



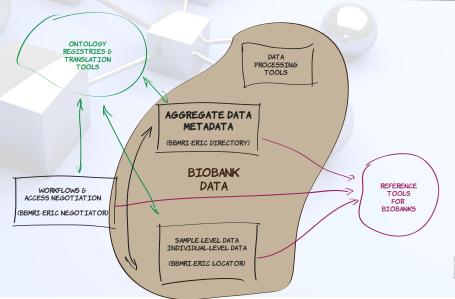
Metadata Services of BBMRI-ERIC: Directory 1.0 and 2.0

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Metadata Services of BBMRI-ERIC





- A researcher wants to identify biobanks of his/her interest: samples, data, access conditions, etc.
 - retrieving samples/data,
 - hosting samples/data.





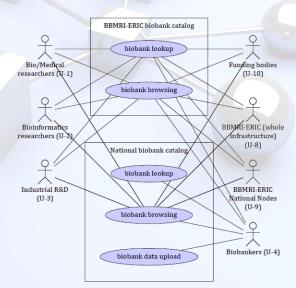
- Limitation of aggregate data: What biobanks might have samples/data of my interest?
 - inability to ask AND questions
 - no guarantee there is an actual combination of parameters,
 - samples/data may not be usable for the given analytical methods,
 - samples/data may come with an informed consent that does not allow their use for a particular research purpose.
- Directory allows for filtering out biobanks that certainly do not have material of interest.
 - input for other tools (e.g., BBMRI-ERIC Sample/Data Negotiator),
 - can be used for manual interactions with biobanks.

- A biobanker wants to
 - promote visibility of their biobank,
 - including all their services: samples + data + expertise,
 - promote access to samples and data,
 - search for similar biobanks,
 - minimize overhead resulting from integration of their biobank with BBMRI-ERIC
 - e.g., manual data input,
 - most biobanks lack an on-line interface (= connector) to their system to publish such information.
- BBMRI-ERIC National Nodes want to promote visibility of their biobanks and services.



- Participants (patients/donors) want to
 - know where their samples/data are hosted,
 - for what research (or what good in general) their samples/data are used.
- Funding agencies want to present the extent of infrastructures they are funding.





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Directory 1.0

http://bbmri-eric.eu/bbmri-eric-directory

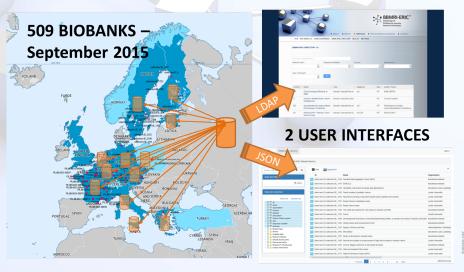
- Released: July 2015
- ► Based on flattened MIABIS 2.0¹
 - merged biobanks and collections for simplicity.
- Machine-readable and human readable interfaces
 - machine readable: LDAP, JSON,
 - human readable: web-based.
- Federated architecture
 - hierarchy of LDAP servers,
 - optional other interfaces for National Nodes that do not want to run their own LDAP (e.g. JSON, CSV),
 - storing data in BBMRI-ERIC LDAP server for National Nodes that do not want to run any infrastructure.

https://github.com/MIABIS/miabis/wiki





On BBMRI-ERIC web: http://www.bbmri-eric.eu/web/guest/bbmri-eric-directory-1.0



Directory 1.0



Mandatory information:

- institutional aspects
 - ID, juridical person, country
 - signed BBMRI-ERIC charter
- basic contact information
 - email
- material types
 - DNA, plasma, serum, urine, saliva, feces, RNA, blood, tissue frozen, FFPE or other fixed tissue, immortalized cell lines, isolated pathogen, other
- available diagnoses
 - preferably in ICD-10, support for other ontologies
 - mandatory only for clinical biobanks

Directory 1.0



Optional information:

- description & URL of biobank,
- detailed contact information
 - person, phone, address, geo-coordinates,
- access conditions for samples and data separately
 - joint-project based, fee-based, other,
 - URL describing policies,
- available data types
 - male samples data, female samples data, biological samples, survey data, imaging data, medical records, national registries, genealogical records, physio biochem measurements, other.



Types of Biobanks in Directory

- Clinical biobanks
 - 136 in BBMRI-ERIC now, estimated 11.6 M samples,
 - problem with very heterogeneous and even unstructured data.



Source: BBMRI-ERIC Directory 1.0



Types of Biobanks in Directory

Population biobanks

- 189 in BBMRI-ERIC now (incl. research studies), estimated 15.4 M samples,
- tend to have more homogeneous/consistent datasets.



Source: BBMRI-ERIC Directory 1.0

Types of Biobanks in Directory

- Other/specific types of biobanks
 - 2 veterinary biobanks
 - ... can still serve human medicine research
 - biobanks containing fetal samples
- Is biobank without biological samples (= "dry biobank") also a biobank?
 - population datasets (e.g., national registries),
 - imaging (e.g., radiology, pathology),
 - definitely of interest, maybe just renaming them to "databanks".

