



WELCOME TO THE

# ETH PHRT

# Swiss Multi-Omics Center

INTEGRATED DIGITIZATION OF CLINICAL BIOSPECIMEN COHORTS ON THE MOLECULAR LEVEL

DNA | RNA | Protein | Metabolites | Lipids

LEARN MORE

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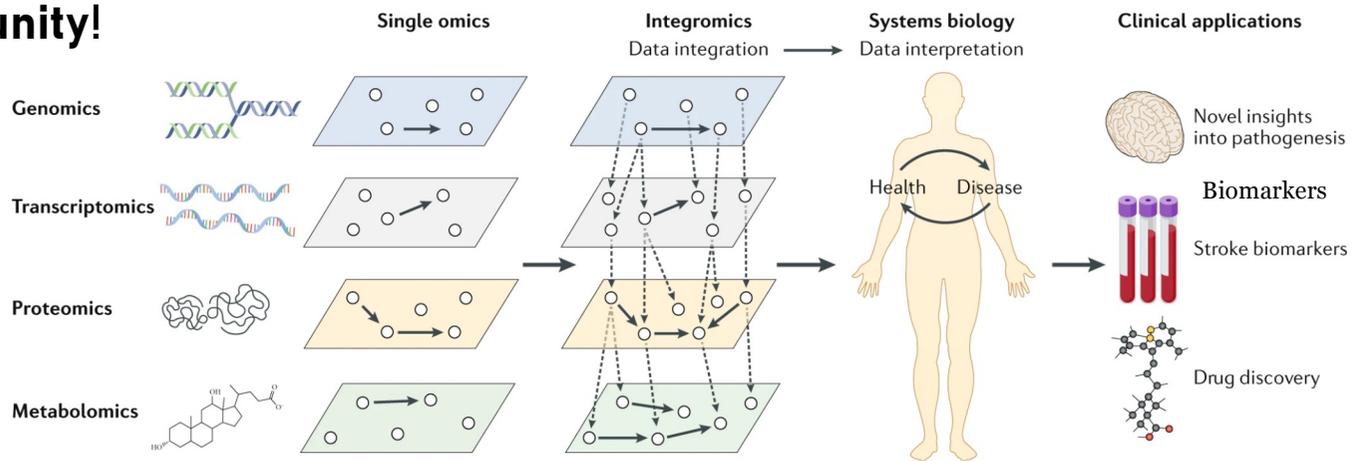
# Personalized health builds on the combination of clinical and molecular data

## Sample digitization is a necessity!

Clinical samples cannot be regenerated. Turning clinical specimens into searchable and reusable digital biobanks paves the road to retrospective and highly-powered studies.

## Sample digitization is an opportunity!

Integrative analyses drive discovery of mechanisms and novel targets.



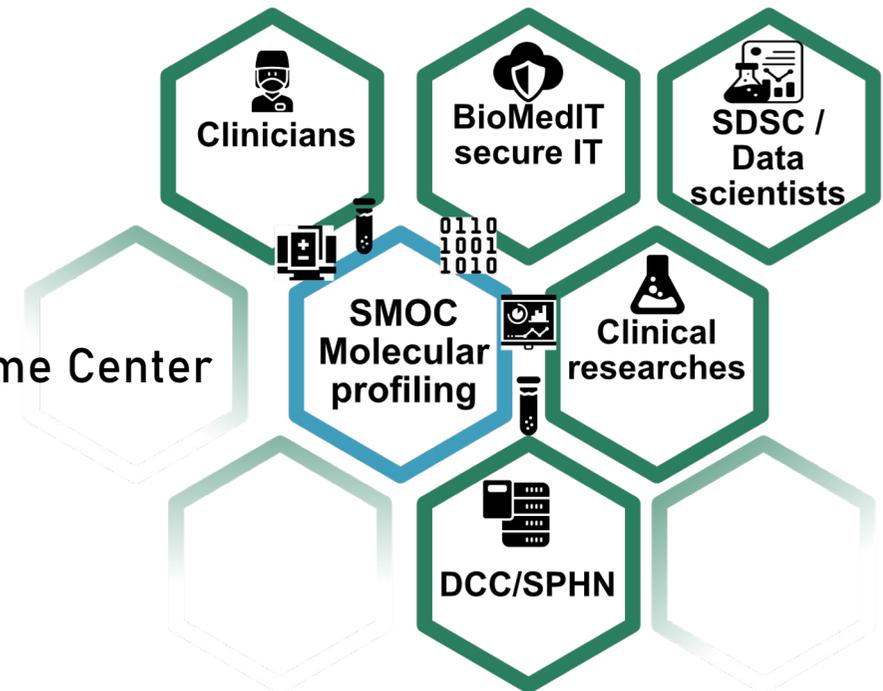
# The ETH PHRT Swiss Multi-Omics Center (SMOC)

