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Cell Networks

*Providing an Insight into your Biological Systems*

## Mechanism-based Applications of Toxicology Ontology

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(presented by Dragana Mitic Potkrajac, PhD)

OpenTox 2011 InterAction Meeting Program,  
9-12 August 2011, Munich



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I. Mechanism-based application requires complex and structured information

II. Ambiguous how detailed information must be:

- in some cases inferred protein-chemical interaction networks are sufficient
- in other cases detailed models of *how* a particular pathway brings about a particular outcome is required



## CCNet's ToxOntology



1. Organizing and managing information about toxic effects
2. Providing as much insight as possible about the mechanisms underlying a toxic effect
3. Helping elucidate underlying biological pathways

Aim is to interpret and explain mechanism of action of drugs/compounds by precisely categorizing terms describing observation/toxicities



## CCNet's ToxWiz™ Ontologies

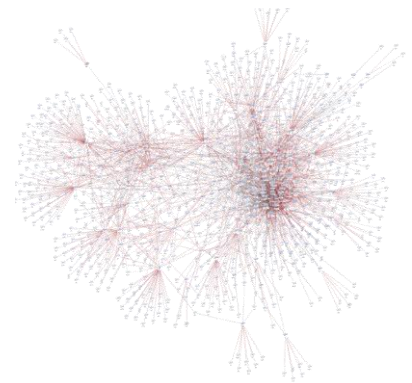


**What are they?**

**What purpose to these ontologies serve?**

**What are the requirements to create this ontology?**

**How do we create, validate and optimize the ontology?**

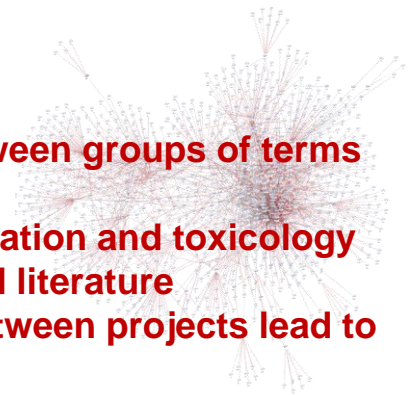


## CCNet's ToxWiz™ Ontologies



### What are they?

- **Controlled vocabularies/terms**
- **Classified into groups & hierarchies**
- **Defined relationships between terms and between groups of terms**
- **A product of over 150 man years of expert curation and toxicology information extraction from safety reports and literature**
- **6 iterations over 8 years for harmonization between projects lead to increased interoperability**
- **912 histopathology terms and 500000 synonyms, over 2 million relationships**



### What purpose to these ontologies serve?

What are the requirements to create this ontology?

How do we create, validate and optimize the ontology?



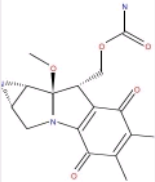
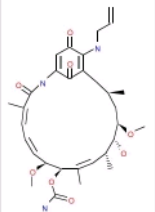
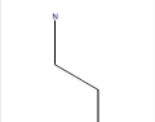
**ToxWiz - Cambridge Cell Networks**

File Search Compare Tools Help

Searching Molecules by Name:  Search

Pathways Properties Legend

- Metabolic pathways
- Signalling pathways
- Toxicology pathways
- Disease clusters
- Toxic endpoint clusters
  - Activity system (e.g. Muscle, bone)
  - Circulatory system
  - Connective and other tissues & cells
  - Development
  - Digestive system (excluding Hepatic)
  - Endocrine system
  - Female Reproductive system
  - Hepatic system
    - A
      - Bile duct carcinoma cluster
      - Bile duct fibrosis cluster
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| Rank | Image   | Molecule Name    | Evidence/Pubmed Abstracts  |
|------|---|------------------|--|
| 1    |    | mitomycin C (2)  | <p><b>PIMD: 12720104</b> Int J Clin Oncol 2003 Apr;8(2):110-2<br/> <a href="#">Liver metastasis of pancreatic cancer managed by intra-arterial infusion chemotherapy combined with</a></p> <p><b>PIMD: 12484005</b> Gan To Kagaku Ryoho 2002 Nov;29(12):2071-3<br/> <a href="#">[A case of intrahepatic bile duct necrosis following hepatic arterial infusion chemotherapy]</a></p> |
| 2    |  | 17-AAG (1)       | <p><b>PIMD: 16081239</b> Toxicol In Vitro 2005 Dec;19(8):1079-88<br/> <a href="#">In vitro detection of differential and cell-specific hepatobiliary toxicity induced by geldanamycin and</a></p>  |
| 3    |  | beta-alanine (1) | <p><b>PIMD: 9328219</b> Toxicology 1997 Oct 19;122(3):193-204<br/> <a href="#">Modulation of taurine levels in the rat liver alters methylene dianiline hepatotoxicity.</a></p>  |

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# CCNet's ToxWiz™ Ontologies



What are they?

