Component specification

Software components:

1. Graphical User Interface (GUI)

o Description: Provides a user-friendly interface for researchers to interact with the software package, facilitating ease of navigation and operation.

2. File Import Function

 Description: Allows users to import data from .csv and .xlsx files seamlessly into the system for processing.

3. Data Selection Function

Description: Enables users to select specific data fields such as Name,
Concentration, and other relevant metrics for analysis.

4. Regulation Analysis Function

 Description: Implements statistical analysis (e.g., t-test) to identify downregulators and up-regulators from the dataset, assisting researchers in pinpointing key metabolites.

5. Data Output Function

 Description: Generates formatted output data that can be easily exported or utilized in further analysis.

6. Clipboard Function

 Description: Provides a clipboard feature for users to copy and paste data directly into MetaboAnalyst or other platforms.

7. Metabolite Separation Function

 Description: Facilitates the separation of different metabolites within the dataset for more focused analysis.

8. Metabolite Identification Function

 Description: Identifies metabolites and converts their names into PubChem IDs, linking to an extensive database for further research.

Interactions to Accomplish Use Cases

- 1. Use Case 1: Metabolite Selection
 - The researcher utilizes the Graphical User Interface (GUI) to import raw data files (.csv or .xlsx).
 - They select relevant parameters such as Name and Concentration using the Data Selection Function.
 - o The researcher then uses the Metabolite Separation Function to filter specific metabolites of interest.
 - o Finally, the Data Output Function generates a formatted list of metabolites, which can be copied using the Clipboard Function for further analysis.

2. Use Case 2: Regulation Analysis

- o The researcher accesses the GUI to import their experimental data.
- They select the relevant variables for analysis, including concentrations of potential regulators, through the Data Selection Function.
- o The researcher initiates the Regulation Analysis Function, which uses a t-test to identify down-regulators and up-regulators in the dataset.
- The results are displayed in the output area, where the researcher can utilize the Clipboard Function to copy the findings directly to MetaboAnalyst or other platforms.

Preliminary Plan

- 11/13: Complete the development of basic functions that yield a usable output compatible with MetaboAnalyst.
- 11/20: Finalize testing of the GUI and ensure all import and selection functions are operational.
- 11/27: Implement the regulation analysis and output functions, ensuring accurate data processing.
- 12/1: Conduct final reviews and testing, addressing any issues, and prepare for project completion.