

FAIR data curation applied to HZB infrastructure

NeXus files, Icat server, python routines for data ingestion

- Overview of data staging workflow at HZB
- NeXus as a community standard for metadata schema and file format
- Icat services
- Use case EMIL
- Development directions and discussion

Consultancy:

- Development in collaboration with researchers of [ad hoc solutions](#) for the data curation
- Curation on data [access](#) and [publication](#) services
- Awareness on [Data staging workflows](#) and data – metadata community [standards](#)

HZB

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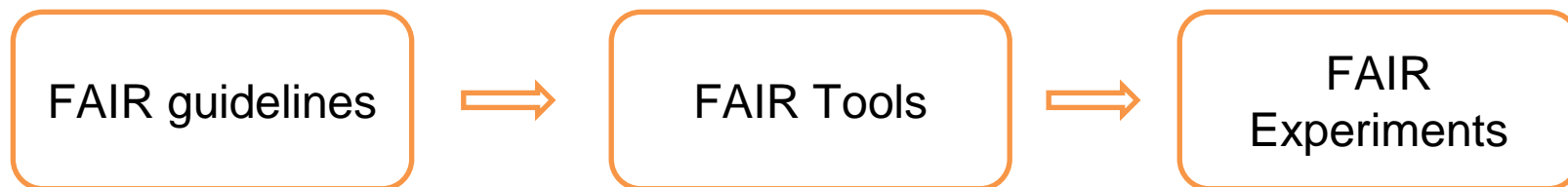
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ICAT

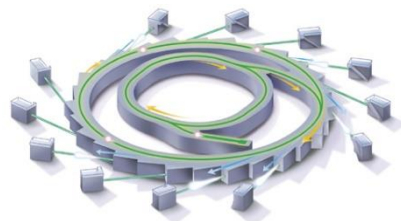
NeXus

MariaDB

minimal automatic metadata
generated from proposal and
measurement
settings



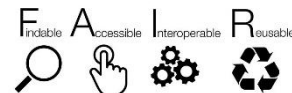
Data ingestion, PID
generation and
provision of services
for the data
discovery



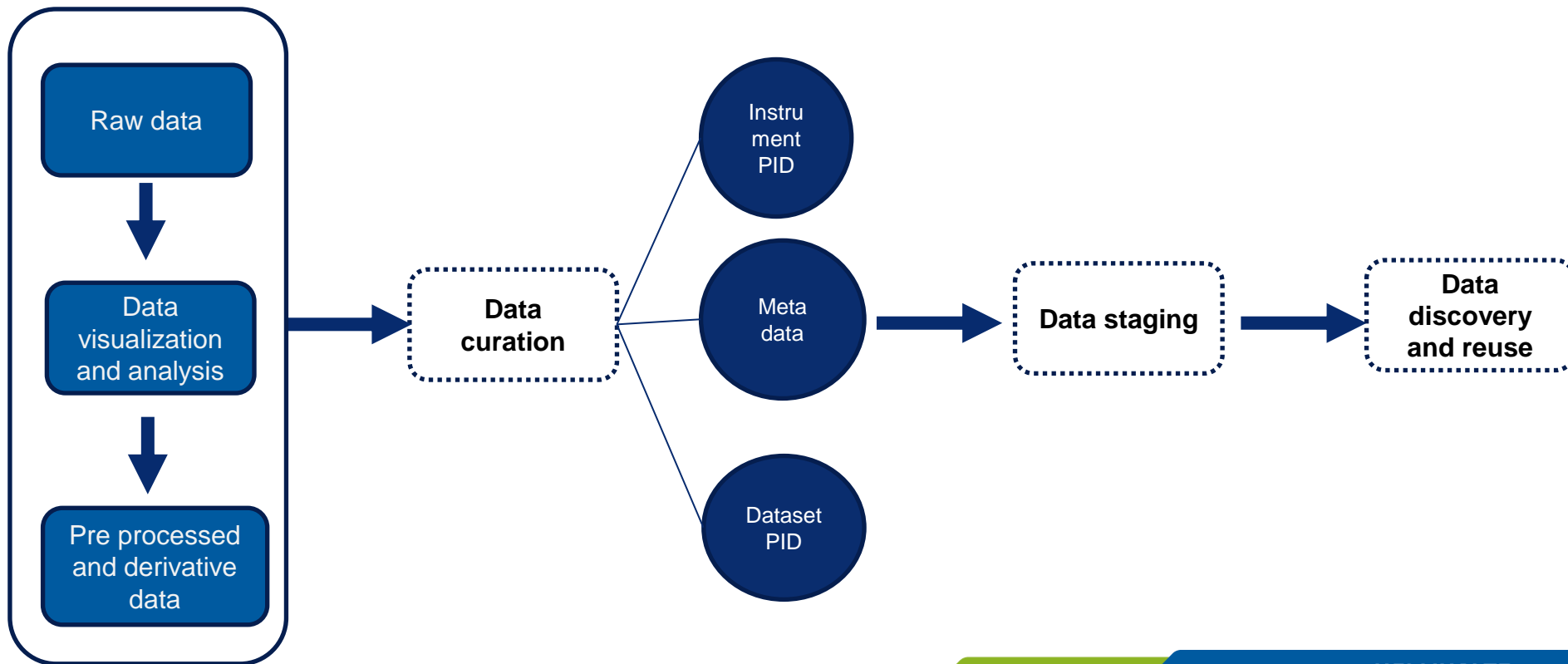
Proposal
submission
and beam time
reservation



