SciMatics SciQSAR model for severe skin irritation in vivo

- 1. QSAR identifier
- 1.1 QSAR identifier (title)

SciMatics SciQSAR model for severe skin irritation in vivo, Danish QSAR Group at DTU Food.

1.2 Other related models

Leadscope Enterprise model for severe skin irritation in vivo, Danish QSAR Group at DTU Food.

MultiCASE CASE Ultra model for severe skin irritation in vivo, Danish QSAR Group at DTU Food.

1.3. Software coding the model

SciQSAR version 3.1.00.

2. General information
2.1 Date of QMRF
January 2015.
2.2 QMRF author(s) and contact details
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2.3 Date of QMRF update(s)
2.4 QMRF update(s)
2.5 Model developer(s) and contact details
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2.6 Date of model development and/or publication

January 2014.

2.7 Reference(s) to main scientific papers and/or software package

Contrera, J.F., Matthews, E.J., Kruhlak, N.L., and Benz, R.D. (2004) Estimating the safe starting dose in phase I clinical trials and no observed effect level based on QSAR modelling of the human maximum recommended daily dose. *Regulatory Toxicology and Pharmacology*, 40, 185 – 206.

SciQSAR (2009) Reference guide: *Statistical Analysis and Molecular Descriptors*. Included within the SciMatics SciQSAR software.

2.8 Availability of information about the model

The training set is non-proprietary and consists of data from the Registry of Toxic Effects of Chemical Substances (RTECS®) database and the Hazardous Substances Data Bank (HSDB) extracted from CHEMBANK (2000), data from Annex I of Directive 67/548/EEC and expert judgments of certain groups of chemicals. The model algorithm is proprietary from commercial software.

2.9 Availability of another QMRF for exactly the same model

3. Defining the endpoint

3.1 Species

Rabbit (for those chemicals that with experimental data, the remaining chemicals where classified based on experts judgment).

3.2 Endpoint

QMRF 4. Human Health Effects

QMRF 4. 4. Skin irritation /corrosion

3.3 Comment on endpoint

A chemical is characterised as a skin irritant if it produces a non-allergic, inflammatory reaction of the skin (i.e. oedema and erythema) at the site of contact on first exposure. The skin irritating chemical is acting directly on the skin cells and decreases the barrier function of the skin. The damage on the epidermal cells initiates a release of inflammatory mediators that attracts inflammatory cells and an inflammatory reaction in the skin can be seen.

In the training set for this model compounds causing no or mild skin irritation were categorised as mild skin irritants. Compounds causing severe skin irritation were categorised as severe skin irritants. The specific classification into mild and severe skin irritants from the four data sources differs and the criteria can be seen under 3.6.

3.4 Endpoint units

No units, 1 for positives and 0 for negatives.

3.5 Dependent variable

Skin irritation *in vivo*, severe or mild.

3.6 Experimental protocol

The training set includes data from RTECS ®, HSDB, Annex I of Directive 67/548/EEC plus expert judgments of certain groups of chemicals. Except for the chemicals classified based on expert judgement chemicals have been tested *in vivo*, according to the experimental protocol for the Draize test (RTECS 2012) or OECD guideline 404 (2002). Detailed descriptions of the experimental protocols can be found in the references.

701 of the training set chemicals were compiled from the RTECS® database (CHEM-BANK 2000). The chemicals in RTECS have been categorised in to three skin irritant categories (RTECS 2012): mild (RTECS code MLD), moderate (RTECS code MOD) and severe (RTECS code SEV). Of the 701 chemicals 291 were categorised as severe skin irritant based on two criteria: 1) they all have the RTECS code SEV, i.e. severe erythema (beet redness) to slight eschar formation (injuries in depth) and severe oedema (raised more

than 1 mm and extending beyond area of exposure), and 2) no requirements on dose or duration of exposure was made. The remaining 410 RTECS chemicals were categorised as mild skin irritants based on the following two criteria: 1) RTECS code MLD, i.e. well defined erythema and slight oedema (edges of area well defined by definite raising), and 2) a requirement of 500 mg and 24 hours exposure.

