

Workshop on PIDs within NFDI

Report of the Working Group “Persistent Identifiers (PID)” of the
Section *Common Infrastructures* of the NFDI

Imprint

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Table of Content

Introduction	5
Presentations	7
PIDs for data repository entries, datasets, and schemas (FAIRmat)	7
PIDs for Instruments (NFDI4Ing)	7
PIDs for Samples (Daphne4NFDI)	7
IGSN (NFDI4Earth)	8
Survey variables (KonsortSWD)	9
PIDs for Catalysis (NFDI4Cat)	10
StrainInfo PIDs for microbial strains (NFDI4Microbiota)	10
PIDs for Cultural Data (NFDI4Culture & RADAR4Culture)	11
PIDs for digital twins of plant genetic resources (FAIRagro)	11
Requests for the PID4NFDI proposal	12
Further reading	13
Appendix	14

Abstract

In order to gain an overview of the current state of the discussion on PIDs and for the identification of use cases for the initiation phase of a PID service within the NFDI basic services, the working group Persistent Identifier of the Section *Common Infrastructures* of the NFDI hosted an online workshop in January 2023. In the course of the workshop, members of nine different NFDI consortia presented the current application of PIDs in their consortia.

Introduction

In recent years, persistent identifiers (PIDs) have been widely accepted in the general scientific community to identify research assets like data objects, general research outputs, or the researchers themselves. Because of this, members of all NFDI consortia use different kinds of PIDs already in their everyday work. As of today the assignment of identifiers in all NFDI consortia is scattered and heterogeneous in terms of actors, services, scope, quality, and costs involved. Having the core function of PIDs in RDM in mind, an analysis of existing gaps and the development of joint solutions in order to serve the needs of the individual communities and the NFDI as a whole are needed.

For these reasons the working group Persistent Identifier within the Section *Common Infrastructures* of the NFDI, with representatives of existing PID services and infrastructures who are already involved in various domain-specific consortia, was initiated. The aim of the working group is to develop a common strategy for the implementation and extension of community-based and broadly used PID services that is closely aligned with the needs of NFDI consortia. This includes the necessary technical & organisational measures as well as the training and educational aspects particularly around standardised and complete metadata ensuring compliance with the FAIR principles.

In order to get a joint overview of the current state of the discussion on PIDs and, as examples, their current applications in the NFDI consortia, an online workshop with PID experts from the NFDI consortia and the working group members was conducted on January 27, 2023, from 09 to 11 a.m. The results of the workshop also served as a basis for the planned application for PID services in the NFDI and for the identification of use cases for the initiation phase of a PID service within the NFDI basic services.

After an introductory presentation on the motivation and objectives of the working group and the planned basic services, members of the following nine NFDI consortia presented the current application of PIDs in their consortia:

- Daphne4NFDI
- FAIRagro
- FAIRmat
- KonsortSWD
- NFDI4Cat
- NFDI4Earth

- NFDI4Ing
- NFDI4Microbiota
- NFDI4Culture & RADAR4Culture

The presentation slot was followed by an open question and discussion slot. Leading questions were:

- What are the developments in the different NFDI communities?
- How can we learn mutual?
- What do I want from the Persistent Identifier working group?
- What are my wishes for Base4NFDI?
- What are the next steps?

The present report documents key findings and discussion points.

Tab. 1: Program of the Workshop, 27.01.2023

Time	Agenda	Speaker
09:00-09:15	Welcoming and Introduction	Philipp Wieder (GWGD)
09:15-09:20	PIDs for data repository entries, datasets, and schemas (FAIRmat)	Markus Scheidgen (HU Berlin, NOMAD)
09:20-09:25	PIDs for Instruments (NFDI4Ing)	Michael Selzer (KIT)
09:25-09:30	PIDs for Samples (Daphne4NFDI)	Rolf Krahl (HZB)
09:30-09:35	IGSN (NFDI4Earth)	Kirsten Elger (GFZ)
09:35-09:40	Survey variables (KonsortSWD)	Janete Bach (GESIS) Peter Mutschke (GESIS)
09:40-09:45	<i>Break</i>	
09:45-09:55	PIDs for Catalysis (NFDI4Cat)	David Linke (Leibniz-Institut für Katalyse e.V.)

09:55-10:00	StrainInfo PIDs for microbial strains (NFDI4Microbiota)	Lorenz Reimer (DSMZ)
10:00-10:05	PIDs for Cultural Data (NFDI4Culture & RADAR4Culture)	Desiree Mayer (SLUB Dresden) Sandra Göller (FIZ Karlsruhe)
10:05-10:10	Digital twins of plant genetic resources (FAIRagro)	Matthias Lange (IPK) Daniel Arend (IPK)
10:10-10:15	Summary	Britta Dreyer (TIB)
10:15-11:00	Questions & Discussion	Moderated by Philipp Wieder

Presentations

PIDs for data repository entries, datasets, and schemas (FAIRmat)

We currently provide DOIs for datasets to our users on request. In the future, we want to add handle-based PIDs on all our entries (~10 million) automatically. We are also interested in PIDs for users, schemas and data-type definitions, samples, instruments, and more. However, we would prefer a singular service and API with abstract metadata over a pluralism of different PID types. The PID metadata should inform about the type/schema and leave the domain specific metadata to the identified (meta-)data.

PIDs for Instruments (NFDI4Ing)

In the context of NFDI4Ing the KIT plan to expand it's "Gerätepool" (means instrument data base) and to implement PIDs for uniquely identifying the scientific instruments included.

PIDs for Samples (Daphne4NFDI)

DAPHNE stands for DAta for PHoton and Neutron Experiments. The project focuses on research with photons and neutrons at large-scale research facilities. As distinguished from most other NFDI consortia, DAPHNE's scope is not a particular field of research, but rather a

class of experimental methods. The users of photon and neutron facilities come from almost all areas of research.

As a result, a huge variety of samples need to be considered. To illustrate this with at least a few examples: we have simple material samples, such as single crystals, crystal powders, or novel functional materials. But we also have complex structures, such as solar cells or batteries. An important case are macromolecular crystals, e.g. proteins. But we also have archeological artefacts, fossils, artwork, e.g. paintings, and even living plants. A special case are “ephemeral samples”: samples that are created, measured and destroyed during one single experiment, such as a liquid jet injected in the measurement chamber.

The accurate description of the sample being measured is an important part in the metadata for the resulting dataset. In many cases, the full history of sample preparation steps constitutes such a description. Assuming that each preparation and characterization step in turn generates a FAIR dataset, what we actually need is a sample PID that ties all these datasets together and allows us to track the sample's history.

Note that the lab that created the sample may be a different institution than the facility performing the measurement. Therefore, an internal sample id of the lab will not be enough. Furthermore the sample may be held in a collection outside the context of the experiment. So we need to be able to integrate existing external IDs.

Samples may be complex structures composed out of individual samples, each having their own history. On the other hand, a sample may be cut into pieces, each part being treated differently in subsequent steps. Therefore a parent / child relation is needed between sample PIDs.

How we should deal with ephemeral samples is still an open discussion.

IGSN (NFDI4Earth)

The International Generic Sample Number (IGSN) is a globally unique, resolvable persistent identifier for physical objects originally developed for earth sciences samples. IGSN identifier link to sample descriptions on the internet supporting sample discovery and enabling the connection between datasets, publications and the originating samples.

Since 2012, IGSNs have been registered as Handles (one namespace: 10273, internal organization managed by adding centrally-managed agent-specific namespaces). IGSN is an addition to the original sample name and shall not replace it. If cited in research articles, IGSNs have the possibility to “outsource” detailed sample descriptions from the manuscript texts. IGSN is already an established PID system with >10 M. registered samples and managed by IGSN e.V., a global, non-profit organization. Recent activities of IGSN e.V. reach out beyond the Geosciences domain (e.g. life sciences, archaeology, Daphne4NFDI).

In October 2021, DataCite and IGSN announced their strategic partnership and from January 2023, IGSNs are registered as DataCite DOIs. Consequently, each DataCite member can assign a DOI-IGSN. IGSN has modular and quite rich metadata with a “common kernel” ideally provided for all samples which can be further customized to serve more subdisciplines or disciplines. There are agreed recommendations to map IGSN metadata to the DataCite Schema, however, only using the DataCite schema (that has been developed for different purposes) would “only” represent a summary of the sample description, because the DataCite Schema is much more generic than IGSN metadata; we still need to find out a convenient model for not losing the richness of the IGSN metadata and sample descriptions in the new world of DataCite DOI-IGSNs.

Links and further information on IGSN:

- IGSN e.V: www.igsn.org
- IGSN GitHub at <https://igsn.github.io>
- IGSN-related project (2022/2023): FAIR WISH - FAIR Workflows to establish IGSN for Samples in the Helmholtz Association - (i) expanding domain-specific metadata for different geo-bio samples, (ii) development of a sample description template. All project output available in the FAIR WISH Zenodo Community: https://zenodo.org/communities/fair_wish
- Latest publication about IGSN: Klump, J., Lehnert, K., Ulbricht, D., Devaraju, A., Elger, K., Fleischer, D., Ramdeen, S. and Wyborn, L., 2021. Towards Globally Unique Identification of Physical Samples: Governance and Technical Implementation of the IGSN Global Sample Number. *Data Science Journal*, 20(1), p.33. DOI: <http://doi.org/10.5334/dsj-2021-033>

Survey variables (KonsortSWD)

Referencing research data and their inherent detailed entities supports FAIR usage. We enhance the state of the art of citing research data, by developing an infrastructure to reference detailed attributes, here initially variables, within such data. By assigning PIDs to these attributes, individual elements of the data files can be referenced and retrieved with the required metadata for machine-actionable and human access. The PIDs will not only enable citeability within scientific papers but also give access for processing the contained data itself, e.g., within script languages like R or python. We provide recommendations for assigning Persistent Identifiers (PIDs) below the study level. The recommendations rely on use case partners' data types and services. However, it would also benefit other institutions, such as data repositories that hold tabular data and want to take advantage of uniquely

identifying their data at a lower granularity level. To this end, the daJra PID registration service was enlarged under the KonsortSWD Measure TA.5-M.1.

PIDs for Catalysis (NFDI4Cat)

In catalysis samples prepared by machines or in a lab play an important role. These samples are often difficult to reproduce even when they are prepared by the same protocol. Therefore, tracking the materials used and knowing for which sample instance catalytic or characterisation measurements were carried out is of huge importance in the community (as confirmed by a survey). Often the work on one sample/material is moreover distributed across labs and institutions which would also benefit from a wider use of PIDs.

Early 2022 NFDI4Cat created an internal working group on PIDs with the goal to make PIDs available to catalysis researchers from all institutions and possibly also to industrial partners which contribute materials to catalysis research. One goal is to make PIDs available “from the beginning”, that is for the synthesis in the lab. No relabelling or renaming should be necessary until publication in the NFDI4Cat RDM portal. We have been developing a handle-based service as part of our shared infrastructure that can be interfaced from local RDM tools like ELNs (electronic lab notebooks) and the central NFDI4Cat repository that is being built. Besides for samples the PIDs will also be useful for other resources applied in the research process such as materials, devices, instruments, models, schemas etc. A layered metadata schema is suggested similar to IGSN that also reflects confidentiality requirements.

StrainInfo PIDs for microbial strains (NFDI4Microbiota)

Within NFDI4Microbiota the database StrainInfo (<https://straininfo.dsmz.de/>) is developed. Straininfo is a database for collecting, matching and providing persistent identifiers for microbial strains. In the field of bacteria and archaea every newly cultivated and described species needs to be deposited as type strain in at least two international culture collections. Each collection is identifying strains with their own stable identifier that is recognizable. Through time, bacterial strains are assigned with multiple identifiers, which again are used in literature and databases to refer to used strains. To resolve if two identifiers from two collections point to the same strain can be cumbersome. Therefore the StrainInfo database collects and matches strain identifiers on a large scale, which is not limited to culture collection numbers but also includes sequence accession number and DOIs. In the long run a new central registry for microbial strains should be established, that offers persistent

identifiers independent from culture collections and thereby can be applied to any microbial strain.

PIDs for Cultural Data (NFDI4Culture & RADAR4Culture)

In NFDI4Culture, authority data IDs are the most widely used PIDs for research data. ORCID, ROR, geoNames and also wikidata are quite commonly used.

To meet these special needs in the 4Culture-Community, RADAR4Culture, the publication service for NFDI4Culture from FIZ Karlsruhe has implemented the possibility to enter GND-IDs for subject headings in the metadata-field “keywords”. This is an important step towards FAIR publications and the linkage between research data. Current discussions are about more controlled fields in metadata schemas of repositories that should require PIDs, the use of PIDs for different data types, e.g. PIDs for complex search queries (with timestamps), DOIs for annotations or specific elements of a publication, such as single entries in work catalogs, and the need of stable URIs for vocabularies and their provision to make them mappable to larger terminologies.

More information:

- About RADAR: <https://radar.products.fiz-karlsruhe.de/en/radarabout/ueber-radar>
- Event of NFDI4Culture:
<https://nfdi4culture.de/news-events/events/forum-data-publication-and-availability-5-persistent-identifiers.html>

PIDs for digital twins of plant genetic resources (FAIRagro)

Agriculture is facing increasing challenges, e.g., growing demand for food with stagnant productivity, climate change, biodiversity loss, and natural resource degradation. One pillar is plant breeding, which can in principle make use of a big data ecosystem (Tian et al. 2021). Around collections and gene banks to preserve PGRFA, the volume of data potentially useful for plant breeding is rapidly increasing due to technological advancements in genome analysis, precision phenotyping, and digitization in the plant breeding value chain. Data is generated, stored, and analyzed by a variety of stakeholders, such as private sector companies, resort research, or scientific research institutions with different primary interests. The types of acquired data generated range from temporally and spatially resolved data to analysis results on quantitative and qualitative traits or molecular characteristics, developed software, and records of field experiments, which may include scoring data on agronomic and breeding traits as well as information on fertilization, crop protection, field and soil conditions, geodata, and weather data. The data differ not only in their

object of study, but also in their type, format, and context of origin. The NFDI consortium FAIRAgro therefore forms in this domain important anchors for the use cases for the infrastructure being developed.

This proposed use case aims to implement a Digital Twin (DT) service to use IGSNs as PUIDs for material samples, complement DOIs as recommended PUIDs for Plant Genetic Resources for Food and Agriculture (PGRFA) (Manzella, Daniele 2016) by individual samples. The unique identification of all samples, materials, collected data, and annotated metadata, as well as their relation to each other, must be performed throughout the lifecycle to enable their integration into a shareable data space. This is where the Digital Twin concept (DT) (Vachalek et al. 2017) comes in as an extension of the Fair Digital Object (FDO) (Wharton 2022). DTs are virtual, digital representations of products or services and consist of a physical part and a virtual representation as well as a bridging link between the two (Portela et al. 2020), thereby also mapping the dynamics of the object. Digital twins are about the **"as-is" state** of an entity, while an FDO is a **historical collection** of data that may affect many entities over a period of time. This concept, originally built on an approach to integrate complex industrial and digital processes, could be complementary to DOIs, as an identifier for PGRFA, towards a solution to identify samples, which is based on a particular material, and could be the basis for FAIR and transparent documentation of the research data lifecycle for plant research and plant breeding data in field and lab.

Requests for the PID4NFDI proposal

During the discussion slot the participants were asked for their wishes and requests for the planned PID basic service (PID4NFDI). The participants highlighted that the development of cook books, guidance, best practices, and the support of open discussions in the NFDI community would be very helpful.

A participant also mentioned that PIDs and confidentiality is not a contradiction and emphasized that one doesn't need to expose any information when minting e.g. a handle.

Metadata richness for PIDs, which was seen closely connected with Interoperability, were also topics of the discussion. At the end of the workshop, a participant made the point to take the discussion further on if there is a need for a unified PID service to avoid the splitting up of the PID landscape.

Further reading

Bertelmann, R., Buys, M., Kett, J., Pampel, H., Pieper, D., Scholze, F., Sens, I., Burger, F., Dreyer, B., Glagla-Dietz, S., Hagemann-Wilholt, S., Hartmann, S., Schrader, A., Schirrwagen, J., Summann, F., & Vierkant, P. (2023). PID Network Deutschland. Netzwerk für die Förderung von persistenten Identifikatoren in Wissenschaft und Kultur. Helmholtz Open Science Office. <https://doi.org/10.48440/os.helmholtz.059>

Bingert, Sven, Brase, Jan, Burger, Felix, Dreyer, Britta, Hagemann-Wilholt, Stephanie, Vierkant, Paul, & Wieder, Philipp. (2022). Concept for Setting up the Persistent Identifier Services Working Group in the NFDI Section "Common Infrastructures" (1.0). Zenodo. <https://doi.org/10.5281/zenodo.6507760>

Klump, J., Lehnert, K., Ulbricht, D., Devaraju, A., Elger, K., Fleischer, D., Ramdeen, S., & Wyborn, L. (2021). Towards Globally Unique Identification of Physical Samples: Governance and Technical Implementation of the IGSN Global Sample Number. In Data Science Journal (Vol. 20). Ubiquity Press, Ltd. <https://doi.org/10.5334/dsj-2021-033>

Appendix

- PID und Basisdienste - Philipp Wieder
- PIDs for data repository entries, datasets, and schemas (FAIRmat) - Markus Scheidgen
- PIDs for Instruments (NFDI4Ing) - Michael Selzer
- PIDs for Samples (Daphne4NFDI) - Rolf Krahl
- IGSN (NFDI4Earth) - Kirsten Elger
- Survey variables (KonsortSWD) - Janete Bach, Claus-Peter Klas & Peter Mutschke
- PIDs for Catalysis (NFDI4Cat) - David Linke
- StrainInfo PIDs for microbial strains (NFDI4Microbiota) - Lorenz Reimer
- PIDs for Cultural Data (NFDI4Culture & RADAR4Culture) - Desiree Mayer & Sandra Göller
- Digital twins of plant genetic resources (FAIRagro) - Daniel Arend

PID und Basisdienste

Ausgangslage für PID Entwicklungen



Vorarbeiten und aktuelle Entwicklungen

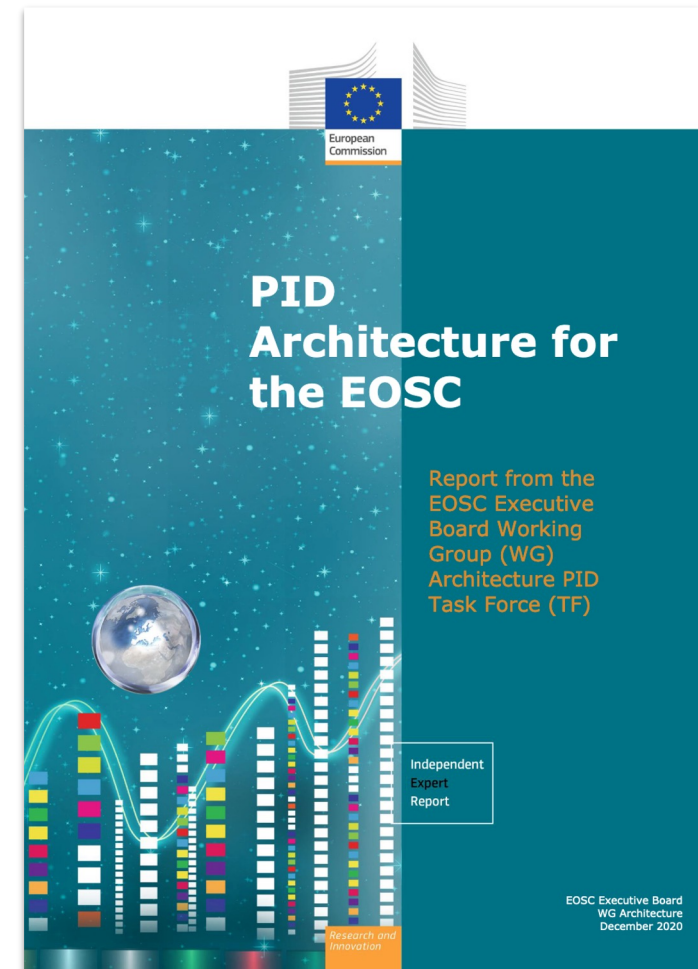
- EOSC
 - Publication: PID Architecture for the EOSC [1]
 - PID Policy and Implementation Task Force
- Verschiedene Gruppen der Research Data Alliance
- Projekten, die EOSC zuarbeiten
 - FREYA (beendet)
 - DICE
 - FAIRCORE4EOSC

NFDI Persistent Identifier Services Working Group (PID WG)

- Wichtigkeit des Themas innerhalb der NFDI unbestritten
- Konzept ist publiziert [2]
- Aktuell 13 Einrichtungen aktiv

27.01.2023

Workshop on PIDs within NFDI



Ziele der PID Working Group



1. Erfassung der PID Use Cases und Anforderungsanalyse
 - Evaluierung der Disziplinen/Konsortien
 - Anforderungsanalyse
 - Abbildung der Anforderungen auf existierende Services
2. Entwicklung NFDI-weiter Konzepte für die Nutzung und Integration relevanter PID Systeme
 - Fokus auf einfacher Nutzung & Integration
 - Interoperabilität von Metadaten
 - Einbettung in “NFDI Service Architektur”

Concept for Setting up the Persistent Identifier Services Working Group in the NFDI Section “Common Infrastructures”

Name of the working group
Persistent Identifier Services

Acronym
infra-pid

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Version 1.00
2022-04-28

<https://zenodo.org/record/6507760#.YxBrbvHP0-R>

Ziele der PID Working Group



3. Wissenstransfer zur Unterstützung Nutzung und Entwicklung von PID Diensten
 - Cook Books und Best Practices
 - Weitergabe von Informationen an Projekte, Entwickler & weitere Stakeholder
4. Entwicklung von Governance- und Lizenzmodellen für die NFDI
 - Auswertung existierender Modelle
 - Definition & Umsetzung von spezifischen Erweiterungen für die NFDI
5. Zusammenarbeit mit deutschen und internationalen Institutionen bzgl. PIDs
 - Beratung von & Zusammenarbeit mit NFDI Konsortien
 - Fokus auf EOSC bzgl. generellen Dienstangebot
 - Kollaboration mit (inter)nationalen Playern

Deliverables



- Sammlung von konsolidierten Anforderungen der NFDI Konsortien
- Einheitlicher, einfacher Zugang zur Nutzung und Registrierung von PIDs
- Schnittstellen zur Nutzung und Integration von PID Services
- Technische (Weiter)entwicklung: z.B. neue PID Typen, einheitliche Metadaten, ...
- Organisatorische Konzepte bzgl. Governance und Lizenzen
- Monitoring der Dienste
- Betriebsmodell und rechtlich geprüftes Rahmenwerk für NFDI-weiten Betrieb

Leitfragen für Heute

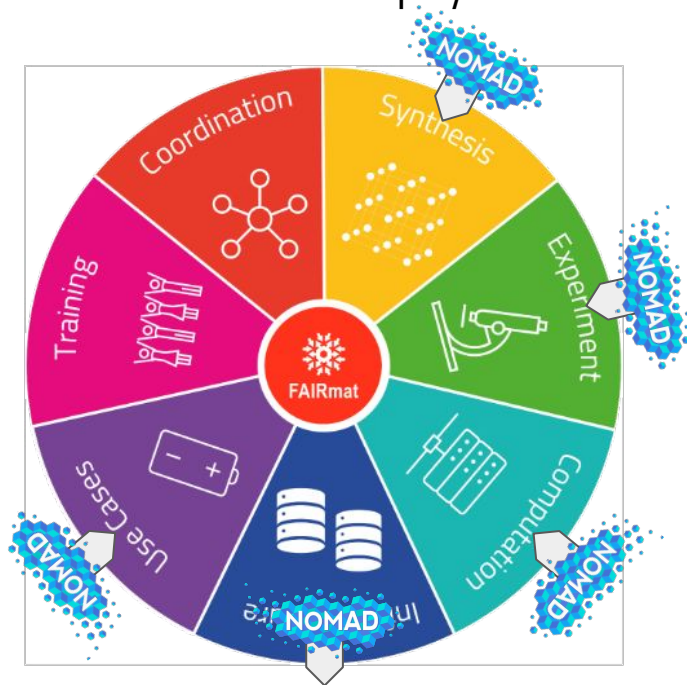
- Welche Entwicklungen gibt es in den verschiedenen NFDI Communities?
- Was kann ich von den anderen lernen?
- Welche Wünsche habe ich an Persistent Identifier AG?
- Welche Wünsche habe ich an Base4NFDI?
- Wie geht es weiter?

