

# QcDM Project User Guide

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## Table of Contents

<b>1. Overview and Installation .....</b>	<b>3</b>
1.1. <i>Install R and RStudio .....</i>	3
1.2. <i>Download QcDM Project.....</i>	4
1.3. <i>Install required R packages.....</i>	5
1.3.1. <i>Install QcDMconverter .....</i>	5
1.3.2. <i>Install QcDMui.....</i>	7
<b>2. User guide for QcDMconverter.....</b>	<b>8</b>
2.1. <i>Test1: Processing a single input file using QcDMconverter .....</i>	8
2.1.1. <i>Launch QcDMconverter .....</i>	9
2.1.2. <i>Process data .....</i>	10
2.1.3. <i>Output from data processing .....</i>	11
2.2. <i>Test2: Processing multiple input files with the same format .....</i>	14
2.2.1. <i>Process data .....</i>	14
2.2.2. <i>Output from data processing .....</i>	15
<b>3. User guide for QcDMui.....</b>	<b>17</b>
3.1. <i>Required data format of QcDMui .....</i>	17
3.2. <i>Launch QcDMui.....</i>	18
3.3. <i>Process data.....</i>	19
3.3.1. <i>Specify cutoffs and exclusion criteria.....</i>	20
3.3.2. <i>Select time period and wards .....</i>	21
3.4. <i>View glucometrics.....</i>	22
<b>4. Running QcDM Project from USB Drive .....</b>	<b>24</b>
4.1. <i>Set up the portable environment .....</i>	24
4.2. <i>Run QcDMconverter with PortableEnv .....</i>	25
4.3. <i>Run QcDMui with PortableEnv.....</i>	26
<b>Bibliography.....</b>	<b>27</b>

## 1. Overview and Installation

The QcDM (Quality care for Diabetes Mellitus) Project provides R-Shiny applications for generating glucometrics measures from point-of-care blood glucose data at three different units of analysis: patient-sample, patient-day, and patient-stay. The application facilitates convenient and flexible generation of glucometrics measures by providing user-friendly graphical user interfaces (GUIs) that runs in a browser on the user's desktop.

The QcDM Project consists of two R-Shiny GUIs and an R package. The first GUI, QcDMconverter, processes raw glucose data files to conform to the data format required by the application. The second GUI, QcDMui, is the main tool of the QcDM Project that uses functions in the [QcDM package](https://github.com/nyilin/QcDM) (<https://github.com/nyilin/QcDM>) to analyse processed glucose data to generate glucometrics measures.

The application operates on Windows and Mac operating systems, and is free under an academic non-commercial license. The current version of this application can be downloaded from a [GitHub repository](https://github.com/nyilin/QcDM_Project) ([https://github.com/nyilin/QcDM\\_Project](https://github.com/nyilin/QcDM_Project)). Installation of this application is described in details in the following sections.

### 1.1. Install R and RStudio

**The QcDM Project runs on R version 3.6.0 or later, but users should avoid version 4.0.0 because it has been reported to have issues with Shiny applications.** Users are recommended to install the latest version of R and RStudio by following the instructions below.

Install R from the installer downloadable from the official website of the R Project.

- For Windows: download the Setup Wizard from <https://cran.r-project.org/bin/windows/base/>. Follow through the installation steps and keep the default options.
- For macOS: download the installer (a pkg file) from <https://cran.r-project.org/bin/macosx/>. Follow through the installation steps and keep the default options.

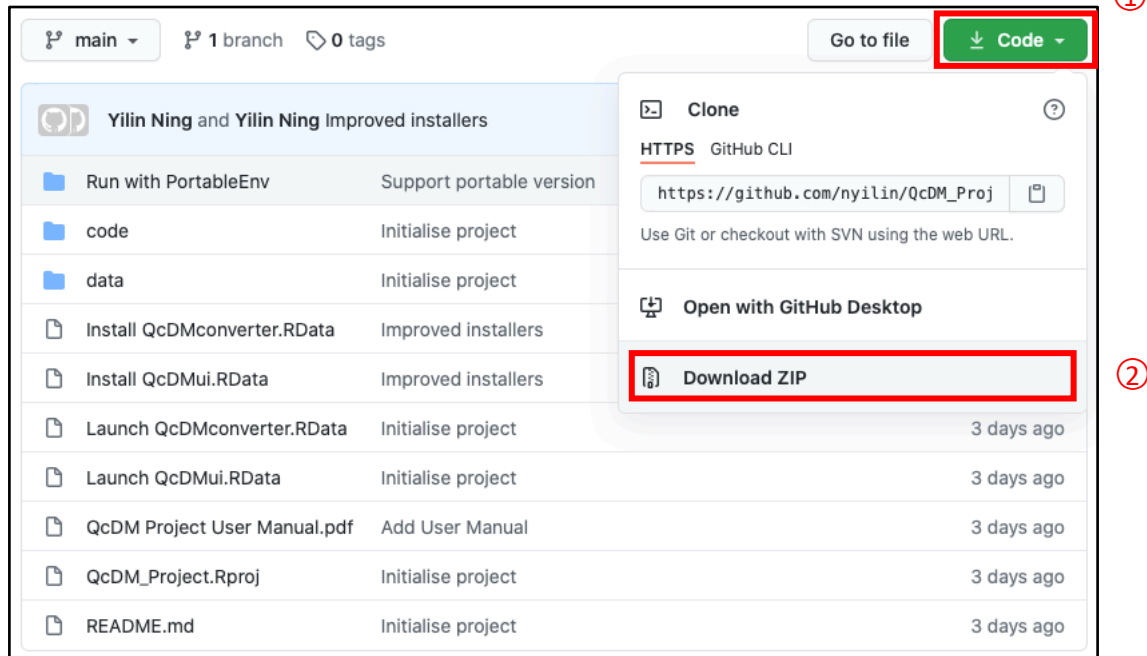
Install RStudio Desktop from the installer downloadable from the official website, <https://rstudio.com/products/rstudio/download/#download>.

For macOS, an additional application, XQuartz, is required, which can be downloaded from the official website: <https://www.xquartz.org>.

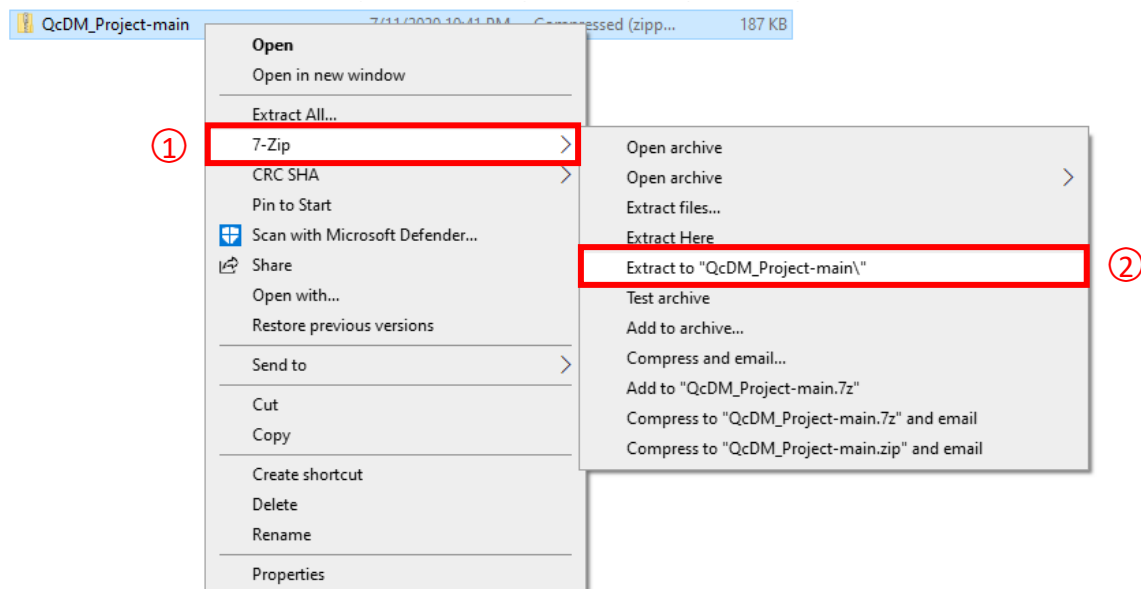
## 1.2. Download QcDM Project

From the [GitHub repository for the QcDM Project](https://github.com/nyilin/QcDM_Project)

([https://github.com/nyilin/QcDM\\_Project](https://github.com/nyilin/QcDM_Project)), download the project as a zip file, named 'QcDM\_Project-main.zip'. **We recommend using Chrome to access the link and download the zip file:**



For Windows, unzip the file by right-clicking on 'QcDM\_Project-main.zip' and selecting the following (and for illustration purposes we assume 7-Zip is used but any software that unzips file can be used):



For macOS, unzip the file by doubling clicking on it.

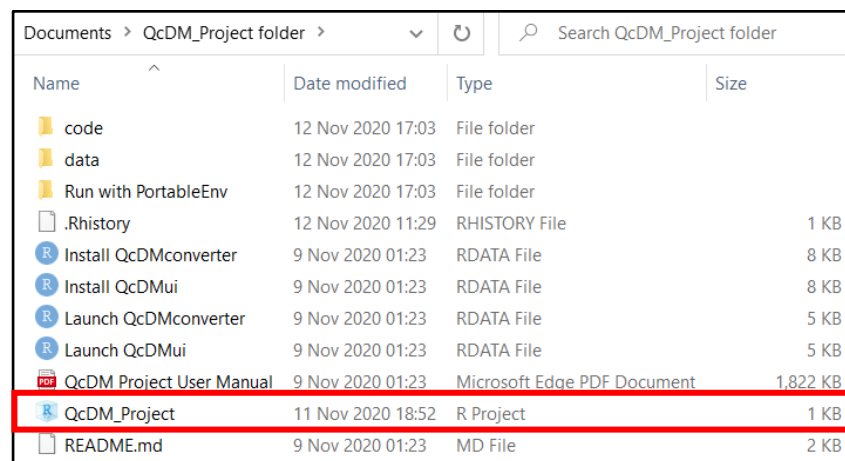
**We recommend users to rename the folder as 'QcDM\_Project folder' and move this folder to the 'Documents' folder. In the rest of this document, we assume users are using the**

Windows operating system and have followed the recommendations to rename and move the QcDM Project folder.