## Mission: Enable multi-omics analyses of clinical samples

- Deliver clinical-grade, population scale, molecular data enriched with expert data analysis
- Democratize access to data (costs, actionability)

## Organization: Integration of existing expert units

- **genomics, transcriptomics @ Health 2030 Genome Center**
- **proteomics @ ETH Zürich**
- **metabolomics, lipidomics @ ETH Zürich**
- Tight collaboration and coordination across Centers
- ~10 PIs/Senior Scientists involved in management



# The ETH PHRT Swiss Multi-Omics Center: Portfolio

PHRT Swiss Multi-Omics Center

## PHRT Clinical Omics Centers

Genomics

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**CGAC**  
Clinical Genomic  
Analysis Center

Proteotyping

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**CPAC**  
Clinical Proteotype  
Analysis Center

Metabolomics & Lipidomics

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**CMAC**  
Clinical Metabolomics  
Analysis Center

**Measure**

Single entry point for digitizing your clinical cohorts at the level of DNA, RNA, Proteins, Metabolites and Lipids

**Integrate**

Integrated FAST track and DEEP track analysis of multi-omics data

**Analyze**

Delivering molecular insights supporting publications and clinical decision making

**Share**

FAIR sharing of primary and derivative data based on ethical consent

 <b>GETTING THE MOST OF YOUR CLINICAL SAMPLES</b> Single entry point for integrated multiomic analysis.	 <b>MOLECULAR INSIGHTS INTO CLINICAL COHORTS</b> Standardised and automated processing of large clinical cohorts.	 <b>SAFE DATA STORAGE AND EXCHANGE</b> Handling of sensitive clinical data according to established <a href="#">SPHN/BioMedIT</a> guidelines.
 <b>CONSULTANCY</b> We will find together the right strategy for digitizing and analyzing you clinical samples.	 <b>FEASIBILITY STUDY</b> Small scale studies for optimizing clinical cohort experimental design and generating proof of concept data.	 <b>SUPPORT WITH GRANT AND PAPER WRITING</b> We will provide relevant text snippets supporting grants and publications.

**One entry point, three centers with expertise across the value chain.  
 Scalable production and analytics.**

# Multi-Omic investigation of rare disease led to therapeutic insights

nature metabolism



Article

<https://doi.org/10.1038/s42255-022-00720-8>

## Integrated multi-omics reveals anaplerotic rewiring in methylmalonyl-CoA mutase deficiency



**Hope for patients with a severe rare disease**  
By combining the results of multiple molecular analyses, scientists can better diagnose the hereditary disease methylmalonic aciduria. There is also hope with regard to therapy.



