

	<b>QMRF identifier (JRC Inventory): Q17-301C-0054</b>
	<b>QMRF Title: VEGA Ready Biodegradation model</b>
	<b>Printing Date: Dec 11, 2019</b>

## 1. QSAR identifier

### 1.1. QSAR identifier (title):

VEGA Ready Biodegradation model

### 1.2. Other related models:

No related models

### 1.3. Software coding the model:

Ready Biodegradation model v1.0.9

The model is based on the OECD 301C Modified MITI (I) Test data and provides a qualitative evaluation (binary classification) of ready biodegradability properties.

<http://www.vega-qsar.eu/>

## 2. General information

### 2.1. Date of QMRF:

11 May 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:

The model was published in 2014 (see 2.7).

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

Lombardo A, Pizzo F, Benfenati E, Manganaro A, Ferrari T, Gini G (2014). A new in silico classification model for ready biodegradability, based on molecular fragments. Chemosphere. 108, 10–16 <http://dx.doi.org/10.1016/j.chemosphere.2014.02.073>

### 2.8. Availability of information about the model:

Freely available through the VEGA web site ([www.vega-qsar.eu](http://www.vega-qsar.eu)) as .pdf file describing the parameters. The training, test and validation sets

are also available (see 9.3).

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

Other QMRF for this model are not available

## **3.Defining the endpoint - OECD Principle 1**

### **3.1.Species:**

Activated sludge (according to OECD 301C Modified MITI (I) Test).

### **3.2.Endpoint:**

2.3.a.Persistence: Biodegradation. Ready/not ready biodegradability 301CModified MITI (I) Test

### **3.3.Comment on endpoint:**

Ready biodegradability tests are screening tests in which a high concentration of the test substance is used and ultimate biodegradation is measured by non-specific parameters under aerobic conditions. In the OECD 301C Modified MITI (I) test, a substance can be considered readily biodegradable if 60% of the substance degrades in 28 days (in terms of ThOD).

### **3.4.Endpoint units:**

Adimensional.

### **3.5.Dependent variable:**

Structural alerts associated to: Readily Biodegradable, possible Readily Biodegradable, NON Readily Biodegradable and possible NON Readily Biodegradable.

### **3.6.Experimental protocol:**

OECD 301C Modified MITI (I) Test. It measures the oxygen uptake in a period of 28 days of a solution of the substance inoculated with activated sludge under aerobic conditions.

### **3.7.Endpoint data quality and variability:**

For the development and testing of the model the datasets were extracted from the OECD QSAR toolbox v 2.0 and from the BIOWIN 5 and 6 models (in EPISuite<sup>TM</sup>) and then combined (see ref. 1 and 2, section 9.2). To validate the model data were extracted from Cheng et al., 2012 (see ref 3, section 9.2). 17 compounds were found with non-concordant data among the OECD QSAR toolbox v 2.0 and the BIOWIN 5 and 6 models (i.e. not classified in the same class: Readily Biodegradable or NON-Readily Biodegradable) and were excluded. All the common data with Cheng et al., 2012 were eliminated and not checked to verify non-concordant data because the model was used only as external validation set.

## **4.Defining the algorithm - OECD Principle 2**

### **4.1.Type of model:**

SAR built using both statistically-related fragments extracted using SARpy (see ref 4, section 9.2) and expert-based fragments.

### **4.2.Explicit algorithm:**

see ref 1, section 9.2

The classification scheme is based on a flow chart (see ref. 1, section 9.2). Basically if at least one

fragment linked with non-ready biodegradability and with high specificity is found, the chemical is classified as NON Readily Biodegradable, otherwise, if no highly specific fragments are found but at least one fragment linked with non ready biodegradability and lower specificity is found, the compound is classified as possible NON Readily Biodegradable. If no fragments linked with non ready biodegradability are found but fragments linked with ready biodegradability are found, the compound is classified as Readily Biodegradable (if highly specific fragments are found) or possible Readily Biodegradable (if only lower specific fragments are found). If no fragments are found, the compound is classified as unknown.

