



ACDC Kick-Off Program

12:45 – 13:00 Welcome & Registration

Introductions

	Opening	Daoud Sie and Katy Wolstencroft
13:00-13:10	Core Facility Sector Plan	Rogier Postma
13:10 – 13:25	The Science Case – Multi-omics and visual omics research at AUMC	Eric Reits
13:25 – 13:45	The ACDC platform	Daoud Sie & Katy Wolstencroft

Research support for Omics & Bioimaging

13:45 – 14:00	A FAIR outlook for AUMC researchers	Elize Vlainic
14:00 – 14:10	Funding Opportunities for Core Facilities in grant applications	Laura Manca

Coffee Break

ACDC Ongoing projects & User stories

14:30 – 14:40	Automating metadata structuring for Metabolights submission	Karen Sap
14:40 – 14:50	Genomics data pipelines with iRODs	Bilgehan Nevruz
14:50 – 15:00	Proteomics and pooling data for analyses	Alex Henneman
15:00 – 15:10	Making Flow Cytometry Flow	Alberto Miranda Bedate
15:10 – 15:20	QuPath Analysis on HPC	Giulia Bergamaschi
15:20 – 15:30	Genomics analysis on HPC	Yongsoo Kim
15:30 – 15:40	Biomero – compute and imaging	Przemek Krawczyk
15:40-15:50	Q & A all project talks	

Wrap-up & Close

15:50 – 16:05	ACDC Roadmap	Katy Wolstencroft and Daoud Sie
16:05 – 16:15	Wrap-up and close	
16:15 – 17:30	Drinks	

Core Facility Sector Plan

Rogier Postma - Managing director division 9



A Healthy Future For All

Multi-year strategy Amsterdam UMC

Today's challenges

- Growing demand for care
- Aging population
- Tight labor market
- Climate change
- New pandemics

Impulses

- Acting today for a better tomorrow
- Health and sustainability are part of our DNA
- Digital transformation for all
- Flexibility for employees

Discovering Tomorrow's Healthcare Together

Core values

- Caring
- Proactive
- Inquisitive

Core tasks

- Research
- Care
- Education & Training
- Valorization

Our ambitions

Our role

- We're taking action to help shape the health-care of the future.
- Stronger networks and alliances.

Our colleagues

- Diversity is our strength.
- Social connection is key.
- We're harnessing technology to change the way we work.
- We trust each other

Our content

- Connecting core tasks for innovation
- Towards a healthy and sustainable future.

What do you mean, rocket science?
There's nothing we can't do together.

Join us!

Imaging & Omics Core Facilities

- **Imaging & Omics Core Facilities**
 - Cellular Imaging
 - Proteomics Core Facility
 - Core Facility Metabolomics
 - Microscopy and Cytometry Core Facility
 - Core Facility Genomics

