Metabolomic Evaluation of SARS-COV-2 Infections

A Data Management Plan created using DMPTool

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Metabolomic Evaluation of SARS-COV-2 Infections

Data Collection

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

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, 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

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

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.
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

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

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 colaborators will have access.

Selection and Preservation

All data on the public repository (metabolomics and patient metadata), 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.

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

Through the 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.

No additional resources needed