Which services are running to implement FAIR data
access?



From data acquisition to data repository



At the measurement station

- **Automatize the metadata capture** with fully integrated tools in the measurement station system
- Identify among the machine metadata **relevant metadata** and **map** to measurement description- metadata schema
- **Optimize** the metadata harvesting **avoiding interference** with data acquisition/measurements

This tools integration is the focus of the collaboration Hub Matter, FDM-HZB and your group.

Before the experiment

Proposal

- Proposal number ID
- Project description and meas. background
- Method
- Instrumentation
- Sample
- Scientific team
- Duration
- Instrument metadata

Facility

Beam line , Detectors

During the experiment

Facility logs

- Beam parameters
- Motor position
- Instrument configuration
- Detector settings
- Sample settings

Users log book

Parameter settings
changes respect proposal
sample description
Run logs

After the experiment

Analysis & derivative data

- Description of pre-process.
- Extraction procedure description
- External link to codes
- Software
- Workflow
- plots

Publication

Subset or full
quality parameter
reference to journal publ.
DOI

Before the experiment

GATE

- Project description,
- authors, People metadata
- sample description,
- time frame, embargo

During the experiment

Elog notebook

- Measurements parameter, screenshot, free text notes
- Instruments logs
- Control system logs

After the experiment

Nexus-python routines, metadata server icat

- Description of pre-processing
- Instrument description
- External link to codes and software
- Licence
- Ancillary information

Recommendations for key elements of metadata by ExPands working group on FAIR data in the PaN community:

- PI and Coll
- Requested instrument
- Sample description
- Facility
- Proposal ID
- Experiment description
- Experimental team /Experiment time
- Data format /raw data or derivative
- Acquisition and pre processing software
- File ID
- Creator
- Publisher
- Release date and use Licence
- Preservation description
- Representation information
- Instrument parameters
- Analysis software
- File generator system
- Lab Notebook should be attached
- Calibration information
- Data relationship
- File identifiers
- PIDs
- Workflow

