

Tutorial:

MINERVA platform for map reviewers

WP1



Funded by the Horizon 2020
Framework Programme of the
European Union



February 2024



Tutorial:

MINERVA platform for map reviewers

University of Liège ONTOX Team



Funded by the Horizon 2020
Framework Programme of the
European Union

Grant Agreement ID: 963845

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Here you will find relevant links:

The ONTOX MINERVA session is maintained by Elixir Luxembourg (<https://minerva.pages.uni.lu/doc/>).

General link for ONTOX's MINERVA: <https://ontox.elixir-luxembourg.org/minerva/>

Physiological maps link:

- Maps and ontologies can be found linked on the menu on our MINERVA instance:
<https://ontox.elixir-luxembourg.org/minerva/>

For previous versions, please refer to the GitHub repository for each map or ontology: <https://github.com/ontox-maps>

If you need credentials for access, please contact us.



SEARCH



OVERLAYS



INFO



CONTENT

DRUG

CHEMICAL

MiRNA

When accessing the ONTOX MINERVA link, you will land on a page like this one. At the time of this tutorial, the menu doesn't provide any link to the maps because this is a public version of our MINERVA platform. Links will appear as soon as we have the map's papers published.

Physiological Maps



Liver Lipid Metabolism
(coming soon)



Liver Bile Acid Secretion
(coming soon)



Neural Tube Closure
(coming soon)



Brain Development
(coming soon)



Nephron Physiology
(coming soon)



Adverse outcome pathways (coming soon)



Liver steatosis



Cholestasis



Neural Tube Closure Defects



Cognitive Function Defects



Kidney Tubular Necrosis



Kidney Crystallopathy



This project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 962640



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Once the maps are publicly available, an orange circle will indicate the map is available. Click on the figure of your choice to visualize the link on the white space just below this text box.

Physiological Maps



Liver Lipid Metabolism



Liver Bile Acid Secretion



Neural Tube Closure
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metamap_v5

SEARCHOVERLAYSIINFO


CONTENTDRUGCHEMICALMiRNA


SEARCH IN CONTENT:
keyword
☐ PERFECT MATCH


1 Phenotype: Liver Lipid Metabolism
Physiological Map


Pathway: Background overview
Liver Lipid Metabolism PM [click here]
Annotations: No annotations


Physiological Maps


 Liver Lipid Metabolism

 Liver Bile Acid Secretion


 Neural Tube Closure
(coming soon)


 Brain Development
(coming soon)


 Nephron Physiology
(coming soon)


 **Physiological Maps**


Adverse outcome pathways
(coming soon)


 Liver steatosis

 Cholestasis


 Neural Tube Closure Defects

 Cognitive Function Defects

 Kidney Tubular Necrosis

 Kidney Crystallopathy


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LEGENDCOMMENTS

CLEAR

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metamap_v5

LEGENDCOMMENTS

CLEAR

SEARCHOVERLAYSINFO

CONTENTDRUGCHEMICALMiRNA

1 Phenotype: Liver Lipid Metabolism Physiological Map

Pathway: Background overview
Liver Lipid Metabolism PM [click here]
Annotations: No annotations

Physiological Maps

1

Liver Lipid Metabolism

Liver Bile Acid Secretion

Neural Tube Closure
(coming soon)

Brain Development
(coming soon)

Nephron Physiology
(coming soon)

ONTOX Physiological Maps

Adverse outcome pathways
(coming soon)

Liver steatosis

Cholestasis

Neural Tube Closure Defects

Cognitive Function Defects

Kidney Tubular Necrosis

Kidney Crystallopathy

ONTOX

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ONTOX

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LIÈGE université

If you are reviewing a map that is not listed here or is still not public, the UL team will send you a link to access it and credentials to log in.

If you arrive on this page, please use the credentials that have been sent to you. If you did not receive these credentials, please contact us.

After entering your login and password, you will be redirected to the next page.

If you use direct links to the maps, you can jump to slide 6.

AUTHORIZATION FORM

LOGIN:

PASSWORD:

LOGIN

RESET PASSWORD

> BACK TO MAP

[+ ADD PROJECT](#) [REFRESH](#)

Show entries

ProjectId	Created	Created by	Name	Status	Edit	Remove
empty	2014-03-27	admin		Ok		
Lipid_Metabolism_PM_20220310	2022-03-10	admin	Lipid_Metabolism_PM	Ok		
Liver_Bile_Production_and_Secretion_20220203	2022-02-21	admin	Liver_Bile_Secretion_PM	Ok		
physmap_108	2022-03-02	admin	Physmap	Ok		
physmap_92	2022-02-23	admin	Physmap	Ok		
physmap_97	2022-02-25	admin	Physmap	Ok		
physmap_99	2022-02-25	admin	Physmap	Ok		
vitD	2022-03-01	admin	VitaminD_PM	Ok		

Showing 1 to 8 of 8 entries

Previous **1** Next

[+ ADD PROJECT](#) [REFRESH](#)

After logging in, this is the page you will see. Here you will be able to choose between the available maps to open.

Don't worry! Only the maps you were asked to review will be available in this list.