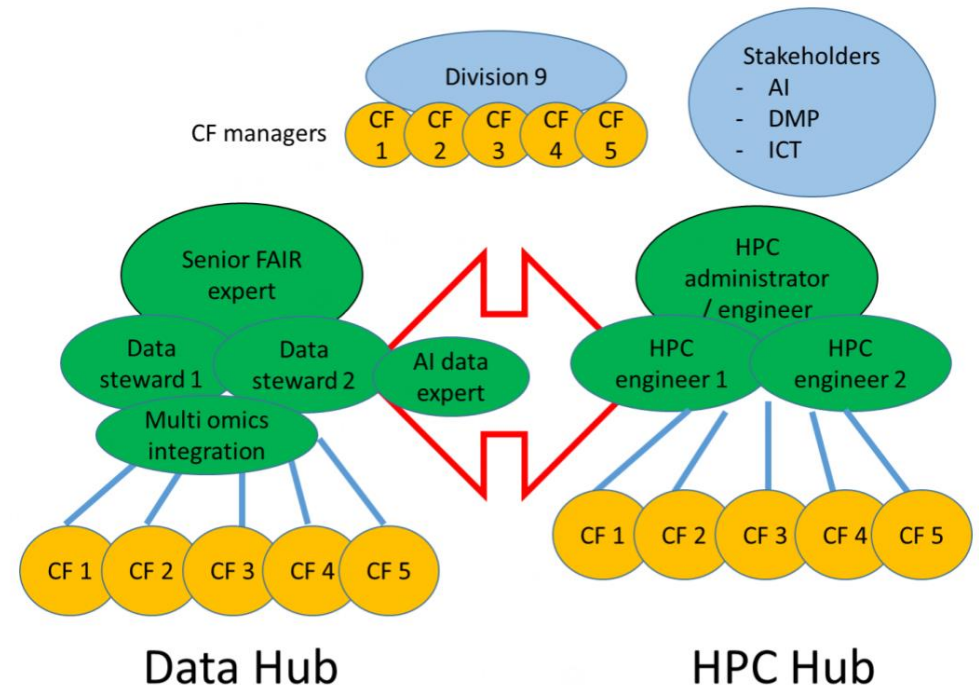


Core facilities Amsterdam UMC outlook 2023

- The rapid developments in advanced imaging and omics technologies and AI data analysis in life sciences are facilitated by the 5 Amsterdam UMC Core Facilities (CFs) covering flow cytometry, microscopy, genomics/transcriptomics, proteomics, and metabolomics. State-of-the-art technologies, professional training and support, optimal quality control, and efficient use of instruments enable scientific breakthroughs by the Amsterdam UMC research community, yet the harmonized core facilities are faced with a number of challenges that need strategic investments.
- We propose to create a 1) Data Hub to integrate CF data (FAIR and AVG compliant) and a 2) High-Performance Computing (HPC) Hub for all CF data, enable pipelines and applicable for scientists. Integrating data from imaging and omics core facilities will catalyze innovations, minimize impact on costs for researchers, and enhance the visibility of Amsterdam UMC in the Dutch and European research infrastructure landscape. Strengthening the Amsterdam UMC core facilities and combining the advanced data-management and AI-driven data analyses, the Amsterdam UMC can take on ambitious challenges in life sciences and beyond, enabling breakthrough impacts in research towards new therapies in personalized and regenerative medicine.

Omics & Imaging Data & Compute Hub

- Integrated data and compute platform
- Consolidate local solutions
- Enable multi-omics and AI research
- For the core facilities and their users



Multi-omics & visual-omics research at Amsterdam UMC

Kick-off - 10-03-2025

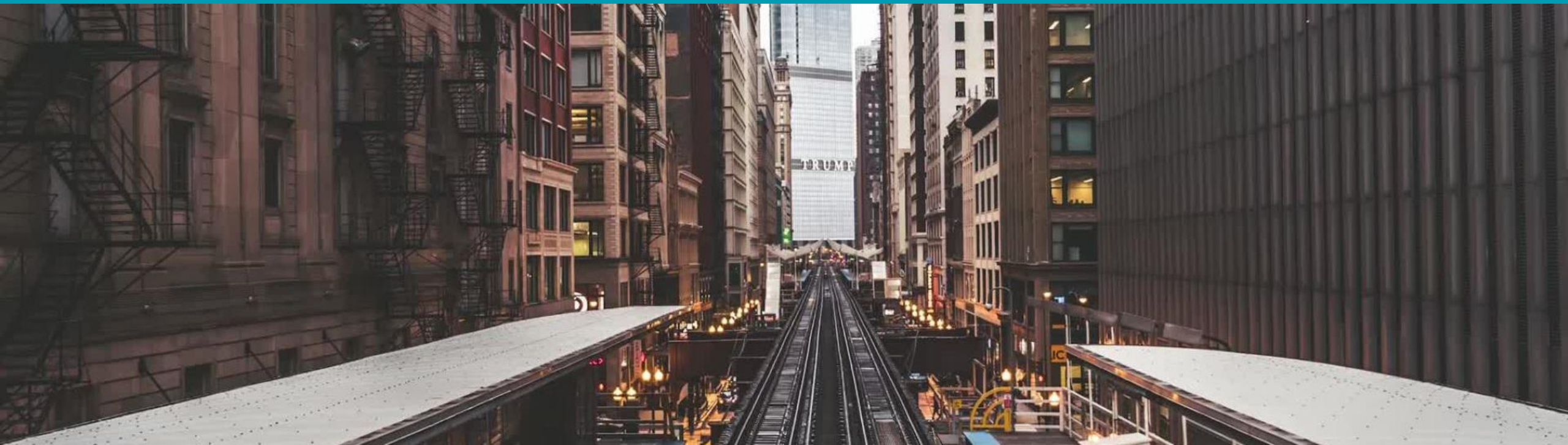
Prof dr. Eric Reits - Facility director Cellular Imaging



The Advanced Compute & Data Core

Dr. Daoud Sie

Dr. Katy Wolstencroft



AC



Daoud Sie

DC



Katy Wolstencroft



Alex Henneman



Bilgehan Nevruz



Karen Sap



Alberto Miranda Bedate

Infrastructure to Provide



FAIR Research Management

Omics, bioimaging and multi-omics
Integration and reuse



High Performance Compute

Omics and bioimaging expertise
For AI and LLM training
For Big Data analysis



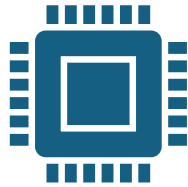
Sustainability

Efficient use of common resources
Continuity and collaboration

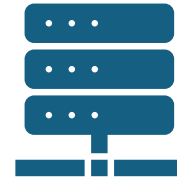
Components of the ACDC Infrastructure



Data: formats, size, storage, archive, backup, accessibility



Compute: CPU, GPU, accessibility/security, software



Metadata: protocols, experiments, models, instruments, raw data, derived data, workflows



Analysis Workflows: provenance, data links, reproducibility

Design Principles



Reuse components

Build on existing software

Adopt community standards



Incremental Approach

Immediate bottlenecks in each CF

Improve the FAIRness of each CF

Pilots first, get feedback, then extend



Minimise extra work

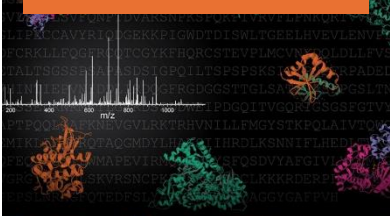
Automation and integration

We like DM and HPC, but its not for everyone!

