Sharing sensitive human genomic, phenotypic, & clinical data

Dr. Mallory Freeberg
Coordinator, European Genome-phenome Archive (EGA)

Tw: @MalloryFreeberg Em: mfreeberg@ebi.ac.uk ORCID: 0000-0003-2949-3921

Bioinformatics for Biobusiness

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What is EMBL-EBI?

World leading source of public biomolecular data

 Our vision is to benefit humankind by advancing scientific discovery and impact through bioinformatics.

 Part of the European Molecular Biology Laboratory (EMBL), Europe's flagship laboratory for the life sciences.



Data resources at EMBL-EBI



Chemicals, molecules and drug discovery

ChEBI
ChEMBL
MetaboLights
Open Targets

SureChEMBL



Genes, genomes and RNA

ArrayExpress

Ensembl
European Nucleotide
Archive
Expression Atlas

HGNC MGnify

Rfam

RNAcentral VectorBase WormBase



Proteins

AlphaFold DB Enzyme Portal InterPro PDBe Pfam

PRIDE

UniProt



Imaging and cellular structure

BioImage Archive Electron Microscopy Data Bank EMPIAR



Genetic variation and disease data

COVID-19 Data Platfo

European Genome-phenome Archive

European variation Archive Mouse informatics

Controlled-access, human data



Literature and knowledge management

BioModels BioSamples BioStudies Complex Portal Europe PMC

GWAS Catalog IntAct

OmicsDI Ontologies

Reactome











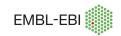












What is special about controlled access data?







Exposure of personal or sensitive data is potentially harmful

Individuals have the right to protection of their personal data

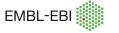
e.g. name, email, genetic characteristics, medical history, disease diagnoses EU General Data Protection Regulations (GDPR)

UK General Data Protection Regulation and Data Protection Act 2018

Finland Data Protection Act (supplement to GDPR)

Research aims

Maximise value
Be good data stewards
Reduce waste



EGA manages sensitive human data



- Mission Permanent archiving and sharing of personally identifiable genetic, phenotypic, and clinical data generated for the purposes of biomedical research projects or in the context of research-focused healthcare systems
- Drive implementation of standards for supporting sharing of controlled-access human data
- FAIR principles
- Community standards



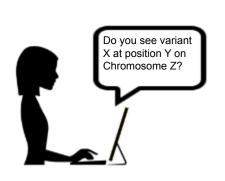


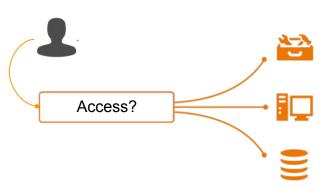


formal collaboration to manage EGA



Challenges for managing controlled access data







Data discovery

How do I find data relevant to me? How do I know these data are useful?

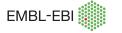
Data access management

How can I access data?

How can I prevent unauthorised access?

Regional data protection regulations

What do I do if my data cannot be shared outside regional borders?





Data Discovery Finding relevant, useful data



Data discovery

How do I find data relevant to me? How do I know these data are useful?

Won't talk about:





- Findable: accessions, indexed
- Accessible: standard access protocols
- o Interoperable: ontologies, vocabularies
- Reusable: rich metadata, data usage license
- Metadata curation and harmonisation

"As open as possible. As controlled as required"

Discovery over public metadata

→ Non-identifiable, aggregated / binned

Discovery over sensitive data without direct access

→ Query tools

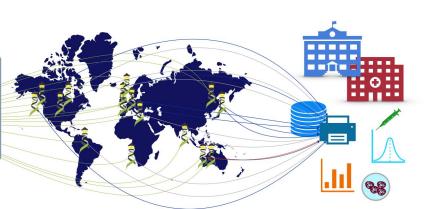


Data Discovery Beacon API for federated discovery

"Yes" (or "no")

Can you provide data about focal deletions in CDKN2A in Glioblastomas from juvenile patients with unrestricted access?

