

Federated EGA: A global network for discovery and access of sensitive human data

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Abstract

With increasingly strict regulations for managing sensitive human omics and healthcare-derived data, solutions are required to ensure research and clinical data can be accessed across national borders. Safely sharing sensitive data is vital for data reuse to support biomedical research and personalised medicine programs. The Federated European Genome-phenome Archive (FE GA) provides a data sharing solution of infrastructure and governance frameworks to support discovery of and secure access to human data globally, while respecting national data protection regulations. Prompted by European initiatives such as the 1+ Million Genomes and European Health Data Space, the FE GA was officially launched in 2022 with a network of nodes in the UK, Spain, Norway, Sweden, Finland, and Germany. The next phase of FE GA will build upon these early successes to expand data sharing both within and outside of Europe. Accelerating this expansion can be achieved through collaboration with initiatives already working towards human data sharing, for example: the European Genome Data Infrastructure (GDI), Data Science for Health Discovery and Innovation in Africa (DS-I Africa), and Australian BioCommons. Ultimately, the FE GA vision is to build a global, interoperable discovery and access network of human data resources, to accelerate disease research and understand and improve human health.

The European Genome-phenome Archive (EGA) securely shares sensitive human data

The EGA is a service for 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 [1]. Data archived at the EGA were collected from individuals whose consent agreements authorise data release only for specific research use to bona fide researchers.