Metabolomics



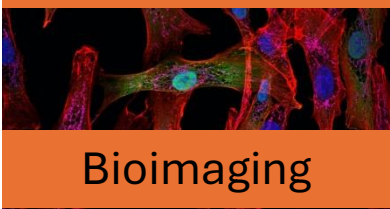
Proteomics



Genomics



Flow Cytometry



Bioimaging

Supporting the Data Life Cycle



Build on existing standards and tools

- AUMC RDM, ICT compute and storage, data privacy, data policy
- FAIRSharing.org – standards registry
- ELIXIR Research Data Management Toolkit – recommended tools

Gap Analysis and Harmonisation

Standards

Data Formats

Repositories

Community

Metabolomics

ISA-TAB
ChEBI

mzML

MetaboLights

MSI

Proteomics

MIAPE-MS
MIAPE-MSI
MIAPE-Quant
PSI-MS

mzML
mzIdentML
mzQuantML

Proteome
Xchange

PSI
HUPO

Genomics

MiXS, MINSEQE
SO

Fastq
Bam
Bed

ENA
EGA

INSDC, HUGO,
GSC

Flow Cytometry

MiFlowCyt
FlowCL

FCS/ACS

Flow Repository

ISAC

Bioimaging

REMBI
Fbbi

OME-TIFF/Zarr

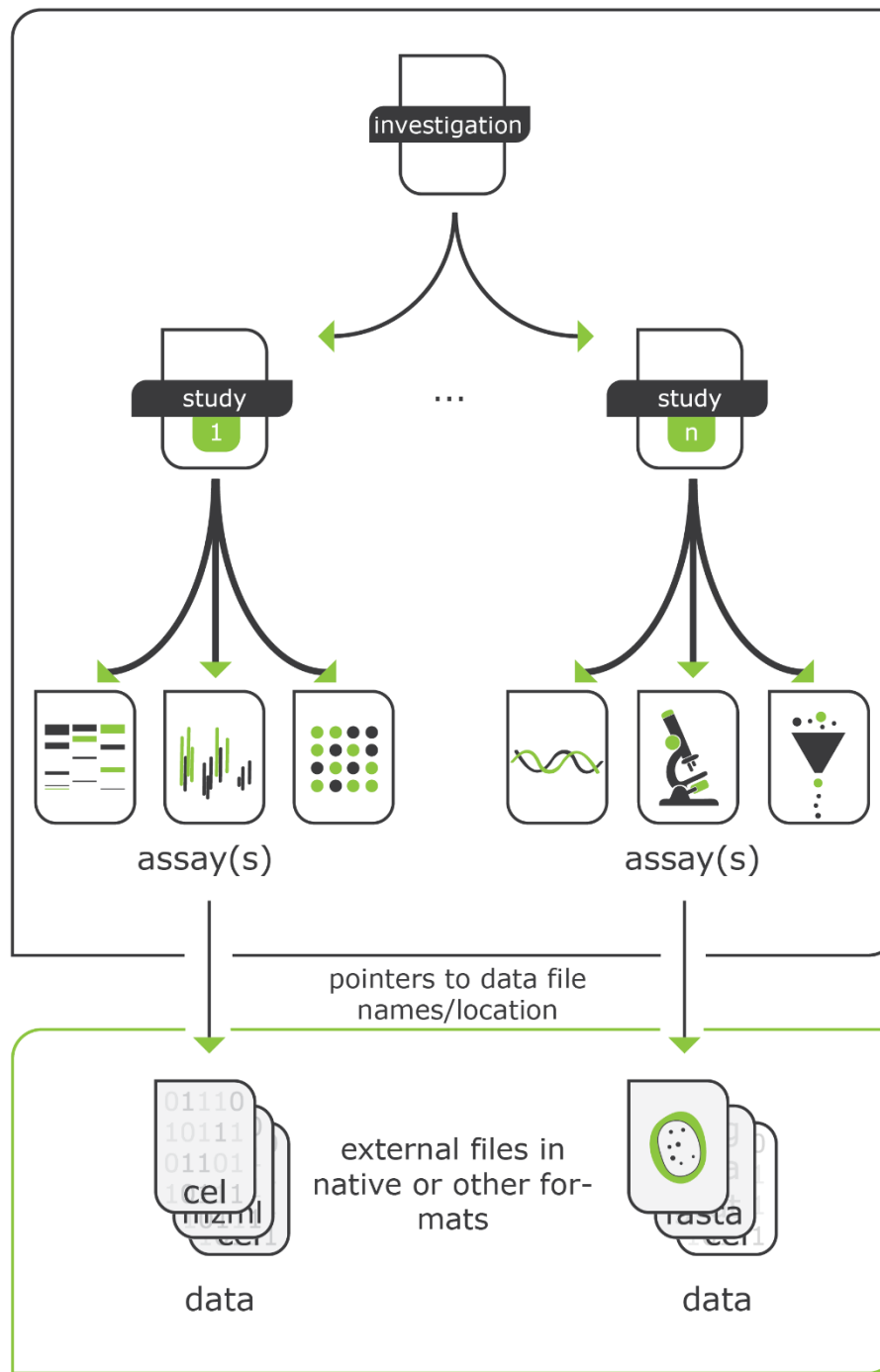
Biolmage
Archive

OME, image.sc

Multi-omics ISA Investigations Studies Assays

Recommended and used by:

- Public repositories, e.g. Metabolights
- ELIXIR and RDMKit
- Health-RI Omics working group
- Netherlands X-Omics



isa

investigation

high level concept to link related studies

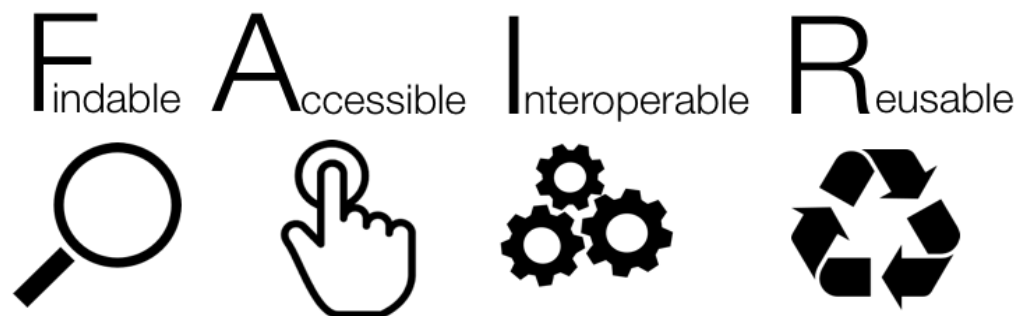
study

the central unit, containing information on the subject under study, its characteristics and any treatments applied.

*a study has associated **assays***

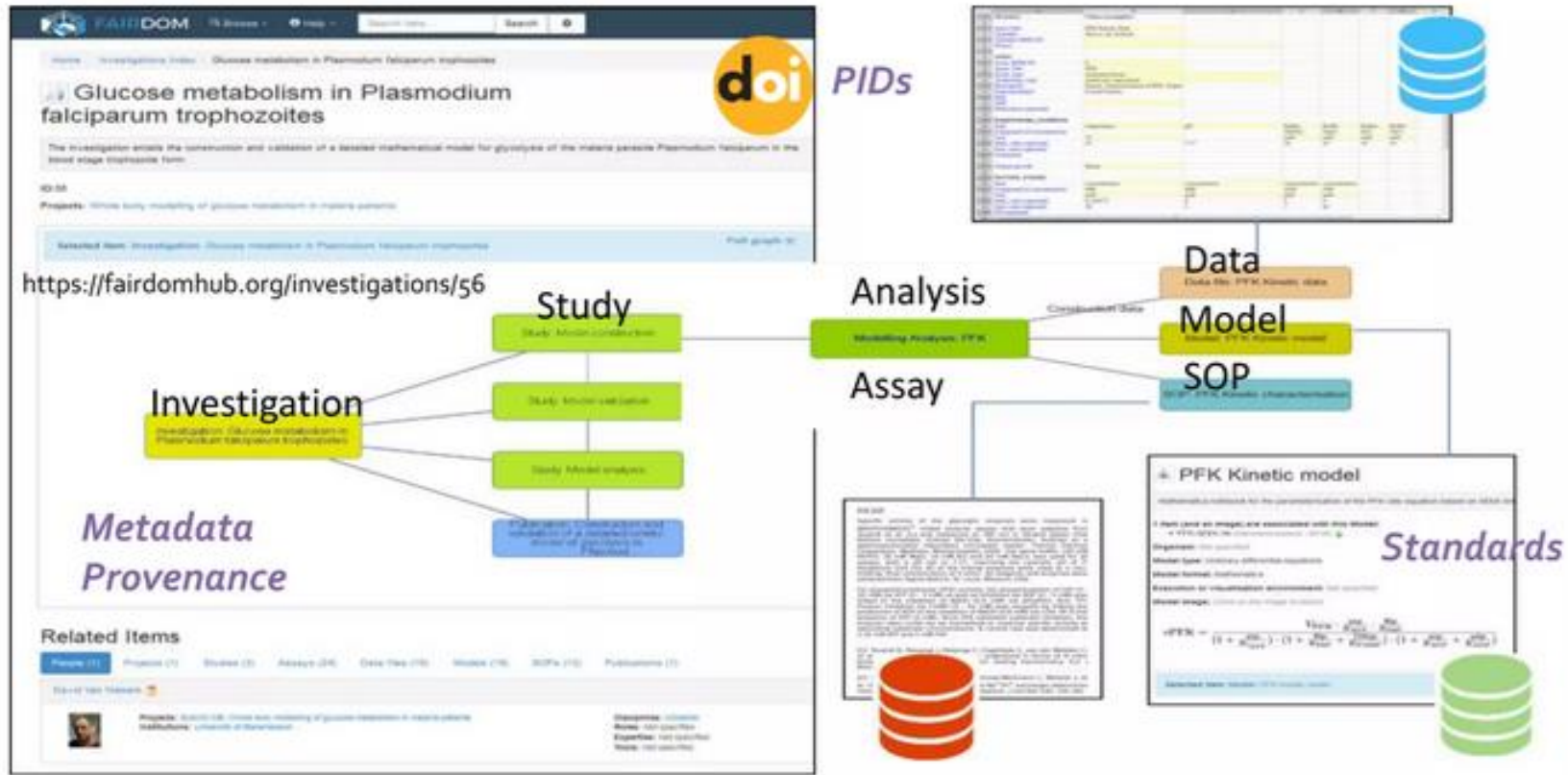
assay

test performed either on material taken from the subject or on the whole initial subject, which produce qualitative or quantitative measurements (data)



