

1 - Binding letter of intent or non-binding letter of intent

This is a **binding** letter of intent as an advance notification for a proposals in 2020

2 - Formal details

Planned name of the consortium

National Research Data Infrastructure for Immunology¹

Acronym of the planned consortium

NFDI4Immuno

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¹ Previously named "NFDI for Adaptive Immune Receptor Repertoires" (NFDI4AIRR)

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3 - Objectives, work programme and research environment

Research area of the proposed consortium (according to the [DFG classification system](#))

Immunology (204-05, 205-18) and related research areas

- Hematology / Oncology (205-14)
- Microbiology / Virology (204-xx)
- Neuropathology (206-07)
- Pathology (205-06)
- Veterinary medicine (207-11, 207-12)
- Theoretical Biology (201-07)

Concise summary of the planned consortium's main objectives and task areas

The immune system plays a fundamental role in health and disease and efficiently protects vertebrate hosts from infections and cancer. However, failures in its regulation can cause autoimmunity, allergy, immunodeficiencies and lymphoid malignancies. To perform the critical task of self/non-self recognition, the adaptive immune system utilizes billions of randomly generated immunoglobulins/antibodies and T-cell receptors (hereafter: adaptive immune receptors), which are expressed by cells of the B-cell and T-cell lineage, respectively. The phenotype, activation state and histo-anatomic location of these cells, as well as the entireness of the adaptive immune receptors of an individual host, commonly referred to as AIRR (adaptive immune receptor repertoire), are reflective of the key processes within the immune system: Diversification, selection, antigen recognition and clonal expansion. A comprehensive understanding of these processes will facilitate mechanistic insights and allow the development of diagnostic markers and novel therapeutic strategies.

To this end, it is necessary to obtain the capability to combine data and metadata from diverse experimental technologies, such as sequencing, cytometry and imaging, which provide complementary observations of these processes. Thus, **the main objective of NFDI4Immuno is to build a network of federated repositories for data describing the state of the immune system in close cooperation with the German immunological community and to provide tools and services that will facilitate integrated data analyses across these repositories.** Over the funding period, NFDI4Immuno will build a federated infrastructure for AIRR and cytometry data, expand existing data models to cover microscopy and transcriptome information and initiate the integration of proteomic data. The resulting network can be expanded to handle additional data types in the future, as determined by the needs of the community. The detailed steps for each of these data types are as follows:

Build a network of federated AIRR-seq repositories: "AIRR-seq" is a heterogeneous set of NGS-based technologies that provide information on the AIRR, i.e., the highly variable regions of adaptive immune receptors. Several partners in the consortium have a long-standing expertise in the generation and analysis of this data type and are among the founding members of the international AIRR Community. As such they have an established involvement in the community's

efforts to provide standards and promote FAIR practices for AIRR-seq data. Building on this expertise and the existing software stack for sharing AIRR-seq data (iReceptor), we plan to roll-out the initial set of data repositories during the first two years. This will create the basic infrastructure for integrating other data types. Data - both from the consortium members and third parties - will be curated and made publicly available throughout the funding period. We plan to give special attention to data from species other than human and mouse, e.g., livestock and non-human primates, which are often not appropriately covered by biomedical repositories.

Enable repositories to manage cytometry data: Cytometry (i.e., both flow and mass cytometry) is the current gold standard in immunology to describe, define and isolate immune cells and provides a rich phenotypic description at single-cell resolution. In spite of its pervasiveness and importance, there is currently no generally accepted way to share these often large, heterogeneous and high-dimensional data sets. To fully utilize the wealth of information in cytometric data, NFDI4Immuno will develop and deploy software stacks that facilitate its storage, annotation and analysis, while providing for close integration with associated AIRR-seq data sets. This is a critical and future-oriented activity, as with the current advent of commercially available platforms that use DNA barcoding of cells and surface markers to combine cytometry with AIRR-seq (e.g., 10X Chromium), the borders between these two technologies become increasingly fluid.

