	QMRF identifier (JRC Inventory): **Q17-22b-0056**
	QMRF Title: **QSARINS model for hydroxyl-mediated tropospheric degradation using online descriptors**
	Printing Date: **Apr 16, 2018**

1.QSAR identifier

1.1.QSAR identifier (title):

QSARINS model for hydroxyl-mediated tropospheric degradation using
online descriptors

1.2.Other related models:

1.3.Software coding the model:

QSPR-Thesaurus

Online Platform of CADASTER project

itetko@vcclab.org

http://www.qspr-thesaurus.eu/home/show.do

MOBYDIGS

Software for multilinear regression analysis and variable subset selection by Genetic Algorithm, ver.
1.0 beta for windows, 2004

Todeschini Roberto, Talete srl, Milan (Italy)

http://www.talete.mi.it/

QSARINS

Software for the development, analysis and validation of QSAR MLR models, ver. 2.2, 2015

Paola Gramatica, email: paola.gramatica@uninsubria.it

http://www.qsar.it/

2.General information

2.1.Date of QMRF:

21/06/2011

2.2.QMRF author(s) and contact details:

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2.3.Date of QMRF update(s):

26/01/2015

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:

2011

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

[1]Roy PP, Kovarich S, Gramatica P (2011). QSAR Model Reproducibility and Applicability: A Case Study of Rate Constants of Hydroxyl Radical Reaction Models Applied to Polybrominated Diphenyl Ethers and (Benzo-)Triazoles, Journal of Computational Chemistry 32, 2386–2396 DOI: 10.1002/jcc.21820

[2]QSPR-Thesaurus http://www.qspr-thesaurus.eu/static/home.do

[3]MOBYDIGS Software for multilinear regression analysis and variable subset selection by Genetic Algorithm, ver. 1.0 beta for windows, 2004 Todeschini Roberto, Talete srl, Milano (Italy). http://www.talete.mi.it/

[4]QSARINS 2.2, 2015. Software for the development, analysis and validation of QSAR MLR models http://www.qsar.it/

2.8.Availability of information about the model:

Non-proprietary. Defined and available algorithm. Training and prediction sets are available in the Supporting Information of the related paper [ref 2; sect 9.2], in the attached sdf files in this QMRF(see Section 9.3) and in the QSARINS database [ref 7,8; sect 9.2].

2.9.Availability of another QMRF for exactly the same model:

None to date.

3.Defining the endpoint - OECD Principle 1

3.1.Species:

Not applicable

3.2.Endpoint:

2.Environmental fate parameters 2.2.b.Persistence: Abiotic degradation in air (Phototransformation). Indirect photolysis (OH-radical reaction, ozone-radical reaction, other)

3.3.Comment on endpoint:

Gas-phase reaction between photochemically produced hydroxyl radicals and organic chemicals at 25 °C and 1 atm for 460 heterogeneous organic chemicals [ref.3; sect.9.2]. The units of the rate coefficient depend on the global order of reaction.

3.4.Endpoint units:

$cm^3 s^{-1} molecule^{-1}$

3.5.Dependent variable:

-log(OH)

3.6.Experimental protocol:

Available at Atkinson, R. J Phys Ref Data 1989, Monograph 1,p1-246. [3]

3.7.Endpoint data quality and variability:

Satisfactory models were also obtained in the past using the same dataset as well as AOPWIN package of EPI Suite have the same training set. The dataset is the famous and widely used Atkinson (1989, [3]) set related to atmospheric reactivity.

4.Defining the algorithm - OECD Principle 2

4.1.Type of model:

QSAR

4.2.Explicit algorithm:

Multiple linear regression QSAR (OLS-Ordinary Least Square)

GA-OLS

The modeling descriptors are: HOMO(Highest occupied molecular orbital energy), D_PathSum(F, rel) (related to the presence of F atoms), G_([Cl, Br, I]) (related to the presence of Cl, Br, I) and SeaC2C2aa (Sum of the bond electro topological values of carbon–carbon aromatic bonds in which the carbons are not substituted). See section 4.3 for a more detailed description of the four modeling descriptors.

