

QSAR Toolbox @ U.S. EPA

Connection with OncoLogic™

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Biochemical Basis of Cancer Development

Carcinogenesis is a multi-step, multi-factorial process that requires both genetic and/or epigenetic cellular changes to progress to invasive cancer.

- 3 Basic Stages of Cancer Formation
 - Initiation
 - Gene mutations
 - Epigenetic changes
 - Promotion
 - Clonal expansion of cancer phenotype (e.g., defect in terminal differentiation, growth control, or decreased sensitivity to apoptosis)
 - Progression
 - Activation of proto-oncogenes or inactivation tumor suppression genes and antimetastatic genes



OncoLogic™: A mechanism-based expert system for predicting carcinogenic potential

- Developed by domain experts in collaboration with expert system developer
- Knowledge from SAR on >10K chemicals
- Class-specific approach to optimize predictive capability
- Considers all relevant factors including biological input when possible
- Predictions with scientific rationale and ordinal ranking
- In January 2021, OncoLogic 8™ was migrated to a newer standalone platform - OncoLogic 9™
- Available for free download from <https://tinyurl.com/oncologic>



OncoLogic™ - Expert System

HOW IT WORKS

- Mimic the thinking and reasoning of human experts using knowledge-based rules for chemical classes to predict cancer concern
 - Assigns a baseline concern level ranging from low to high for each structure class
 - Evaluates how substituents on the chemical may affect carcinogenicity
 - Concern level changes accordingly

BENEFITS

- Expedites the decision-making process
- Allows sharing of knowledge
- Reduces error and inconsistency
- Formalizes knowledge rules for cancer hazard identification
- Allows non-experts to reach scientifically supportable conclusions



OncoLogic™ - Concern Levels

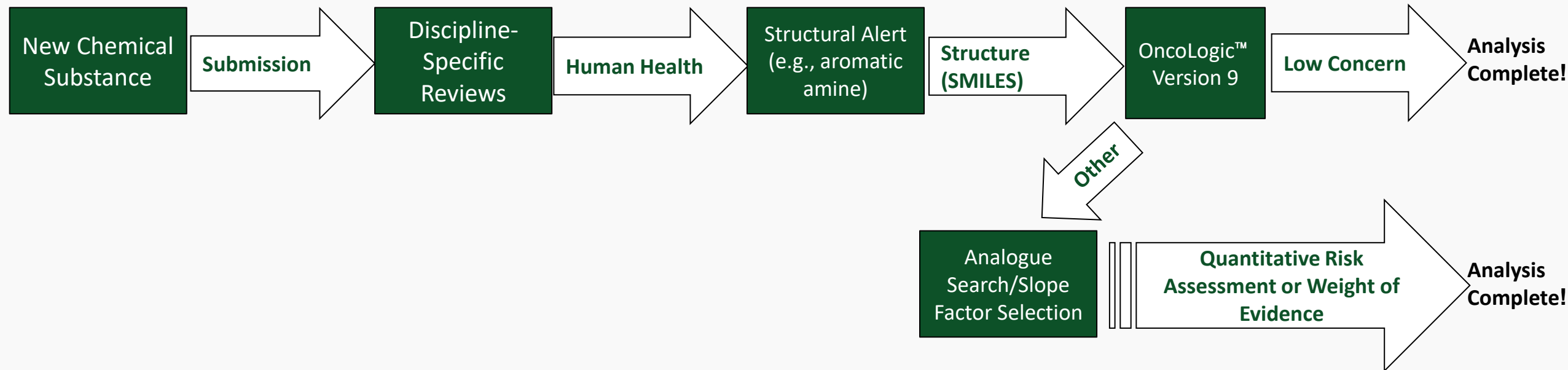
OncoLogic™ Concern	Definition
Low	Unlikely to be carcinogenic
Marginal	Likely to have equivocal carcinogenic activity
Low – Moderate	Likely to be weakly carcinogenic
Moderate	Likely to be a moderately active carcinogen
Moderate – High	Highly likely to be a moderately active carcinogen
High	Highly likely to be a potent carcinogen