Integrate other data types: The spatial and temporal information provided by microscopy complements AIRR-seq and cytometry data, while transcriptome data will allow a broader assessment of cellular phenotypes and activation states. Our main focus here will be the extension of the immunological data model to facilitate thorough integration of these data types and cross-referencing to data sets stored by other NFDI consortia (NFDI4Bioimage and GHGA, respectively) or the European Nucleotide Archive (ENA).

Initiate developments for proteome data: The majority of immune effector molecules are proteins, which are however only superficially measured by the methods described above. We will therefore initiate pilot projects with experts in the field of proteomics and antigen receptor reactivity to evaluate the inclusion of such data types in the second half of the funding period.

These data-centric activities will be translated into the following five task areas:

- *Data curation* - Provide consistent metadata annotation to existing data sets and make them available through the consortium's repositories. Additionally, organize the consortium's Quality Assurance (QA) Panel, which will develop and maintain guidelines for data and metadata quality and perform periodic audits among the federated repositories.
- *Interoperability* - Harmonize metadata descriptors and ontologies with other NFDI consortia to facilitate fast and easy cross-consortia queries for data sets. Develop and standardize programmatic interfaces (APIs) for advanced computational workflows and exchange and interconnection of data within and beyond the NFDI.
- *Community involvement and outreach* - Disseminate and promote the adoption of FAIR practices by the immunological community, to support the cultural change towards Open Science. Engage stakeholders outside of the national scope of NFDI with the long-term goal to connect NFDI4Immuno at the international level, e.g., with EOSC, IEDB, ImmPort, Human Vaccines Project, iReceptor Plus and the AIRR Community.

- *Repository DevOps* - Develop and operate the consortium's federated repository infrastructure.
- *Application development and support* - Build novel user-friendly applications both for data analysis utilizing NFDI4Immuno's comprehensive federated infrastructure, as well as simplified (meta-)data submission to NFDI4Immuno repositories. Provide support and training to users with a primarily experimental background enabling them to appropriately utilize the resources provided by the consortium.

The planned governance structure will consist of the Executive Board (Speaker + two co-Speakers), the General Assembly (all partners) and the Community Council, a committee representing the diverse NFDI4Immuno user communities and their needs. The Executive Board will manage the organizational, financial and legal concerns of the consortium. It will implement a unified strategy to ensure the long-term availability and usability of the consortium's data and code.

Brief description of the proposed use of existing infrastructures, tools and services that are essential in order to fulfill the planned consortium's objectives

Hardware and network infrastructure: In general terms, NFDI4Immuno is designed as a federated repository infrastructure, which will substantially lower the requirements for storage and compute capacity at the individual partner sites. Therefore, it is by default expected that partners can provide these resources as part of their contribution to the consortium. However, especially for large third-party data sets, we are planning to utilize centralized storage services like the Helmholtz Data Federation (HDF). These data sets might also require additional network bandwidth as well as some compute resources. Therefore, the consortium is currently in the process of estimating the potential resource requirements and is in contact with the Helmholtz Infrastructure for Federated ICT Services (HIFIS) for further input regarding this topic.

Software: NFDI4Immuno will build on the freely available iReceptor software stack, the core of which is licensed under LGPL3 (GNU Lesser General Public License). Notably, the consortium partners at DKFZ are currently part of the Horizon 2020-funded "iReceptor Plus" project, that aims to substantially enhance iReceptor's capabilities over the next three years. Therefore, the necessary know-how for further development and operation of the platform will already be present in NFDI4Immuno.

Biobanking: Many pathological processes involve the immune system, thus biobanked samples of diseased tissues are an attractive resource for immunological analysis. Therefore, we consider it valuable to link immunological data at NFDI4Immuno that derives from samples of the German Biobank Alliance (GBA) with its respective metadata stored by the GBA. To this end, we will work together with the German Biobank Node (GBN), which coordinates the GBA and is part of BBMRI-ERIC, to develop and implement such linkage on the sample level.

Interfaces to other proposed NFDI consortia: brief description of existing agreements for collaboration and/or plans for future collaboration

NFDI4Microbiota: The continuous interaction between the immune system of vertebrate hosts and their microbiota makes this consortium a natural partner. To facilitate the parallel analysis of both, we plan on harmonizing metadata and ontologies describing the host, as well as identifying and enhancing formalized descriptions of sampling procedures.

