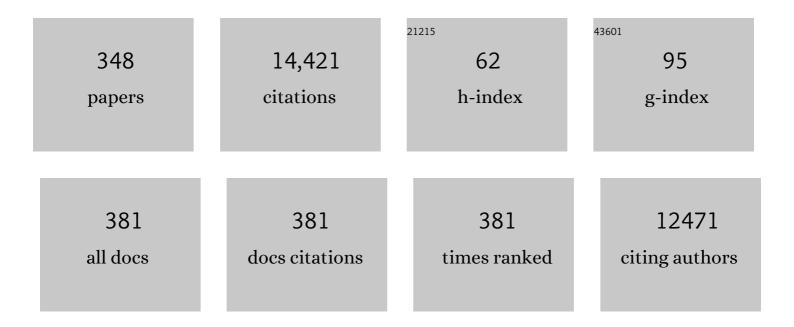
David H Russell

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Temperature Regulates Stability, Ligand Binding (Mg ²⁺ and ATP), and Stoichiometry of GroEL–GroES Complexes. Journal of the American Chemical Society, 2022, 144, 2667-2678.	6.6	18
2	Variable-Temperature Native Mass Spectrometry for Studies of Protein Folding, Stabilities, Assembly, and Molecular Interactions. Annual Review of Biophysics, 2022, 51, 63-77.	4.5	18
3	Cupric Ions Selectively Modulate TRAAK–Phosphatidylserine Interactions. Journal of the American Chemical Society, 2022, 144, 7048-7053.	6.6	4
4	Characterization of lipid carbon–carbon double-bond isomerism via ion mobility-mass spectrometry (IMS-MS) combined with cuprous ion-induced fragmentation. International Journal of Mass Spectrometry, 2022, 479, 116889.	0.7	6
5	THE IMS PARADOX: A PERSPECTIVE ON STRUCTURAL ION MOBILITYâ€MASS SPECTROMETRY. Mass Spectrometry Reviews, 2021, 40, 280-305.	2.8	29
6	A SPECIAL ISSUE DEDICATED TO THE OUTSTANDING SCIENTIFIC CAREER OF PROF. MICHAEL L. GROSS. Mass Spectrometry Reviews, 2021, 40, 161-161.	2.8	0
7	Selective regulation of human TRAAK channels by biologically active phospholipids. Nature Chemical Biology, 2021, 17, 89-95.	3.9	24
8	Editorial: Focus on Ionization Technologies Used in MS: Fundamentals and Applications, Honoring Dr. Sarah Trimpin, Recipient of the 2019 ASMS Biemann Medal. Journal of the American Society for Mass Spectrometry, 2021, 32, 616-617.	1.2	0
9	Molecular assemblies of the catalytic domain of SOS with KRas and oncogenic mutants. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	15
10	Variable-Temperature Electrospray Ionization for Temperature-Dependent Folding/Refolding Reactions of Proteins and Ligand Binding. Analytical Chemistry, 2021, 93, 6924-6931.	3.2	33
11	Thermal Analysis of a Mixture of Ribosomal Proteins by vT-ESI-MS: Toward a Parallel Approach for Characterizing the Stabilitome. Analytical Chemistry, 2021, 93, 8484-8492.	3.2	8
12	Protons Are Fast and Smart; Proteins Are Slow and Dumb: On the Relationship of Electrospray Ionization Charge States and Conformations. Journal of the American Society for Mass Spectrometry, 2021, 32, 1553-1561.	1.2	17
13	Editorial: Focus on Protein Footprinting, Honoring Michael Gross, Recipient of the 2020 John B. Fenn Award for a Distinguished Contribution in Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2021, 32, 1565-1566.	1.2	Ο
14	Implementing Digital-Waveform Technology for Extended <i>m</i> / <i>z</i> Range Operation on a Native Dual-Quadrupole FT-IM-Orbitrap Mass Spectrometer. Journal of the American Society for Mass Spectrometry, 2021, 32, 2812-2820.	1.2	9
15	Entropy in the Molecular Recognition of Membrane Protein–Lipid Interactions. Journal of Physical Chemistry Letters, 2021, 12, 12218-12224.	2.1	10
16	Native IM-Orbitrap MS: Resolving what was hidden. TrAC - Trends in Analytical Chemistry, 2020, 124, 115533.	5.8	33
17	Structural Analysis of 14-3-3-ζ-Derived Phosphopeptides Using Electron Capture Dissociation Mass Spectrometry, Traveling Wave Ion Mobility Spectrometry, and Molecular Modeling. Journal of Physical Chemistry B, 2020, 124, 461-469.	1.2	5
18	Development of native MS capabilities on an extended mass range Q-TOF MS. International Journal of Mass Spectrometry, 2020, 458, 116451.	0.7	13

