Metadata at BBMRI

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About biobanks...

- Repositories of biospecimen (human) for use in research
- Different: types of biobanks, services, processes, quality controls, regulations (consent), data management (LIMS)
- Knowledge discovery about human diseases
- Towards personalized (precision) medicine





BBMRI.EU - Overview

- First research infrastructure funded by EC
- Main aim: Creation of a distributed bio-research infrastructure and network of biobanks with operational units in the members states
- Preparatory phase: February 2008 January 2011 (~5 M€)
- 54 partners from 33 countries, largest infrastructure
- 7 WPs
- WP5 Database harmonization and IT-infrastructure
 - Officially 11 partners from 8 countries
- Continuation: BBMRI-ERIC to start in the second half of 2013
 - BBMRI-ERIC Inauguration Conference Sept.16-17, 2013 in Graz/Austria



Managing resources for the future of biomedical research

BBMRI.EU - Data collection

- 1) Biobank questionnaires
- 2) WP5 metadata model
- 3) WP5 minimum dataset



Managing resources for the future of biomedical research

1) BBMRI.EU questionnaires

- 1. Core (13 pages)
- 2. Funding (1 page)
- 3. ELSI (4 pages)
- 4. IT (10 pages)
- 5. Sample collections (4 pages)



Managing resources for the future of biomedical research

1) BBMRI.EU questionnaires



www.bbmriportal.eu

•315 biobanks registered

•20 704 864 samples (DNA, boold, serum, tissue, cell lines, etc.)

Network	38
Core	325
Collection	669
Costs and Funding	137
Resources & Methods	98
Legal, Ethics and Governance	96
Biobank IT-solutions	89
Outcome of research using biological	
resources	309
Detailed description of biological samples	80



2) BBMRI.EU WP5 metadata model

- Hub-and-spokes
- Metadata model for(national) biobank hubs

- Needs further development
- Too complex?



3) BBMRI.EU WP5 minimum dataset

Data describing biobanks

Definition

BiobankAcronym NameOfBiobank Institution URL

Country ContactName ContactData

Data describing studies

Definition	Allowed values	Explanation
NameOfStudy	Free text in any language	
EnglishStudyName	Free text in English	Translation of study name in English
ContactName	Free text in English	
ContactData	Free text in English	Address, Phone (E.164, No. 905 – 1.IV.2008), e.g., +46 8
		524 877 59, Mail
KindOfStudy	Population-based, specific-disease, broad-spectrum of diseases	If "specific-disease", note ICD10
CategoriesOfDataCollected	[ClinicalDataAvailable, Diagnosis, Health information, Physiological/biochmical measures, Sociodemographic char., Socioeconomic char., Life habits/Behav., Physical environment]	Can be several values

Data describing subjects/cases/samples within biobanks

DefinitionAllowed valuesAgeGroupInterval [a,b], a>0, b<200, b>=a

Explanation

a and b should be selected so that k-anonymity is guaranteed. Age group of donor at time for sample collection, number of age groups determined by biobank

Address, Phone (E.164, No. 905 - 1.IV.2008), e.g., +46 8

Gender of subject Type of sample. From the BBMRI core question.

Date when sample was harvested There exists clinical data related to the sample

Genomics, proteomics etc Can be several values

GenderMale, Female, OtherGender of
SampleTypeSampleTypeDNA, cDNA/RNA, whole blood, blood cells isolates, serum, plasma, fluids, tissues cryo,
tissues paraffin-imbedded, cell-linesType of sample of
SampleDateSampleDateISO-standard (8601) time formatDate when
There existClinicalDataAvailableYes/NoThere existOrganCategoryFrom the BBMRI Detailed descr bio samplesEnomics,
RestrictionsOnSampleUseNone, Consent participant, IRB approval, Approval of owner of collectionCan be sev

NOTES:

Time stamp and version control are part of the metadata schema and upload services

Allowed values

Free text in English

Free text in English

Free text in English

Free text in English

ISO-standard (3166 alpha2), two letter code

ASCII

Explanation

524 877 59, Mail

Data pyramid for biobank information



BBMRI Nordic

- Nordic initiative BioBanking and Molecular Research Infrastructure (<u>BBMRI Nordic</u>) funded by the Nordic Research Council since 2010
- Collaborative network between national biobanking infrastructures in the Nordic countries
- BBMRI.se in Sweden, BBMRI.fi in Finland, BBMRI.no in Norway, Biobank Denmark, researchers from Iceland, Estonia and Faroe-Islands
- Biobank catalogs: first step towards biobank data sharing among the countries member of the network



BBMRI.se

- Similar structure as BBMRI.EU + "Biobank Technology" WP6
- 21 M€ from the Swedish Research Council and the medical universities
- Sweden was leading WP5 during BBMRI.eu preparatory phase

WP5 specific activities:

- Data discovery
- Data integration
- Standards



Questionnaire -minimum dataset for biobanks (MIABIS)

Datum för ifyllande av enkät				Om ja, ange vilken -oppigsd	ata. 🔲 Genomics 💷 Transcriptomics	Proteomics Metabolomics		Övriga unngifter om da	ta förkninnad med nrov	rsamlin <i>oon/st</i> udion
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Proviamlingens ID el.	_		_	Då provinsamlingen ej är p	abörjad eller pågående ifylle	planerad information (enligt	-	 Biddata. Vilka av följande ämnesområ. 	Annat, specificera:	a val mbiliza)?
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Minimum Information About Blobank data Sharing (MIABIS)



Norlin et al. A Minimum Data Set for Sharing Biobank Samples, Information, and Data: MIABIS. Biopreservation and Biobanking. August 2012, 10(4): 343-348.

