

The German Human Genome-Phenome Archive (GHGA)

A national research data infrastructure for human omics data

MIRACUM-DIFUTURE Symposium 2023 October 10, 2023

Oliver Kohlbacher GHGA Co-Director

In cooperation with



The GHGA Vision: Enabling Genomic Medicine



Main Aims:

Establishment of a nationally coordinated, interdisciplinary infrastructure integrating genome research and healthcare

- From Researchers for Research-

Funding: NFDI - The National Research Data Infrastructure Initiative



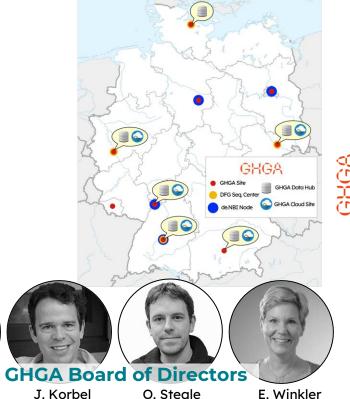
- * FAIR access to research data for all and across all domains of science.
- Funded with 900 M€ for 10 years
- 26+1 consortia selected for funding in 3 cohorts
- Wide variety of topics, including lifes sciences (NFDI4Health, NFDI4Microbiota, NFDI4Bioimage, NFDI4Immuno) with strongest interaction
- BASE4NFDI-Consortium starting in March 2023 ⇒ Development of common Base-Services



The GHGA Consortium



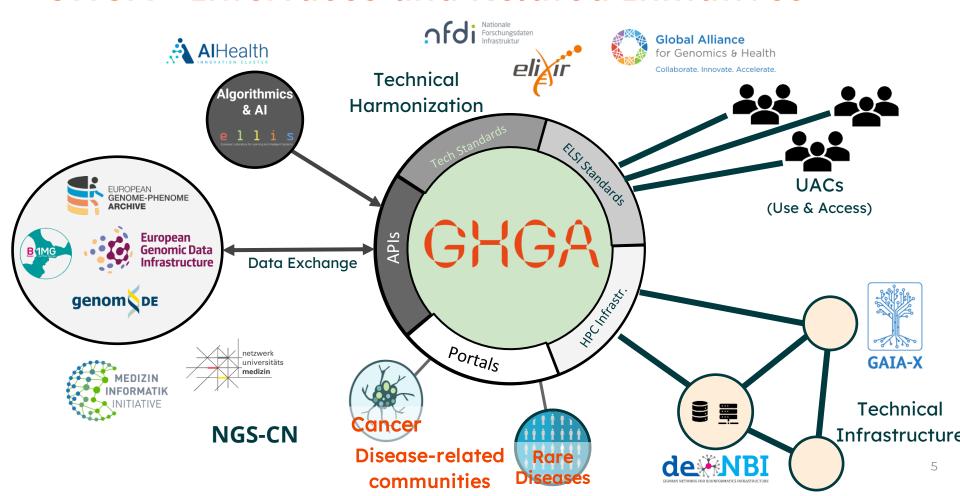
- Funded since 10/2020 as one of the initial nine first-round NFDI consortia
- Leading Institutions in genomic medicine
 - Representing 10 universities, 6
 Helmholtz & 5 other research
 institutions (17 funded partners)
- Network of seven data hubs co-located with major academic sequencing centers (TÜ, K, DD, KI, HD, M, B)
- Connected to national cloud infrastructure
 - ⇒ de.NBI cloud
- Funding from end of 2020 to 2025



O. Kohlbacher (Univ. Tübingen)

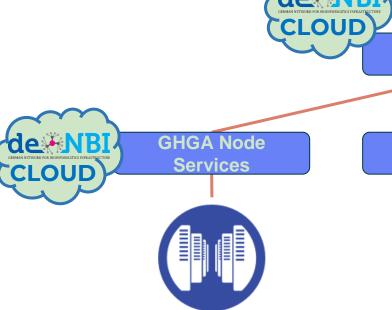
J. Korbel (EMBL) O. Stegle E. Winkler (DKFZ & EMBL) (Univ. Heidelberg)

