

# David C Muddiman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8242200/publications.pdf>

Version: 2024-02-01

327  
papers

13,315  
citations

27035

58  
h-index

49824

91  
g-index

340  
all docs

340  
docs citations

340  
times ranked

12799  
citing authors

#	ARTICLE	IF	CITATIONS
1	The development and application of matrix assisted laser desorption electrospray ionization: The teenage years. <i>Mass Spectrometry Reviews</i> , 2023, 42, 35-66.	2.8	40
2	Understanding the electrospray ionization response factors of per- and poly-fluoroalkyl substances (PFAS). <i>Analytical and Bioanalytical Chemistry</i> , 2022, 414, 1227-1234.	1.9	8
3	Utilizing liquid chromatography, ion mobility spectrometry, and mass spectrometry to assess INLIGHT <sup>®</sup> -labeled derivatized N-linked glycans in biological samples. <i>Analytical and Bioanalytical Chemistry</i> , 2022, 414, 623-637.	1.9	6
4	Simultaneous Measurement of Striatal Dopamine and Hydrogen Peroxide Transients Associated with L-DOPA Induced Rotation in Hemiparkinsonian Rats. <i>ACS Measurement Science Au</i> , 2022, 2, 120-131.	1.9	2
5	Phosphorylation-dependent proteome of Marcks in ependyma during aging and behavioral homeostasis in the mouse forebrain. <i>GeroScience</i> , 2022, 44, 2077-2094.	2.1	1
6	Optimized C-Trap Timing of an Orbitrap 240 Mass Spectrometer for High-Throughput Screening and Native MS by IR-MALDESI. <i>Journal of the American Society for Mass Spectrometry</i> , 2022, 33, 328-334.	1.2	16
7	Development and validation of a high resolving power absolute quantitative per- and polyfluoroalkyl substances method incorporating Skyline data processing. <i>Rapid Communications in Mass Spectrometry</i> , 2022, 36, e9295.	0.7	4
8	Mass Spectrometry Imaging of N-Linked Glycans in a Formalin-Fixed Paraffin-Embedded Human Prostate by Infrared Matrix-Assisted Laser Desorption Electrospray Ionization. <i>Journal of Proteome Research</i> , 2022, 21, 243-249.	1.8	13
9	Sequential paired covariance for improved visualization of mass spectrometry imaging datasets. <i>Journal of Mass Spectrometry</i> , 2022, 57, .	0.7	3
10	Highlighting Functional Mass Spectrometry Imaging Methods in Bioanalysis. <i>Journal of Proteome Research</i> , 2022, 21, 1800-1807.	1.8	9
11	Glycerate from intestinal fructose metabolism induces islet cell damage and glucose intolerance. <i>Cell Metabolism</i> , 2022, 34, 1042-1053.e6.	7.2	10
12	An adaptive teosinte <i>mexicana</i> introgression modulates phosphatidylcholine levels and is associated with maize flowering time. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	21
13	GlycoHunter: An Open-Source Software for the Detection and Relative Quantification of INLIGHT-Labeled N-Linked Glycans. <i>Journal of Proteome Research</i> , 2021, 20, 1855-1863.	1.8	5
14	Three-dimensional (3D) imaging of lipids in skin tissues with infrared matrix-assisted laser desorption electrospray ionization (MALDESI) mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 2793-2801.	1.9	20
15	In situ detection of fatty acid C=C positional isomers by coupling on-tissue mCPBA epoxidation with infrared matrix-assisted laser desorption electrospray ionization mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2021, 35, e9119.	0.7	9
16	Investigations of $\beta$ -carotene radical cation formation in infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Rapid Communications in Mass Spectrometry</i> , 2021, 35, e9133.	0.7	2
17	Spatially resolved metabolomic characterization of muscle invasive bladder cancer by mass spectrometry imaging. <i>Metabolomics</i> , 2021, 17, 70.	1.4	12
18	Enzyme Complexes of Ptr4CL and PtrHCT Modulate Co-enzyme A Ligation of Hydroxycinnamic Acids for Monolignol Biosynthesis in <i>Populus trichocarpa</i> . <i>Frontiers in Plant Science</i> , 2021, 12, 727932.	1.7	5