Split models

Models were developed from three different training set of 191, 230, 230 compounds respectively based on structural similarity analysis (K-ANN, K-means) and random by sorting the response.

K-ANN

$\log(OH)=3.95(\pm0.68)-0.67(\pm0.07)HOMO+1.42(\pm0.25)D\_PathSum(Fe, rel)+0.06(\pm0.01) SeaC2C2aa+0.41(\pm0.09)G\_([Cl, Br, I])$

Random

$-\log(OH)=3.84(\pm0.69)-0.69(\pm0.07)HOMO+1.30(\pm0.26)D\_PathSum(Fe, rel)+0.48(\pm0.10)G\_([Cl, Br, I])+0.06(\pm0.01) SeaC2C2aa$

K means

$-\log(OH)=3.47(\pm0.70)-0.73(\pm0.07)HOMO+1.23(\pm0.23)D\_PathSum(Fe, rel)+0.07(\pm0.01) SeaC2C2aa +0.44(\pm0.12)G\_([Cl, Br, I])$

Full

Model developed on all available experimental data (training set of 460 compounds)-$\log(OH)=3.83(\pm0.48)-0.69(\pm0.05)HOMO+1.26(\pm0.17)D\_PathSum(F, rel)+0.43(\pm0.07)G\_([Cl, Br, I]) +0.06(\pm0.01) SeaC2C2aa$

4.3.Descriptors in the model:

[1]HOMO dimensionless Highest occupied molecular orbital energy. This descriptor characterizes the susceptibility of a molecule toward the attack by the electrophile OH radical, more reactive chemicals having higher HOMO energy

[2]D_PathSum(F, rel) dimensionless AMBIT Fragment [3], positively correlated to the response, is related to the presence of F atoms in the molecule.

[3]G_([Cl, Br, I]) dimensionless AMBIT Fragment [3], positively correlated to the response, is related to the presence of some halogen atoms (Cl, Br, I) in the molecule.

[4]SeaC2C2aa dimensionless E-state index (Sum of the bond electro topological values of carbon–carbon aromatic bonds in which the carbons are not substituted) [4], inversely correlated with the modeled response. The chemicals with higher number of hydrogen atoms can be more attacked by the hydroxyl radical and are, for this reason, more reactive

4.4.Descriptor selection:

In this study different 2D-descriptors (E-state, ALogPS, Molprint fragment, AMBIT Descriptors, GSFragment, ISIDA fragments etc) available at

CADASTER web (http://www.qspr-thesaurus.eu/static/home.do) were calculated, and were pruned by deleting descriptors with less than 2 unique values as well as a correlation of 0.95. In addition we added ETA descriptors [ref.6; sect.9.2], obtaining a large pool of 1023 input descriptors.

Furthermore three quantum-chemical descriptors (Highest Occupied Molecular Orbital (HOMO) and Lowest Unoccupied Molecular Orbital (LUMO) energies, HOMO-LUMO gap were added to above pool of descriptors. Therefore input sets of 1026 (Online and MOPAC) descriptors underwent the subsequent selection for the best modeling variables. The Genetic Algorithm-Variable Subset Selection (GA-VSS), by Ordinary Least Squares regression (OLS), included in MOBYDIGS (and now reproduced in QSARINS [ref 7,8; sect 9.2]), was applied to select only the best combination of descriptors from the input pool: 4 descriptors selected from 1026.

4.5. Algorithm and descriptor generation:

Multiple linear regression (MLR) and variable selection by GA-VSS were performed by Ordinary Least Squares regression (OLS) in order to develop the model. The Genetic Algorithm-Variable Subset Selection (GA-VSS), included in MobyDigs (and verified with the one included in QSARINS [ref 12,13; sect 9.2]), was applied to select only the best combination of descriptors from input pool. Descriptors were generated according to the appropriate uploading format available on CADASTER web. Quantum chemical descriptors were calculated by the semi empirical molecular orbital program MOPAC (AM1 method for energy minimization) in the software HYPERCHEM version 7.03.

