

## Reviewer Report

**Title:** Accessible and reproducible mass spectrometry imaging data analysis in Galaxy

**Version:** Original Submission    **Date:** 6/13/2019

**Reviewer name:** Jane Armstrong

### Reviewer Comments to Author:

This is an interesting paper that provides new tools for the Galaxy platform and that will be of use to biomedical researchers generating and analysing MSI data. The authors demonstrate the value of the toolset that they have developed by reanalysing a published dataset and they provide open access to source code and additional training documentation. Developing open source tools is commendable and I therefore support publication.

Minor Revisions:

Section describing the new MSI toolset: Making more of a distinction in the text between the newly developed tools and existing Galaxy tools would be helpful to the reader. I also suggest adding 'tool(s)' to each subheading within this section i.e. 'Quality control and visualization tools:'

Figure 1: The figure title should make it clear that these are the newly developed Galaxy tools: the use of 'associated' is ambiguous. The figure legend could be improved by clarifying that the MSI data analysis steps are in dark blue and the new tools are shown in pale blue.

Sectioning describing Accessibility and training: I found navigation to the training sets difficult. The link to the Galaxy Training Network/repository is to the home page and the reader then has to click through several pages to find each of the 3 examples. The names are not obvious from the manuscript and I had to click through a number of other examples to find them. The training material is excellent but just difficult to find.

Availability of supporting data: I did not seem to have the necessary permission to access the datasets provided in the last link.

Add FAIR to the list of abbreviations.

### Level of Interest

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Please indicate the quality of language in the manuscript: Choose an item.

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