GHGA - Interfaces and Related Initiatives



GHGA Architecture

- Federated (GHGA Central & Data Hubs)
- Cloud-native
- Infrastructure agnostic
- · Hexagonal microservice architecture



Data Hub A

GHGA Central Services

GHGA Node

Services

Data Hub B



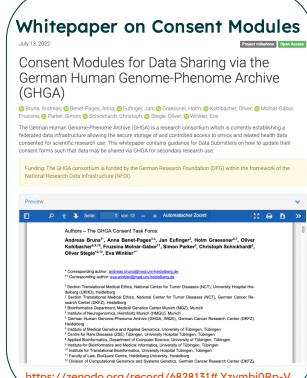


Project Status



Legal and Ethical Questions, Patient Engagement

Patient Engagement PaGODa Study: "Patientenperspektiven auf Governance-Strukturen von Datenbanken" Apondo, Winkler et al. White paper with concrete Explore task-specific & Involvement in Patient recommendations preference-sensitive Involvement Consultant Training Assessment of possible governance of patients within GHGA: models based on patient preferences on GHGA & financial



Legal Foundations



Molnár-Gábor, F., Kaldowski, L., Schenk, A., (2022): Alteinwilligungen im Forschungskontext., Zeitschrift für Datenschutz

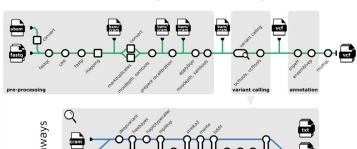


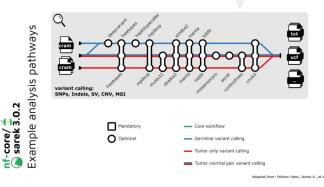
https://zenodo.org/record/6828131#.Yzvmbi0Rp-V

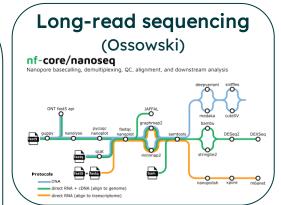
Workflow Standardization

Short-read variant calling

- Germline (Nahnsen)
- Somatic (Hübschmann)







Aberrant Expression Aberrant Splicing Mono-allelic Expression

RNA-seq Variant Calling

RNA-seq for RD (Gagneur)

Integration into communities

 Co-development with nf-core



- Training (Stanford, Broad)
- Benchmarking (NGS-CN)



Outreach and Training

Also on Spotify, YouTube & iTunes ghga.de/codedeslebens







Stay up to date:
GHGA Newsletter
See www.ghga.de or









Training

Webinar Series





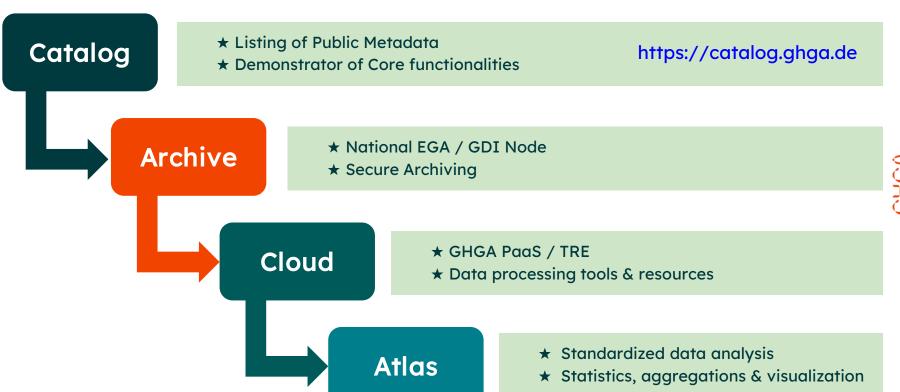
Courses / Collaborations



FAIR in (biological) practice

30.05 - 02.06.2022

Project Phases and Features















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Who we are \rightarrow How we work \rightarrow Jobs \rightarrow

The German Human Genome-Phenome Archive

We are building a **secure national omics data infrastructure**, enabling the use of human genome data for research purposes while preventing data misuse.