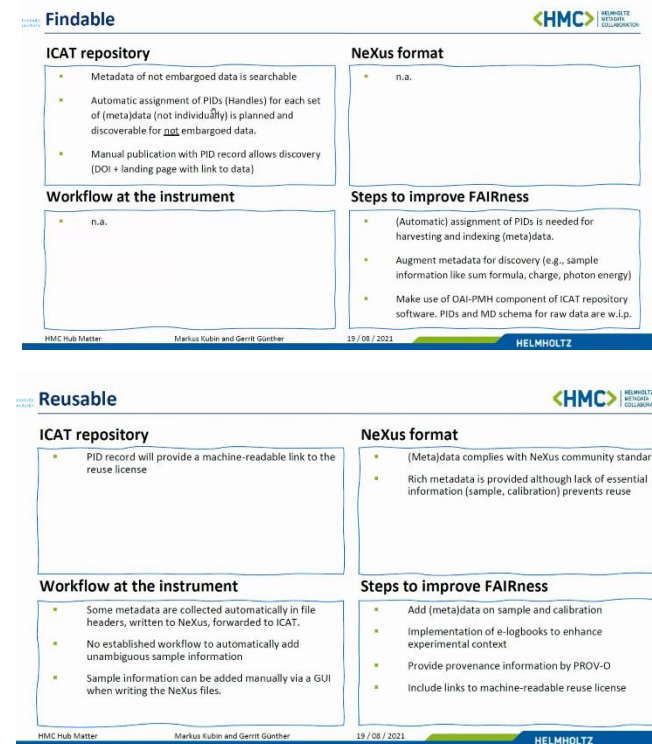
Expands.eu

Tools: elog book+ Nexus writer + ICAT

- **Linking** publication and datasets
- Nexus for rich metadata (from proposal and from measurements setting)
- **Providing unique identification** for data sets
- **Restricted access** to the research data under HZB data policy
- **Access to data and metadata** via searchable online catalogue (ICAT)

Services maintenance and tools development
FDM+Hub Matter

Current implementation of FAIR data



The image shows two slides from a presentation. The top slide is titled 'Findable' and the bottom slide is titled 'Reusable'. Both slides are part of a presentation by HMC Hub Matter, Markus Rubin and Gerrit Günther, dated 19 / 08 / 2023. The slides are organized into four quadrants: ICAT repository, NeXus format, Workflow at the instrument, and Steps to improve FAIRness. The 'Findable' slide shows the current state, while the 'Reusable' slide shows the future state.

Findable	Reusable
ICAT repository <ul style="list-style-type: none">Metadata of not embargoed data is searchableAutomatic assignment of PIDs (Handles) for each set of (meta)data (not individually) is planned and discoverable for <u>not</u> embargoed data.Manual publication with PID record allows discovery (DOI + landing page with link to data)	ICAT repository <ul style="list-style-type: none">PID record will provide a machine-readable link to the reuse license
NeXus format <ul style="list-style-type: none">n.a.	NeXus format <ul style="list-style-type: none">(Meta)data complies with NeXus community standardRich metadata is provided although lack of essential information (sample, calibration) prevents reuse
Workflow at the instrument <ul style="list-style-type: none">n.a.	Workflow at the instrument <ul style="list-style-type: none">Some metadata are collected automatically in file headers, written to NeXus, forwarded to ICAT.No established workflow to automatically add unambiguous sample informationSample information can be added manually via a GUI when writing the NeXus files.
Steps to improve FAIRness <ul style="list-style-type: none">(Automatic) assignment of PIDs is needed for harvesting and indexing (meta)data.Augment metadata for discovery (e.g., sample information like sum formula, charge, photon energy)Make use of OAI-PMH component of ICAT repository software. PIDs and MD schema for raw data are w.i.p.	Steps to improve FAIRness <ul style="list-style-type: none">Add (meta)data on sample and calibrationImplementation of e-logbooks to enhance experimental contextProvide provenance information by PROV-OInclude links to machine-readable reuse license

- Similar to dataset DOI or ORCID but identifying instruments
- PaN facilities have a complex of measuring stations
- Internal manufactured stations and not standard instrument
- Versioning to track the evolution of the measuring unit
- Interlinking PIDs at each stage in the process to provide traceability
- **Further questions ?**

Rolf Krahl, HZB is the reference person for this development

[6]

SISSY I @EMIL PIDs

Instrument DOI:

- <https://doi.org/10.5442/ni000018>

Instrument landing page

- https://www.helmholtz-berlin.de/pubbin/igama_output?modus=einzel&sprache=en&gid=1978

Publication details

- <https://commons.datacite.org/doi.org/10.5442/ni000018>

PID interlinking improves / facilitate the data discovery

- Logs of all the operations (sample preparation) on the sample embed the sample history.
- This log file is preserved and attached to the metadata

PID interlinking improves /
facilitate the data discovery

- Potential assignement of a PID ?

