

QSAR Application Toolbox

Workflow

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Outlook

Description

General scheme and workflow
Basic functionalities
Forming categories

Philosophy

QSAR TOOLBOX

- Toolbox helps registrants and authorities to
 - ✓ Use the methodologies to group chemicals into categories and
 - ✓ Refine and expand the categories approach
 - ✓ Provide a mechanistic transparency of the formed categories
 - ✓ Fill data gaps by read-across, trend analysis and (Q)SARs
 - ✓ Ensure uniform application of read-across
 - ✓ Support the regulatory use of (Q)SAR approach
 - ✓ Improve the regulatory acceptance of (Q)SAR methods

Philosophy

QSAR TOOLBOX

• (Q)SAR Toolbox is a central tool for non-test data in ECHA and OECD

• ECHA and OECD coordinate the Toolbox development

Description

QSAR TOOLBOX

- Toolbox is not a QSAR model and hence can not be compared with other QSAR models
- Training sets (categories) for each prediction are defined dynamically as compared to other (Q)SAR model which have rigid training sets
- Each estimated value can be individually justified based on:
 - ✓ Category hypothesis (justification) and consistency
 - ✓ Quality of measured data and
 - ✓ Computational method used for grouping and data gap filling

Description

QSAR TOOLBOX

- It is developed under the continuing peer review of:
 - ✓ Member state countries,
 - ✓ Regulatory agencies
 - ✓ Chemical industry and NGOs
- The predictions are getting acceptance by toxicologists and regulators due to the:
 - ✓International peer review process for developing the system and
 - ✓ Mechanistic transparency of the results
- The system is freely available and maintained in the public domain by OECD

History

Phase I - The first version (2005 - 2008)

- Emphasizes technological proof-of-concept
- Released in 2008
- Developed by LMC

Phase II (2008 - 2012)

- More comprehensive Toolbox which fully implements the capabilities of the first version
- Second version was released in 2010
- Developed by LMC, subcontractors: LJMU, Lhasa Ltd and TNO
- Third version was released in 2012
- Maintenance 2013 v. 3.2 in January 2014
- Version 3.3.1 was released in December 2014
- Version 3.3.2 was released in February 2015
- Version 3.3.3 is expected to be released in June 30, 2015

Phase III (2013 – 2019 + 4 years maintenance)

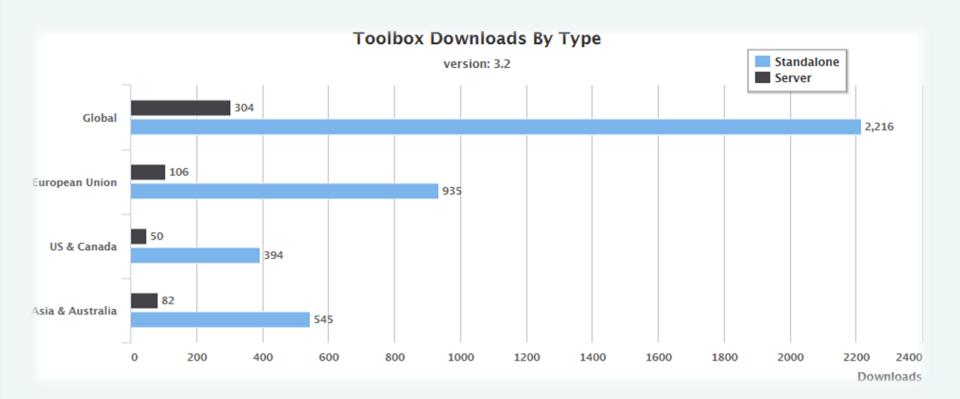
- Significant focus on streamlining + new IT platform
- Knowledge and data rationalization and curation
- Implementation of Ontology, ADME
- Implementation of AOPs
- Version 4.0 will be released in spring of 2016
- Developed by LMC, subcontractors: LJMU and Lhasa Ltd

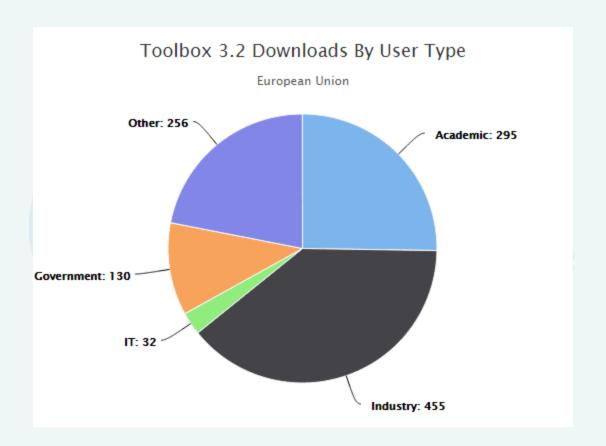
Main Government Donators

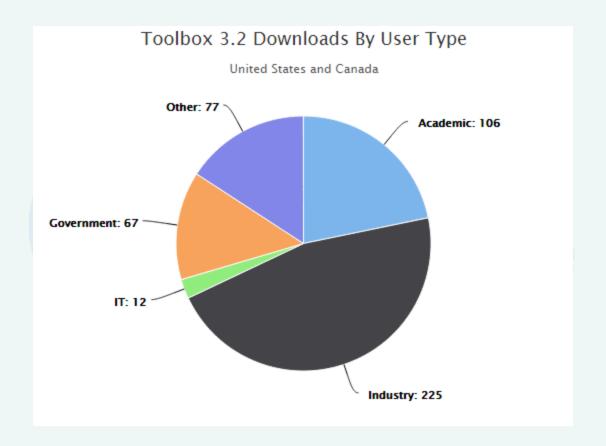
- •OECD
- •European chemicals agency
- •US EPA
- •Environment Canada
- •Health Canada
- •NITE Japan
- •NIES Japan
- •Danish EPA
- •UBA Germany
- •NICNAS Australia
- •DEWNA Australia
- •ISS Italy

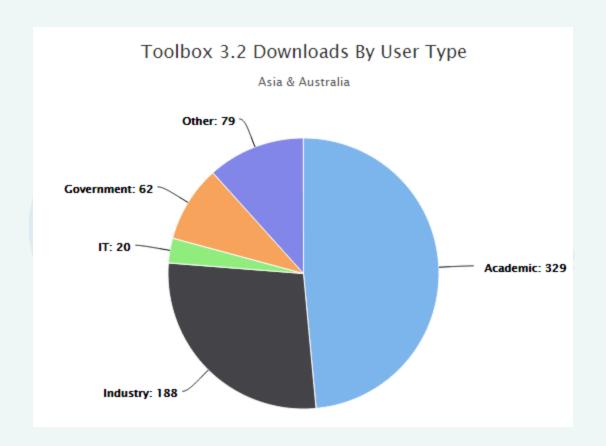
Main Industry Donators

- •L'Oreal
- •DuPont
- •Givaudan
- Dow chemicals
- •BASF
- ExxonMobil
- •3M
- •Firmenich SV
- •SRC, Syracuse
- •Unilever