Findable	Accessible	Interoperable	Reusable
<p>F1: (Meta)data with unique PID</p> <p>F2: Rich Metadata</p> <p>F3: Metadata include ID of data</p> <p>F4: (Meta)data indexed in searchable resource</p>	<p>A1: (Meta)data retrievable by ID, protocol is open, authentication and authorisation where necessary</p> <p>A2: Metadata are accessible, even when data is not</p>	<p>I1: Shared formal language for knowledge representation</p> <p>I2: (Meta)data use FAIR vocabularies</p> <p>I3: (Meta)data include qualified reference to other (meta)data</p>	<p>R1: (Meta)data are richly described with:</p> <ul style="list-style-type: none"> - Data usage license - Clear provenance - Meet relevant community metadata standards

FAIRDOM SEEK: Experimental Context



Federated Catalogue, Integrated View

fairdomhub.org

FAIRDOM-SEEK software is:

- Natively ISA
- Open-source software
- [BSD License](#).
- Source code on [GitHub](#)
- Recommended in RDMKit
- Widely adopted
- **A repository and a registry**

F
indable



A
ccessible



I
nteroperable



R
eusable

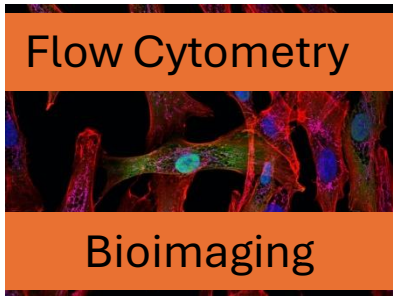
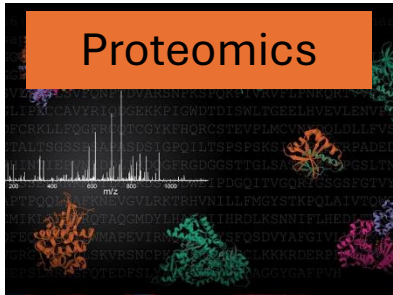
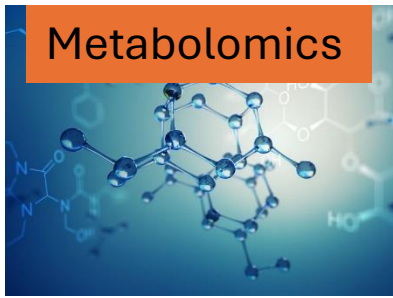


Reuse metadata
templates
Guided by ACDC
Helped by AI agents

Infrastructure provides!

Decide who can see
your data and in what
circumstances

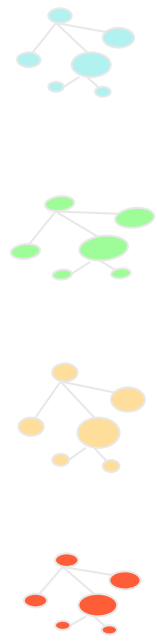
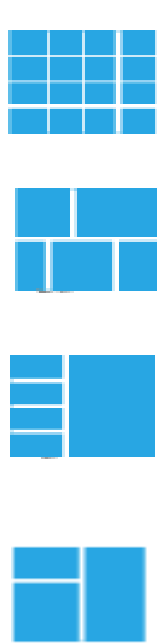
Link data and analyses
Licenses
Public repositories