Considerations : sample charactr are changing while treated, measurements and sample creation can be really distinguished? What is the sample preservation after the measurements ?

NeXus Format

- Nexus for rich and easy to access metadata.

Why NeXus ? <https://www.nexusformat.org/>

- HDF5/NeXus used as institutional standard at neutron, x-ray and muon facilities
- Each facility diversify the dictionary limiting the immediate re-usability.
- NeXus files may help to improve the situation.
- HDF5 format and a tree structure for metadata representative of the complexity of PaN data
- Built in Vocabulary for research community interoperability
- Geometry of the beamline, sample stages, orientation and description of detectors, exposure time, beamline calibration info, scan description
- To store multiple related data set create more entries

Hierarchy in Nexus

Classes (dictionary)

Groups

Levels

Attribute

MultiD array and scalars

NeXus Implementation@ESRF

NXroot

Top level. One per file.

NXentry

One group per measurement

NXinstrument

Describe the instrument.

Only one per NXentry

measurement (@NXcollection)

Flattened view of everything measured

Only one per NXentry

sample (@NXsample)

Define the physical state of the sample during the scan

NXdata

The data to be plotted.

One NXdata group per plot

user (@NXuser)

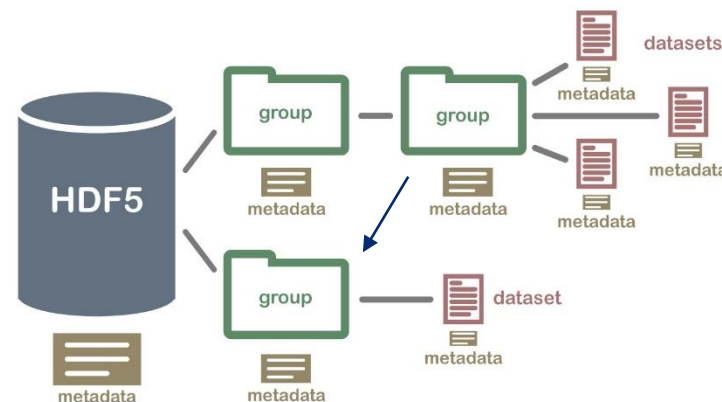
Details of a user, i.e., name, affiliation, email address, etc

NXsubentry

Data or links to data for particular analysis

NeXus structure allows links and pointing to data stored in other parts of the group

- HDF5 format and a tree structure for metadata representative of the complexity of PaN data
- Allows chunked storage and slices reading
- Metadata can be attached
- The I/O can be faster than contiguous data files
- Compression
- Can be prefixed or open database size
- Heterogeneous database with links
- Platform independent
- Suitable for massive databases with a datatype and dataspace definition per dataset



HDF5 structure allows links and pointing to data stored in other parts of the group

- New applications can be defined
- A rich set of tools for verification and file validation (e.g. nxvalidate) and easy data extraction
- Process data must contain Nxprocess group
- Nxentry is mandatory root element

