# QSAR TOOLEOX

The OECD QSAR Toolbox for Grouping Chemicals into Categories

# OECD QSAR Toolbox v.4.1

Examples illustrating customized search (Query Tool) in Toolbox

# **Outlook**

## • Background

- Objectives
- Overview of Query tool
- Query tool window
- The exercise
- Workflow process
- Save QT searches

# Background

 This is a step-by-step presentation designed to take the user through the functionalities of Query tool (QT) engine implemented in Toolbox

# **Outlook**

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# **Objectives**

# This presentation demonstrates a number of functionalities of the Query tool (QT):

- Identifying chemicals with specified structural fragments
- Identifying chemicals answering specific structural fragment criteria combined with parametric ranges
- Identifying chemicals answering specific combination of data, structural fragments and parametric data

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# **Overview of Query tool**

**<u>Goal</u>**: Searching for chemicals by structure, sub fragments, phys-chem properties and experimental data

#### Chemical identifier search

- CAS
- Name
- Molecular structures

#### Parameter search

- Calculated 2D parameters
- Calculated 3D parameters

#### Data search

• Search for data and metadata within imported databases

#### • Extended search

- Profiling
- Substructure
- Similarity

# **Overview of Query tool** Prerequisites

•The Query tool functionality search for single structures matching desired criteria.

•The Query tool functionality search for chemicals within the selected databases and inventories only



# **Overview of Query tool** Procedure for defining query

# **Step 1:** Select databases/inventories of interest in the *Data* module



# **Step 3**: Specify criteria for searching structures



#### Step 2: Click Query button



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# **Query tool window**

• The Query tool is easily accessible on the Toolbox input panel



Close

Execute

# **Outlook**

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# **The Exercise**

- In this exercise we will demonstrate the following queries:
  - Subfragment search for:
    - Substituted diphenyl amine (Example 1)
    - Aliphatic halogens (Example 2)
  - Combination of Environmental Data (BCF) and 3D parameters (Dmax) (Example 3)
  - Combination of predefined category (Aldehydes) and ecotox data (LC50<1mg/l) (Example 4)
  - Combination of positive Ames; positive Carcinogenicity data and Subfragment search (*Epoxides*) (*Example 5*)
  - Combination of Skin sensitization data (EC3) and predefined category (*Aldehydes*) (*Example 6*)

# **Outlook**

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## Subfragment search for identifying chemicals Substituted diphenyl amine

#### Example 1

Search for structures that meet the structural requirements:

• Substituted diphenyl amine



N(c1cccc1)c1ccccc1

#### Example 1

#### Substituted diphenyl amine



#### N(c1ccccc1)c1ccccc1









QT procedure starts with selecting databases/inventories used for searching the desired criteria:

1. **Select** HPVC OECD inventory located under Data section. No databases have been selected in this exercise; 2. Click **Query** button located under **Input** section; The message informs the user that search will be performed on selected databases or inventories only 3. Click **Yes**; 4. Query panel appears; 5. Select **SubFragment** panel;

#### Example 1

fragment;

#### Substituted diphenyl amine



#### Example 1

#### Substituted diphenyl amine



#### N(c1ccccc1)c1ccccc1

Subfragments:	Options Search mode: All v Exact match Exact connectivity
View mode: Facade	<ul> <li>Navigation mode: Cascade </li> <li>xplore</li> </ul>

5. Click the pencil button to draw a single bond; 6. Click near to one of the C atom from the benzene ring to draw a single bond; 7.Click **N** atom from the templates; 8. Put the selected N atom over the C atom from the single bond; 9. Repeat step 5 and draw a single bond to the N atom from NH2 group;



#### Example 1

#### Substituted diphenyl amine



#### N(c1ccccc1)c1ccccc1

Subfragments:	Options Search mode: All v Exact match Exact connectivity
View mode: Facade	Navigation mode: Cascade



10. Left click over benzene ring from the template in order to take benzene ring 11. Left click over the C atom in order to define second benzene ring to the NH group 12. Click **OK**.