BioBanking and Molecular Resource Infrastructure of Sweden



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MIABIS - Eurocourse

MIABIS

Eurocourse



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Some comments...

- We are aiming to have a stable data model for biobank data sharing (MIABIS, omiabis)
- Descriptive metadata about omics data from biobanked samples hasn't been formalized
- Important efforts on data sharing (<u>GEO</u>, <u>EBI-EGA</u>)
- Lot of ontologies! (<u>BioPortal</u>)
- Interesting project for bio-resources identification (PID) (<u>BRIF</u>) (<u>eagle-i</u>) (<u>ORCID</u>)
- EU's new Personal data protection legislation is a big issue
- Informatics integration of biobanking, research and clinics is still in an early phase

BioBanking and Molecular Resource Infrastructure of Sweden

Big data and Biobanking

- The volume of data produced by genomics research (e.g. NGS) is increasing at a higher rate than Moore's Law predicts
 - First human genome took a decade to complete and 3 years for data analysis at the cost of \$3 billions.
 - An entire human genome has **3.3 billions of base pairs**. It can be sequenced and analyzed in a few hours for a few thousand dollars.
 - Base pairs are read in short sequences and then assembled. It is done multiple times to produce an accurate sequence (raw file format, up to 30 terabytes).
 - A proteomics experiment can create more rows of data than a traditional row-based DB can handle
- Omics data formats vary depending on the technic and software (e.g. open-XML data format (proteomics), SFF, Fasta, fastq (NGS))
- Storage and analysis of omics data require a lot of store capacity and computing power
- An ongoing EU project will be a model for omics data storage and analysis (BiobankCloud: <u>http://www.biobankcloud.com/</u>)
- e-Science for Cancer Prevention and Cure <u>eCPC</u>



Big data and Biobanking: Sweden

- BBMRI.se is working towards a secure long term storage and analysis of big data from research studies on human samples
- A register over studies in Swedish bio-medical research institutions was launched in November 2012:
 - BBMRI.se Sample Collection Register
 - To date, over 80 studies have been registered from which ~32 are conducting omics experiments on ~276074 donors. If just a single sequence of each donor will be stored, ~276074 * 30 terabytes of storage will be needed (this a very small fraction of the real amount of data)
 - Another major issue is the omics data analysis
 - Up to date, researchers find their own solutions for omics data storage and analysis



Big data and Biobanking: Big issues

- The omics data resulting from research on biobanked samples is expected to increase dramatically over the next years. Some important matters need to be taken into consideration:
 - Secure long term storage capability (versioning genomes, etc.)
 - Tracking of omics data use and reuse (analysis results)
 - Fragmentation of knowledge among different omics
 - Standardization of omics data representation for data sharing
 - Standardization of omics data representation for analysis and interpretation
 - Personal data privacy protection
 - Some omics data can lead to the identification of the sample donor



Can EUDAT help to build a catalogue for BBMRI?

- Main user: Researcher
- Main aim: Search for bio-resources availability
 How deep to go? (any bio-resource, sample, sample data)
- Sample data at the metadata and aggregated level
- No personal data protection issues
- Centralized catalogue?
 - Biobanks, researchers, vendors... upload data (complicated)
- Distributed catalogue?
 - National catalogues upload data to EUDAT catalogue



EUDAT Catalogue for BBMRI?



Search for sample & sample data

BBMRI.se BioBanking and Molecular Resource Infrastructure of Sweden

Search for any Bio-resource

Conclusions

- Projects as EUDAT would be of great benefit for BBMRI
 - storage infrastructure for omics data
 - Relevant information about sample and sample data availability all over Europe
 - Easy to use (dropbox, eage-i)
- BBMRI does not have a stable metadata model for biobanking but there is sufficient work done in that direction
- Networking biobanks and biomedical research institutions big challenge
- Implementation of a BBMRI catalogue should be divided in several stages, from the general information about biobanks down to the specific data about samples and data analysis
- Interdisciplinary catalogue: social statistics, bio-statistics, modeling, environment, ethics and law, medical terms (SNOMED, ICD codes, etc.)
- Omics data storage and analysis is a bottleneck in biomedical research due to the sizes and the diversity of data formats and analysis methods

Thanks! Tack! Gracias!

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