

NeIC Heilsa: The Roadmap to Sensitive Data Archiving & Federated EGA

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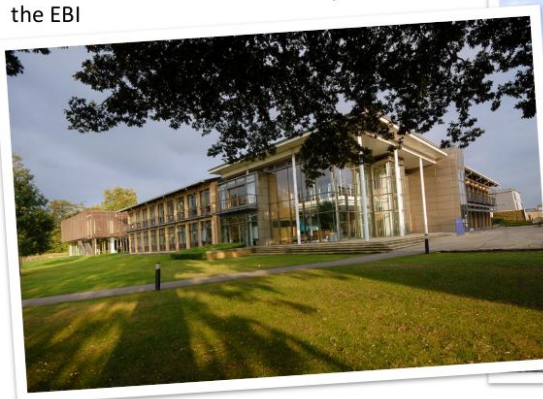
NeIC Heilsa WP1 co-lead

EGA brief history

- The European Genome-phenome Archive (EGA) is a **service for permanent archiving and sharing** of all types of **personally identifiable genetic and phenotypic data** resulting from **biomedical research projects**.

How the EGA is managed

The EGA was launched in 2008 by the EBI



In 2012, EBI and CRG started working together to establish EGA as a joint venture, which has grown in the context of ELIXIR

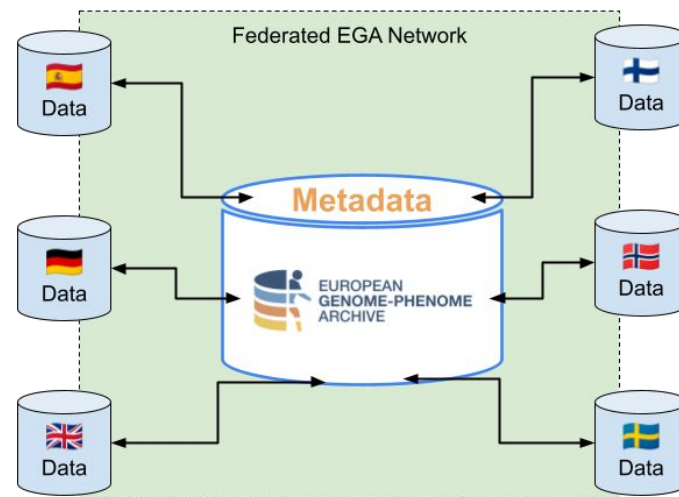


What is Federated EGA

“The **Federated European Genome-phenome Archive** (EGA) is a **distributed network of repositories** for **sharing human-omics data and phenotypes**.

Typically a **node** would be an **organization or project** that **hosts human genetic data** so that the data can remain **within a jurisdiction**. ”

<https://ega-archive.org/federated> (source of the text)