#### Example 1

Substituted diphenyl amine



N(c1ccccc1)c1ccccc1

<b>(</b> ) s	earch									-		×
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Sub	fragmen	ts:										
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1. Click **Add** button; 2. The defined query appears on logic panel; 3. Double click over the query or click

**Execute** button to execute it.

#### Example 1

Substituted diphenyl amine



#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens

- **R** Any atom except H **Hal** – Cl, Br, I
- •HPVC OECD inventory has

been selected

•Search for discrete chemicals



1. Select the **HPVC OECD** inventory (no database has been selected in this case, only HPVC OECD) located under *Data* section; 2. Click **Query** button under *Input* section; 3. Click **Yes** 

#### Example 2

Search for structures that meet the structural requirements:

<ul> <li>Aliphatic halogens</li> </ul>	Search     CAS Name Data Darameters SubFragment Datagony Sir	ND OR NOT Delete Clear
Ŗ	Subfragments: 2	
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H L	Smarts ~	×
<b>R</b> – Any atom except H <b>Hal</b> – Cl, Br, I	Add Edit Remove	[] \$ Rpt
•HPVC OECD inventory has been selected •Search for discrete chemicals	Left click on any marked atom to exp	Rpc Exh
1. Query panel appears; 2. <b>Go</b> to SubFragment panel; 3. <b>Click</b> "Add" button; 4. "SMARTS Editor" window appears;	S F P	OK Cancel
		LXecute

#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens



**R** – Any atom except H Hal – Cl, Br, I

•Define **Enumeration** fragment with three members (halogen atoms)

- 1. Left click the **C** symbol from the template and then left click in the drawing panel;
- 2. Left click the **pencil** button and draw a single bond;
- 3. Select the "[]" (enumeration) button;
- 4. Click over one of the carbon atoms;



#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens



**R** – Any atom except H Hal – Cl, Br, I

•Define **Enumeration** fragment with three members (halogen atoms)

5. Click the **selection tool**; 6. Click the new object; 7. *Object explorer* panel appears; 8. Right click over the *OR* expression node and select **Add child**;



25

#### Example 2

Search for structures that meet the structural requirements:



The OECD QSAR Toolbox for Grouping Chemicals into Categories

SMARTS Editor

#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens



**R** – Any atom except H Hal – Cl, Br, I

•Define **Enumeration** fragment with three members (halogen atoms)

**Hal** substituent is ready when all elements are selected (1). Now we can continue with defining of the rest substituents of the carbon atom (see the general structure above).



#### Example 2

Search for structures that meet the structural requirements:



#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens



```
R – Any atom except H
Hal – Cl, Br, I
```

•Define fragment R including any type atom except H atom

1. Select the pencil button; 2. Draw two single bonds to the C atom;



29

#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens

R - C - Hal

```
R – Any atom except H
Hal – Cl, Br, I
```

•Define fragment R including any type atom except H atom

Click Selection tool; 4. Select C atom;
 Select "Any atom" from the Element drop-down menu;



#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens

**R** – Any atom except H Hal – Cl, Br, I

•Define fragment R including any type atom except H atom

6. Select **Any** from the *Aromatic or aliphatic* drop-down menu; 7.Repeat the steps from 3 to 6 for the second carbon atom



#### Example 2

Search for structures that meet the structural requirements:



#### Example 2

Search for structures that meet the structural requirements:

• Aliphatic halogens



**R** – Any atom except H **Hal** – Cl, Br, I

•Define skeleton of target compound

1. The defined general structure of the aliphatic halogens appears on the separate plot; 2. Click **Add** button; 3. Double click the query to perform the search

CAS Name Data Parameters SubFragment Category Similarity	AND	OR	NOT	Delete	Clear
Clear All					
Subfragments:     [CI,Br,I][CH]([*])[*]       Options       Search mode:       All       Exact match       Exact connectivity		3			
iew mode: Facade	2				
2 Update Add	Save	Load			
				Execute	Close

