

Integrating Predictive Toxicology Applications & Resources

An OpenTox Workshop

30 May, 2010

Potsdam, Germany

Workshop Overview

Barry Hardy

**Douglas Connect
OpenTox Project Coordinator**

Workshop Overview

1

- Introduction & Overview

2

- Stakeholder Perspectives

3

- Infrastructure Requirements & Interoperability
- OpenTox Framework Design

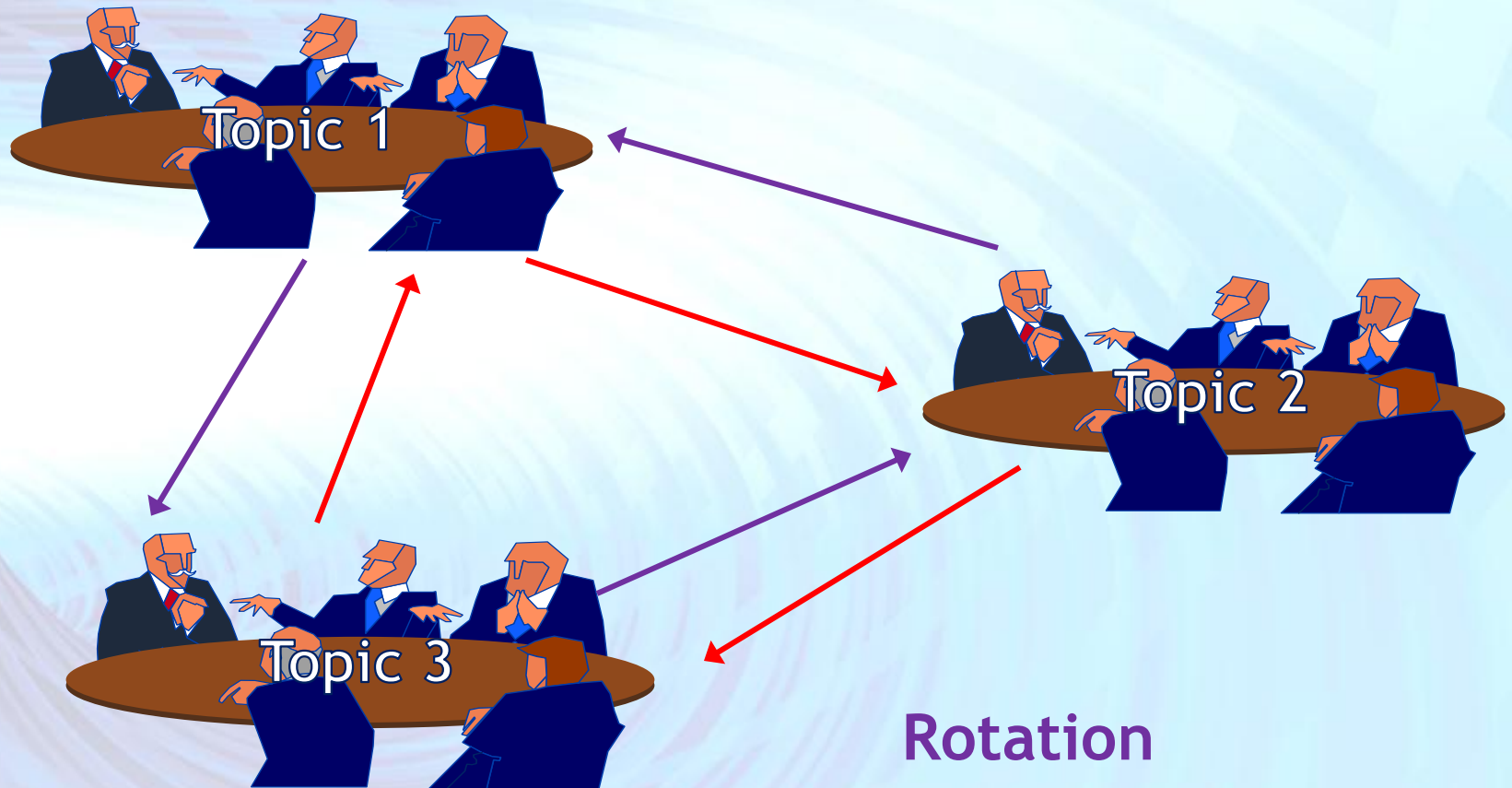
4

- OpenTox Integrating Application Demonstrations
- Extensibility, Sustainability & Future Directions

5

- Knowledge Café Discussions
- Concluding Discussion & Actions

Knowledge Café Discussion Format



Development of Strategies for Interoperable Resources & Applications in Predictive Toxicology



Eliminate
traditional
circus acts,
animals



Create
theatrical
themes,
storylines,
new acts



Reduce
dangerous acts,
traditional
humour,
transport costs

**Blue
Ocean**

Raise
tent standards,
artistic
sophistication,
ticket prices!



Based on Blue Ocean Strategy,
Kim & Mauborne 2006

Perspectives

Robert Kavlock (EPA, US)

Carl Westmoreland (Unilever, UK)

Emilio Benfenati (Mario Negri Institute, Italy)

Egon Willighagen (Uppsala University, Sweden)

Jeffrey Wiseman (Pharmatropé, US)

Michael Schwarz (University of Tuebingen, Germany)

Requirements in Predictive Toxicology Infrastructure

Barry Hardy

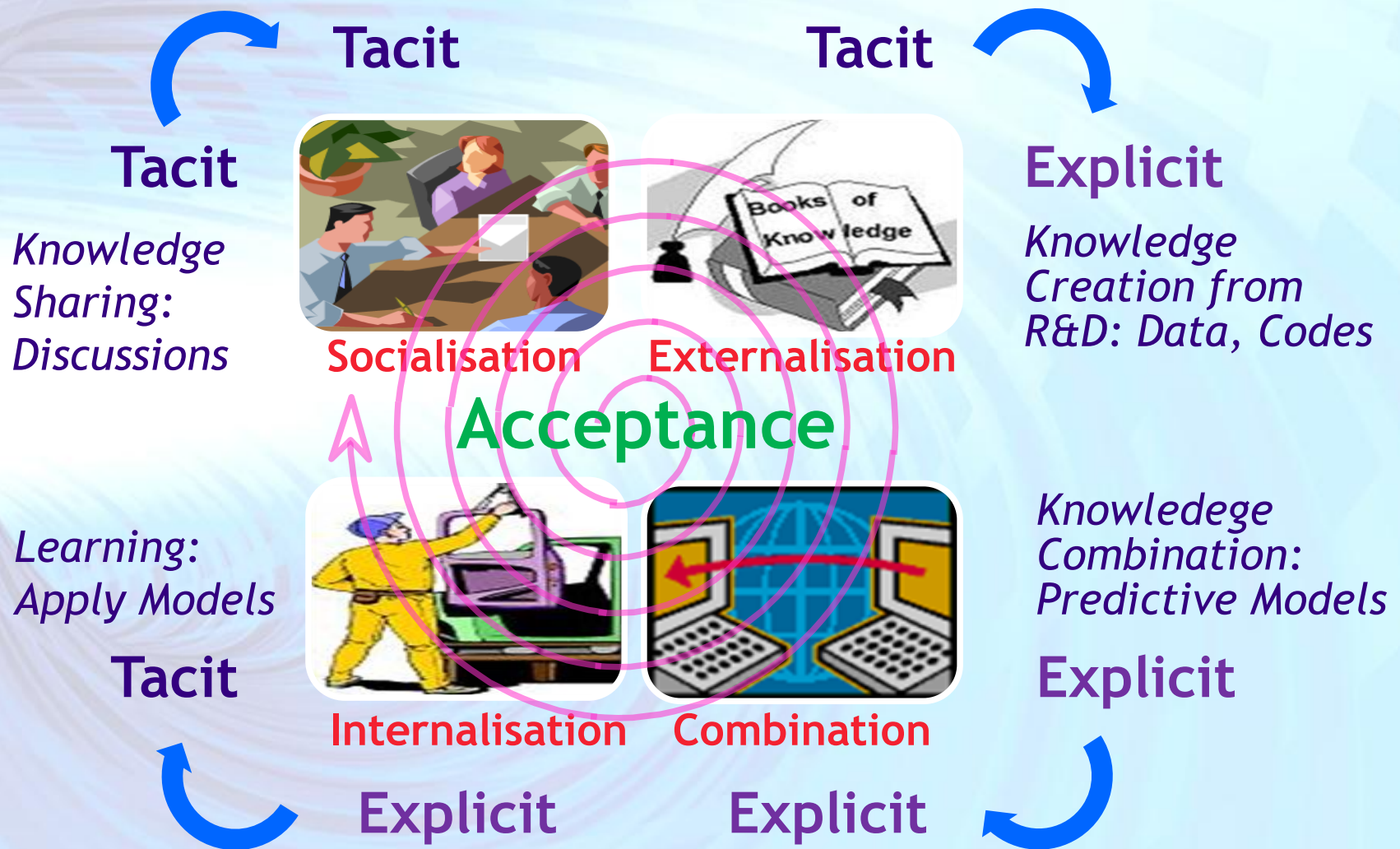
**Douglas Connect
OpenTox Project Coordinator**

Need for communications in the community overcoming different languages and vocabularies



Explaining the rules of different games on a
conservation project trip in the Caprivi, Namibia

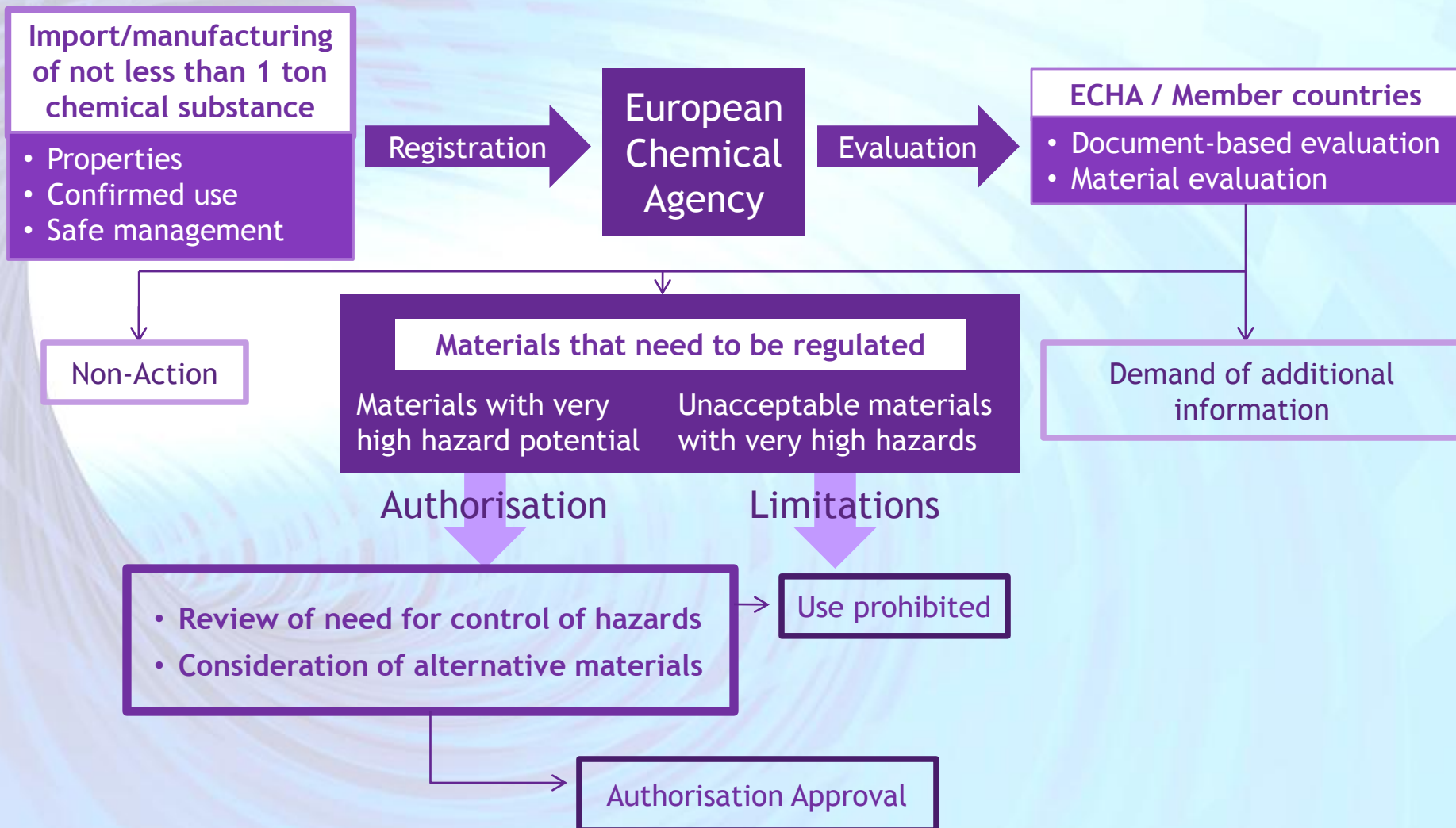
SECI Model for Knowledge Management



REACH



REACH Registration



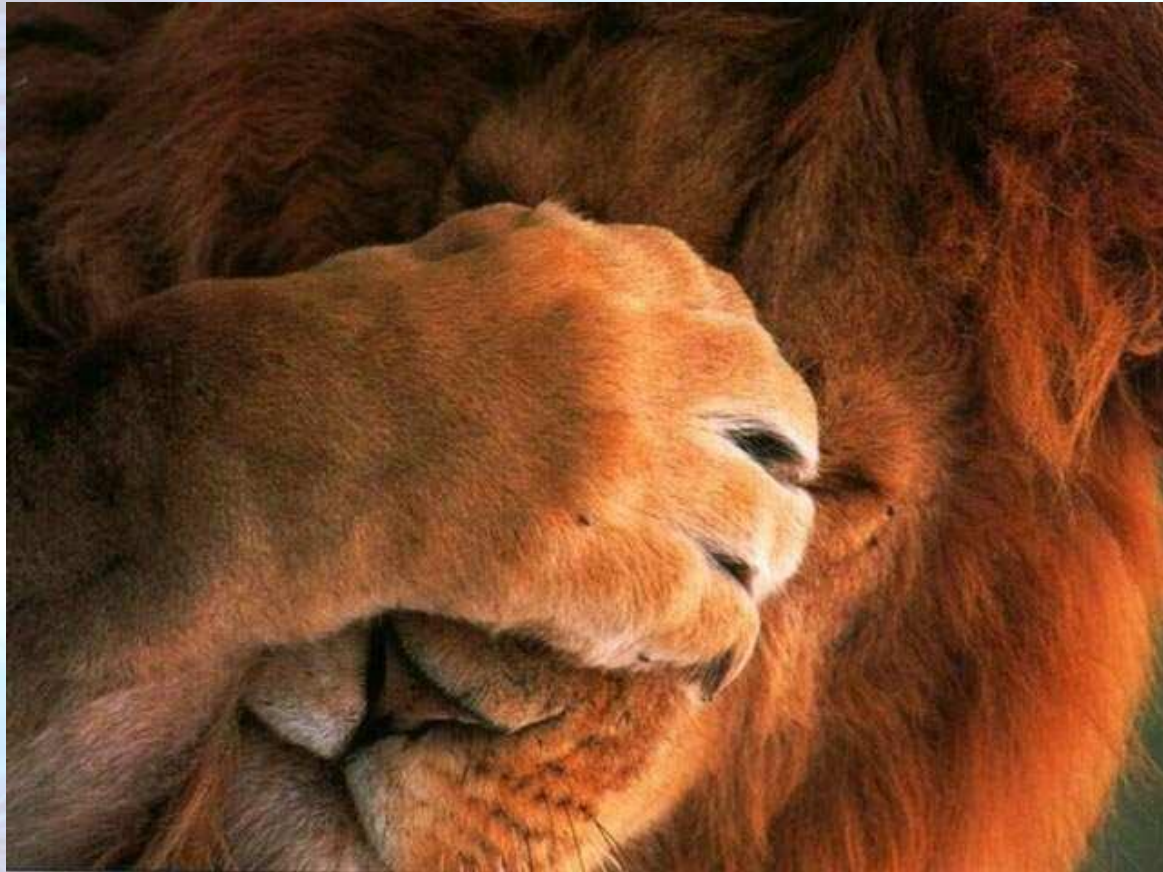
What is our situation today when we say ...

**Let's build an
integrated predictive
toxicology application!**

**Let's collaborate on an
analysis!**



Faced with such an integration challenge ...



... we tend to look away from some of the problems ...

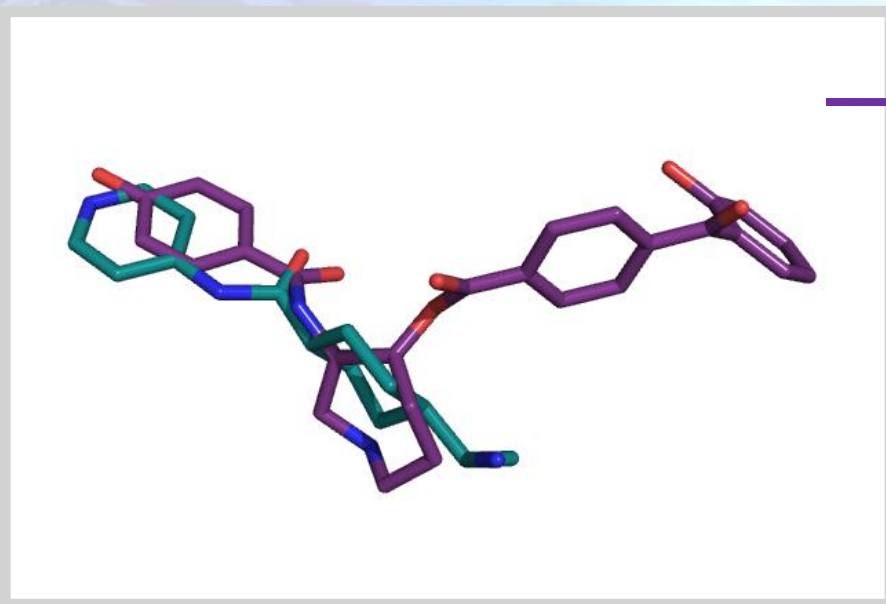
Faced with such an integration challenge ...



... we tend to look away from some of the problems and from the need to collaborate more closely.

Collaborative Predictive Toxicology Challenge

Input Structure



VO

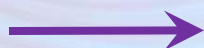


Out - Toxic or Not?