### 1.3. Install required R packages

A few R packages are required to use the two GUIs in the QcDM Project, and they can be installed by using the RData files in the 'QcDM\_Project folder' following the instructions below.

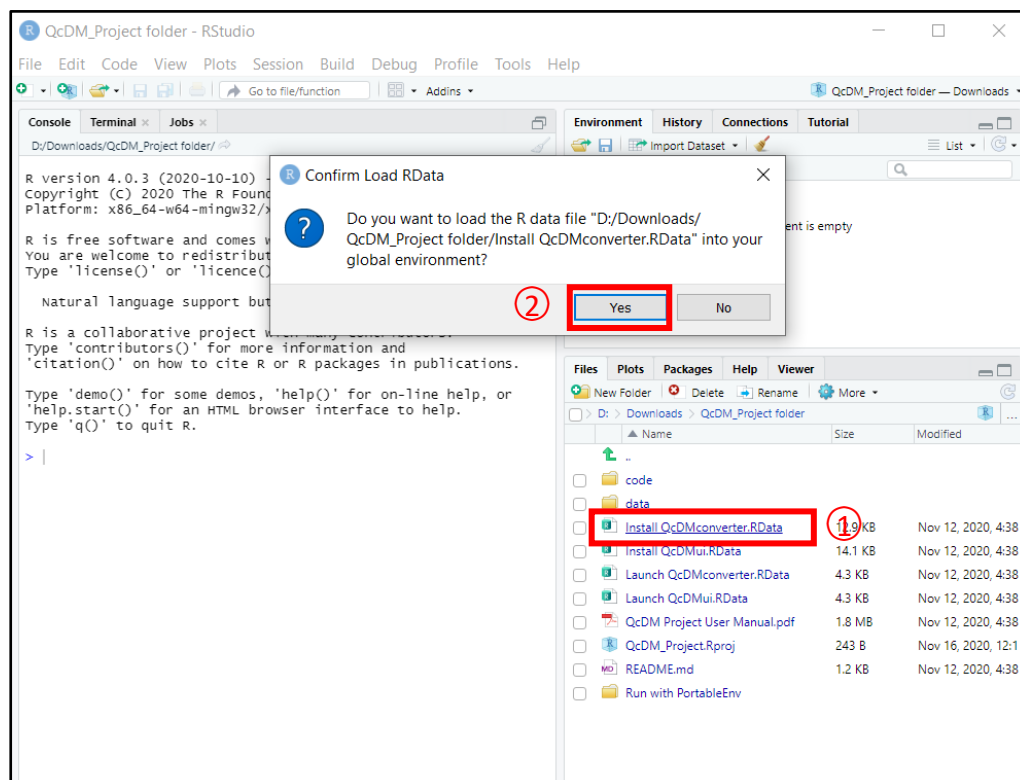
The screenshot below shows all the subfolders and files contained in the 'QcDM\_Project folder'. Firstly, users need to open RStudio by doubling-clicking 'QcDM\_Project.Rproj' in the 'QcDM\_Project folder':



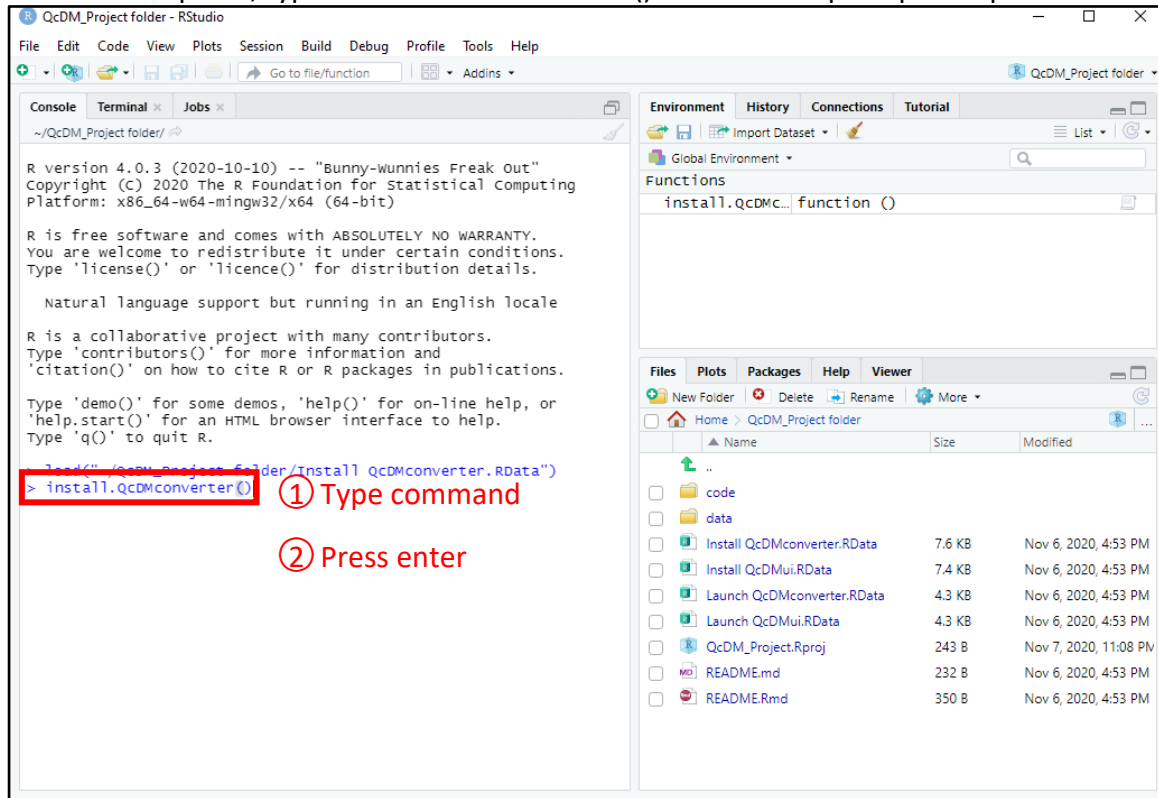
Name	Date modified	Type	Size
code	12 Nov 2020 17:03	File folder	
data	12 Nov 2020 17:03	File folder	
Run with PortableEnv	12 Nov 2020 17:03	File folder	
.Rhistory	12 Nov 2020 11:29	RHISTORY File	1 KB
Install QcDMconverter	9 Nov 2020 01:23	RDATA File	8 KB
Install QcDMui	9 Nov 2020 01:23	RDATA File	8 KB
Launch QcDMconverter	9 Nov 2020 01:23	RDATA File	5 KB
Launch QcDMui	9 Nov 2020 01:23	RDATA File	5 KB
QcDM Project User Manual	9 Nov 2020 01:23	Microsoft Edge PDF Document	1,822 KB
QcDM_Project	11 Nov 2020 18:52	R Project	1 KB
README.md	9 Nov 2020 01:23	MD File	2 KB

#### 1.3.1. Install QcDMconverter

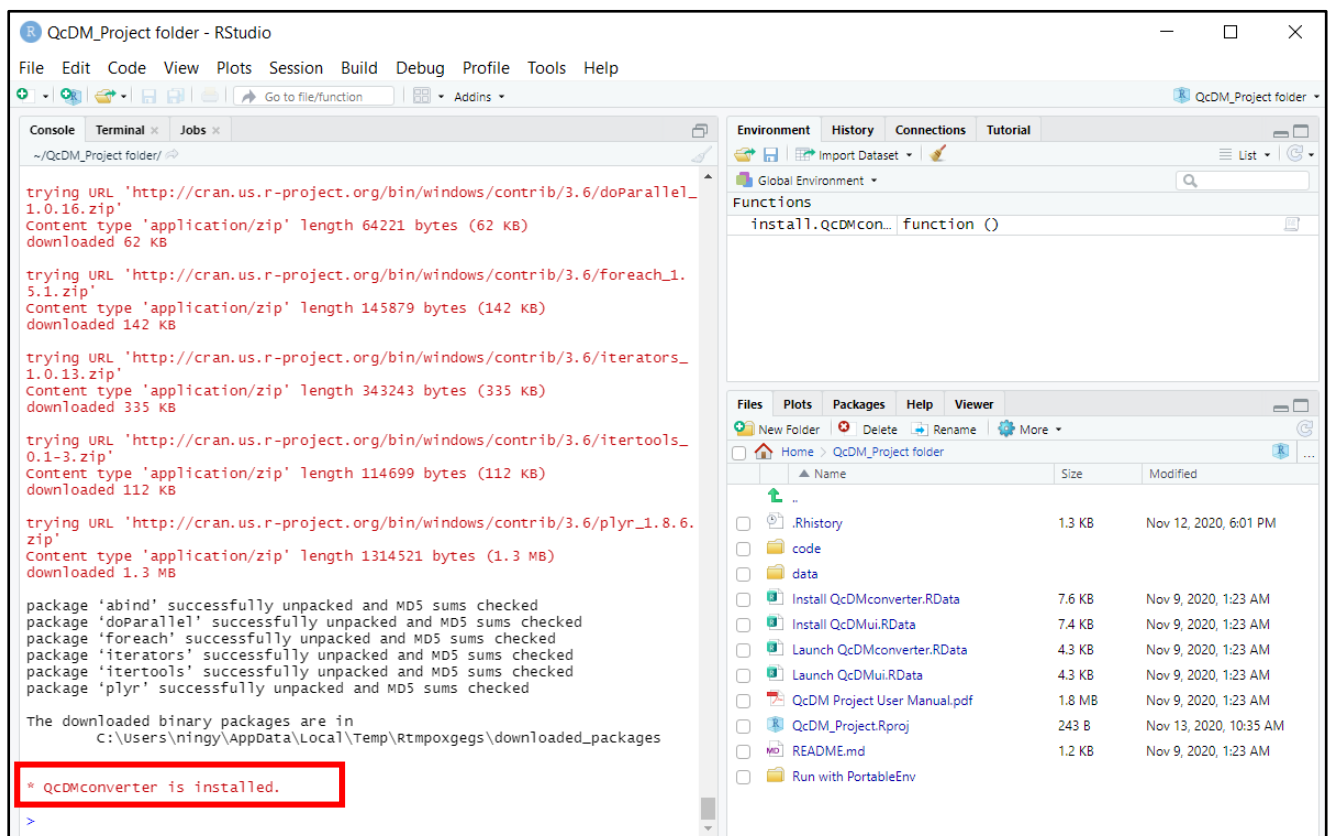
In the File panel of RStudio, click 'Install QcDMconverter.RData', and click 'Yes' in the pop-up window.