# Subfragment search for identifying chemicals

#### Aliphatic halogens

#### Example 2

• Aliphatic halogens



## Subfragment search for identifying chemicals BCF and (Dmax)

#### Example 3

•Structures search criteria

- Endpoint is BCF  $\geq$  10 L/kg bdwt
- 3D parameter minimum value

of Diameter maximum >15 Å

## Subfragment search for identifying chemicals BCF and (Dmax)

#### Example 3

- •Structures search criteria
  - Endpoint is BCF  $\geq$  10 L/kg bdwt
  - 3D parameter minimum value of Diameter maximum >15 Å



- 1. Select databases including BCF data (Bioaccumulation Canada; Bioaccumulation fish CEFIC LRI and Bioconcentration NITE). No inventory has been selected in this exercise;
- 2. Click Query button;
- 3. Confirm that the searches perform on selected databases and inventories;
- 4. Go to Data panel;
- 5. Expand the endpoint tree and select BCF;
- 6. Specify qualifier "≥" 10 in the *Mean value* field;
- Define the scale "Bioaccumulation" and appropriate unit (L/kg bdwt);
- 8. Click Add button;


### Example 3

- •Structures search criteria
  - Endpoint is BCF  $\geq$  10 L/kg bdwt
  - 3D parameter minimum value

of Diameter maximum >15 Å



- 2. Select **Diameter maximum** from the popup list;
- Specify qualifier "≥" 15 in the *Expression* field;
- 4. Click Add button;



### **Example 3**

- Structures search criteria
  - Endpoint is BCF  $\geq$  10 L/kg bdwt
  - 3D parameter minimum value of Diameter maximum >15 Å

Both queries should be linked together by logical AND. For this purpose both queries should be selected first. How to do this:

1. Right click over the first query to select it (the selected boundary should become orange colored);



Selected auerv



(S) Not selected query

- 2. When the both queries are selected, click "AND" button;
- 3. The two queries are combined by logical "AND";
- 4. Double click "AND" query or click "Execute" button to execute the search;





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Maximum donor delocalizability	Copy path		Not calculated	Not calculated	Not calculated	Not
Planarity	Superior		Not calculated	Not calculated	Not calculated	Not
Planarity conjugate				Not calculated	Not calculated	Not
VdW surface	Sort	· · · · ·	1 Ascending	Not calculated	Not calculated	Not
VdW surface DPSA 5	Calculate/extract all paran	neters for all chemicals	Descending	Not calculated	Not calculated	Not
VdW surface DPSA	Calculate/extract all 3D pa	rameters for all chemicals	Descending	Not calculated	Not calculated	Not
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LC 50 and predefined category (Aldehydes)

### Example 4

•Structures search criteria

- Endpoint: LC 50 <1 mg/l
- Fish: P.promelas
- Effect: *Mortality*
- Predefined category: *aldehydes*

- Structures search criteria
  - Endpoint: LC 50 <1 mg/l
  - Fish: *P.promelas*
  - Effect: *Mortality*
  - Predefined category aldehydes

	Query
Databases	
ptions 🖌	Query tool
f Select All Unselect All Invert	The query was created with the following data sources:
	Bioaccumulation Canada
Ecotoxicological Information	Bioaccumulation fish CEFIC LRI
Aquatic ECETOC	Bioconcentration NITE
🗸 Aquatic Japan MoE	
	Would you like to restore them?
ECHA CHEM	from you me to restore them
ECOTOX	

- 1. Select databases related to LC50 data (Aquatic ECETOC; aquatic Japan MoE; Aquatic OASIS; ECOTOX). No inventories has been selected in this case;
- 2. Click Query tool button;
- Select "No";
- 4. Go to Data panel;
- 5. Type in filter the name of searched endpoint (LC50);
- Select Aquatic Toxicity and expand the endpoint tab;
- 7. Check **LC50**

SubFragment	Category	arity	AND	OK
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LC 50 and predefined category (Aldehydes)

- •Structures search criteria
  - Endpoint: LC 50 <1 mg/l
  - Fish: *P.promelas*
  - Effect: *Mortality*
  - Predefined category: *aldehydes*

