



Sensitive data archiving in Nordics and Federated EGA

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NeIC 2019 - Open Science with Sensitive Data workshop

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NordForsk



Slides adapted from other presentations by

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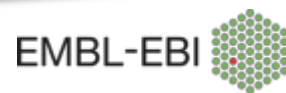
Tryggve2 major deliverables

1. **Sensitive data archiving**
2. Production quality processing services
3. Homogenized user experience
 - User mobility
 - Workflow mobility
 - Data mobility
4. Nordic use cases
 - Research
 - Infrastructure development
5. ELIXIR AAI
6. IT Security
7. ELSI Topics



EGA

The EGA was created by the EBI, in 2008, as an extension of the ENA...

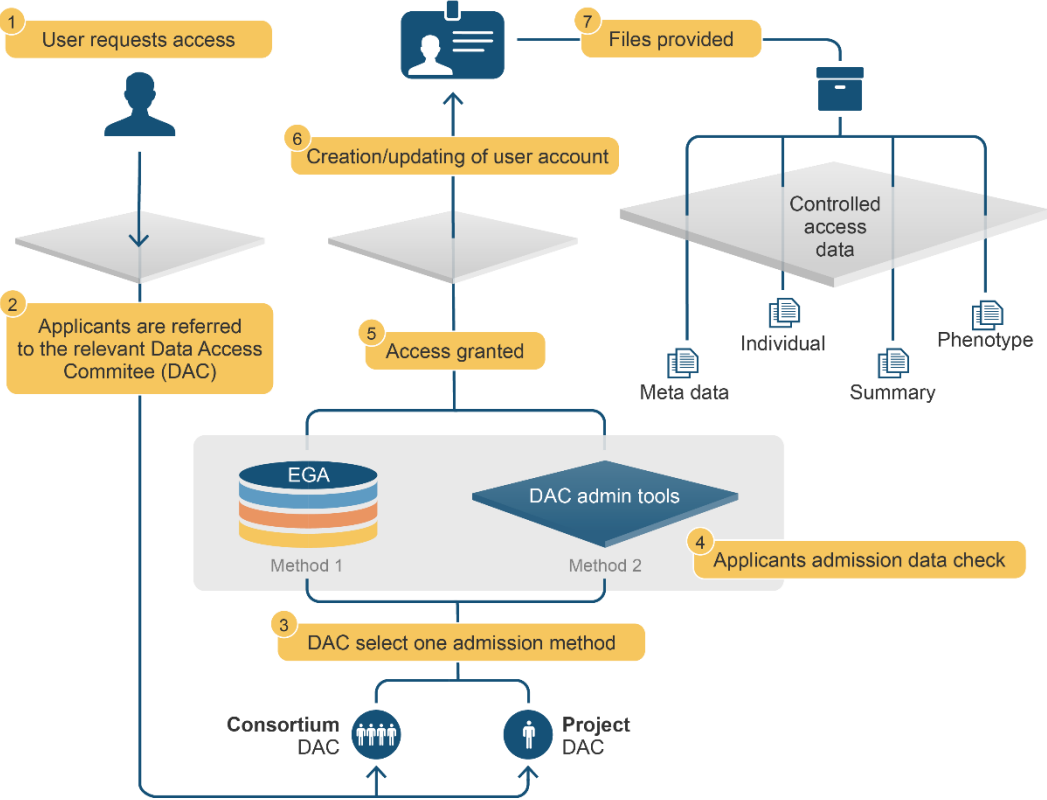


EGA Mission Statement

"The European Genome-phenome Archive (EGA) seeks to accelerate data sharing for human genetic and phenotype data, consistent with local ethical agreements and international standards, to enable molecular medicine."

What is the EGA?

The EGA is a resource for permanent secure archiving and sharing of all types of potentially identifiable biomolecular and phenotypic data resulting from biomedical research projects.



Data is provided by research centers and health care institutions.

Access is controlled by Data Access Committees.