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19	First-Principles Collision Cross Section Measurements of Large Proteins and Protein Complexes. Analytical Chemistry, 2020, 92, 11155-11163.	3.2	24
20	Discovery of Potent Charge-Reducing Molecules for Native Ion Mobility Mass Spectrometry Studies. Analytical Chemistry, 2020, 92, 11242-11249.	3.2	21
21	Evidence for Many Unique Solution Structures for Chymotrypsin Inhibitor 2: A Thermodynamic Perspective Derived from vT-ESI-IMS-MS Measurements. Journal of the American Chemical Society, 2020, 142, 17372-17383.	6.6	26
22	Ag ⁺ Ion Binding to Human Metallothionein-2A Is Cooperative and Domain Specific. Analytical Chemistry, 2020, 92, 8923-8932.	3.2	19
23	Molecular Mechanism of ISC Iron–Sulfur Cluster Biogenesis Revealed by High-Resolution Native Mass Spectrometry. Journal of the American Chemical Society, 2020, 142, 6018-6029.	6.6	30
24	Structural Analysis of the Effect of a Dual-FLAG Tag on Transthyretin. Biochemistry, 2020, 59, 1013-1022.	1.2	7
25	Tracking the Structural Evolution of 4-Aminobenzoic Acid in the Transition from Solution to the Gas Phase. Journal of Physical Chemistry B, 2020, 124, 2081-2087.	1.2	24
26	Melting of Hemoglobin in Native Solutions as measured by IMS-MS. Analytical Chemistry, 2020, 92, 3440-3446.	3.2	20
27	Collision-Induced Unfolding Studies of Proteins and Protein Complexes using Drift Tube Ion Mobility-Mass Spectrometer. Analytical Chemistry, 2020, 92, 7218-7225.	3.2	18
28	Solvent Mediation of Peptide Conformations: Polyproline Structures in Water, Methanol, Ethanol, and 1-Propanol as Determined by Ion Mobility Spectrometry-Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2019, 30, 77-84.	1.2	19
29	Development and Evaluation of a Reverse-Entry Ion Source Orbitrap Mass Spectrometer. Journal of the American Society for Mass Spectrometry, 2019, 30, 192-198.	1.2	21
30	Intrinsic GTPase Activity of K-RAS Monitored by Native Mass Spectrometry. Biochemistry, 2019, 58, 3396-3405.	1.2	25
31	Topological Characterization of Coordination-Driven Self-assembly Complexes: Applications of Ion Mobility-Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2019, 30, 1654-1662.	1.2	15
32	Topological Analysis of Transthyretin Disassembly Mechanism: Surface-Induced Dissociation Reveals Hidden Reaction Pathways. Analytical Chemistry, 2019, 91, 2345-2351.	3.2	22
33	Substance P in Solution: Trans-to-Cis Configurational Changes of Penultimate Prolines Initiate Non-enzymatic Peptide Bond Cleavages. Journal of the American Society for Mass Spectrometry, 2019, 30, 919-931.	1.2	13
34	Focus on Ion Mobility Spectrometry, Honoring Gert von Helden, Martin F. Jarrold, and David E. Clemmer, Recipients of the 2018 John B. Fenn Award for a Distinguished Contribution in Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 2019, 30, 893-897.	1.2	0
35	67TH ASMS Conference on Mass Spectrometry and Allied Topics. Journal of the American Society for Mass Spectrometry, 2019, 30, 1-295.	1.2	Ο
36	Variable-Temperature ESI-IMS-MS Analysis of Myohemerythrin Reveals Ligand Losses, Unfolding, and a Non-Native Disulfide Bond. Analytical Chemistry, 2019, 91, 6808-6814.	3.2	23

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37	New insights into the metal-induced oxidative degradation pathways of transthyretin. Chemical Communications, 2019, 55, 4091-4094.	2.2	18
