OECD QSAR Toolbox v.4.4

Step by step example on how to implement AOP information from Effectopedia Wizard to Toolbox
Outlook

• Background
• Specific Aims
• Overview of Effectopedia AOP
• Migrating the Effectopedia AOP to Toolbox
• Building a new AOP-based profiler in Toolbox for predictive purposes
• Application of migrated AOP for collecting weight-of-evidences
This is a step-by-step presentation designed to take the Toolbox user through the workflow on how to implement and use an already developed Effectopedia AOP within the Toolbox.

Note: This tutorial only applies to those who run Toolbox in standalone/single user mode.
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Specific aims

This presentation demonstrates:

• Migration of an AOP built in the Effectopedia platform to the Toolbox;*
• configuring of the migrated AOP using functionalities of the Toolbox in order to be used for predictive purposes;
• the capabilities of implemented AOP for collecting weights of evidences – run an example chemical.

* Note: The AOP used in the current example is just an illustrative example and it is not OECD approved.
Outlook

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• **Overview of the used Effectopedia AOP**

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Overview of the used Effectopedia AOP

- An example AOP for assessing endocrine disruption (ED) of chemicals developed in Effectopedia platform will be used;

- The descriptive AOP was implemented based on an AOP cited by P. Schmieder et al, 2004;

- Graphical illustration of the AOP is shown on next slide.
Overview of the used Effectopedia AOP

Visualization of an example AOP cited by Schmieder*

*An AOP cited by P. Schmieder et al, 2004
Overview of the used Effectopedia AOP

Visualization of AOP built in Effectopedia

• Each AOP node is associated with detailed information for definition and relevant AOP information (such as: test, references, associated pathways etc.)

• Short explanation of the Effectopedia AOP nodes is illustrated on the next few slides
Overview of the used Effectopedia AOP
Visualization of AOP built in Effectopedia

Node correspondence is given in green highlight.
Overview of the used Effectopedia AOP
Visualization of AOP built in Effectopedia

MIE
Key events
Adverse outcome

In vitro assays
In vivo assays
Overview of the used Effectopedia AOP

Visualization of AOP built in Effectopedia

MIE

Definition of MIE

In vitro assays associated with MIE node

Definition of measurement

Associated tests

References

Continues on next slide
Overview of the used Effectopedia AOP

Visualization of AOP build in Effectopedia

 KE1

In vitro assays associated with KE1

Definition

-associated tests

Associated pathways

References

 Continues on next slide
Overview of the used Effectopedia AOP
Visualization of AOP build in Effectopedia

In vitro assays associated with KE2

Definition

Associated tests and pathways

References

KE2

KE2 definition
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Migrating the Effectopedia AOP to Toolbox

• The basic migration principle is the AOP nodes from Effectopedia AOP to coincide with the AOPs nodes from Toolbox (TB)

• Please note that this migration should be performed with good knowledge of the AOP and the key events in order to select the correct nodes in the Toolbox.
The most important TB fields needed to establish the correspondence between both type of nodes are:

- Endpoint (data to be available for the endpoint of interest)
- Data unit (unit of the endpoint data to coincide with unit of the data in Toolbox)
- Profilers (profilers related to MIEs to be available)

Preliminary analysis of the corresponding TB endpoints and profilers is performed for this purpose. Data is not available for all the Effectopedia AOP nodes. The AOP nodes mapped with the Toolbox nodes (data is available for the endpoints) are illustrated on the next few slide.
Migrating the Effectopedia AOP to Toolbox

Correspondence between nodes

**MIE**
- Receptor Binding
- ER Binding

**KE1**
- Liver Cells
- Altered Protein Expression
- Vitellogenin

**KE2**
- Liver
- Altered proteins
- Gonad
- Ova-testis;
  Sex-reversed;
  Fecundity

**KE4**
- Sex reversal;
- Altered behavior;
- Repro.
- Skewed Sex Ratios;
  Yr Class

*An AOP cited by P. Schmieder et al, 2004*
Migrating the Effectopedia AOP to Toolbox

• The migration process is a sequence of two simple steps illustrated on the next two slides.
**Step 1:** Activating the Effectopedia wizard

1. Once the Toolbox is opened, enter a chemical in the document (does not matter which one - this will not affect the exercise. The reason for this action is the endpoint tree to be visible on the screen); In this example we enter a chemical with CAS# 66-25-1;
2. Right-click over the endpoint tree;
3. Select **Activate Effectopedia Wizard**.
Migrating the Effectopedia AOP to Toolbox

