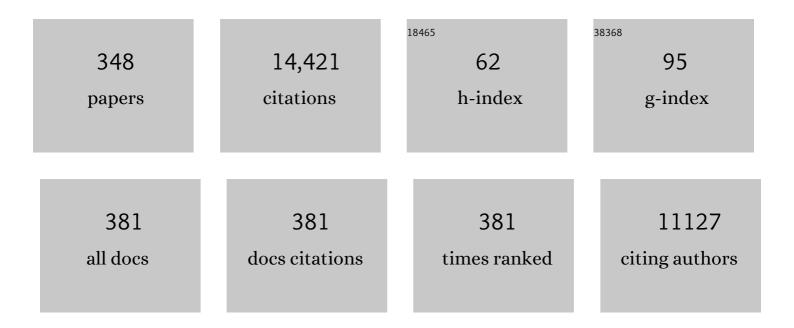
David H Russell

List of Publications by Year in descending order

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DAVID H RUSSEU

#	Article	IF	CITATIONS
1	Size-Selected (2â^10 nm) Gold Nanoparticles for Matrix Assisted Laser Desorption Ionization of Peptides. Journal of the American Chemical Society, 2005, 127, 5304-5305.	6.6	370
2	Anion Template Effect on the Self-Assembly and Interconversion of Metallacyclophanes. Journal of the American Chemical Society, 2005, 127, 12909-12923.	6.6	335
3	Ion mobility–mass spectrometry: a new paradigm for proteomics. International Journal of Mass Spectrometry, 2005, 240, 301-315.	0.7	282
4	Proteolysis in Mixed Organicâ^'Aqueous Solvent Systems:  Applications for Peptide Mass Mapping Using Mass Spectrometry. Analytical Chemistry, 2001, 73, 2682-2685.	3.2	266
5	A Universal Strategy for Proteomic Studies of SUMO and Other Ubiquitin-like Modifiers. Molecular and Cellular Proteomics, 2005, 4, 56-72.	2.5	195
6	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
7	Determination of the protein composition of the occlusion-derived virus of Autographa californica nucleopolyhedrovirus. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 9797-9802.	3.3	170
8	Peptidomics of CNS-associated neurohemal systems of adultDrosophila melanogaster: A mass spectrometric survey of peptides from individual flies. Journal of Comparative Neurology, 2004, 474, 379-392.	0.9	170
9	Fine-Tuning the Ring-Size of Metallacyclophanes:  A Rational Approach to Molecular Pentagons. Journal of the American Chemical Society, 2001, 123, 773-774.	6.6	164
10	Petroleum Crude Oil Characterization by IMS-MS and FTICR MS. Analytical Chemistry, 2009, 81, 9941-9947.	3.2	164
11	Tandem quadrupole Fourier-transform mass spectrometry of oligopeptides and small proteins Proceedings of the National Academy of Sciences of the United States of America, 1987, 84, 620-623.	3.3	157
12	Thermal Denaturation:Â A Useful Technique in Peptide Mass Mapping. Analytical Chemistry, 2000, 72, 2667-2670.	3.2	154
13	Coupling High-Pressure MALDI with Ion Mobility/Orthogonal Time-of-Flight Mass Spectrometry. Analytical Chemistry, 2000, 72, 3965-3971.	3.2	152
14	Saccharomyces cerevisiae THI4p is a suicide thiamine thiazole synthase. Nature, 2011, 478, 542-546.	13.7	149
15	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
16	Neuropeptidomics of the Mosquito <i>Aedes aegypti</i> . Journal of Proteome Research, 2010, 9, 2006-2015.	1.8	141
17	Biodegradation of triclosan by a wastewater microorganism. Water Research, 2012, 46, 4226-4234.	5.3	139
18	High-resolution Mass Spectrometry and Accurate Mass Measurements with Emphasis on the Characterization of Peptides and Proteins by Matrix-assisted Laser Desorption/Ionization Time-of-flight Mass Spectrometry. Journal of Mass Spectrometry, 1997, 32, 263-276.	0.7	135

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19	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
20	lsotopic labeling investigation of the oxygenation of nickel-bound thiolates by molecular oxygen. Journal of the American Chemical Society, 1992, 114, 4601-4605.	6.6	128
21	Solid Phase Syntheses of Oligoureas. Journal of the American Chemical Society, 1997, 119, 1556-1564.	6.6	128
22	Silver Nanoparticles as Selective Ionization Probes for Analysis of Olefins by Mass Spectrometry. Analytical Chemistry, 2008, 80, 6796-6799.	3.2	121
23	Polycarbonates Derived from Glucose via an Organocatalytic Approach. Journal of the American Chemical Society, 2013, 135, 6826-6829.	6.6	117
24	Determining Membrane Protein–Lipid Binding Thermodynamics Using Native Mass Spectrometry. Journal of the American Chemical Society, 2016, 138, 4346-4349.	6.6	116
25	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
26	Characterization of Poly(amidoamine) Dendrimers and Their Complexes with Cu2+by Matrix-Assisted Laser Desorption Ionization Mass Spectrometry. Macromolecules, 2001, 34, 3567-3573.	2.2	109