> CDKN2A:DEL size<1Mb granularity:record ncit:C3058 DUO:000004 HP:0003621



Beacon defines an open standard for federated discovery of human genomic, phenotypic, and clinical data in biomedical research and clinical applications

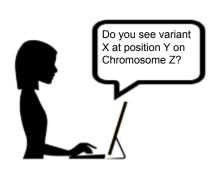
Beacon v2 enables enhanced querying over clinical and phenotypic information, curated variant annotations, and data use conditions

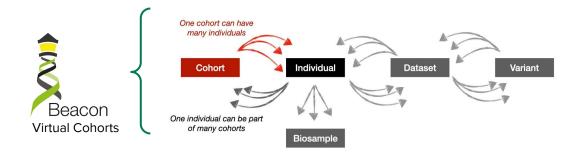






Data Discovery Enhanced discovery features

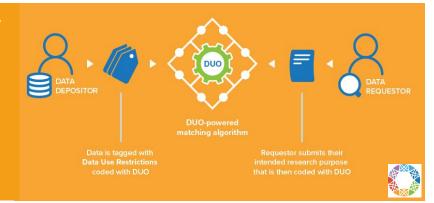




Data discovery

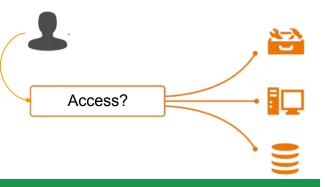
How do I find data relevant to me? How do I know these data are useful? Data Use Ontology
(DUO) to describe
data use conditions
and restrictions

Automate matching with research purposes





Data Access How to manage access safely



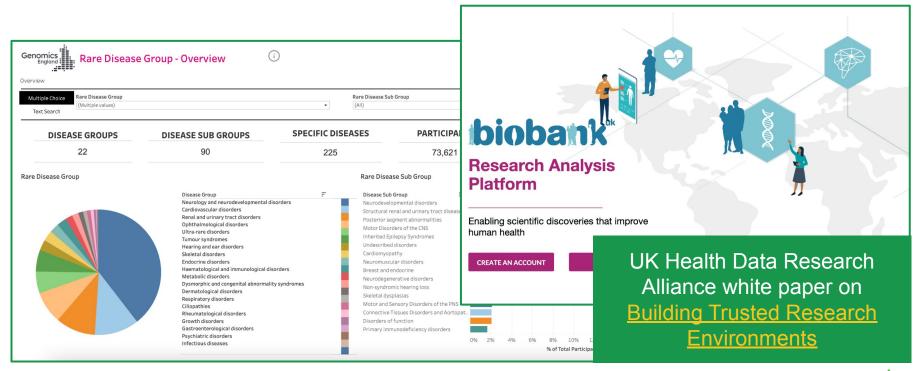
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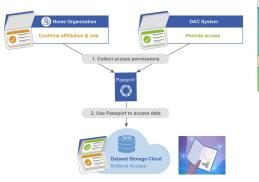


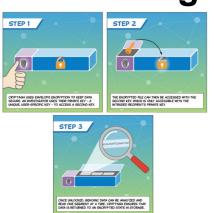
Data Access Bring analyses to the data

Trusted research environments (TREs) / Secure Data Environments (SDEs)



Data Access Adopt modern, automated, scalable processes for federating data access







Standardised researcher identities and authorised data

GA4GH Passports / Visas

Secure file container format

GA4GH Crypt4GH

Machine computable data use conditions and restrictions

GA4GH DUO

Interoperable authentication and authorisation systems

ELIXIR Life Science Login / GA4GH AAI







Challenges for managing controlled access data

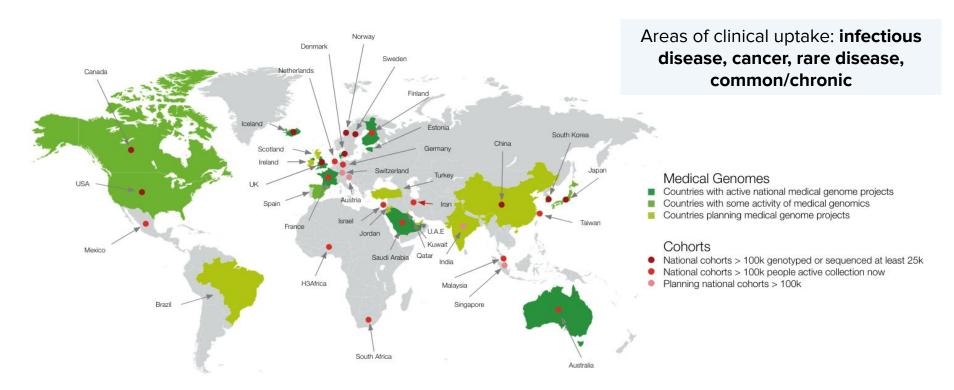


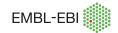
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What do I do if my data cannot be shared outside regional borders?

