

CITATION REPORT

List of articles citing

MASH Suite: a user-friendly and versatile software interface for high-resolution mass spectrometry data interpretation and visualization

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| # | Paper | IF | Citations |
|----|---|------|-----------|
| 62 | Impact of Phosphorylation on the Mass Spectrometry Quantification of Intact Phosphoproteins. | | |
| 61 | Top-down proteomics in health and disease: challenges and opportunities. <i>Proteomics</i> , 2014 , 14, 1195-2108 | 10.8 | 135 |
| 60 | Top-down proteomics reveals concerted reductions in myofilament and Z-disc protein phosphorylation after acute myocardial infarction. <i>Molecular and Cellular Proteomics</i> , 2014 , 13, 2752-64 | 7.6 | 81 |
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| 58 | Middle-down mass spectrometry enables characterization of branched ubiquitin chains. <i>Biochemistry</i> , 2014 , 53, 4979-89 | 3.2 | 61 |
| 57 | Middle-Down and Top-Down Mass Spectrometric Analysis of Co-occurring Histone Modifications. <i>Current Protocols in Protein Science</i> , 2014 , 77, 23.7.1-23.7.28 | 3.1 | 25 |
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| 2 | Top-Down Characterization and Intact Mass Quantitation of a Monoclonal Antibody Drug from Serum by Use of a Quadrupole TOF MS System Equipped with Electron-Activated Dissociation. | | 0 |
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