27	Fast-atom-bombardment-tandem mass spectrometry studies of alkali-metal ions of small peptides. Analytical Chemistry, 1986, 58, 1076-1080.	3.2	107
28	An evaluation of Fourier transform mass spectrometry for high mass applications. Mass Spectrometry Reviews, 1986, 5, 167-189.	2.8	103
29	Liquid sample introduction for matrix-assisted laser desorption ionization. Analytical Chemistry, 1993, 65, 2534-2537.	3.2	94
30	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
31	An electrostatic focusing ion guide for ion mobility-mass spectrometry. International Journal of Mass Spectrometry, 2004, 239, 43-49.	0.7	92
32	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
33	Evaluation of matrix-assisted laser desorption ionization-time-of-flight mass measurement accuracy by using delayed extraction. Journal of the American Society for Mass Spectrometry, 1996, 7, 995-1001.	1.2	89
34	<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
35	Fast-atom bombardment-tandem mass spectrometry studies of organo-alkali-metal ions of small peptides. Competitive interaction of sodium with basic amino acid substituents. Analytical Chemistry, 1988, 60, 1818-1824.	3.2	86
36	Reproducibility and quantitation of matrix-assisted laser desorption ionization mass spectrometry: Effects of nitrocellulose on peptide ion yields. Biological Mass Spectrometry, 1993, 22, 544-550.	0.5	86

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37	Distinguishing between Phosphorylated and Nonphosphorylated Peptides with Ion Mobilityâ^'Mass Spectrometry. Journal of Proteome Research, 2002, 1, 303-306.	1.8	86
38	Tailoring Nanoparticle Surface Chemistry to Enhance Laser Desorption Ionization of Peptides and Proteins. Nano Letters, 2007, 7, 3023-3025.	4.5	86
39	Guest Packing Motifs within a Supramolecular Nanocapsule and a Covalent Analogue. Journal of the American Chemical Society, 2013, 135, 4314-4324.	6.6	86
40	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
41	The Caulobacter crescentus phage phiCbK: genomics of a canonical phage. BMC Genomics, 2012, 13, 542.	1.2	85
42	Surface-Induced Dissociation on a MALDI-Ion Mobility-Orthogonal Time-of-Flight Mass Spectrometer:Â Sequencing Peptides from an "In-Solution―Protein Digest. Analytical Chemistry, 2001, 73, 2233-2238.	3.2	83
43	Trafficking of ODV-E66 is mediated via a sorting motif and other viral proteins: Facilitated trafficking to the inner nuclear membrane. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 8372-8377.	3.3	83
44	Peak capacity of ion mobility mass spectrometry: the utility of varying drift gas polarizability for the separation of tryptic peptides. Journal of Mass Spectrometry, 2004, 39, 361-367.	0.7	83
45	Interkingdom responses of flies to bacteria mediated by fly physiology and bacterial quorum sensing. Animal Behaviour, 2012, 84, 1449-1456.	0.8	83
46	Peak capacity of ion mobility mass spectrometry:. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 782, 385-392.	1.2	82
47	Cesium ion desorption ionization with Fourier transform mass spectrometry (FTMS). Analytical Chemistry, 1984, 56, 578-581.	3.2	80
48	The Molecular Basis of Sugar Sensing in Drosophila Larvae. Current Biology, 2013, 23, 1466-1471.	1.8	78
49	A collision cross-section database of singly-charged peptide ions. Journal of the American Society for Mass Spectrometry, 2007, 18, 1232-1238.	1.2	77
50	Are proton transfer reactions of excited states involved in UV laser desorption ionization?. Organic Mass Spectrometry, 1992, 27, 827-830.	1.3	76
51	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
52	Mass spectrometry. Analytical Chemistry, 1992, 64, 467-502.	3.2	74
53	Analysis of Phosphorylated Peptides by Ion Mobility-Mass Spectrometry. Analytical Chemistry, 2004, 76, 6727-6733.	3.2	72
54	A genetically encoded photocaged Nε-methyl-l-lysine. Molecular BioSystems, 2010, 6, 1557.	2.9	72