EPA TSCA New Chemicals Workflow

- Refines class-based structural alerts to better predict cancer hazards

OncoLogic™ Concern
Low
Marginal
Low – Moderate
Moderate
Moderate – High
High





Integration of OncoLogic 9™ with QSAR Toolbox

- For analysis of some chemical classes (e.g., organophosphates), OncoLogic 9™ requires the user to answer a series of questions for the prediction of carcinogenic potential
- The aim of the current project is to reduce the need for expert knowledge through the integration of QSAR Toolbox with OncoLogic 9™
- The data and knowledge that exists in QSAR Toolbox would provide automatic answers to the questions asked by OncoLogic 9™
- Currently, QSAR Toolbox can automatically answer questions related to predicting concern for organophosphorus compounds



Integration of OncoLogic 9™ with QSAR Toolbox

Questions that may be asked by OncoLogic 9™ for organophosphorus compounds, which have been integrated with QSAR Toolbox:

1. Select the results of the *IN VIVO* Genotoxicity Testing
 - Answers: Positive, Negative, Unknown
2. Select the results of the *IN VITRO* Genotoxicity Testing
 - Answers: Positive, Negative, Unknown
3. Is the chemical highly toxic?
 - Answers: Yes, No, Unknown
4. Is rapid hydrolysis or detoxification expected?
 - Answers: Yes, No, Unknown



Integrating QSAR Toolbox with OncoLogic 9™ for Classification of Genotoxicity

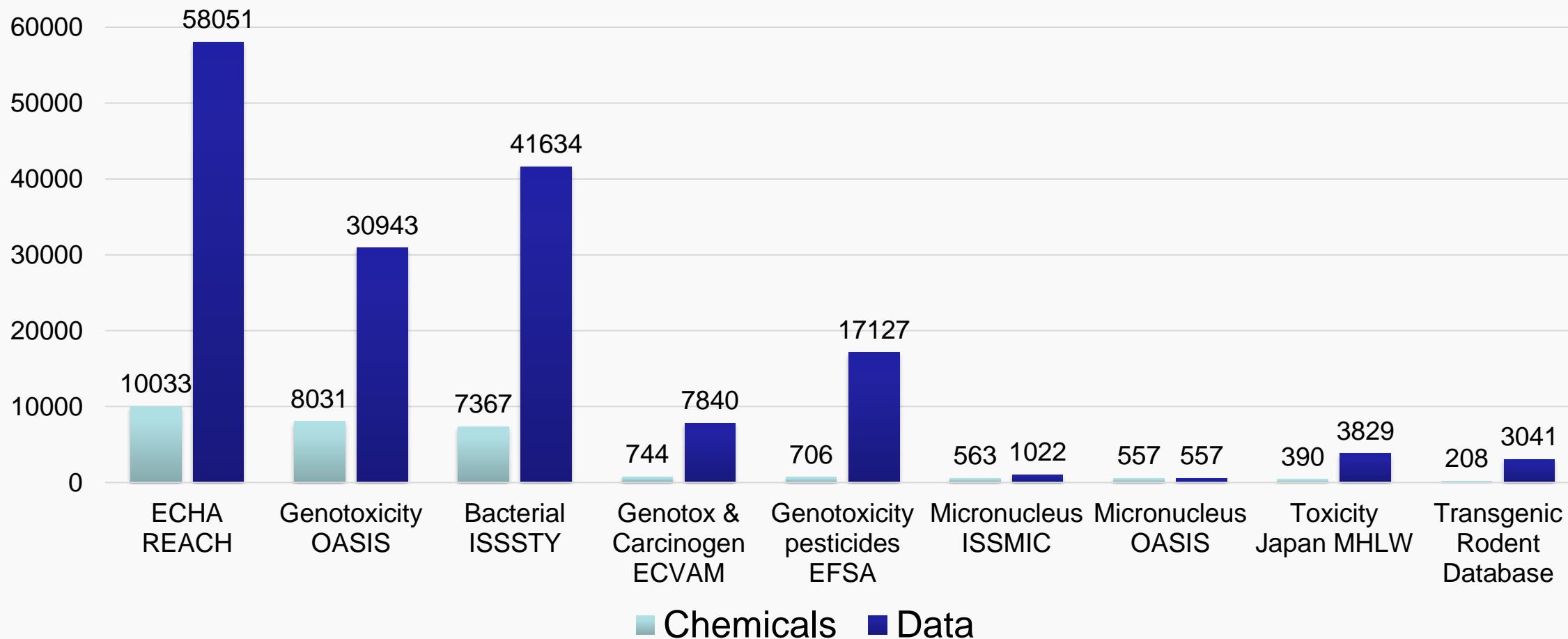
- QSAR Toolbox databases provide *in vivo* and *in vitro* genotoxicity data
- The system provides an automatic answer to the requested information about genotoxicity and extracts experimental data from QSAR Toolbox databases
- Answers are based on:
 - Positive – worst-case scenario considered (at least one positive experimental data point was identified)
 - Negative – only negative data was found
 - Unknown – i.e., equivocal (technically compromised, inconclusive, etc.) or the chemical has no data
- The answer is explained in the final report