List of attributes

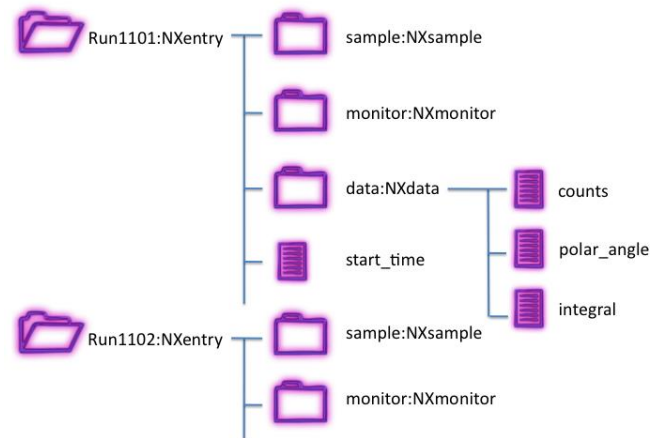
- <http://definition.nexusformat.org/nxdl/3.1/>
- <https://github.com/nexusformat/definitions/blob/main/nxdlTypes.xsd>

- Classes definitions in xml– mapping

Schema of the Nexus classes and attributes

- <http://definition.nexusformat.org/nxdl/nxdl.xsd>
- NXDL files must adhere to the specifications of the NeXus XML Schema, as defined in *nxdl.xsd* and *nxdlTypes.xsd*.

```
xmllint --noout --schema nxdl.xsd base_classes/NXentry.nxdl.xml  
base_classes/NXentry.nxdl.xml validates
```



- Expands (expands.eu), Panosc (panosc.eu) , CERIC-ERIC EOSC work on tools and standards for the interoperability of the PaN community data and connection to the European Open Science Cloud

- <https://manual.nexusformat.org/utilities.html#data-analysis>
- A number of Python routines to process X-ray photons emission data in hdf5
- XRF spectroscopy PyMCA
- <https://gitlab.elettra.eu/panosc/xrffitvis/>
- Xrayutilities for conversions spec hdf5
- IGOR pro can upload HDF5 www.wavemetric.com
- ORIGIN lab (+HDF5Browser App)
- DAWN
- Matlab
- Spec2hdf5 available tools (silx.org)
- Spec2nexus (<https://spec2nexus.readthedocs.io>)
- PyMCA (<http://pymca.sourceforge.net/>)
- NeXpy (<http://nexpy.github.io/nexpy/>)

NeXus structure and playground

- Ingestion workflow
 - Access to the tools by virtual machine
-
1. data collection
 2. identification of the instrument dictionary
 3. sample data info collection
 4. parameters attribution
 5. local data saving
 6. icat repository ingestion

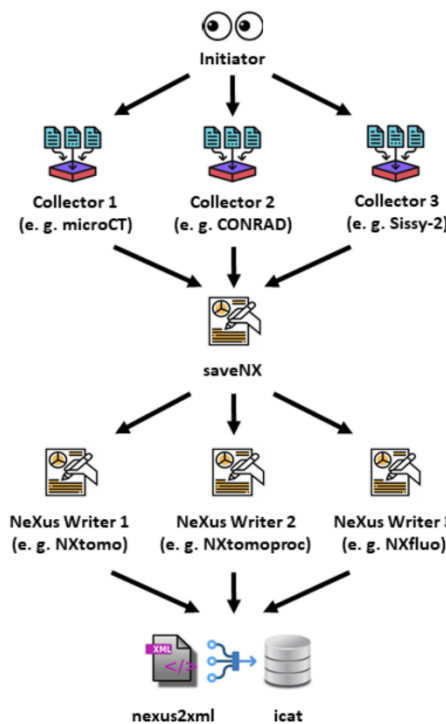
Contact person: Gerrit Günther, HMC

<https://gitlab.helmholtz-berlin.de/jaf/nexuswriter/-/blob/master/nexusCore/icatRepo/nexus2xml.py>

Nexus application for describing XAS

<https://github.com/nexusformat/definitions/blob/main/applications/NXxas.nxdl.xml>

■ Sketch for the ingestion



Initiator: The entry point that starts a collector. There are various initiators ranging from command-line interface to GUI.

Collector: An experiment specific module that collects data from different sources and assigns values to a python dictionary {...}

saveNX: A distributor that starts the appropriate NeXus Writer according to the NeXus definition schema of an entry; it may start different writer routines for a file.