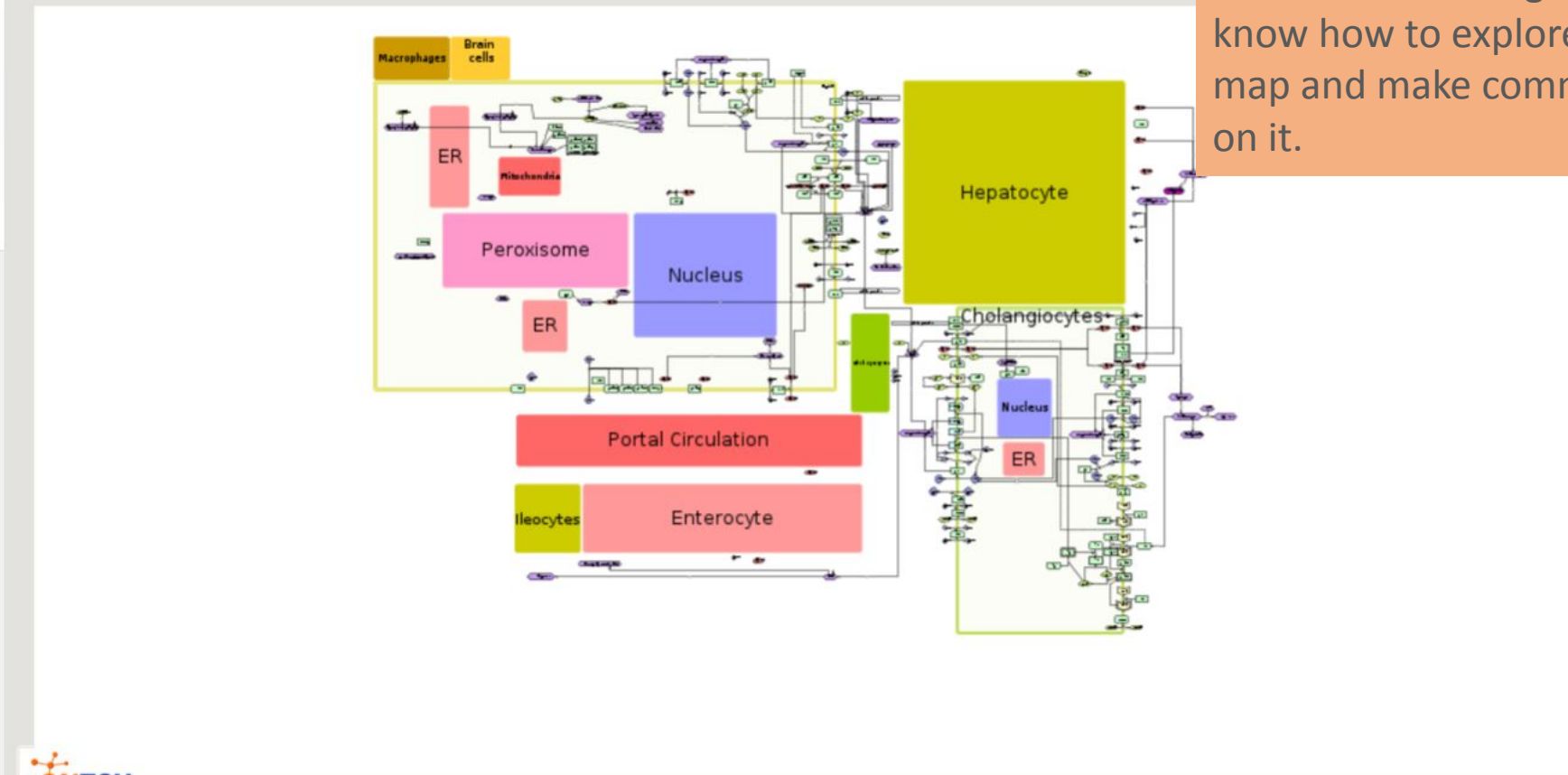


SEARCH IN CONTENT:

keyword

☐ PERFECT MATCH

+
 -



And here we are, with the map to be reviewed.

Let's take a look at the MINERVA platform environment and get to know how to explore the map and make comments on it.



SEARCH



OVERLAYS



INFO



CONTENT

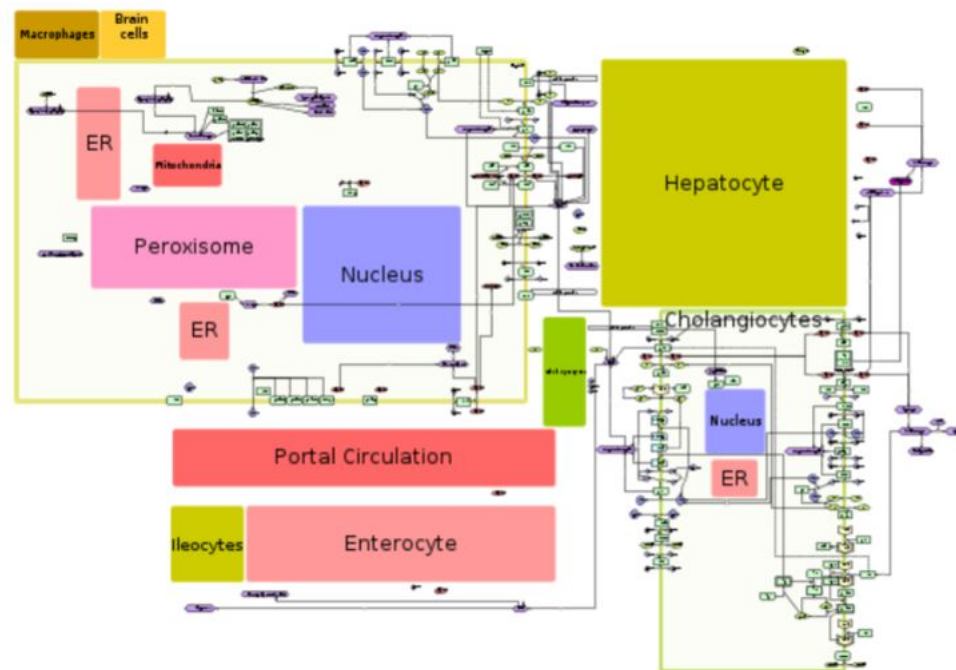
DRUG

CHEMICAL

MiRNA

First, let's make it easier to look at the map.

