

From physiological maps to disease ontology maps using a systems biology approach

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ONTOX is part of the ASPIS cluster that represents Europe's effort towards the sustainable, animal-free and reliable chemical risk assessment of tomorrow.

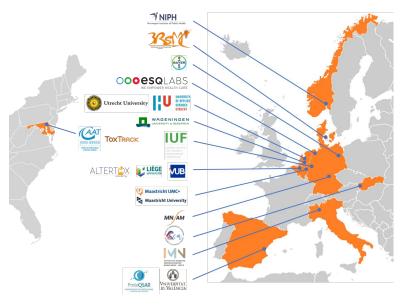


This project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 963845

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The ONTOX project

Ontology-driven and artificial intelligence-based repeated dose toxicity testing of chemicals for next generation risk assessment





Goal: development of an animal-free and human-relevant strategy for the prediction of chemical-induced toxicity

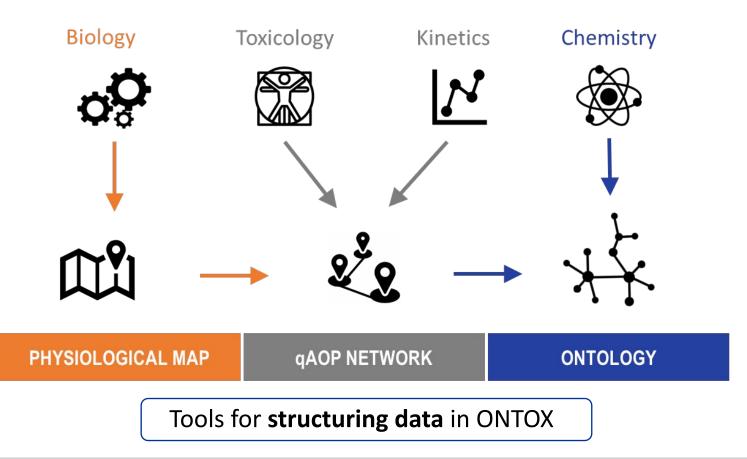
Focus: liver, kidneys and developing brain



Rationale: Rely as much as possible on available data, models and methods

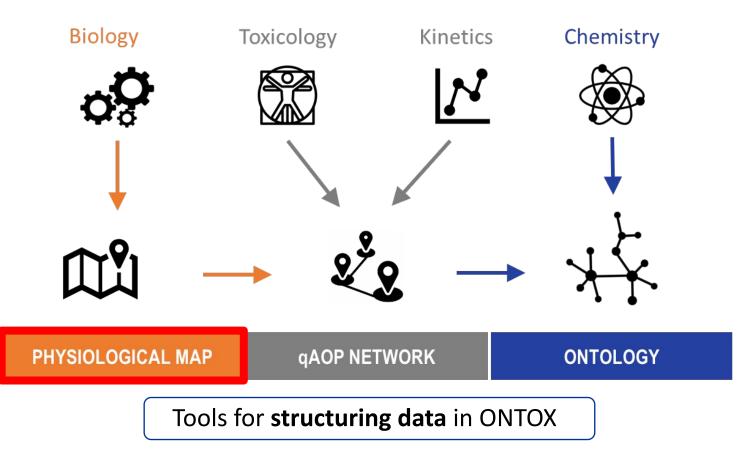
Toolbox: *in vitro* & *in silico* methods





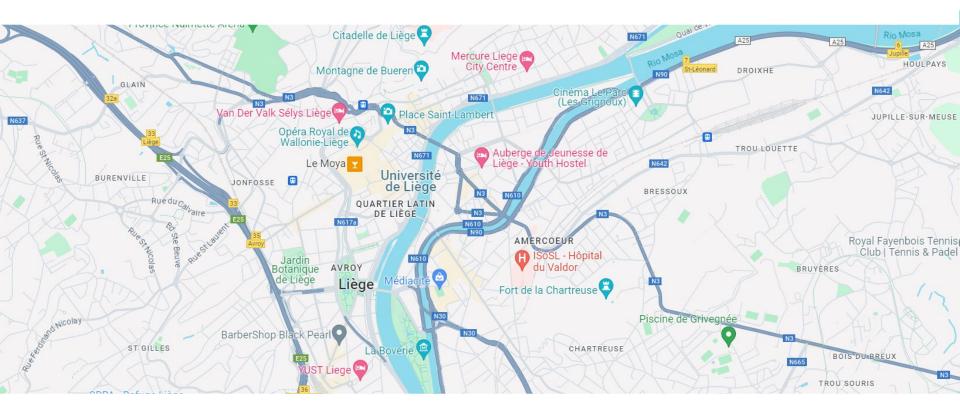
Vinken, et al. 2021 - 10.1016/j.tox.2021.152846