What purpose do the ontologies serve?

- To facilitate prediction of toxic effects – *prospective analysis*
- To help explain causes of toxic effects – *retrospective analysis*
- To elucidate modes of action and create hypothesis for MOA
- To support extraction process of knowledge relevant to toxicology from toxicology reports and literature
- To enable integration and transfer of findings to clinical observations

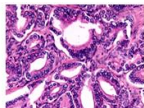
What are the requirements to create this ontology?

How do we create, validate and optimize the ontology?

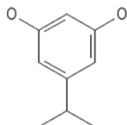


# Ontology supported discovery

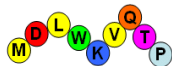
Pathology



Structure



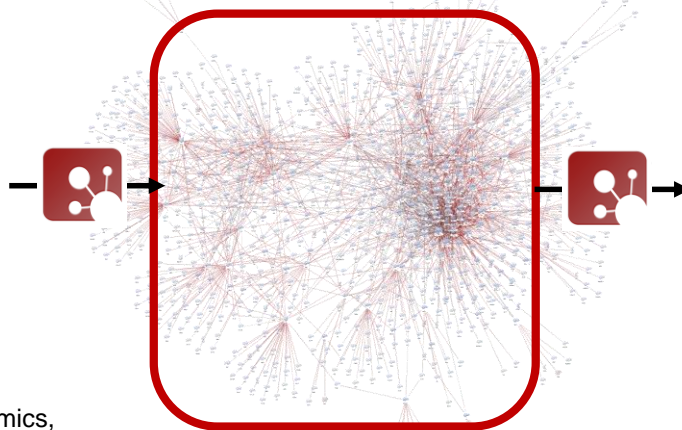
Target/receptor



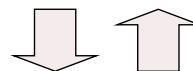
Genomics, proteomics, metabolomics



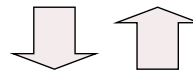
Toxwiz  
Knowledge Database  
& Browser



Toxicity & pharmacology prediction



Mechanistic hypotheses



Biomarker identification

150 man-years of expert curation, including over 1000 catalogued toxic endpoints & associated mechanistic hypotheses