38	Hydration of Guanidinium Ions: An Experimental Search for Like-Charged Ion Pairs. Journal of Physical Chemistry Letters, 2019, 10, 1349-1354.	2.1	9
39	Selective binding of a toxin and phosphatidylinositides to a mammalian potassium channel. Nature Communications, 2019, 10, 1352.	5.8	27
40	Substance P in the Gas Phase: Conformational Changes and Dissociations Induced by Collisional Activation in a Drift Tube. Journal of the American Society for Mass Spectrometry, 2019, 30, 932-945.	1.2	8
41	A Focus Honoring Carol Robinson's Election to the National Academy of Sciences. Journal of the American Society for Mass Spectrometry, 2019, 30, 1-3.	1.2	3
42	Influence of water and enzyme SpnF on the dynamics and energetics of the ambimodal [6+4]/[4+2] cycloaddition. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E848-E855.	3.3	57
43	Melting proteins confined in nanodroplets with 10.6 μm light provides clues about early steps of denaturation. Chemical Communications, 2018, 54, 3270-3273.	2.2	18
44	Collision-Induced Unfolding of Partially Metalated Metallothionein-2A: Tracking Unfolding Reactions of Gas-Phase Ions. Analytical Chemistry, 2018, 90, 11856-11862.	3.2	22
45	Conformationally Regulated Peptide Bond Cleavage in Bradykinin. Journal of the American Chemical Society, 2018, 140, 9357-9360.	6.6	25
46	Fourier Transform-Ion Mobility-Orbitrap Mass Spectrometer: A Next-Generation Instrument for Native Mass Spectrometry. Analytical Chemistry, 2018, 90, 10472-10478.	3.2	59
47	Ions from Solution to the Gas Phase: A Molecular Dynamics Simulation of the Structural Evolution of Substance P during Desolvation of Charged Nanodroplets Generated by Electrospray Ionization. Journal of the American Chemical Society, 2017, 139, 2981-2988.	6.6	49
48	Melting Proteins: Evidence for Multiple Stable Structures upon Thermal Denaturation of Native Ubiquitin from Ion Mobility Spectrometry-Mass Spectrometry Measurements. Journal of the American Chemical Society, 2017, 139, 6306-6309.	6.6	86
49	Characterizing the <i>Conformationome</i> : Toward a Structural Understanding of the Proteome. Accounts of Chemical Research, 2017, 50, 556-560.	7.6	53
50	Rapid capillary mixing experiments for the analysis of hydrophobic membrane complexes directly from aqueous lipid bilayer solutions. Analyst, The, 2017, 142, 310-315.	1.7	2
51	ESI-IM-MS and Collision-Induced Unfolding That Provide Insight into the Linkage-Dependent Interfacial Interactions of Covalently Linked Diubiquitin. Analytical Chemistry, 2017, 89, 10094-10103.	3.2	14
52	Investigation of the mechanism of the SpnF-catalyzed [4+2]-cycloaddition reaction in the biosynthesis of spinosyn A. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10408-10413.	3.3	38
53	Defining Noncovalent Ubiquitin Homodimer Interfacial Interactions through Comparisons with Covalently Linked Diubiquitin. Journal of the American Chemical Society, 2016, 138, 16588-16591.	6.6	13
54	"Wet―Versus "Dry―Folding of Polyproline. Journal of the American Society for Mass Spectrometry, 2016, 27, 1037-1047.	1.2	19

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55	Gas and Liquid Phase Diffusivities of Isomeric Metal Complexes Derived from Multifold Ring-Closing Metatheses: Ion Mobility Mass Spectrometry Trumps DOSY NMR. Organometallics, 2016, 35, 2071-2075.	1.1	15