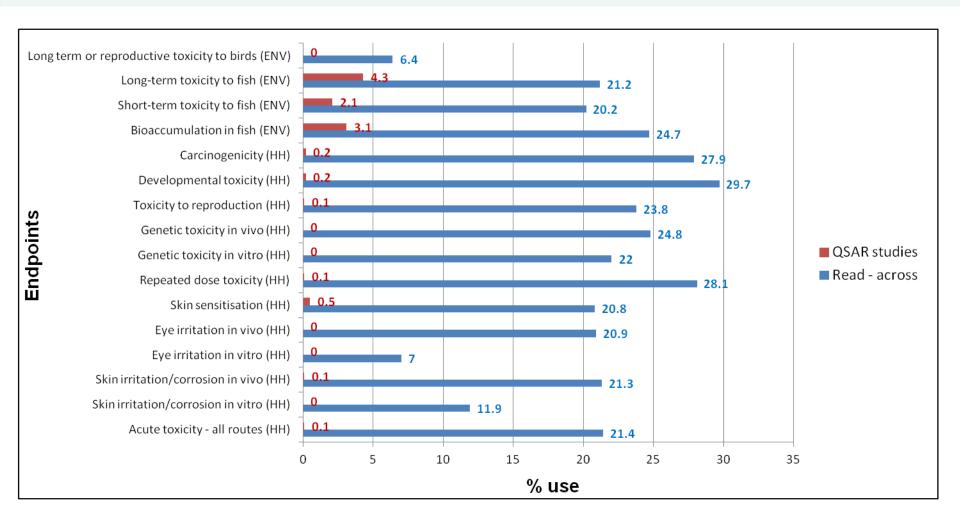






QSAR and read-across based submissions to the ECHA

For existing substances at or above 1000 tpa* (2011 ECHA report)





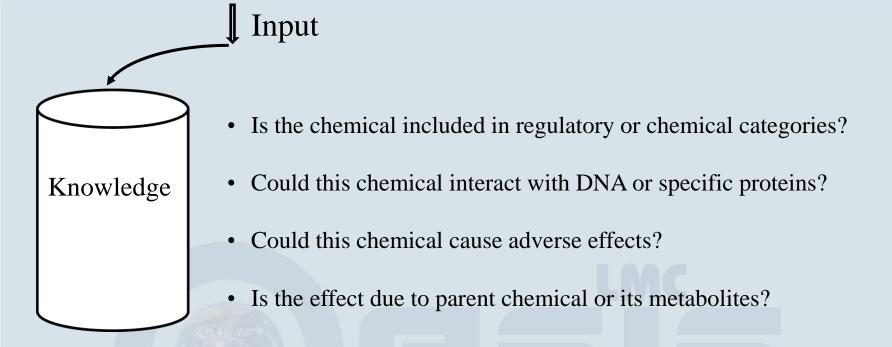
Outlook

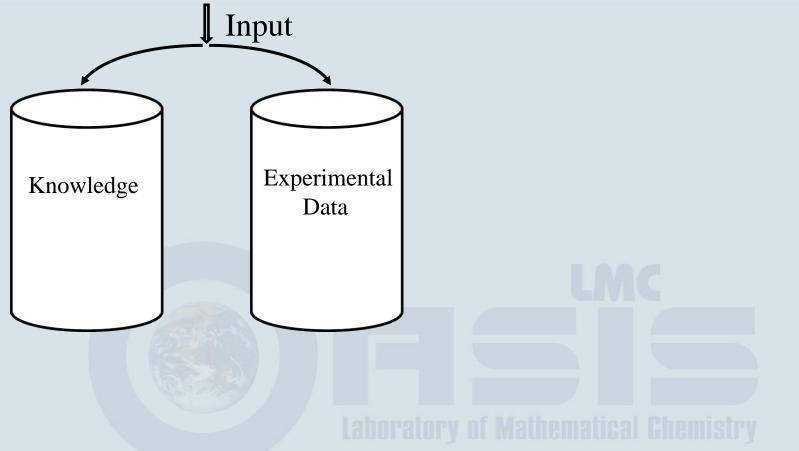
Description
General scheme and workflow
Basic functionalities
Forming categories

General Scheme

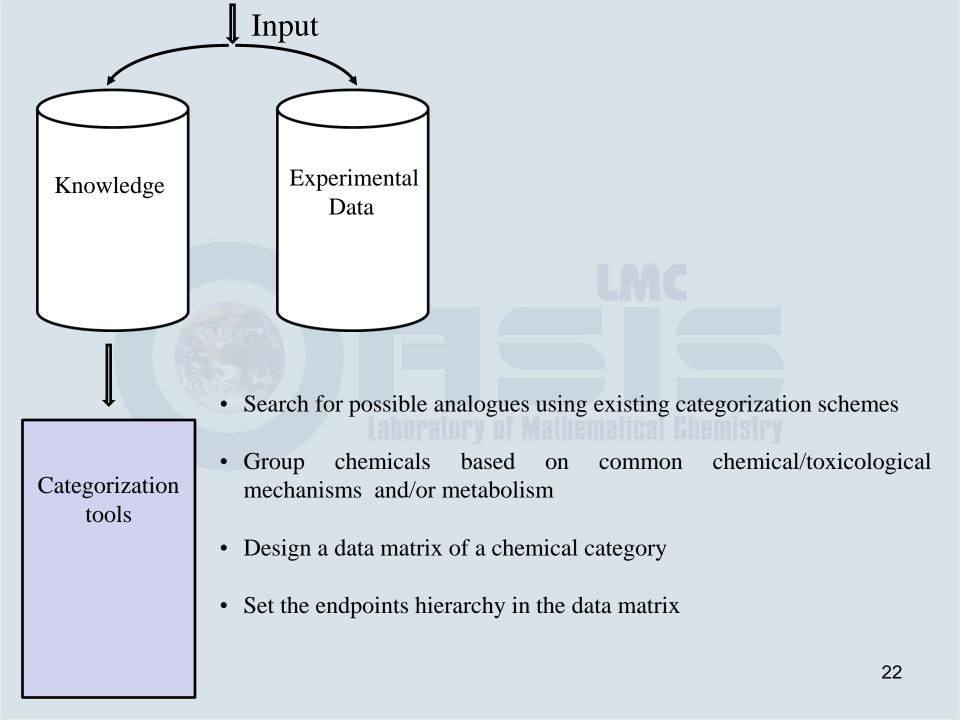
- •Theoretical knowledge,
- •Empiric knowledge,
- •Computational methods,
- •Information technologies.