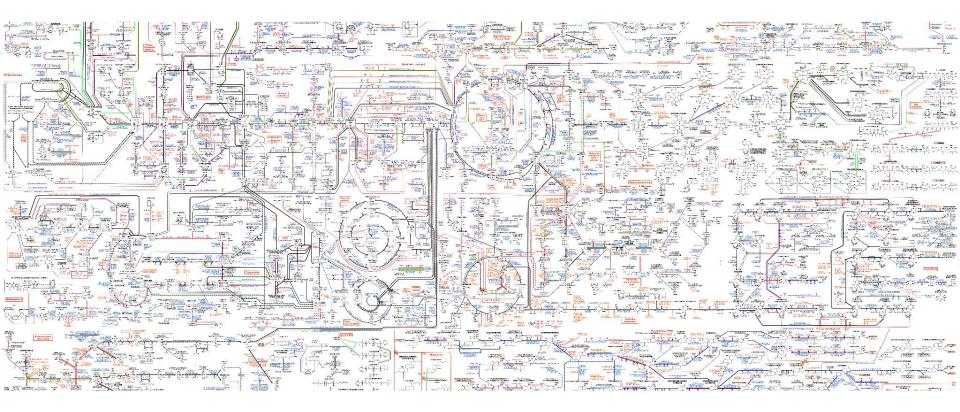




Vinken, et al. 2021 - 10.1016/j.tox.2021.152846

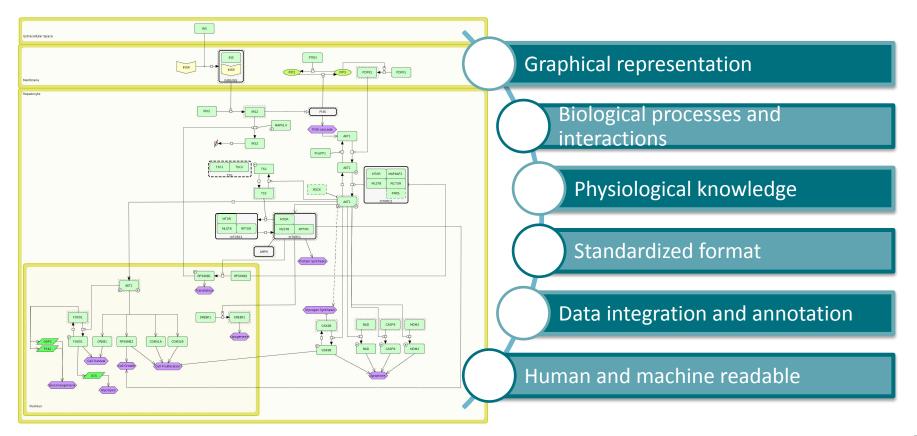






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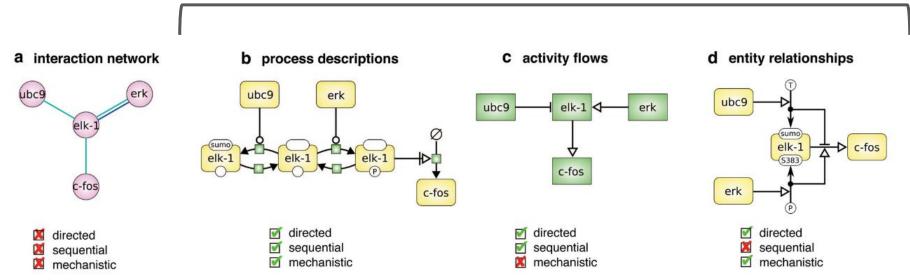
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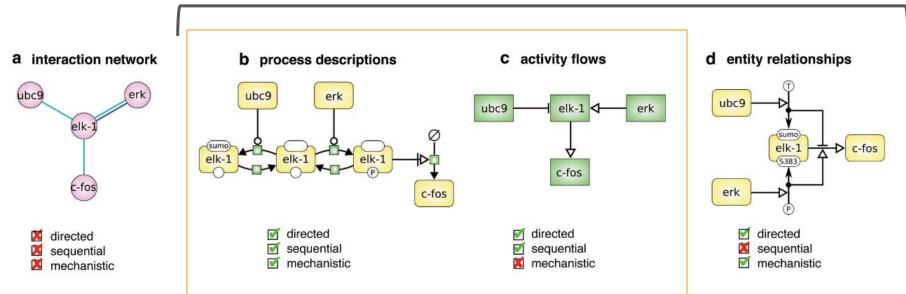






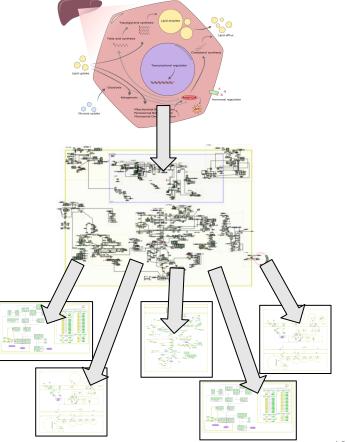






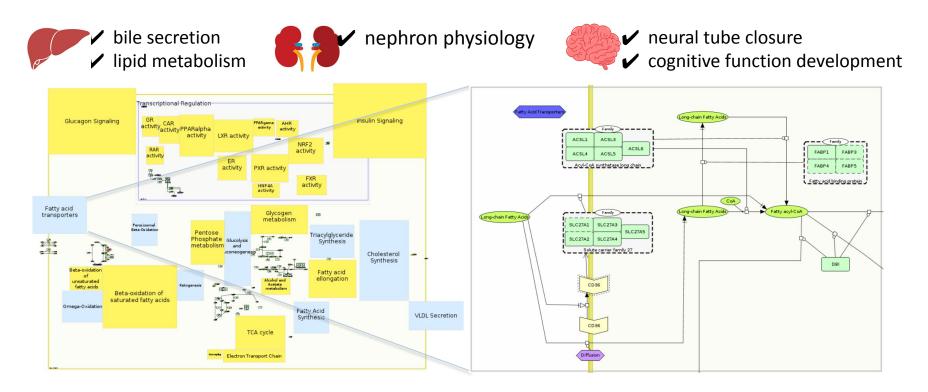
- Conceptual model: used to navigate on the main map and submaps.
- Main map: different PMs presents different requirements for the main map (ex: Brain map has a cell-cell interaction using Activity Flow SBGN and graphical representation; Liver maps have submaps integrated in a big Process Description SBGN map).
- Submaps: they include detailed representation of undisturbed pathways and AOPs.





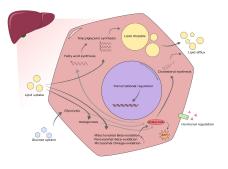
Our current maps

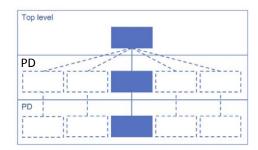






Liver maps





Liver Lipid Metabolism Physiological Map

Pathway	Location on the map
Cholesterol Biosynthesis	Submap and main map
Pathway	
Fatty Acid and Cholesterol	Submap and main map
Transporters	
Glucose metabolism	Main map
(hepatocytes)	
Fatty acid omega-oxidation	Main map
Peroxisomal beta-oxidation	Main map
Triacylglyceride Synthesis	Main map
Mitochondrial Metabolism	Submap
Pathways	
Gene regulatory network	Main map
Glucagon signaling	Submap and main map
Insulin signaling	Submap and main map

Main curator and domain expert



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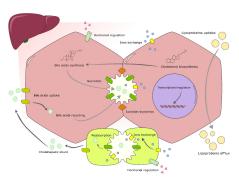


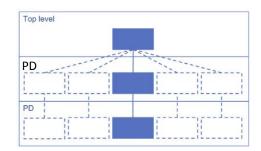
Julen Sanz Serrano

Vrije Universiteit Brussel (VUB) - Belgium



Liver maps





Liver Bile Secretion Physiological Map

Pathway	Location on the map	
Cholesterol Biosynthesis Pathway	Submap and main map	
Fatty Acid and Cholesterol Transporters	Submap and main map	
Mitochondrial Metabolism Pathways	Submap	
Apoptosis Pathway	Submap	
Autophagy Pathway	Submap	
Bile acids biosynthesis	Main Map	
Canaliculi dynamics pathways	Main Map	
Bile salts circulation	Main Map	
Bile salts uptake	Main Map	
Cholehepatic shunt	Main Map	
Glucagon signaling	Submap and main map	
Insulin signaling	Submap and main map	
Gene regulatory network	Main Map	

