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26. September 2013

Dear Professor Theodoridis,

Thank you very much for your invitation to contribute a paper to the Metabolomics special issue of the Journal of Chromatography B and for your very kind extension of the deadline. Enclosed you find our manuscript entitled “Straightforward interpretation of metabolomics, proteomics, transcriptomics, and genomics data by comprehensive visualization and pathway enrichment using a single software tool”.

The manuscript presents an extended version of the Java-based tool InCroMAP, which is able to perform an integrated enrichment analysis and pathway-based visualization of multi-omics data. The tool was originally designed for the evaluation of genomics, transcriptomics, and proteomics data from various platform types. The novel, extended version presented in the manuscript supports the analysis of annotated metabolomics data, thereby complementing the tool for comprehensive systems biology data evaluation.

We think this manuscript is of great interest to the readers of the Journal of Chromatography B.

Thank you for considering our work. Please feel free to correspond with me by e-mail (lars.rosenbaum@uni-tuebingen.de).

Sincerely,

Lars Rosenbaum