Databases Available from QSAR Toolbox

- Databases including *in vivo/in vitro* genotoxicity data are:
 - [ECHA REACH](#)
 - [Genotoxicity & Carcinogenicity ECVAM](#)
 - [Genotoxicity OASIS](#)
 - [Genotoxicity pesticides EFSA](#)
 - [Bacterial ISSSTY](#)
 - [Micronucleus ISSMIC](#)
 - [Micronucleus OASIS](#)
 - [Toxicity Japan MHLW](#)
 - Transgenic Rodent Database

Distribution of *In Vitro/In Vivo* Genotoxicity Data Across QSAR Toolbox Databases



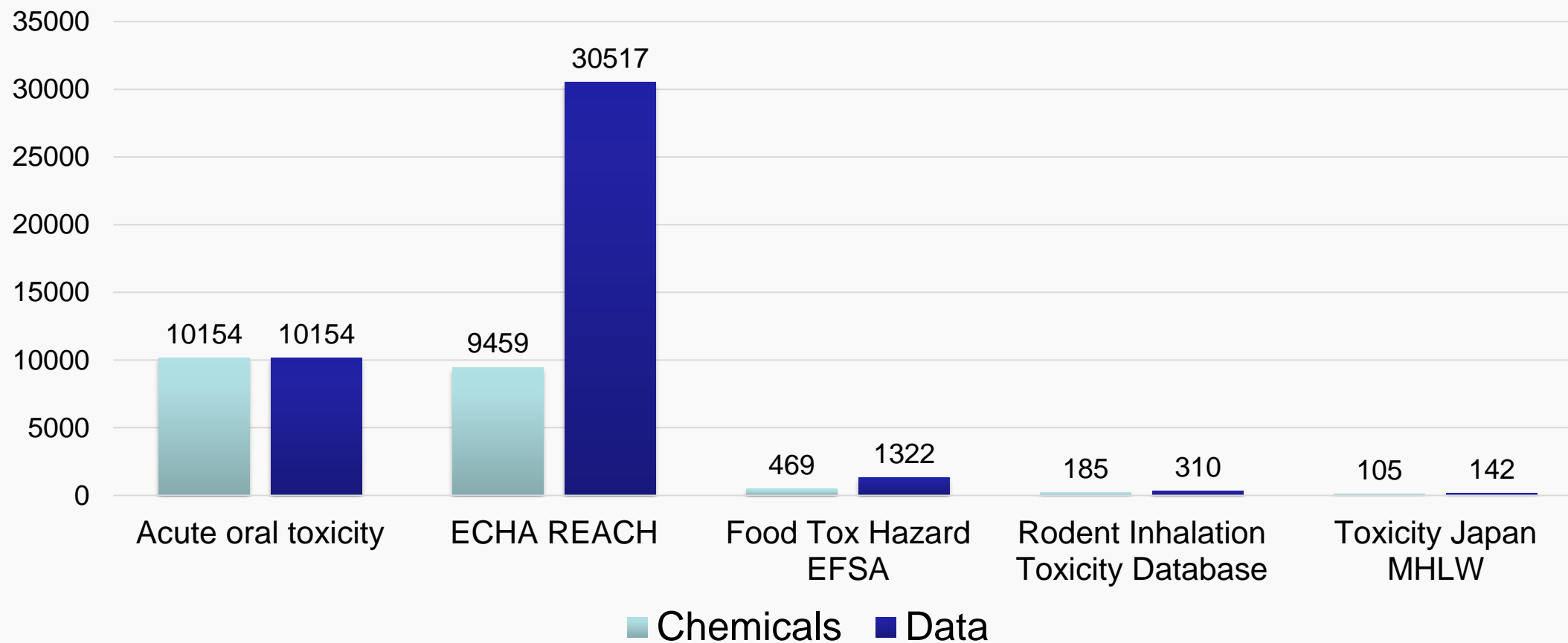


Integrating QSAR Toolbox with OncoLogic 9™ for Classification of Toxicity

- OncoLogic 9™ incorporates toxicity data into overall classification for level of concern
- Potential answers: Yes, No, Unknown
- QSAR Toolbox can extract documented **acute toxicity** data from the following Toolbox databases:
 - Acute Oral Toxicity Database
 - [ECHA REACH](#)
 - [Food Tox Hazard EFSA](#)
 - Rodent Inhalation Toxicity Database
 - [Toxicity Japan MHLW](#)



Distribution of Acute Toxicity Data Across QSAR Toolbox Databases





Classification of Chemicals as Highly Toxic Based on LD₅₀ or LC₅₀ Extracted from QSAR Toolbox