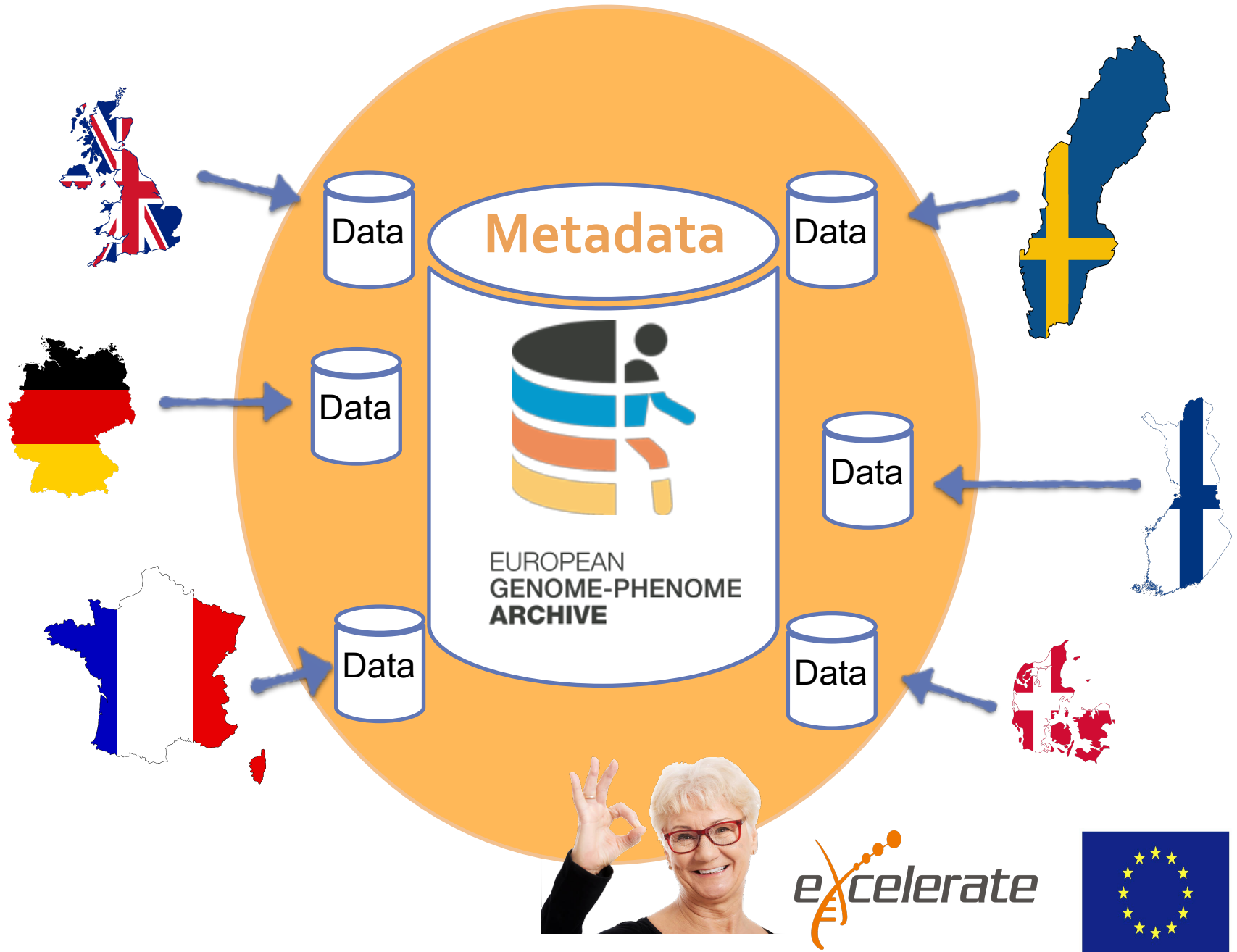
Data requesters are researchers from other research or health care institutions.