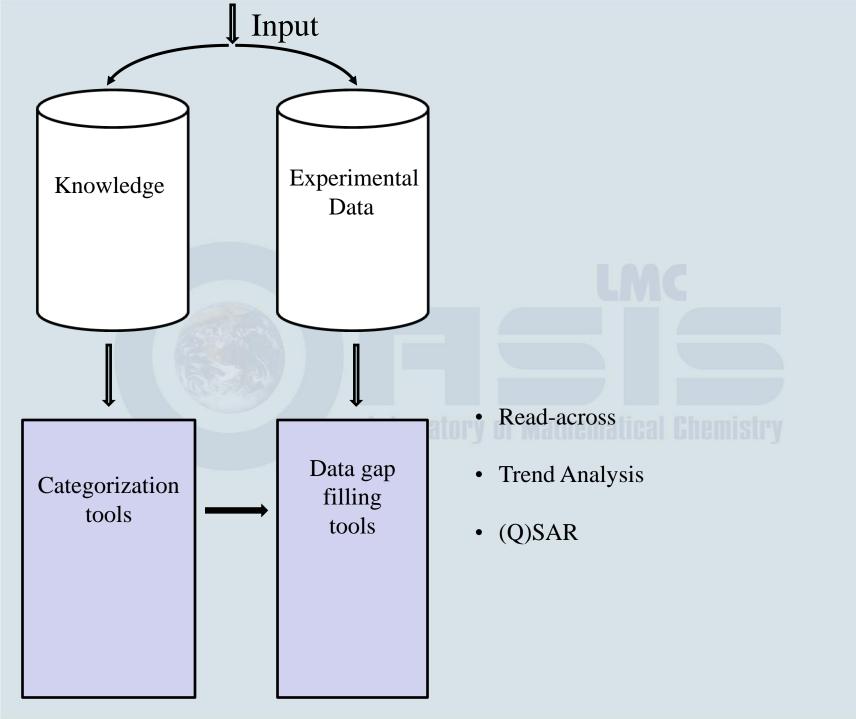


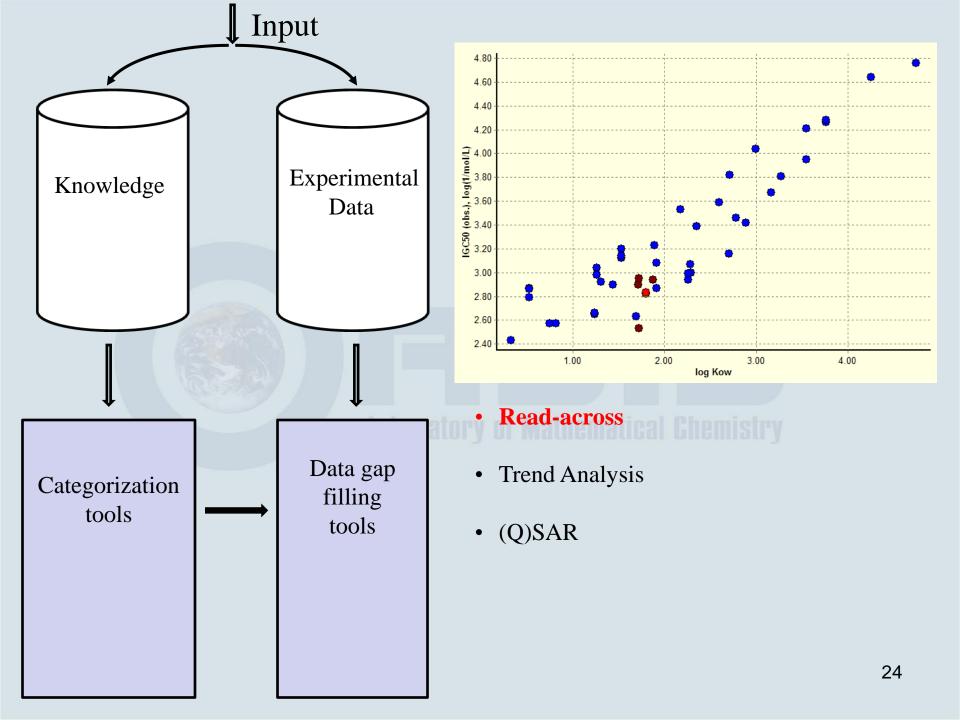


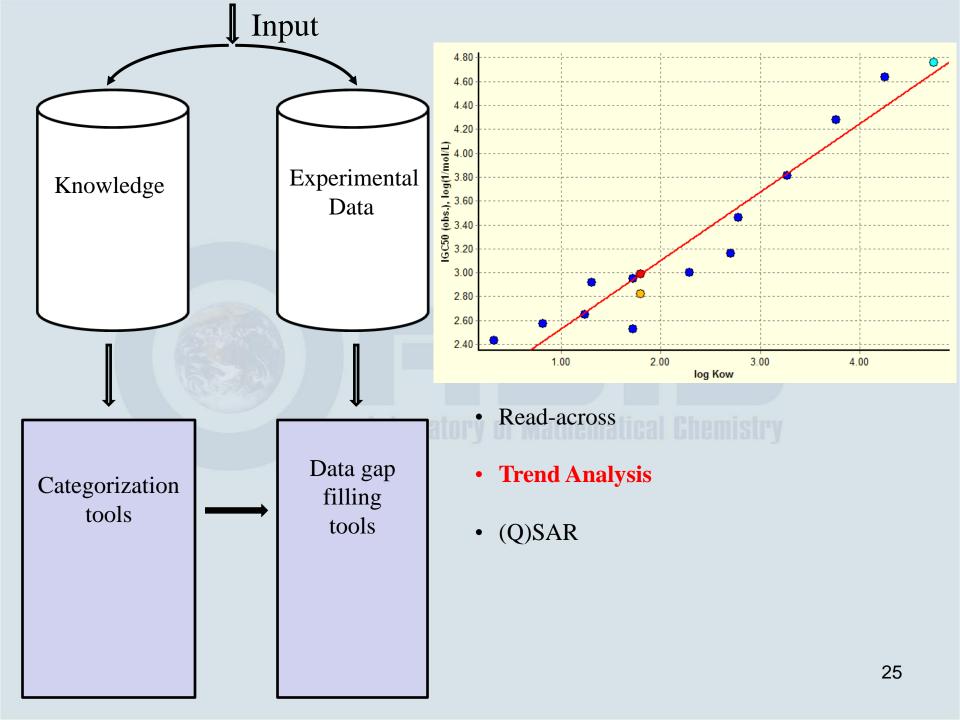


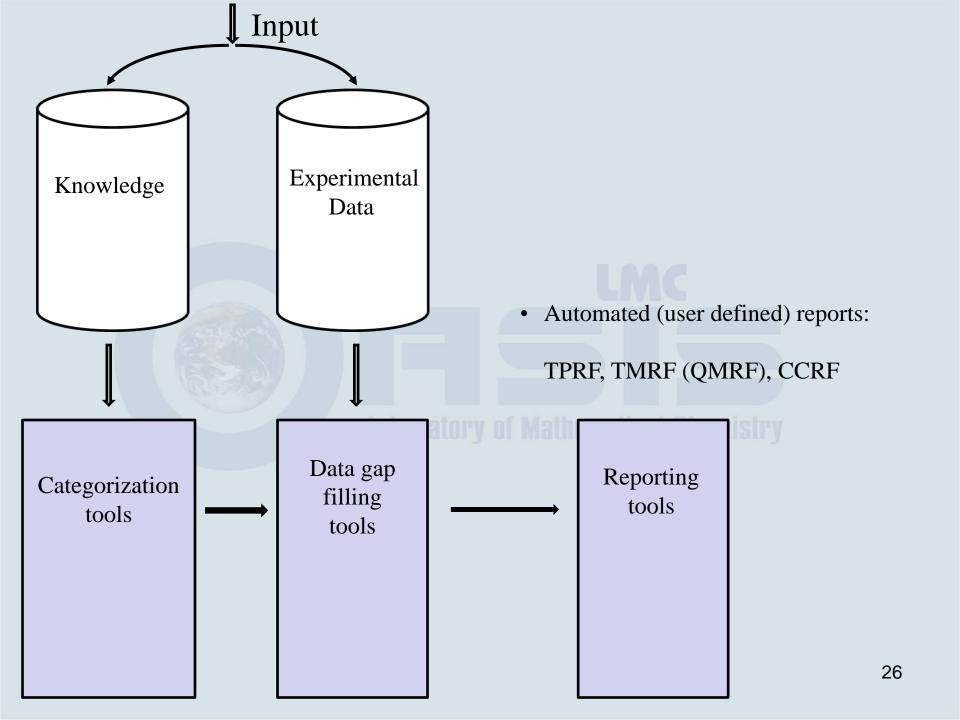
- Are data available for assessed endpoints of target chemical?
- Is information for the data sources available?