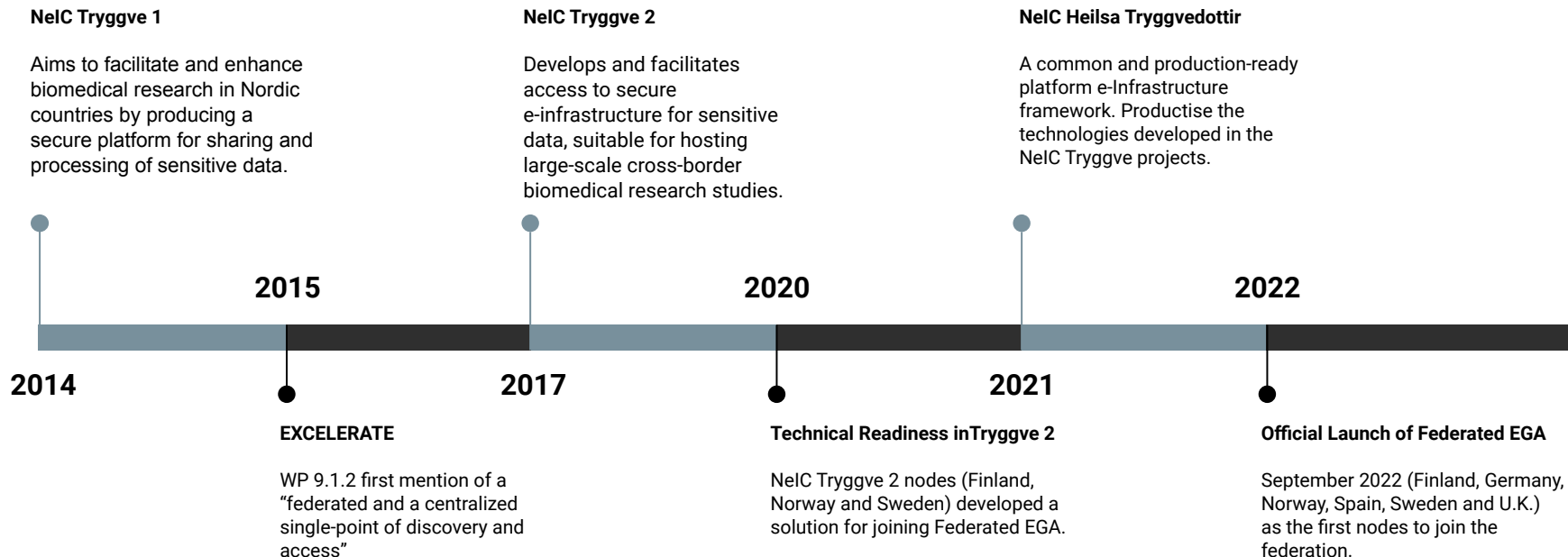
Benefits for research community

- Unified accession identifiers and versioning scheme for identifying and referencing data conforming to the FAIR principles;
 - EGA accession identifiers can be used in scientific publications by researchers;
- Use of international standards (crypt4GH, beacon, HTSGET etc.);
- Unified brand for storage, discovery and access to human genetic and phenotype data.