Step2: Open the Effectopedia AOP

1. Click **Open Effectopedia AOP** and browse to the example folder (by default it is located here: C:\Program Files (x86)\Common Files\QSAR Toolbox 4.4\Config\Examples). An example AOP is available;
2. Select the “AOP Erbinding_New_5-Copy.aopz” file;
3. Click **Open**;

The AOP is loaded in the editor (see next slide).
Migrating the Effectopedia AOP to Toolbox

Migrated Effectopedia AOP to the Effectopedia editor
Outlook

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• **Building a new AOP-based profiler in Toolbox for predictive purposes**
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After migration of the AOP into the TB, the process continues with configuring of the AOP nodes;

Configuring is a mandatory process in order for the migrated AOP to be used for predictive purposes. Each node must be configured separately.

AOP node is associated with configuration of the following TB objects:
- Databases
- Profilers
- Simulators
- Endpoints
- Thresholds

Each AOP nodes has three states:
- Passed – chemical meets the criteria of the node
- Not passed – chemical does not meet the criteria of the node
- Not checked – rules are not applied to the chemicals
In order for the migrated AOP to be used for predictive purpose (pass/not pass) each node must be configured in terms of:

- databases/profilers (specify which databases/profilers to be used in AOP)
- selected endpoints (specify which endpoints from the selected database to be used for pass/not pass state)
- thresholds for:
  - data (specify data threshold above/less which the node to be pass or not passed)
  - profilers (specify which categories from a selected profiler to be considered for pass or not pass state)

As mentioned before, preliminary analysis of the profilers and endpoint data corresponding to the nodes of the Effectopedia AOP is done.

Mapping is possible for four out of five AOP nodes (MIE, KE1, KE2 and AO nodes).

The user should be aware of the different endpoint data and possible unit conversions in order to be able to arrange the nodes correctly.

Configuring of each node with respect to the mentioned TB objects will be shown in the next slides.
Configuring the migrated AOP

Effectopedia AOP Editor

Migration of an AOP to Toolbox is possible via the Effectopedia Editor

Name of the AOP

Nodes of the AOP

Panel with description the selected node

Save button to save the configured AOP

Settings for configuring the nodes
Configuring the migrated AOP

Node: **MIE – Estrogen receptor binding**

Solely for the purposes of the current tutorial it is proposed the following TB objectives to be considered for configuring MIE node as they are related to the Estrogen receptor binding endpoint:

1) **Relevant databases:**
   - Receptor Mediated Effects;
   - ToxCastDB;
   - Toxicity to reproduction.

2) **Relevant endpoints (associated with Toxicity to reproduction):**
   - Relative ERBA
   - EC50 <OR> IC50
   - AC50

3) **Relevant Profilers:**
   - Estrogen Receptor Binding
   - rtER Expert System – USEPA

4) **Relevant thresholds for pass/not pass (positive/negative)**
   - for profilers:
     - categories associated with activity should be passed
     - categories associated with negative should be not passed
   - for data: threshold for endpoints configured in point 2) need to be defined:
     - Data bigger than 0.001% relative ER (or AC50<0.000021 mg/l*) should be passed
     - Data less than 0.001% (or AC50>0.000021 mg/l*) should be not passed

*AC50 <2.10E-05 mg/l – arbitrary taken threshold*
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

1) Relevant databases:
- Receptor Mediated Effects;
- ToxCastDB;
- Toxicity to reproduction

1. Click the node (MIE);
2. Click **Select databases** button;
3. Select the following databases from the list: **ToxCast;** **Receptor Mediated Effects;** **Toxicity to reproduction**;
4. Click **OK**. Configuring of the databases is done.
Configuring the migrated AOP

**Node:** MIE – Estrogen receptor binding

Configuring MIE node:

2) **Relevant endpoints:**
   - Relative ERBA
   - EC50 <OR> IC50
   - AC50

For defining the endpoints a specific endpoint path should be added, see the sequence of step illustrated on the right.

1. Keep the first node (MIE) selected;
2. Click **Add** to define the endpoint tree position;
3. Select **Toxicity to reproduction**;
4. Click **Next**;
5. Select **Relative ERBA** for endpoint;
6. Click **Finish (6)**.

The first endpoint is configured, continue with second one.

**Note:** The AOP window is always minimized during the configuration of the nodes.

Continues on next slide
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

2) Relevant endpoints:
   - Relative ERBA
   - EC50 < OR > IC50
   - AC50

For defining the endpoints a specific endpoint path should be added, see the sequence of step illustrated on the right.

Repeat steps 2 and 3 from previous slide (the first step is done already);
4. Click button for defining a list;
5. Click Select;
From the predefined list with endpoints select EC50 (6,7) and IC50 (8,9);
10. Click OK;
11. Click Finish (11).
The second endpoint is configured, continue with the last one.

