

Kevin Jeanne Dit Fouque

List of Publications by Year in descending order

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papers

670
citations

623734

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34
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docs citations

34
times ranked

655
citing authors

#	ARTICLE	IF	CITATIONS
1	AT-hook peptides bind the major and minor groove of AT-rich DNA duplexes. <i>Nucleic Acids Research</i> , 2022, 50, 2431-2439.	14.5	6
2	Nanomolar affinity of EF-hands in neuronal calcium sensor 1 for bivalent cations Pb ²⁺ , Mn ²⁺ , and Hg ²⁺ . <i>Metallomics</i> , 2022, 14, .	2.4	6
3	Trapped Ion Mobility Spectrometry, Ultraviolet Photodissociation, and Time-of-Flight Mass Spectrometry for Gas-Phase Peptide Isobars/Isomers/Conformers Discrimination. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 1267-1275.	2.8	12
4	Exploring the Conformations and Binding Location of HMGA2-DNA Complexes Using Ion Mobility Spectrometry and 193 nm Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 1092-1102.	2.8	2
5	Exploring the Conformational and Binding Dynamics of HMGA2-DNA Complexes Using Trapped Ion Mobility Spectrometry-Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 1103-1112.	2.8	4
6	Structural Insights from Tandem Mass Spectrometry, Ion Mobility-Mass Spectrometry, and Infrared/Ultraviolet Spectroscopy on Sphingonodin I: Lasso vs Branched-Cyclic Topoisomers. <i>Journal of the American Society for Mass Spectrometry</i> , 2021, 32, 1096-1104.	2.8	4
7	Exploring structural signatures of the lanthipeptide prochlorosin 2.8 using tandem mass spectrometry and trapped ion mobility-mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 4815-4824.	3.7	9
8	Proteoform Differentiation using Tandem Trapped Ion Mobility, Electron Capture Dissociation, and ToF Mass Spectrometry. <i>Analytical Chemistry</i> , 2021, 93, 9575-9582.	6.5	18
9	A Bifunctional Leader Peptidase/ABC Transporter Protein Is Involved in the Maturation of the Lasso Peptide Cochochodin I from <i>Streptococcus suis</i> . <i>Journal of Natural Products</i> , 2021, 84, 2683-2691.	3.0	11
10	Trapped Ion Mobility Spectrometry of Native Macromolecular Assemblies. <i>Analytical Chemistry</i> , 2021, 93, 2933-2941.	6.5	32
11	Effective discrimination of gas-phase peptide conformers using TIMS-ECD-ToF MS/MS. <i>Analytical Methods</i> , 2021, 13, 5216-5223.	2.7	6
12	Substrate Sequence Controls Regioselectivity of Lanthionine Formation by ProcM. <i>Journal of the American Chemical Society</i> , 2021, 143, 18733-18743.	13.7	19
13	Following Structural Changes by Thermal Denaturation Using Trapped Ion Mobility Spectrometry-Mass Spectrometry. <i>Journal of Physical Chemistry B</i> , 2020, 124, 6257-6265.	2.6	11
14	Salt bridges govern the structural heterogeneity of heme protein interactions and porphyrin networks: microperoxidase-11. <i>RSC Advances</i> , 2020, 10, 33861-33867.	3.6	0
15	Dynamics of the E. coli β -Clamp Dimer Interface and Its Influence on DNA Loading. <i>Biophysical Journal</i> , 2019, 117, 587-601.	0.5	12
16	Exploring the Conformational Space of Growth-Hormone-Releasing Hormone Analogues Using Dopant Assisted Trapped Ion Mobility Spectrometry-Mass Spectrometry. <i>Journal of Physical Chemistry B</i> , 2019, 123, 6169-6177.	2.6	8
17	Recent advances in biological separations using trapped ion mobility spectrometry - mass spectrometry. <i>TrAC - Trends in Analytical Chemistry</i> , 2019, 116, 308-315.	11.4	52
18	Effective Liquid Chromatography-Trapped Ion Mobility Spectrometry-Mass Spectrometry Separation of Isomeric Lipid Species. <i>Analytical Chemistry</i> , 2019, 91, 5021-5027.	6.5	64