Main curator and domain expert



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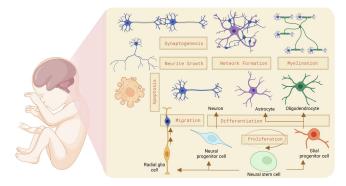


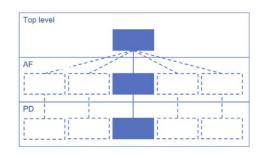
Julen Sanz Serrano

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Brain map





Brain Development Physiological Map

Pathway	Location on the map	
Cell-cell interaction map	Main map	
Neuron development	Submap	
Oligodendrocyte development	Submap	
Astrocyte development	Submap	
Radial glia development	Submap	

Main curator and domain expert



Luiz Ladeira

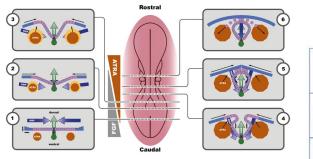
University of Liège - Belgium



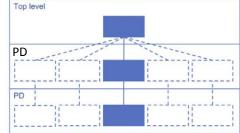
Eliška Kuchovská

IUF – Leibniz-Institut für umweltmedizinische Forschung GmbH - Germany

NTC map







Neural Tube Closure Physiological Map

Pathway	Location on the map
BMP signaling pathway	Submap
SHH signaling pathway	Submap
FGF signaling pathway	Submap
Wnt signaling pathway	Submap
Gene regulatory network	Submap
ATRA metabolism	Submap
Folate metabolism	Submap

Main curator and domain expert



Alessio Gamba

University of Liège - Belgium

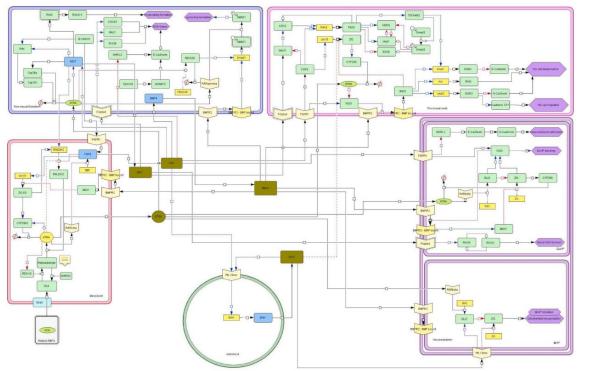


Job Berkhout

Utrecht University (UU) - The Netherlands The Centre for Health Protection of the Dutch National Institute for Public Health and the Environment (RIVM)



NTC map



Utrec

Harm Heusinkveld

Utrecht University (UU) - The Netherlands The Centre for Health Protection of the Dutch National Institute for Public Health and the Environment (RIVM)



An ontology for developmental processes and toxicities of neural tube closure

Check for spclates

tube closure Harm J. Heusinkveld^{***}, Yvonne C.M. Staal^{**}, Nancy C. Baker^b, George Daston^{*},

Harm J. Heusinkveld^{a, s}, Yvonne C.M. Staal^a, Nancy C. Baker^b, George Da Thomas B. Knudsen^d, Aldert Piersma^a

- ⁸ Centre for Health Protection, National Institute for Public Health and the Environment (RIVM), Bilthoven, the Netherlands
- ⁶ Leidos, Research Triangle Park NC 27711, USA
 ⁶ Global Product Stewardship, The Proceer & Gamble Company, Cincinnati, OH USA
- ^d Center for Computational Toxicology and Exposure, U.S. Environmental Protection Agency, Research Triangle Park NC 27711, USA

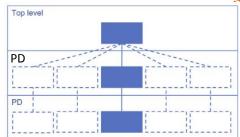
https://ontox.elixir-luxembourg.org/minerva/index.xhtml?id=Neural_Tube_Closure_PM_v1



Kidney map

Nephron Physiological Map

Pathway	Location on the map	
Renin-Angiotensin pathway	Submap and main map	
Mitochondrial metabolism pathways	Submap	
Transporters dynamics pathways	Main map	
Endocytosis pathway	Submap	
Post translational modifications	Submap and main map	
Thyroid hormones signaling pathway	Submap and main map	
TLR signaling pathway	Submap and main map	
HTR2C signaling pathway	Submap and main map	
Apoptosis	Submap	
NLRP3 inflammasome	Submap	



Main curator and domain expert



Alessio Gamba

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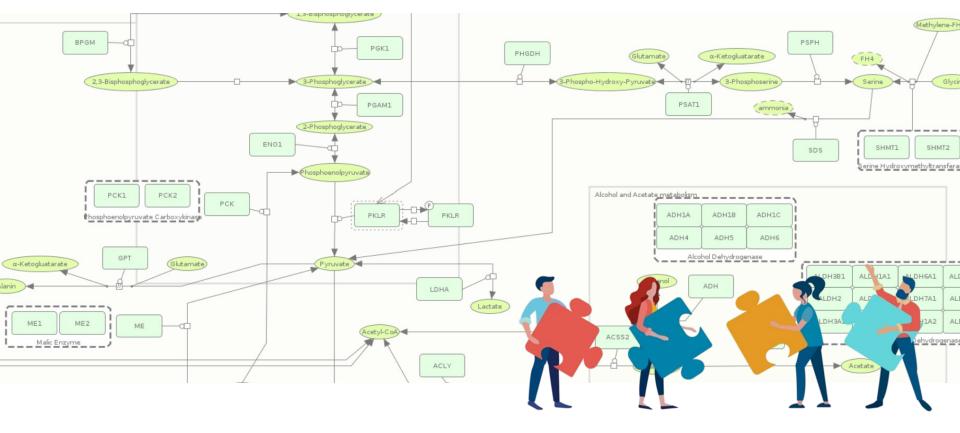