In the Console panel, type 'install.QcDMconverter()' after the '>' prompt and press Enter.



Installation is successful when a message '\* QcDMconverter is installed' is displayed in the Console panel (please ignore 'warning message' if any). Do not close the RStudio window as it will be used to install another R package (QcDMui).



### 1.3.2. Install QcDMui

Similar to the steps in the previous section, click on 'Install QcDMui.RData' in the File panel and click 'Yes' in the pop-up window.

In the Console panel, type 'install.QcDMui()' after the '>' prompt and press Enter.

Installation is successful when a message '\* QcDMui is installed' is displayed in the Console panel.

Close the RStudio window after successfully installing QcDMconverter and QcDMui. If prompted to select whether to save workspace image or history when closing the window, there is no need to save either of them.

## 2. User guide for QcDMconverter

QcDMconverter is an R-Shiny application that processes raw input glucose data based on the requirements of the QcDMui to simplify the analytical steps when generating glucometrics measurements.

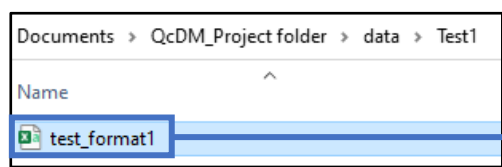
The following information is required from an input glucose data file for the generation of glucometrics:

- an indicator for hospital stays experienced by each patient,
- the date-time stamp associated with each glucose measurement,
- the value of each glucose measurement,
- the ward location associated with each glucose measurement.

New input glucose data may be provided as a single comma-separated version (CSV) file (e.g., see Test1 described in Section 2.1), or as multiple CSV files that follow the same format (e.g., see Test2 described in Section 2.2). QcDMconverter extracts essential information from input file(s) and converts them to a common format that conforms to the requirements by QcDMui.

### 2.1. Test1: Processing a single input file using QcDMconverter

To illustrate the use of QcDMconverter, we simulated glucose data with 4459 rows for 100 unique hospital stays in July 2020, randomly assigned into 4 ward locations (i.e., A, B, C and D). The unit of measurement of the glucose readings is mmol/L. For illustrative purposes, 6 of the 4459 rows had non-numerical glucose readings. This data, which conforms to the required data format by QcDMui (described in details in Section 3.1), is saved as a CSV file named 'test\_format1.csv' in the 'Test1' folder, which is in the 'data' subfolder of 'QcDM\_Project folder'.



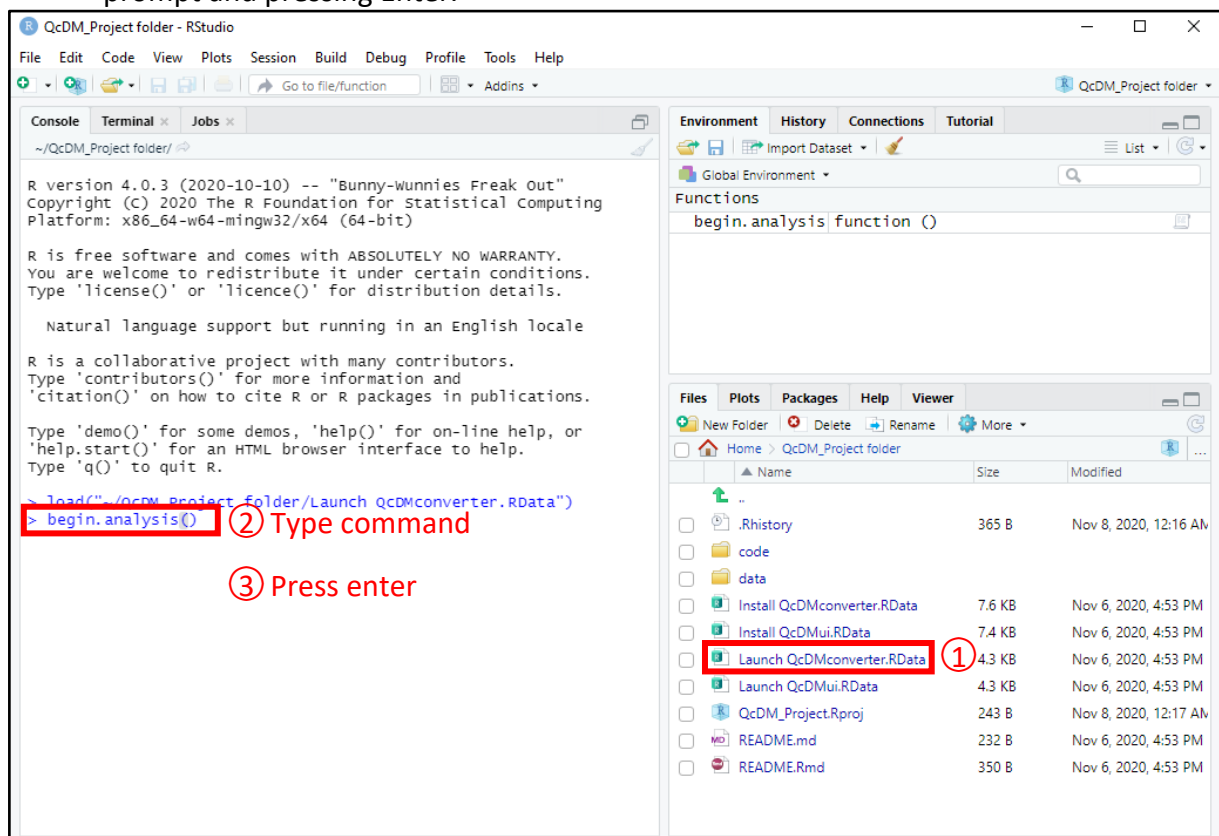
	A	B	C	D
1	ADMISSION.ID	RESULT	RESULT.DATE	LOCATION
2	8226	5.7	07/01/2020 00:07	D
3	4194	6.3	07/01/2020 00:03	A
4	143	3.6	07/01/2020 00:08	C
5	6363	19	07/01/2020 00:06	C
6	3931	8.8	07/01/2020 00:04	C
7	4105	16.9	07/01/2020 00:08	C
8	270	7.1	07/01/2020 00:09	D
9	5598	7.5	07/01/2020 00:05	D
10	6709	7.7	07/01/2020 00:07	B



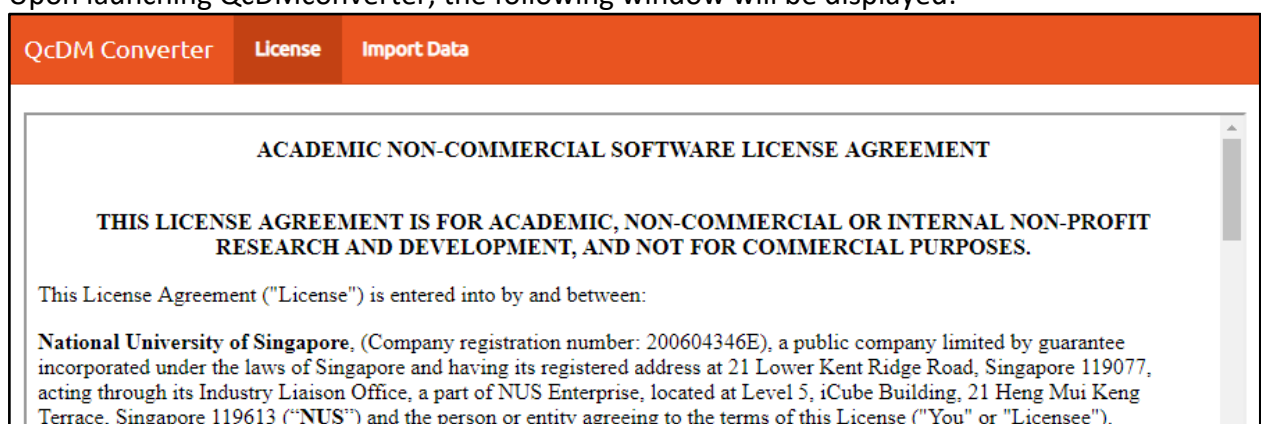
### 2.1.1. Launch QcDMconverter

To ensure QcDMconverter can be launched properly by using 'Launch QcDMconverter.RData' in the 'QcDM\_Project folder', follow the instructions below.

- Goto 'QcDMui\_project folder', double click 'QcDM\_Project.Rproj' to open RStudio.
- Click 'Launch QcDMconverter.RData' in the File panel, click 'Yes' in the pop-up window.
- In the Console panel, execute command 'begin.analysis()' by typing it after the '>' prompt and pressing Enter.

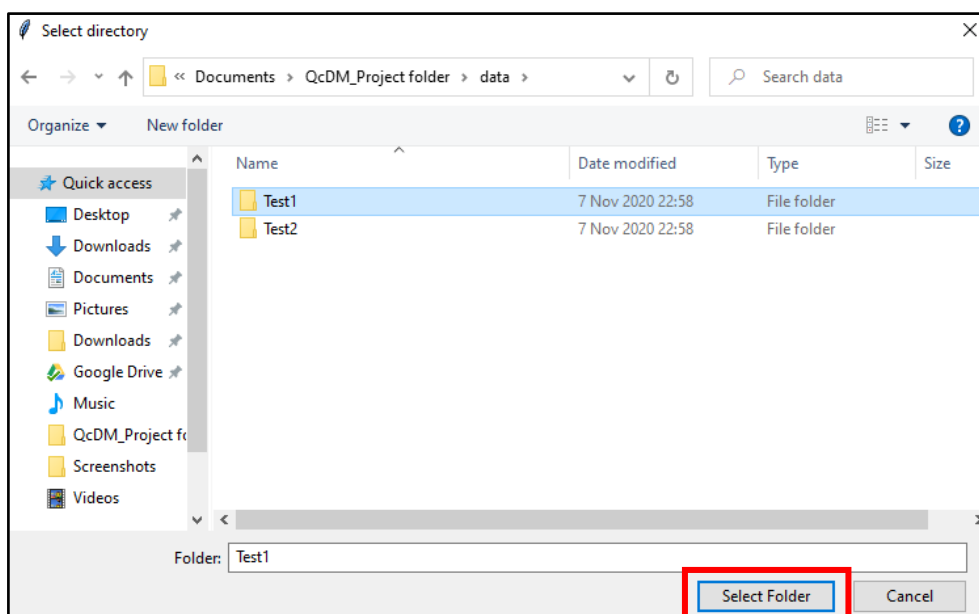


Upon launching QcDMconverter, the following window will be displayed:



### 2.1.2. Process data

Click on the 'Import Data' tab to begin data processing. The following window will pop out for users to locate the 'Test1' folder:



Click 'Select Folder' to proceed, and the following dashboard will be displayed. Use it to select appropriate options for reading in the input file, and click 'Next' to proceed.