## Case Study: Phthalic Acid and Testis Degeneration

### How Ontologies permit faster interpretation

#### Parameters and Statistics

#### Toxic endpoint clusters

| Rank | Molecular Mechanism   | Type                     | Molecules Inside                                | Molecules Next To   |
|------|---|--------------------------|---|---|
| 1    | <a href="#">Testis atrophy induction cluster</a>            | Male Reproductive system | 3: Dipentyl phthalate, dibutyl phthalate, Di... |   |
| 2    | <a href="#">Testis atrophy association cluster</a>          | Male Reproductive system | 4: Dipentyl phthalate, Mehp, dibutyl phtha...   | 1: Dioctyl phthalate  |
| 3    | <a href="#">Testis toxicities association cluster</a>       | Male Reproductive system | 4: Dipentyl phthalate, Mehp, dibutyl phtha...   | 1: Dioctyl phthalate  |
| 4    | <a href="#">Leydig cell hyperplasia cluster</a>             | Male Reproductive system | 7: Palatinol IC, Benzyl butyl phthalate, Mo...  | 4: phthalic acid, Dioctyl phthalate, Palatinol N, Didp          |
| 5    | <a href="#">Leydig cell toxicities cluster</a>              | Male Reproductive system | 7: Palatinol IC, Benzyl butyl phthalate, Mo...  | 4: phthalic acid, Dioctyl phthalate, Palatinol N, Didp          |
| 6    | <a href="#">Testis atrophy cluster</a>                      | Male Reproductive system | 9: Monoocetyl phthalate, Ergoplast FDC, Di...   | 4: Palatinol N, Benzyl butyl phthalate, BUTYL ISODECYL PH...    |
| 7    | <a href="#">Heart fibrosis inhibition cluster</a>           | Circulatory system       |   | 4: Mehp, dibutyl phthalate, Dioctyl phthalate, phthalic acid    |
| 8    | <a href="#">Hepatocyte hyperplasia induction cluster</a>    | Hepatic system           | 1: Dioctyl phthalate                            | 3: Mehp, dibutyl phthalate, phthalic acid                       |
| 9    | <a href="#">Leydig cell hyperplasia association cluster</a> | Male Reproductive system | 1: phthalic acid                                | 1: Dioctyl phthalate  |
| 10   | <a href="#">Spinal cord atrophy induction cluster</a>       | Nervous system           |   | 10: Palatinol IC, DIETHYL PHTHALATE, Palatinol N, Benzyl ...    |
| 11   | <a href="#">Hepatocyte hyperplasia association cluster</a>  | Hepatic system           | 1: Dioctyl phthalate                            | 3: Mehp, dibutyl phthalate, phthalic acid                       |
| 12   | <a href="#">Leydig cell toxicities association cluster</a>  | Male Reproductive system | 1: phthalic acid                                | 1: Dioctyl phthalate  |
| 13   | <a href="#">Leydig cell hyperplasia induction cluster</a>   | Male Reproductive system | 1: phthalic acid                                | 1: Dioctyl phthalate  |
| 14   | <a href="#">Liver steatosis inhibition cluster</a>          | Hepatic system           |   | 4: Mehp, dibutyl phthalate, Dioctyl phthalate, phthalic acid    |
| 15   | <a href="#">Hepatocyte hypertrophy association cluster</a>  | Hepatic system           |   | 4: Mehp, dibutyl phthalate, Dioctyl phthalate, phthalic acid    |
| 16   | <a href="#">Bile duct inflammation cluster</a>              | Hepatic system           |   | 7: Dioctyl phthalate, dibutyl phthalate, Mehp, phthalic acid... |

***Significantly over-represented molecular mechanism groups are shown in Red/Orange***

***Most significantly over-represented Organ Systems are: Reproductive System & Male Reproductive System***



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## CCNet's ToxWiz™ Ontologies



What are they?

What purpose do they serve?

What are the requirements to create this ontology?

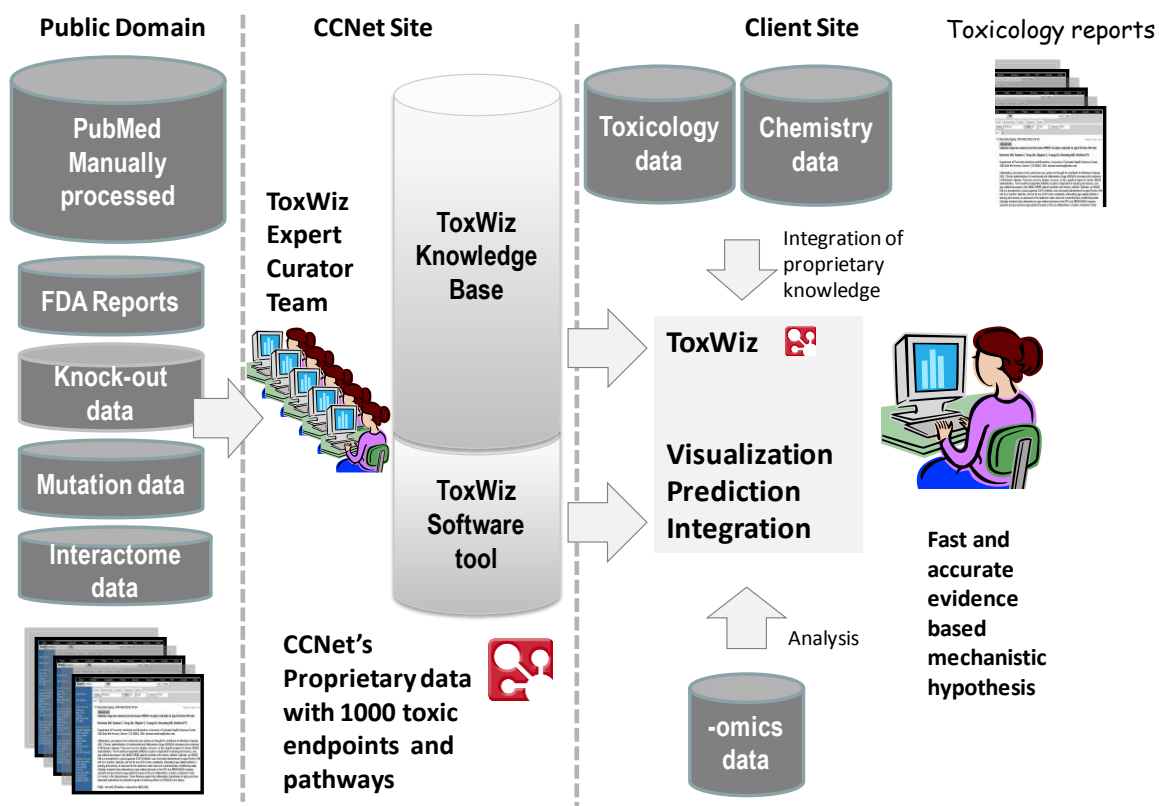
- To capture adequately toxicology test results in pre-clinical testing
- To classify and define a spectrum of histopathology findings
- To capture long term toxicity
- To capture all available knowledge from the literature and toxicology reports and deal with human way of interpreting and recording the findings
- To be interoperable with other ontology efforts