56	Determining Membrane Protein–Lipid Binding Thermodynamics Using Native Mass Spectrometry. Journal of the American Chemical Society, 2016, 138, 4346-4349.	6.6	116
57	The Influence of Lipid Bilayer Physicochemical Properties on Gramicidin A Conformer Preferences. Biophysical Journal, 2016, 110, 1826-1835.	0.2	21
58	Long-Lived Intermediates in a Cooperative Two-State Folding Transition. Journal of Physical Chemistry B, 2016, 120, 12040-12046.	1.2	19
59	Following a Folding Transition with Capillary Electrophoresis and Ion Mobility Spectrometry. Analytical Chemistry, 2016, 88, 10933-10939.	3.2	7
60	Cryogenic Ion Mobility-Mass Spectrometry: Tracking Ion Structure from Solution to the Gas Phase. Accounts of Chemical Research, 2016, 49, 1421-1428.	7.6	37
61	Cis→Trans Isomerization of Pro ⁷ in Oxytocin Regulates Zn ²⁺ Binding. Journal of the American Society for Mass Spectrometry, 2016, 27, 1376-1382.	1.2	10
62	Increasing Ubiquitin Ion Resistance to Unfolding in the Gas Phase Using Chloride Adduction: Preserving More "Native-Like―Conformations Despite Collisional Activation. Analytical Chemistry, 2016, 88, 5934-5940.	3.2	24
63	Ion Mobility-Mass Spectrometry Reveals the Energetics of Intermediates that Guide Polyproline Folding. Journal of the American Society for Mass Spectrometry, 2016, 27, 22-30.	1.2	37
64	Cu2+-Phosphatidylserine Binding and its Implications for Protein-Membrane Interactions. Biophysical Journal, 2015, 108, 181a.	0.2	0
65	Focus on the 20-Year Anniversary of SEQUEST. Journal of the American Society for Mass Spectrometry, 2015, 26, 1797-1798.	1.2	6
66	Water-Mediated Dimerization of Ubiquitin Ions Captured by Cryogenic Ion Mobility-Mass Spectrometry. Journal of Physical Chemistry Letters, 2015, 6, 4947-4951.	2.1	25
67	How Closely Related Are Conformations of Protein Ions Sampled by IM-MS to Native Solution Structures?. Journal of the American Society for Mass Spectrometry, 2015, 26, 1433-1443.	1.2	93
68	Effects of charge states, charge sites and side chain interactions on conformational preferences of a series of model peptide ions. Analyst, The, 2015, 140, 6933-6944.	1.7	14
69	Unfolding of Hydrated Alkyl Diammonium Cations Revealed by Cryogenic Ion Mobility-Mass Spectrometry. Journal of the American Chemical Society, 2015, 137, 8916-8919.	6.6	18
70	Configurationally-Coupled Protonation of Polyproline-7. Journal of the American Chemical Society, 2015, 137, 8680-8683.	6.6	21
71	From Solution to Gas Phase: The Implications of Intramolecular Interactions on the Evaporative Dynamics of Substance P During Electrospray Ionization. Journal of Physical Chemistry B, 2015, 119, 4693-4698.	1.2	18
72	Probing the Electron Capture Dissociation Mass Spectrometry of Phosphopeptides with Traveling Wave Ion Mobility Spectrometry and Molecular Dynamics Simulations. Journal of the American Society for Mass Spectrometry, 2015, 26, 1004-1013.	1.2	14

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73	Reaction of Human Cd ₇ metallothionein and <i>N</i> -Ethylmaleimide: Kinetic and Structural Insights from Electrospray Ionization Mass Spectrometry. Biochemistry, 2015, 54, 6021-6028.	1.2	20
74	Elucidation of Conformer Preferences for a Hydrophobic Antimicrobial Peptide by Vesicle Capture-Freeze-Drying: A Preparatory Method Coupled to Ion Mobility-Mass Spectrometry. Analytical Chemistry, 2015, 87, 578-583.	3.2	14