31 HSDB chemicals (CHEM-BANK 2000) were included in the training set and all were categorised as severe skin irritants, i.e. highly irritating or corrosive, according to HSDB criteria.

56 chemicals from Annex I of Directive 67/548/EEC (EU classifications 1967) were included in the training set as severe skin irritants. These chemicals have EU classifications R34 (causes burns) or R35 (causes severe burns).

The remaining 48 chemicals in the training set were classified in to mild (8) or severe (40) skin irritants based on expert judgments. This group of chemicals consisted of presumably not irritating chemicals that the model was otherwise confused by and where experimental data could not be found (i.e. set to be mild skin irritants), together with some well-known severe skin irritants.

3.7 Endpoint data quality and variability

As data originates from multiple sources and different criteria have been used for classification in to mild and severe irritants a certain degree of variability in data is expected.

4. Defining the algorithm

4.1 Type of model

This is a categorical (Q)SAR model based on calculated molecular descriptors, and if available the modeller's own or third-party descriptors or measured endpoints can be imported and used as descriptors.

4.2 Explicit algorithm

This is a categorical (Q)SAR model made by use of parametric discriminant analysis to create a linear discriminant function (see 4.5). The specific implementation is proprietary within the SciQSAR software.

4.3 Descriptors in the model

Molecular connectivity indices

Molecular shape indices

Topological indices

Electrotopological (Atom E and HE-States) indices

Electrotopological bond types indices

SciQSAR software provides over 400 built-in molecular descriptors. Additionally, SciQSAR makes it possible to import the modeller's own or third-party descriptors or use measured endpoints as custom descriptors.

4.4 Descriptor selection

The initial descriptor set is manually chosen by the model developer from the total set of built-in descriptors. Furthermore, the set of descriptors applied in the modelling by the program is on top of this selection determined by thresholds for descriptor variance and number of nonzero values likewise defined by the model developer.

57 descriptors were selected from the initial pool of descriptors by the system and used to build the model.

4.5 Algorithm and descriptor generation

For a binary classification problem SciQSAR uses discriminant analysis (DA) to make a (Q)SAR model. SciQSAR implements a broad range of discriminant analysis (DA) methods including parametric and non-parametric approaches. The classic parametric method of DA is applicable in the case of approximately

normal within-class distributions. The method generates either a linear discriminant function (the within-class covariance matrices are assumed to be equal) or a quadratic discriminant function (the within-class covariance matrices are assumed to be unequal). When the distribution is assumed to not follow a particular law or is assumed to be other than the multivariate normal distribution, non-parametric DA methods can be used to derive classification criteria. The non-parametric DA methods available within SciQSAR include the kernel and *k*-nearest-neighbor (kNN) methods. The main types of kernels implemented in SciQSAR include uniform, normal, Epanechnikov, bi-weight, or tri-weight kernels, which are used to estimate the group specific density at each observation. Either Mahalanobis or Euclidean distances can be used to determine proximity between compound-vectors in multidimensional descriptor space. When the kNN method is used, the Mahalanobis distances are based on the pooled covariance matrix. When the kernel method is used, the Mahalanobis distances are based on either the individual within-group covariance matrices or the pooled covariance matrix. (Contrera *et al.* 2004)

If the data outcome is continuous, regression analysis is used to build the predictive model. Within SciQSAR several regression methods are available: ordinary multiple regression (OMR), stepwise regression (SWR), all possible subsets regression (PSR), regression on principal components (PCR) and partial least squares regression (PLS). The choice of regression method depends on the number of independent variables and whether correlation or multicollinearity among the independent variables exists: OMR is acceptable with a small number of independent variables, which are not strongly correlated. SWR is used under the same circumstances as OMR but with greater number of variables. PSR is used for problems with a great number of independent variables. PCR and PLS are useful when a high correlation or multicollinearity exist among the independent variables. (SciQSAR 2009)

To test how stable the developed models are, SciQSAR have built-in cross-validation procedures (see 6.).