Quellen

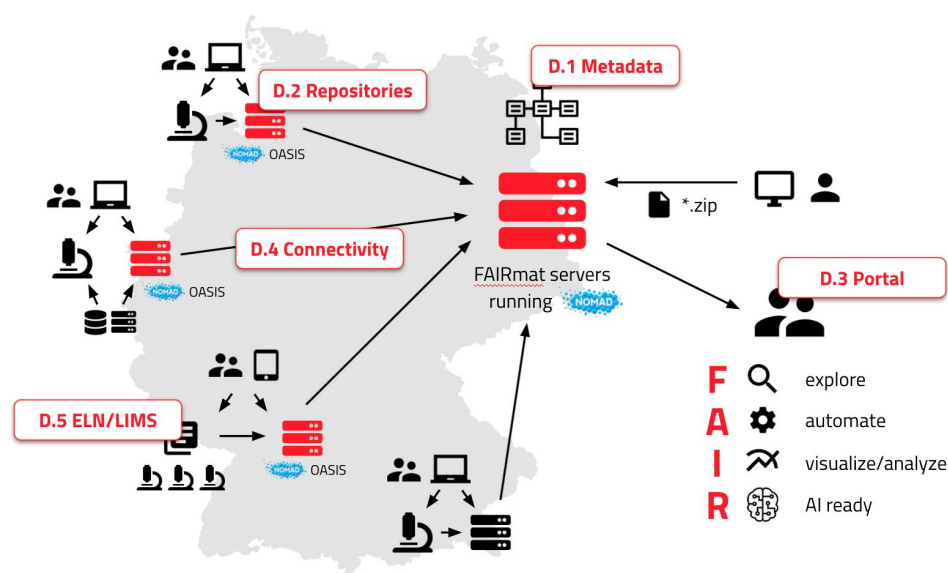
- [1] PID architecture for the EOSC: <https://s.gwdg.de/iWKgoE>
- [2] Concept for Setting up the Persistent Identifier Services Working Group in the NFDI Section “Common Infrastructures”: <https://doi.org/10.5281/zenodo.6507759>
- [3] Nationale Forschungsdateninfrastruktur: Förderung von NFDI-weiten Basisdiensten: <https://s.gwdg.de/APBrFz>

What is FAIRmat and NOMAD?

FAIRmat is the NDFI consortium to build a FAIR federated data infrastructure for solid state physics



NOMAD is a web-based software for FAIR research data management in materials science



What we need from PID and PID services

- a **singular** PID service to issue and resolve PIDs for data from all domains
- meaningful and flexible metadata schema for PID metadata
 - dublin-core-style metadata
 - metadata to describe the type of data (e.g. via schema-pid)
 - metadata to describe non-human programs and instruments as source/"author" (e.g. via source-pid)
 - metadata to describe the repository (e.g. via repository-pid)
- we want to use PIDs for
 - repository entries (simulations, samples, instruments, ELNs, measurement data, documents, ...); $\sim 10^7$ of them
 - user curated datasets; $\sim 10^3$ of them
 - schemas and data-type definitions; $\sim 10^3$ of them
- ideally the service would be able to "promote" PIDs to become DOIs resolveable through the respective existing DOI services (datacite, etc.)
- we want to transition from issuing DOIs on demand to issuing PIDs automatically
- data and PIDs should go through a life-cycle together
 - might end in deletion (ephemeral PIDs) or long term archival
 - PID metadata and what it resolves to has to be changeable throughout the whole life-cycle

Instrument data base, meta data and related topics

A short introduction to the setup of a KIT Gerätepool and a brief outlook on further developments and possible fields of application



The KIT instrument data base („Gerätepool“)

- Background - Why do we need a Gerätepool?
- Who is responsible for the development of the Gerätepool?
- First bundle of instruments
- First round of questions to the instrument experts
- Where and how will the Gerätepool be realised?
- Future goals

The KIT instrument data base („Gerätepool“)

- First round of questions -

Erfassung von Gerätedaten

AG Gerätepool

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Prof. Dr. Christian Kübel (INT) christian.kuebel@kit.edu +49 721 608-28970

Dr. Matthias Mail (INT) matthias.mail@kit.edu +49 721 608-28909

Anlage: xxxxx

Anlagenbezeichnung: _____

Gerätebezeichnung: _____

(vom Hersteller)

Hersteller: _____

Erstzugang am: _____

Gerätezustand: Originalzustand

Upgrade _____

Beschreibung

Generalüberholt _____

Modifiziert _____

Beschreibung

Datum

Software für das Gerät: _____

Institut: _____

Standort: _____

Raum: _____

The KIT instrument data base („Gerätepool“)

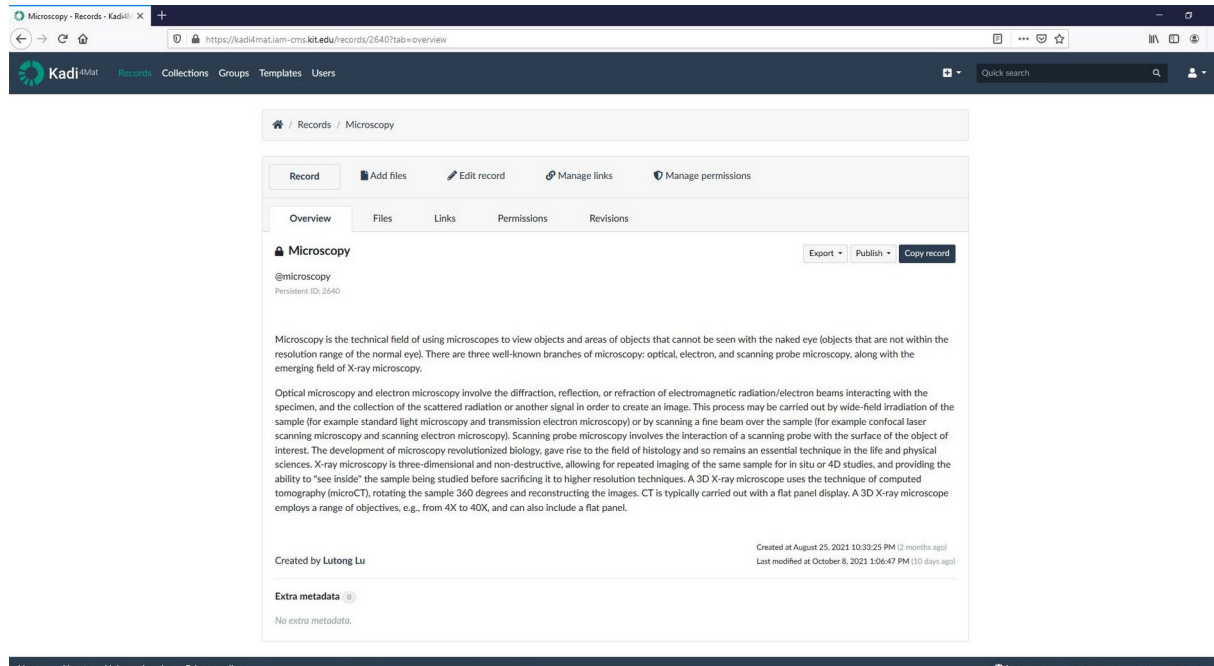
- First bundle of instruments -



Excel ribbon: Einfügen, F K U, Zwischenablage, Schriftart, Ausrichtung, Zahl, Bedingte Formatierung, Als Tabelle formatieren, Zellenformatvorlagen, Einfügen Löschen Format, Zellen, Sortieren und Suchen und Filtern, Auswählen, Bearbeiten

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P
Anlage	Anlagenbezeichnung	Anlagenbezeichnung	Gerätebezeichnung (vom Hersteller)	Hersteller	Geräteklasse	Subklasse	Erstzugang am	Gerätezustand	Leistungsbeschreibung 1	Leistungsbeschreibung 2	Software für das Gerät	Institut	Standort	Raum	Gerätebetreiber (Geräteverantwortlicher)
2	40002275	Zubehör Tandem-Massenspektrometrie-System mit Ioni	-Erweiterung: Probenaufgabenmodul				19.02.2016					IFG	0330	167	Frank Kirschhöfer
3	40002373	Bio-AFM Rastersondenmikroskop					12.02.2010					IFG	0330		Dr. Hartmut Gilemann
4	40002379	UHV - NEKAFS - Apparatur - gebraucht -					03.02.2010					IFG	0330		Dr. Alessi Nefedov
5	40002382	FT-IR-Spektrometersystem					14.01.2010					IFG	0330	314	Stefan Heissler
6	40002387	Diffraktometersystem D8					22.03.2010					IFG	0330	068	Dr. Peter Weidler
7	40003043	ToF-SIMS	Time of Flight Secondary Ion Mass Spectrometer				03.09.2010					IFG	0321	313	Dr. Alexander Velle
8	40003094	FTIR-Mikroskop					10.06.2011					IFG	0330	314	Stefan Heissler
9	40003097	Hochenergie-Röntgen-Diffraktometer D8 DISCOVER					17.06.2011					IFG	0330	005	Dr. Peter Weidler
10	40003116	Quarz Micro Kristallwaage Q-Sense E4 QCM-D System					17.10.2011					IFG	0330	067	Frank Kirschhöfer
11	40003191	UVVIS Spektralphotometer Cary 5000					05.06.2012					IFG	0330	314	Stefan Heissler
12	40003354	DVS	Dynamic Vapor Sorption Analyzer				30.01.2014					IFG	0330	068	Dr. Lars Heinke
13	40003373	Röntgendiffraktometer D8					14.04.2014					IFG	0330	53	Franz König
14	40003599	Gaschromatograph 597B MSD System Agilent					31.05.2017					IFG	0330	166	Frank Kirschhöfer
15	40003653	ESI-QTOF Massenspektrometer					22.12.2017					IFG	0330	168	Frank Kirschhöfer
16	60104587	Dynamisches Kontaktwinkelmessgerät und Tensiometer					08.08.2011					IFG	0330	068	Ingo Fischer
17	60104840	AKTApurifier 10 mit Zubehör					20.12.2011					IFG	0330	127	Prof. Dr.-Ing. Matthias Franz
18	60107826	HPLC-Komplettanlage Agilent 1100					04.12.2013					IFG	0330	166	Frank Kirschhöfer
19	60108177	FT-IR-Spektrometer Typ Vertex 70					17.12.2013					IFG	0330	054	Dr. Peter Thissen
20	60108309	Complete Xnano Instrument System	Indirect nano plasmonic sensing				27.03.2014					IFG	0330	213	Dr. Hartmut Gilemann
21	60111473	Probe System MPS150 mit System Sourcormeter 2635B	Leitfähigkeitsmesssystem				31.12.2015					IFG	0330	353	Dr. Lars Heinke
22	60112632	GE Äkta Purifier100 FPLC Anlage,					31.12.2016					IFG	0330	230	Prof. Dr.-Ing. Matthias Franz
23	60112669	STARe System TGA 2 Vorführgerät	Thermogravimetric analysis				02.11.2016					IFG	0330	068	Dr. Matthias Schwozter
24	60113447	EMXnano Bench - Top Demo System					13.10.2017					IFG	0330	368A	Peter Krolla
25	60113474	AZURA Lab SMB System Knauer	LC-System				27.10.2017					IFG	0330	126	Prof. Dr.-Ing. Matthias Franz
26	60114138	QCM 3T-analytik					26.02.2018					IFG	0330	385	Dr. Peter Weidler
27	60114139	QCM 3T-analytik					26.02.2018					IFG	0330	385	Dr. Peter Weidler
28	60114184	Zubehör zu Mikroskop Zeiss Asioplan	gehört zu FA 60071176, neue ND = 4 Jahre				13.12.2017					IFG	0330	268	Peter Krolla
29	60116676	Malvern Zetasizer Nano-ZS	- gebraucht -				12.07.2019					IFG	0330	168	Frank Kirschhöfer
30	60117183	Rasterkraftmikroskop Nanosurf NaioAFM					13.12.2019					IFG	0330	065	Dr. Hartmut Gilemann
31	40003753	Konfokales Laserscanning-Mikroskop Zeiss					14.02.2019					IFG-NPB	30.43	706	Martin Bastmeyer, Prof. Dr.
32	40003778	Spectroscopic Imaging Nulling Ellipsometer					01.09.2018					IFG-NPB	0330	202	Dr. Melke König-Edel
33	60118621	Solar Ellipsometer V3-03-T (gebraucht)					03.07.2015					IFG-NPB	0330	202	Dr. Melke König-Edel
34	60112644	Digitalmikroskop (Steuergerät & Grafiksystem)					06.12.2016					IFG-NPB	0330	206	Dr. Melke König-Edel

The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat



The screenshot shows a web browser window displaying the Kadi4Mat interface. The URL is <https://kadi4mat.iam-cms.kit.edu/records/2640?tab=overview>. The page title is "Microscopy - Records - Kadi4Mat". The interface includes a navigation bar with "Records", "Collections", "Groups", "Templates", and "Users". Below the navigation bar, there are tabs for "Overview", "Files", "Links", "Permissions", and "Revisions". The "Overview" tab is active, showing a record for "Microscopy" created by "Lutong Lu". The record includes a description of microscopy, its history, and its applications. The record was created on August 25, 2021, and last modified on October 8, 2021.

Kadi4Mat is the Karlsruhe Data Infrastructure for Materials Science, a software for managing research data with the aim of combining new concepts with established technologies and existing solutions.

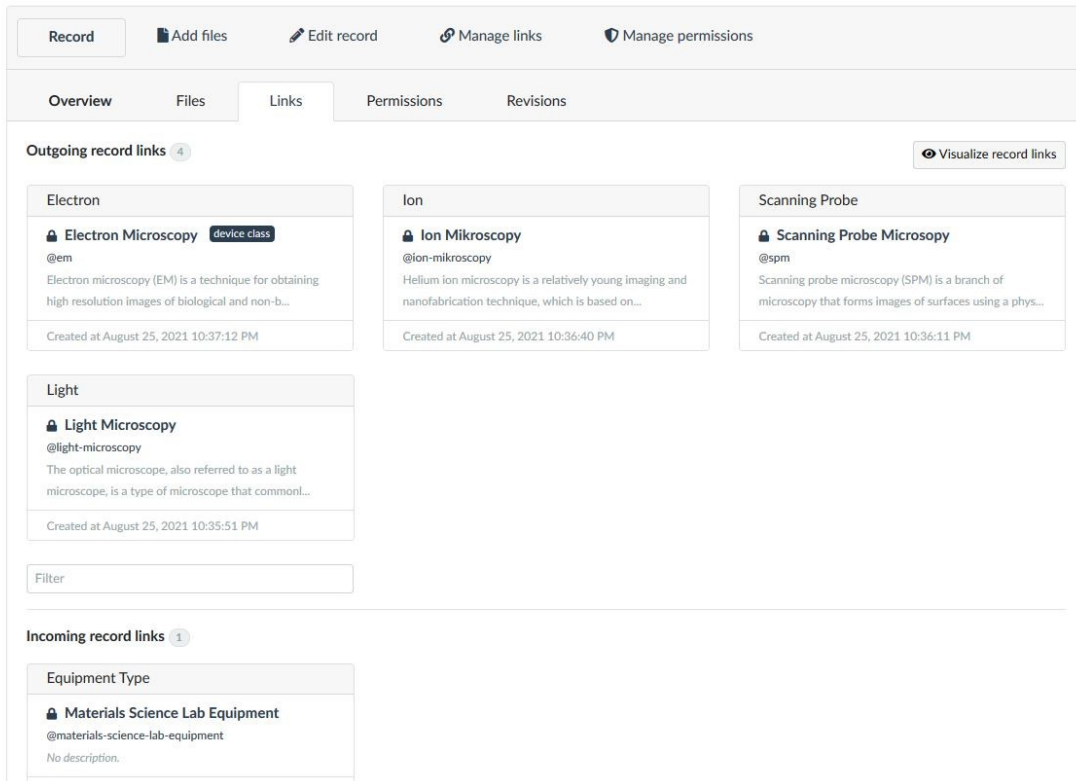
<https://kadi4mat.iam-cms.kit.edu/about>

The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat - Overview



Kadi4Mat is the **Karlsruhe Data Infrastructure for Materials Science**, a software for managing research data with the aim of combining new concepts with established technologies and existing solutions.

<https://kadi4mat.iam-cms.kit.edu/about>



The screenshot displays the Kadi4Mat interface for managing record links. At the top, there are navigation buttons: Record, Add files, Edit record, Manage links, and Manage permissions. Below this is a tabbed interface with 'Overview', 'Files', 'Links', 'Permissions', and 'Revisions'. The 'Links' tab is active, showing 'Outgoing record links' (4) and a 'Visualize record links' button. The outgoing links are categorized into three groups: Electron, Ion, and Scanning Probe. Each group contains a record card with a lock icon, a title, a user handle, a description, and a creation timestamp. Below the outgoing links is a 'Filter' input field. At the bottom, there is a section for 'Incoming record links' (1) with a card for 'Materials Science Lab Equipment'.

Record Add files Edit record Manage links Manage permissions

Overview Files Links Permissions Revisions

Outgoing record links 4 Visualize record links

Electron

🔒 **Electron Microscopy** device class
@em
Electron microscopy (EM) is a technique for obtaining high resolution images of biological and non-b...
Created at August 25, 2021 10:37:12 PM

Ion

🔒 **Ion Mikroskopy**
@ion-mikroskopy
Helium ion microscopy is a relatively young imaging and nanofabrication technique, which is based on...
Created at August 25, 2021 10:36:40 PM

Scanning Probe

🔒 **Scanning Probe Microscopy**
@spm
Scanning probe microscopy (SPM) is a branch of microscopy that forms images of surfaces using a phys...
Created at August 25, 2021 10:36:11 PM

Light

🔒 **Light Microscopy**
@light-microscopy
The optical microscope, also referred to as a light microscope, is a type of microscope that common...
Created at August 25, 2021 10:35:51 PM

Filter

Incoming record links 1

Equipment Type

🔒 **Materials Science Lab Equipment**
@materials-science-lab-equipment
No description.

The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat - Searching



Kadi4Mat is the Karlsruhe Data Infrastructure for Materials Science, a software for managing research data with the aim of combining new concepts with established technologies and existing solutions.

<https://kadi4mat.iam-cms.kit.edu/about>

Search Sort by Relevance Search extras

Create new record

Select a record template

No results found

10 100

Hide public records

Filter by collection

Include child collections

Filter by type

Filter by tag

Filter by MIME type

49 results found

Microscopy @microscopy Created 2 months ago
Last modified 10 days ago

Microscopy is the technical field of using microscopes to view objects and areas of objects that cannot be seen with the naked eye (objects that are not within the resolution range of the normal eye). There are three well-known branches of microscopy: optical, electron, and scanning probe microscopy...

Created by **Lutong Lu**

Electron Microscopy device class @em Created 2 months ago
Last modified 10 days ago

Electron microscopy (EM) is a technique for obtaining high resolution images of biological and non-biological specimens. It is used in biomedical research to investigate the detailed structure of tissues, cells, organelles and macromolecular complexes. The high resolution of EM images results from t...

Created by **Lutong Lu**

EDX @edx Created 14 days ago
Last modified 14 days ago

No description.

