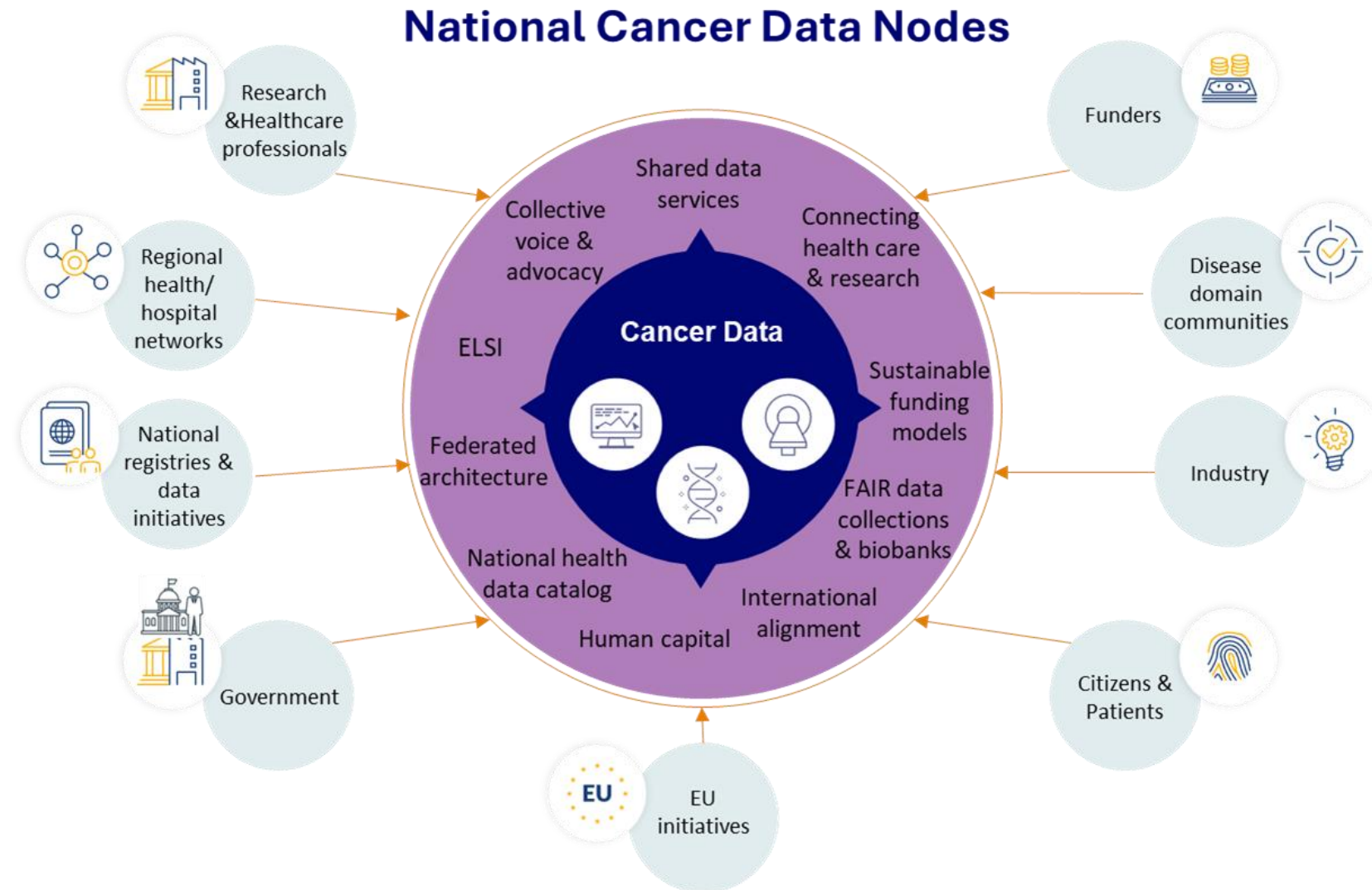




Dutch National Cancer Data Node (NCDN-NL)

2025-12-08

A warm welcome to the whole system in this room

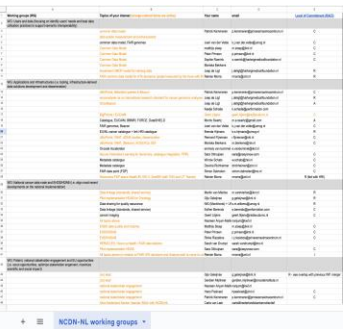


EOSC4Cancer and Node meetings

2021-11-02 (Heidag)
2022
2023-09-17
2024-04-12
2024-July-September
CANDLE proposal prep

Node WGs established

2024-10-18



Working group	Chair	Members	Start date	End date
Common data model	Dr. [Name]	Dr. [Name], Dr. [Name], Dr. [Name]	2024-10-18	2025-12-08
Data linkage	Dr. [Name]	Dr. [Name], Dr. [Name], Dr. [Name]	2024-10-18	2025-12-08
cBioportal	Dr. [Name]	Dr. [Name], Dr. [Name], Dr. [Name]	2024-10-18	2025-12-08
Genomic data	Dr. [Name]	Dr. [Name], Dr. [Name], Dr. [Name]	2024-10-18	2025-12-08
ESFRI-EHDS	Dr. [Name]	Dr. [Name], Dr. [Name], Dr. [Name]	2024-10-18	2025-12-08

CANDLE granted

2024-12-17
(9:46am CET)

Node meetings, WG meetings & CANDLE kick off

2025-01-31
2025-06-17
2025-06-24
2025-July-Dec
(Common data model,
data linkage, cBioportal
symposium, share
genomic data, EOSC-
ESFRI-EHDS)
2025-12-08

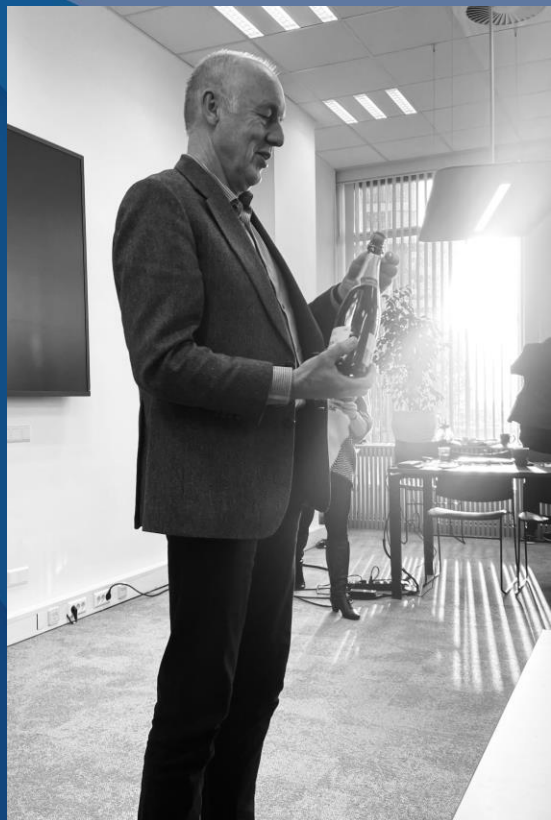
NCDN-NL=EU best practices
acknowledged by EC and MS

- 2025-11-27~28
- eCancer thematic working group meeting with 25 MS and three DGs: (EHDS blueprint DC, bring EU initiative together on national level)
 - Cyprus EU presidency event, 2026-04-23, NCDN roadshow
 - EC cancer mission projects support 2026-09-29~10-01



Symposium: cBioPortal - Making cancer data accessible for all

Tuesday 11 November 2025
09:00 - 17:00
Netherlands Cancer Institute
Symposium



Go together: go faster and further

Guiding principles

- EHDS blueprint and realization of functions of NCDNs
- 'Whole system in the room' =user-driven
- *Productize* functions tailored for users' needs

EU opportunities

- CANDLE
- ECHoS collaboration
- EUnetCCC
- JA-PCM
- New EU funding opportunities

Events

- Cyprus Presidential event
- EU-funded project NCDN supports
- CANDLE related events
- NCDN meetings, WGs

CANDLE



NCDN-NL

New year,
fresh start,
endless
possibilities.
2026
Make 2026
your masterpiece!

National Cancer Data Nodes

National Cancer Data Nodes (NCDNs) are federated hubs, responsible for standardising and harmonising data procedures within their country/region, and providing the digital tools for research collaboration – all aligned with national EHDS roles. These nodes will link to the UNCAN.eu platform (built by UNCAN-Connect). They are crucial to consider national conditions when building a European cancer data infrastructure. During the project, CANDLE will further conceptualise the nodes and their users.



Austria



Belgium



Cyprus

Czech
Republic

Denmark



Finland



France



Germany



Greece



Ireland



Italy



Lithuania



Luxembourg



Malta



Netherlands



Norway



Poland



Portugal



Slovenia



Spain

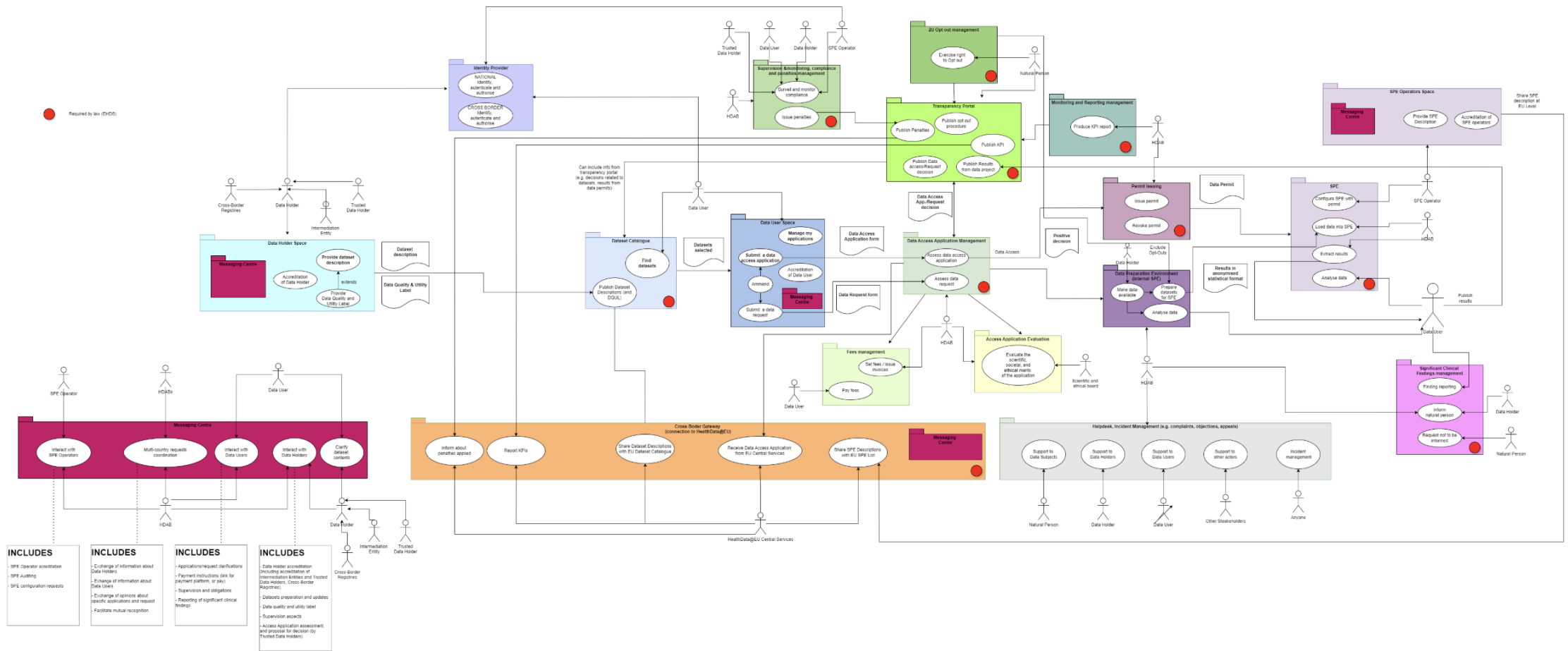


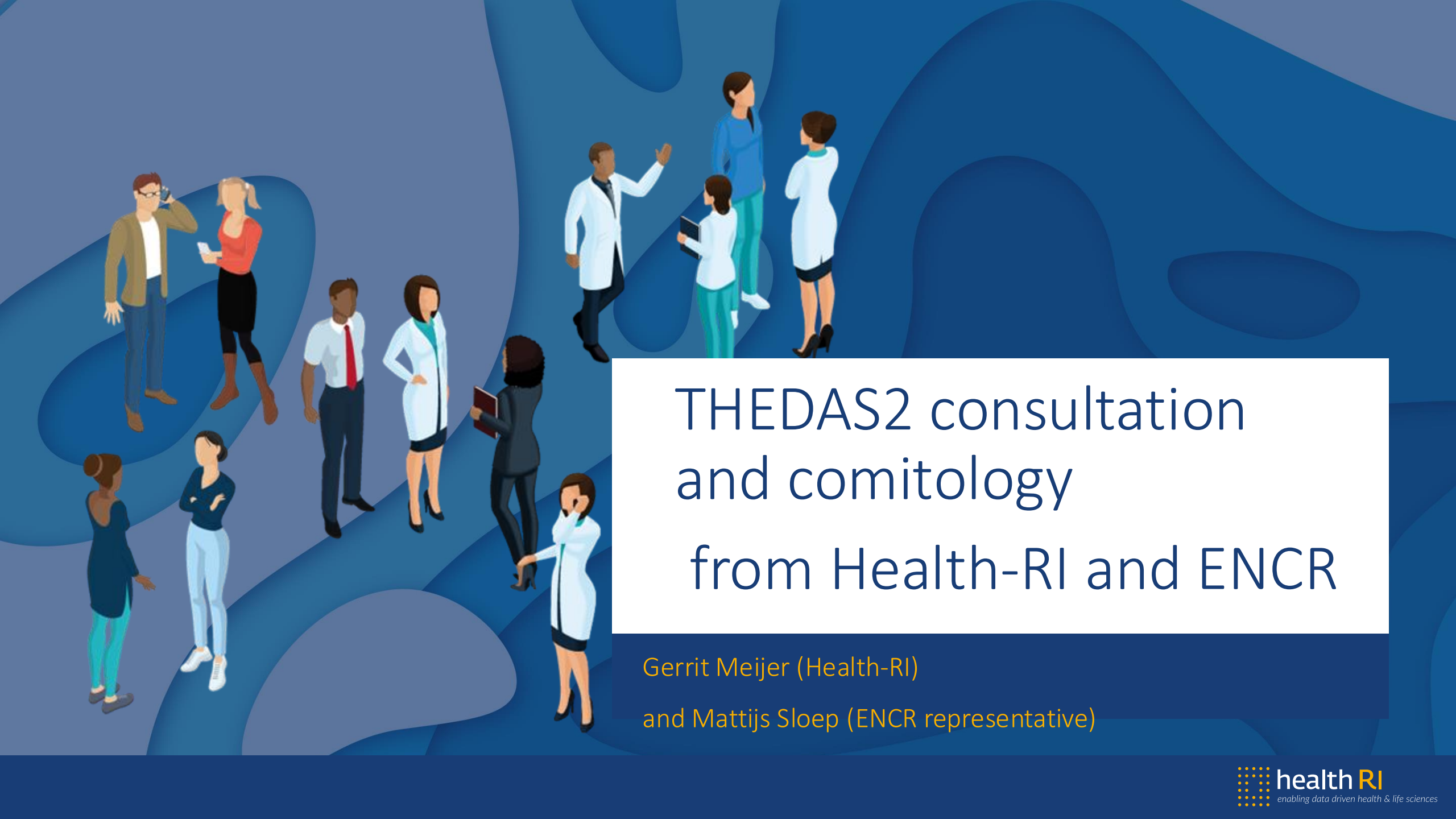
Sweden



Switzerland

Backbone





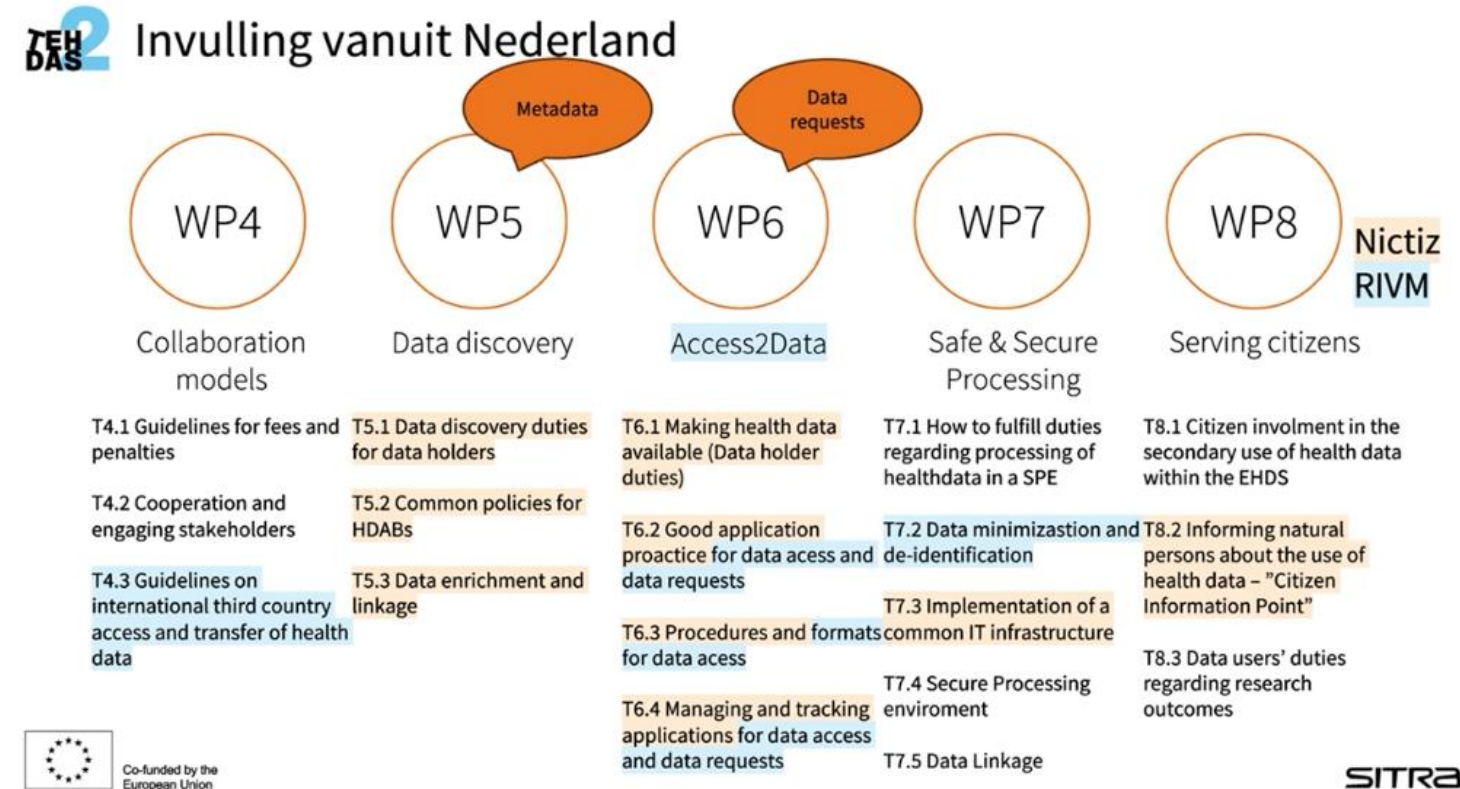
THEDAS2 consultation and comitology from Health-RI and ENCR

Gerrit Meijer (Health-RI)
and Mattijs Sloep (ENCR representative)



TEHDAS2 basis voor uitvoeringshandelingen EHDS

- Europees project
- 30 Europese landen
- NL: Nictiz & RIVM, Health-RI nauw betrokken
- Doel: EC adviseren over guidelines (niet verplichtend) en technische specificaties (wel verplichtend) voor secundair gebruik van gezondheidsdata



Comitologieproces

Implementing Acts under the EHDS

- Lidstaten en Europese Commissie overleggen over lagere wetgeving EHDS (uitvoeringshandelingen):
- Nederland heeft maar 3-4% van de stemmen in Europa en sluit daarom bondjes met landen (Duitsland, Estland, Finland oa.), Nederland heeft keuzes gemaakt op welke onderwerpen (extra) ingezet wordt (in vet)
- Elke uitvoeringshandeling 3x in vergadering (totale doorloop 4 – 10 mnd)
 - 1^e vergadering: discussiestuk EC
 - Initiële standpuntbepaling
 - Herijking obv discussiestuk
 - Schriftelijk commentaar
 - 2^e vergadering: non-paper EC
 - Herijking obv non-paper
 - Schriftelijk commentaar
 - 3^e vergadering: stemming
- Nederlandse inbreng voorbereid door “Tafel Europa” van VWS: VWS DICIO, NICTIZ, RIVM, NEN, ZIN, IGJ en Health-RI

Health-RI goed betrokken in hele proces

- Sneak-preview van TEHDAS2 guidelines, voordat deze gepubliceerd worden
- Leveren van input bij publieke consultaties
- Motiveren van netwerk om actieve input te leveren bij publieke consultatie
- Gesprekken met RIVM en NICTIZ naar aanleiding van NL input
- Participatie in diverse TEHDAS2 workshops
- Participatie in “Tafel Europa” VWS voor comitologie
- NL specifiek: juridische sessies



ENCR - the European Network of Cancer Registries

Matthijs Sloep - Tehdas2 public consultations

The ENCR Network



ENCR Recommendations

2024

Recommendations for coding tumours of
the central nervous system (CNS)



ENCR Recommendations

2025

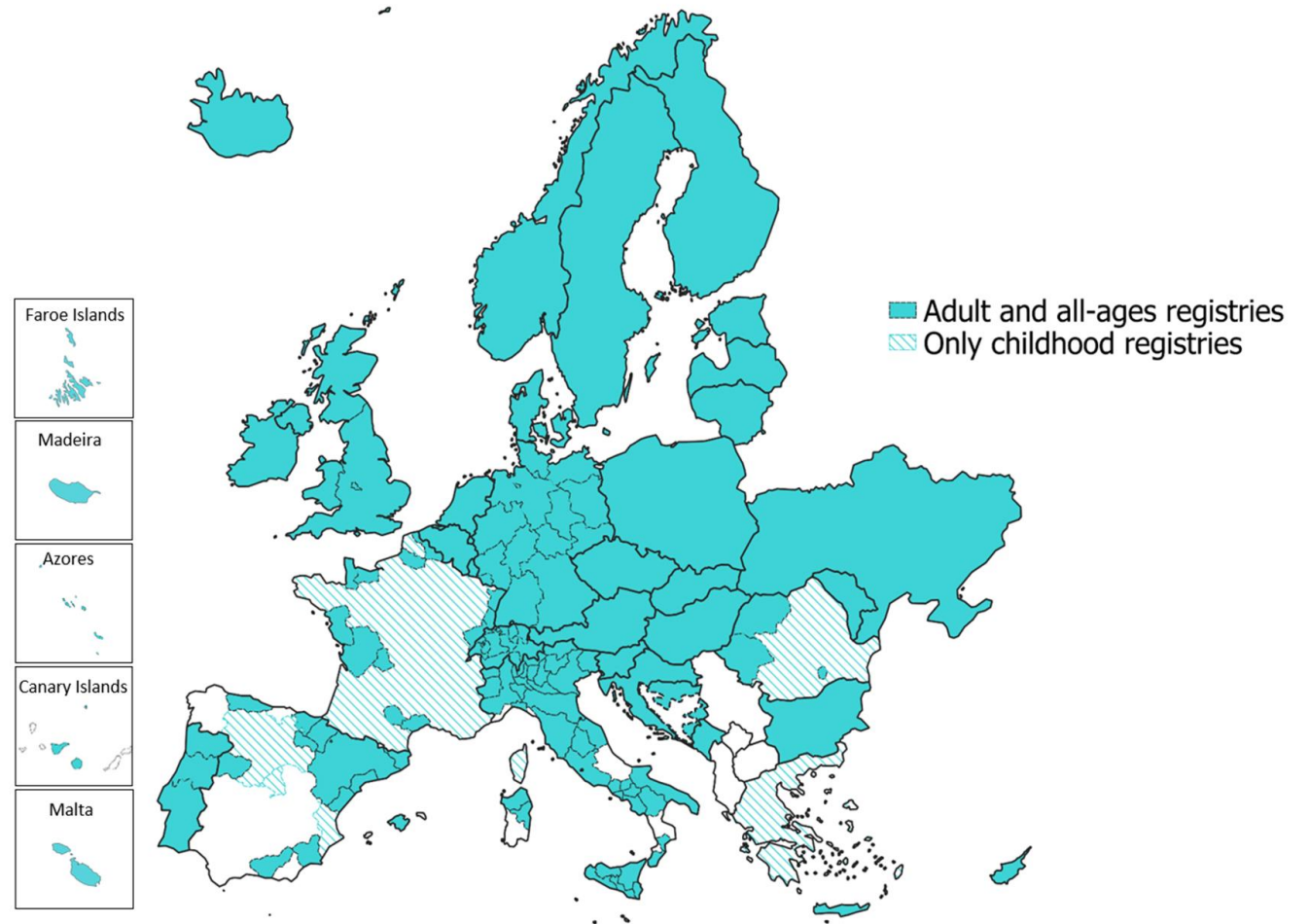
Recording Recurrence, Progression and
Transformation Episodes



ENCR Recommendations

2025

Treatment Data Recording (phase I)



European Network
of Cancer Registries



European
Commission

CancerWatch Joint action



JA CancerWatch

Improving Quality and Timeliness of Cancer Registry Data
feeding into the European Cancer Information System

EC contribution: € 13.000.000

Duration: Sept 2025 – Aug 2028 (36 months)

Project Lead: Norwegian Institute of Public Health

Number of beneficiaries: 92

Who are the beneficiaries: Cancer Registries, Public Health
Organizations, Ministries of Health, Cancer Centers.

IARC and JRC partake in the project



Tehdas2 - Second public consultation



Project  [Public consultations](#) [News](#) [Events](#) [Get involved](#) [Results](#)

[Home](#) / [News](#) / [Second public consultation on guidelines for effective secondary use of health data across Europe concludes](#)

01.12.2025

Second public consultation on guidelines for effective secondary use of health data across Europe concludes

Between 30 September and 30 November 2025, TEHDAS2 invited stakeholders from across Europe to provide feedback on eleven draft guidelines and technical specifications that will shape the implementation of the European Health Data Space (EHDS). The consultation period has now closed, with all feedback gathered for review.

 **1. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON FEES RELATED TO THE EHDS REGULATION** ([Click to view](#))

 **2. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON PENALTIES FOR NON-COMPLIANCE RELATED TO THE EHDS REGULATION** ([Click to view](#))

 **3. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON MINIMUM CATEGORIES AND LIMITATIONS ON THE REUSE OF HEALTH DATA** ([Click to view](#))

 **4. DRAFT GUIDELINE FOR DATA HOLDERS ON MAKING PERSONAL AND NON-PERSONAL ELECTRONIC HEALTH DATA AVAILABLE FOR REUSE** ([Click to view](#))

 **5. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON THE PROCEDURES AND FORMATS FOR DATA ACCESS** ([Click to view](#))

 **6. DATA ACCESS APPLICATION MANAGEMENT SYSTEM (DAAMS) – DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES** ([Click to view](#))

 **7. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON DATA MINIMISATION, PSEUDONYMISATION, ANONYMISATION AND SYNTHETIC DATA** ([Click to view](#))

 **8. DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES ON THE IMPLEMENTATION OF THE COMMON IT INFRASTRUCTURE** ([Click to view](#))

 **9. DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES ON THE IMPLEMENTATION OF SECURE PROCESSING ENVIRONMENTS** ([Click to view](#))

 **10. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON IMPLEMENTING OPT-OUT FROM THE SECONDARY USE OF HEALTH DATA** ([Click to view](#))

 **11. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON IMPLEMENTING THE OBLIGATION OF NOTIFYING THE NATURAL PERSON ON A SIGNIFICANT FINDING FROM THE SECONDARY USE OF HEALTH DATA** ([Click to view](#))

Tehdas2 - Second public consultation

1. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON FEES RELATED TO THE EHDS REGULATION (Click to view)

2. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON PENALTIES FOR NON-COMPLIANCE RELATED TO THE EHDS REGULATION (Click to view)

3. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON MINIMUM CATEGORIES AND LIMITATIONS ON THE REUSE OF HEALTH DATA (Click to view)

- Expand billable tasks
- No. Patients
- Different pricing, invoices
- Cross border requests
- Ethical framework

Tehdas2 - Second public consultation

4. DRAFT GUIDELINE FOR DATA HOLDERS ON MAKING PERSONAL AND NON-PERSONAL ELECTRONIC HEALTH DATA AVAILABLE FOR REUSE (Click to view)

5. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON THE PROCEDURES AND FORMATS FOR DATA ACCESS (Click to view)

6. DATA ACCESS APPLICATION MANAGEMENT SYSTEM (DAAMS) – DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES (Click to view)

7. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON DATA MINIMISATION, PSEUDONYMISATION, ANONYMISATION AND SYNTHETIC DATA (Click to view)

- Guidance
- Fast track
- Reidentification
- Small data holders

Tehdas2 - Second public consultation



8. DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES ON THE IMPLEMENTATION OF THE COMMON IT INFRASTRUCTURE ([Click to view](#))



9. DRAFT TECHNICAL SPECIFICATION FOR HEALTH DATA ACCESS BODIES ON THE IMPLEMENTATION OF SECURE PROCESSING ENVIRONMENTS ([Click to view](#))



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11. DRAFT GUIDELINE FOR HEALTH DATA ACCESS BODIES ON IMPLEMENTING THE OBLIGATION OF NOTIFYING THE NATURAL PERSON ON A SIGNIFICANT FINDING FROM THE SECONDARY USE OF HEALTH DATA ([Click to view](#))

- CDM guidance
- EHDS/GDPR
- Opt out bias
- Significant findings

First public consultation results



[Project](#)  [Public consultations](#) [News](#) [Events](#) [Get involved](#) [Results](#)

Results

TEHDAS2 is developing guidelines and technical specifications to enable seamless use of health data across Europe under the upcoming European Health Data Space (EHDS). The results of this work are organised by target groups and shared here.





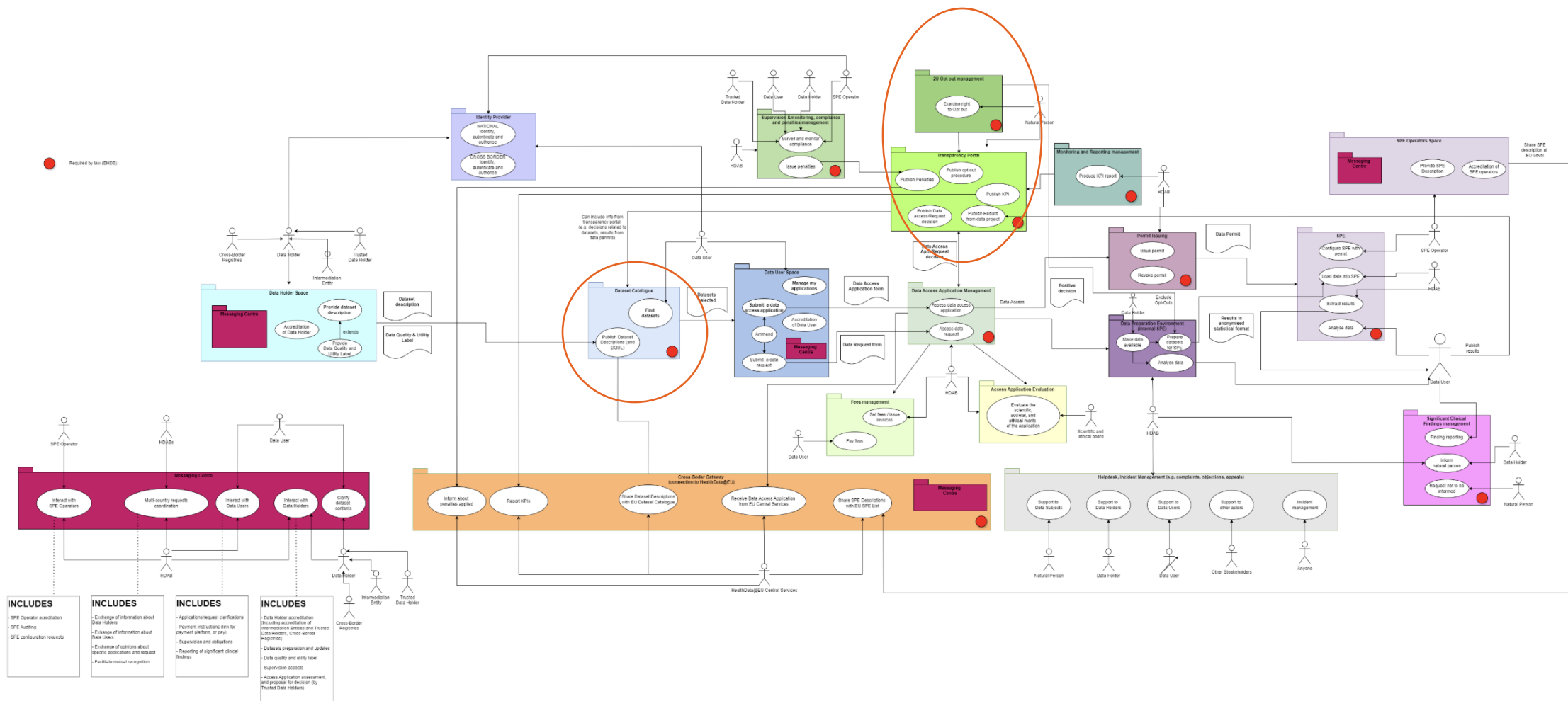
HERACLES's concluding messages:

- the oncology catalogue
- infrastructure from a FAIR and privacy perspective, aligned with the work of HDAB, EHDS, and Health-RI

Sarah van Drumpt (TNO)

Erik Cornelisse (TNO)

Backbone





A privacy preserving dataspace for cancer research
HERACLES and Dataspaces

Objective and content

This presentation introduces the HERACLES project and explains how it relates to European and National initiatives.

Table of content:

- General introduction to HERACLES and its purpose
- The scope and description of the HERACLES infrastructure
- The architecture and how data space standards are applied
- How the HERACLES infrastructure can contribute to Health-RI & HDAB



Introduction on why HERACLES

Complication

- **Health related data is fragmented** across various systems in different formats and cannot always be used in a **privacy-friendly way** and **without medical confidentiality concerns**, hence hindering data use for innovation, research and policy making.

Question

- How to come to a '**learning health system**' based on privacy-friendly use of data, where insights gained directly or through scientific research contribute to improving healthcare?

Answer

- The answer lies in a **combination of governance** and **supporting technology** where sensitive data stored at different sources are used for science, innovation and policy-making without compromising privacy.



Impact

Quality of Diagnosis

Quality of Treatment

Cost reduction in
healthcare

Trustworthy ecosystem

Actors

Pharma

Hospital

Infrastructure
service provider

Health
insurance

National
Fund raiser

Patient
organizations

Netherlands Cancer
Organization

Research
Institutes

Applications

Ovarium cancer
insight counter

Lung cancer
insight counter

Datasets

IKNL
NKR



Pharmo
INSZO-STIZON



RadboudUMC
EPD



UMCG



CZ

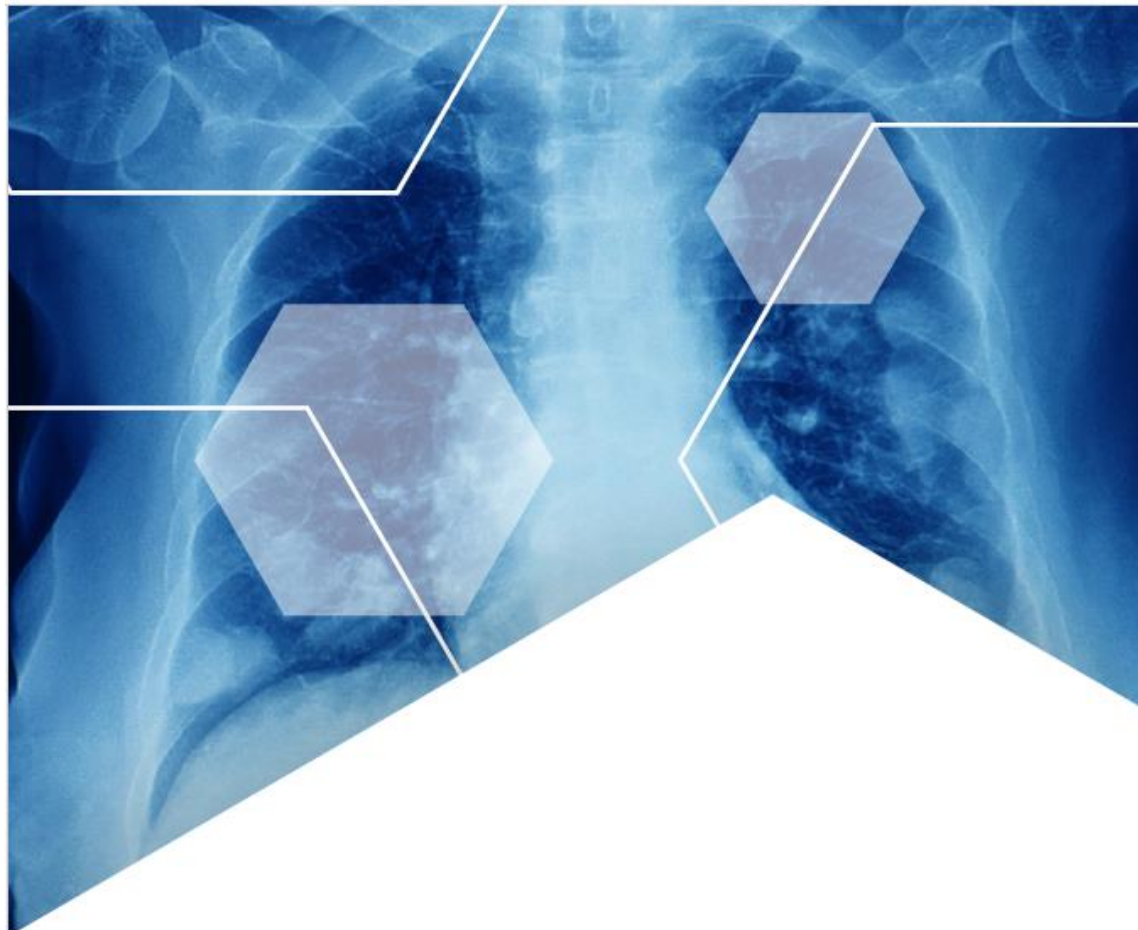


Infrastructure

Health Data Space

Data Sharing Meta model, including business models and governance structure

Secure Gateway, identity provisioning, clearing house, registry/broker
servers, computing, OS, storage, network



Van mythe naar realiteit de hobbelige weg naar een lerend gezondheidssysteem

Lessen van het HERACLES project uit de praktijk

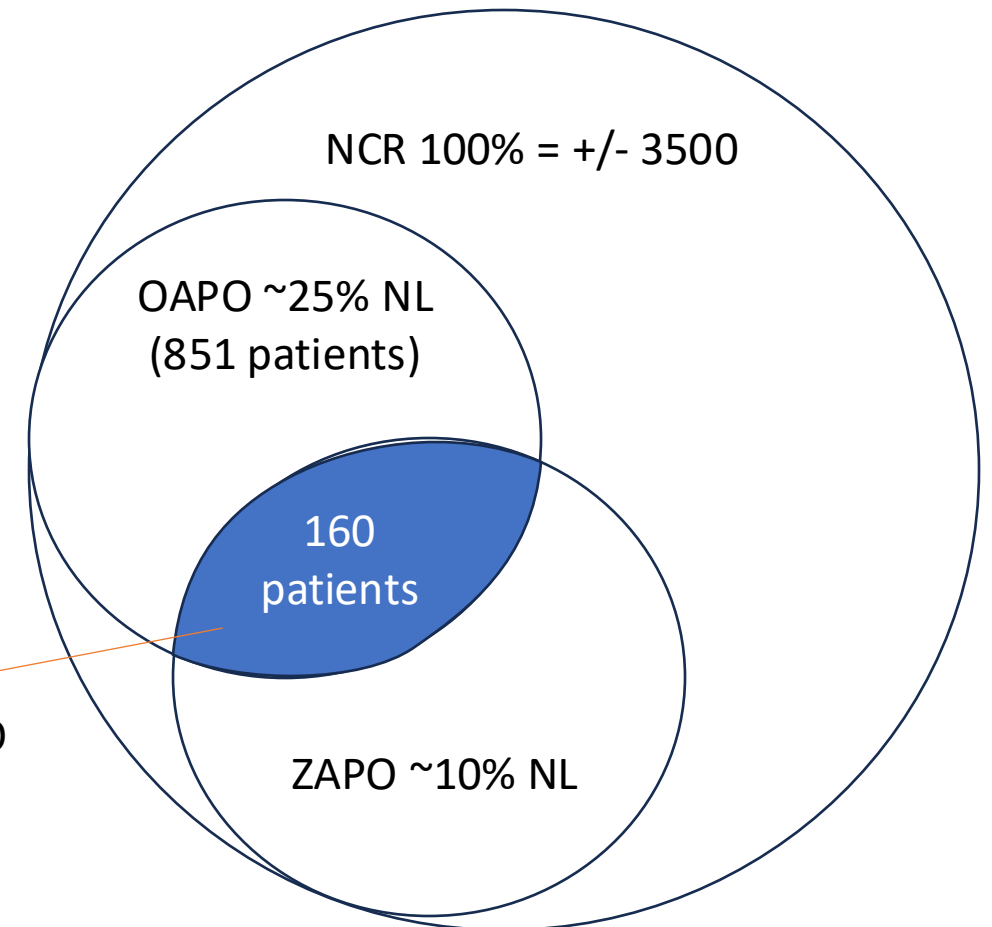
Results

- Number of patients:
 - NCR – OAPO 851 patients
 - NCR – OAPO – ZAPO 160 patients
 - NCR – OAPO – ZAPO: inclusion criteria 96 patients
- Additionally: 36 patients with no anti-cancer treatment (from Pharmo data)

15 no treatment

21 treatment registered NCR

NCR+OAPO+ZAPO

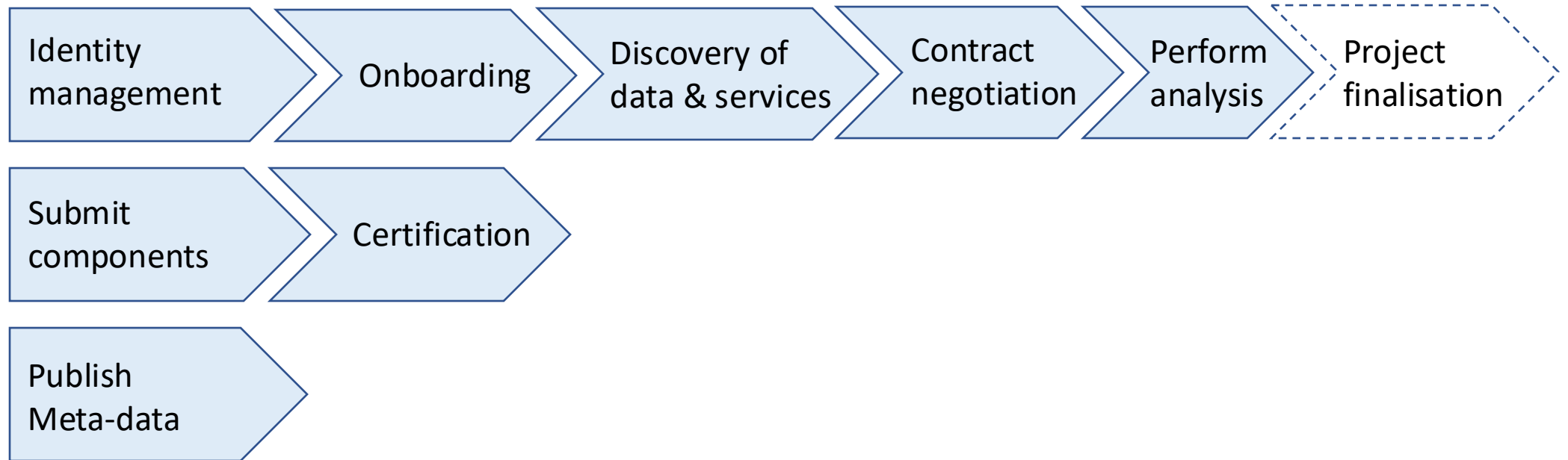


High level infrastructure process flow

EHDS

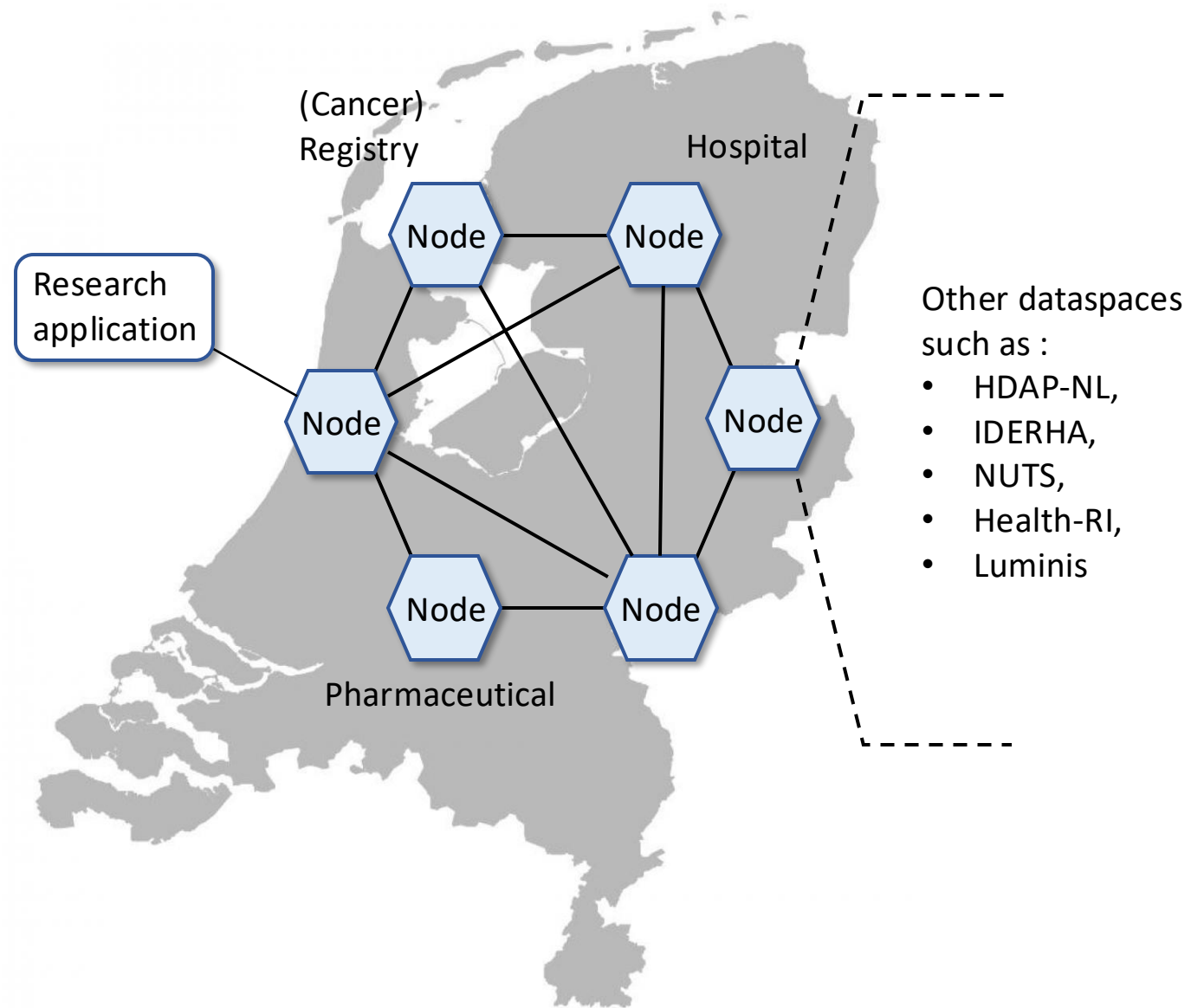


HERACLES

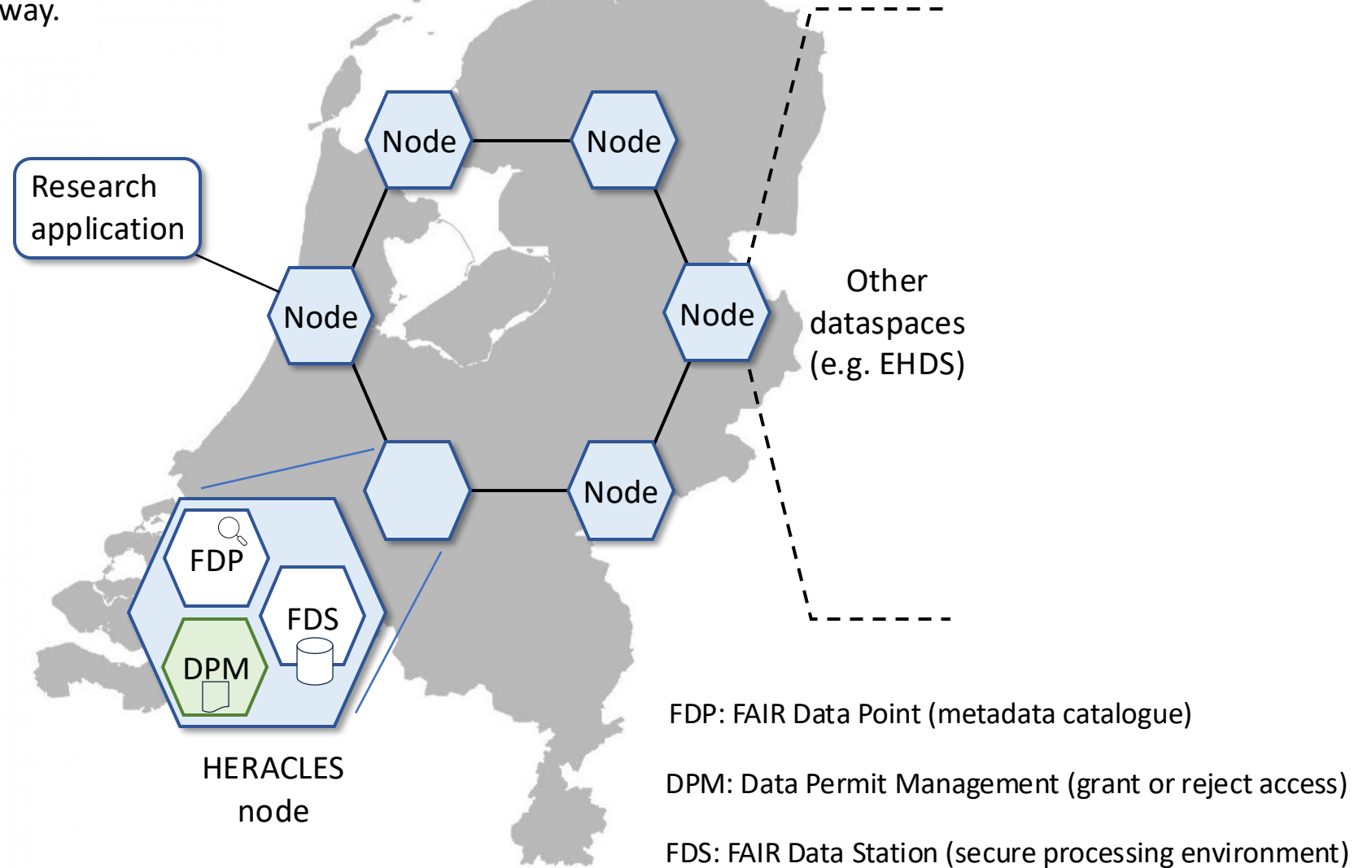




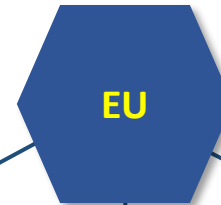
HERACLES



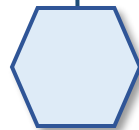
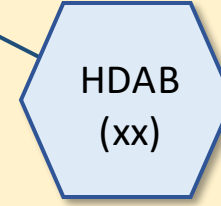
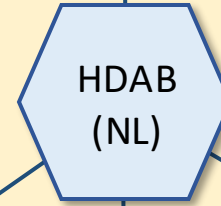
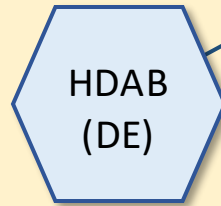
Components operate across nodes as a federation of systems in a harmonized and standardized way.



Centrale Europese node

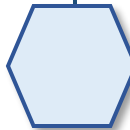


Centrale nationale node



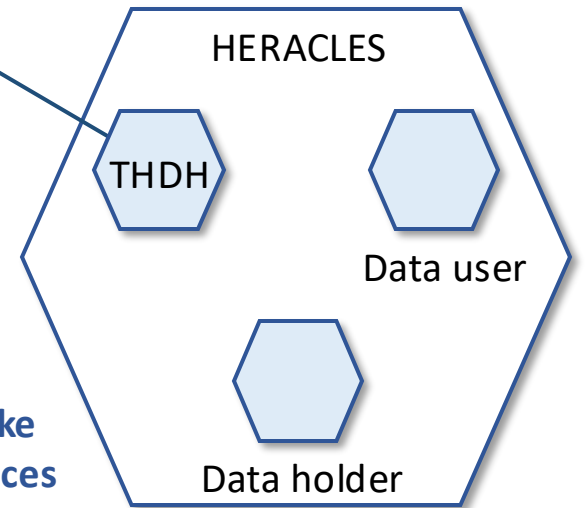
Data holder

Trusted Health
Data Holder



Data holder

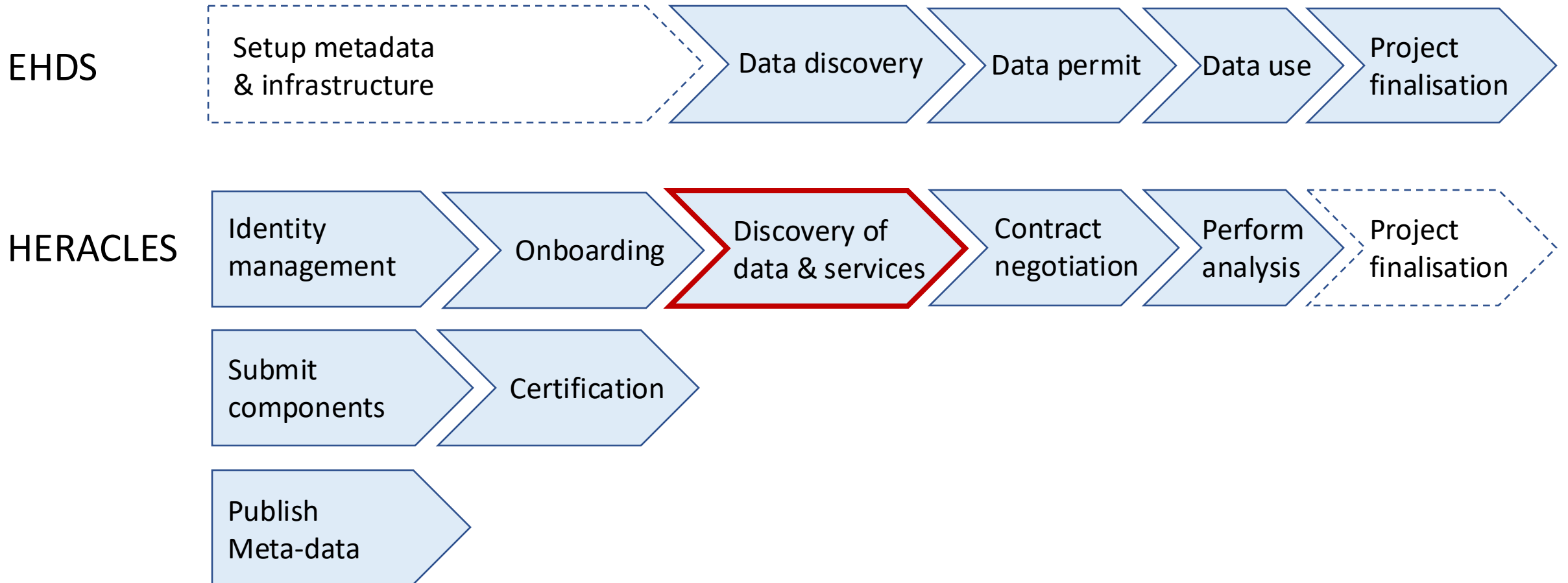
Topic
specifieke
dataspaces



HERACLES

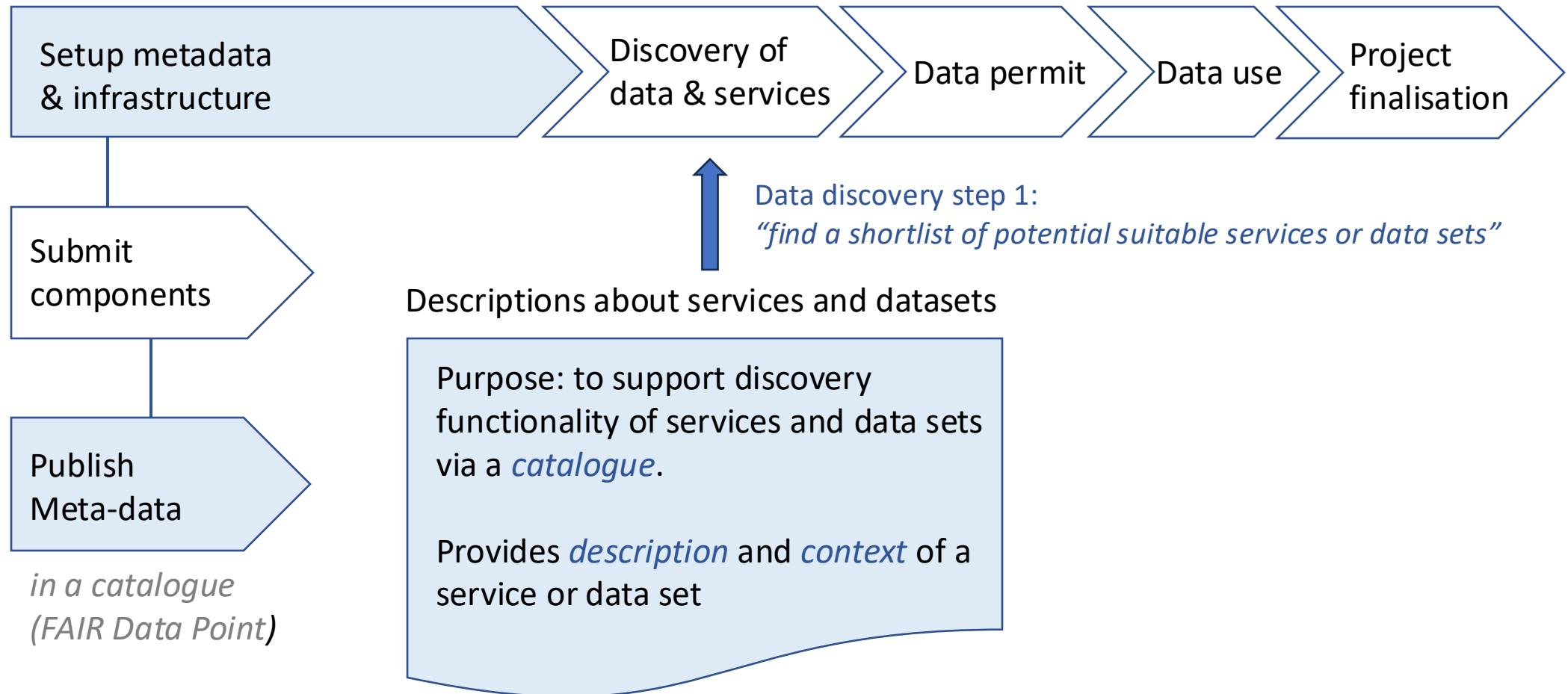
High level infrastructure process flow

Compliant to EHDS



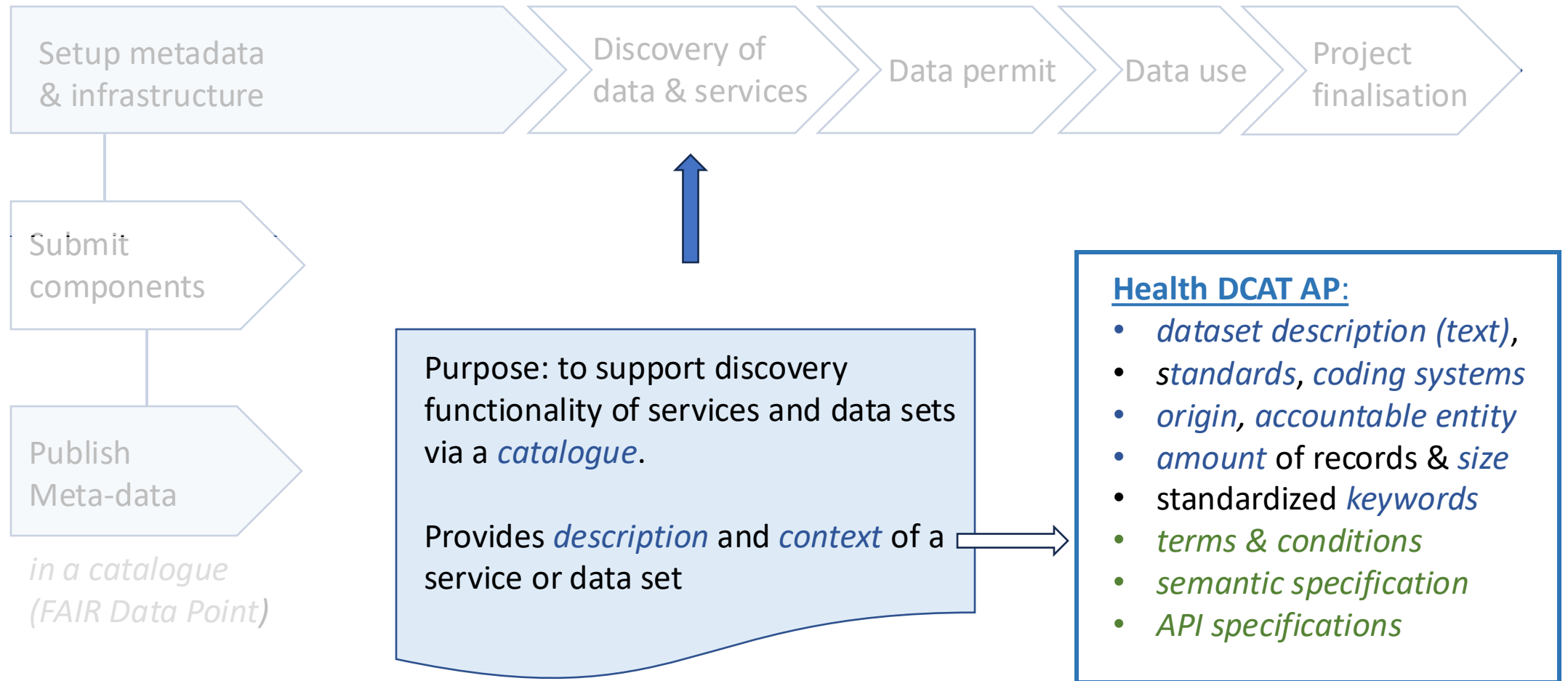
High level infrastructure process flow

Create & publish service and data descriptions (*step 0*)



High level infrastructure process flow

Create & publish service and data descriptions (*step 0*)



Discovery of data & services

Find suitable data & services via a FAIR Data Point (FDP)

Step 1 – “find a shortlist of potential suitable services and data sets”

Data set attributes:

- *dataset description (text),*
- *standards, coding systems*
- *origin, accountable entity*
- *amount of records & size*
- *standardized keywords*

Step 2 - “functional verification to determine which services and data are suitable based on information about the data”

Data attribute/reference:

- *Semantic/methodological specification (incl. units, measurement platform)*
- *quality per attribute:*
 - *% of missings*
 - *% of outliers ?*
 - *amount + std. deviation*

Data services (pre-train):

- *Provide sample data*
- *Overview of data distributions*

Step 3 - “define (sub)dataset request and test compliancy against terms and conditions, and also check technical feasibility.