- ☐ LD50
- ☐ Liver Toxicity
- ☐ Secondary Metabolites
- ☐ Bioavailability
- ☐ Mutagenicity
- ☐ Carcogenicity
- ☐ Reproductive Toxicology
- ☐ Skin Irritation
- ☐ Aqua Toxicity
- ☐ Combined predictions for arrays of multiple end points

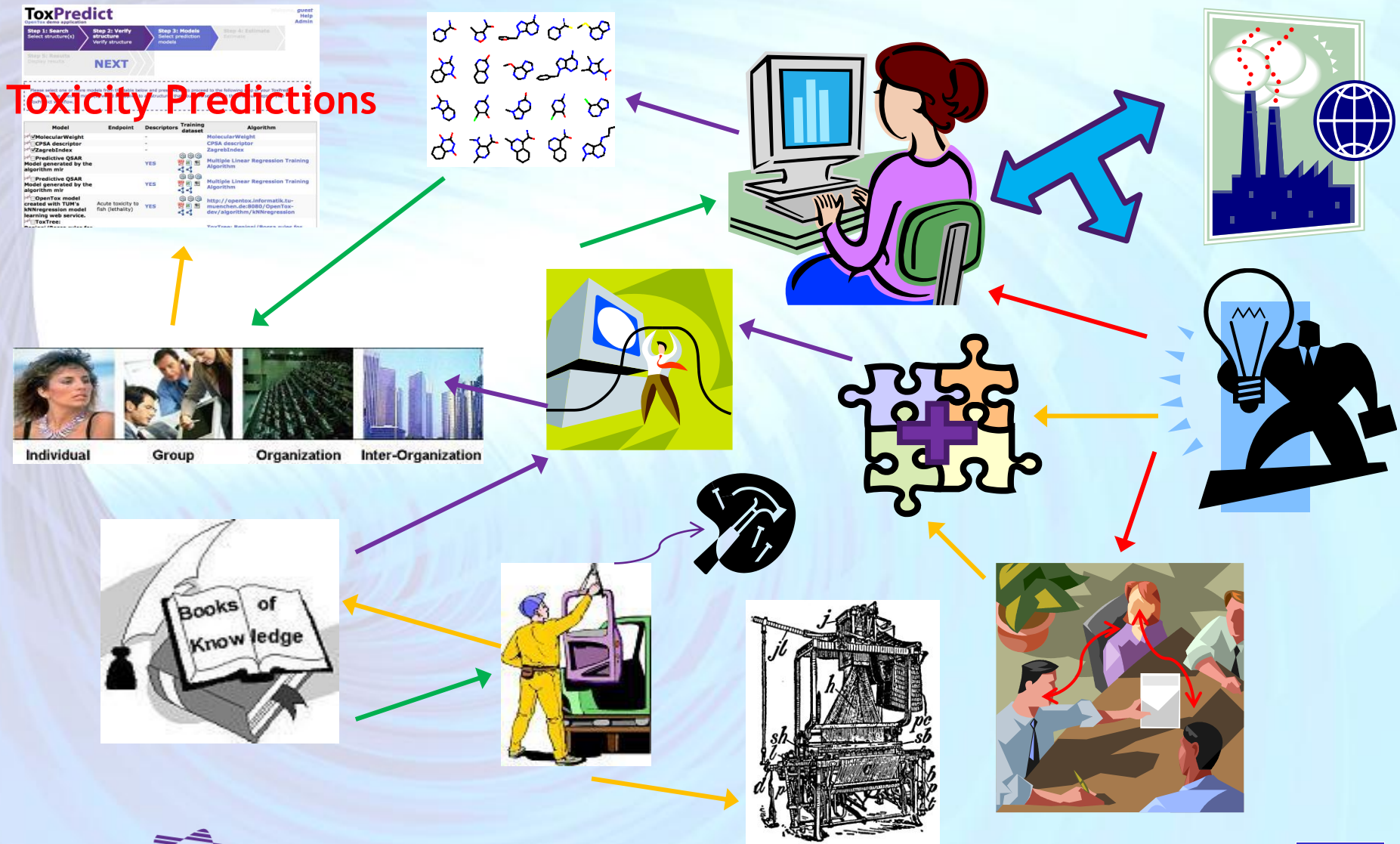


Driver



Increasing demands on industry to satisfy safety evaluation and risk assessment required by REACH legislation.

Accelerating Knowledge Flows in Predictive Toxicology



Compelling Needs of Users

Integrated Testing

in silico

in vitro

TTC

Read
Across

Category
Formation

REACH Reporting
(QPRF, QMRF)

Applicability
Domain

Validation

Human
Data

Compelling Needs of Users

Multidisciplinary R&D

Good Support of Flexible Applications

Transparency -
Not Black Box!

Mechanistic
rationale

QSAR &
Expert
Systems

Workflows

Automated
Integration

Applicability
Domain

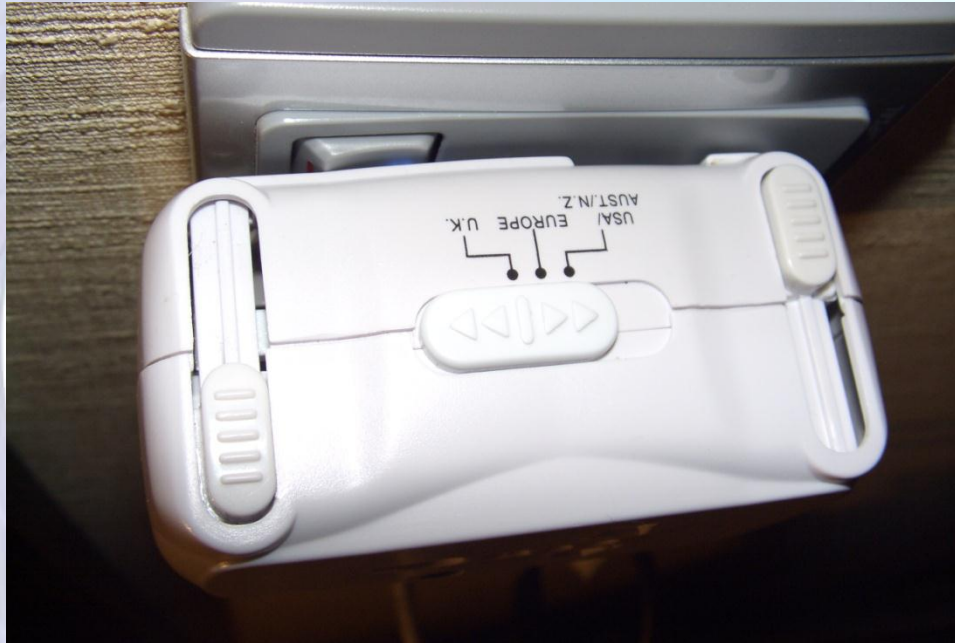
Categories

Systems
Biology

Challenges to Integrated Resources & Applications

- Database silos
- Missing information
- Varying quality
- Hard to integrate data
- Hard to integrate models
- No common framework
- Lack of standards
- Lack of validation
- Complex subject
- Application difficult
- Lack of transparency
- Interdisciplinary collaboration

Absence of Interoperability creates Problems



Adaptor Challenge in Jeddah, 2008

Interacting Components create Solutions



Adaptor Solution in Jeddah, 2008

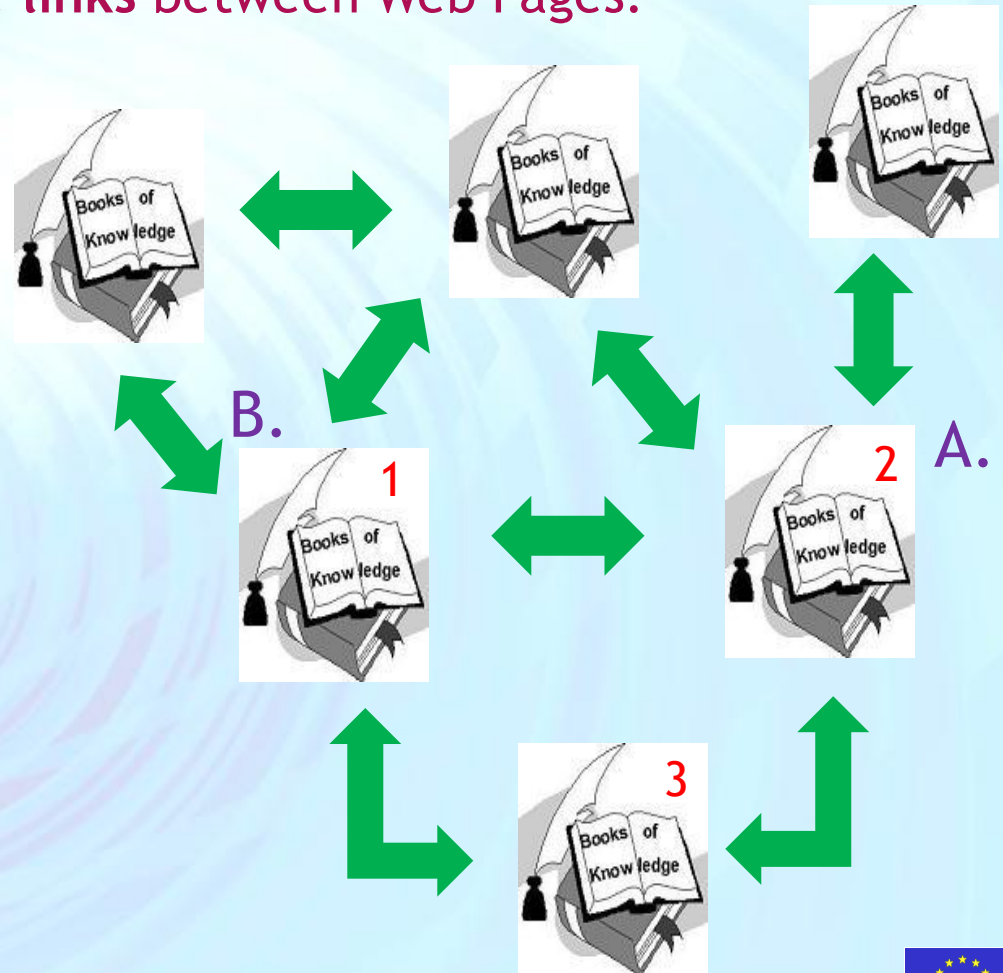
Value is in Linking

The key idea of Google's founders in creating their search engine:
There is useful knowledge in the **links** between Web Pages.

Page Ranking

A page is ranked higher in a search if:

- A. it has more connections to it than other pages
- B. the pages connecting to it have higher ranking themselves



Linked Data enables Knowledge Creation, Combination and Analysis

Linked Data is a term used to describe the exposing, sharing, and connecting of data on the Semantic Web using:

URIs a generic means to identify entities in the world

HTTP a simple yet universal mechanism for retrieving resources

RDF a generic graph-based data model with which to structure and link data

Linked Data needs:

1. Provision of a **URI** that describes a Data Resource
2. Use of **HTTP** to retrieve useful data from the **URI**
3. A Data Format described with standardised semantics (so relationships are enabled) e.g. **RDF**
4. Data should provide links to other Data (through **URIs**)

Linked Data approach can also be applied to other resource types e.g., for algorithms or models as done in OpenTox...



DBpedia = Linked Data approach applied to Wikipedia

Solution created by Linked Open Data, Web Applications and Crowdsourcing



Haiti Earthquake Crisis Response (2010)

wiki.openstreetmap.org

Interoperability & Vocabulary

Dogs

- Collie
- Labrador



Cats

- Siamese
- Persian



Birds

- Sparrow
- Owl



Interoperability & Vocabulary



Interoperability & Ontology

Org A

Collie

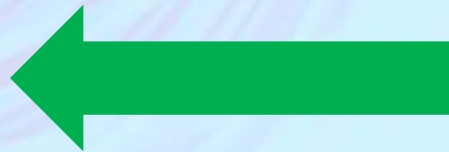


Siamese



Org B

Collie



Siamese

OpenTox Approach

A diagram on the left side of the slide consists of three concentric semi-circles. The outermost semi-circle is the largest, the middle one is smaller, and the innermost one is the smallest. They are all colored purple. The text 'Framework', 'Diverse Access', and 'Interoperability' is positioned to the right of the outer, middle, and inner semi-circles respectively.

Framework

- Toxicity Data (Linked)
- *in silico* models
- Validation & Reporting
- Interpretation aids

Diverse Access

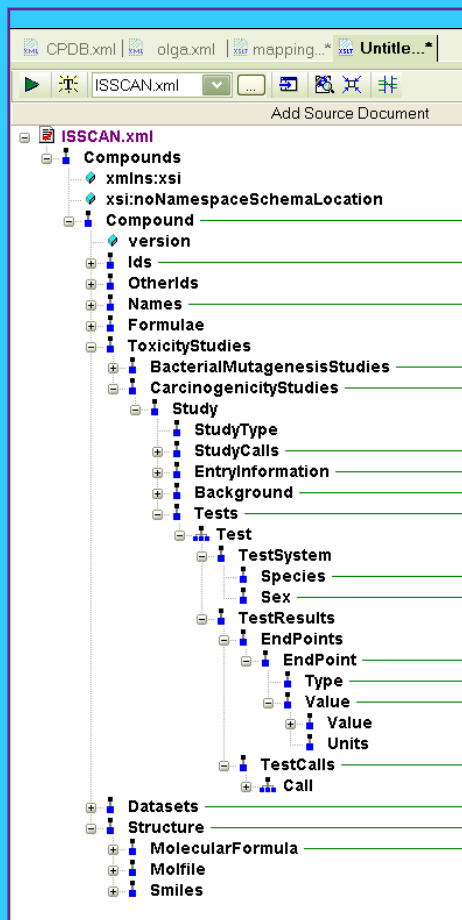
- Toxicologists
- Computational Scientists
- Interfaces for new algorithm development & integration

Interoperability

- Promote Standards
- Core Open Source Components
- Support Ontologies & Integration of Multiple Resources

Toxicological Endpoint Ontology Development

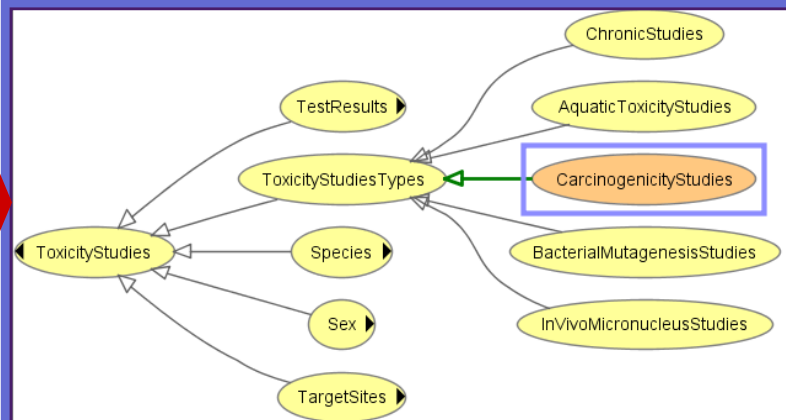
ToxML schema



Other publicly available resources:
DSSTox, GoReni (ITEM), ISSCAN ...

OpenTox Toxicological Endpoint Ontology

Ontology Development



Re-use of terms defined in
neighbouring ontologies (e.g. OBO)

Collaborative
Protégé
Environment

OpenToxipedia



Barry Hardy Log out Quicktools Site Setup Help

Site Map Accessibility Contact Data

Search Site

Home Toxicity Prediction OpenTox Blog People Partners Development OpenToxipedia

User Guidance Latest Entries A B C D E F G H I J K L M N O P Q R S T U V W
X Y Z by Categories Entries OpenToxipedia

You are here: Home » OpenToxipedia

Contents View Edit Rules Sharing History

Actions Display Add new... State: Published

OpenToxipedia

by Barry Hardy — last modified Sep 03, 2009 01:09 PM

OpenTox Community Resource for Toxicology Vocabulary and Ontology

OpenTox is supporting the creation and curation of OpenToxipedia, a community-based predictive toxicology knowledge resource. All members of the community are welcome to provide entries, suggested definition edits or additional information to entries in the resource.

OpenTox is supporting the application and development of the **ToxML** standard for representation of toxicology data, the **OECD principles for (Q)SAR model validation**, and the use of the **OECD HT** standard for regulatory reporting purposes.

OpenToxipedia provides here a Vocabulary Resource of toxicology terminology. We hope you find the resource useful and consider contributing to terms and their content.