Continues on next slide
Configuring the migrated AOP

Node: **MIE – Estrogen receptor binding**

Configuring MIE node:

2) **Relevant endpoints:**
   - Relative ERBA
   - EC50 <OR> IC50
   - **AC50**

AC50 endpoints is associated with specific metadata information such as: assay; provider and species. For more details see the sequence of step illustrated on the right.

Repeat steps 1 and 2 from slide 29;
3. Select **ToxCast**;
4. Select endpoint **AC50**;
5. Consecutively add the following metadata: assay provider: **Novascreen**; assay: **Novascreen Bovine ER**; Test organism (species): **Bos Taurus**;
6. Click **Finish**.

The AC50 endpoint associated with assay “Novascreen Bovine ER” is configured. Moreover two additional assays related to AC50 endpoint need to be defined – see next slide.

Continues on next slide
Configuring the migrated AOP

**Node:** MIE – Estrogen receptor binding

Configuring MIE node:

2) **Relevant endpoints:**
   - Relative ERBA
   - EC50 <OR> IC50
   - **AC50**

AC50 endpoints is associated with specific metadata information such as: assay; provider and species. For more details see the sequence of step illustrated on the right.

Repeat steps 1 and 2 and 3 from previous few slides including selecting ToxCast level. Once ready with selection of AC50 endpoint add the following metadata:

4a. assay: **Novascreen Human ER**; species: **Homo sapiens** and one more assay:

4b. assay: **Novascreen Mouse ERα**; species: **Mus musculus**.

The difference between previous selection is the assay and test organism(species) fields;

5. Click **Finish**.
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

All the information added to nodes is displayed in the panel. The next step is to specify the relevant profilers and to add thresholds for profilers and data (see next slides).
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

3) Relevant profilers:
   - *Estrogen Receptor Binding*
   - *rtER Expert System – USEPA*

1. Keep the first node selected;
2. Click **Select profilers**;
3. Select the profilers from the list: *Estrogen Receptor Binding; rtER Expert System – USEPA*;
4. Click **OK**.
Configuring the migrated AOP

Node: **MIE – Estrogen receptor binding**

Configuring MIE node:

4) **Threshold for the relevant profilers:**

**Profiler: Estrogen Receptor binding**

- **Passed:**
  - Strong binder, NH2 group;
  - Strong binder, OH group;
  - Very strong binder, OH group;
  - Moderate binder, NH2 group;
  - Moderate binder, OH group;
  - Weak binder, NH2 group;
  - Weak binder, OH group

- **Not passed:**
  - No OH, NH-group
  - Non binder, impaired OH or NH2 group
  - Non binder, MW>500
  - Non binder, non cyclic structure
  - Non binder, without OH or NH2 group
  - Auxiliary categories (e.g. Profiling timeout)

First node (MIE) is selected (1); Click Add (2) in the threshold section; Click on **Profiler** (3) radio button; Select “**Estrogen Receptor Binding**” profiler from the list (4); Select all categories related to the “pass” condition of the node (usually these are all categories related with activation). Select all by hold **Ctrl button** and click the **arrow button** (6), to move them in the panel “Passed”. Repeat the same procedure with all categories associated with “not pass” condition of the node. They are moved to the panel “Not passed” (7); Click **OK** (8).

Note: A category can be removed from the list by double click.
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

4) Threshold for the relevant profilers:

Profiler: rtER Expert System – USEPA

- **Passed:**
  - all categories except No alert found and Auxiliary categories (e.g. Profiling timeout)

- **Not passed:**
  - No alert found category
  - Auxiliary categories (e.g. Profiling timeout)

Repeat the same procedure as described in the previous slide, but related to the profiler "rtER Expert System – USEPA". The snapshot illustrates already configured profiler. Keep in mind that all categories except "No alert found" and auxiliary categories (e.g. Profiling timeout) are moved to “Passed” panel. Once ready click OK.
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

4) Data thresholds:

ER data is associated with different scales. In this respect configuration is performed for each individual scale:

Scale: “Estrogen Binding Affinity(USA)”
  - Passed: RBA>0.00001%
  - Not passed: RBA<0.00001%

1. Select node MIE;
2. Click Add in the threshold section;
3. Select radio button “Ordinal”.
4. Select scale “Estrogen Binding Affinity(USA)” from the list with units.
5. Select “RBA>0.00001%” from the drop-down menu (6).
7. Click “V” button to moved it to the “Passed” panel (8).
Configuring the migrated AOP