Compliancy references:

- *terms & conditions (policies)*
- *semantic specification*
- *API specifications*
- *Resource specifications*

Compliancy (query) services:

- *Request “amount” information about specific selection of the dataset*

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- **Context -> QUANTUM label**

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Compliancy (query) services:

- *Request “amount” information about specific selection of the dataset*

HOME

 Fair Data Point Browse Search > [Catalog](#)

Pharmo

Pharmo connector

Datasets

[Treatment of primary and recurrent ovarian cancer](#) ★★☆☆☆

DCAT Dataset

Dataset for studying the patient journey of epithelial ovarian cancer (EOC) patients, focusing on systemic therapies, treatment patterns, predictive factors, and identification of recurrence/progression. Based on linked data from the Netherlands Cancer Registry (NCR) and the PHARMO Data Network, including pharmacy, laboratory, and hospital data.

[Lung Cancer Early Diagnosis Prediction Model Dataset](#)

DCAT Dataset

Dataset for developing a prediction model to identify individuals at risk of Small Cell Lung Cancer (SCLC) and Non-Small Cell Lung Cancer (NSCLC) based on clinical and patient characteristics from general practitioner records. Aimed at enabling earlier diagnosis and improving treatment outcomes through timely referral.

Conforms to

 [Catalog Shapes](#)

Modified

26-09-2025

Issued

26-09-2025

License

Publisher

did:web:pharmo.heracles.dataspac.es

Catalog identifier

<https://pharmo.heracles.dataspac.es/fdp/api/catalog/4c35f92a-89e3-40ac-82b4-28d8c588fb6a#identifier>

HOME

[Fair Data Point](#)
[Browse](#)
[Search](#)

Distributions

[TSG Federated Learning Data Plane](#)

DCAT Distribution

Protocol: tsq:FL

Table structure

Patientmatrix

852 records —

Columns

patient_id

Type: string

Description: Unique patient ID (PHARMO)

key_nkr_spec

Type: string

Description: Patient key number specific to this request

key_eid

Type: string

Description: Tumor key number

startfu

Type: date

Description: Start data collection of specific patient

endfu

Type: date

Description: Day-month-year end of follow-up in the PHARMO Data Network (death, end of study period [31-12-2021], or end of follow-up in PHARMO)

CED

Type: date

Description: Cohort entry date (i.e., first diagnosis of EOC)

oapo_avail

Type: integer

Description: Availability of out-patient pharmacy data (1=Yes)

zapo_avail

Type: integer

Description: Availability of in-patient pharmacy data (1=Yes)

kln_avail

Type: integer

Description: Availability of clinical laboratory data (1=Yes)

hosp_avail

Type: integer

Description: Availability of hospital data (1=Yes)

Keywords

Coding System

- [SNOMED CT](#)
- [ICD-10 ID](#)
- [WCIA](#)
- [ATC code](#)

Health Theme

- [cancer](#)
- [ovarian cancer](#)

Number of Records

138224

Number of Unique Individuals

851

Quality Certificate

- [Quantum label \(pdf\)](#)



HOME

 Fair Data Point Browse [Search](#)[Search](#)

Found 6 results.

[PharmaCure Solutions](#)

DCAT Catalog

PharmaCure Solutions connector

Issued 26-09-2025 **Modified** 26-09-2025<https://pharmacure.heracles.dataspace.es/fdp/api/catalog/b3c48b2f-89f2-46b9-9aae-73141b24109f>[Astra Zeneca](#)

DCAT Catalog

Astra Zeneca connector

Issued 26-09-2025 **Modified** 26-09-2025<https://astrazeneca.heracles.dataspace.es/fdp/api/catalog/0c45e2f7-7e8f-4b1c-b151-d9d2331ebb82>[DataGuard Innovations](#)

DCAT Catalog

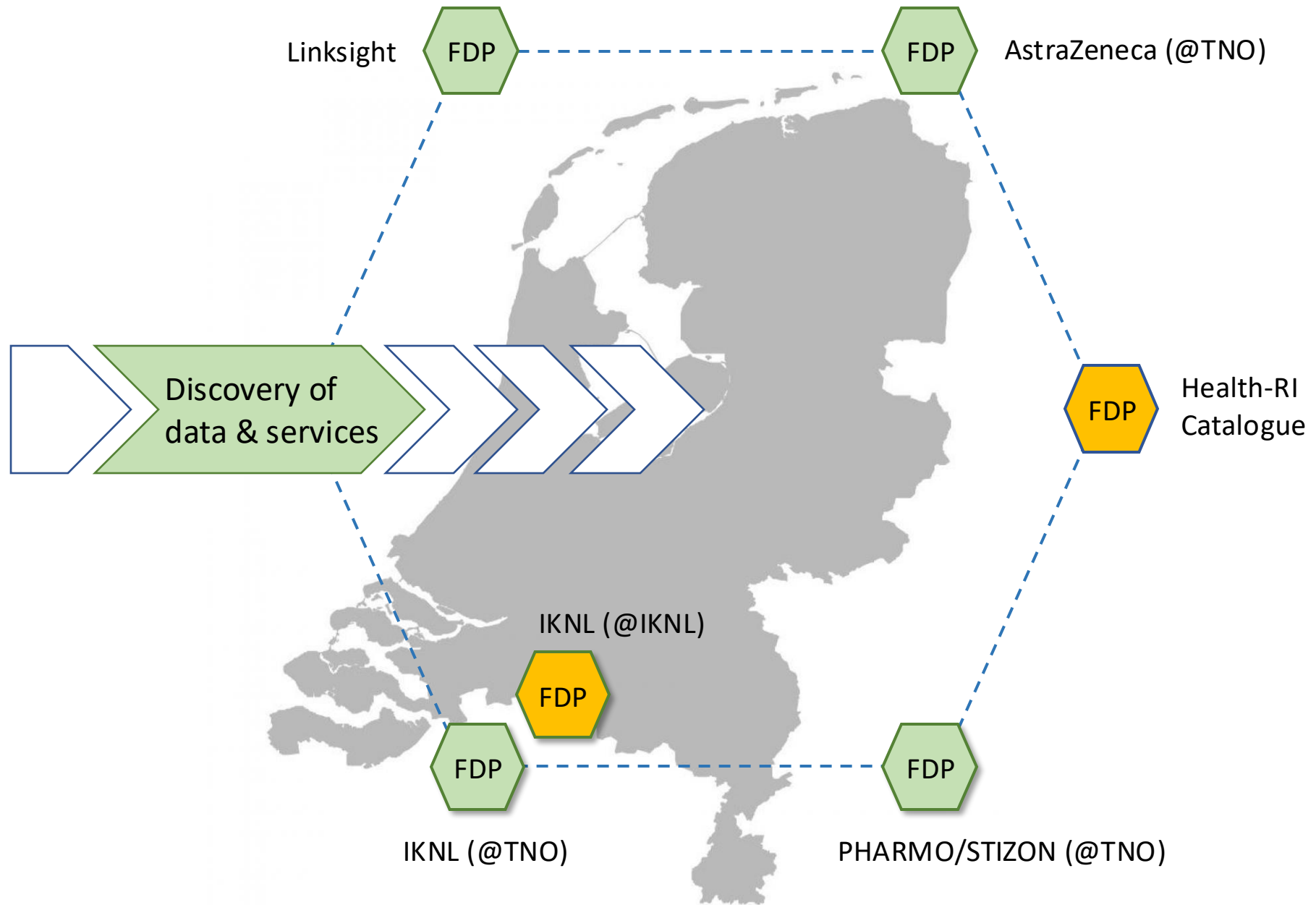
DataGuard Innovations connector

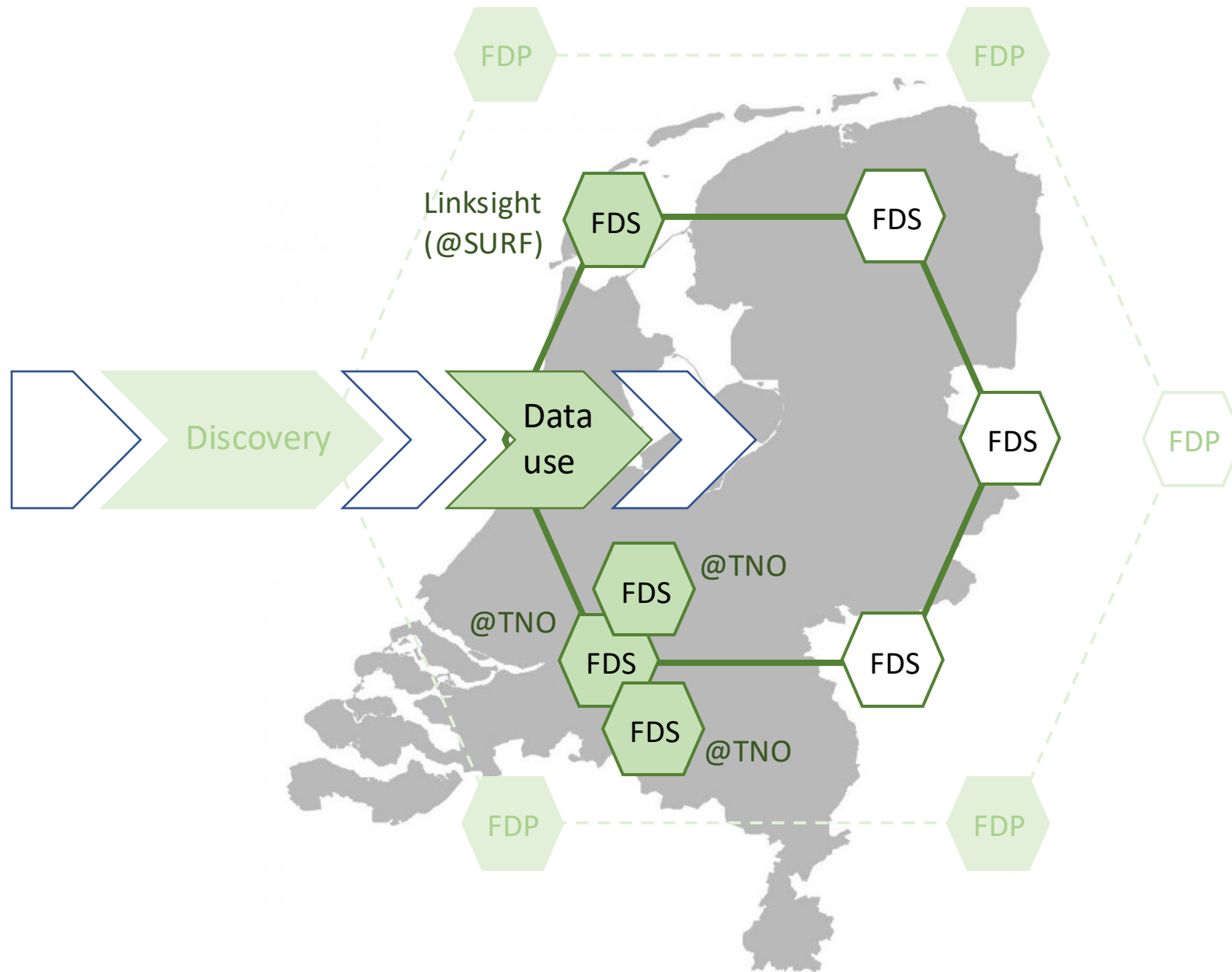
Issued 26-09-2025 **Modified** 26-09-2025<https://dataguard.heracles.dataspace.es/fdp/api/catalog/b1a14ba6-93ce-44d6-991c-fa160e7f2400>[IKNL NCR Synthetic Dataset](#)

DCAT Dataset

A synthetic dataset that mimics a part of the Netherlands Cancer Registry (NCR) is available for research purposes. This dataset does not contain data on real patients. It enables researchers to use

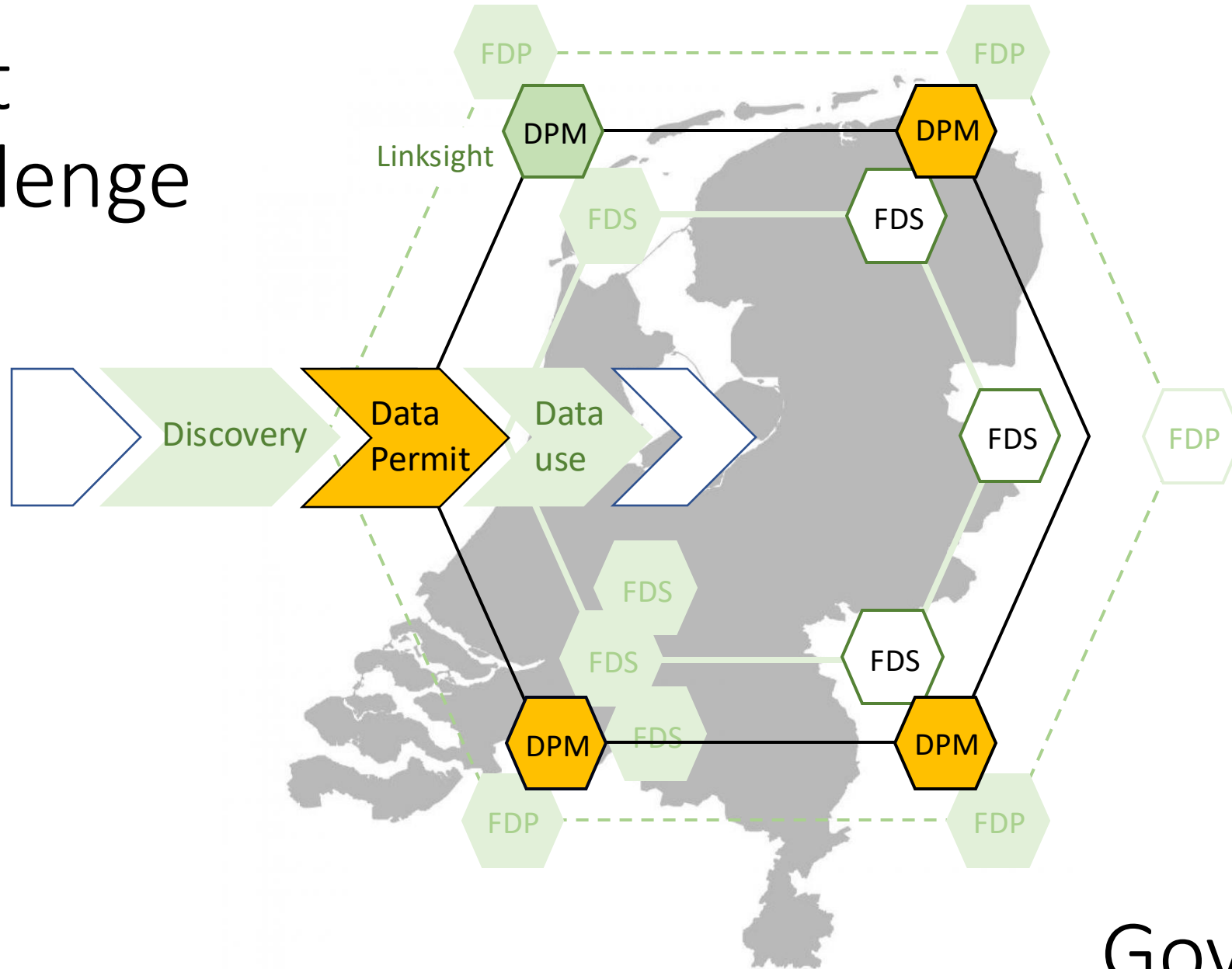






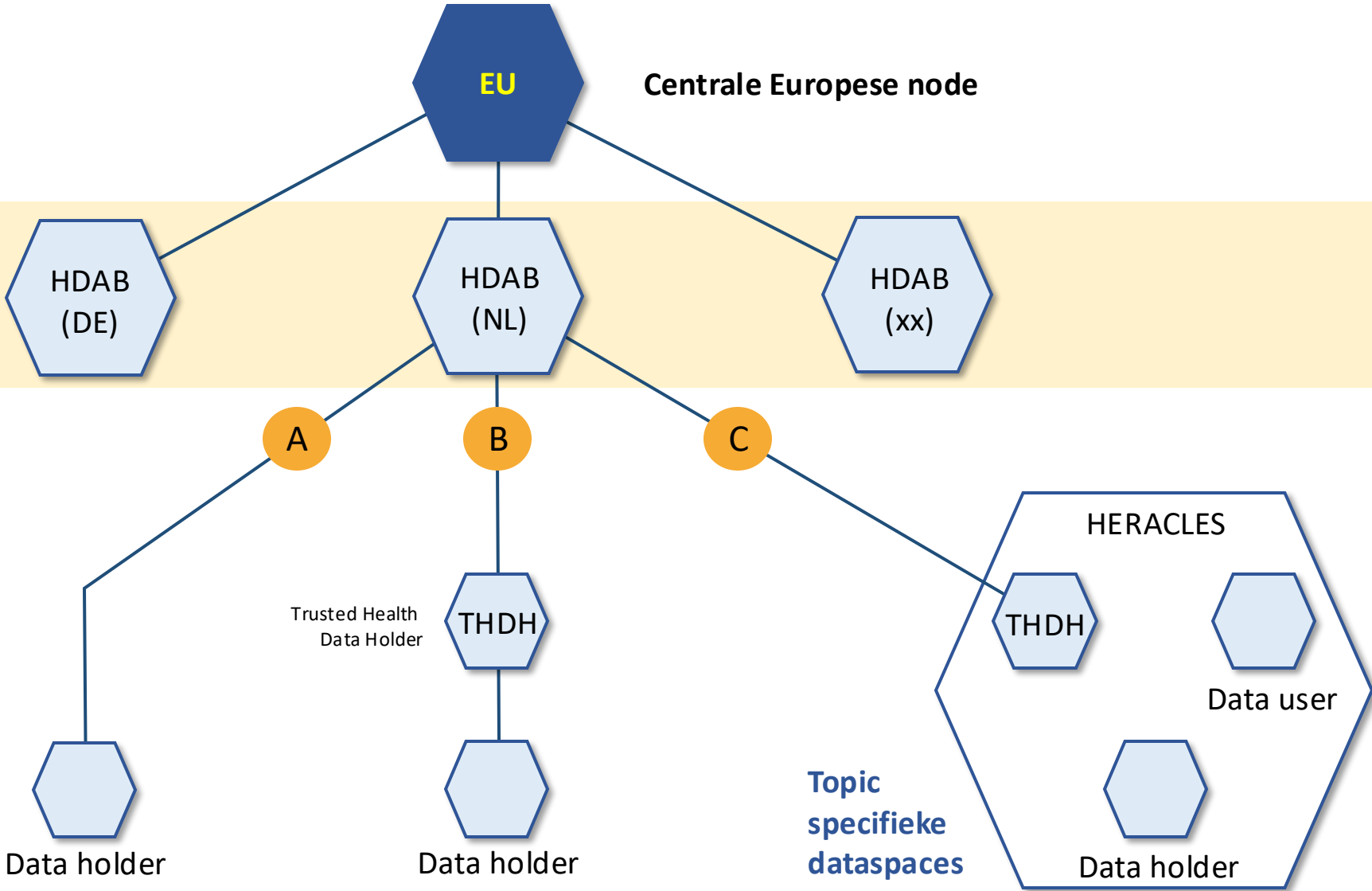
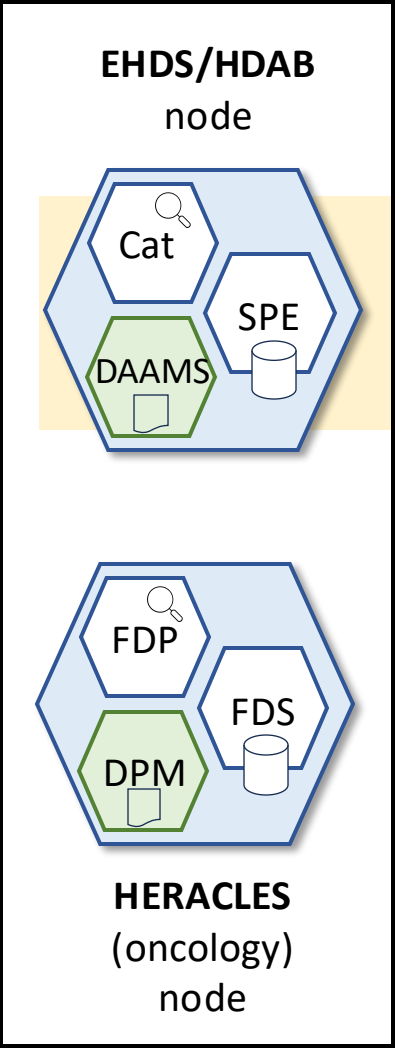
PET

Next challenge



Governance

Compliant functionalities



Summary



From a technical infrastructure point of view, HERACLES:

- Proposed a technical architecture based on EU standards to support the implementation of **EHDS** – including the use of Health DCAT AP and QUANTUM label.
- Demonstrated which/how software components can be implemented according to the proposed architecture - FDPs, FDSs with MPC and SPEs
- Is still sharing knowledge and experiences with **THEDAS2**, **Health-RI** and **HDAP-NL** to accelerated the development of a national (self-learning and improving) data sharing infrastructure and to avoid reinventing the wheel.



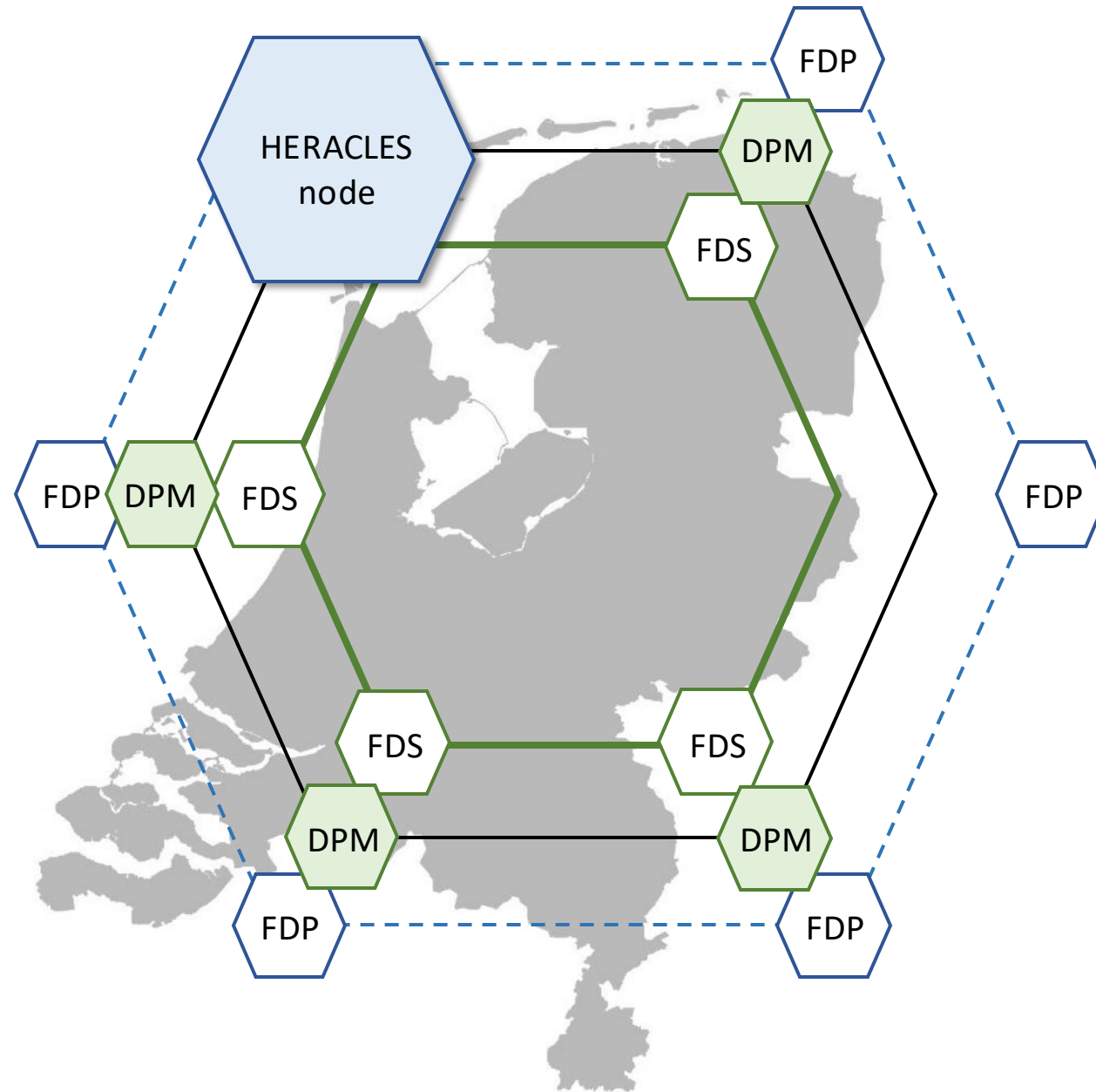
Contact

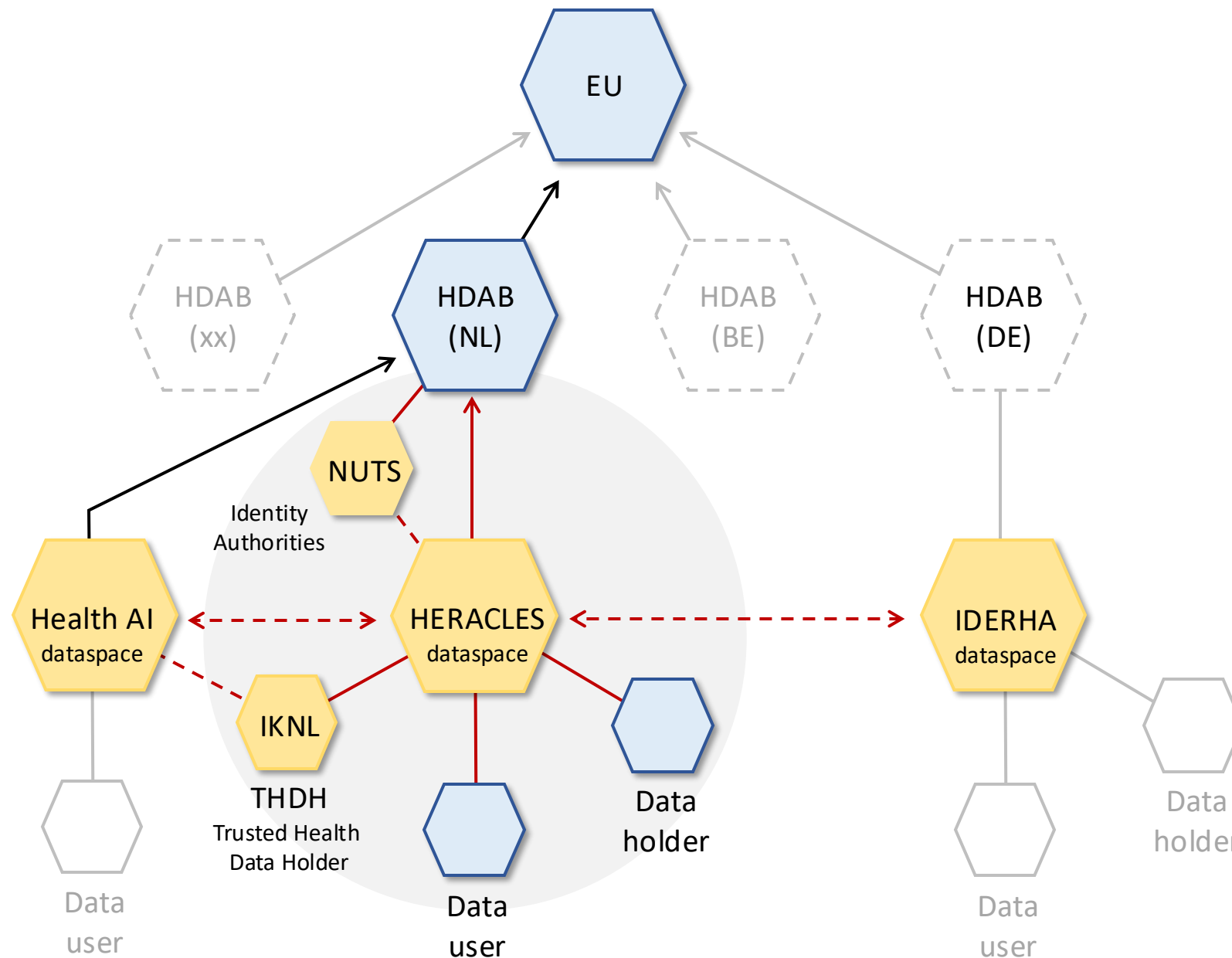
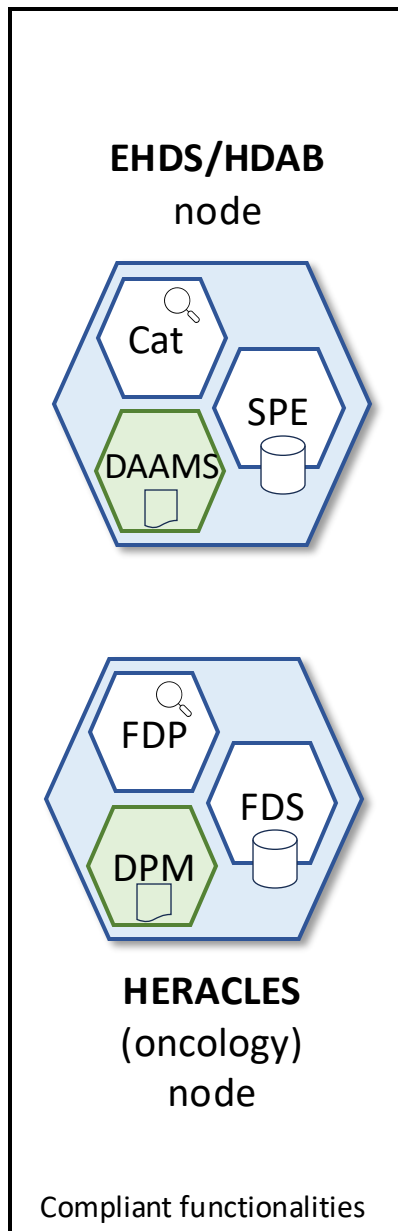
☎ +31 6 15 67 48 72

✉ Erik.Cornelisse@tno.nl

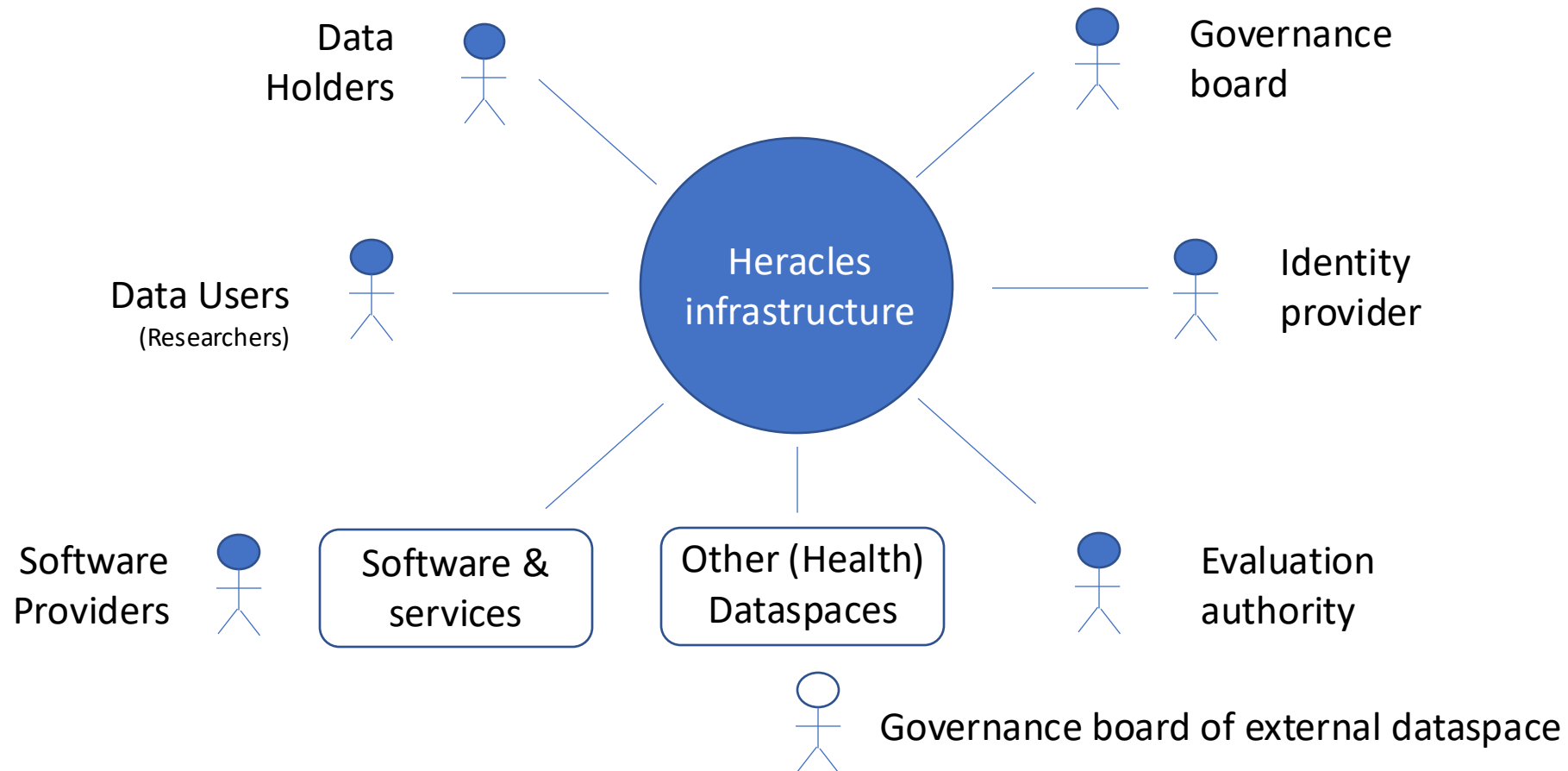
💻 www.tno.nl

🌐 www.linkedin.com/in/erik-cornelisse-7757141





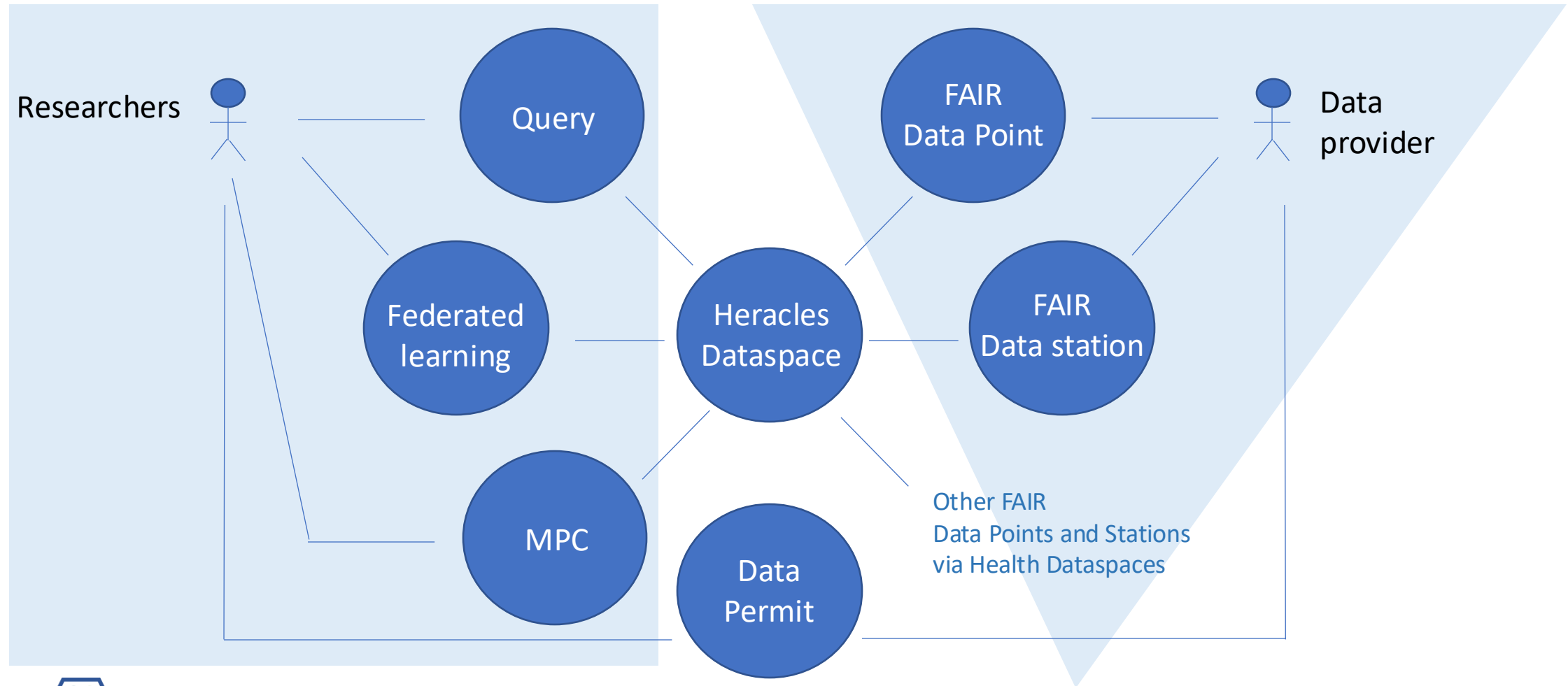
HERACLES infrastructure context diagram



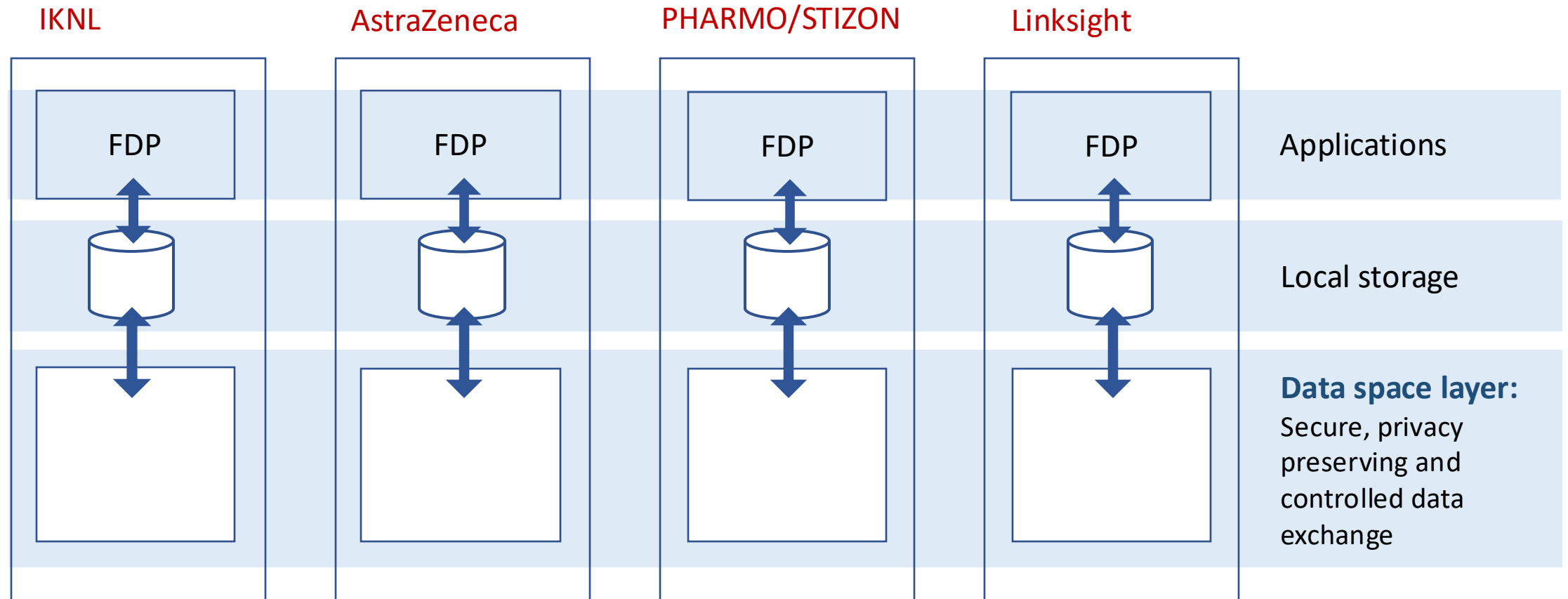
HERACLES infrastructure decomposition

Research application side

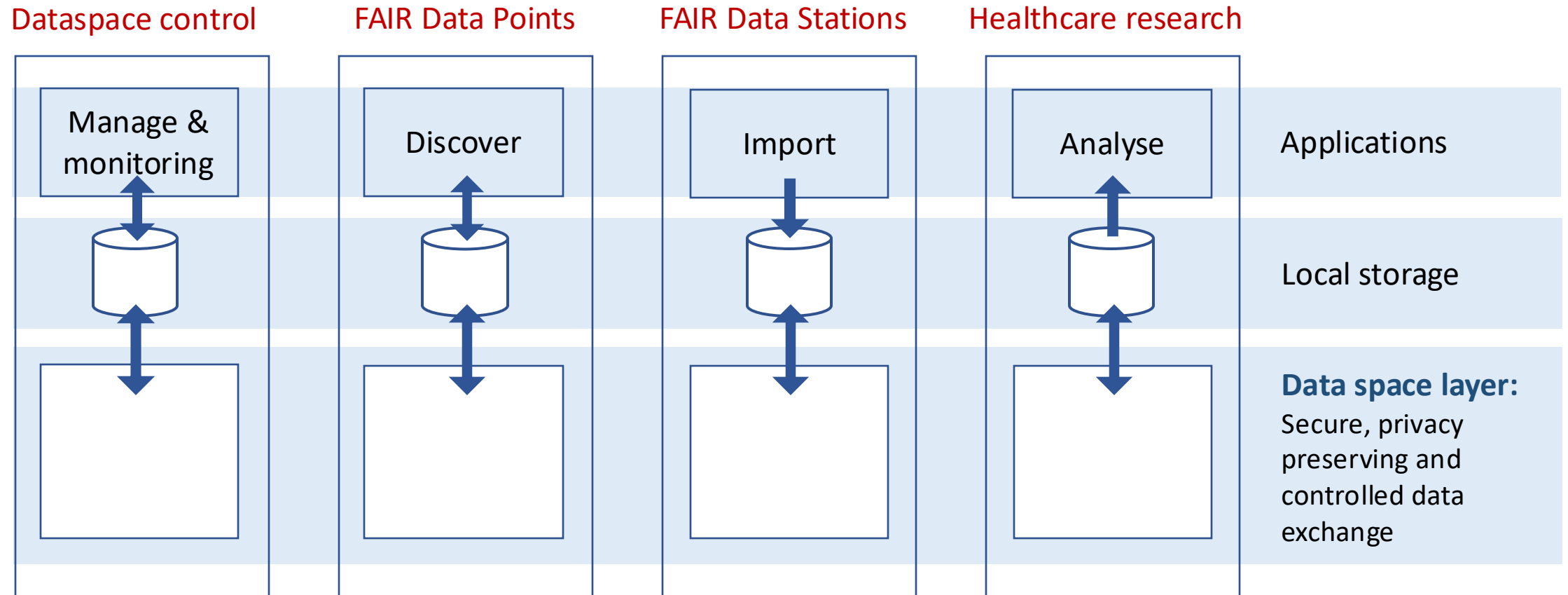
FAIR data application side



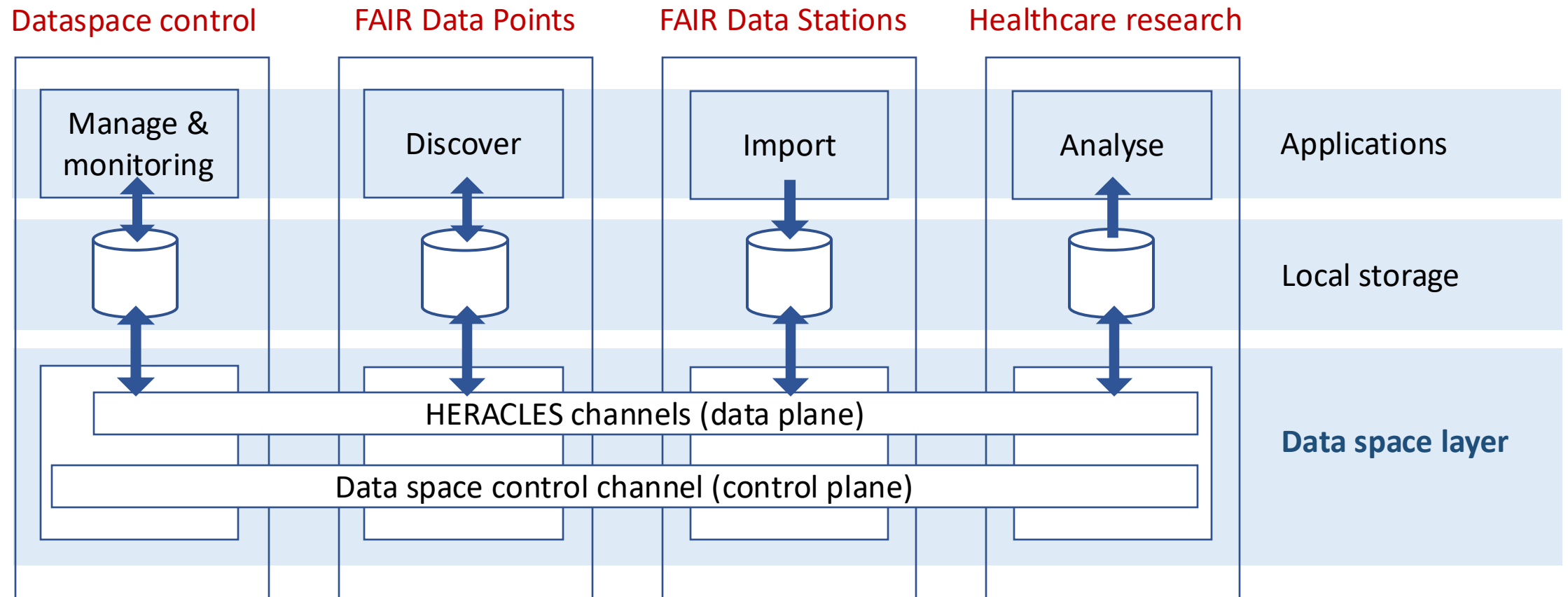
Federated Catalogue (FAIR Data Point)



Four types of services



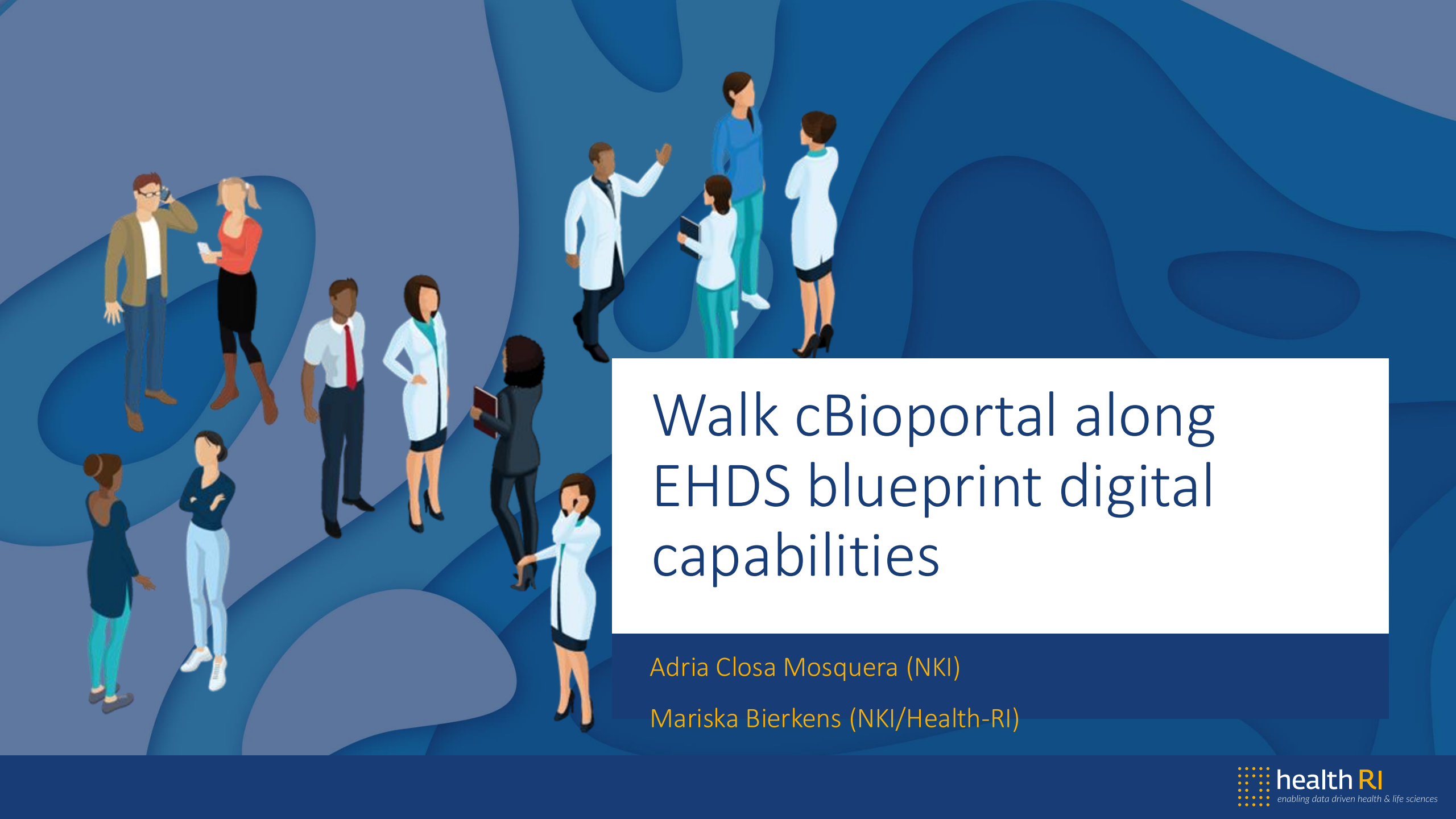
Connected by standards (DSP, DCP, DCAT, VC ...)





Coffee break

Coffee break
10:45-11:00

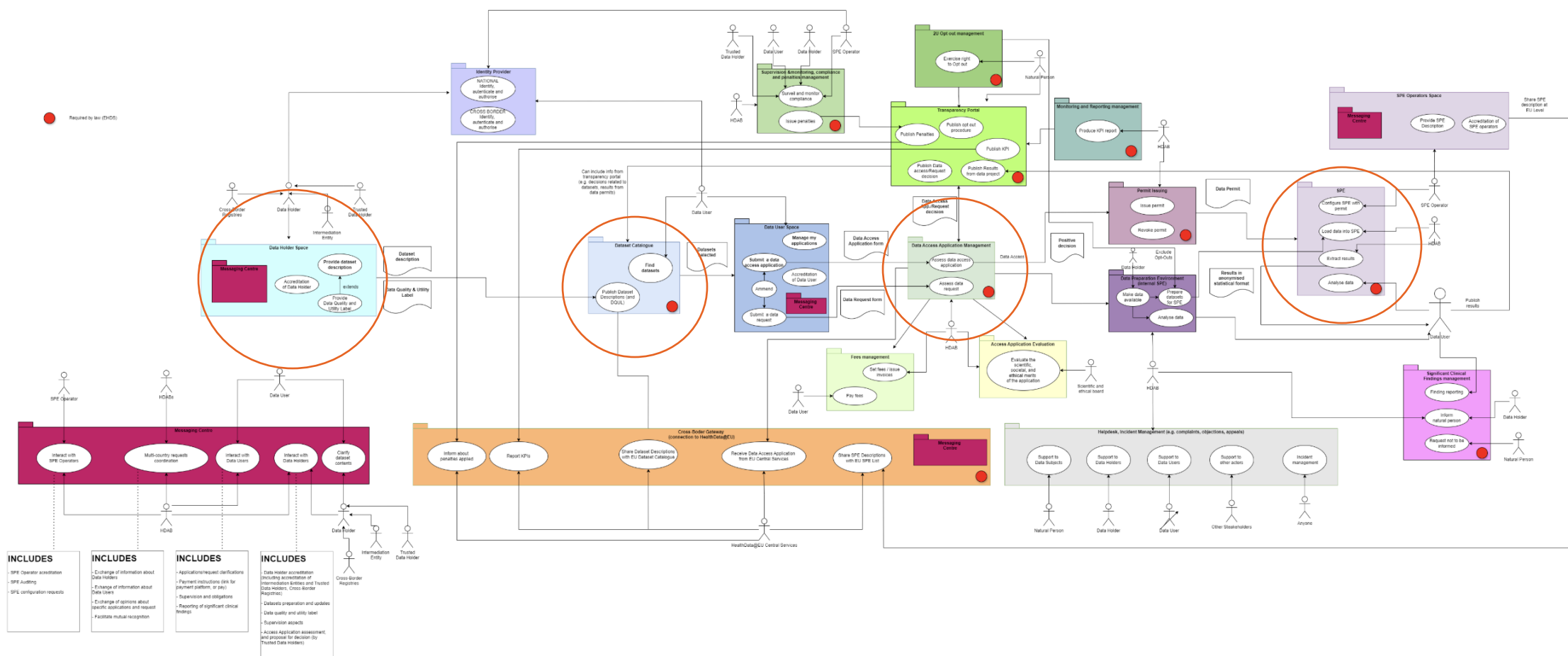
An illustration of a diverse group of healthcare professionals, including doctors, nurses, and researchers, standing and interacting in a modern, blue-toned environment. The background features large, abstract, wavy shapes in various shades of blue. The professionals are dressed in a mix of business casual and clinical attire, such as lab coats and scrubs. Some are holding devices like smartphones or tablets, suggesting a focus on digital health.