**QcDM Converter**   **License**   **Import Data**

Please select a text file to upload and use the radio buttons to format the file into columns. Click Next when done.

**Data folder**

C:/Users/ningy/Documents/QcDM\_Project folder/data/Test1

**Select File**

test\_format1.csv

☒ Header

**Separator**

☒ Comma

☐ Semicolon

☐ Tab

ADMISSION.ID	RESULT	RESULT.DATE	LOCATION
8226	5.7	07/01/2020 00:07	D
4194	6.3	07/01/2020 00:03	A
143	3.6	07/01/2020 00:08	C
6363	19	07/01/2020 00:06	C

C:/Users/ningy/Documents/QcDM\_Project folder/data/Test1/test\_format1.csv

**Next**

Select appropriate options based the format of the file selected, then click 'Next'

Use the following tab to configuration additional data options based on the input file selected, then click 'Import' to begin data processing. **Note that this will overwrite existing output file that was previously generated from the same input file.**

**QcDM Converter**   **License**   **Import Data**

**A) Admission ID**  
ADMISSION.ID

**B) Result Date**  
RESULT.DATE

**B1) Date Format**  
mdy

**B2) Time Format**  
hm

**C) Result Value**  
RESULT

**D) Location**  
LOCATION

**Import**

Please Configure QcDM using the controls on the left and click the 'Import' button  
Please monitor the job status using the progress bar below. If an error occurs information will be shown here.  
After clicking 'Import', existing files processed from the same input files selected will be overwritten.

Configure the GUI based on the format of the file selected, then click 'Import'

After clicking 'Import', a progress bar will be displayed at the lower right corner of the GUI

Processing Data Files Processing File: 1 of 1

The following text message will be displayed at the top of this tab to indicate the completion of data processing:

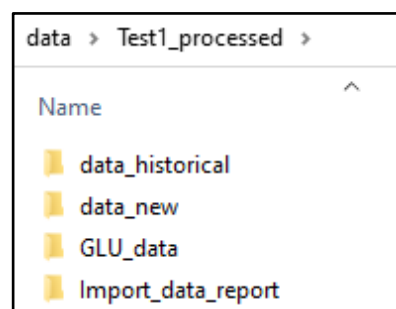
```
Single File Processing Mode:

Processing succeeded for the following files:
SUC: C:/Users/ningy/Documents/QcDM_Project folder/data/Test1/test_format1.csv

1 of 1 processed successfully
```

### 2.1.3. Output from data processing

After performing the data processing steps in Section 0, a folder named 'Test1\_processed' is created in the same folder as 'Test1', which contains four subfolders:



Subfolder 'data\_new' contains a file named 'Jul2020\_test\_format1.csv', which was created from the input 'test\_format1.csv', now with the month and year of the glucose data appended to the beginning of its file name, and with the date-time stamp converted to

format 'yyyy-mm-dd hh:mm:ss'. As seconds are missing in the date-time stamp in the input data, they are assigned to be '11' by QcDMconverter after data processing.

	A	B	C	D
1	ADMISSION.ID	RESULT.DATE	RESULT	LOCATION
2	8226	2020-07-01 00:07:11	5.7	D
3	4194	2020-07-01 00:03:11	6.3	A
4	143	2020-07-01 00:08:11	3.6	C
5	6363	2020-07-01 00:06:11	19	C
6	3931	2020-07-01 00:04:11	8.8	C
7	4105	2020-07-01 00:08:11	16.9	C
8	270	2020-07-01 00:09:11	7.1	D
9	5598	2020-07-01 00:05:11	7.5	D
10	6709	2020-07-01 00:07:11	7.7	B

When proceeding to the generation of glucometrics measurements, QcDMui will look for input glucose data in the 'data\_new' subfolder (see Section 3). Subfolders 'data\_historical' and 'GLU\_data', which are empty now, will be used later by QcDMui to archive input data and store processed data, respectively.

Subfolder 'Import\_data\_report' stores PDF reports that will be created by QcDMconverter. There will be two reports per input file, with the names of the corresponding input appended to the end of file names:

'Overall summary\_test\_format1.pdf' provides basic summary statistics for the input glucose values and date-time stamps to facilitate early detection of data entries errors. For example, the 6 entries in 'test\_format1.csv' with non-numerical glucose values are reported:

'test\_format1.csv':

	A	B	C	D
1	ADMISSION.ID	RESULT	RESULT.DATE	LOCATION
2	3588	d9.6	07/05/2020 11:45	B
3	4194	c7.7	07/05/2020 23:51	A
4	4558	a4.4	07/05/2020 06:06	C
5	3588	>33.5	07/09/2020 12:09	B
6	4194	> 33.3	07/10/2020 19:51	A
7	4194	> 33.3	07/10/2020 20:08	A

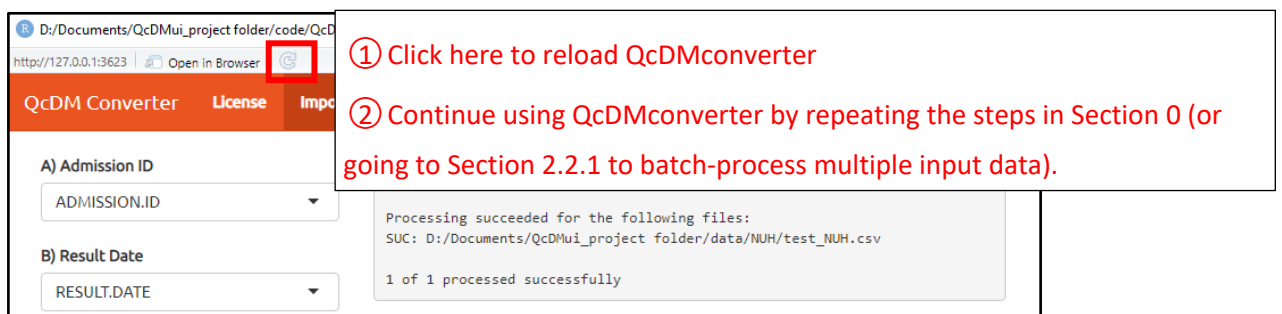
'Overall summary\_test\_format1.pdf':

2) Report of invalid readings		
Invalid BG values		
Label	N (%)	
> 33.3	2 (0.0)	
>33.5	1 (0.0)	
a4.4	1 (0.0)	
c7.7	1 (0.0)	
d9.6	1 (0.0)	
Invalid BG timestamps		
There is no invalid BG timestamp in the input data.		

'Detailed\_summary\_test\_format1.pdf' provides more details on the input glucose values and date-time stamps in each ward location.

Invalid glucose values and date-time stamps in the input glucose data will be recoded to 'NA' (which indicates missing data in R) by QcDMconverter in the processed data. If this is inline with users' intention (e.g., the non-numerical glucose values are codes used to indicate entries without glucose measurements and hence should be coded as missing), users may proceed to generate glucometrics from the processed data using QcDMui (see Section 3). If the invalid readings identified in the report reflect errors in data entry, users may want to make the necessary corrections to the raw input data and repeat the steps in Section 2.1.2.

After the completion of data processing for Test1, users may reload QcDMconverter to process other input data:



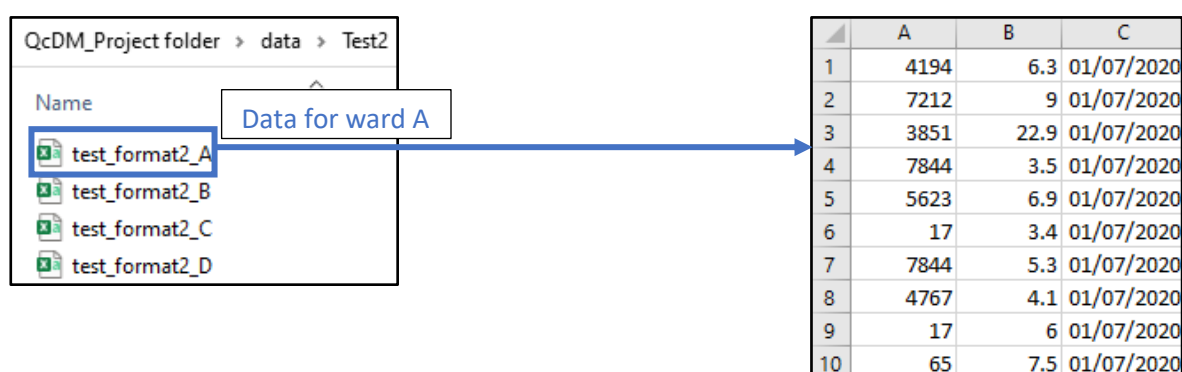
The screenshot shows the QcDM Converter web application running in a browser. The address bar shows the URL `http://127.0.0.1:3623`. The application has a header with 'QcDM Converter', 'License', and 'Import' buttons. Below the header, there are two sections: 'A) Admission ID' with a dropdown menu showing 'ADMISSION.ID', and 'B) Result Date' with a dropdown menu showing 'RESULT.DATE'. To the right of these sections, there is a status box that says 'Processing succeeded for the following files: SUC: D:/Documents/QcDMui\_project folder/data/NUH/test\_NUH.csv' and '1 of 1 processed successfully'. A red box highlights the 'Reload' button (a circular arrow icon) in the top right corner of the application. Two red annotations are present: '① Click here to reload QcDMconverter' pointing to the 'Reload' button, and '② Continue using QcDMconverter by repeating the steps in Section 0 (or going to Section 2.2.1 to batch-process multiple input data).' pointing to the status box.

Users may also close the QcDMconverter window to return to the RStudio window, where they can proceed to the generation of glucometrics measurements using QcDMui (see Section 3), or close the RStudio window now and proceed later. If prompted to select whether to save workspace image when closing the RStudio window, there is no need to save either of them.