How do we create, validate and optimize the ontology?



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# Toxicity information gathering and use

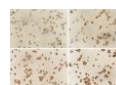
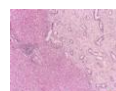
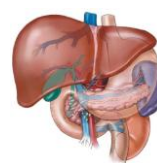


# Ontology of toxic endpoints



Endpoints are organized into categories: Example liver

| Category     | Example Toxic Endpoint Cluster | Includes                            | Description                                |
|--------------|--------------------------------|-------------------------------------|--|
| System       | Hepatotoxicity                 | Gall bladder, liver, hepatic system | General observations (e.g. clinical, etc.) |
| Organ/tissue | Liver toxicities               | Liver                               | Pathology report for any toxicity in liver |
| Organ/tissue | Liver hypertrophy              | Liver                               | Pathology report of specific toxicity      |
| Cells        | Hepatocyte neoplasia           | Cells or cell-lines                 | Result from cell-based assays              |

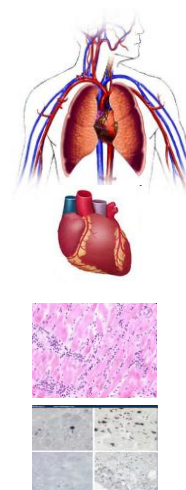


# Ontology of toxic endpoints



Endpoints are organized into categories: Example heart

| Category     | Example                  | Includes              | Description                                |
|--------------|--------------------------|-----------------------|--|
| System       | Cardiotoxicity           | Cardiovascular system | General CV health                          |
| Organ/tissue | Heart toxicities         | Heart                 | Pathology report for any toxicity in heart |
| Organ/tissue | Heart ischaemia          | Heart                 | Pathology report of specific toxicity      |
| Cells        | Cardiac myocyte necrosis | Cells or cell-lines   | Result from cells or cell-assays           |



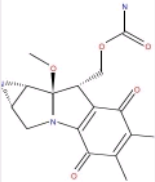
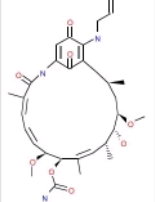
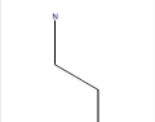
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    - K

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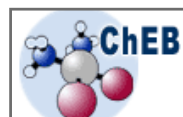
Interoperability with other ontology efforts:  
Existing ontologies but not for toxicology, although could be implemented.  
Important for all: if we know what for to use it, it make sense.



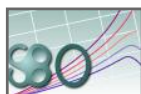
Biological Pathway Exchange



The Open Biological and Biomedical Ontologies *Pathway ontology*



*The Protein Ontology*



Systems Biology Ontology



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## Creation, validation and optimization of CCNet's Ontology

### CCNet Ontology of toxic endpoints: starting point

1. Hierarchy of organs & tissues for model organisms derived by consultation of various sources (e.g. NCBI, ENSEMBL, UniProt)

System – Organ – Components / Tissue / Cell-type  
e.g. Hepatic system – Liver – Bile duct / Kupffer Cell

N.B. includes synonyms (Biliary = Bile duct, etc.)