Walk cBioportal along EHDS blueprint digital capabilities

Adria Closa Mosquera (NKI)

Mariska Bierkens (NKI/Health-RI)

Backbone



One Portal, One Standard: Transforming Translational Research Data for an EHDS-Future with cBioPortal

*Empowering clinical research with efficient
data management workflows.*

Adria Closa, PhD

Bioinformatician & Data Steward

Department of Pathology

The Translational Gastrointestinal Oncology group

The Netherlands Cancer Institute

Email: a.c.mosquera@nki.nl

08 December 2025



Biology degree (2007-2011)
Master Advance Genetics
(2011-2012)
*Autonomous University of
Barcelona (UAB)
Catalonia*



PhD in Medicine (2012-2017)
*University of Barcelona (UB)
Catalan Institute of Oncology
(ICO-IDIBELL)
Catalonia*



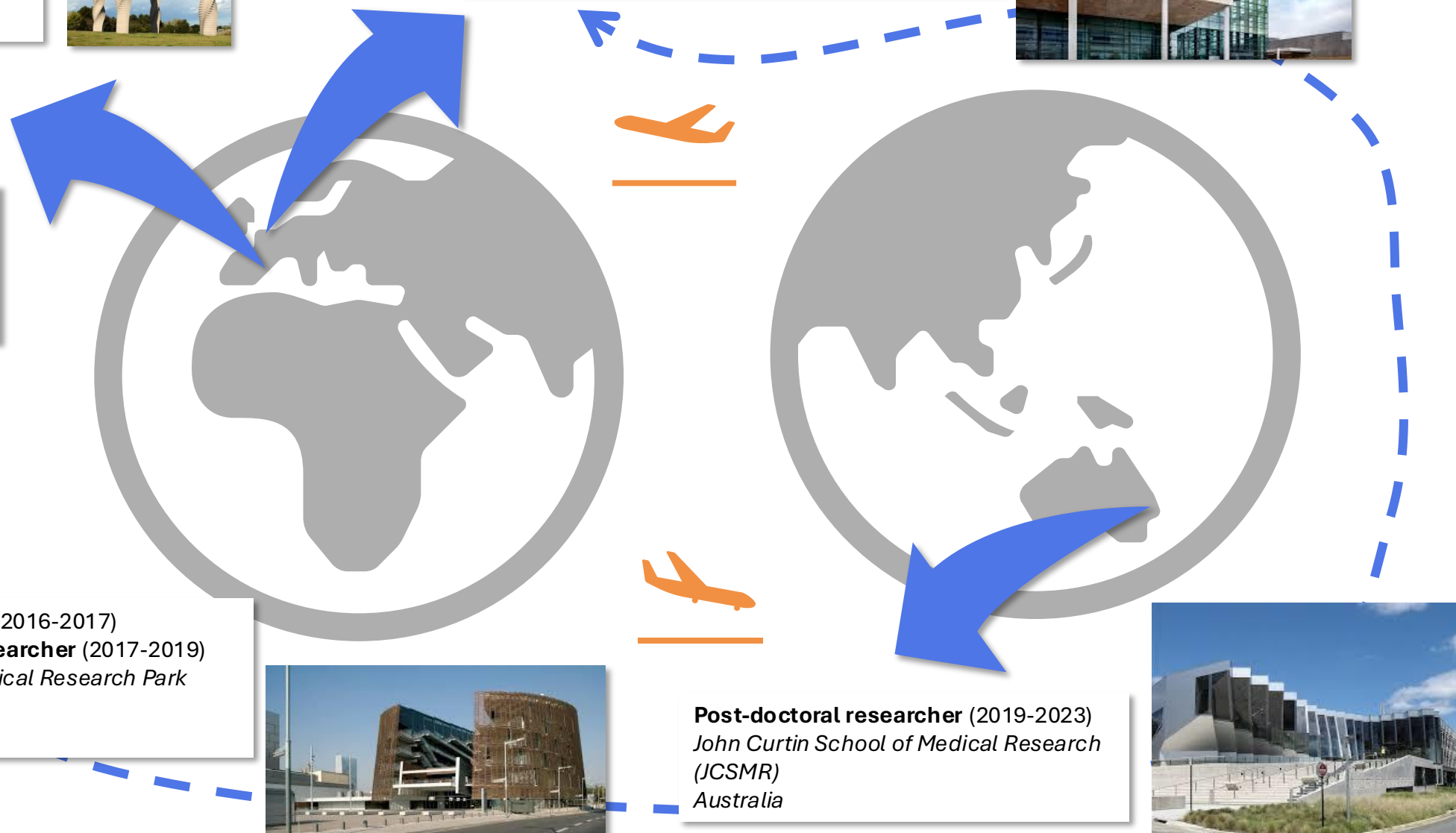
Bioinformatician (2016-2017)
Post-doctoral researcher (2017-2019)
*Barcelona Biomedical Research Park
(PRBB)
Catalonia*

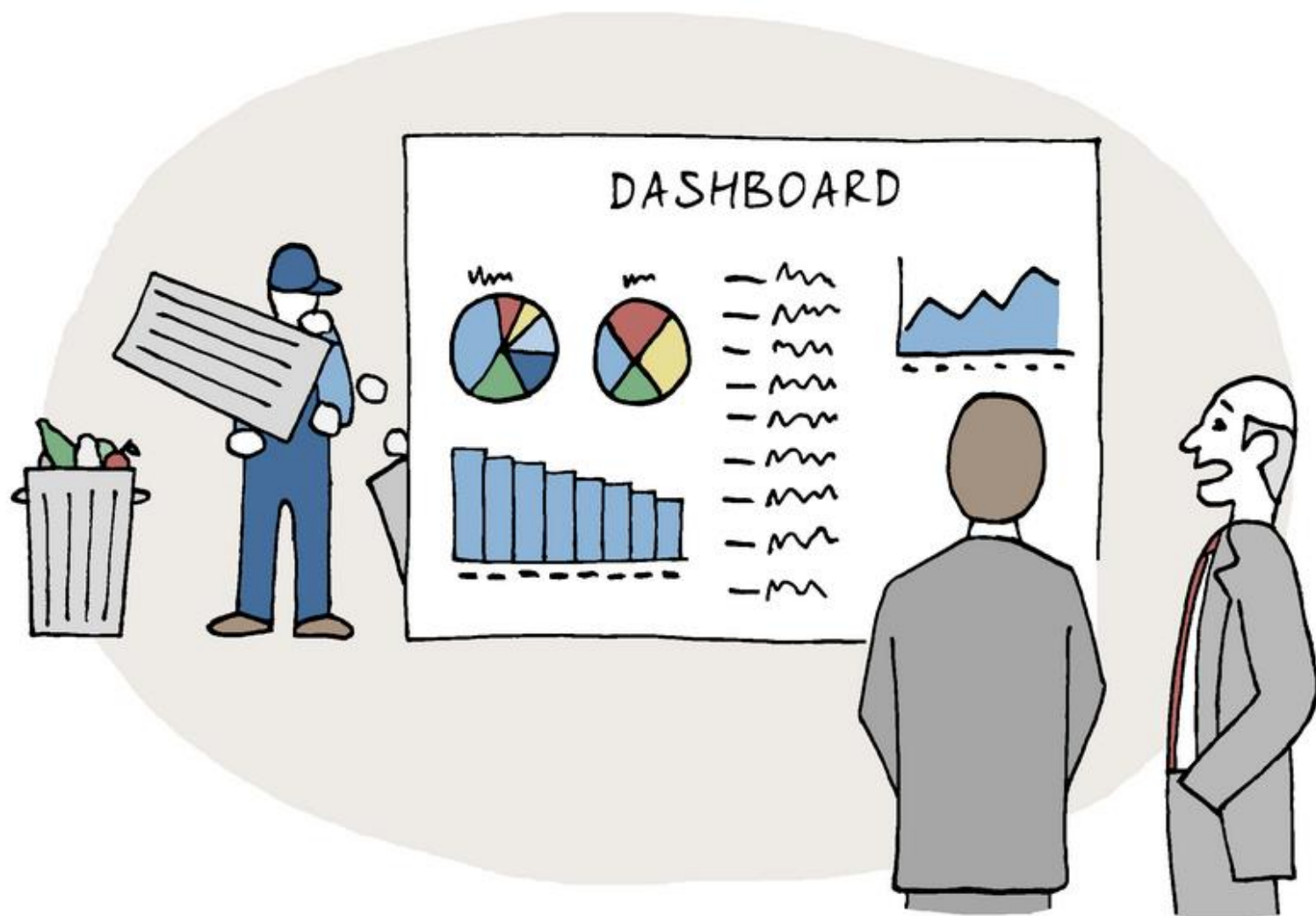


Bioinformatician / data steward (2023-Present)
*Netherlands Cancer Institute (NKI)
The Netherlands*

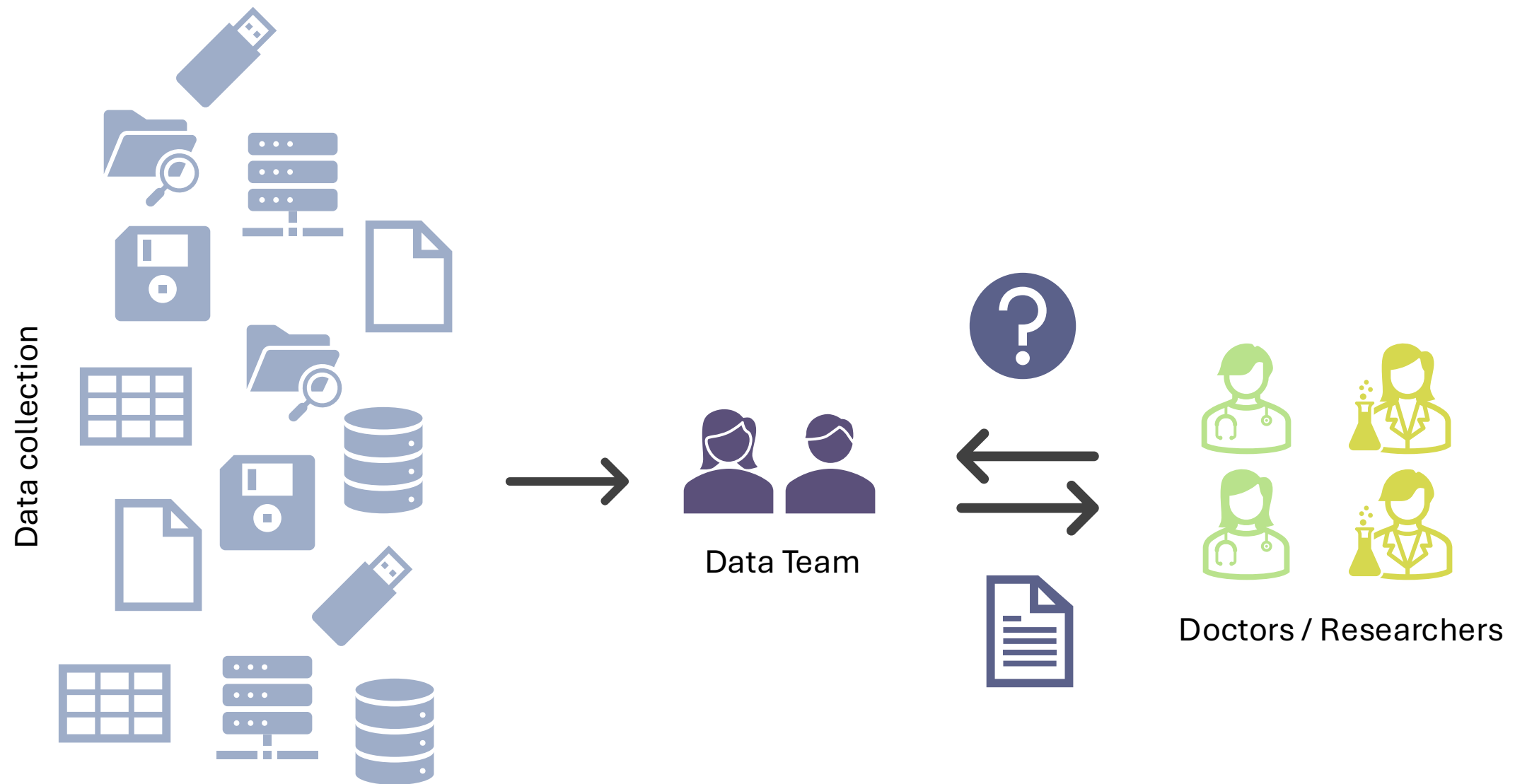


Post-doctoral researcher (2019-2023)
*John Curtin School of Medical Research
(JCSMR)
Australia*



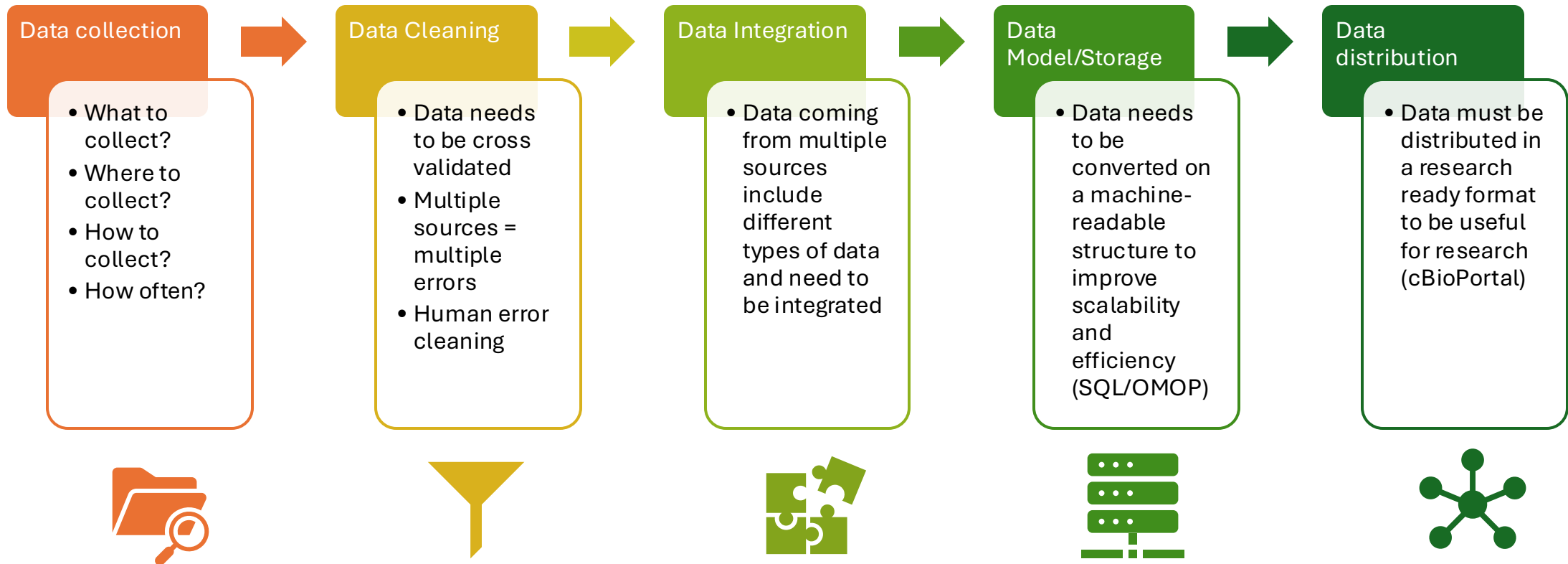


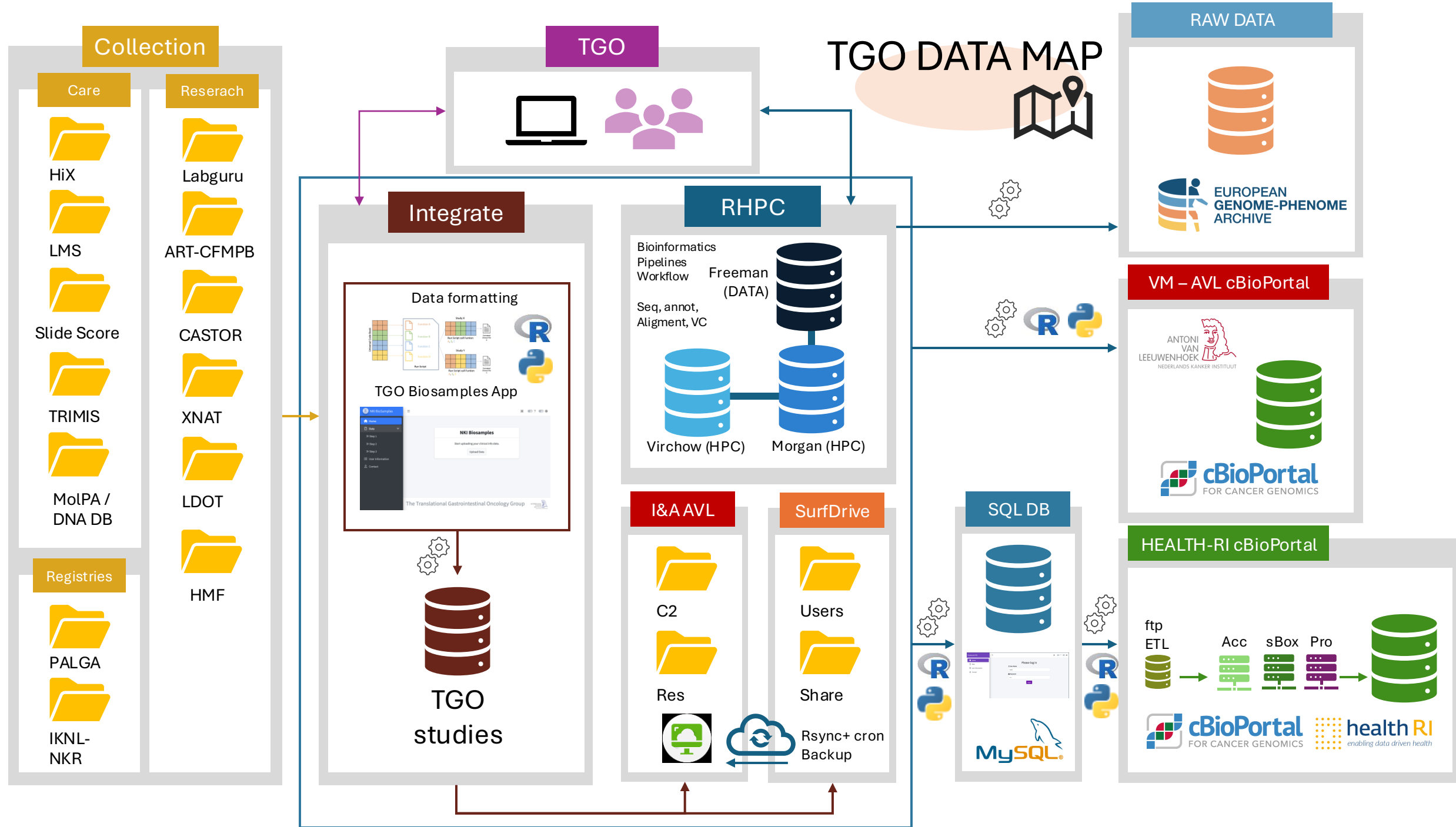
DO WE TRUST THIS DATA?



*“The **main goal of a data management team** is to make sure that data across the organization is **reliable, accessible, secure, and usable** so the researchers can make informed decisions and generate value.”*

TGO DATA MANAGEMENT JOURNEY





The diagram illustrates the integration of various data capturing platforms into a unified analysis and visualization framework. It is organized into three main horizontal sections: Data Capturing Platforms, Integration, Analysis & Visualisation, and Data integration platform.

Data Capturing Platforms: This section lists seven platforms, each with a representative icon:

- Informed Consent:** Icon of three people.
- Clinical Characteristics:** Icon of a hospital building and a person, labeled "CS".
- Imaging:** Icon of a film strip with a crosshair.
- Biosamples & Derivatives:** Icon of a petri dish and test tubes.
- Pathology:** Icon of a microscope and a tissue section.
- Molecular Profiling:** Icon of a DNA double helix.
- PROMs (Patient Reported Outcome Measures):** Icon of a checklist.

Below these platforms are two clusters of overlapping circles representing data types:

- Left Cluster:** Disease Profile & Events, Outcomes, Treatment, Survival.
- Right Cluster:** Copy Number Alterations, Structural Variant, DNA Methylation, Protein, Mutations, mRNA Expression, Biological Pathways.

Integration, Analysis & Visualisation: This central section features the cBioPortal logo and a large blue arrow pointing right. Above the arrow are icons for Mutation, Copy Number, Gene Expression, DNA Methylation, MicroRNA, RPPA, and Clinical Data. Below the arrow are icons for a Kaplan-Meier survival plot, a bar chart, and a pie chart labeled "Tumor Grade" with values 316, 98, and 121.

Data integration platform: This section on the left shows a 3D visualization of data integration. The vertical axis is labeled "Platforms" and the horizontal axis is labeled "Genes / Loci". The depth axis is labeled "Samples" and lists various cancer types: BRCA, BLCA, COAD, GBM, HNSC, KIRC, LAML, LUAD, LUSC, OV, PAAD, and UCEC. The data is represented as a stack of heatmaps and a table.

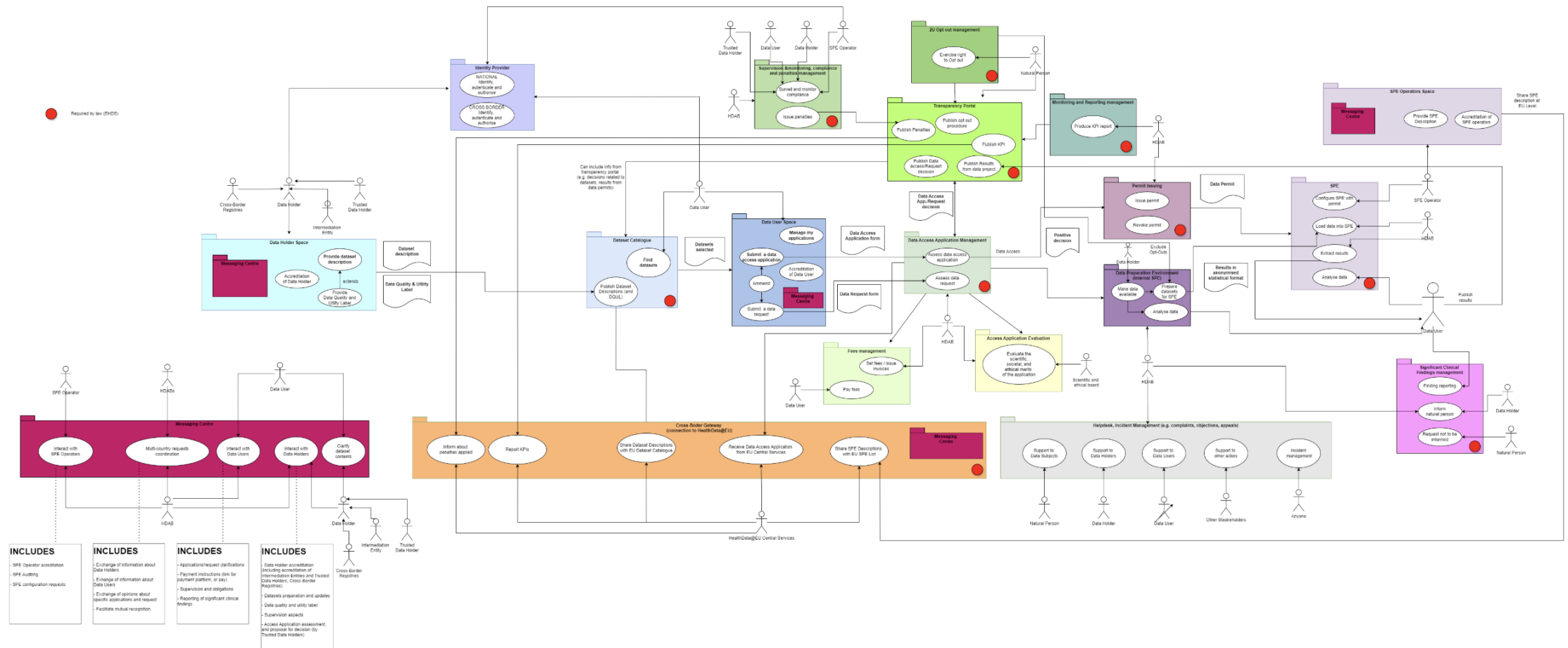
On the right side of the Data integration platform section are two boxes:

- Biological discovery** (green box)
- Clinical interpretation** (blue box)

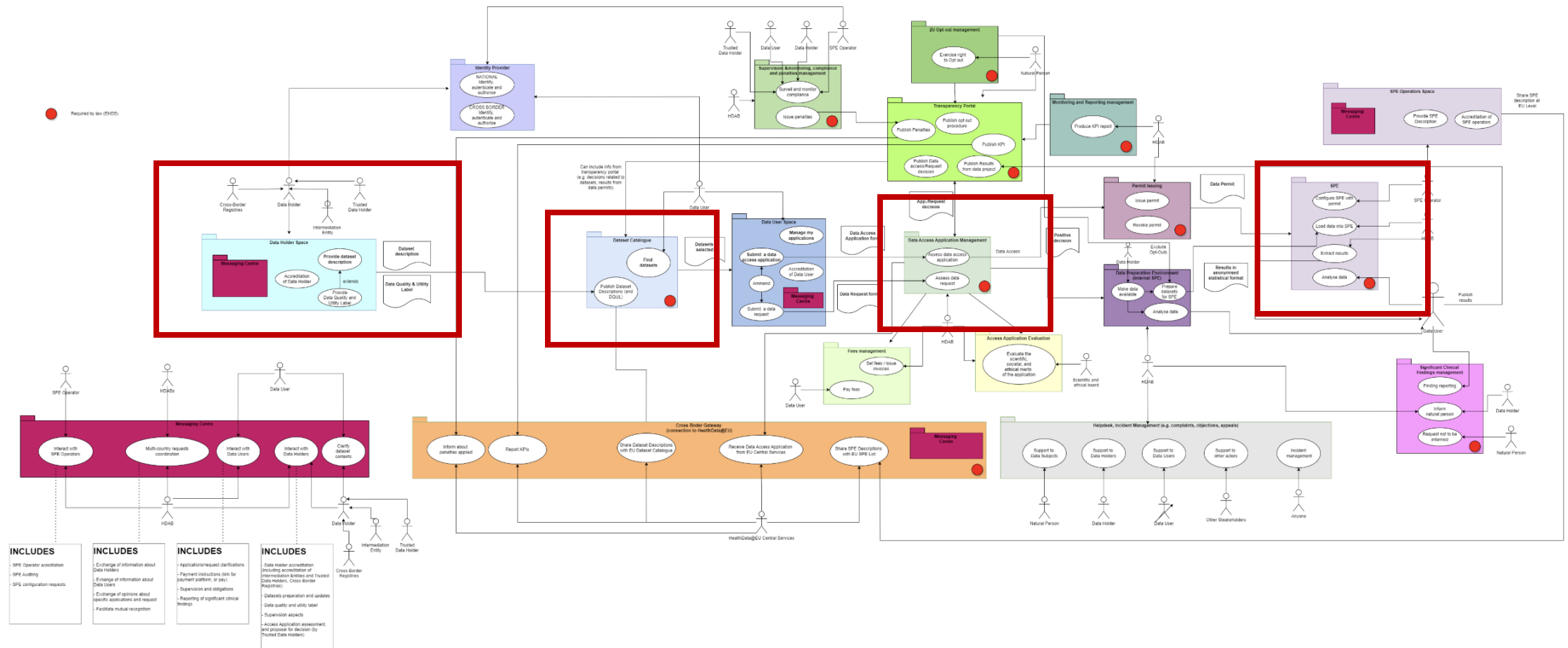
Adapted from TCGA Nat Genet 2014

EHDS digital business capabilities (DBC) blueprint

Where does cBioPortal fit in this DBC in the Netherlands?

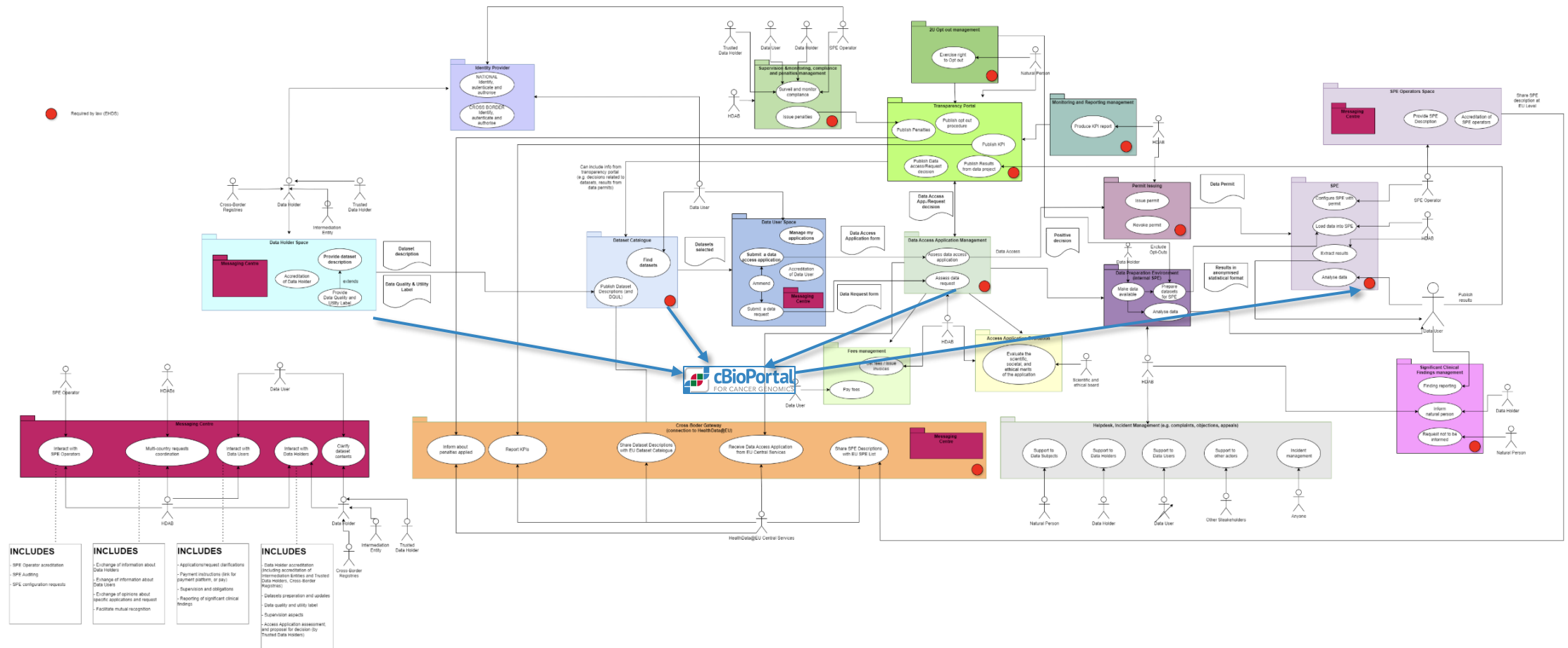


Where does cBioPortal fit in this DBC in the Netherlands?



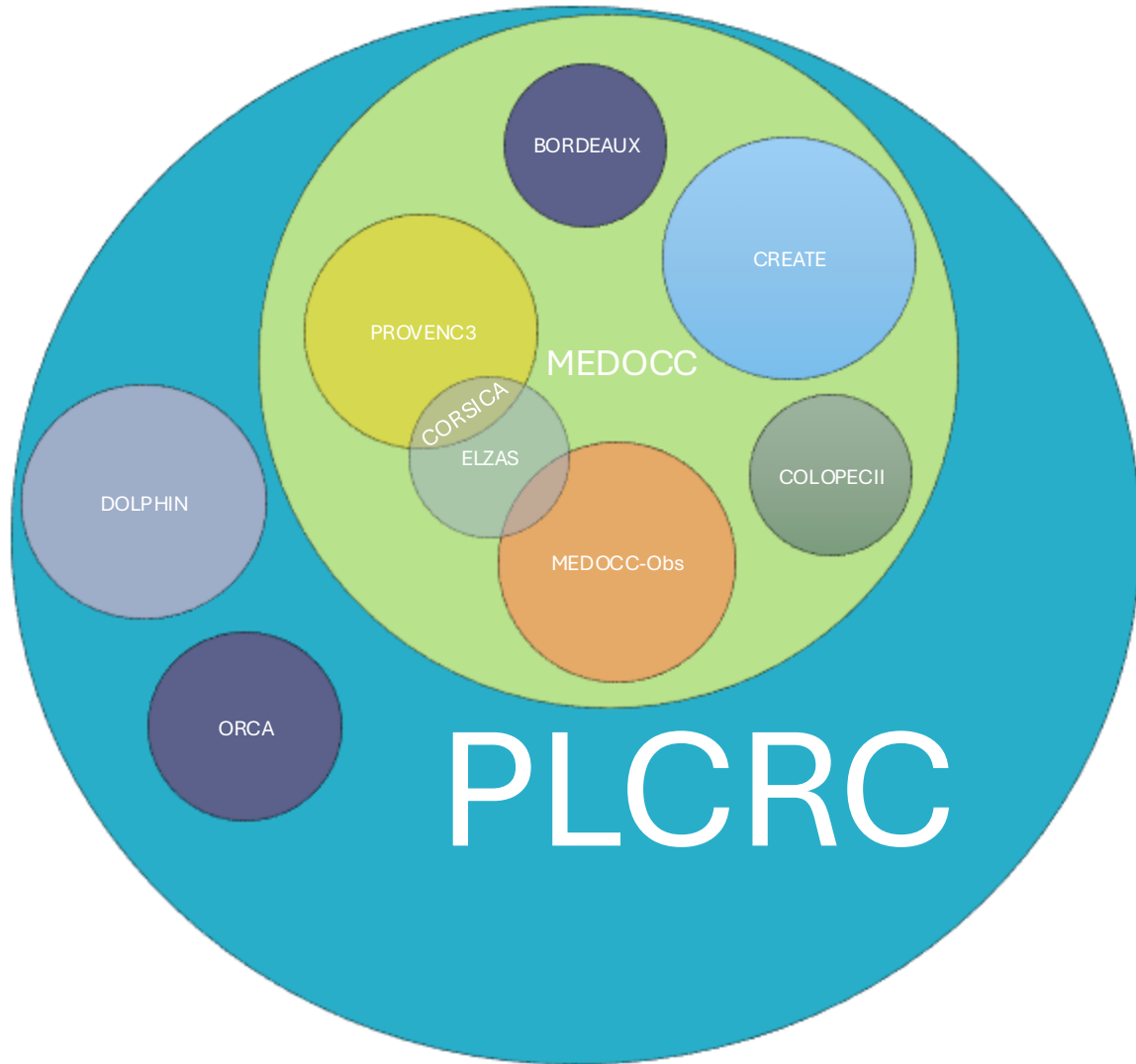
EHDS digital business capabilities (DBC)

Where does cBioPortal fit in this DBC in the Netherlands?



Real use case: mapping the PLCRC data workflow to the EHDS blueprint

PROSPECTIEF
LANDELIJK
CRC COHORT



1 centrale plek voor RWD van darmkankerpatiënten in Nederland



**70 Participating Centers &
> 25,000 Participating Patients**

What is PLCRC?

The Prospective National CRC cohort (PLCRC) provides an infrastructure for collecting clinical data, patient-reported outcomes and storing body material such as blood and tumor tissue from patients with small intestine, colon, rectal and anus cancer. This happens after obtaining the patient's consent.

Main goal?

The goal of the cohort is to facilitate scientific research to improve the prognosis and quality of life of (future) patients.

NKI biobank infrastructure is used to collect and store blood and tissue samples from donors that are part of the different sub-studies inside the PLCRC umbrella.

Executive board PLCR members:

- Prof. Dr. Miriam Koopman, internist oncologist, UMCU
- Dr. Robert Jan Kwakman, internist oncologist, UMCU
- Dr. Anna van Tetering, program manager PLCRC
- Dr Frederieke van der Baan, project leader PLCRC

PLCRC needs and requirements for study monitoring



Weekly monitoring of blood collections and tissues for MEDOCC and CrEATE patients.



Global statistics of inclusion and status of patients, blood and tissue availability and tumor stage distribution.



Structured and clean ready data for data analysis and research.



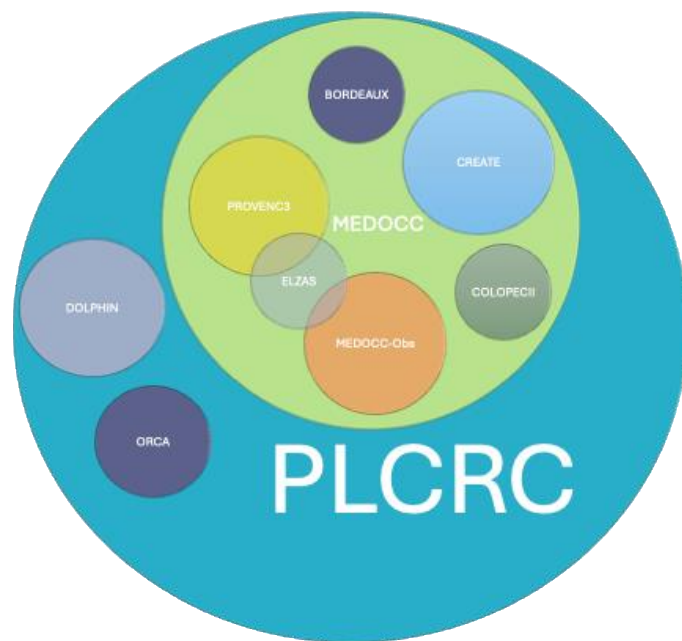
An approach to communicating study data to clinicians and partners involved in the MEDOCC during the weekly UMCU meetings.



Prediction of the next blood collection moment per patient to be able to prepare the next appointment request.

Can we make all PLCRC studies available in cBioPortal at national level?





The cBioPortal for Health-RI provides **visualization, analysis** and **download** of various studies.

Please adhere to [the TCGA publication guidelines](#) when using TCGA data in your publications.

Please cite [Gao et al. *Sci. Signal.* 2013](#) & [Cerami et al. *Cancer Discov.* 2012](#) when publishing results based on cBioPortal.

Query

Please cite [cBioPortal](#)

Select Studies for Visualization & Analysis: 33 studies available (166951 samples) Data type Search...

Study	Count	Study	Count
Bowel	23	Brain/CNS	1
Breast	2	Lymph	5
Ovary	1	Pan-cancer	1

☐ Select all listed studies (33) [Help](#)

Bowel

Colorectal Adenocarcinoma

- ☐ B15PON 108 samples
- ☐ B16PON 52 samples
- ☐ CAIRO5 8069 samples
- ☐ CAIRO5 (Clin Cancer Res 2023) 260 samples
- ☐ CAIRO5 (Lancet Oncol 2023) 521 samples
- ☐ CAIRO6 5032 samples
- ☐ DOLPHIN 13751 samples
- ☐ DeCoDe WP5 911 samples
- ☐ DeCoDe WP6 507 samples
- ☐ NGS-ProToCol Colorectal 78 samples
- ☐ PLCRC Cohort Management 17625 samples
- ☐ PLCRC-BORDEAUX 1797 samples
- ☐ PLCRC-CORSICA 3848 samples
- ☐ PLCRC-CREATE 9824 samples
- ☐ PLCRC-ELZAS 6855 samples
- ☐ PLCRC-MEDOC 66608 samples
- ☐ PLCRC-MEDOC-OBS 5070 samples
- ☐ PLCRC-PROVENC3 6067 samples
- ☐ PLCRC-PROVENC3 (March 2025) 528 samples
- ☐ decode_wps 366 samples
- ☐ mtFIT (Lancet Oncol 2024) 13187 samples

COLON ADENOCARCINOMA

- ☐ Colorectal cancer stool proteomics study 310 samples
- ☐ MOCCAS 3734 samples

Brain/CNS

Diffuse Glioma

- ☐ Low-Grade Gliomas (UCSF, Science 2014) 61 samples

Breast

Breast Invasive Carcinoma

- ☐ Test study es_0 842 samples

Invasive Breast Carcinoma

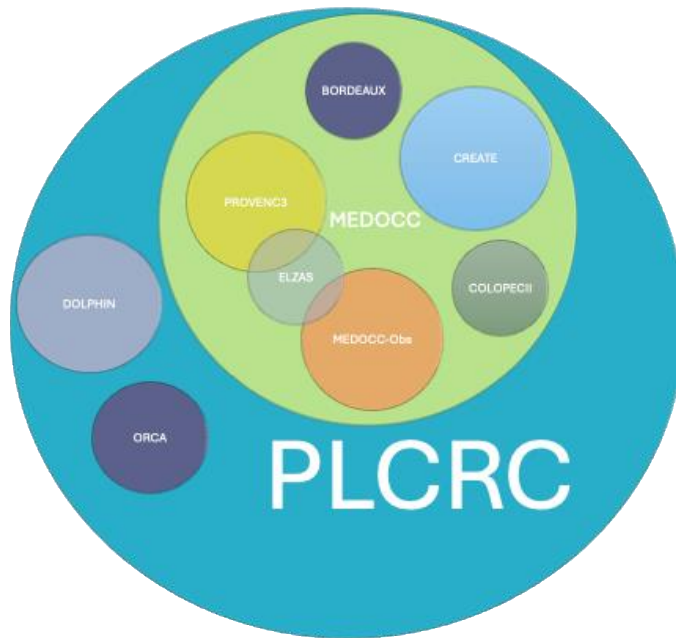
- ☐ MAPK on resistance to anti-HER2 therapy for breast cancer (MSK, Nat Commun. 2022) 145 samples

33 studies available (166951 samples)

Q Query By Gene

OR

🔍 Explore Selected Studies



PLCRC-MEDOC

PLCRC (Prospective Dutch ColoRectal Cancer cohort, www.PLCRC.nl) is a nationwide infrastructure initiative to facilitate colorectal cancer research, with participation of over 50 hospitals that are open for patient accrual. PLCRC-MEDOC (Molecular Early Detection of Colon Cancer) is a substudy of PLCRC, in which a.o. blood samples are being collected from patients with localized (stage I, II, III) disease, for various translational research studies. The main purpose of this cBioPortal study request is to obtain an overview of PLCRC-MEDOC patient accrual and longitudinal blood sample collection. Last update 28-02-25

Click gene symbols below or [Query](#)

Summary Clinical Data

Selected: 4,231 patients | 61,852 samples [Custom Selection](#) [Charts](#) [Groups](#) [Settings](#)

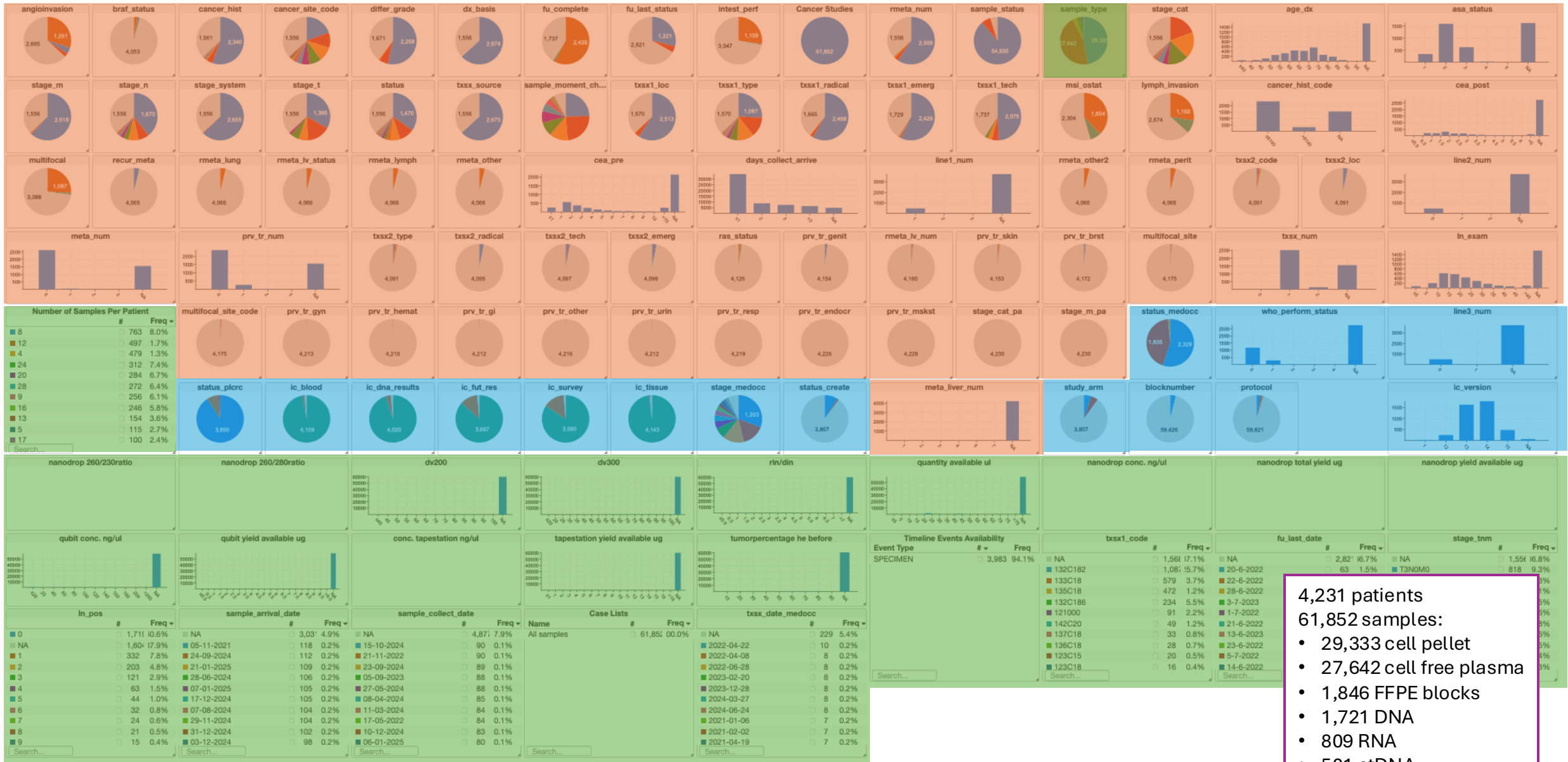


PLCRC-MEDOCC

PLCRC (Prospective Dutch ColoRectal Cancer cohort, www.PLCRC.nl) is a nationwide infrastructure initiative to facilitate colorectal cancer research, with participation of over 50 hospitals that are open for patient accrual. PLCRC-MEDOCC (Molecular Early Detection of Colon Cancer) is a substudy of PLCRC, in which a.o. blood samples are being collected from patients with localized (stage I, II, III) disease, for various translational research studies. The main purpose of this cBioPortal study request is to obtain an overview of PLCRC-MEDOCC patient accrual and longitudinal blood sample collection. Last update 28-02-25

Summary Clinical Data

Selected: 4,231 patients | 61,852 samples



4,231 patients
61,852 samples:

- 29,333 cell pellet
- 27,642 cell free plasma
- 1,846 FFPE blocks
- 1,721 DNA
- 809 RNA
- 501 ctDNA

IKNL-NKR

LDOT

ART-CFMPB

Attribute	Value
age_dx	57
angioinvasion	No
asa_status	2
cancer_hist	Adenocarcinoma, NOS
cancer_hist_code	8140
cancer_site_code	C187
cea_post	0.5
cea_pre	0.5
differ_grade	Moderately Differentiated
dx_basis	Histology of Primary Tumor
fu_complete	No
ic_blood	Yes
ic_dna_results	Yes
ic_fut_res	Yes
ic_survey	Yes
ic_tissue	Yes
ic_version	13
intest_perf	No
ln_exam	15
ln_pos	0
lymph_invasion	No
meta_num	0
rmsi_ostat	No
multifocal	No
prv_tr_num	0
rmeta_num	0
stage_cat	2a
stage_m	M0
stage_medocc	T3N0M0
stage_n	N0
stage_system	Pathology
stage_t	T3
stage_tnm	T3N0M0
status	Included
status_create	Included
status_medocc	Included
status_plcrc	Included
study_arm	Intervention
txsx_date_medocc	2020-08-17
txsx_num	1
txsx_source	NKR Dec 2023
txsx1_code	133C18
txsx1_emerg	No, elective
txsx1_loc	Organ
txsx1_radical	Microscopically radical resection
txsx1_tech	Conventional scopic
txsx1_type	Sigmoid resection
Number of Samples Per Patient	30

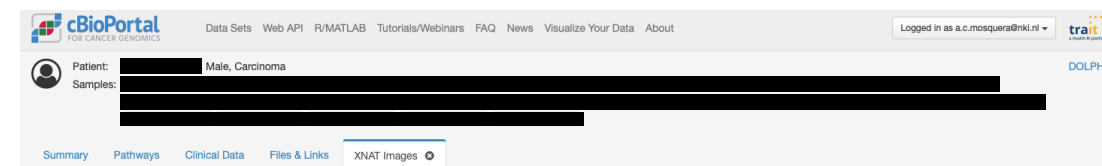
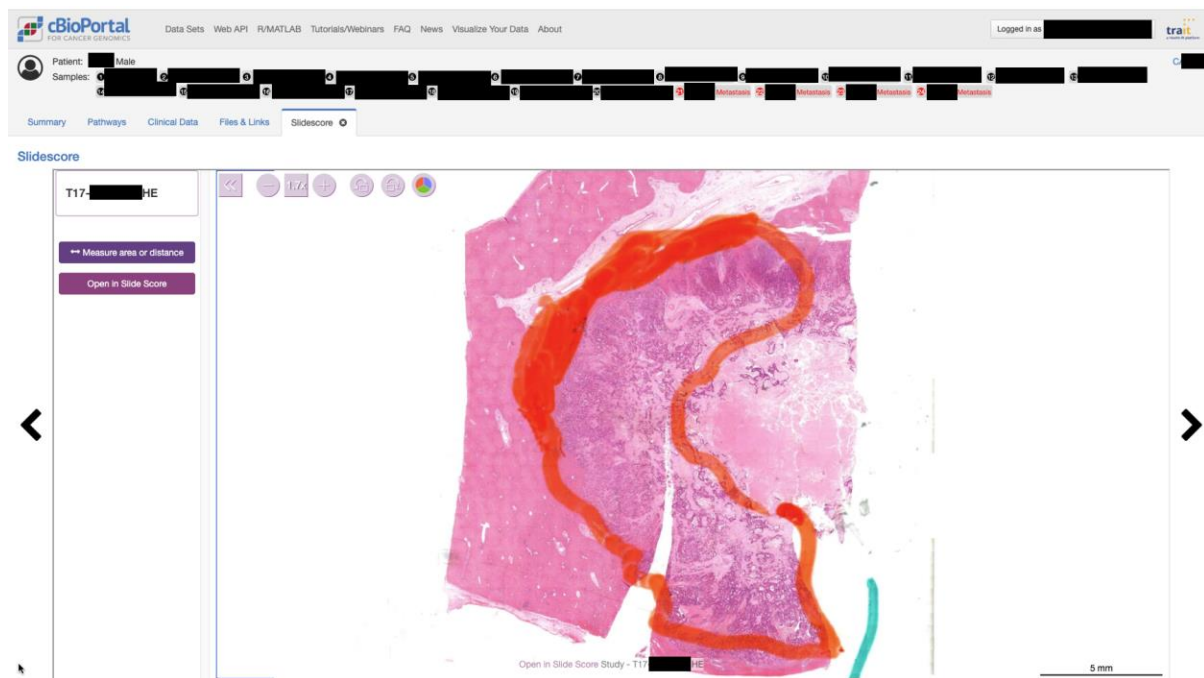
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days_collect_arrive	1	1	2	2	4	1	1	4	1	1	1	4	2	1
sample_arrival_date	01-03-2022	01-03-2022	04-09-2020	04-09-2020	17-08-2020	01-03-2022	23-07-2021	17-08-2020	24-02-2021	01-03-2022	07-03-2023	17-08-2020	10-11-2021	23-07-2021
sample_collect_date	28-02-2022	28-02-2022	02-09-2020	02-09-2020	13-08-2020	28-02-2022	22-07-2021	13-08-2020	23-02-2021	28-02-2022	06-03-2023	13-08-2020	08-11-2021	22-07-2021
sample_moment_checked	TFSU18m	TFSU18m	2Post	2Post	T1Pre	TFSU18m	TFSU12m	T1Pre	TFSU6m	TFSU18m	T7FU3y	T1Pre	T4FU12m_O	T4FU12m
sample_status	Available	Available	Used	Available	Available	Available	Available	Available	Available	Available	Available	Available	Available	Available
sample_type	Cell Pellet	Cell-free Plasma	Cell-free Pellet	Cell Pellet	Cell Pellet	Cell-free Plasma	Cell Pellet	Cell-free Plasma	Cell-free Plasma	Cell Pellet	Cell-free Plasma	Cell Pellet	Cell Pellet	Cell Pellet

Timeline Data

Specimen

START_DATE	STOP_DATE	EVENT_TYPE	SPECIMEN_TYPE	SAMPLE_ARRIVAL_DATE	SAMPLE_TYPE	SAMPLE_ID
0		SPECIMEN	Blood	17-08-2020	Cell-free Plasma	LMS-12345-4539036
0		SPECIMEN	Blood	17-08-2020	Cell Pellet	LMS-12345-4539032
0		SPECIMEN	Blood	17-08-2020	Cell-free Plasma	LMS-12345-4539035
0		SPECIMEN	Blood	17-08-2020	Cell Pellet	LMS-12345-4539033
4		SPECIMEN	Tissue	21-08-2020	FFPE block	T20-70073
18		SPECIMEN	Blood	04-09-2020	Cell-free Plasma	LMS-12345-4550561
18		SPECIMEN	Blood	04-09-2020	Cell Pellet	LMS-12345-4550558
18		SPECIMEN	Blood	04-09-2020	Cell-free Plasma	LMS-12345-4550562
18		SPECIMEN	Blood	04-09-2020	Cell Pellet	LMS-12345-4550559
191		SPECIMEN	Blood	24-02-2021	Cell-free Plasma	LMS-12345-4647236
191		SPECIMEN	Blood	24-02-2021	Cell Pellet	LMS-12345-4647233
191		SPECIMEN	Blood	24-02-2021	Cell-free Plasma	LMS-12345-4647235
191		SPECIMEN	Blood	24-02-2021	Cell Pellet	LMS-12345-4647232
340		SPECIMEN	Blood	23-07-2021	Cell Pellet	LMS-12345-4715662
340		SPECIMEN	Blood	23-07-2021	Cell Pellet	LMS-12345-4715663
340		SPECIMEN	Blood	23-07-2021	Cell-free Plasma	LMS-12345-4715666
340		SPECIMEN	Blood	23-07-2021	Cell-free Plasma	LMS-12345-4715665
450		SPECIMEN	Blood	10-11-2021	Cell Pellet	LMS-12345-4751959
450		SPECIMEN	Blood	10-11-2021	Cell-free Plasma	LMS-12345-4751961
450		SPECIMEN	Blood	10-11-2021	Cell-free Plasma	LMS-12345-4751962
450		SPECIMEN	Blood	10-11-2021	Cell Pellet	LMS-12345-4751958
561		SPECIMEN	Blood	01-03-2022	Cell Pellet	LMS-12345-4784369
561		SPECIMEN	Blood	01-03-2022	Cell-free Plasma	LMS-12345-4784371
561		SPECIMEN	Blood	01-03-2022	Cell Pellet	LMS-12345-4784368
561		SPECIMEN	Blood	01-03-2022	Cell-free Plasma	LMS-12345-4784372
932		SPECIMEN	Blood	07-03-2023	Cell Pellet	BB000083897
932		SPECIMEN	Blood	07-03-2023	Cell-free Plasma	BB000083896
932		SPECIMEN	Blood	07-03-2023	Cell Pellet	BB000083898
932		SPECIMEN	Blood	07-03-2023	Cell-free Plasma	BB000083901

Slidescore and XNAT

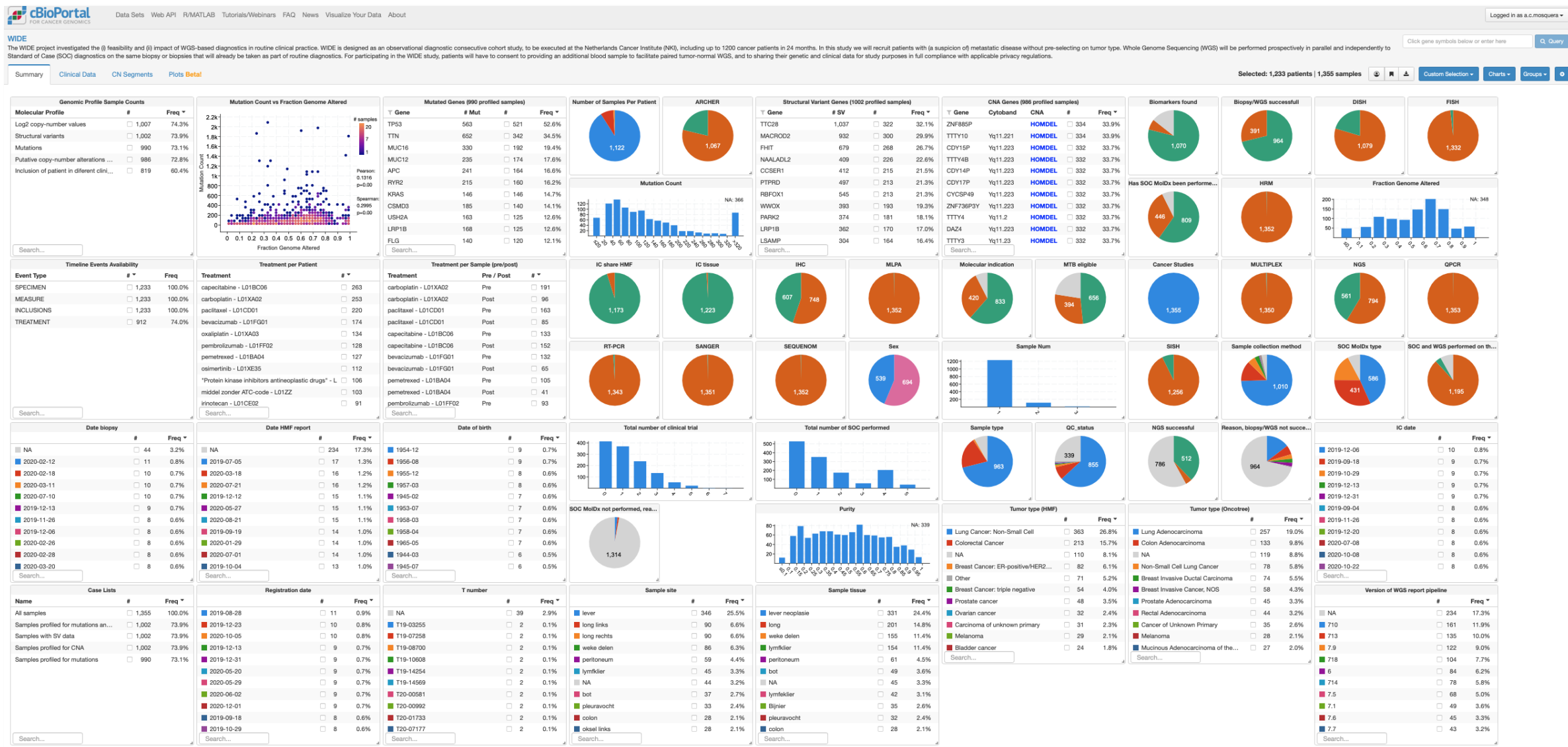


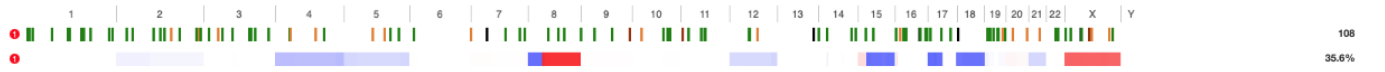
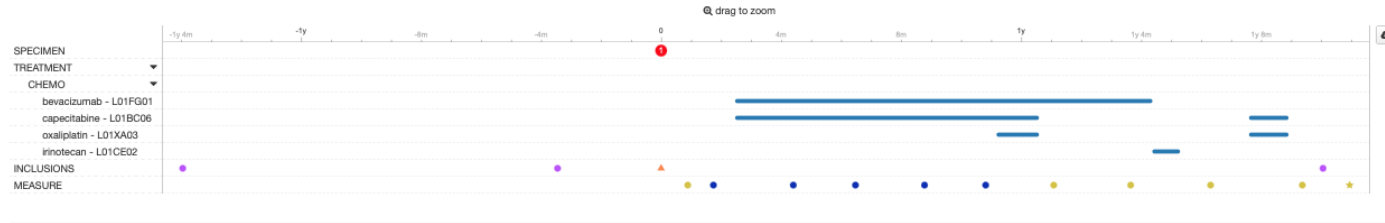
XNAT Images PICNIC0015317 | Patient XNAT Images

[Open in new window](#)



WIDE: Integration of ALEA data + sequencing data from HMF in NKI cBioPortal





108 Mutations (page 1 of 11)

Gene	Protein Change	Annotation	Mutation Type	Allele Freq	Copy #	Cohort	COSMIC
NRAS	G12S		Missense	0.24	ShallowDel	2%	926
PIK3CA	E545K		Missense	0.28	ShallowDel	11%	1431
RAF1	S257L		Missense	0.26	ShallowDel	<1%	12
TP53	R273H		Missense	0.39	ShallowDel	53%	1312
CTCF	S224Cfs*5		FS del	0.27	ShallowDel	<1%	
PHOX2B	X143_splice		Splice	0.14	ShallowDel		
APC	R213*		Nonsense	0.35	ShallowDel	17%	56
APC	X216_splice		Splice	0.13	ShallowDel	17%	36
AR	Q91del		IF del	0.05	ShallowDel	8%	
FANCA	R517Q		Missense	0.13	ShallowDel	3%	

Showing 1-10 of 108 Mutations [Show more](#)

29 Structural Variants (page 1 of 3)

Gene 1	Gene 2	Status	Annotation	Variant Class	Event Info	Connection Type
C9ORF3	FANCC	s		DEL	NA	NA
CBLB	CBLB	s		DUP	NA	NA
SMAD3	SMAD3	s		DEL	NA	NA
GPHN	GPHN	s		DEL	NA	NA
TRIP11	TRIP11	s		DUP	NA	NA
ADAM2	ADAM2	s		INV	NA	NA
ARHGAP39	ARHGAP39	s		DUP	NA	NA
ASPH	HEATR5A	s		LINE	NA	NA
ASPH	KRT8	s		LINE	NA	NA
DPH6-AS1	DPH6-AS1	s		DUP	NA	NA

Showing 1-10 of 29 Structural Variants [Show more](#)

5 Copy Number Alterations

Gene	CNA	Annotation	Cytoband	Cohort
SMAD3	DeepDel		15q22.33	2%
NMTR5-TGA3-1	DeepDel		17p11.2	6%
MTRNR2L1	DeepDel		17p11.2	6%
MTND2P13	DeepDel		17p11.2	6%
MTND1P15	DeepDel		17p11.2	6%

Showing 1-5 of 5 Copy Number Alterations

Patient timeline:

- Specimen collection
- Treatments received by the patient during study
- Inclusions in clinical trials
- Different tumor measurements

Mutation data

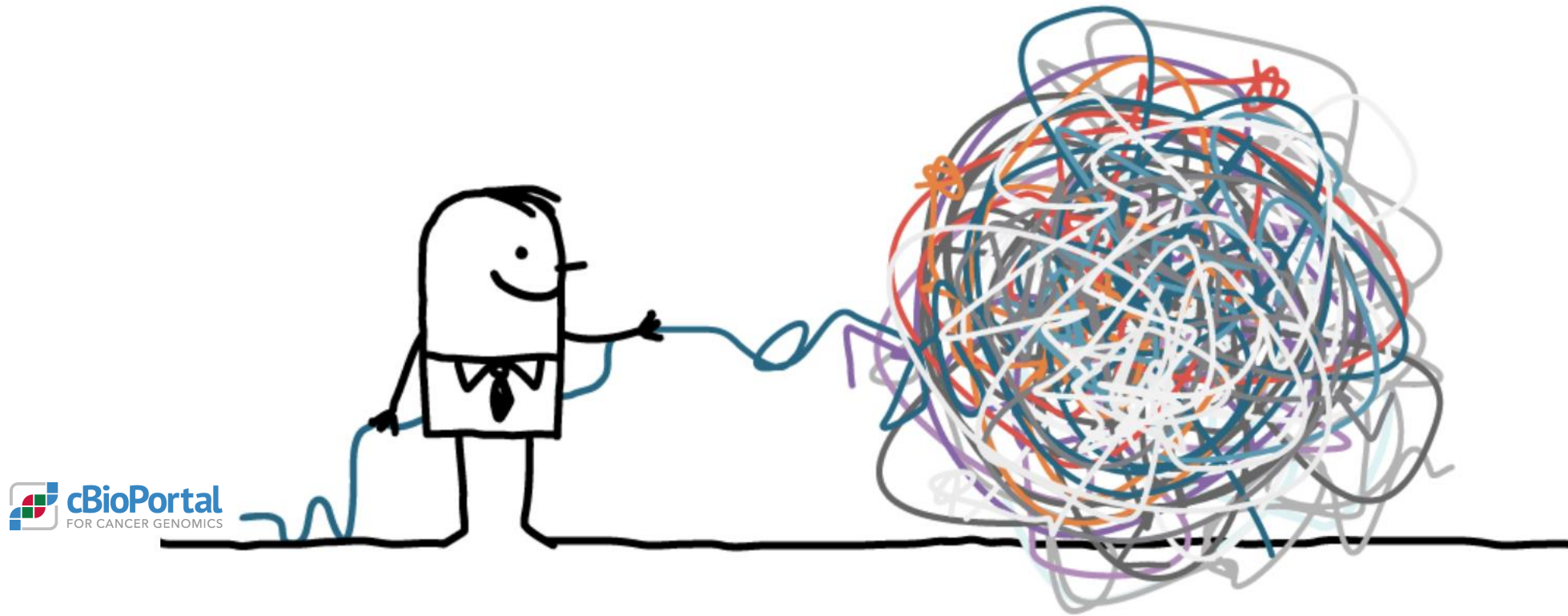
Segmented copy number

Annotated mutation data linked to multiple reference data bases (OncoKB, My Caner Genome, Cancer Hotspots)

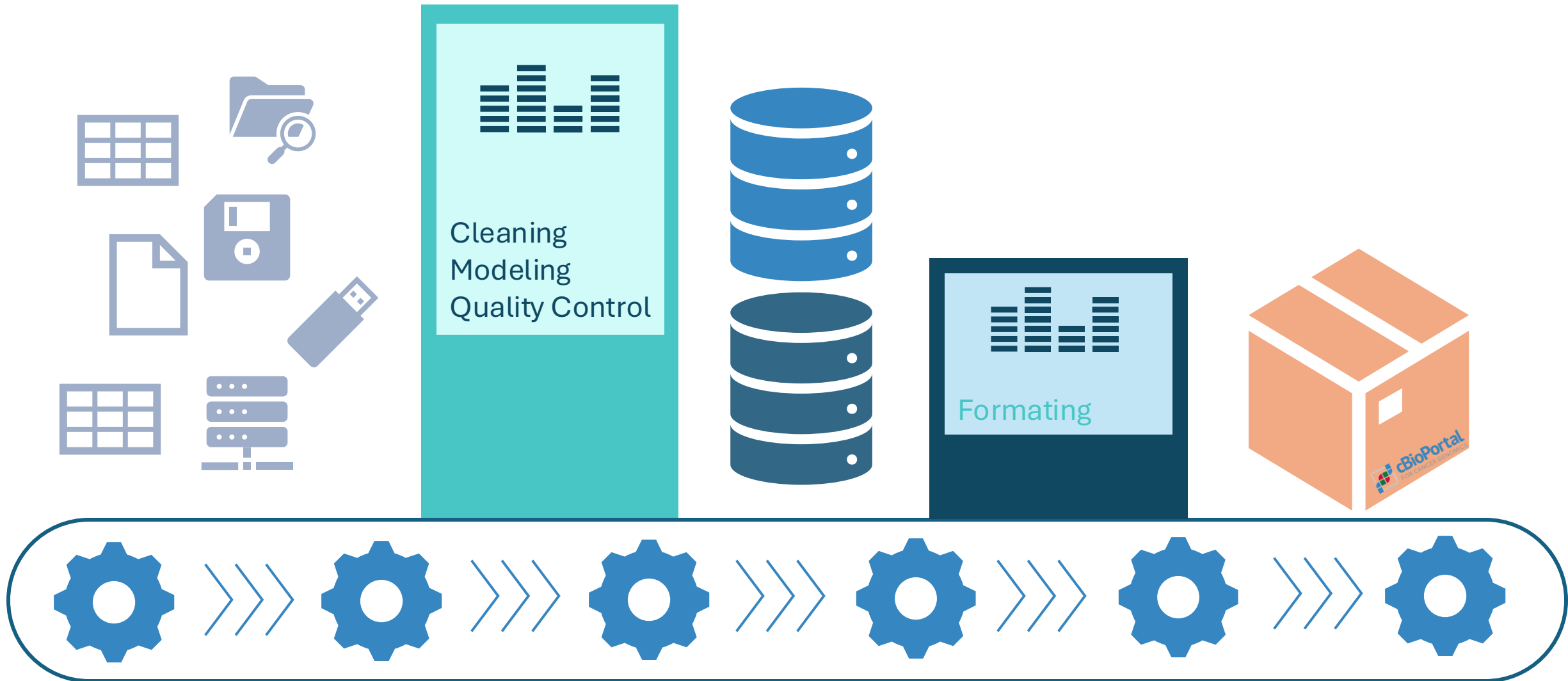
Structural Variants (DUP, DEL, INV, etc)

Copy Number Alterations

From unstructured data to cBioPortal?