<https://ega-archive.org>



*ex*celerate







WP9

9.1.2

EMBL-EBI



D-1 Sensitive data archiving - Key Targets



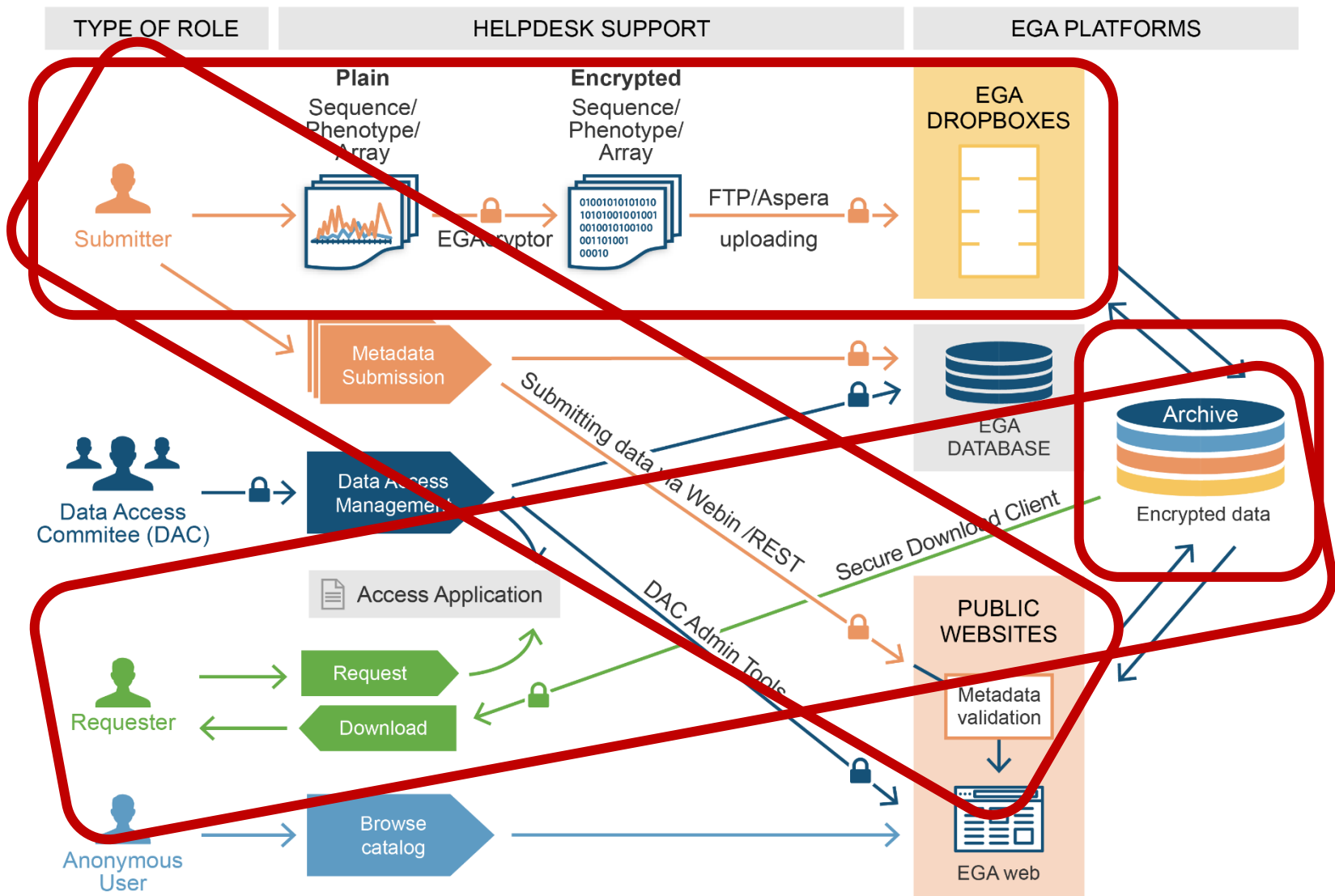
- Develop a **code base** for archiving of sensitive data
 - *Compatible with the needs of ELIXIR-Excelerate WP 9.1.2*
 - Packaged for convenient deployment
 - **Support core processes & other relevant processes**
 - “Data in” & “Data out” (through several mechanisms)
 - Federated querying
 - *Also for data that will **not** be exposed through EGA*
- Establishment of **service instances** in member nodes