African research represented in EGA

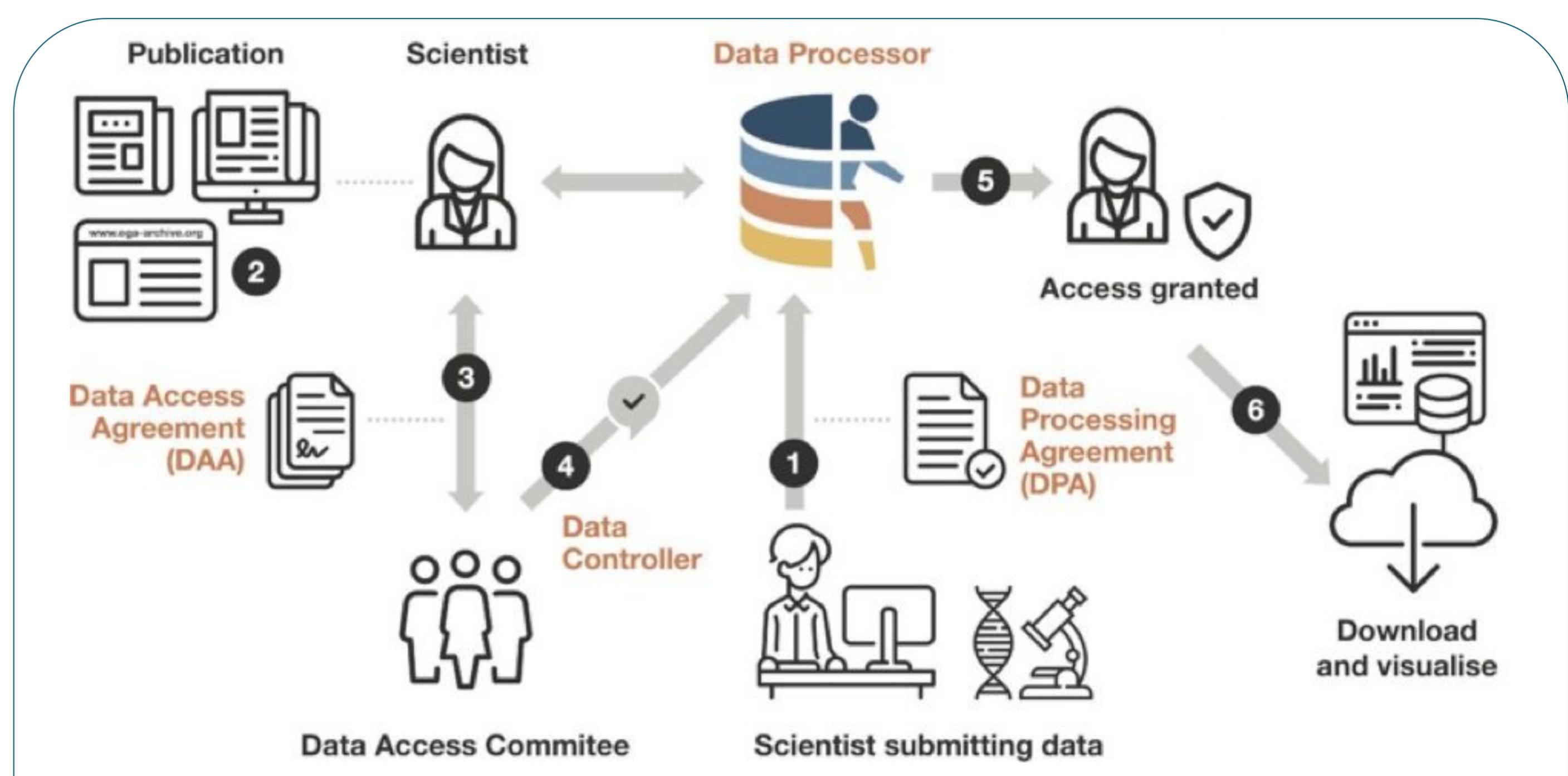
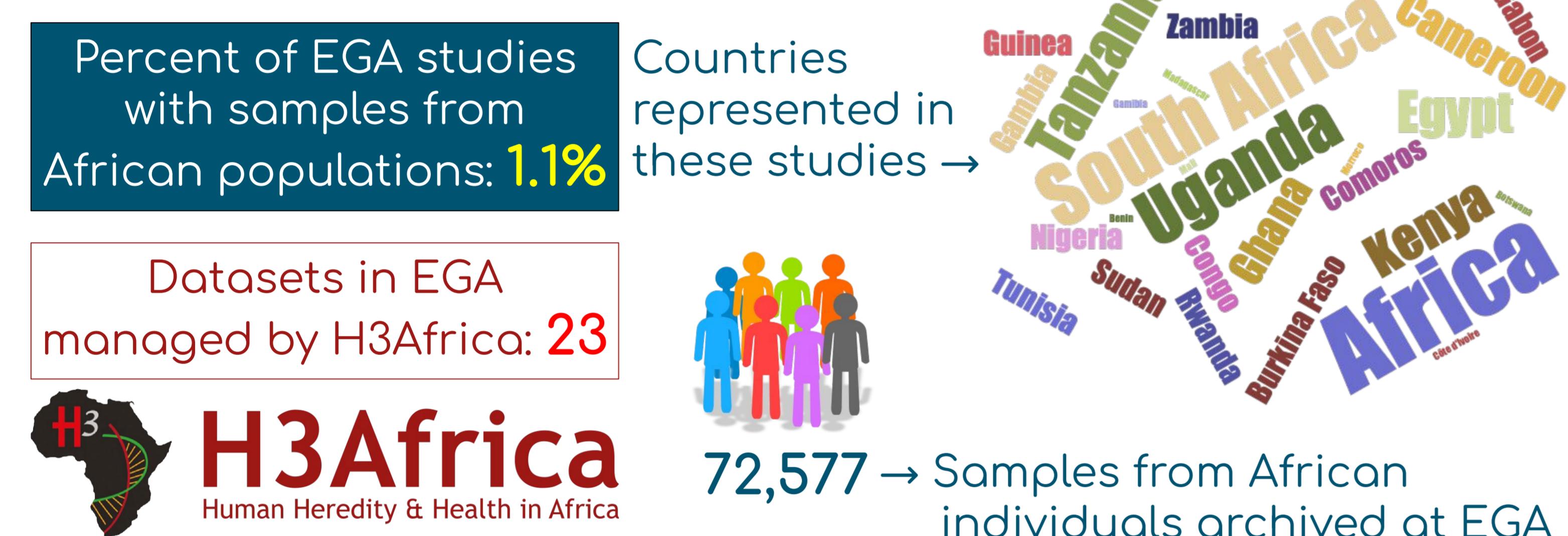


Figure 1: EGA facilitates the submission, discovery, access, and distribution of sensitive human data following strict protocols. Researcher submits controlled access human data to EGA after signing a Data Processing Agreement (1). EGA processes, archives, and releases the dataset to be findable. Another researcher discovers data of interest at the EGA (2). They contact the Data Access Committee (DAC) for the data of interest and agree to the terms of data reuse by signing a Data Access Agreement (3). The DAC informs EGA that access is approved (4). The EGA grants access to the researcher (5) who can then download and visualise the data (6).

Federated EGA as a global network for sensitive data discovery and access

Vision: A global resource for discovery and access of sensitive human omics and associated data consented for secondary use, through a network of national human data repositories to accelerate disease research and improve human health.