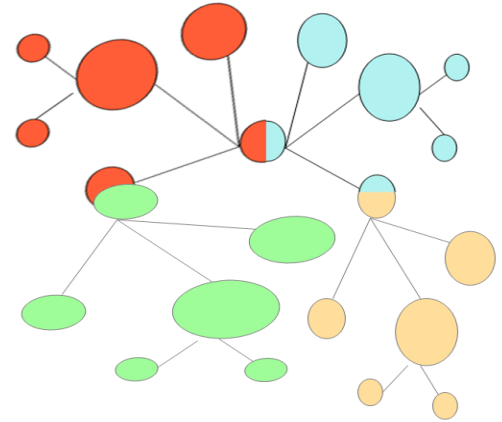
Catalogue of Findable Research Assets



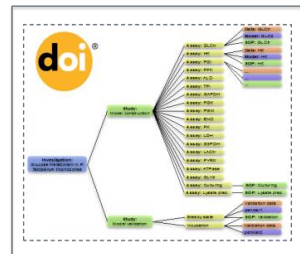
Accessibility Permissions and Tiering



Interoperable research assets and results. RDF FAIR Data Point

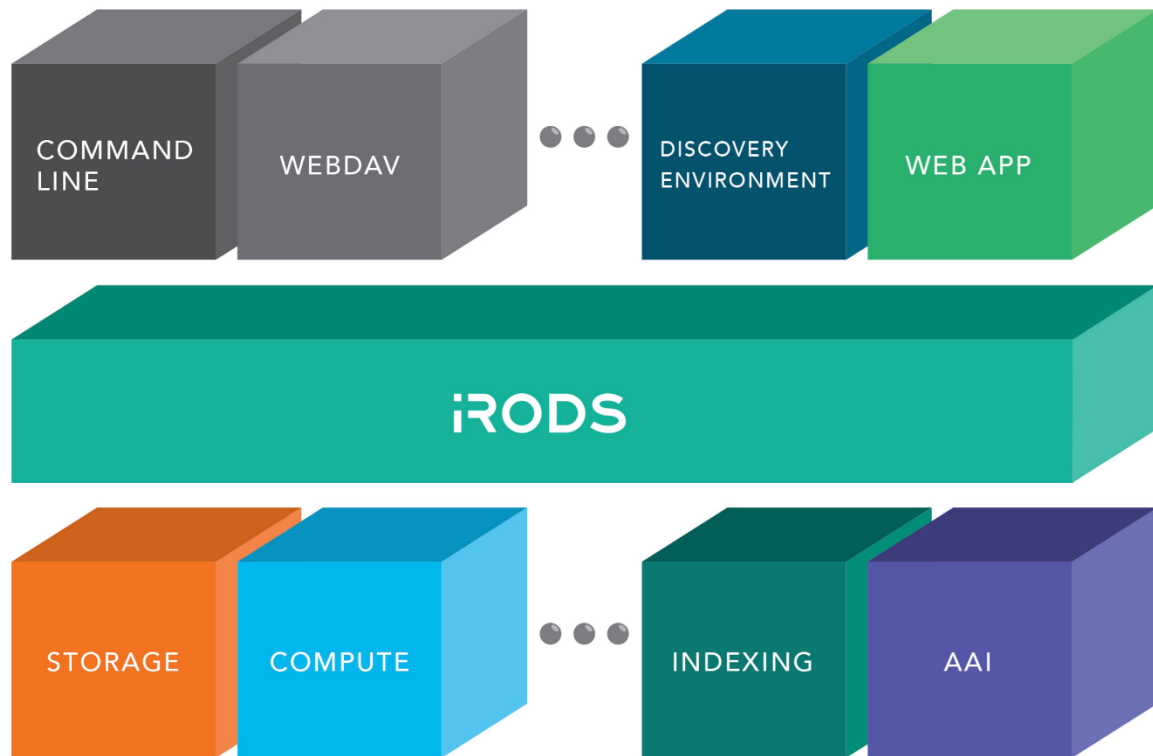


Reuse: Package, integrate and publish and archive



Reuse of Data Infrastructure

iRODS: Integrated Rule Oriented Data System



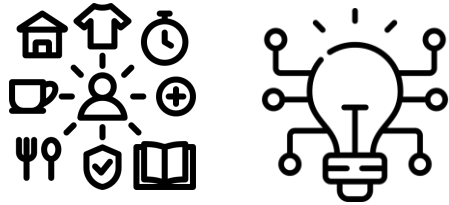
Open-Source and widely adopted

- **Data Virtualisation** – automated ingest, backup and tiering
- **Data discovery** – simple metadata associated with data objects
- **Workflow automation** – rules engine determines actions triggered
- **Secure collaboration** – permissions and federation

High Performance Compute Core



- Granted innovation budget by board of directors
 - *"Impuls Digitalisering en Technologische Innovatie"*



- Pre-funding to:
 - Invest in base HPC infrastructure
 - Respond to user needs
 - Respond to innovations in the field