Data modeling and standardization allows automation

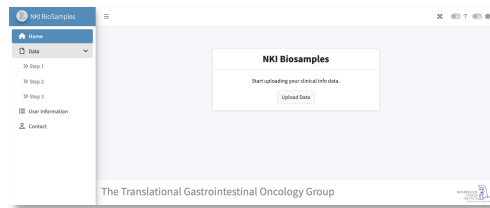


TGO data workflows summary

- External vs internal patients
- Blood samples
- Tissue samples
- Isolations (blood/tissue)
- Seq data
- Molecular data
- Clinical data
- Informed consent
- Imaging data



TGO Biosamples App



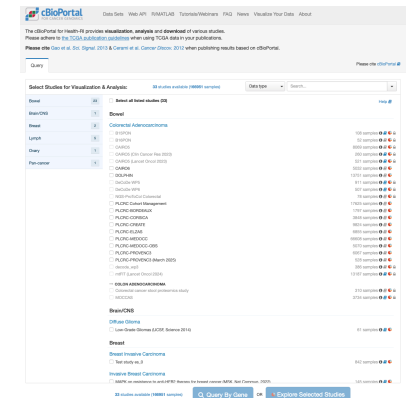
- Detection errors registration biomaterial and patient data.
- Integrates biomaterial info, collections time points, IC and minimal set of clinical variables.
- Allows active longitudinal follow up of patients per study.
- Dashboard to explore the data.

- Centralized all data in a db.
- Combine Biosample app output with IKNL-NKR clinical data and molecular data.



- The cBioPortal File Generator extracts data from the SQL database to generate cBio files ready to import data to cBioPortal

- Data is shared with researchers and doctors in cBioPortal to easily explore de data in a user-friendly environment



Collecting data

Cleaning/Curation

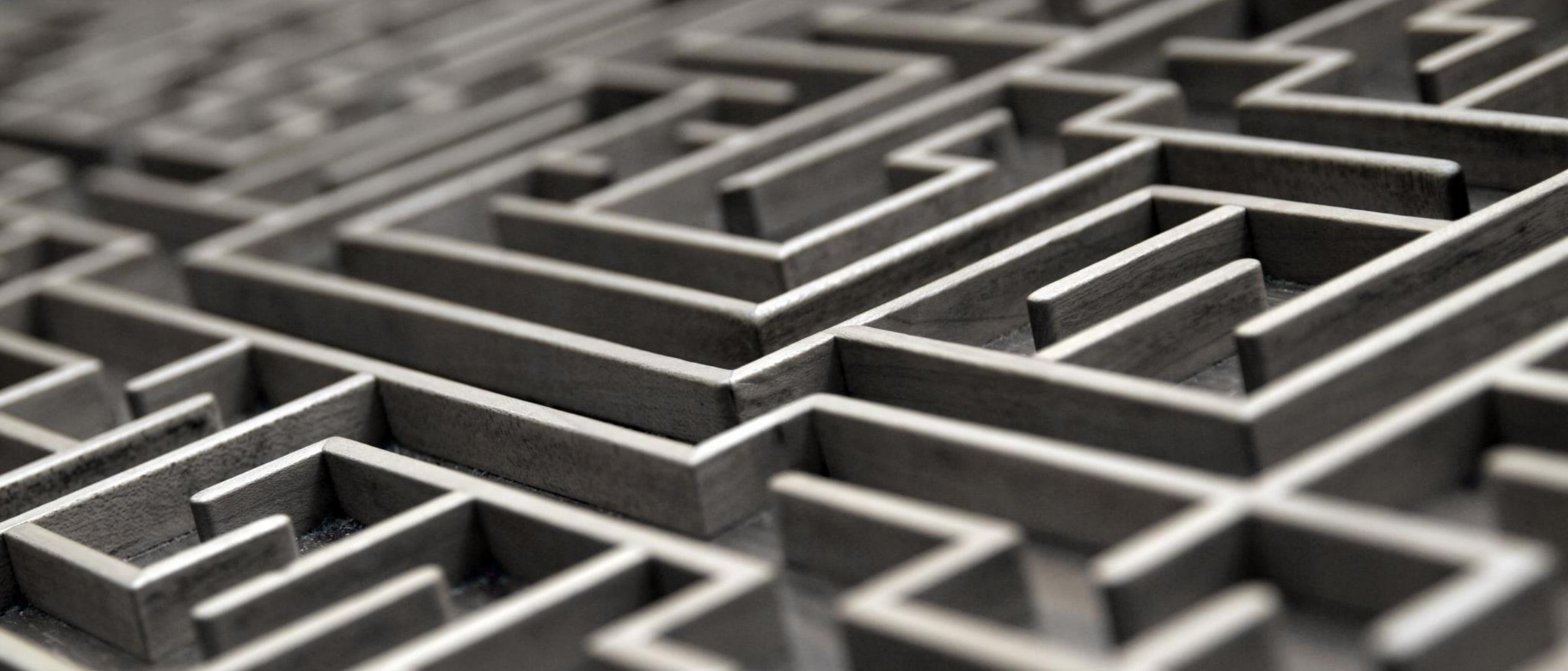
Data modeling

Data formatting

Data Distribution

DATA COLLECTION

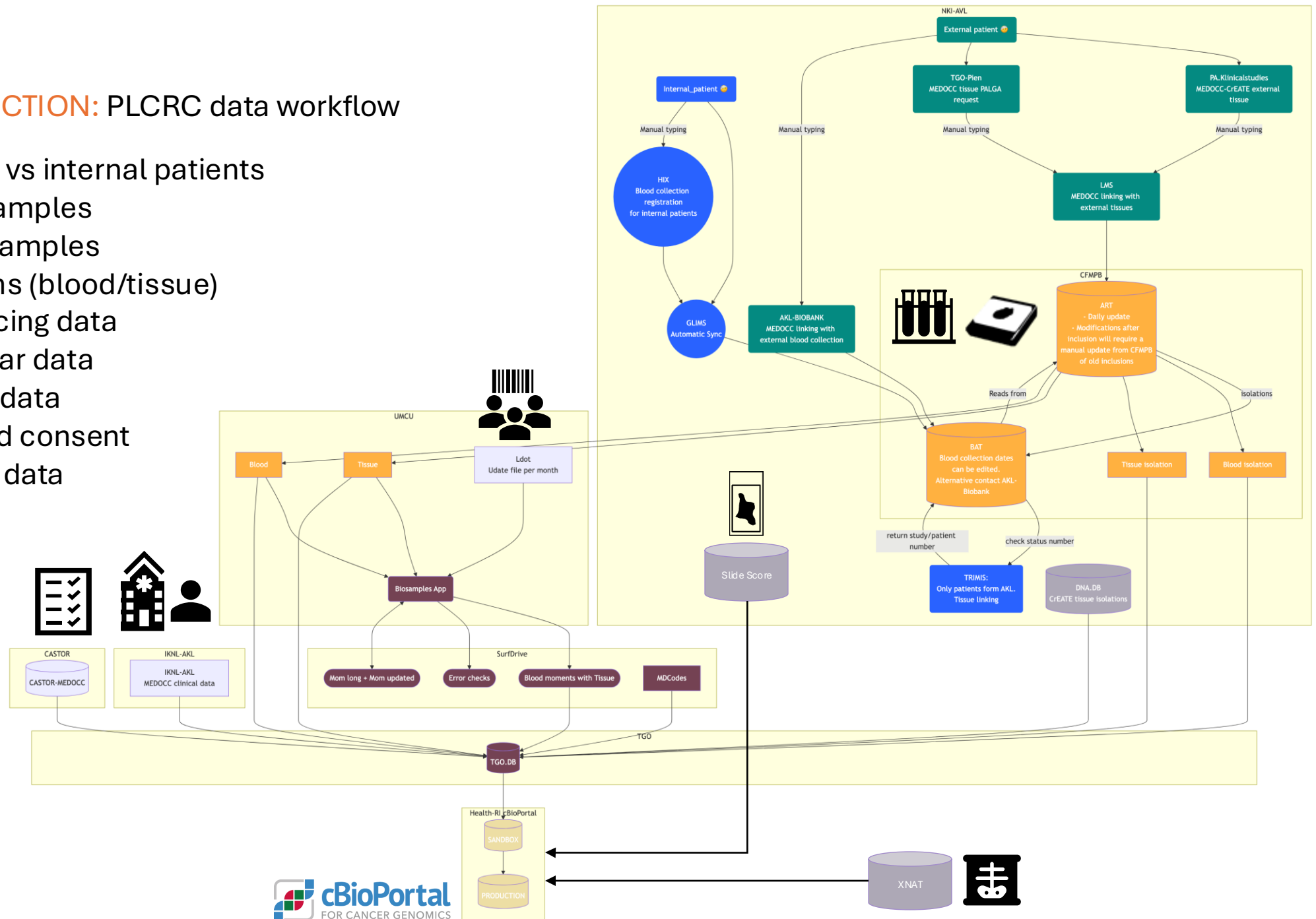




It's a challenging maze

DATA COLLECTION: PLCRC data workflow

- External vs internal patients
- Blood samples
- Tissue samples
- Isolations (blood/tissue)
- Sequencing data
- Molecular data
- Clinical data
- Informed consent
- Imaging data





Challenges:

- Care data is **not structured for research** use.
 - Example: different file formats, lack of standardization, Dutch vs English, etc.
- Data is **fragmented** (information is spread in multiple EDC systems and files).
 - Example: data comes from a lot of different places, some data overlaps but does not match, complementary data needs to be integrated, different ids depending on the origin of the data, etc.
- Data is **not monitored** in the different capture systems and **no validation** rules inside EDC systems (accumulation of **human errors**).

Dit bloed is afgenomen voor PLCRC/MEDOCC

PLCRC MEDOCC

Let op: MEDOCC-ID op dit formulier moet overeenkomen met MEDOCC-ID op buizen!

Alles volledig invullen!

MEDOCC-ID	MEDOCC - 243 - [redacted]
Geboortedatum en -jaar (mm/jjjj)	01 - 1952
Datum afname (dd/mm/jjjj)	10 - 10 - 2024
Afnamemoment aankruisen	☐ Voor Operatie ☐ Na Operatie (Algem. - Check na operatie) ☒ Follow Up
Ziekenhuis	Reinier de Graaf Ziekenhuis

Beschrijven met veldnummers en dit formulier, zo spoedig mogelijk verzenden naar NIO-AVL.
Zie aanwijzingen van dit formulier voor informatie en instructies.

Na te vullen door verzending MD-811

Datum ontvangst:

Paraf:

PLC/MEDOCC afnameformulier

Wersie 1.7 maart 2023



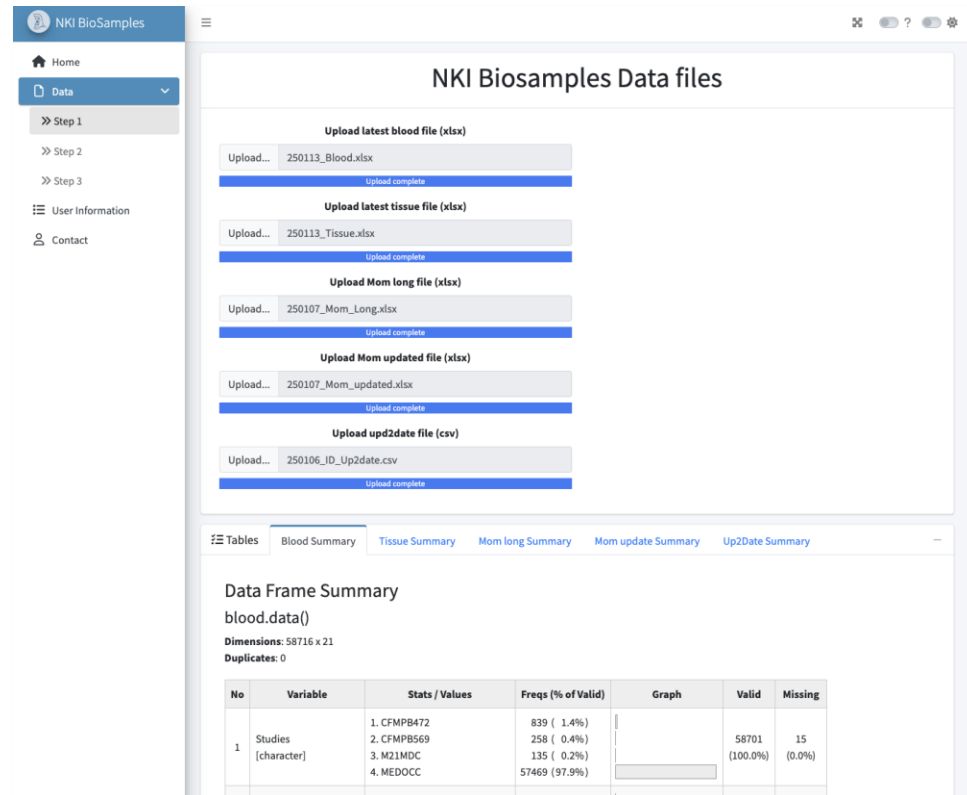
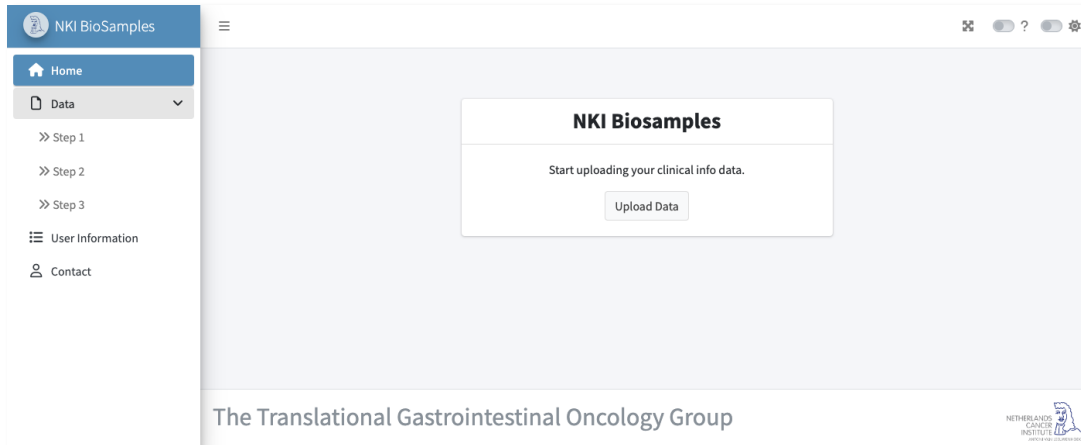
Error type	Description	Examples
ID formats	Medocc ids format is MEDOCC-XXX-XXXX or XXX-XXX	MED12300000123, MEDOCC-123-123, MED-123-123, MED-123-0123
Wrong ids	MEDOCC ids that does not exist on the study	MEDOCC-296-0020 vs MEDOCC-296-0019
IDs missing	All blood collections and tissues included on the study must have a MEDOCC id, sometimes is missing	
Intake moments incorrect	Intake moment should be registered for each blood intake. The variables to use are Na operatie, Voor operatie or Follow-up.	1954, preoperatief, preop, 2-3 weken postoperatief, 3 weken postoperatief, Eerste postoperatieve controle, MED-123-004 (postop afname), MED-123-005 (Controle-arm) postoperatief, post-op (MED-316-001), postoperatief 2-3 weken na operatie, volgt, 31-10-2024, MED-687-002 (t=6 mnd), 6 maanden postop, MED-567-151, vandaag
Incorrect dates	The arrival date of the sample to the hospital should always be the same or later than the collection date	Arrival dates happening before the collection, futuristic dates, dates from XIX century, dates does not match with the collection moment.
Decimal problems	Concentration problems how "." or "," are used depending on the machine.	Ng/ul concentration ranking from 1-10000 depending on the decimals used, also happening with 260/280 or 260/230 ratios.

DATA CLEANING AND CURATION: TGO Biosamples App



TGO Biosamples App

- Error report application
- Generate document for weekly follow up



Check list

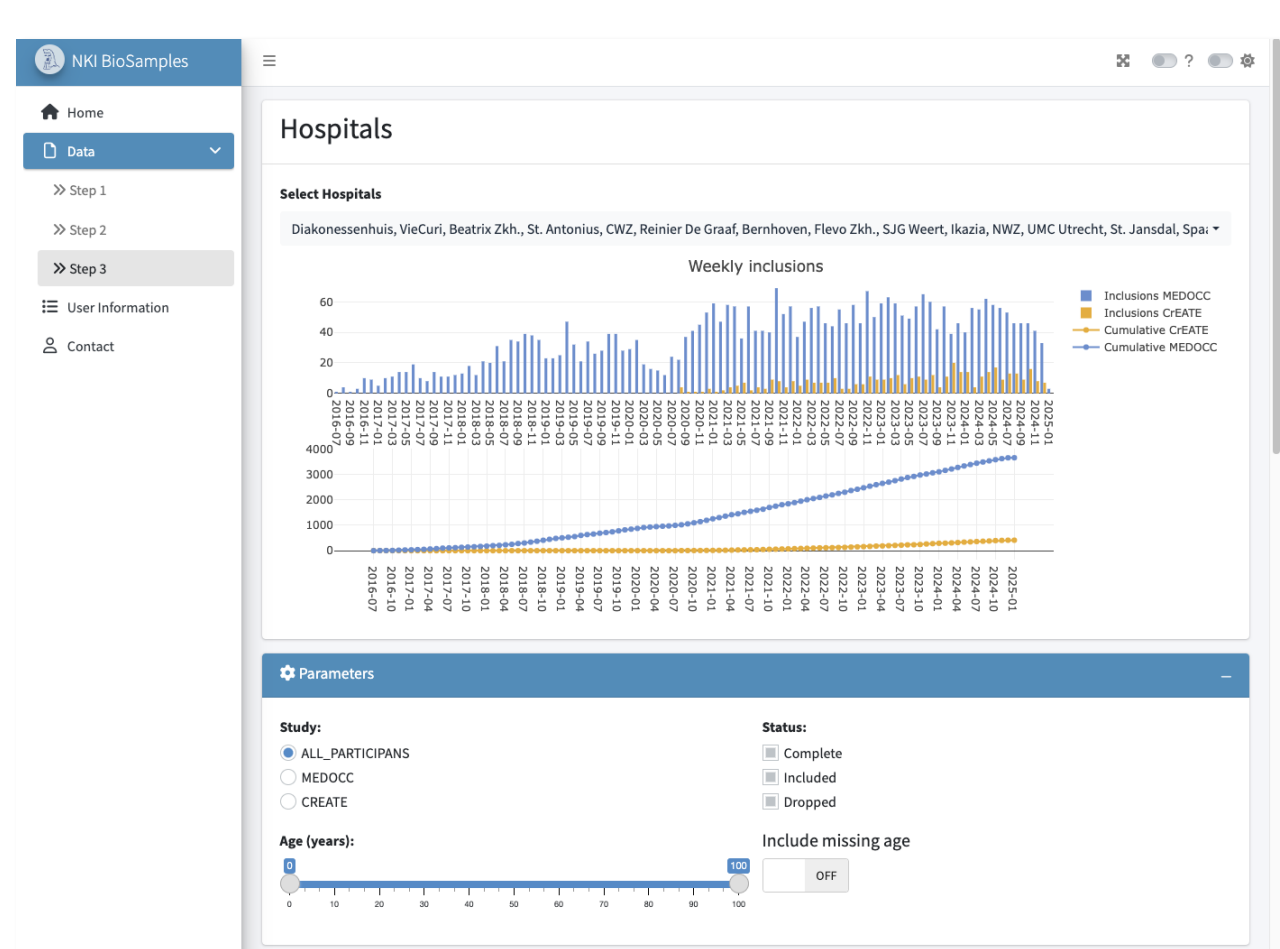
- Total number of Tube ID with NA 0 .
- Length unique MEDOCC ID 4540 .
- Length unique HHH_PPP 4540 .
- Hospital missing 0 .
- Collect date with NA 0 .
- Arrival date with NA 0 .
- Number of samples with an irregular interval date 21 .
- Sample types: Cell Pellet : 32482 Cell-free Plasma : 30828 NA : 0 .
- Collect Moment in ART: T12Peri : 3909 T1Pre : 14338 T2Post : 14124 T7_3FU : 30939 NA : 0 .

	HHH_PPP	Hospital_Abb	BIRTH_MY	STATUS_MEDOCC	STATUS_MEDOCC_DASHBOARD	TXSX_DATE_MEDOCC	BIRTH_DATE	BIRTH_COMPARE	IC_DATE	IC_VER
1			11-1943	Included	Complete	2021-07-14		1943	Similar	-2021
2			12-1948	Dropped	Dropped	2022-07-01		1948	Similar	-2022
3			11-1953	Included	Included	2022-07-04		1953	Similar	-2022
4			07-1945	Included	Included	2022-07-12		1945	Similar	-2022
5			01-1966	Included	Complete	2021-12-07		1966	Similar	-2021
6			09-1939	Included	Complete	2019-04-29		1939	Similar	-2019
7			01-1967	Dropped	Dropped	2021-07-05		1967	Similar	-2021
8			03-1955	Included	Included	2022-06-13		1955	Similar	-2022
9			08-1941	Included	Included	2022-05-04		1941	Similar	-2022
10			08-1958	Included	Complete	2021-12-17		1958	Similar	-2021

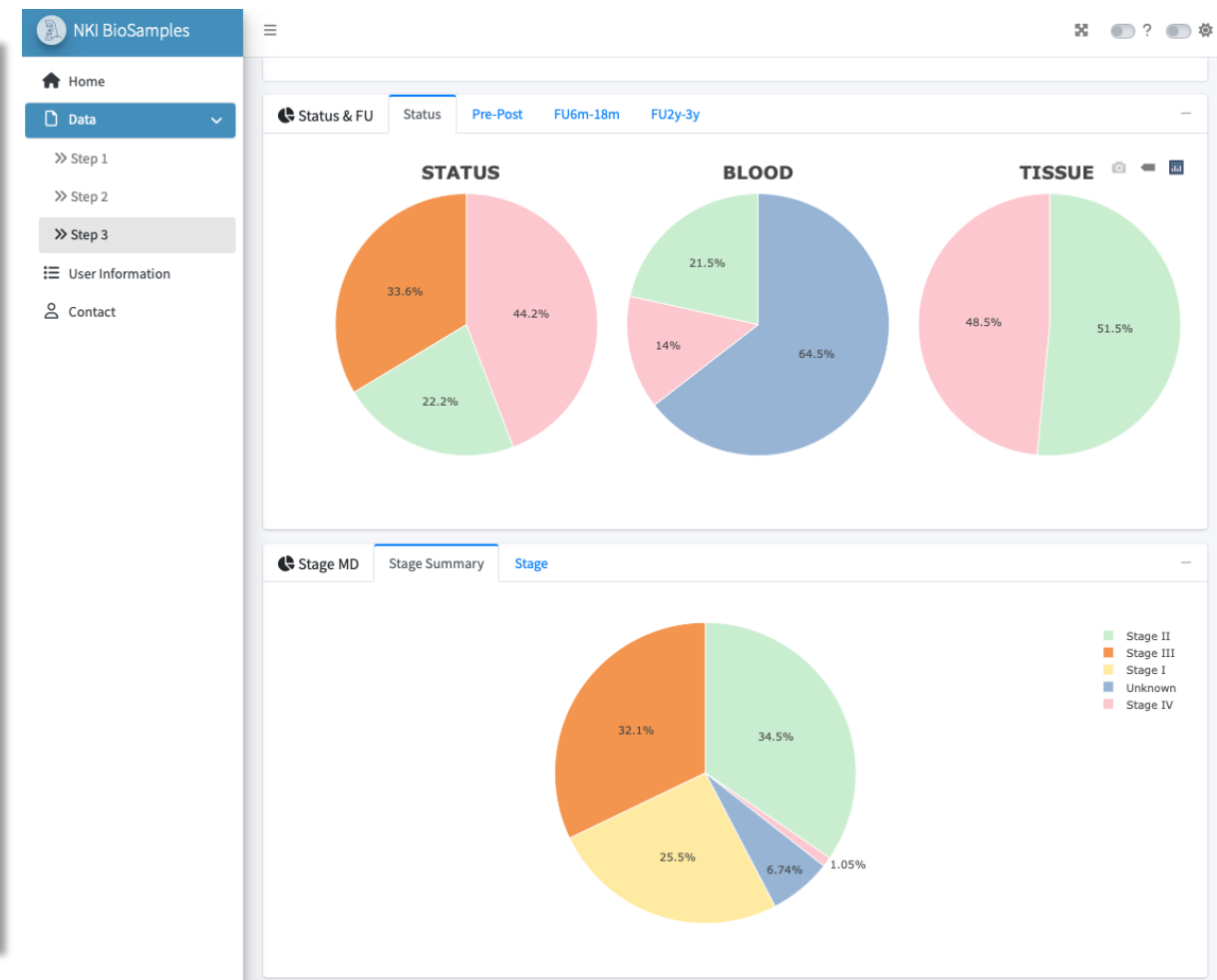
Showing 1 to 10 of 4,602 entries

Previous 1 2 3 4 5 ... 461 Next

Download Biosamples Download CheckDates Download Mom Updated Download Mom Long



- Follow up inclusion per study
- Parameters to filter at Study, Status, Age and Hospital level

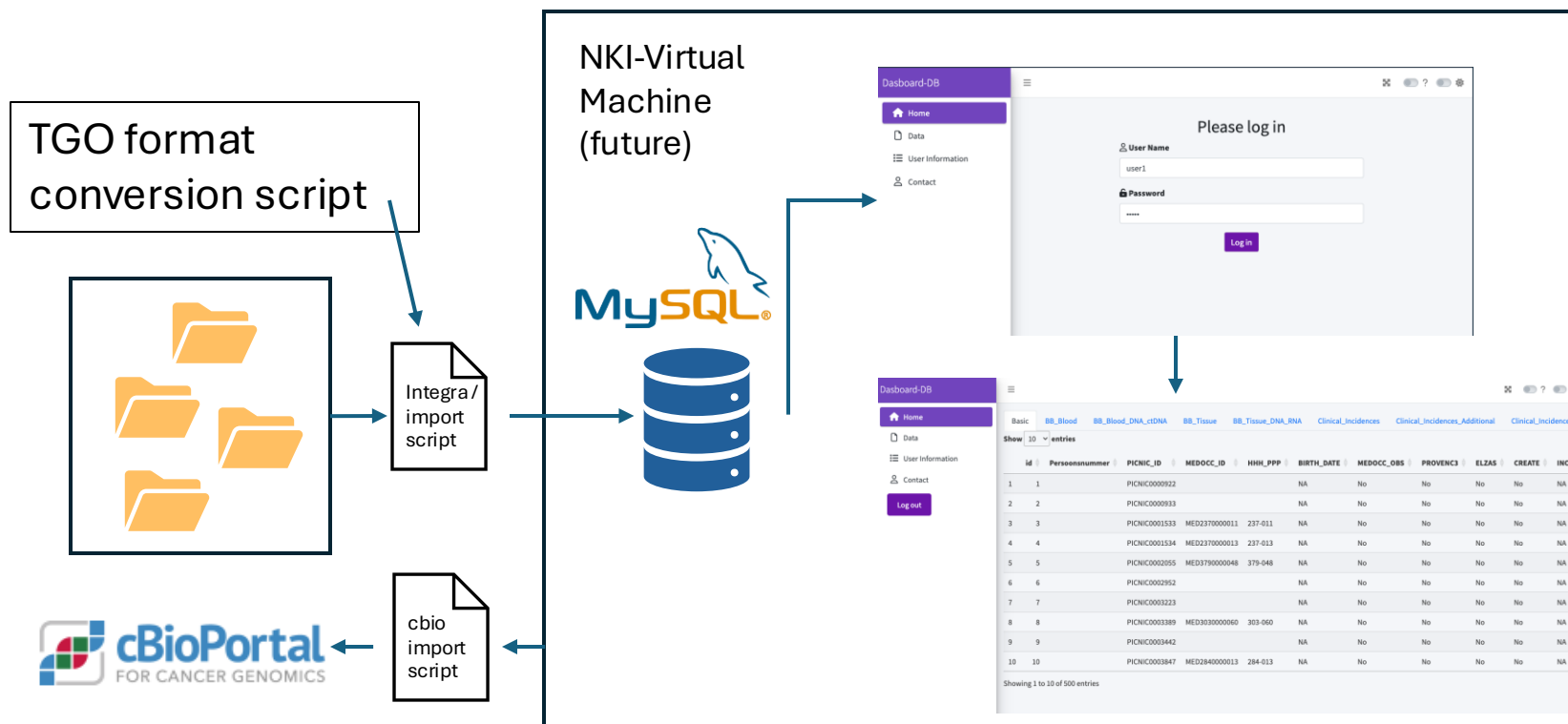


- Distribution of patient status
- Blood collection
- Tissue Collection
- Time points
- Stage Summary

MODELING AND STORAGE: The TGO database



TGO-Study-DB (prototype)



TGO-Study-DB

Home

Data

User Information

Contact

Please log in

User Name

user1

Password

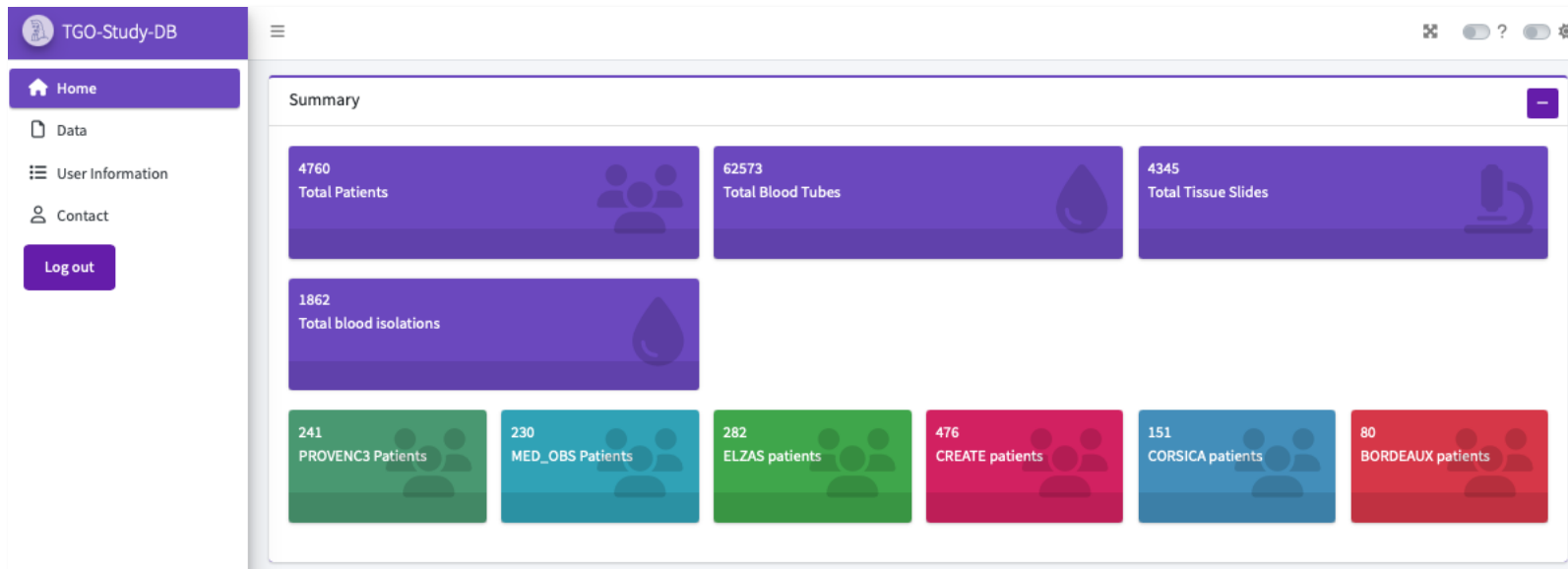
Log in

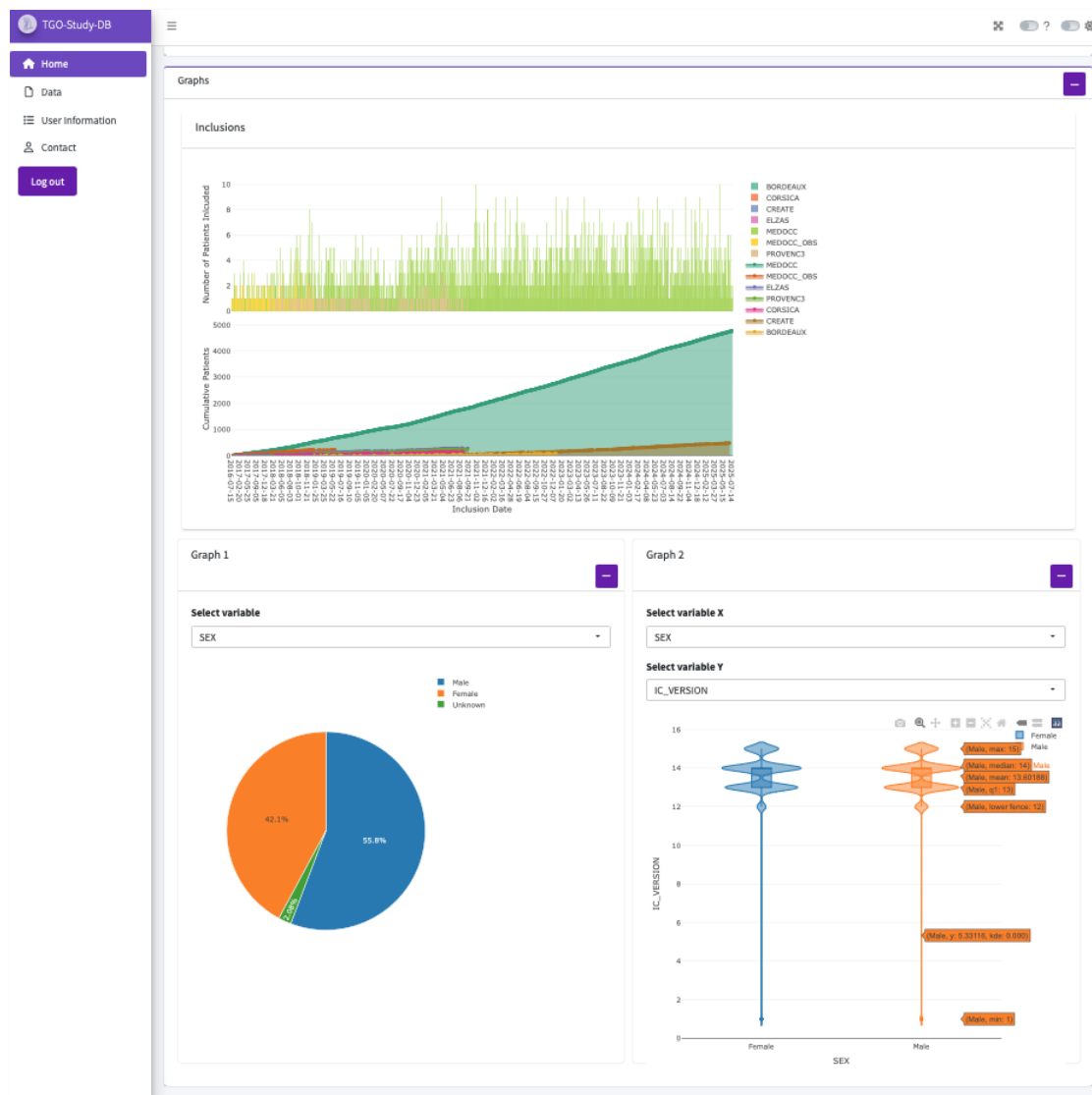
The Translational Gastrointestinal Oncology Group

NETHERLANDS
CANCER
INSTITUTE

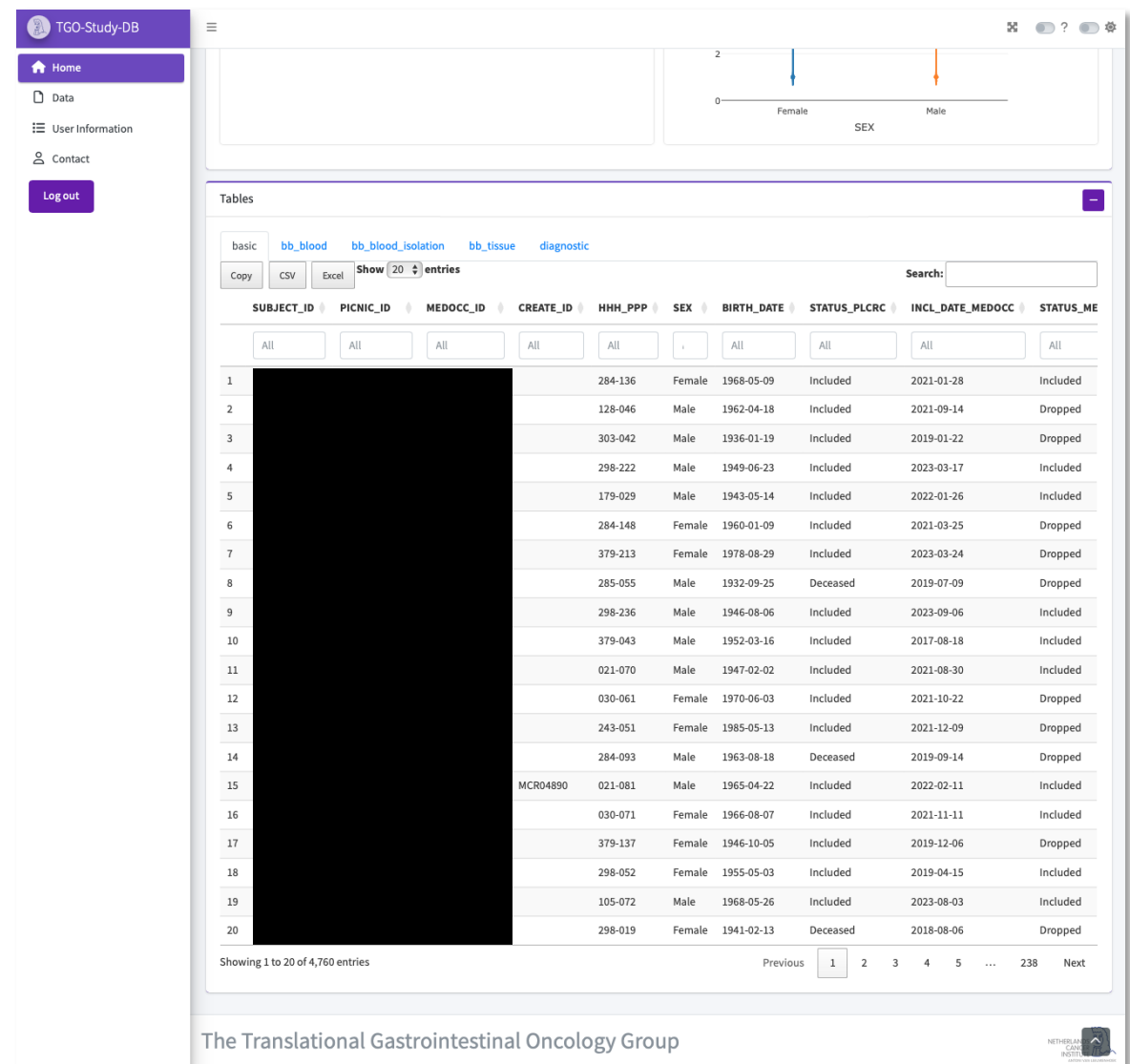
TGO-Study-DB (prototype phase)

- User management to control access
- Connected to a live MySQL database
- Summary page to give general info of the data collected in the DB



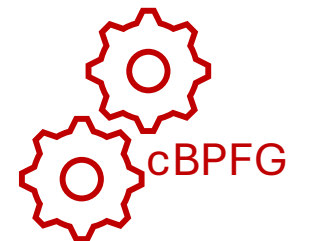


- Small dashboard to explore data.
- Inclusion graph to check the inclusion status per sub-study since the beginning of the study.
- Give tools to create graph base on variables collected on the database.



- Easy access and visualization of the data collected in the database (allow to apply filters per variable).
- Export buttons to share the data with researcher under request.

FORMATTING AND DISTRIBUTION: cBioPortal File Generator (cBPFG)



cBioPortal File Generator (cBPFG)

Data types cBioPortal:

- Meta data
- Clinical patient
- Sample data

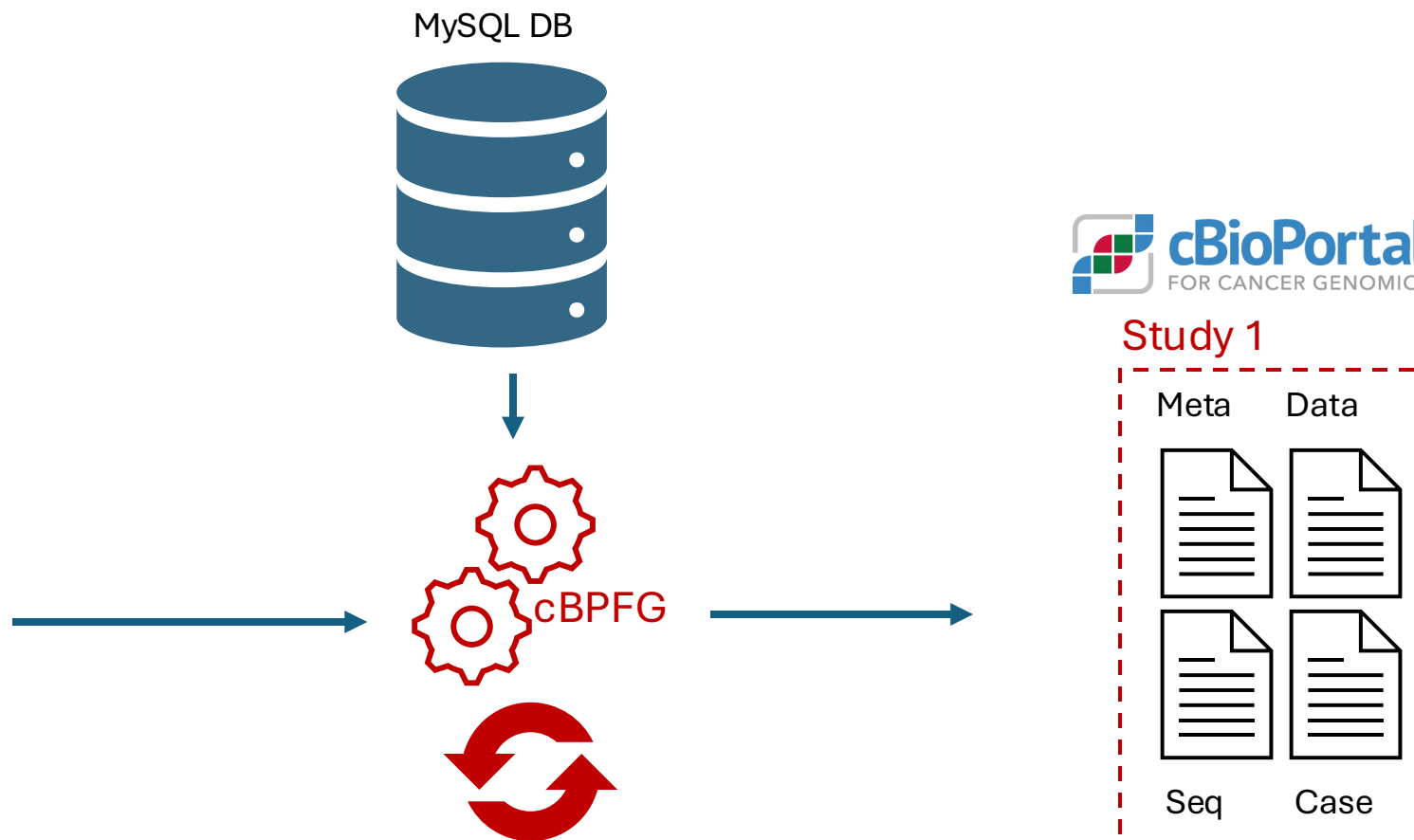
Mandatory

- Timelines
- Sequencing data
 - Mutations
 - Discrete/continuous CNV
 - SV
 - Segmented
 - Expression
 - Methylation
 - Fusion
 - Protein

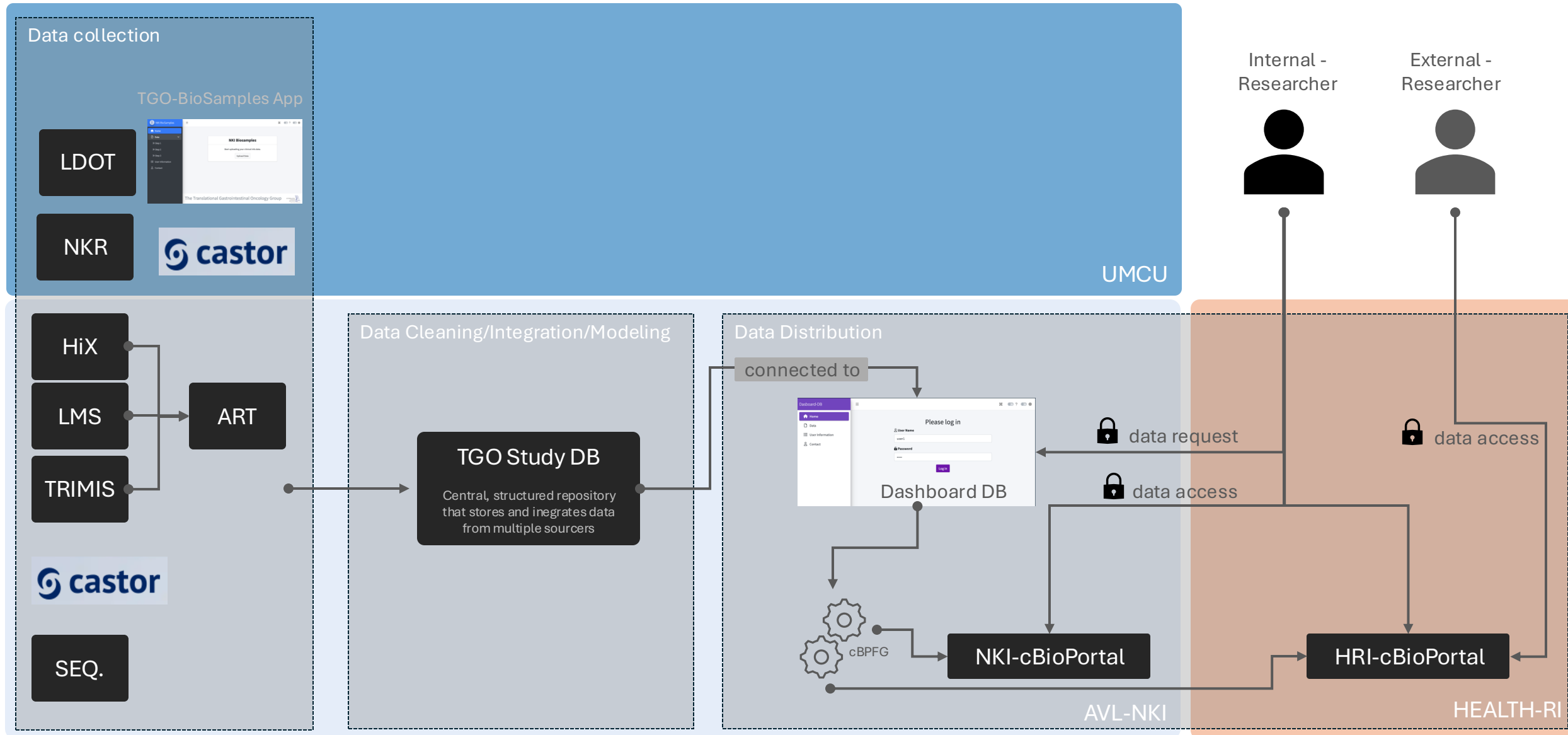
Optional

- Case list

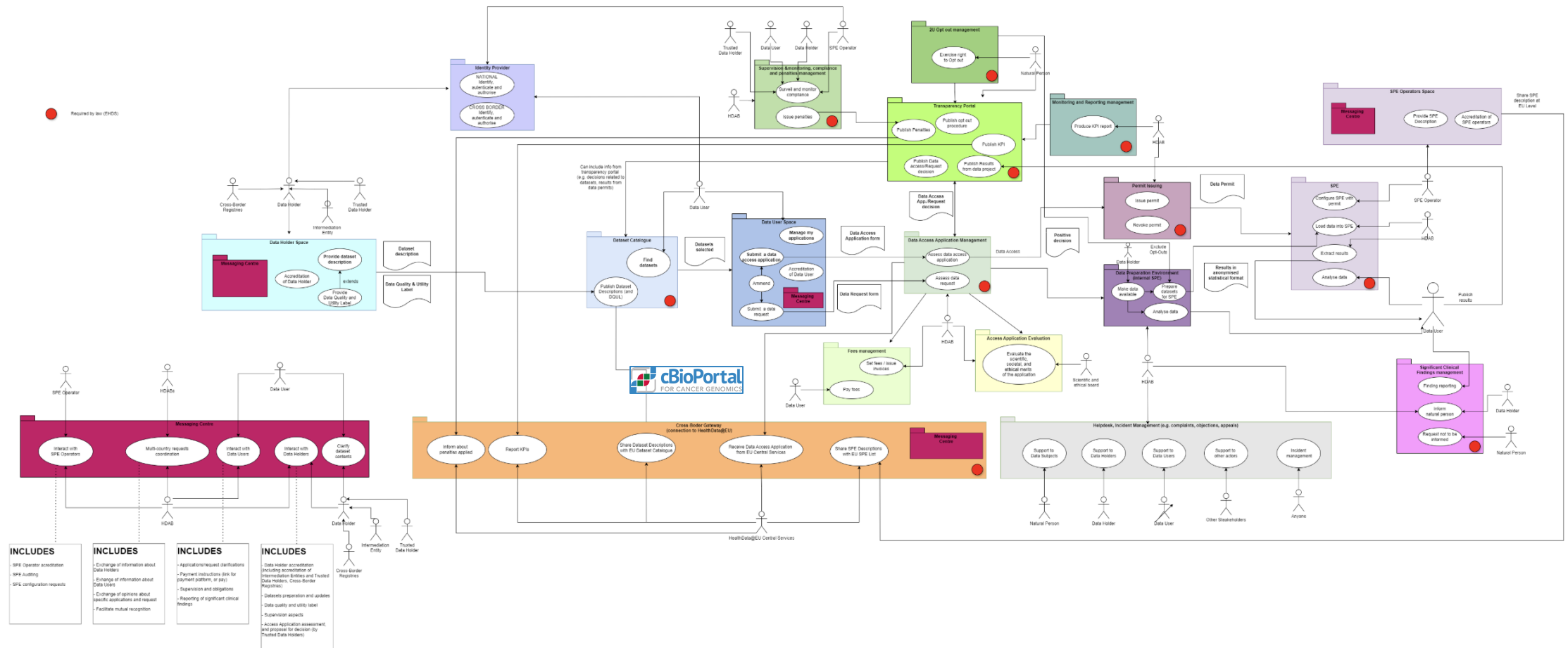
Mandatory



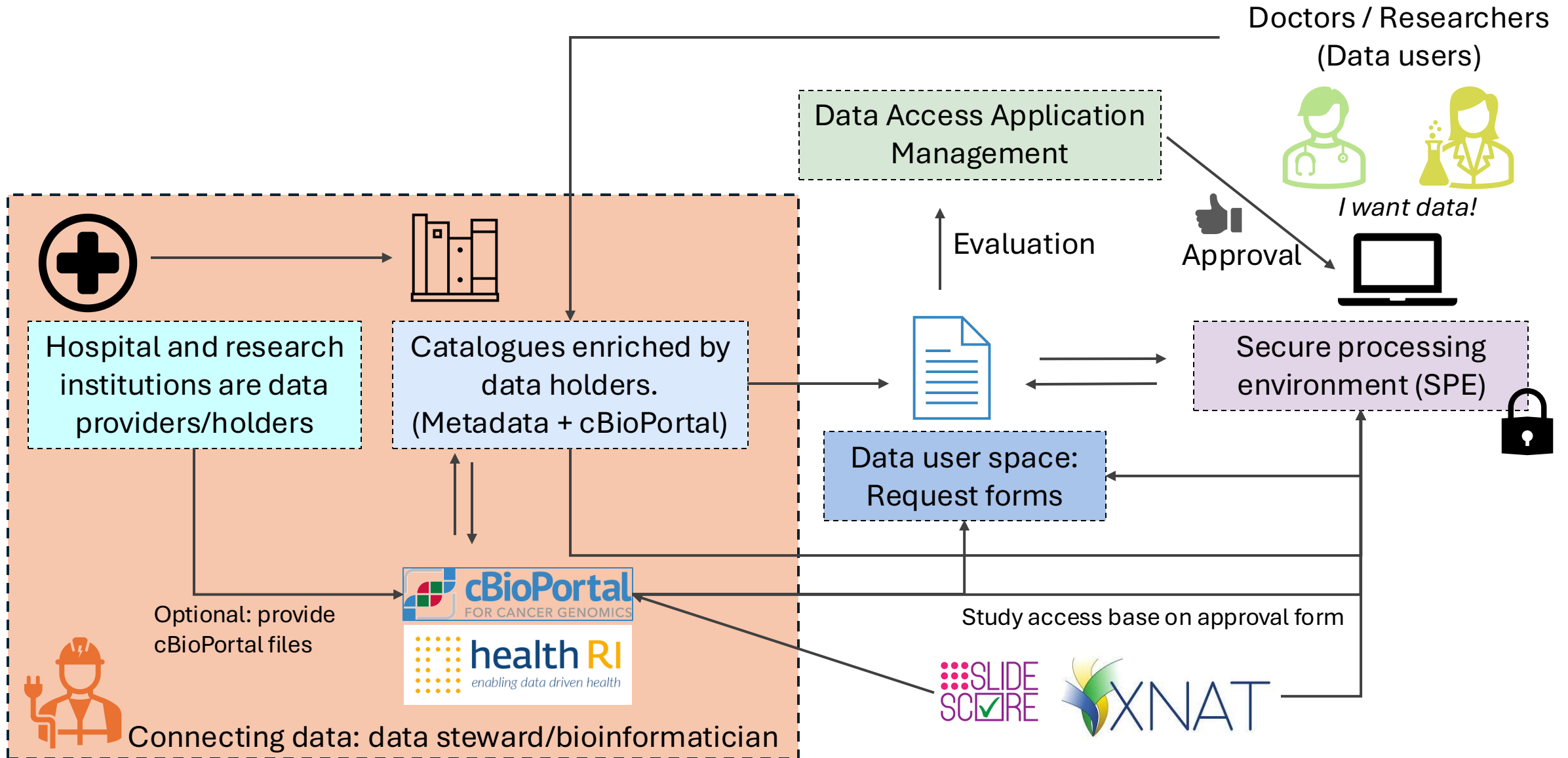
PLCRC data workflow status in 2025



How would the EHDS business capabilities blueprint align with the PLCRC data workflow?



Summary: Deployment of cBioPortal in the EHDS DBC blueprint



How would the EHDS business capabilities blueprint fit with the translational research data journey?

Data holder collecting study data.

Responsible to **clean** and **model** the data to provide a **good quality** datasets.

Generate cBioPortal ready files

Catalogue + cBioPortal:

cBioPortal is used as a catalogue.
PLCRC datasets are uploaded at the national Health-RI cBioPortal and researchers can request access to the datasets

Data access application management:

Health-RI are already managing any data request via forms coming from cBioPortal (TopDesk)

Secure processing environment (SPE).

cBioPortal can be accessed via browser with a user account. But easy to connect to any SPE

“PLCRC’s translational research substudies demonstrate how national infrastructures (IKNL, PALGA, etc.) and the Health-RI platform can deliver FAIR, high-quality research data at scale. This blueprint is easily transferable to other studies and offers a solid foundation for the expansion of the EHDS”

Is cBioPortal a good tool for EHDS?

Pros	Cons
Central place for clinical, sample and molecular data.	Gene centric, limited functionalities for isoform or SV.
Visualization of additional data from other tools via resource linking (imaging data).	Annotation dependent on external sources, needs to be limited the access in close environments.
Can be used to explore data sets but at the same time work as a catalogue.	Expertise is needed to create the cBioPortal ready files from raw data.
Build in a variety of analysis tools, can be even connect to to external tools like Galaxy.	Automatization can be implemented but each data holder needs to model the data before or work with data steward/Bioinformatician to model the data an make it available
User friendly, no need technical skills to use the tool.	
Open-source tool witch high capacity for customization, expandability end scalability.	
Allows user management access.	

Data Team:

*Mariska Bierkens
Lana Meiqari
Adria Closa
Iris Huitink*

Research Team:

*Gerrit Meijer
Remond Fijneman
Beatriz Carvalho
Meike de Wit
Steven Ketelaars
Denise van Steijn
Renske de Wit
Romy Bos*

Technicians Team:

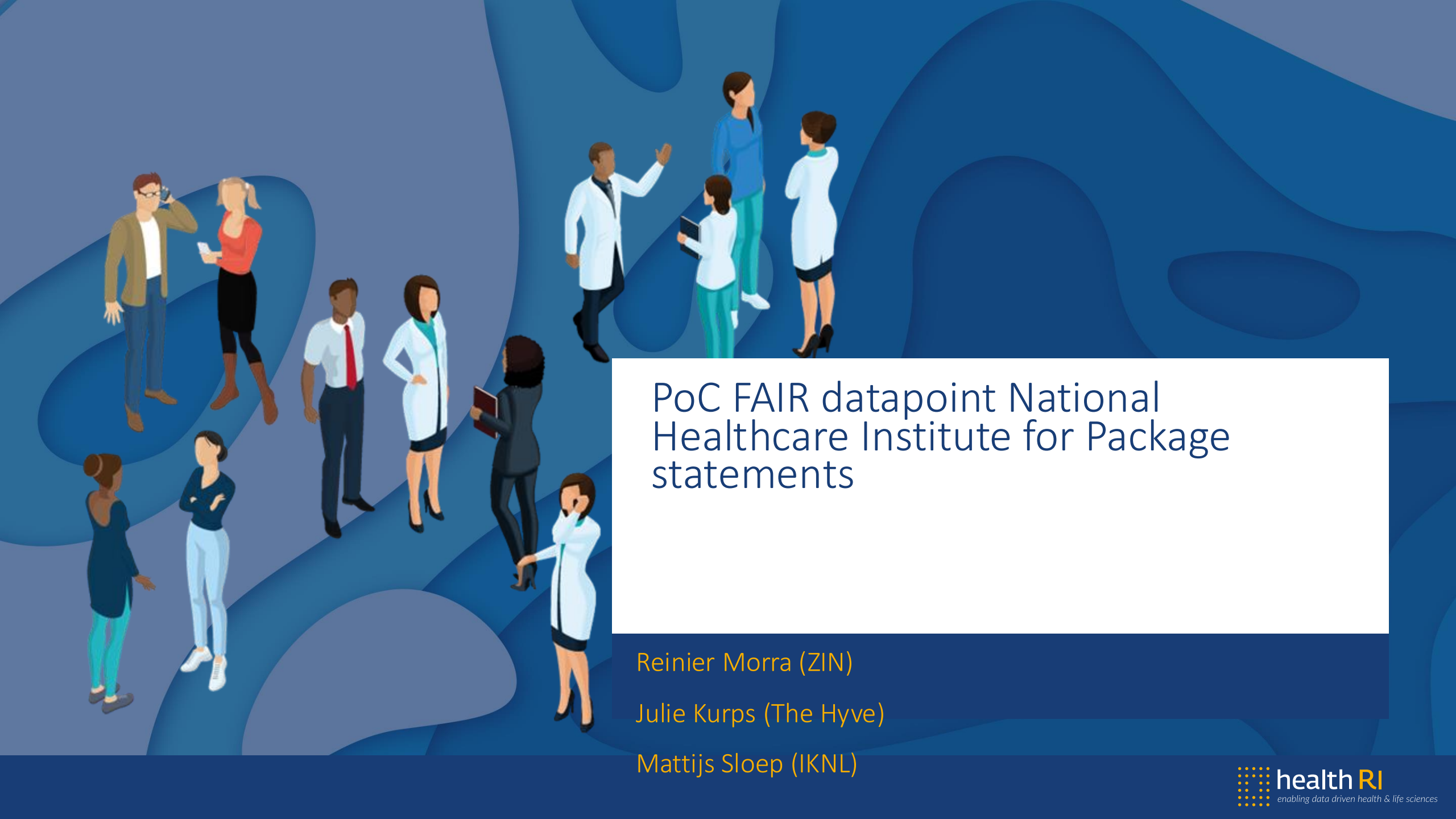
*Pien Delis-van Diemen
Marianne Tijssen
Margriet Lemmens
Anne Bolijn*



Contact: a.c.mosquera@nki.nl
Room H.02.009



aclosa
NKI-TGO-group

An illustration of a group of diverse healthcare professionals in a meeting. There are ten people in total, including men and women of various ethnicities. Some are wearing white lab coats, while others are in business casual attire. They are standing on a blue background with large, abstract, wavy shapes. One man is on a phone, another is holding a clipboard, and others are gesturing or looking at each other.

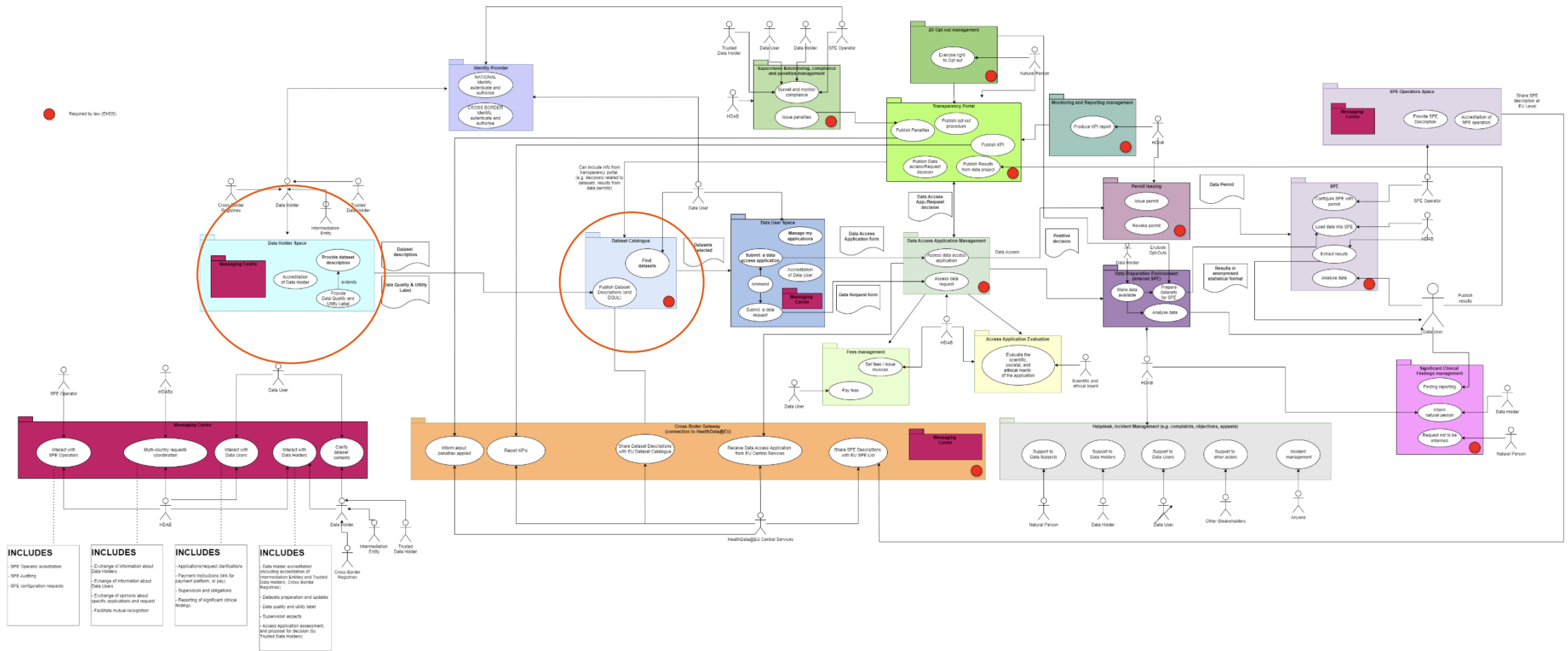
PoC FAIR datapoint National Healthcare Institute for Package statements

Reinier Morra (ZIN)

Julie Kurps (The Hyve)

Mattijs Sloep (IKNL)

Backbone



Research project FAIR package registry

- Introductie onderzoek: Doelstelling, problem, aanpak, PoC
- Resultaten
 - Modelling
 - Infrastructuur PoC FAIR Pakketregister
 - Behoeft onderzoek op basis interviews
 - Use cases:
 - Signalering/Nieuw bewijs
 - Evalueren & Monitoren met IKNL
 - BIA Evaluatie
- Inpassing binnen het Zorginstituut en IKNL
 - Mogelijke rol/relatie met cyclisch pakketbeheer
 - Kansen en risico's
 - FDP voor data en documenten van het Zorginstituut
- Conclusies, discussie & vervolg

DISCLAIMER

Het is een onderzoeksproject.
Intern nog geen besluit genomen om het
FAIR Pakketregister in productie te nemen.