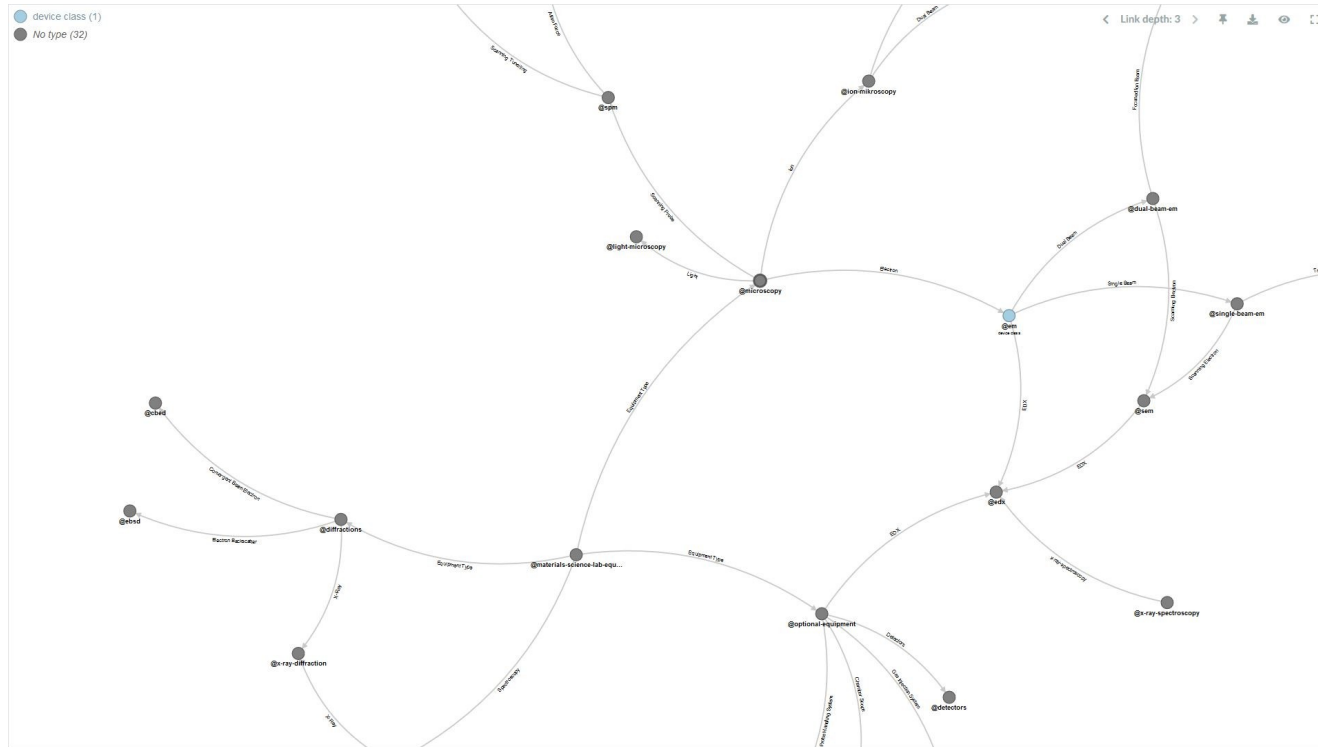
Created by **Matthias Mail**

Scanning Electron Microscopy @sem Created 2 months ago
Last modified 14 days ago

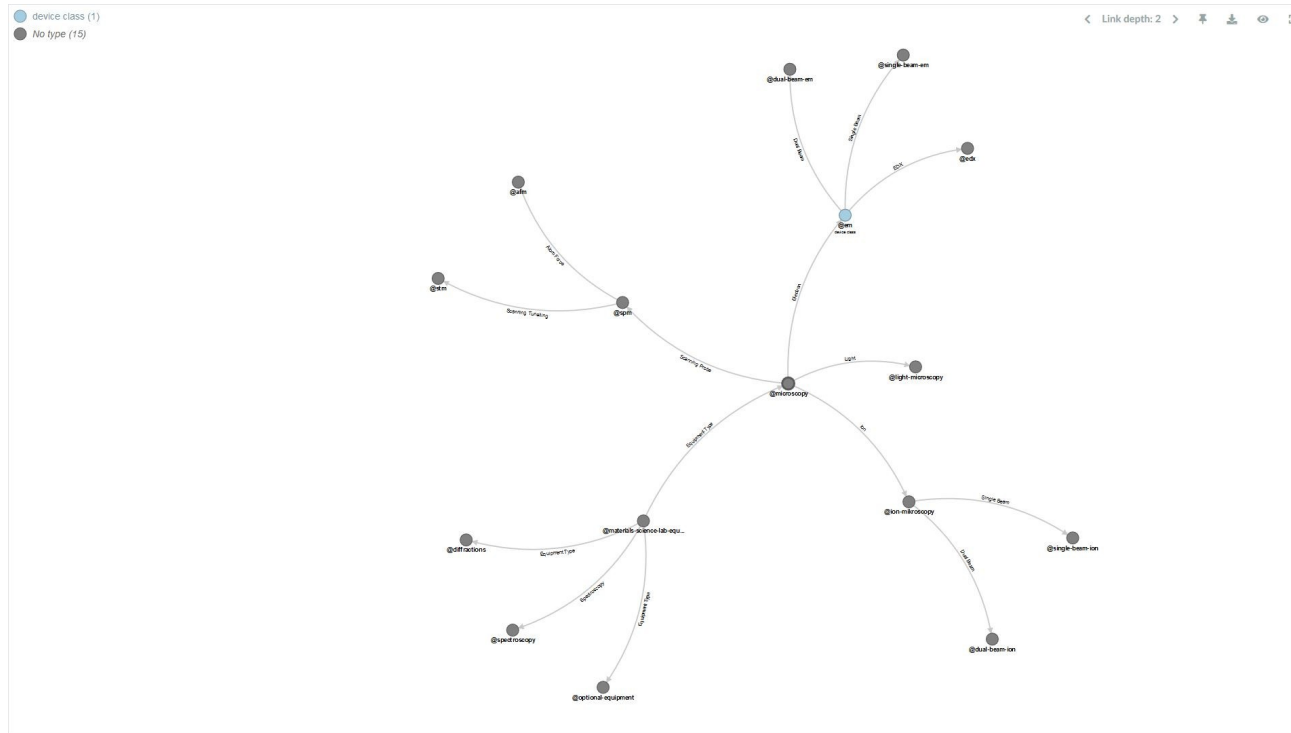
A scanning electron microscope (SEM) is a type of electron microscope that produces images of a sample by scanning the surface with a focused beam of electrons. The electrons interact with atoms in the sample, producing various signals that contain information about the surface topography and compos...

Created by **Lutong Lu**

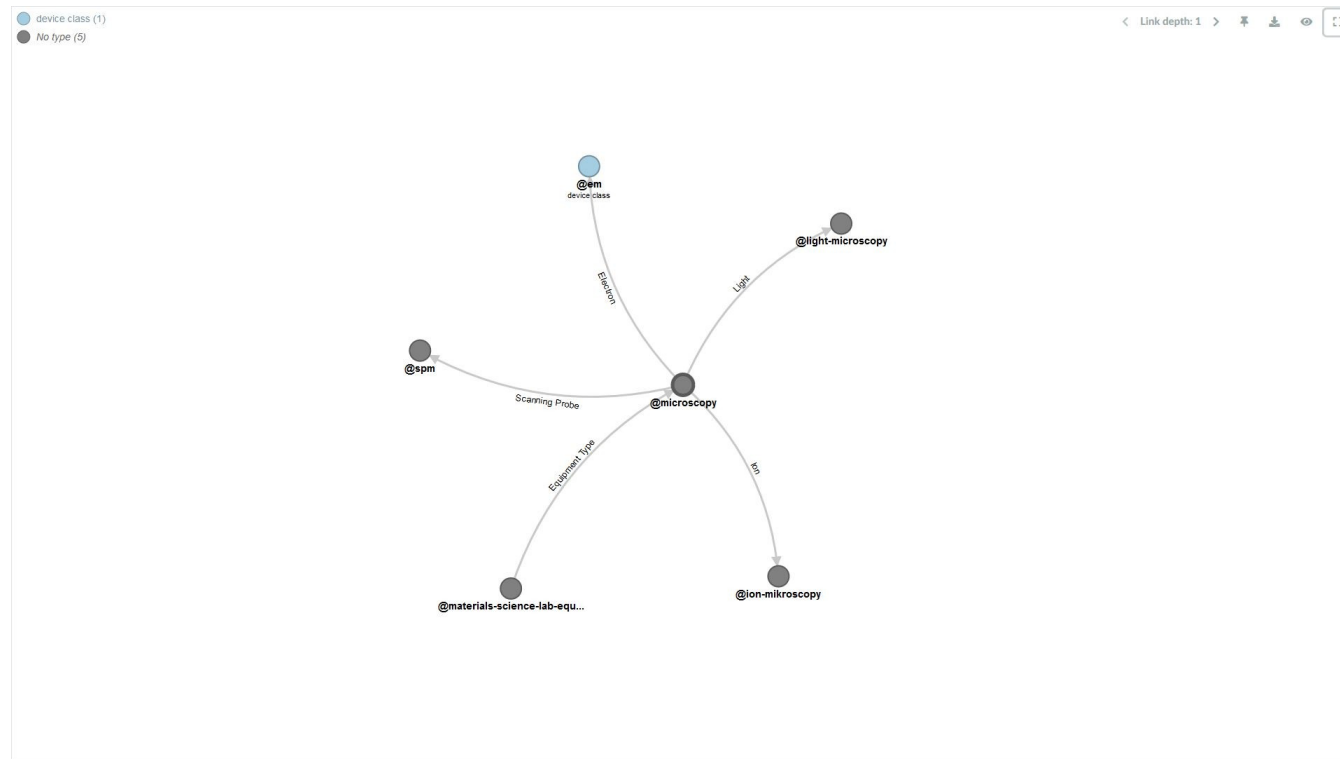
The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat - Searching



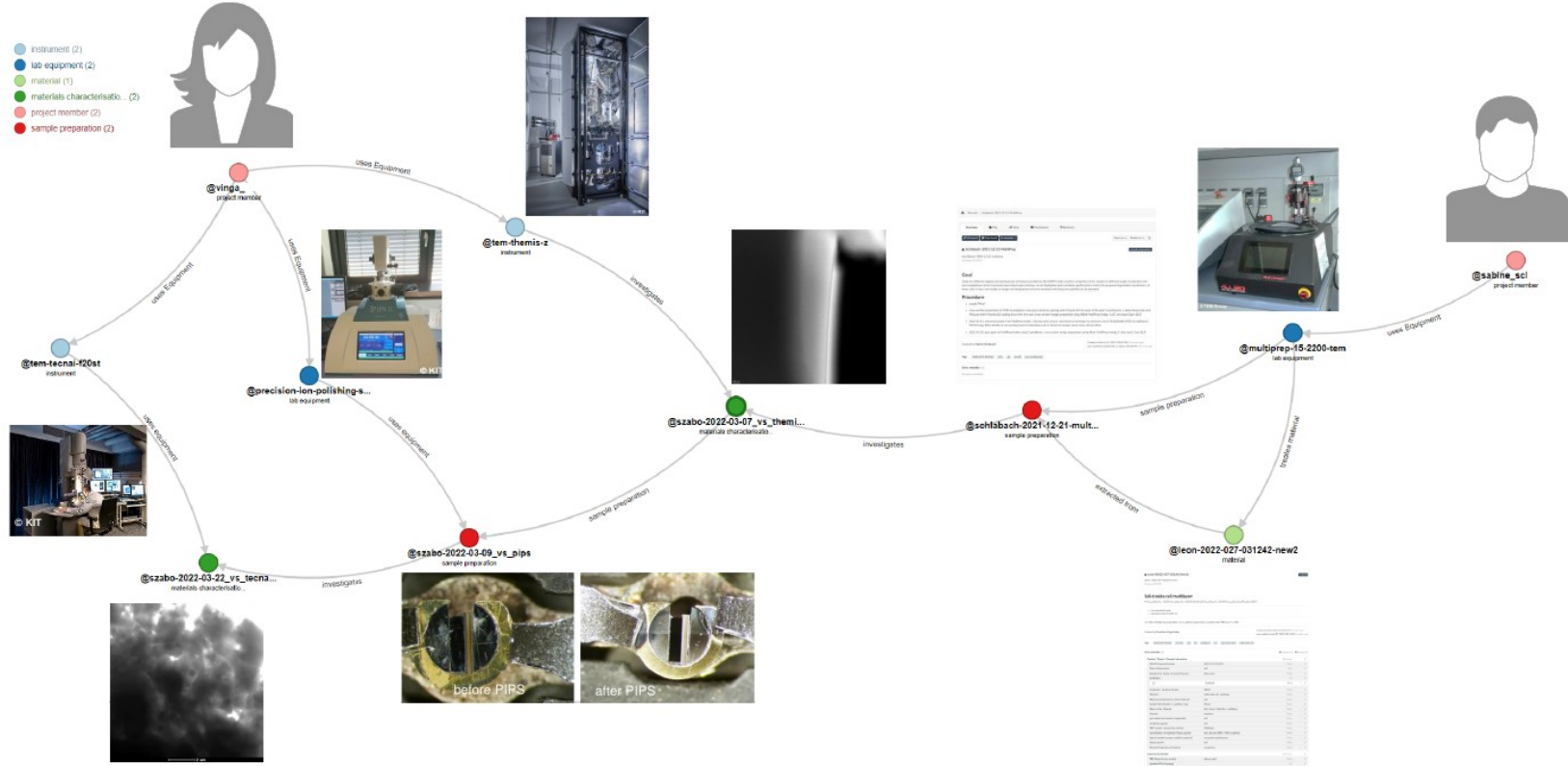
The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat - Searching



The KIT instrument data base („Gerätepool“) - Inside Kadi4Mat - Searching



Usage of Instrument descriptions in the research process



The KIT instrument data base („Gerätepool“)

- Future goals -

- Including more instruments
 - Older
 - Smaller/Cheaper ones
 - etc....
- Full description of all instruments
- Using the database for other topics, e.g.
 - Ontology development
 - Correlative approaches
 - Electronic Lab Notebooks
 - etc.



- Thank you for listening
- Any questions?

PIDs for Sample in DAPHNE4NFDI

Rolf Krahl 

Workshop on PIDs within NFDI, 27 January 2023



- DAta from PHoton and Neutron Experiments (DAPHNE4NFDI): photons and neutrons at large-scale research facilities.
- Scope of DAPHNE is a class of experimental methods, not a particular field of research. We have research from all areas.
- Accurate description of the sample being measured is an important part in the metadata for the resulting dataset.
- Huge diversity of samples.

Note

Discussion on sample PIDs in DAPHNE is at the very beginning. No conclusive concept yet. I can only present my own personal view on the matter here.

We have all sorts of different samples. Some examples:

- single crystals, crystal powder, novel functional materials
- complex structures: solar cells, batteries
- macromolecular crystals, e.g. proteins
- archeological artefacts, fossils
- artwork, e.g. paintings
- living plants

Diversity of live cycles:

- Most samples are specifically created for the particular experiment, often in the user's home lab.
- Samples may have a complex history of preparation steps.
- Samples may be split in parts, each part being treated differently in subsequent steps.
- Samples may be held in a collection outside the context of the experiment.
- Operando measurements: samples may be changing during the experiment.
- Ephemeral samples: samples may be created, measured and destroyed during one experiment, e.g. the sample may be a liquid jet injected in the measurement chamber.

Why do we need sample PIDs, what are the requirements:

- The full history of sample preparation steps constitutes the sample description.
- If each preparation and characterization step generates a dataset, if all these datasets are FAIR and if they are all linked together by referencing the sample PID, we get the full picture.
- If the same sample is measured in more than one independent investigations, we want to be able to combine the results.
- We need a parent / child relation for sample PIDs.
- We need to be able to integrate existing external ID, e.g. from collections holding objects.
- Open discussion: ephemeral samples.

IGSN

International Generic Sample Numbers

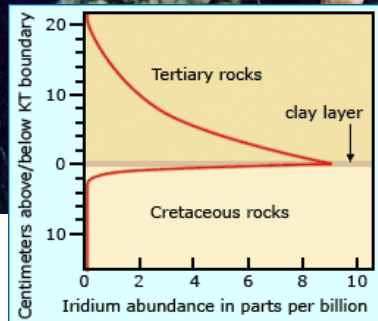
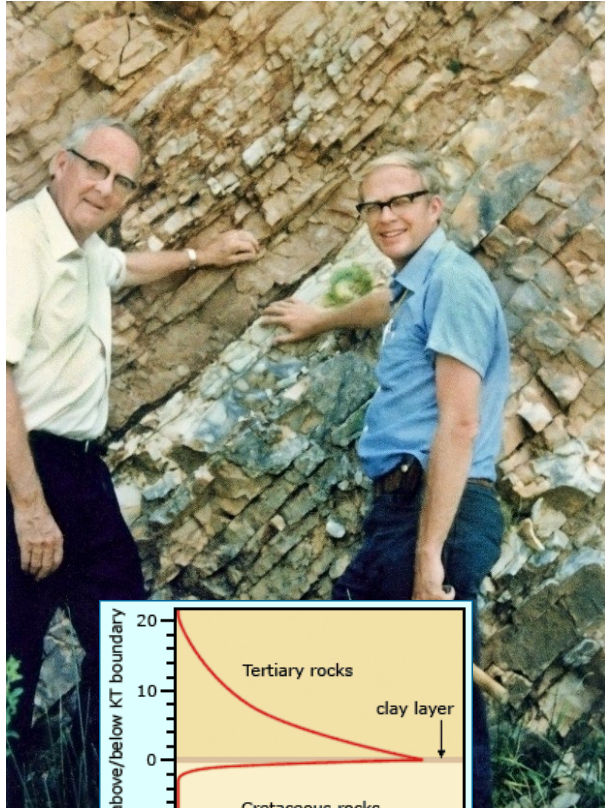
uniquely identifying your samples

Kirsten Elger 

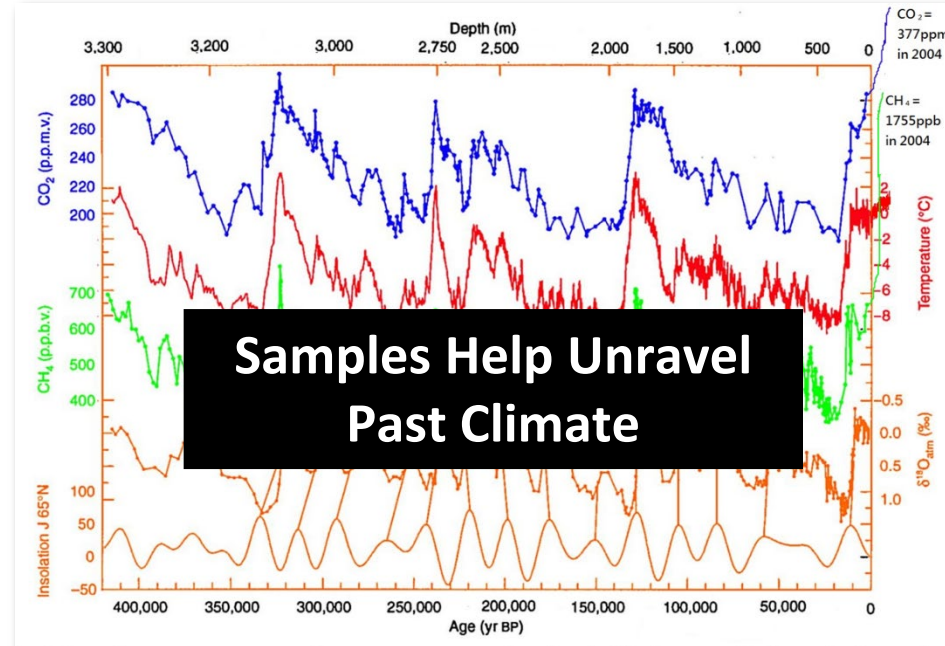
GFZ German Research Centre for Geosciences, Library and Information Services
Vice president of the IGSN e.V.



Samples & Science



Samples Record Unique Events in History



Archaean zircons in Miocene oceanic hotspot rocks establish ancient continental crust beneath Mauritius

Lewis D. Ashwal, Michael Wiedenbeck & Trond H. Torsvik

Nature Communications 8,

Received: 20 September 2016

Article
doi:10.

Samples Provide Access to the Inaccessible

2017

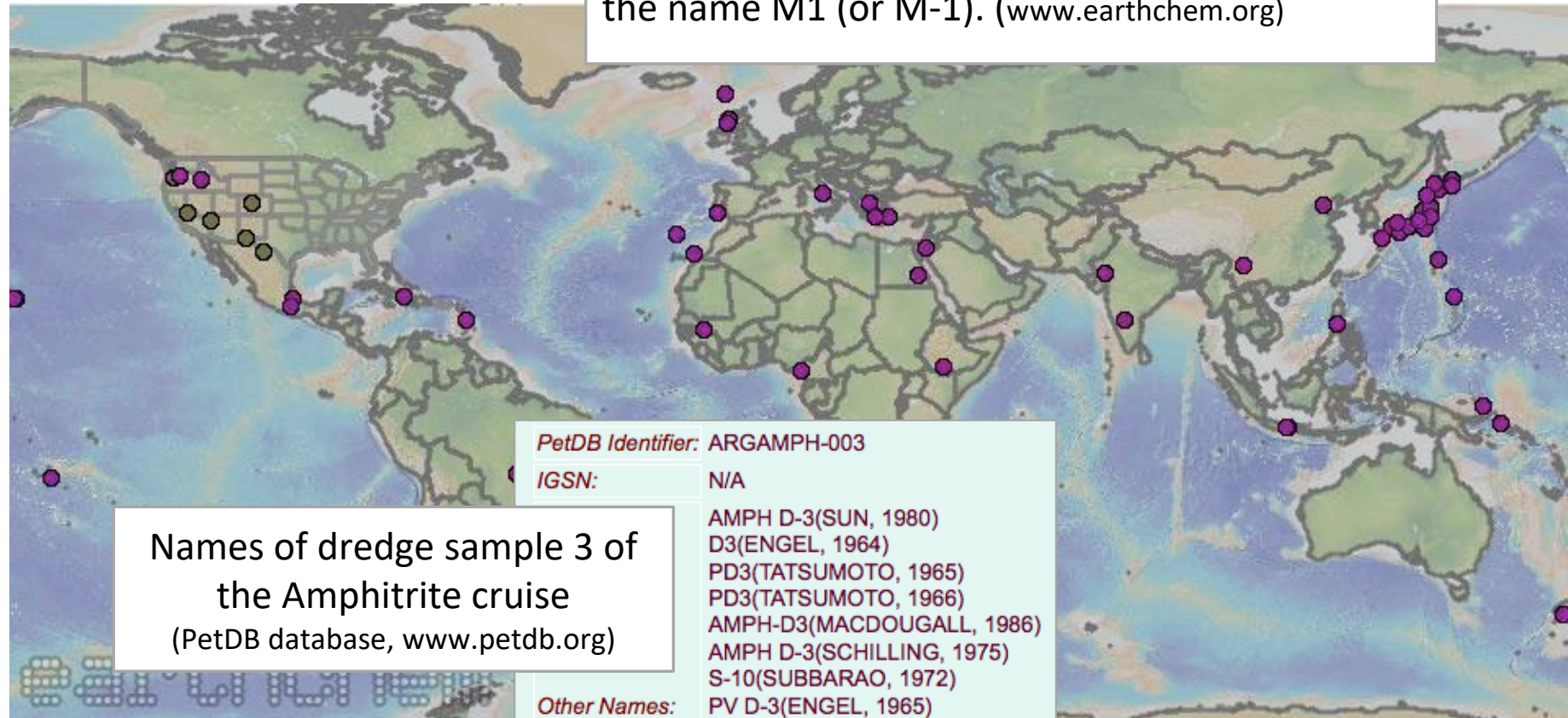
Credit: K. Lehnert (Lamont)

Why IGSN? Unambiguous names

EarthChem Portal

- One-stop shop for geochemical data (at LDEO)
- gives users the ability to search various federated databases simultaneously
- <https://www.earthchem.org/>

The EarthChem Portal includes 75 samples with the name M1 (or M-1). (www.earthchem.org)



Credit: K. Lehnert (Lamont)

IGSN International Generic Sample Number

A globally unique and persistent identifier (PID) for physical objects (originally derived from the Earth Sciences)

- guaranteed to be **unique** via a centralized control mechanism (unique name spaces, Handle identifier)
- resolves to **virtual sample representations** (sample metadata profiles) managed at federated IGSN Allocating Agents.