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55	Label-Free Biosensing with Lipid-Functionalized Gold Nanorods. Journal of the American Chemical Society, 2011, 133, 4182-4185.	6.6	72
56	Oligonucleotide analysis with MALDI–ion-mobility–TOFMS. Analytical and Bioanalytical Chemistry, 2002, 373, 612-617.	1.9	70
57	Field-corrected ion cell for ion cyclotron resonance. Analytical Chemistry, 1990, 62, 520-526.	3.2	67
58	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
59	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
60	Aerosol Matrix-Assisted Laser Desorption Ionization Mass Spectrometry. Journal of the American Society for Mass Spectrometry, 1994, 5, 1-9.	1.2	65
61	Identification of Individual Proteins in Complex Protein Mixtures by High-Resolution, High-Mass-Accuracy MALDI TOF-Mass Spectrometry Analysis of In-Solution Thermal Denaturation/Enzymatic Digestion. Analytical Chemistry, 2001, 73, 2558-2564.	3.2	65
62	Mass spectrometry. Analytical Chemistry, 1988, 60, 294-342.	3.2	63
63	Sexually dimorphic metabolism of branched-chain lipids in C57BL/6J mice. Journal of Lipid Research, 2004, 45, 812-830.	2.0	63
64	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
65	Ratiometric Pulsed Alkylation/Mass Spectrometry of the Cysteine Pairs in Individual Zinc Fingers of MRE-Binding Transcription Factor-1 (MTF-1) as a Probe of Zinc Chelate Stability. Biochemistry, 2001, 40, 15164-15175.	1.2	60
66	Observation of Conserved Solution-Phase Secondary Structure in Gas-Phase Tryptic Peptides. Journal of the American Chemical Society, 2002, 124, 4214-4215.	6.6	60
67	The structure of an insect chymotrypsin. Journal of Molecular Biology, 2000, 298, 895-901.	2.0	59
68	Ion mobility-mass spectrometry applied to cyclic peptide analysis: Conformational preferences of gramicidin S and linear analogs in the gas phase. Journal of the American Society for Mass Spectrometry, 2004, 15, 870-878.	1.2	59
69	Biochemical and Functional Analyses of the Human Toll-like Receptor 3 Ectodomain. Journal of Biological Chemistry, 2007, 282, 7668-7678.	1.6	59
70	Structure and Function of the Virulence-Associated High-Temperature Requirement A of <i>Mycobacterium tuberculosis</i> . Biochemistry, 2008, 47, 6092-6102.	1.2	59
71	Molecular Weight Distributions of Asphaltenes and Deasphaltened Oils Studied by Laser Desorption Ionization and Ion Mobility Mass Spectrometry. Analytical Chemistry, 2008, 80, 8592-8597.	3.2	59
72	Hepatic phenotype of liver fatty acid binding protein gene-ablated mice. American Journal of Physiology - Renal Physiology, 2009, 297, G1053-G1065.	1.6	59

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73	Fourier Transform-Ion Mobility-Orbitrap Mass Spectrometer: A Next-Generation Instrument for Native Mass Spectrometry. Analytical Chemistry, 2018, 90, 10472-10478.	3.2	59
74	Target-gas excitation accompanying collisional activation of large polyatomic ions. Journal of the American Chemical Society, 1986, 108, 6174-6179.	6.6	58
75	Template Effect for O2Addition acrosscis-Sulfur Sites in Nickel Dithiolates. Journal of the American Chemical Society, 1996, 118, 1791-1792.	6.6	57
76	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
77	Identification of the Naturally Occurring Flavin of Nitroalkane Oxidase from Fusarium oxysporum as a 5-Nitrobutyl-FAD and Conversion of the Enzyme to the Active FAD-containing Form. Journal of Biological Chemistry, 1997, 272, 5563-5570.	1.6	56
78	Pro-sterol Carrier Protein-2. Journal of Biological Chemistry, 2000, 275, 25547-25555.	1.6	56
79	Liver Proteome Analysis in a Rodent Model of Alcoholic Steatosis. Journal of Proteome Research, 2009, 8, 1663-1671.	1.8	56
80	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
81	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
82	The structure of gas-phase bradykinin fragment 1–5 (RPPGF) ions: An ion mobility spectrometry and H/D exchange ion-molecule reaction chemistry study. Journal of the American Society for Mass Spectrometry, 2005, 16, 893-905.	1.2	53