Node: **MIE – Estrogen receptor binding**

Configuring MIE node:

4) **Data thresholds:**

ER data is associated with different scales. In this respect configuration is performed for each individual scale:

**Scale: “Estrogen Binding Affinity (USA)”**

- **Passed:**
  - RBA > 0.00001%

- **Not passed**
  - RBA < 0.00001%

Select “**RBA<0.00001%**” (1) from the drop-down menu (2). Click “**V**” button (3) to moved it to the **Not Passed** panel (4). Finally click **OK** (5).
Configuring the migrated AOP

Node: MIE – Estrogen receptor binding

Configuring MIE node:

4) Data thresholds:
ER data is associated with different scales. In this respect configuration is performed for each individual scale.

Scale: Estrogen Binding Affinity (OASIS II):

Passed:
- 0.001 < RBA < 0.1%
- 0.1 < RBA < 10%
- 0 < RBA < 0.001%
- RBA > 10%

Not passed:
- Not active

Scale: "Mass concentration":

Passed:
- (0 - 0.000021] mg/L

Not passed:
- (0.000021 - 100] mg/L

Repeat steps from 1-2 from previous slide. The "Ordinal" radio button should be selected, when scale "Estrogen Binding Affinity (OASIS II)" is configured and "Ratio" when scale "Mass concentration" is used. Then add the members of the scale to passed and not passed panels as shown on the picture. Keep in mind for sign of the ranges when define members of the "Mass concentration scale" (the box “Open” should be checked when the interval is open “]”) and not checked when the interval is closed “[”). Data in “Mass concentration” use “mg/l” unit. Once ready click OK. Configuring the node with the thresholds is ready.
Configuring the migrated AOP

Congratulations! Node MIE is configured

Relevant databases:
- Receptor Mediated Effects
- ToxCastDB
- Toxicity to reproduction (ER)

Associated endpoint tree positions:
- Human Health Hazards > Toxicity to Reproduction
- Endpoint=Relative ERBA
- Human Health Hazards > Toxicity to Reproduction
- Endpoint=EC50 < OR > IC50
- Human Health Hazards > ToxCast
- Assay=Novascreen
- Assay=Novascreen Bovine ER
- Test organisms (species) = Bos taurus
- Endpoint=AC50
- Human Health Hazards > ToxCast
- Endpoint=AC50
Configuring the migrated AOP

Node KE1 - Cellular response: Altered protein expression: increasing Vitellogenin synthesis

The following TB objectives need to be considered for configuring KE1 node in order to work correctly:

1) Relevant databases:
   - ECOTOX;

2) Relevant endpoints (related to Acute aquatic toxicity endpoint tree):
   - LOEC:
     - Effect: Biochemistry
     - Measurement: Vitellogenin
     - Trend: Increasing
   - NOEC:
     - Effect: Biochemistry
     - Measurement: Vitellogenin
     - Trend: Increasing

3) Relevant Profilers:
   - No relevant profilers

4) Relevant thresholds for pass/not pass (positive/negative)
   - Chemicals with data in the range [0 - 1) mg/L should be passed
   - Chemicals with data in the range [1 - 1000] mg/l should be not passed

Note: The final outcome of the configuring the nodes is illustrated in the forthcoming slides. For configuring the nodes please use the steps illustrated on the previous slides.
Configuring the migrated AOP

Node KE1 - Cellular response: Altered protein expression: increasing Vitellogenin synthesis

Add the relevant database

Add the relevant endpoint tree positions. Keep in mind two additional metadata fields called "Measurement" and "Trend" are added to the list with default fields.
Configuring the migrated AOP

Node KE1 - Cellular response: Altered protein expression: increasing Vitellogenin synthesis

Add the relevant thresholds by using “Add” button. The scale is “Mass concentration”, so it is “ratio” type. Add the both ranges to the respective panels (pass/not passed). The defined settings appears in the panel.
Configuring the migrated AOP

Node KE2 - Tissue/Organ: Liver Altered proteins

The following TB objectives need to be considered for configuring KE2 node in order to work correctly:

1) Relevant databases:
   • ECOTOX

2) Relevant endpoints (related to Acute aquatic toxicity endpoint tree):
   • LOEC:
     • Effect: Reproduction
     • Measurement: Fecundity
     • Trend: Decreasing
   • NOEC:
     • Effect: Reproduction
     • Measurement: Fecundity
     • Trend: Decreasing

3) Relevant Profilers:
   • No relevant profilers

4) Relevant thresholds for pass/not pass (positive/negative) (same as in KE1)
   • Chemicals with data in the range [0 - 1) mg/L should be passed
   • Chemicals with data in the range [1 - 1000] mg/l should be not passed

