

European infrastructure for translational medicine



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Our mission: Accelerate the translation of research discoveries into patient benefit



We support, using our capacities and expertise, academia, industry, patients and policy makers.

EATRIS acts as facilitator for the development and adoption of Personalised Medicine in Europe and globally.

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Who we are : A LAAAAARGE COMMUNITY



Facilities, resources and services to support translational research & Personalized Medicine



Five Scientific Platforms: dynamizing research

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ADVANCED THERAPY MEDICINAL PRODUCTS

Tissue engineering, Gene therapy, Cell therapy, GMP facilities, vector design & production

BIOMARKERS

Biobank facilities, Multiplexed immunostaining, Deep genome sequencing

IMAGING AND TRACING

(pre-clinical) PET imaging, GMP tracer development and production, (Ultra) high field MRI, Optical and hybrid imaging SMALL MOLECULES

Advanced screening (also in 3D cultures), Development of xenograft and *in vivo* models, Drug (re-)formulation, (Pre-)clinical validation nanomedicines VACCINE, INFLAMMATION AND IMMUNE MONITORING

Antigen characterisation, Vaccine formulation, Process development

What we do to facilitate the progress of Personalised Medicine





Provide access to research services and expertise to industry, academia and research funders, particularly at the preclinical level

Develop **new research tools** that facilitate patient stratification

Train the next generation of scientific experts in Personalised Medicine

Improve the translational **research eco-system** to accelerate the progress of Personalised Medicine

EATRIS PLUS: Catris a flagship project for creating new services and resources for Personalised Medicine R&D

EATRIS-Plus is aiming to enhance EATRIS' long-term sustainability by maturing key capacities of the infrastructure and offering access to scientific tools and services to support Personalised Medicine.

- 20 partners (including 14 EATRIS nodes)
- 2 non-member countries represented: Ireland & Germany
- 4 years of funding (2020-2023) €4,9 million
- Project runs until December 2023

Find out more: eatris.eu/eatrisplus



Reflecting & Addressing the Infrastructure's Objectives



provide access to infrastructure tools and resources novel academia-industry collaboration models



strengthen operational practices, quality excellence and financial performance



drive patient empowerment through active involvement in the research process



build lasting partnerships with key stakeholders at global level



WP1: Multi-omic technologies (Checz Rep)



Whole Genome Sequencing, IMTM/UP, CZ
Epigenetic modifications DNA- UU, SE
Metabolomic analysis- RUMC, NL
Proteome analysis- IMTM/UP, CZ
Transcriptome RNA- FIMM/UH, FI
MicroRNA sequencing- FIMM/UH, FI
MicroRNA qRT-PCR, IRYCIS, ES

WP2: Data Stewardship & Integration (The Netherdlands)

> WP3: Quality Assessment (Finland)

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WP 1

Multi-omic technologies for Personalised Medicine

- Mapping and benchmarking EATRIS' omic technical capacities
- Establishment and validation of reference tools for multi-omic technologies
- Samples processing and analysis using multi-omics methods, definition of quality criteria for data outputs
- Developing a repository for mapping reference values for age-matched and gender-match stratified populations
- Analysing a healthy cohort that will be the basis for the normal reference database that relates to the personalised medicine

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WP 2

Data stewardship and integration of omic research in Personalised Medicine

WP 3

Quality Assessment for multi-omic technologies

- FAIRification of data and analysis pipelines
- Design workflows for joint analysis of cross-omics data from different cohorts
- Demonstrate added value of the cross-omics analysis to Personalised Medicine

- Establishment of reference materials and agreement on SOPs
- Inter- and intra-laboratory quality assessment
- Assembling EATRIS-Plus Multi-omic Toolbox
- Establishing the EATRIS Quality Certificate

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PM ORIENTED



An open access resource, containing:

- SOPs
- Guidelines for best practices
- Reference materials
- Quality parameters
- Data analytical tools
- Criteria for reference values
- Troubleshooting guidelines
- Repository of multi-omic data

To enable high-quality research in the context of patient stratification and accelerate PM solutions implementation.