Federated EGA - History briefly

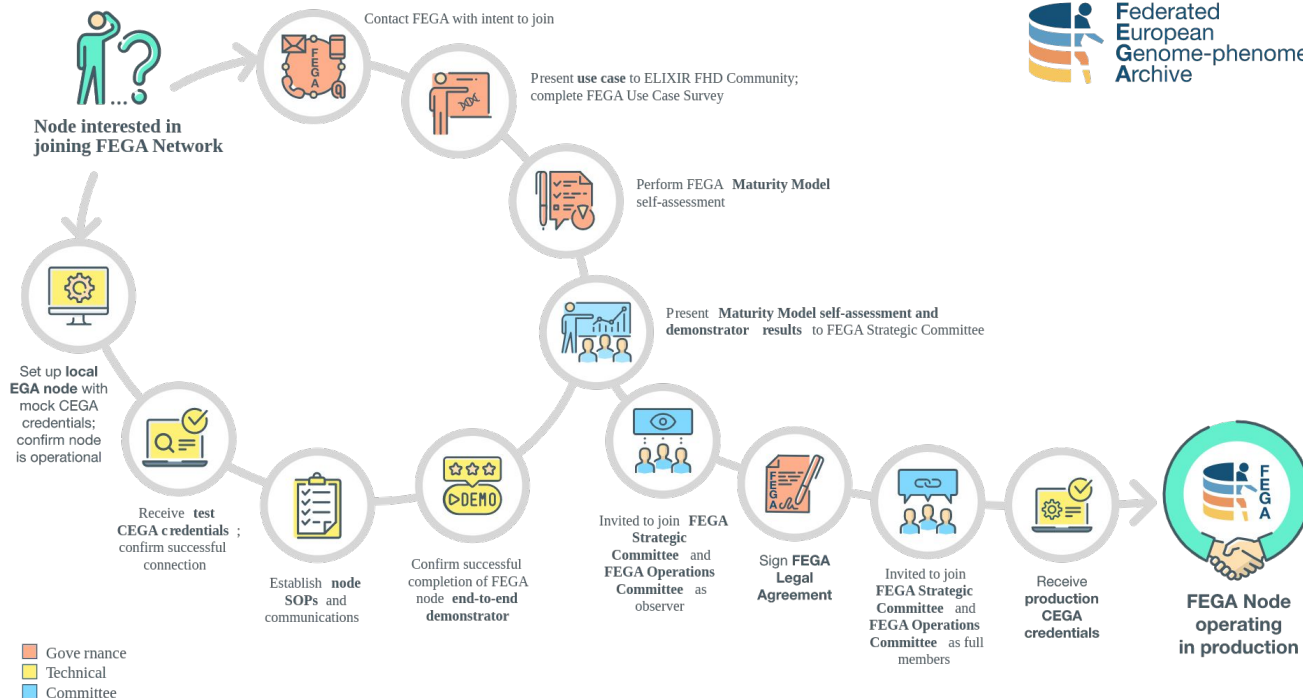
- “Portable submission toolkit” or “mini-EGA in-a-box” mentioned in the EXCELERATE grant (2015-2019) <https://elixir-europe.org/about-us/how-funded/eu-projects/excelerate>
 - *Subtask 9.1.2: Portable submission toolkit. This task is composed of data format definitions and software components, a “mini-EGA in-a-box” will allow **increased local control** and coordination of data collection, and allow early validation of standardized data and metadata formats. This implementation provides the practical means for distributed projects to collect access-controlled human biomedical data in a manner that maintains a coordinated data model and dataset registry, enabling **federated and a centralized single-point of discovery and access.***
- In March 2016 (AHM Barcelona) when polling the ELIXIR Nodes, the need **for *not* moving the sensitive files from “local jurisdictions”** arise as the main driver for it
 - ... but still willing to share data for research
- Known over the years as “*mini-EGA in-a-box*”, “*Federated EGA DROPBOX*”, “*Local EGA*”, “*EGA sensitive data archive*”, “*FEGA*”, “*LEGA*” etc.
- As the ELIXIR funding for the task was minimal, some Elixir/Nordic Nodes step in to move the project forward

Federated EGA - Timeline*



* from a NeIC involvement perspective

Federated EGA - Journey*



* as illustrated by <https://ega-archive.github.io/FEAGA-onboarding>

NeIC Heilsa Tryggvedottir - Work Packages

- WP1: Federated cross-border analysis, sensitive data archives, and roadmaps
- WP2: Creating and sharing genome/phenome test datasets

NeIC Heilsa Tryggvedottir - joining Federated EGA

- A code base for archiving of sensitive data
 - Documentation available at: <https://neic-sda.readthedocs.io/en/latest/>
 - Packaged to be convenient for setting up Federated EGA, or stand-alone instances for archiving of sensitive data
 - Technical questions about the solution <https://gitter.im/neicnordic/sda-pipeline>
- Common operating procedures for (Federated EGA) nodes
 - Documentation
 - Processes
 - Roles and responsibilities
 - Security

NeIC Heilsa Tryggvedottir - Outreach

- The Nordic ELIXIR nodes are engaged in multiple international projects and collaborations, such as:
 - 1+MG <https://digital-strategy.ec.europa.eu/en/policies/1-million-genomes>
 - B1MG <https://b1mg-project.eu/>
 - ELIXIR Federated human data community <https://elixir-europe.org/communities/human-data>
 - ELIXIR Converge <https://elixir-europe.org/about-us/how-funded/eu-projects/converge>
 - EOSC-Nordic <https://www.eosc-nordic.eu/about-eosc-nordic/>
 - EuroHPC - Lumi <https://www.lumi-supercomputer.eu/eurohpcju/>
 - GA4GH <https://www.ga4gh.org/>



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Thank you!