Note: The final outcome of the configuring the nodes is illustrated in the forthcoming slides.
Configuring the migrated AOP

Node KE2 - Tissue/Organ: Liver Altered proteins

Relevant database: “ECOTOX”

Relevant endpoints: LOEC and NOEC

Relevant thresholds

Associated endpoint tree positions: Ecotoxicological Information#Aquatic Toxicity
End point=LOEC
Effect=Reproduction
Measurement=Fecondity
Trend=Decreasing

Information#Aquatic Toxicity
Effect=Reproduction
Measurement=Fecondity
Trend=Decreasing

Thresholds:
Scale name: Mass concentration;
Passed: [0 - 1] mg/L;
Not passed: [1 - 1000] mg/L;
Scale type: RatioScale
Configuring the migrated AOP

Node KE3 - Individual: Sex reversal

• This Effectopedia AOP node is not configured, because there are no Toolbox data available and relevant profilers.

• Node: keep it as it is.
Configuring the migrated AOP

Node AO - Population: Skewed sex ratios

The following TB objectives need to be considered for configuring AO node in order to work correctly:

1) Relevant databases:
   • ECOTOX

2) Relevant endpoints (related to Acute aquatic toxicity endpoint tree):
   • LOEC:
     • Effect: Population
     • Measurement: Sex ratio
     • Trend: Change
   • NOEC:
     • Effect: Population
     • Measurement: Sex ratio
     • Trend: Change

3) Relevant Profilers:
   • No relevant profilers

4) Relevant thresholds for pass/not pass (positive/negative) (same as in KE1 and KE2)
   • Chemicals with data in the range [0 - 1) mg/L should be passed
   • Chemicals with data in the range [1 - 1000] mg/l should be not passed

Note: The final outcome of the configuring the nodes is illustrated in the forthcoming slides.
Configuring the migrated AOP

**Node AO - Population: Skewed sex ratios**

- **Relevant database:** “ECOTOX”

- **Relevant endpoints:** LOEC and NOEC

- **Relevant thresholds**
Configuring the migrated AOP

All AOP nodes are configured in Toolbox and ready to be used for predictive purposes. Finish the process by saving the AOP

1. Click Save button (1);
2. Browse to the folder on your computer (2) (remember the folder, you will need it (see next slide)), gave name of the custom AOP (3);
3. Click Save (4);
4. Close the window of the AOP editor (5).
Configuring the migrated AOP

In order to use already configured AOP you need to:

1) paste the saved AOP file (*.aoptb) in the following installation folder: C:\Program Files (x86)\QSAR Toolbox\QSAR Toolbox 4.4\Toolbox Client\Bin\AOP (by default)
2) restart the Toolbox client
3) start the Classical User Interface (3)
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Application of the migrated AOP

Target chemical:
CAS: 104-40-5
Name: 4-Nonylphenol

Usage: laundry and dish detergents, pesticides, care body products and plastics

1. Enter the target by CAS;
2. Blue AOP sign appears on the “Toxicity to reproduction” level and right-click on the sign
3. Select Activate AOP (3) in the pop-up menu.
Once the AOP tree appears follow the steps:
1. Click on node MIE;
2. Go to Profiling module;
3. Select both green profilers;
4. Click Apply.
1. Go to **Data** module;
2. Select databases highlighted in green;
3. Click **Gather**;
4. Click **OK**;
5. A pop-up message informs about the number of collected data. Click **OK**.
1. Click node KE1: Cellular response: Altered proteins
2. Go to Data module;
3. Select the database highlighted in green;
4. Click Gather;
5. Click OK;
6. A pop-up message informs about the number of collected data. Click OK.
Application of the migrated AOP

- Nodes KE1 is colored in red because experimental data is found for the target chemical in ECOTOX;
- KE2 and AO are also colored in red, because experimental data is found for the target chemical in the same database (ECOTOX);
- As a result four AOP nodes (MIE; KE1; KE2 and AO) are passed (e.g. the boundary rules implemented in the nodes are “passed”)

![Diagram of the migrated AOP process with nodes colored and passed status indicated.]
Application of the migrated AOP

Outcome of AOP application

• Migrated AOP nodes have been applied to the target chemical. Each node has been executed individually.

• As a result four out of five AOP nodes are passed (MIE; KE1; KE2 and AO).

• In other words, an ER binding alerts related to MIE and experimental data for almost all KEs has been found for the target chemical.

• Based on the outcome of the AOP nodes, it could be concluded that the target may elicit positive estrogen receptor binding effect.