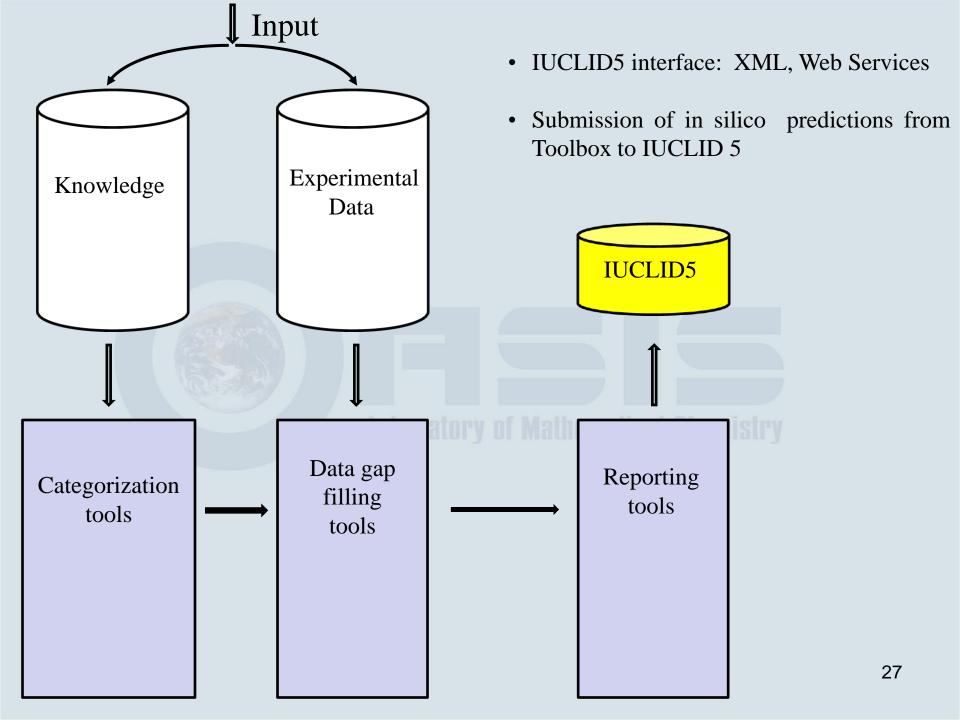


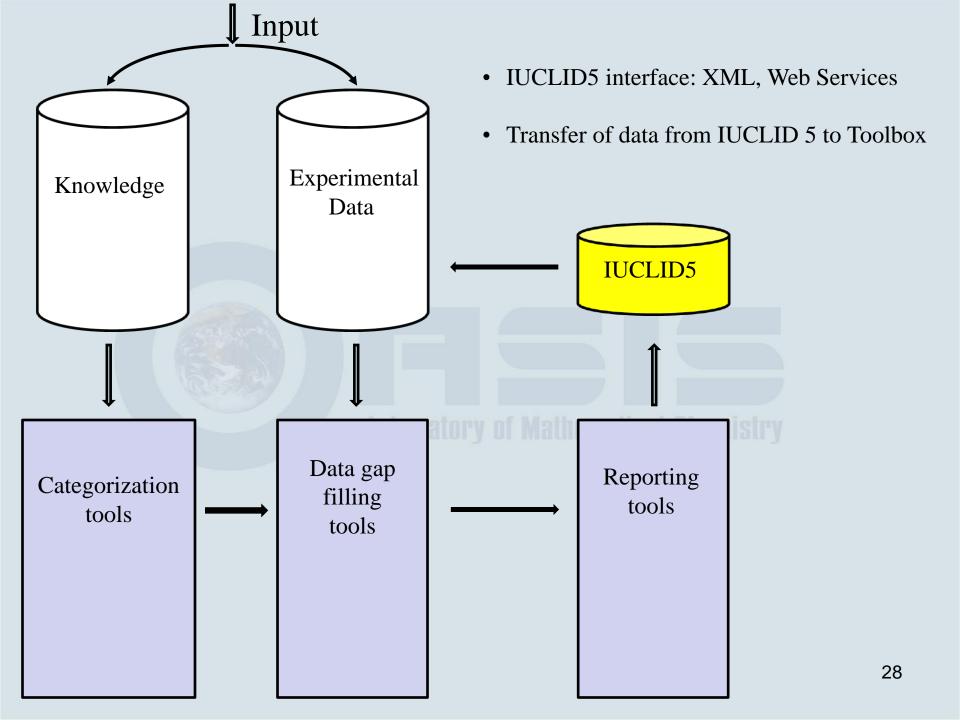


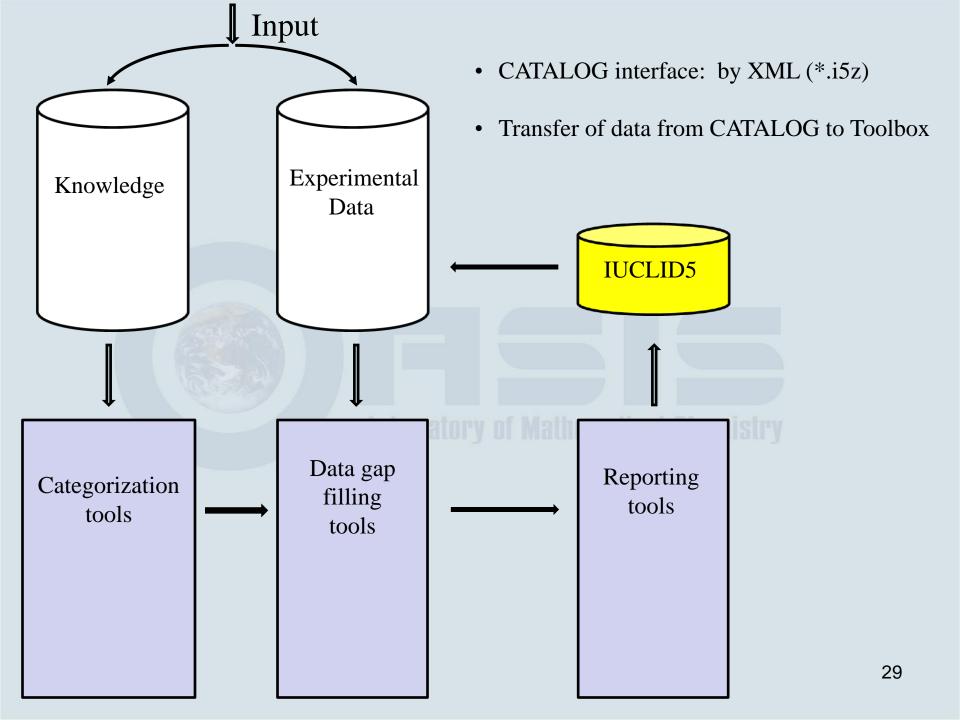




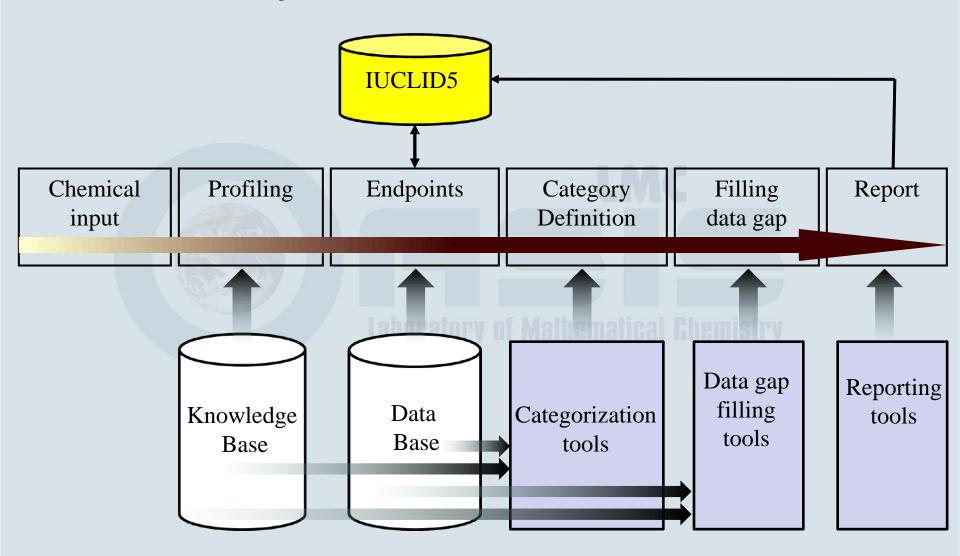








System Workflow

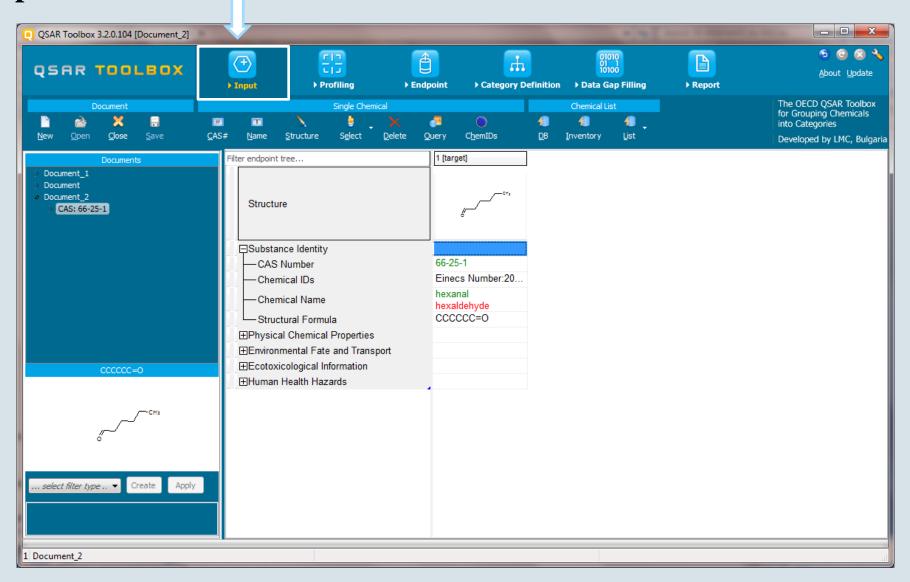




Outlook

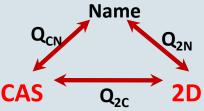
Description
General scheme and workflow
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Input



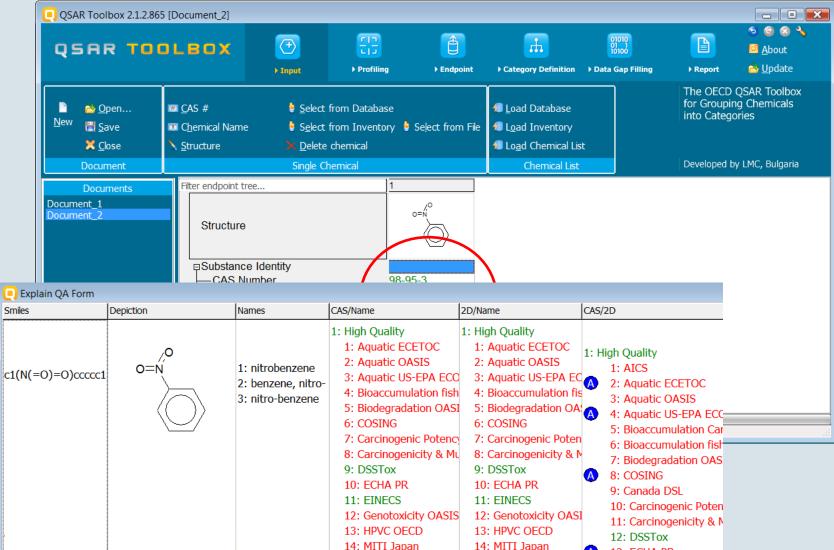
Input

QA of chemical structures

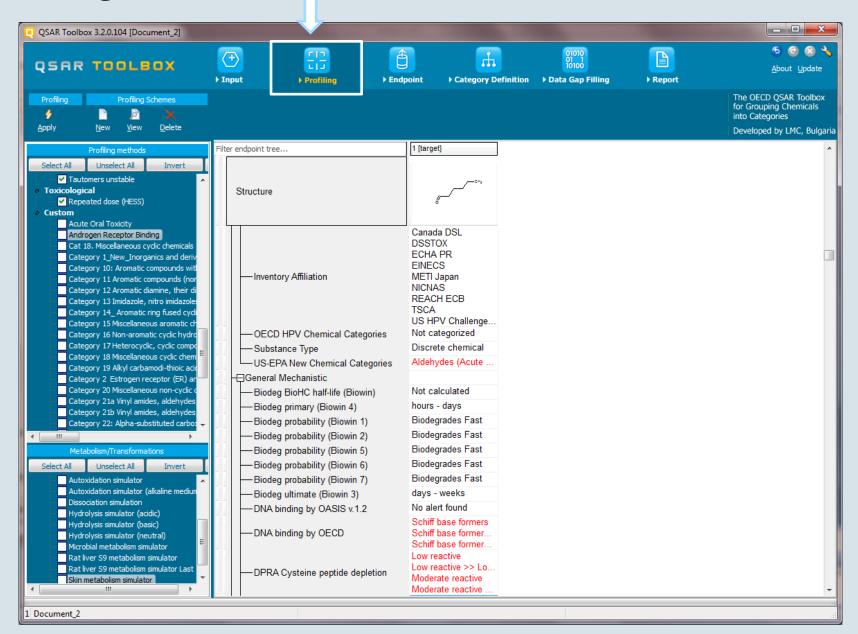


13: ECHA PR

15: Phys-chem FDISH



15: Phys-chem FDISHIT



- Predefined
- General Mechanistic
- Endpoint Specific
- Empiric
- Custom



- Predefined
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- Endpoint Specific
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- ✓ Database Affiliation
- ✓ Inventory Affiliation
- ✓ OECD HPV Chemical Categories
- ✓ Substance Type
- US-EPA New Chemical Categories



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General Mechanistic