83	Longitudinal Surface Plasmon Resonance Based Gold Nanorod Biosensors for Mass Spectrometry. Langmuir, 2010, 26, 6066-6070.	1.6	53
84	Characterizing the <i>Conformationome</i> : Toward a Structural Understanding of the Proteome. Accounts of Chemical Research, 2017, 50, 556-560.	7.6	53
85	Detection of femtomole and sub-femtomole levels of peptides by tandem magnetic sector/reflectron time-of-flight mass spectrometry and matrix-assisted laser desorption ionization. Journal of the American Society for Mass Spectrometry, 1991, 2, 91-94.	1.2	52
86	A method for removal of N-BOC protecting groups from substrates on TFA-sensitive resins. Tetrahedron Letters, 1998, 39, 7439-7442.	0.7	52
87	Fragmentation chemistry of [M + Cu]+ peptide ions containing an N-terminal arginine. Journal of the American Society for Mass Spectrometry, 2000, 11, 626-638.	1.2	51
88	Development of a Fourier-transform ion cyclotron resonance mass spectrometer-ion mobility spectrometer. Review of Scientific Instruments, 2000, 71, 4078.	0.6	51
89	Site-selective chemical protein glycosylation protects from autolysis and proteolytic degradation. Carbohydrate Research, 2009, 344, 1508-1514.	1.1	51
90	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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91	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
92	Cultivation of lipid-producing bacteria with lignocellulosic biomass: Effects of inhibitory compounds of lignocellulosic hydrolysates. Bioresource Technology, 2014, 161, 162-170.	4.8	50
93	Sequence and side-chain specific photofragment (193 nm) ions from protonated substance P by matrix-assisted laser desorption ionization time-of-flight mass spectrometry. Journal of the American Society for Mass Spectrometry, 1999, 10, 1038-1040.	1.2	49
94	Cryogenic Ion Mobility-Mass Spectrometry Captures Hydrated Ions Produced During Electrospray Ionization. Journal of Physical Chemistry A, 2013, 117, 953-961.	1.1	49
95	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
96	Matrix-assisted laser desorption ionization hydrogen/deuterium exchange studies to probe peptide conformational changes. Journal of the American Society for Mass Spectrometry, 1999, 10, 719-731.	1.2	48
97	Proteomic Analysis of 3T3-L1 Adipocyte Mitochondria during Differentiation and Enlargement. Journal of Proteome Research, 2011, 10, 4692-4702.	1.8	48
98	A fundamental introduction to ion mobility mass spectrometry applied to the analysis of biomolecules. Journal of Biomolecular Techniques, 2002, 13, 56-61.	0.8	48
99	Profile and Flight Time Analysis of Bovine Insulin Clusters as a Probe of Matrix-assisted Laser Desorption/Ionization Ion Formation Dynamics. , 1997, 32, 714-722.		47
100	Detection of High-Mass Biomolecules in Fourier Transform Ion Cyclotron Resonance Mass Spectrometry: Theoretical and Experimental Investigations. Analytical Chemistry, 1994, 66, 1583-1587.	3.2	46
101	Membrane Charge and Curvature Determine Interaction with Acyl-CoA Binding Protein (ACBP) and Fatty Acyl-CoA Targeting. Biochemistry, 2002, 41, 10540-10553.	1.2	45
102	Detection of mass 16241 ions by Fourier-transform mass spectrometry. Analytical Chemistry, 1986, 58, 483-485.	3.2	44
103	Mass spectrometry. Analytical Chemistry, 1990, 62, 268-303.	3.2	44
104	Comparative studies of 193-nm photodissociation and TOF-TOFMS analysis of bradykinin analogues: The effects of charge site(s) and fragmentation timescales. Journal of the American Society for Mass Spectrometry, 2006, 17, 721-729.	1.2	44
105	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
106	Aerosol Matrix-Assisted Laser Desorption Ionization for Liquid Chromatography/Time-of-Flight Mass Spectrometry. Analytical Chemistry, 1994, 66, 1601-1609.	3.2	43
107	Isolation and characterization of two distinct forms of liver fatty acid binding protein from the rat. Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids, 1999, 1436, 413-425.	1.2	43