2. Set of histopathological observational keywords & principles  
(by consultation with toxicologists, e.g. Frank Bonner)

e.g. Hypertrophy, Hyperplasia, Neoplasia, Carcinoma, Necrosis, Fibrosis, Inflammation, etc.

3. Several principles of progression in time/pathology

e.g. Carcinoma is a subset of Neoplasia, Hyperplasia precedes Neoplasia

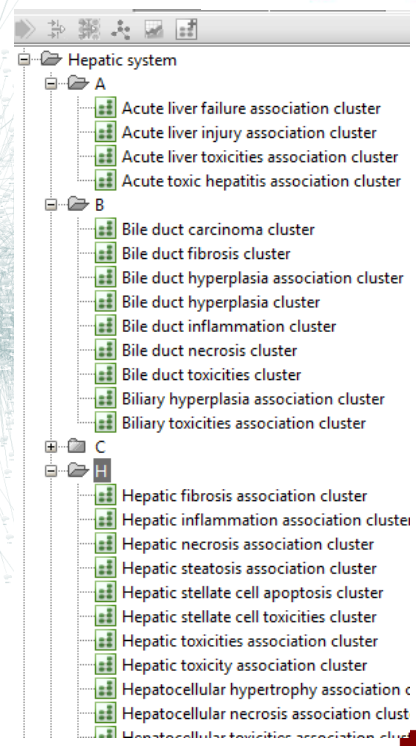
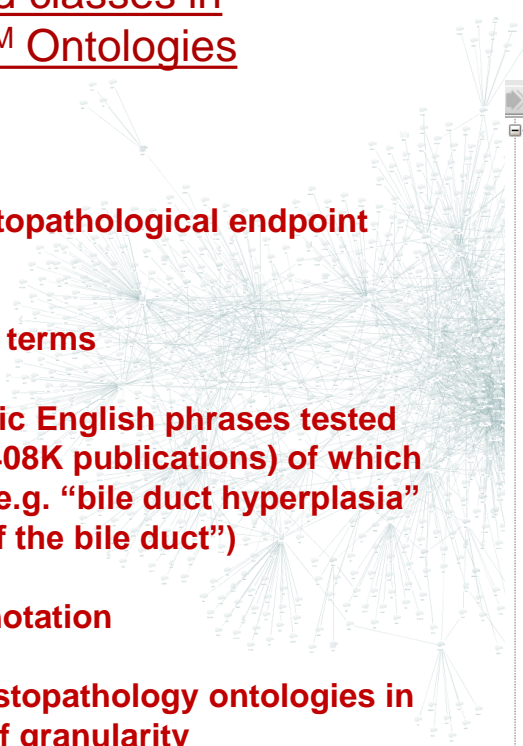


## Connections and classes in CCNet ToxWiz™ Ontologies



### An integrated set

- **92 high-level histopathological endpoint descriptors**
- **912 tissue/organ terms**
- **500000 systematic English phrases tested in the literature (408K publications) of which 2786 are in use (e.g. “bile duct hyperplasia” = “hyperplasia of the bile duct”)**
- **Basis set for annotation**
- **Thousands of histopathology ontologies in different levels of granularity**



## Developing the ontology



Using ontologies always leads to changes and improvements.

Over two years several changes were necessary according to customer feedback

### 1. Finer definitions of certain pathologies:

- e.g. Carcinoma
  - Adenocarcinoma
  - Squamous cell carcinoma
  - Metastatic carcinoma
  - Etc.

### 2. Some other broad terms to cover pathological observations, eg. primary & secondary as qualifiers

Generally: terms are added in order to capture better what experimentalists are saying & capturing.