Devon Barnes

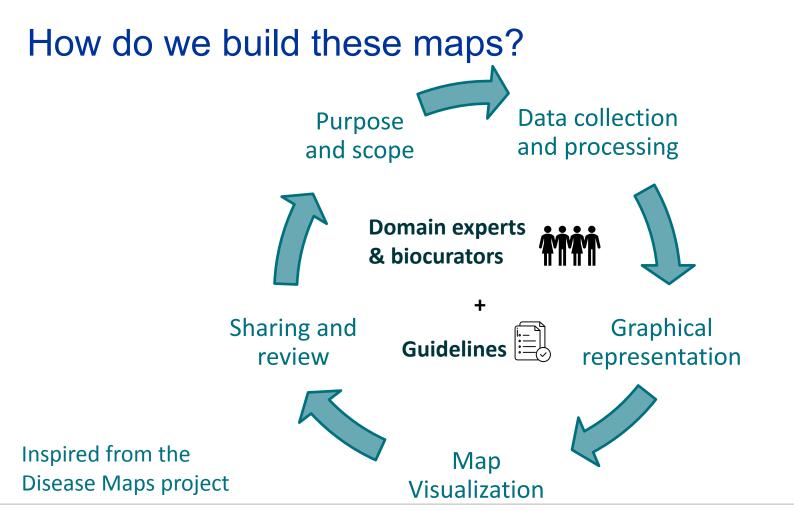
Utrecht University (UU) - The Netherlands



Assembling the puzzle

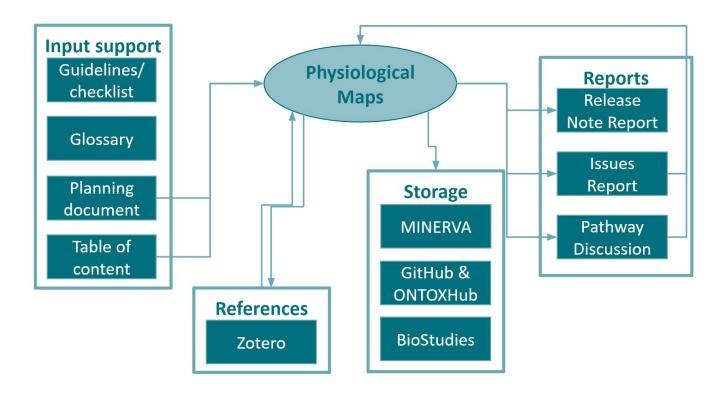








Documentation structure





Planning phase

Modular structure:

- Use of submaps (= smaller modules)
- Improve data visualization
- Simplify review and curation processes
- ✓ Modules can be submitted to WikiPathways: accelerate curation

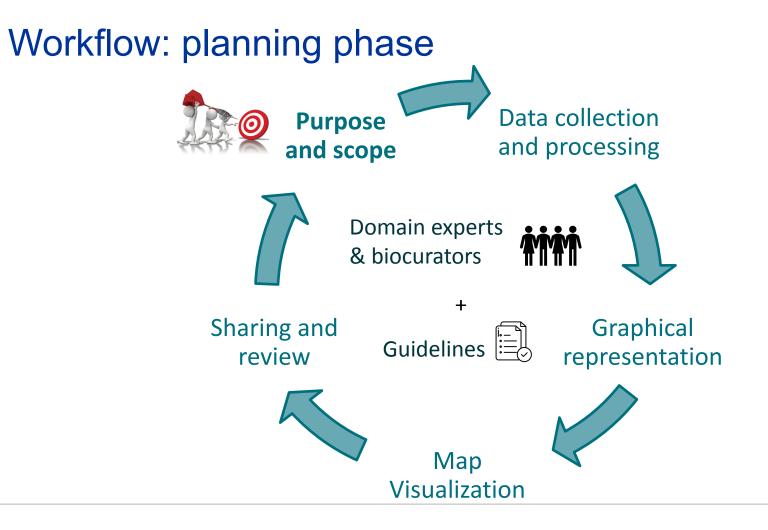
• Annotation:

Manual & automated annotation (guidelines, MINERVA)

• Documentation:

- ✔ Curation guidelines
- Quality control checklist
- Storage:
 - MINERVA, GitHub, BioStudies





Purpose and scope



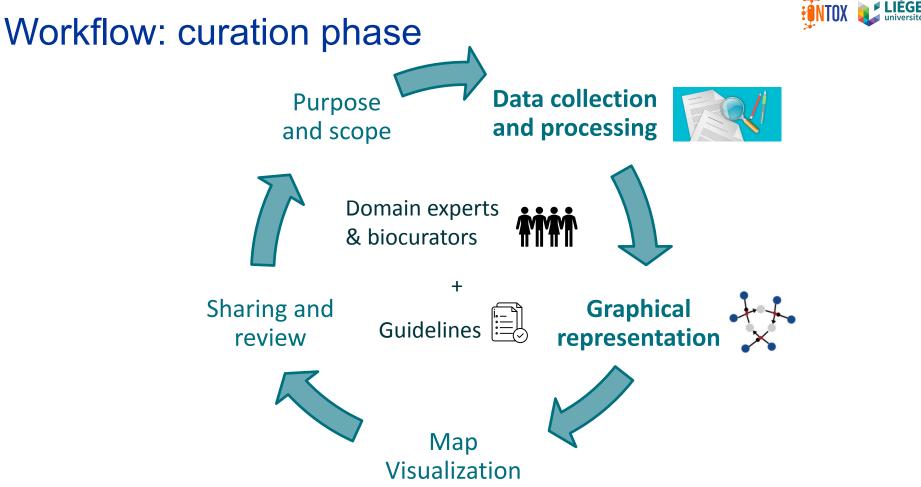
Plan for the map:

• Who? List of curators and domain experts

- Why? Intended use of the map
- What?
 - Map content (pathways, molecules)
 - ✔ First references
- How?
 - ✔ Level of granularity
 - ✔ Sustainability/storage

Table of content:

Pathway	Diagrams	Diagrams	
	Resource	Access	
Cholesterol Biosynthesis Pathway	Reactome, WP, and literature	<u>10.3180/R-HSA-1</u> <u>91273.7</u> , <u>WP4718</u>	Pathway revised. Stable version. Comments are in the <u>Discussion</u> <u>file</u> .
Fatty Acid and Cholesterol Transporters	Wikipathways Wikipathways KEGG	<u>WP5061</u> <u>WP5304</u> <u>map04979</u>	Pathway revised. Stable version. Comments are in the <u>Discussion</u> <u>file</u> . Lipoprotein uptake and secretion are included.
Mitochondrial Metabolism Pathways	PDMap, COVID19Map, and literature	PDMap COVID19Map	Pathway revised. Stable version. Comments are in the <u>Discussion</u> <u>file</u> . Oxidative stress and fatty acid biosynthesis are included.





