#### User manual of the GUI version of PCprophet

## 1. Monitor resolution

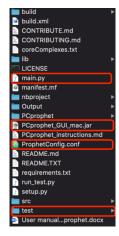
If you are running the GUI version of PCprophet using Apple products, such as laptop or PC, there is no requirement for the resolution as the resolution of GUI has been optimised. However, if you are using Windows system, the GUI version works best with the resolution of 1920\*1080.

### 2. Installing PCprophet and Java

In order to run the GUI version of PCprophet, the command line version must be successfully installed. Please follow the instructions for installing the command line version of PCprophet and required dependencies on our GitHub repository (<a href="https://github.com/anfoss/PCprophet">https://github.com/anfoss/PCprophet</a>). Besides, Java (version >= 1.8) needs to be installed and the Java paths need to be appropriately configured. To test if Java has been successfully installed, please try 'java', 'javac' to see if the help document can display in the console, as demonstrated in the following figures:

#### 3. Folder structure

In general, after unzipping the file, the structure of PCprophet GUI folder will be shown as the following:



In this folder, there are four important files/folders, including (i) PCprophet\_GUI\_win.jar, which is the main executable file of the GUI version of PCprophet; (ii) main.py, which is the main program of PCprophet; (iii) ProphetConfig.conf, which contains all parameter settings for PCprophet; and (iv) the test folder, which is for input files and sample description file. As PCprophet is able to deal with coelution profiles of different replicates and conditions, we need users to provide a descriptive file for these replicates and conditions, the details of which will be described in the following section. In addition, there is no need to manually change the ProphetConfig.conf file. Once the parameters are configured via the GUI version of PCprophet, the parameters will be written and updated to the ProphetConfig.conf file by the main.py, for your future reference. Note that during the prediction by PCprophet, a temporary folder (termed 'tmp') will be generated to store the intermediate files during the prediction and analysis. It is suggested that before running PCprophet, this folder should be deleted.

## 4. <u>Updating the PCprophet folder</u>

As the GUI version is dependent on the original command line version of PCprophet, it is therefore crucial to keep the "PCprophet" folder (see above figure) up to date. To do so, before running the GUI version, please copy and paste the most up to date PCprophet source code folder to into the GUI folder. Note that this folder doesn't exist in the initial GUI folder, please copy the "PCprophet" folder to the GUI folder before running.

# 5. Running the GUI version of PCprophet using provided example

To run the GUI version of PCprophet, please use the console of your operating system and locate to the PCprophet folder. For example, in Windows, when the console (i.e. the Command Prompt) has started, type the 'cd' command to relocate to the PCprophet folder. In this example, the folder is in "D:\test\PCprophet GUI win":

```
Command Prompt

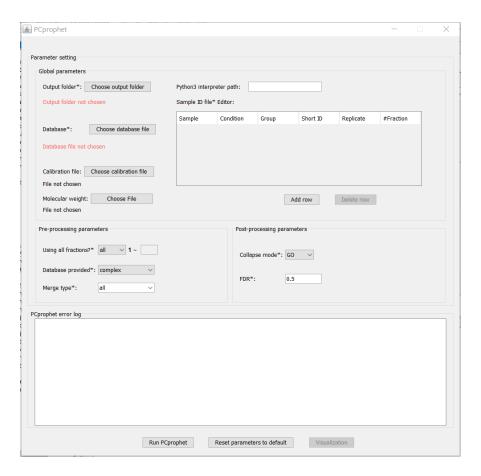
Microsoft Windows [Version 10.0.18363.1139]

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C:\Users\chenl>D:

D:\>cd test\PCprophet_GUI_win
```

Then type "java -jar PCprophet\_GUI\_win.jar" to initiate the GUI version of PCprophet. The main interface should display as shown in the following figure:



All the mandatory parameters are marked with a '\*' and must be provided. As can been from the figure, some of the parameters have been provided using their default values. Please refer to our GitHub repository for the detailed explanation of these parameters.

Due to the fact that the macOS system also has built-in Python (version 2.x), PCprophet may have problem using the right version of Python (i.e. version >3.0). Therefore, we would like users to provide the location of the Python 3 interpreter. This path can be easily obtained by the console of Windows and macOS. For example, in Windows, simply type 'where python3', the path of the Python 3 interpreter will be shown as follows:

```
Command Prompt

Microsoft Windows [Version 10.0.18363.1139]

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C:\Users\chenl>where python

C:\Users\chenl\anaconda3\python.exe

C:\Users\chenl>
```

Users can just copy the content in the red rectangle and paste into the text box of 'Python 3 interpreter path'.