NeXus Writer: A module that reads the python dictionary {...} and writes its content to the NeXus file according to a specific NeXus schema.

nexus2xml: An interface to HZB's icat to satisfy its demands: structures the produced files in folders, reads their content and writes a summary of searchable terms to a xml file before starting the ingestion.

Test the tools: NeXus writer and icat ingestion

- Notebook to optimize the data reuse with annotations, protocols
- The data in the notebook are under embargo
- Software esrf notebook
- **Web interface**

<https://icat.helmholtz-berlin.de/datahub/>

Time stamped in chronological order
can be used by users or software

- Contact person: Rolf Krah

The screenshot shows a web browser at the URL <https://icat.helmholtz-berlin.de/datahub/investigation/7482/events>. The page has a dark header with 'Data Portal' and 'My Data' links, and a 'Log out Luigia Cristiano' button. Below the header, there's a breadcrumb trail: '- / gate1:1910002-XX-1.1-P Test proposal to try out ingestion workflows with LISE/M.'. A 'Logbook' section is visible with buttons for '+ New', 'Take a photo', 'View', and 'PDF'. There are search filters for 'Everywhere' and a search bar showing '9 found'. The logbook entries are grouped by date:

- November 1st 2021**
 - 12:10:33 test2
- July 8th 2021**
 - 11:46:59

this		having	some		
is		columns	and	rows	
a					
table					
- July 2nd 2021**

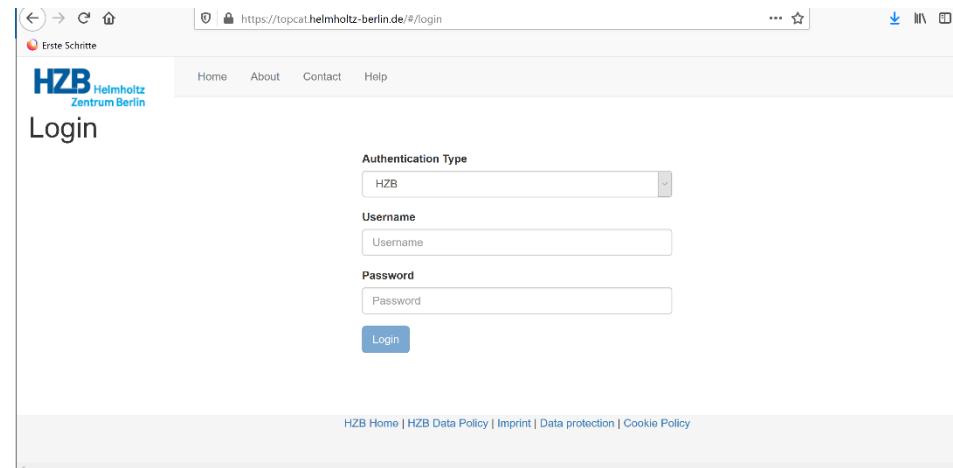
- Umbrella ID ; <https://www.umbrellaid.org/>
- Keycloak log in services are in implementation <https://www.keycloak.org/>

- Provide access to the data and is a DBMS developed at PaN facilities icatproject.org
- The data request means data extracted from tape and allocated on disk
- The software uses a **schema** or metadata, a web interface to the database
- **Dataservice** for data upload and download
- **Web interface** to search the data (topcat)

<https://topcat.helmholtz-berlin.de/#/login>

- Further questions ?

Rolf Krah



The screenshot shows a web browser window with the URL <https://topcat.helmholtz-berlin.de/#/login>. The page features the HZB Helmholtz Zentrum Berlin logo and a navigation bar with links for Home, About, Contact, and Help. The main heading is "Login". Below this, there is a form with the following fields: "Authentication Type" (a dropdown menu currently showing "HZB"), "Username" (a text input field), and "Password" (a text input field). A blue "Login" button is positioned below the password field. At the bottom of the page, there is a footer with links: "HZB Home | HZB Data Policy | Imprint | Data protection | Cookie Policy".