4.6. Software name and version for descriptor generation:

QSPR-Thesaurus

Online Platform of CADASTER project. Descriptors generated from SMILES, available in QSARINS (QSARINS-Chem). This enables an end user to regenerate the descriptors for a new compound.

itetko@vcclab.org

http://www.qspr-thesaurus.eu/static/home.do

4.7. Chemicals/Descriptors ratio:

Split Model:

47.75 (191 chemicals / 4 descriptors)

57.50 (230 chemicals / 4 descriptors)

65 (230 chemicals / 4 descriptors)

Full model: 115 (460 chemicals / 4 descriptors)

5. Defining the applicability domain - OECD Principle 3

5.1. Description of the applicability domain of the model:

Quantitative measures of a model applicability domain (AD) are needed to evaluate the degree of extrapolation and for the identification of problematic compounds.

Response and descriptor space

Range of experimental –log(OH) values: 9.44 - 15.7

Range of descriptors values:

HOMO: (-)7.3- (-)13.68

D_PathSum(F, rel): 0-1.667

G_([Cl, Br, I]): 0-4

SeaC2C2aa:0-24.03211

The chemical space of the model includes alkanes, alkenes, alcohols, halogenated chemicals, amines, aromatics, and other functional groups.

5.2. Method used to assess the applicability domain:

AD was verified by the leverage approach [9] (for the structural domain), and by the identification of response outliers (compounds with cross-validated standardized residuals greater than 2.5 standard deviation units).

Graphically, the plot of hat values (h) versus standardized residuals, i.e. the Williams plot, verified the presence of response outliers and training set chemicals that are structurally very influential in determining model parameters (compounds with leverage value (h) greater than 3p/n (h*), where p is the number of the model variables plus one, and n is the number of the objects used to calculate the model). For our model h* is equal to 0.033 (number of variables in the model are four and total number of compounds is 460)

For new compounds without experimental data, leverage can be used as a quantitative measure for evaluating the degree of extrapolation: for compounds with a high leverage value (h > h*), that are structural outliers, predictions should be considered less reliable. In QSARINS the Insubria graph allows chemicals to be identified for which the predictions are inter- or extrapolated by the model.

5.3. Software name and version for applicability domain assessment:

QSARINS 1.0 (verified also on version 2.2)

Software for the development, analysis and validation of QSAR MLR models, ver. 2.2, 2015

Paola Gramatica, email: paola.gramatica@uninsubria.it

http://www.qsar.it/

5.4. Limits of applicability:

Some common compounds have been found as outliers or influential in all the models:

Outliers for response (standardized residuals>2.5 standard deviation units):

Overestimated: triethyl phosphate (61) and 2-(chloromethyl)-3-chloro-1-propene (403)

Underestimated: bromomethane (18), dimethylsulfide (37), diethyl sulfide (263), ethyl methyl sulfide (353), 3-methyl-1,2 butadiene (342).

Outliers for structure (Hat cut off=0.033):

fluorinated chemicals: 1,1,2,2-tetrachloroethene (232), 1,1-dichloro-2,2,2-trifluoroethane (262), 1,1,1,2,2-pentafluoroethane (265), hexafluorobenzene (267), 1-chloro-1,2,2,2-tetrafluoroethane (414) and propylpentafluorobenzene (457)

6.Internal validation - OECD Principle 4

6.1.Availability of the training set:

Yes

6.2.Available information for the training set:

CAS RN: Yes

Chemical Name: Yes

Smiles: Yes

Formula: Yes

INChI: No

MOL file: No

6.3.Data for each descriptor variable for the training set:

All

6.4.Data for the dependent variable for the training set:

All

6.5.Other information about the training set:

Three different splitting procedures were adopted, two based on structural similarity analysis (K-ANN, K-means) and one random by sorting the response, in order to propose models that have a demonstrated high performance in predicting external chemicals of different typology, avoiding the bias derived from an unique split. The number of training set in three divisions (K-ANN, Random, K-Means) are 191, 230 and 230 respectively.