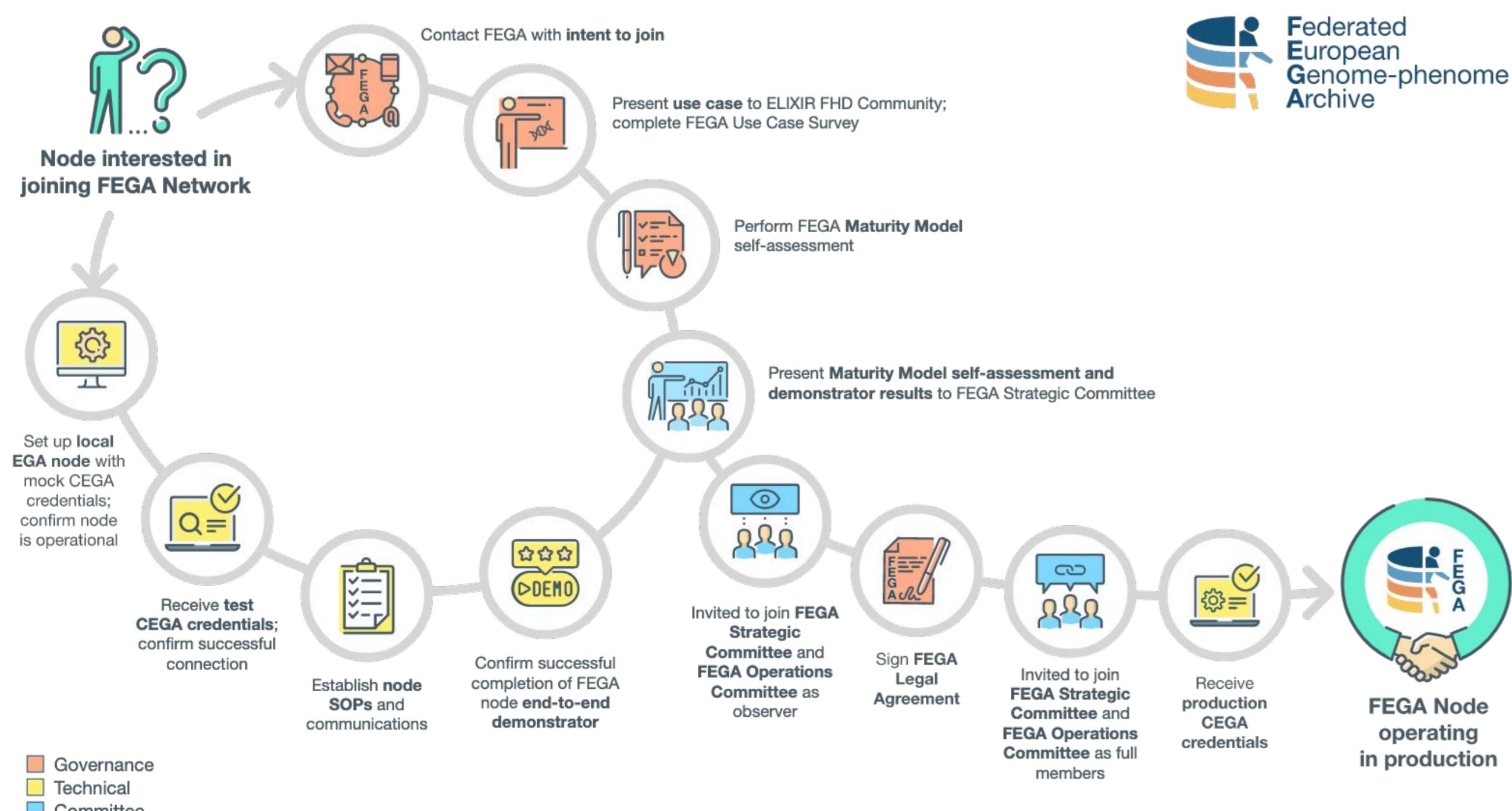
Value for Researchers

- Common standards and interfaces → interoperability
- Common phenotypic & clinical data models → (meta)data harmonisation
- Federated user identity → simpler, faster data access

