[Federated] EGA

Providing global discovery and access for sensitive human data

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Sharing sensitive data key to enable science & health





Exposure of personal or sensitive data is potentially harmful Individuals have the right to protection of their personal data

Governments pass legislation that codifies data protection rights

Support aims of research & health care initiatives

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 development of and implement community standards for supporting FAIR data sharing



Global Alliance for Genomics & Health Collaborate. Innovate. Accelerate.

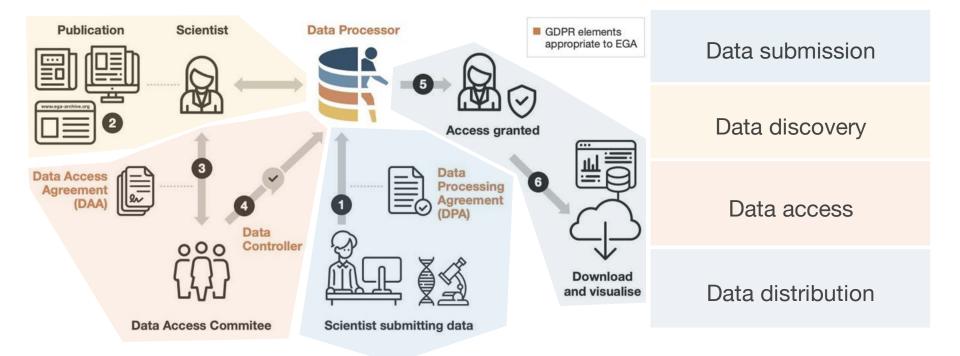


2008: EGA launched at EMBL-EBI

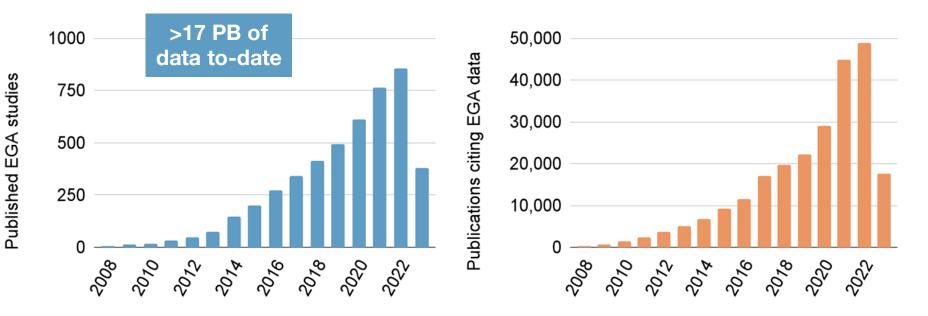


2013: CRG & EMBL-EBI enter formal collaboration to co-manage EGA

EGA data access model



EGA grows and remains key to life science research



Impact of EGA in personalised medicine

– Common diseases

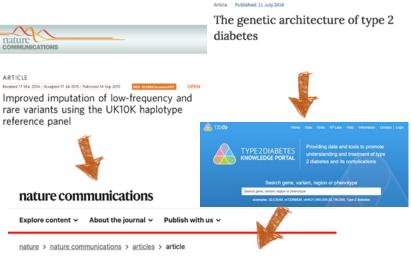
- UK10K (EGAS00001000713)
- WGS GoT2D (<u>EGAS00001001459</u>)
- WES GoT2D (EGAS00001001460)

Revision and reanalysis of existing genetic data by applying more powerful analytic techniques and resources generated.

A total of 70,127 subjects (70KforT2D) were retained for downstream analysis

Up to 7 loci were identified that were not previously reported as associated with T2D. Providing more candidate causal variants.

