## Table 7: MS Data Analysis - Feedback and Suggestions to Improve Software Tools

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## Scope:

Liquid chromatography-mass spectrometry (LC-MS) is an essential tool for identification, and structure elucidation for a wide array of molecules from small organic molecule, to large biotherapeutics. The multi-dimensional data matrix obtained from LC-MS/MS (or MS/MS) analyses makes the data interpretation complex. The roundtable discussion is intended to establish better guidance with regard to using data analysis software, and application of LC-MS. A better understanding of current practices to handle LC-MS (and MS/MS) data will be achieved, and experience in the field will be exchanged.

## **Questions for Discussion:**

- 1. Could mass spectrometer manufacturers provide a better guidance to the different software available to the end users? Could trainings on these software depending on the applications be provided earlier in the process of the acquisition of a mass spectrometer?
- 2. How to analyze the LC-MS/MS data: software or manual interpretation? Raw data preprocess: noise removal, peak detection, etc., list of software, manual analysis tools.
- 3. Could the following tools be integrated in current software: i) integration of HTML format to generate live data, with the possibility to zoom into the details of a mass spectrum or an LC trace after data were processed, ii) Integration of tools to generate publishable high-quality figures.
- 4. What kinds of information can be achieved from LC-MS/MS analysis and how to achieve? Sequence coverage, identification, structure elucidation, quantitation. What is currently available to analyze automatically small organic molecules?
- 5. LC-MS/MS and Machine Learning integration.
- 6. What are critical areas of improvement for MS data analysis software?
- 7. How to deal with false positives and false negatives?

## **Discussion Notes:**

This table did not have attendees.