# Mallory A Freeberg, PhD

Who I Am Community Builder. Data Sharing Expert. Bioinformatics & Metadata Enthusiast. Open Science Advocate.

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Professional Experience

#### EMBL European Bioinformatics Institute, Cambridge, UK

Coordinator, European Genome-phenome Archive

Jun 2021 - present

- Contribute expertise to help establish the Federated EGA Network for transnational discovery of and access to sensitive human data (blog post)
- Engage with external partners and stakeholders to ensure data submission, discovery, and distribution services align with community standards
- Manage up to14-person team responsible for operating production data archive services (publication)

Project Lead, UK Biobank, European Genome-phenome Archive

Aug 2019 - May 2021

- Oversaw data flow to archive 500,000 whole human genomes totalling 12PB of data
- Built and led 3-person team to successfully distribute 4PB of data to cloud-based analysis platform

Senior Bioinformatician, Human Cell Atlas Data Coordination Platform

Oct 2017 - Jul 2019

- · Coordinated data wrangling work for 6-person team across UK- and US-based institutions
- · Performed curation and ingestion of flagship HCA datasets into the Platform
- Developed HCA metadata standard for cellular-resolution transcriptomics data (publication)

# Johns Hopkins University, Baltimore, MD USA

Trainer, Galaxy Training Network, The Galaxy Project

Jan 2017 - Sep 2017

- Designed and delivered seven Galaxy training workshops in three countries (publication)
- Created training materials and computational workflows to support reproducible transcriptomics research (contributions)

Postdoctoral Research Fellow, Department of Biology

Jun 2015 - Sep 2017

- Investigated post-transcriptional gene regulatory networks using neural networks
- · Piloted Oxford Nanopore direct RNA sequencing protocols to catalog transcriptomes

Education

## PhD Bioinformatics, University of Michigan, USA

May 2015

Department of Computational Medicine and Bioinformatics

Dissertation: Computational analysis of the post-transcriptional gene regulatory network (thesis)

## BSc Bioinformatics, Saint Vincent College, USA

May 2009

Herbert M. Boyer School of Natural Science, Mathematics, and Computing

Dissertation: Functional annotation of non-coding elements in the Amphioxus genome

Selected Publications Corvo A, Matalonga L, Spalding D, et al. (2023) Remote visualization of large-scale genomic alignments for collaborative clinical research and diagnosis of rare diseases. *Cell Genomics*. DOI:10.1016/j.xgen.2022.100246

Thakur M, Bateman A, Brooksbank C, Freeberg MA, et al. (2022) EMBL's European Bioinformatics Institute (EMBL-EBI) in 2022. *Nucleic Acids Res.* DOI:10.1093/nar/gkac1098

Freeberg MA, Fromont L, D'Altri T, et al. (2021) The European Genome-phenome Archive in 2021. *Nucleic Acids Res.* DOI:10.1093/nar/gkab1059

Rehm HL, Page AJH, Smith L, et al. (2021) GA4GH: International policies and standards for data sharing across genomic research and healthcare. *Cell Genomics* 1. DOI:10.1016/j.xgen.2021.100029

Füllgrabe A, George N, Green M, et al. (2020) Guidelines for reporting single-cell RNA-seq experiments. *Nat Biotechnol* 38:1384-6. DOI:10.1038/s41587-020-00744-z (arXiv preprint DOI:1910.14623v1)

⇒Full list of publications can be found on Google Scholar.

Selected Invited Talks Federated EGA: Providing global discovery and access for sensitive human data. *ISMB/ECCB*; Jul 2023; Lyon, FR (slides)

The Federated EGA: Global discovery and access for sensitive human data. *GA4GH 10th Plenary*; Sep 2022; Barcelona, ES (slides)

FAIR principles and promoting openness in the life sciences. *NIHR Statistics Group Workshop*; Feb 2022; virtual (slides)

The European Genome-phenome Archive Metadata Model. *GHGA Workshop on Metadata in Biomedical Genome Research*; Jun 2021; virtual (slides)

Selected Conferences & Workshops Demonstrating federated EGA services for sensitive data discovery and access. *ELIXIR All Hands*; Jun 2023; Dublin, IE (workshop)

Resource considerations for establishing and operating a federated human data node. *ELIXIR All Hands*; Jun 2022; Amsterdam, NL (poster)

Managing single cell transcriptomics data. EMBL-EBI Training Course; Jul 2019; Cambridge, UK (course)

The Data Coordination Platform: Making the Human Cell Atlas data easily accessible. *The Biology of Genomes*; May 2018; Cold Spring Harbor, NY USA (poster)

Leadership Roles Co-lead, GA4GH "Future of VCF" Working Group (Mar 2022 - Present)

Guide discussions and assessment of solutions for scaling the Variant Call File format for representing genetic variants across large-scale studies

Organiser, ELIXIR Federated Human Data Community (Feb 2022 - Present)

Set agenda and priority topics for FHD Community meetings, co-organise FHD Community Day event, coordinate federate human data activities across ELIXIR members

Chair, Federated EGA Operations Committee (Jan 2021 - Present)

Lead Committee in fulfilment of responsibilities, coordinate with Committee members to make progress towards shared goals and milestones

Teaching & Mentoring

Mentor, general (2020 - Present)

Mentor staff and peers to support them achieving their career and professional development goals

Mentor, Open Life Science (Jul 2020 - Present)

Mentor participants on open science practices as part of a 16-week mentorship and training program to empower open science ambassadors

UK STEM Ambassador (Jan 2018 - Present)

Give talks, demonstrate science experiments, share STEM career guidance

Training Courses ELIXIR LEadership And Diversity mentoring programme

Managing Managers

Emotional Intelligence and Mindfulness

Talented Women's Impact Programme

EICAT Complementary Scientific Skills Training - Scientific Project Management

Handling Conflict and Difficult Conversations

2023-present
2021
2022
2020
2019-2020
2019-2020
2019-2020
2019