## 2.2. Test2: Processing multiple input files with the same format

QcDMconverter can batch-process multiple files in the folder if they follow the same format. To illustrate this mode of usage, 'test\_format1.csv' is split into 4 CSV files and saved to the folder 'Test2' (which is in the 'data' subfolder of 'QcDM\_Project folder'), where each file contains all the entries corresponding to the same ward.

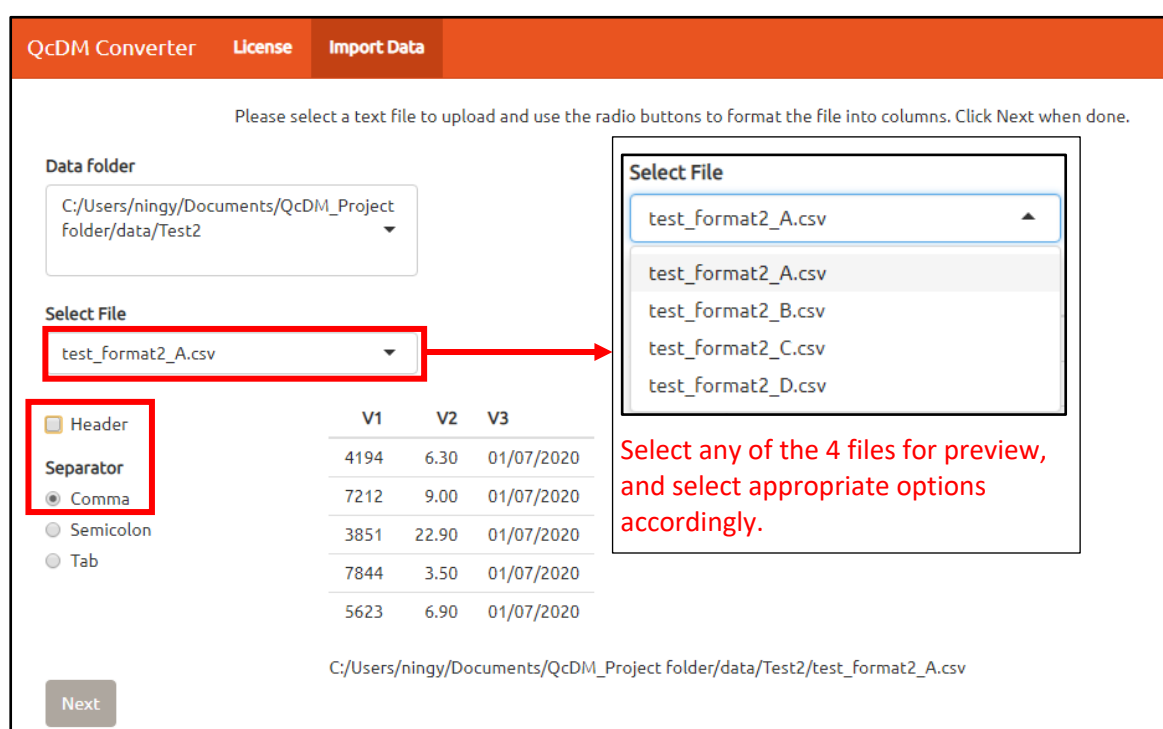
For illustrative purpose, each file now contains three columns, without column names, for the admission ID, glucose values and date stamp for the glucose measurements (without hours or minutes), respectively. Such file format is used in the Yale Glucometrics project when generating glucometrics (1). As the ward location is no longer included as a data column, it must be specified as the last few characters in the file names, preceded by '\_ '.



	A	B	C
1	4194	6.3	01/07/2020
2	7212	9	01/07/2020
3	3851	22.9	01/07/2020
4	7844	3.5	01/07/2020
5	5623	6.9	01/07/2020
6	17	3.4	01/07/2020
7	7844	5.3	01/07/2020
8	4767	4.1	01/07/2020
9	17	6	01/07/2020
10	65	7.5	01/07/2020

### 2.2.1. Process data

After launching the QcDMconverter GUI as described in Section 0 and selecting the 'Test2' folder using the GUI, users can select any of the 4 files in this folder for preview, in order to inform the specification of options when reading in these files. Make sure you uncheck 'Header'.



Please select a text file to upload and use the radio buttons to format the file into columns. Click Next when done.

Data folder  
C:/Users/ningy/Documents/QcDM\_Project folder/data/Test2

Select File  
test\_format2\_A.csv

Header  
Separator  
☒ Comma  
☐ Semicolon  
☐ Tab

V1	V2	V3
4194	6.30	01/07/2020
7212	9.00	01/07/2020
3851	22.90	01/07/2020
7844	3.50	01/07/2020
5623	6.90	01/07/2020

C:/Users/ningy/Documents/QcDM\_Project folder/data/Test2/test\_format2\_A.csv

Next

Select any of the 4 files for preview, and select appropriate options accordingly.

After clicking 'Next', the GUI displays an additional option E upon detecting multiple files in the folder selected. Users may choose to process one file each time, or process all files in the folder by selecting 'Yes', provided that all files follow the same format.

The screenshot shows the 'Import Data' tab of the QcDM Converter GUI. The configuration options are as follows:

- A) Admission ID:** V1
- B) Result Date:** V3
- B1) Date Format:** dmy
- B2) Time Format:** Not available
- C) Result Value:** V2
- D) Location:** <<From Filename>> (highlighted with a red box)
- E) Process all files in this folder with these formats:** Yes (highlighted with a red box)

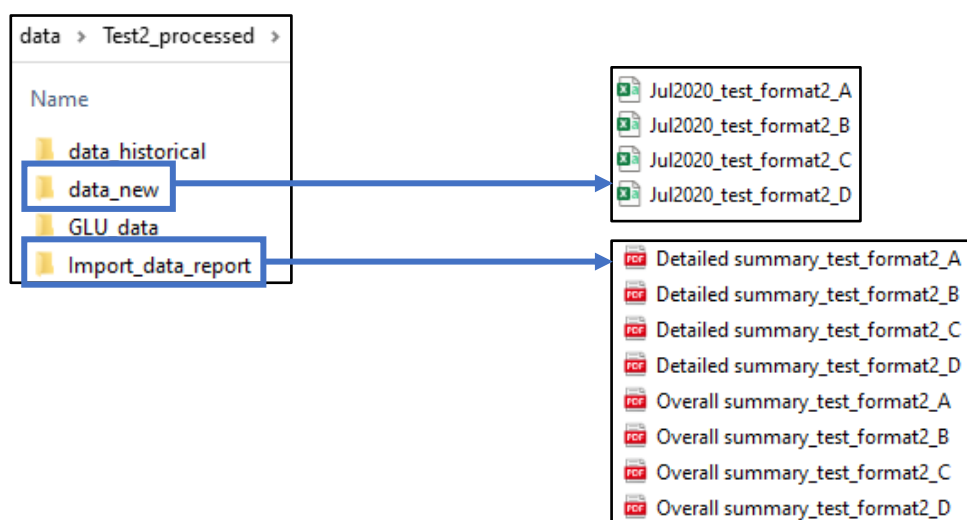
Callout for D) Location: Ward location is now specified as part of the file name instead of as a column in the data.

Callout for E) Process all files in this folder with these formats: When the GUI detects multiple files in the selected folder, users have this additional option of batch-processing all files using the same configuration.

After configuration, click 'Import' (under option E) to start processing.

### 2.2.2. Output from data processing

As described in Section 2.1.3, a folder named 'Test2\_processed' is created by QcDMconverter, with the following four subfolders:



Despite the different data formats adopted in Test1 and Test2, the processed data files in the 'data\_new' folder follow the same format:

	A	B	C	D
1	ADMISSION.ID	RESULT.DATE	RESULT	LOCATION
2	4194	2020-07-01 11:11:11	6.3	A
3	7212	2020-07-01 11:11:11	9	A
4	3851	2020-07-01 11:11:11	22.9	A
5	7844	2020-07-01 11:11:11	3.5	A
6	5623	2020-07-01 11:11:11	6.9	A
7	17	2020-07-01 11:11:11	3.4	A
8	7844	2020-07-01 11:11:11	5.3	A
9	4767	2020-07-01 11:11:11	4.1	A
10	17	2020-07-01 11:11:11	6	A

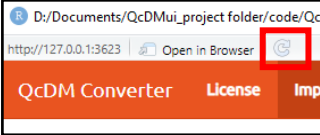
As time stamps are not available in the input data files in Test2, they are assigned as 11:11:11 AM after the data processing.

As described in Section 2.1.3, the overall summary reports generated for each input file is useful for detecting invalid entries in glucose values and date-time stamp. For example, 3 glucose values in the input file for ward A are reported to be non-numerical in 'Overall summary\_test\_format2\_A.pdf':

<b>2) Report of invalid readings</b>							
Invalid BG values							
	<table> <tr> <th>Label</th><th>N (%)</th></tr> <tr> <td>&gt; 33.3</td><td>2 (0.3)</td></tr> <tr> <td>c7.7</td><td>1 (0.2)</td></tr> </table>	Label	N (%)	> 33.3	2 (0.3)	c7.7	1 (0.2)
Label	N (%)						
> 33.3	2 (0.3)						
c7.7	1 (0.2)						
Invalid BG timestamps							
There is no invalid BG timestamp in the input data.							

More detailed summaries of glucose values and date-time stamps for each ward can be found in their respective detailed summary reports. Users may proceed to use QcDMui (see Section 3) provided information you are fine with the information provided in the pdf files.

After the completion of data processing for Test2, users may reload QcDMconverter to process other input data:



① Click here to reload QcDMconverter.

② Continue using QcDMconverter by repeating the steps in Section 0 or 2.2.1.

Users may also close the QcDMconverter window to return to the RStudio window, where they can proceed to the generation of glucometrics measurements using QcDMui (see Section 3), or close the RStudio window now and proceed later. If prompted to select whether to save workspace image when closing the RStudio window, there is no need to save either of them.

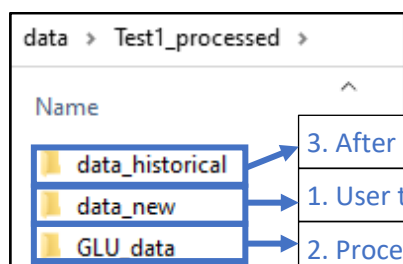


### 3. User guide for QcDMui

QcDMui is an R-Shiny graphical user interface developed for convenient and flexible generation of glucometrics measures from point-of-care glucose data.