For that, click on "Overlays".



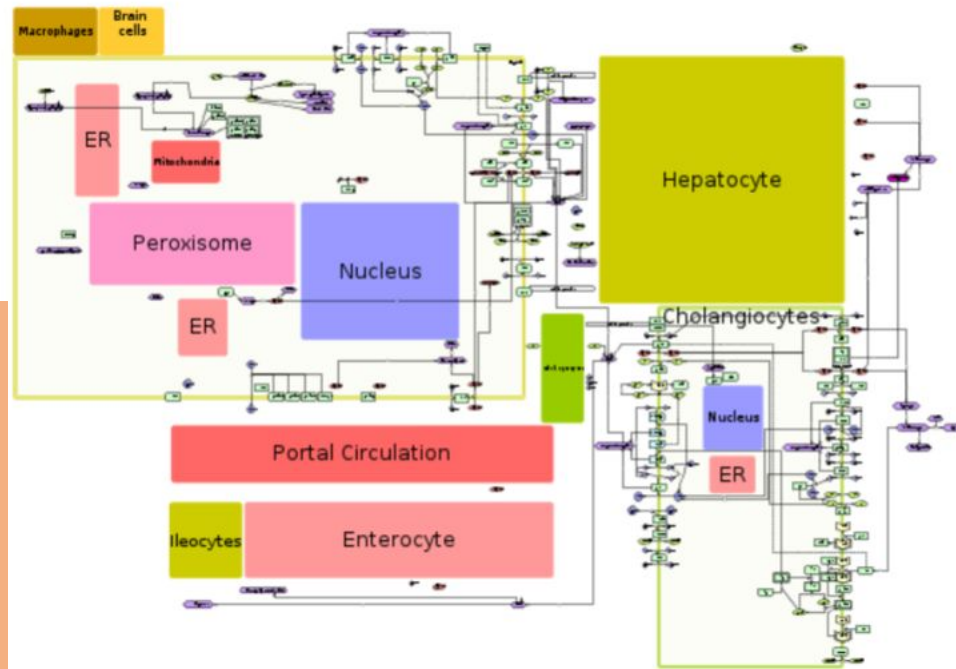
BACKGROUNDS:

Name	View
Pathways and compartments	🔍
Network	🔍
Empty	🔍



Now, to visualize the whole map without having the compartment masks, you can choose the “Network” mode of visualization by clicking on the magnifier icon.

It is not mandatory to use this resource, but it will make the viewing cleaner.



Home

Liver Lipid Metabolism

Menu

Lock

User

February_2024

SHOW OVERVIEW

LEGEND

COMMENTS

CLEAR

Physiological Map

SEARCH

OVERLAYS

SUBMAPS

CONTENT

DRUG

CHEMICAL

SEARCH IN CONTENT:

keyword

☐ PERFECT MATCH

Some maps might contain an overview model that can be accessed on the button highlighted above.

The diagram is a metabolic map of liver lipid metabolism. It features several interconnected pathways represented by colored boxes and lines. Yellow boxes include 'Glucagon Signaling', 'Insulin Signaling', 'Glycogen metabolism', 'Pentose Phosphate metabolism', 'Beta-oxidation of saturated fatty acids', 'Fatty acid elongation', and 'Fatty Acid Synthesis'. Blue boxes include 'Transcriptional Regulation', 'Fatty acid transporters', 'Cholesterol Synthesis', and 'Lipoprotein Secretion'. A central 'TCA cycle' box is also present. Lines connect these boxes, indicating the flow of metabolites and regulatory interactions. For example, 'Glucagon Signaling' and 'Insulin Signaling' both influence 'Transcriptional Regulation'. 'Fatty acid transporters' lead to 'Beta-oxidation of saturated fatty acids', which then feeds into the 'TCA cycle'. 'Fatty acid elongation' and 'Fatty Acid Synthesis' are part of the 'Fatty acid' pathway, which also leads to 'Lipoprotein Secretion'. 'Cholesterol Synthesis' is a separate pathway that also contributes to 'Lipoprotein Secretion'.