Cross-border team

- Tryggve partners
 - ELIXIR Nodes: Finland, Norway, Sweden
- Central EGA
 - CRG (ELIXIR-ES) & EMBL-EBI

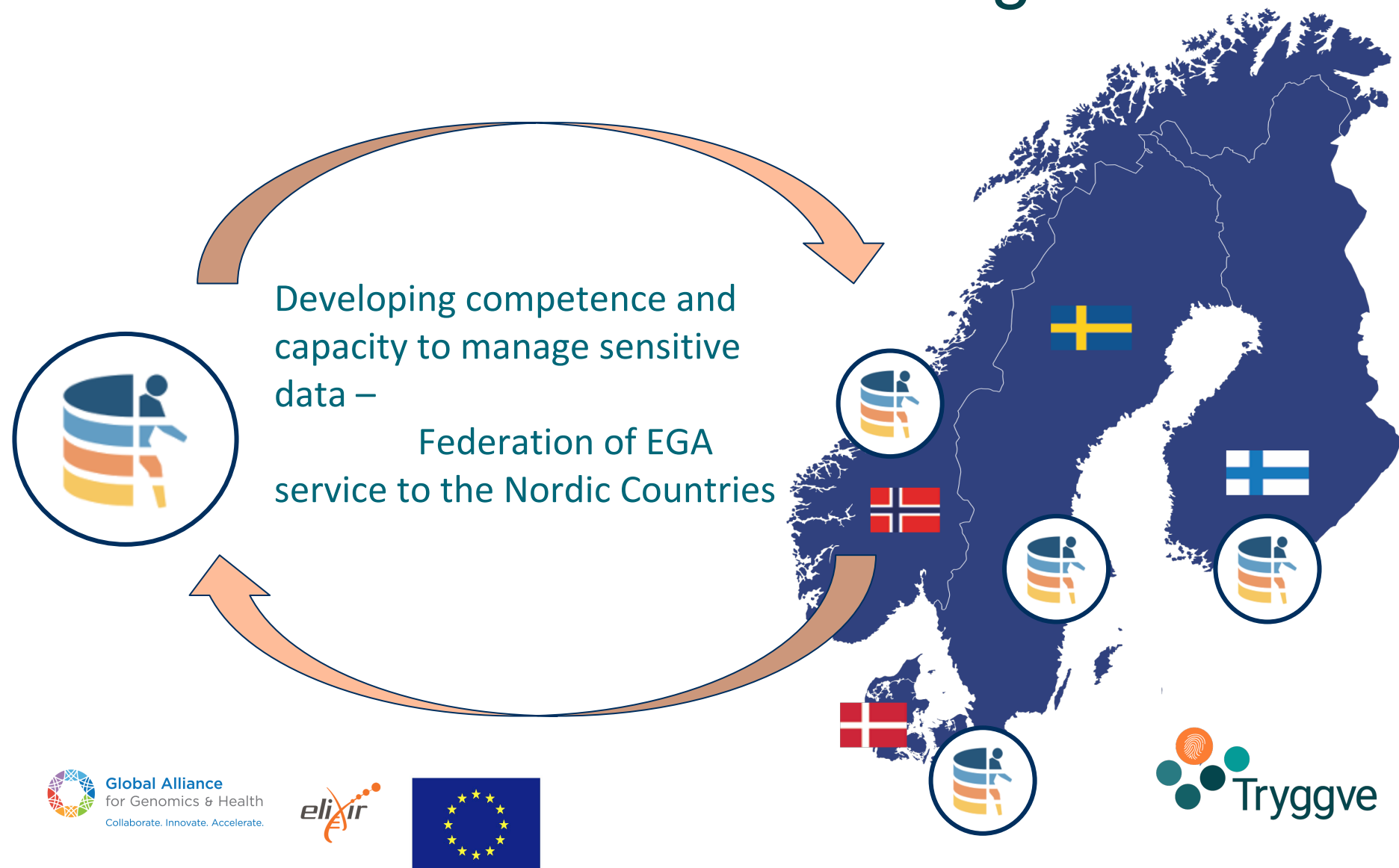




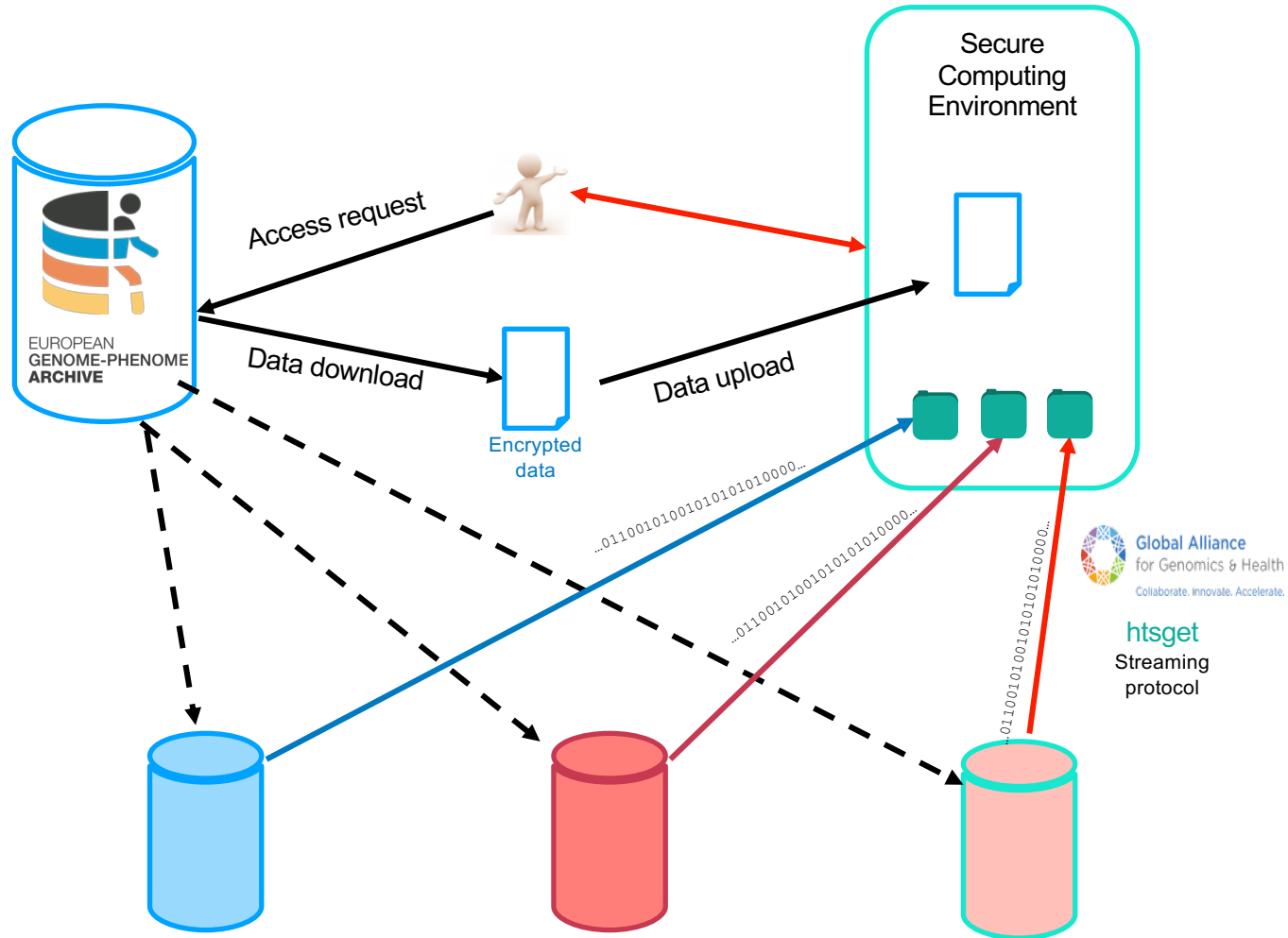
D-1 Sensitive data archiving - milestones