#### 4.3.Descriptors in the model:

- [1]1,2-dichlorobenzene c1ccc(c(c1)Cl)Cl Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [2]1-ethyl-3-methylbenzene c1cc(cc(c1)CC)C Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [3]1,2-dichloroethane C(C(Cl))Cl Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [4]2-chloroaniline Nc1ccc(cc1Cl) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [5]diphenylmethanone c1ccccc1C(=O)c2ccccc2 Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [6]dimethoxyphosphinic acid O=P(OC)(OC)O Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [7]cyclohex-4-ene-1,2-dicarbaldehyde O=CC1CC=CCC1C(=O) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [8]benzene-1,3-diamine Nc1ccc(c(N)c1) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [9]1-bromopropane CCCBBr Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [10]3-chlorophenol Oc1ccc(c(c1)Cl) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [11]fluorine F Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [12](1-phenylethyl)benzene c1ccc(cc1)C(c2ccccc2)C Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [13]methoxy(sulfanylidene)phosphinous acid P(=S)(OC)O Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [14]N-phenylaniline c1ccc(cc1)Nc2ccc(cc2) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [15]naphthalen-1-amine c1ccc2c(c1)cccc2N Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [16]1-methylnaphthalene Cc1cccc2ccccc12 Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [17]benzyltrimethylamine C(c1cccc(c1))N(C)C Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [18]2-methylnonane CCCCCCC(C)C Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability
- [19]1,3-benzothiazole c1nc2ccccc2s1 Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[20]tin [Sn] Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[21]methanimine C=N Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[22]1,4-diethylbenzene c1cc(ccc1C(C))C(C) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[23]propoxybenzene CCCOc1ccccc1 Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[24]2-ethyl-1-methoxyhexane O(C)CC(CC)CCCC Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[25]2-chlorobenzaldehyde O=Cc1c(cccc1Cl) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[26]1-ethyl-2-methylbenzene c1ccc(c(c1)CC)C Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[27]3-chloroaniline Nc1ccc(c(c1)Cl) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[28](2-methoxyethyl)(propyl)amine CCCNCCOC Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[29]2,4-dimethylpent-1-ene C=C(C)CC(C)(C) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[30]bromobenzene c1ccc(cc1)Br Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[31]methylcarbamic acid O=C(O)NC Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[32]1,1,1-trichloroethane CC(Cl)(Cl)Cl Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[33]chloroethene C(=CCl) Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[34]dithioperoxol SS Fragment in ruleset 1: highly specific fragments linked with non-ready biodegradability

[35]benzylbenzene C(c1ccccc1)c2ccccc2 Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[36]1-chloro-2-methylbenzene c1ccc(c(c1)C)Cl Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[37]naphthalene c1cccc2ccccc12 Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[38]bromine Br Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[39]3-methylaniline Nc1ccc(c(c1)C) Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[40]hydroxylamine ON Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[41]chlorobenzene c1ccc(c(c1))Cl Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[42]benzenesulfonic acid c1ccc(cc1)S(=O)(=O)O Fragment in ruleset 2: lower specific fragments linked with non-ready biodegradability

[43]pentan-2-amine C(N)(C)CC(C) Fragment in ruleset 2: lower specific fragments linked with non-

ready biodegradability

[44]2-hydroxyethyl acetate O=C(OCCO)C Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[45]propanedial O=CCC(=O) Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[46]N-(2-hydroxyethyl)formamide O=C(N(CCO)) Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[47]1-propoxynonane O(CCCCCCCCC)CCC Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[48]2,6-dimethylhepta-1,5-diene CC(=C)CCC=C(C)C Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[49]2-methoxybutane CCC(OC)C Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[50](dodecyloxy)phosphonous acid P(O)(O)OCCCCCCCCCCCC Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[51]benzyl formate O=C(OCc1ccccc1) Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[52]6-oxohexanoic acid O=C(O)CCCC(=O) Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[53]docosane CCCCCCCCCCCCCCCCCCCC Fragment in ruleset 3: highly specific fragments linked with ready biodegradability

[54]ridecan-1-ol OCCCCCCCCCCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[55]butan-2-one O=C(C)CC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[56]3-methoxyprop-1-ene C(OC)C=C Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[57]2-methylhept-2-ene C(C)CCC=C(C)C Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[58]methyl propanoate O=C(OC)CC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[59]butyl formate C(=O)OCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[60]1-ethoxybutane CCOCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[61]octan-1-ol OCCCCCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[62]tridecane CCCCCCCCCCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[63]propanoic acid CCC(=O)O Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[64]benzoic acid O=C(O)c1ccc(cc1) Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[65]butan-1-ol OCCCC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[66]acetamide O=C(N)C Fragment in ruleset 4: lower specific fragments linked with ready

biodegradability

[67]acetaldehyde O=CC Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[68]ethane-1,2-diol OCCO Fragment in ruleset 4: lower specific fragments linked with ready biodegradability