- ☑ Biodeg BioHC half-life (Biowin)
- ☑ Biodeg primary (Biowin 4)
- ☑ Biodeg probability (Biowin 1)
- ☑ Biodeg probability (Biowin 2)
- ✓ Biodeg probability (Biowin 5)
- ☑ Biodeg probability (Biowin 6)
- ☑ Biodeg probability (Biowin 7)
- ☑ Biodeg ultimate (Biowin 3)
- ✓ DNA binding by OASIS v.1.1
- ✓ DNA binding by OECD
- ✓ DPRA Cysteine peptide depletion
- ✓ DPRA Lysine peptide depletion
- Estrogen Receptor Binding
- ✓ Hydrolysis half-life (Ka, pH 7)(Hydrowin)
- ✓ Hydrolysis half-life (Ka, pH 8)(Hydrowin)
- Hydrolysis half-life (Kb, pH 7)(Hydrowin)
- ✓ Hydrolysis half-life (Kb, pH 8)(Hydrowin)
- ✓ Hydrolysis half-life (pH 6.5-7.4)
- Ionization at pH = 1
- ✓ Ionization at pH = 4
- ✓ Ionization at pH = 7.4
- ✓ Ionization at pH = 9
- ✓ Protein binding by OASIS v1.1
- ✓ Protein binding by OECD
- ✓ Protein binding potency
- Superfragments
- ✓ Toxic hazard classification by Cramer (original)
- ✓ Toxic hazard classification by Cramer (with extensions)
- ✓ Ultimate biodeg