75	Fluorescent Probes for Tracking the Transfer of Iron–Sulfur Cluster and Other Metal Cofactors in Biosynthetic Reaction Pathways. Journal of the American Chemical Society, 2015, 137, 390-398.	6.6	21
76	From Solution to the Gas Phase: Factors That Influence Kinetic Trapping of Substance P in the Gas Phase. Journal of Physical Chemistry B, 2014, 118, 14336-14344.	1.2	34
77	Cultivation of lipid-producing bacteria with lignocellulosic biomass: Effects of inhibitory compounds of lignocellulosic hydrolysates. Bioresource Technology, 2014, 161, 162-170.	4.8	50
78	Size-to-Charge Dispersion of Collision-Induced Dissociation Product Ions for Enhancement of Structural Information and Product Ion Identification. Analytical Chemistry, 2014, 86, 4791-4798.	3.2	5
79	Characterizing Intermediates Along the Transition from Polyproline I to Polyproline II Using Ion Mobility Spectrometry-Mass Spectrometry. Journal of the American Chemical Society, 2014, 136, 12702-12711.	6.6	91
80	Metal-Induced Conformational Changes of Human Metallothionein-2A: A Combined Theoretical and Experimental Study of Metal-Free and Partially Metalated Intermediates. Journal of the American Chemical Society, 2014, 136, 9499-9508.	6.6	67
81	Evolution of Hydrogen-Bond Networks in Protonated Water Clusters H ⁺ (H ₂ 0) _{<i>n</i>} (<i>n</i> = 1 to 120) Studied by Cryogenic Ion Mobility-Mass Spectrometry. Journal of Physical Chemistry Letters, 2014, 5, 1825-1830.	2.1	31
82	Mechanistic Consequences of Chiral Radical Clock Probes: Analysis of the Mononuclear Non-Heme Iron Enzyme HppE with 2-Hydroxy-3-methylenecyclopropyl Radical Clock Substrates. Journal of the American Chemical Society, 2014, 136, 2944-2947.	6.6	10
83	The Molecular Basis of Sugar Sensing in Drosophila Larvae. Current Biology, 2013, 23, 1466-1471.	1.8	78
84	Combining Chemical Labeling, Bottom-Up and Top-Down Ion-Mobility Mass Spectrometry To Identify Metal-Binding Sites of Partially Metalated Metallothionein. Analytical Chemistry, 2013, 85, 3229-3237.	3.2	43
85	The Periodic Focusing Ion Funnel: Theory, Design, and Experimental Characterization by High-Resolution Ion Mobility-Mass Spectrometry. Analytical Chemistry, 2013, 85, 9543-9548.	3.2	12
86	From Solution to the Gas Phase: Stepwise Dehydration and Kinetic Trapping of Substance P Reveals the Origin of Peptide Conformations. Journal of the American Chemical Society, 2013, 135, 19147-19153.	6.6	133
87	<i>Cis</i> – <i>Trans</i> Isomerizations of Proline Residues Are Key to Bradykinin Conformations. Journal of the American Chemical Society, 2013, 135, 3186-3192.	6.6	89
88	CAPA-gene products in the haematophagous sandfly Phlebotomus papatasi (Scopoli) – vector for leishmaniasis disease. Peptides, 2013, 41, 2-7.	1.2	5
89	Sol–Gel-Derived Silver-Nanoparticle-Embedded Thin Film for Mass Spectrometry-Based Biosensing. Langmuir, 2013, 29, 6502-6507.	1.6	35
90	Coupling Supported Lipid Bilayer Electrophoresis with Matrix-Assisted Laser Desorption/Ionization-Mass Spectrometry Imaging. Analytical Chemistry, 2013, 85, 6047-6052.	3.2	12

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91	Guest Packing Motifs within a Supramolecular Nanocapsule and a Covalent Analogue. Journal of the American Chemical Society, 2013, 135, 4314-4324.	6.6	86
92	Cryogenic Ion Mobility-Mass Spectrometry Captures Hydrated Ions Produced During Electrospray Ionization. Journal of Physical Chemistry A, 2013, 117, 953-961.	1.1	49