Guidance for Vocabulary Resource entries

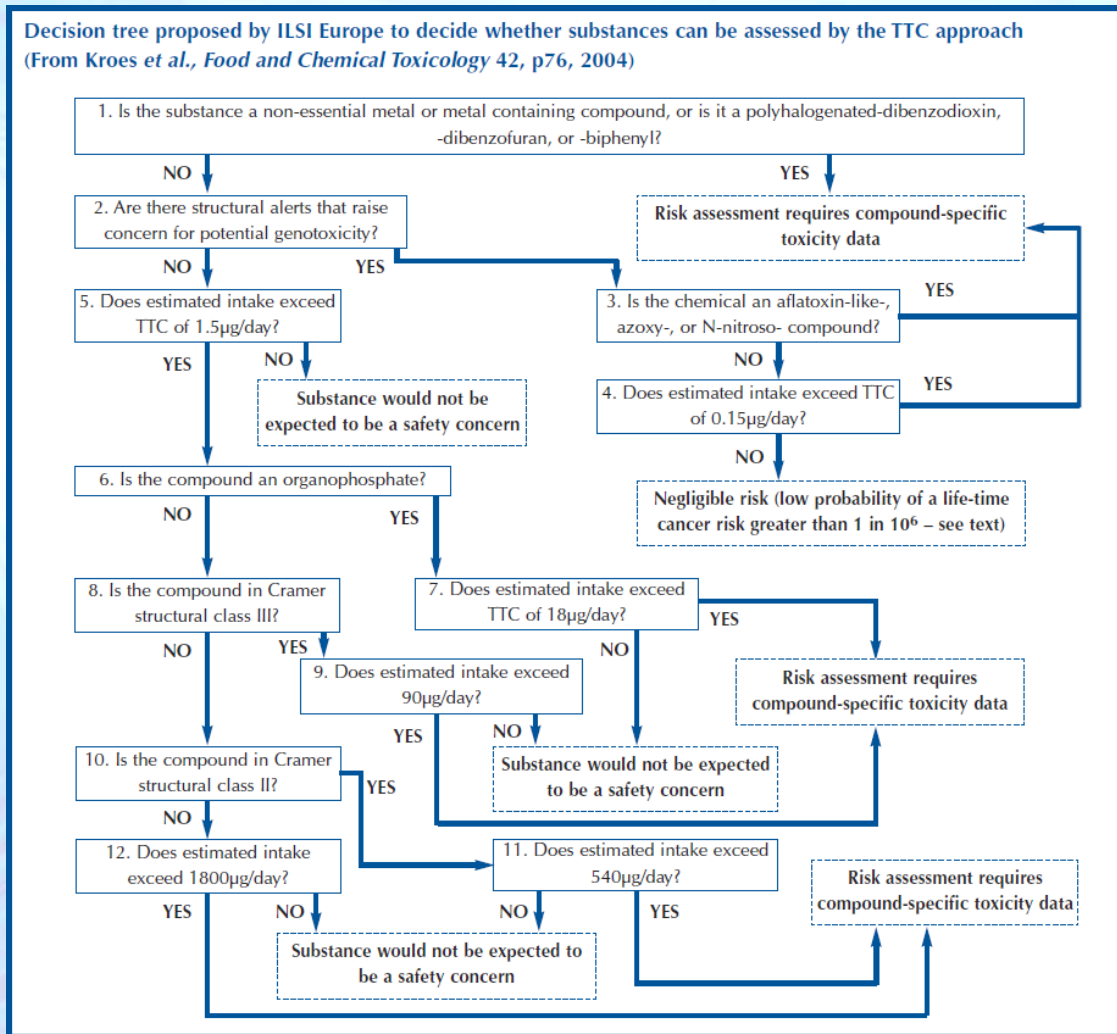


www.opentox.org/opentoxipedia



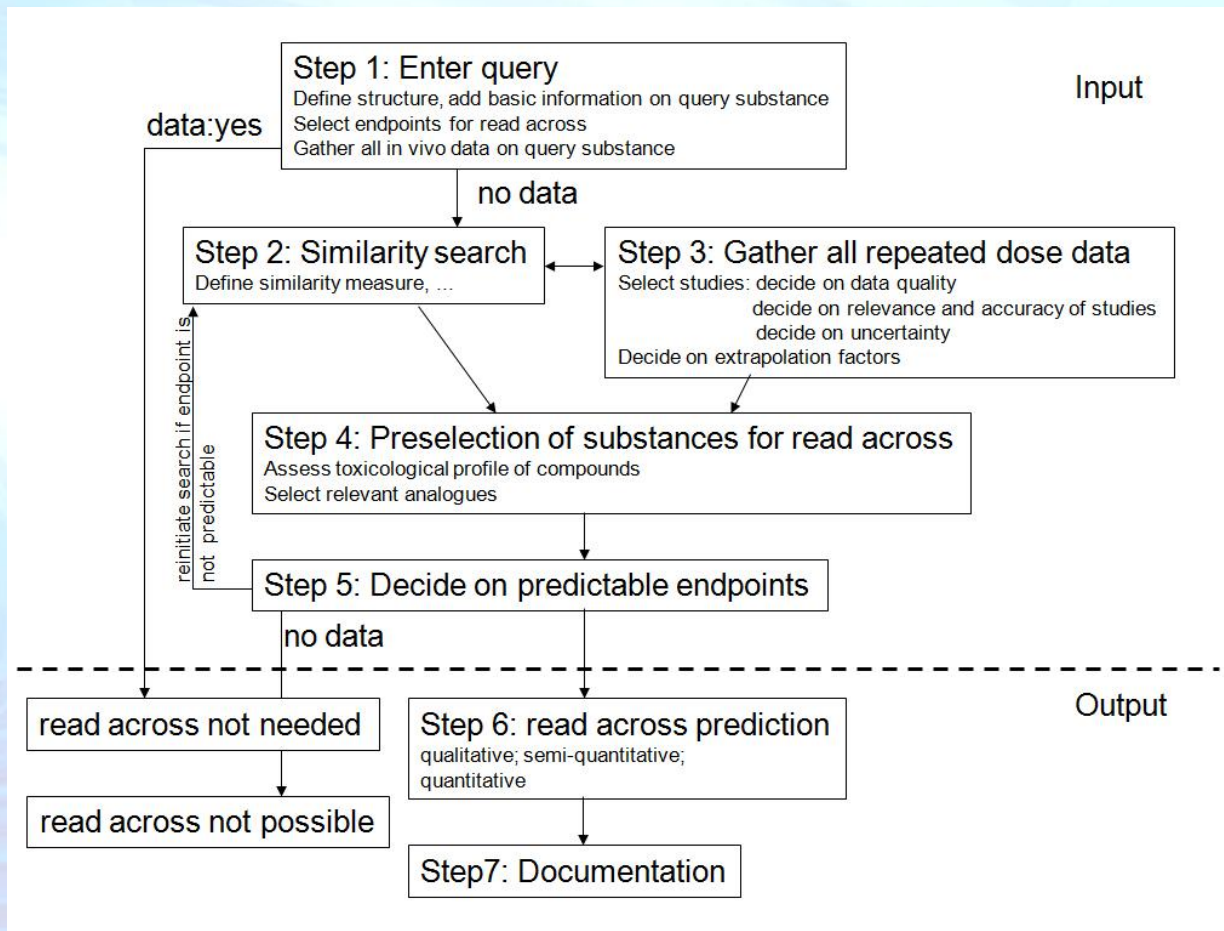
OpenTox Use Case Example

Implement Threshold of Toxicological Concern (TTC) using ILSI decision tree guidelines



OpenTox Use Case Example

Support Read Across for Repeated Dose Toxicity



Step 1: Search
Select structure(s)


**Step 2: Verify
structure**
Verify structure

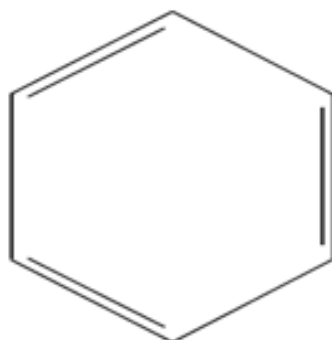
Step 3: Models
Select prediction
models

Step 4: Estimate
Estimate

Step 5: Results
Display results

This page lists your ToxPredict workflow results for the structure(s) you have selected and the model prediction(s) you have chosen to run. You could also retrieve the ToxPredict report in various other formats, e.g. [SDF](#), [CML](#), [SMI](#), [PDF](#), [CSV](#), [ARFF](#), [RDF/XML](#) or [RDF/N3](#).

Download as 



CAS RN
EINECS
IUPAC name
Synonym

71-43-2
200-753-7
benzene
(6)annulene; benzine; Benzol; Benzolene;
bicarburet of hydrogen; carbon oil; Coal naphtha;
cyclohexatriene; mineral naphtha; motor benzol;
nitration benzene; Phene; Phenyl hydride;
pyrobenzol.

Synonym
Synonym
Synonym
Quality label

21742.0
Benzene
benzene
OK

MolecularWeight  **MolecularWeight**

MW

78.1112

	OECD Principle	OpenTox addresses Validation Principles by...
1	Defined Endpoint	providing a unified source of well defined and documented toxicity data with a common vocabulary
2	Unambiguous Algorithm	providing transparent access to well documented models and algorithms as well as to the source code
3	Defined Applicability Domain	integrating tools for the determination of applicability domains during the validation of prediction models
4	Goodness-of-fit, robustness and predictivity	providing scientifically sound validation routines for the determination of errors and confidences
5	Mechanistic interpretation (if possible)	integrating tools for the prediction of toxicological mechanisms and the recording of opinions and analysis in reports

The OpenTox Framework Design

Stefan Kramer

Technical University of Munich

OpenTox Approach

A diagram on the left side of the slide consists of three concentric semi-circles. The outermost semi-circle is the largest, the middle one is smaller, and the innermost one is the smallest. They are all purple with white outlines. The text 'Framework', 'Diverse Access', and 'Interoperability' is aligned with the middle, outer, and inner semi-circles respectively.

Framework

- Toxicity Data (Linked)
- *in silico* models
- Validation & Reporting
- Interpretation aids

Diverse Access

- Toxicologists
- Computational Scientists
- Interfaces for new algorithm development & integration

Interoperability

- Promote Standards
- Core Open Source Components
- Support Ontologies & Integration of Multiple Resources

Strategic Context/Goals

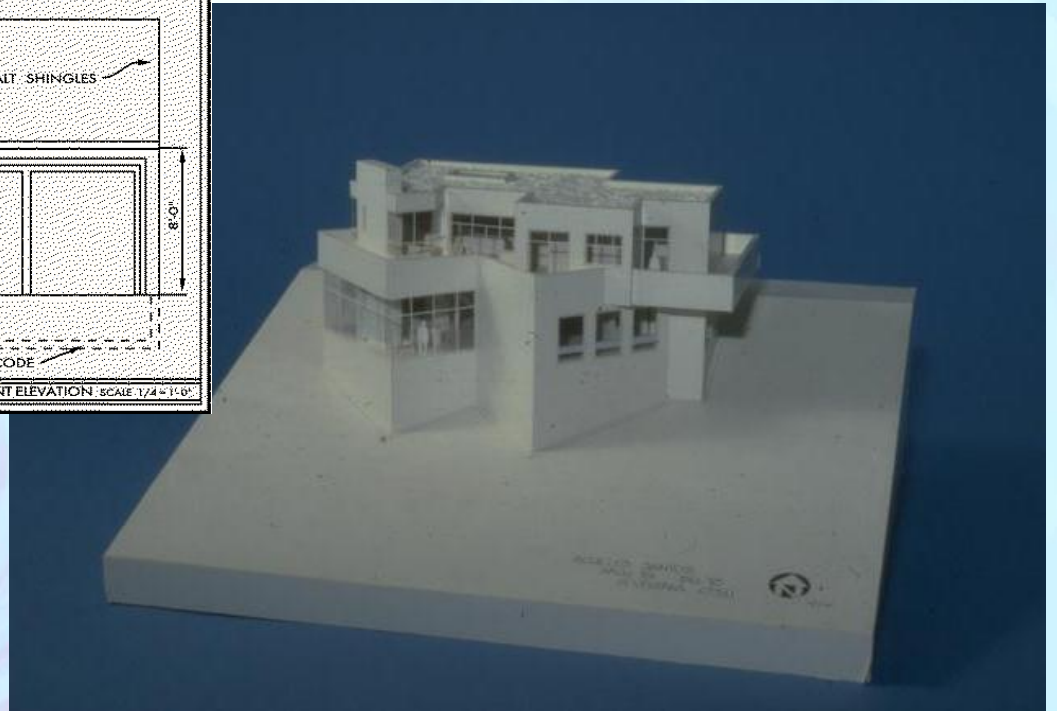
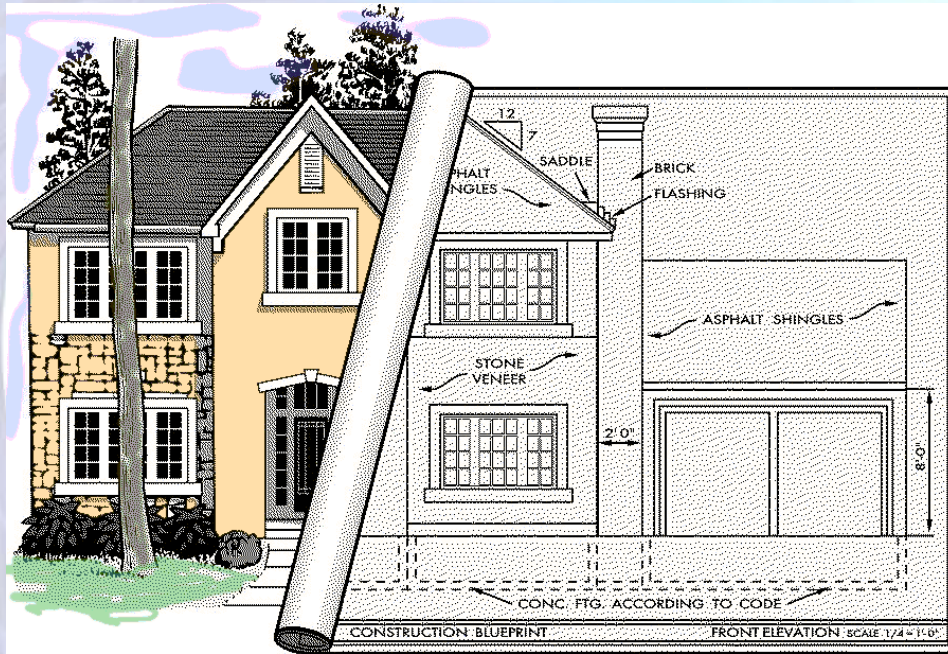
- REACH: possible reduction of test animals by using existing experimental data in conjunction with QSAR
- Also practical needs: **reporting** and form filling
- By the OECD principles, a number of requirements to a framework like OpenTox arise

	OECD Principle	OpenTox addresses by...
1	Defined Endpoint	providing a unified source of well defined and documented toxicity data with a common vocabulary
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Consequences for Requirements on OpenTox

User Requirements		Software Requirements
Unambiguous data	⇒	<i>formal way of representing information about data</i>
Unambiguous access	⇒	<i>well-defined interfaces</i>
Transparency of computational tools	⇒	<i>formal way of representing information about methods, well-defined interfaces</i>
Variety of user groups	⇒	<i>simplicity and modularity of design</i>
Need to integrate various resources (e.g., databases, prediction methods, models, ...) to make meaningful predictions	⇒	<i>distributed architecture, interoperability</i>
Need to integrate biological information	⇒	<i>again, modularity of design, extensibility</i>

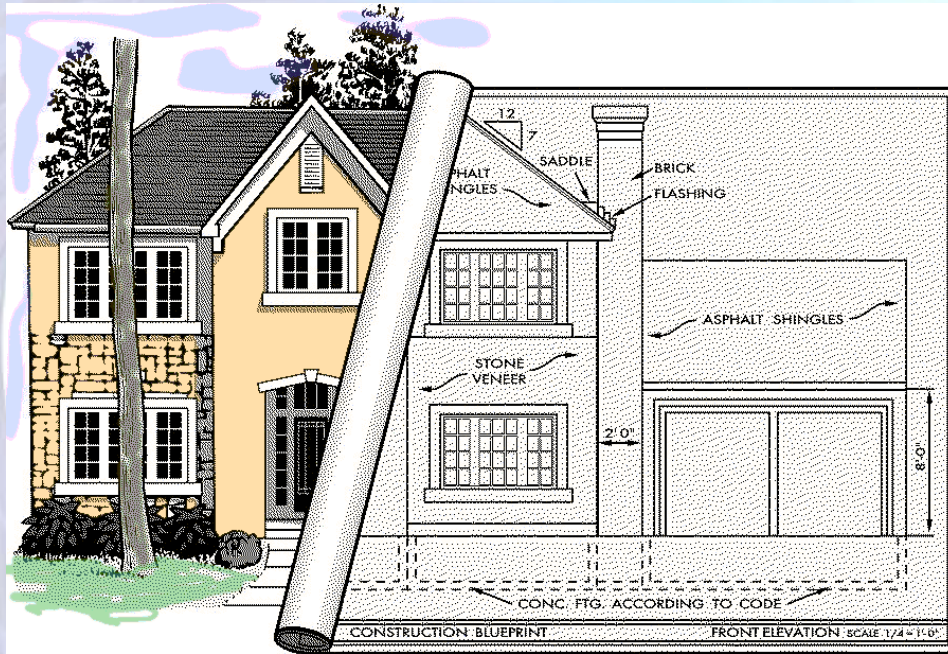
Software Architecture



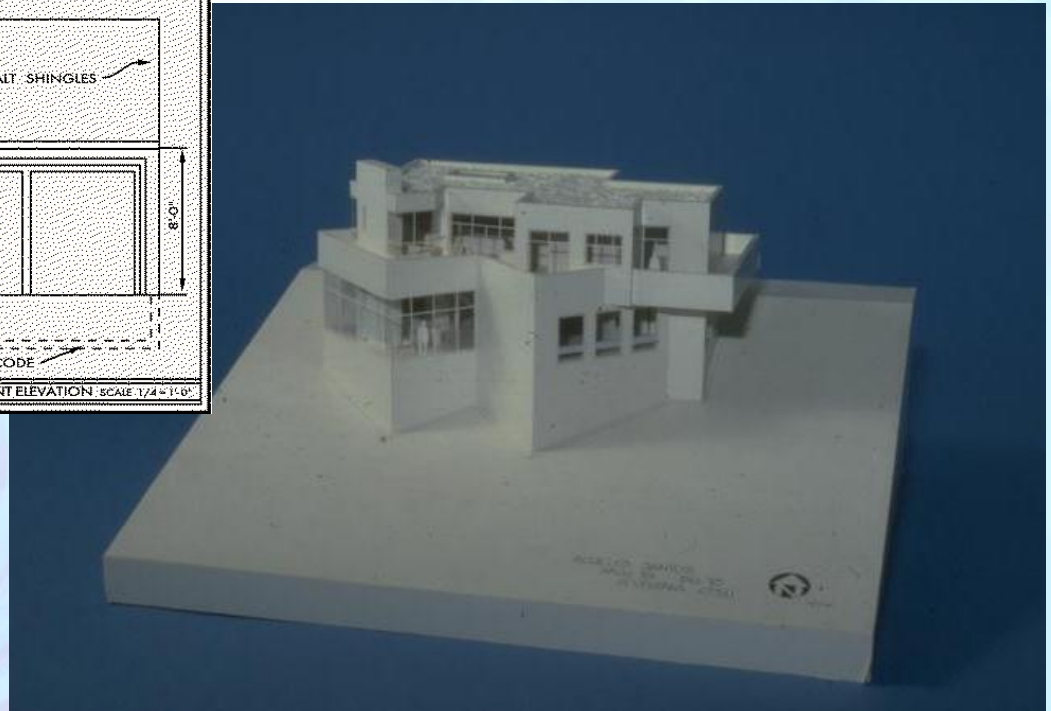
Software Architecture



Software Architecture



To solve a wide range of modelling tasks: needs buildings blocks (modularity) and well-defined ways of combining them



Technological choices...

Representational State Transfer (REST): What and Why?

What?

- Architectural style for distributed information systems on the Web
- Simple interfaces, data transfer via **hypertext transfer protocol (HTTP)**, stateless client/server protocol
 - GET, POST, PUT, DELETE
- Each **resource** is **addressed** by its own **web address**

Why?

- **Lightweight** approach to **web services**
- **Simplifies/enables** development of **distributed systems**
- (More or less) language independent/installation-free

Ontologies: What and Why?

What?

- Formal, shared conceptualization of a domain

Why?

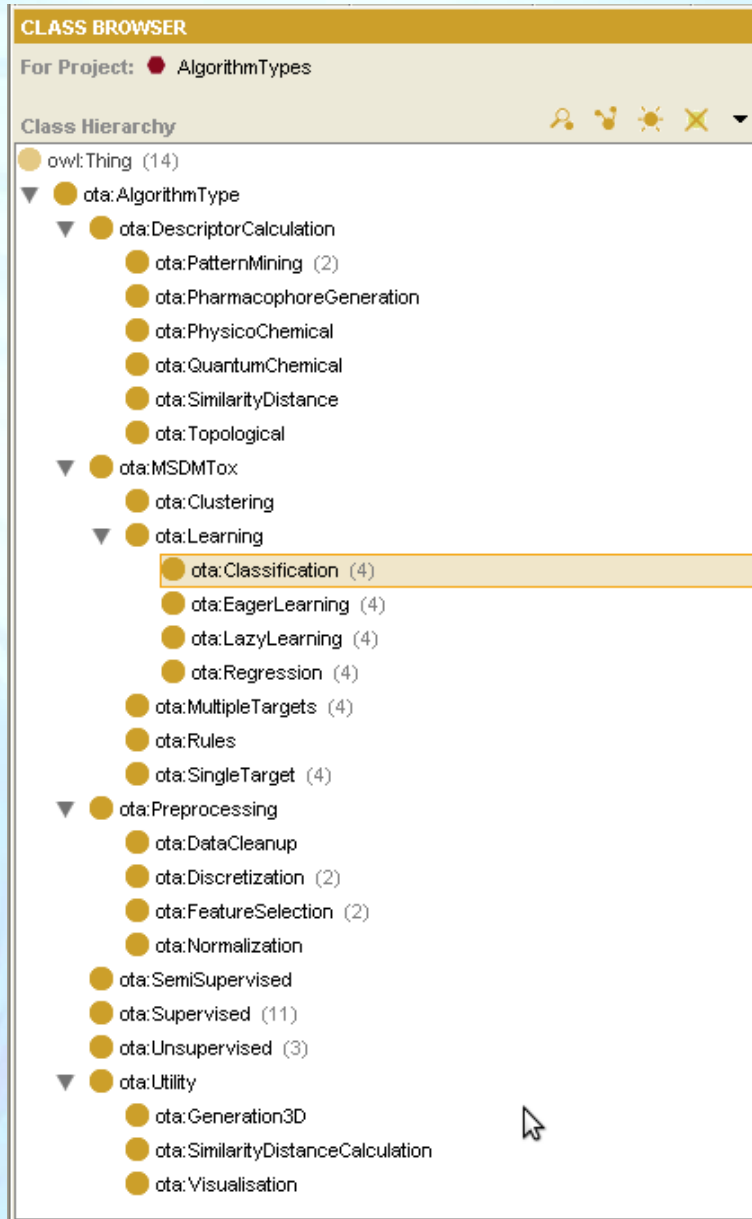
- Distributed services **need** to be able to “talk to each other”, i.e. have a **common understanding** of endpoints, any type of property, methods, etc.