Value for Federated EGA nodes

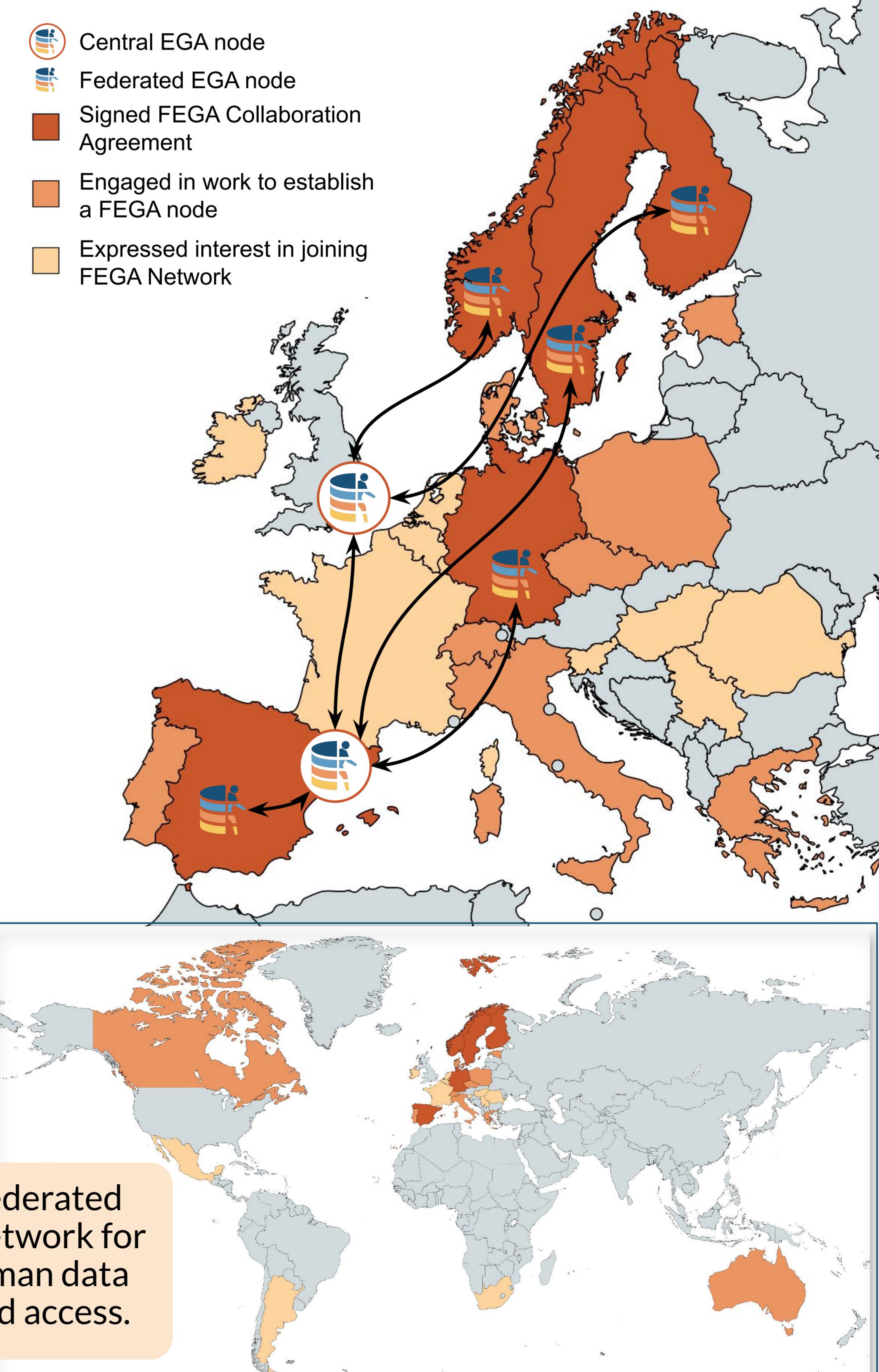
- Peer collaboration network of expertise & knowledge
- Increased data reuse via FE GA discovery network
- Access to common standard operating procedures
- Collaboration on technical implementations
- Increased interoperability with similar initiatives
- Access to shared training and outreach materials

How to join Federated EGA Network



about Federated EGA!

Federated EGA in 2023



References & Resources

- [1] <https://doi.org/10.1093/nar/gkab1059>
- Establishing a FE GA Node: <https://ega-archive.github.io/FE GA-onboarding/>
- FE GA updates @ ELIXIR Federated Human Data Community: <https://elixir-europe.org/communities/human-data>

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