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19	Evidence of <i>Cis</i> / <i>Trans</i> -Isomerization at Pro7/Pro16 in the Lasso Peptide Microcin J25. <i>Journal of the American Society for Mass Spectrometry</i> , 2019, 30, 1038-1045.	2.8	12
20	Structural signatures of the class III lasso peptide BI-32169 and the branched-cyclic topoisomers using trapped ion mobility spectrometry–mass spectrometry and tandem mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2019, 411, 6287-6296.	3.7	20
21	Measuring the Integrity of Gas-Phase Conformers of Sodiated 25-Hydroxyvitamin D3 by Drift Tube, Traveling Wave, Trapped, and High-Field Asymmetric Ion Mobility. <i>Analytical Chemistry</i> , 2019, 91, 4092-4099.	6.5	13
22	Microheterogeneity of Topoisomerase IA/IB and Their DNA-Bound States. <i>ACS Omega</i> , 2019, 4, 3619-3626.	3.5	9
23	General rules of fragmentation evidencing lasso structures in CID and ETD. <i>Analyst, The</i> , 2018, 143, 1157-1170.	3.5	27
24	Insights from ion mobility-mass spectrometry, infrared spectroscopy, and molecular dynamics simulations on nicotinamide adenine dinucleotide structural dynamics: NAD ⁺ vs. NADH. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 7043-7052.	2.8	14
25	Linear and Differential Ion Mobility Separations of Middle-Down Proteoforms. <i>Analytical Chemistry</i> , 2018, 90, 2918-2925.	6.5	43
26	Metal ions induced secondary structure rearrangements: mechanically interlocked lasso vs. unthreaded branched-cyclic topoisomers. <i>Analyst, The</i> , 2018, 143, 2323-2333.	3.5	12
27	Identification of Lasso Peptide Topologies Using Native Nano-electrospray Ionization-Trapped Ion Mobility Spectrometry–Mass Spectrometry. <i>Analytical Chemistry</i> , 2018, 90, 5139-5146.	6.5	34
28	Structural Motif Descriptors as a Way To Elucidate the Agonistic or Antagonistic Activity of Growth Hormone–Releasing Hormone Peptide Analogues. <i>ACS Omega</i> , 2018, 3, 7432-7440.	3.5	8
29	Fast and Effective Ion Mobility–Mass Spectrometry Separation of <i>d</i> -Amino-Acid-Containing Peptides. <i>Analytical Chemistry</i> , 2017, 89, 11787-11794.	6.5	76
30	Signatures of Mechanically Interlocked Topology of Lasso Peptides by Ion Mobility–Mass Spectrometry: Lessons from a Collection of Representatives. <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 315-322.	2.8	17
31	Characterization of Intramolecular Interactions of Cytochrome <i>c</i> Using Hydrogen–Deuterium Exchange-Trapped Ion Mobility Spectrometry–Mass Spectrometry and Molecular Dynamics. <i>Analytical Chemistry</i> , 2017, 89, 8757-8765.	6.5	35
32	IRMPD Spectroscopy: Evidence of Hydrogen Bonding in the Gas Phase Conformations of Lasso Peptides and their Branched-Cyclic Topoisomers. <i>Journal of Physical Chemistry A</i> , 2016, 120, 3810-3816.	2.5	15
33	Gas-phase conformations of capistruin – comparison of lasso, branched-cyclic and linear topologies. <i>Rapid Communications in Mass Spectrometry</i> , 2015, 29, 1411-1419.	1.5	11
34	Ion Mobility–Mass Spectrometry of Lasso Peptides: Signature of a Rotaxane Topology. <i>Analytical Chemistry</i> , 2015, 87, 1166-1172.	6.5	48