IGSN: GRO000188

IGSN: GRO000188
Sample Name: 12PTX04
Other Name(s): Rhodes at Pawtuxet
Sample Type: Individual Sample
Parent IGSN: Not Provided

Description	
Material:	Liquid>aqueous
Classification:	Not Provided
Field Name:	Not Provided
Description:	Not Provided

IGSN: GMY00007W

IGSN: GMY00007W
Sample Name: TN182_47_002
Other Name(s):
Sample Type: Individual Sample
Parent IGSN: GMY00001B

Description	
Material:	Rock
Classification:	Igneous>Plutonic>Mafic
Field Name:	gabbro, hornblende gabbro
Description:	mafic plutonic rock

Benefits of IGSN

- unambiguously cite and track physical samples on a global scale:
 - allows previously impossible **linking of samples to data and publications**,
 - allows previously impossible **linking** and integration of **sample-based observations across data systems**, and
 - paves the road towards advanced data mining of sample-based data.
- persistently link to online digital representations of samples (landing pages)
 - builds a federated global sample catalog.
 - are citable in scholarly literature



The screenshot shows the IGSN landing page for sample SO273-31D-10. At the top left is the GFZ logo (Helmholtz Centre Potsdam). The page is divided into several sections: 'General Identifiers' (Program: SO273, Expedition: SO273, Type: Individual Sample, Name: SO273-31D-10, IGSN: GFSO273N31, Parent IGSN: GFSO273N21, Release Date: N/A), 'Sampling Location' (Latitude: -43.8543, Longitude: 38.88, Coordinate System: WGS84, Elevation: -728, Location Type: Ridge, Location Name: Southwest Indian Ridge), 'Geology' (Material: Rock, Rock Classification: Igneous>Volcanic>Mafic, Rock Type: Plag-Basalt, Rock Description: light (most of the primary mineralogy preserved), weathering: light (most of the primary mineralogy preserved), shape: subangular, Manganese crust (mm): 0.1, Lithology: Plag-Basalt, Average grain size: aphanitic, Texture: highly phryic), 'Sampling' (Drilling Method: dredging, Platform Type: Ship, Platform Name: Sonne, Chief Scientist: Prof. Dr. Jürgen Koepke, Start Date: 2020-03-20T23:06:56+00:00, End Date: 2020-03-21T00:55:02+00:00), 'Sample Family' (a tree view of sub-samples SO273-31D-1 through SO273-31D-6), 'Location Map' (a map of the Southwest Indian Ridge with a red dot at the sample location, scale 0-1000 km), and 'Publications & Datasets'.

IGSN Landing Page



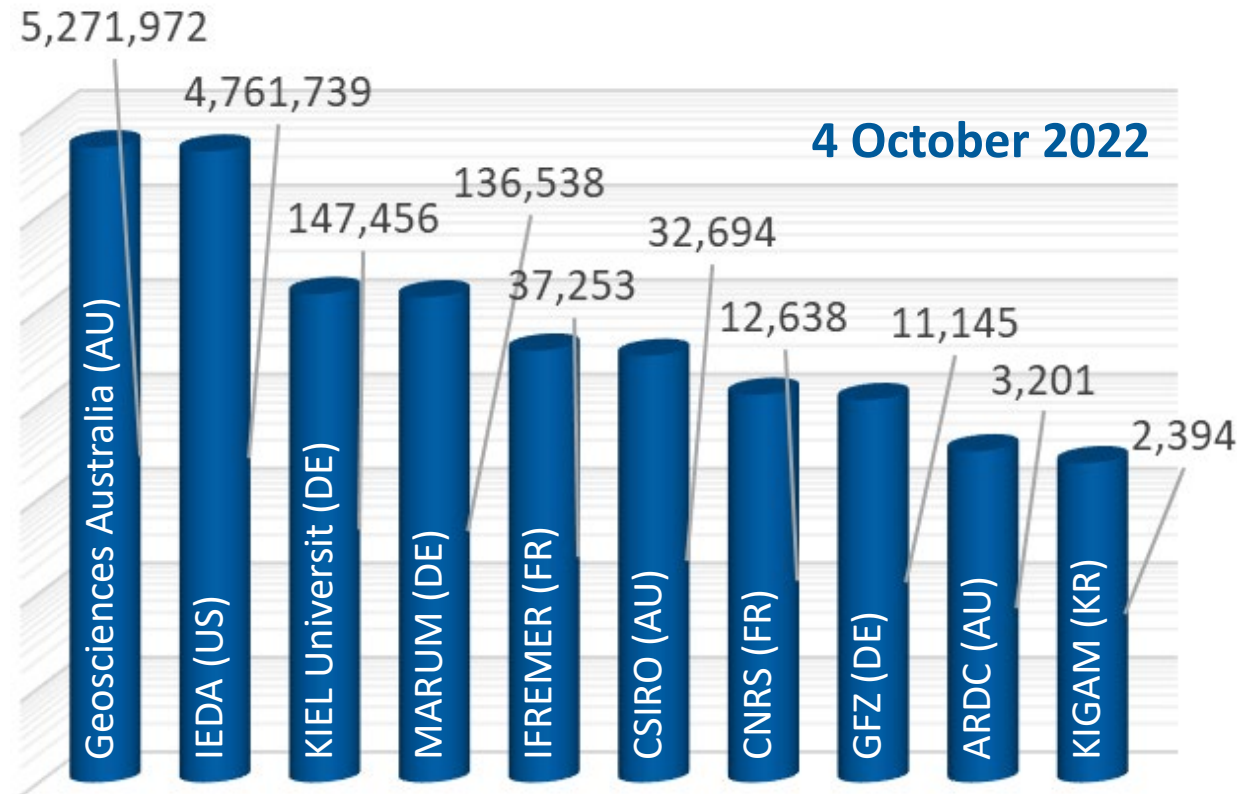
<https://www.igsn.org/>

IGSN e.V.

- established in 2011, registered as non-profit organization in Germany
- Members are organizations that want to provide registration and catalogue services (Allocating Agents)
 - currently 16 members,
 - 6 affiliate members and
 - 10 active Allocating Agents



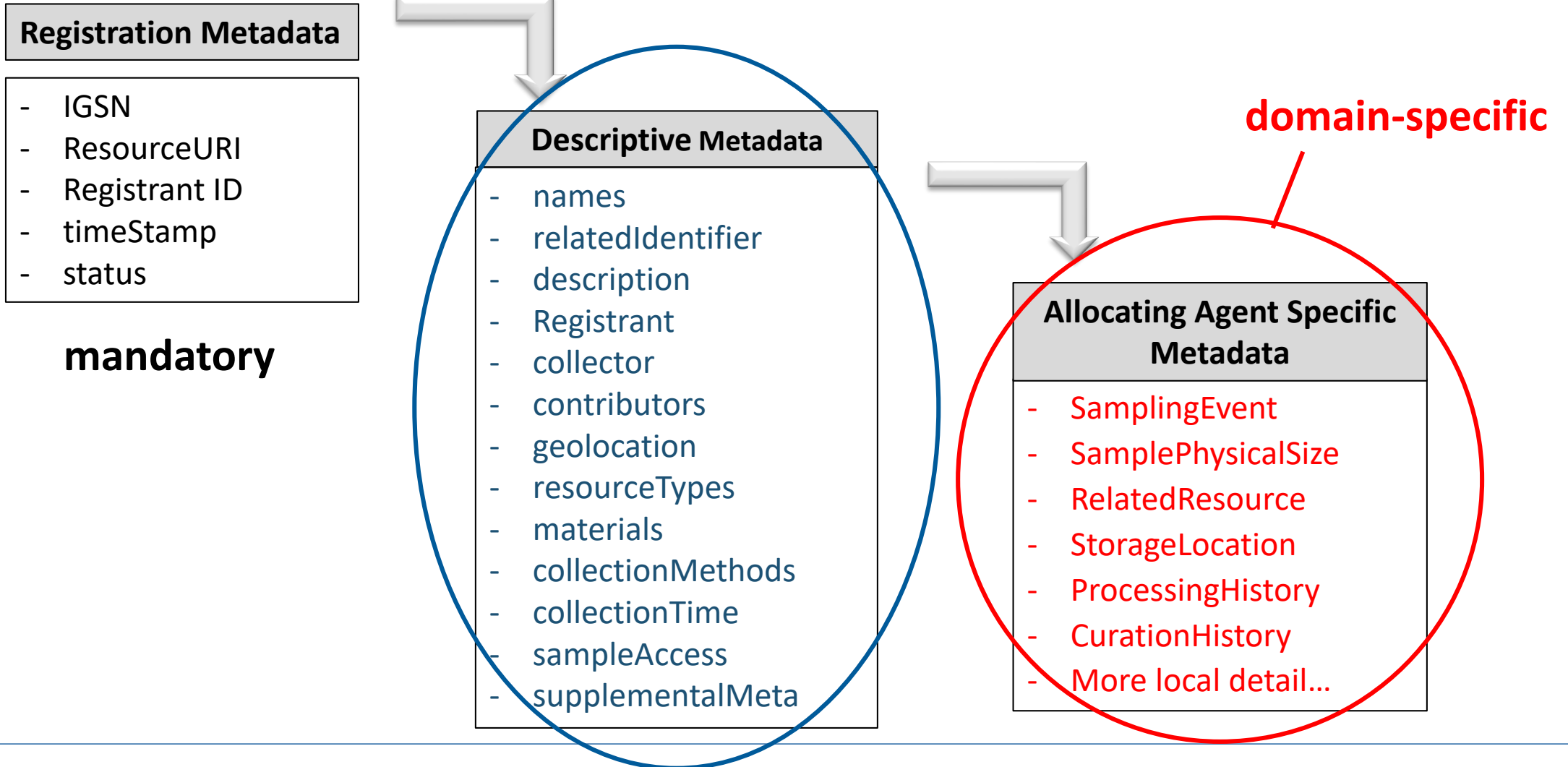
Number of IGSN by Allocating Agent



4 October 2022

Total = 10,417,030 registered IGSNs

IGSN Metadata Levels



Common Kernel: recommended for data discovery

IGSN metadata schema (XML)

```
<?xml version="1.0" encoding="UTF-8"?>
- <resource type="Sample" xsi:schemaLocation="http://pmd.gfz-potsdam.de/igsn/schemas/description/1.3 http://pmd.gfz-potsdam.de/igsn/schemas/description/1.3/resource.xsd" xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xmlns="http://pmd.gfz-potsdam.de/igsn/schemas/description/1.3">
  <identifier type="IGSN">SSDPRR01ECV2101</identifier>
  <name>RR01_1_A_3</name>
  <parentIdentifier type="IGSN">SSDPRR01EH40001</parentIdentifier>
- <registrant>
  <name>GFZ Data Services</name>
  - <affiliation>
    <identifier type="URL">https://ror.org/04z8jg394</identifier>
    <name>GFZ Potsdam</name>
  </affiliation>
</registrant>
- <collector>
  <identifier type="ORCID">0000-0003-2776-0846</identifier>
  <name>Christopher Juhlin</name>
  - <affiliation>
    <identifier type="URL">https://www.ror.org/048a87296</identifier>
    <name>Uppsala University</name>
  </affiliation>
</collector>
- <contributors>
  - <contributor type="ProjectLeader">
    <identifier type="ORCID">0000-0003-2776-0846</identifier>
    <name>Christopher Juhlin</name>
  </contributor>
  - <contributor type="Other">
    <name>Riksriggen, Engineering Geology</name>
  </contributor>
</contributors>
- <geoLocations>
  - <geoLocation>
    <geometry type="Point" sridType="4326">15.255724 59.224578</geo
  </geoLocation>
</geoLocations>
- <resourceTypes>
  <resourceType>http://vocabulary.odm2.org/samplingfeaturetype/spec
</resourceTypes>
- <materials>
  <material>http://vocabulary.odm2.org/medium/rock/</material>
</materials>
- <collectionMethods>
  <collectionMethod>Corer:Rock</collectionMethod>
</collectionMethods>
```

IGSN Description Metadata is intended to describe the core elements of a specimen. The set of attributes is seen as the "birth certificate" of a specimens and should not contain stateful attributes, where possible. The base document for the development are the notes from the [IGSN Metadata Kernel Workshop](#) held in Los Angeles in September 2015.

IGSN Descriptive Metadata Elements

ID	Element	A/C	Occ	Definition	Description and instructions
	resource	Root element			IGSN "birth certificate" for a physical sample, associated feature, or collection
1	identifier	C	1	string	The Identifier is a unique string that identifies a resource. IGSN (International GeoSample Number) registered by an IGSN member. Format should be: "10273/foo"
1.1	identifierType	A	1	include/identifierType.xsd	currently only type=IGSN is supported
2	name		1...1, not nillable	string	Text string for people to understand what is identified. What would typically be presented in a user interface. Collector's or contributor's local/field name used to name the specimen; not globally unique but typically unique within a set of specimens.
3	alternateIdentifiers		0...1		Other formal identifiers for this resource, in addition to the IGSN.
3.1	alternateIdentifier	C	0...n	string	An identifier or identifiers other than the primary Identifier applied to the resource being registered. This may be any

IGSN technical (2012-2022)

- Central registry using the handle.net system
- Handle namespace: 10273
- One central Handle server operated at GFZ
- Uniqueness of IGSNs organized via different, centrally registered, namespaces (for allocating agents) and individual sub-namespaces
- IGSN resolve via <https://igsn.org/<IGSN-number>> or <https://hdl.handle.net/10273/<IGSN-number>>“ to IGSN Landing Pages

The future of IGSN (from 1.1.2023 on)

- IGSNs will be registered as DataCite DOIs (resource type „physical object“)
- IGSN metadata schema will be mapped to DataCite schema (Guidelines available)
- Each DataCite Member or DataCite Consortium Member can assign DataCite „DOI IGSNs“
- DataCite recommends to use individual DOI prefixes for IGSN DOIs

Partnership between IGSN and DataCite

📅 October 28, 2021 👤 Matt Buys and Kerstin Lehnert

<https://doi.org/10.5438/7270-1155>

Earlier this year, [DataCite and IGSN](#) announced their roadmap towards a partnership to support the [global adoption](#), implementation, and use of physical sample identifiers. Today, we are pleased to share the announcement of the partnership agreement.

DataCite is a community-led organisation with a vision to connect research and identify knowledge. We have been providing the means to create, find, cite, connect, and use research across 48 countries globally since 2009. In addition to DataCite core services, we support the scaling efforts of several identifier communities through governance, sustainability, insurance, and technical implementation.

IGSN e.V. is an international, non-profit organization with more than 20 members, which has operated a central registration system for IGSNs since 2011. The IGSN is a globally unique and persistent identifier for physical samples. The core purpose of IGSN is to enable transparent and traceable connections between research activities and objects, including samples, collections, instruments, grants, data, publications, people and organizations.

IGSN and DataCite have a common purpose, and a close relationship in the future will provide mutual benefit to our shared vision of connecting research and identifying knowledge. The partnership brings years of experience across our organizations and communities to scale sample community engagement, develop sample identifier practice standards, and increase adoption globally.

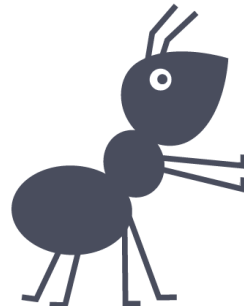
KonsortSWD



Consortium for the
Social, Behavioural, Educational
and Economic Sciences

NFDI4 PID Working
Group 2023

Presentation
27 January 2023



KonsortSWD PID registration Service

Workshop on PIDs within NFDI

Janete Saldanha Bach
Claus-Peter Klas
Peter Mutschke

GESIS – Leibniz Institute for the Social Sciences

Janete Bach



Janete Saldanha Bach is a Researcher at GESIS – Leibniz Institute for the Social Sciences, based in the Knowledge Technologies (KTS) Department, team FAIR Data and Human Information Interaction, working in the consortia KonsortSWD Project of the National Research Data Infrastructure (NFDI). She holds a Ph.D. and a Master's degree in Science and Technology Studies (STS) and a bachelor's degree in Information Science. Her research expertise is in Open Science, especially in research data management and data reuse in the Social Sciences. She is currently involved in consortium KonsortSWD, Task Area 5 Measure 1 - developing the conceptual framework for the PID registration service at a variable level and Task Area 5 Measure 2 Enhancing data findability.

Claus-Peter Klas

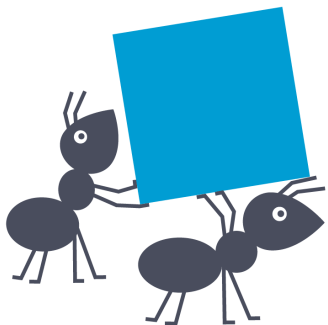


Claus-Peter Klas is lead of the Data & Service Engineering team in the department Knowledge Technologies for the Social Sciences of GESIS. He received his PhD in computer science at the University of Duisburg-Essen and was a postdoctoral researcher in the Department of Multimedia and Internet Applications, Faculty of Mathematics and Computer Science, University of Hagen, Germany. His research focuses on information retrieval, interactive information retrieval, information systems, databases, digital libraries, preservation and grid and cloud architectures. He developed the software Daffodil founded on a nation research project and worked in national and European research projects such as The European Film Gateway, SHAMAN (Sustaining Heritage Access through Multivalent ArchiviNg) and Smart Vortex (Scalable Semantic Product Data Stream Management for Collaboration and Decision Making in Engineering). He is currently responsible for several infrastructure projects within GESIS, such as da|ra, SowiDataNet or Missy, all concerned with providing information and data for social scientists. In addition, he lead the measure PID Services in the national research infrastructure project NFDI. In his team, they are developing an open source DDI suite to support getting DDI into operation.

Peter Mutschke



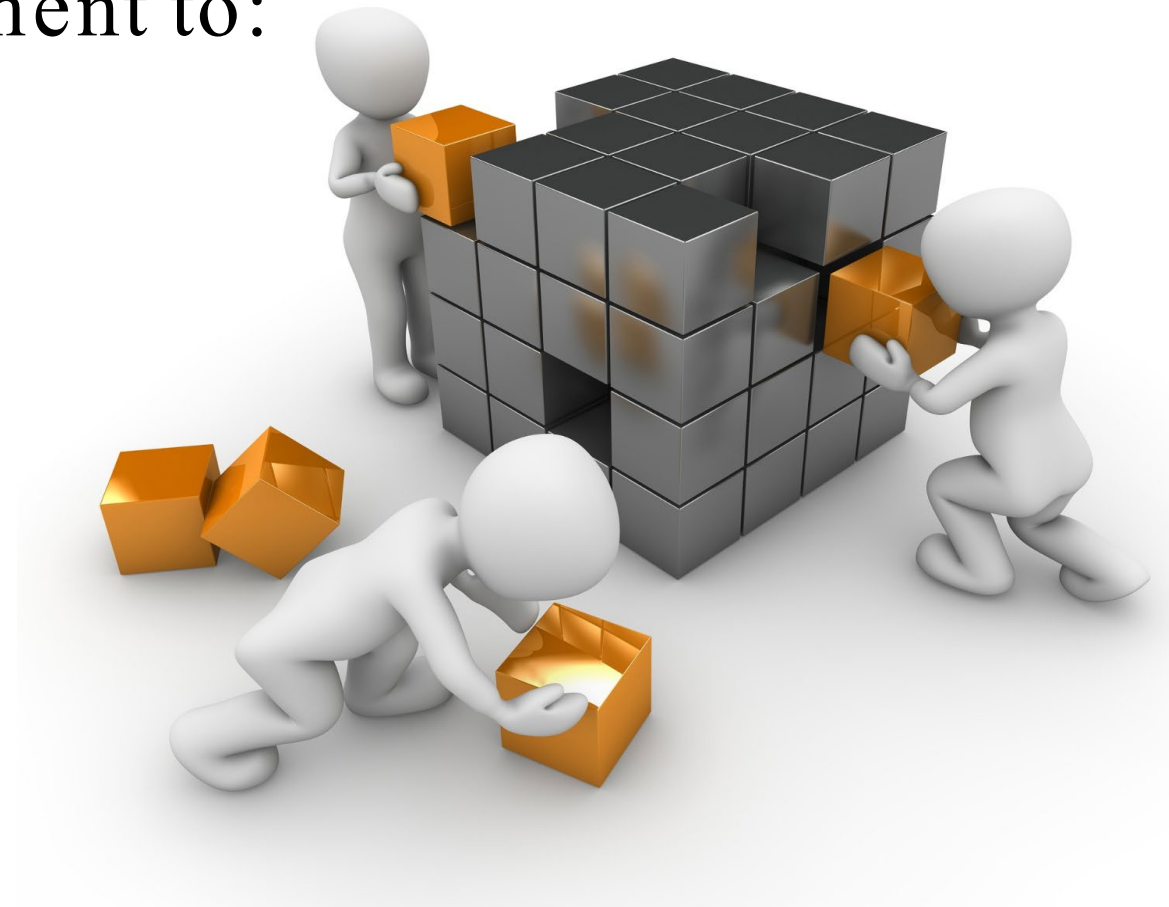
Peter Mutschke is deputy head of the department "Knowledge Technologies for the Social Sciences (KTS)" and leader of the team "FAIR Data and Human Information Interaction" of KTS. His research interests include Information Retrieval, Network Analysis and Open Science. He worked in a number of national and international research projects, such as the DFG projects DAFFODIL and IRM and the EU projects WeGov, SENSE4US, OpenMinTeD and MOVING. Peter served as a member of the management committee of the Leibniz research alliance "Science 2.0/Open Science" from 2013-2021. He founded and coordinates the GO FAIR Implementation Network "Cross-Domain Interoperability of Heterogeneous Research Data (Go Inter)", and he is member of the steering committee of the FAIR Digital Objects Forum (fairdo.org) where he also co-chairs a working group on semantics. He is currently involved in consortia KonsortSWD, NFDI4DataScience and BERD@NFDI of the National Research Data Infrastructure (NFDI). ORCID: <https://orcid.org/0000-0003-3517-8071>.