- 1. Open Metadata field
- 2. Select "**Test organism (species)**" from thedrop-down menu;
- 3. Click Add;
- 4. Select *Pimephales promelas* from the dropdown menu;
- 5. Click Add;
- 6. Go back to **Metadata** field and select **Effect** from the drop-down menu;
- 7. Click Add;
- 8. Use filter to find **"Mortality**" from the appeared drop-down menu;
- 9. Click Add



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### Example 4

- Structures search criteria
  - Endpoint: LC 50 <1 mg/l
  - Fish: P.promelas
  - Effect: *Mortality*
  - Predefined category: *aldehydes*

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4	Filter: mas Mass concentra Mass Mass fraction Concentration i Administered d	tion n body (mass) ose(mass)	Update		Save	Load	]		

- 1. Open **Data** panel;
- Open qualifiers from Mean value and select "<";</li>
- 3. Type "1" in the blank field;
- 4. Use Filter to find "Mass concentration";
- 5. Select "**mg/l**" from the list with units;
- 6. Click Add;

 $\mathbf{v}$ 

- •Structures search criteria
  - Endpoint: LC 50 <1 mg/l
  - Fish: *P.promelas*
  - Effect: *Mortality*
  - Predefined category: *aldehydes*

E Search					_	×
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Target	- I`					
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Down Up	_ 2	2				 
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Acyl halide						
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Aldehyde	1					
Aldimine						
Aldoxime derivatives						
Aliphatic amine, primary						 
and a second	<u> </u>	4				
Combine profiles Invert						
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ndpointPath/EndpointDefinition :						
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Text metadata: Test organisms (species) is Updat	te					
Iext metadata: Effect is 'Mortality'	1	Save	Load			
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- 1. Open Category panel;
- 2. Use *Filter* to find **Organic functional groups** profiler from the list with profilers;
- 3. Find "Aldehyde";
- 4. Click "**Up**" to move the selected category from panel "*Profilers*" to the panel "*Target*"
- 5. The selected category appears in the panel "Target"

### Example 4

### •Structures search criteria

- Endpoint: LC 50 <1 mg/l
- Fish: P.promelas
- Effect: *Mortality*
- Predefined category: *aldehydes*

1. Click Add button;

2. The query with predefined category appears on the logic panel.

The both queries should be combined by logical "AND". Follow the steps explained on slide # 39

- 3. Right click over the both queries to highlight them;
- 4. Click **AND** button;
- The two queries are combined by logical "AND";
- 6. Click **Execute** button in order to execute the Query



LC 50 and predefined category (Aldehydes)

#### **Example 4**

QSAR TOOLEOX	Input Profiling Data	ry definition   Data Ga	p Filling > Repo	rt			
Data Import Export Gather Import IUCLID6 IUCLID6							
Documents	Filter endpoint tree Structure	Lag	2	3	4	5	6 ~_0_0
	Structure info     Parameters     Physical Chemical Properties     Environmental Fate and Transport     Ecotoxicological Information     Human Health Hazards						
Options ▲     f Select All Unselect All Invert							

The Query tool (QT) identified 6 chemicals matching the desired criteria. The list of identified chemicals appear automatically on the data matrix. Number of found chemicals is marked in the name of the node (1). In order to check the correctness of the performed query search the user should gather ecotox data and profile according to OFG Profiler. The last two steps are presented on the next two slides.

LC 50 and predefined category (Aldehydes)



LC 50 and predefined category (Aldehydes)

QSAR TOOLBOX	nput Profiling Data Catego	ory definition	ap Filling ► Repr	prt			
Documents     Document 1     Query Tool: 6	Filter endpoint tree Structure Structure Parameters Physical Chemical Properties Environmental Fate and Transport Ecotoxicological Information Human Health Hazards	1 	2 OO M: 0.3 (0.26+0.34) m	3 	4 M: 0.845 mg/L	5 M: 0.767 mg/L	6 
Profiling methods      Options	Organic functional groups	Aldehyde Alkyl (hetero)arenes Alkyl-, alkenyl- and z Aryl Ether tert-Bubd	Aldehyde Aryl Aryl halide Ether	Aldehyde Alkyl halide Aryl	Aldehyde Aryl Aryl halide Phenol	Aldehyde Aryl Aryl halide Phenol	Aldehyde Aryl Ether