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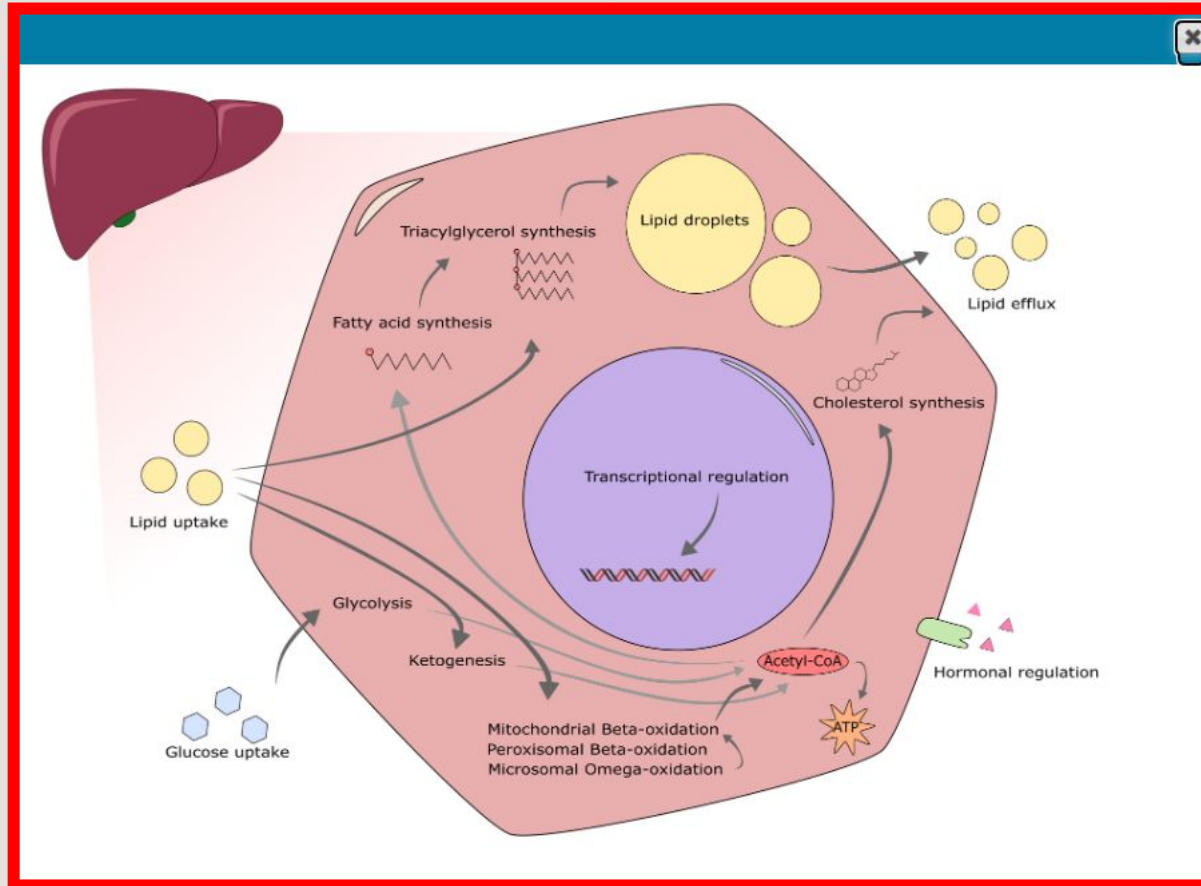
15

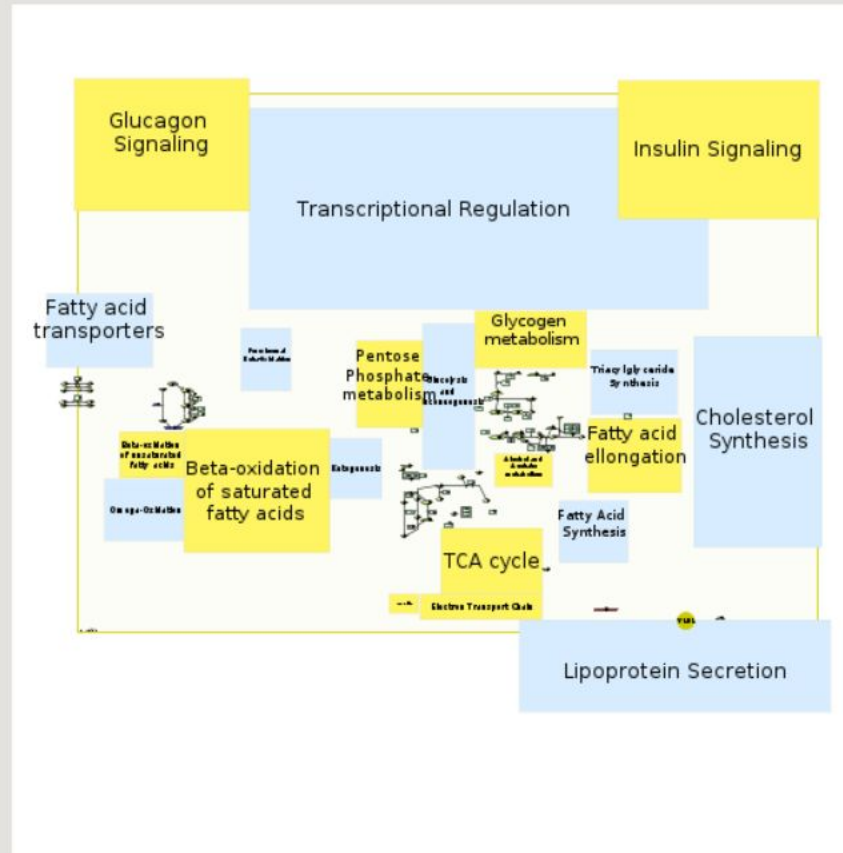
This is especially helpful when navigating big maps. It works as a mini-map, and contain clickable links that will redirect you to places on the molecular interactions maps - main map and submaps.