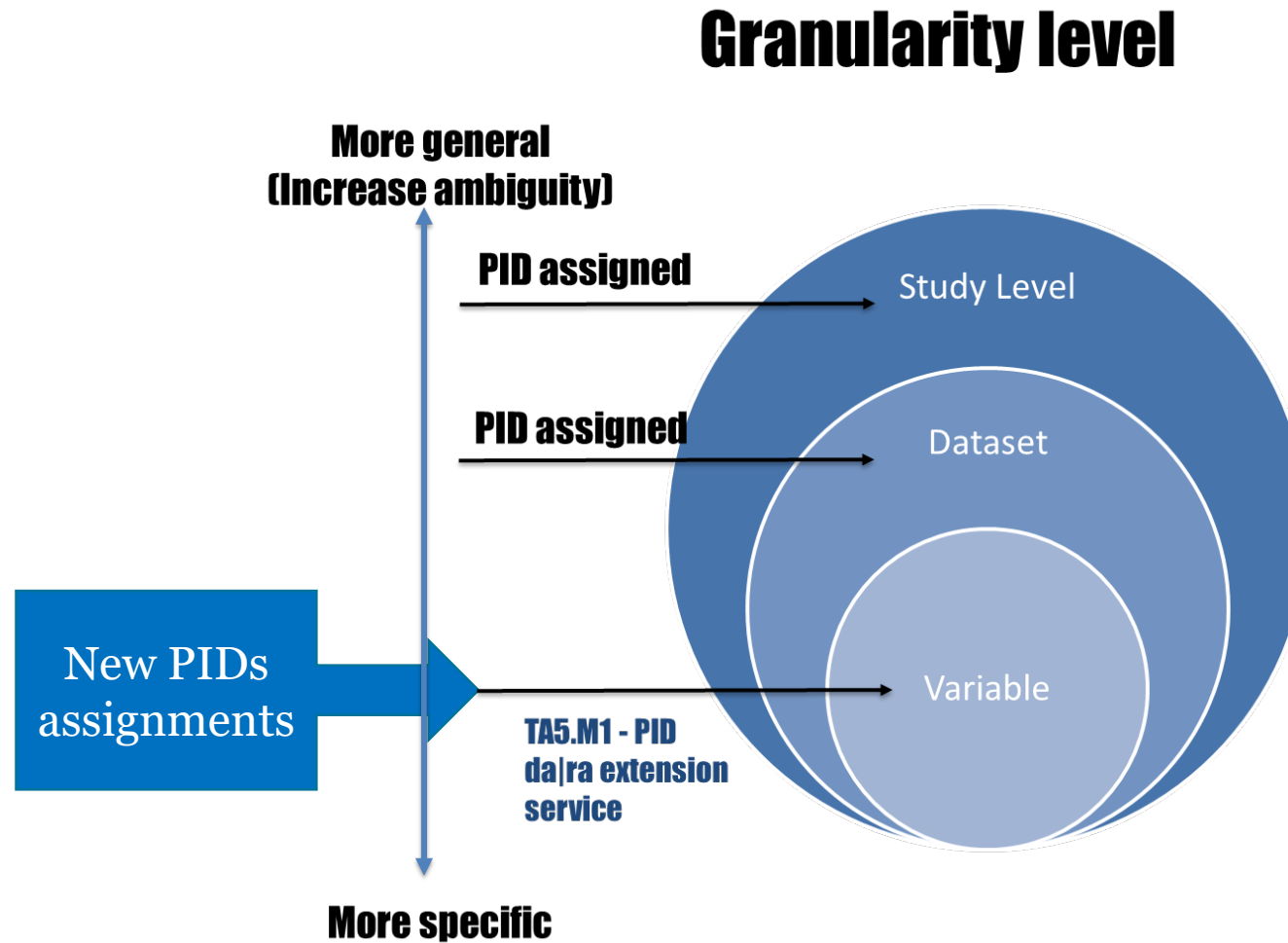


Agenda

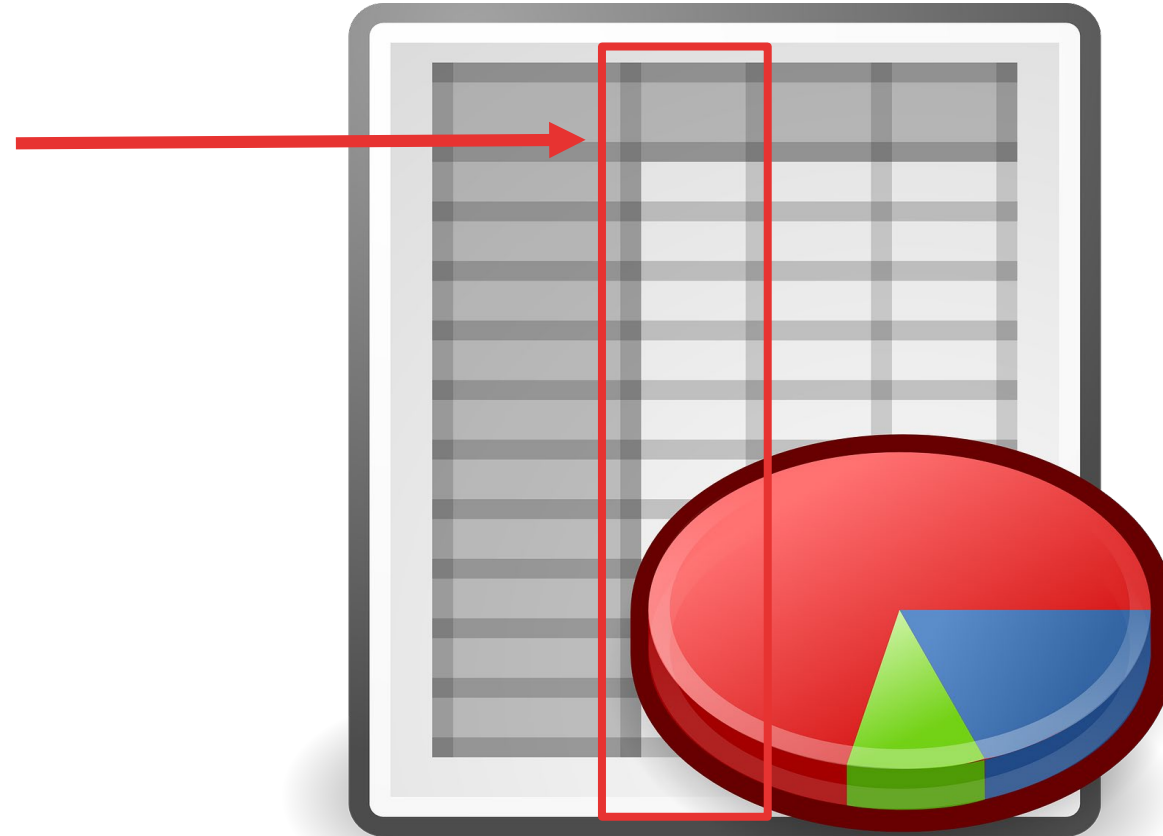
- The PID Registration service for variables
 - Goal
 - The Research data granularity levels
 - Data formats: initial approach and future use
 - PID registration service provider
 - Use cases

- Identify survey variables, **using one identifier**- the PID - will simplify FAIR data management to:
 - to boost subsequent citation,
 - get direct (meta)-data access, and
 - promote data reuse.

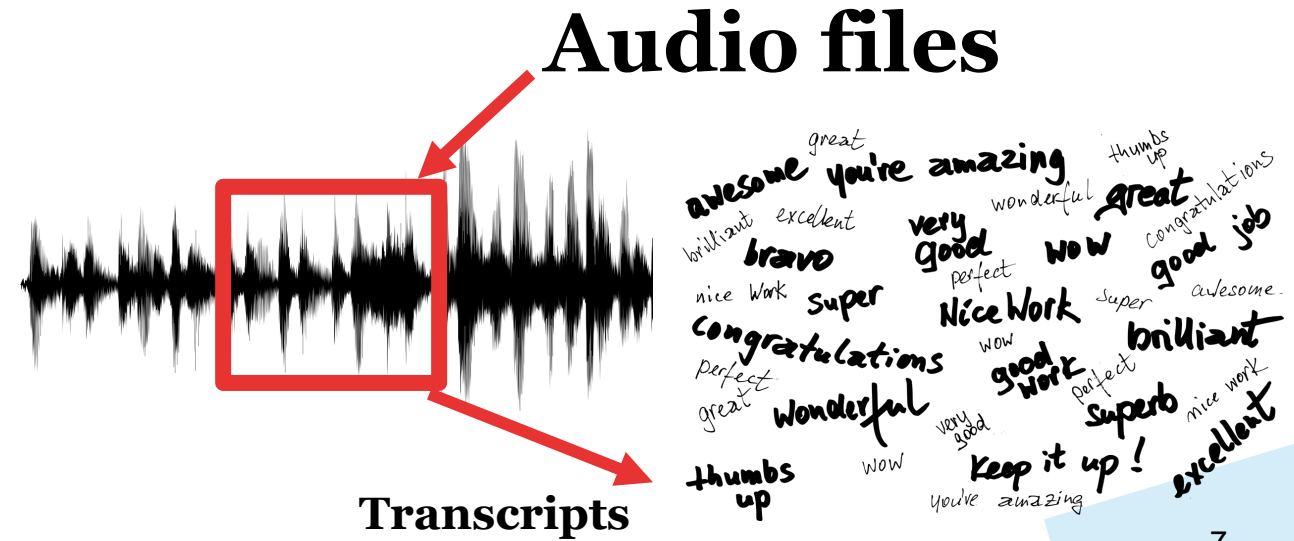
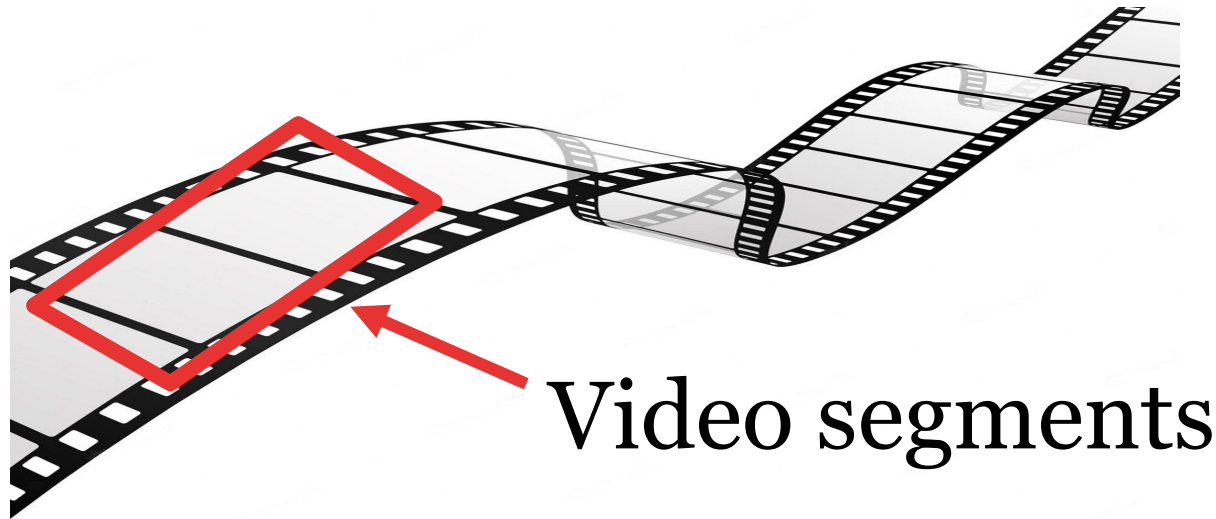




Variables in
tabular data
format



Data formats possible in the future



- This is a **da|ra** service **widening** and assigns a PID with **Handle** standard (ePIC);
- The service will be upgraded to **handle PIDs on variable** level;

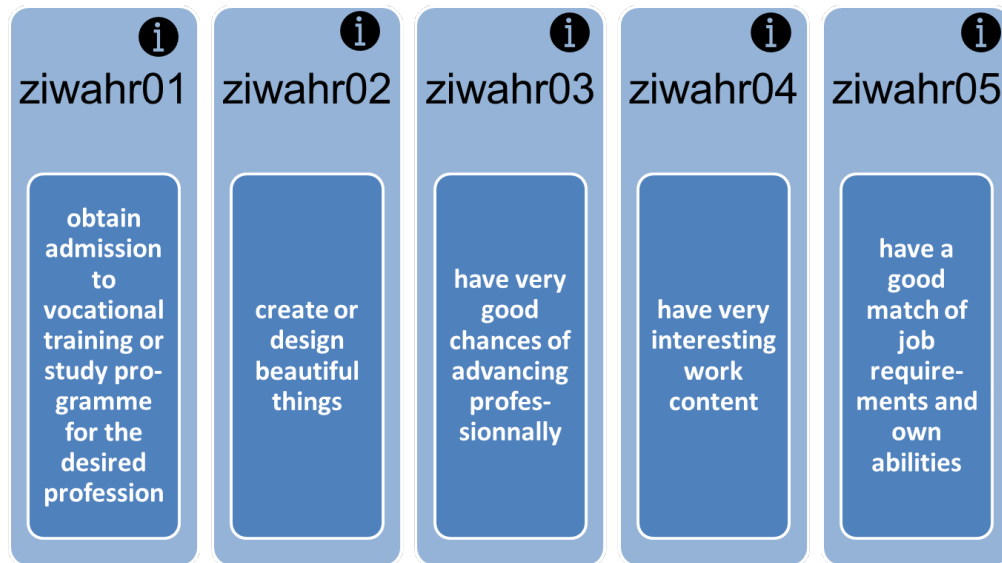
da|ra



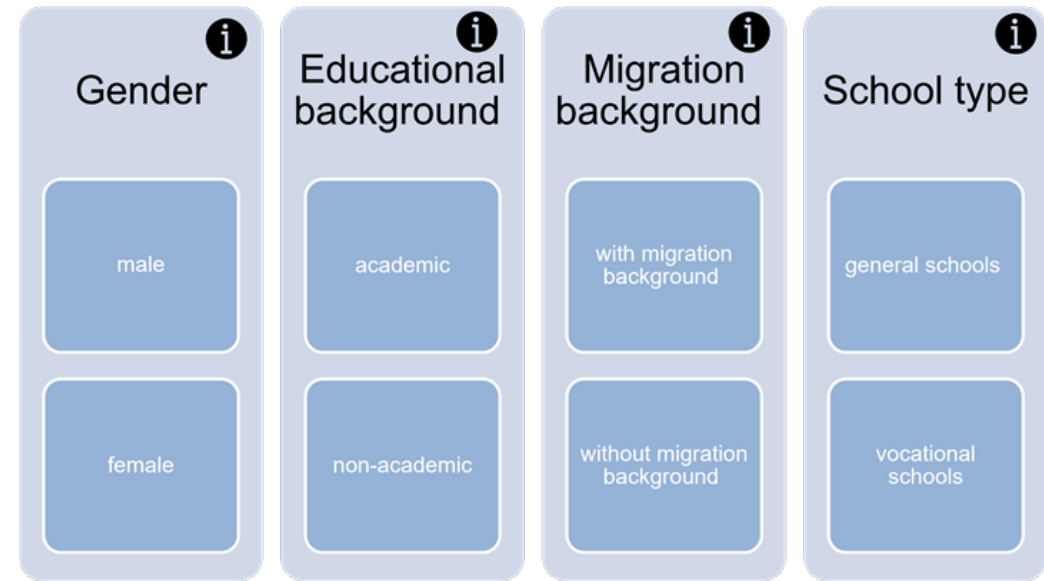


Higher Education Analytical Data System (HEADS) project at the DZHW needs a standard of data citation is to make its results widely usable and citable, particularly the entire information packages that comprise a central reporting variable (“indicator”) and the related multivariate analyses conducted in HEADS.

- PIDs for each variable, i.e., for each indicator or differentiation variables



 Persistent Identifier



 Persistent Identifier

DZHW

Deutsches Zentrum für Hochschul- und Wissenschaftsforschung

■ PIDs for information packages

Students with university entrance qualifications in 2015 half a year before graduation

Indicator: Career goals - probability of realization

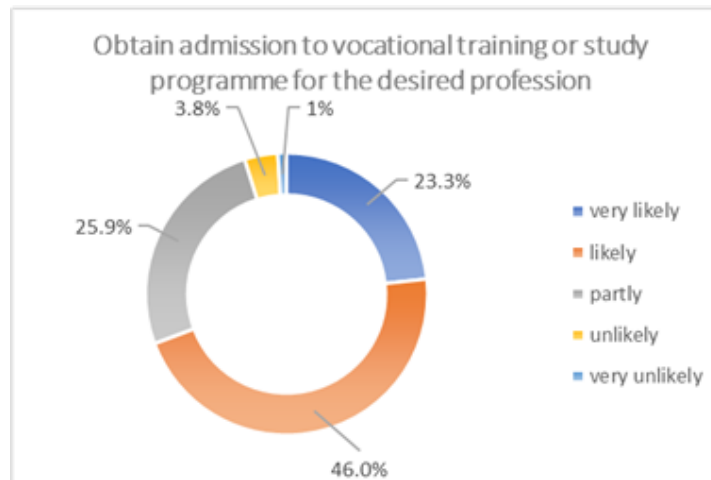
Question text: Everyone has goals. How likely do you think it is to achieve the following career goals?

Statement

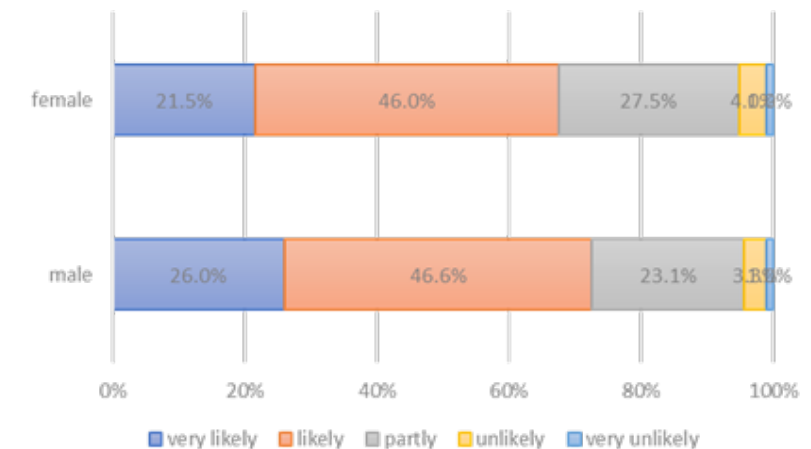
obtain admission to vocational training or study programme for the desired profession (total value)

Feature

Gender



Obtain admission to vocational training or study programme for the desired profession - by gender



Options for differentiation variables

Gender
Educational background
Migration background
School type



- PIDs for more than 500,000 variables from 6,500 national and international studies covering various topics in the Social Sciences, Economics, and Behaviour Sciences.



HaSpaD

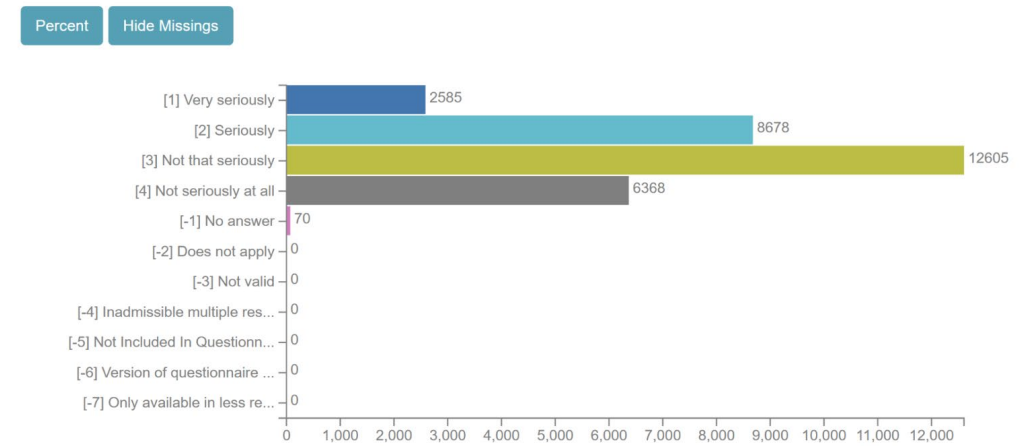
- PIDs for variables from harmonization tools and services
- Automatic access to variable data using scripts:
 - researchers are responsible for getting access to the datasets directly from the data providers;
 - With unique identifiers assigned, the data could be automatically accessed;
 - it makes it easier to use dozens of harmonised variables of the same topic from numerous diverse instruments.



Provide information on all household members: Germans living in the former eastern and western German states, foreign citizens, and immigrants residing in Germany. Some topics include household composition, occupational biographies, employment, earnings, health and satisfaction indicators.

- PIDs for 101.574 variables
 - available from 560 datasets,
 - distributed in 21.280 questions, and
 - 309 instruments

☰ bip/bip_171: Interest in Politics



Variable graph: *bip/bip_171* Interest in Politics



Qualiservice consists primarily of qualitative interview transcripts and context data in text, videos, and description data.

- PIDs are assigned at the file level for disambiguating similar data types and file naming;
- Provides a direct way of citing, identifying, and getting the target file.

Janete Saldanha Bach, Claus-Peter Klas and Peter Mutschke. 2023. KonsortSWD PID registration Service. In Workshop on PIDs within NFDI. *NFDI4 PID Working Group*, 27 Jan. 2023. 16 slides.