#### 3.1. Required data format of QcDMui

QcDMui requires a specific data format and folder structure for the input glucose data files. Specifically, QcDMui expects a data folder to have the following subfolders:



1. User to put glucose CSV file(s) in the format generated by QcDMconverter.
2. Processed monthly data will be organized in this subfolder by ward, year and month.
3. After processing, file(s) in 'data\_new' will be moved here for archiving purposes.

For easy management, users are recommended to provide glucose data by month, and indicate the calendar time of each file using the first seven characters of the file name, where the first three characters are the abbreviated month, and the next four characters correspond to the four-digit calendar year (e.g., 'Jul2020' for glucose data in July 2020).

As illustrated in the snapshot below, each data file in 'data\_new' should contain at least four variables:

	A	B	C	D
1	ADMISSION.ID	RESULT	RESULT.DATE	LOCATION
2	17	3.4	07/01/2020 01:15	A
3	17	6	07/01/2020 02:16	A
4	17	3.4	07/01/2020 07:50	A
5	17	5.8	07/01/2020 09:34	A
6	17	6.2	07/01/2020 11:40	A
7	17	7	07/01/2020 17:21	A
8	17	7.5	07/01/2020 22:18	A
9	17	7.4	07/02/2020 07:00	A
10	17	7.1	07/02/2020 12:05	A

- ADMISSION.ID: the identifier for a specific hospital stay experienced by a patient.
- RESULT.DATE: the date-time stamp associated with each glucose measurement, where:
  - only '/' or '-' is used to separate date, month and year, and no mix of the two symbols is allowed in a single time variable (see below for an example)
  - ':' is used to separate hour, minute and second, and the order is always 'hh:mm:ss' (see below for an example). Second can be missing, but not minute or hour. This is to allow us to perform distributional analytics for assessing timeliness of glucose measurements (2).
- RESULT: the value of glucose measurement. It can be in mg/dL or mmol/L, but not a mix of the two units. If 95% of glucose values are greater than 33.3, then the unit of these values is taken as mg/dL, otherwise, the unit will be taken as mmol/L.
- LOCATION: the ward location where the glucose measurements were taken.

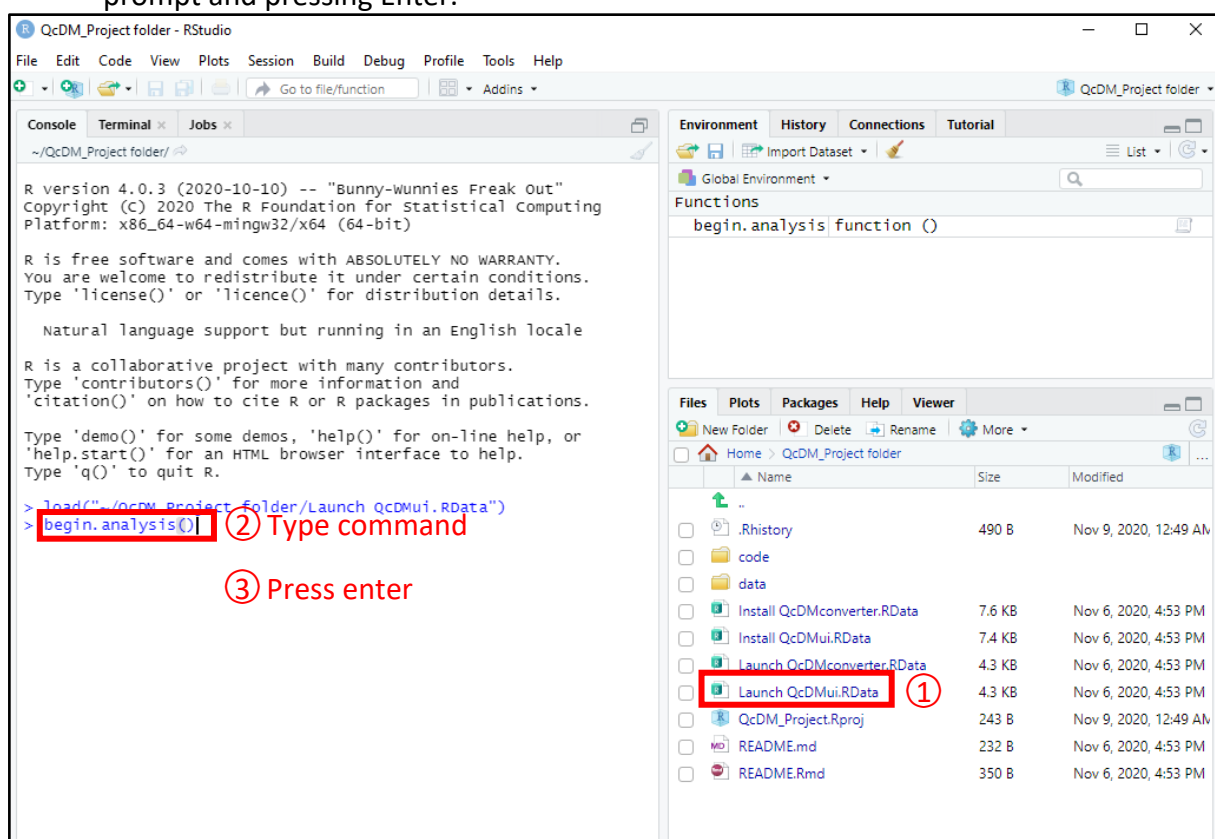
**Users are strongly recommended to process raw input data files using QcDMconverter, as described in Section 2, to create the required folder structure and conform to the required data format to facilitate their usage of the QcDMui.**

The next two sections illustrate the usage of QcDMui and the corresponding output using example data provided in folder 'Test1', which is available in the 'data' subfolder of 'QcDM\_Project folder'. We assume users have followed the instructions in Section 2.1 to create a folder 'Test1\_processed' that contains the required subfolders and processed data.

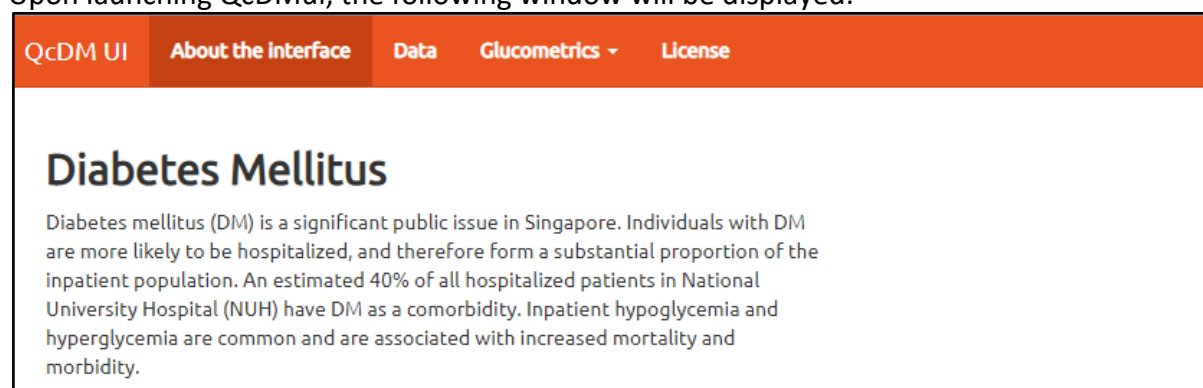
### 3.2. Launch QcDMui

QcDMui can be launched by using 'Launch QcDMui.RData' in the 'QcDM\_Project folder' following the instructions below.

- Goto 'QcDM\_Project folder', double click 'QcDM\_Project.Rproj' to open RStudio.
- Click 'Launch QcDMui.RData' in the File panel, click 'Yes' in the pop-up window.
- In the Console panel, execute command 'begin.analysis()' by typing it after the '>' prompt and pressing Enter.



Upon launching QcDMui, the following window will be displayed:



An introduction to the tool and a brief instruction on how to use it is available under the tab 'About the interface'.

### 3.3. Process data

Users can use the 'Data' tab to specify the glucose data for which glucometrics measurements need to be generated.

The following window will pop out upon clicking on the 'Data' tab for users to locate the 'Test1\_processed' folder. Click 'Select Folder' to proceed. Confirm the selection by clicking 'Confirm data folder' in the GUI.

The screenshot displays the QcDM UI interface. At the top, there is a navigation bar with tabs: 'QcDM UI', 'About the interface', 'Data', 'Glucometrics', and 'License'. The 'Data' tab is currently selected.

Below the navigation bar, the 'Data Folder' section shows a dropdown menu with the path 'C:/Users/ninyg/Documents/QcDM\_Project folder/data/Test1\_processed'. A red box highlights the 'Confirm data folder' button, which is labeled with a circled '2'.

To the right of the 'Data Folder' section, there is a 'Time range' section with 'From' and 'To' fields for Year and Month. The 'From' Year is set to 2020 and the 'To' Year is set to 2020. The 'From' Month is set to 11 and the 'To' Month is set to 11. A red box highlights the 'Confirm time range' button.

Below the 'Time range' section, there is a 'Wards' section with a radio button for 'All wards with data in the selected time range' and a radio button for 'Selected wards'. The 'Selected wards' radio button is selected. A red box highlights the 'Confirm choice' button.

Below the 'Wards' section, there is a 'Number of unique patients in each selected ward and month' section with a text input field and a 'Confirm ward' button. A red box highlights the 'Confirm ward' button.

Below the 'Confirm ward' button, there is an 'Exclusion criteria' section with two checkboxes and input fields. The first checkbox is 'Exclude patient stays having fewer BG measurements than the number specified below' with an input field set to 2. The second checkbox is 'Exclude patient stays having length of stay less than or equal to the number specified below in terms of hours' with an input field set to 24.

At the bottom of the 'Data' tab, there is a 'Confirm' button and a 'Process complete.' message.

The following two subsections describe the 'Data' tab in detail.

