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To the editors of

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**Revised manuscript “Integrated enrichment analysis and pathway-centered visualization of metabolomics, proteomics, transcriptomics, and genomics data by using the InCroMAP software”, submitted under manuscript number JCB-13-943**

Dear Mrs. ten Napel, Dear Professor Theodoridis,

Thank you very much for your very kind offer for resubmission of our manuscript and your very valuable suggestions to improve the content.

We added an additional figure (new Fig. 2c) to Figure 2 showing additional metabolites to illustrate the InCroMAP data evaluation further and modified the text accordingly. We are aware that not so many metabolites are visible in Figure 2, but we consciously selected this data set to demonstrate the power of InCroMAP because it is includes a rare combination of a variety of omics data, like metabolite levels, mRNA, miRNA, protein expression and DNA methylation.

I apologize for the delay of my answers based on my high workload in my new job in the industry. I hope that the revised version of our manuscript is now acceptable for publication and thank you again for your very kind consideration.

Sincerely yours,

Lars Rosenbaum

(on behalf of the authors)