX Models

| Model          | Endpoint   | Balanced model size       |
|----------------|--|---------------------------|
| INHTOX_CAT1    | Inhalation Toxicity GHS category 1 (<100 ppm)      | 1674 (P:838/N:836)        |
| INHTOX_CAT2    | Inhalation Toxicity GHS category 2 (100-500 ppm)   | 1188 (P:597/N:591)        |
| INHTOX_CAT3    | Inhalation Toxicity GHS category 3 (500-2500 ppm)  | 1562 (P:833/N:729)        |
| INHTOX_CAT4    | Inhalation Toxicity GHS category 4 (2500-5000 ppm) | 529 (P:264/N:265)         |
| INHTOX_RATLC50 | Inhalation Toxicity, Acute, Rat 4 hours LC50       | 2744 (P:1345/M:43/N:1356) |

P=Number of positives; N=Number of negatives; M= Number of marginals



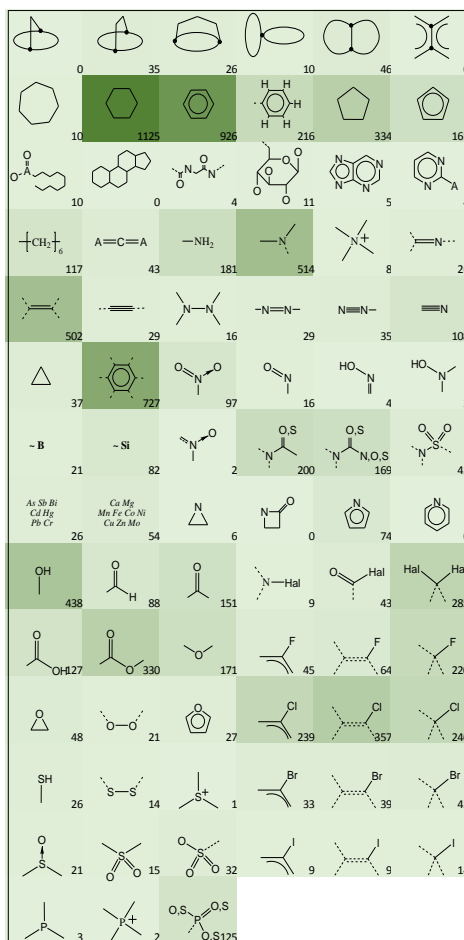
# Predictive Performance of InhTOX Balanced Models

| Model          | Leave group out mode % Off/Times | Balanced model |             |                   |                   |          | External validation, correct classification |
|----------------|----------------------------------|----------------|-------------|-------------------|-------------------|----------|---|
|                |                                  | Sensitivity    | Specificity | Positive Accuracy | Negative Accuracy | Coverage |   |
| INHTOX_CAT1    | 10/10                            | 52.1           | 71.7        | 63.8              | 61.5              | 64.4     | 75.1  |
| INHTOX_CAT2    | 10/10                            | 50.8           | 72.7        | 62.3              | 62.6              | 62.3     | 80.5  |
| INHTOX_CAT3    | 10/10                            | 70.5           | 71.0        | 74.6              | 66.6              | 64.7     | 100.0                                       |
| INHTOX_CAT4    | 10/10                            | 60.4           | 71.8        | 66.9              | 65.5              | 61.7     | 78.0  |
| INHTOX_RATLC50 | 10/10                            | 69.5           | 75.1        | 72.3              | 72.6              | 66.8     | n/a   |

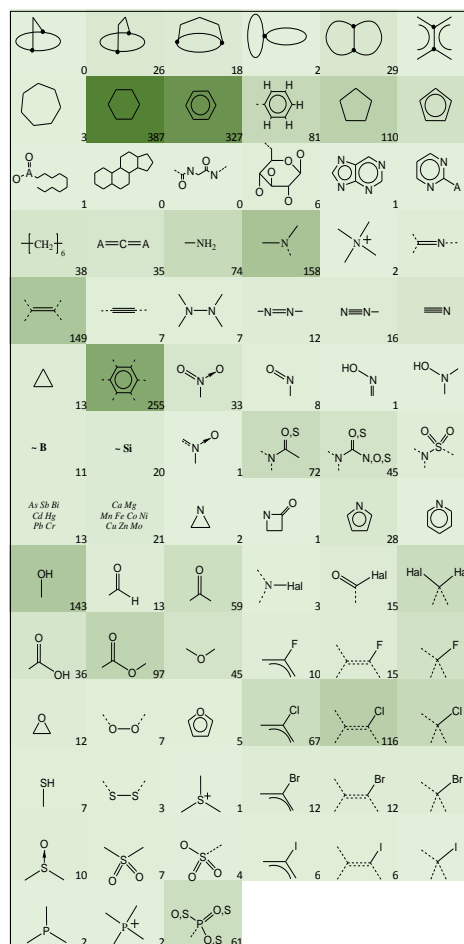
- If a query chemical raises an out of domain call in a GHS balanced model but has no unknown fragments in a corresponding INHTOX\_RATLC50 model, the out of domain call is considered as negative.