Ontologies

- Standards: **OWL** as representation language and **SPARQL** as query language
- There are many ongoing biological ontology projects
- Our strategy: use existing work and standards wherever possible
- However, there are new ontology needs for OpenTox applications, e.g. for algorithms, toxicological endpoints

OpenTox
Ontology Working Group



OpenTox: Databases

Chemical compounds - Mozilla Firefox

File Edit View History Bookmarks Tools Help

http://apps.ideaconsult.net:8180/ambit2/query/smarts?type=smiles&search=[*]OC(=O)[#6;H1]=[#6;H1]c1cccc1&t

Disable Cookies CSS Forms Images Information Miscellaneous Outline Resize Tools View Source Options

Chemical compounds

ToxPredict TTC Depiction Datasets Chemical compounds Similarity Substructure Algorithms References Features Templates Models Ontology RDF playground Help

ambit

SMARTS

Keywords

Search for substructure and properties
This site and AMBIT REST services are under development!

Retrieve data

Search results SMARTS [*]OC(=O)[#6;H1]=[#6;H1]c1cccc1 Download as Max number of hits: 100

#	Compound	ECHA REGISTRATION DATE	ECHA CasRN	ECHA EC	ECHA Names	ECHA SYNON Names	ECHA SYNON Names	ECHA SYNON Names	ECHA SYNON Names	ECHA SYNON Names	ECHA SYNON Names
1		30.11.2010	78-37-5	201-110-3	linalyl cinnamate						
2		30.11.2010	90-50-6	201-999-8	3,4,5-trimethoxycinnamic acid						

Default
Identifiers
Datasets
Models
Endpoints
All descriptors
pKa
Molecule size
Electronic descriptors (PM3 optimized structure)
Electronic descriptors (original structure)
Toxtree: Cramer rules
☒ http://apps.ideaconsult.net

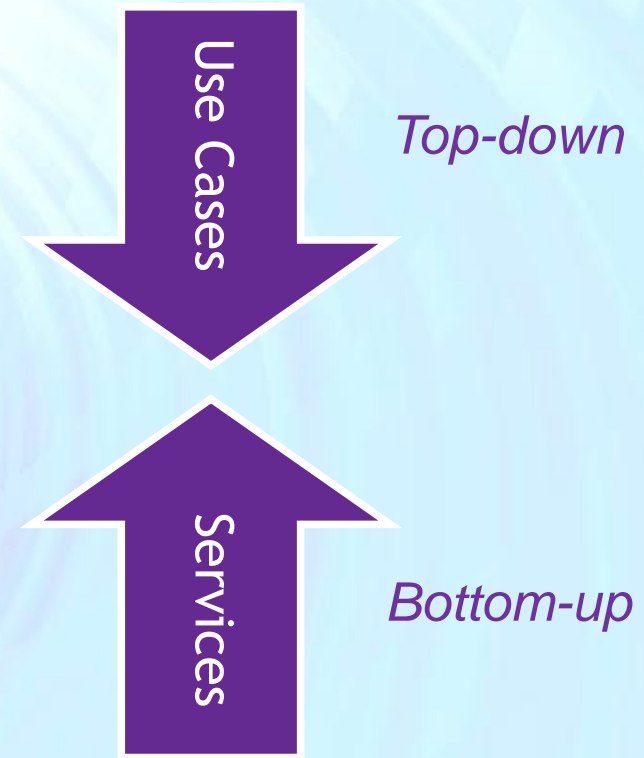
OpenTox: Databases

Dataset	OK	Probably OK	Probably ERROR	Unknown	Probably ERROR%
ECHA list of pre-registered substances	N/A	N/A	N/A	N/A	N/A
Chemical Identifier Resolver	67779	5314	3638	3471	4.75%
ChemIDplus	64802	7986	921	1745	1.24%
ChemDraw	17918	1147	502	478	2.57%
JRC PRS list	61332	4833	4022	2880	5.83%
ISSCAN	931	50	98	62	9.40%
CPDBAS	778	37	0	693	0%
DBPCAN	60	2	0	147	0%
EPAFHM	281	5	0	331	0%
KIERBL	102	1	0	175	0%
IRISTR	346	16	0	177	0%
FDAMDD	213	19	1	983	0.08%
ECETOC skin irritation	158	12	0	5	0%
Skin sensitisation (LLNA)	160	7	4	38	1.95%
Bioconcentration factor (BCF) Gold Standard Database	N/A	N/A	N/A	N/A	N/A

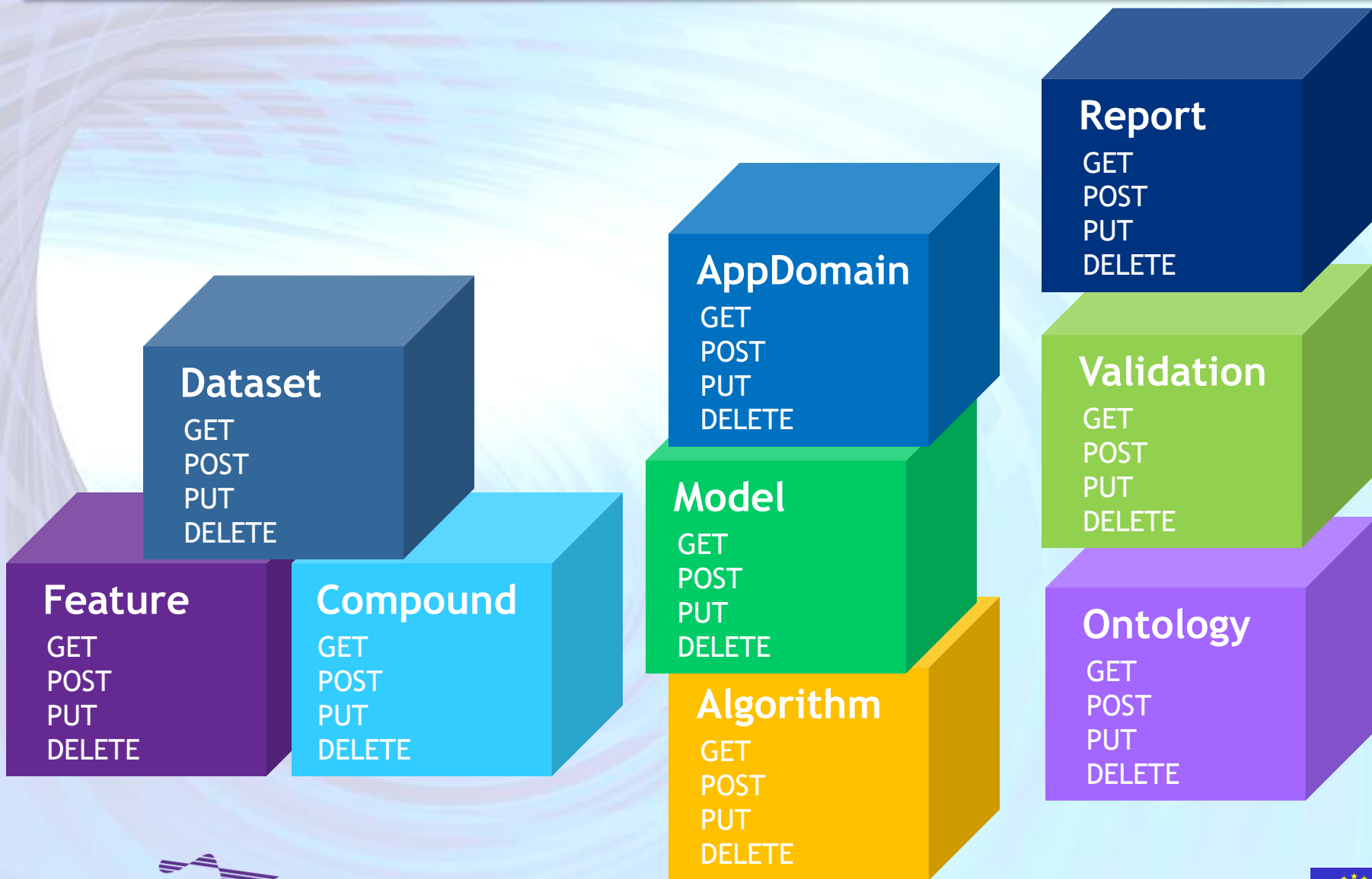
Integrating diverse data sources into OpenTox ontology

Development Process

- Bidirectional
 - from the services (building blocks) to the use cases, and, vice versa,
 - from the use cases to the services
- Use cases
 - ToxPredict (shown later)
 - ToxCreate (shown later)
 - Read-across
 - ToxCast
 - ...



Overview of Application Programming Interfaces



Interface Definitions

Description	Method	URI	Parameter	Result	Status codes
Get available feature URIs for a compound	GET	/compound/{cid}/feature	?feature_uris[]="URI to features" (optional)	Returns representation of the features as uri-list or RDF All available features are returned, if no parameter is specified.	200,404,503
Create a new feature value	POST	/compound/{cid}/feature	?feature_uri="URI to feature" (mandatory, single feature)&value=the_value	URI of the compound with the new feature, e.g. /compound/{id}?feature_uris[]=the-new-feature	200,400,503
Update a new feature value	PUT	/compound/{cid}/feature	?feature_uri="URI to feature" (mandatory, single feature)&value=the_value		200,400,404,503
Delete specified features from the compound	DELETE	/compound/{cid}/feature	?feature_uris[]="URI to features" (optional)		200,400,404,503

Interface Definitions

Description	Method	URI	Parameters	Result	Status codes
get description of a specific feature definition	GET	/feature/{id}	–	URI-list or RDF representation of a feature.	200,404,503
create a new feature	POST	/feature	Content-type ="any-of-RDF-types", content=RDF-representation	URI of the new feature definition.	200,400,404,503
update feature	PUT	/feature/{id}	Content-type ="any-of-RDF-types", content=RDF-representation	–	200,400,404,503
delete feature	DELETE	/feature/{id}	–	–	200,400,404,503
get a list of available feature definitions	GET	/feature	?query=URI-of-the-owl:sameAs-entry	URI list or RDF of features found by the query or all available, if query is empty. Returns all features, for which owl:sameAs is given by the query.	200,404,503

Interface Definitions

Description	Method	URI	Parameters	Result	Status codes
Get a list of available datasets	GET	/dataset	Query parameters (optional, to be defined by service providers).	List of URIs or RDF for the metadata only.	200,404,503
Get a dataset	GET	/dataset/{id}	–	Representation of the dataset in a supported MIME type.	200,404,503
Query a dataset	GET	/dataset/{id}	compound_uris[] and/or feature_uris[] to select compounds and features; further query parameters may be defined by service providers.	Representation of the query result in a supported MIME type.	200,404,503
Get metadata for a dataset	GET	/dataset/{id}/metadata	–	Representation of the dataset metadata in a supported MIME type.	200,404,503
Get a list of all compounds in a dataset	GET	/dataset/{id}/compounds	–	List of compound URIs.	200,404,503
Get a list of all features in a dataset	GET	/dataset/{id}/features	–	RDF or List of feature URIs (pointing to feature definitions/ontologies).	200,404,503

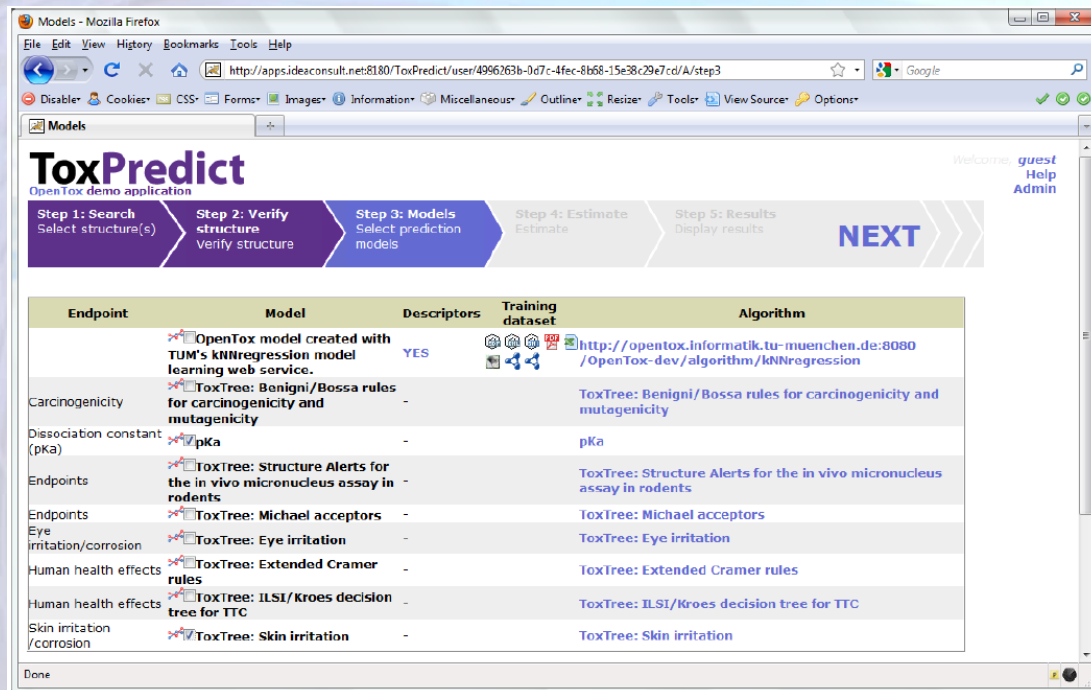
Interface Definitions

Description	Method	URI	Parameters	Result	Status codes
Get a list of all available models	GET	/model	(optional) ?query=URI-of-the-owl:sameAs-entry	List of model URIs or RDF representation. If query specified, returns all models, for which owl:sameAs is given by the query.	200,404,503
Get the representation of a model	GET	/model/{id}	-	Representation of the model in a supported MIME type.	200,404,503
Delete a model	DELETE	/model/{id}	-	-	200,404,503
Apply a model to predict a dataset	POST	/model/{id}	dataset_uri result_dataset=dataseturi dataset_service=datasetserviceuri	URI of created prediction dataset (predictions are features), task URI for time consuming computations.	200,202,400,404,500,503
Apply a model to predict a compound	POST	/model/{id}	compound_uri	Prediction in a supported MIME type; task URI for time consuming computations.	200,202,400,404,500,503

Interface Definitions

Description	Method	URI	Parameters	Result	Status codes
Retrieve SPARQL query results	GET	/ontology	?query=SPARQL_QUERY (mandatory)	RDF representation of the query results.	200,404,500
Predefined query to retrieve all models	GET	/ontology/models		RDF representation of all models.	
Predefined query to retrieve all endpoints	GET	/ontology/endpoints		RDF representation of all endpoints.	
Predefined query to retrieve all algorithms	GET	/ontology/algorithms		RDF representation of all algorithms.	
Submit SPARQL query and/or OpenTox service URL	POST	/ontology	uri[]=URL of a OpenTox RDF resource query=SPARQL_QUERY	RDF representation of the query results, if query is specified. if uri[] is specified, the server retrieves a RDF representation and adds it to the RDF storage, thus making it available for the subsequent queries.	200,404,500,502

What you can do with it ...



Endpoint	Model	Descriptors	Training dataset	Algorithm
	OpenTox model created with TUM's kNNregression model learning web service.	YES	http://opentox.informatik.tu-muenchen.de:8080/OpenTox-dev/algorithm/kNNregression	
Carcinogenicity	ToxTree: Benigni/Bossa rules for carcinogenicity and mutagenicity	-		ToxTree: Benigni/Bossa rules for carcinogenicity and mutagenicity
Dissociation constant (pKa)	pKa	-		pKa
Endpoints	ToxTree: Structure Alerts for the in vivo micronucleus assay in rodents	-		ToxTree: Structure Alerts for the in vivo micronucleus assay in rodents
Endpoints	ToxTree: Michael acceptors	-		ToxTree: Michael acceptors
Eye irritation/corrosion	ToxTree: Eye irritation	-		ToxTree: Eye irritation
Human health effects	ToxTree: Extended Cramer rules	-		ToxTree: Extended Cramer rules
Human health effects	ToxTree: ILSI/Kroes decision tree for TTC	-		ToxTree: ILSI/Kroes decision tree for TTC
Skin irritation/corrosion	ToxTree: Skin irritation	-		ToxTree: Skin irritation

Simple building of predictive toxicology applications based on well-established methods and databases

What you can do with it ...



The screenshot shows the ToxPredict web application interface in a Mozilla Firefox browser window. The URL is <http://apps.ideaconsult.net:8180/ToxPredict/user/496263b-0d7c-4fec-8b68-15e38c29e7cd/A/step3>. The interface includes a navigation bar with steps: Step 1: Search (Select structure(s)), Step 2: Verify structure (Verify structure), and Step 3: Models (Select prediction models). Below this is a table with columns: Endpoint, Model, and Descriptors.