Our Mission

GHGA Metadata Catalog

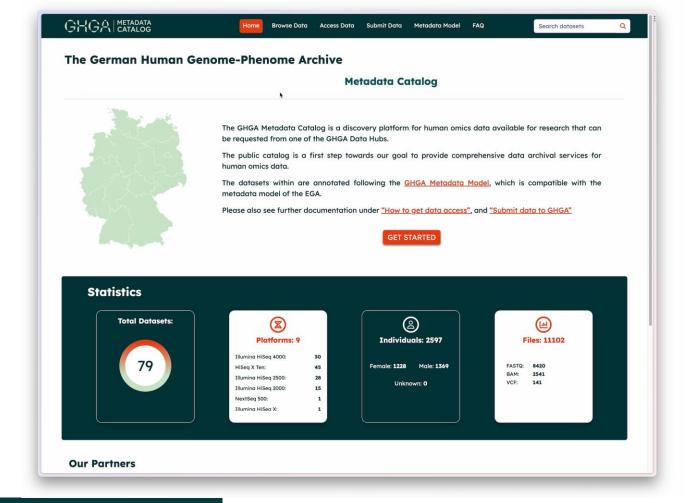
A public frontend for the discovery of human omics study data from German research institutions.

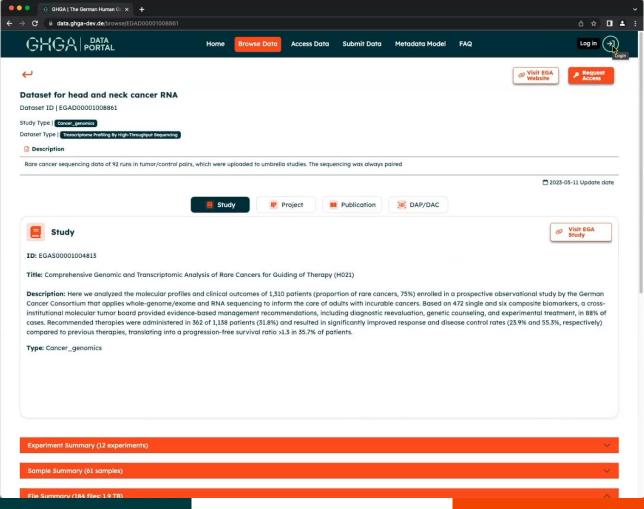
Consent Tools

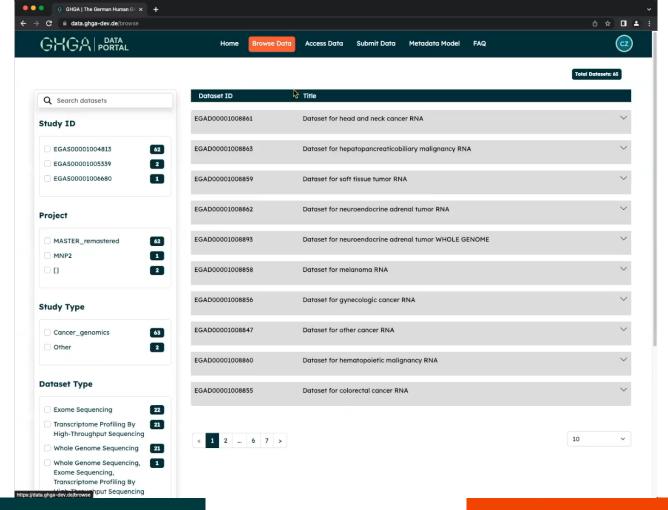
GHGA has developed different tools to help clinicians, researchers and institutions wanting