UNIVERSITÄTS-  
**KINDERSPITAL  
ZÜRICH**

### News & views

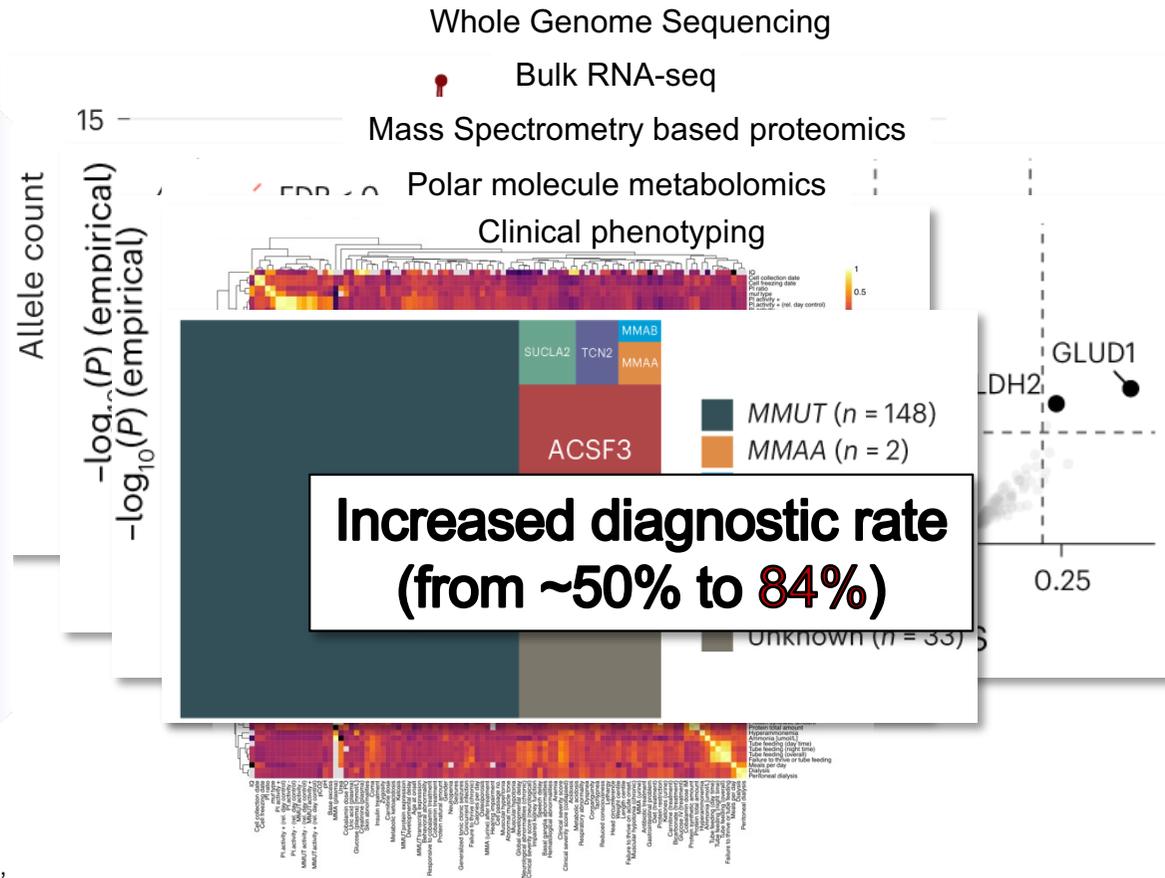
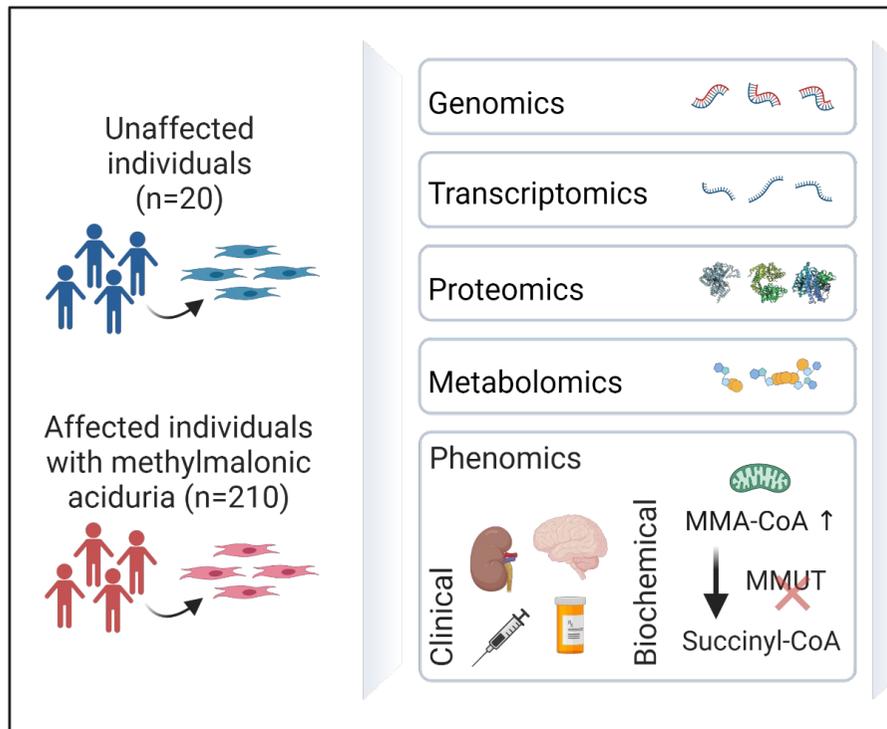
Mechanisms of disease