### 3.3.1. Specify cutoffs and exclusion criteria

The screenshot shows the 'Data' tab of the QcDM UI. At the top, there is a navigation bar with 'QcDM UI', 'About the interface', 'Data' (selected), 'Glucometrics', and 'License'. Below the navigation bar, the 'Data folder' section shows a dropdown menu with the path 'C:/Users/ningy/Documents/QcDM\_Project folder/data/Test1\_processed' and a 'Confirm data folder' button. The main section is divided into two parts, both highlighted with red boxes. The first part, 'Hypo cutoff', 'Hyper cutoff', 'Target range', and 'Hyperglycemia index (HGI)', contains input fields for specifying cutoff values. The second part, 'Exclusion criteria', contains checkboxes and input fields for specifying exclusion criteria. At the bottom, there is a 'Confirm' button and a 'Process complete.' message. A red box highlights the 'Confirm' button, and a text box next to it says 'Click on 'Confirm' button before proceeding to the panel on the right.'

Hypo cutoff	Hyper cutoff	Target range	Hyperglycemia index (HGI)
Please specify the first hypo cutoff (mmol/L): 4	Please specify the first hyper cutoff (mmol/L): 14	Please specify the lower value of the target range (mmol/L): 4	Please specify the cutoff for HGI (mmol/L): 10
Please specify the second hypo cutoff (mmol/L): 3	Please specify the second hyper cutoff (mmol/L): 20	Please specify the lower value of the target range (mmol/L): 10	
Please specify the third hypo cutoff (mmol/L): 2.5	Please specify the third hyper cutoff (mmol/L): 24		

**Exclusion criteria**

☐ Exclude patient stays having fewer BG measurements than the number specified below  
2

☐ Exclude patient stays having length of stay less than or equal to the number specified below in terms of hours  
24

After selecting the right data folder and parameters, please click on 'Confirm' button below before proceeding to the panel on the right.

**Confirm**  
Process complete.

Click on 'Confirm' button before proceeding to the panel on the right.

This part allows users to customize cutoff values for hypoglycemia, hyperglycemia, target range and hyperglycemia index. Units of glucose measurement used here (mmol/L or mg/dL) will be inferred from input data\*.

This part allows users to customize exclusion criteria.

\*: If 95% of glucose values are greater than 33.3, then the unit of these values is taken as mg/dL, otherwise, the unit will be taken as mmol/L.

After specifying appropriate cutoff values and exclusion criteria, click on 'Confirm' button. QcDMui will import data in the 'data\_new' subfolder, split them by month and ward, and

organize the processed data into the respective subfolders within the 'GLU\_data' subfolder. A text message 'Process complete.' will be displayed under the 'Confirm' button upon completion of data processing. Users can then proceed to the panel on the right to select time period and wards for the generation of glucometrics.

### 3.3.2. Select time period and wards

**Time range:**

From To  
Year Year  
2020 2020  
Month Month  
7 7  
Confirm time range

**Wards**

☐ All wards with data in the selected time range ☒ Selected wards  
Confirm choice  
Please select wards to view (multiple selection allowed)  
A Multiple selection allowed  
Confirm ward

**Number of unique patients in each selected ward and month:**

Show 10 entries Search:

	Wards	Jul 2020
1	A	15

Showing 1 to 1 of 1 entries Previous 1 Next

Users are required to use these two parts to specify the period and ward for glucometrics generation, based on the availability of data in the 'GLU\_data' subfolder.

After selecting the ward and time range, a summary of the selected data by ward and month will be presented here, so that users can modify their selection if the selected data is not available.

### 3.4. View glucometrics

After processing glucose data in 'Data' tab, click on 'Glucometrics' tab and then go to 'Summary Statistics' tab to view the following tables that summarize the glucometrics information:

Data Summary			
<b>Summary of BG measurements</b>			
		<b>Summary</b>	
Total (N)			603
Minimum (mmol/L)			3.3
5-th percentile (mmol/L)			4.4
25-th percentile (mmol/L)			5.4
75-th percentile (mmol/L)			9.9
95-th percentile (mmol/L)			16.4
Maximum (mmol/L)			31
Non-numeric values (N)			3
Exclusion Summary			
Total number of patient-stays			15
No exclusion criterion was selected.			
Glucometrics			
<b>Summary</b>			
Location			A
Period			01 Jul 2020 to 31 Jul 2020
	<b>Patient-sample</b>	<b>Patient-day<sup>#1</sup></b>	<b>Patient-stay<sup>#2</sup></b>
Count	600	141	15
<b>GLYCEMIC CONTROL</b>			
Hyperglycemia			
Percent with glucose $\geq 14$ mmol/L	56 (9.3%)	25 (17.7%)	6 (40%)
Percent with glucose $\geq 20$ mmol/L	15 (2.5%)	7 (5%)	2 (13.3%)
Percent with glucose $\geq 24$ mmol/L	8 (1.3%)	3 (2.1%)	1 (6.7%)
Hyperglycemia index (HGI): AUC ( $> 10$ mmol/L)/LOS (in hours)*			
	Median (IQR)		0 (1.8)
	Mean (SD)		1.6 (2.7)
Other metrics			
Percent with glucose $\geq 4$ and $< 10$ mmol/L	439 (73.2%)	110 (78%)	11 (73.3%)
Glucose (mmol/L)*			
	Median (IQR)	6.7 (4.9)	7 (4.9)
	Mean (SD)	8.2 (4.1)	8.7 (4)
Average patient-day mean glucose for a patient-stay (mmol/L)*			
	Median (IQR)		7 (4.9)
	Mean (SD)		8.7 (4)
<b>HYPOGLYCEMIA</b>			
Percent with glucose $< 4$ mmol/L	13 (2.2%)	10 (7.1%)	7 (46.7%)
Percent with glucose $< 3$ mmol/L	0 (0%)	0 (0%)	0 (0%)
Percent with glucose $< 2.5$ mmol/L	0 (0%)	0 (0%)	0 (0%)
Percent of patient-stays with a recurrent hypoglycemia ( $< 4$ mmol/L) day (10-240 mins)			2 (13.3%)
<b>GLYCEMIC VARIABILITY</b>			
Standard deviation: SD (mmol/L) <sup>2*</sup>			
	Median (IQR)	1.2 (2)	2 (1.9)
	Mean (SD)	1.6 (1.3)	2.4 (1.9)
J-index (mmol/L) <sup>2*</sup>			
	Median (IQR)	22.5 (37.9)	31.2 (57.6)
	Mean (SD)	41.5 (48.4)	52 (50.2)



\*:The summary statistics of the glucometrics for patient-day means and patient-stay means.

<sup>#1</sup>: 17 (12.06%) patient-days were removed for calculation of patient-day SD and J-index.

<sup>#2</sup>: 1 (6.67%) patient-stays were removed for calculation of patient-stay SD and J-index.

Summary Statistics report is saved in the folder: 'C:/Users/ningy/Documents/QcDM\_Project folder/data/Test1\_processed/Glucometric\_report' as a CSV file named 'Test1\_processed\_2020\_11\_16\_12\_38\_17.csv' and as a HTML report named 'Test1\_processed\_2020\_11\_16\_12\_38\_17.html'.

Tables above are also saved as a CSV file and a HTML report in the 'Glucometric\_report' folder for later use:

data > Test1_processed > Glucometric_report	
Name	^
	Test1_processed_2020_11_16_12_38_17
	Test1_processed_2020_11_16_12_38_17

The HTML report can also be saved as a PDF document by opening it using the Chrome browser:

**Glucometrics**

Summary statistics  
2020-11-16  
Data Summary

Summary of BG measurements	
Total (N)	
Minimum (mmol/L)	
5-th percentile (mmol/L)	
25-th percentile (mmol/L)	
75-th percentile (mmol/L)	
95-th percentile (mmol/L)	
Maximum (mmol/L)	31
Non-numeric values (N)	3

Exclusion Summary

Total number of patient-stays	15
No exclusion criterion was selected.	

Context menu options:

- Back (Alt+Left Arrow)
- Forward (Alt+Right Arrow)
- Reload (Ctrl+R)
- Save as... (Ctrl+S)
- Print... (Ctrl+P)
- Cast...
- Translate to English
- View page source (Ctrl+U)
- Inspect (Ctrl+Shift+I)

Callout box instructions:

- ① Right click on this page
- ② Click "Print..."
- ③ At the bottom-right corner of the pop-up window, click "Print"

After viewing glucometrics tables, users may reload QcDMui to process other glucose data:

~ /QcDM\_Project folder/code/QcDMui - Shiny

http://127.0.0.1:7854 | Open in Browser

QcDM UI | About the Interface | Data Summary

Callout box instructions:

- ① Click here to reload QcDMui
- ② Continue using QcDMui by repeating the steps in Section 3.3.

Users may also close the QcDMui window to return to the RStudio window. When closing the RStudio window, if users are prompted to select whether to save workspace image, there is no need to save either of them.

## 4. Running QcDM Project from USB Drive

In the previous three sections, we have described how to install and use the QcDM Project on a desktop with a Windows or Mac operating system. Alternatively, the QcDM Project can be installed into a USB Drive and used in a plug-and-play approach on any computer with the **Windows** operating system. This portable approach is not available for the Mac operating system.

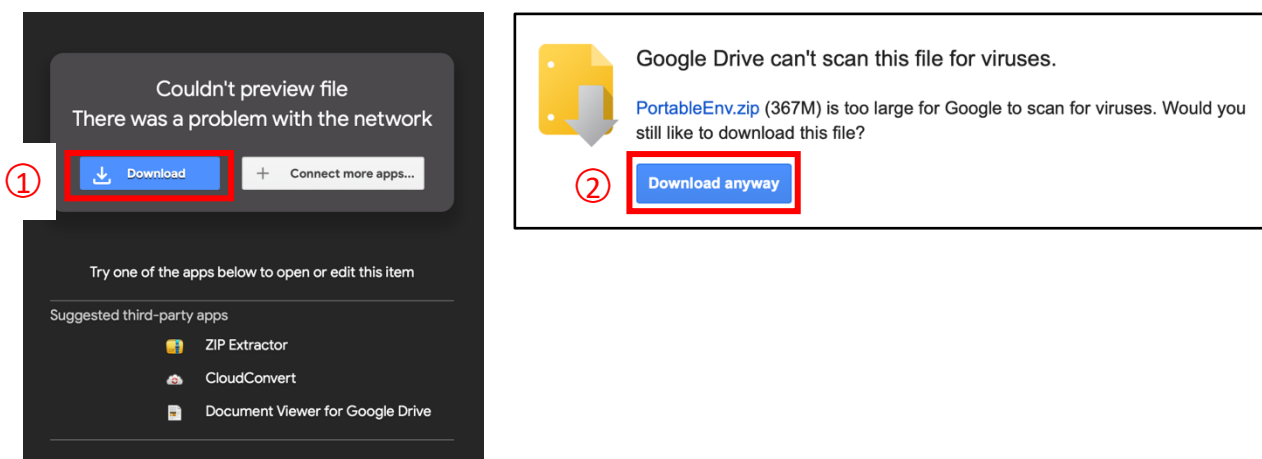
### 4.1. Set up the portable environment

The portable environment for running the QcDM Project, which includes R-portable (version 3.6.3) and two software (Pandoc and TinyTeX) for generating PDF reports.