- **2018-12** – Deployment of core functionality code base in a production-like environment demonstrated - (2019-02)
- **2019-05** – At least one sensitive data archiving instance technically ready to join the Federated EGA

Nordic Vision for Data Management

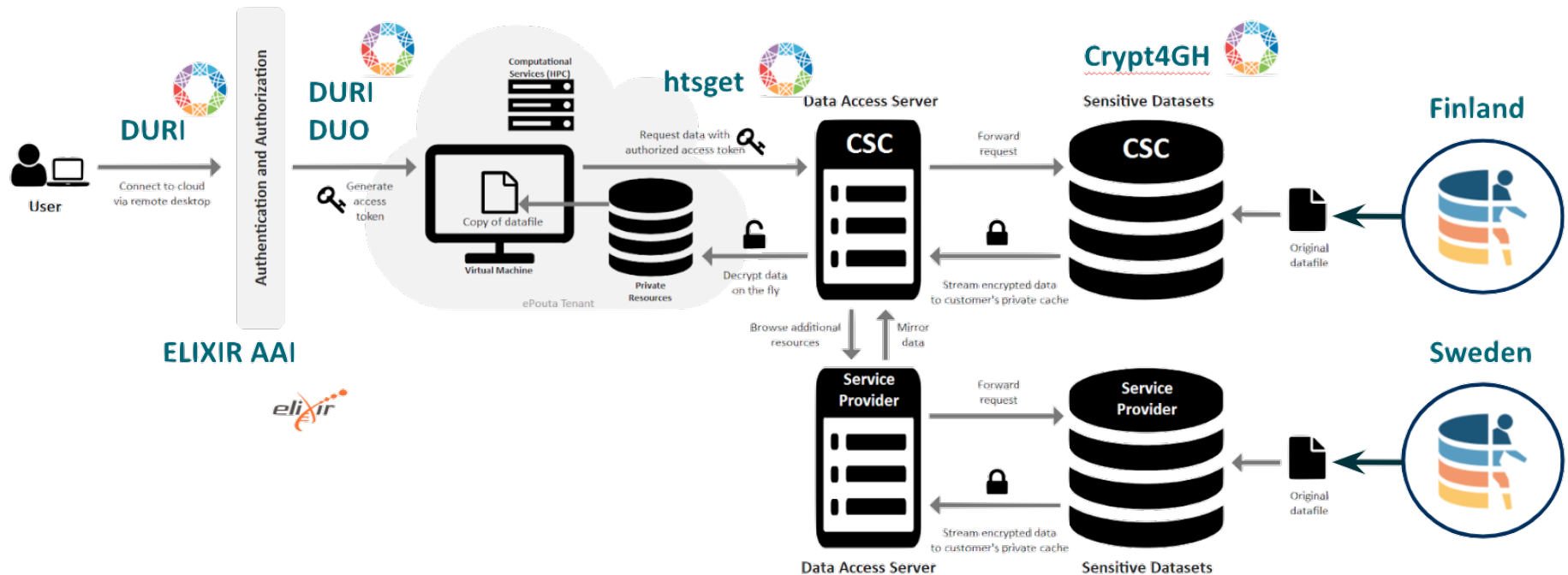


Future ways of accessing data



NorTwinCan use case

Possible technical solution



The GA4GH Mission



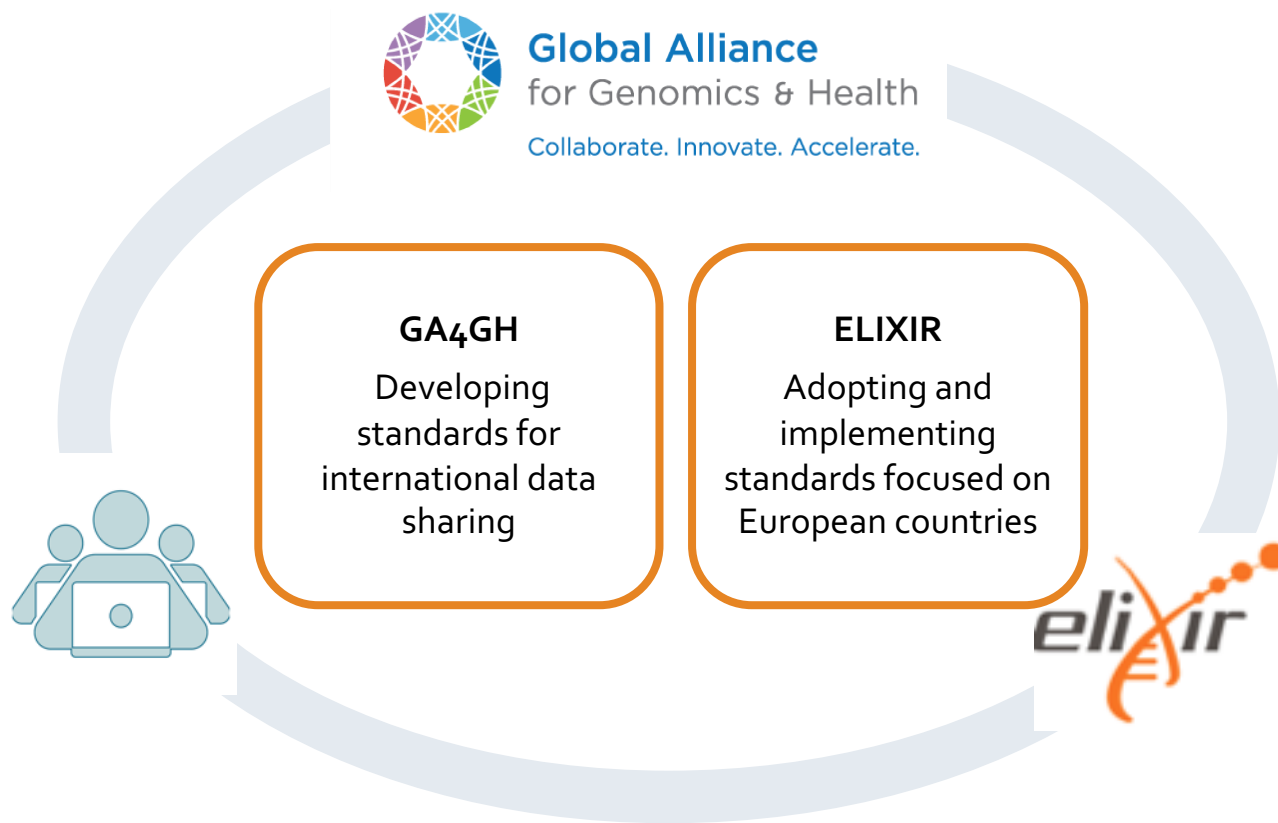
LAUNCH of GA4GH in 2013

The Global Alliance for Genomics and Health aims to accelerate progress in genomic science and human health by developing **standards** and framing policy for **responsible** genomic and health-related data **sharing**.

Core Principles of Data Sharing

- Enable international data sharing
- Promote sharing across the translational continuum (discovery research, clinical trials, clinics, diagnostic labs, industry)
- Encourages technology-enabled federated approaches (bring analysis to the data)
- Promote interoperability: scientific, technical, and ethical
 - Adopt standards, have transparent documentation
 - Standardized file formats, bioinformatics & variant calling protocols, variant annotation
 - Consent policies to ensure data can be shared internationally