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Endpoint Specific

- ✓ Acute aquatic toxicity classification by Verhaar
- ✓ Acute aquatic toxicity MOA by OASIS
- ✓ Aquatic toxicity classification by ECOSAR.
- ☑ Bioaccumulation metabolism alerts
- ☑ Bioaccumulation metabolism half-lives
- Biodegradation fragments (BioWIN MITI)
- ✓ Carcinogenicity (genotox and nongenotox) alerts by ISS
- DNA alerts for AMES, MN and CA by OASIS v.1.1
- Eye irritation/corrosion Exclusion rules by BfR.
- Eye irritation/corrosion Inclusion rules by BfR.
- ☑ in vitro mutagenicity (Ames test) alerts by ISS
- ☑ in vivo mutagenicity (Micronucleus) alerts by ISS
- ✓ Keratinocyte gene expression
- ✓ Oncologic Primary Classification
- ✓ Protein binding alerts for skin sensitization by OASIS v1.1
- ✓ rtER Expert System ver.1 USEPA
- ☑ Skin irritation/corrosion Exclusion rules by BfR.
- ☑ Skin irritation/corrosion Inclusion rules by BfR.

Profiling

- Predefined
- General Mechanistic
- Endpoint Specific
- Empiric
- Custom



Empiric

- ✓ Chemical elements
- ✓ Groups of elements
- Lipinski Rule Oasis
- ✓ Organic functional groups
- ✓ Organic functional groups (nested)
- ✓ Organic functional groups (US EPA)
- ☑ Organic functional groups, Norbert Haider (checkmol)
- ▼ Tautomers unstable

Toxicological

Repeated dose (HESS)

Custom

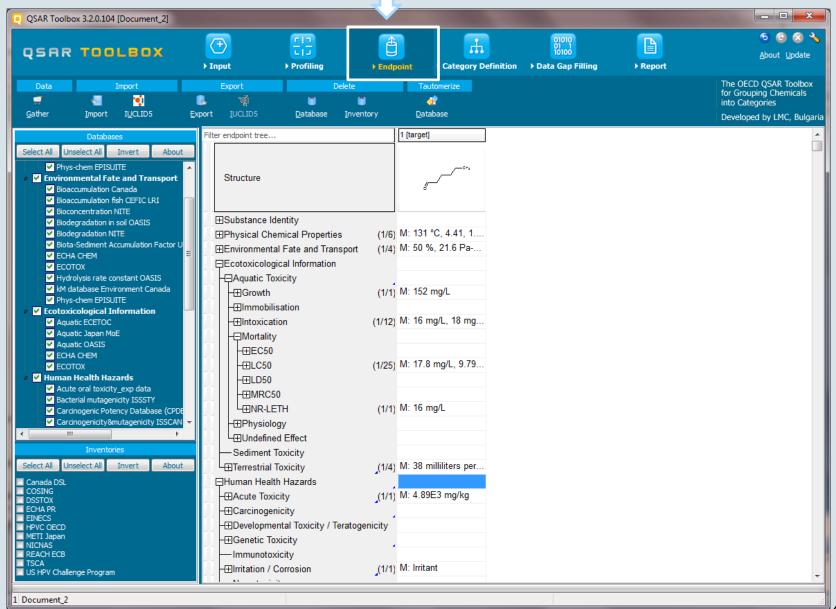
- ✓ aldehydes
- Anilines

Profiling

- Predefined
- General Mechanistic
- Endpoint Specific
- Empiric
- Custom

Molecular transformations

Metabolism/Transformations Documented Observed Mammalian metabolism Observed Microbial metabolism Observed Rat In vivo metabolism Observed Rat Liver S9 metabolism ■ Simulated Autoxidation simulator Autoxidation simulator (alkaline medium) Dissociation simulation Hydrolysis simulator (acidic) Hydrolysis simulator (basic) ■ Hydrolysis simulator (neutral) Microbial metabolism simulator Rat liver S9 metabolism simulator Skin metabolism simulator



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- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazard



- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazard

- Physical Chemical Properties
 - Chemical Reactivity COLIPA
 - ECHA CHEM
 - Experimental pKa
 - ☐ GSH Experimental RC50
 - Phys-chem EPISUITE



- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazard



Environmental Fate and Transport

- Aquatic US-EPA ECOTOX
- Bioaccumulation Canada
- Bioaccumulation fish CEFIC LRI
- Bioconcentration NITE
- Biodegradation in soil OASIS
- Biodegradation NITE
- Biota-Sediment Accumulation Factor US-EPA
- ECHA CHEM
- Hydrolysis rate constant OASIS
- kM database Environment Canada
- Phys-chem EPISUITE
- Terrestrial US-EPA ECOTOX

aboratory of Mathematical Chemistry

- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazard



- Aquatic ECETOC
- Aquatic Japan MoE
- Aquatic OASIS
- Aquatic US-EPA ECOTOX
- ECHA CHEM
- ☐ Terrestrial US-EPA ECOTOX

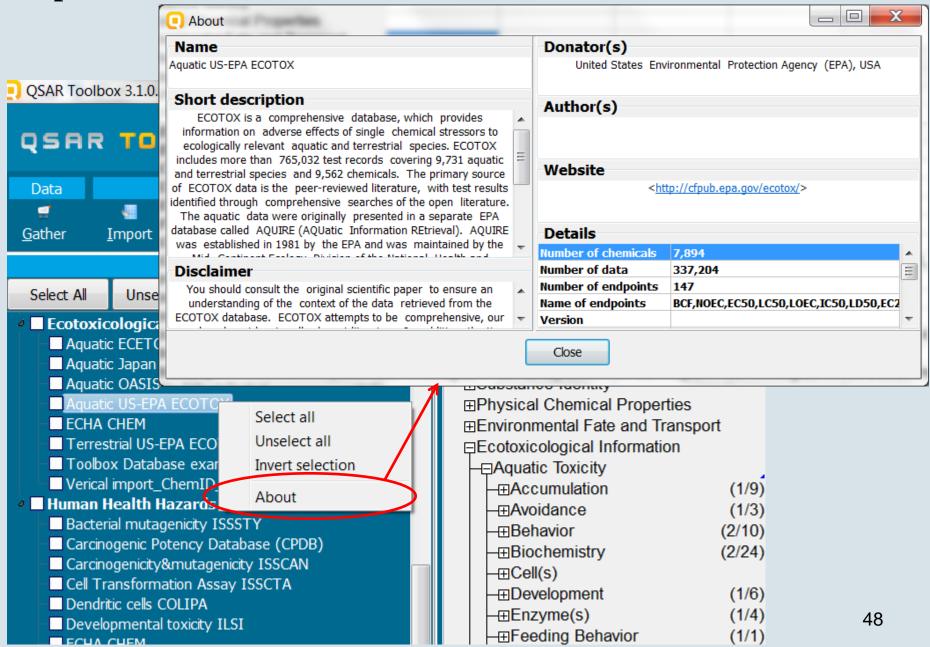


- Physical Chemical Properties
- Environmental Fate and Transport
- Ecotoxicological Information
- Human Health Hazard