Preview of Directory 2.0

- Directory 2.0 introduces new entities:
 - collections,
 - biobanks now only represent the institutional aspects,
 - collections contain all the sample-related information,
 - biobank networks,
 - contact information,
 - to avoid contact duplication for biobanks, collections, and biobank networks.



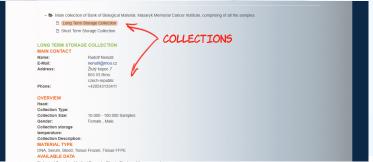
Preview of Directory 2.0

- 4 dc=directory.dc=bbmri-eric.dc=eu [webinar.bbmri-eric.eu] ou=biobanknetworks nu=hinhanks BIOBANKS i 🥮 c=at 💶 🥮 c=be = C=cz 🖶 🧀 biobankID=bbmri-effc:ID:CZ_CUNI_HK biobankID=bbmri-eric:ID:CZ CUNI LF1 COLLECTIONS biobankID=bbmri-eric:ID:CZ_CUNI_PILS biobankID=bbmri-eric:ID:CZ MMCI = CollectionID=bbmri-eric:ID:CZ_MMCI:collection:all_samples + CollectionID=bbmri-eric:ID:CZ_MMCI:collection:all_samples:LTS + CollectionID=bbmri-eric:ID:CZ MMCI:collection:all samples:STS biobankID=bbmri-eric:ID:CZ UPOL LF

Preview of Directory 2.0



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Preview of Directory 2.0

- Expected public release in next BBMRI-ERIC Newsletter (scheduled December 18, 2015).
- Ongoing data curation and updates.

- Support for linked data and semantics
 - modularized data model of MIABIS 2.0
 - clarifying definition of terms
 - problem with "minimum datasets":
 - what is really mandatory?
 - non-IT experts have sometimes a bit flexible view of word "mandatory"
 - MIABIS 2.0 to be ontologized
 - MIABIS (1.0) has undergone this ⇒ OMIABIS¹ & OBIB²
- Achieving interoperability though standardization: ISO TC 276 (WG1–WG5)
 - TC 276 WG5: Data Integration

http://www.ontobee.org/ontology/OMIABIS

²http://obofoundry.org/ontology/obib.html

- Support for linked data and semantics
 - some ontologies are very difficult to translate: e.g., ICD-10 ↔ SNOMED CT transform is context dependent
 - depends on level of detail: the more fine grained, the more difficult.
 - Support for different diagnoses ontologies in Directory 1.0
 - with syntactic search only (regex/wildcard patterns).
- Support for other data types
 - e.g., imaging biobank (working group between MIABIS and DICOM with ESR).



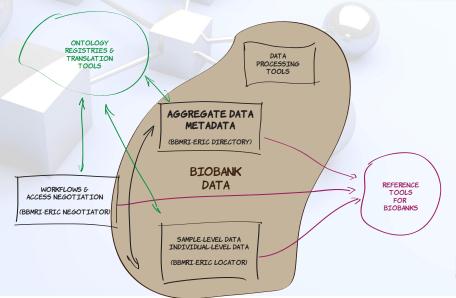
- Support for capabilities of biobanks
 - increasing amount of requests to host samples/data for external projects
- Support for studies
 - collections provide 1:N mapping to samples
 - studies will support M:N mapping
 - one sample can participate in multiple studies
- Support for quality management
 - on multiple levels of aggregation,
 - starting with declaring SOP availability.

- ▶ BBMRI-ERIC Directory can serve as a identifier translation/mapping service:
 - possible introduction of attributes for other identifier types for each entity.
- BBMRI-ERIC considers also other types of identifiers (under discussion):
 - DOI: assignment to biobanks/collections, plus possible dynamic assignment to sample sets once used for publication,
 - ongoing discussion with DataCite for BBMRI-ERIC to become DOI assigner.
 - Marker papers for biobanks/collections,
 - ORCID for affiliated people,
 - identifiers assigned by other communities.

- Metadata services and Directory become WP1 of CS IT
- Moving Directory onto the Molgenis platform
- ► Further development led by Morris Swertz (BBMRI.nl)
- ► Integration of BBMRI-LPC data
 - at the end of BBMRI-LPC project.

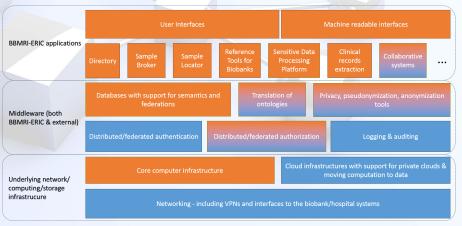


Toward Other Services of BBMRI-ERIC





Toward Other Services of BBMRI-ERIC





Thank you for your attention! Q?/A!

http://www.bbmri-eric.eu/ <petr.holub@bbmri-eric.eu>