The 'Sample file ID editor' is designed for users to provide the basic information of their experiment. A file named 'sample\_ids.txt' will be generated based on the input to this editor and the file will be saved to the test folder, where all the co-elution files of the experiment should be stored as

well. The following figure demonstrate the information we have put for the example test file (i.e. test frac.txt) we provided in the 'test' folder:



There are six columns of information to provide by clicking the 'Add row' button in the 'Sample' column, the co-elution file of current condition and replicate should be provided using the relative file path: "./test/test\_frac.txt". Notice that the "test" folder is the one for storing the replicate and condition information. Please make sure that the file name is correct otherwise PCprophet will have problems recognising and locating the file. All the co-elution files should be described using this editor. After you click the 'Run PCprophet' button, you will be able to see a 'sample\_ids.txt' file has been generated in the 'test' folder:



And the 'sample ids.txt' file should content the following information:



After the 'Run PCprophet' button has been clicked, the prediction process will be initiated. When the process is over (either successfully finished or interrupted by errors), a dialog box will show as follows:



The 'PCprophet error log' was designed to pass the error message to the users to indicate anything that goes wrong during the prediction. However, the warning messages, which do not affect the prediction and analysis processes, are also shown here:

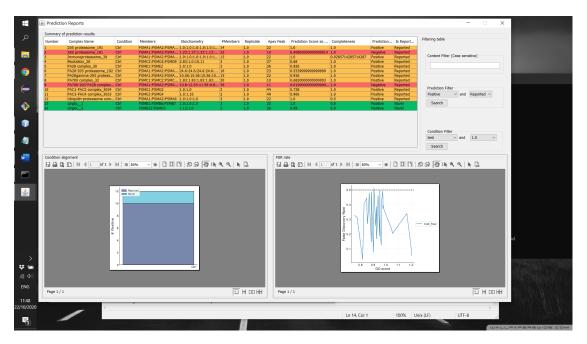
```
PCprophet error log

=====Starting running PCprophet at 11:10:49.644=====
self.cor.append(np.hstack((D / np.sqtt(ssAs * ssBs), np.zeros(9) + np.nan)))
D:PCprophet_GUI_winPCprophet\collapse.py:312: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

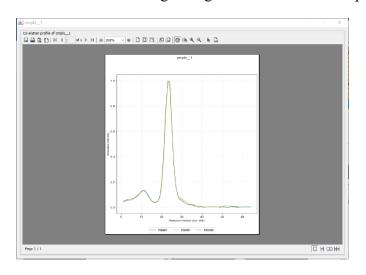
See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html$return:
tmp["ID"] = "cmplx_" + str(count)
=====PCprophet has finished running at 11:24:48.928=====
```

We recommend users not to close the program when PCprophet is still running at the backstage. This is due the mechanism of Java invoking other non-Java programs (Python program in our case). If you would like to kill the running PCprophet, simply go to the Task Manager (for Windows) or Activity Monitor (for macOS). Based on the security consideration, the current GUI version of PCprophet does not provide the option for parallel processing, which will be offered in our future versions once the security issue is tackled.

After PCprophet has successfully finished the prediction, users can click the 'Visualization' button to review the prediction results:



Users can further filter the prediction results using the options provided in the top-right panel. If users would like to come back to all the results after filtering, simply go to the text input area of 'Content Filter' and hit the 'backspace' key on the keyboard. Users can also click any row in the table to closely examine the co-elution profile of selected complex. For example, if the 'cmplx\_1' (i.e. #13) is clicked, users will be able to see the following dialog box for its co-elution profiles:



## 6. About the current version of PCprophet GUI

The current version of PCprophet GUI is able to perform PCprophet prediction and demonstrate the protein complex prediction results. The display of protein complex differential analysis results is currently not included. We will include this function in the future version of GUI. For complex differential analysis, please refer to the 'DifferentialComplexReport.txt' and 'DifferentialProteinReport.txt' in the output folder for more details.

## 7. Contact

This GUI is still a beta version. For questions, bug reports and suggestions, please feel free to contact Dr Andrea Fossati (Andrea.Fossati@ucsf.edu) and Dr Chen Li (Chen.Li@monash.edu).