

Critical Care Project Report

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1 Introduction

The critical care patient data will be the cornerstone of many future medical and health services research. However, from data to research, there is still a huge gap to cover. Data has to be converted to a query-able form and we also need to know more about the data quality before it is ready for use. The mission of our UCL research software group (RSDG) is to provide an open source software to bridge this gap and deliver the ready-to-use data alongside with the data manipulation tools to the researchers.

A pipeline which converts the XML datasets to the query-able RData format has been built. It also allows us to export the selected data from RData to CSV for Excel. In the pipeline data quality check will be conducted guiding by a business readable data quality check dictionary. The non-validated data will be either modified or removed according to the criterion. Subsequently, an auto-generated data quality report will be created to report the key fields missingness and data sanity by site/unit. By the end of the pipeline the researchers will be able to receive a R table which is query-able and validated. We also have demonstrated the ability of deriving variables, such as calculating SOFA score from physiology data and identifying sepsis. We believe the paper about this will be ready soon.

Beyond the functionalities, we are keen on the usability and sustainability of the software as we believe an robust, understandable and accessible software is the first step of concrete science. All of our development are published on GitHub which is visible by all the allowed users. We made the installation of our software as easy as possible. The code are self-explanatory and well documented, which provides a great accessibility to the users and future developer. Testings are conducted after any modification of code automatically on Travis CI system. We believe the software is a gateway of the data which may play a great part in the future success of the community.

2 The Data Process Pipeline

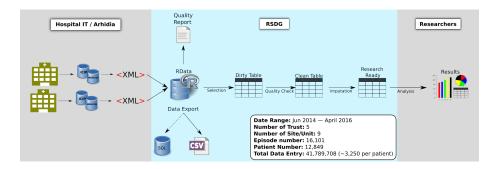


Figure 1: The Data Process Pipeline

2.1 The XML data and the Data Safe Haven

Data associated with more than 12,000 patients from 5 NHS trusts are available in XML format. The data were drawn from the ICIP and Medicuse database which contains demographic, drugs, laboratory, nursing and physiology data. The de-identifiable data are processed locally and the identifiable data is on the UCL Data Safe haven (IDSH). In order to create the pipeline of on IDSH, we ordered and configured a UNIX virtual environment. Due to the limit capacity of IDSH, it is still useful to keep a local de-identifiable copy for development purpose. Our pipeline is designed to be portable on multiple platforms. It can be executed on both local environments and IDSH with either de-identifiable or identifiable data. ## XML parser

The XML parser in R combines and restructures the XML files into a newly defined R data structure **ccRecord**, which significantly improved the clarity of the data by organising data under tables and removing the redundancy of the XML files. **ccRecord** is designed as a flexible, simplified, and query-able data structure for critical care data. Data in **ccRecord** format will be eventually stored in a RData file which is about 500 times smaller than that of the XML files. In addition the data provenance is recored by each episode, henceforth we will be able to tell which file does each episode data comes from and when it has been parsed. The selected data fields can be exported as a CSV file for the Excel users.

It is worth mentioning that a C++ equivalent XML parser, which has a much better performance comparing to the R parser, is developed but not deployed due to the time constraint of the project. To incorporate the C++ parser into the pipeline will be a potential future improvement.

2.2 Auto-generated quality report

Data can have defects in many different ways. Therefore a synthetic quality assessment which may allow us to report back to the source of the data can be extremely useful. An automatic generated data quality report which reflects to quality issues have been developed. Based on the report, we will be able to tell the major missingness issues and some data sanity issues and the basic information such as the duration, sites, number of episodes, and number of patient.

2.3 Data validation and cleaning

Data validation and cleaning functions have been incorporated in the ccdata package. Data sanity check is conducted in many aspects regarding to the intrinsic property of the individual fields. There are three major validation, the numeric range, the text category, and the missingness. The users are required to fill a yaml form to guide the validation check. We will be able to flag the level of the data sanity and opt out the data that do not make sense or inform the users according to the yaml dictionary. We

will again go through the yaml dictionary in the next session. After this stage, we are able to deliver a "cleaned" and query-able R table to the researchers.

3 The R package: ccdata

The ccdata R package is the centralised toolset we developed for pipelining and data manipulation. The package bundles not only all the R/C++ code but also documentation and tests together, which makes the code sharing easier. The ccdata package is portable in almost all platforms where R environment is provided. It can be installed effortlessly on Windows or Unix based systems. Although it is necessary to have some further tidy-ups in the subsequent development cycle, in the current stage, the main part of the R code is well documented and properly tested.

In order to prepare the first paper, many data manipulation processes are needed. Instead of making some one-off scripts, we incorporated the data manipulation processes as re-usable functions in our ccdata package. The frequently functions such as detecting unique patient and spell and data imputation can be called from the package, which reduced the duplication of work in the future. The package is still growing as the research goes on.

3.1 Data Selection and cleaning in the yaml configuration

Notwithstanding the fact that to be able use and program with ccdata package offers a great flexibility in data analysis, the pipeline can be performed without any programming knowledge. The users can run the pipeline by only filling the yaml configuration file. The yaml form is straight forward and self-explanatory for non-programmers. The user selected data fields with the the cleaning process can be provided by the end of the pipeline in a single CSV file which can be open in MS Excel.

Here is an example of the yaml data cleaning configuration of heart rate, in which three filters **nodata**, **range**, **missingness** are presented in the following data selection and cleaning configuration.

```
NIHR_HIC_ICU_0108:
    shortName: hrate
    dataItem: Heart rate
    distribution: normal
    decimal_places: 0

# filter1: do not use the episode where hrate cannot be found.
    nodata:
        apply: drop_episode

# filter2: mark all the values based on reference range (traffic colour) remove entries where the range check is not fullfilled.
    range:
        labels:
        red: (0, 300)
```



```
amber: (0, 170)
green: (50, 150)
apply: drop_entry

# filter3: compute the item missing rate on given cadences; in this case, we compute the daily (red) and hourly (amber) missing rate, and only accpet episodes of which hourly missing rate (amber) is lower than 30%.

missingness:
    labels:
        red: 24
        amber: 1
    accept_2d:
        amber: 70
apply: drop_episode
```

4 Summary

The RSDG got involved in the critical care data project since January 2016. In the last six months, we participated the paper preparation with the critical care team, meanwhile developed the re-usable and sustainable software tools for the data pipeline and data manipulation. In the potential next phase of this project, we will focus on the linkage of the data of other HIC groups and external sources. We will continue our support to the researchers by providing sustainable software tools.