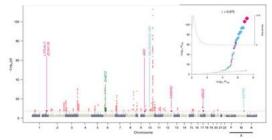
refs: <u>Nature Comms (2015)</u>, <u>Nature (2016)</u>, <u>Nature Comms (2018)</u>



lature

Article | Open Access | Published: 22 January 2018

Re-analysis of public genetic data reveals a rare Xchromosomal variant associated with type 2 diabetes



European 1+MG initiative sets vision for data

sharing Countries that have signed the 1+MG Declaration since 2018 Austria elixir Belgium Bulgaria Croatia Cyprus **Czech Republic** Estonia Finland Germany Greece API Hungary Italy Federated EGA Latvia Lithuania Luxembourg Malta Netherlands Norway Portugal Slovenia Spain Sweden Towards access to at least 1 million UK sequenced genomes in the EU by 2022 European



Beyond One Million Genomes Project





More genomics data generated in healthcare context



Areas of clinical uptake: infectious disease, cancer, rare disease, common/chronic

Medical Genomes

- Countries with active national medical genome projects
- Countries with some activity of medical genomics
- Countries planning medical genome projects

Cohorts

- National cohorts > 100k genotyped or sequenced at least 25k
- National cohorts > 100k people active collection now
- Planning national cohorts > 100k

Federated EGA collaboration officially launched in September 2022



ELIXIR Europe

@EGAarchive announced the first signings of the Collaboration Agreement
between national Nodes and Central EGA,
an important step in the formation of a #FederatedEGA.

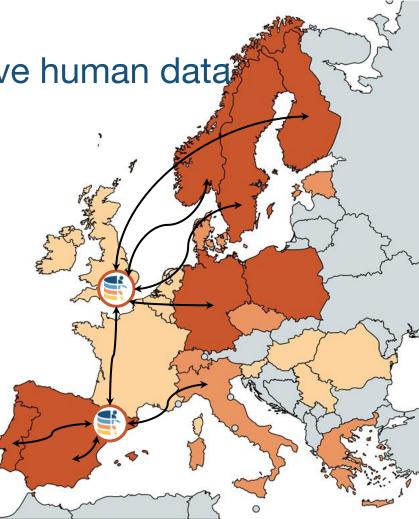
Read our joint news release with @emblebi & @CRGenomica: elixireurope.org/news/federated...

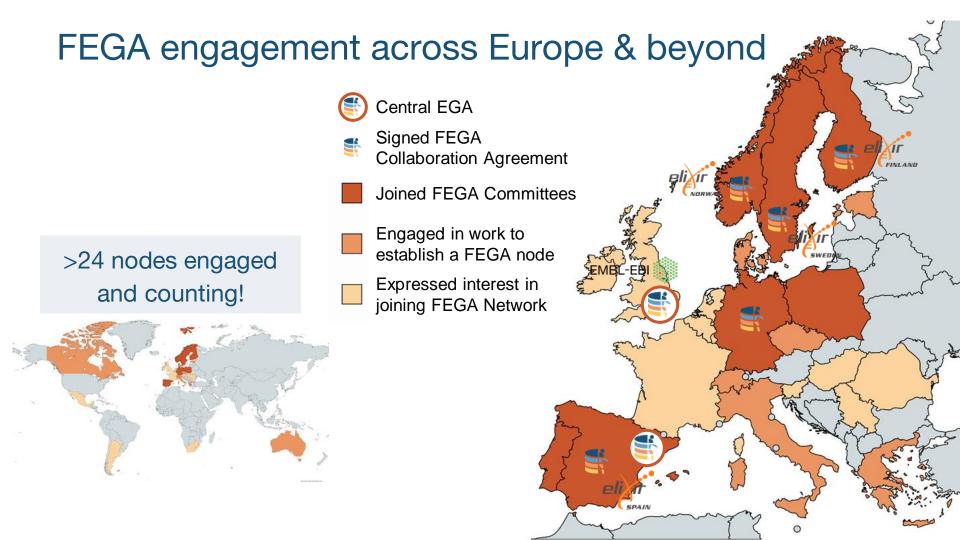


Federated EGA:

discovery & access for sensitive human data

Federated EGA strives to support the discovery of and secure access to human data globally, while respecting national data protection regulations, with the goal of accelerating disease research and understanding and improving human health.