6.6.Pre-processing of data before modelling:

Transformation to logarithmic units and multiplied by -1 to obtain positive values

6.7.Statistics for goodness-of-fit:

Here we have three different training set for input set of descriptors. Therefore we are reporting the statistical fittings of all the models.

$n_{Training}$=191, R^2=0.847, Ra^2= 0.844, s=0.39, F=257.10

$n_{Training}$=230, R^2=0.814, Ra^2= 0.810, s=0.44, F=245.49

$n_{Training}$=230, R^2=0.813, Ra^2= 0.810, s=0.47, F = 244.49

6.8.Robustness - Statistics obtained by leave-one-out cross-validation:

$n_{Training}$=191, Q^2_{LOO}=0.834; $n_{Training}$=230, Q^2_{LOO}=0.803; $n_{Training}$=230, Q^2_{LOO}=0.803

High value of Q^2_{LOO} (leave-one-out) means that the models, when verified for this technique of internal validation, are robust.

6.9.Robustness - Statistics obtained by leave-many-out cross-validation:

$Q^2_{LMO}$was not calculated, since we calculated Q^2_{BOOT}(see 6.11).

6.10.Robustness - Statistics obtained by Y-scrambling:

R^2_{ys}= 0.018-0.020, Q^2_{ys}=0.011-0.019

(Values are in range for three splitting). The low values of Y-scrambled $R^2$and Q^2mean that the proposed models are not given by chance.

6.11.Robustness - Statistics obtained by bootstrap:

Split Models

$n_{Training}$=191, Q^2_{BOOT}=0.822; $n_{Training}$=230, Q^2_{BOOT}=0.795; $n_{Training}$=230, Q^2_{BOOT}=0.795

Full Model

Q^2_{BOOT}=0.797.The high value of Q^2_{BOOT} means that the models are robust and stable.

6.12.Robustness - Statistics obtained by other methods:

No information available

7.External validation - OECD Principle 4

7.1.Availability of the external validation set:

Yes

7.2.Available information for the external validation set:

CAS RN: Yes

Chemical Name: Yes

Smiles: Yes

Formula: Yes

INChI: No

MOL file: No

7.3.Data for each descriptor variable for the external validation set:

All

7.4.Data for the dependent variable for the external validation set:

All

7.5.Other information about the external validation set:

We have distributed our dataset into training and prediction set using three different splitting procedures. One based on response and two are based on structural similarity analysis confirming well balance in the training and prediction set both in response and structure. The number of external validation set in three divisions (K-ANN, Random, K-means) are 269, 230 and 230 respectively.

7.6.Experimental design of test set:

The random by response splitting was obtained by ordering the chemicals according to their descending kinetic constant value, and then putting the most and the least reactive in the training set and one out of every two chemicals in the prediction set (50% of the full dataset). This splitting guarantees that the prediction set spans the entire range of the experimental measurements and is numerically representative of the dataset.

The splitting of the data set realized by Kohonen Artificial Neural Network (K-ANN) takes advantage of the clustering capabilities of K-ANN, allowing the selection of a structurally meaningful training set and a representative prediction set[ref 10; sect 9.2].

Another approach for splitting into training and prediction sets is by using K-means clustering which ensures that the similarity principle can be employed for grouping chemicals and splitting them in balanced

training and prediction sets [ref 11; sect 9.2].

7.7. Predictivity - Statistics obtained by external validation:

$n_{Prediction}$=269, Q^2_{F1}[ref 12; sect 9.2]=0.778,
Q^2_{F2}[ref 13; sect 9.2]=0.775, Q^2_{F3}[ref
14; sect 9.2]=0.745, RMSE=0.49, CCC [ref 15,16; sect 9.2]=0.876

$n_{Prediction}$=230, Q^2_{F1}[ref 12; sect
9.2]=0.796, Q^2_{F2}[ref 13; sect 9.2]=0.795, Q^2_{F3}[ref
14; sect 9.2]=0.786, RMSE=0.47, CCC[ref 15,16; sect 9.2]=0.891$n_{Prediction}$=230, Q^2_{F1}[ref 12; sect

9.2]=0.795, Q^2_{F2}[ref 13; sect 9.2]=0.793, Q^2_{F3}[ref
14; sect 9.2]=0.829, RMSE=0.44, CCC [ref 15,16; sect 9.2]=0.892

The high values of external Q^2, calculated in different ways
(see references for more details), and CCC show that the proposed models
are predictive for new chemicals. In fact, the models show good results
when applied to the chemicals not used during the model development
(chemicals in the prediction sets).