PID Service report
<https://doi.org/10.5281/zenodo.6397367>

Service demo application
<https://youtu.be/fm8T-hlhsXg>

The service is part of KonsortSWD project deliverable, NFDI funding number 442494171



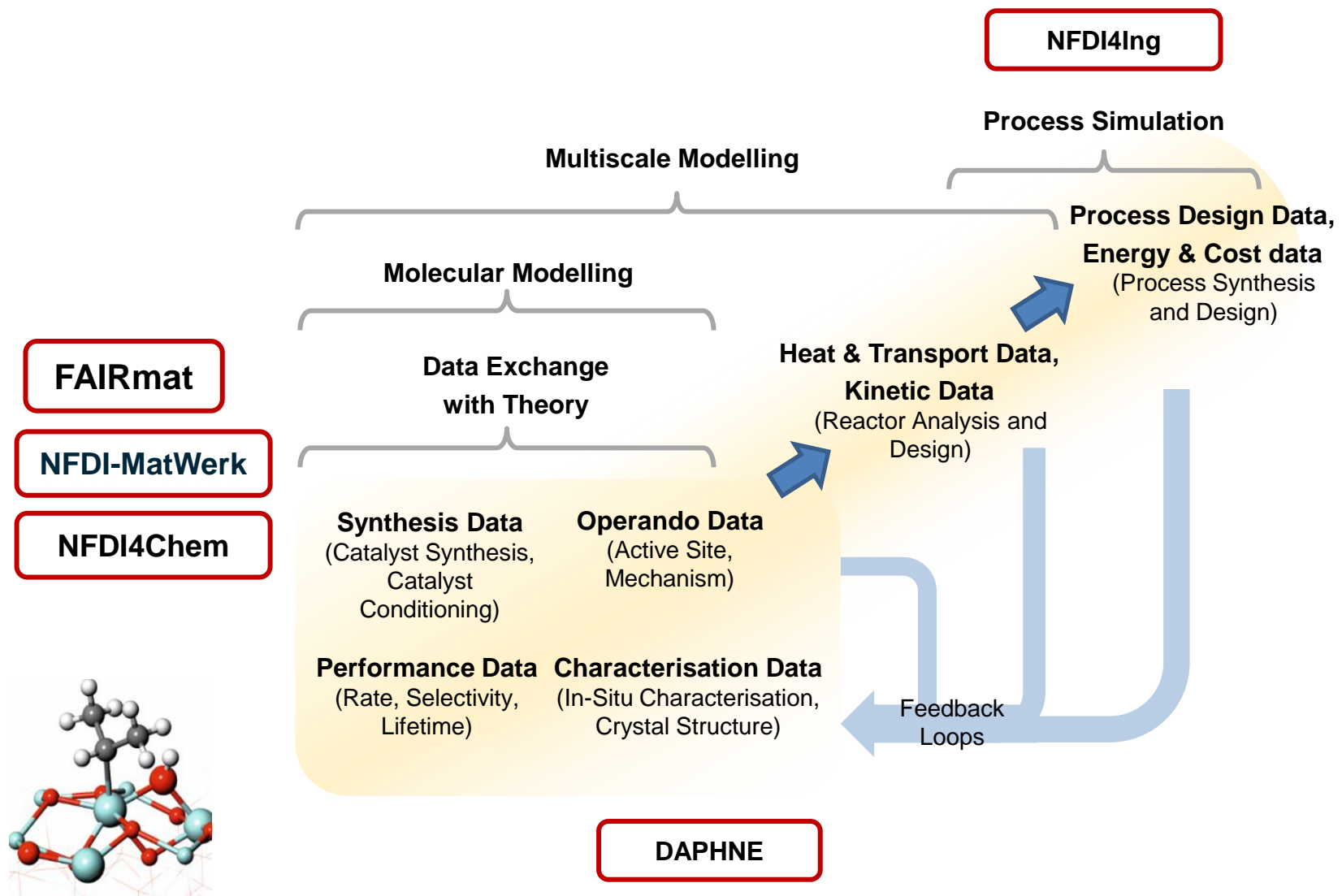
PIDs for Catalysis (NFDI4Cat)

2023-01-27, David Linke (LIKAT)





NFDI4Cat's goal: A Data Value Chain in Catalysis



Syngas Plant
Linde Engineering





- PIDs are an important part of FAIR principles
 - » F1. (Meta)data are assigned a globally unique and **persistent identifier**
 - » A1. (Meta)data are **retrievable by their identifier** using a standardized communication protocol
 - » I3. (Meta)data include **qualified references** to other (meta)data
 - » R1. (Meta)data are richly described with a plurality of accurate and relevant attributes
 - R1.2. (Meta)data are associated with detailed **provenance**
- Stable links to resources help to address the “reproducibility crisis” in science
- Links between data and other resources get more and more important to advance science.
- Long term preservation (findability) of the links to scientific resources must be assured.
 - » Identifying resources only by its url in a storage system is too fragile.
 - » Storage systems (“repositories”) will evolve/change over time.
- **Stable references (PIDs) are needed that are resolvable via a commonly agreed process.**

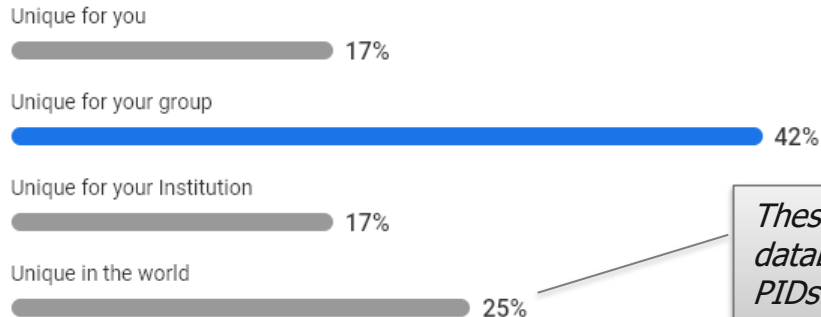


Online-Survey on PIDs from NFDI4Cat Consortium Meeting

Libriz

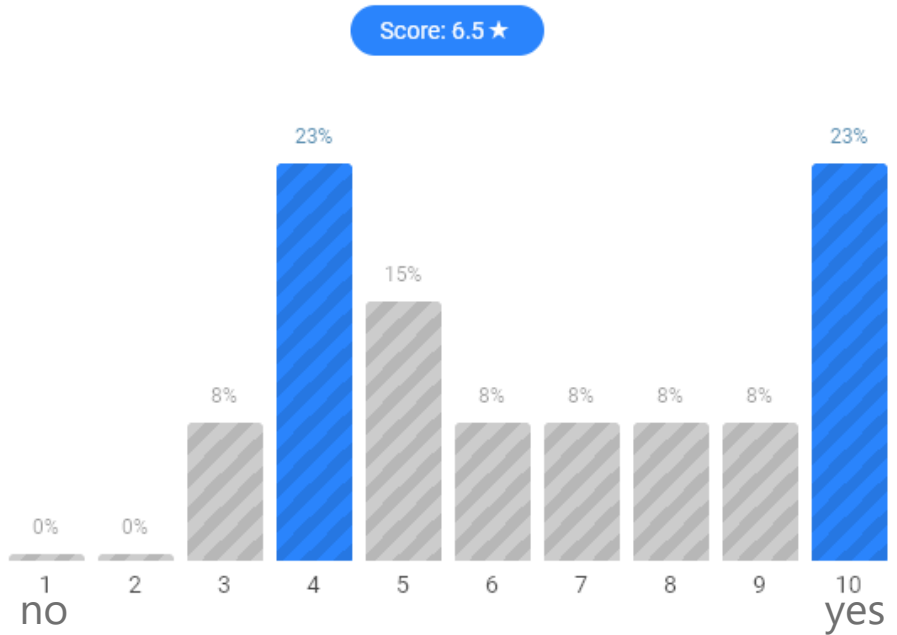
Survey from Feb-2022

How unique are the IDs you use?

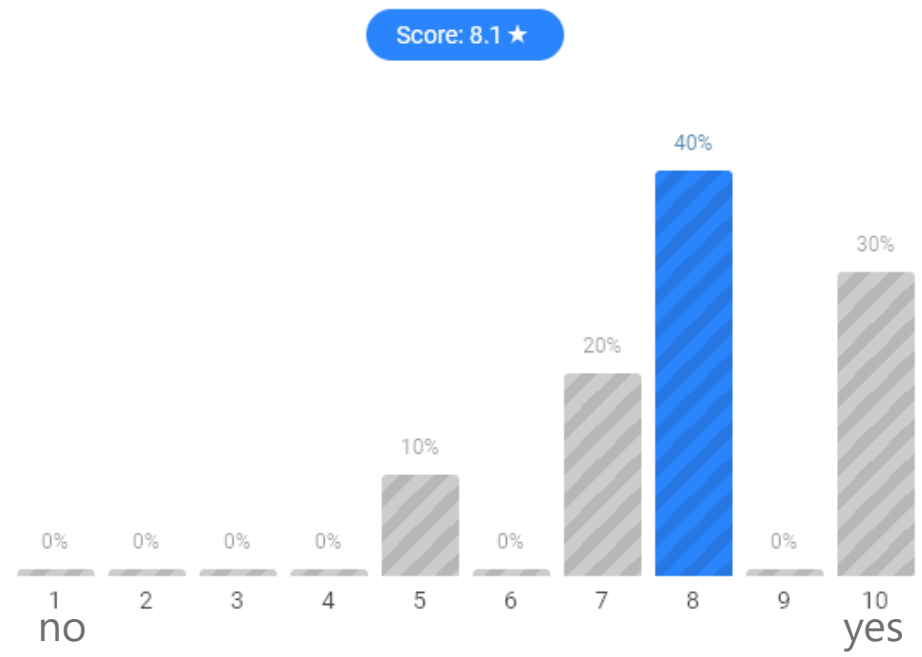


These are UUID4 database keys, not PIDs.

Will the sample IDs and all information associated with the IDs remain accessible and findable after you leave?



Would you switch to using Catalyst Ressource IDs if you can create and access them easily?



Are you interested in being part of a working group that promotes this topic?



=> As a result, a working group on PIDs was formed in March 2022.

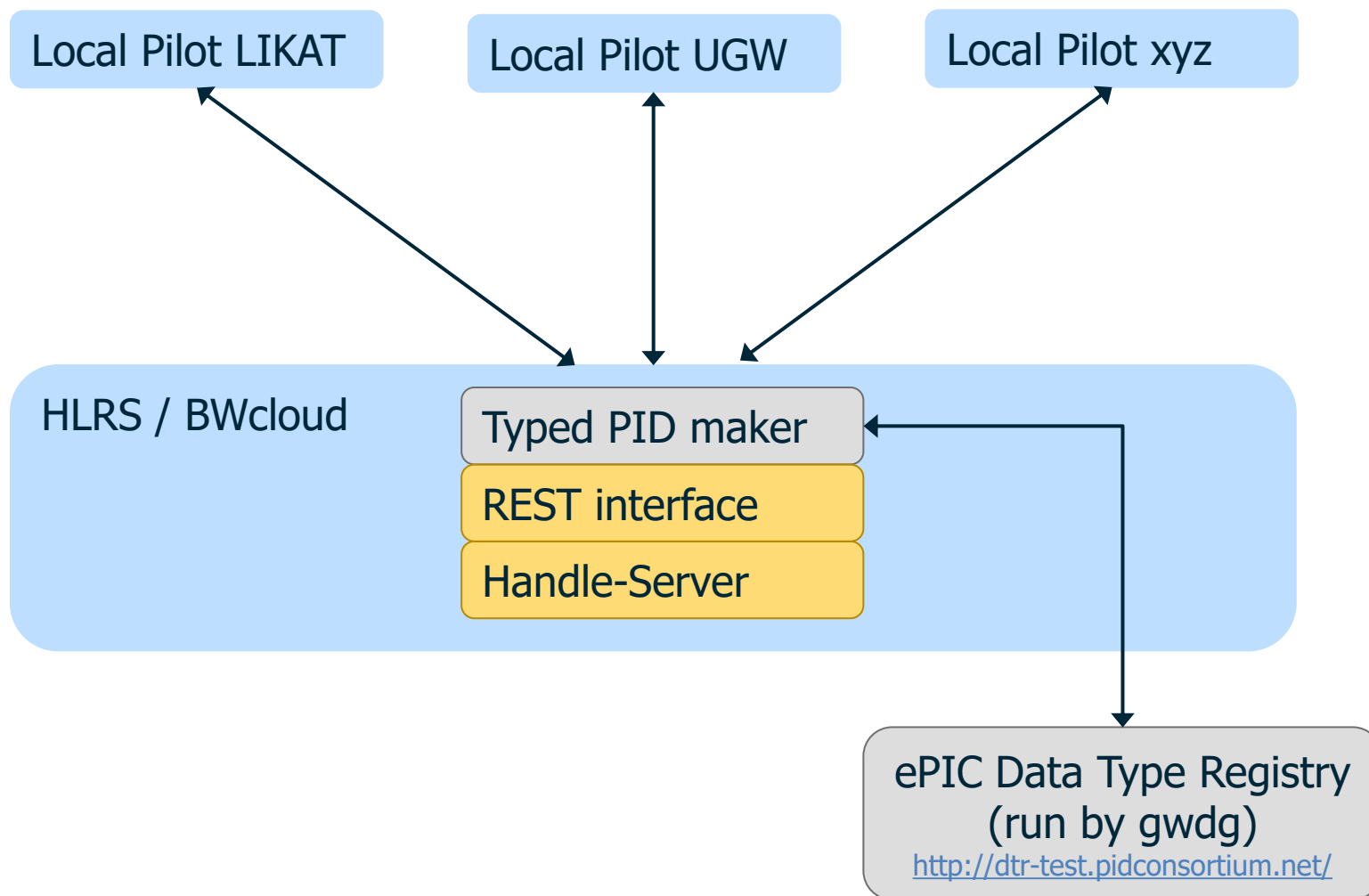


- There are already well established systems that should be used:
 - » **DOIs** for publications, software, data collections
 - » **ORCID**s for identifying researchers
 - » **ROR**s for identifying institutions (e.g. <https://ror.org/029hg0311>)
- **There is no established PID system for other resources like samples, materials, devices, instruments, models etc.** which
 - » supports **assigning PIDs in the early research phase** and can later be cited,
 - » can be adjusted (by NFDI4Cat) to the metadata requirements for catalysis resources,
 - » can cope with metadata requirements that differ between the resources,
 - » has a low barrier of entry,
 - » and low costs per PID.
- **IGSN & RRID cannot cover catalysis needs** but are excellent examples to learn from.



Proposed Solution: handle-system-based PIDs

Libriz



Several different institutional RDM systems (e.g. ELNs) will get PIDs from our central handle server.

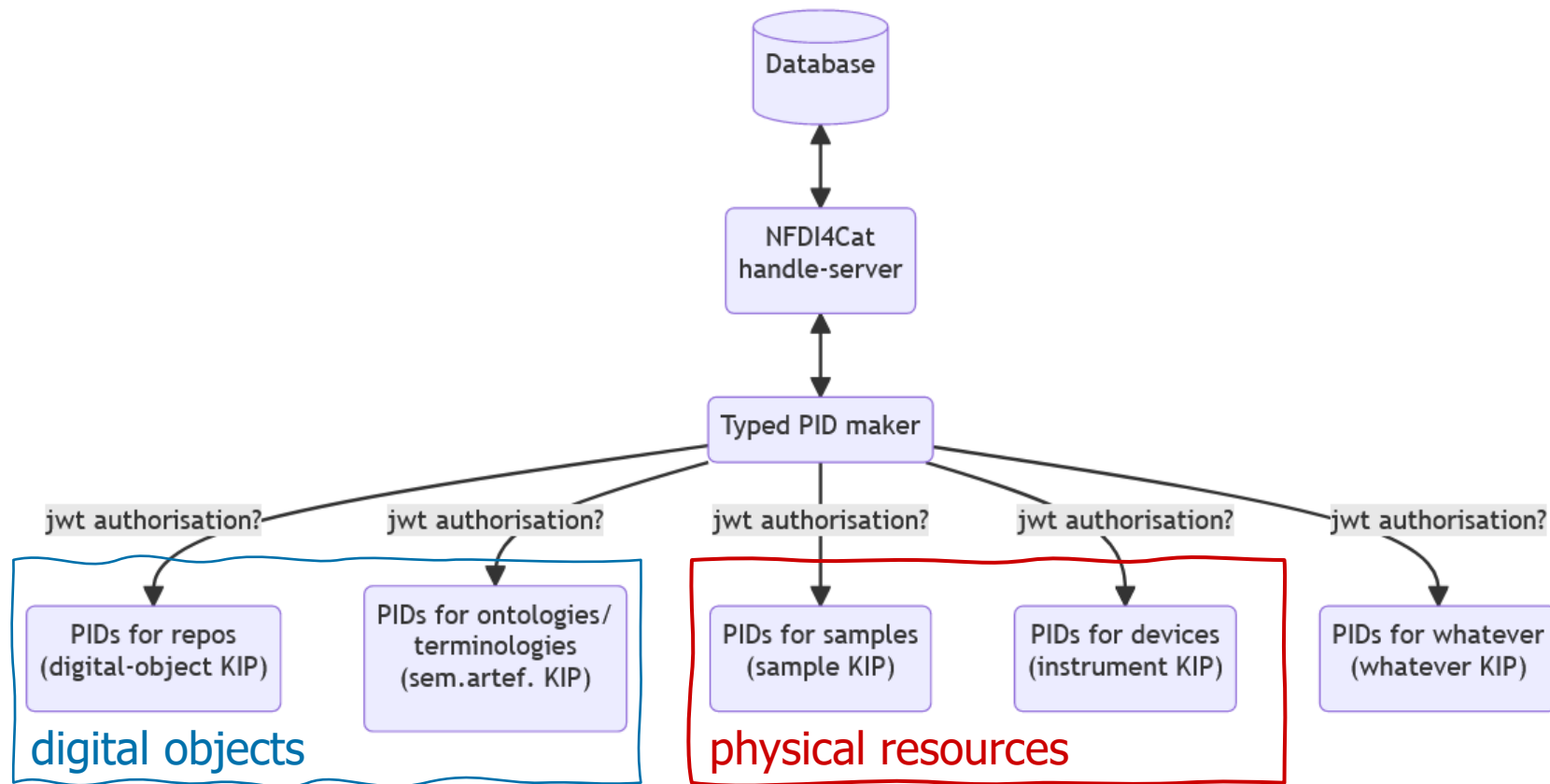
Researchers can use these PIDs from the first lab book entry to final publication.

We will test **Typed PID Maker** from KIT/SCC for

- PID schema validation
- Authorization

Fall-back alternative:

Add fastapi REST interface to B2Handle/pyHandle (from ePIC)



Open: How many different types of PIDs do we want/need?

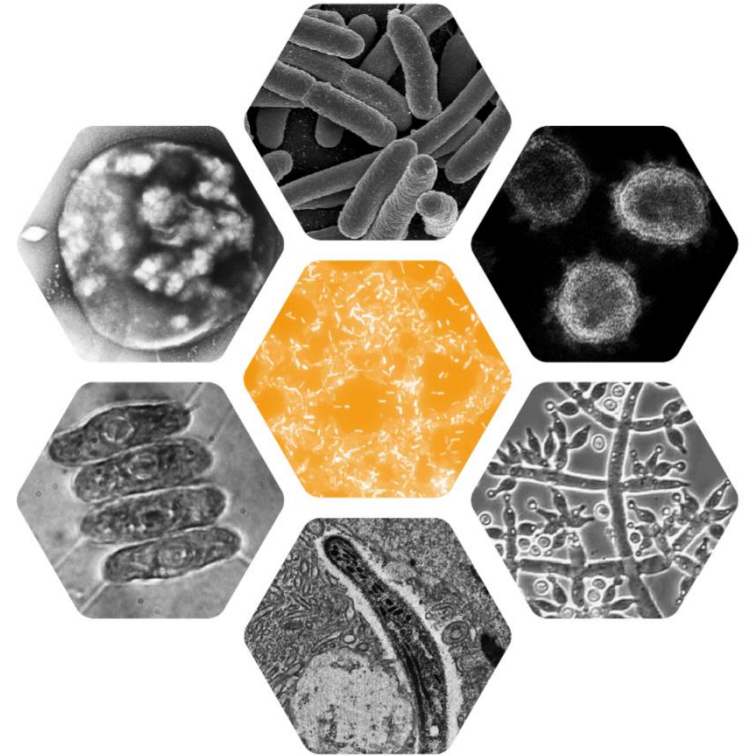
It is probably sensible to support at least two: one for physical resources & one for digital objects.



- In principle PIDs could be used instead of urls.
 - » Idea was discussed with some participants in FAIR-impact event & members of RDA Fair-Digital Objects group
 - Idea is not new (implemented in a French biomedical project).
 - Most(?) experts consider this as out-of-scope use of PIDs.

- Current plan: use **w3id.org** as provider for permanent urls.
 - » <https://w3id.org/nfdi4cat/> has been registered.
 - » We are going to use IDs as suffix-parts of the url.
 - e.g. for the photo-catalysis vocabulary
<https://w3id.org/nfdi4cat/voc/photocatalysis/concept12345>

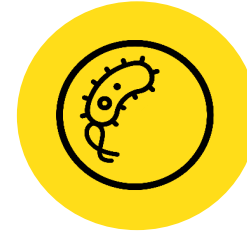
StrainInfo: PIDs for microbial strains



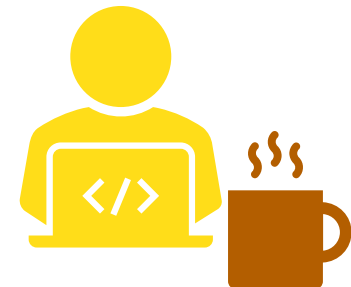
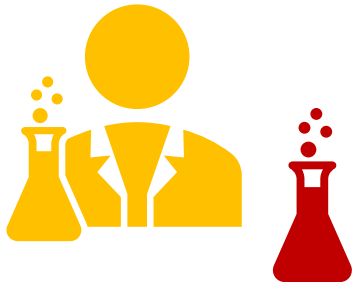
Version 2023-01-27

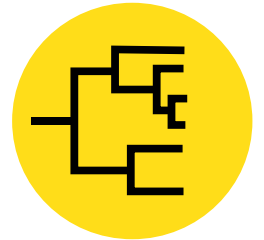
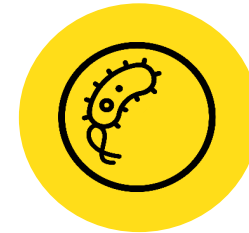
Lorenz Reimer
DSMZ

NFDI4 
MICROBIOTA



 Find a microbial strain and information about it?