For this model, the kernel method was used.

4.6 Software name and version for descriptor generation

SciQSAR version 3.1.00.

4.7 Descriptors/chemicals ratio

In this model 57 descriptors were used. The training set consists of 836 compounds. The descriptor/chemical ratio is 1:14.7 (57:836).

5. Defining Applicability Domain

5.1 Description of the applicability domain of the model

The definition of the applicability domain consists of two components; the definition in SciQSAR and the inhouse further refinement algorithm on the output from SciQSAR to reach the final applicability domain call.

1. SciQSAR

The first criterion for a prediction to be within the models applicability domain is that all of the descriptor values for the test compound can be calculated by SciQSAR. If SciQSAR cannot calculate each descriptor value for the test chemical no prediction value is given by SciQSAR and it is considered outside the model's applicability domain.

2. The Danish QSAR group

The Danish QSAR group has applied a stricter definition of applicability domain for its SciQSAR models. In addition to the applicability domain definition made by SciQSAR a second criterion has been applied for predictions generated from (Q)SAR models with a binary endpoint. For each prediction SciQSAR calculates the probability (p) for the test compound's membership in one of the two outcome classes (positive or negative). The probability of membership in a class is a measure of how well training set knowledge is able to discriminate a positive prediction from a negative prediction within the nearest space of the subject compound-vector. The probability of membership value is also a measure of the degree of confidence of a prediction. The Danish QSAR group uses this probability for a prediction to further define the model's applicability domain. Only positive predictions with a probability equal to or greater than 0.7 and negative predictions with a probability equal to or less than 0.3 are accepted. Positive predictions with a probability between 0.5 and 0.7 as well as negative predictions with a probability between 0.3 and 0.5 are considered outside the model's applicability domain. When these predictions are wed out the accuracy of the model in general increases at the expense of reduced model coverage. Furthermore, as SciQSAR does not define a structural domain, only predictions which were within either Leadscope structural domain (defined as at least one training set chemical within a Tanimoto distance of 0.7) or CASE Ultra structural domain (no unknown fragments for negatives and maximum 1 unknown fragment for positives) were defined as being inside the SciQSAR applicability domain.

5.2 Method used to assess the applicability domain

The system does not generate predictions if it cannot calculate each descriptor value for the test compound.

Only positive predictions with probability equal to or greater than 0.7 and negative predictions with probability equal to or less than 0.3 were accepted.

5.3 Software name and version for applicability domain assessment

SciQSAR version 3.1.00.

5.4 Limits of applicability

The Danish QSAR group applies an overall definition of structures acceptable for QSAR processing which is applicable for all the in-house QSAR software, i.e. not only SciQSAR. According to this definition accepted structures are organic substances with an unambiguous structure, i.e. so-called discrete organics defined as: organic compounds with a defined two dimensional (2D) structure containing at least two carbon atoms, only certain atoms (H, Li, B, C, N, O, F, Na, Mg, Si, P, S, Cl, K, Ca, Br, and I), and not mixtures with two or more 'big components' when analyzed for ionic bonds (for a number of small known organic ions assumed not to affect toxicity the 'parent molecule' is accepted). Structures with less than two carbon atoms or containing atoms not in the list above (e.g. heavy metals) are rendered out as not acceptable for further QSAR processing. Calculation 2D structures (SMILES and/or SDF) are generated by stripping off accepted organic and inorganic ions. Thus, all the training set and prediction set chemicals are used in their non-ionized form. See 5.1 for further applicability domain definition.

6. Internal validation
6.1 Availability of the training set
Yes
6.2 Available information for the training set
CAS
SMILES
6.3 Data for each descriptor variable for the training set
No
6.4 Data for the dependent variable for the training set
All

6.5 Other information about the training set

836 compounds are in the training set: 418 positives and 418 negatives.

6.6 Pre-processing of data before modelling

The chemicals were classified in to mild or severe skin irritants based on the criteria described under 3.6.