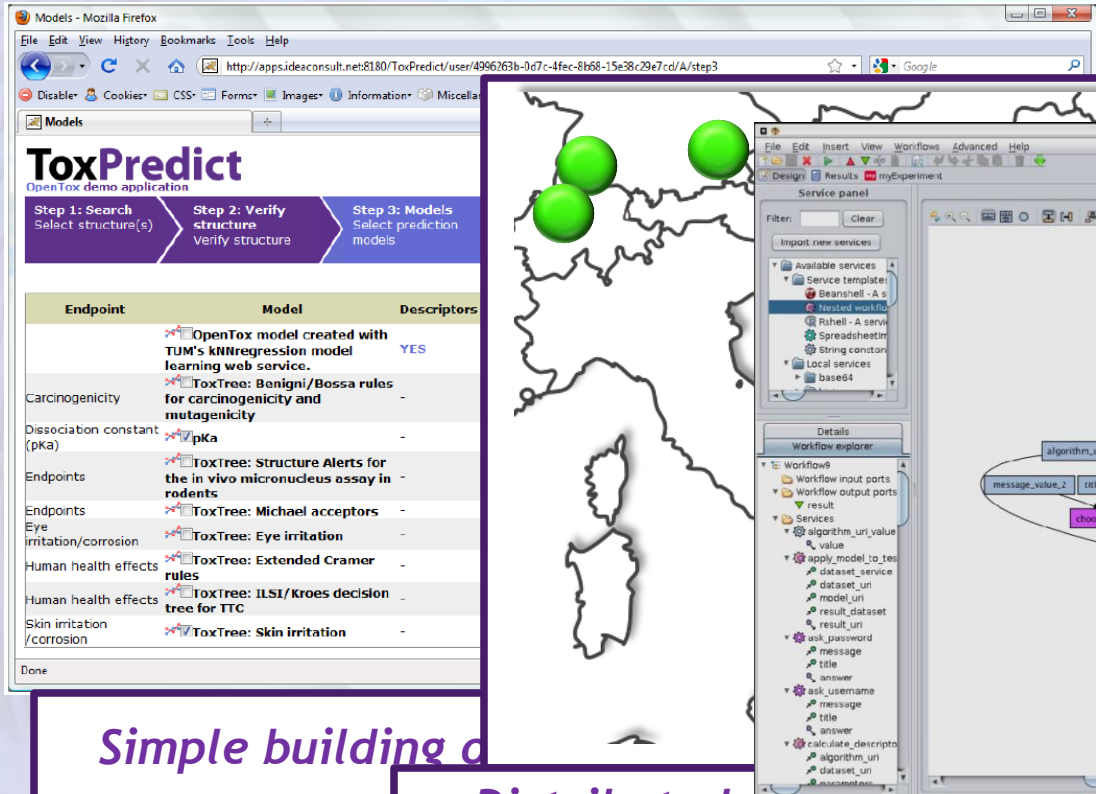
Endpoint	Model	Descriptors
	OpenTox model created with TUM's kNNregression model learning web service.	YES
Carcinogenicity	ToxTree: Benigni/Bossa rules for carcinogenicity and mutagenicity	-
Dissociation constant (pKa)	pKa	-
Endpoints	ToxTree: Structure Alerts for the in vivo micronucleus assay in rodents	-
Endpoints	ToxTree: Michael acceptors	-
Eye irritation/corrosion	ToxTree: Eye irritation	-
Human health effects	ToxTree: Extended Cramer rules	-
Human health effects	ToxTree: ILSI/Kroes decision tree for TTC	-
Skin irritation/corrosion	ToxTree: Skin irritation	-

Below the table is a "Done" button. To the right of the browser window is a map of Europe with five green circular markers placed in various locations: two in Northern Europe (UK/Ireland area), one in Central Europe (Germany/Poland area), and two in Southern Europe (Spain/Italy area).

Simple building of applications methods and

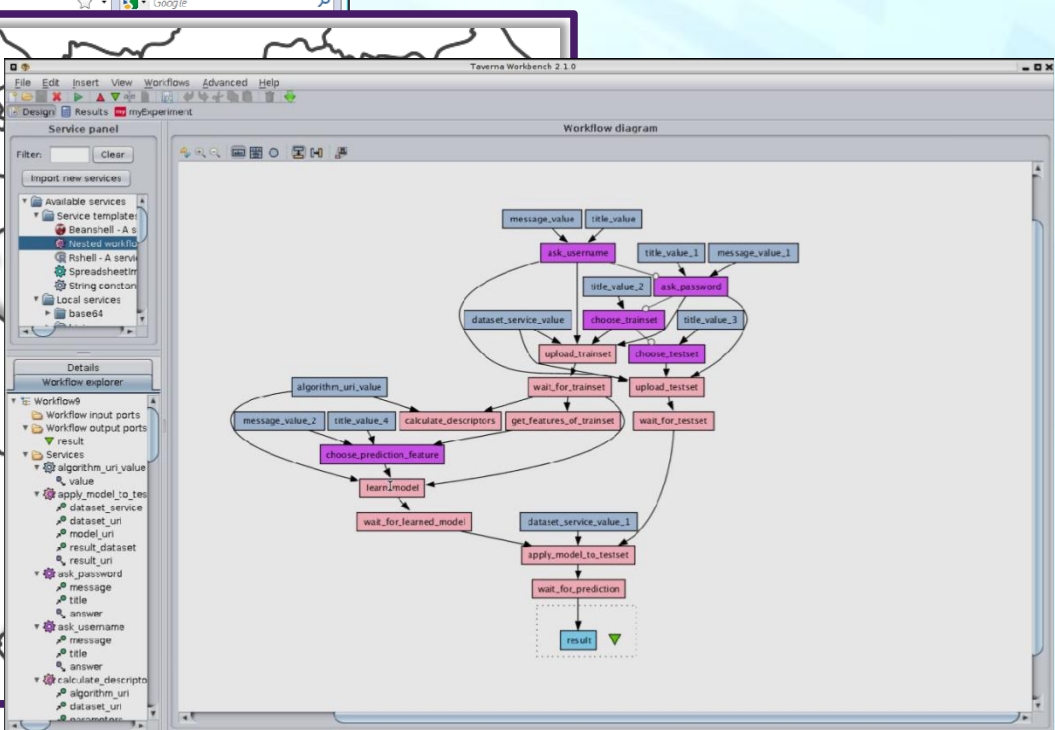
Distributed applications, integrating wide range of data, models, prediction methods

What you can do with it ...



The screenshot shows the ToxPredict web application in a Mozilla Firefox browser. The URL is <http://apps.ideaconsult.net:8180/ToxPredict/user/496763b-0d7c-4fec-8b68-15e38c29e7cd/A/step3>. The interface has three steps: Step 1: Search, Step 2: Verify, and Step 3: Models. A table lists various endpoints and the models used for prediction.

Endpoint	Model	Descriptors
Carcinogenicity	OpenTox model created with TUM's kNN regression model learning web service.	YES
Dissociation constant (pKa)	ToxTree: Benigni/Bossa rules for carcinogenicity and mutagenicity	-
Endpoints	ToxTree: Structure Alerts for the in vivo micronucleus assay in rodents	-
Endpoints	ToxTree: Michael acceptors	-
Eye irritation/corrosion	ToxTree: Eye irritation	-
Human health effects	ToxTree: Extended Cramer rules	-
Human health effects	ToxTree: ILSI/Kroes decision tree for TTC	-
Skin irritation/corrosion	ToxTree: Skin irritation	-



The screenshot shows the Taverna Workflow editor interface. The workflow diagram includes steps such as 'ask_username', 'ask_password', 'choose_trasnet', 'upload_trasnet', 'wait_for_trasnet', 'get_features_of_trasnet', 'wait_for_trasnet', 'choose_prediction_feature', 'learn_model', 'wait_for_learned_model', 'dataset_service_value_1', 'apply_model_to_testset', and 'wait_for_prediction'. The workflow is designed to integrate various services and data sources for computational biology.

Simple building of
applications
methods and

Distributed of
wide range of
methods

Integration into workflow systems for
computational biology

ToxCast

- Focus on prediction of toxicological endpoints
- Use *in vitro* data to predict *in vivo* endpoints
- Multi label approach
 - approximately 400 labels, 300 instances, 1600 features

Structure	<i>In vitro</i>			<i>In vivo</i>		
	ncgc_ar_agonist	ncgc_ar_antagonist	...	chr_rat_thyroid	ch_rat_liver	...
Abamectin	1	0	...	0	0	...
Acephate	0	0	...	0	1	...
Acetamiprid	1	1	...	1	0	...
...

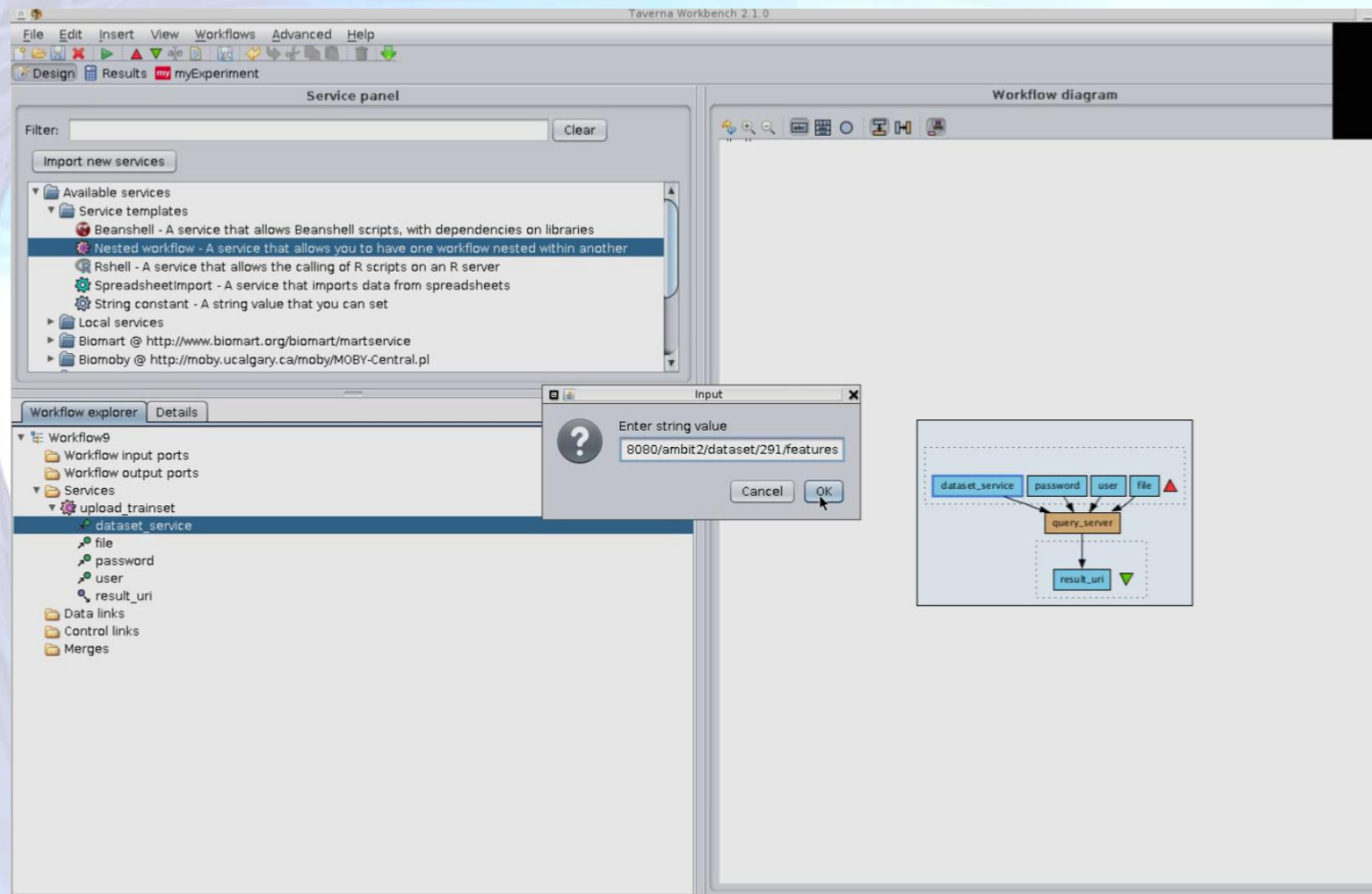
ToxCast: Approach and Further Steps

- Use all data available at this point, including data points with **missing values**
- Take into account interdependencies between *in vivo* endpoints (**multi-label classification**)
 - improves upon predicting endpoints individually
- Use **applicability domain** on *in vitro* data to avoid unjustified predictions
 - improves upon predicting everything
- Results comparable to other multi-label datasets
- Further steps: take into account descriptions of assays to link into biological databases

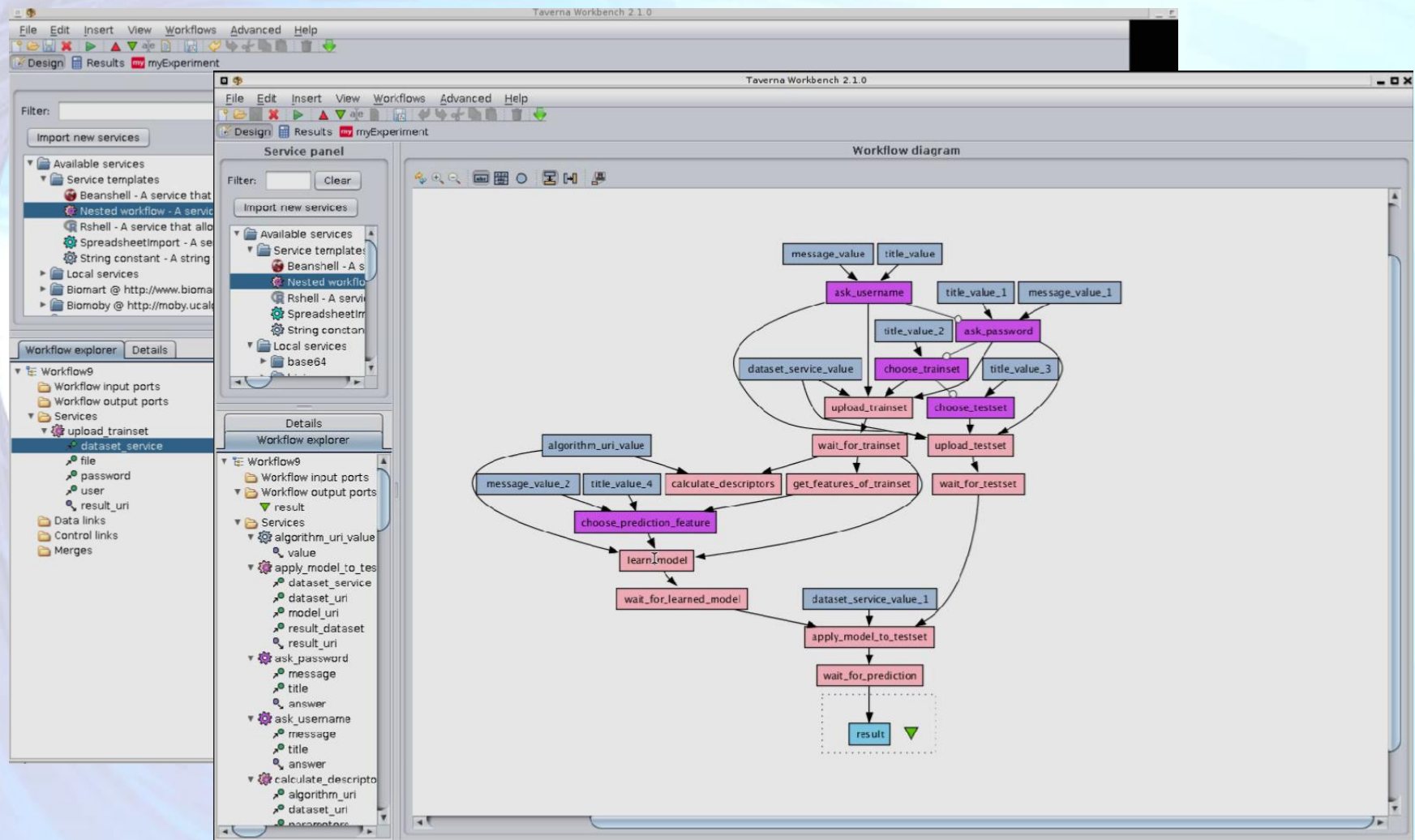
Taverna Workflow System

- Standard for workflows in computational biology
- OpenTox services can be integrated into Taverna workflows
- Allows for the integration of biological information
 - pathway data, ...
 - mechanistic explanation

Taverna Workflow System



Taverna Workflow System



Consequences for Requirements on OpenTox

User Requirements		Software Requirements
Unambiguous data	⇒	<i>formal way of representing information about data</i>
Unambiguous access	⇒	<i>well-defined interfaces</i>
Transparency of computational tools	⇒	<i>formal way of representing information about methods, well-defined interfaces</i>
Variety of user groups	⇒	<i>simplicity and modularity of design</i>
Need to integrate various resources (e.g., databases, prediction methods, models, ...) to make meaningful predictions	⇒	<i>distributed architecture, interoperability</i>
Need to integrate biological information	⇒	<i>again, modularity of design, extensibility</i>

Strategic Context/Goals

- REACH: possible reduction of test animals by using existing experimental data in conjunction with QSAR
- Also practical needs: **reporting** and form filling
- By the OECD principles, a number of requirements to a framework like OpenTox arise

	OECD Principle	OpenTox addresses by...
1	Defined Endpoint	providing a unified source of well defined and documented toxicity data with a common vocabulary
2	Unambiguous Algorithm	providing transparent access to well documented models and algorithms as well as to the source code
3	Defined Applicability Domain	integrating tools for the determination of applicability domains during the validation of prediction models
4	Goodness-of-fit, robustness and predictivity	providing scientifically sound validation routines for the determination of errors and confidences
5	Mechanistic interpretation (if possible)	integrating tools for the prediction of toxicological mechanisms and the recording of opinions and analysis in reports

Summary and Future Work

- Comprehensive framework for predictive toxicology that allows you to address a wide range of tasks
 - interface definitions, services (also for: validation, **reporting**, ...), use cases
 - **interoperability** and **extensibility by design**
- Further work on other use cases: e.g.
 - Read Across (technical infrastructure available)
 - ToxCast
 - pathway prediction service: developing a service for predicting transformation products
 - Synergy pilot
 - ...