Vandaag bedoeld om methodiek en
lessen te delen en te horen hoe jullie
toepasbaarheid zien.

Doelstelling

Context: rol van het Zorginstituut in cyclisch pakketbeheer

datagedreven systeem om pakketuitspraken te monitoren en te evalueren (real world data)

Pakketuitspraken stroomlijnen

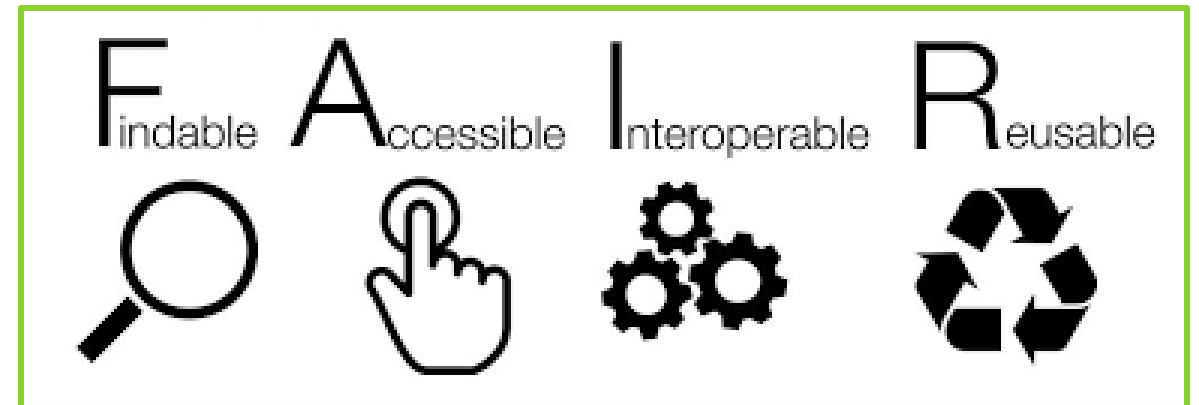
om informatie beter beschikbaar maken

Focus
vandaag

FAIR: machine-readable, machine-actionable

PoC: Borstkanker use cases:

- Signalering
- Real World Data (IKNL) voor een lerend zorgsysteem
- Budget-Impact-Analysis (BIA)



Focus vandaag

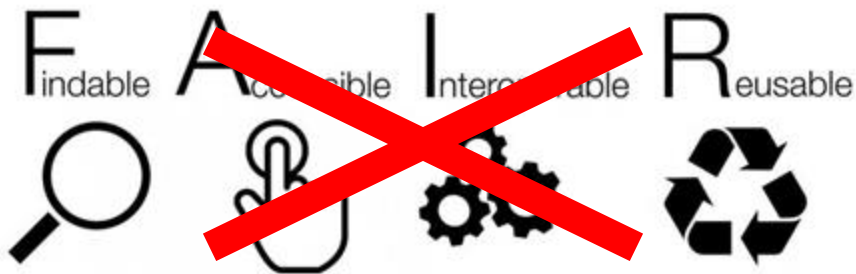
Aanzet cyclisch pakketbeheer met FAIR pakketregister: Wens voor meer monitoren en evalueren passende zorg



Het probleem

Pakketbesluiten niet FAIR

- Gepubliceerd op de website van het Zorginstituut, niet goed doorzoekbaar
- Als PDF documenten:
 - Ongestructureerde data: kan moeilijk hergebruikt of geanalyseerd worden
- Niet FAIR!

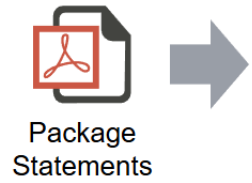


Praktijkdata beperkt beschikbaar voor M&E

- RCT niet altijd representatief voor de praktijk
- Momenteel geen / beperkte data toets of een uitspraak te monitoren en evalueren is op basis praktijk data
- Behoeftte aan zowel overall survival (OS) en kwaliteit van leven (QoL) data
- Voorbeeld: PARP remmers

Modelling

FAIRification process



Consulting experts

Identify relevant (meta)data

Datum 10 juli 2024

Betreft Pakketadvies herbeoordeling sacituzumab govitecan (Trodelvy®)

Onze referentie

2024026955

Geachte mevrouw Agema,

Zorginstituut Nederland adviseert u hierbij over de uitkomst van zijn herbeoordeling van het farmaco-economisch dossier van sacituzumab govitecan (Trodelvy®) voor de behandeling van volwassen patiënten met inoperabele of gemetastaseerde triple-negatieve borstkanker (mTNBC)¹ die twee of meer eerdere systemische therapieën hebben gekregen, waaronder één lijn taxaanbevattende therapie en ten minste één voor gevorderde ziekte. Dit advies volgt op het eerdere, positieve vergoedingsadvies d.d. 15 juli 2022. Het Zorginstituut liet u daarin weten dat sacituzumab govitecan bij de eerdergenoemde indicatie voldeed aan de stand van de wetenschap en praktijk, en in het verzekerde pakket kon worden opgenomen indien bij prijsonderhandelingen een korting van tenminste 75% werd overeengekomen.

Concept Statements

Pakketuitspraak (PU)

- Een PU (focus van dit werk) verstaan we
 - Vanuit het Zorginstituut:
 - Standpunten (open pakket);
 - Pakketadviezen voor voornamelijk geneesmiddelen (gesloten pakket, sluis);
 - Een Zvw geschil kan een aanleiding zijn voor een [advies](#) (buiten scope)
 - Duidingen (veelal in de eerstelijns zorg) onder welk wetsdomein zorg valt zijn buiten scope.
 - Vanuit Zorgverzekeraars Nederland (buiten scope, behalve BIA-gedeelte)
 - ZN maakt o.a. gebruik van de commissie BOM als er geen PU is vanuit ZIN. Zie [hier](#) voor borstkanker de actuele adviezen.
- Behandelt een vraag voor vergoeding (= 'vergoedingsvraag', kan ook door ZIN geagendeerd worden) van een behandeling vanuit het basispakket
 - Geneesmiddel vergoedingssysteem (GVS) geneesmiddel, of
 - Medisch specialistisch geneesmiddel (voor Medisch Specialistische Zorg, MSZ)
 - Ook voor medisch-specialistische behandelingen (AFT pakketuitspraak) = duiding
- De vergoedingsvraag kan beantwoord worden met een standpunt en eventueel gevolgd met een pakketadvies.
 - De beantwoording van vergoedingsvraag begint met een duiding.
 - Een duiding is een toets op SWP (stand van Wetenschap en Praktijk), [en](#) [eventueel andere pakketcriteria](#)
 - SWP: Zorg voldoet aan SWP als het in voldoende mate bewezen effectief is in vergelijking met de standaardbehandeling of gebruikelijke behandeling (relatieve effectiviteit)
 - Een duiding leidt altijd tot een standpunt.
 - Aanvullend op een standpunt kan ook een pakketadvies gegeven worden.
 - Bij een pakketadvies worden (volgens op de SWP-toets -> SWP is een knock-out criterium) ook de andere pakketcriteria gewogen.
 - De 4 pakketcriteria zijn:
 - 1 noodzakelijkheid (moet je je ertegen verzekeren, en hoe hoog is de ziektelast?),
 - 2 effectiviteit (is SWP, om deze te toetsen wordt het

MR Morra, R.

[Overzicht sluismiddelen waarover het Zorginstituut heeft geadviseerd | Zorginstituut Nederland](#)

GM Gothlin, M.P.

Een duiding (beoordeling) is een proces, met een standpunt of advies als resultaat.

GM Gothlin, M.P.

Meestal, ook andere pakketcriteria kunnen getoetst worden.

E eelke

of?

MR Morra, R.

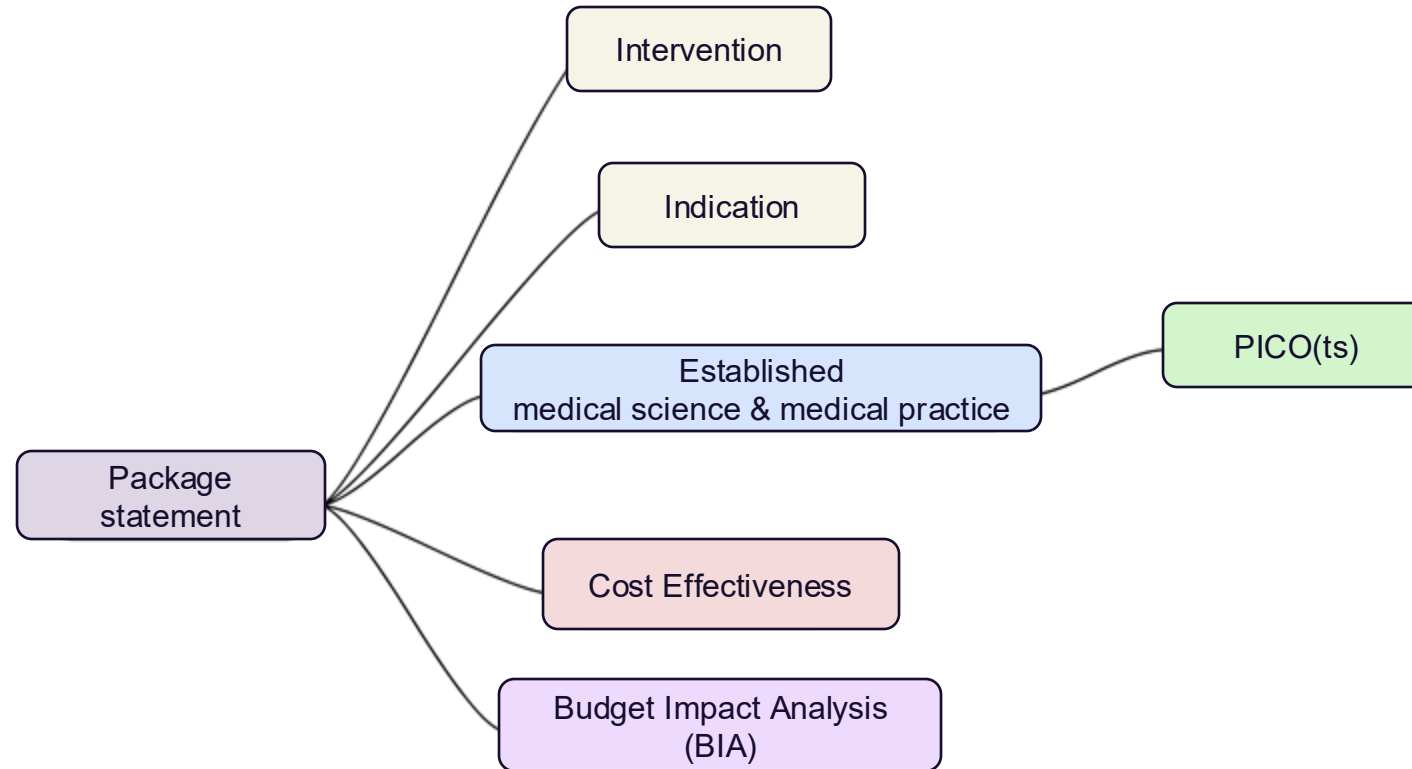
Een advies over een sluisgeneesmiddel van het Zorginstituut gaat altijd samen met een advies over een mogelijke prijsonderhandeling en afspraken over [See more](#)

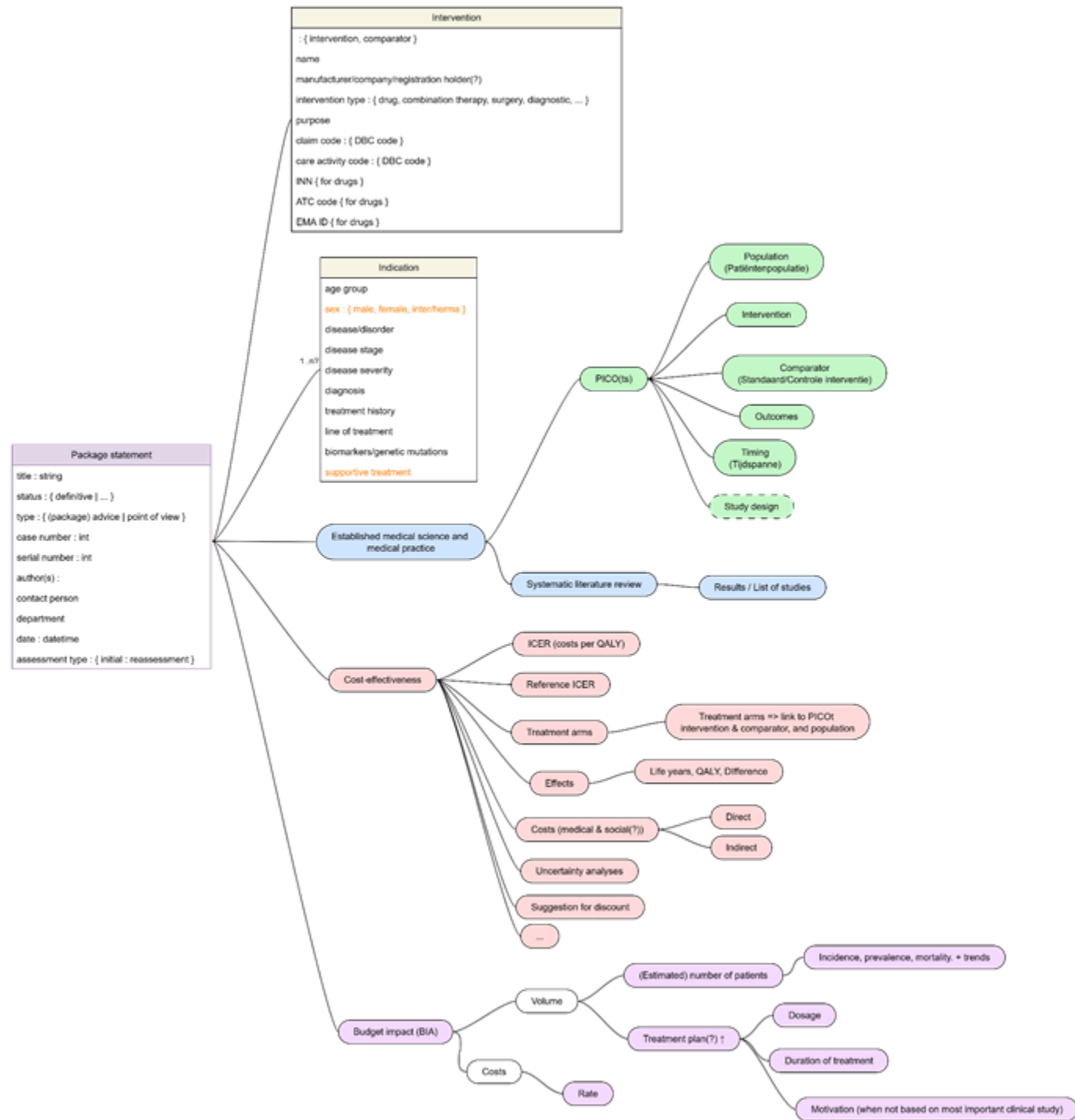
GM Gothlin, M.P.

Is het zorg zoals de beroepsuitoefenaars plegen te bieden?

✓ Resolved



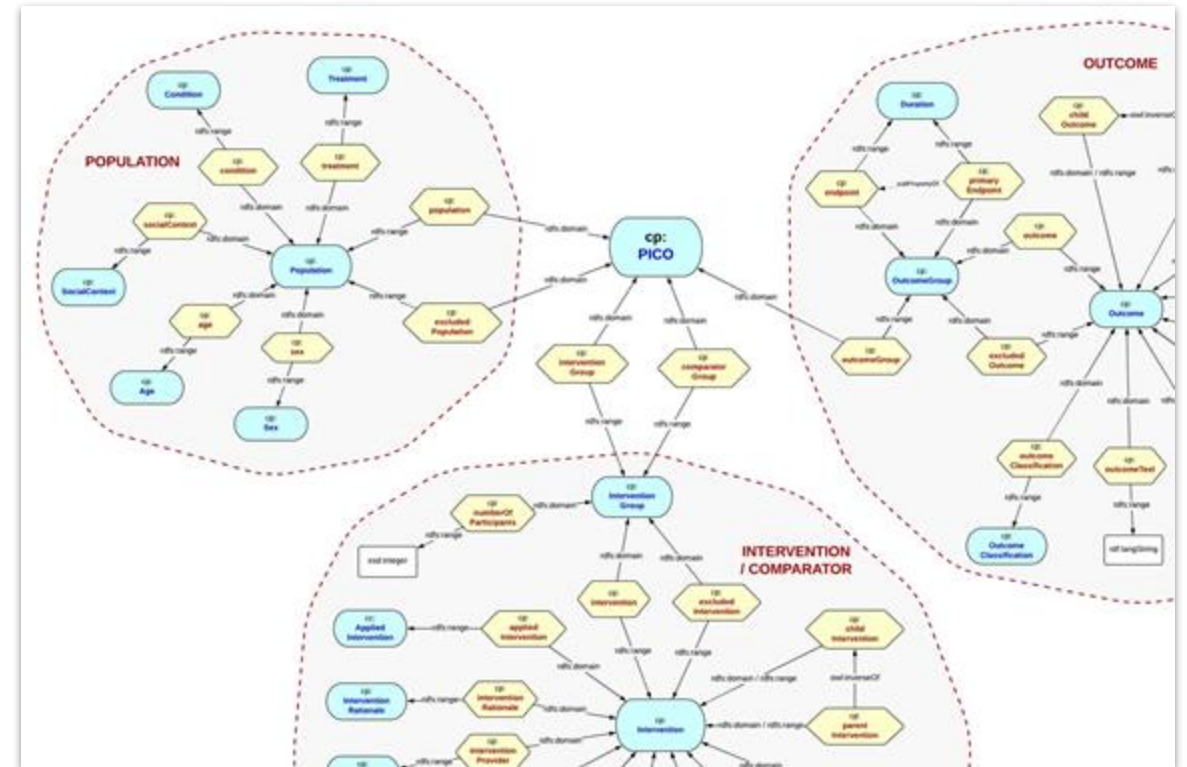






Re-use of existing standards

- Explore ontologies and vocabularies to represent (parts of) the information in the package statements
- Examples:
 - Cochrane PICO ontology
 - Ontology of biomedical investigations for study data
 - DataQube ontology for data sets



(Example: PICO ontology)

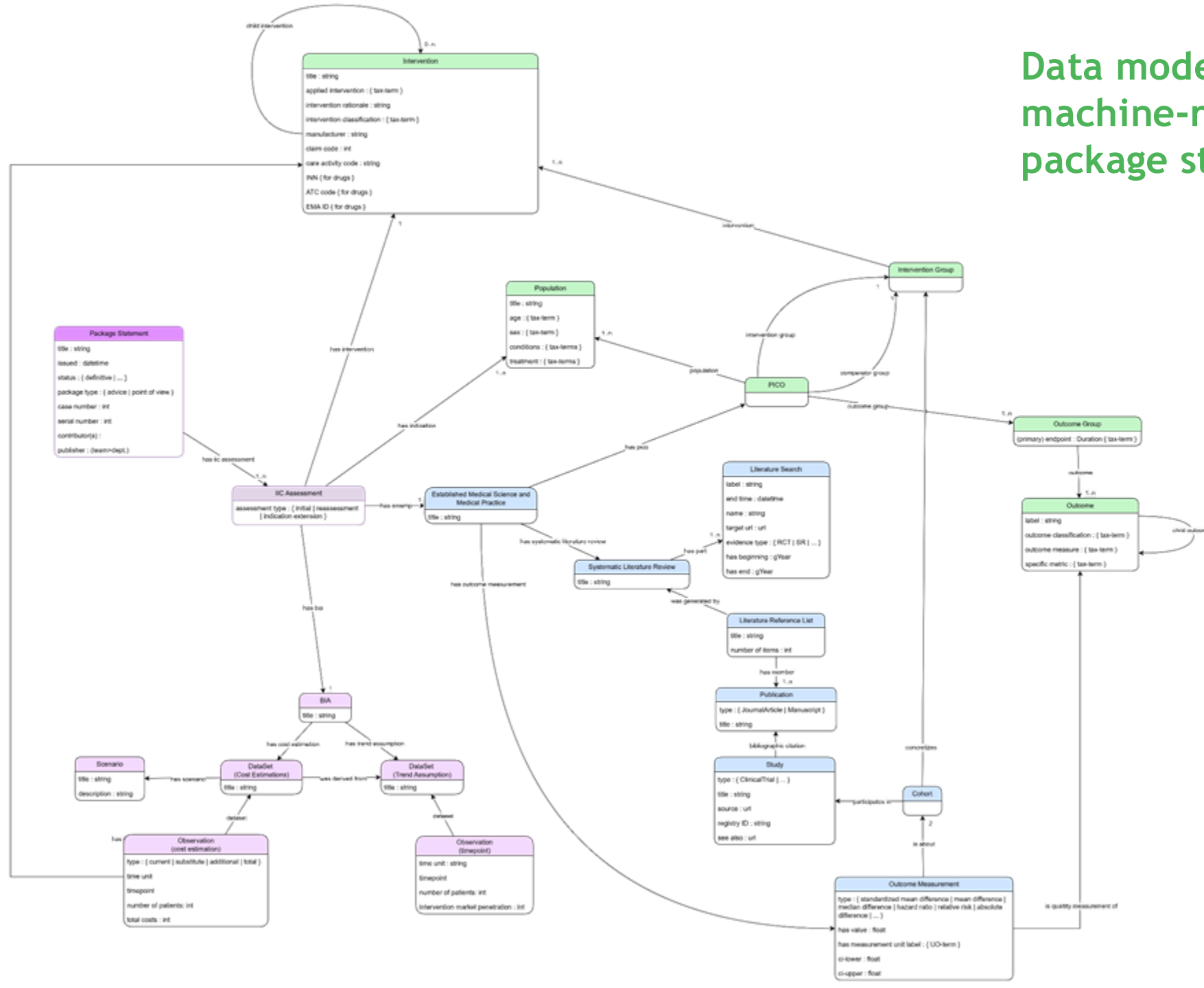
Keuze proces ontologiën - hergebruik waar mogelijk

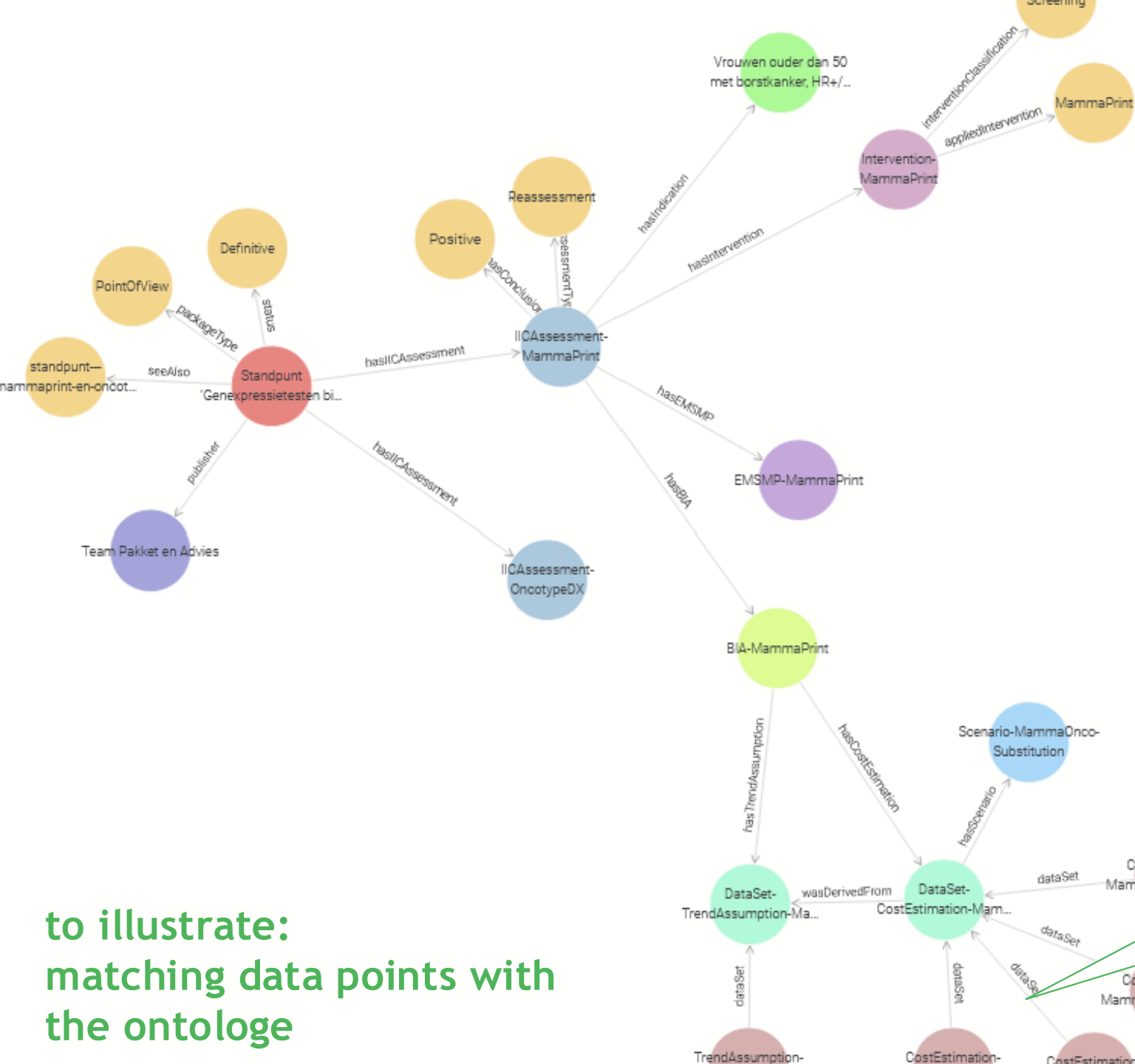
Onderdeel	Overweging	Keuze	Afgevallen
Interventie, Indicatie, en de PICO	1. Dekt de lading 2. Semantische representatie i.t.t. OMOP en SDTM 3. Gerenommeerd instituut 4. Eenvoud in gebruik voor zoekvragen i.t.t. ERGO	Cochrane PICOts + eigenschappen voor de use cases, zoals bijvoorbeeld prijs en codes voor de interventie	• OMOP (Observational Medical Outcomes Partnership) • SDTM (Study Data Tabulation Model) • Eligibility Rule Grammar and Ontology (ERGO)
Taxonomie	1. Aansluiting op de Cochrane PICO ontologie 2. beschikbaar is als Linked Data met verwijzingen naar andere vocabulaires, zoals MedDRA en UMLS (die weer verwijzingen hebben naar o.a. SNOMED-CT) 3. Meer van de benodigde termen bevatte dan bijvoorbeeld SNOMED-CT	Applicatie-specifieke taxonomy op basis termen Cochrane Core vocabulaire + de extra benodigde termen	Eligibility Rule Grammar and Ontology (ERGO)
Literatuur-onderzoek	Geen 1 omvatte ontologie die de informatie over het literatuuronderzoek kan weergeven, zoals zoekopdracht en resultaten (als lijst weten-schappelijke artikelen die verwijzen naar studies)	Documenten: Provenance ontologie (PROV-O) Activiteiten en uitkomsten: Schema.org Literatuurreferenties: FABiO	BIBO (Bibliographic Ontology) [Had ook gekund]

Keuze proces ontologiën - hergebruik waar mogelijk

Onderdeel	Overweging	Keuze	Afgevallen
Uitkomstmat en en Studie	- Inzet STATO omdat verschil nodig tussen de uitkomstmaat uit de PICO ontologie en de waarde van een uitkomst zoals gerapporteerd in een specifieke studie of in meta-analysen in de context van een pakketuitspraak. - Voor het definiëren van studies en relevante groepen zijn klassen uit OBI gebruikt omdat dit een veelgebruikte en up-to-date ontologie is.	- STATO ontologie (statistics ontology) met verwijzing naar de onderzochte 'kwaliteit', d.w.z. de PICO uitkomstmaat. - Ontology of Biomedical Investigations (OBI)	Ontology of Clinical Research (OCRe)
Budget Impact Analyse (BIA)	Geen geschikte ontologieën gevonden.	Zelf ontwikkeld met Data Qube vocabulaire, dat weer gebaseerd is op het Statistical Data and Metadata Exchange (SDMX) informatiemodel	-
Overig	Properties die 2x voorkomen: - uit een externe ontologie met een te restrictief domein en bereik - Ander domein / bereik	Properties met dezelfde naam, maar andere namespace, zijn gedefinieerd.	-

Data model / Schema machine-readable package statements





InterventionRationale
 Interventie beslissing om chemotherapie achterwege te laten

Interventie
 Interventie moleculaire diagnostiek genexpressietest op basis van 70 genen, MammaPrint®

InterventieId
 https://w3id.org/zinl/fpr-o#careActivityCode

InterventieClaimCode
 https://w3id.org/zinl/fpr-o#claimCode
 xsd:integer

InterventieMarketingAuthorizationHolder
 https://w3id.org/zinl/fpr-o#marketingAuthorizationHolder

InterventieCostType
 https://w3id.org/zinl/fpr-o#costType
 additional

InterventieNumberOfPatients
 https://w3id.org/zinl/fpr-o#numberOfPatients
 808 xsd:integer

InterventieTimepoint
 https://w3id.org/zinl/fpr-o#timepoint
 default

InterventieTotalCosts
 https://w3id.org/zinl/fpr-o#totalCosts
 -4251723 xsd:float

to illustrate:
 matching data points with
 the ontologie

Demo

- Coupling with the national health data catalog
- FDP

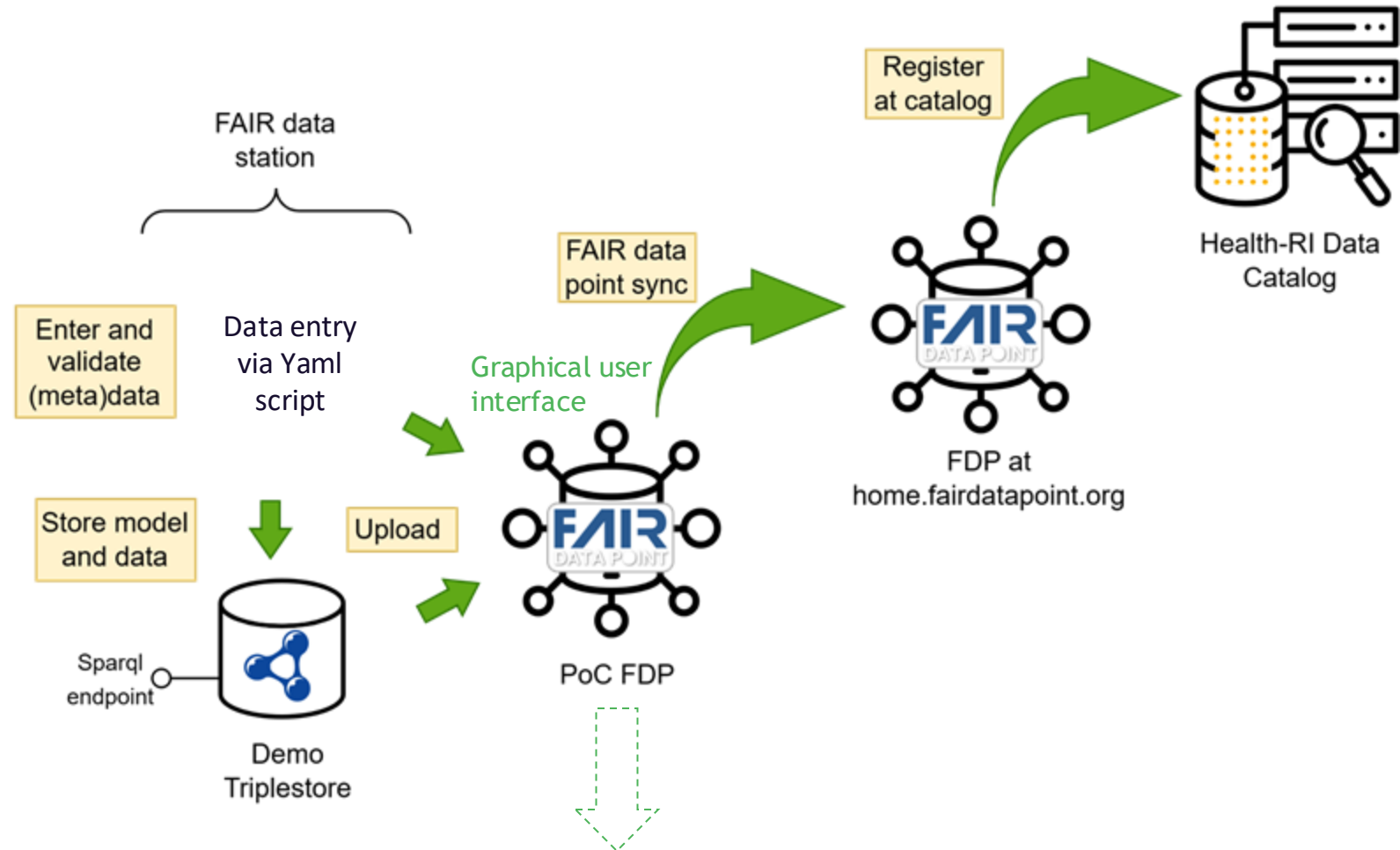
Infrastructure PoC FAIR Packet statement registry

PoC Components

- 1) Queryable model of 3 package statements in breast cancer domain
- 2) FAIR data station (triplestore with Sparql endpoint) met data and semantic model
- 3) FAIR data point in which the package statements are registered with relevant metadata
- 4) Entry in Health-RI national health data catalog
- 5) PURL's on basis of w3id.org

Next steps:

- 6) Create entry form and templates
> currently data entry via scripts from YAML files, which have been manually created
- 7) Publication on open-overheid en open.minVWS



<https://www.open-overheid.nl/>
<https://open.minvws.nl/>

PURL's:

- Link to the FDP: <https://w3id.org/zin/fdp>
- The FAIR Package Registry ontology: <https://w3id.org/zin/fpr-o>
- The taxonomy (part Cochrane & part new terminology): <https://w3id.org/zin/fpr-tax>
- Examples from 3 package statements: <https://w3id.org/zin/package-statements/af>

FDP Homepage Zorginstituut met metadata en link naar datacatalogi Zorginstituut

FAIR Data Point

Metadata for machines

Search FAIR Data Point...
Advanced

Log in

[PoC] FAIR Data Point - National Health Care Institute (Zorginstituut Nederland)

[PoC] FAIR Data Point - Zorginstituut Nederland

Conforms to

- FAIR Data Point Profile

Metadata modified

20-10-2025

Metadata issued

12-09-2025

Title

- [PoC] FAIR Data Point - National Health Care Institute (Zorginstituut Nederland)

Theme

- Health

Publisher

https://fdp-zin.dci.thehyve.nl#publisher

License

http://purl.org/NET/rdflicense/cc-zero1.0

Language

- English

Keyword

- FAIR
- Proof-of-concept
- Catalogi with our public datasets

Show more

FAIR Data Point

Metadata for machines

Search FAIR Data Point...
Advanced

Log in

Identifier

https://fdp-zin.dci.thehyve.nl/

End point url

https://fdp-zin.dci.thehyve.nl/

End point description

https://fdp-zin.dci.thehyve.nl/

Description

- [PoC] FAIR Data Point - Zorginstituut Nederland
- [PoC] FAIR Data Point - National Health Care Institute

Creator

- _g_L23C1720

Contact point

_g_L36C539

Application profile

- FAIR Data Point Profile

Access rights

public

Metadata identifier

identifier

ttl

rdif+xml

json-ld

Catalogs

Package statements - National Health Care Institute [TEST]

Metadata records for machine-readable package statements

HEAL

Issued 17-09-2025 Modified 20-10-2025



7711187



Metadata records for machine-readable package statements

Conforms to	• Catalog Profile
Title	• Package statements - National Health Care Institute [TEST] • Pakketuitspraken - Zorginstituut Nederland [TEST]
Themes	• Health
Temporal coverage	• g_L55C673
Release date	17-09-2025
Publisher	g_L27C784
Modification date	08-10-2025
Licence	http://purl.org/NET/rdflicense/cc-zero1.0
Language	• English • Dutch

FDP Pagina PU met rdf data

FAIR Data Point

https://fdp-zin.dci.thehyve.nl/dataset/7c19c8c0-abd8-4e20-a3c5-45332ac6bbde

ZIN PlekZIN M365OWIZ D8KIK-V Gar OvKIK-V KRTimeTellNS ZakSyncKIK-VDMNWEGIZHealth-RIZINLagerweijiStandaarden.nlNTS SNOWMEDTZW

FAIR Data Point
Metadata for machines

Search FAIR Data Point...Log inAdvanced

[PoC] FAIR Data Point - National Health Ca... / Package statements - National Health... / Standpunt 'Genexpressie Testen bij vr...

Standpunt 'Genexpressie Testen bij vrouwen ouder dan 50 jaar en vroeg stadium borstkanker'

Standpunt 'Genexpressie Testen bij vrouwen ouder dan 50 jaar en vroeg stadium borstkanker'

Conforms to

- Dataset Profile

Version0.0.1

Title

- Standpunt 'Genexpressie Testen bij vrouwen ouder dan 50 jaar en vroeg stadium borstkanker'
- Point-of-view 'Gene Expression Testing in Women Over 50 with Early-Stage Breast Cancer'

Theme

- Health

Statusunder development

Purpose

- Public Benefit

Publisher_g_L25C1717

Population coverage

- Women older than 50 with Breast Cancer, HR+/HER2-, N0 or N1, High risk based on AO! or Predict

Language

- English

Keyword

- Age more than 50
- Breast cancel
- Gene expression test
- Show more

Identifierhttps://graph-zin.dci.thehyve.nl/repositories/mamma-onco#

Health theme

- https://www.wikidata.org/wiki/Q128581
- https://www.wikidata.org/wiki/Q7843332

Geographical coverage

- Netherlands

Documentation

- https://www.zorginstituutnederland.nl/documenten/2023/10/24/standpunt---mammaprnt-en-oncotype-dx-vergoede-zorg-voor-bepaalde-groep-vrouwen

Distribution

- RDF serialization of Mammaprnt/Oncotype DX package statement

Description

- Standpunt 'Genexpressie Testen bij vrouwen ouder dan 50 jaar en vroeg stadium borstkanker'
- Point-of-view 'Gene Expression Testing in Women Over 50 with Early-Stage Breast Cancer'

Creator_g_L34C2471

Contact point_g_L45C3407

Conforms toDataset Profile

Applicable legislationhttp://data.europa.eu/eli/reg/2025/327/oj

Access rightspublic

ttlrdf+xmljson-id

DistributionsSample DistributionsDataset SeriesAnalytics Distributions

Distributions

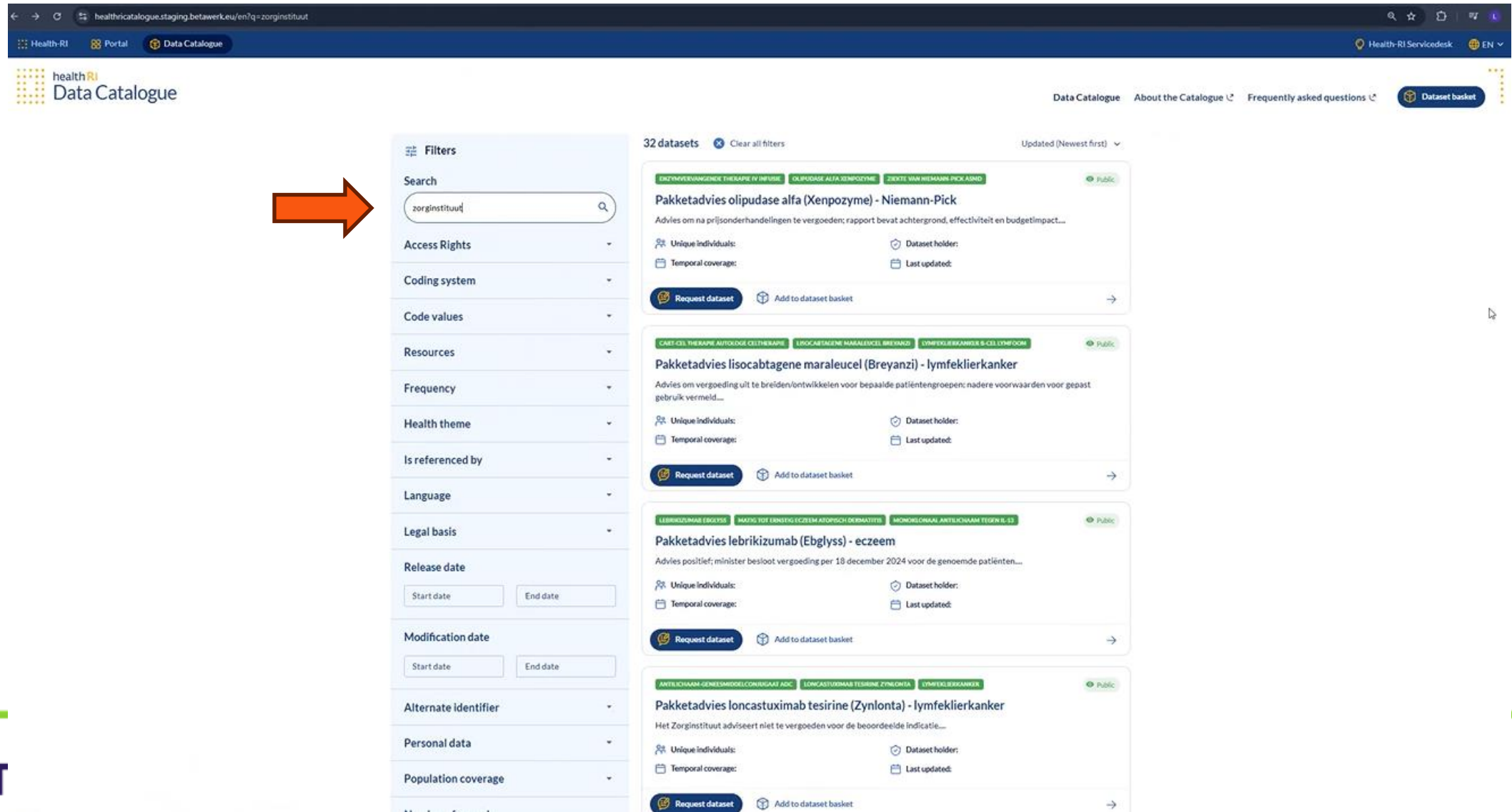
RDF serialization of Mammaprnt/Oncotype DX package statement

RDF serialization of Mammaprnt/Oncotype DX package statement

Issued 18-09-2025Modified 20-10-2025Media Typehttps://www.iana.org/assignments/media-types/text/turtle

3

Data gepubliceerd in Health-RI Datacatalogus (acceptatie omgeving)



The screenshot displays the Health-RI Data Catalogue interface. On the left, a sidebar contains a 'Filters' section with a search bar and various filter categories: Access Rights, Coding system, Code values, Resources, Frequency, Health theme, Is referenced by, Language, Legal basis, Release date (with start and end date inputs), Modification date (with start and end date inputs), Alternate identifier, Personal data, and Population coverage. An orange arrow points from the search bar in the filters to the search results. The main content area shows '32 datasets' with a 'Clear all filters' button and a 'Updated (Newest first)' dropdown. The first three dataset entries are visible:

- Pakketadvies olipudase alfa (Xenpozyme) - Niemann-Pick**
Advies om na prijsonderhandelingen te vergoeden; rapport bevat achtergrond, effectiviteit en budgetimpact...
Unique individuals: Dataset holder: Last updated:
Temporal coverage: Request dataset Add to dataset basket
- Pakketadvies lisocabtagene maraleucel (Breyanzi) - lymfeklierkanker**
Advies om vergoeding uit te breiden/ontwikkelen voor bepaalde patiëntengroepen; nadere voorwaarden voor gepast gebruik vermeld...
Unique individuals: Dataset holder: Last updated:
Temporal coverage: Request dataset Add to dataset basket
- Pakketadvies lebrikizumab (Ebglyss) - eczeem**
Advies positief; minister besloot vergoeding per 18 december 2024 voor de genoemde patiënten...
Unique individuals: Dataset holder: Last updated:
Temporal coverage: Request dataset Add to dataset basket

The bottom of the page shows the footer with the Health-RI logo and the page number 124.

PU in Health-RI Datacatalogus (acceptatie omgeving) voorbeeld

The screenshot displays the Health-RI Data Catalogue interface. The top navigation bar includes links for Health-RI, Portal, and Data Catalogue, along with a search icon and a language selector set to EN. The main header features the Health-RI logo and the title 'Data Catalogue'. On the right, there are links for 'Data Catalogue', 'About the Catalogue', 'Frequently asked questions', and a 'Dataset basket' button.

The central content area shows a dataset profile with the following sections:

- Home Publisher:** E-Mail: info@zin.nl
- Description:** Standpunt 'Genexpressie Testen bij vrouwen ouder dan 50 jaar en vroeg stadium borstkanker'. A link to 'View distributions' is provided.
- Health theme:** breast cancer, triple-negative breast cancer
- Keywords:** Age more than 50, Breast cancer, Gene expression test, HER-, HR, Hormone Receptor Positive Tumor, Human Epidermal Growth Factor 2 Negative Carcinoma Of Breast, Mammaprint, Molecular diagnostic, N0 Category, N1 Category, Oncotype DX.
- Population coverage:** Women older than 50 with Breast Cancer, HR+/HER2-, N0 or N1, High risk based on AOI or Predict
- Geographical coverage:** Netherlands
- Purpose:** Public Benefit
- Conforms to:** Dataset Profile
- Distributions:** RDF serialization of Mammaprint/Oncotype DX package statement

On the right side, a light blue box contains additional metadata:

- Identifier:** <https://graph.zin.dci.thelhyve.nl/repositories/mamma-onco#>
- Access rights:** Public
- Legal basis for publishing:** <http://data.europa.eu/eli/reg/2025/327/oj>
- Contact details:** (Redacted)
- Documentation:** <https://www.zorginstituutnederland.nl/documenten/2023/10/24/standpunt--mammaprint-en-oncotype-dx-vergoede-zorg-voor-bepaalde-groep-vrouwen>
- Dataset languages:** English
- Theme(s):** Health

The footer of the page features the Health-RI logo.

Behoefte onderzoek op basis interviews

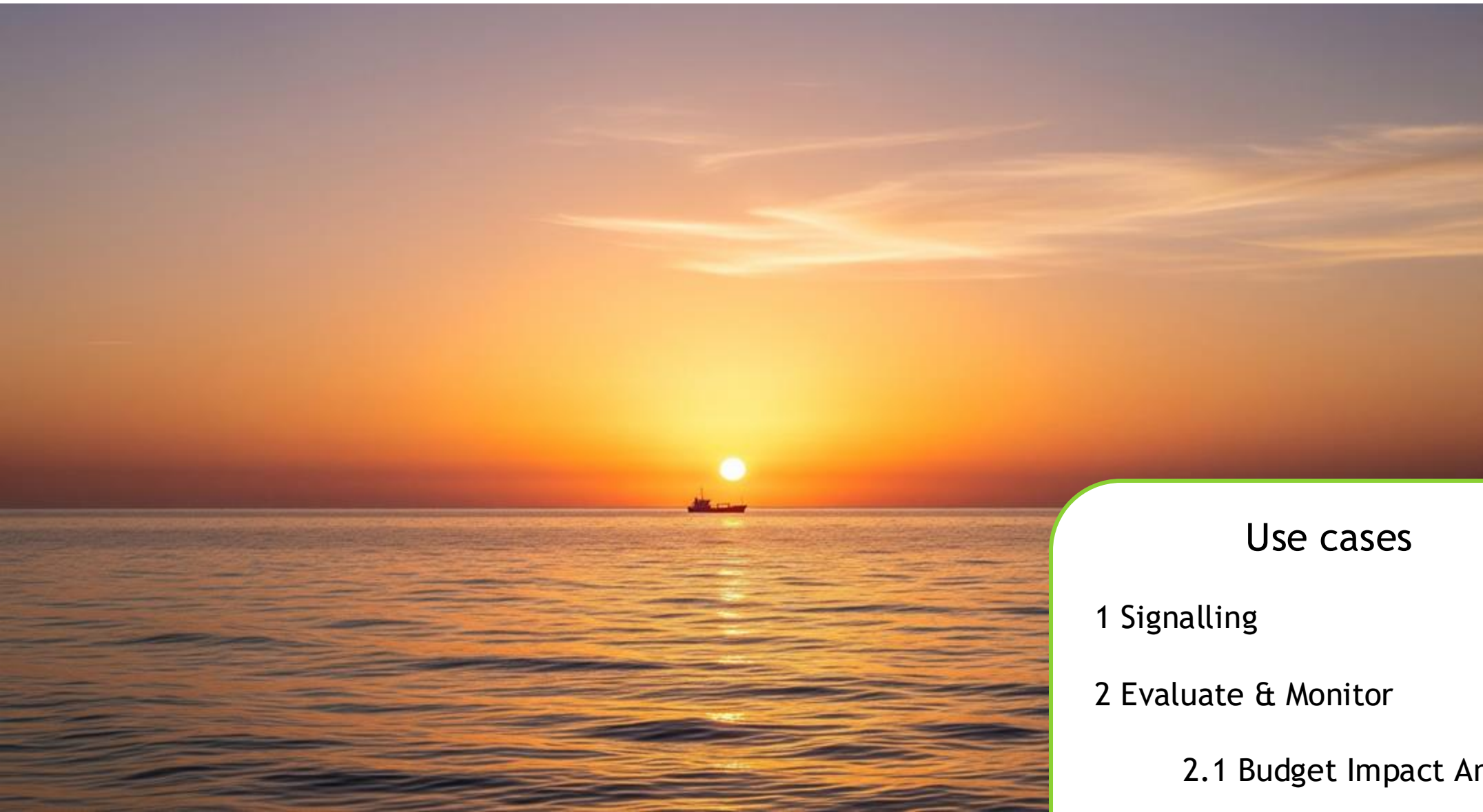
Resultaten Interviews

Internal and external stakeholders

Wishlist:

- FAIR data to query as first step of process automation
 - focus on **findability and search**
 - focus on **machine-readability**
- Coupling with RWD
- Future outlook
 - Which data will we need to evaluate new statements

Use cases



Use cases

1 Signalling

2 Evaluate & Monitor

2.1 Budget Impact Analysis

2.2 Real-World Data - IKNL

Signalling

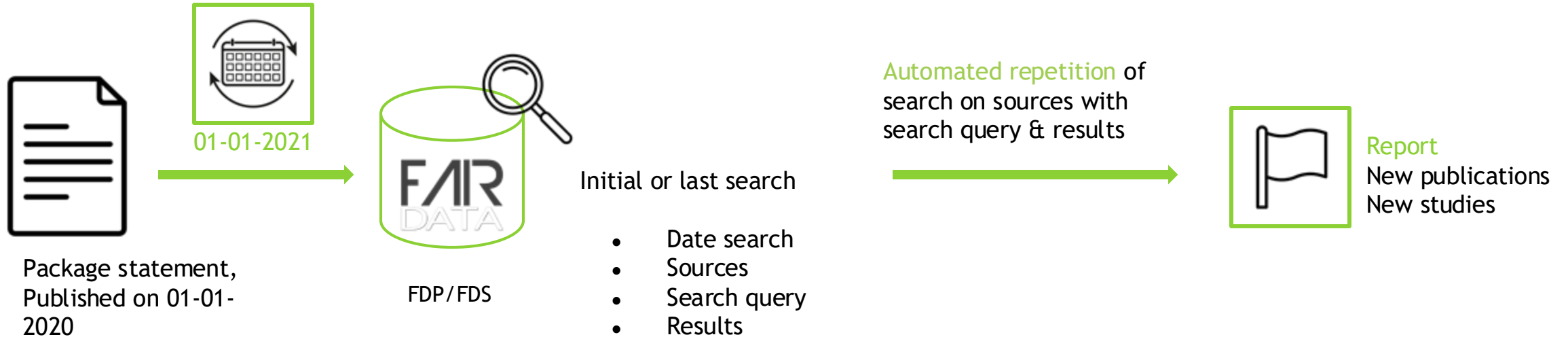
- What's on the horizon? -



Is there new evidence that prompts us to reevaluate?

