

## Plan Overview

---

*A Data Management Plan created using DMP Tool*

**Title:** Metabolomic Evaluation of SARS-COV-2 Infections

**Creator:** Danilo Bernardineli

**Affiliation:** Non Partner Institution

**Principal Investigator:** Danilo Bernardineli

**Data Manager:** Danilo Bernardineli

**Funder:** Digital Curation Centre (dcc.ac.uk)

**Funding opportunity number:** 50621

**Template:** Digital Curation Centre

**Last modified:** 07-08-2024

### Copyright information:

The above plan creator(s) have agreed that others may use as much of the text of this plan as they would like in their own plans, and customize it as necessary. You do not need to credit the creator(s) as the source of the language used, but using any of the plan's text does not imply that the creator(s) endorse, or have any relationship to, your project or proposal

---

# Metabolomic Evaluation of SARS-COV-2 Infections

## Data Collection

---

### What data will you collect or create?

Tabular metabolomics dataset for 450 patients

- \* 2 samples per patient
  - \* The first one is on the onset of internment
  - \* The second one is on discharge or aggravation of the condition
- \* Each sample contains 99 primary metabolites
- \* Format: CSV
- \* Expected minimum volume: 900 rows / 100 columns

Tabular patient metadata with categorizations

- \* Complication degree
  - \* No complications
  - \* Complications with no intensive support
  - \* Complications with intensive support
- \* Age range
  - \* Between 10 and 39 years old
  - \* Between 40 and 59 years old
  - \* Between 60 and 79 years old
  - \* Older than 80 years
- \* Format: CSV
- \* Expected minimum volume: 450 rows / 3 columns

### How will the data be collected or created?

The metabolomics dataset will be acquired through sputum collection and the standard protocol around it.

The patient metadata will be collected through administrative questionnaires by the associated hospitals, and transformed into an tabular CSV file.

Once retrieved, the files are going to be upload to an public Git repository at [https://gitlab.com/ion\\_ia/covid19\\_metabolomics](https://gitlab.com/ion_ia/covid19_metabolomics), which has version control by default. Any changes on the files, or a posterior QA processes, will be documented on the commit log and the CHANGELOG.md at the root directory.

## **Documentation and Metadata**

---

### **What documentation and metadata will accompany the data?**

The data will accompany an README.md containing the context and directions for it. Also, the public repository containing the data will have all pertinent documentation, like the analysis code, the research project, and the published content around it.

Also, regarding metadata, we'll adopt an multiple header standard, were the first lines of the CSV files indicates variable name, group and subgroup.

## **Ethics and Legal Compliance**

---

### **How will you manage any ethical issues?**

The ethics around the retrieval of data will be regulated through an ethical agreement between the researchers and the health providers.

As for data sensitivity, it will be handled through tokenization and anonimization of all possible patient identifiers, like bar-codes and internal ids. The dictionary for mapping those tokens will be stored on an private repository.

### **How will you manage copyright and Intellectual Property Rights (IP/IPR) issues?**

The non-sensitive data will be donated to the Brazilian public health provider (SUS - Sistema Único de Saúde), and all the generated data and analysis code will be published in an to-be defined public commons license, like MIT License.

## **Storage and Backup**

---

### **How will the data be stored and backed up during the research?**

The data will be stored and backed up in an Gitlab repository, which contemplates and takes care of all storage, security and backup issues.

## **How will you manage access and security?**

The repository with the metabolomics and metadata will be public, however the token dictionary for identifying the patients will be secured on an private repository on which only selected IONMedicine collaborators will have access.

## **Selection and Preservation**

---

### **Which data are of long-term value and should be retained, shared, and/or preserved?**

All data on the public repository (metabolomics and patient metdata), is to be retained and shared on long-term. We vouch for storing it for at least 5 years, and the data is going to be aggregated into large-scale metabolomics datasets, like MetabolomeXchange, allowing for contribution on large-scale aggregated studies.

As for the data on the private repository (patient identity dictionary), they are going to be retained for 1 year, with an deletion scheduled for afterwards.

### **What is the long-term preservation plan for the dataset?**

The data will be stored on Gitlab, which is an consolidated provider of code and data repositories. No efforts, costs or time required.

## **Data Sharing**

---

### **How will you share the data?**

Through the [https://gitlab.com/ion\\_ia/covid19\\_metabolomics](https://gitlab.com/ion_ia/covid19_metabolomics) git repository, and through pushing it on data finder mechanisms, like DataCite, Kaggle, and others. We intent to get an persistent identifier through DataCite.

The data will be available as soon we have it. If needed, modifications to the data will be handled through the Git versioning.

### **Are any restrictions on data sharing required?**

No

## **Responsibilities and Resources**

---

### **Who will be responsible for data management?**

The responsible for the data management will be the Artificial Intelligence division of IONMedicine. Tasks will be divided accordingly.

The creation and maintenance of the data repositories, as well as the publishing of it on several data providers like Kaggle, will be attribution of Danilo L. Bernardineli

**What resources will you require to deliver your plan?**

No additional resources needed

---