108	Analysis of protein mixtures by matrix-assisted laser desorption ionization-ion mobility-orthogonal-time-of-flight mass spectrometry. International Journal of Mass Spectrometry, 2002, 219, 253-267.	0.7	43

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109	Comparative peptidomics of four related hemipteran species: Pyrokinins, myosuppressin, corazonin, adipokinetic hormone, sNPF, and periviscerokinins. Peptides, 2008, 29, 162-167.	1.2	43
110	The neuropeptidomics of Ixodes scapularis synganglion. Journal of Proteomics, 2009, 72, 1040-1045.	1.2	43
111	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
112	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
113	Divergent pathways for the addition of dioxygen to sulfur in nickel cis-dithiolates: an isotopomeric analysis. Inorganic Chemistry, 1993, 32, 4171-4172.	1.9	42
114	Structure and Function of Normal and Transformed Murine Acyl-CoA Binding Proteins. Archives of Biochemistry and Biophysics, 1998, 350, 201-213.	1.4	42
115	Improvement of Resolution, Mass Accuracy, and Reproducibility in Reflected Mode DE-MALDI-TOF Analysis of DNA Using Fast Evaporationâ^'Overlayer Sample Preparations. Analytical Chemistry, 2000, 72, 3860-3866.	3.2	42
116	Determination of copper binding sites in peptides containing basic residues: a combined experimental and theoretical study. International Journal of Mass Spectrometry, 2001, 204, 31-46.	0.7	42
117	Accurate mass measurement of DNA oligonucleotide ions using high-resolution time-of-flight mass spectrometry. Journal of Mass Spectrometry, 2002, 37, 357-371.	0.7	41
118	A Facile Method to Synthesize Histones with Posttranslational Modification Mimics. Biochemistry, 2012, 51, 5232-5234.	1.2	40
119	Keto-enol tautomerism of gas-phase ions. Structure of reactive 1,3-cyclohexadien-5-one radical cations. Journal of the American Chemical Society, 1978, 100, 6133-6137.	6.6	39
120	Formation of ionic transition-metal carbonyl cluster fragments by ion-molecule reactions. 1. The chromium hexacarbonyl and iron pentacarbonyl systems. Journal of the American Chemical Society, 1985, 107, 3762-3768.	6.6	39
121	Laser desorption studies of high mass biomolecules in Fourier-transform ion cyclotron resonance mass spectrometry Proceedings of the National Academy of Sciences of the United States of America, 1992, 89, 5701-5704.	3.3	39
122	Binding of DNA Purine Sites to Dirhodium Compounds Probed by Mass Spectrometry. Inorganic Chemistry, 2004, 43, 6177-6187.	1.9	39
123	Laser ion beam photodissociation studies of model amino acids and peptides. Journal of the American Chemical Society, 1989, 111, 1161-1171.	6.6	38
124	Effects of Matrix Structure/Acidity on Ion Formation in Matrix-Assisted Laser Desorption Ionization Mass Spectrometry. Journal of the American Chemical Society, 1997, 119, 2534-2540.	6.6	38
125	Improving mass spectrometric sequencing of arginine-containing peptides by derivatization with acetylacetone. , 1997, 32, 1337-1349.		38
126	A High Repetition Rate (1 kHz) Microcrystal Laser for High Throughput Atmospheric Pressure MALDI-Quadrupole-Time-of-Flight Mass Spectrometry. Analytical Chemistry, 2003, 75, 648-654.	3.2	38

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127	Structure and Function of the Sterol Carrier Protein-2 N-Terminal Presequence. Biochemistry, 2008, 47, 5915-5934.	1.2	38
128	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
129	Cesium desorption ionization studies of .betacyclodextrin by Fourier transform mass spectrometry. Journal of the American Chemical Society, 1985, 107, 5652-5657.	6.6	37
130	Investigation of the dynamics of matrix-assisted laser desorption/ionization ion formation using an electrostatic analyzer/time-of-flight mass spectrometer. Journal of Mass Spectrometry, 1999, 34, 684-690.	0.7	37
131	Resolution equations for high-field ion mobility. Journal of the American Society for Mass Spectrometry, 2004, 15, 1320-1324.	1.2	37
