

Harmonising the way human omics data is described in different settings – making data sharing easier – with the GHGA Metadata Model

<https://www.ghga.de/resources/metadata-model>



Helping data producers to describe their submitted data and to retrieve data of interest



FAIR metadata by utilizing established and widely used ontologies and vocabularies



Implementation of GHGA Metadata Catalog using the Linked Data Modelling Language (LinkML)



Data submission in a standardised manner using GHGA's submission spreadsheet

TRACK ON GITHUB

The GHGA Metadata Model is stored and accessible in a public GitHub repository. Here, you can track every new release of the model.



Subscribe to our newsletter to learn about new metadata developments.