OpenTox Application Demonstrations

Nina Jeliazkova (Ideaconsult, Bulgaria)

Christoph Helma (In Silico Toxicology, Switzerland)

Andreas Karwath (Albert-Ludwigs Univ. Freiburg, Germany)

Use Cases

ToxPredict

- offers easy access for toxicological hazard estimation of a chemical structure
- for non-QSAR specialists
- a simple yet easy-to-use user interface

ToxCreate

is aimed at researchers in

- life sciences and toxicology
- QSAR experts
- people interested in machine learning/statistics
- pharmaceutical industry R&D

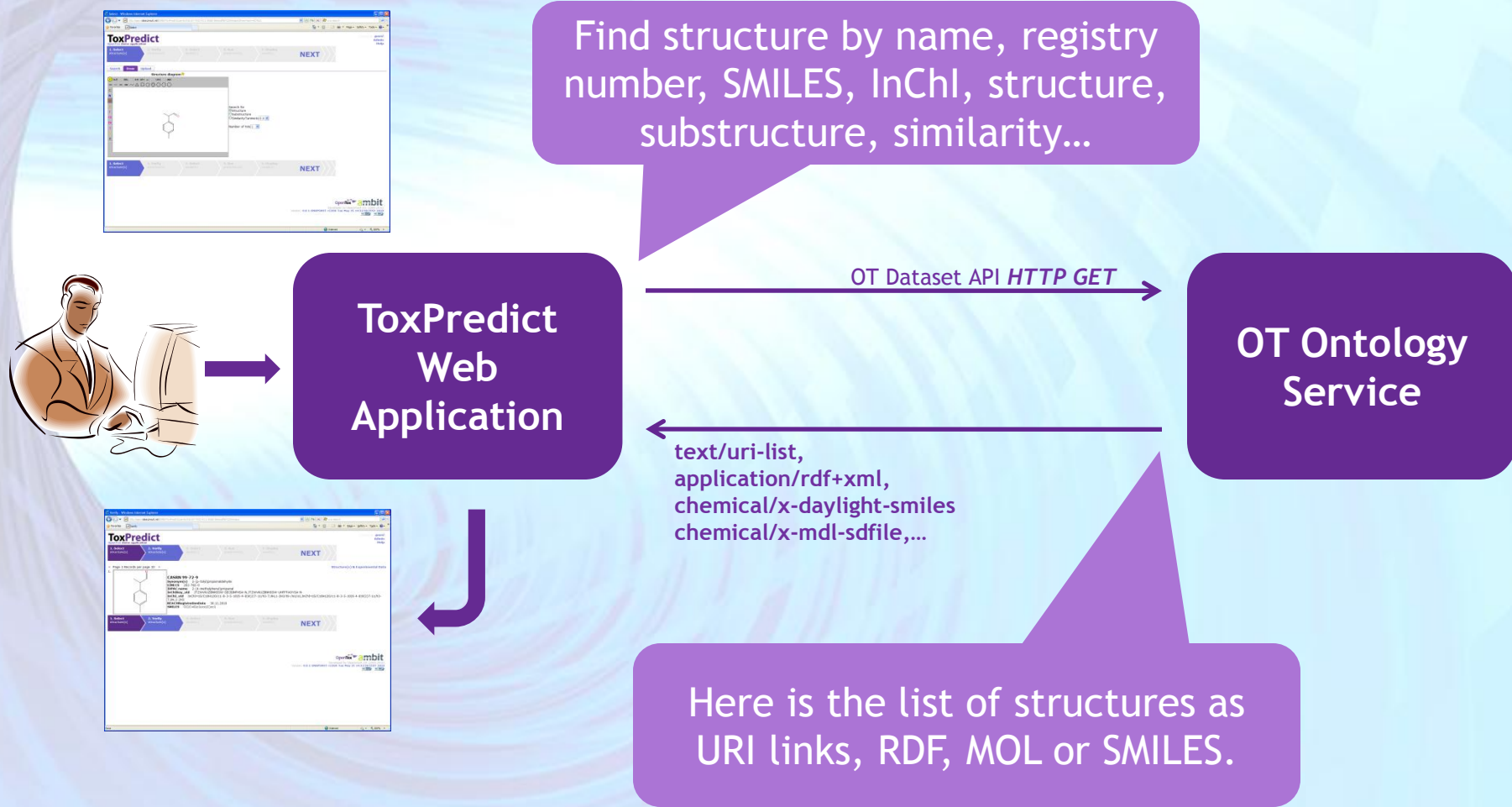
ToxPredict

The use case can be divided into the following five steps:

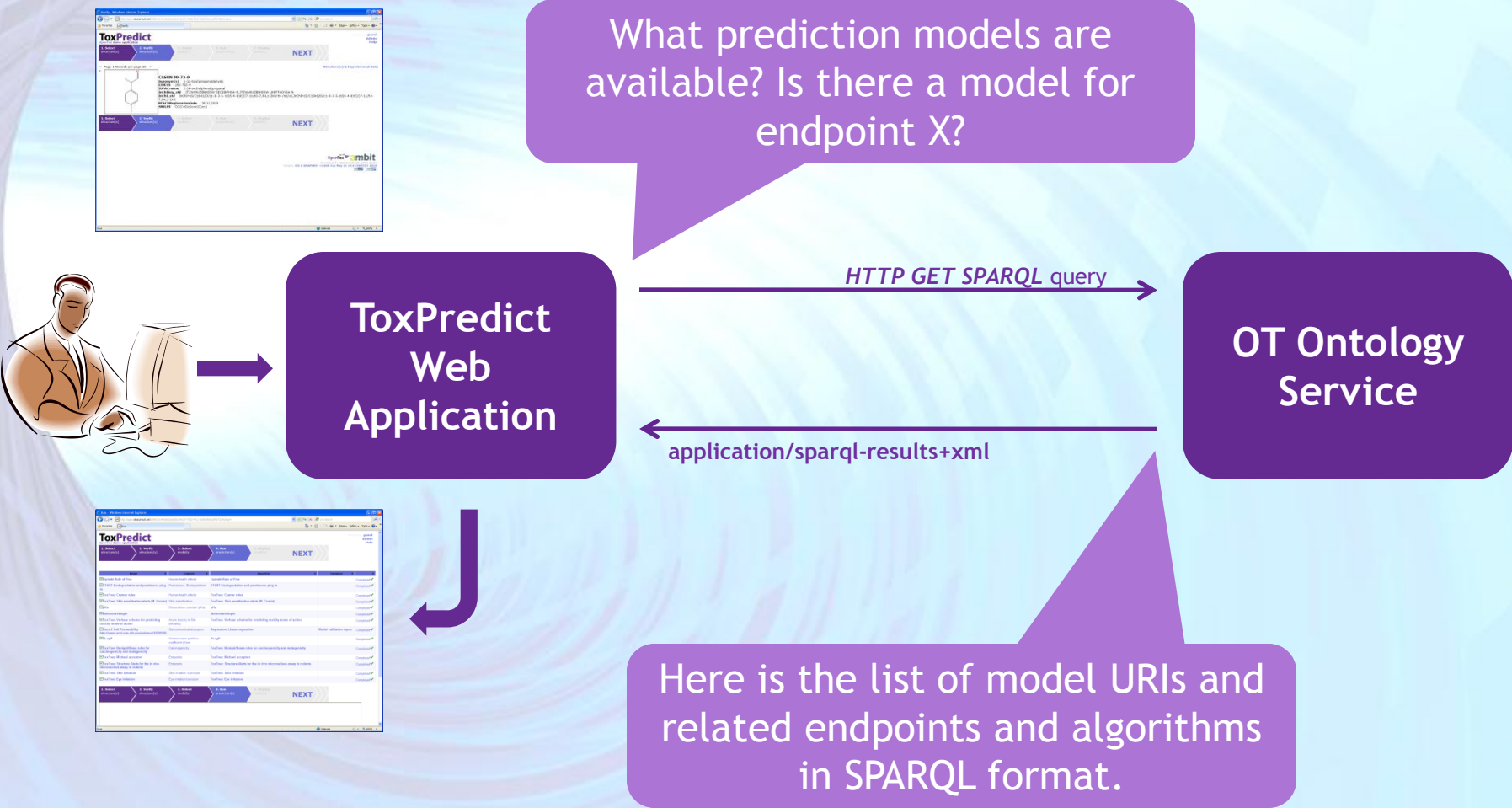
1. Enter/select a chemical compound
2. Display selected/found structures
3. Select models
4. Perform the estimation
5. Display the results

→ Live Demo: www.toxpredict.org

ToxPredict: Step 1 (behind the scenes)



ToxPredict: Step 3 (behind the scenes)



ToxCreate: intended audience

Toxicologists with

- access to toxicological data
- basic computer skills
- little or no knowledge of QSAR algorithms

Goal:

- build and validate prediction models from user provided training data

→ Live demo www.toxcreate.org

Future Development

Inclusion of further QSAR algorithms

- build and validate multiple models
- automated selection of the best model (based on validation results)

GUI refinement:

- based on user feedback

Expert interface:

- access to all model building parameters
- command line/GUI versions

Behind the Scenes of ToxCreat

ToxCreat

CreateInspectPredictAbout

This service is for testing purposes only - once a week all models will be deleted. Please send bug reports and feature requests to our [issue tracker](#).

This service creates **lazar** classification models (more model building algorithms will follow) from your uploaded datasets. Here are [instructions](#) , for creating training datasets in Excel.

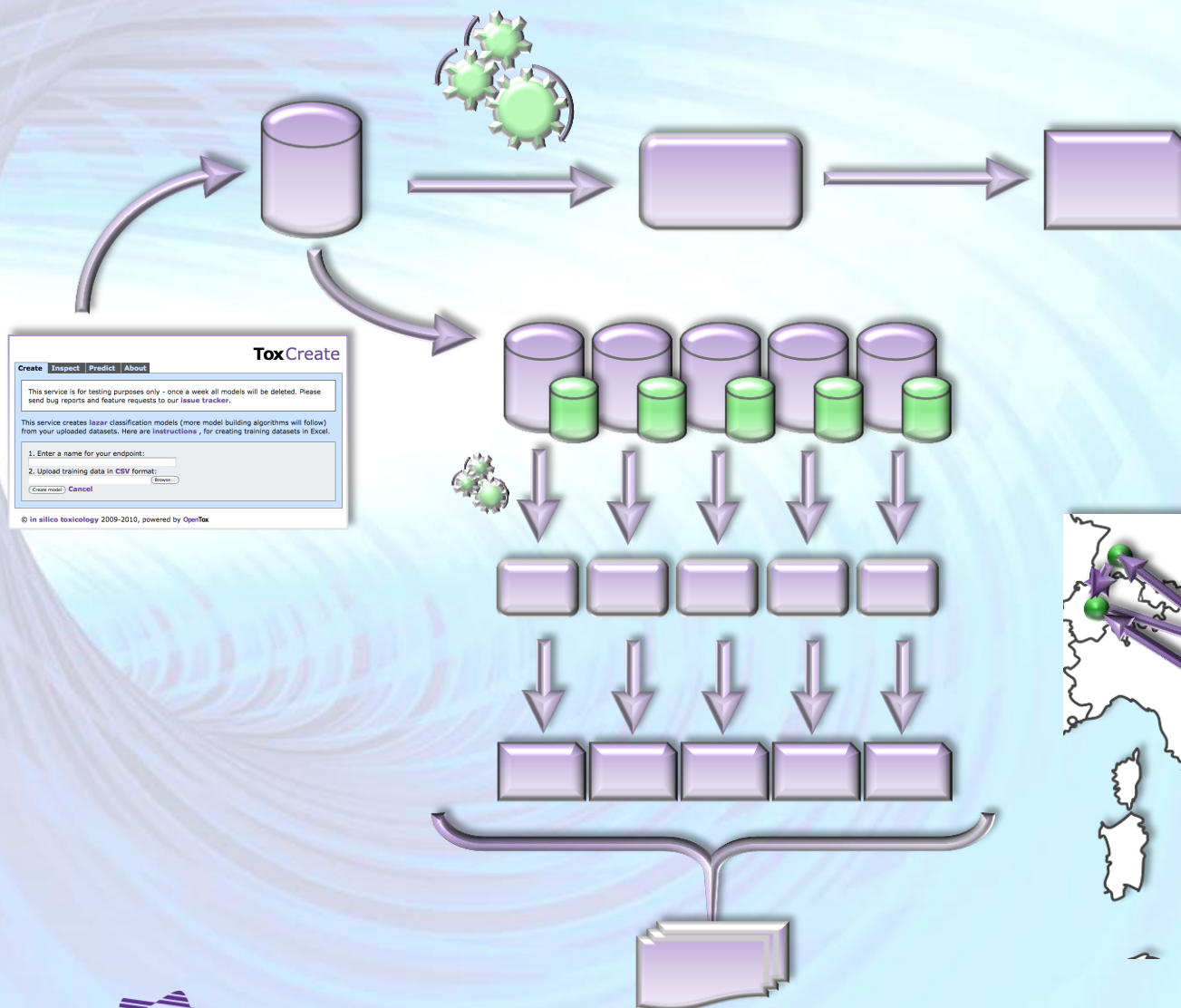
1. Enter a name for your endpoint:

2. Upload training data in **CSV** format:

© in silico toxicology 2009-2010, powered by [OpenTox](#)

OpenTox

Behind the Scenes of ToxCreat



Current State of the System

Web services online:

AlgorithmWS: NTUA, TUM, IDEA, IST

ModelWS: NTUA, TUM, IDEA, IST

FeatureWS: NTUA, TUM, IDEA, IST, ALU-FR

CompoundWS: NTUA, TUM, IDEA, IST

ValidationWS: ALU-FR

DatasetWS: NTUA, TUM, IDEA, IST, ALU-FR

Collaboration, Sustainability & Future Directions

Barry Hardy

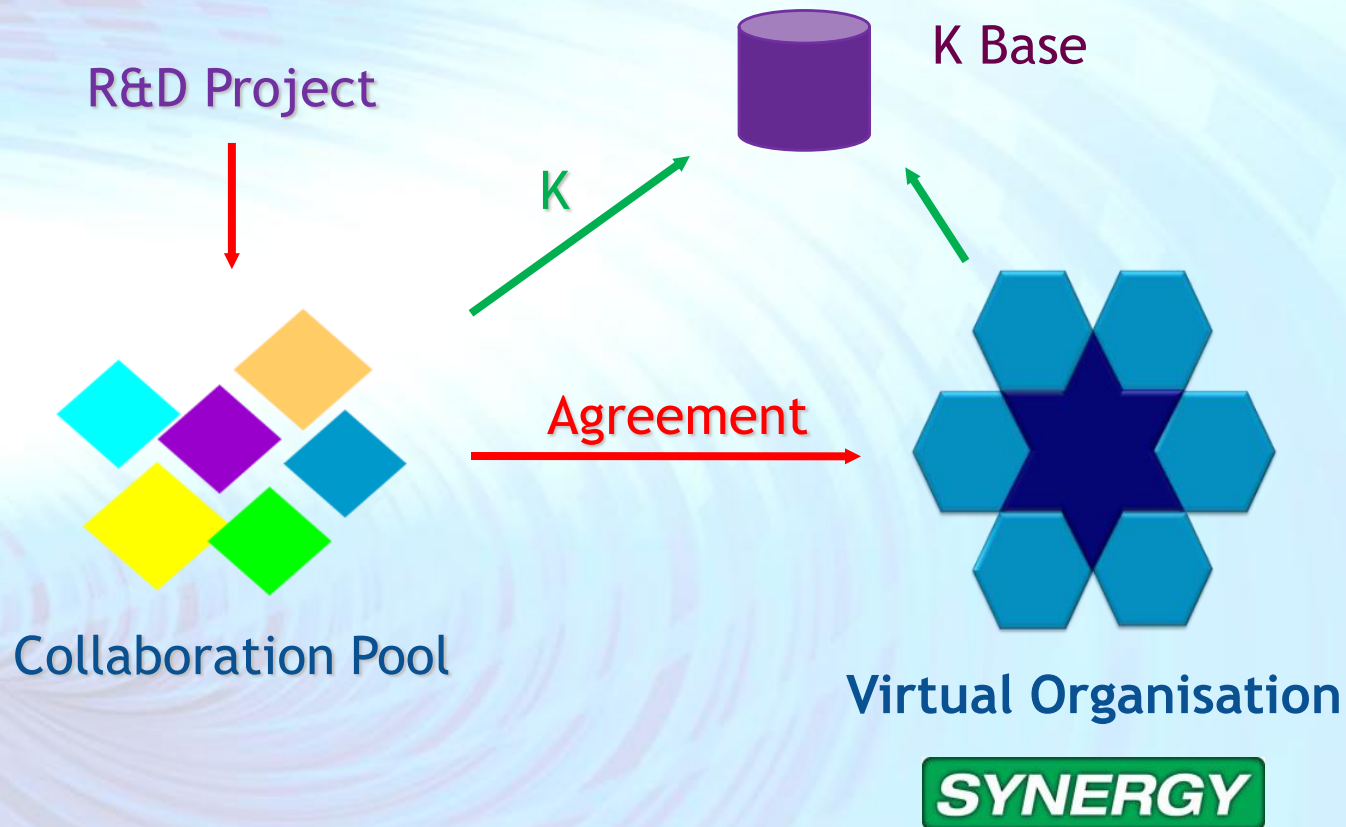
**Douglas Connect
OpenTox Project Coordinator**

Our Drivers - Taking on Technical, Cultural and “Other” Challenges of the Unexpected

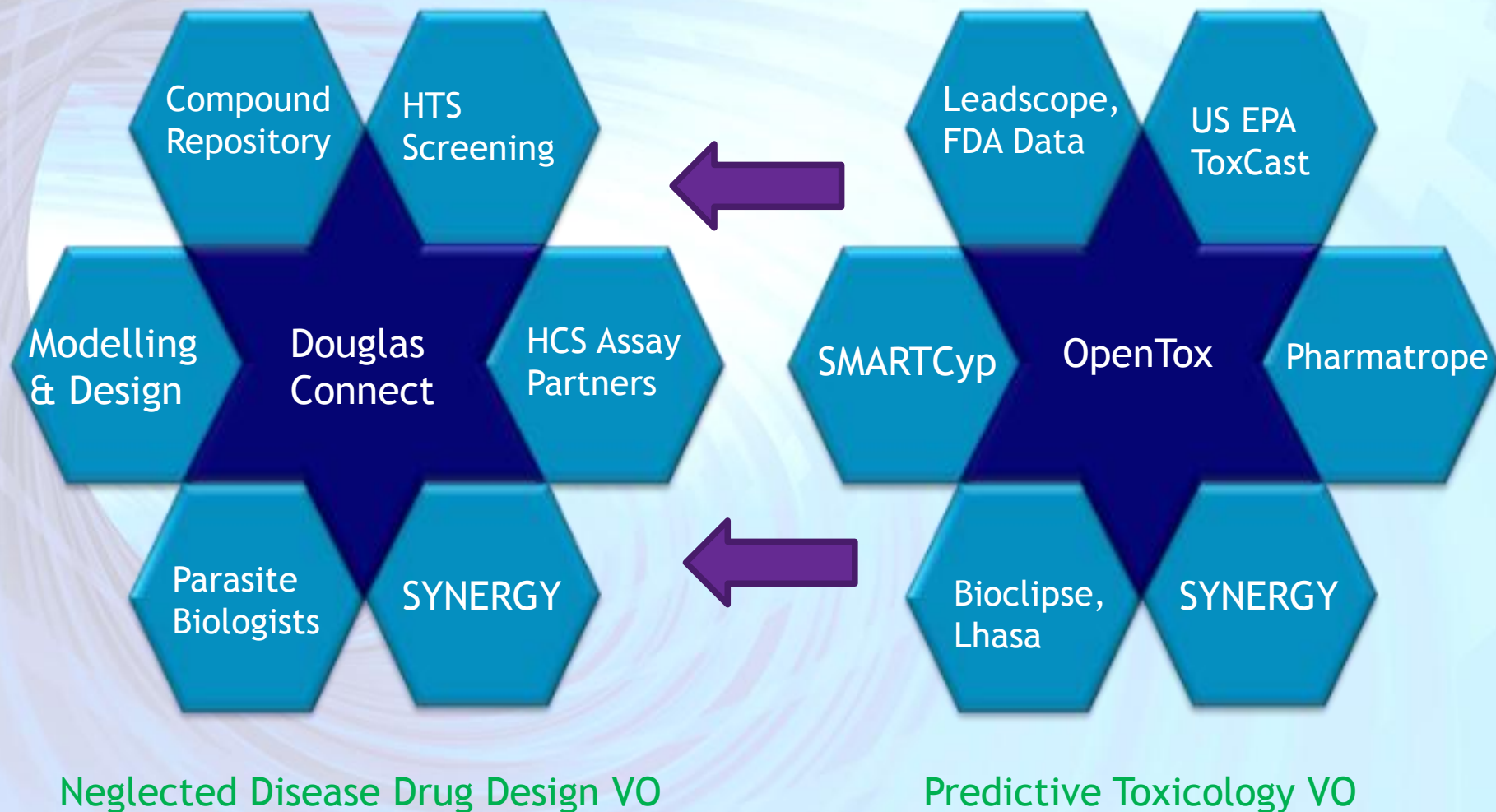


[Visit with Lions at Mukuni Reintroduction Project, Livingstone, Zambia](#)