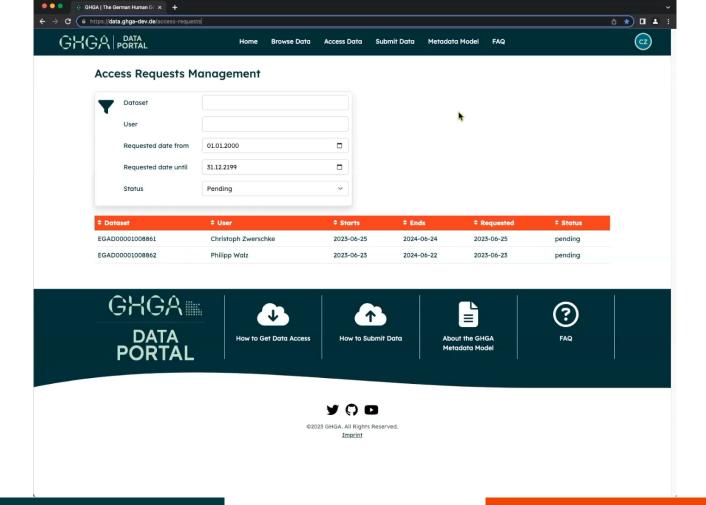
GHGA Lecture Series

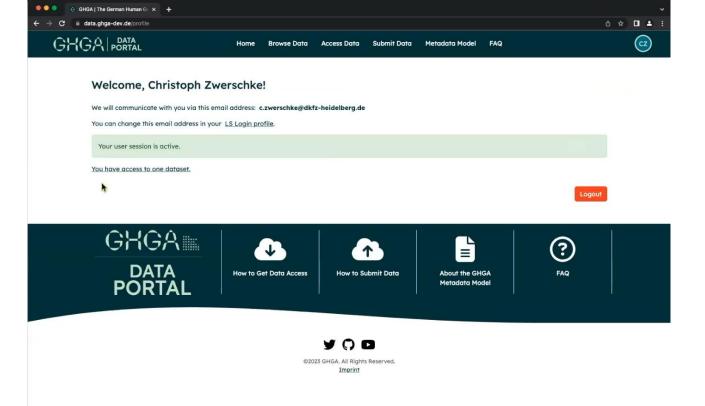
"Advances in Data-Driven Biomedicine" diving into fascinating world of datadriven medicine and their





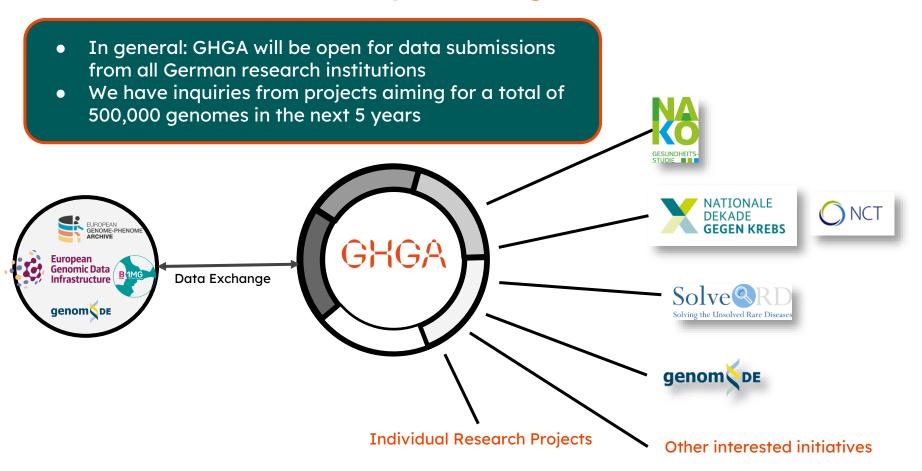






Perspective

Projects that are planning to use GHGA



Perspective

Archive

- GHGA Archive will be in production in Q1/2024
- Data hubs are already starting to store genomic raw data but not yet within the legal framework of GHGA
- Operational status of archive will then require just a change of metadata no additional data transfer

Cloud

- Secure processing environments (SPE) suitable for person-related data are available through the de.NBI Cloud
- Data transfer processes are being established to re-encrypt and stage data

genomDE

- Several GHGA Data Hubs are preparing to act as Genome Data Centers within genomDE
- Local legal constructs (contract data processing) are being prepared to also process non-consented data

Thanks to the whole GHGA team!



Partner Institutions









German Biobank Node











































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