[69]propan-1-amine NCCC Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[70]sulfanone S(=O) Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[71]heptanes CCCCCC Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[72]anisole O(c1ccccc1)C Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[73]butane CCCC Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[74](chloromethyl)benzene C1(ccccc1)C(Cl) Fragment in ruleset 5: lower specific fragments linked with ready biodegradability extracted from unknown

[75]diazene N=N Fragment in ruleset 6: expert-based fragments linked with non-ready biodegradability

[76]halogenated ring structure [R][Cl,F,Br,I] Fragment in ruleset 6: expert-based fragments linked with non-ready biodegradability

[77]carbonyl bound to aromatic structure [a][C;D2]=O Fragment in ruleset 7: expert-based fragments linked with ready biodegradability

[78]formonitrile C#N Fragment in ruleset 7: expert-based fragments linked with ready biodegradability

#### 4.4.Descriptor selection:

Full details are explained in the literature (see ref. 1 in section 9.2). Briefly, SARpy was run on the training set to extract rulesets 1-4. All the rules were checked removing all the fragments that had a likelihood ratio  $< 2$  and/or a true positive rate (TP)  $< 70\%$ . The rules with TP = 100% were considered highly specific, the other balanced. SARpy was run again on the compounds of the training set that could not be predicted using rulesets 1-4. 2 rulesets were extracted (one linked with readily and one with non-readily biodegradable compounds). After the check (performed with the same rules explained above) only one ruleset remained, ruleset 5. The compounds of the training set were checked and 2 new rulesets were extracted (one linked with readily and one with non-readily biodegradable compounds): rulesets 6 and 7. These last fragments were all confirmed with studies from the literature.

#### 4.5.Algorithm and descriptor generation:

Fragments both statistically or expert based. Details provided in section 4.3

#### 4.6.Software name and version for descriptor generation:

SARpy

A general software that automatically extracts knowledge from a dataset and detects the molecular structural fragments associated with the activity of interest.

#### 4.7.Chemicals/Descriptors ratio:

582 chemicals (training set)/78 fragments = 7.5

### 5.Defining the applicability domain - OECD Principle 3

#### 5.1.Description of the applicability domain of the model:

The model is applicable to heterogeneous chemicals. In the implementation of the model several pieces of information are given to evaluate if a prediction is reliable (chemical falling in the Applicability Domain or not). The information about the Applicability Domain (AD) is combined into a unique index called Global Applicability Domain Index (ADI). ADI values range between 0 and 1. ADI >0.8 means that the compound is in the AD, ADI < 0.65 means that the compound is out of the AD, a value between 0.65 and 0.8 means that the compound is possibly out of the AD.

#### 5.2.Method used to assess the applicability domain:

The Applicability Domain of the model is defined by considering several parameters as described below:

1. Similar molecules with known experimental value. This index takes into account how similar are the first three most similar compounds found. Values near 1 mean that the predicted compound is well represented in the dataset used to build the model. Defined intervals are:  $1 \geq \text{index} > 0.85$  strongly similar compounds with known experimental value in the training set have been found;  $0.85 \geq \text{index} > 0.7$  only moderately similar compounds with known experimental value in the training set have been found;  $\text{index} \leq 0.7$  no similar compounds with known experimental value in the training set have been found.
2. Accuracy of prediction for similar molecules. This index takes into account the error in prediction for the three most similar compounds. Values near 0 mean that the predicted compounds falls in an area of the model's space where the model gives reliable predictions, otherwise the greater is the value, the worse the model behaves. Defined intervals are:  $1 \geq \text{index} > 0.8$  accuracy of prediction for similar molecules found in the training set is good;  $0.8 \geq \text{index} > 0.5$  accuracy of prediction for similar molecules found in the training set is not optimal;  $\text{index} \leq 0.5$  accuracy of prediction for similar molecules found in the training set is not adequate.
3. Concordance for similar molecules. This index takes into account the difference between the predicted value and the experimental values of the three most similar compounds. Values near 0 mean that the prediction made disagrees with the values found in the model's space, thus the prediction could be unreliable. Defined intervals are:  $1 \geq \text{index} > 0.8$  similar molecules found in the training set have experimental values that agree with the predicted; value  $0.8 \geq \text{index} > 0.5$  some similar molecules found in the training set have experimental values that disagree with the predicted value;  $\text{index} \leq 0.5$  similar molecules found in the training set have experimental values that

disagree with the predicted value.