 *Streptomyces hiroshimensis*

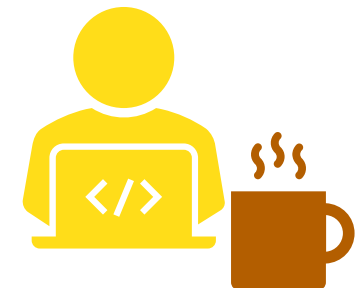
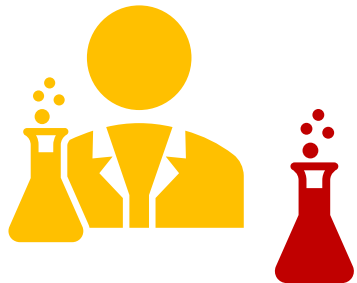
 *Actinomyces biverticillatus*

 *Streptomyces fervens*

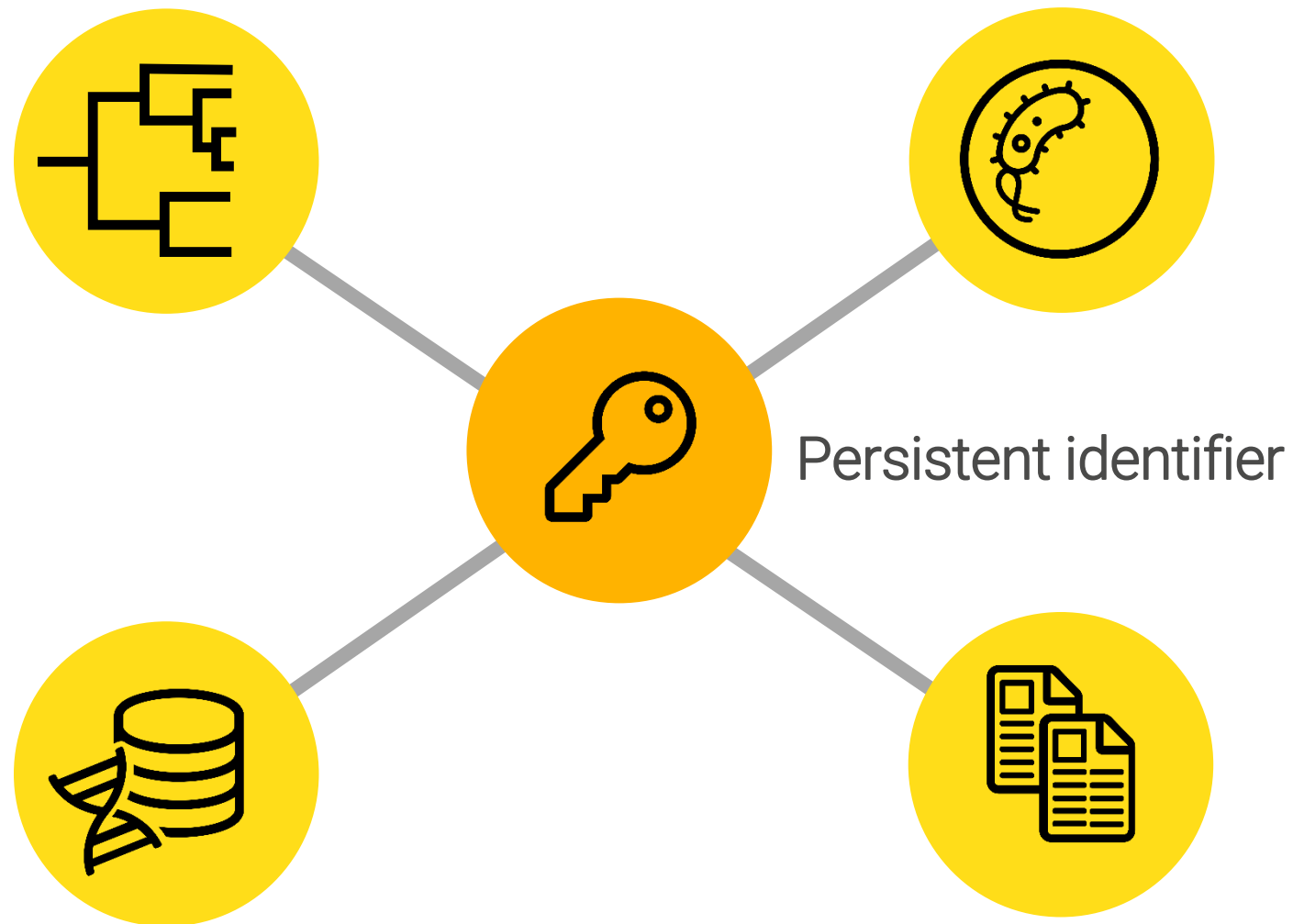
 *Streptomyces spitsbergensis*

 *Streptoverticillium salmonicida*

 *Streptomyces fervens subsp. melrosporus*



How to connect distributed data?





Home > Microbe Products > Bacteriology and Archaea > Anaerobes > 15703

Bifidobacterium adolescentis Reuter

15703™

📄 DOWNLOAD GENOME

LEARN ABOUT OUR ENHANCED AUTHENTICATION INITIATIVE >

Bifidobacterium adolescentis strain E194a (Variant a) is a whole-genome sequenced type strain that was isolated from the intestine of an adult. It has applications in bioinformatics and is a microbiome standard component.

🌟 96/100 Bioz Stars [229 Product Citations](#)

Product category	Bacteria
Product type	Anaerobe
Strain designation	E194a (Variant a)
Type strain	Yes
Genome sequenced strain	Yes
Isolation source	Intestine of adult
Applications	Bioinformatics
Product format	Freeze-dried
Storage conditions	2°C to 8°C

SEE ADDITIONAL PRODUCT INFORMATION >

<https://www.atcc.org/products/15703>

JCM Catalogue

Bifidobacterium adolescentis Reuter 1963

Taxonomy in NCBI database: [Bacteria](#); [Actinobacteria](#); [Actinomycetia](#); [Bifidobacteriales](#); [Bifidobacteriaceae](#).

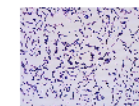
1275^T <-- ATCC 15703 <-- G. Reuter E194a (Variant a).
Accessioned in 1982.
=ATCC 15703 =BCRC 14606 =CCM 4987 =CCUG 17359 =CCUG 18363
=CCUG 45213 =CECT 5781 =CGMCC 1.2190 =CIP 64.59 =DSM 20083
=KCTC 3216 =LMG 10502 =NCTC 11814 =VTT E-981074.

Type strain [596].

Medium: [13](#); Temperature: 37°C; Anaerobic; Rehydration fluid: [663](#).



[open link in new window](#)



[open link in new window](#)

Source: Intestine of adult [110].

Biochemistry/Physiology: [3093,3126,4424].

Cell wall: Lys(Orn)-Asp [010,3126].

Enzyme electrophoretic profile: [3126,4425].

G+C (mol%): 58.9 (Tm) [181].

DNA-DNA relatedness: [023,3126].

Phylogeny: 16S rRNA (M58729), 16S rRNA gene (AB437354, LC071806), *hsp60* (AF210319) [5374].

DNA typing: RELP [4423], rRNA gene restriction pattern [4597], genome-wide tiling array [9039].

Other taxonomic data: Protein profile [1865].

Genome sequence: AP009256, UAQG00000000.

More information: Species-specific primer [4521].

NCBI Taxonomy ID: 1680.

Genomic DNA is available from RIKEN BRC-DNA Bank: JGD 07423.

Publication(s) using this strain [A04015, A04018, A04079, A04152, A04168, A04196, A05011, A05036, A05048, A06033, A07171, B07183, A08019, A08201, A08244, A08278, A08304, A09019, A09026, A09198, A09201, A10090, A10101, A10294, A11008, A11225, A11236, A11237, A11252, A11297, A12287, A12408, A12422, A12441, A13116, A13170, A13553, A14435, A15097, A15541, A16131, A17103, A17549, A18008, A18010, A18152, A18169, A18418, A18569, A19161, A19175, A19522, A19531, A20144, A20166, A20446, A20562, A21209, A21447, A21524, A21549].

Patent publication(s) using this strain [WO2007/020884, WO2007/114378, WO2007/099997, WO2007/023588, JP2007-112805A, JP2008-074768A, WO2008/062555, JP2009-125055A, JP2009-084215A, JP2009-242430A, WO2009/028253, WO2011/027875, WO2012/005240, JP2012-180288A, JP2012-180373A, JP2013-042749A, JP2014-055194A, 2015-198638, 2015-96555, 2016-028595, WO2016/103699, CN105671120, 2017-216979, 2019-94307, 2019-92469, 2020-132560, 2020-92704].

https://www.jcm.riken.jp/cgi-bin/jcm/jcm_number?JCM=1275



Open Access | Published: 21 April 2017

Bifidobacterium adolescentis (DSM 20083) and *Lactobacillus casei* (Lafti L26-DSL): Probiotics Able to Block the In Vitro Adherence of Rotavirus in MA104 Cells

Karem Prunella Fernandez-Duarte, Nury Nathalia Olaya-Galán, Sandra Patricia Salas-Cárdenas, Jazmin Lopez-Rozo & I

SpringerLink

Probiotics and I

2806 Accesses

Original Paper | Published: 22 January 2004

Abstract

Rotavirus is the most common cause of gastroenteritis in young children. Even though there are several strategies for its prevention, they are constantly being updated in order to demonstrate their effectiveness in rotavirus infection. In MA104 cells, the treatment with probiotics showed

Physico-chemical and transglucosylation properties of recombinant sucrose phosphorylase from *Bifidobacterium adolescentis* DSM20083

L. A. M. van den Broek, E. L. van Boxtel, R. P. Kievit, R. Verhoef, G. Beldman & A. G. J. Voragen

Applied Microbiology and Biotechnology 65, 219–227 (2004) | Cite this article

1013 Accesses | 60 Citations | 6 Altmetric | Metrics

Abstract

Clones of a genomic library of *Bifidobacterium adolescentis* were grown in minimal medium with sucrose as sole carbon source. An enzymatic fructose dehydrogenase assay was used to identify sucrose-degrading enzymes. Plasmids were isolated from the positive colonies and sequence analysis revealed that two types of insert were present, which only differed with respect to their orientation in the plasmid. An open reading frame of 1,515 nucleotides with high homology for sucrose phosphorylases was detected on these inserts. The gene was designated *SucP* and encoded a protein of 56,189 Da. *SucP* was heterologously expressed in *Escherichia coli*, purified, and characterized. The molecular mass of *SucP* was 58 kDa, as estimated by SDS-PAGE, while 129 kDa was found with gel permeation, suggesting that the native enzyme was a dimer. The enzyme showed high activity towards sucrose and a lower extent towards α -glucose-1-phosphate. The transglucosylation properties were investigated

<https://link.springer.com/article/10.1007/s12602-017-9277-7>

<https://link.springer.com/article/10.1007/s00253-003-1534-x>

ORIGINAL RESEARCH article

Front. Microbiol. 08 April 2022
Sec. Food Microbiology
<https://doi.org/10.3389/fmicb.2022.860014>

Study on the Biochemical Characterization and Selectivity of Three β -Glucosidases From *Bifidobacterium adolescentis* ATCC15703

Yanbo Hu¹, Liy

¹ School of Food Science and Engineering
² School of Life Sciences, Changchun

Three β -glucosidases from *Bifidobacterium adolescentis* ATCC 15703 were overexpressed in *Escherichia coli* and purified. Affinity chromatography, SDS-PAGE, and specific activity assays suggested that BaBg1A, BaBg1B, and BaBg1C were β -glucopyranoside (β -D-glucopyranoside) (β -D-GlcPyran) β -1,3-glucosidase and β -D-GlcPyran β -1,3-glucosidase and β -D-GlcPyran β -1,3-glucosidase, respectively. The highest bioconversion activity was observed with BaBg1A, which was further studied by HPLC and HPLC-MS. BaBg1A was the most active enzyme, and it hydrolyzed both β -D-GlcPyran and β -D-GlcPyran into the rare ginsenosid Rg_1 and Rg_2 , where it hydrolyzed both β -D-GlcPyran and β -D-GlcPyran. BaBg1A was not active on Rb1a

<https://www.frontiersin.org/articles/10.3389/fmicb.2022.860014/full>

Food Science & Nutrition

Open Access

ORIGINAL RESEARCH | Open Access | CC BY

Combining of transcriptome and metabolome analyses for understanding the utilization and metabolic pathways of Xylo-oligosaccharide in *Bifidobacterium adolescentis* ATCC 15703

Jian Yang, Qilong Tang, Lei Xu, Zhijiang Li, Yongqiang Ma, Di Yao

First published: 30 September 2019 | <https://doi.org/10.1002/fsn3.1194> | Citations: 9

SECTIONS

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Abstract

A combination of transcriptome and metabolome analyses was applied to understand the utilization and metabolism of Xylo-oligosaccharide (XOS) in *Bifidobacterium adolescentis* 15703 as well as identifying the key regulatory-related genes and metabolites. Samples of cultures grown on either XOS or xylose were collected. The transcript and metabolite profiles were obtained from high-throughput RNA-sequencing data analysis and UHPLC system. Compared with xylose, XOS highly promoted the

<https://onlinelibrary.wiley.com/doi/10.1002/fsn3.1194>



Datasets Taxonomy **Genome** Gene Command-line tools Documentation

Datasets / Genome / ASM1042v1

Genome assembly ASM1042v1

[Download](#) [datasets](#) [curl](#)

Reference sequence	RefSeq GCF_000010425.1
Submitted sequence	GenBank GCA_000010425.1
Taxon	Bifidobacterium adolescentis ATCC 15703
Strain	ATCC 15703
Submitter	Gifu University, Life Science Research Center, Japan
Date	Dec 5, 2006

View the [legacy Assembly page](#)

Assembly statistics

These statistics describe the RefSeq genome sequence GCF_000010425.1

Genome size	2.1 Mb
Number of chromosomes	1
Number of scaffolds	1
GC percent	59
Assembly level	Complete Genome

https://www.ncbi.nlm.nih.gov/data-hub/genome/GCF_000010425.1/

Datasets Taxonomy **Genome** Gene Command-line tools Documentation

Datasets / Genome / 49964_F01

Genome assembly 49964_F01

[Download](#) [datasets](#) [curl](#)

Reference sequence	RefSeq GCF_900445615.1
Submitted sequence	GenBank GCA_900445615.1
Taxon	Bifidobacterium adolescentis
Strain	NCTC11814
WGS project	UAQG01
Submitter	SC
Date	Jun 28, 2018

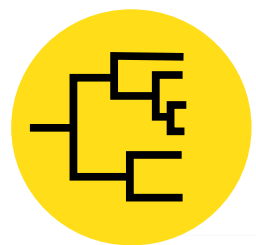
View the [legacy Assembly page](#)

Assembly statistics

These statistics describe the RefSeq genome sequence GCF_900445615.1

Genome size	2.2 Mb
Number of contigs	57
Contig N50	238.5 kb
Contig L50	3
GC percent	59
Assembly level	Contig

https://www.ncbi.nlm.nih.gov/data-hub/genome/GCF_900445615.1/



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Species *Bifidobacterium adolescentis*

- ① **Name:** *Bifidobacterium adolescentis* Reuter 1963 (Approved Lists 1980)
- ① **Category:** Species
- ① **Proposed as:** sp. nov.
- ① **Etymology:** ad.o.le.scen.tis. L. n. *adolescens* -tis, adolescent; L. gen. masc
- ① **Gender:** neuter
- ① **Type strain:** [ATCC 15703](#); [BCRC 14606](#); [CCRC 14606](#); [CCUG 17359](#); [CCUG 17359](#)
- ① See detailed strain information at [BacDive](#)
- ① Conduct genome-based taxonomy at [TYGS](#)
- ① **16S rRNA gene:** M58729 [Analyse](#) [FASTA](#) [ENA](#) [SINA](#)
- ① **Effective publication:** Reuter G. Vergleichenden Untersuchung über die Bifidobakterien. *Infektionskrankheiten und Hygiene. Abteilung I* 1963; **191**:486-507.
- ① **IJSEM list:** Skerman VBD, McGowan V, Sneath PHA. Approved lists of bacteria
- ① **Nomenclatural status:** validly published under the ICNP
- ① **Taxonomic status:** correct name
- ① **Risk group:** 1
- ① ▼ **Synonyms:**

Name	Kind
Bifidobacterium stercoris Kim et al. 2010	heterotypic synonym, validly

Entrez
PubMed
Nucleotide
Protein
Genome
Structure

Search for as complete name lock

Display levels using filter: none

Bifidobacterium adolescentis ATCC 15703

Taxonomy ID: 367928 (for references in articles please use NCBI:txid367928)

current name
Bifidobacterium adolescentis ATCC 15703
 equivalent: [Bifidobacterium adolescentis str. ATCC 15703](#)

NCBI BLAST name: **high G+C Gram-positive bacteria**

Rank: **strain**

Genetic code: [Translation table 11 \(Bacterial, Archaeal and Plant Plastid\)](#)

Lineage (*full*)

[cellular organisms](#); [Bacteria](#); [Terrabacteria group](#); [Actinobacteria](#); [Actinomycetia](#); [Bifidobacteriales](#); [Bifidobacteriaceae](#); [Bifidobacterium](#); [Bifidobacterium adolescentis](#)

Comments and References:

genome sequence
Determination of the DNA genome sequence of this strain has been or is being determined either in whole or in part.

<https://www.ncbi.nlm.nih.gov/data-hub/taxonomy/1680/>

How to connect distributed data?



Persistent identifier:

- Culture collection identifier
- INSDC sequence accession no.
- DOIs (mostly in literature)

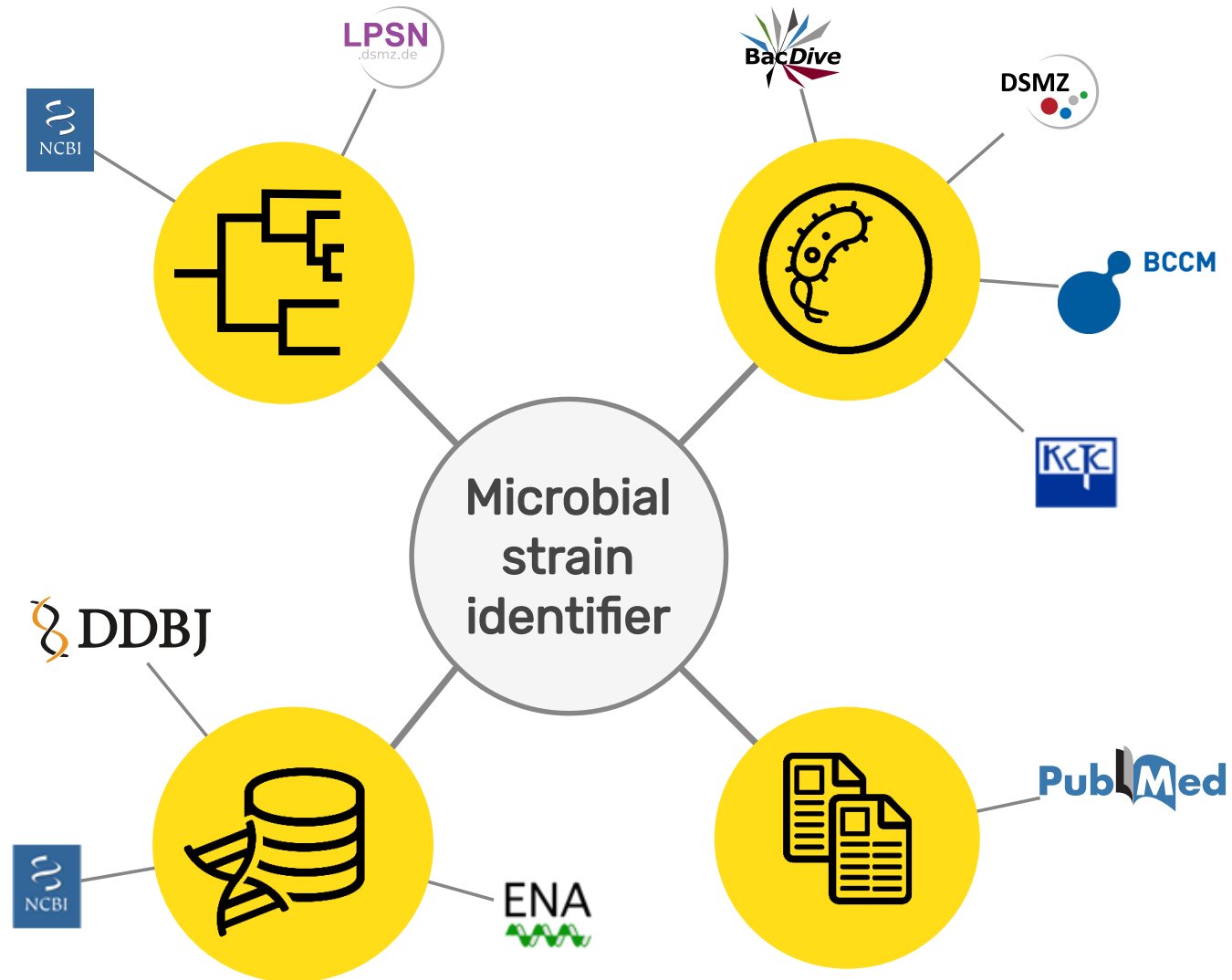


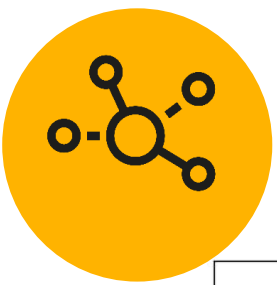
Problem to solve:

- Collect and connect them!
- Build a central database/registry

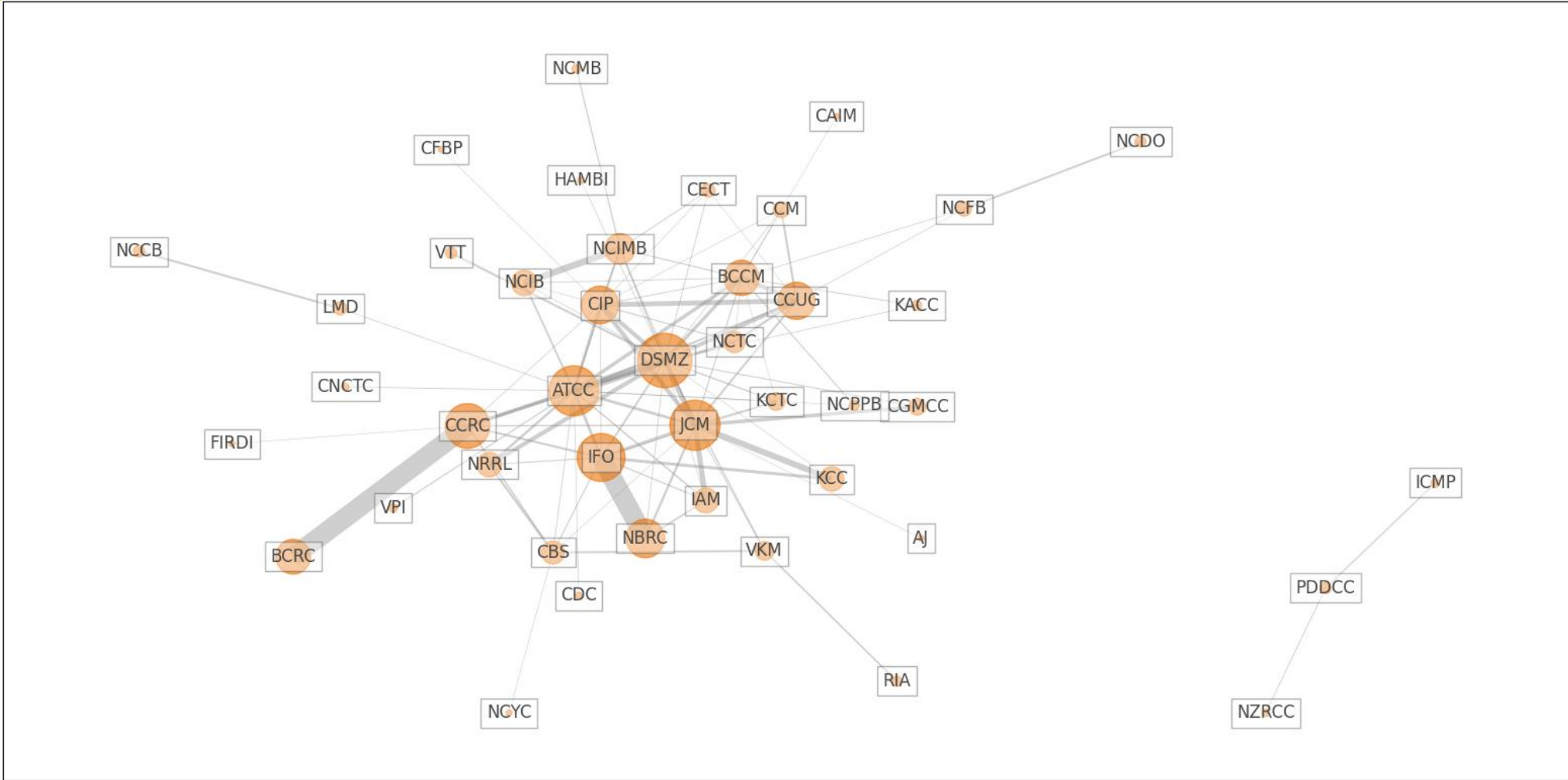


Use relations to connect data





Reveal the relations





Provide a Webinterface & an API

StrainInfo ARCHIVE

HOME > PASS

Culture ID: 5339 | Strain ID: 1234 | Type strain: ✓ | Species: Bifidobacterium adolescentis | Seq. acc. no.: F124596, ...