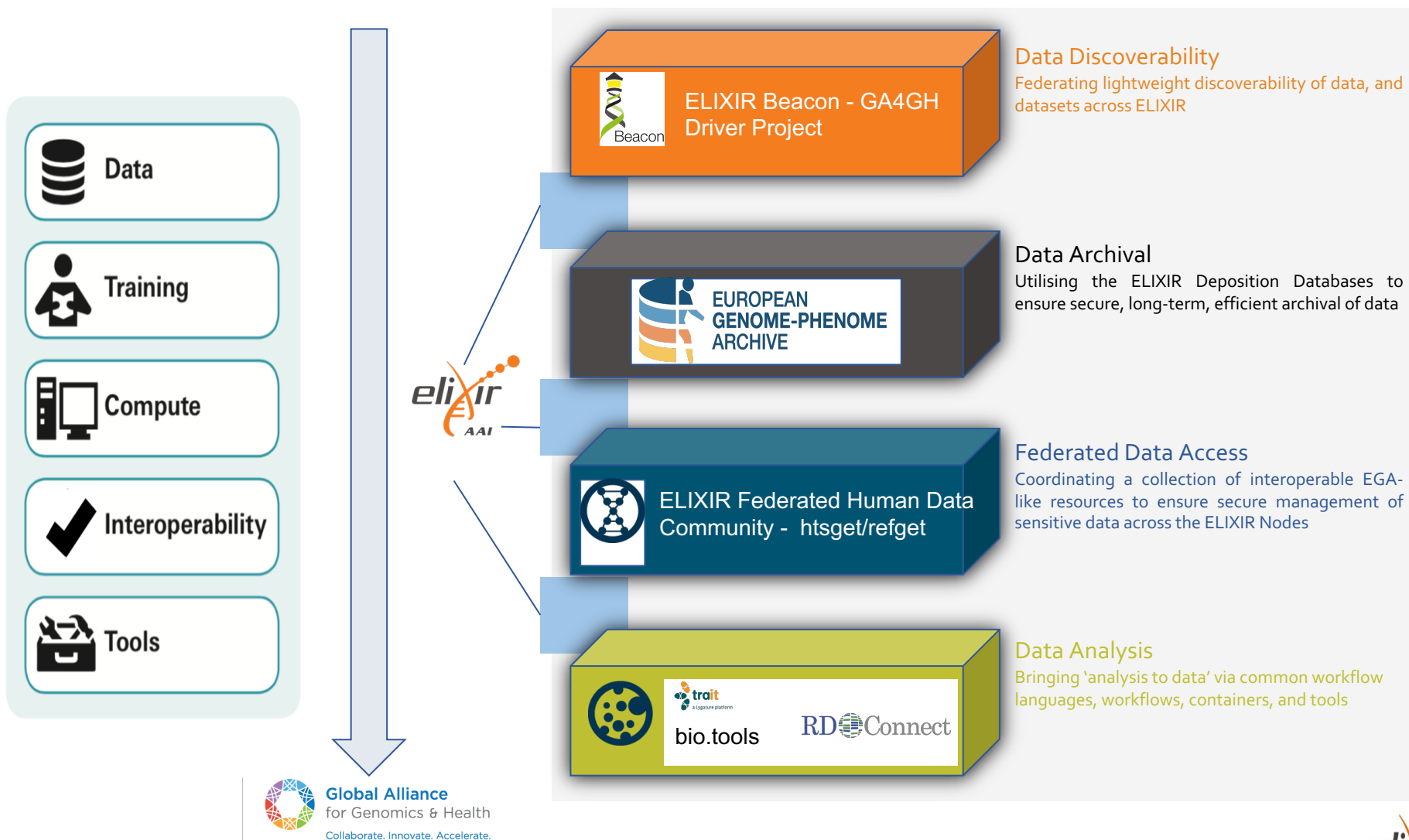
GA4GH and ELIXIR have formed strategic partnership



Working towards GA4GH standards, APIs and toolkits to be used throughout ELIXIR Nodes for human data discovery and access



ELIXIR Human Genomics & Translational Data - technically



ELIXIR Federated Human Data (2019-21)



ELIXIR Federated Human Data (2019-21)

- Many human data initiatives across Europe
- ELIXIR human data community to coordinate approaches
 - Metadata, phenotypes, AAI, ELSI, discovery



ELIXIR

- M
- EL
- CO
-

DECLARATION OF COOPERATION

Towards access to at least 1 million sequenced genomes in the European Union by 2022

Česká republika

and

Eesti Vabariik

and

Reino de España