4. Atom Centered Fragments similarity check. This index takes into account the presence of one or more fragments that are not found in the training set, or that are rare fragments. First order atom centered fragments from all molecules in the training set are calculated, then compared with the first order atom centered fragments from the predicted compound; then the index is calculated as following: a first index RARE takes into account rare fragments (those who occur less than three times in the training set), having value of 1 if no such fragments are found, 0.85 if up to 2 fragments are found, 0.7 if more than 2 fragments are found; a second index NOTFOUND takes into account not found fragments, having value of 1 if no such fragments are found, 0.6 if a fragments is found, 0.4 if more than 1 fragment is found. Then, the final index is given as the product  $RARE * NOTFOUND$ . Defined intervals are: index = 1 all atom centered fragment of the compound have been found in the compounds of the training set;  $1 > index \geq 0.7$  some atom centered fragment of the compound have not been found in the compounds of the training set or are rare fragments;  $index < 0.7$  a prominent number of atom centered fragments of the compound have not been found in the compounds of the training set or are rare fragments

#### **5.3. Software name and version for applicability domain assessment:**

VEGA Ready Biodegradation model version 1.0.9

This is included in the stand alone version of VEGA: VEGANic v 1.0.9

<http://www.vega-qsar.eu/>

#### **5.4. Limits of applicability:**

The model was built only for organic chemical. The user should verify the ADI (see section 5.1 and 5.2).

### **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: No

Smiles: Yes

Formula: No

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:**

For some chemicals of the training, test and validation sets are available continuous values (Biological Oxygen Demand, BOD%) and binary values (readily or non-readily biodegradable), for other only the binary



classification. The model is based on the binary values.

#### **6.6.Pre-processing of data before modelling:**

Continuous values are converted in binary values: if BOD (28 d) 60% then the compound was classified as readily biodegradable, otherwise non-readily biodegradable. If BOD is available for period lower than 28d, only BOD values 60% were considered and the compound were classified as readily biodegradable.

When multiple values were available, the compounds with values not in agreement were deleted; otherwise the mean value was used to classify the compound.

All the chemical structures were manually checked deleting doubtful compounds, mixture, inorganic compounds and tautomers.

#### **6.7.Statistics for goodness-of-fit:**

The readily biodegradable compounds were considered as active.

Accuracy = 92.2%

Sensitivity = 94.8%

Specificity = 89.8%

MCC = 0.85

FP rate = 0.10

FN rate = 0.05

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

No leave-one-out cross-validation obtained

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

No leave-many-out cross-validation obtained

#### **6.10.Robustness - Statistics obtained by Y-scrambling:**

No Y-scrambling statistics obtained

#### **6.11.Robustness - Statistics obtained by bootstrap:**

No bootstrap statistics obtained

#### **6.12.Robustness - Statistics obtained by other methods:**

No statistics obtained with other methods obtained

### **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: No

Smiles: Yes

Formula: No

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:**

To test the model two datasets were used: a test set of 146 compounds, obtained from the same source of the training set (1 and 2, section 9.2), and an external set of 874 new compounds obtained from 1 and 3, section 9.2. This second, large set was available after the development of the model. All the data were pre-processed as explained in section 6.6.

### **7.6. Experimental design of test set:**

The dataset were randomly split into training and test set with respectively the 80% and the 20% of the compounds.

### **7.7. Predictivity - Statistics obtained by external validation:**

The readily biodegradable compounds were considered as active.

Test set:

Accuracy = 81.7%

Sensitivity = 87.3%

Specificity = 76.9%

MCC = 0.64

FP rate = 0.23

FN rate = 0.13

External set:

Accuracy = 76%

Sensitivity = 73.1%

Specificity = 83.6%

MCC = 0.51

FP rate = 0.27 FN rate = 0.16 External set in AD:

Accuracy = 80.7% Sensitivity = 75.6%

Specificity = 90.7%

MCC = 0.63 FP rate = 0.24

FN rate = 0.09

### **7.8. Predictivity - Assessment of the external validation set:**

MCC values always > 50% and the other statistics prove that the model is predictive. The use of the applicability domain improves the results.

Test set: 119 compounds (81.5%) have an Atom Centered Fragment evaluation of 1 (= all atom centered fragment of the compound have been found in the compounds of the training set).

External set: 617 compounds (70.1%) have an Atom Centered Fragment evaluation of 1 (= all atom centered fragment of the compound have been found in the compounds of the training set).