Curation phase

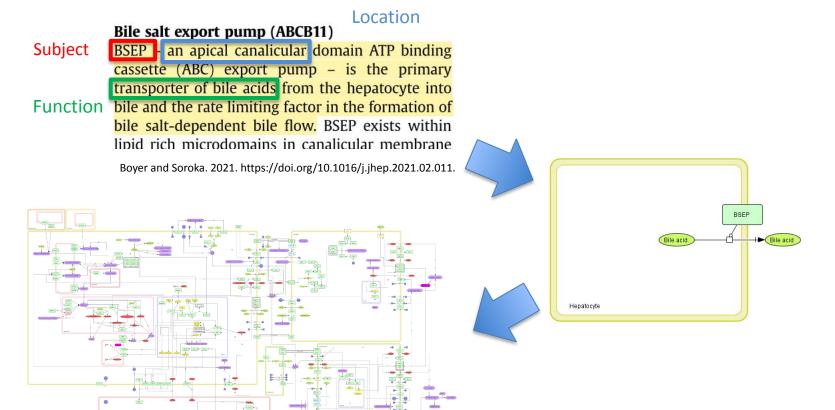
- Data resources:
 - 🖌 🛛 Books, research papers, review articles, databases 🤃 💞 🜾 reactome
 - Other maps (e.g. from Disease Maps project)
 - Tools for systematic review sysrev

Graphical representation:

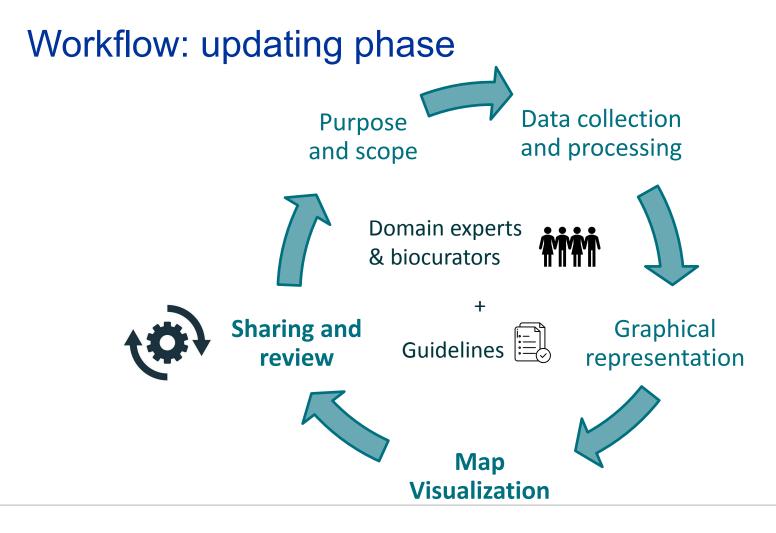
- ✔ Graphical standard Systems Biology Graphical Notation (SBGN):
 - Visual language for biochemical interaction networks
 - Standardized representation
- ✔ Diagram editor: CellDesigner
 - SBGN-compatible networks
 - Data format: Systems Biology Markup Language (SBML)
 - Free and open
 - Manual annotation



Curating literature



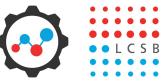




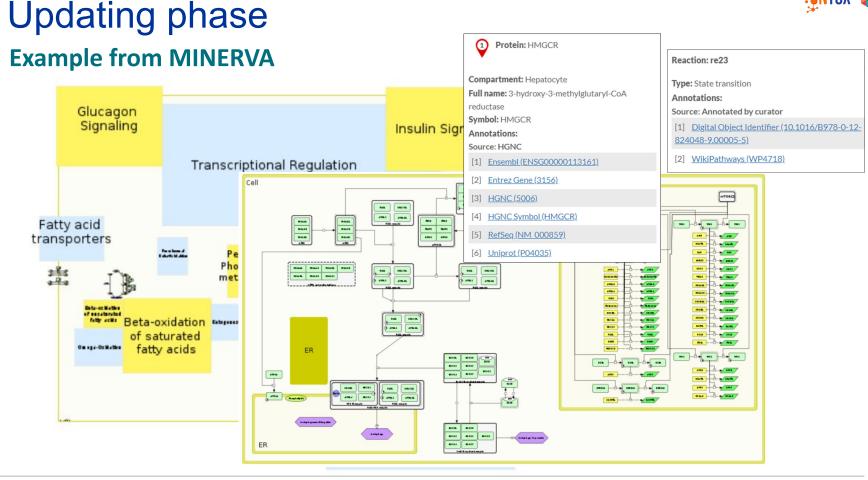
Updating phase

- Visualization and annotation tool MINERVA platform:
 - Molecular INteraction NEtwoRks VisuAlization
 - ✔ User-friendly interface
 - Content exploration (e.g. searching for drug targets), overlay of experimental datasets
 - Automated annotation using identifiers (e.g. ChEBI, Ensembl, Uniprot, Gene Ontology)
 - ✔ Conversion in various formats (SBML, GPML, etc.)







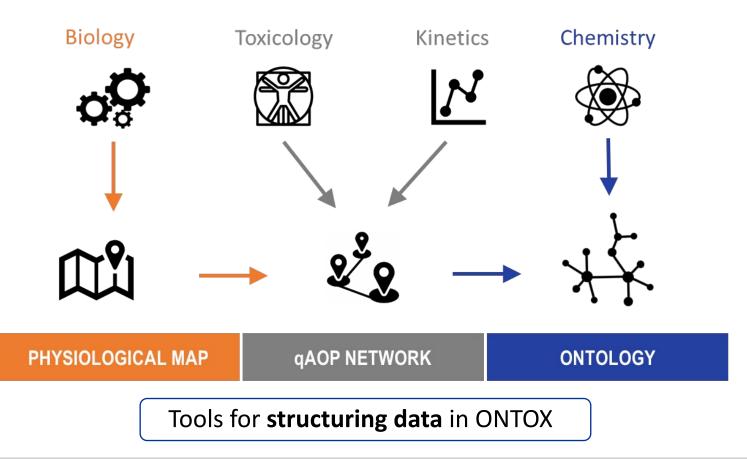


Updating phase

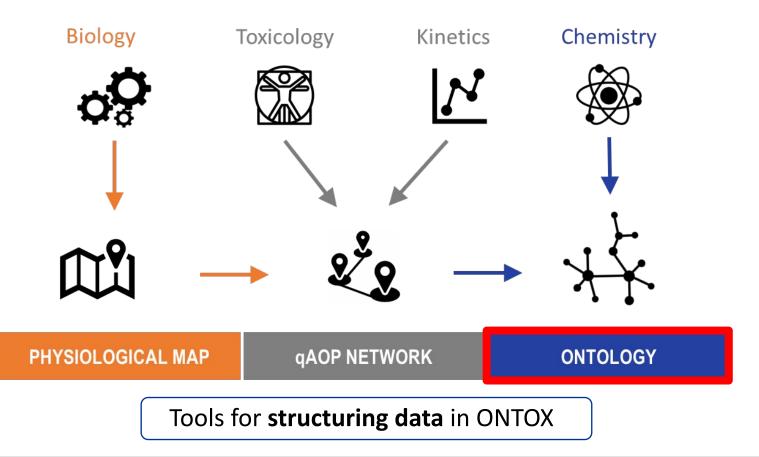
- Sharing and review:
 - Maps need a continuous update
 - Commenting facilitated by the MINERVA Platform
 - Storage on GitHub and BioStudies
 - Collaboration between domain experts and curators
 - Create a **bridge between systems biology and toxicology communities**