■ Human Health Hazards

- Bacterial mutagenicity ISSSTY
- Carcinogenic Potency Database (CPDB)
- Carcinogenicity&mutagenicity ISSCAN
- Cell Transformation Assay ISSCTA
- Dendritic cells COLIPA
- Developmental toxicity ILSI
- ECHA CHEM
- Estrogen Receptor Binding Affinity OASIS
- Eye Irritation ECETOC
- Genotoxicity OASIS
- Keratinocyte gene expression Givaudan
- Micronucleus ISSMIC
- Micronucleus Oasis
- MUNRO non-cancer EFSA
- Rep Dose Tox Fraunhofer ITEM
- Repeated Dose Toxicity HESS
- Rodent Inhalation Toxicity Database
- Skin irritation
- Skin sensitization
- Skin sensitization ECETOC
- Terrestrial US-EPA ECOTOX
- Toolbox Database example
- Toxicity Japan MHLW
- ToxRefDB US-EPA
- Yeast estrogen assay database University of Tennessee-Knoxville (USA)



Task C - ECHA CHEM database

C. Expansion of the repertoire of databases including data content normalization

Development of methodology to explore ECHA CHEM database

QSAR Toolbox datand ECHA CHEM contribution

Endpoint tree	QSAR Toolbox (without ECHA CHEM)		
	Substances	Data	
Phys-chem properties	38 000	71 000	
Environmental	4 500	58 000	
Ecotoxicological	8 500	510 000	
Human health	15 000	550 000	

ECHA CHEM contribution		
Substances	Data	
5 700 (15%)	85 000 (120%)	
4 000 (88%) 28 000 (48%)		
4 600 (54%)	4%) 95 000 (18%)	
5 400 (36%)	140 000 (25%)	

Total	55 000	1 200 000

6 200 (9%)	350 000 (29%)
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QSAR Toolbox databases

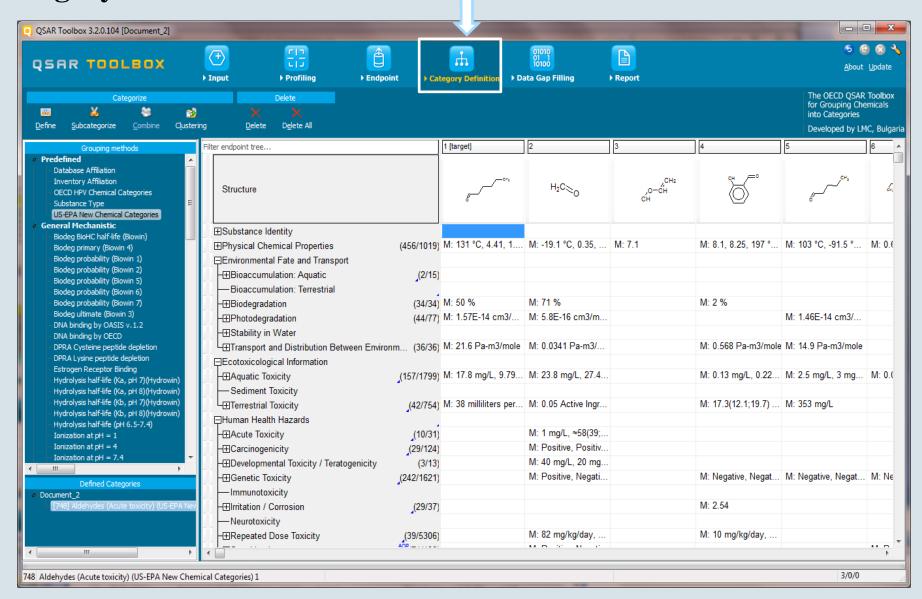
Contribution of REACH dissemination data ECHA CHEM database

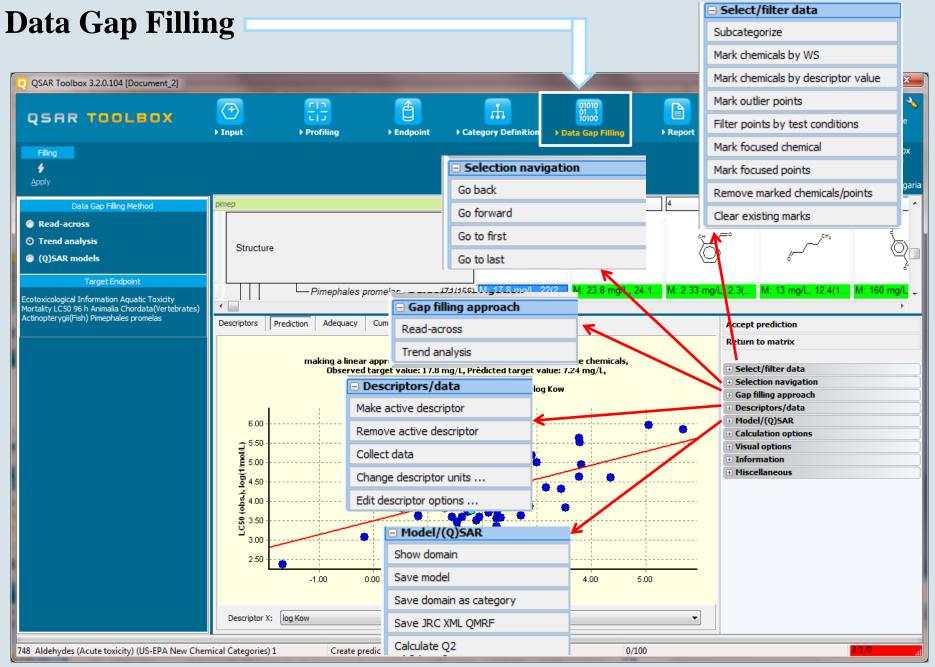
Endpoint tree	ECHA CHEM		Total	
Enapoint tree	Substances	Data	Substances	Data
Substance Identity	6188	301 868	92 854	1 538 014
Physical Chemical Properties	5 738	86 453	41931	156 528
Environmental Fate and Transport	4 001	25 759	7581	85 585
Ecotoxicological Information	4 626	94 009	11 895	608 065
Human Health Hazards	5 426	95 647	18 813	687 836