### **7.9. Comments on the external validation of the model:**

The test set is balanced (52.7% NON Readily Biodegradable and 47.3% Readily Biodegradable). The external set is not balanced (72.5% NON Readily Biodegradable and 27.5% Readily Biodegradable). In both conditions the model is predictive.

## **8. Providing a mechanistic interpretation - OECD Principle 5**

### **8.1. Mechanistic basis of the model:**

The model is based on structural alerts. The majority of them are statistically based, and some have mechanistic explanation:

- Azo group. The initial step of the biodegradation of azo dyes is cleavage of the azo group. This reaction is catalysed by the enzyme azoreductase, which is inhibited by molecular oxygen and the Ready Biodegradability tests are conducted under aerobic condition. For this reason this structural alert is considered associated to chemicals NON Readily Biodegradable.
- All compounds containing halogen atoms. The presence of halogenated organic compounds in the environment is recent, so the enzymes that have evolved to metabolize these compounds are considered to be in a relatively early stage of development. The initial conversion of non-toxic compounds yields toxic products (e.g. the monooxygenase-catalysed oxidations of xenobiotics performed by various microorganisms). For this reason this structural alert is considered associated to chemicals NON Readily Biodegradable.
- Aromatic aldehydes (defined as a carbonyl group linked to any aromatic ring). Several reactions convert aromatic compounds into intermediates, which are subject to ring-cleavage and subsequent funnelling into the Krebs cycle. For this reason this structural alert is considered associated to chemicals Readily Biodegradable.
- Nitrile group. They can be degraded by several strains of bacteria, fungi and plants under aerobic conditions. For this reason this structural alert is considered associated to chemicals Readily Biodegradable.

For details and full reference, see ref.1 in section 9.2.

#### **8.2.A priori or a posteriori mechanistic interpretation:**

A posteriori mechanistic interpretation.

#### **8.3.Other information about the mechanistic interpretation:**

For details and full reference, see ref. 1 in section 9.2

### **9.Miscellaneous information**

#### **9.1.Comments:**

No additional comments

#### **9.2.Bibliography:**

- [1]Lombardo A, Pizzo F, Benfenati E, Manganaro A, Ferrari T, Gini G (2014). A new in silico classification model for ready biodegradability, based on molecular fragments. Chemosphere. 108, 10-16 <http://dx.doi.org/10.1016/j.chemosphere.2014.02.073>
- [2]Toropov AA, Toropova AP, Lombardo A, Roncaglioni A, De Brita N, Stella G, Benfenati E (2012). CORAL: the prediction of biodegradation of organic compounds with optimal SMILES-based descriptors. Central European Journal of Chemistry. 10(4), 1042-1048 DOI: 10.2478/s11532-012-0031-4
- [3]Cheng F, Ikenaga Y, Zhou Y, Yu Y, Li W, Shen J, Du Z, Chen L, Xu C, Liu G, Lee PW, Tang Y (2012). In silico assessment of chemical biodegradability. Journal of Chemical Information Modeling. 52(3), 655–669 doi: 10.1021/ci200622d
- [4]Ferrari T, Gini G, Bakhtyari NG , Benfenati E (2011). Mining toxicity structural alerts from SMILES:

a new way to derive structure-activity relationships. In: Proc. IEEE SSCI 2011: Symposium Series on Computational Intelligence – CIDM 2011, 120–127. 10.1109/CIDM.2011.5949444

**9.3.Supporting information:**

qmrf480_VEGA_Training_582.sdf	<a href="http://qsar.db.jrc.ec.europa.eu/qmrf/protocol/Q17-301C-0054/attachment/A1082">http://qsar.db.jrc.ec.europa.eu/qmrf/protocol/Q17-301C-0054/attachment/A1082</a>
VEGA_Test_146	<a href="http://qsar.db.jrc.it:80/qmrf/download_attachment.jsp?name=qmrf480_VEGA_Test_146.sdf">http://qsar.db.jrc.it:80/qmrf/download_attachment.jsp?name=qmrf480_VEGA_Test_146.sdf</a>
VEGA_External_798	<a href="http://qsar.db.jrc.it:80/qmrf/download_attachment.jsp?name=qmrf480_VEGA_External_798.sdf">http://qsar.db.jrc.it:80/qmrf/download_attachment.jsp?name=qmrf480_VEGA_External_798.sdf</a>

**10.Summary (JRC QSAR Model Database)**

**10.1.QMRF number:**

Q17-301C-0054

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