NFDI4Agri: Global health requires maintaining a close interaction between human health and veterinary disciplines. For zoonotic infections, including current pandemics, intertwining disease epidemiology and immune response patterns helps understanding infection dynamics and mammalian host defense. We will cooperate with NFDI4Agri in harmonizing metadata structures and ontologies and will develop data curation systems for domain-specific (i.e., veterinary immunological and epidemiological) metadata.

InnoMatSafety: InnoMatSafety is the other consortium besides NFDI4Immuno that plans to manage a significant amount of cytometry data. Therefore, we aim to establish a platform for regular knowledge exchange in respect to the technical aspects of data management and storage. In the best case, this could lead to a shared backend storage infrastructure with domain-specific metadata layers on top of it. In addition, we will evaluate the potential shared requirements for domain-specific (i.e., immunological) metadata.

NFDI4Bioimage: Microscopy data is one of the key primary data types in immunological research. As NFDI4Bioimage aims to provide generic and domain-spanning tools and services for the storage and management of microscopy and photonics-based imaging data, NFDI4Immuno is interested in utilizing these resources. To ensure the findability of imaging data describing the immunological processes that are the central focus of NFDI4Immuno, we will cooperate in developing and harmonizing metadata structures and ontologies.

GHGA: We consider GHGA as the NFDI's central infrastructure for generic sequencing data from human subjects and will interact closely with regard to the deposition of such data (e.g., transcriptome) within their repositories and cross-referencing and accessing information between our consortia. Generic sequencing data outside of the scope of GHGA (i.e., majority of studies with non-human subjects) will be linked via ENA and we hope to develop the necessary programmatic abstraction layers together with GHGA.

NFDI4Health: The centralized metadata storage and catalog proposed by NFDI4Health could be a complementary top-layer to NFDI4Immuno's federated data repositories. To this end, we plan to develop the required interfaces for data exchange and will make our expertise in domain-specific metadata and ontologies available to the NFDI4Health consortium.

NFDI4RSE: The development of novel applications for data analysis and management and the continuous maintenance of existing software tools are central components of NFDI4Immuno. Therefore we plan to collaborate closely with NFDI4RSE to implement best practices ensuring the quality and sustainability of our code base. Furthermore, we welcome NFDI4RSE's proposition to establish resources to teach "Software Carpentry", as we consider this a critical step to raise overall bioinformatic capabilities of the immunological community.

NFDI4Life Umbrella: We share NFDI4Life's assessment that the creation of an intermediary structure for life science-specific coordination will be of benefit to the consortia involved and the NFDI as a whole. Therefore we support the proposal to establish such a coordinative council.

4 - Cross-cutting topics

Please identify cross-cutting topics that are relevant for your consortium and that need to be designed and developed by several or all NFDI consortia.

We consider the cross-cutting topics identified in the Leipzig-Berlin Declaration² as a common work assignment for the whole NFDI, which will also constitute an important process of organizational development. Therefore, NFDI4Immuno explicitly supports the task areas described in the Declaration and their suggested implementations. We identified the following cross-cutting topics and services as being especially relevant for our consortium:

- Open data and metadata standards
- Harmonization and standardization of ontologies
- Supporting FAIR and Open Science practices within the communities
- Data literacy and RDM curricula for scientists
- Guidelines for quality assurance
- Free and Open Source research software
- Management of personal and/or confidential data
- Common authentication and authorization infrastructure
- Integration of the NFDI into European RDM infrastructures (EOSC)

Please indicate which of these cross-cutting topics your consortium could contribute to and how.

NFDI4Immuno can contribute its expertise regarding the development of open community data standards, as well as the extension and harmonization of domain-spanning ontologies. Several members of the planned consortium have already engaged in similar activities in the context of data standards development by the AIRR Community.

² Bierwirth M *et al.* Leipzig-Berlin-Erklärung zu NFDI-Querschnittsthemen der Infrastrukturentwicklung.
DOI: [10.5281/zenodo.3895209](https://doi.org/10.5281/zenodo.3895209)