- Privacy, security & compliancy
 - To support both research & diagnostics

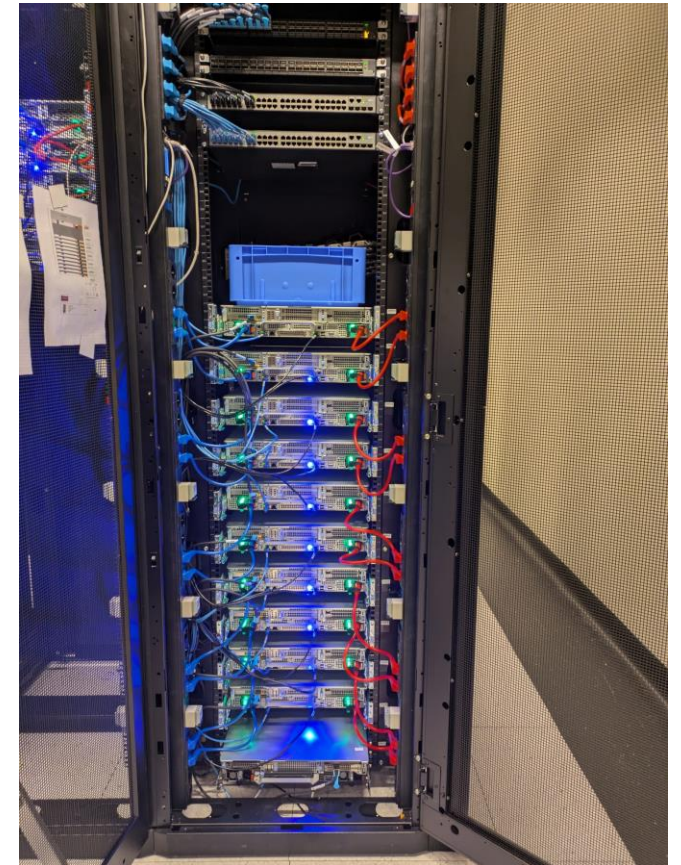


- Sustainable
 - Consolidate siloed HPC resources



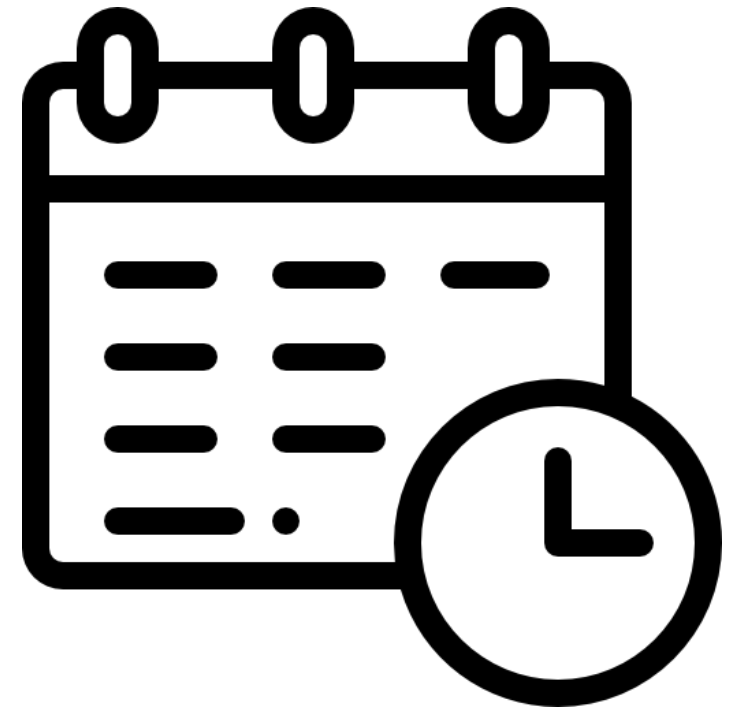
Boelelaan MedFac

Meibergdreef Datacenter



Planning

- **31th of March 2025** : Release new HPC cluster
 - Validated with diagnostic pipelines
 - Validated with standardised pipelines
 - Start migration of users
 - Consolidation of siloed HPC Resources



Support & admin team:

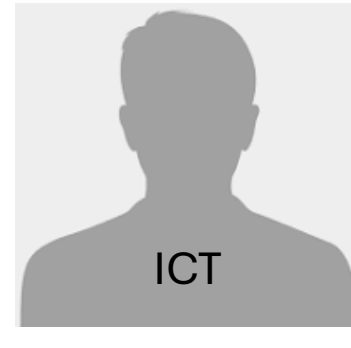
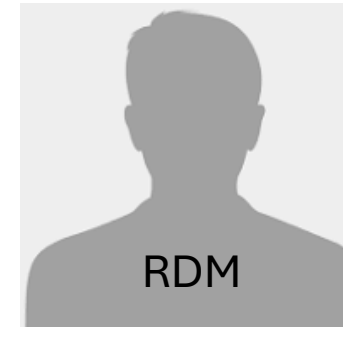
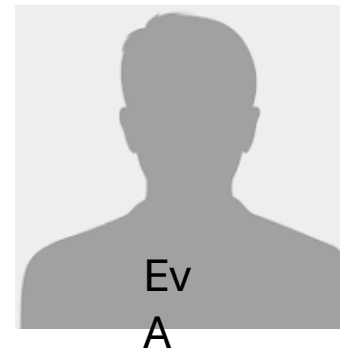
- Collaboration of many departments:
 - **ACDC**
 - Human Genetics
 - Radiology & Nuclear Medicine
 - ICT
 - Research Support / RDM
 - Zorg Support (EvA / ICT)
- To provide:
 - User group meetings
 - Updates & feedback (bi-directional)
 - MS Teams page
 - Training & education