☐ PERFECT MATCH

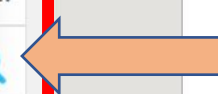


Powered by MINERVA Platform (v16.4.0)



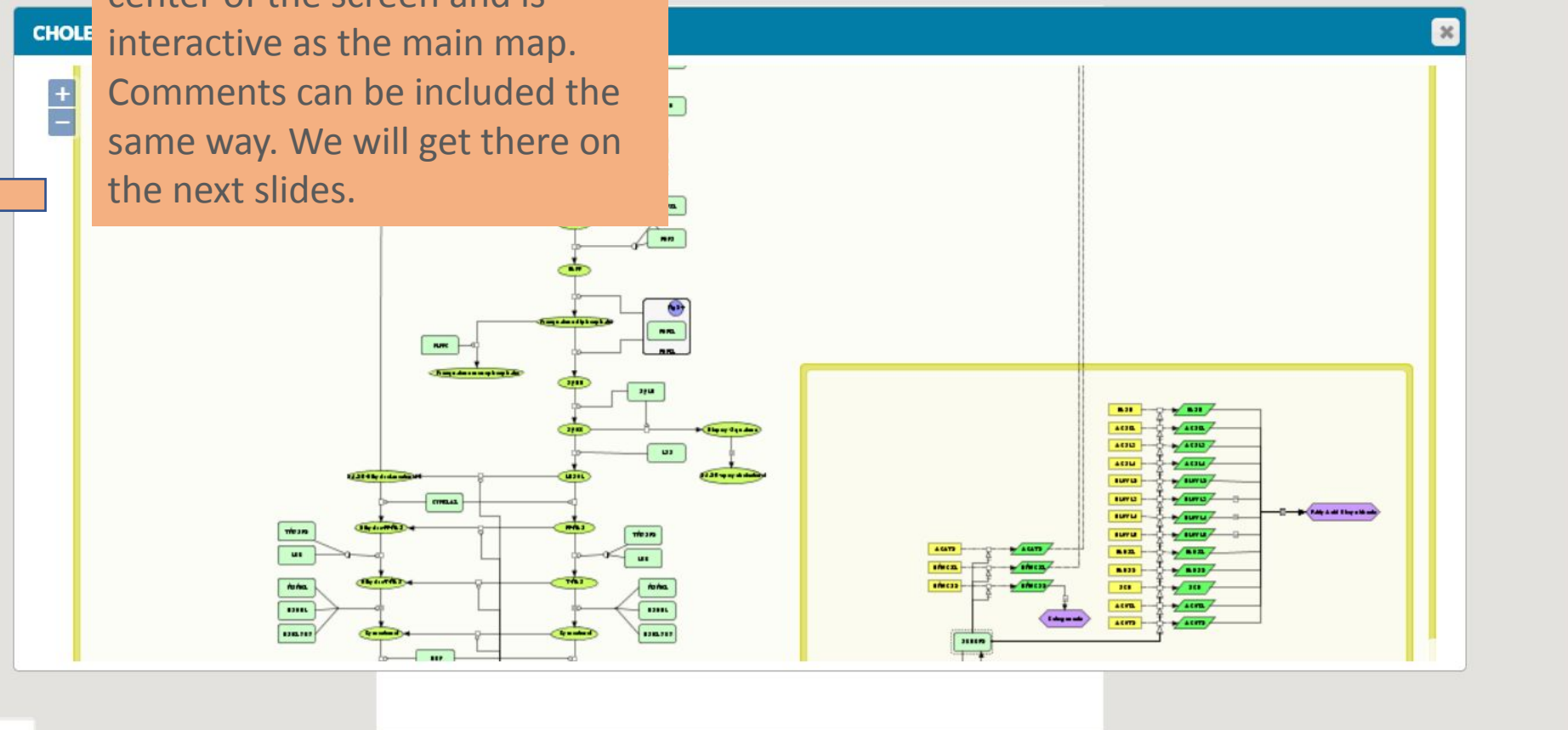


Name	Find anchor	View
Cholesterol_biosynthesis		
FA_and_lipoproteins_transp		
Glucagon_signaling		
Glucose_metabolism		
Insulin_signaling		
Mitochondrial_metabolism		



Click on the magnifier button to open a submap.

The submap will open on the center of the screen and is interactive as the main map. Comments can be included the same way. We will get there on the next slides.



