Plan Overview

A Data Management Plan created using DMPTool

DMP ID: https://doi.org/10.48321/D19061

Title: The University of Chicago Consortium for Food Allergy Research Clinical Research Center

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Funder: National Institutes of Health (nih.gov)

Funding opportunity number: RFA-AI-22-076

Grant: https://grants.nih.gov/grants/guide/rfa-files/RFA-AI-22-076.html

Template: NIH-Default DMSP

Project abstract:

Food Allergy (FA) is a common chronic condition associated with life-threatening allergic reactions in all age groups. As an estimated 8% of children and 11% of adults in the United States are affected by FA, a significant need exists to expand the body of knowledge around prevention, diagnosis and treatment across all age groups. One of the most critical needs in FA is the development of minimally and/or non-invasive diagnostic tests and algorithms that provide both highly sensitive and highly specific results. The most commonly used tests for FA, percutaneous skin prick testing (SPT) and allergen-specific IgE (sIgE), often fail to accurately predict clinical reactivity. The doubleblind placebo controlled oral food challenge (DBPCFC) therefore remains the gold standard test for confirming FA, determining reactivity thresholds, and understanding reaction severity. The DBPCFC has several limitations, however, as it requires hours to complete, causes significant morbidity, and has substantial site to site variability despite efforts to standardize protocols. These limitations result in high procedural costs, long wait times, and paralyzing anxiety that make this test impractical in many clinical settings. Without accurate, inexpensive, and widely available diagnostics, FA clinical trials are impeded by the high associated cost and reluctance by potential volunteers to incur the added risk. In addition, the accuracy of epidemiologic studies is limited. Thus, we believe developing novel diagnostic tools that accurately detect the presence of disease, the threshold of reactivity, the severity of reactions, the response to therapy and the early detection of anaphylaxis is an essential step to improving the lives of individuals with FA. The objective of this proposal, therefore, is to develop novel diagnostic methods for IgE-mediated FA through already established collaborations at the UChicago in cooperation with and to support the Consortium for Food Allergy Research (CoFAR). The central hypothesis is that repurposing established platforms with track records of success developed by scientists at UChicago outside the field of FA will allow for the rapid identification of novel biomarkers of FA and anaphylaxis providing the fundamental first step to the development of

improved FA diagnostic platforms. Further, we anticipate the generation of large datasets in our aims and believe the creation of a FA data commons to organize and share these datasets, as well as all datasets generated by CoFAR, is fundamental to efficient discovery in FA. In order to achieve our objective, we will perform highly sensitive 5mC profiling utilizing LABS-seq on cfDNA from a cross-sectional cohort of human subjects with peanut or egg allergy and use non-invasive laser Doppler to measure cutaneous blood flow during SPT and during OFC and determine when changes in blood flow occur with respect to allergen SPT and food doses. Finally, we propose the creation of a FA data commons across CoFAR sites to store all data generated from the Consortium.

Start date: 01-01-2024

End date: 12-31-2030

Last modified: 01-19-2024

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The University of Chicago Consortium for Food Allergy Research Clinical Research Center

Data source	Number of Subjects	Condition	File Format	File Size (approx)
LABS-seq	130	Peanut Allergy	.fastq and .txt	500 gigabytes
LABS-seq	130	Egg Allergy	.fastq and .txt	500 gigabytes
Doppler ultrasound	250	Food Allergy	.vmr and .txt	40 gigabytes

In this proposed project, the cleaned, clinical/demographic spreadsheet data for all variables will be shared openly. The rationale for sharing only cleaned data is to foster ease of data reuse.

All LABS-seq and Doppler data will be preserved and shared.

To facilitate interpretation and reuse of the data, a README file and data dictionary will be generated and deposited into a repository along with all shared datasets. The data dictionary will define and describe all variables in the dataset.

In addition, protocols and data collection instruments will be shared and linked to the relevant datasets.

Raw seq (.fastq) is a simple text file and can be opened with a text editor. The following R analytics packages are openly available to reuse the genomic data include, EnrichR, ClusterProfiler, Reactome PA, and MethylKit. Other associated analytic code will be provided through GitHub. Doppler (.txt) data can be analyzed in Excel or MatLAB.

Data sharing policies will adhere to the principles of FAIR (Findable, Accessible, Interoperable, and Reusable) data management.

Epigenomic data will follow MINSEQ-E

Doppler data will follow best practices,

Epigenome data will be stored in GEO.

If our proposed NIAID Food Allergy Data Commons is funded, our Doppler data will be permanently housed in this Data Commons.

If our proposed NIAID Food Allergy Data Commons is not funded, our Doppler data will be deposited and shared via Knowledge@UChicago (https://knowledge.uchicago.edu/), which provides metadata, ensures long-term access, and registers a digital object identifier (DOI) for each dataset to facilitate discoverability and citation. Additionally, the dataset(s) will be openly licensed and made publicly available as soon as possible or at the time of associated publication. As the University of Chicago's institutional repository, Knowledge@UChicago is supported collaboratively by the University's Library and IT Services. It is built on a cloud-based platform maintained by a service provider named TIND. Knowledge@UChicago uses an open archival information system (OAIS) compliant approach to preservation, which is complemented by fixity checking, redundancy backup, and storage of archival packages on geographically separated servers.

Data will be retained for a minimum of 10 years following project completion, in accordance with the University of Chicago's data retention policy. After this period, data may be archived in institutional repositories or other appropriate long-term storage facilities to ensure continued access and preservation.

The data will be made identifiable in the specified data repositories via persistent unique identifiers and multiple keywords to maximize search results. Dataset DOIs will be included in associated publications.

Data will be made available at the time of associated publication. The data deposited in GEO and the Food Allergy Data Commons (upon creation) will be available indefinitely. Data stored at the Center for Research Informatics at the University of Chicago will be stored for a minimum of three years.

There are no anticipated factors or limitations that will affect the access, distribution or reuse of the scientific data generated by the proposal.

Access to shared data will not be controlled.

All shared data will be deidentified

The PI, Dr. Christina Ciaccio, in collaboration with coinvestigators, will be responsible for overseeing the data management plan and ensuring compliance during the project. The PI will be responsible for updating and revising the Data Management and Sharing Plan when necessary and reporting compliance annually.

Planned Research Outputs

Dataset - "Doppler Data"

.vmr and .txt file of Doppler data generated during anaphylaxis

Dataset - "Metagenomic data of food allergy patients"

LABS-seq data of peanut allergy and egg allergy patients and controls

Planned research output details

Title	Туре	Anticipated release date	access	Intended repository(ies)	Anticipated file size	License	Metadata standard(s)	May contain sensitive data?	May contain PII?
Doppler Data	Dataset	2027-12-29	Open	Knowledge@UChicago	40 GB	Creative Commons Attribution Non Commercial Share Alike 4.0 International		No	No
Metagenomic data of food allergy patients		2027-12-30	Open	Gene Expression Omnibus	500 GB	Attribution Non Commercial Share Alike	Sequence (MIxS)	No	No