## Anaplerosis in action

Pamela Sara E. Head & Charles P. Venditti

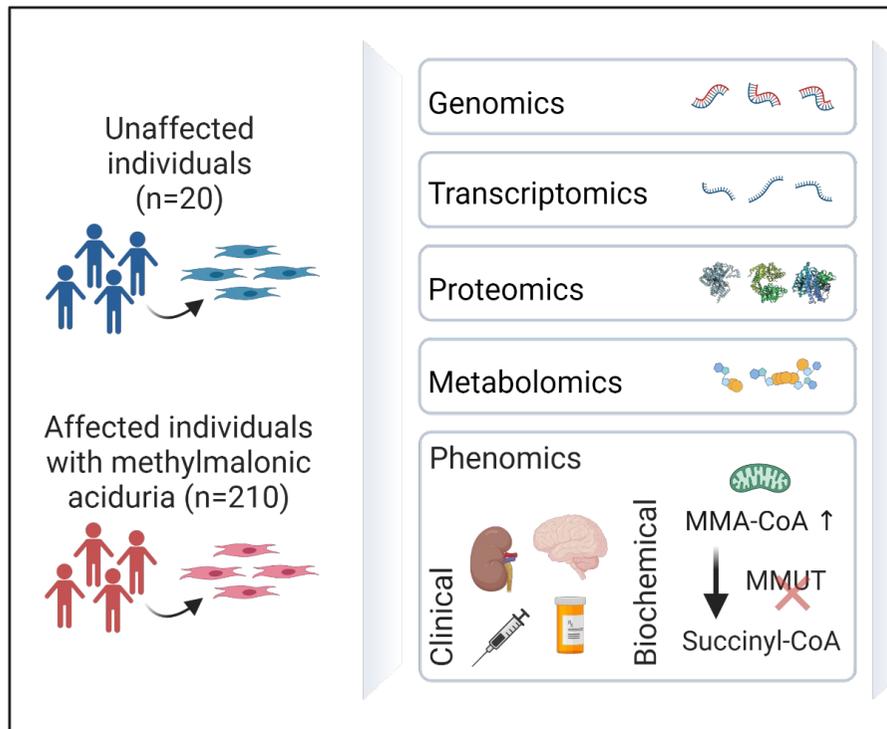
Investigation of multi-omic changes and their effects on regulation of metabolic pathways confirm anaplerotic deficiencies in methylmalonic acidemia, strengthening the need for future therapies aimed at replenishing intermediates of the tricarboxylic acid cycle.

# Integrated multi-omics analysis in methylmalonic aciduria

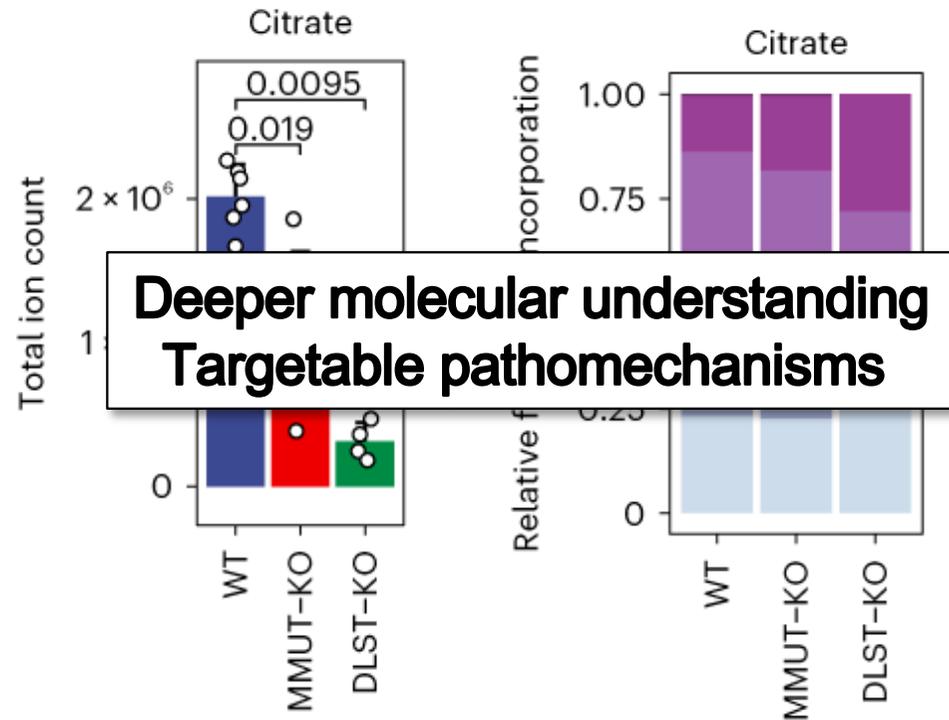


Forny, P., Bonilla, X., Lamparter, D., Shao, W., Plessl, T., Frei, C., Bingisser, A., Goetze, S., van Drogen, A., Harshman, K., Pedrioli, P.G.A., Howald, C., Poms, M., Traversi, F., Bürer, C., Cherkaoui, S., Morscher, R.J., Simmons, L., Forny, M., Xenarios, I., Aebersold, R., Zamboni, N., Räscher, G., Dermitzakis, E.T., Wollscheid, B., Baumgartner, M.R., Froese, D.S., 2023. *Nature Metabolism*.