93	An Experimental Study of the Solvent-Dependent Self-Assembly/Disassembly and Conformer Preferences of Gramicidin A. Analytical Chemistry, 2013, 85, 7826-7833.	3.2	23
94	Posttranslational modification of CENP-A influences the conformation of centromeric chromatin. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 11827-11832.	3.3	114
95	Crystal Structure of Mycobacterium tuberculosis Polyketide Synthase 11 (PKS11) Reveals Intermediates in the Synthesis of Methyl-branched Alkylpyrones. Journal of Biological Chemistry, 2013, 288, 16484-16494.	1.6	21
96	Polycarbonates Derived from Glucose via an Organocatalytic Approach. Journal of the American Chemical Society, 2013, 135, 6826-6829.	6.6	117
97	A Facile Method to Synthesize Histones with Posttranslational Modification Mimics. Biochemistry, 2012, 51, 5232-5234.	1.2	40
98	Damping Factor Links Periodic Focusing and Uniform Field Ion Mobility for Accurate Determination of Collision Cross Sections. Analytical Chemistry, 2012, 84, 2818-2824.	3.2	19
99	How Alkali Metal Ion Binding Alters the Conformation Preferences of Gramicidin A: A Molecular Dynamics and Ion Mobility Study. Journal of Physical Chemistry A, 2012, 116, 689-696.	1.1	29
100	Neuropeptides of the cotton fleahopper, Pseudatomoscelis seriatus (Reuter). Peptides, 2012, 34, 39-43.	1.2	5
101	Biodegradation of triclosan by a wastewater microorganism. Water Research, 2012, 46, 4226-4234.	5.3	139
102	Imaging secondary metabolism of Streptomyces sp. Mg1 during cellular lysis and colony degradation of competing Bacillus subtilis. Antonie Van Leeuwenhoek, 2012, 102, 435-445.	0.7	50
103	The Caulobacter crescentus phage phiCbK: genomics of a canonical phage. BMC Genomics, 2012, 13, 542.	1.2	85
104	Interkingdom responses of flies to bacteria mediated by fly physiology and bacterial quorum sensing. Animal Behaviour, 2012, 84, 1449-1456.	0.8	83
105	Characterization of Structural Changes of Metallothionein by Ion Mobility-Mass Spectrometry (IM-MS): Metal-Free Vs. Metallated Forms. Biophysical Journal, 2012, 102, 57a.	0.2	0
106	Proteomic Methods for Biomarker Discovery in a Rat Model of Alcohol Steatosis. Methods in Molecular Biology, 2012, 909, 259-277.	0.4	3
107	Evidence for Radical-Mediated Catalysis by HppE: A Study Using Cyclopropyl and Methylenecyclopropyl Substrate Analogues. Journal of the American Chemical Society, 2012, 134, 16171-16174.	6.6	17
108	Mineralization of Acephate, a Recalcitrant Organophosphate Insecticide Is Initiated by a Pseudomonad in Environmental Samples. PLoS ONE, 2012, 7, e31963.	1.1	21

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109	Ion Mobility-Mass Spectrometry (IM-MS) for Top-Down Proteomics: Increased Dynamic Range Affords Increased Sequence Coverage. Analytical Chemistry, 2012, 84, 3390-3397.	3.2	61
110	Elucidation of chemical structures of pink-red pigments responsible for â€`pinking' in macerated onion (Allium cepa L.) using HPLC–DAD and tandem mass spectrometry. Food Chemistry, 2012, 131, 852-861.	4.2	9
111	The de novo engineering of pyrrolysyl-tRNA synthetase for genetic incorporation of l-phenylalanine and its derivatives. Molecular BioSystems, 2011, 7, 714.	2.9	76
112	Number of Solution States of Bradykinin from Ion Mobility and Mass Spectrometry Measurements. Journal of the American Chemical Society, 2011, 133, 13810-13813.	6.6	142