In development ! Will is giving you technical details on this collaborative work

Read out routine of the **sample database** using the time of the measurement to extract the corresponding sample history.

A **manual** integration vs **automatic** readout is possible

Notes, comments and images can be attached,

Sample information from proposal is also possible

Information from elogbook can also be attached

Python routines to ingest the data do not interfere with the measurements workflow

Reference to Guenther et al. ICALEPCS 21

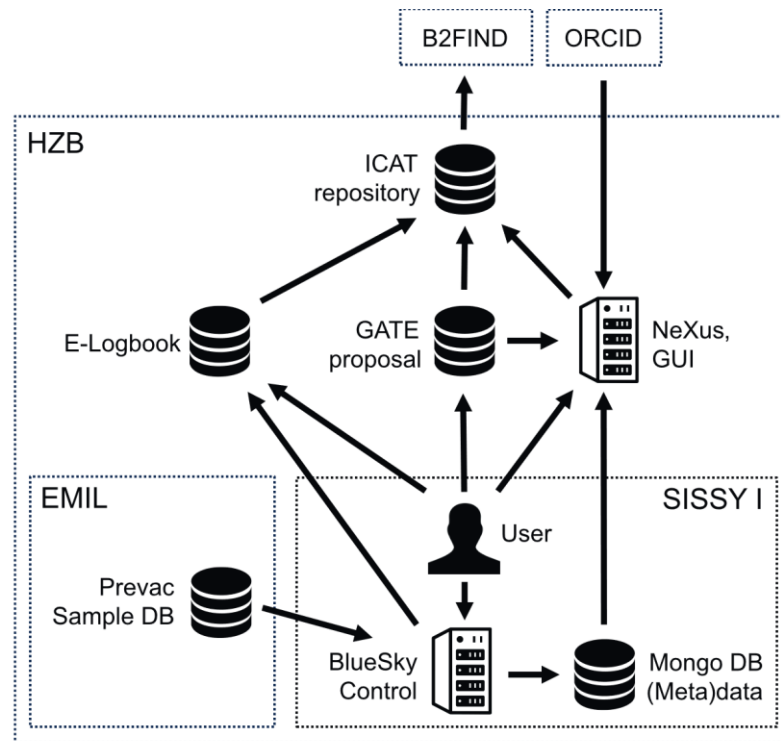
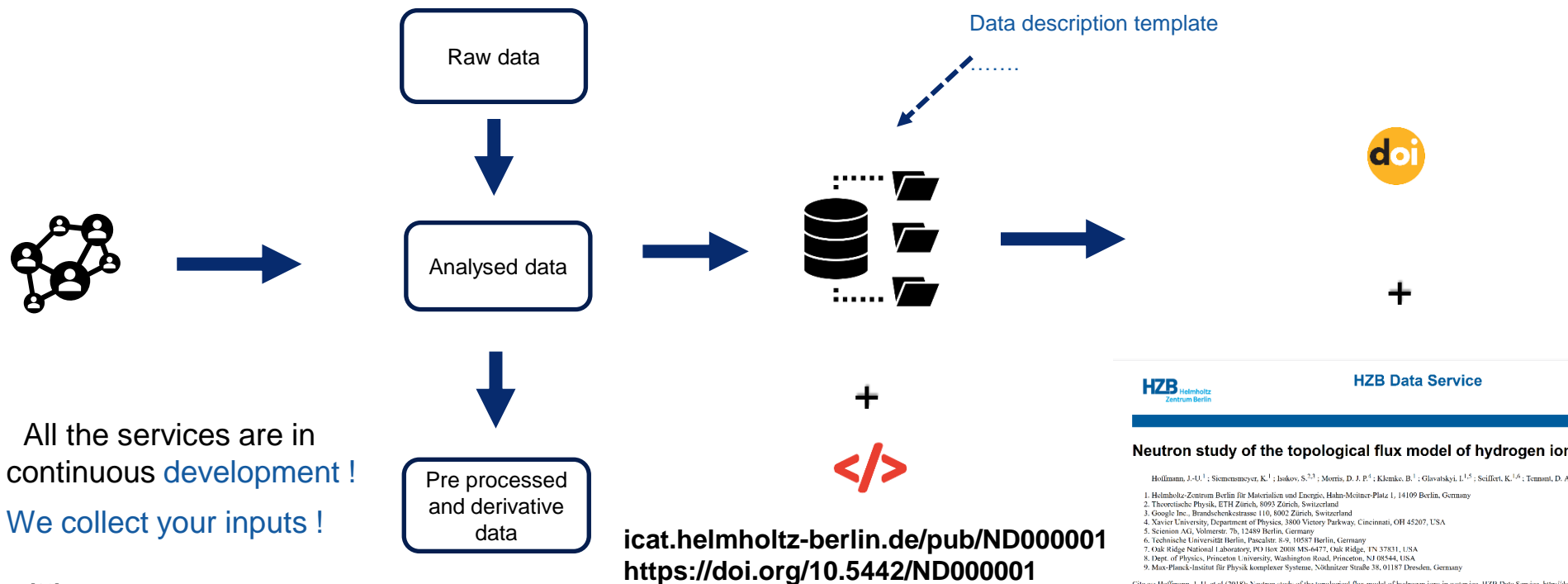