# Distribution of Main Structural Features in InhTox Models

## RATLC50



## Cat1

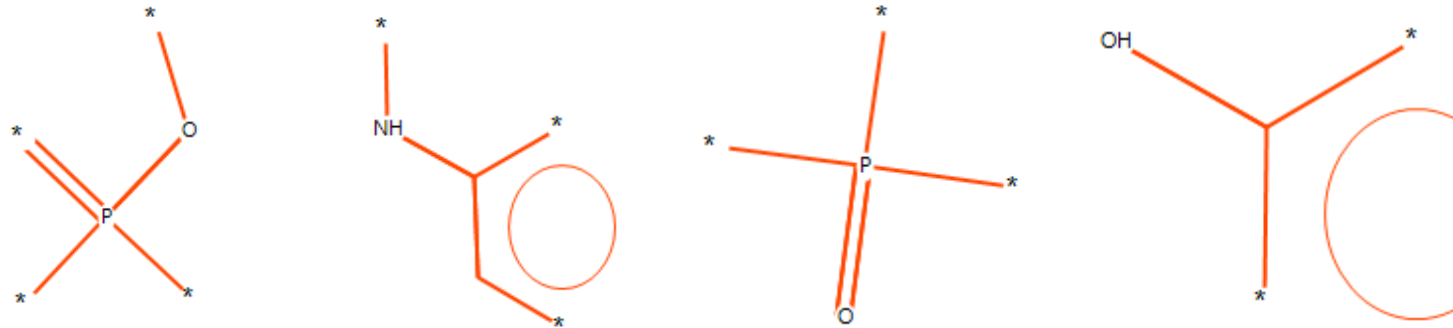


Structural feature representation is maintained between the categorical and LC50 models

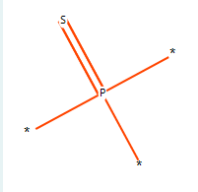
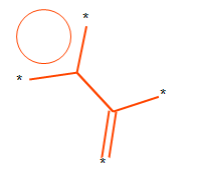
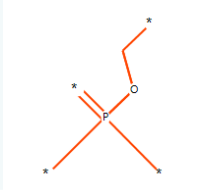
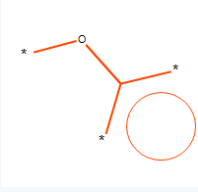
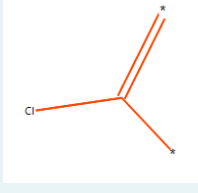

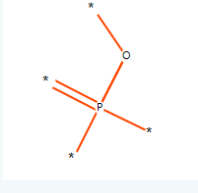
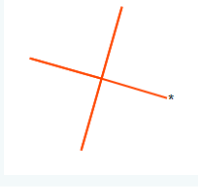
# Analysis of Structural Alerts in Final InhTox Models

- There is a clear distinction between alerts, identified for different GHS categories. The distinction is based on the mechanistic mode of action
- LC50 model identifies more modes of actions, compare to the GHS models

Top alerts LC50 model



# Top alerts for GHS models

| Model       | Alert   | Positives | Negatives | Model       | Alert   | Positives | Negatives |
|-------------|---|-----------|-----------|-------------|---|-----------|-----------|
| InhTOX_CAT1 |    | 37        | 12        | InhTOX_CAT3 |    | 93        | 27        |
| InhTOX_CAT1 |    | 35        | 12        | InhTOX_CAT3 |    | 92        | 27        |
| InhTOX_CAT2 |   | 41        | 14        | InhTOX_CAT4 |   | 13        | 6         |
| InhTOX_CAT2 |  | 30        | 12        | InhTOX_CAT4 |  | 10        | 3         |

# Conclusions

- Combining a similarity-based downsampled (Q)SAR model and a supporting database of additional chemicals provides more sensitive predictions of inhalation toxicity, without a significant decrease in structural coverage.
- The sensitivity and negative predictivity of the new Inhalation Toxicity models demonstrate a reliable QSAR method for predicting acute inhalation toxicity of chemical substances, only using their structures

# Questions?

Please forward your inquiries to:

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