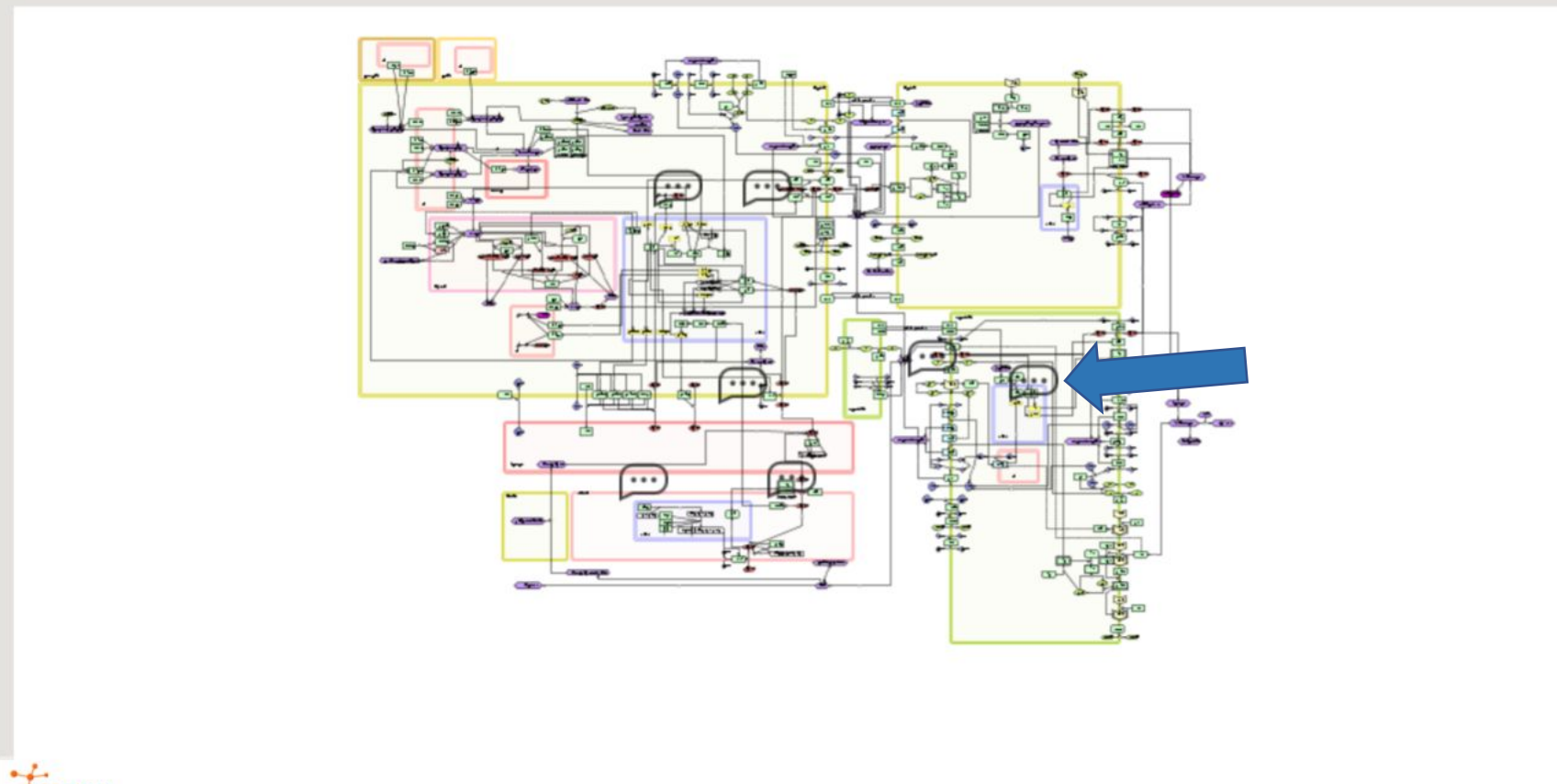
BACKGROUNDS: ?

Name	View
Pathways and compartments	
Network	
Empty	

USER-PROVIDED OVERLAYS:

No	Name	View	Data	Edit
No data available in table				
<div>Add overlay</div> <div> Refresh</div>				

In the top bar, we have one box that allows the display of the comments (orange arrow) in the map (blue arrow), so you can see where other contributors made comments and also click on the comments to open and read them.



Liver_Bile_Secretion_PM Feb.03_2022

SEARCH OVERLAYS INFO

BACKGROUNDS:

Name	View
Pathways and compartments	
Network	
Empty	

USER-PROVIDED OVERLAYS:

No	Name	View	Data	Edit
No data available in table				

Add overlay Refresh

Powered by MINERVA Platform (v16.0.7)

There is also a box opening a legend panel with the current version of the CellDesigner graphical notation (the notation used in the map construction).

For your reference, there is an image with the full panel in the next slide.

☒ LEGEND ☐ COMMENTS

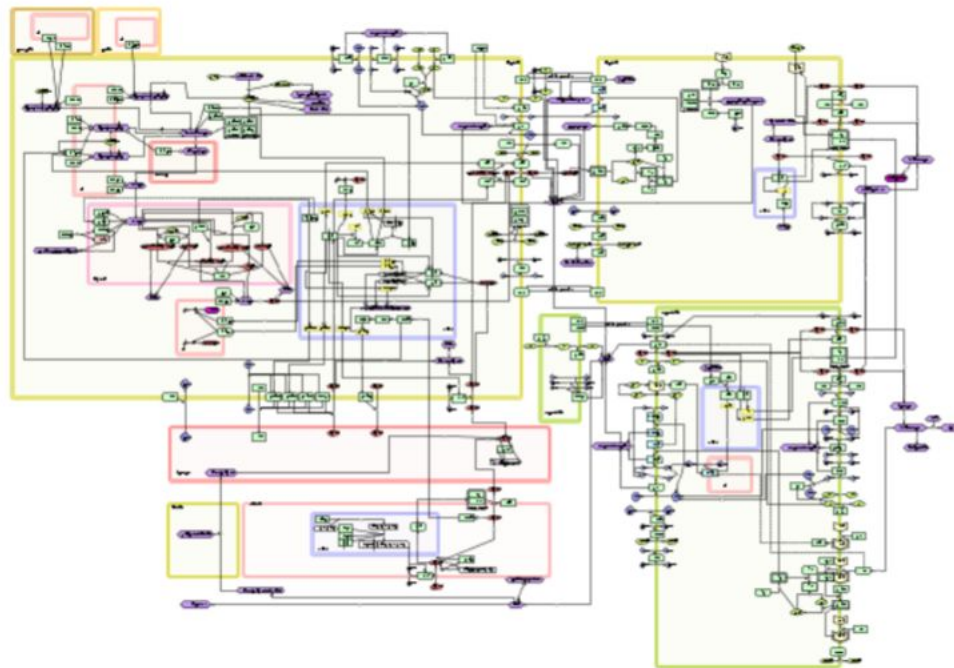
BACKGROUNDS

Name
Pathways and co
Network
Empty

To navigate through the map, you can use the zoom buttons or your mouse scroll. You can also use your mouse to click and drag to change locations. It is the same mechanism for using Google Maps, for example.

USER-PROVIDED OVERLAYS:

No	Name	View	Data	Edit
No data available in table				
Add overlay		Refresh		



BACKGROUNDS:

Next, you can choose where to pin the comment (in the "type" menu), and there is a place to type your comment.