- 1. Go to the *Profiling* module;
- 2. Check Organic functional group profile;
- 3. Click Apply;
- 4. All 6 chemicals have "Aldehyde" fragment within its structure;

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

### Example 5

- Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive

#### AND

- Endpoint: Summary carcinogenicity
- Species: Rat
- Route: gavage
- Data: Positive
- Predefined category: *Epoxides*

# Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

### Example 5

- Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive

#### AND

- Endpoint: Summary carcinogenicity
- Species: Rat
- Route: gavage
- Data: Positive
- Predefined category: Epoxides



Before application of Query tool, the user should select databases which includes required data. Inventories are not selected in this particular case. 1. Select the following databases:

- Bacterial mutagenity ISSSTY
- Carcinogenicity potency Database (CPDB)
- Carcinogenicity&mutagenicity ISSCAN
- Genotoxicity OASIS
- Toxicity Japan MHLW
- 2. Click Query button;
- 3. Select **No** to not restore the databases used in the previous example.

# Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

### Example 5

- •Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive

#### AND

- Endpoint: Summary carcinogenicity
- Species: Rat
- Route: gavage
- Data: Positive
- Predefined category: *Epoxides* 
  - 1. Click on the **Data** panel;
  - 2. Use filter to find searched endpoint;
  - 3. Select Gene mutation;



## Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment



## Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

### **Example 5**

- Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive
  - Endpoint: Summary carcinogenicity
  - Species: Rat
  - Route: gavage
  - Predefined category: *Epoxides*

1. Before defining the second query click **Clear All** button;

2. Open endpoint path and select *Carcinogenicity* node;

3. Select Summary carcinogenicity;



## Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment



# Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

- •Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive
  - Endpoint: Summary carcinogenicity
  - Species: Rat
  - Route: gavage
  - Predefined category: *Epoxides*

- 1. Click *Category* panel;
- 2. Select Carcinogenicity (genotox and nongenotox) alerts by ISS;
- 3. Select category **Epoxides and aziridines** (Genotox);
- 4. Click **Up** to move the selected category to the panel Target;
- 5. Click Add button;
- 6. The query appears on the logic panel;



## Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

- •Structures search criteria
  - Endpoint: Gene mutation
  - Species: S.thyphimurium
  - Data: Positive
  - Endpoint: Summary carcinogenicity
  - Species: Rat
  - Route: gavage
  - Predefined category: *Epoxides*

- 1. Right click over the three queries (see slide # 39);
- 2. Click **AND** button;
- 3. The three queries are combined by logical "AND";
- 4. Click **Execute** button;



### Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

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## Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including Epoxide fragment

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Cell Transformation Assay ISSCTA DBIU6 Dendritic cells COLIPA Developmental & Reproductive Toxicity (DART) Developmental toxicity ILSI Inventories Options f Select All Unselect All Invert Canada DSL	Genetic Toxicity	(6/92) M: Equivocal M: Positive M: Positive M: Positive M: Positive	M: Positive M M: Positive M M: Positive M M: Positive M M: Positive M	1: Positive 1: Positive 1: Positive 1: Positive	M: Positive M: Positive M: Positive M: Positive	M: Positive M: Positive M: Positive M: Positive M: Positive	M: Positive M: Positive M: Positive M: Positive
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### Subfragment search for identifying chemicals

Chemicals with Ames and Carcinogenicity positive data including \_\_\_\_\_ Epoxide fragment