132	Functional Analysis of RNA Binding by the Hepatitis C Virus RNA-dependent RNA Polymerase. Journal of Biological Chemistry, 2005, 280, 38011-38019.	1.6	37
133	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
134	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
135	Mass and energy selective ion partitioning in a two-section Fourier-transform ion cyclotron resonance spectrometer cell. Analytical Chemistry, 1989, 61, 53-57.	3.2	36
136	Synthesis, characterization and antibacterial activity of FeIII, CoII, CuII and ZnII complexes probed by transmission electron microscopy. Journal of Inorganic Biochemistry, 2010, 104, 1214-1223.	1.5	36
137	Pulsed-Alkylation Mass Spectrometry for the Study of Protein Folding and Dynamics:  Development and Application to the Study of a Folding/Unfolding Intermediate of Bacterial Luciferase. Biochemistry, 2001, 40, 15153-15163.	1.2	35
138	Optimization of a matrix-assisted laser desorption ionization-ion mobility-surface-induced dissociation-orthogonal-time-of-flight mass spectrometer: simultaneous acquisition of multiple correlated MS 1 and MS 2 spectra. International Journal of Mass Spectrometry, 2001, 212, 519-533.	0.7	35
139	The influence and utility of varying field strength for the separation of tryptic peptides by ion mobility-mass spectrometry. Journal of the American Society for Mass Spectrometry, 2005, 16, 158-165.	1.2	35
140	Mass spectrometric assignment of Leu/Ile in neuropeptides from single neurohemal organ preparations of insects. Peptides, 2005, 26, 2151-2156.	1.2	35
141	Sol–Gel-Derived Silver-Nanoparticle-Embedded Thin Film for Mass Spectrometry-Based Biosensing. Langmuir, 2013, 29, 6502-6507.	1.6	35
142	Ion-molecule reaction chemistry of various gas-phase C6H6 radical cations. Journal of the American Chemical Society, 1977, 99, 3603-3609.	6.6	34
143	Electrocyclic ring opening of the cyclobutene radical cation. Journal of the American Chemical Society, 1979, 101, 2082-2086.	6.6	34
144	Effects of the Water Content in the Sample Preparation for MALDI on the Mass Spectra. Analytical Chemistry, 1998, 70, 4527-4533.	3.2	34

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145	Ratiometric Pulsed Alkylation Mass Spectrometry as a Probe of Thiolate Reactivity in Different Metalloderivatives ofStaphylococcus aureuspI258 CadCâ€,‡. Biochemistry, 2004, 43, 3824-3834.	1.2	34
146	Gas-Phase Conformations of Proteolytically Derived Protein Fragments:  Influence of Solvent on Peptide Conformation. Journal of Physical Chemistry B, 2004, 108, 15321-15331.	1.2	34
147	Isolation and characterization of two disintegrins inhibiting ADP-induced human platelet aggregation from the venom of Crotalus scutulatus scutulatus (Mohave Rattlesnake). Toxicology and Applied Pharmacology, 2006, 212, 59-68.	1.3	34
148	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
149	Fragmentation of vitamin B12 during 337 nm matrix-assisted laser desorption ionization. Biological Mass Spectrometry, 1994, 23, 205-211.	0.5	33
150	Sub-Femtomole Peptide Detection in Ion Mobility-Time-of-Flight Mass Spectrometry Measurements. Journal of Proteome Research, 2003, 2, 427-430.	1.8	33
151	Native IM-Orbitrap MS: Resolving what was hidden. TrAC - Trends in Analytical Chemistry, 2020, 124, 115533.	5.8	33
152	Variable-Temperature Electrospray Ionization for Temperature-Dependent Folding/Refolding Reactions of Proteins and Ligand Binding. Analytical Chemistry, 2021, 93, 6924-6931.	3.2	33
153	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
154	Evolution of Hydrogen-Bond Networks in Protonated Water Clusters H ⁺ (H ₂ O) _{<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
155	An evaluation of the analytical utility of the photodissociation of fast ion beams. Mass Spectrometry Reviews, 1990, 9, 405-451.	2.8	30
156	Proteome analysis of Escherichia coli K-12 by two-dimensional native-state chromatography and MALDI-MS. Molecular Microbiology, 2003, 47, 383-396.	1.2	30
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