Automated, periodic search request to identify new data/new studies/new publications for:

- Individual package statements
- Specific PICOs
- Final outcome measure available instead of surrogate outcome measure



Evaluate & Monitor: Budget Impact Analysis

- What's on the horizon? -

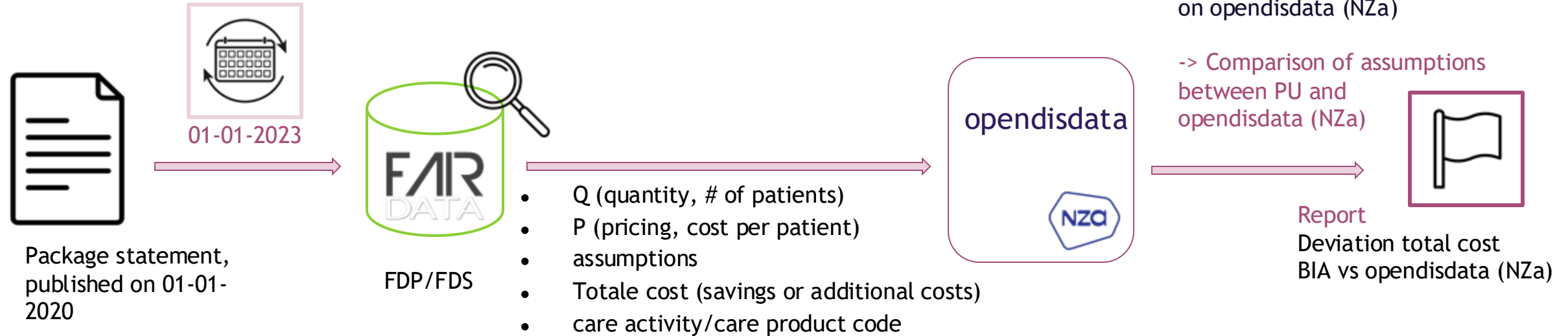


Cyclic package management

Evaluate assumptions and estimates in BIA part of the package statement

- Automated
- Periodically
- Leveraging e.g., opendisdata (NZa)

Application - fact check leveraging opendisdata



Evaluate & Monitor: RWD

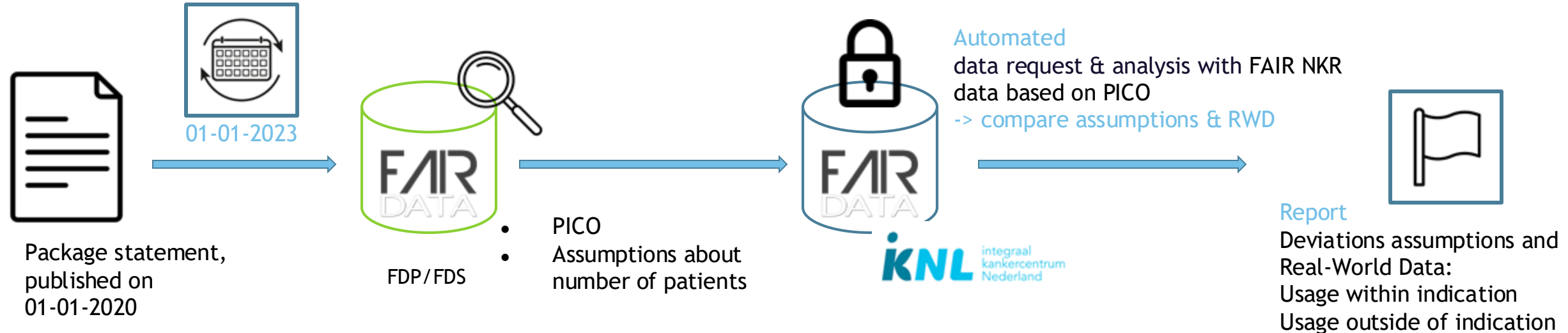
- What's on the horizon? -



Cyclic package management

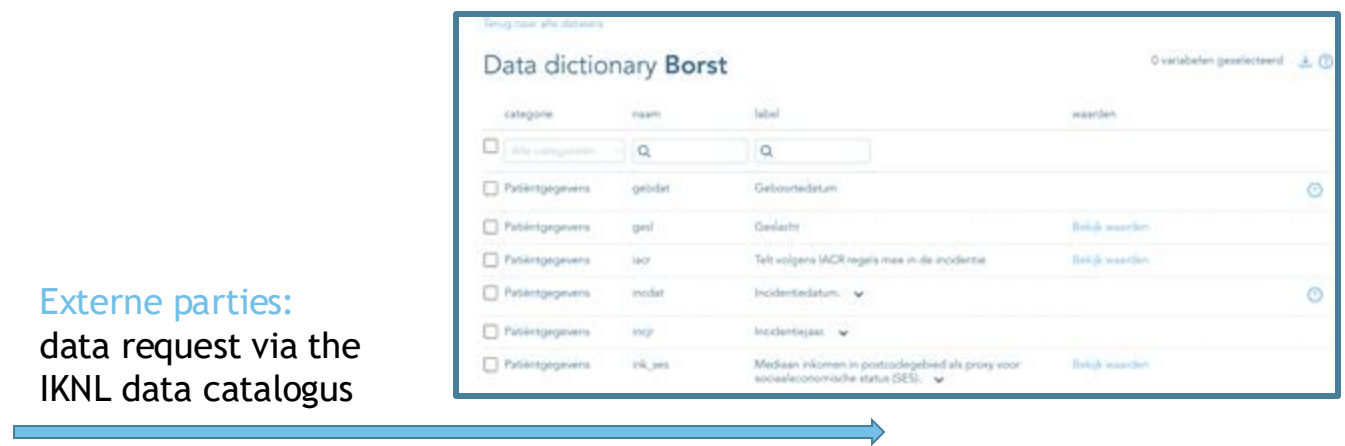
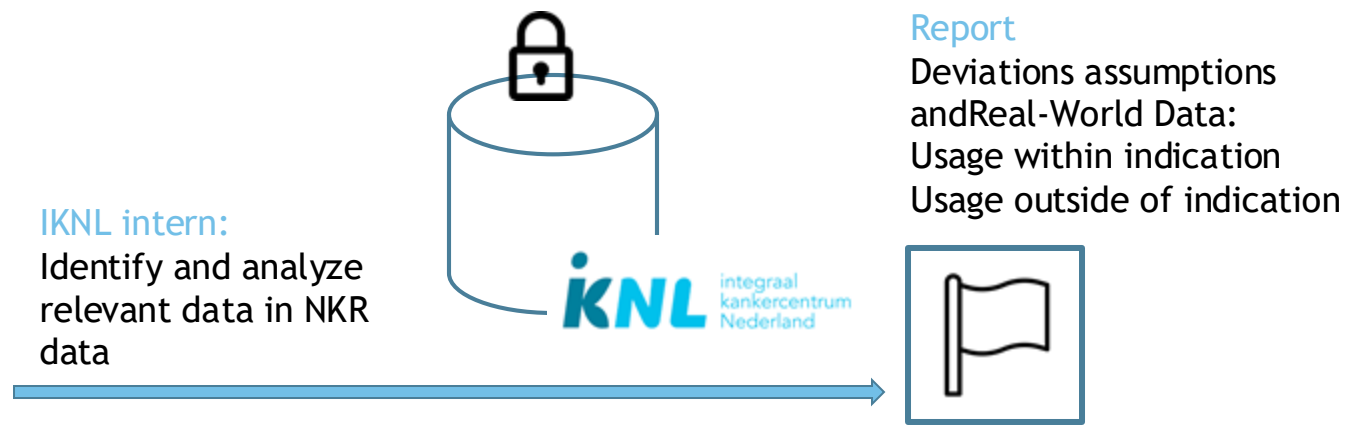
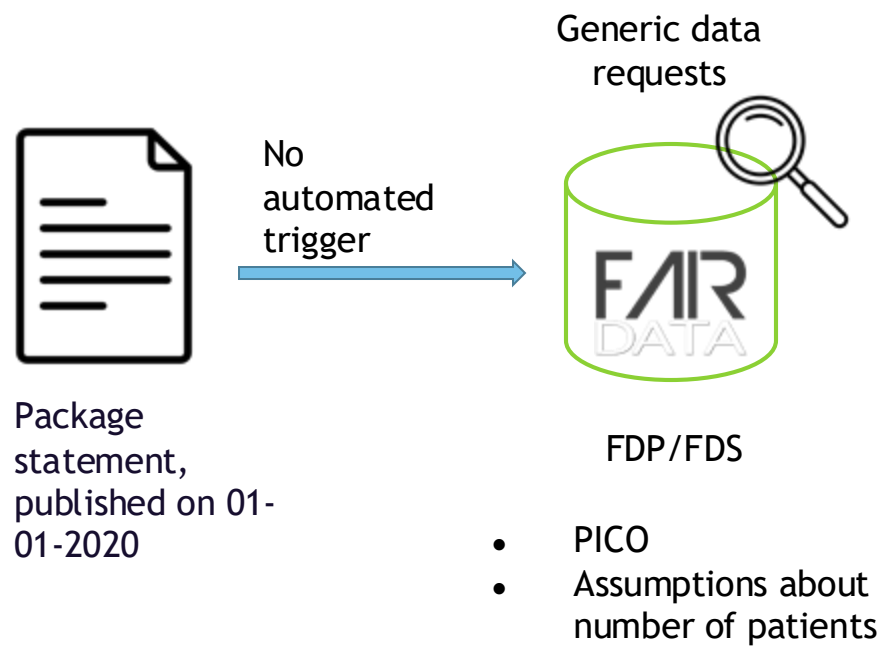
- 1) Evaluate assumptions and estimates about indications and number of patients
 - Automated
 - Periodically
 - Leveraging RWD (NKR data)
- 2) Data availability check (PARP blockers)

Application - RWD fact check leveraging NKR data



Evaluate & Monitor: RWD

- Status Quo -



FAIR Package statement

Intervention
title : Chemotherapie op basis van MammaPrint uitslag
applied intervention : { Genetic test advised Chemotherapy }
intervention rationale : Pas chemotherapie toe naar aanleiding van hoge risicoschatting MammaPrint
intervention classification : { Screening (?) }

Population
title : Vrouwen ouder dan 50 met borstkanker, HR+/HER2-, N0, hoog risico op basis van AO! of Predict
age : { Age more than 50 years }
sex : { Female }
conditions : { Breast cancer, Estrogen Receptor Positive Tumor, Progesterone Receptor Positive Tumor, Human Epidermal Growth Factor 2 Negative Carcinoma Of Breast, N0 Category, High risk based on AO! or Predict }
treatment : { Surgery }

DataSet (Trend Assumption)
title : Geschatte trend voor MammaPrint en/of Oncotype DX

Observation (timepoint)
time unit : N/A
timepoint: default
number of patients: 808
intervention market penetration : 1.0

dataset



IKNL Real-world data

Terug naar alle datasets

Data dictionary **Borst**

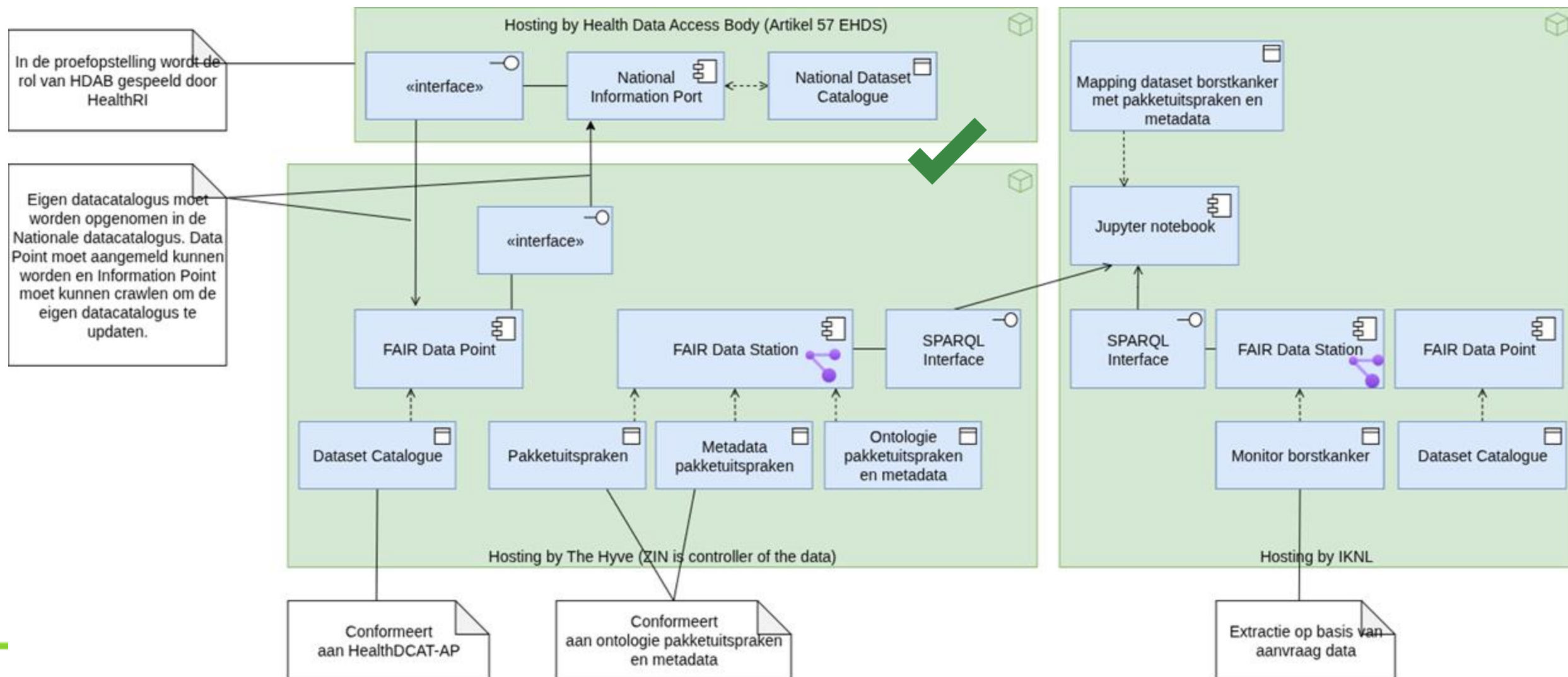
categorie	naam	label
☐ Alle categorieën		
☐ Patiëntgegevens	gebdat	waarden
☐ Patiëntgegevens	gesl	
☐ Patiëntgegevens	iacr	

Verberg waarden

Waarde	Omschrijving
0	nee
1	ja, uitslag: zeer laag risico
2	ja, mammaprint, uitslag: laag risico
3	ja, mammaprint, uitslag: hoog risico
8	ja, mammaprint, uitslag: onbekend
11	ja, oncotype DX, uitslag: laag risico
12	ja, oncotype DX, uitslag: intermediair risico
13	ja, oncotype DX, uitslag: hoog risico
19	ja, oncotype DX, uitslag: onbekend

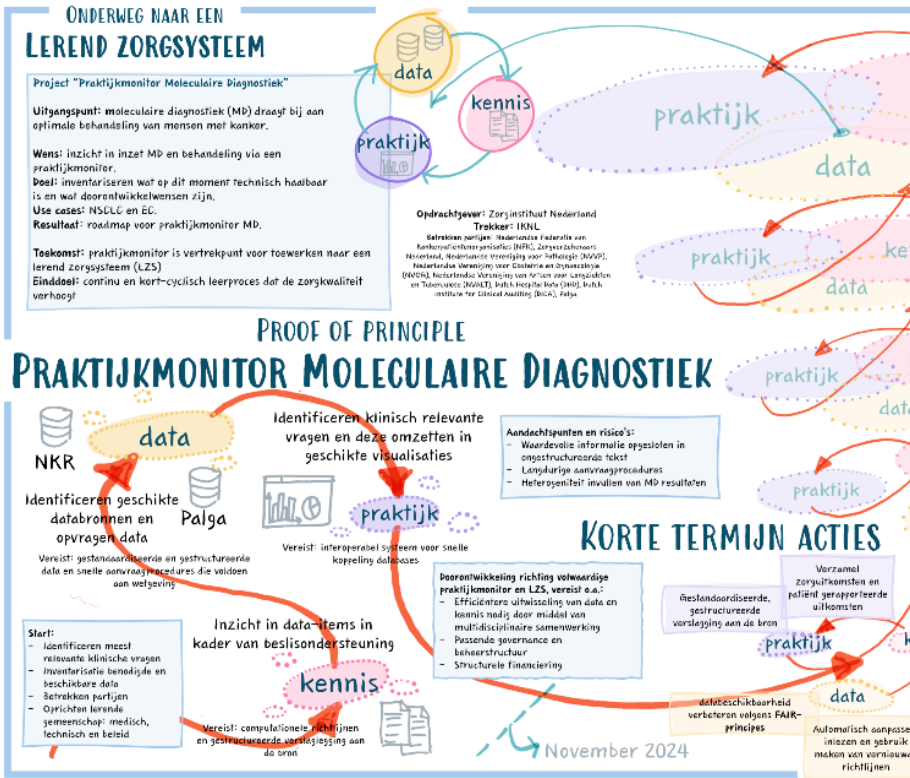
Evaluate & Monitor: RWD

- What's on the horizon? -



Roadmap lerende zorgsysteem

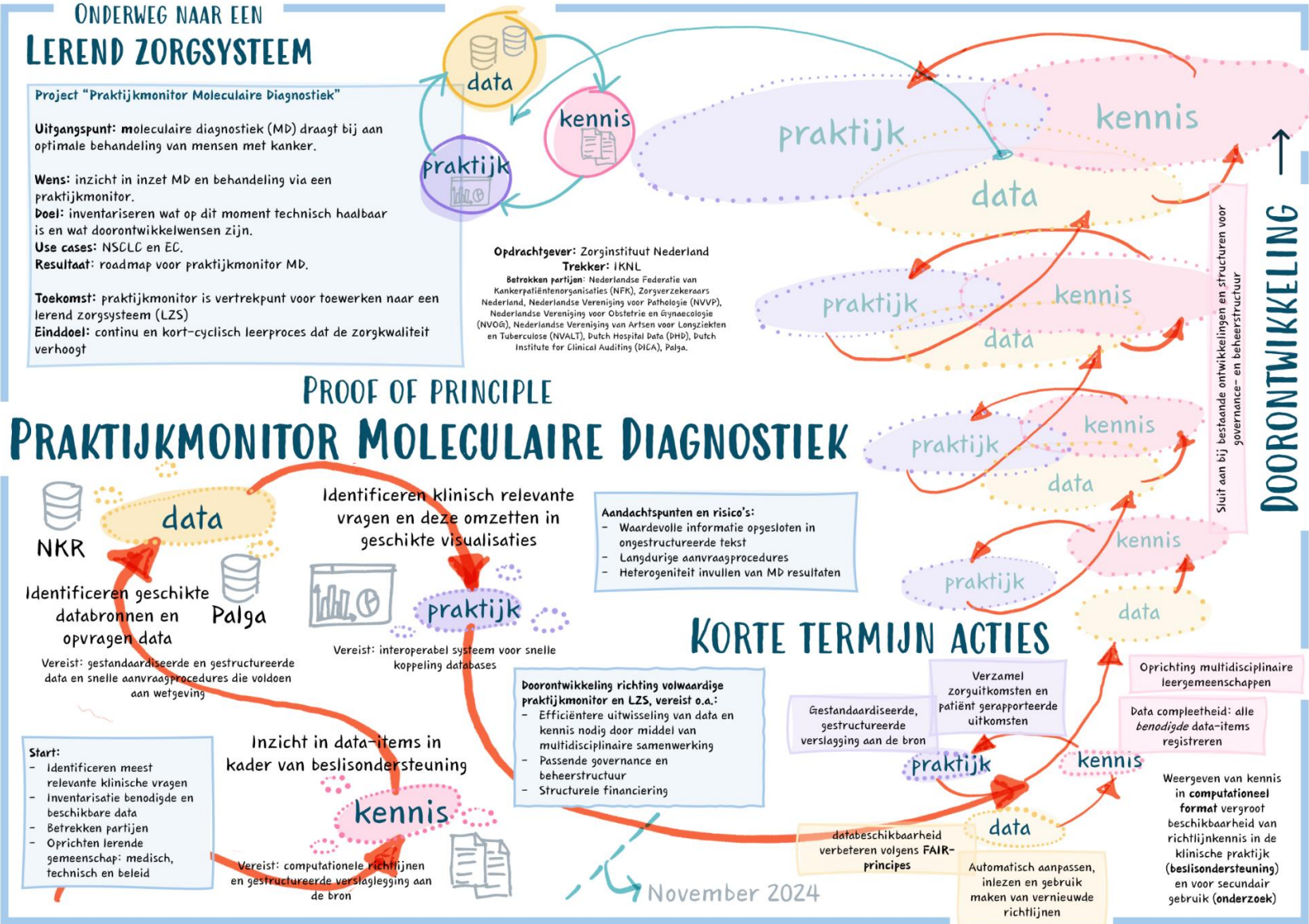
Visie en strategie secundair datagebruik verbeteren
Databeschikbaarheid voor een lerend zorgsysteem

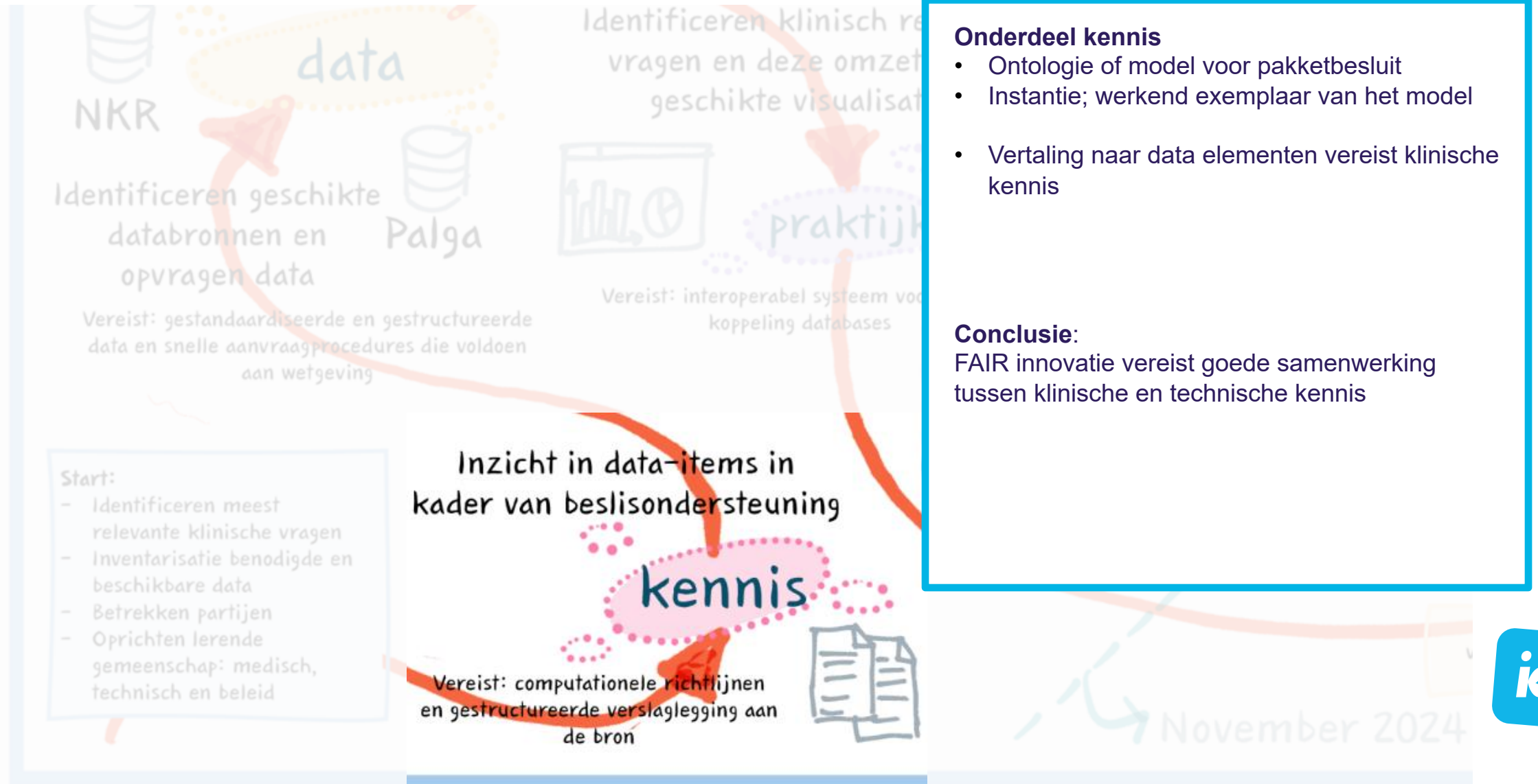


msen (foto: © Bert Roos)

de landelijke moleculaire-pathologie-lab, maar ondanks geïntegreerde gegevens blijkt het toch niet zo makkelijk om alle benodigde data te verzamelen.







Toekomst: praktijkmonitor is vertrekpunt voor toewerken naar een lerend zorgsysteem (LZS)
Einddoel: continu en kort-cyclisch leerproces dat de zorgkwaliteit verhoogt

Nederland, Nederlandse Vereniging voor Pathologie (NVVP), Nederlandse Vereniging voor Obstetrie en Gynaecologie (NVOG), Nederlandse Vereniging van Artsen voor Longziekten en Tuberculose (NVALT), Dutch Hospital Data (DHD), Dutch Institute for Clinical Auditing (DICA), Palga.

PRAKTIJKMONITOR



Vereist: gestandaardiseerde en gestructureerde data en snelle aanvraagprocedures die voldoen aan wetgeving

Onderdeel data

- Niet alle relevante data items zijn beschikbaar
- Wens voor FAIR by design voor cyclische pakketbeheer
- Stel eisen aan RWD

Conclusie:

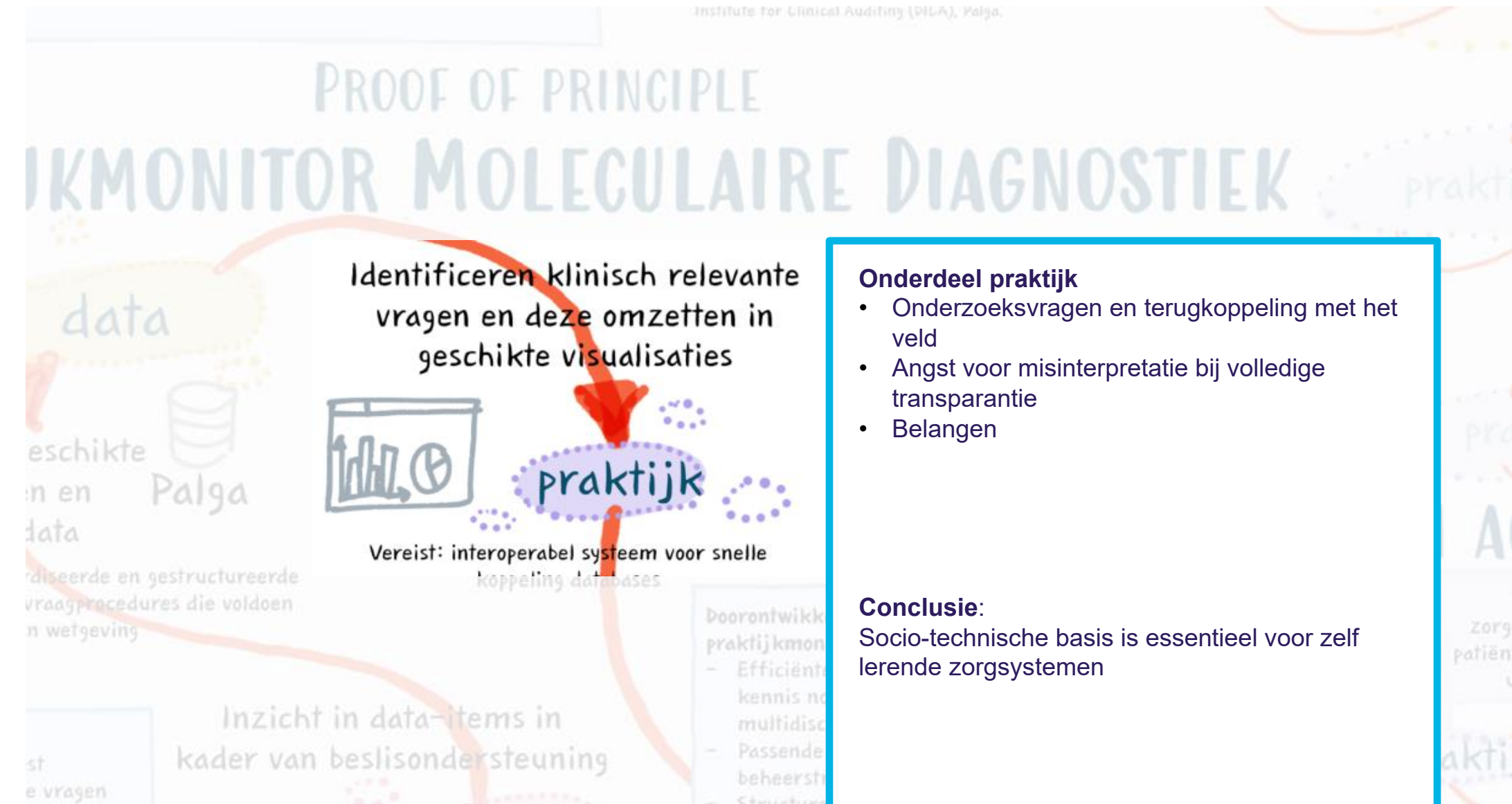
Cyclisch pakketbeheer vereist voorbereiding o.a. door een datatoets voor uitbrengen pakketbesluit waar wens voor evalueren en monitoren bestaat

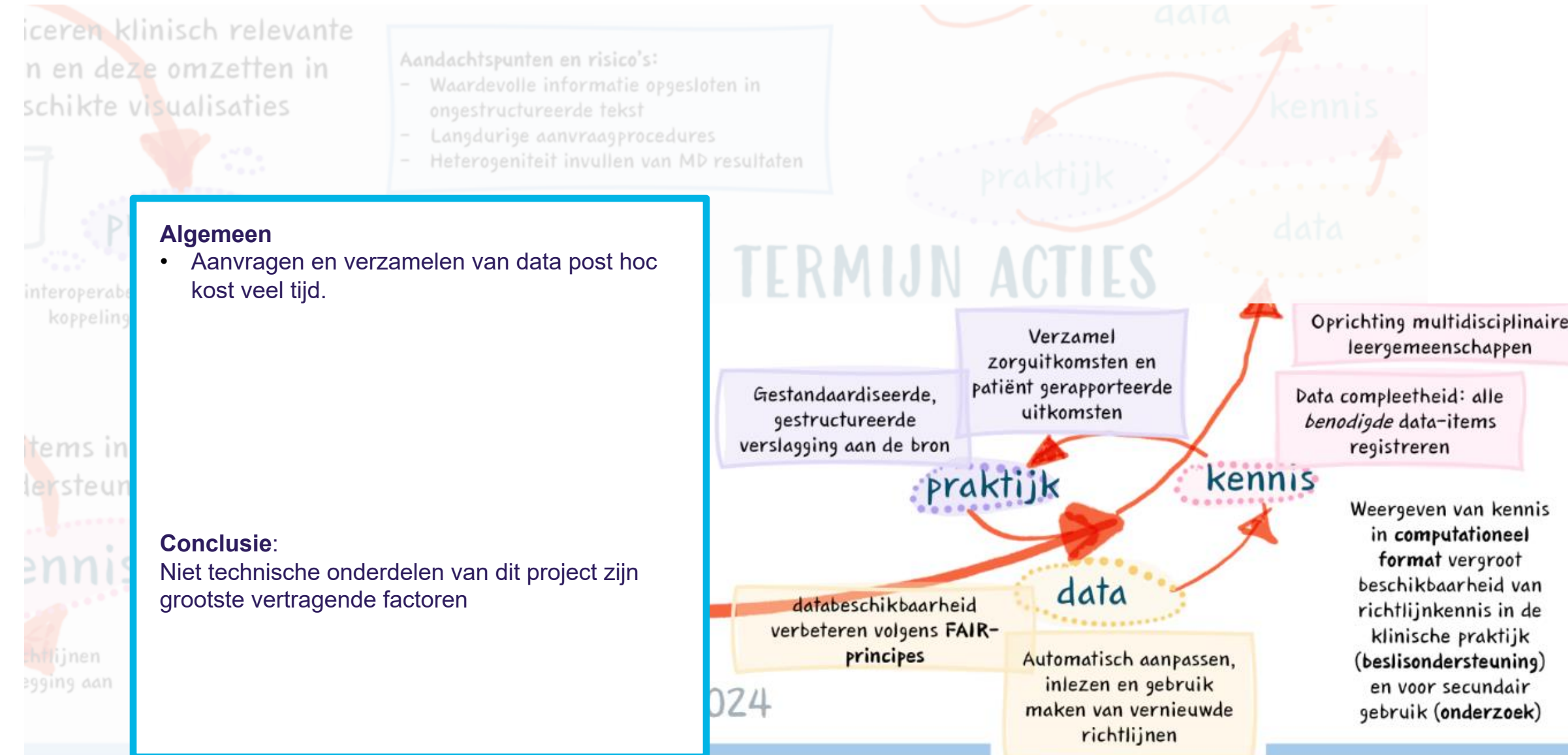
DIAGNOSTIE

spunten en risico's:
devolle informatie opgesloten in
structureerde tekst
durige aanvraagprocedures
ogeniteit invullen van MD resulta

KORTE TI

ng volwaardige
praktijkmonitor en LZS, vereist o.a.:
- Efficiëntere uitwisseling van data en
kennis nodig door middel van





Opdrachtgever: Zorginstituut Nederland

Trekker: IKNL

Betrokken partijen: Nederlandse Federatie van

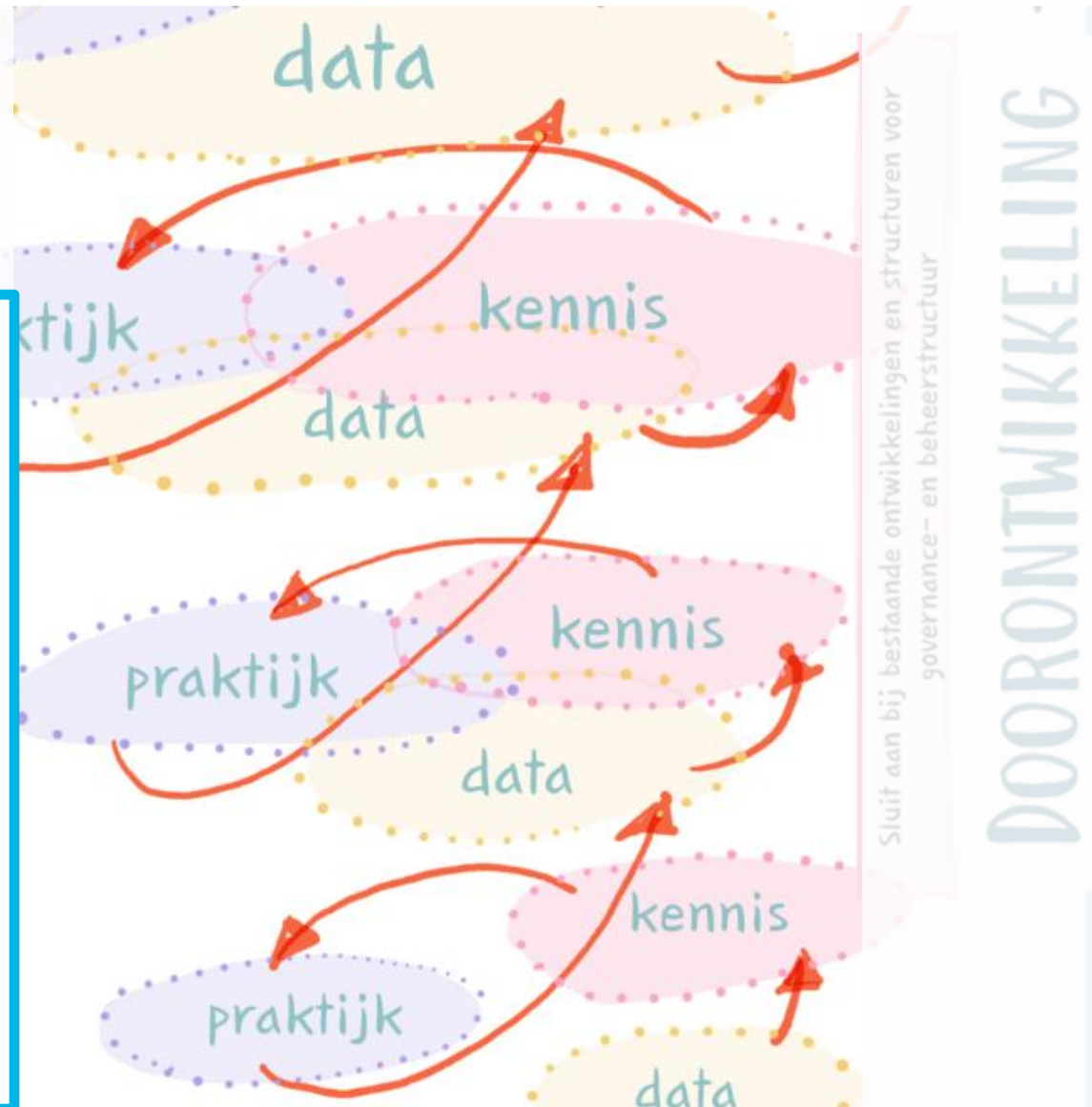
Kanker
Nederland
Nederland
(NVOG), Ne
en Tuber
In

Relevantie voor IKNL en lerend zorgsysteem

- Iteraties zijn ultieme resultaat en doel
- FAIR Pakketregister met machineleesbare pakketbesluiten zijn heel relevant voor IKNL
- Veel kansen tot verdere samenwerking en borging in relevante initiatieven (OncoNext, dure geneesmiddelen, kwaliteitsregistraties etc.)

Conclusie:

Zelf lerende zorgsystemen zijn relevant en veel mogelijkheden tot opvolging.



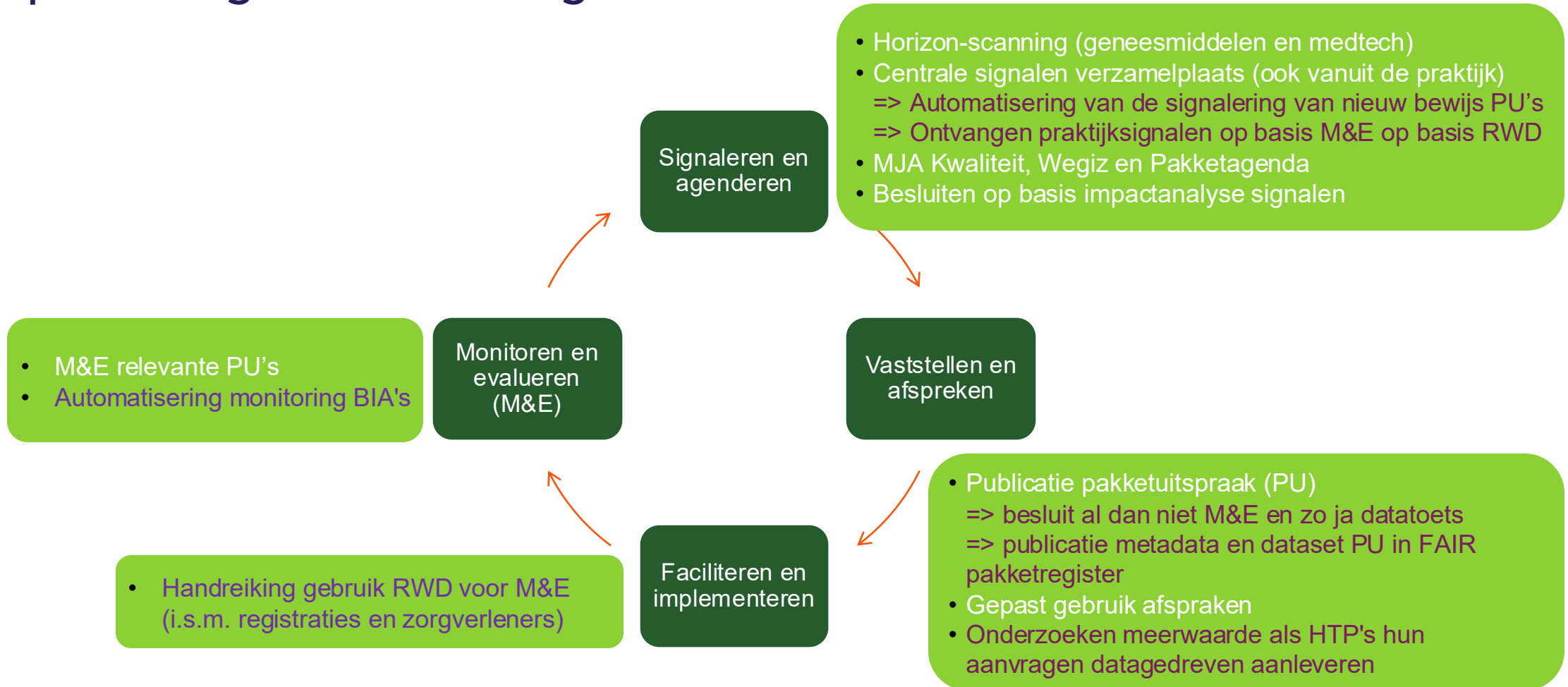
Inpassing binnen het Zorginstituut en IKNL

DISCLAIMER

Het is een onderzoeksproject.
Intern nog geen besluit genomen om het
FAIR Pakketregister in productie te nemen.

Vandaag bedoeld om methodiek en
lessen te delen en te horen hoe jullie
toepasbaarheid zien.

Aanzet cyclisch pakketbeheer met FAIR pakketregister - Rol Zorginstituut



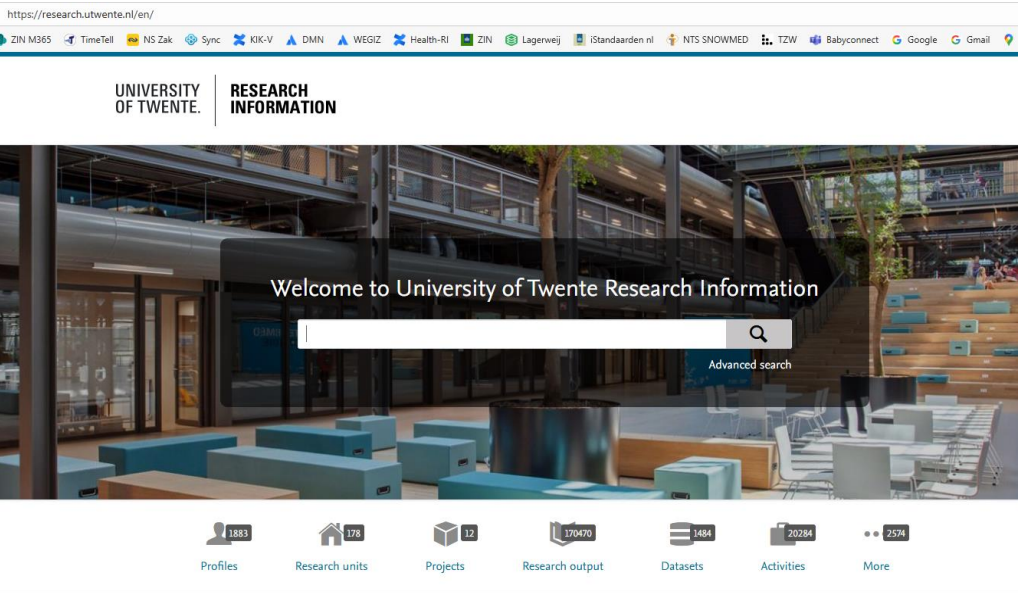
Aanzet cyclisch pakketbeheer met FAIR pakketregister - Rol VWS & veldpartijen



Kansen en risico's van een FAIR pakketregister

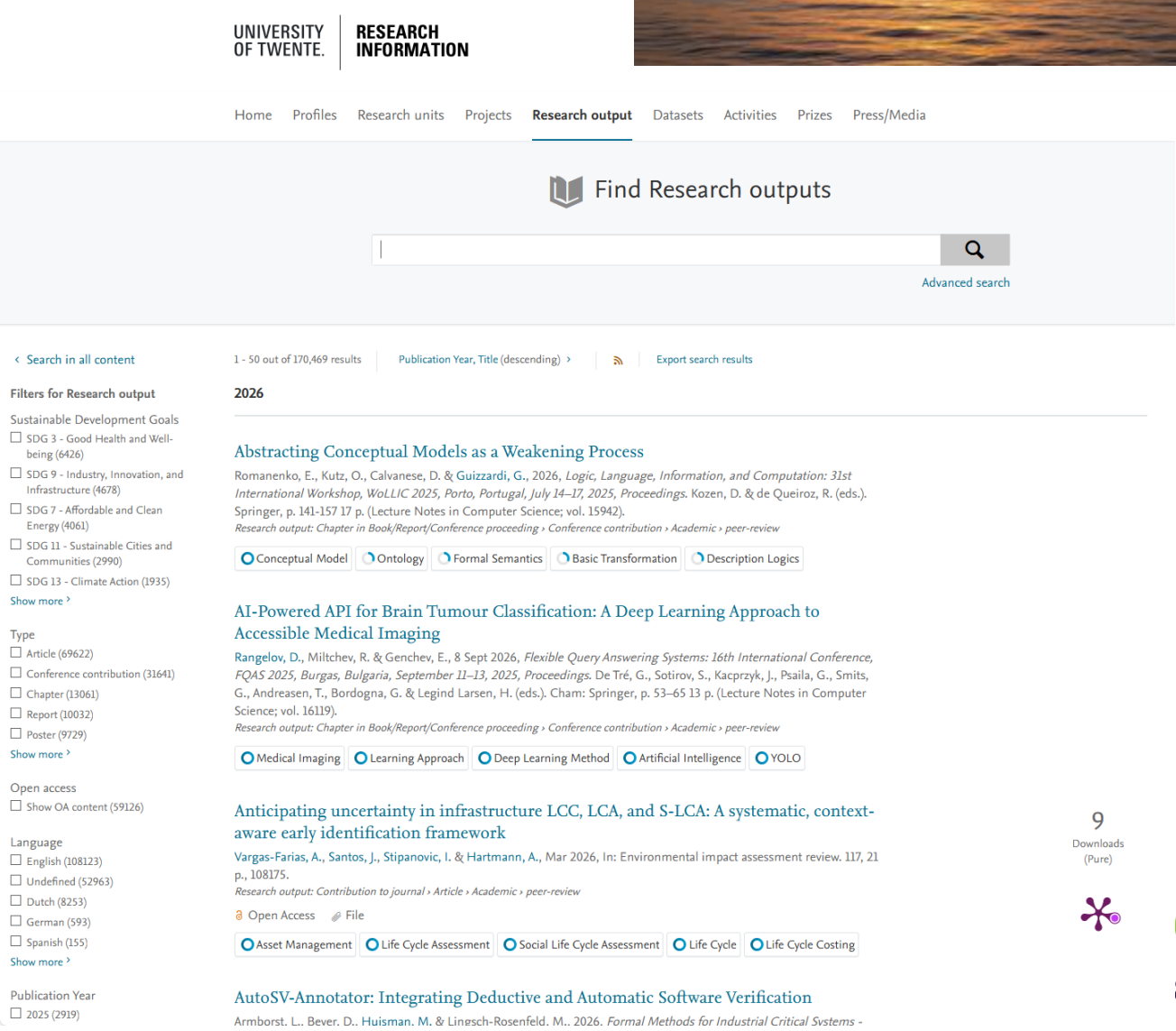
Stakeholder	Kans	Bedreiging	Nog verkennen
Zorginstituut - Inhoud	• Automatisering signalering • Automatisering BIA herberekening • Sneller signalen uit het veld door data toets vooraf en automatisering E&M door registraties	• HTP heeft makkelijker toegang tot alle uitspraken om daar analyses op te doen t.b.v. rechtzaken	• Hoe proces gaat lopen • Onderzoeken meerwaarde als HTP's hun aanvragen datagedreven aanleveren
Zorginstituut – Data	• Evalueren en monitoren op basis RWD • FAIR publicatie onze data en documenten (gewenst vanuit EHDS, WOO)		• Hoe proces gaat lopen - ook (de inspanning) op data vlak • Relatie met OMOP en <u>Pharmaceutical ResearchProtocol (Vulcan UDP) v1.0.0-ballot</u>
VWS	• Inzicht in effectiviteit op basis RWD ter beoordeling of prijsarrangement (nog) passend is • Transparantie over pakketbesluiten naar burger		
IKNL – Inhoud	• Inzicht in effectiviteit bij geëxcludeerde subpopulaties • RWD validatie van effectiviteit studies	• Methodologische risico's - Misinterpretatie van resultaten (e.g. regio's, ziekenhuizen)	• Wenselijkheid en haalbaarheid van verzamelen relevante variabelen • Juridische basis
IKNL - Data	• NKR data gebruik voor HTA en lerende zorgsystemen voor snellere impact • Koppelen van registratie data met relevante bronnen	• NKR is een onafhankelijke bron, geen kwaliteitsregistratie of beleidsinstrument	• Relatie met OMOP CDM en andere standaarden • Synergie met EHDS en FAIR initiatieven
Zorgverleners / aanbieders	• Snel informatie vinden over vergoede behandelingen • Snel inzicht over effectiviteit bij onderbelichte subpopulaties	• Mis-interpretatie van behandelkeuzes gebaseerd op klinische kennis en ervaring	• Proces, implementatie van data in FDP in EPDs • Koppelen aan (machineleesbare) richtlijnen
Health technology providers (HTP)	• Datagedreven aanleveren aanvragen • Beter overzicht op alle uitspraken		

Mogelijkheden: "Bibliotheek" gebaseerd op FDP



Welcome to University of Twente Research Information

We use this website to showcase information about our researchers and their research output, activities, press/media, prizes and research units at the University of Twente (UT). The University of Twente is a modern, entrepreneurial university, leading in the area of new technologies and a catalyst for change, innovation and progress in society. We work on the technologies of the future (ICT, biotechnology, nanotechnology), combining technological knowledge and innovation with behavioural and social academic research. Our strength lies in the combination of excellent science, entrepreneurship and international orientation. Our researchers focus on themes with a large societal impact: health, water, green energy, safety and education.



Mogelijkheden:
Tijdslijn van gerelateerde
pakketbesluiten
gebaseerd op FDP



HAS

HAUTE AUTORITÉ DE SANTÉ

Médicament

Vaccination

Dispositif

Évaluation économique

Moyens d'information

Agenda

Ex : diabète, antalgique, alzheimer, prothèse de hanche ...

Dans tout le site

RECHERCHE AVANCÉE

Industriels > Médicament > YERVOY (ipilimumab)

ÉCOUTER

AJOUTER À MA SÉLECTION

Historique des avis

Informations technique

YERVOY (ipilimumab)

MÉDICAMENT - Mis en ligne le 16 oct. 2025

Substance active (DCI) : ipilimumab

Historique des avis (16)

14/09/2025

OPDIVO/YERVOY (nivolumab/ipilimumab) - Cancer colorectal (CRC)

L'essentiel: Avis favorable au remboursement, uniquement dans « le traitement de première ligne chez des patients adultes atteints de cancer colorectal métastatique ». L'avis est assorti d'un ASMR 1.

ASMR : 1 2 3 4 5

12/12/2024

OPDIVO (nivolumab) et YERVOY (ipilimumab) - Cancer colorectal

Refus d'autorisation d'accès précoce des spécialités OPDIVO (nivolumab) et YERVOY (ipilimumab) en association, dans l'indication de cancer colorectal métastatique.

→

22/11/2023

OPDIVO/YERVOY (nivolumab/ipilimumab) - Mélanome avancé

Inscription dans le sous-groupe B-RAP muté. L'essentiel: Avis favorable au remboursement, uniquement « en 3ème ligne chez les patients adultes atteints de mélanome avancé ». L'avis est assorti d'un ASMR 1.

ASMR : 1 2 3 4 5

Informations techniques

Code ATC L01XC11
L01FX04

Exploitant BRISTOL-MYERS SQUIBB

Présentation • YERVOY 5 mg/mL, solution à diluer pour perfusion
1 flacon en verre de 10 mL (CIP : 34009 580 877 0 7)
1 flacon en verre de 40 mL (CIP : 34009 580 878 7 5)

Toutes nos publications sur

- Cancer de l'appareil digestif
- Cancer de l'appareil respiratoire
- Cancer de l'appareil urinaire
- Cancer de la peau et des tissus conjonctifs
- Enfants - Adolescents
- Traitement médicamenteux

. Conclusies, discussie & vervolg

Geslaagde 1e stap op weg naar automatisering van datagedreven cyclisch pakketbeheer dat naar meer smaakt

- Technisch & informatiekundig geslaagd.
- Enthousiasme voor het concept zowel bij het Zorginstituut als IKNL en zowel vanuit de inhoud als de informatiekundigen
- De volgende stap is adoptiebesluit door het Zorginstituut in samenspraak met VWS en veldpartijen

Stap 1 op weg naar geautomatiseerd cyclisch pakketbeheer

- We hebben 3 pakketuitspraken '100%' FAIR gemaakt*!
 - Machine-leesbaar
 - Volgens standaarden
 - Gepubliceerd volgens FAIR principes
- Bewezen hoe het zou kunnen werken (use cases)
- Kennismodelering: rijke expressie met zowel meerdere onderlinge relaties en op meerdere plekken structuur varianten (o.a. BIA) (en soms uitzonderingen)
- De menselijke evaluatie is lastig te standaardiseren
- Vrij uitgebreid datamodel waardoor opstellen yml file met de data makkelijker is dan inrichten entry forms voor de verschillende varianten
- Data beschikbaar op FAIR manier

Vervolgstappen

- Besluit of we hiermee verder gaan, zo ja:
 - Oude besluiten met basis metadata publiceren (deel 2024 gedaan, ander deel onderweg)
 - Bekijken welke besluiten zinvol zijn om deels of volledig uit te werken tbv signalering, evalueren en monitoring, herbeoordeling
 - Evaluatie in hoeverre AI ingezet kan worden bij het coderen van de (oude) pakketbesluiten
 - Inschatting benodigde inzet en maken projectplan
- Kijken of het aansluit bij
 - [Home - Pharmaceutical Research Protocol \(Vulcan UDP\) v1.0.0-ballot](#)
We kunnen vragen voor de initiele beoordelingen of HTP's op basis hiervan aanleveren.
Dan ontvangen we de studies ook als data
 - Internationale samenwerking met HTA bodies
 - Pakketuitspraken van zorgverzekeraars
- Publicatie

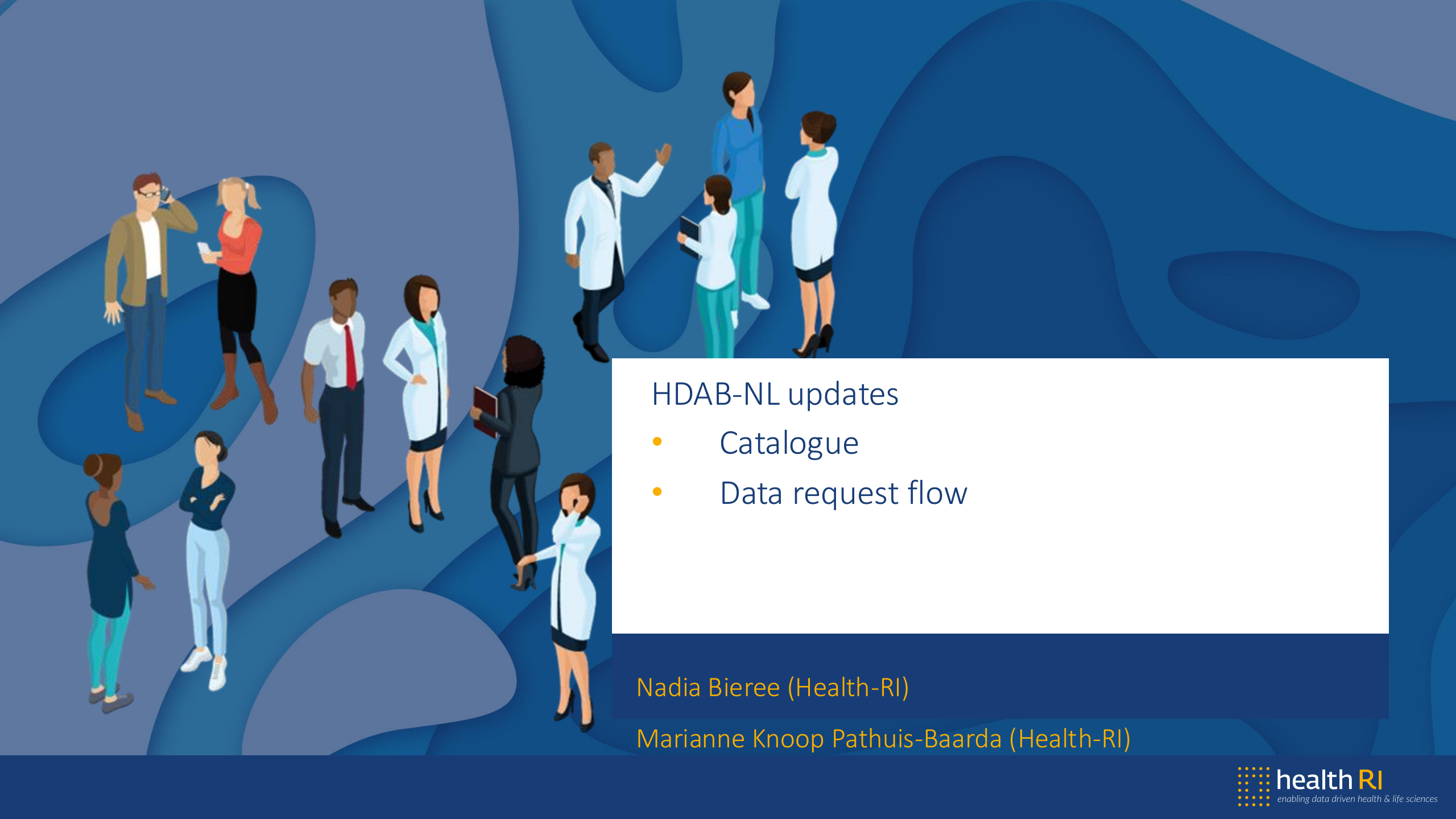


The Hyve

Enabling open science



12:30-13:00



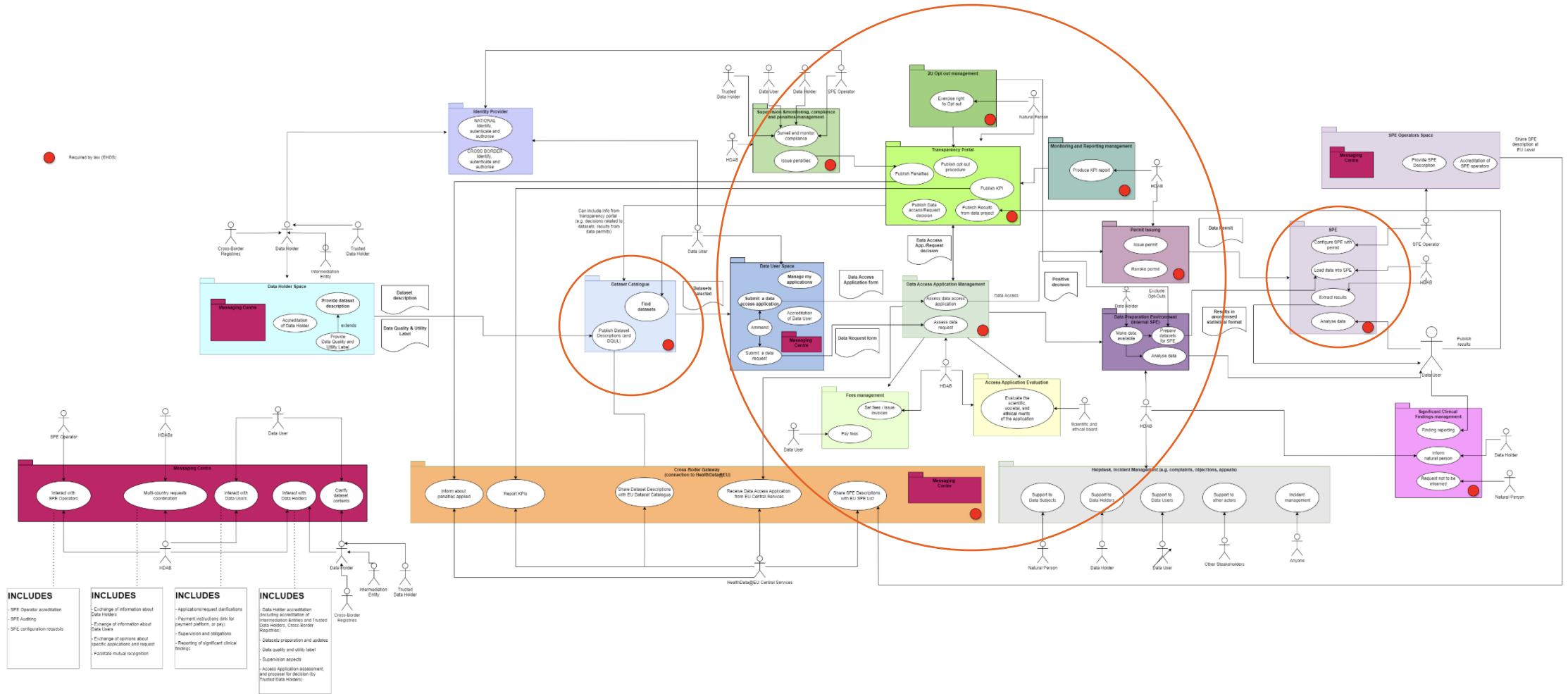
HDAB-NL updates

- Catalogue
- Data request flow

Nadia Bieree (Health-RI)

Marianne Knoop Pathuis-Baarda (Health-RI)

Backbone



HDAB-NL & Catalogue



Marianne Knoop Pathuis - Baarda

Senior Productmanager for the Catalogue at Health-RI

Lead Work Package 6 – Catalogue in HDAB-NL Direct Grant project

Marianne.knoop@health-ri.nl

Our direct grant work packages in 3 phases



1) De uiteindelijke vorm van operationalisatie is nog nader te bepalen

Applications Minimal Viable Products



Data Request Service - for a first version of a Data Access Application Management System (DAAMS), the system that can process data requests and data access (permit) requests.

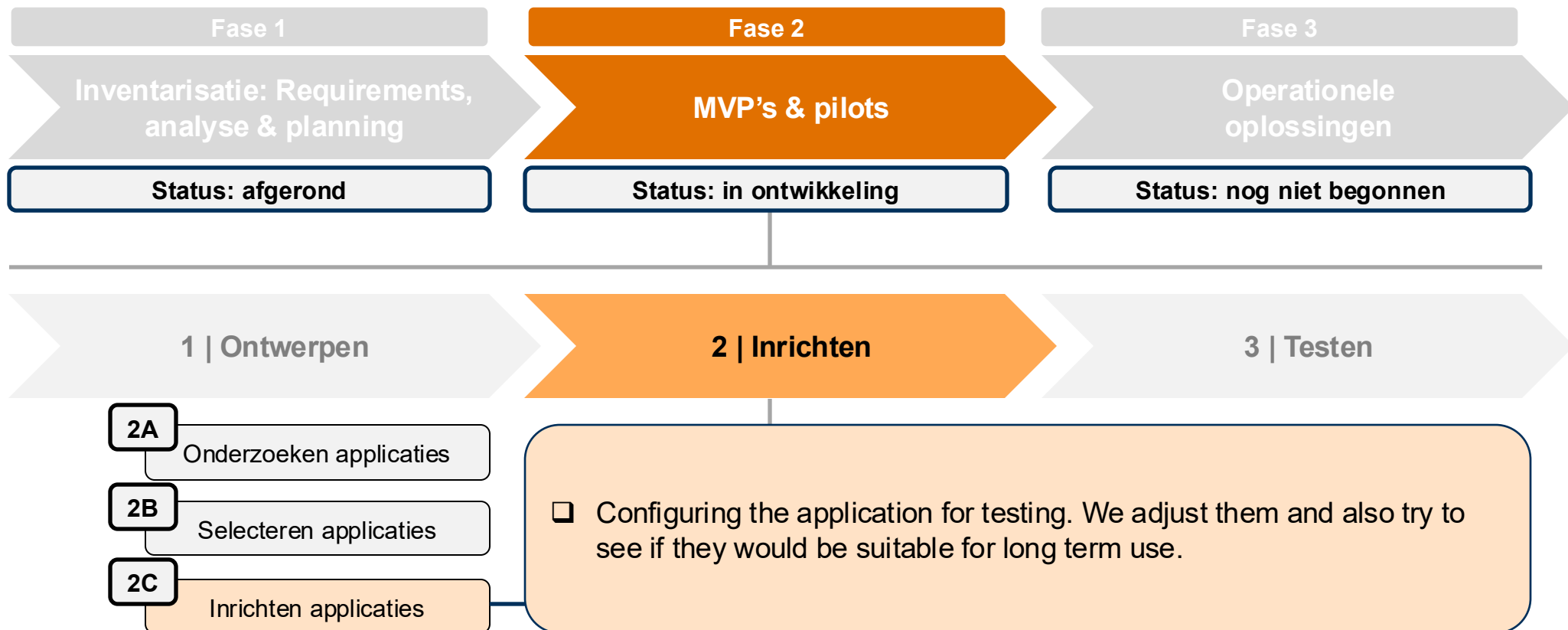


National Health Data Catalogue - For the first version of the national dataset catalogue.



Microdata environment - for a first version of the Secure Processing Environment (SPE), also testing other SPE's (o.a..anDREa, SURF SANE)

In the configuration phase we have researched and chosen applications. Now they are being configured for testing.



Onderzoek plannen

Onderzoek uitvoeren

Onderzoek afronden

Data user

Onderzoeksvraag
opstellen

Data zoeken en
vinden

Data aanvragen

Data analyseren

Data beschikbaar
stellen voor
hergebruik

Toetsing laten uitvoeren
en goedkeuring verkrijgen

Resultaten
publiceren

Gezondheidsdata- infrastructuur secundair gebruik

Nationale
gezondheidsdata
catalogus

Data-
Aanvraagservice

Netwerk van veilige
verwerkings-
omgevingen

Voorzien van generieke functies

Datahouder

Data vindbaar maken

Data aanvraag
verwerken

Data klaarzetten voor
veilige verwerking

Data beschikbaar stellen

Burger

Gezondheidswinst
ondervinden

Transparantie
verkrijgen

Zeggenschap
uitoefenen

Catalogue



catalogus.healthdata.nl

- Considerations HDAB-NL
 - Available for pilot
 - Transferrable to HDAB
 - Modular approach
 - CKAN back-end
 - FDP
 - FE
- CKAN
 - Open source (pro and con)
 - DCAT-AP (but not fully v3)
 - GDI momentum
- HealthDCAT-AP fields added

Filters

Search

Search...

Access Rights

Coding system

Code values

Resources

Frequency

Health theme

Is referenced by

Language

Legal basis

Release date

Start date

End date

440 datasets Clear all filters

Updated (Newest first)

COVID-19 PHASE EDUCATION NATIONAL OTHER PEOPLE BY AGE PEOPLE BY EDUCATION PEOPLE WITH MIGRATION BACKGROUND

Online academisch onderwijs als gevolg van COVID-19 crisis: Voor wie werkt het (niet) en welke factoren kunnen dit verklaren?

Online academisch onderwijs als gevolg van COVID-19 crisis: Voor wie werkt het (niet) en welke factoren kunnen dit verklaren?...

Unique individuals: Dataset holder: Femke Hilverda
Temporal coverage: September 1, 2020 - August 31, 2022 Last updated: September 16, 2021

Request dataset Add to dataset basket

EHR ELECTRONIC HEALTH RECORDS PRIMARY CARE ROUTINE HEALTHCARE DATA

Non-public

Nivel Primary Care General Practices 2011

Electronic health record data from general practices spread throughout the Netherlands. General practice care comprises regular healthcare provision during normal practice hours, excluding out-of-hours primary care, emergency care and hospital admissions...

Unique individuals: Dataset holder: Nivel
Temporal coverage: Last updated:

Request dataset Add to dataset basket

CADDementia

The Challenge on Computer-Aided Diagnosis of Dementia based on structural MRI data. By addressing comparability, generalizability, and clinical applicability, we aim to take a step forwards to clinical use of computer-aided diagnosis (CAD) methods for dementia...

Catalogue

- Details added, filters added
- Some newer HealthDCAT fields will follow after HealthDCAT-AP stability

Home

Hematological malignancies

Request dataset

Add to dataset basket

View my dataset basket

Publisher

Name: IKNL: the Netherlands Comprehensive Cancer Organisation

E-Mail: info@iknl.nl

Description

The data from the hemato-oncology registry of the Netherlands Cancer Registry (NKR) are used to provide insight into the incidence and survival (cancer epidemiology), diagnostic strategies, primary treatment, and the coherence of policy (i.e., the extent to which guidelines are followed) as well as the impact of treatment recommendations for various hematological cancer types.

Health theme oncology**Is referenced by** https://iknl.nl/getmedia/b28517e6-2145-4bad-97a1-3415e8903cd3/hemato_oncologischezorg-in-NL_digitaal.pdf

Keywords

B-cell lymphoma

Follicular lymphoma

Haemato oncology

Leukemia

Lymph node cancer

Multiple myeloma

Myelodysplastic syndromes

Myeloproliferative disorders

Plasma cell tumors

Population coverage Adults aged 18–100 diagnosed with haematological cancer in the Netherlands since 1989**Geographical coverage** Netherlands

Identifier

<http://www.fdp-acc.healthdata.nl/catalog/2d7ab6b8-7edf-4c6a-8227-129d8c877867>

Access rights

Restricted

Legal basis for publishing

```
{ "value": "http://data.europa.eu/eli/reg/2025/327/oj", "label": "http://data.europa.eu/eli/reg/2025/327/oj" }
```

Contact details

IKNL: the Netherlands Comprehensive Cancer Organisation

Documentation

<https://iknl.nl/getmedia/b7276fdb-be51-4f1d-aa3a-e20311af8c4f/RANK-Bestandsindeling-tumorsoort-2025.pdf>https://iknl.nl/getmedia/ce4ead48-ba30-4b10-8df8-955e4c6f1c18/NKR_Itemset_Hemato-oncologie_ALL_april2022_IKNL.pdf

Other responsible organisation(s)

IKNL: the Netherlands Comprehensive Cancer Organisation

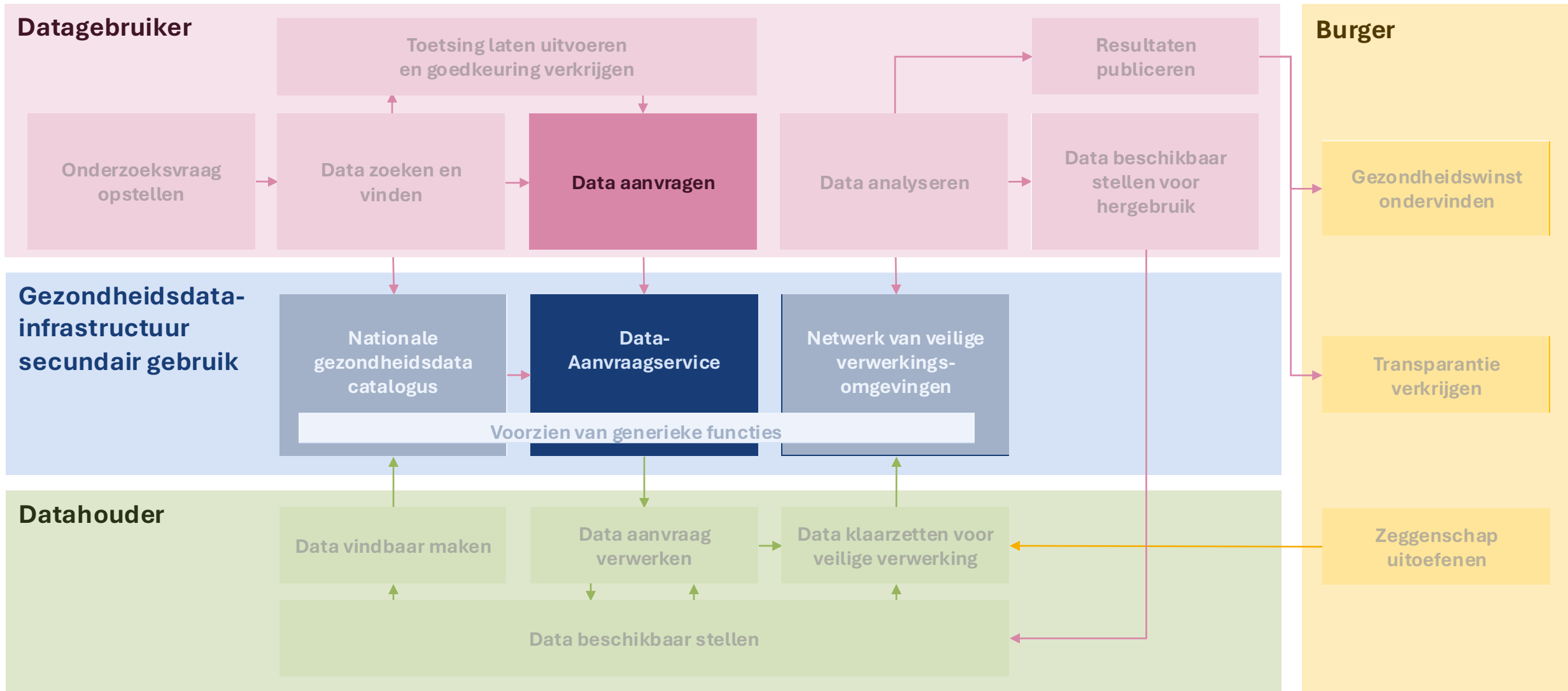
DAAMS

Secundair datagebruik Health-RI: het proces

Onderzoek plannen

Onderzoek uitvoeren

Onderzoek afronden



DAAMS-NL Solution Approach

MVP

- The DAAMS solution will be developed as a Minimum Viable Product (MVP) to validate core functionality and gather feedback from real-world use cases before scaling to a full production system. The MVP approach enables iterative development, early stakeholder engagement, and risk mitigation through phased delivery.