Answers	Oral (gavage, capsule, drinking water, feed, unspecified)	Dermal	Inhalation (aerosol, gas)	Inhalation (vapor)	Inhalation (dust, mist)
	Mass fraction (mg/ kg)	Mass fraction (mg/kg)	Volume concentration (ppm)	Mass concentration (mg/L)	Mass concentration (mg/L)
Yes	≤ 5	≤ 50	≤ 100	≤ 0.5	≤ 0.05
No	> 5	> 50	> 100	> 0.5	> 0.05
Unknown	Not “yes” or “no” category, no data available				



Classification of Hydrolysis Based on Calculation of Half-Life

- Question: "Is rapid hydrolysis or detoxification expected?"
- Potential Answers: Yes, No, Unknown
- QSAR Toolbox can model hydrolysis half-life values or search for experimental kinetic hydrolysis half-life data, classify chemicals into the following categories, and use the category to automatically answer the question in OncoLogic 9™:
 - very slow (>100 days)
 - slow (30 - 100 days),
 - moderate (10-30 days)
 - fast (5-10 days)
 - very fast (1-5 days)
 - extremely fast (< 1 day)

No

Yes

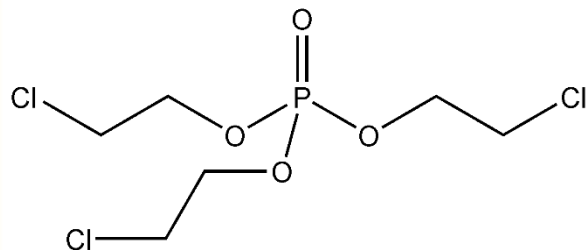
Example for the Classification of *In Vivo* Genotoxicity