# Integrated multi-omics analysis in methylmalonic aciduria



## Disturbed TCA Metabolism



**Deeper molecular understanding  
 Targetable pathomechanisms**

Forny, P., Bonilla, X., Lamparter, D., Shao, W., Plessl, T., Frei, C., Bingisser, A., Goetze, S., van Drogen, A., Harshman, K., Pedrioli, P.G.A., Howald, C., Poms, M., Traversi, F., Bürer, C., Cherkaoui, S., Morscher, R.J., Simmons, L., Forny, M., Xenarios, I., Aebersold, R., Zamboni, N., Rätsch, G., Dermitzakis, E.T., Wollscheid, B., Baumgartner, M.R., Froese, D.S., 2023. *Nature Metabolism*.

# The ETH PHRT Swiss Multi-Omics Center in the Swiss Health Ecosystem

Catalyst for

*translation of basic science into clinics*

*longitudinal multi-omics*

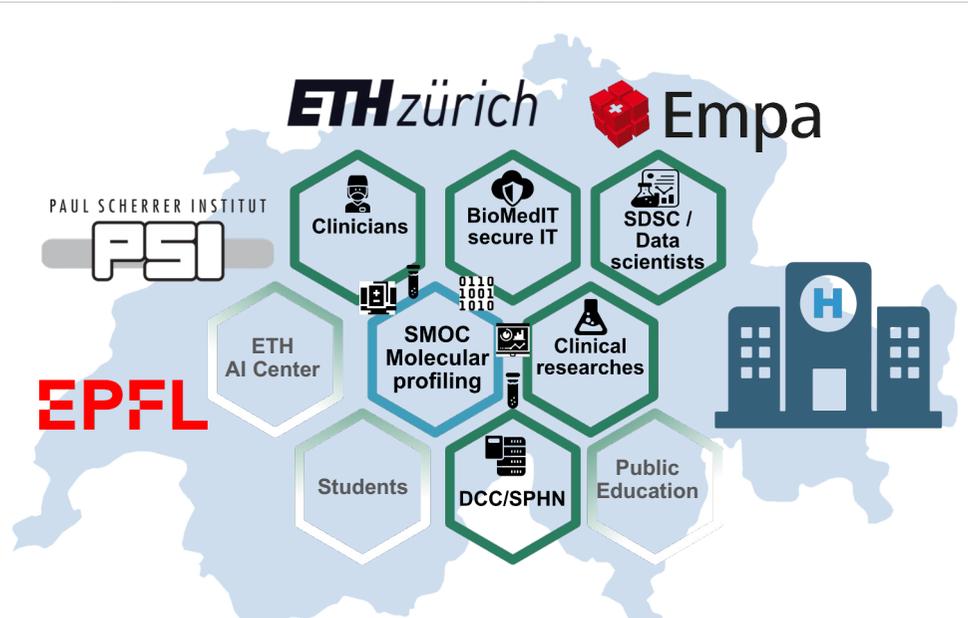
*interventional trials*

*mechanistic and functional understanding of disease*

*new predictive diagnostics*

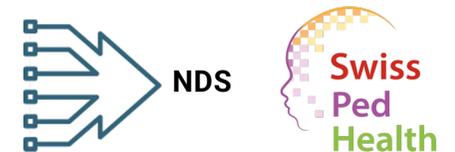
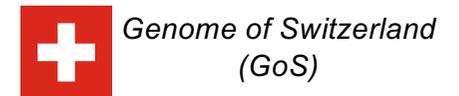
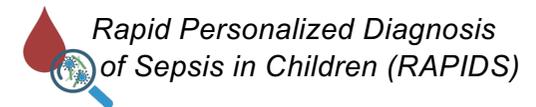
*novel treatment approaches*

Digital Switzerland Strategy & BFI-Botschaft 25–28

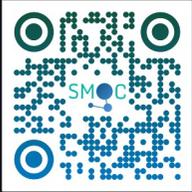


ETH domain infrastructure nourishing Swiss PH ecosystem

Enabler of



Building on the strategic investments made by the SFA-PHRT and SPHN initiatives to establish a sustainable infrastructure for deep human molecular analysis.



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**“Turning clinical specimens into searchable and reusable digital biobanks drives the development of multimodal data analysis technologies supporting clinical decision-making in Switzerland”.**

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