Only structures acceptable for SciQSAR were used in the final training set. That is, only discrete organic chemicals as described in 5.4 were used. In case of replicate structures, one of the replicates was kept if all the compounds had the same activity and all were removed if they had different activity. No further structures accepted by the software were eliminated (i.e. outliers).

6.7 Statistics for goodness-of-fit

SciQSARs own internal performance test of the model gave the following Cooper's statistics for predictions within the applicability domain as defined by SciQSAR (i.e. the first criterion described in 5.1):

- Sensitivity (true positives / (true positives + false negatives)): 100%
- Specificity (true negatives / (true negatives + false positives)): 97.6%
- Concordance ((true positives + true negatives) / (true positives + true negatives + false positives + false negatives)): 98.8%

6.8 Robustness – Statistics obtained by leave-one-out cross-validation
Not performed.
6.9 Robustness – Statistics obtained by leave-many-out cross-validation
SciQSAR's own internal 10-fold cross-validation (10*10% out) procedure was used for predictions within the applicability domain as defined by SciQSAR (i.e. the first criterion described in 5.1). As the probability domain was not applied (i.e. the second criterion described in 5.2) the accuracy of the predictions when applying this domain can be expected to be higher than reflected in these cross-validation results. This gave the following Cooper's statistics:
 Sensitivity (true positives / (true positives + false negatives)): 77.3% Specificity (true negatives / (true negatives + false positives)): 71.3% Concordance ((true positives + true negatives) / (true positives + true negatives + false positives + false negatives)): 74.3%
6.10 Robustness - Statistics obtained by Y-scrambling
Not performed.
6.11 Robustness - Statistics obtained by bootstrap
Not performed.
6.12 Robustness - Statistics obtained by other methods Not performed.

- 7. External validation
- 7.1 Availability of the external validation set
- 7.2 Available information for the external validation set
- 7.3 Data for each descriptor variable for the external validation set
- 7.4 Data for the dependent variable for the external validation set
- 7.5 Other information about the training set
- 7.6 Experimental design of test set
- 7.7 Predictivity Statistics obtained by external validation
- 7.8 Predictivity Assessment of the external validation set
- 7.9 Comments on the external validation of the model

External validation has not been performed for this model.

8. Mechanistic interpretation

8.1 Mechanistic basis of the model

The SciQSAR software provides over 400 calculated physico—chemical, electrotopological E-state, connectivity and other molecular descriptors. The descriptors selected for the model may indicate modes of action that are obvious for persons with expert knowledge about the endpoint.

8.2 A priori or posteriori mechanistic interpretation

A posteriori mechanistic interpretation. The descriptors selected for the model may provide a basis for mechanistic interpretation.

8.3 Other information about the mechanistic interpretation

9. Miscellaneous information

9.1 Comments

The model can be applied to predict if a chemical causes severe or mild skin irritation in vivo.

9.2 Bibliography

EU classifications (1967) Directive 67/548/EEC- classification, packaging and labelling of dangerous substances. Annex I of Directive 67/548/EEC is available on: http://www.reach-compliance.eu/french/legislation/docs/launchers/launch-annex-1-67-548-EEC.html

CHEM-BANK (2000) Databanks of potentially hazardous chemicals: RTECS, OHMTADS, CHRIS, HSDB, IRIS, TSCA, NPG and ERG2000. USA. CHEM-BANKTM, CD-ROM, SilverPlatter International N.V., November 2000.

OECD guideline 404 (2002) OECD Guidelines for the Testing of Chemicals No. 404: Acute Dermal Irritation/Corrosion. Organisation for Economic Cooperation and Development; Paris, France. Available online at: http://www.oecd-ilibrary.org/environment/oecd-guidelines-for-the-testing-of-chemicals-section-4-health-effects 20745788.

RTECS (2012) Comprehensive guide to the RTECS. Registry of Toxic Effects of Chemical Substances 2012. Available on: http://ccinfoweb.ccohs.ca/rtecs/search.html

9.3 Supporting information