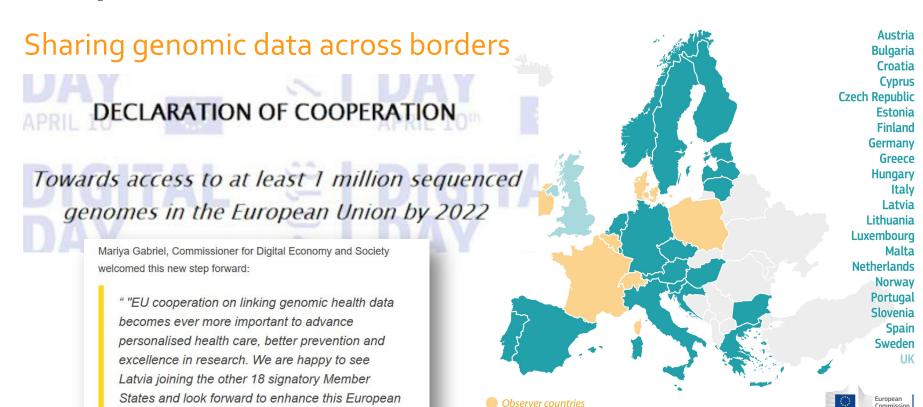


Genomics & healthcare is a global effort





European 1+ Million Genomes Declaration



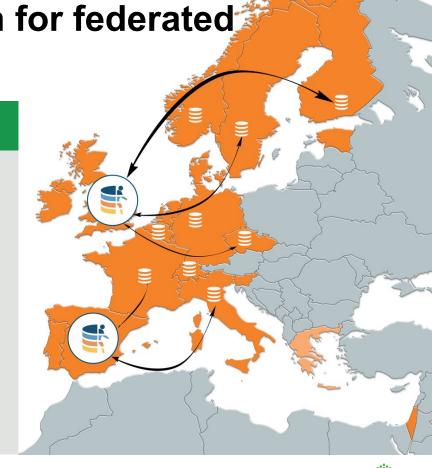
collaboration. " "

Infrastructure & coordination for federated human data access

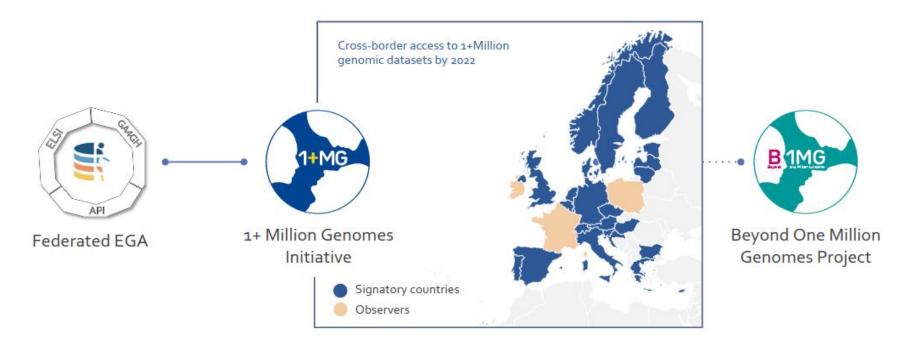
Federated EGA Network provides:

- Federated network of human data repositories and regional hubs
- Platform for discovery, access, distribution of human data
- Coordination and standards for discovery and access
- Community of experts to build capacity across Europe

ELIXIR Nodes engaged through the ELIXIR Federated Human Data Community



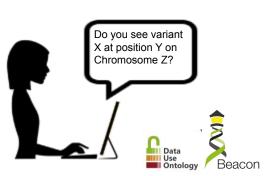
Federated Human Data initiatives in Europe

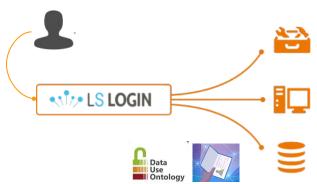


Building towards genomic data infrastructure



Solutions for managing controlled access data







Data discovery
FAIR data, Beacon, DUO

How do I find data relevant to me? How do I know these data are useful? Data access management

TREs, Passports/Visas, Crypt4GH, DUO, AAI

How can I access data?

How can I prevent unauthorised access?

Regional data protection regulations
1+MG, Federated EGA, B1MG

What do I do if my data cannot be shared outside regional borders?



Thank you

Thanks to:

- EGA teams at EMBL-EBI and CRG
- Collaborators at ELIXIR and GA4GH
- Collaborators with Federated EGA