Joining FEGA network follows established pathway Federated Contact FEGA with intent to join European Genome-phenome EG Archive Present use case to ELIXIR FHD Community; complete FEGA Use Case Survey - ACA Node interested in ioining FEGA Network Perform FEGA Maturity Model self-assessment 3 Present Maturity Model self-assessment and 11 Mond R demonstrator results to FEGA Strategic Committee Set up local EGA node with mock CEGA 0 V credentials: مٰك confirm node Q= is operational 습습습 FEG 3 $(\checkmark$ DEMO 222 AR @= Receive test Invited to join FEGA CEGA credentials: Strategic confirm successful Committee and Sign FEGA C.CO. Confirm successful connection **FEGA** Operations Invited to join Legal completion of FEGA Receive Establish node Committee as **FEGA Strategic** Agreement SOPs and node end-to-end production **FEGA Node** observer Committee and demonstrator CEGA communications operating **FEGA Operations** credentials Committee as full in production Governance members Technical

Committee

FEGA Maturity Model enables self-assessment



Federated EGA Maturity Level Model

The vision

Level definitions

FEGA MM

The maturity model is envisioned as a mechanism to drive engagement and disengagement of nodes being part of the EGA federation. This model is divided into different domains and subdomains with a 5-levels scale. The first two levels are deemed to drive the initial interactions of an organization willing to join the Federated EGA ecosystem. Levels 3 and 4 aim to establish clear guidelines to reach a fully operational Federated EGA Node. Level 5 is designed to facilitate periodic review of the different indicators as well as to ensure the adoption of the most up-to-date technical and legal specifications.

[1] Governance, Strategy & Sustainability	0
[2] Legal	0
[3] Data & metadata management	0
[4] Technical Infrastructure	0
[5] Operations Support	0
[6] Communications, Community Building, and Engagement	0



https://inab.github.io/fega-mm/

Guidance for nodes considering joining FEGA

Federated European Genome-phenome Archive	Docs » Establishing a Federated EGA Node Federated European Genome-phenome Archive	Le 2. Explore implemented standards Central EGA largely adhere to GA4GH standards. Specific standards already implemented are summarised below:					
Establishing a Federated EGA Node	(Search docs	Standard	Purpose	Specification Version	Supported Version	Implementation	Publication/Preprint
What am I doing here? What is Federated EGA?	Establishing a Federated EGA Node TOPICS	Beacon	Supports discovery of genomic variants and individuals.	v1.0.1	v0.3	Specification, Documentation, Endpoint	N/A
How do I start?	Maturity Model	Crypt4GH	Enables direct byte- level compatible random access to encrypted genetic data stored in community standards (e.g. CRAM, VCF).		v1.0	Specification, Documentation, Endpoint	DOI
What does the journey look like?	Data and Metadata Management 1. Learn about data security best practices B 2. Explore implemented standards						
Acknowledgements License							
Aaturity Model	3. Understand data definitions and	Data Use Ontology	ogy semantically tag	2021-02-23 r	3 2021-02- 23	Specification, Documentation, Endpoint	DOI
ata and Metadata Management	flow	(DUO)					
utreach and Training	4. What's next?						
echnical and Operational	Outreach and Training						
overnance and Legal	Technical and Operational Governance and Legal		on a researcher's authorization level or intended use.				

https://ega-archive.github.io/FEGA-onboarding/

What's next for EGA and Federated EGA?

- Build on human data sharing knowledge & experiences of experts across Europe
- → Accelerate additional national nodes joining Federated EGA
- Continue advocating for community standards via Driver Projects and ELIXIR::GA4GH Strategic Partnership
- → Support transnational access to human data across Europe & beyond via ELIXIR-coordinated initiatives/projects e.g. 1+MG, B1MG, GDI

Acknowledgements: Federated EGA





Thank you!

Questions?











Additional resources

- EGA: <u>https://ega-archive.org/</u>
- Joining FEGA: <u>https://ega-archive.github.io/FEGA-onboarding/</u>
- GA4GH: <u>https://www.ga4gh.org/</u>
- ELIXIR: <u>https://elixir-europe.org/</u>
- GDI: <u>https://gdi.onemilliongenomes.eu/</u>