Humane Genetica

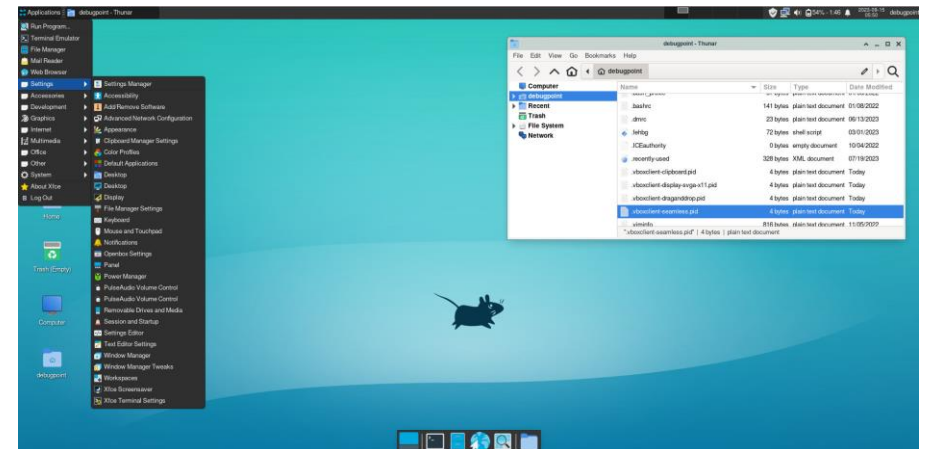
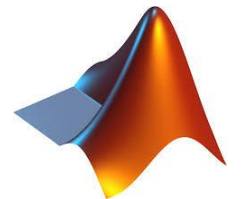


Radiologie en Nucleaire Geneeskunde



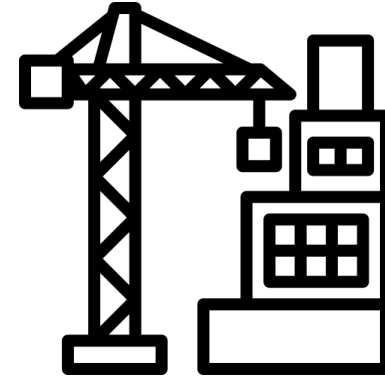
Applications

- Terminal interface
 - MobaXterm (SSH client with X server)
- Desktop interface
 - Remote XFCE desktop (RDP)
- Web interface
 - OpenOnDemand
 - VisualStudio Code / Pycharm (websockets)
 - Matlab (webVNC)
 - RStudio / Posit (websockets)
 - Jupyter notebook (websockets)
 - Remote XFCE desktop (webVNC)

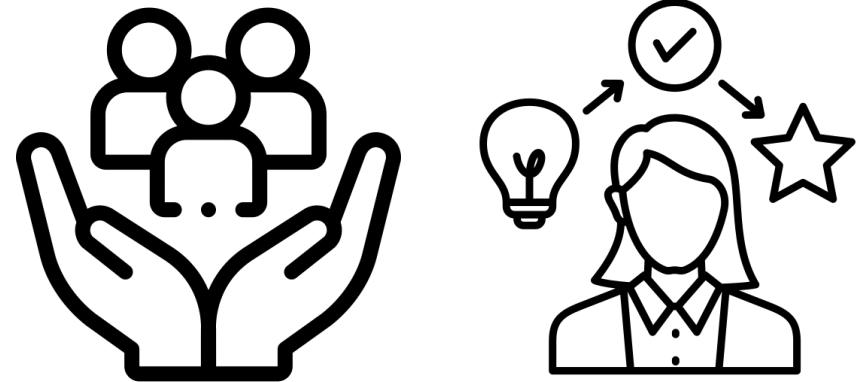


In summary

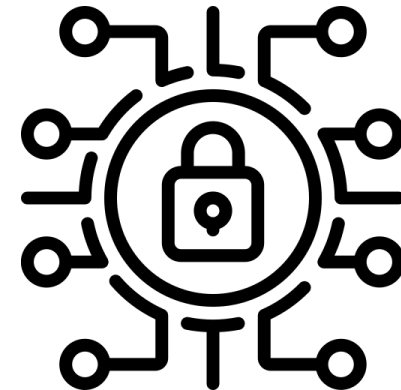
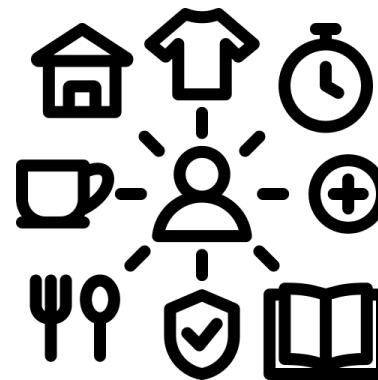
- High Performance Compute Infrastructure
- Data Management Infrastructure



- Expertise
- Support community



- Applications – user interface
- Safety & security
- Privacy & compliancy



Today's speakers





**BiImaging, Genomics,
Proteomics
Przemek Krawczyk**

**DMP - Elize Vlainic RDM
Grant Support – Laura Manca
All CFs**

**Proteomics, Metabolomics,
BiImaging, RDM
Karen Sap**

**Metabolomics, Proteomics,
Bioimaging
Karen Sap & Przemek Krawczyk**

**Genomics, Flow Cytometry
Alberto Miranda Bedate
RDM, Privacy**

**Genomics and Flow Cytometry
Bilgehan Nevruz and Giulia
Bergamaschi**

**Genomics, Proteomics and Flow
Cytometry: Bilgenhan Nevruz,
Yongsoo Kim, Alex Henneman &
Giulia Bergamaschi**