Low-code

- To accelerate implementation and ensure flexibility, the MVP will be built using a low-code platform (Microsoft Dynamics). This choice supports rapid prototyping, easier maintenance, and adaptability to evolving requirements, while reducing dependency on extensive custom development.

Re-use

- The DAAMS system that is to be developed, will be based on the existing Data Request Service application developed by Health-RI, which already provides foundational capabilities for managing data access requests.

ea73c039-bfde-442c-b84e-019462099300 - Unsaved

Request · Data access

Edward Robinson
Owner

Application Business Pro...
Active for 7 days

Received

Pre-Screening

Assessment & Cost Determination

Data Permit/Request Approval

Provisioning (16 Hrs)

Active

Application

Pre-screening

Assessment

Cost determination

Data Permit/Request Approval

Related

Checklists for HDAB employees

Form assist

Data Access Application

Date added

12-11-2025

15:37

Title

Access Application submitted from Insomnia3

Application ID

ea73c039-bfde-442c-b84e-019462099300

Application type

Data Access Application

User ID

sanne.de.vries

Application created

22-9-2025

12:30

Application modified

2025-09-22T10:30:00Z

Version

2

Timeline

Search timeline

Enter a note...

Highlights

Section 6. Description of the data needed

Country ID

NL

HDAB Contacts

Health-RI data steward team

How Data From Different Sources Will Be Linked

Using pseudonymized patient identifiers across datasets.

Person Profile Data Date

22-9-2025

12:30

How the Study Cohort Is Formed

Registry-based selection

Has the study cohort been formed based on information of study participants?

Yes

Does the informed consent cover the requested registry extractions?

Yes

Consent and Information Letter Attachment

Choose File

No file chosen

Confirmation That Data

Nee

↑
JSON request received from
HealthData@EU portal parsed as case

And much more! More
information?



Edward Robinson
Edward.Robinson@health-ri.nl

Mind the gap

- Prior to Data permit application / data request, a series of interactions is needed between data user and data holder:
 - What is the fit-gap with the research question?
 - What (part of) the data is suitable?
 - What to fill in in the request or permit application – taking into account data minimization? (sharing the forms to both data user and data holder?)

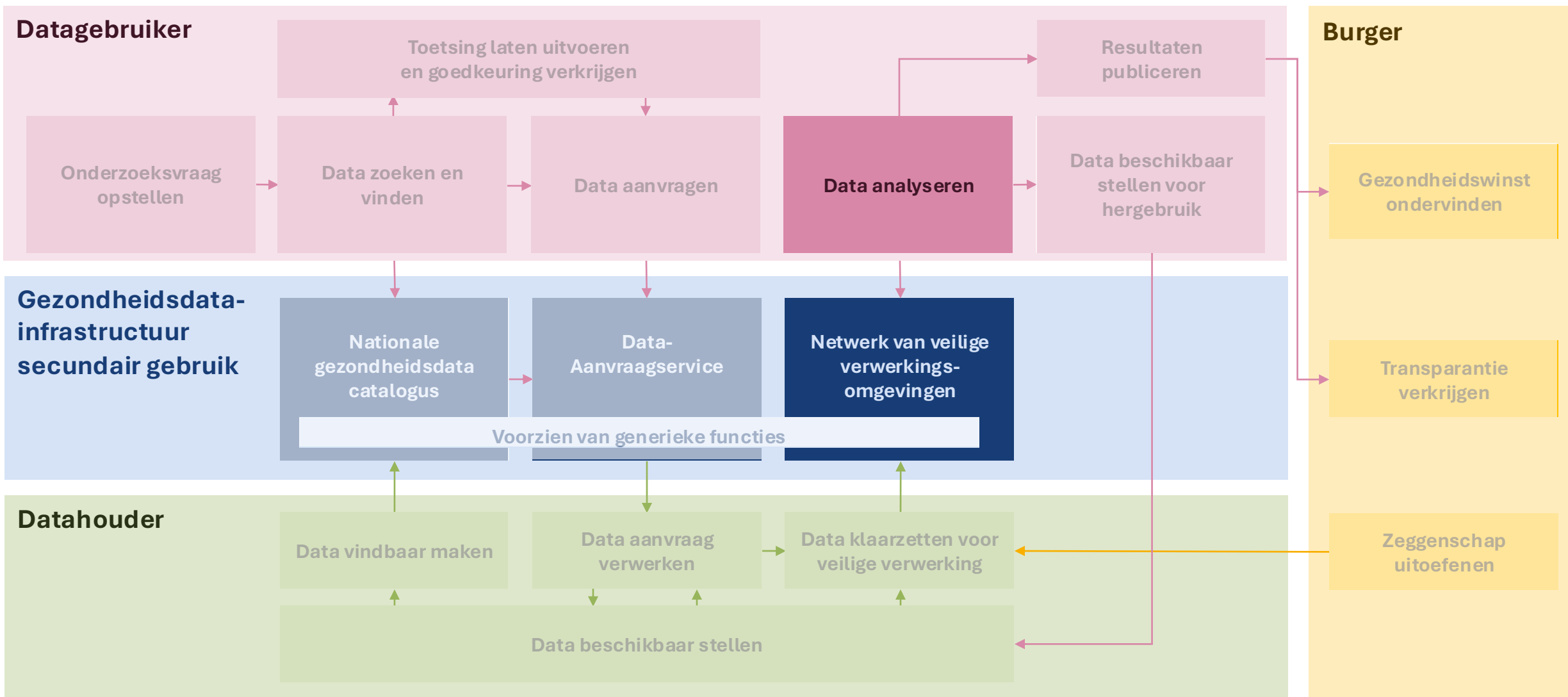
SPE

Secundair datagebruik Health-RI: het proces

Onderzoek plannen

Onderzoek uitvoeren

Onderzoek afronden



Secure Processing Environment

Testing with already available products:

- Microdata environment (National Bureau of Statistics)
- SURF SANE
- AnDREa

In the end working towards a network of validated SPEs, nationally and internationally

To test: How will data preparation work cross-border, and with the European SPE?

Next Steps

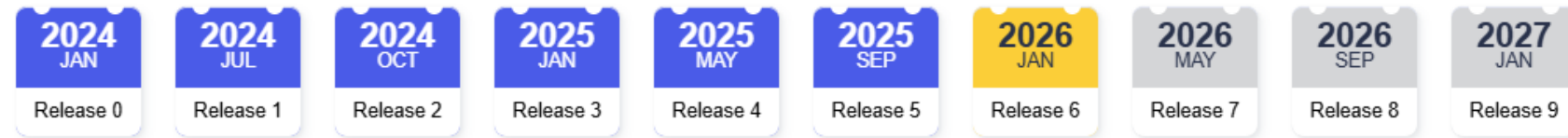
In January and February we will test the process with 7 selected use cases, using the applications as far as they can already be connected. Mocking the interfaces where needed.



1) 'Real life usecases' moeten overeenkomen met realistische scenario's die in de praktijk kunnen plaatsvinden. Tegelijk moeten ze geschikt zijn voor het testen van een **minimal** viable product en hoeven ze niet overeen te komen met 'de meest complexe scenario's die we kunnen bedenken'

HealthData@EU

Delivery Timeline



Objective of Release 6

Release 6 will focus on enabling seamless interaction between the HealthData@EU Central Platform and Member States' Data Access Application Management Systems (DAAMS). This release will introduce key interoperability features to support application status exchange, bidirectional communication, and decision management.

Planned features

The planned features include:

- Enabling the HealthData@EU Central Platform to receive detailed DAAMS status updates via the National Contact Points (NCPs).
- Supporting message exchange between users and Health Data Access Bodies (HDABs) through the platform and NCPs.
- Allowing users to modify submitted Data Access or Data Request Applications upon HDAB request.
- Displaying final application decisions (positive or negative) transmitted by the NCP on behalf of the HDAB directly in the user's profile.
- Creating a European registry of Data Access and Data Request decisions.
- Allowing users to request amendments to a positive Data Access Decision (Data Permit).
- Allowing users to submit appeals following negative Data Access or Data Request decisions.



Primair

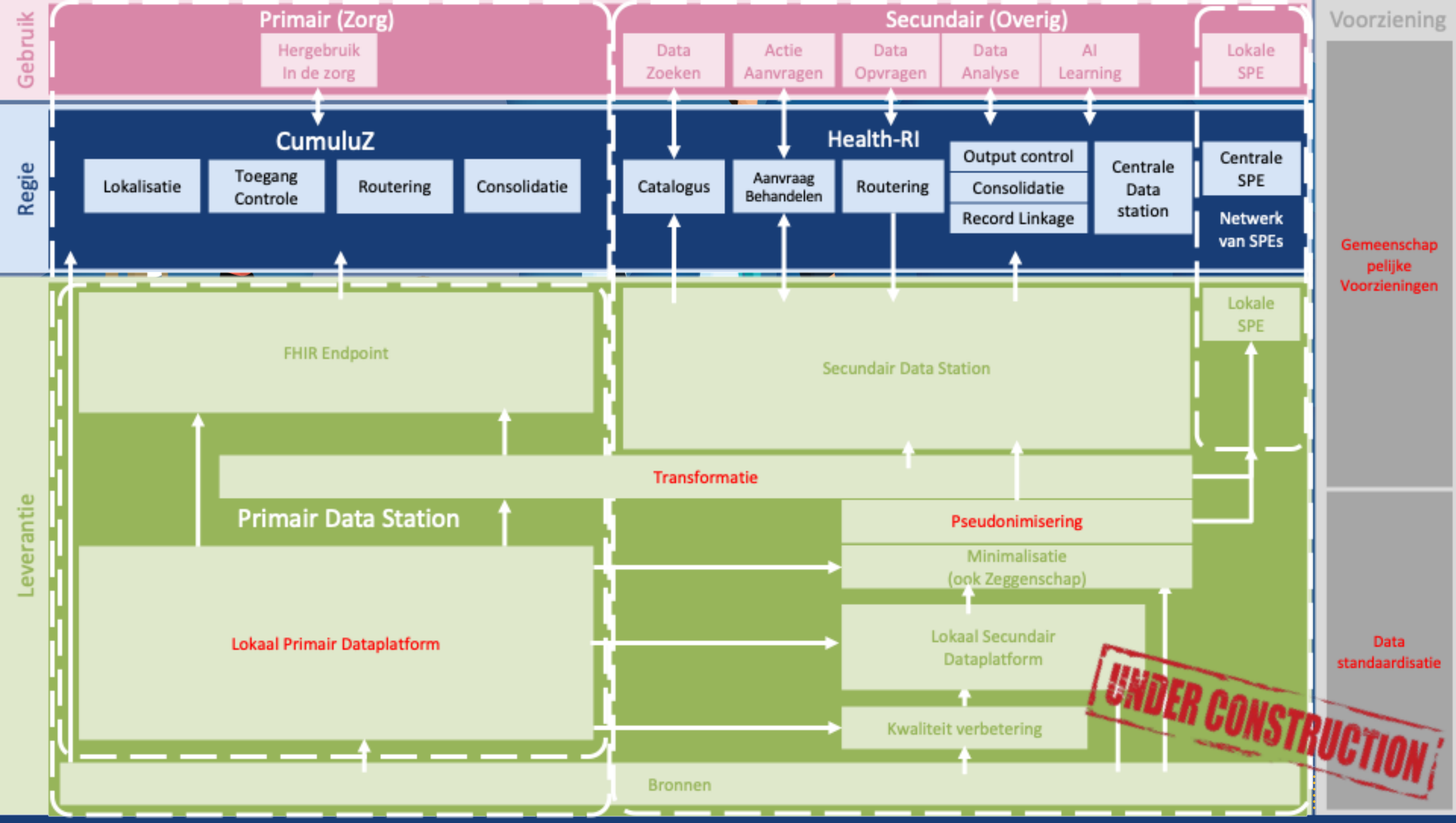


health RI

enabling data driven health & life sciences

Secundair

Landelijk Dekkend Netwerk



Questions?

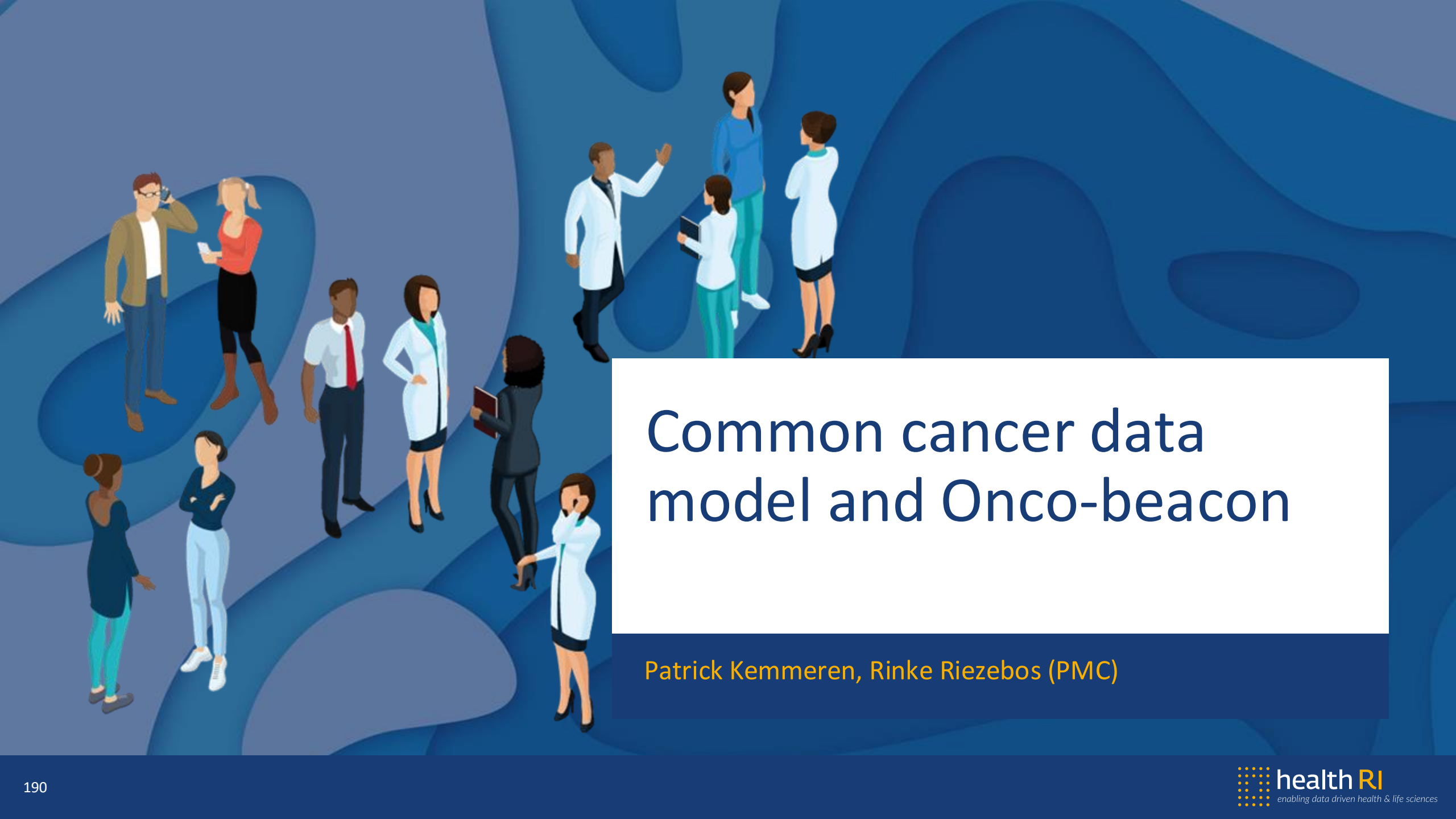


Marianne Knoop Pathuis - Baarda

Senior Productmanager for the Catalogue at Health-RI

Lead Work Package 6 – Catalogue in HDAB-NL Direct Grant project

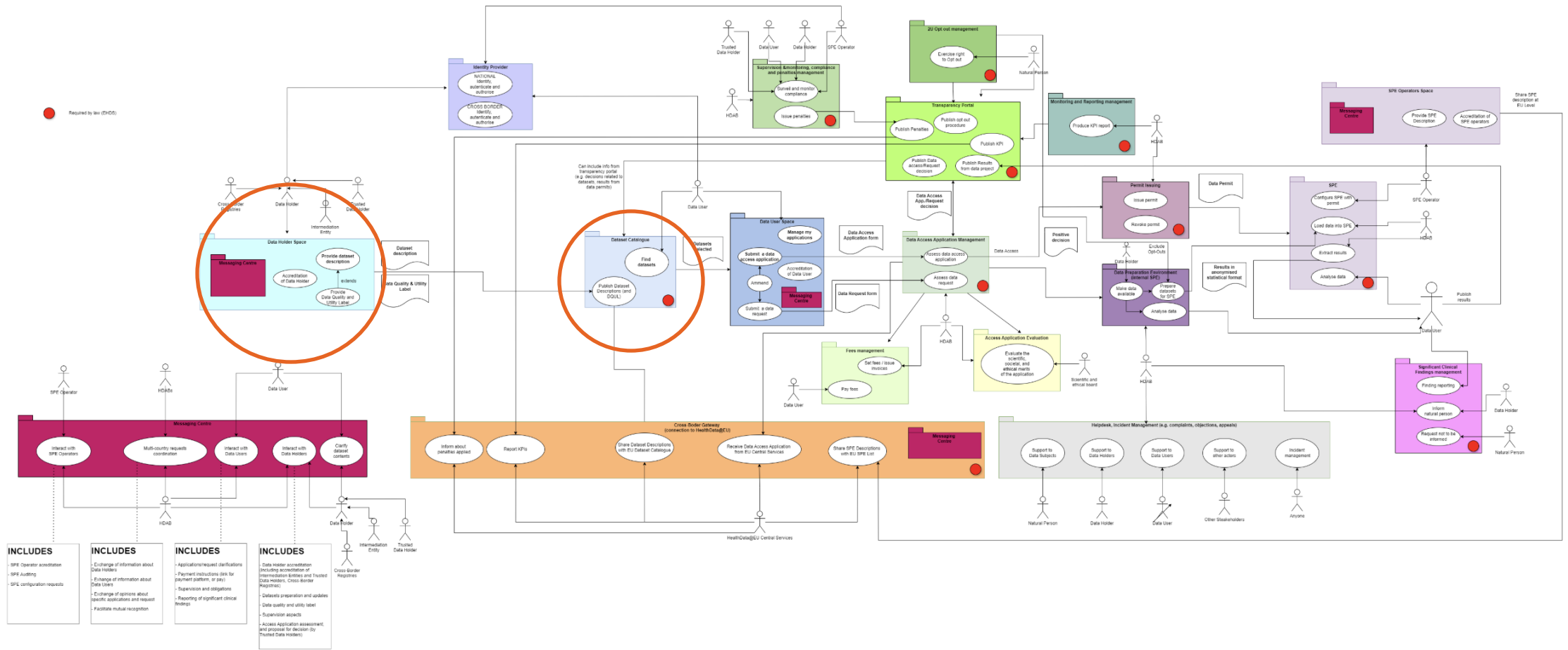
Marianne.knoop@health-ri.nl

An illustration of a diverse group of healthcare professionals, including doctors, nurses, and researchers, standing in a modern, blue-toned environment. They are dressed in white lab coats, scrubs, and business attire, engaged in various activities like talking on a phone, holding a clipboard, or gesturing during a discussion. The background features large, abstract blue shapes.

Common cancer data model and Onco-beacon

Patrick Kemmeren, Rinke Riezebos (PMC)

Backbone



Overarching aim

Establish recommendations and agreements on the use of a **common data interoperability model** (CIM) to foster data interoperability for **secondary use** in **oncology**



Key principles

- Is focused on secondary use of oncology data
- Is an umbrella model that reuses existing internationally established models
 - openEHR, FHIR, OMOP, CDISC
- Supports single and multi patient/sample queries
- Is never going to be perfect, 80/20 rule is key
- Supports a few selected use-cases for practical usage

User stories should drive the use-cases

As a **clinician**, I want a data-driven tool that offers personalized treatment recommendations using clinical guidelines, patient data, trial eligibility, and outcomes from similar cases, to inform my decision for tumor board discussions.

As a **(biomedical) researcher**, I want to use sequencing data in combination with medical imaging, histopathology and clinical data to investigate which molecular and cellular characteristics relate to different cancer types, outcome and quality of life.

As an **epidemiologist**, I want

Standardise and communicate



1

Modeleringstandaarden

Welke en hoe passen we deze toe?

De meest gebruikte (inter)nationale modeleringstandaarden* en hoe deze worden toegepast in het Máxima.

*Preklinische standaarden worden later toegevoegd

1.1

OpenEHR

Focust zich op het vastleggen breed scala aan generieke en specialistische klinische gegevens voor langdurige opslag.

- Hoge dekking breed toepasbaar
- Consistente structuur
- Flexibel & robuust



1.2

FHIR

Richt zich op structuur voor uitwisseling en deling klinische en administratieve gegevens om interoperabiliteit te bevorderen.

- Hoge interoperabiliteit
- Gestandaardiseerde formats
- 80/20 regel voor dekking



1.3

Zibs

Richt zich op zorginformatie binnen de Nederlandse zorg, met als doel eenduidige en herbruikbare gegevensmodellen

- Nederlandse focus
- Niet volledig dekkend
- Opgenomen in de Wegiz



1.4

OMOP

Focust zich op een uniform datamodel voor het vastleggen van gegevens ten behoeve van observationele studies en onderzoeksdoeleinden.

- Gestandaardiseerd datamodel
- Ondersteunt onderzoek
- Hoge interoperabiliteit



1.5

CDISC

Focust zich op het uniform vastleggen en uitwisselen van klinische studiegegevens in farmaceutisch en biomedisch onderzoek.

- Efficiëntie in gegevens analyse
- Regelgevende acceptatie
- Strikte structuur



Mapping is (deels) mogelijk

Sommige standaarden kun je mappen voor eenduidige en bruikbare uitwisseling van informatie.

↔ Deels
↔ Volledig

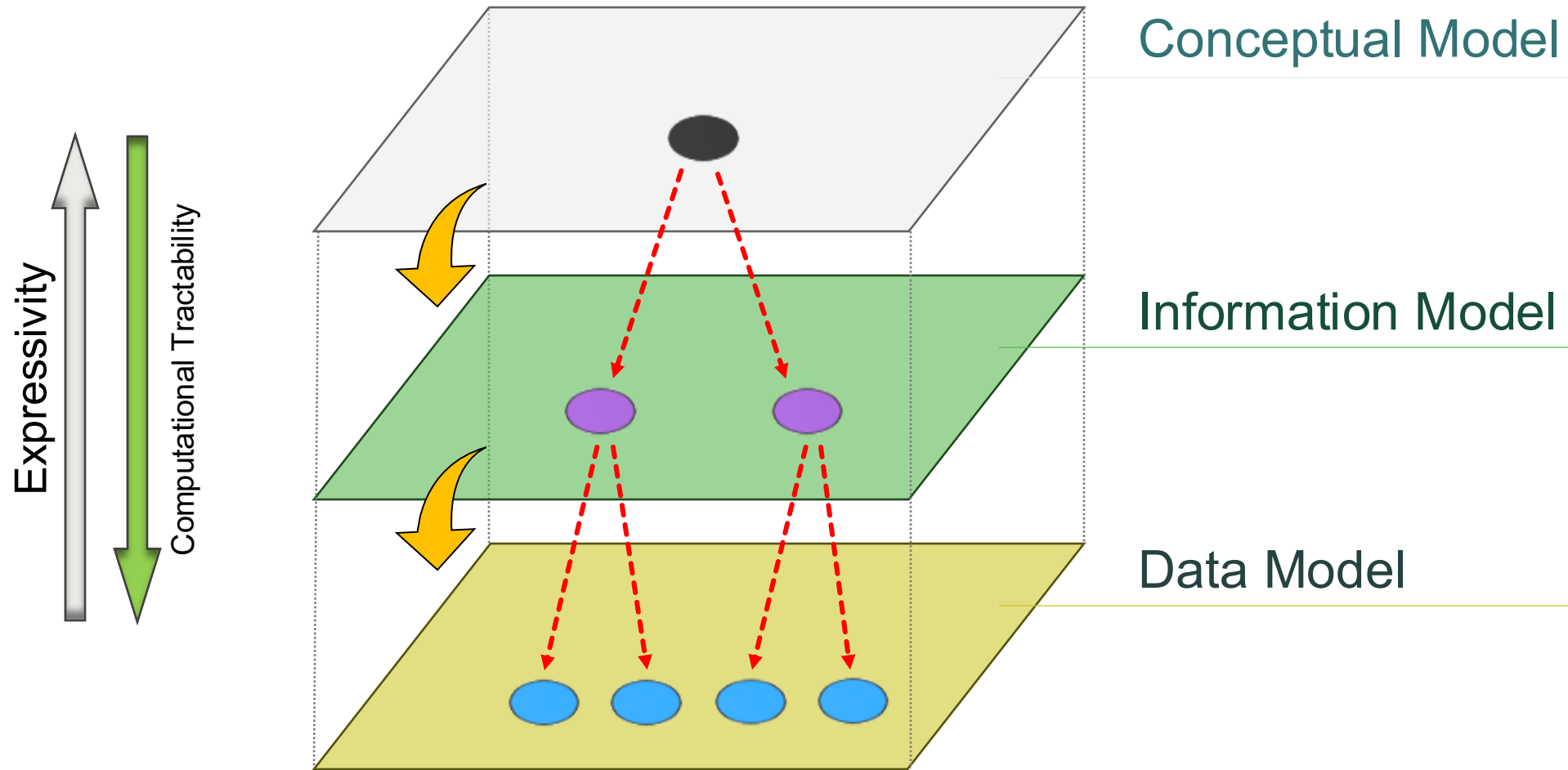


Wat gebruiken we in het Máxima?

Afhankelijk van het doel gebruiken we één of meerdere standaarden.

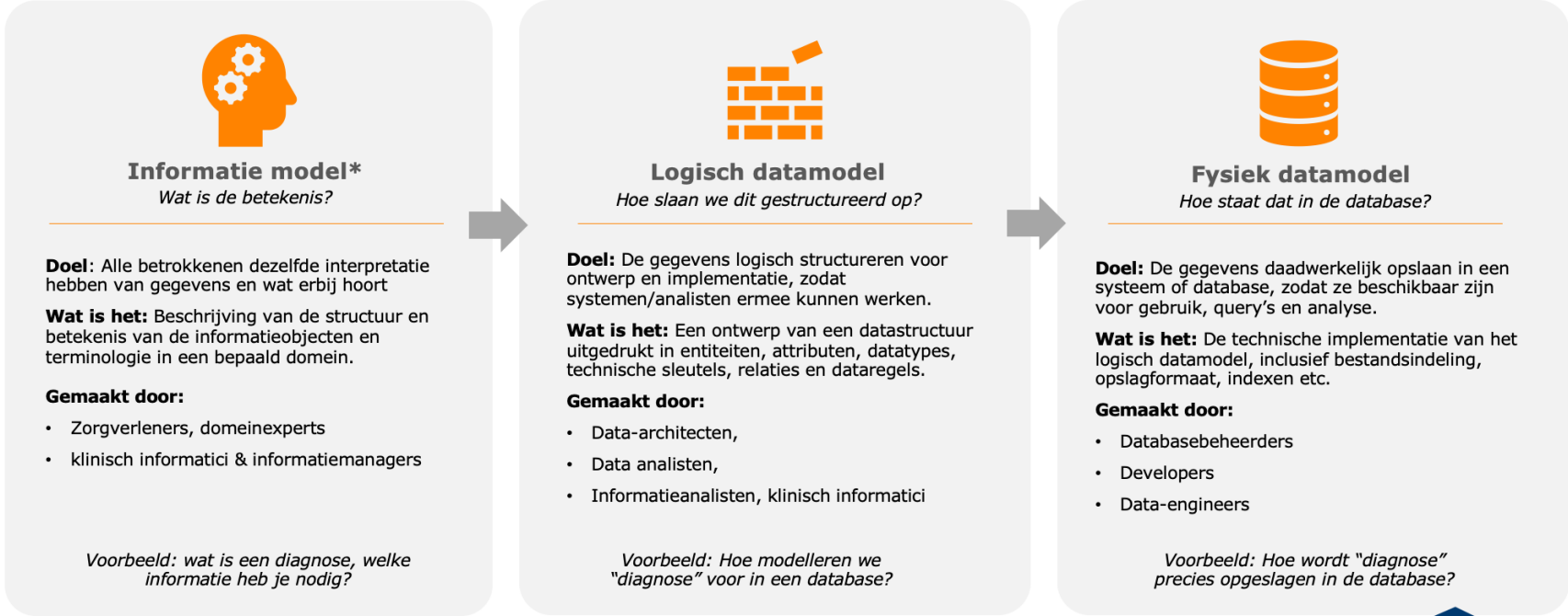
- **OpenEHR:** wordt o.a. gebruikt door TDC en IDT voor structureren van klinische data en ontwikkeling van evaluatie en beslisalgoritmen
- **CDISC:** worden gebruikt voor Intended-to-File studies die o.a. ingediend worden bij FDA
- **Zibs:** t.b.v. VIPP 5 programma voor uitwisseling in bgz

Model-driven development



Van domeinbegrip tot implementatie: 3 modelleerlagen

Elke laag bouwt voort op de vorige: het informatiemodel geeft betekenis, het logisch model vertaalt dit naar datastructuur, en het fysiek model zorgt voor opslag en vindbaarheid.



Focus van projecten:



Ter info – voor nu buiten scope

*Lijkt erg op het begrip conceptueel datamodel, voor simpliciteit deze laag niet meegenomen

An inhouse example

Way forward – gekozen oplossingsrichting

Gekozen oplossingsrichting:



Centraal informatie model

Wat is de betekenis?

KIS als centraal informatie model

- **KIS wordt uitgebreid** met bouwstenen/templates benodigd preklinisch (bijv. biomaterial)
- Valideren en/of aanpassen bouwstenen met nieuwe stakeholders
- Betrekken/uitbreiden KIS team met nieuwe stakeholders



In lijn met elkaar



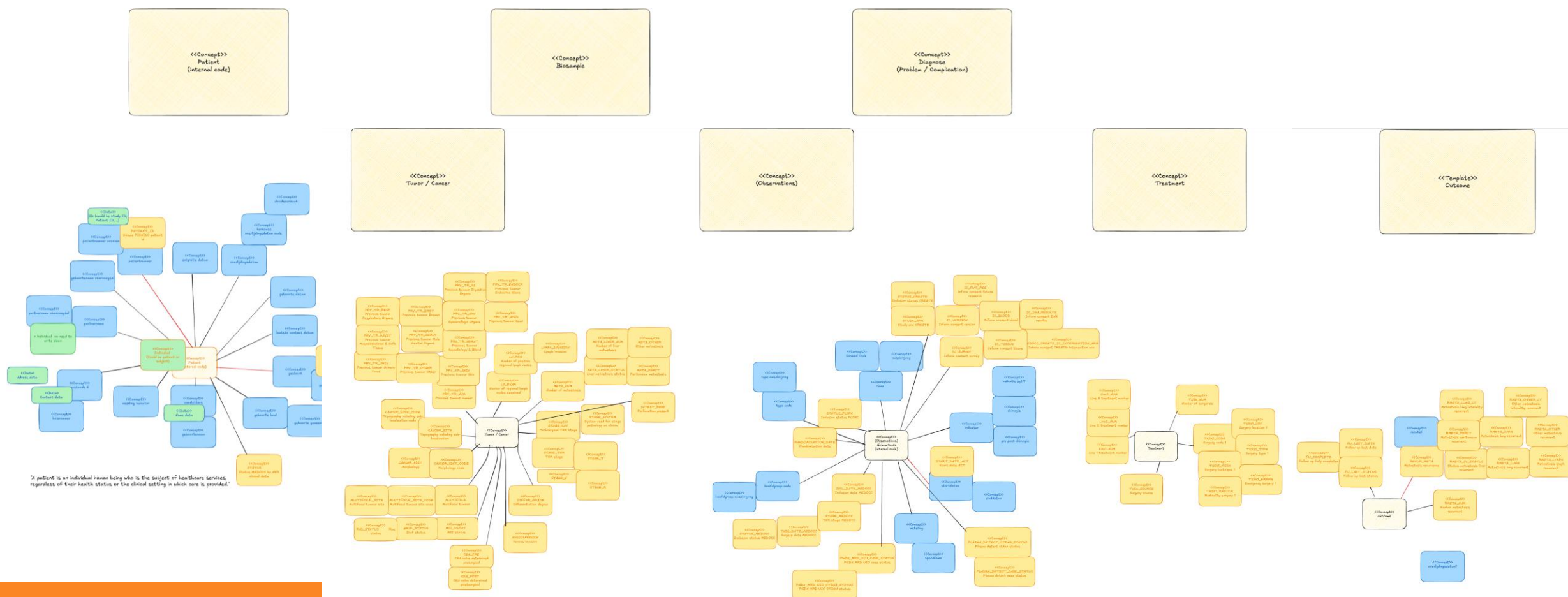
Common data model

Hoe slaan we dit gestructureerd in de database?

Pragmatisch naar één common data model (logisch)

- Eén logisch datamodel voor kinderoncologie (Máxima)
- **CSR wordt uitgebreid** met KIS als kaders voor één centraal datamodel (logisch)
- Pragmatische insteek, uitgangspunt alle entiteiten centraal. Mocht dit niet passend zijn, altijd nog de mogelijkheid om via mappings aparte entiteiten te definiëren.

Alignment oncology core concepts

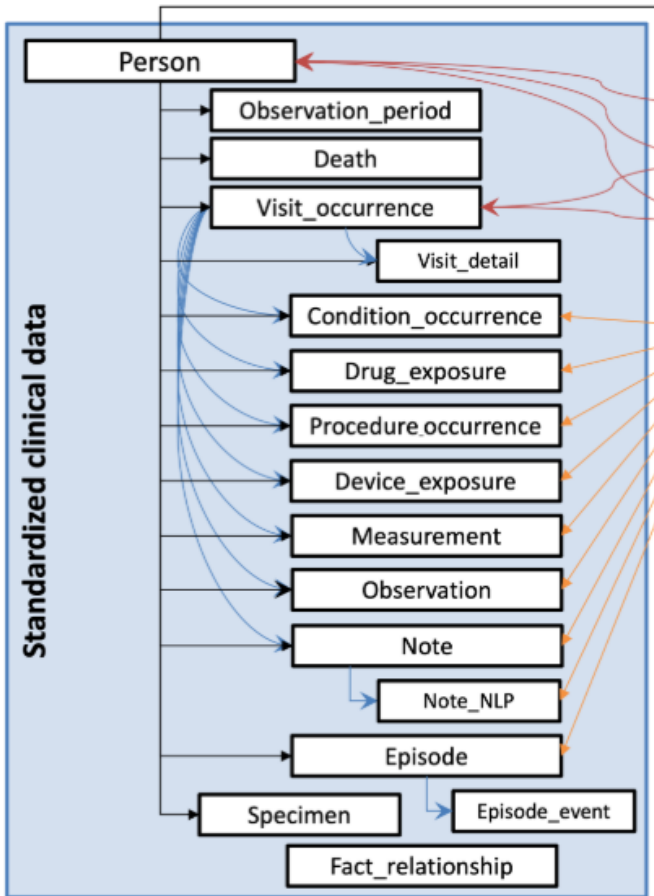


Focus on core concepts & mapping to existing standards

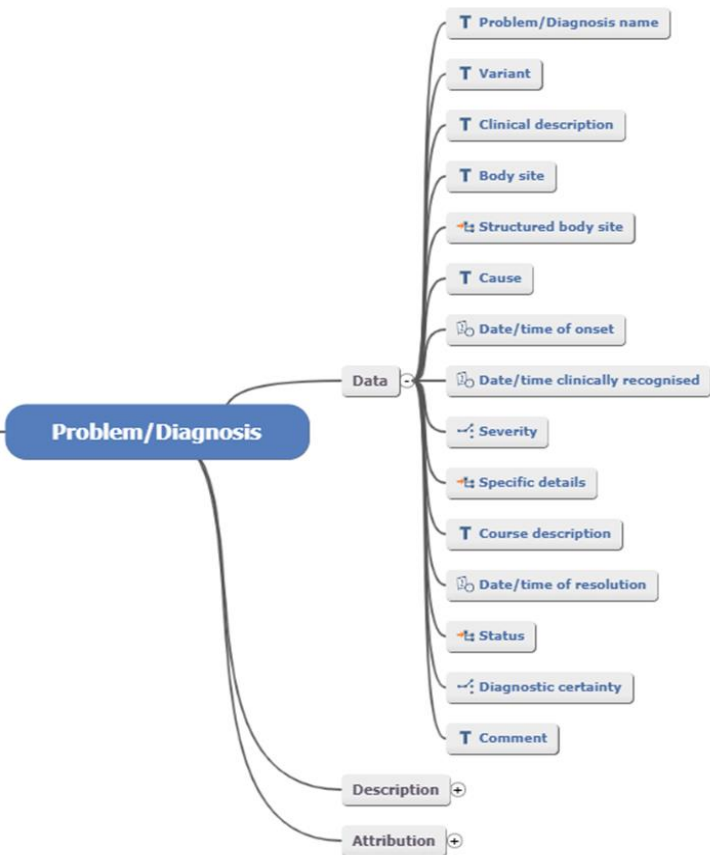
Highest level of classes in the data model							
1+MG-MDC		OMOP		mCODE FHIR		AACR GENIE	
Domain	Item description	Domain	Item description	Domain	Item description	Domain	Item description
1_SUBMITTER	Information about the data submitter						
2_PATIENT	Cancer patient's personal information and demographics	PERSON	(Information about the person)	Patient Information Group	Patients demographics	Patient Demographics	Patient demographics, sequencing biosample information) + Social Determinants of Health
3_SEQUENCE	Details of the DNA sequence (useful for grouping purposes based on technology and protocols and to account for technical differences and possible biases)	EPISODE	(aggregates lower-level clinical events into a higher-level abstraction representing clinically and analytically relevant disease phases, outcomes and treatments.)	Disease Characterization Group	The mCODE Disease Characterization group includes data elements specific to the diagnosis and staging of cancer.	Cancer Diagnosis (by Dx)	Cancer diagnosis and tumour staging
4_SAMPLE	Details about the source (i.e., normal tissue, primary cancer, metastasis) of the analysed genomic material	EPISODE EVENT	(connects qualifying clinical events to the appropriate EPISODE entry.)	Health Assessment Group	The mCODE Assessment group contains information related to the patient's general health, before and during treatment.	Imaging	Type of imaging technique, disease status according to imaging
5_DISEASE_CANCER	Cancer disease description (histology, staging, and grading) to be reported for each cancer-related event	CONDITION OCCURRENCE	(records of Events of a Person suggesting the presence of a disease or medical condition stated as a diagnosis, a sign, or a symptom, which is either observed by a Provider or reported by the patient.)	Genomics Group	mCODE includes genomics-related data elements needed to inform cancer assessment and treatment options.	Clinical Visits	Clinical visits - follow-up
6_DISEASE_COMORBIDITY	Co-occurring medical conditions other than cancer	PROCEDURE OCCURRENCE	(activities or processes ordered by, or carried out by, a healthcare provider on the patient with a diagnostic or therapeutic	Cancer Treatments Group	The Treatment group includes reporting of procedures and medications used to treat a cancer patient, or relevant to that treatment. Treatments fall into three classes: medications, surgery, and radiotherapy.	Procedures	Procedure type, intent
7_TREATMENT	Therapeutic interventions for cancer and comorbidities	MEASUREMENT	(structured values, numerical or categorical, obtained through systematic and standardized examination or testing on a Person or Person's sample - predominately lab tests with a few exceptions, like blood pressure or function tests.	Outcomes Group	Recording patient outcomes in mCODE involves disease status, tumor size, and date of death.	Treatments	Treatment types, medications, toxicities
8_TREATMENT_CANCER_SURGERY	Surgical procedures for cancer	DRUG EXPOSURE	(drug exposure)			Labs and Biomarkers	Lab test used, results
9_TREATMENT_CANCER_CHEMOTHERAPY	Chemotherapy for cancer					Outcomes	Follow-up information, outcome data
10_TREATMENT_CANCER_RADIATION_THERAPY	Radiation therapy for cancer						
11_TREATMENT_CANCER_HORMONAL_THERAPY_AND_BIOLOGICAL_THERAPY	Hormonal therapy and biological therapy (including immunotherapy, targeted therapy...) for cancer						
12_ENVIRONMENTAL_EXPOSURE	Exposure to environmental factors for which an association with cancer is established (or reported)						
13_BIOMARKER	Biomarkers with clinical value						

Identificeren van informatiebouwblokken

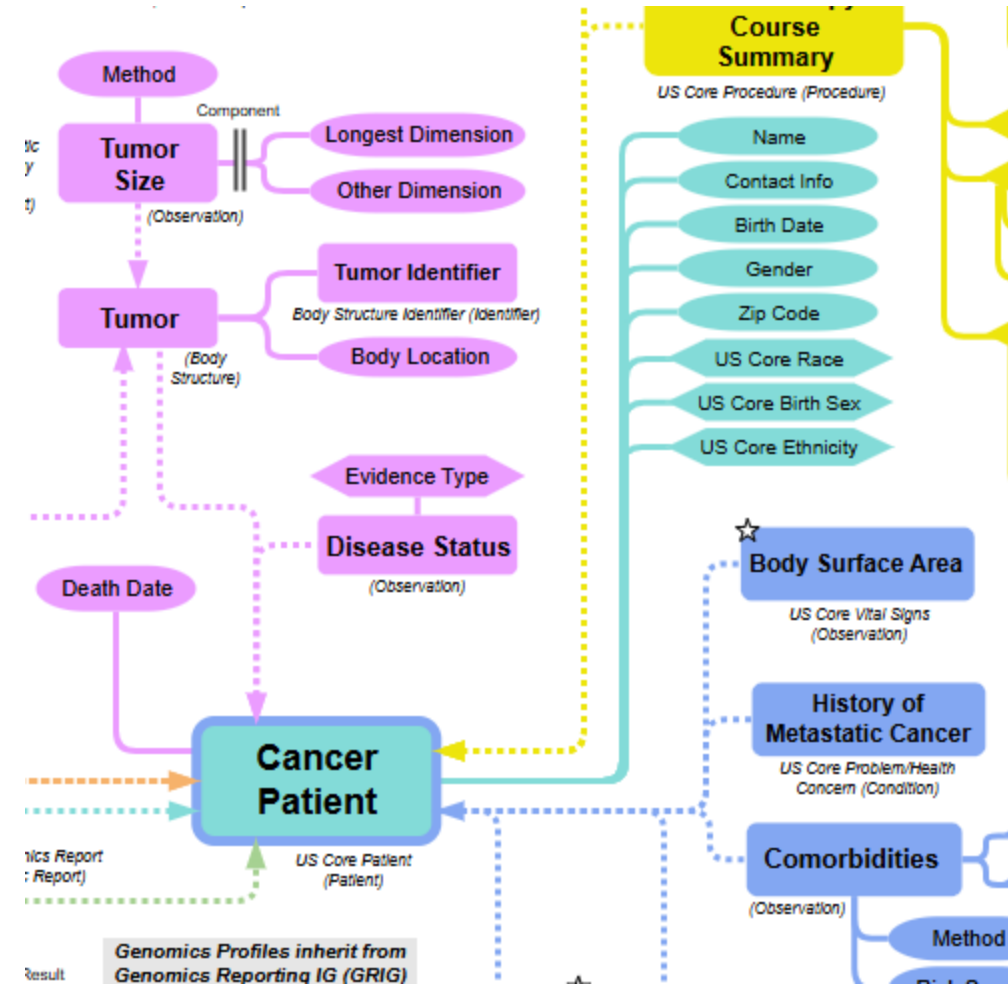
OMOP Common Data Model



openEHR



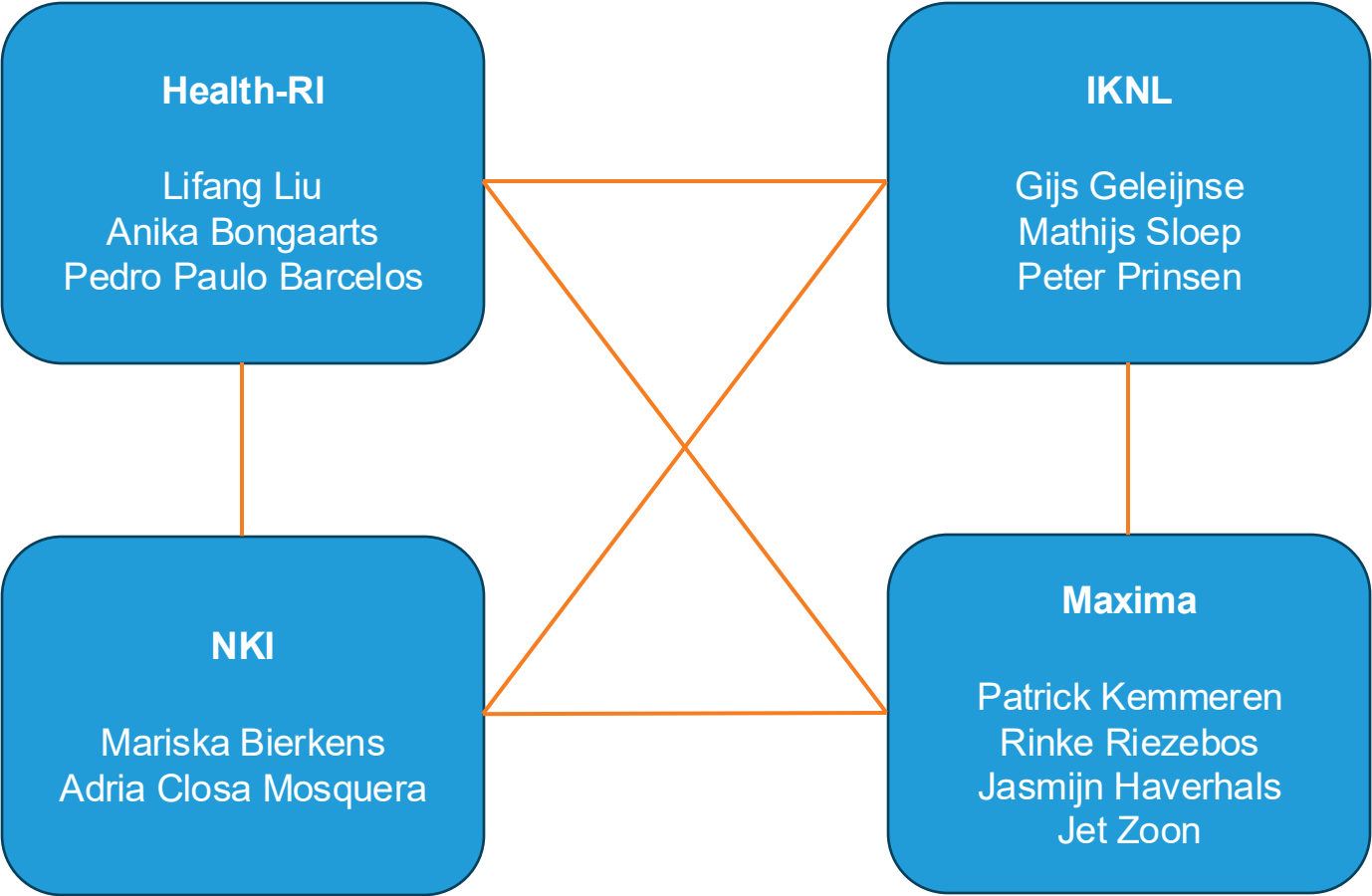
mCODE STU 4



Next steps

- Version 1.0 (almost) ready to be further shared
- Mapping (and overarching) to existing international standards: openEHR, HL7-FHIR, MG1+, OMOP
- Run actual use-case with NKI, IKNL & PMC

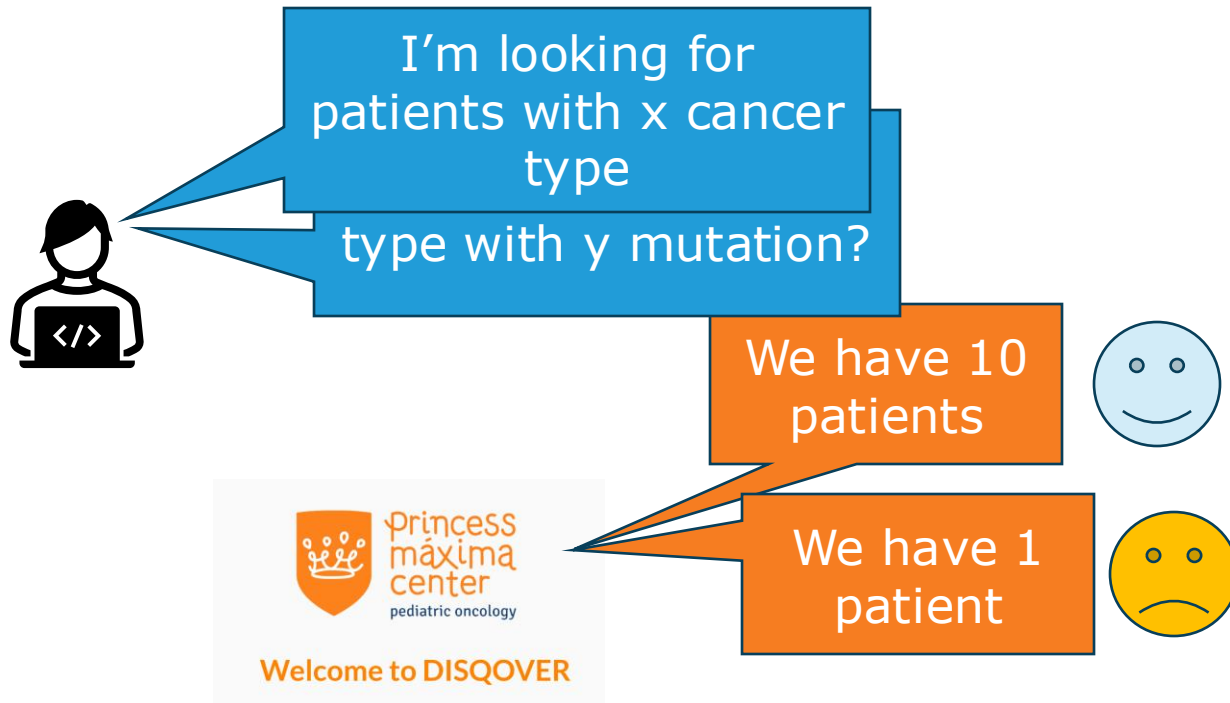
Core CIM working group





OncoBeacon: a federated catalog for integrated clinical and molecular cancer data

Hinri Kerstens



What would the answer have been if we could ask the question globally?

Federated data, the cancer use case

Cancer is a leading cause of disease related death worldwide.

- Most drugs don't work well
 - ~30% response rate is considered good
 - molecular mechanisms are not fully understood
- We need statistical power to find new biomarkers for (non)response
 - clinical trials hampered due to the low number of (known) cases
 - data lives in silos
 - need integrated clinical and molecular data

Survival rates have increased over the past decades but are still low in particular for some very rare malignancies.

Aim: build a (inter)national (NL) beacon genomics network

Context:

- no top down national (govt) omics agenda
- grass-roots 'data-holder' initiative supported by GDI/Health-RI

Outcomes:

- Discover and integrate datasets
- Scalable searching
- Harmonised data structures & ontology usage

Finding federated data? Beacon

An API specification for genomic/phenoclinical data discovery while ensuring privacy, developed by GA4GH

- request/response format
- biological data format



Global Alliance
for Genomics & Health
Collaborate. Innovate. Accelerate.



Beacon

Beacon Components

"Beaconized" dataset

Cohorts >50 (3 required)

- Cohort metrics
- Cohort design
- ...

Individuals >80 (2 required)

- Sex
- Diseases
- PhenotypicFeatures
- ...

Biosamples >50 (3 required)

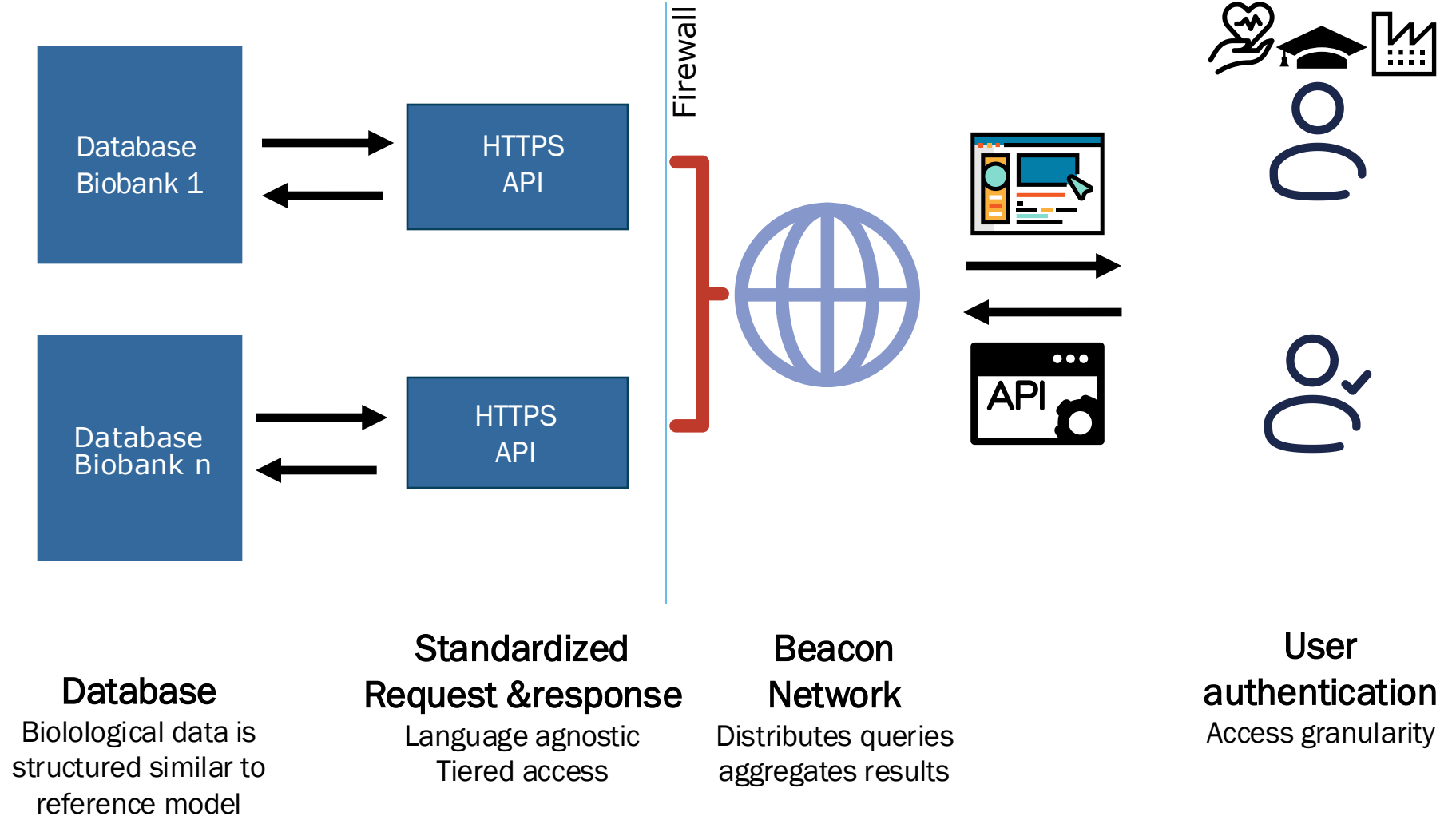
- Origin (tissue)
- Status (tumor/normal)
- Individual
- ...

Analyses 10 (3 required)

- Pipeline
- Biosample
- ...

genomicVariations >150

- Variant type
- Case level data
- Individual
- ...



One specification, multiple implementations

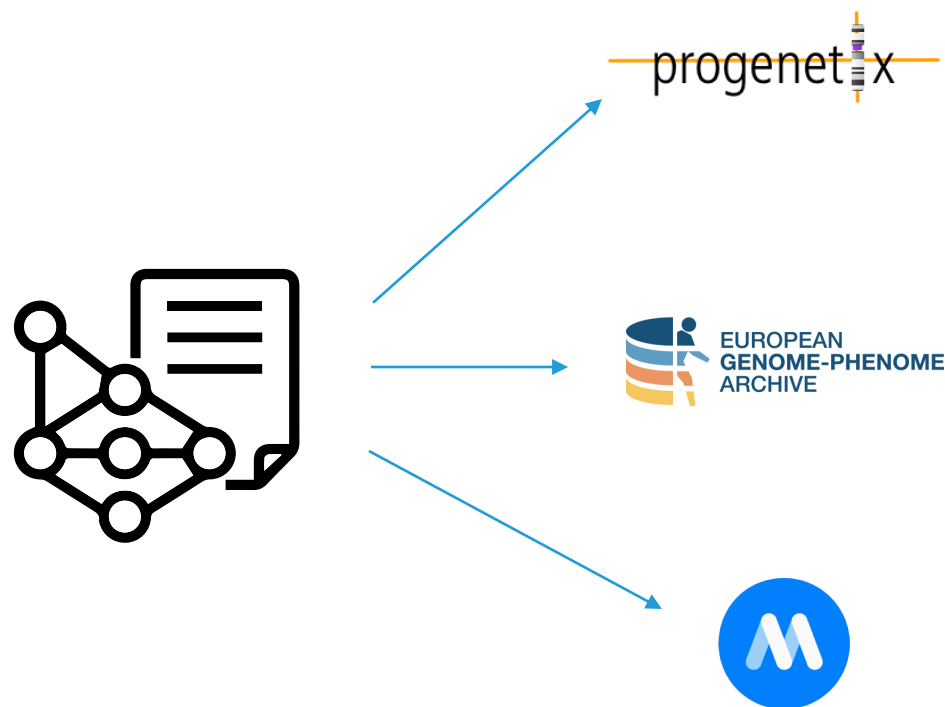


Preference based on technical requirements



	Progenetix	BeaconV2-pi	Molgenis
Permission granularity	Yes	Yes	Yes
Data access granularity	Experimental	Dataset level	Database level
Federated authentication	No	Yes	Yes
Coding style	Less structured	Structured, object oriented	Structured, object oriented
Code comments	Lacking	Lacking	Few
Unit Testing (framework)	Few	Yes	Yes
Logging/exception handling	Few	Yes, stepwise	Yes
Documentation	Incomplete	Near complete	Near complete
Ingestor/Beaconizer	Yes	beacon2-ri-toolsv2	No
Containerized deployment	No	Yes	Yes
Configurability	Hard	Easy	Intermediate

And to what extent platforms adhere to the specifications



Synthetic data

Beacon platforms



Standard queries at
all complexity levels



Not perfect
Closely matches
expectations/specifications
Code lends itself to further
development/improvement

The ability to map cancer genomics attributes

Feature	Progenetix	BeaconV2-pi	Molgenis
Tumor (neoplastic)	biosampleStatus	biosampleStatus	PathologicalState
ICDO morphology	icdoMorphology	histologicalDiagnosis?	IndividualsDiseases.diseaseCode
ICDO topography	icdoTopography	sampleOriginType?	sampleOriginType?
Pathological stage	pathologicalStage	pathologicalStage	
Tumor grade	tumorGrade	tumorGrade	
Primary/relapse/metastasis	tumorProgression	tumorProgression	IndividualsDiseases.stage
Cancer of unknown primary	histologicalDiagnosis.value	histologicalDiagnosis?	
Bone marrow transplant status	interventionsOrProcedures*	interventionsOrProcedures*	
Reference sample (blood,skin,..)	biosampleStatus.value	biosampleStatus?	
Age at diagnosis	interventionsOrProcedures*	interventionsOrProcedures*	IndividualsDiseases.ageAtDiagnosis**
Pre-treatments	treatments	treatments	
FFPE/FF + duration?	sampleStorage	sampleStorage	
Molecular subtype	measurements	measurements	
Mutation signature	measurements	measurements	
Homologous recombination def.	measurements	measurements	
Tumor drivers	measurements	measurements	
Tumor purity	measurements	measurements	

The Beacon schema

The schema used in Beacon heavily relies on the use of hierarchical ontologies.