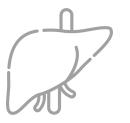




Vinken, et al. 2021 - 10.1016/j.tox.2021.152846

Ontology case-studies





Lipid metabolismBile secretion



Nephron

- Steatosis
- Cholestasis

- Tubular necrosis
- Cholestasis

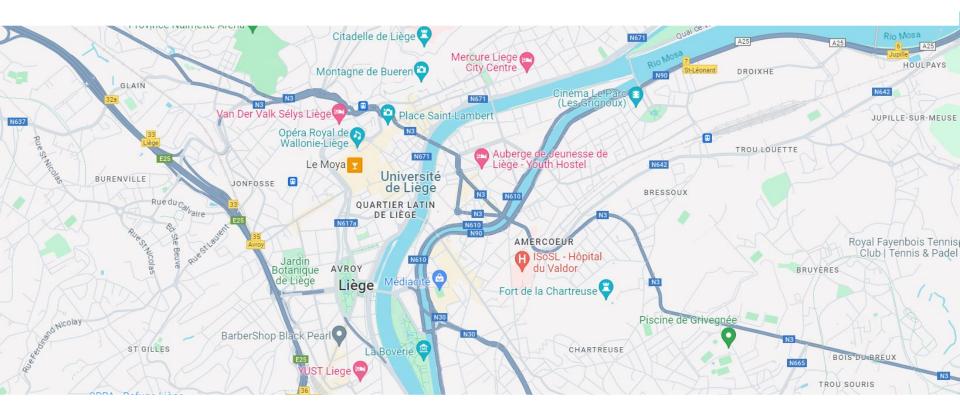


Neural tube closureBrain development

- Neural tube closure defects
- Cognitive function defects



The concept of ontology maps



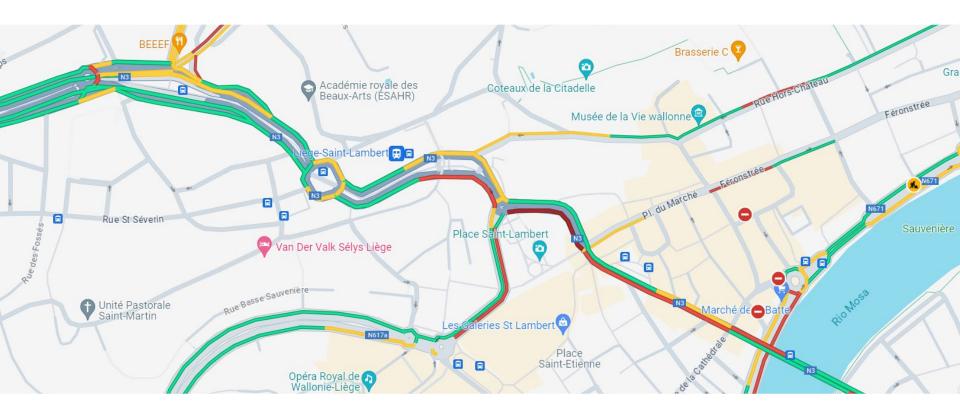


The concept of ontology maps

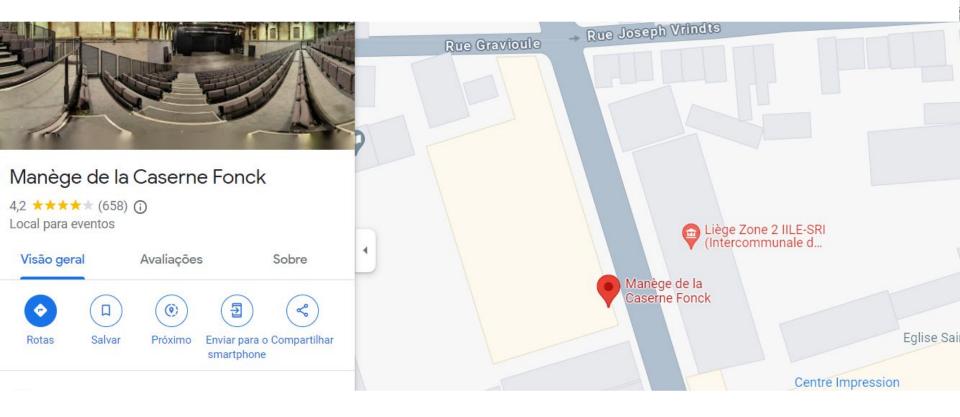