113	Efficient Electrophoretic Method to Remove Neutral Additives from Protein Solutions Followed by Mass Spectrometry Analysis. Analytical Chemistry, 2011, 83, 2814-2818.	3.2	4
114	Label-Free Biosensing with Lipid-Functionalized Gold Nanorods. Journal of the American Chemical Society, 2011, 133, 4182-4185.	6.6	72
115	Molecular Dynamics and Ion Mobility Spectrometry Study of Model β-Hairpin Peptide, Trpzip1. Journal of Physical Chemistry A, 2011, 115, 4427-4435.	1.1	23
116	Studies of Histidine As a Suitable Isoelectric Buffer for Tryptic Digestion and Isoelectric Trapping Fractionation Followed by Capillary Electrophoresis–Mass Spectrometry for Proteomic Analysis. Analytical Chemistry, 2011, 83, 8108-8114.	3.2	7
117	Proteomic Analysis of 3T3-L1 Adipocyte Mitochondria during Differentiation and Enlargement. Journal of Proteome Research, 2011, 10, 4692-4702.	1.8	48
118	Conformation and Self-Assembly of the Transmembrane Peptide Gramicidin A: Insights from ion Mobility Spectrometry and Molecular Dynamics. Biophysical Journal, 2011, 100, 387a.	0.2	0
119	Increased ion transmission in IMS: A high resolution, periodic-focusing DC ion guide ion mobility spectrometer. International Journal of Mass Spectrometry, 2011, 301, 166-173.	0.7	29
120	Effect of Cysteic Acid Position on the Negative Ion Fragmentation of Proteolytic Derived Peptides. Journal of the American Society for Mass Spectrometry, 2011, 22, 31-37.	1.2	2
121	A Mass-Selective Variable-Temperature Drift Tube Ion Mobility-Mass Spectrometer for Temperature Dependent Ion Mobility Studies. Journal of the American Society for Mass Spectrometry, 2011, 22, 1134-45.	1.2	67
122	Negative Ion Fragmentation of Cysteic Acid Containing Peptides: Cysteic Acid as a Fixed Negative Charge. Journal of the American Society for Mass Spectrometry, 2011, 22, 1622-1630.	1.2	8
123	Gas-phase ion dynamics in a periodic-focusing DC ion guide (Part II): Discrete transport modes. International Journal of Mass Spectrometry, 2011, 303, 154-163.	0.7	12
124	Saccharomyces cerevisiae THI4p is a suicide thiamine thiazole synthase. Nature, 2011, 478, 542-546.	13.7	149
125	A study of ion-neutral collision cross-section values for low charge states of peptides, proteins, and peptide/protein complexes. International Journal of Mass Spectrometry, 2010, 298, 111-118.	0.7	44
126	A Facile System for Genetic Incorporation of Two Different Noncanonical Amino Acids into One Protein in <i>Escherichia coli</i> . Angewandte Chemie - International Edition, 2010, 49, 3211-3214.	7.2	189

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127	Highâ€ŧhroughput method for onâ€ŧarget performic acid oxidation of MALDIâ€deposited samples. Journal of Mass Spectrometry, 2010, 45, 157-166.	0.7	7
128	Amino acid influence on copper binding to peptides: Cysteine versus arginine. Journal of the American Society for Mass Spectrometry, 2010, 21, 522-533.	1.2	43
129	Combining isoelectric point-based fractionation, liquid chromatography and mass spectrometry to improve peptide detection and protein identification. Journal of the American Society for Mass Spectrometry, 2010, 21, 1612-1619.	1.2	17
130	Synthesis, characterization and antibacterial activity of Felll, Coll, Cull and Znll complexes probed by transmission electron microscopy. Journal of Inorganic Biochemistry, 2010, 104, 1214-1223.	1.5	36