#	ARTICLE	IF	CITATIONS
19	Multiple Infusion Start Time Mass Spectrometry Imaging of Dynamic SIL-Glutathione Biosynthesis Using Infrared Matrix-Assisted Laser Desorption Electrospray Ionization. <i>Journal of Proteome Research</i> , 2021, , .	1.8	8
20	Enhancing Metabolomic Coverage in Positive Ionization Mode Using Dicationic Reagents by Infrared Matrix-Assisted Laser Desorption Electrospray Ionization. <i>Metabolites</i> , 2021, 11, 810.	1.3	1
21	Multimodal Mass Spectrometry Imaging of Rat Brain Using IR-MALDESI and NanoPOTS-LC-MS/MS. <i>Journal of Proteome Research</i> , 2021, , .	1.8	8
22	NIST Interlaboratory Study on Glycosylation Analysis of Monoclonal Antibodies: Comparison of Results from Diverse Analytical Methods. <i>Molecular and Cellular Proteomics</i> , 2020, 19, 11-30.	2.5	87
23	Development of a relative quantification method for infrared matrix-assisted laser desorption electrospray ionization mass spectrometry imaging of Arabidopsis seedlings. <i>Rapid Communications in Mass Spectrometry</i> , 2020, 34, e8616.	0.7	12
24	Three-Dimensional Imaging with Infrared Matrix-Assisted Laser Desorption Electrospray Ionization Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 292-297.	1.2	14
25	Determination of Optimal Electrospray Parameters for Lipidomics in Infrared-Matrix-Assisted Laser Desorption Electrospray Ionization Mass Spectrometry Imaging. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 319-325.	1.2	14
26	Investigating host-pathogen meta-metabolic interactions of <i>Magnaporthe oryzae</i> infected barley using infrared matrix-assisted laser desorption electrospray ionization mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 139-147.	1.9	5
27	Lipidomic profiling of single mammalian cells by infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 8211-8222.	1.9	20
28	3D Imaging and metabolomic profiling reveal higher neuroactive kavalactone contents in lateral roots and crown root peels of <i>Piper methysticum</i> (kava). <i>GigaScience</i> , 2020, 9, .	3.3	9
29	Comparative Proteomic Analysis of Wild Type and Mutant Lacking an SCF E3 Ligase F-Box Protein in <i>Magnaporthe oryzae</i> . <i>Journal of Proteome Research</i> , 2020, 19, 3761-3768.	1.8	1
30	Infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI) mass spectrometry imaging analysis of endogenous metabolites in cherry tomatoes. <i>Analyst, The</i> , 2020, 145, 5516-5523.	1.7	18
31	Enhanced protocol for quantitative N-linked glycomics analysis using Individuality Normalization when Labeling with Isotopic Glycan Hydrazide Tags (INLIGHT)â„¢. <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 7569-7579.	1.9	11
32	Metabolite Profiling Reveals Predictive Biomarkers and the Absence of $\hat{1}^2$ -Methyl Amino- <i>l</i> -alanine in Plasma from Individuals Diagnosed with Amyotrophic Lateral Sclerosis. <i>Journal of Proteome Research</i> , 2020, 19, 3276-3285.	1.8	18
33	A Versatile Platform for Mass Spectrometry Imaging of Arbitrary Spatial Patterns. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 2547-2552.	1.2	19
34	Direct Analysis of Native <i>N</i> -Linked Glycans by IR-MALDESI. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 1759-1762.	1.2	12
35	Methods for Cryosectioning and Mass Spectrometry Imaging of Whole-Body Zebrafish. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 768-772.	1.2	19
36	Coupling IR-MALDESI with Drift Tube Ion Mobility-Mass Spectrometry for High-Throughput Screening and Imaging Applications. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 642-650.	1.2	22