The concept of ontology maps



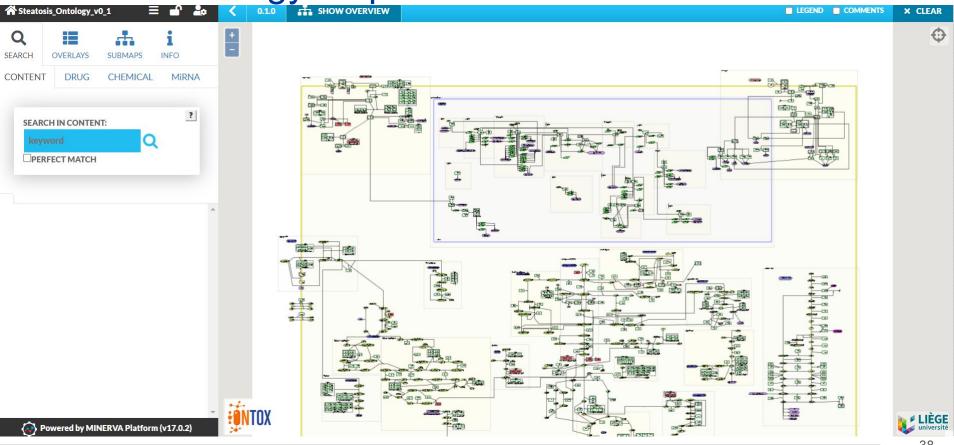


The concept of ontology maps

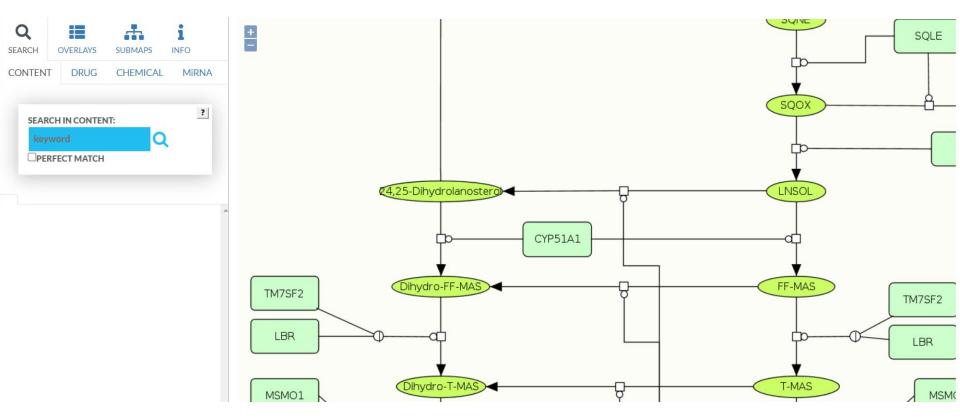


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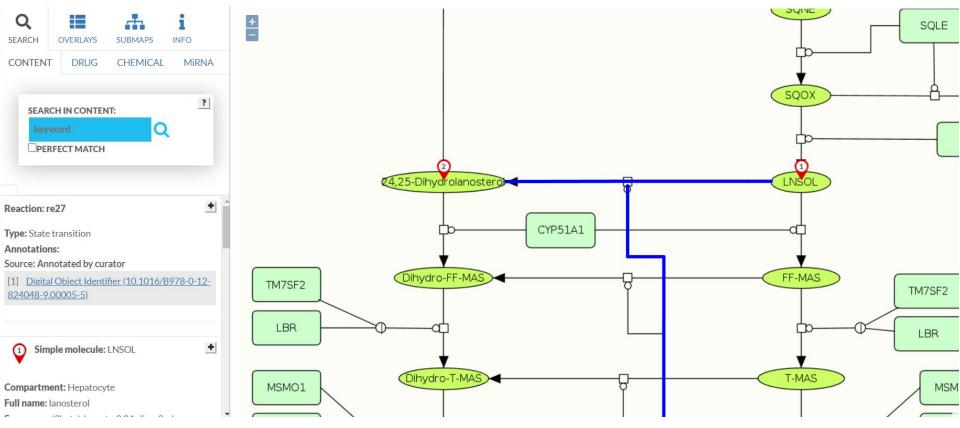




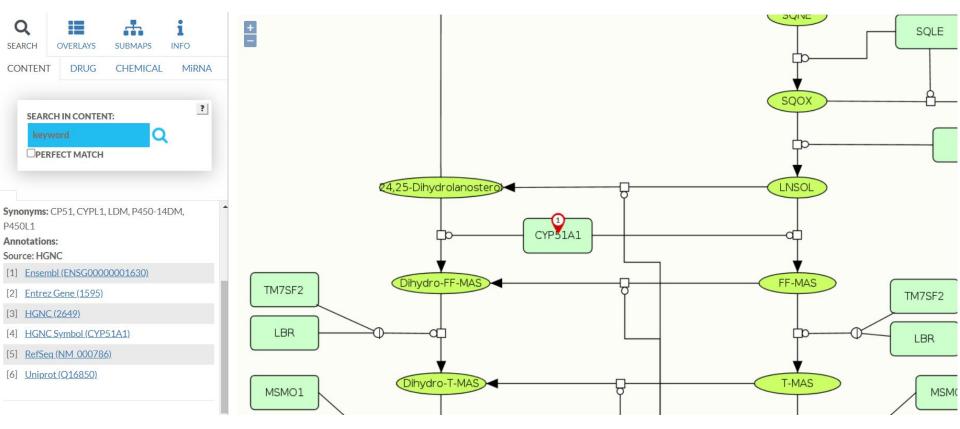














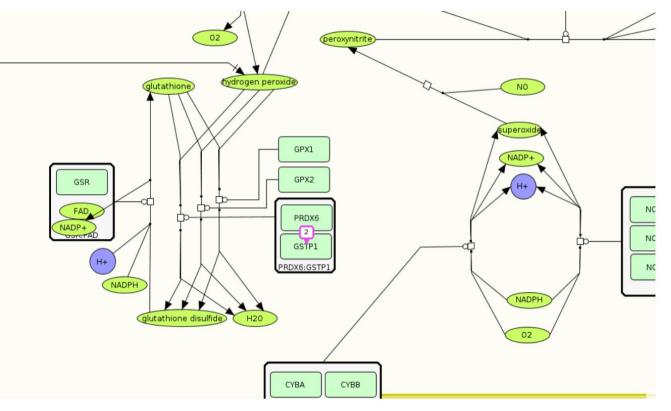
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Q		-	i
SEARCH	OVERLAYS	SUBMAPS	INFO
CONTEN	T DRUG	CHEMICAL	MiRNA
SEARCH FOR TARGETS OF:			?
paracetamol			