OncoLogic™ 9 report

Chemical class	Level of concern	
Organophosphorus Compounds		
Trialkyl (thio)phosphates	Moderate	

OncoLogic Justification Report



JUSTIFICATION

Organophosphorus (OP) compounds are organic compounds containing phosphorus (P) as a constituent. The principal carcinogenic OP compounds are:

- i) alkyl phosphates, phosphonates, phosphoramidates and their thio derivatives (phosphorothioates, phosphorothiolates, thiophosphonates, thiophosphoramidates.
- ii) phosphoramides, and
- iii) phosphorus-containing nitrogen mustards

The structural features that may affect carcinogenicity of alkyl phosphates, phosphonates, phosphoramidates, and their related thio derivatives include:

- i) nature of alkyl groups (size, degree of branching, halogenation) which may serve as an alkylating moiety.
- ii) presence or absence of electron-withdrawing group which may enhance alkylating activity.
- iii) ease or resistance to detoxifying hydrolysis which may abolish alkylating activity.
- iv) potentially carcinogenic hydrolysis product(s).

Phosphoramides and certain phosphorus-containing nitrogen-mustards (e.g. cyclophosphamides) may require metabolic activation to alkylating intermediates.

The baseline level of concern for this unsubstituted trialkyl

phosphate-type compound, where R1 is ethyl, R2 is ethyl, R3 is ethyl, is LOW-MODERATE.

There are no or inadequate carcinogenicity data on this compound. The concern level derived for this compound is based on structure-activity relationship analysis indicative of potential alkylating activity.

The single chloro, bromo, or iodo substituent on the beta position of R1 is expected to increase the level of concern.

The single chloro, bromo, or iodo substituent on the beta position of R2 is expected to increase the level of concern.

The single chloro, bromo, or iodo substituent on the beta position of R3 is expected to increase the level of concern.

Therefore the level of concern is raised to UNKNOWN.

Because the compound is genotoxic IN VIVO, and its overall toxicity is not high, the carcinogenicity concern based on its functional properties is HIGH-MODERATE.

The concern level based on SAR analysis differs from the level based on the functional properties of the compound. The concern level assigned for the compound is the average of the levels derived from SAR analysis and consideration of the functional properties.

The final level of concern for this trialkyl phosphate-type compound is MODERATE.

The following questions have been automatically answered

by QSAR Toolbox:
 Question: "Select the results of the IN VIVO Genotoxicity Testing."
 Answer: "Positive"

 Question: "Is the compound highly toxic?"
 Answer: "No"

Explain Answers

The concern based on structure-activity relationship consideration for this trialkyl phosphate-type compound is LOW-MODERATE

The concern based on the functional properties of this trialkyl phosphate-type compound is HIGH-MODERATE

Considering both the SAR analysis and the functional properties of the trialkyl phosphate-type compound, the average level of concern is used. The final level of concern is MODERATE

The effect of any highlighted substituents is uncertain.

Explanation of QSAR Toolbox Classification of *In Vivo* Genotoxicity



for the compound is the average of the levels derived from SAR analysis and consideration of the functional properties.

The final level of concern for this trialkyl phosphate-type compound is MODERATE.

The following questions have been automatically answered by QSAR Toolbox:

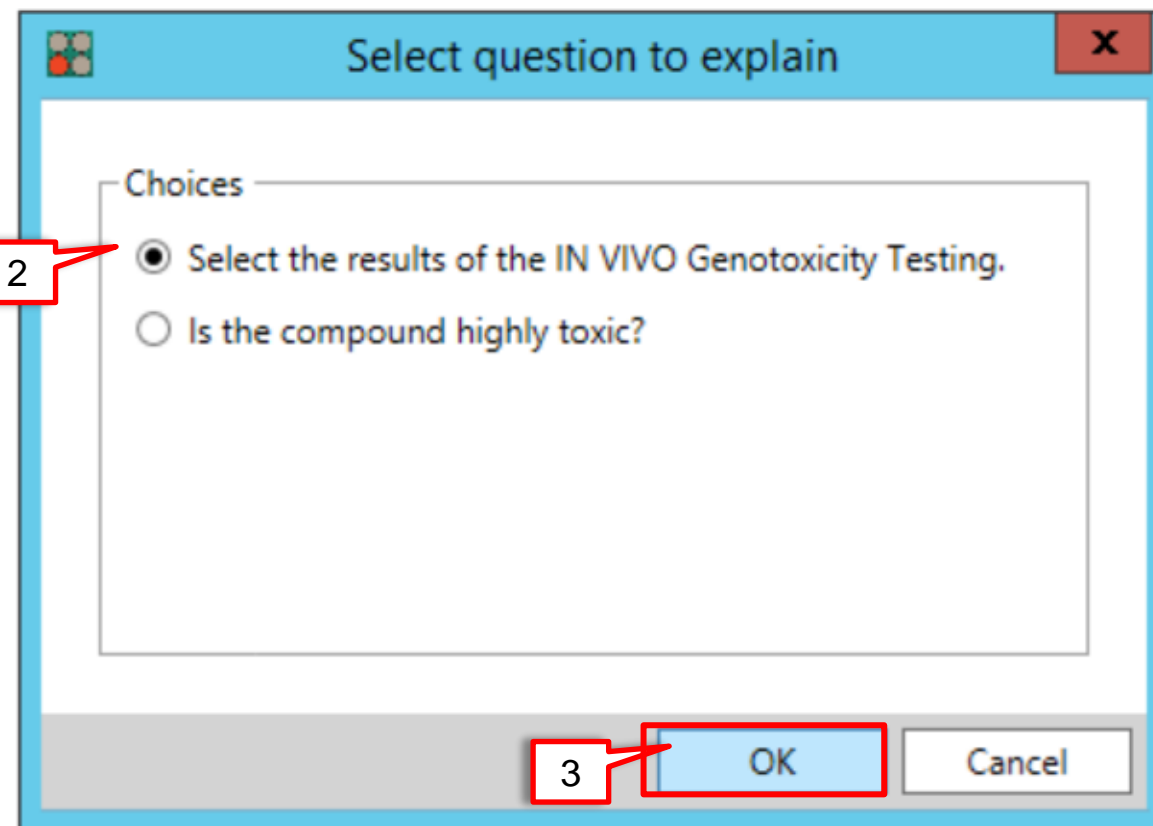
Question: "Select the results of the IN VIVO Genotoxicity Testing."

Answer: "Positive"

Question: "Is the compound highly toxic?"

Answer: "No"

1 **Explain Answers**



QSAR Toolbox Explanation



Explanation for: *in vivo* genotoxicity data -> Positive *in vivo* data found

The screenshot displays the QSAR Toolbox interface with the following components:

- Chemical Structure:** ClCCOP(=O)(OCCl)OCCl
- Category tree:** [2] Positive *in vivo* data found. A diagram shows two puzzle pieces (1 and 2) being combined into a third piece, with an 'OR' label below.
- Query details:** [2] Data Query Metabolism. A list of filters with checkboxes and counts:

Filter	Count
<input checked="" type="checkbox"/> ECHA REACH	280
<input checked="" type="checkbox"/> <i>in vivo</i> mammalian somatic and germ cell study: gene mutation	11
<input checked="" type="checkbox"/> ECHA REACH	953
<input checked="" type="checkbox"/> <i>in vivo</i> mammalian somatic cell study: cytogenicity / bone marrow chromosome aberration	3316
<input checked="" type="checkbox"/> ECHA REACH	19
<input checked="" type="checkbox"/> <i>in vivo</i> mammalian somatic cell study: cytogenicity / erythrocyte micronucleus	
<input checked="" type="checkbox"/> ECHA REACH	
<input checked="" type="checkbox"/> <i>in vivo</i> mammalian somatic cell study: gene mutation	
<input checked="" type="checkbox"/> ECHA REACH	
<input type="checkbox"/> Immunotoxicity	
<input type="checkbox"/> Irritation / Corrosion	
<input type="checkbox"/> Neurotoxicity	
<input type="checkbox"/> Repeated Dose Toxicity	
<input type="checkbox"/> Sensitisation	
<input type="checkbox"/> ToxCast	
<input type="checkbox"/> Toxicity to Reproduction	
<input type="checkbox"/> Toxicokinetics, Metabolism and Distribution	
- Explanation:** Result: Evaluation result: True. All values: Detail: Mean Min v: Max v Unit (Scale). Positive: Complete data table:

Endpoint path	Assigned SMILES	C
Human Health Hazards	False	G
Genetic Toxicity		

QSAR Toolbox Categories

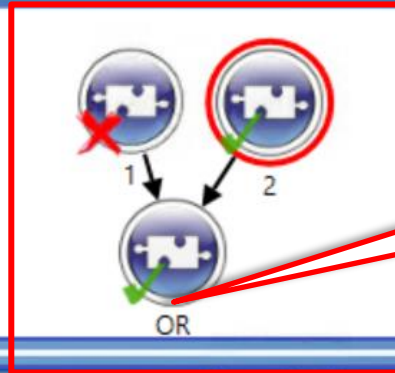


The defined categories for negative, positive, and unknown data

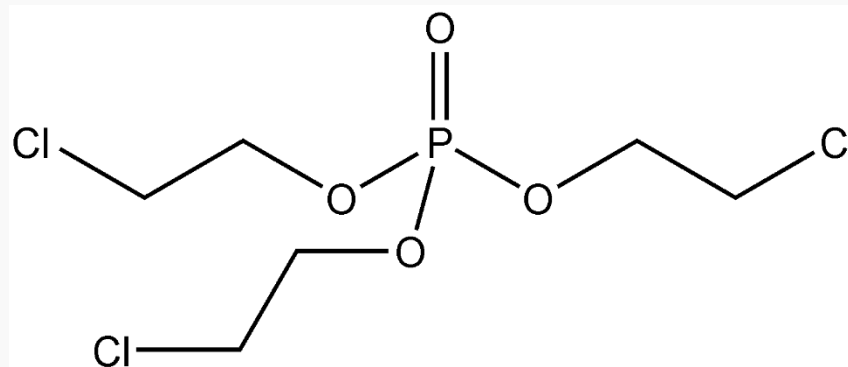
Categories

Filter:

- in vivo genotoxicity data
 - Negative in vivo data found
 - Positive in vivo data found**
 - Unknown in vivo data



Data boundaries requiring chemicals to have 'positive' *in vivo* data according to **one of two genotoxicity scales embedded in Toolbox**



QSAR Toolbox Explanation



Categories

Filter:

- in vivo genotoxicity data
 - Negative in vivo data found
 - Positive in vivo data found**
 - Unknown in vivo data

ClCCOP(=O)(OCCl)OCCl

Explanation

Result

Evaluation result: True

All values

Detail: Mean Min v: Max v Unit (Scale)

Positive

Complete data

Endpoint path	Assigned SMILES	C
Human Health Hazards		
Genetic Toxicity	False	G

Definition Properties Training Set Literature MetaInfo Table Custom Captions Scheme

Category tree

[2] Positive in vivo data found

ADD
DEL
AND

Query details

[2] Data Query Metabolism

Save

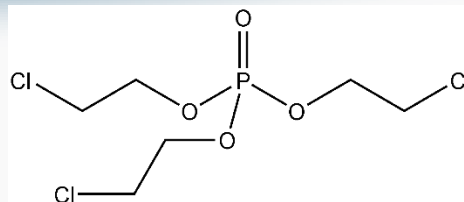
Filter:

- ECHA REACH 280
- in vivo mammalian somatic and germ cell study: gene mutation 11
- ECHA REACH 953
- in vivo mammalian somatic cell study: cytogenicity / bone marrow chromosome aberration 3316
- ECHA REACH 19
- in vivo mammalian somatic cell study: cytogenicity / erythrocyte micronucleus
- ECHA REACH
- in vivo mammalian somatic cell study: gene mutation
- ECHA REACH
- Immunotoxicity
- Irritation / Corrosion
- Neurotoxicity
- Repeated Dose Toxicity
- Sensitisation
- ToxCast
- Toxicity to Reproduction
- Toxicokinetics, Metabolism and Distribution

Metadata

Type of method

Databases Searched by QSAR Toolbox



Selected databases with genotoxicity data used for searching experimental data

Results derived after performing the data search

Expanded results with metadata

Query details

[2] Data Query Metabolism

Save

Filter:

- ECHA REACH
- in vivo mammalian somatic and somatic cell study: gene mutation
- ECHA REACH
- in vivo mammalian somatic cell study: cytogenicity / bone marrow chromosome aberration
- ECHA REACH
- in vivo mammalian somatic cell study: cytogenicity / erythrocyte micronucleus
- ECHA REACH
- in vivo mammalian somatic cell study: gene mutation
- ECHA REACH

Close

280

11

953

3316

19

Metadata

Type of method

Available data in the respective database

Defined type of method

QSAR Toolbox Explanation Summary



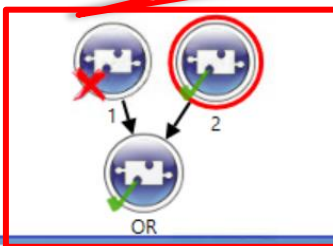
The defined categories for negative, positive, and unknown data

in vivo genotoxicity data

- Negative in vivo data found
- Positive in vivo data found**
- Unknown in vivo data

ClCCOP(=O)(OC)OCCCl

Data boundaries requiring chemicals to have 'positive' *in vivo* data according to **one of two genotoxicity scales embedded in Toolbox**



Selected databases with genotoxicity data used for searching experimental data

Results derived after performing the data search

Category tree

[2] Positive in vivo data found

Query details

[2] Data Query Metabolism

Filter:

- ECHA REACH
- in vivo mammalian somatic and germ cell study: gene mutation
- ECHA REACH
- in vivo mammalian somatic cell study: cytogenicity / bone marrow chromosome aberration
- ECHA REACH
- in vivo mammalian somatic cell study: cytogenicity / lymphocyte micronucleus
- ECHA REACH
- in vivo mammalian somatic cell study: gene mutation
- ECHA REACH
- Immunotoxicity
- Irritation / Corrosion
- Neurotoxicity
- Repeated Dose Toxicity
- Sensitisation
- ToxCast
- Toxicity to Reproduction
- Toxicokinetics, Metabolism and Distribution

Metadata

Type of method

Expanded results with metadata

Result

Evaluation result: True

Endpoint	Path	Assigned SMILES
Human Health Hazards	Genetic Toxicity	False

Available data in the respective database

Defined type of method



Recent Updates

- 1. New version of OncoLogic™ 9.2**
 - Automatic prediction of alkylating ability for an organophosphorus compound
 - Automatic prediction of chelating ability for an organophosphorus compound
- 2. OncoLogic™ 9 Add-In for QSAR Toolbox**
 - Evaluation of the carcinogenic potential of multiple compounds using batch mode
 - Add-In is used to extract information from QSAR Toolbox to automatically answer OncoLogic™ questions
- 3. QSAR Toolbox update (QSAR Toolbox 4.5 Service Pack 1)**



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THANK YOU!

Questions?

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