Please, don't forget to identify yourself with your e-mail, so that we can reach you to discuss if needed.

ADD COMMENT

Type

Protein: NHE3

Pinned

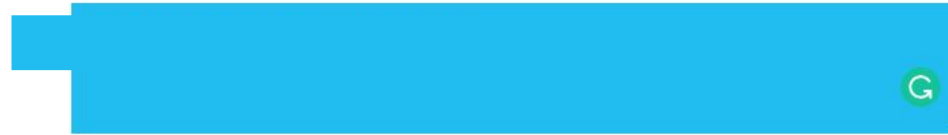


Email:

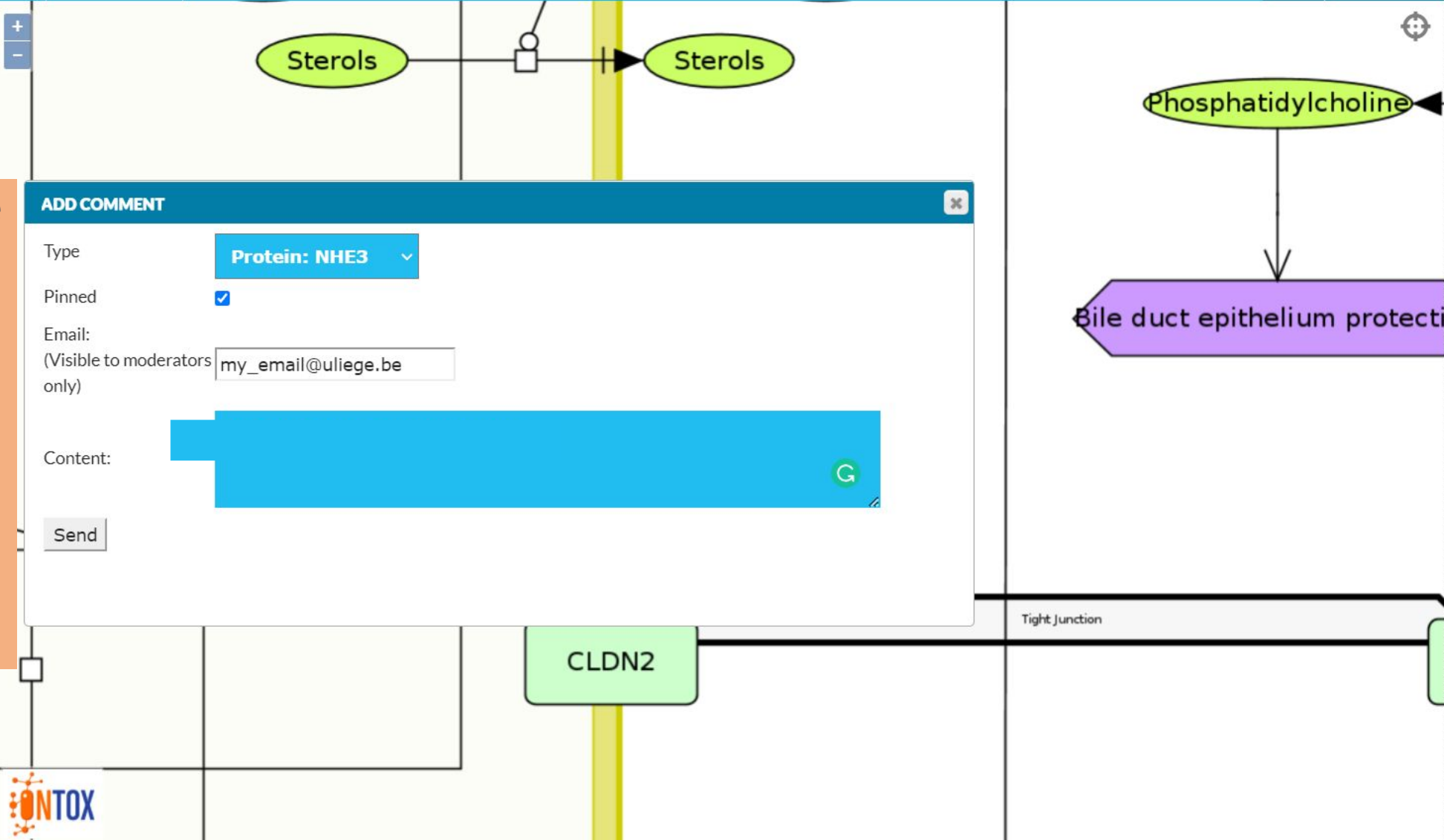
(Visible to moderators only)