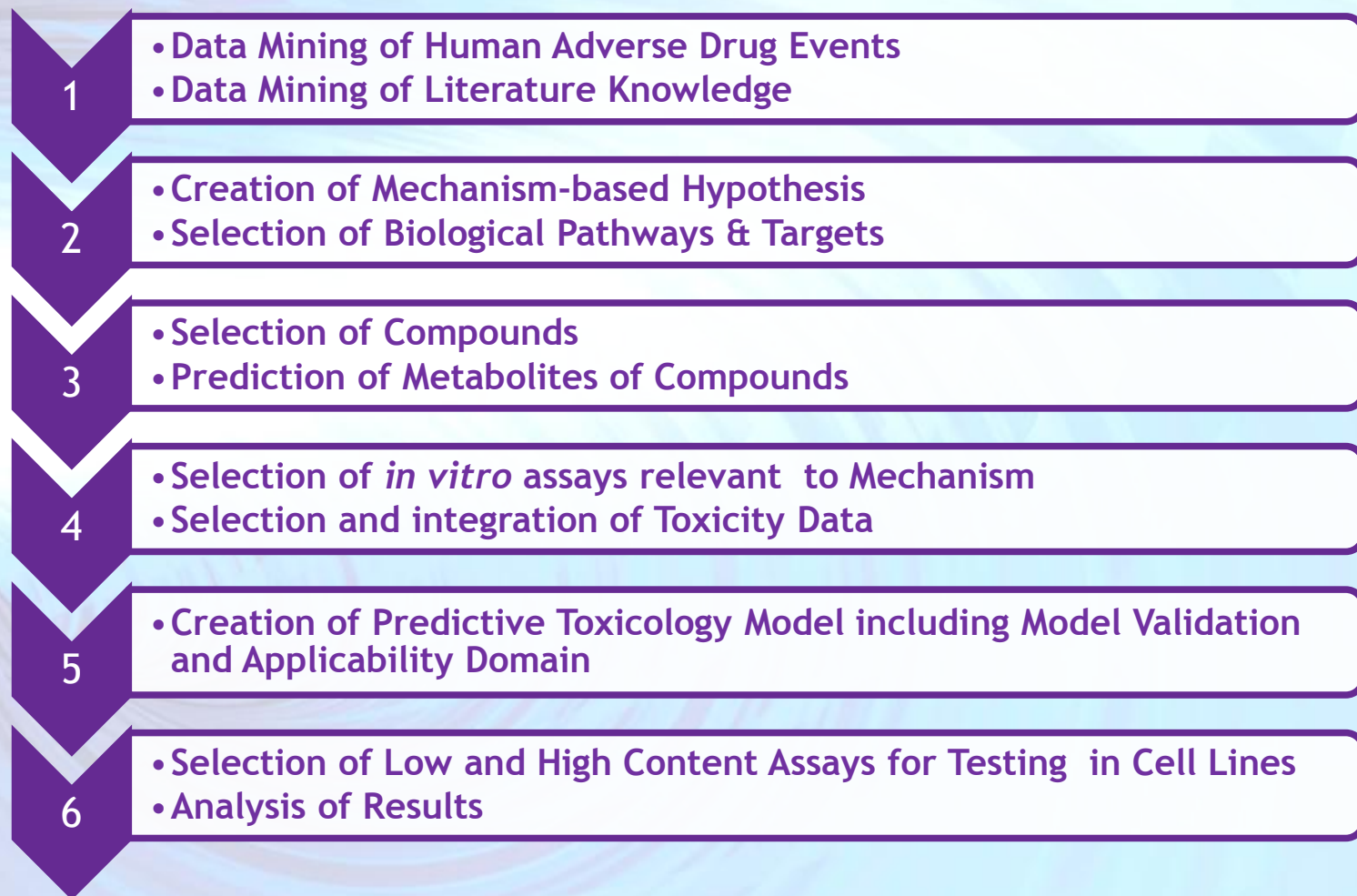
Virtual Organisation Pilots



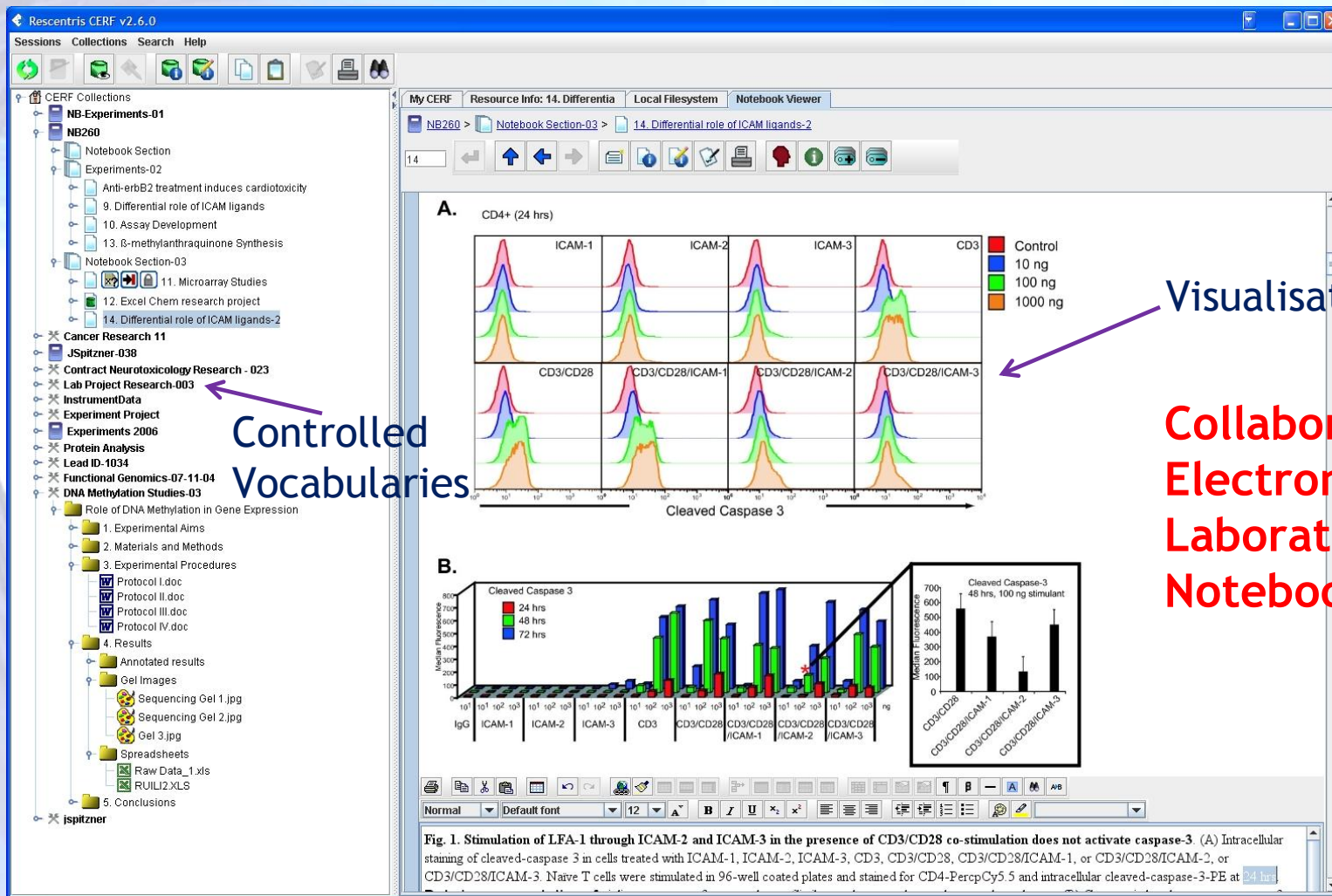
Virtual Organisation Pilots



OpenTox - Synergy Predictive Toxicology VO Pilot Strategy Development & Case Study



Recording of Collaborative R&D

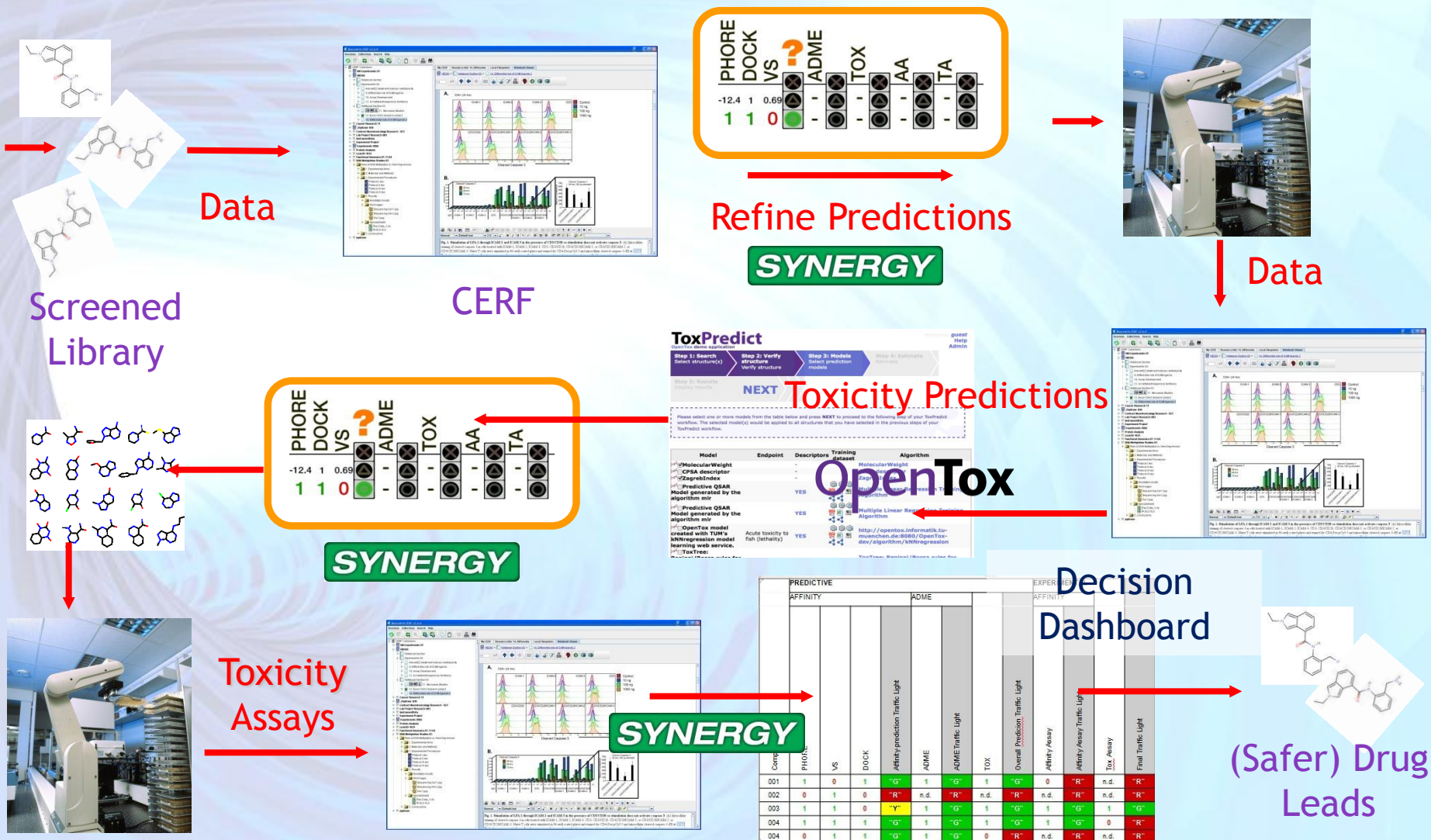


Controlled
Vocabularies

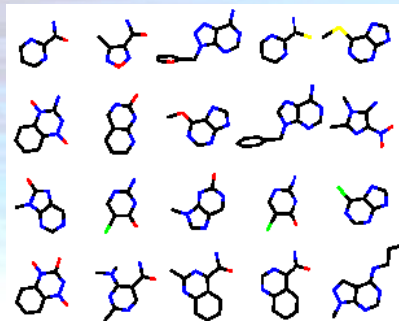
Visualisation

Collaborative
Electronic
Laboratory
Notebook (ELN)

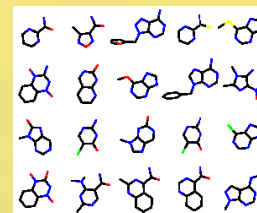
Synergy Collaboration Pilots



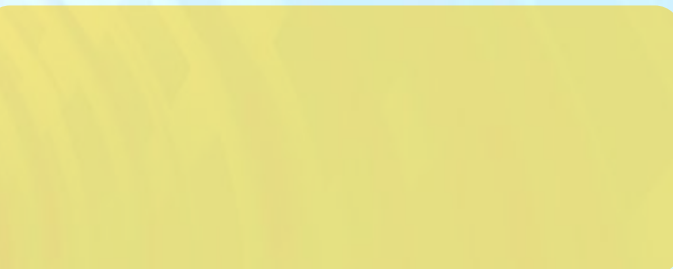
1. A library of compounds is entered to the ELN



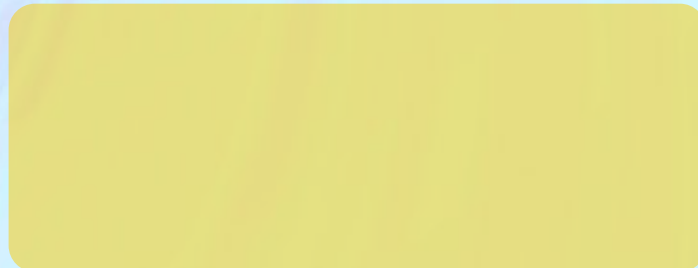
ELN



Synergy

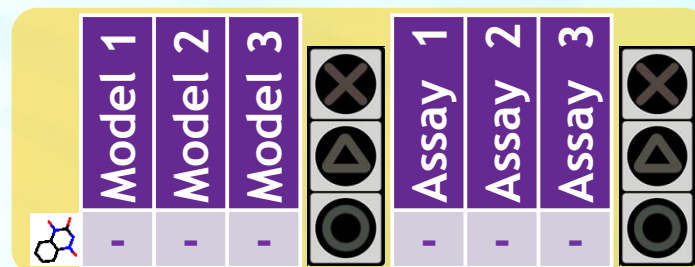


OpenTox

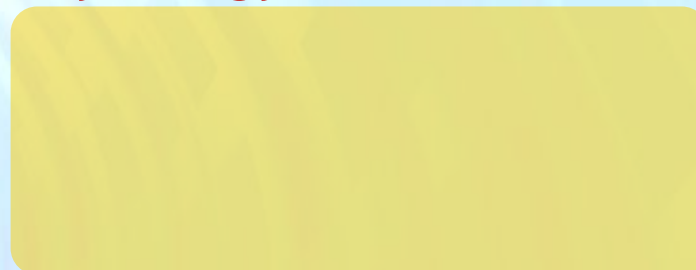


2. Each compound is assigned a data structure in ELN

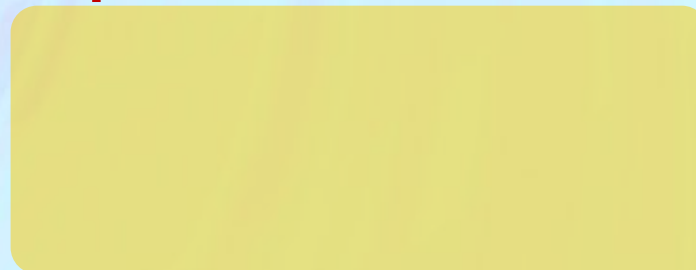
ELN



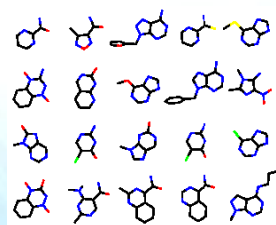
Synergy



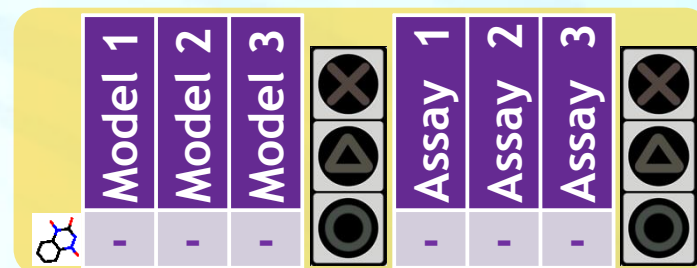
OpenTox



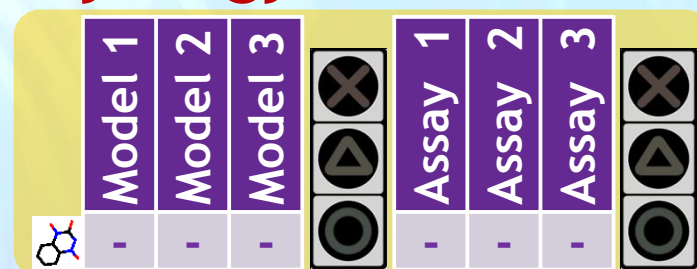
3. ELN passes compounds to OpenTox and SYNERGY



