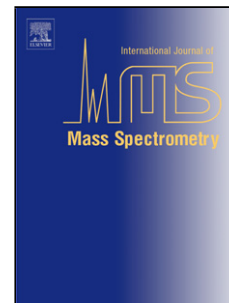


Accepted Manuscript

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PII: S1387-3806(17)30269-5
DOI: <http://dx.doi.org/10.1016/j.ijms.2017.08.014>
Reference: MASPEC 15857

To appear in: *International Journal of Mass Spectrometry*

Received date: 23-5-2017
Revised date: 10-8-2017
Accepted date: 22-8-2017

Please cite this article as: Lukasz G.Migas, Aidan P.France, Bruno Bellina, Perdita E.Barran, *ORIGAMI*: A software suite for activated ion mobility mass spectrometry (aIM-MS) applied to multimeric protein assemblies, *International Journal of Mass Spectrometry*<http://dx.doi.org/10.1016/j.ijms.2017.08.014>

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ORIGAMI: A Software Suite for Activated Ion Mobility Mass Spectrometry (aIM-MS) Applied To Multimeric Protein Assemblies

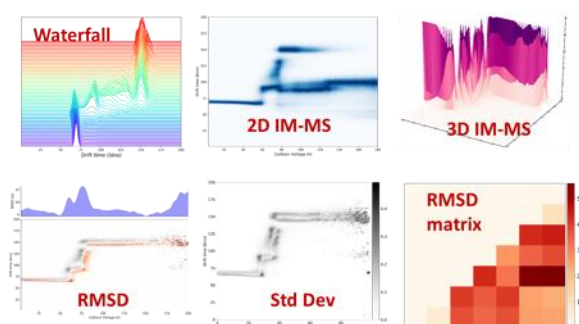
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Graphical abstract



Highlights

- New methodology for controlling acquisition parameters and faster data analysis following activation of ions separated by ion mobility mass spectrometry.
- Software package capable of simultaneous analysis of multiple MassLynx .raw files.
- Visualization of the change in mobilities for both parent and product ions following activation.
- Easy extraction, data processing and extensive plotting tools.

Abstract

We present here a software suite (**ORIGAMI**) that facilitates the rapid acquisition and analysis of ion mobility data following collisional activation. **ORIGAMI** was developed for use on Waters Synapt instruments where data acquisition is achieved by interfacing WREnS (Waters Research Enabled Software) and MassLynx. Two components are presented, the first is **ORIGAMI^{MS}** which enables the activation of ions via a sequential ramping of collision voltage prior to ion mobility analysis. We

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