Follow the instructions in Section 1.2 to download the QcDM Project as zip file from the GitHub repository, unzip it, rename the resulting folder as 'QcDM\_Project folder', and move this folder to the root directory of the USB Drive.

Download the portable environment as a zip file named 'PortableEnv.zip' from the following link:

<https://drive.google.com/file/d/1bPk8pwZ2HaCDbXhP8e67wcVzWNXs4R76/view?usp=sharing>.



Unzip the file (see Section 1.2 for instructions) to a folder named 'PortableEnv', and move this folder to the root directory of the USB Drive.

The root directory of the USB Drive should contain the following two folders, which take up approximately 1.8 GB on the disk:

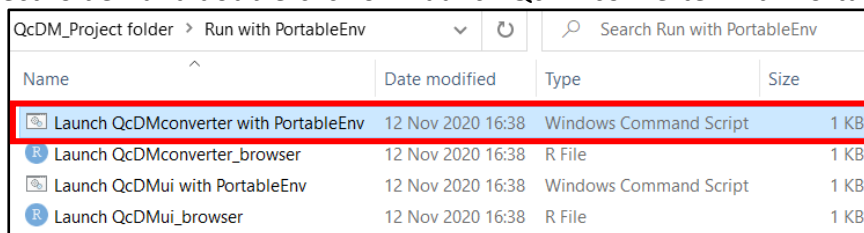
USB Drive (E:)			Search USB D
Name	Date modified	Type	
Other files	12 Nov 2020 18:03	File folder	
QcDM_Project folder	12 Nov 2020 18:14	File folder	
PortableEnv	12 Nov 2020 18:19	File folder	

The portable environment has been configured by installing the R packages and software required by the QcDM Project, therefore users do not need to go through the installation steps in Section 1.1 and 1.3.



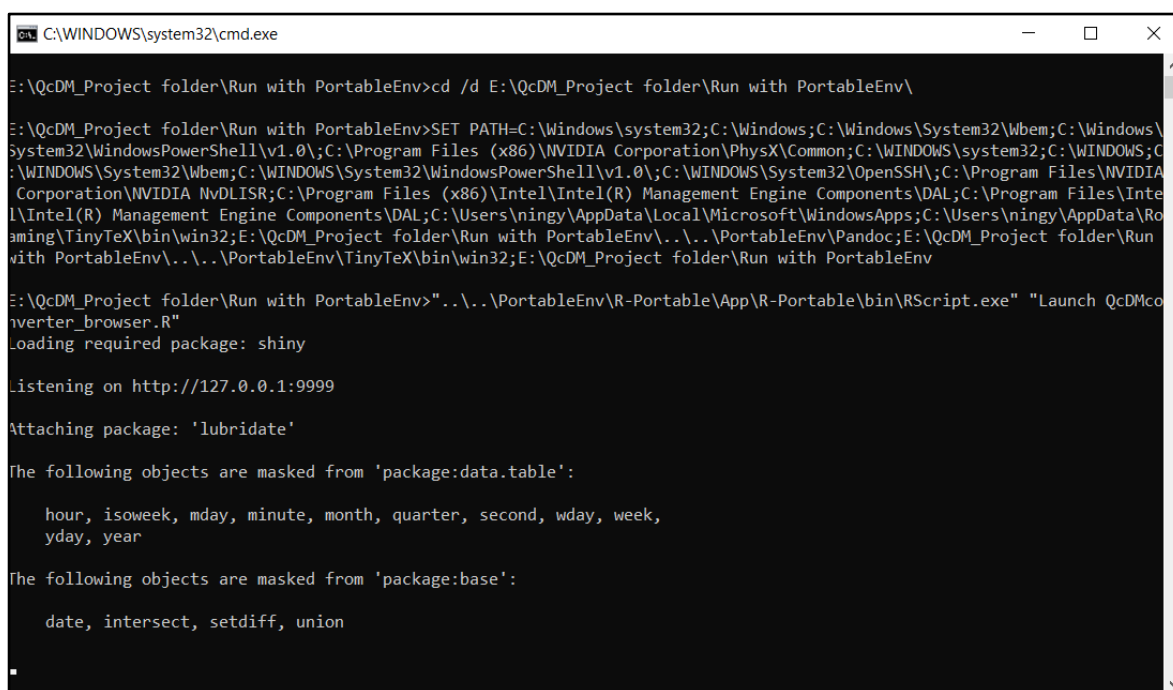
## 4.2. Run QcDMconverter with PortableEnv

To launch QcDMconverter from the USB Drive, go to the 'Run with PortableEnv' subfolder of 'QcDM\_Project folder' and double-click on 'Launch QcDMconverter with PortableEnv.cmd':



Name	Date modified	Type	Size
Launch QcDMconverter with PortableEnv	12 Nov 2020 16:38	Windows Command Script	1 KB
Launch QcDMconverter_browser	12 Nov 2020 16:38	R File	1 KB
Launch QcDMui with PortableEnv	12 Nov 2020 16:38	Windows Command Script	1 KB
Launch QcDMui_browser	12 Nov 2020 16:38	R File	1 KB

The following command line window will pop-up:



```
C:\WINDOWS\system32\cmd.exe

E:\QcDM_Project folder\Run with PortableEnv>cd /d E:\QcDM_Project folder\Run with PortableEnv\

E:\QcDM_Project folder\Run with PortableEnv>SET PATH=C:\Windows\system32;C:\Windows;C:\Windows\System32\Wbem;C:\Windows\System32\WindowsPowerShell\v1.0\;C:\Program Files (x86)\NVIDIA Corporation\PhysX\Common;C:\WINDOWS\system32;C:\WINDOWS;C:\WINDOWS\System32\Wbem;C:\WINDOWS\System32\WindowsPowerShell\v1.0\;C:\WINDOWS\System32\OpenSSH\;C:\Program Files\NVIDIA Corporation\NVIDIA NvDLISR;C:\Program Files (x86)\Intel\Intel(R) Management Engine Components\DAL;C:\Program Files\Intel\Intel(R) Management Engine Components\DAL;C:\Users\ningy\AppData\Local\Microsoft\WindowsApps;C:\Users\ningy\AppData\Local\Microsoft\WindowsApps;E:\QcDM_Project folder\Run with PortableEnv\..\..\PortableEnv\Pandoc;E:\QcDM_Project folder\Run with PortableEnv\..\..\PortableEnv\TinyTeX\bin\win32;E:\QcDM_Project folder\Run with PortableEnv

E:\QcDM_Project folder\Run with PortableEnv>"..\..\PortableEnv\R-Portable\App\R-Portable\bin\RScript.exe" "Launch QcDMconverter_browser.R"
Loading required package: shiny

Listening on http://127.0.0.1:9999

Attaching package: 'lubridate'

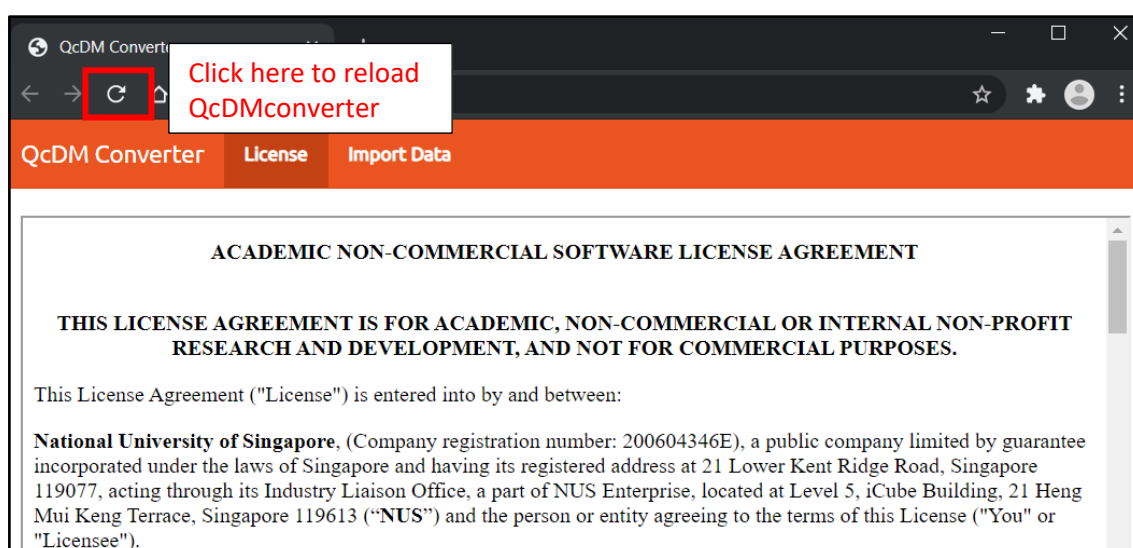
The following objects are masked from 'package:data.table':

    hour, isoweek, mday, minute, month, quarter, second, wday, week,
    yday, year

The following objects are masked from 'package:base':

    date, intersect, setdiff, union
```

and then QcDMconverter will open in the default web browser (in this example Chrome):



Users can follow the instructions in Section 2.1.2 or 2.2.1 to process glucose data.



Users can follow the instructions in Section 3.3 and 3.4 to process data and view glucometrics.

After successful completion of glucometrics generation, users may reload QcDMui and analyse additional glucose data by clicking the reload button of the web browser (as shown in the screenshot above). To close QcDMui, first close the QcDMui window in the web browser, then close the command line window.

## Bibliography

1. Thomas P, Inzucchi SE. An internet service supporting quality assessment of inpatient glycemic control. In: Journal of Diabetes Science and Technology. SAGE Publications Inc.; 2008. p. 402–8.
2. Chen Y, Kao SL, Tai E-S, Wee HL, Khoo EYH, Ning Y, et al. Utilizing distributional analytics and electronic records to assess timeliness of inpatient blood glucose monitoring in non-critical care wards. BMC Med Res Methodol. 2016;16(1).