Pro 4 Custom profile Apply View New Delete	Profiling     Data     Category definition	Data Gap Filling	► Report				
Ocuments ▲ Â Document 1 Q Query Tool: 6	Filter endpoint tree  Structure  Structure info Parameters Physical Chemical Properties	1	2	3 >(O)>	4 0 0	5	6 H3C
2 Select All Unselect All Invert About Filter: Carci Close Close Carcinogenicity (genotox and nongenotox) alerts by ISS	Environmental Fate and Transport     Ecotoxicological Information     Human Health Hazards     (6/176)     Profile     Substance type     Endpoint Specific     Carcinogenicity (genotox and nongenotox) al	M: 34.7 mg/kg bdwt	M: 2.96 mg/kg bdwt Discrete chemical Aliphatic halogens (Ger Epoxides and aziridines Structural alert for genc	M: 118 mg/kg bdwt/ Discrete chemical Epoxides and aziridines Structural alert for genc	M: 24.3 mg/kg bdwt Discrete chemical Epoxides and aziridines Structural alert for genc	M: 21.3 mg/kg bdwt Discrete chemical Epoxides and aziridines Structural alert for geno	M: 74.4 mg/kg bdwt Discrete chemical Epoxides and aziridines Structural alert for genc

- 1. Go to Profiling;
- 2. Click Unselect All;
- 3. Use filter and select Carcinogenicity (genotox and nongenotox) alerts by ISS;
- 4. Click Apply;
- 5. All 6 chemicals have "Epoxides and Aziridines" fragment within its structure.

Chemicals with Skin sensitization positive data including Aldehyde fragment

### Example 6

•Structures search criteria

- Endpoint: EC3
- Data: Positive
- Predefined category: alpha, betaunsaturated aldehydes

# Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

- •Structures search criteria
  - Endpoint: *EC3*
  - Data: Positive
  - Predefined category: alpha, betaunsaturated aldehydes



- 1. Select databases including skin sensitization data. No inventories has been selected in this case;
- 2. Click Query tool button;
- 3. Select **No** to confirm that the databases in previous example not be used;
- 4. Click on *Data* panel;
- 5. Use filter to find searched endpoint;
- 6. Check EC3;
- 7. Go to Data panel;
- 8. Select scale Skin sensitization II (ECETOC);
- 9. Select Positive;
- 10.Click Add button;

	Search	– 🗆 X
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5	Skin sensitisation II (ECETOC)	
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	Negative	
	Positive	
		9
	Update	
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		10 Execute Close

# Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

- •Structures search criteria
  - Endpoint: EC3
  - Data: Positive
  - Predefined category: alpha, betaunsaturated aldehydes

- 1. Click Category panel;
- Type in *Filter* to find the profile "Protein binding alerts for skin sensitization by OASIS";
- Select category Michael addition >>alpha, beta-Unsaturated carbonyl compounds>>alpha, beta-Aldehydes and move the selected category to the panel Target;
- 4. Click Add button;
- 5. The query appears on the logic panel.



# Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

- •Structures search criteria
  - Endpoint: EC3
  - Data: Positive
  - Predefined category: alpha, betaunsaturated aldehydes



- 2. Click AND button;
- 3. The two queries are combined together by logical "*AND*";
- 4. Click **Execute** button or double click And;



## Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

#### **Example 6**

QSAR TOOLBOX	Input     ▶ Profiling     ▶ Data     ▶ Cates	gory definition   Data Gap	o Filling ► Repo	rt						X 0 %
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	Structure info     Parameters     Physical Chemical Properties     Environmental Fate and Transport     Ecotoxicological Information     Human Health Hazards     Acute Toxicity									
Databases      Options 4      f     Select All     Unselect All     Invert      Rep Dose Tox Fraunhofer ITEM      Repeated Dose Toxicity HESS      Rodent Inhalation Toxicity Database      Shu Irritation	Bioaccumulation     Carcinogenicity     Developmental Toxicity / Teratogenicity     Genetic Toxicity     Immunotoxicity     Irritation / Corrosion     Neurotoxicity									
Skin Sensitization     Skin sensitization ECETOC     ToxCastDB     Toxicity Japan MHLW     Toxicity to reproduction (ER)     ToxRefDB US-EPA	Protoinduced coacity     Repeated Dose Toxicity     Sensitisation AW SW AOP     ToxCast     Toxicity to Reproduction									

The Query tool (QT) identified 14 chemicals matching the desired criteria. In order to check the correctness of the query the user should gather experimental data for skin sensitization and profile the identified chemicals according to the "*Protein binding alerts for skin by OASIS*" profiler used in the query boundary. The last two steps are presented on the next two slides.

## Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

			_							
QSAR TOOLBOX		0101 01 1010								
Data Import Export Gather 2 106 IUCUD6	Input	efinition 🕨 Data Gap	p Filling 1 2 3 4	Go to the <i>Data</i> Databases are Click <b>OK</b> ;	module; already s	elected, c	lick <b>Gath</b>	<b>ier</b> data;		
Documents  A  Document 1	Filter endpoint tree		5	. Data appear oi	n data ma	atrix ansv	vering th	e following		
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	- Human Health Hazards									
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Rodent Inhalation Toxicity Database	Neurotoxicity									
Skin Initiation	Photoinduced toxicity									
Skin sensitization ECETOC	Repeated Dose Toxicity									
ToxCastDB	Sensitisation AW SW AO	P								
Toxicity to reproduction (ER)	Skin									
ToxRefDB US-EPA				M. De cities			5			
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### Subfragment search for identifying chemicals

Chemicals with Skin sensitization positive data including Aldehyde fragment

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Documents     Documents     Documents     Profiling me     Documents     Document	Filter endpoint tree Structure  Structure  Structure info Parameters	1 2 4 н <sub>1</sub> с	<ol> <li>Click</li> <li>Perf</li> <li>sele</li> <li>Che</li> <li>Che</li> <li>All</li> <li>Prot</li> </ol>	Click <b>Apply</b> ; Perform right click over the <i>Protein binding alerts</i> and select "Profile statistic"; Check the box next to <i>Group by category</i> . All 14 chemicals are alpha-beta aldehydes according to Protein binding alerts for skin sensitization profiler.						
<ul> <li>Keratinocyte gene expression</li> <li>Oncologic Primary Classification</li> <li>Protein binding alerts for Chromosomal aberral</li> <li>Protein binding alerts for skin sensitization acc</li> <li>Protein binding alerts for skin sensitization by i</li> <li>Frotein binding alerts for skin sensitization by i</li> <li>Frotein binding version by i</li> <li>Frotein binding version by i</li> <li>Frotein binding alerts for skin sensitization by i</li> <li>Frotein binding alerts for skin sensitization by i</li> <li>Frotein binding version binding alerts for skin sensitization by i</li> <li>Frotein binding alerts for skin sensitization by i</li> <li>Frotein binding version binding interversion control binding interversion context sensitization control binding interversion for the sensitization control binding interversion control binding interversion for the sen</li></ul>		M: Positive M: Positive M: Positive M: Positive M: Positive Michael Addition > M Michael Addition >> M Michael	Addition 4 Addition 4 Addition 4 Addition 4 Addition 5 Addition 5	rofile Statistic Siroup by category Category Michael Addition >> Michael addition on alpha Michael Addition >> Michael addition on alpha Michael Addition >> Michael addition on conju Michael Addition >> Michael addition on conju Michael Addition >> Michael addition on conju	Count % 7	14 Michael Addition a compounds >> alg Save to smi   Print   Add	Michael addition on al ha,beta-Aldehydes     din new doc     2     5910-85-0	pha,beta-Unsaturated carbonyl		
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# **Outlook**

- Background
- Objectives
- Overview of Query tool
- Query tool window
- The exercise
- Workflow process
- Save QT searches

# **Saving the prediction result**

- This functionality allows storing/restoring the current state of Toolbox documents including loaded chemicals, experimental data, profiles, predictions etc., on the same computer. The functionality is implemented based on saving the sequence of actions that led to the current state of the Toolbox document and later executing these actions in the same sequence in order to get the same result(s).
- Saving/Loading the file with TB prediction is shown on the next screenshots

# **Saving the QT requests**



# **Open saved file**