Related cultures

Parent
ATCC 15703

Children
CCUG 18363 | NCTC 11814 | NCAIM B.01822 | VTT E-981074

Strain

LMG 10502	ATCC 15703	CCTM 3063	CCUG 17359	CCUG 18363	CUETM 89-14	JCM 1275	Kilian AK 4
LMG 11036	LMG 11037	NCFB 2204	NCTC 11814	Reuter E194a	LMG 11037.1	E194a (Varian...	AK4
E194a	CCUG 45213	NCDO 2204	CCRC 14606	NCIMB 702204	CIP 64.59	LMG 11036 t2T	LMG 11036 t1T
LMG 11037 t2T	ATCC 15703 b...	LMG 10502TQ...	LMG 10502QC...	KCTC 3216	AS 1.2190	CCM 4987	BF32,
E 194A	G. Reuter E19...	CECT 5781	strain AS 1.21...	BO			
CGMCC 1.2190	VTT E-981074	VTT E-001750	E-981074				

Sequences

accession number	culture u	description	seq. length
AB116269	JCM 1275	Bifidobacterium adolescentis gene for 16S rRNA, partial sequence, strain: JCM 1275	480
AB198732	ATCC 15703	Bifidobacterium adolescentis rpoB gene for RNA polymerase beta chain, partial cds, strain: ATCC 15703	129
AB198733	ATCC 15703	Bifidobacterium adolescentis rpoB gene for RNA polymerase beta chain, partial cds, strain: ATCC 15703, rifaximin resistant clone	129
AF124596	DSM 20083	Bifidobacterium adolescentis alpha-galactosidase (aga) gene, complete cds	2403
AF210319	JCM 1275	Bifidobacterium adolescentis heat shock protein 60 (hsp60) gene, completecds	2031
AF261676	ATCC 15703	Bifidobacterium adolescentis strain ATCC 15703 L-lactate dehydrogenasegene, partial cds	312
AF275881	E-981074	Bifidobacterium adolescentis clone nru-1 16S ribosomal RNA gene, partialsequence	1451
AF275882	E-981074	Bifidobacterium adolescentis clone nru-5 16S ribosomal RNA gene, partialsequence	1448
AF358444	DSM 20083	Bifidobacterium adolescentis alpha-glucosidase (aglA) gene, partial cds	2043
AF384979	CIP 64.59	Bifidobacterium adolescentis pyruvate kinase gene, partial cds	331

Publications

title	authors	journal	year
Gram-positive bacteria with a high DNA G+C content are characterized by a common insertion within their 23S rRNA genes	Roller C, Ludwig W, Schleifer KH	J Gen Microbiol	1992
Numerical classification of Streptomyces and related genera	Williams ST, Goodfellow M, Alderson G, Wellington EM, Sneath PH, Sackin MJ	J Gen Microbiol	1983
Specific PCR and	Satokari RM, Vaughan EE, Akkermans AD, Saarela M, de Vos WM	Appl Environ Microbiol	2001
terium based on	Jian W, Zhu L, Dong X	Int J Syst Evol Microbiol	2001
terium species by	Requena T, Burton J, Matsuki T, Munro K, Simon MA, Tanaka R, Watanabe K, Tannock GW	Appl Environ Microbiol	2002

Visit <https://straininfo.dsmz.de/>

Thank you!



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<https://straininfo.dsmz.de/>

www.nfdi4microbiota.de
[@nfdi4microbiota](https://twitter.com/nfdi4microbiota)



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27th January 2023 | Workshop on PIDs within NFDI

PIDs in NFDI4Culture

Sandra Göller | FIZ Karlsruhe
Desiree Mayer | SLUB Dresden

DFG Deutsche
Forschungsgemeinschaft

Gefördert durch die Deutsche Forschungsgemeinschaft (DFG) - 441958017

Overview:

- Which are the most important PIDs?
- Authority Data as PIDs
- PIDs in RADAR4Culture
- Status quo in
NFDI4Culture/Desiderata/Discussions



Most important PIDs in NFDI4Culture

Highly sustainable and reliable
Controlled Standards from National
Libraries

Authority
Data, e.g.:



{ BnF



Highly sustainable and reliable
Open to public / community driven

Structured
Data, e.g.:



metadata

(research)-data

PIDs in Repositories:

- make relations between publications and research data clear, controlled fields in the meta data schema are necessary.
- PIDs in controlled fields also enable search within a repository and from outside

Usage of PIDs within NFDI4Culture Repositories – An Example with RADAR4Culture

[RADAR4Culture](#)

- publishing service for research data on tangible and intangible cultural assets
- provided by FIZ Karlsruhe - Leibniz Institute for Information Infrastructure






[RADAR Metadata Schema](#)

- based on the [DataCite metadata schema kernel 4.4](#)
- [Metadata Schema v9.1](#): 10 mandatory metadata fields (including 6 mandatory fields for the DOI registration), 13 optional metadata fields
- combination of free text fields and options for standardised or normed entries:
 - individuals (**ORCID**)
 - organisations (**ROR**)
 - funding organisations (**Crossref Funder Registry**)
 - keywords (**Gemeinsame Normdatei, GND**)

In Discussion

- GND-IDs as additional alternative for ORCID
- further authority data / vocabulary IDs for keywords
- vocabulary IDs for subject areas

Dataset: Example

RADAR Metadata	Content	Statistics	Technical Metadata
Creator/Author:	Göller, Sandra  https://orcid.org/0000-0003-4553-3671  FIZ Karlsruhe – Leibniz Institute for Information Infrastructure		
Title:	Example		
Keywords:	 Forschungsdaten		
Publishers:	 FIZ Karlsruhe – Leibniz Institute for Information Infrastructure		
Production year:	2023		
Subject areas:	Information Technology		
Resource type:	Dataset		
Publication year:	2023		
Rights holders:	 FIZ Karlsruhe – Leibniz Institute for Information Infrastructure		
Funding:	Deutsche Forschungsgemeinschaft		

Status quo NFDI4Culture:

- DOIs for publications; URN for digitizations (individual pages with permalinks)

Desiderata

- More controlled fields in metadata schemas of repositories should require PIDs
- More terms in subject headings or improved access to create subject headings in authority data (e.g. in GND)

Recent Discussions:

- use of multiple PIDs for individual objects
- permanent referencing of artworks and their components
- Transparency about possible PIDs in repositories
- handling of changes in metadata due to new research results
- Use of PIDs for different data types, e.g. PIDs for complex search queries (with timestamps)
- DOIs for Annotations or specific elements/ single entries in workcatalogues
- Stable URIs for vocabularies are required for mappings





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Thank you!

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FAIRragro

UseCase



“PID for digital twins in plant genetic resources”

within NFDI FAIRragro

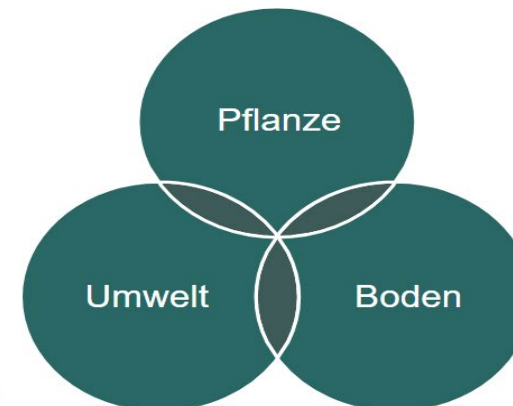
Daniel Arend, Matthias Lange
Leibniz Institute of Plant Genetics and Crop Plant Research

FAIRe Daten für die Agrosystemforschung

Skalen



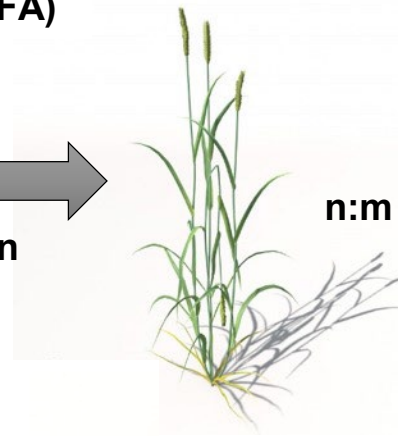
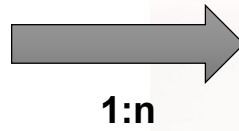
Themen



(Ewert et al. FAIRragro FAIR Data Infrastructure for Agrosystems - Proposal 2021, Zenodo, DOI: 10.5281/zenodo.7528172)

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Plant Genetic Resources for Food and Agriculture (PGRFA)

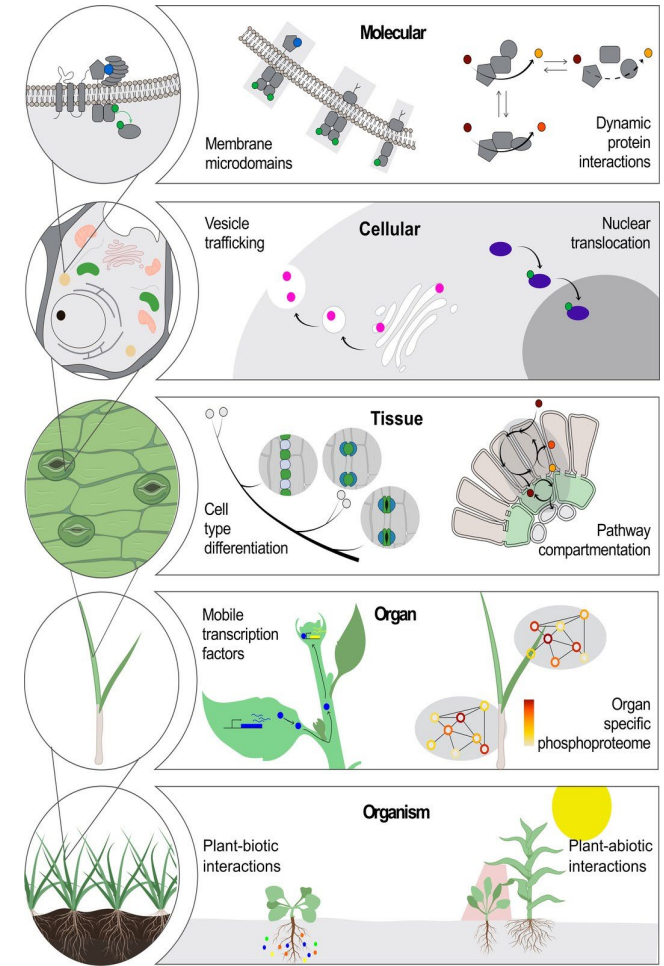


n:m

Plant Anatomy

Processes

Functions



(Plant Cell Atlas Consortium et al., Cell Biology, 2021)

PGRFA Materials



Food and Agriculture Organization of the United Nations



The International Treaty ON PLANT GENETIC RESOURCES FOR FOOD AND AGRICULTURE

• PUIDs

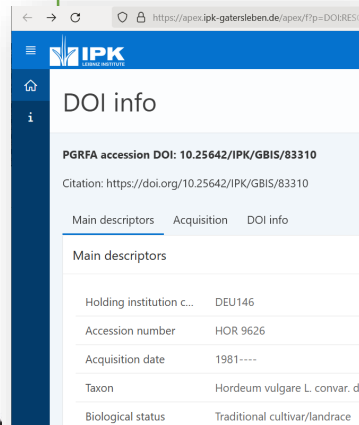
- Genbanks specific: accession numbers e.g. HOR 9626 / IPK Gatersleben
- Global PUIDs: DOIs e.g. 10.25642/IPK/GBIS/8331

• meta data

- darwin core/data cite
- MCPD - multi crop passport descriptor

• research data Infrastructures

- National => GLIS – Global Information System
- Germany -> in NFDI services ecosystem (FAIRAgro, DataPlant, 4BioDiversity)



Digital Twins for PGRFA

• PUIDs for samples

- samples
- domain specific databases e.g. sequences: EMBL-ENA, e!DAL-PGP
- proprietary global resolver: Identifier.org <https://identifiers.org/ena.embl:PRJEB40589>



• meta data - standard: MIAPPE+

- sample property
- environment
- management
- research data infrastructures
- institutional repos e.g. LIMS, DBs, file stores -> FAIR activation by FAIRAgro



<https://www.ipk-gatersleben.de/forschung/genbank/genbankdokumentation>

#FuturePlants

M. Lange, D. Arend, use case workshop NFDI section Infra - AG PID, 2023-01-27

www.ipk-gatersleben.de

Digital Twins in plant research data ecosystem



Raw Image Data

Spectrum: images taken at visible light, static fluorescence, near-infrared wavelengths, NMR images, CT images
Angles: top, several side views

Image-Derived Traits

Architecture: plant height, projected leaf area, leaf angles, growth rate
Color: average leaf hue, green to brown ratio, variance in leaf color
Intensity: static fluorescence, near-infrared emitted radiation

Environmental Data

Shoot environment: air temperature, humidity, light intensity, CO₂ concentration
Root environment: soil temperature, water content, nutrition levels, pH

Metadata

Plant: species, genotype, seed origin
Conditions: soil and container type, watering regime, experiment location
Measurements: observation units, measurement methods, sensor types

isatab
miappe

Sample flow

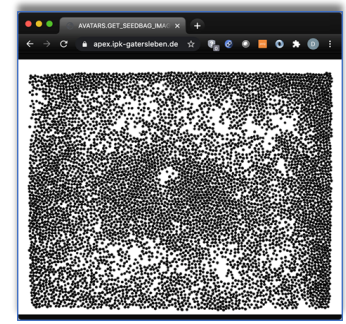
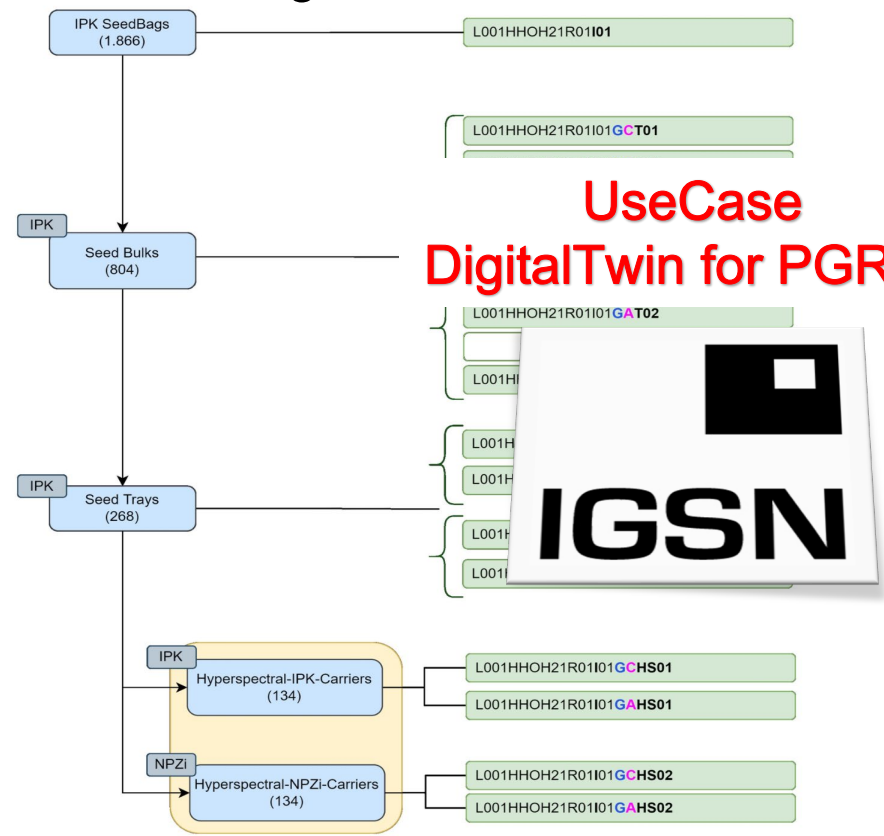


image source: [AVATARS project](#) IPK, NPZI

Digital Twins



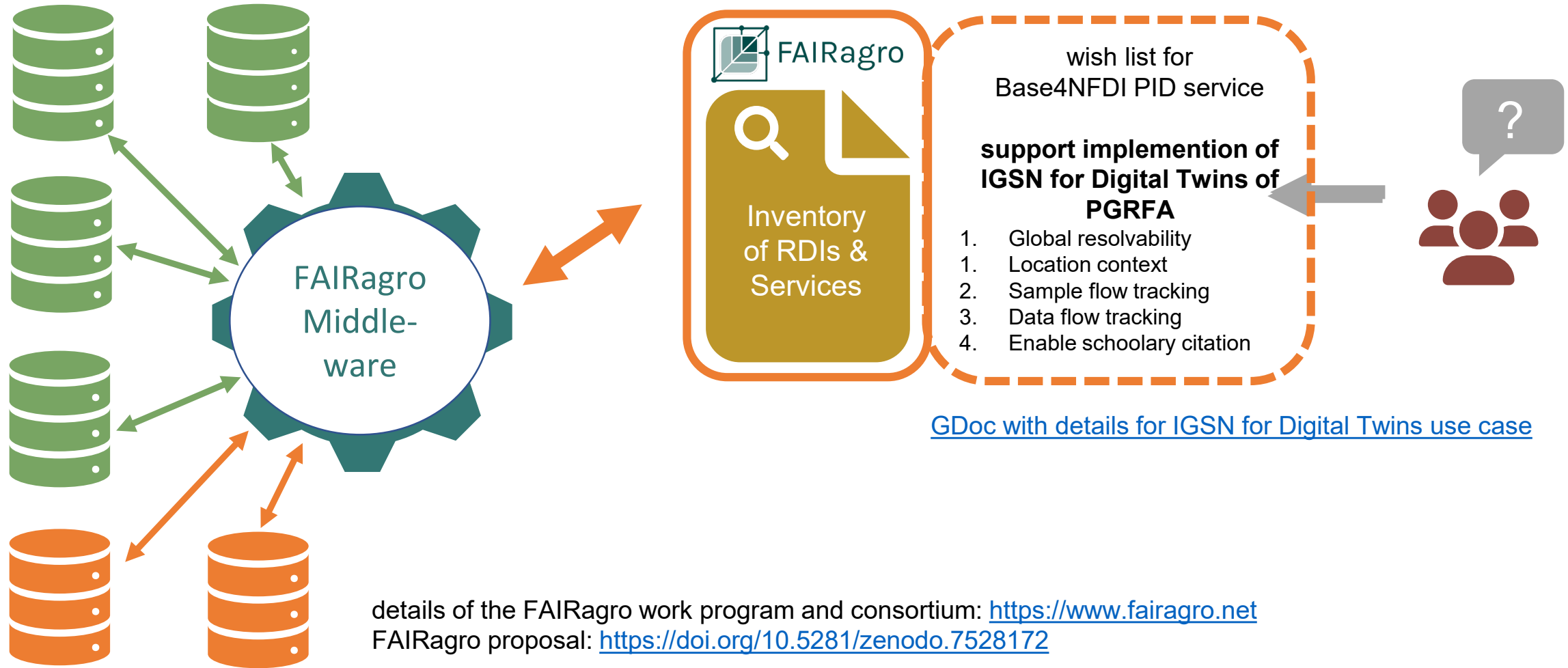
C004HMOO20R06I01ST01

genotype | field plot | season | physical object

(Rey-Mazon, NPZI; Plant 2030 status seminar; 2022)

(Arend et al. 2022 The Plant Journal; DOI: 10.1111/tpj.15804)

Integration into FAIRagro – Inventory and Search Portal in Task Area 4



Research Data Infrastructures (RDIs) for PGRFA