131	Gas-phase ion dynamics in a periodic-focusing DC ion guide. International Journal of Mass Spectrometry, 2010, 296, 36-42.	0.7	25
132	Genetic incorporation of an aliphatic keto-containing amino acid into proteins for their site-specific modifications. Bioorganic and Medicinal Chemistry Letters, 2010, 20, 878-880.	1.0	56
133	Neuropeptidomics of the Mosquito <i>Aedes aegypti</i> . Journal of Proteome Research, 2010, 9, 2006-2015.	1.8	141
134	Factors That Influence Helical Preferences for Singly Charged Gas-Phase Peptide Ions: The Effects of Multiple Potential Charge-Carrying Sites. Journal of Physical Chemistry B, 2010, 114, 809-816.	1.2	31
135	Longitudinal Surface Plasmon Resonance Based Gold Nanorod Biosensors for Mass Spectrometry. Langmuir, 2010, 26, 6066-6070.	1.6	53
136	Two capa-genes are expressed in the neuroendocrine system of Rhodnius prolixus. Peptides, 2010, 31, 408-411.	1.2	17
137	Investigation of the interaction of iron(iii) complexes with dAMP by ESI-MS, MALDI-MS and potentiometric titration: insights into synthetic nuclease behavior. Dalton Transactions, 2010, 39, 5094.	1.6	9
138	A genetically encoded photocaged Nε-methyl-l-lysine. Molecular BioSystems, 2010, 6, 1557.	2.9	72
139	A convenient method for genetic incorporation of multiple noncanonical amino acids into one protein in Escherichia coli. Molecular BioSystems, 2010, 6, 683.	2.9	56
140	A Cryogenic-Temperature Ion Mobility Mass Spectrometer for Improved Ion Mobility Resolution. , 2010, , 137-151.		1
141	Hepatic phenotype of liver fatty acid binding protein gene-ablated mice. American Journal of Physiology - Renal Physiology, 2009, 297, G1053-G1065.	1.6	59
142	A dual time-of-flight apparatus for an ion mobility-surface-induced dissociation-mass spectrometer for high-throughput peptide sequencing. International Journal of Mass Spectrometry, 2009, 287, 39-45.	0.7	10
143	A novel approach to collision-induced dissociation (CID) for ion mobility-mass spectrometry experiments. Journal of the American Society for Mass Spectrometry, 2009, 20, 907-914.	1.2	22
144	A new copper containing MALDI matrix that yields high abundances of [Peptide + Cu]+ ions. Journal of the American Society for Mass Spectrometry, 2009, 20, 1263-1271.	1.2	14

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145	The contributions of molecular framework to IMS collision cross-sections of gas-phase peptide ions. Journal of the American Society for Mass Spectrometry, 2009, 20, 1593-1602.	1.2	22
146	A mechanistic study of the H/D exchange reactions of protonated arginine and arginine-containing Di- and tripeptides. Journal of the American Society for Mass Spectrometry, 2009, 20, 2049-2057.	1.2	9
147	The neuropeptidomics of Ixodes scapularis synganglion. Journal of Proteomics, 2009, 72, 1040-1045.	1.2	43
148	Site-selective chemical protein glycosylation protects from autolysis and proteolytic degradation. Carbohydrate Research, 2009, 344, 1508-1514.	1.1	51
149	Anion Effects on Ionization Efficiency Using Cold Nanoparticles as Matrices for LDI-MS. Journal of Physical Chemistry C, 2009, 113, 1641-1647.	1.5	23
150	On the Structure Elucidation Using Ion Mobility Spectrometry and Molecular Dynamics. Journal of Physical Chemistry A, 2009, 113, 8221-8234.	1.1	50
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