Multi-Omic Toolbox

- Multiomic Toolbox WG: WP1, WP2, WP3
- Consolidating contents: "in house" and complementary
- Organizing contents for easy information access and flow : categories, subheads, icons, searching tools, questions...
- Harmonization with other EATRIS Tools
- Attending Broad range of potential users: different interest, different paths
- Dinamyc tool: prototype for "challenging" by users and relevant stakeholders
- Linking other EATRIS Plus resources: Education and training & Regulatory & QC & Patient Engagement

Add Value in Personalized Medicine based on OMICs approaches



1st EATRIS-Plus Multi-omics Stakeholder Group workshop

March 2021

Frontiers | Frontiers in Molecular Biosciences

- Moving beyond genomics
- Evaluating new technologies
- Discussion topics Data standardization
 - Capture variability omics data at source

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- Data privacy and regulatory aspects
- Implementation in clinical care



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TABLE 3 Summary of challenges and recommendations from the EATRIS + multi-omics workshop (March 2021).

| Challenge | Recommendation | |
|---|---|--|
| Moving beyond genomics | Communicate and educate on the pros and cons of other omics technologies such as proteomics, metabolomics and lipidomics | |
| | Develop multi-modal data integration models that showcase the added value of multi-omics approaches Personlized Medicine | |
| New technologies, new challenges | • Share lessons-learned, failures and successes when evaluating new technologies in Personalized Medicine | |
| | Evaluate the added value of Artificial Intelligence and Digital health in Personalized Medicine, particularly combination with multi-omics data | |
| Data standardisation | Adopt international standards of health data and models including the FAIR principles of data stewardsl (e.g., OMOP, FHIR, CDISC) | |
| | Define criteria for quantity, quality and FAIR levels of data prior to multi-modal data analyses for a speci objective in Personalized Medicine | |
| | Work with flexible and dynamic mathematical models to adapt to changing data collections in Personaliz Medicine | |
| Variability in omics data at source | • Use internationally recognised laboratory standards and standard operating procedures for omics analys | |
| | Adopt and apply quality assurance and control schemes for laboratories, such as the EATRIS Certificate Commitment to Quality | |
| | • Include confounding factors such as population diversity in biological systems in the multi-modal data analy | |
| Data privacy and regulatory aspects | • Consider ethical, legal, societal aspects when designing multi-omics Personalized Medicine studies | |
| | Comply with international standards on data security, including the General Data Protection Regulation personal data | |
| | • Report of the successes and failures of implementations from the European landscape | |
| Implementation of Personalized Medicine in routine clinical care | Consider well prior to multi-omics Personalized Medicine implementation: 1) the benefits, 2) the risks, associated ethical and social aspects, 4) room for innovation | |

Check for updates

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RECEIVED 21 June 2022 ACCEPTED 08 September 2022 Tackling the translational challenges of multi-omics research in the realm of European personalised medicine: A workshop report

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FAIRification of multi-omics metadata

Workshop at the 21st European Conference on Computational Biology (ECCB) Organized by Gary Saunders, Anna Niehues, Emanuela Oldoni

| | The FAIRification process – two use cases | The EATRIS-Plus multi-omics demonstrator cohort, Anna Niehues The iCAN Digital Precision Cancer Medicine project, Veronika Suni | Bottlenecks and challenges Finding and choosing relevant data |
|-------------------|--|--|--|
| | Tools, services, and resources to facilitate multi-omics data FAIRification | FAIRsharing: discover and curate an ecosystem of research standards and databases, Allyson Lister 10 years of ISA: Lessons learned from the community and recent developments, Philippe Rocca-Serra | and metadata standards Increased amount of data standards, and data sets submitted to repositories |
| | FAIRification practices at omics data repositories | MetaboLights and FAIRification, Claire O'Donovan The PRIDE database: Enabling FAIR practices for proteomics data, Deepti J. Kundu and Juan Antonio Vizcaino | Necessary next steps Address increased needs for database management and curation Key messages |
| $\langle \rangle$ | Interactive tools to capture and share metadata using templates | Sharing study meta-data for biologists: the Phenotype database as solution, Jildau Bouwman FAIR Genomes and MOLGENIS as a FAIRification platform, Joeri van der Velde | Reach out to communities, tool developers, and service providers to communicate needs of researchers and find solutions together |

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Panel discussion

Many thanks to our committed members

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* The contribution in Norway is shared between University of Oslo (UiO), University of Bergen (UiB), Norwegian University of Science and Technology (NTNU), the Arctic University of Norway (UiT) and the four Regional Health Authorities in Southeastern, Western, Central and Northern Norway



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Translational Trends