Figure 2: Visualization of the data infrastructure connecting the SISSY I instrument with services inside and outside the HZB





- **Federated service** for data discovery
(~20 PaN research facilities involved)
Interlinking PIDs at each stage in the data process
to provide traceability
EOSC data catalogues services and analysis services
- Development of **research group- community dictionary** to implement as standard.
- Define **application** for a general representation of the beamlines measurement and specific for each end station or research group
- **Enable eelogbook access to external data authors**
- **Actions:** gather ideas, test new implementation, optimizing and updating the tools

- Further **tailoring of tools**
- Test the workflow in action
- **HZB in ExPaNDS and Pansoc that work to bridge the national facilities into EOSC and OpenAire**
Initiatives focusing on the definition of ontologies and vocabularies for experimental techniques
- External users: remote desktop applications_ VISA, FastX and Jupyter notebook providing a recipe for data analysis

- 10. <https://zenodo.org/record/4424770#.YXKNwedCRzp>
- 2. I. Boscaro, Clarke, F. Cesmat, K. Roarty. 2020. Expands vision and roadmap. 10.5281/zenodo.4424770
- 1. B. Matthews, Expands symposium for librarians, 2021
- 6. Rolf Krahel Berlin 2018; Persistent Identification of Instruments WG
www.rd-alliance.org/groups/persistent-identification-instrument-wg
- Rolf Krahel workshop for Research data management at HZB, 2019
<https://www.helmholtz-berlin.de/media/media/spezial/events/datenmanagement/5-hzb-icat.pdf>
- 17. Collins et al., 2021. ExPaNDS ontologies v1.0.
10.5281/zenodo.4806026

Thanks.

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Helmholtz-metadata.de/en/pages-helpdesk
Hmc-matter@helmholtz-berlin.de

There were great efforts put in the last two decades in the research community to elaborate a common standard for high data-rate macromolecular crystallography (HDRMX). This agreed “Gold Standard” builds on the NeXus/HDF5 NXmx application definition and the International Union of Crystallography (IUCr) imgCIF/CBF dictionary, and it is compatible with major data-processing programmes and pipelines. Here we demonstrate the EuXFEL data packed into a NeXus file, which is fully compliant with the Gold Standard by design, since it is built directly from HDRMX NeXus definitions. We use open-source software developed both by community (cctbx) and in-house (extra-data).

Small files generated at beamline better to be encapsulated in larger HDF5 for the ingestions in tapes.

In the framework of national and international initiatives work on homogeneous solutions that take nevertheless in account the diversity of each infrastructure

Need of have granted access to the data from the external