Beacon term (179)

BiosampleStatus

"Ontology value from Material
Entity term (BFO:0000040)"

Properties

id, label



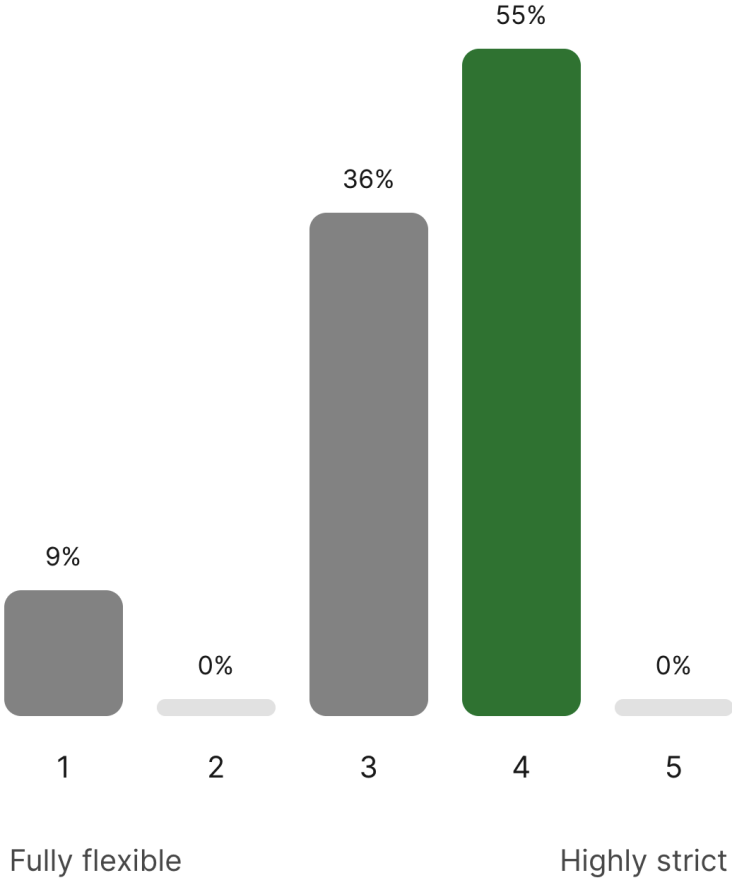
```
"id": "EFO:0009654", "label": "reference sample"  
"id": "EFO:0009656", "label": "neoplastic sample"
```

Structured but **not** always **harmonized** data

Harmonization of cancer genomics attributes in Beacon

Score: ★ 3.4

Should we be more flexible or stricter?



Structured and **more harmonized** data

Adding and harmonizing cancer data attributes

Adding to and harmonizing the (onco)Beacon schema involved three steps

- Inventory of oncology-specific data attributes not represented by Beacon terms
- Use generic concepts from the standard beacon model to describe these attributes
- Apply (require) specific ontology classes in attribute ids and values

Specific attribute	Beacon concept	Ontology classes
Tumor Mutational Burden	assayCode date (optional) measurementValue	NCIT:C181335 - Tumor Mutation Burden Assessment string NCIT:C188348 - Mutations per Megabase

Collecting data in the Princess Máxima Center

Collected information from 50 patients (59 samples) in the iTHER* studies.

- Structured and curated data sources.
- Manual extraction from pathology reports, under review.
- Automated, partially validated analysis workflows, needs review.

Current data provides an overview of what we can report; the values are not yet guaranteed to be accurate.

* DOI: 10.1016/j.ejca.2022.09.001: Implementation of paediatric precision oncology into clinical practice: The Individualized Therapies for Children with cancer program 'iTHER'

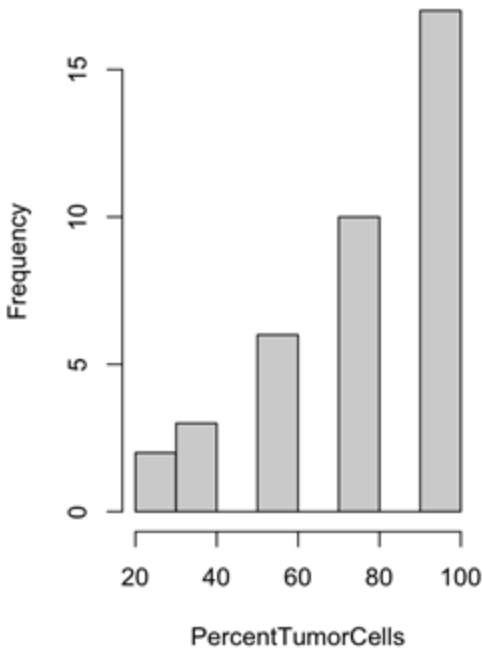
Processing data in the Princess Máxima Center

Attribute	Scope	Terms extracted (269)	Id queried	Query result
sampleStatus	biosample	✓	EFO:0009655	✓
histologicalDiagnosis	biosample	✓	ICDO3:89633	✓
sampleOriginDetail	biosample	✓	ICDO3:C402	✓
sampleOriginType	biosample	✓	OBI:0001479	✓
sampleStorage	biosample	✓	NCIT:C185338	✓
pathologicalStage	biosample	✓	NCIT:C27971	✓
tumorProgression	biosample	✓	NCIT:C84509	✓
pathologicalTnmFinding	biosample	✓ After fix in beaconV2-pi	NCIT:C48700,NCIT:C48706,NCIT:C48724	✓
Tumor Grade	biosample	✓	NCIT:C14162	✓
Tumor Purity	biosample	✓	NCIT:C159484	✓
PreoperativeChemotherapy	individual	✓	NCIT:C68770	✓

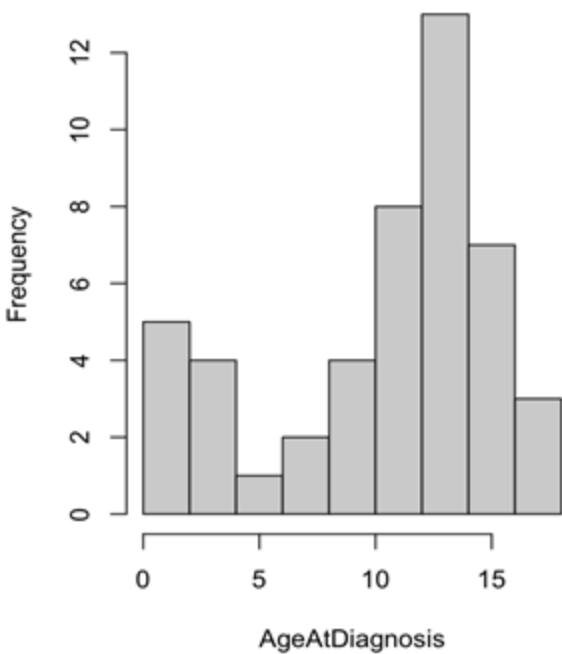
Query filtering in the Princess Máxima Center

Attribute	Scope	Query + Filtering	Result (count)	Query result
Cancer Cellularity Measurement	biosample	NCIT:C123556 > 89	59	✗
Age At Diagnosis	individual	NCIT:C156420 > P14Y	50	✗

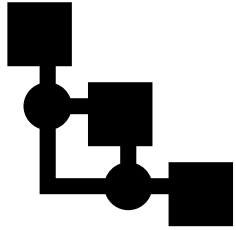
Histogram of PercentTumorCells



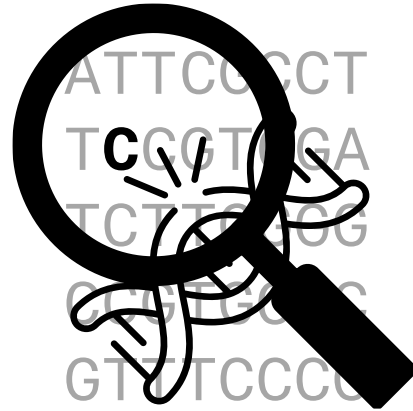
Histogram of AgeAtDiagnosis



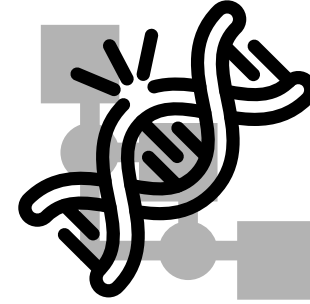
OncoBeacon adds (to) the oncological context



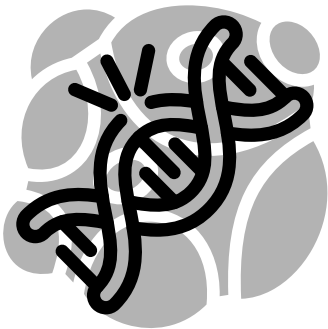
Further specification of existing attributes using ontology terms



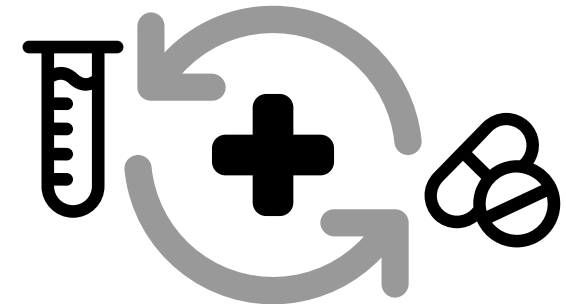
Contextual attributes that help researchers correctly interpret the genetic data



Standards for molecular subtyping



Attributes describing molecular drivers of tumor initiation and tumor genesis



Attributes placing samples in the context of the patient's treatment trajectory

Next steps

Community:

- Feedback to cancer and beacon working groups
- Cancer community to define minimal standards
- OMOP to Beacon mapper/integrator

We:

- Solve query issues
- Deploy OncoBeacons (international)
- Be transparent about the clinical to Beacon field mappings
- Test federated queries on the beacons

Acknowledgements

Patrick Kemmeren
Alex Staritsky
Marcel Santoso
Jasper van Dalum

PI, Princess Máxima Center for Pediatric Oncology
ETL development
Beacon platform deployment, testing, debugging
Cloud infrastructure development

Joep de Ligt
Stefan Dentre
Matthijs van Niekerc
Sophie Roerink

Lead Data, Hartwig medical Foundation
Beacon implementation
Cloud infrastructure
Clinical data mapping

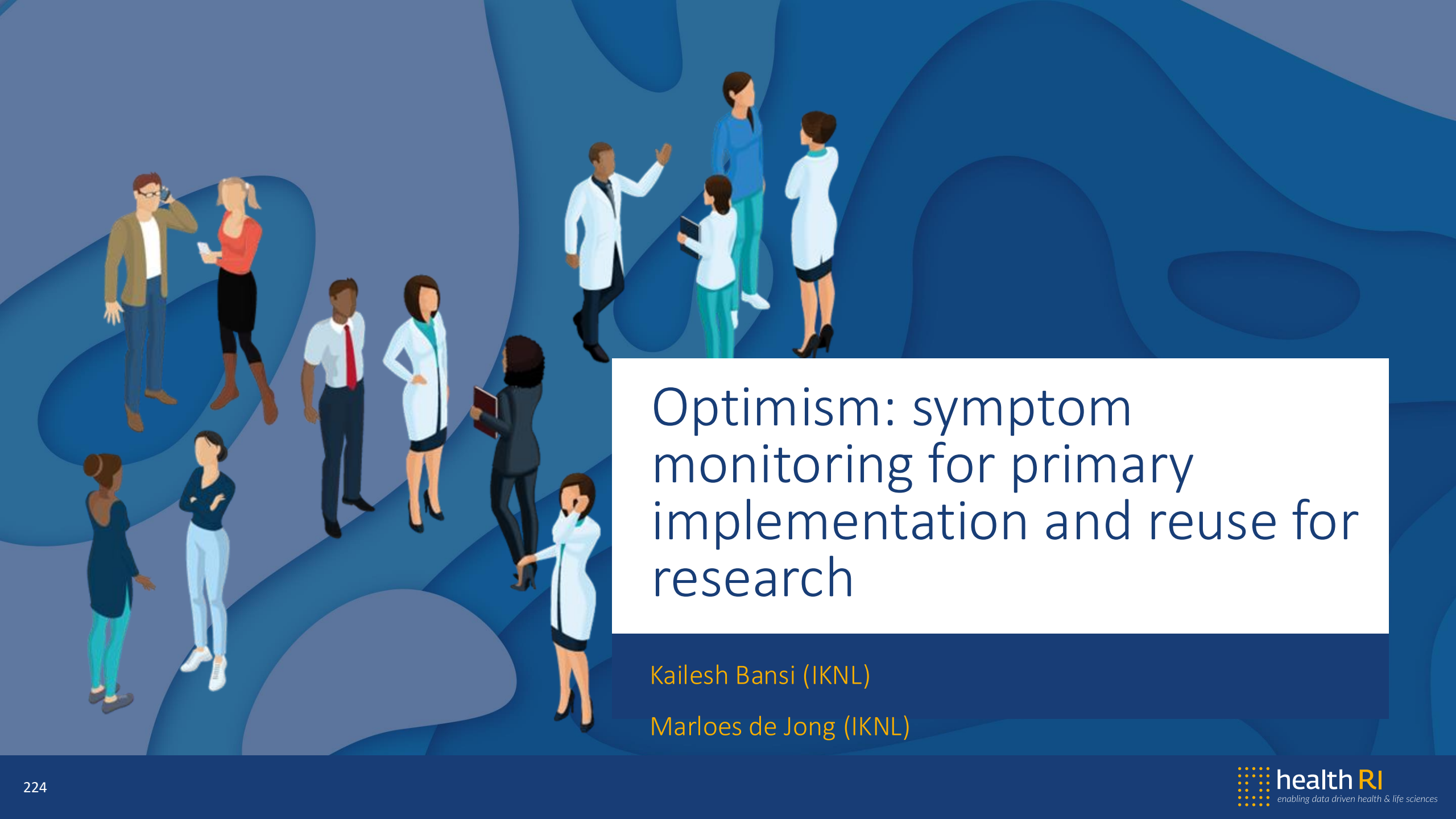
Jordi Rambla & Michael Baudis
Oriol López-Doriga

Beacon insights
Beacon v2 Production Implementation developer



Coffee break

Coffee break
14:30-14:45

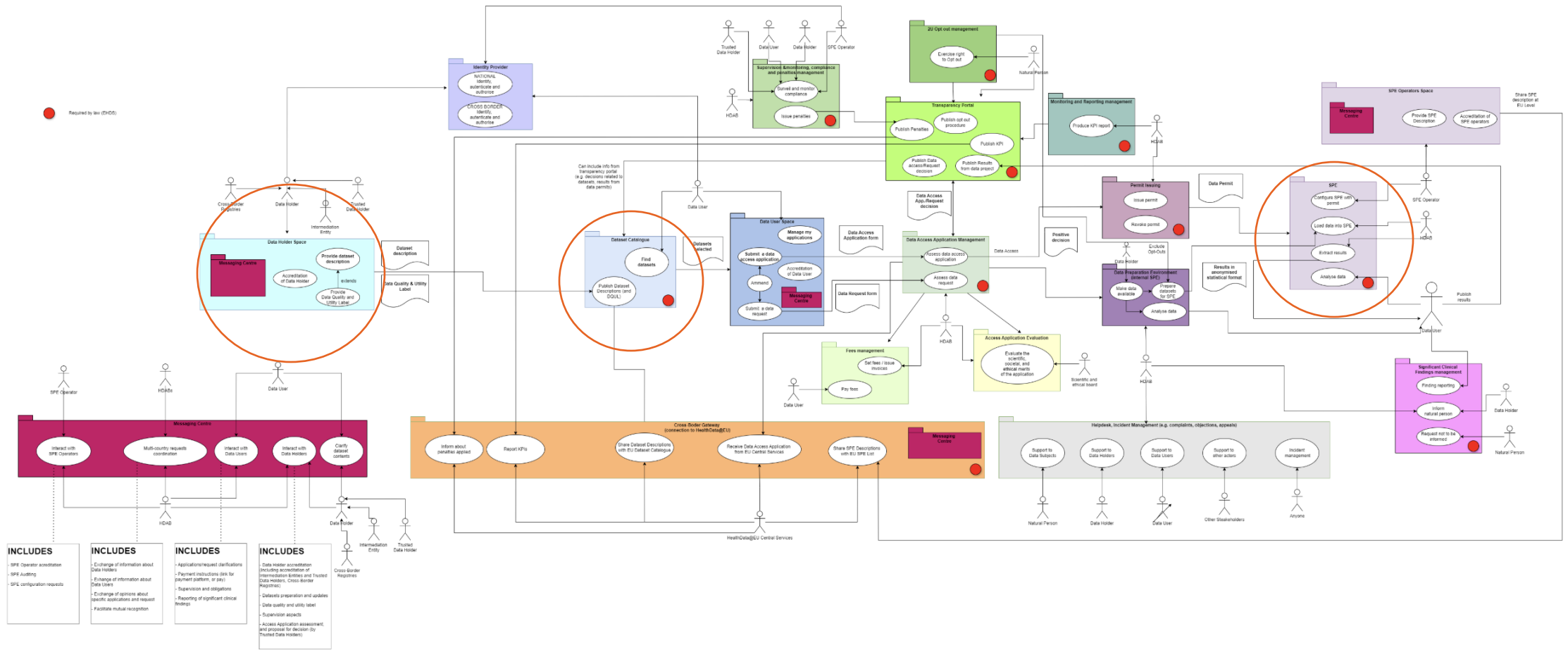


Optimism: symptom monitoring for primary implementation and reuse for research

Kailesh Bansi (IKNL)

Marloes de Jong (IKNL)

Backbone



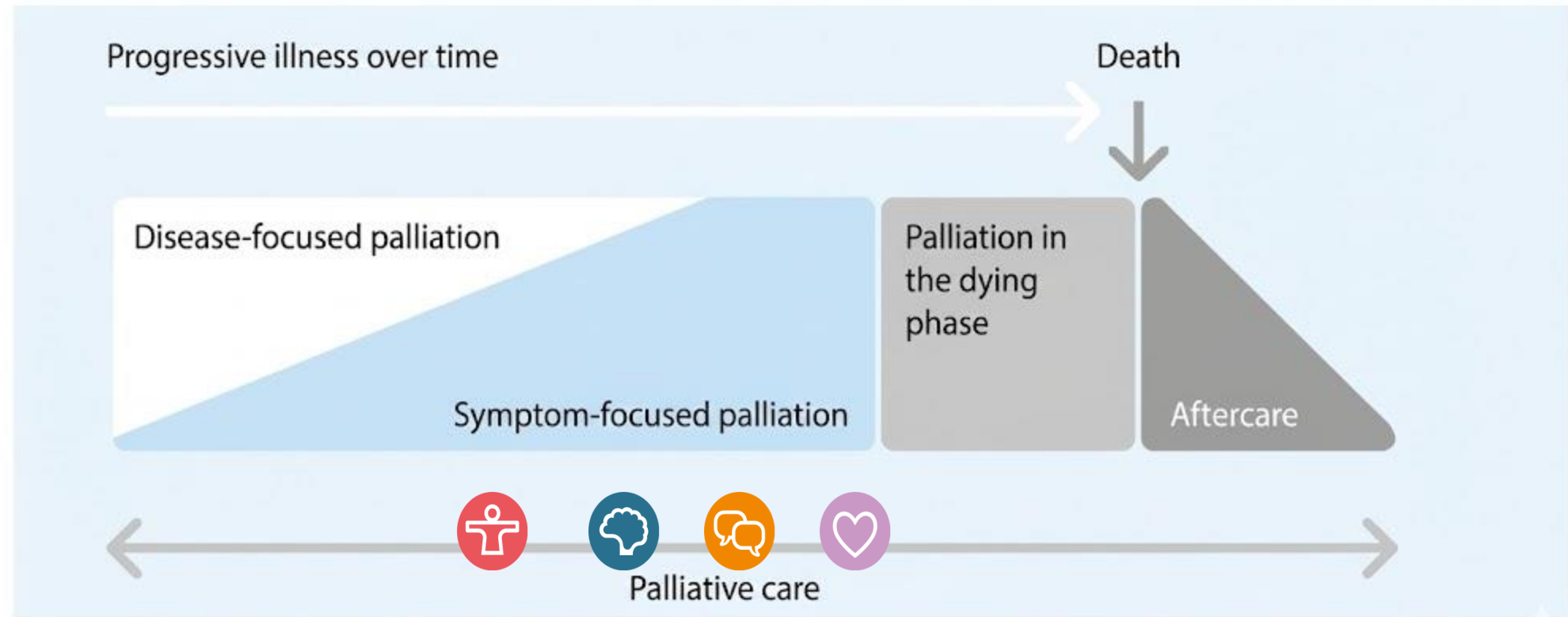
OPTIMISM →

**Development and implementation of
structural symptom monitoring for patients
with incurable cancer**

**Marloes de Jong, IKNL
Kailesh Bansi, IKNL**

HRI Oncology node, 8 december 2025

Phases in palliative care



model by Joanne Lynn and David Adamson

4 dimensions/pillars of palliative care



Physical

- *Pain*
- *Shortness of breath*
- *Fatigue*



Psychological

- *Anxiety*
- *Frustration*
- *Depressive feelings*



Social

- *Loneliness*
- *Tensions in relations*
- *Loss of work*



Spiritual

- *Questions about meaning*
- *Existential fears*



Aim **OPTIMISM**

OPTIMIzation of symptom management through implementation of structural
Symptom **M**onitoring

Establish a scalable **symptom monitoring system** to improve patient and
caregiver **quality of life**
and facilitate **research**

System's applicability extends
beyond oncology and palliative care

Academic hospitals / Dutch Centers of Expertise in Palliative Care & IKNL
Project duration: 60 months
Budget: € 3.957.790,-

Ontwikkeling en implementatie van structurele symptoom monitoring voor patiënten met ongeneeslijke kanker

start binnenkort

Onderzoekssamenvatting

In 2021 zijn in Nederland 45.863 mensen aan kanker overleden, wat het belang van palliatieve zorg voor een grote groep patiënten met kanker benadrukt. Patiënten met ongeneeslijke kanker ervaren vaak verschillende klachten zoals pijn, vermoeidheid, kortademigheid, obstipatie en angst. Deze klachten kunnen gelijktijdig voorkomen. Naarmate de ziekte vordert, nemen het aantal en de ernst van de klachten toe. Zelfs bij milde klachten kunnen deze een aanzienlijke invloed hebben op de kwaliteit van leven en dagelijkse activiteiten van patiënten. In de dagelijkse zorg is het een uitdaging om de aanwezige klachten van een patiënt goed in kaart te kunnen brengen. Daardoor is het bepalen van klachten vaak niet optimaal.

Voorgestelde oplossing

De implementatie van symptoommonitoring kan bijdragen aan het verminderen van klachten. Symptoommonitoring houdt in dat patiënten aangeven hoe ernstig zij hun klachten ervaren, zodat zorgverleners snel kunnen ingrijpen om de klachten te verminderen. Dit heeft diverse voordelen, zoals minder last van klachten, minder bezoeken aan de spoedeisende hulp en minder ziekenhuisopnames,

Projectgegevens

- / **Projectnummer:** 15985
- / **Projectleider:** prof. dr. An Reyners
- / **Projectteam:** dr. Jurrian van der Werf (IKNL)
- dr. Rudolf Fehrmann (University Medical Center Groningen) - dr. Everlien de Graaf (UMC Utrecht)
- / **Instituut:** UMC Groningen
- / **Startdatum:** 1 juli 2024
- / **Kankersoort:** Algemeen (geen specifieke kankersoort)
- / **Looptijd:** 60 maanden
- / **Projectbudget:** € 3.957.790,-

OPTIMISM: OPTIMization of symptom management through implementation of structural Symptom Monitoring



OPTIMISM – why?

Patients with advanced cancer have complex care needs

- On average, 6 simultaneously occurring symptoms
- As the disease progresses, the number of symptoms and severity increase
- Significant impact on quality of life and daily activities, even at low intensity
- Challenge to properly map all present complaints!

Most reported symptoms:



Pain



Fatigue



Dyspnea



Loss of appetite



Anxiety



Depressed mood

Why symptom monitoring in palliative care?



At end of life:
average of
6 concurrent symptoms
of which
4 clinically relevant

Most reported symptoms:



Pain



Fatigue



Dyspnea



Loss of appetite



Anxiety



Depressed
mood

If **not consistently assessed**, may fail to recognize or underestimate severity and impact, leading to **suboptimal care**.

+

Often assessed retrospectively, relying on **patient recall** rather than **real-time evaluation**

=

Challenge to **provide timely and effective interventions**

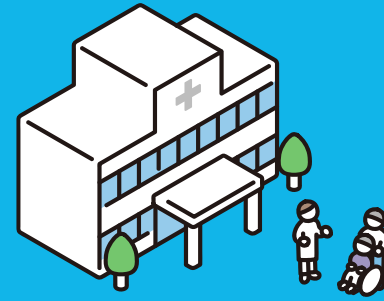
Possible benefits



Reduced symptom burden



Less anxiety



Fewer emergency room visits and hospital admissions



Improved quality of life

Successful example:

Ontario, Canada, with the Edmonton Symptom Assessment System (ESAS)

OPTIMization of quality of life by individualized Symptom Management: national implementation of structural symptom monitoring and realization of a research data-hub

OPTIMISM →

→ Systematic approach

Where patients actively and structurally report symptoms (USD-4D)

→ Infrastructure

For (inter)national research on symptom burden and quality of life

SONCOS normeringsrapport 2025

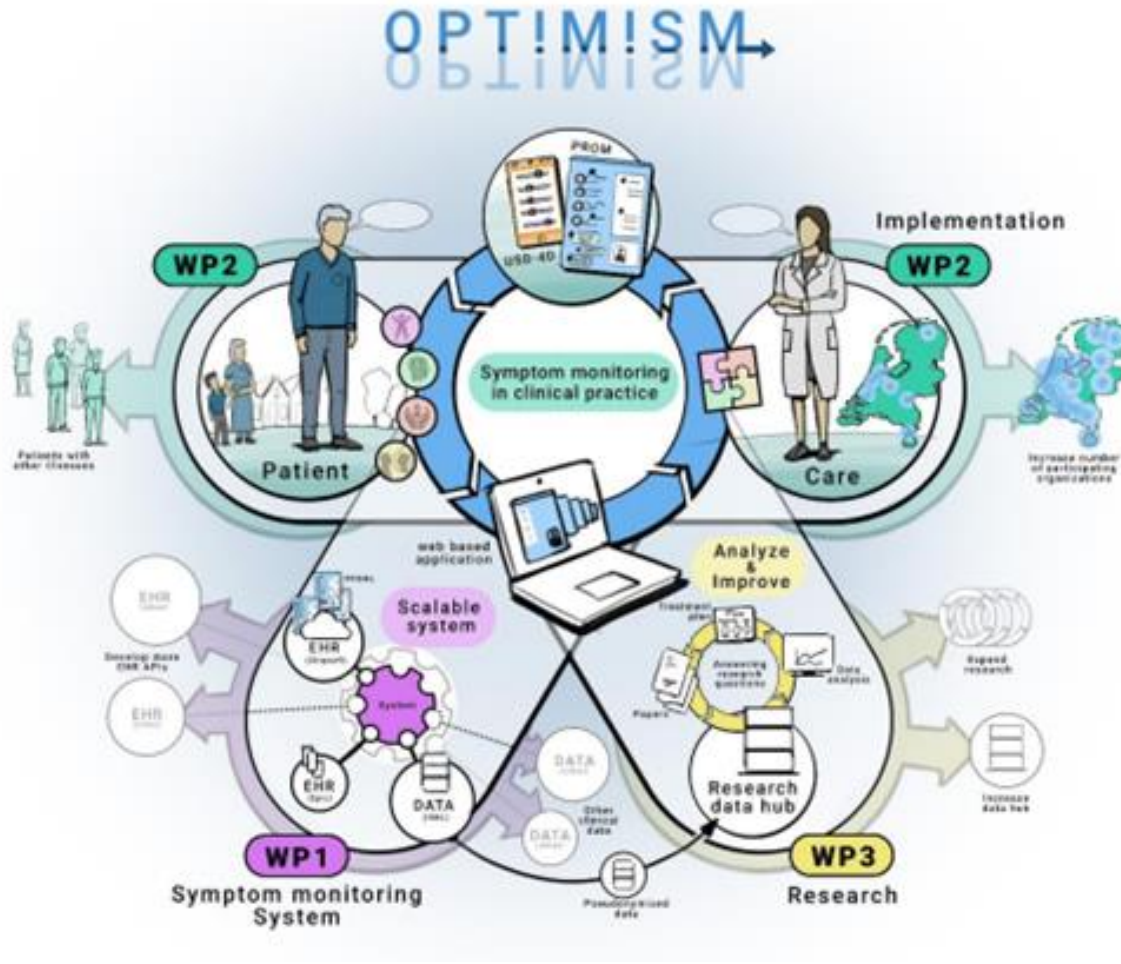
“Symptom monitoring is an integral part of palliative care, as part of proactive care planning.”

“Attention to symptom monitoring and symptom treatment on the four dimensions (physical, psychological, social, and spiritual) is in any case present for all patients with a life expectancy of <1 year.”

Preliminary inventorisatio**n** of hospitals



Plan of investigation: 3 work packages



WP1 Development and deployment

- Symptom monitoring system
- Goal architecture for scalability
- Data research hub

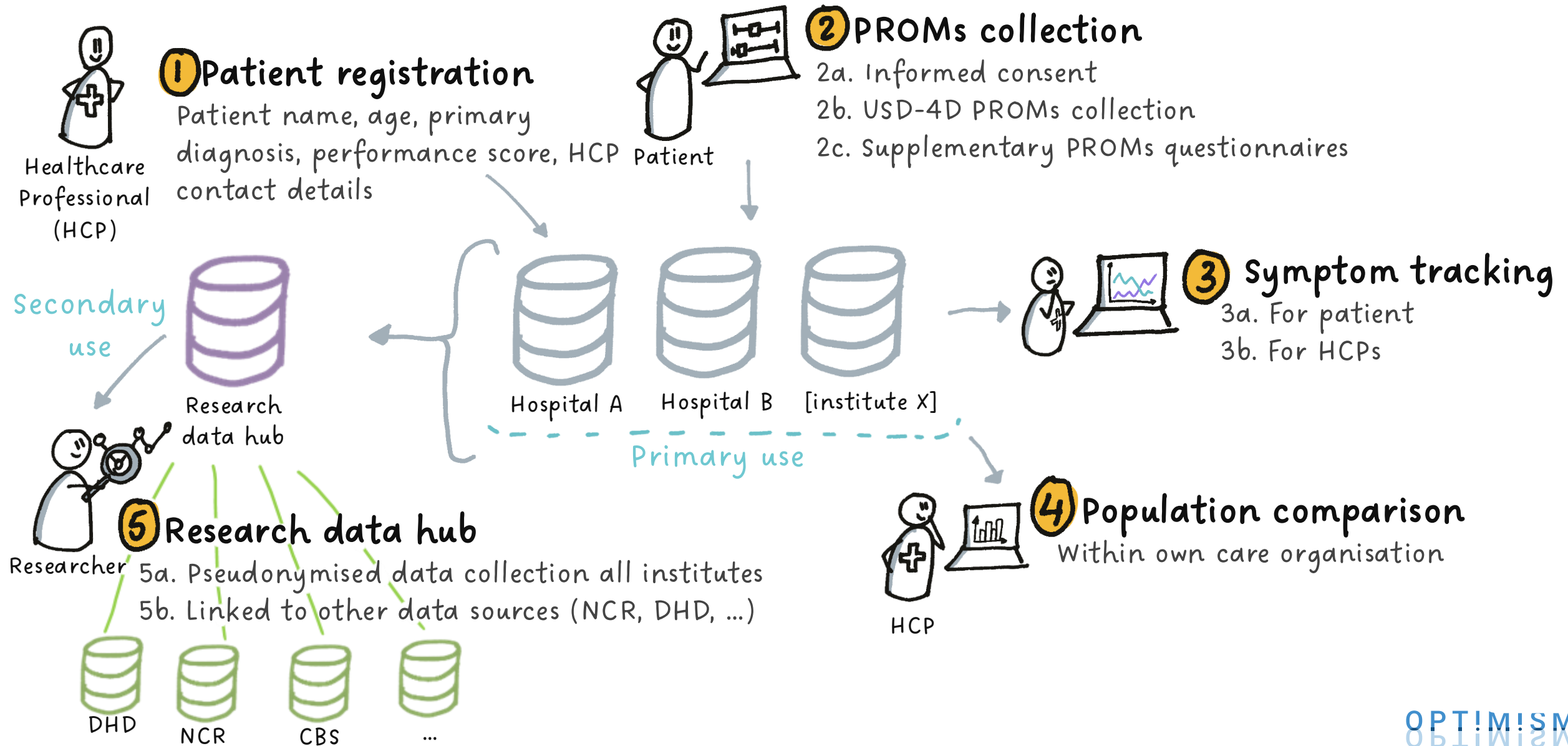
WP2 Implementation in daily clinical oncology practice

- Starts in academic hospitals
- Additional care settings

WP3 Research on

- Implementation strategies
- Obtained symptom data

Development of the symptom monitoring system



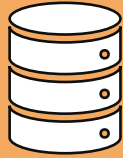
Lessons learned from pilot implementation

- Difficulty finding entry points (implementation and data export)
- Individual approaches within hospitals and departments:
 - Continuous improvement
 - Value-driven care
 - Appropriate care
- Varying data export:
 - FHIR
 - CSV
 - XML

Researcher's need



Data must be findable (currently lacking regarding symptoms)



Data must be interoperable (a lot of time spent on data harmonization)

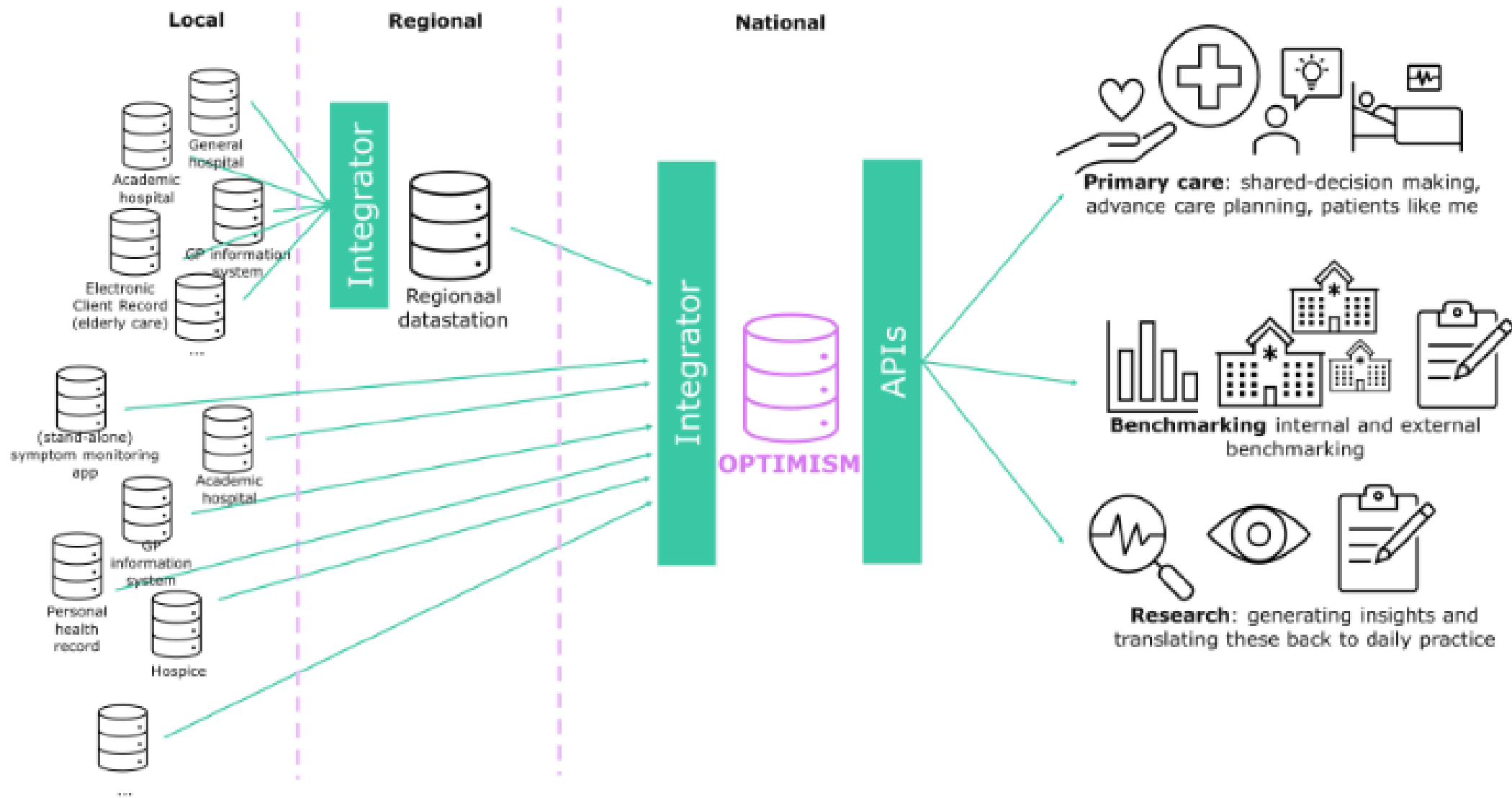


Data must be of quality (completeness of data)



Real-world data (currently often retrospective)

Technical components



Lessons learned from pilot implementation

- Difficulty finding entry points (implementation and data export)
- Individual approaches within hospitals and departments:
 - Continuous improvement
 - Value-driven care
 - Appropriate care
- Varying data export:
 - FHIR
 - CSV
 - XML

Optimism & Dutch Oncology Node

How would Optimism fit in an oncology node:

- Where do you see opportunities?
- How can we support datalinkage? For instance genomics
- For reuse?
- For a learning healthsystem?

OPTIMISM →

- Any questions?

- E-mailadres: OPTIMISM@UMCG.nl





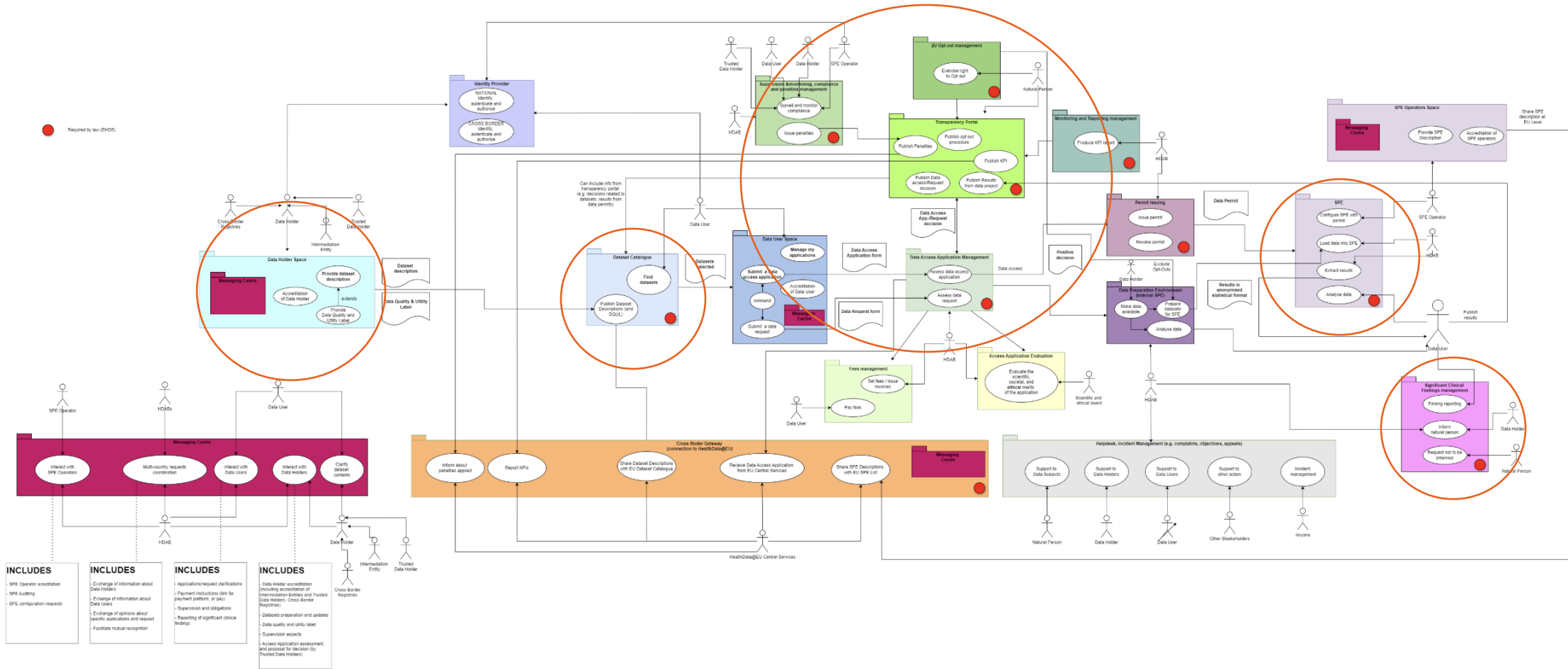
A mini workshop Nederlandse Kanker Agenda: Delayed diagnosis, and needs for data infrastructure

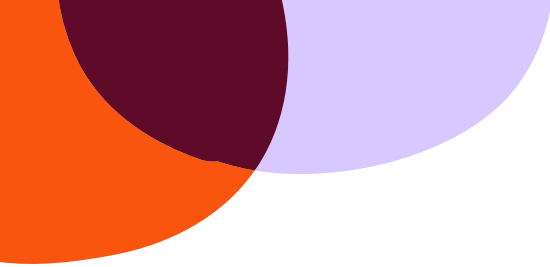
Femke Jacobs (IKNL)

Tilja van den Berg (NKC)

Warnyta Minnaard (NKC)

Backbone

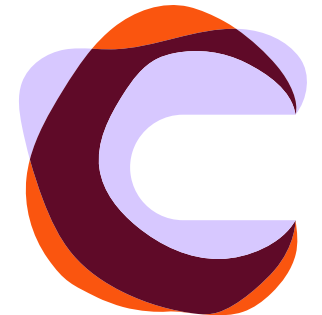




NEDERLANDSE KANKER AGENDA

ACCELERATION TEAM DIAGNOSTICS

Workshop on delayed diagnosis, and needs for
data infrastructure



**Nederlands
Kanker
Collectief**

One plan for the Netherlands

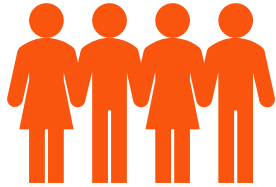


24
nov 2023

Netherlands Cancer Agenda (NKA)

- 20 ambitious & clear goals for 2032 which are apolitical, independent of resources/funding
- Direct and proactive approach with action plans
- Continuous monitoring and adjustment/alignment
- Endorsed by the government

Working together as the Dutch Cancer Collective



Mission – why we do what we do

Together, we reduce the impact of cancer on society.



How we work together

Connect – organisations, knowledge, disciplines, and life before, during, and after cancer

Strengthen – combining forces, embracing perspectives, creating integrated solutions

Accelerate – keeping a clear focus, removing obstacles together, move quickly



What we do

Work together on the 20 goals of the Agenda

Conceptual framework

Prevention

Life before cancer

Life with cancer

Life after cancer

1. Smoking behaviour

Patient
journey

8. Early detection

9. Diagnostics

12. Rare cancers

Quality of life

15. Late effects

17. Work and cancer

20 goals



Instruments

Organisation of care & support

Research, innovation & implementation

Data infrastructure

Education

Information & communication

Policy

How do we accelerate together?



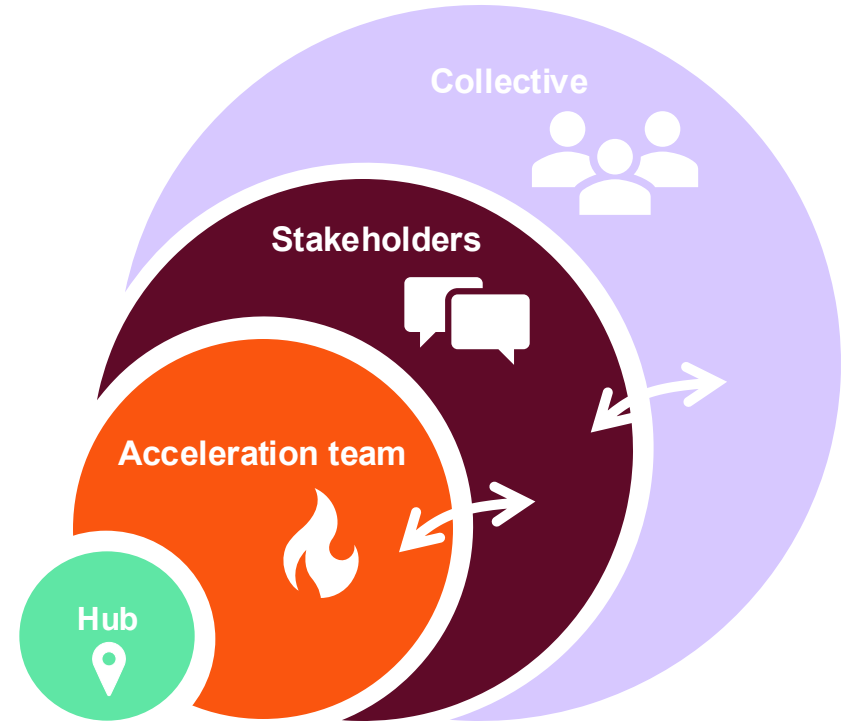
Acceleration teams

Who
Focus

Experts, leaders
Content - unravelling the problem together, arriving at concrete actions and implementing them

Role
Input

Action holder developing and implementing solutions
Expertise, time and resources to realise the goal



Who
Focus

How the hub complements

Supporting project manager
Process focus - brings structure, monitors and facilitates the end goal and the process: pace, effectiveness, etc
Facilitator, process supervisor and initiator
Way of working, neutrality, 'interspace'



Diagnostics

By 2032, people with cancer will have a personalised diagnosis (more) quickly with minimal burden

01.

What do we want to change?

By 2032, people with cancer will have a personalised diagnosis (more) quickly with minimal burden

We want to make the right diagnosis as soon as possible so that appropriate treatment can be started quickly. This takes into account the wishes, values and preferences of people with cancer. Currently, there are differences in outcomes and lead times of diagnostic processes between hospitals. Data on these are not available for all tumour types. The use of technology and innovations can make the diagnostic pathway more efficient. This potential is currently insufficiently exploited

The use of comprehensive molecular diagnostics has proven added value for certain patient groups, for example for Cancer of Unknown Primary (CUP) and lung cancer. This is not known for all other patient groups. There are differences in the offer of various forms of comprehensive molecular diagnostics across hospitals.

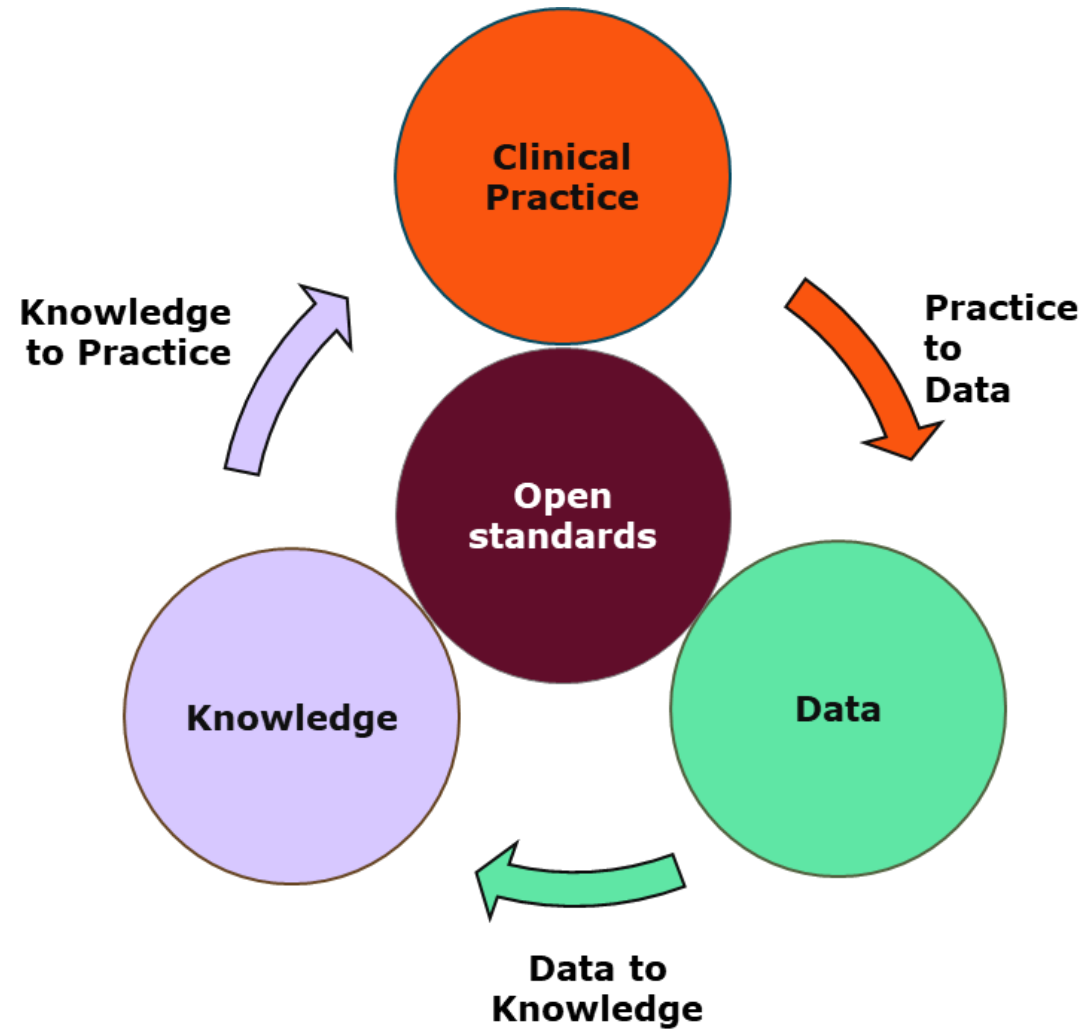
02.

Activities for 2026

Activities <i>What will we do in 2026</i>	<i>What this contributes to</i>
Activity 1: Diagnostic patient journey and milestones The diagnostic process can be experienced as a black box. In this patient journey, we identify phases and critical moments to best support the person suspected of having cancer.	Lack of information, shared decision-making and point of contact
Activity 2: Coordinated, integrated and streamlined diagnostics Analysis of the opportunities to better align the various components of diagnostics, so that they take place in a coordinated, integrated and streamlined manner.	Diagnosis takes a long time
Activity 3: Bringing together initiatives for a learning health system Analysis of initiatives and stakeholders involved in developing a learning health system to identify gaps and potential for acceleration.	Data and insights are not sufficiently utilized
Activity 4: Analysis of diagnostic turnaround times Desk research and (possibly) additional data analysis of turnaround times per tumor group, stage, and/or technique to gain insight into bottlenecks and opportunities for acceleration.	Diagnosis takes a long time
Activity 5: Analysis of practice variation in diagnostics Desk research and (possibly) additional data analysis of practice variation by tumor group, stage, and/or technique in deviation from guidelines and innovations to gain insight into the scope of the problem and to assess what is desirable and undesirable, and where opportunities for improvement lie.	Inequality and inefficiency in diagnostic pathways create differences in the quality and accessibility of care

ACTIVITIES FOR 2026

Activity 3: bringing together initiatives for a learning health system



Activity 3: bringing together initiatives for a learning health system

Goals



Define the content of a LHS

Define to which urgent/relevant questions a LHS should contribute



Identify existing initiatives

Identifying relevant Dutch initiatives (and affiliated parties) that are working on (components of) a learning health system. We are interested in initiatives that are collecting, analysing and/or standardizing data on cancer diagnostics, aiming to enable healthcare evaluation.



Support and connect existing initiatives

Use the strengths of the Dutch Cancer Collective to support, connect and improve existing initiatives

Discussion – the role of DCC

- What role do you think the Dutch Cancer Collective can play in bringing existing initiatives together?
- What support is still missing in the field?
- How can we truly help parties working on elements of a learning health system to move forward?

Discussion – learning health systems

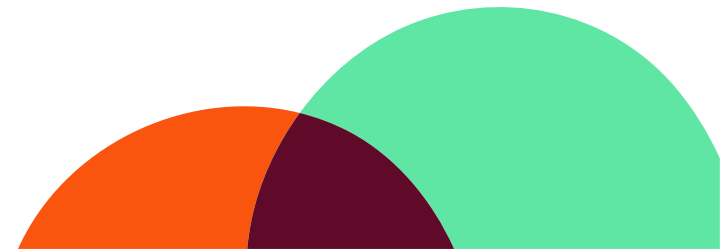
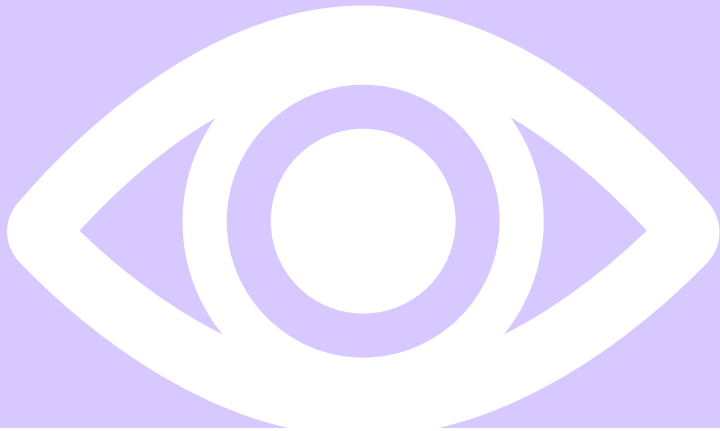
- What role do you think the Dutch Cancer Collective can play in bringing existing initiatives together?
- What support is still missing in the field?
- How can we truly help parties working on elements of a learning health system to move forward?
- What initiatives and/or stakeholders working on learning health systems in oncology should we be aware of?
 - Focusing on the Netherlands, and perhaps international initiatives that can be an inspiration for us
- What is their scope? What questions are the systems aiming to answer?

Would you be interested to brainstorm further with us to drive this activity forward?

Questions? Want to get involved?

Reach out to us via: warnyta@nederlandsankercollectief.nl &

tilja@nederlandsankercollectief.nl

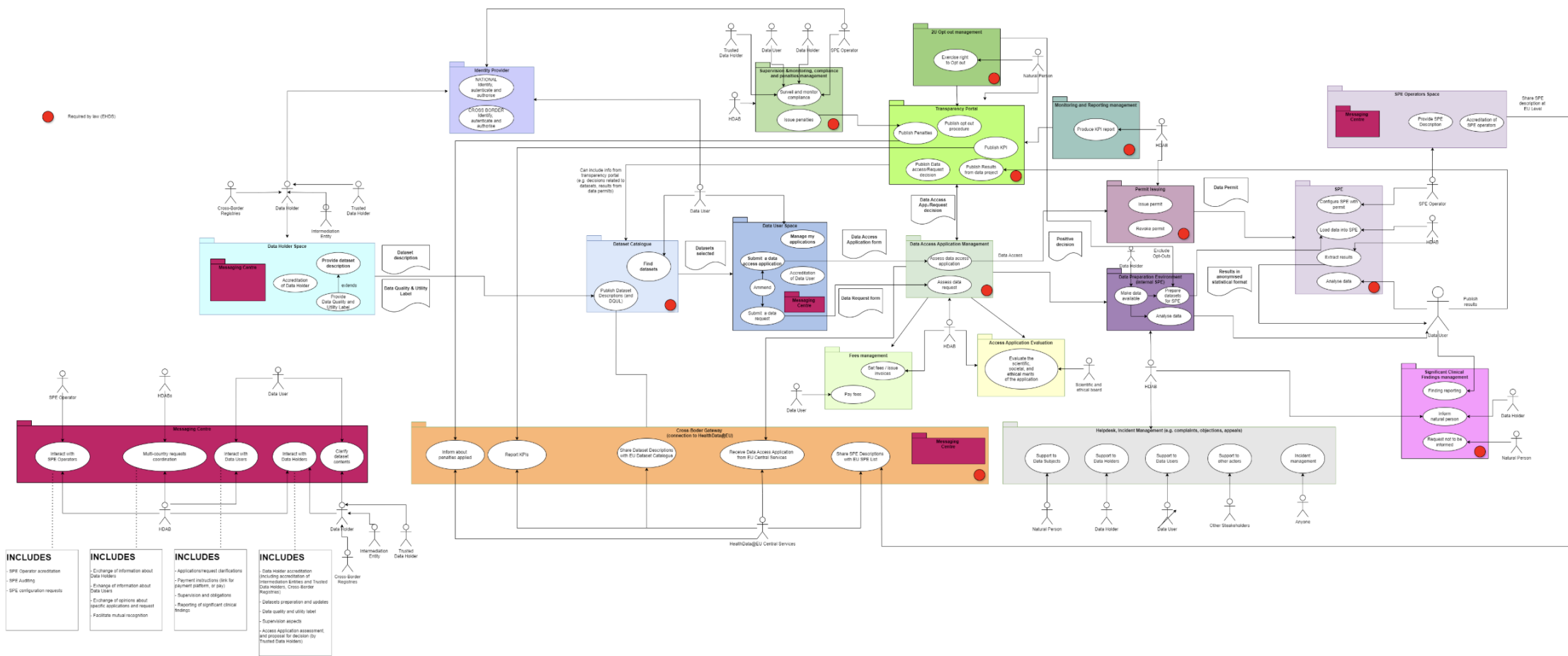




An (extended) panel discussion: Our NCDN-NL

ALL node members

Backbone





Carla van Laer (NKC)



Patrick Kemmeren(PMC)



Joep de Ligt (HMF)



Carla Pieterman (KWF)



Joppe Tra (HRI)



Anika Bonggarts (IKNL)



Eva Meize (IKNL)



Lifang Liu (HRI)

Q1: What is your vision and expectation about the Dutch National Cancer Data Node (NCDN-NL)?

Nr of comments	What is your vision and expectation about the Dutch National Cancer Data Node (NCDN-NL)?
1	To show what is possible by bringing the right data together
2	A nation-wide data findability infrastructure for oncology data. beyond the metadata catalogus, a more advanced and robust infrastructure with oncology-specific items that can be used for exploration journeys.
3	Stimulating datadriven learning healthcare system for oncology for the whole patient pathway so. incl. prevention, palliative care and berievement care
4	The NCDN should be THE place for experts and thought leaders from data holders and other stakeholders that take a tactical and strategic approach to improve data availability, data quality and data usage in order to create faster and better insights.
6	A 'living lab' on how we can operationalize the EHDS/HDAB ambitions, and a Trusted Health Data Holder for the oncology domain
7	Work as a coordinator between EHDS and data holders giving support to them on how model and transfer data, and make data available for others.
8	I envision the NCDN-NL as a national enabling infrastructure that makes it easier, faster, and safer to use cancer data for impact: improving patient outcomes, reducing inequalities, accelerating research, and informing policy. The node should not become "another platform," but rather a connector and simplifier across existing datasets, standards, and initiatives in alignment with European initiatives, projects & networks
9	To bring together the required infrastructure, tooling and harmonization of data for oncology research and care and act as a linking pin for international harmonization and sharing of data.

Q2: What are the priorities/priority projects in your organization in the coming year(s)? Are there any challenges and opportunities related to data? if so, what are they?

Nr of comments	What are the priorities/priority projects in your organization in the coming year(s)? Are there any challenges and opportunities related to data? if so, what are they?
1	Making the learning care system a reality, this is mostly a data standardisation and sharing problem (for now)
2	See our digital strategy, https://www.prinsesmaximacentrum.nl/storage/configurations/prinsesmaximacentrumnl/files/digitale_strategie/idt_strategie_nl.pdf And our main multi-year strategy, for example paragraph 2.3 (page 29) on data
3	Datadriven evaluation and monitoring of package decisions And broader data availability for oncology
4	For IKNL and DICA: create a well integrated portfolio of data and information products. Increase the data availability and data quality. Biggest challenges: 1. Data availability from hospitals (by far the biggest challenge) 2. Data linkage 3. Data standardization and addressing the semantic layer.
6	Help out with (technical) innovations that support sovereign, automated, verifiable and privacy preserving data analysis for secondary use. Related questions: How to make data FAIR? How to verify your computation? How to preserve privacy? How to come to an equal level playing field, with a healthy market around it (incl. public/private partners)
7	Centralise and harmonise data to be able to share data for secondary use.
8	Implementing the (action plans of) Netherlands Cancer Agenda through coordinated action across care, research, data, and lived experience.
9	Standardize and harmonize clinical, phenotypical and genomic data; develop federated query possibilities; cloud-enabled secure processing environments.

Q3: What do you see that NCDN-NL can help you?

Nr of comments ▾

What do you see the NCDN-NL can help you? ▾

- 1 Help organisations realise this is possible and needed now
- 2 National alignment on standards. Making sure everyone uses the same ontologies.
- 3 Information modelling to come to shared models to facilitate interoperability
Stimulate datastatoms which make cancerdata available both at healthcare providers as well as (research)institutions
In co-creation :-)
- 4 Create a business case and urgency to improve data availability and the application of information standards in hospitals.
- 5 Exchange of ideas, jointly work on projects
- 6 Application area to test innovations
- 7 I hope NCDN-NL can help on model clinical/research data to make it FAIR and manage the data request via EHDS
- 8 The node can help by offering a unified entry point to discover and access data, shared standards and reusable agreements, and technical support for federated queries and secure analytics. It lowers operational barriers through interoperable workflows, provides structured pathways to move NKC use cases into practice, ensures trusted federated governance, and aligns the Netherlands with European data infrastructures.

Q4: How do you like to work with the node to solve challenges and maximize opportunities?

Nr of comments ▾

How do you like to work with the node to solve challenges and maximize opportunities? ▾

1 getting things done

2 We can contribute with expertise and experience

3 Present work, collect feedback and see how we can co-create together

4 Both in general meetings and in separate projects. Focus less on knowledge sharing and focus more on results and strategy.

6 Discuss pains, brainstorm for solutions, openness to work together on pilots

7 Deployment of a national cBioPortal instance connected to all data holders and help them to model the data in cBioPortal format. Make cBioPortal accessible via SPE environments. Provide support on modelling data to be able to use it for secondary purposes.

8 I would like to work with the node in a co-creative, iterative, and transparent way—focused on real use cases, clear governance, and early involvement of professionals and patients. Collaboration should connect to existing national initiatives, avoid duplication, and start with high-value, feasible projects that build trust. Preferred collaboration includes joint exploration of use cases (e.g. from the action plan diagnostic), short learning cycles that integrate technical and organisational alignment

Action list

(see more on the meeting agenda)

Take home message

- NCDN-NL is a community for all of us. Come, share and collaborate.
- NCDN-NL brings the Dutch consensus on standards, tools for cancer data.
- Maximize opportunities EHDS offers
- Maximize benefits from national and international expertise (e.g. NCDN-network)
- Maximize EU funding opportunities (CANDLE, EUnetCCC, JA-PCM as focal points)