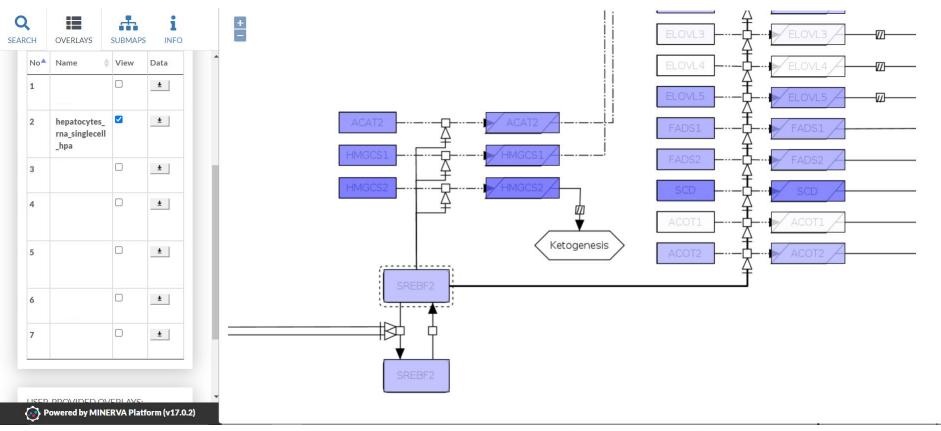
paracetamol

Drug: Acetaminophen

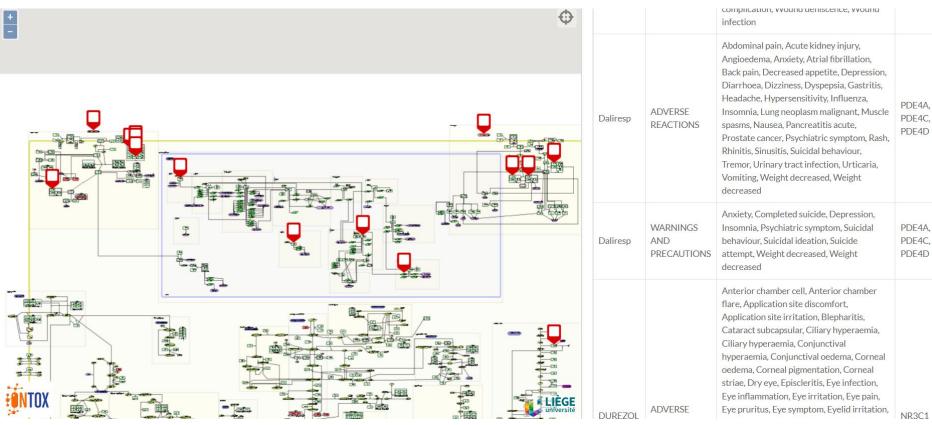
Description: Acetaminophen (paracetamol), also commonly known as _Tylenol_, is the most commonly taken analgesic worldwide and is recommended as first-line therapy in pain conditions by the World Health Organization (WHO) [A176318]. It is also used for its antipyretic effects, helping to reduce fever [F4124]. This drug was initially approved by the U.S. FDA in 1951 [FDA label] and is available in a variety of forms including syrup form, regular tablets, effervescent tablets, injection, suppository, and other forms [L5756, L5774, F4124], [FDA label]. Acetaminophen is often found combined with other drugs in more than 600 over the counter (OTC) allergy medications, cold medications, sleep



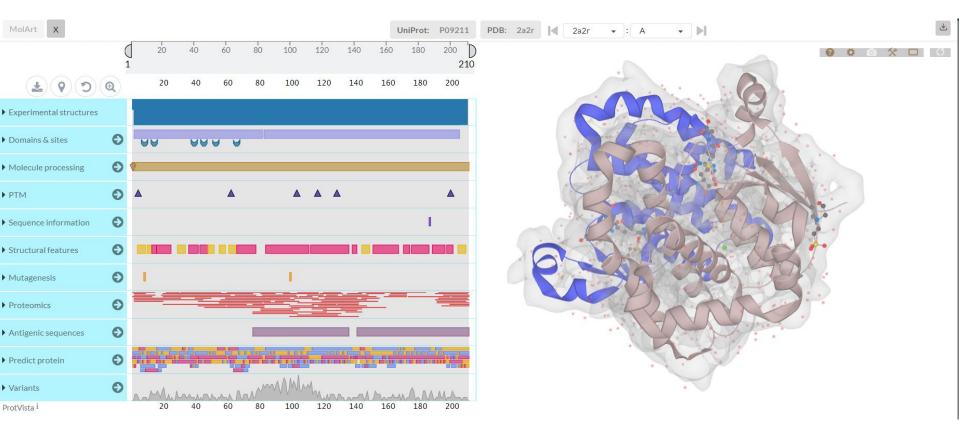




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Multi-layered ontology maps



Integration with adverse outcome pathways (AOPs)

- ✔ Use PMs as benchmarks to **fill gaps** in AOPs
- ✔ AOP networks designed using SBGN representation
 - Data integration between physiological maps and AOPs
 - □ Standardization of AOPs and interoperability
- Integration with other data (chemical, kinetic, omics, etc.)
 - Plug-ins displaying tables with annotated information
 - Overlay of data (e.g. omics, drug databases)

Conclusions

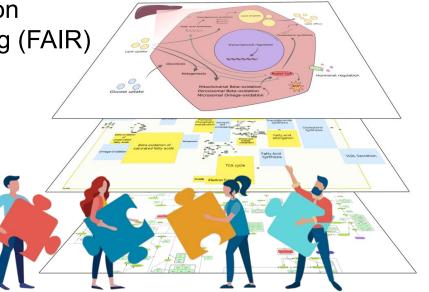
From physiological maps...

- Graphical representation of biological processes
- Standardized notation
- Guidelines for data curation and annotation
- Tool for knowledge integration and sharing (FAIR)

... to ontology maps:

- Founded on PMs
- AOPs, chemical, kinetic, omics data
- Mechanistic risk assessment

Large-scale community effort





Resources

systems medicine disease maps

Disease maps for specific diseases



Liver steatosis C

Cholestasis ★☆☆☆☆ Resource: Liver Lipid Metabolism Physiological Map Contact: Luiz Ladeira, University of Liège, Liège, Belgium

Resource: Liver Bile Secretion Physiological Map Contact: Luiz Ladeira, University of Liège, Liège, Belgium

Cross-disease projects

Neural tube closure defects ★★★★☆ Resource: Neural Tube Closure Physiological Map Contact: Alessio Gamba, University of Liège, Liège, Belgium Publications: PubMed

Heusinkveld, et al. 2021. Doi: 10.1016/j.reprotox.2020.09.002

Kidney crystallopathy and tubular necrosis ★☆☆☆☆ Resource: Nephron Physiological Map Diseases: kidney crystallopathy, tubular necrosis Contact: Alessio Gamba, University of Liège, Liège, Belgium

Cognitive function defects ★☆☆☆☆ Resource: Brain Development Physiological Map Contact: Luiz Ladeira, University of Liège, Liège, Belgium

Resources





Acknowledgments



Prof. Liesbet Geris Dr. Bernard Staumont Dr. Alessio Gamba













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IUF LEIBNIZ-INSTITUT FÜR UMWELT-MEDIZINISCHE FORSCHUNG

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Prof. Roos Masereeuw **Devon Barnes** Manoe Janssen



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Marc Teunis

Job Berkhout



Prof. Thomas Hartung Tom Luechtefeld

Hubert Dirven Tim Hofer Malene Lislien Oddvar Myhre

Agata Impellizzeri Graciela López Soop

ALTERT SX wards alternatives to animal testing

François Busquet Mario Cecilia Matteo Piumatti

Non-ONTOX partners



Marek Ostaszewski Ahmed Hemedan Alexander Mazein



Acknowledgments

The ONTOX team



Thank you for your attention



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