7.8. Predictivity - Assessment of the external validation set:

The response range value of training sets for three splitting is
[9.44-15.7] and the prediction set response range are [9.5-14.77],
[9.6-14.77], [9.6-14.6] respectively for K-ANN, Random and K-means
clustering procedure.
Therefore, the three splittings guarantee a balanced distribution of
chemicals in training and prediction sets regarding the response.
HOMO:
Training set
K-ANN [(-)8.12- (-)13.31]
Random [(-)8.12- (-)13.31]
K-means [(-)8.19- (-)13.31]

Prediction set
[(-)7.3- (-)13.68] for all three splits

D_PathSum(F,rel):

Training set[0- 1.666667] for all three splits
Prediction set[0- 1.595238] for all three splits

G_([Cl,Br,I]):
Training setK-ANN [0- 4]Random [0- 3]K-means [0- 3]

Prediction set[0-4] for all three splits

SeaC2C2aaTraining set[0- 24.03211] for all three splitsPrediction setK-ANN [0- 23.89256]Random [0- 20.36159]K-means [0- 21.21273]. Therefore, the three splittings guarantee a balanced distribution of
chemicals in training and prediction sets regarding the structure.The prediction sets are all large and

representative of the training
sets, therefore the models can be reliably applied to the external sets.

7.9.Comments on the external validation of the model:

Models selected by GA from three different splitting procedures (K-ANN, K-means, random) demonstrated high performance in predicting external chemicals of different typology avoiding the bias derived from unique split.

8.Providing a mechanistic interpretation - OECD Principle 5

8.1.Mechanistic basis of the model:

The model was developed by statistical approach. No mechanistic basis was defined a priori.

8.2.A priori or a posteriori mechanistic interpretation:

A posteriori mechanistic interpretation.

The descriptor combination appeared from the input of online pool of descriptors is HOMO, SeaC2C2aa, D_pathSum(F, rel), D_pathSum(F, rel). Highest occupied molecular orbital (HOMO) energy. This descriptor characterizes the susceptibility of a molecule toward the attack by the electrophile OH radical, more reactive chemicals having higher HOMO energy.The E-state index SeaC2C2aa (Std coeff. = 0.266) is the sum of the bond electro topological values of carbon–carbon aromatic bonds in which the carbons are not substituted. This descriptor is inversely correlated with the modeled response in the univariate model. The chemicals with higher number of hydrogen atoms can be more attacked by the hydroxyl radical and are, for this reason, more reactive. The remaining two descriptors D_pathSum(F, rel) and G_(Cl, I, Br), both positively correlated to the response, are the AMBIT descriptors and are counts of the number of halogen atoms in the molecules. Molecules with more halogen atoms tend to have less reactivity.

8.3.Other information about the mechanistic interpretation:

No information available

9.Miscellaneous information

9.1.Comments:

The model is transparent in its reproducibility by the above mentioned descriptor freely available online. In order to predict chemicals without experimental activity it is suggested to use the full model developed from all available (n=460) chemicals with wider domain of applicability.

The statistical quality of the full model
n=460, R^2=0.806, Q^2_{LOO}=0.801, Q^2_{BOOT}=0.797, RMSEtr=0.45, RMSECV=0.45

9.2.Bibliography:

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9.3.Supporting information:

10.Summary (JRC QSAR Model Database)

10.1.QMRF number:

Q17-22b-0056

10.2.Publication date:

2017-09-27

10.3.Keywords:

QSPR-Thesaurus;hydroxyl;tropospheric degradation;QSARINS;INSUBRIA;

10.4.Comments:

old# Q47-19-49-479