... but this does not mean that ontology standard is a moving target. The changes and improvements only increase the granularity and are captured by higher categories





### 3. Changes in order to synchronize the key objectives with other running projects and initiatives

#### **INHAND**

International Harmonization of Nomenclature and Diagnostic Criteria for Lesions in Rats and Mice



eTRIKS:  
ontology synchronization among all EU projects and Initiatives



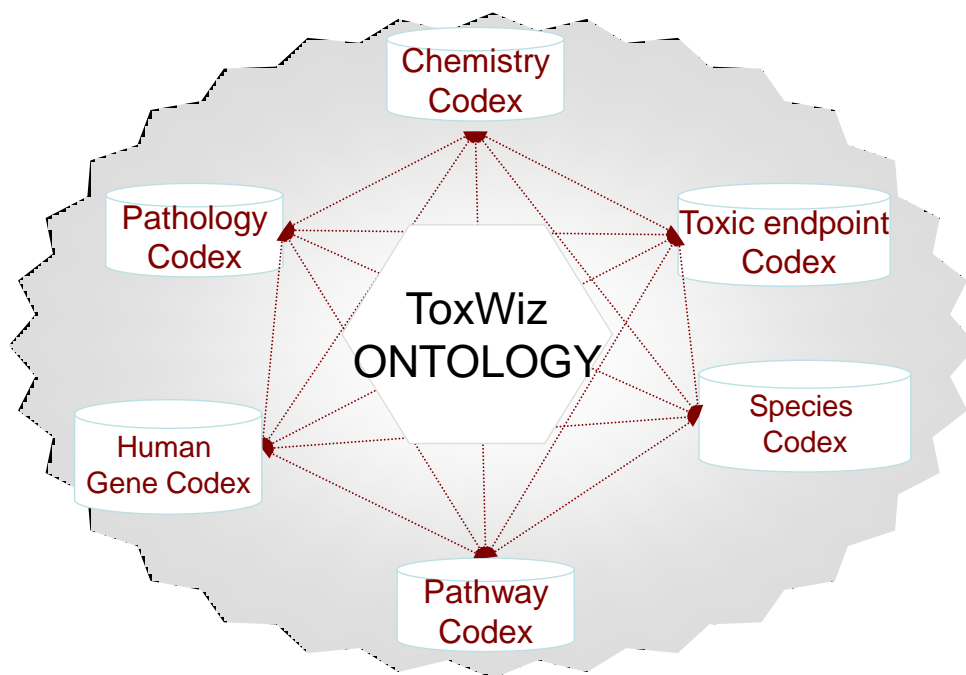
## CCNet Expert Databases: 150 man years of expert curation: ToxWiz CODECES

\* 17 million processed literature references, 4 million catalogued references supporting each database entry, 10 million relationships

| <b>Codex Database</b> | <b>Content</b>   |
|-----------------------|--|
| Chemistry Codex       | 50 000 chemical structures with all reported targets (drugs, industrials, metabolites, toxins) |
| Pathology Codex       | 1000 pathologies linked to 100.000 molecules   |
| Toxic Endpoint Codex  | 1300 toxic-endpoints linked to 100.000 molecules   |
| Pathway Codex         | 500 pathways (signalling, metabolic)   |
| Gene Codex            | 20 000 human targets/proteins/genes  |
| Species Codex         | cross species genes for 17 species   |



## Interoperability achieved via harmonization of ontology



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## Summary



➤ to our knowledge the only one currently developed with capability to support a mechanistic interpretation

### ToxWiz ontologies

- Derived from 150 man years of expert curation of safety information
- Describes histopathological terms with 912 terms
- Classifies further 500 000 terms used by pathologists
- Designed with relevant structure to describe MoA hypothesis and biological pathways
- Modeled on over 8 years of information extraction from safety reports

### Benefits

- Extracting and mapping information about chemical structures related to
  - Toxicity
  - Disease
  - Hypothesis
- Allows exchange of data between user groups
- Deals with the different ways pathologists report
- Aids information classification for QSAR prediction
- Supports –omics interpretations
- Supports in-vitro findings

➤ CCNet base ontology framework as open source, with proprietary specialized add on modules' ?



## A few words on Ontologies

*Ontology is the philosophical discipline which aims to understand how things in the world are divided into categories and how these categories are related together. This is exactly what information scientists aim for in creating structured, automated representations, called 'ontologies,' for managing information in fields such as science, government, industry, and healthcare.*

Applied Ontology: An Introduction, Katherine Munn, Barry Smith (Eds.)

*Often an ontology of the domain is not a goal in itself. Developing an ontology is akin to defining a set of data and their structure for other programs to use.*

Ontology Development 101: A Guide to Creating Your First Ontology,  
Noy & McGuinness, Stanford

Key point: Ontologies only make sense if they are used for something. Gene-ontology (GO) is perhaps the best example in the life sciences. They can seem mysterious until you use them.