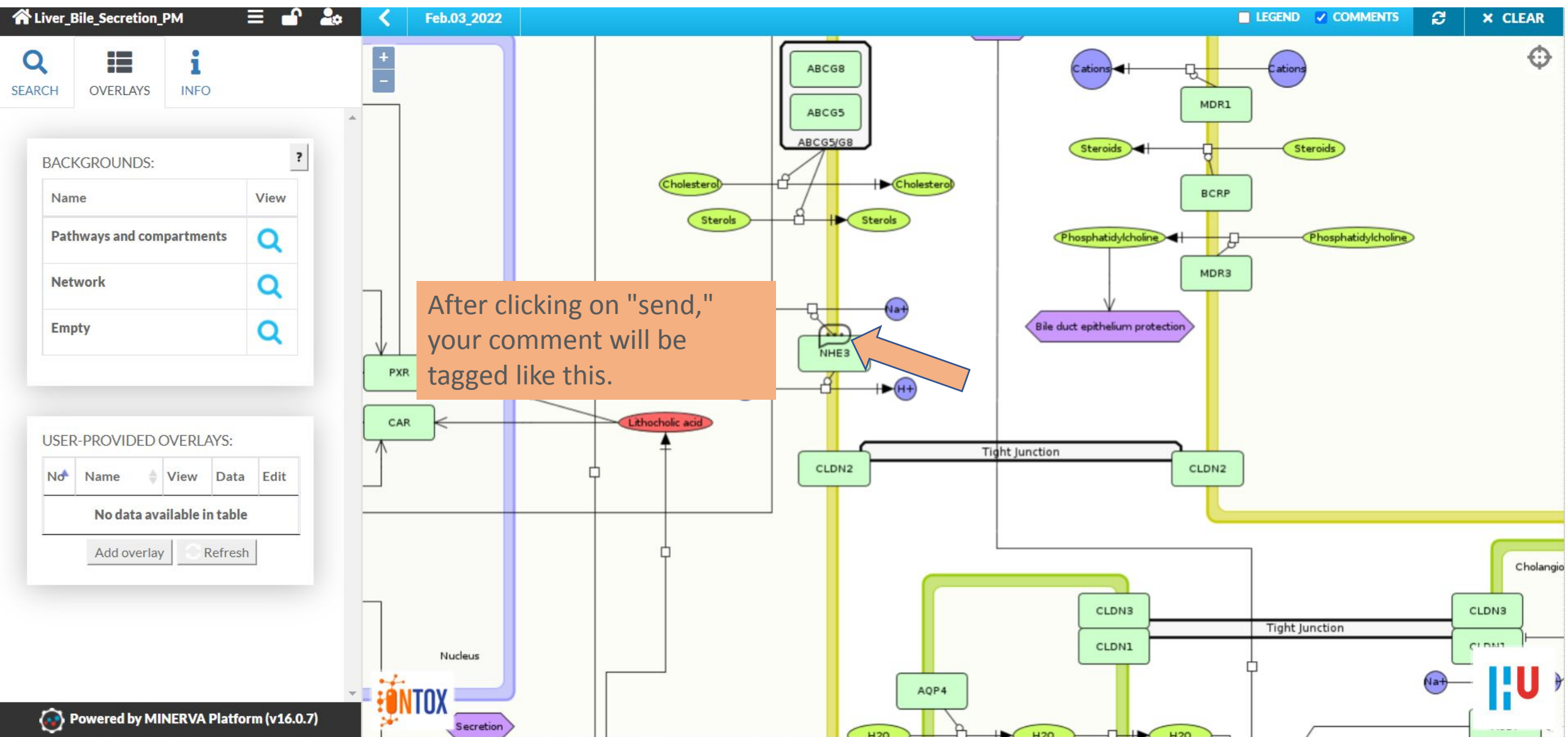
my_email@uliege.be

Content:



Send





Lipid_Metabolism_PM Mar.10_2022 LEGEND COMMENTS CLEAR

SEARCH OVERLAYS INFO

CONTENT DRUG CHEMICAL MiRNA

SEARCH IN CONTENT: keyword ☐ PERFECT MATCH

1 **Phenotype:** Triacylglyceride synthesis

Pathway: Triacylglyceride synthesis

Annotations:

Source: Annotated by curator

[1] [WikiPathways \(WP325\)](#)

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ONTTOX

GPAT

acyl dihydroxyacetone phosphate

fatty acyl CoA

DGAT

phosphatidic acid

phospholipids

diacylglycerol

MOGAT

monoacylglycerol

triacylglycerol

triacylglyceride synthesis

You might notice that when you click on some nodes, information appears in the left-hand menu.

These are the node's annotations, and they could have links to other databases, like the Wikipathways in this example. Then, you can click on the link (orange arrow) to access the complementary information.



Some considerations to take into account when reviewing a map:

The physiological maps are not homogeneous: this means that different mechanisms described could have very different levels of details, and this is not necessarily a problem. If you feel that a mechanism could benefit from more details, please do not hesitate to comment.

Including a reference in your comment is of great help for us doing the upgrades of the maps. The DOI or PubMed ID of a paper might be sufficient for us to find the information, but the more details you give us, the more we benefit from your comments.

And of course, you can always send your suggestions and questions also by email.

Thank you!



Useful links:

ONTOX's Physiological Maps Curation Guidelines:

https://github.com/ontox-maps/guides_and_documentation/tree/main/guidelines

MINERVA user manual:

https://minerva.pages.uni.lu/doc/user_manual/v15.0/index/

Basic exploration tools:

https://minerva.pages.uni.lu/doc/user_manual/v15.0/index/#basic-exploration-tools

Contacts:

University of Liège Team

Liesbet Geris - liesbet.geris@uliege.be

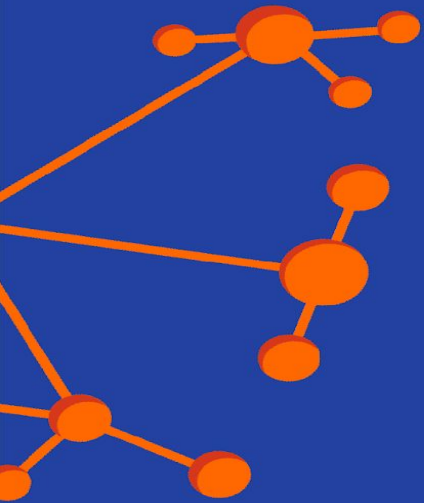
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