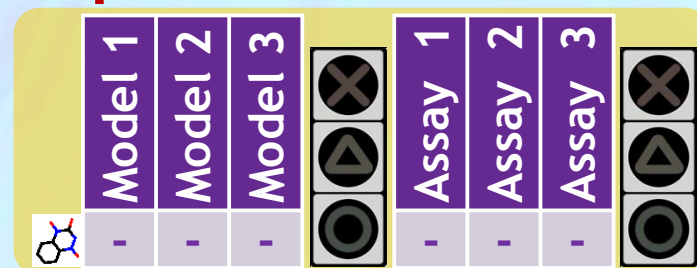
ELN



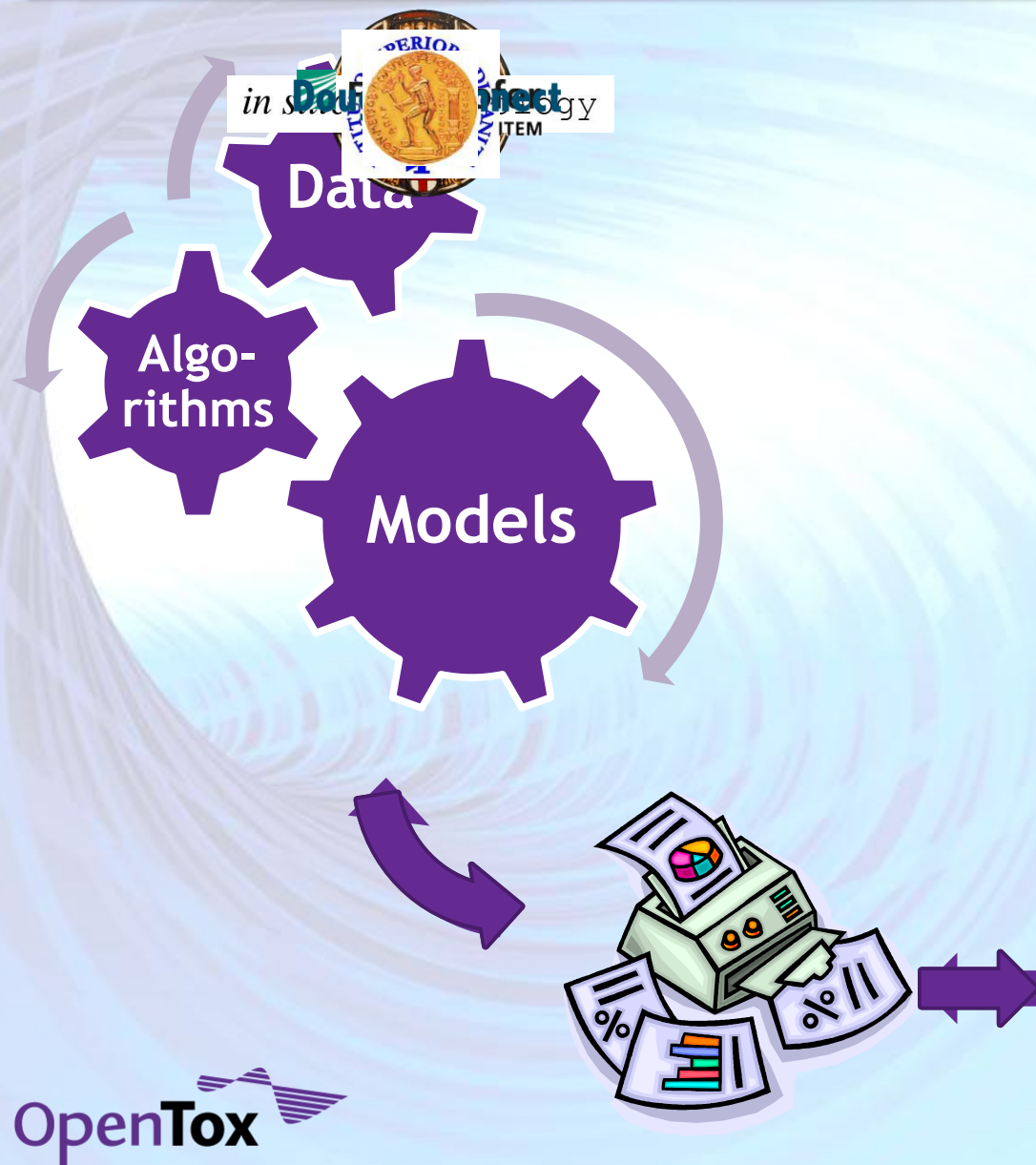
Synergy










OpenTox










4. OpenTox computes toxicity predictions










ELN

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	-	-	-		-	-	-	
	-	-	-		-	-	-	
	-	-	-		-	-	-	

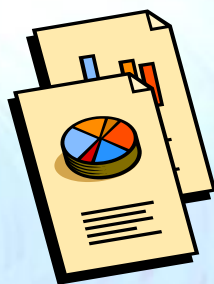
Synergy

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	-	-	-		-	-	-	
	-	-	-		-	-	-	
	-	-	-		-	-	-	

OpenTox

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	1	0	1		-	-	-	
	-	-	-		-	-	-	
	-	-	-		-	-	-	

5. OpenTox sends back a report to ELN



ELN

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	1	0	1		-	-	-	

Synergy

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	-	-	-		-	-	-	

OpenTox

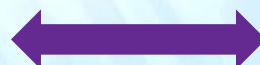
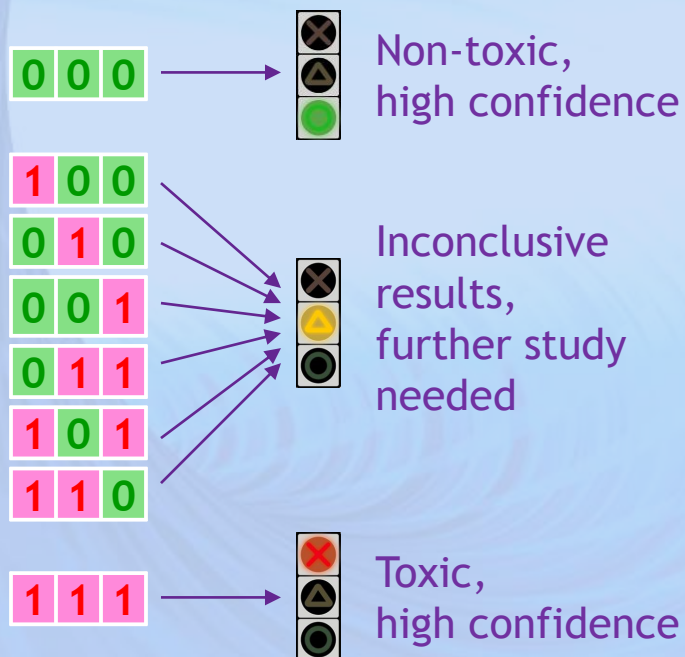
	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	1	0	1		-	-	-	

6. ELN sends the results to SYNERGY

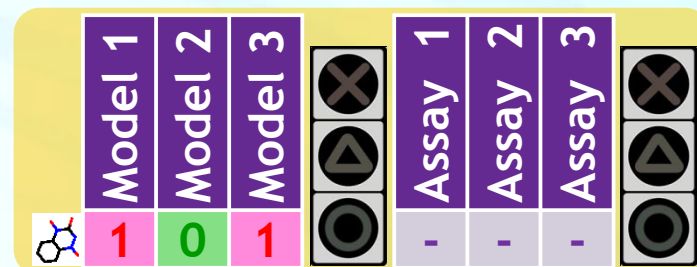


7. SYNERGY applies the Recommendation Rules

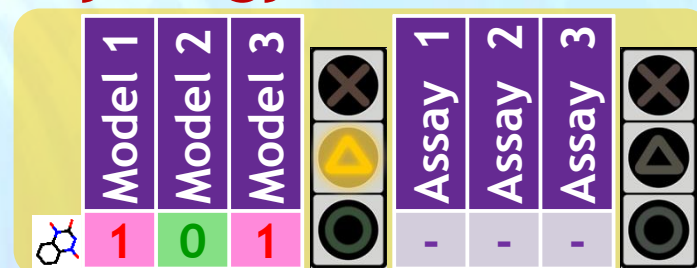
Recommendation Rules:



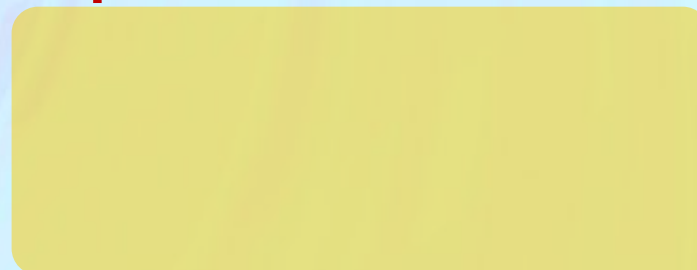
ELN



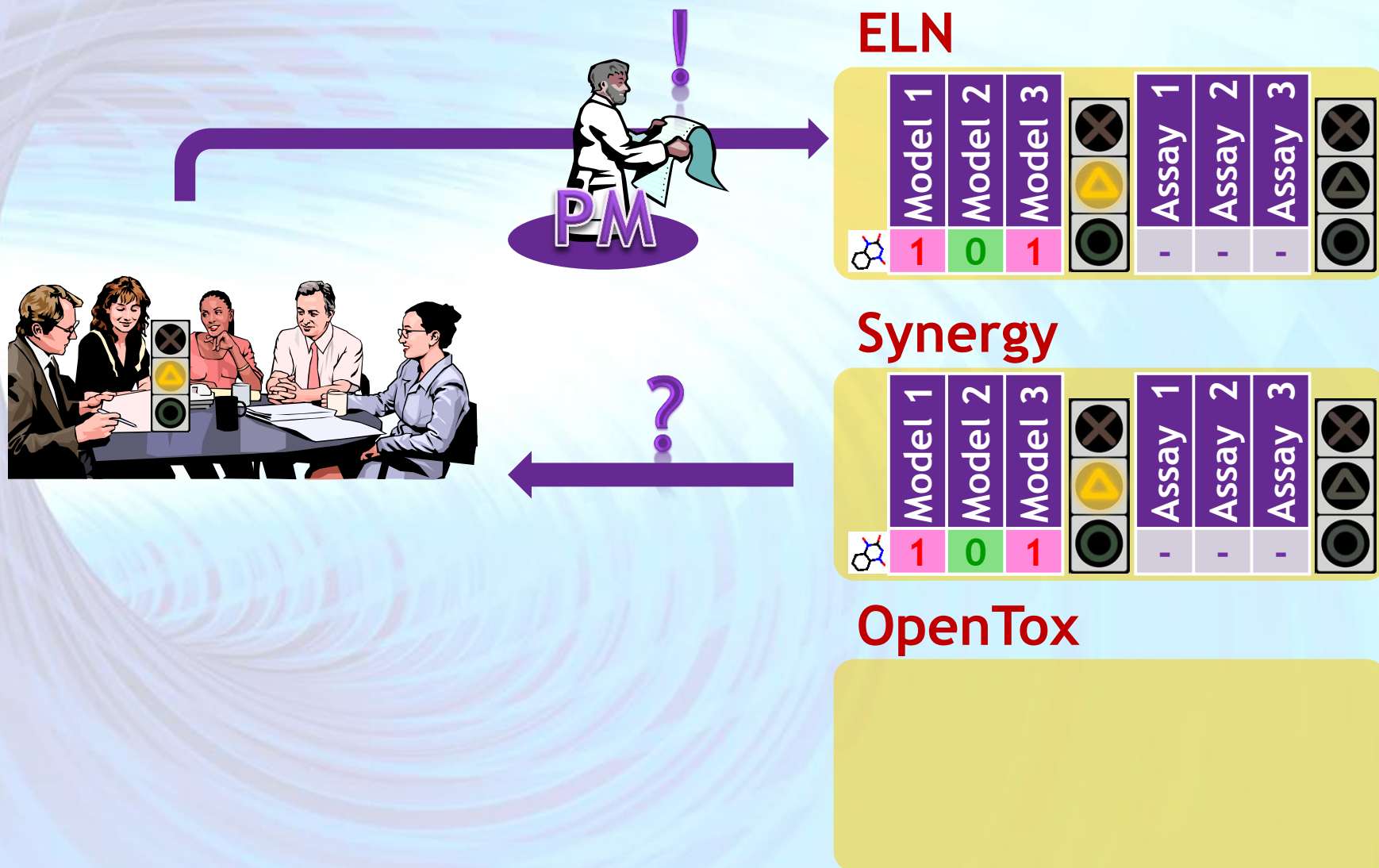
Synergy



OpenTox


















8. Inconclusive data → SYNERGY calls a meeting















9. Experimental assays confirm toxicity



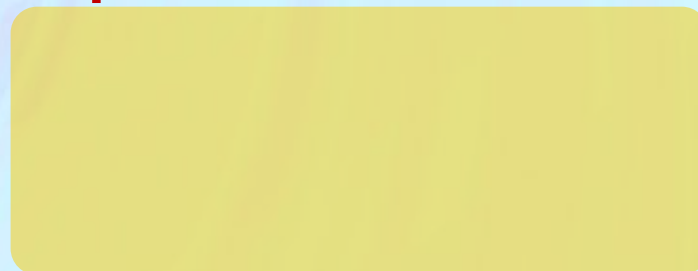
ELN

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	1	0	1		-	1	1	
								
								

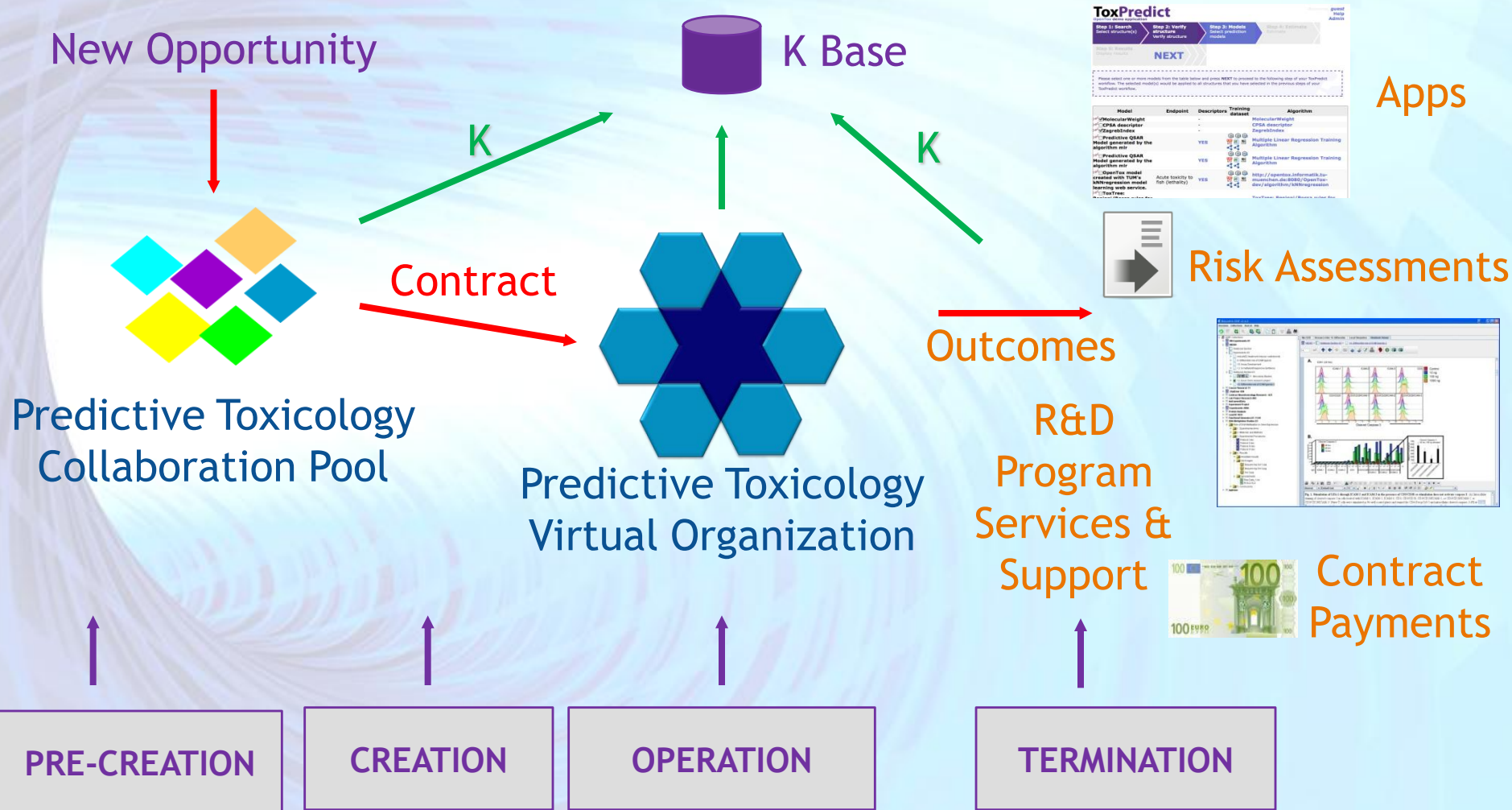
Synergy

	Model 1	Model 2	Model 3		Assay 1	Assay 2	Assay 3	
	1	0	1		-	-	-	
								
								

OpenTox



Sustainability Model



ToxPredict

Step 1: Search
Select structure(s)

Step 2: Verify structure
Verify structure

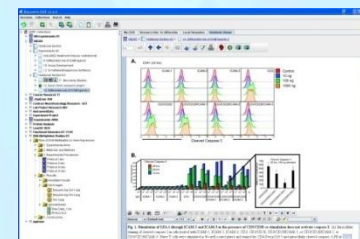
Step 3: Model
Using prediction model

Step 4: Results
Results

NEXT

Please click one or more results from the table below and click **Next** to proceed to the following step of your ToxPredict workflow. The selected model(s) will be applied to all structures that you have selected in the previous steps of your ToxPredict workflow.

Model	Endpoint	Description	Training dataset	Algorithm
✓ PhenolWeight	-	-	WaterfordMethyl	-
✓ CPA descriptor	-	-	CPA descriptor	-
✓ Organophosphate	-	-	Organophosphate	-
✓ Predictive QSAR	-	-	-	-
Model generated by the algorithm mlr	YES	Multiple Linear Regression Training	Algorithm	-
✓ Predictive QSAR	-	-	-	-
Model generated by the algorithm mlr	YES	Multiple Linear Regression Training	Algorithm	-
✓ Report the model	-	-	-	-
Report the model created with ToxPredict	YES	Report the model created with ToxPredict	Report the model created with ToxPredict	-
✓ Report the model	-	-	-	-
Report the model created with ToxPredict	YES	Report the model created with ToxPredict	Report the model created with ToxPredict	-



Acknowledgements - OpenTox Partners

In Silico Toxicology,
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Switzerland

Albert Ludwigs University
Freiburg, Germany

Ideaconsult,
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di Sanità, Italy

Technical University
of Munich, Germany



National Technical
University of Athens,
Greece

Fraunhofer Institute
for Toxicology &
Experimental Medicine,
Germany

David Gallagher, UK

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Academy of Medical
Sciences, Russia

Seascape Learning,
India

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Andreas Maunz
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Georgia Melagraki
Antreas Afantitis
Pantelis Sopasakis
David Gallagher
Vladimir Poroikov
Dmitry Filimonov
Alexey Zakharov
Alexey Lagunin

Tatyana Gloriovova
Sergey Novikov
Natalia Skvortsova
Sunil Chawla
Steve Bowlus
Indira Ghosh
Surajit Ray
Gaurav Singhai
Om Prakash
Sylvia Escher
Sara Weiss
Helvi Grimm

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- University of North Carolina
- EC Environment Directorate General
- Organisation for Economic Cooperation & Development
- CADASTER
- Bayer Healthcare

Final words...

For more information, visit

www.opentox.org

Contact me:

barry.hardy@douglasconnect.com

**Many thanks for your
attention!**



OpenTox - An Open Source Predictive Toxicology Framework, www.opentox.org, is funded under the EU Seventh Framework Program: HEALTH-2007-1.3-3 Promotion, development, validation, acceptance and implementation of QSARs (Quantitative Structure-Activity Relationships) for toxicology, Project Reference Number Health-F5-2008-200787 (2008-2011).