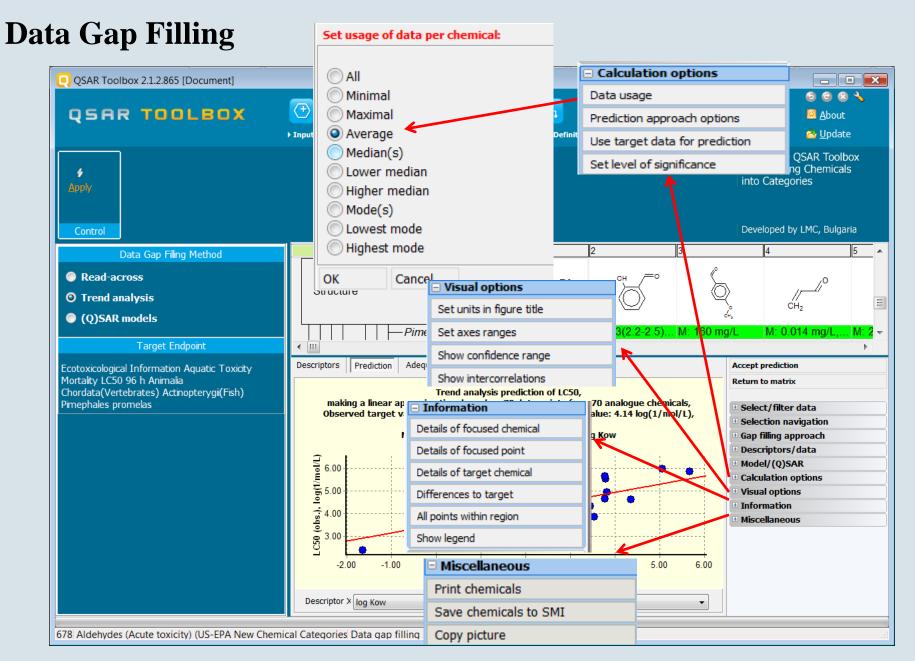
Inventories

Inventories	Number of chemicals
AICS	38760
COSING	1314
DSSTOX	8606
ECHA PR	142619
EINECS	72561
HPVC EU	4843
METI Japan	16133
REACH ECB	74074
US HPV Challenge Program	9125
TSCA 2005	65709
Canada DSL	22017
Total number	197015

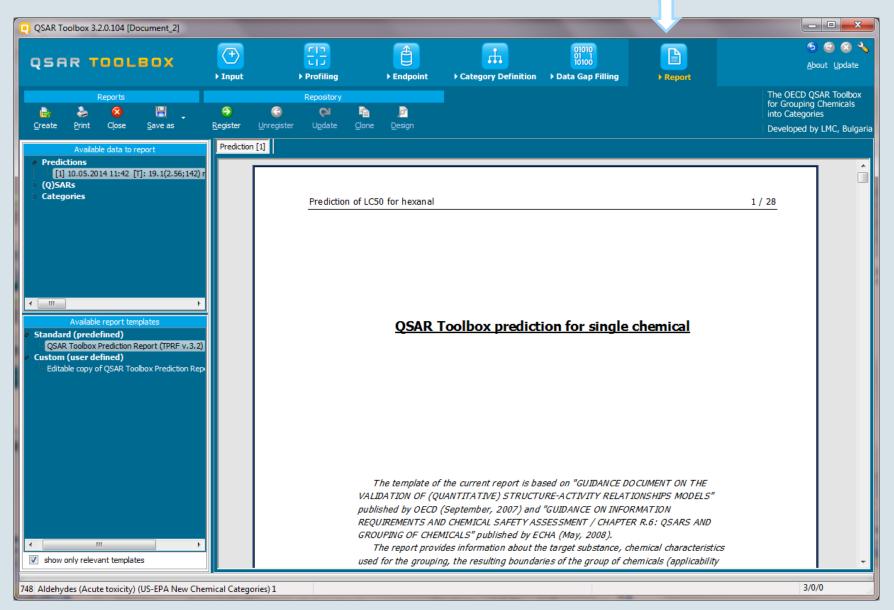
Category definition







Report





Outlook

Description
General scheme and workflow
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Forming categories

How to build categories? Laboratory of Mathematical Chemistry

Basic guidance for category formation

Recomended categorization phases:

- 1. Phase I. Endpoint non-specific structure-related profilers (primary categorization)
- 2. Phase II. Endpoint specific profilers (for subcategorization) based on endpoint driving interaction mechanisms
- 3. Phase II. Additional structure-related profilers, to further eliminate dissimilar chemicals (to increase the consistency of category)

Recommended Categorization Phases

Phase I. Structure based

- US EPA Categorization
- OECD Categorization
- Organic functional group
- Structural similarity
- ECOSAR

Repeating Phase I due to Multifunctionality of chemicals

Broad grouping
Endpoint Non-specific

Recommended Categorization Phases

Phase I. Structure based

- US EPA Categorization
- OECD Categorization
- Organic functional group
- Structural similarity
- ECOSAR

Repeating Phase I due to Multifunctionality of chemicals

Phase II. Mechanism based

- DNA binding mechanism
- Protein binding mechanism
- Mode of action –acute aquatic toxicity
- Genotoxicity/carcinogenicity
- Cramer rules
- Verhaar rule
- Skin/eye irritation corrosion rules

Metabolism accounted for

Broad grouping
Endpoint Non-specific

Subcategorization Endpoint Specific

athematical Chemistry

Recommended Categorization Phases

Phase I. Structure based

- US EPA Categorization
- OECD Categorization
- Organic functional group
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Repeating Phase I due to Multifunctionality of chemicals

Phase II. Mechanism based

- DNA binding mechanism
- Protein binding mechanism
- Mode of action –acute aquatic toxicity
- Genotoxicity/carcinogenicity
- Cramer rules
- Verhaar rule
- Skin/eye irritation corrosion rules

Metabolism accounted for

Phase III. Eliminating dissimilar chemicals

Apply Phase I categorization

Broad grouping
Endpoint Non-specific

Subcategorization Endpoint Specific

lathematical Chemistry

Subcategorization Endpoint Specific

Basic guidance for category formation

Suitable categorization phases:

- 1. Phase I. Endpoint non-specific structure-related profilers (primary categorization)
- 2. Phase II. Endpoint specific profilers (for subcategorization) based on endpoint driving interaction mechanisms
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Performing categorization:

- 1. The categorization phases should be applied successively
- 2. The application order of the phases depend on the specificity of the data gap filling performed (data availability, endpoint specificity)
- 3. More categories of same phase could be used in forming categories
- 4. Some of the phases could be skipped if consistency of category members is reached
- 5. Subcategorization should be applied at Data gap filling stage

Example – Ecotoxicological Information

Grouping methods

General Mechanistic

DNA binding by OASIS

DNA binding by OECD

Estrogen Receptor Rinding

Protein binding by OASIS

Protein binding by OECD

Protein Rinding Potency

Superfragments

Toxic hazard classification by Cramer (original)

Toxic hazard classification by Cramer (with extensions)

Endpoint Specific

Acute aquatic toxicity classification by Verhaar

Acute aquatic toxicity MOA by OASIS

Aquatic toxicity classification by ECOSAR

Bioaccumulation - metabolism alerts

Bioaccumulation - metabolism half-lives

Biodegradation fragments (BioWIN MITI)

Eye irritation/corrosion Exclusion rules by BfR

Eye irritation/corrosion Inclusion rules by BfR

Micronucleus alerts by Benigni/Bossa

Mutagenicity/Carcinogenicity alerts by Benigni/Bossa

Oncologic Primary Classification

Skin irritation/corrosion Exclusion rules by BfR

Skin irritation/corrosion Inclusion rules by BfR

Empiric

Chemical elements

Groups of elements

Lipinski Rule Oasis

Organic functional groups

Organic functional groups (nested)

Organic functional groups (US EPA)

Organic functional groups, Norbert Haider (checkmon)

Structure similarity

Custom

Aldehydes

Acute Aquatic Toxicity

Mode of Action

Structural similarity (not for narcotics)

Data Gap Filling Approach

Trend analysis, External QSARs

Example – Human Health Hazard

Grouping methods

General Mechanistic

DNA binding by OASIS

DNA binding by OECD

Estrogen Receptor Binding

Protein binding by OASIS

Protein binding by OECD

Protein Binding Potency

Superfragments

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Toxic hazard classification by Cramer (with extensions)

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Structure similarity

Custom

Aldehydes

Mutagenicity/Carcinogenicity

Mechanism

Structural similarity

Data Gap Filling Approach

Trend analysis, External SARs