#	ARTICLE	IF	CITATIONS
37	Effects of Prenatal Exposure to a Mixture of Organophosphate Flame Retardants on Placental Gene Expression and Serotonergic Innervation in the Fetal Rat Brain. <i>Toxicological Sciences</i> , 2020, 176, 203-223.	1.4	37
38	Analysis of neurotransmitters in rat placenta exposed to flame retardants using IR-MALDESI mass spectrometry imaging. <i>Analytical and Bioanalytical Chemistry</i> , 2020, 412, 3745-3752.	1.9	12
39	Discovery and quantification of bioactive peptides in fermented cucumber by direct analysis IR-MALDESI mass spectrometry and LC-QQQ-MS. <i>Food Chemistry</i> , 2019, 271, 715-723.	4.2	43
40	Heterogeneous antiretroviral drug distribution and HIV/SHIV detection in the gut of three species. <i>Science Translational Medicine</i> , 2019, 11, .	5.8	38
41	Quantitative proteomic analysis of tomato genotypes with differential cadmium tolerance. <i>Environmental Science and Pollution Research</i> , 2019, 26, 26039-26051.	2.7	17
42	Internal Energy Deposition in Infrared Matrix-Assisted Laser Desorption Electrospray Ionization With and Without the Use of Ice as a Matrix. <i>Journal of the American Society for Mass Spectrometry</i> , 2019, 30, 2380-2391.	1.2	27
43	Systematic evaluation of repeatability of IR-MALDESI-MS and normalization strategies for correcting the analytical variation and improving image quality. <i>Analytical and Bioanalytical Chemistry</i> , 2019, 411, 5729-5743.	1.9	18
44	Label-Free Quantitative Proteomics of Enriched Nuclei from Sugarcane ( <i>Saccharum</i> ssp) Stems in Response to Drought Stress. <i>Proteomics</i> , 2019, 19, e1900004.	1.3	21
45	Artemisinin Biosynthesis in Non-glandular Trichome Cells of <i>Artemisia annua</i> . <i>Molecular Plant</i> , 2019, 12, 704-714.	3.9	62
46	Mass spectrometry imaging (MSI) of fresh bones using infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Analytical Methods</i> , 2019, 11, 5929-5938.	1.3	15
47	Discriminating normal regions within cancerous hen ovarian tissue using multivariate hyperspectral image analysis. <i>Rapid Communications in Mass Spectrometry</i> , 2019, 33, 381-391.	0.7	4
48	CAD1 and CCR2 protein complex formation in monolignol biosynthesis in <i>Populus trichocarpa</i> . <i>New Phytologist</i> , 2019, 222, 244-260.	3.5	43
49	A novel integrated strategy for the detection and quantification of the neurotoxin $\beta^2$ -N-methylamino-l-alanine in environmental samples. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 2597-2605.	1.9	12
50	Improving wood properties for wood utilization through multi-omics integration in lignin biosynthesis. <i>Nature Communications</i> , 2018, 9, 1579.	5.8	162
51	Characterization of a novel miniaturized burst-mode infrared laser system for IR-MALDESI mass spectrometry imaging. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 2395-2402.	1.9	25
52	Characterization of the Spectral Accuracy of an Orbitrap Mass Analyzer Using Isotope Ratio Mass Spectrometry. <i>Analytical Chemistry</i> , 2018, 90, 1897-1906.	3.2	30
53	Demonstration of hydrazide tagging for O-glycans and a central composite design of experiments optimization using the INLIGHT reagent. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 1409-1415.	1.9	4
54	Quantitative mass spectrometry imaging of glutathione in healthy and cancerous hen ovarian tissue sections by infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Analyst</i> , 2018, 143, 654-661.	1.7	36

#	ARTICLE	IF	CITATIONS
55	MSiReader v1.0: Evolving Open-Source Mass Spectrometry Imaging Software for Targeted and Untargeted Analyses. <i>Journal of the American Society for Mass Spectrometry</i> , 2018, 29, 8-16.	1.2	193
56	IR-MALDESI method optimization based on time-resolved measurement of ion yields. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 963-970.	1.9	12
57	Direct analysis of terpenes from biological buffer systems using SESI and IR-MALDESI. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 953-962.	1.9	8
58	IR-MALDESI mass spectrometry imaging of underivatized neurotransmitters in brain tissue of rats exposed to tetrabromobisphenol A. <i>Analytical and Bioanalytical Chemistry</i> , 2018, 410, 7979-7986.	1.9	19
59	Evaluation of Digital Image Recognition Methods for Mass Spectrometry Imaging Data Analysis. <i>Journal of the American Society for Mass Spectrometry</i> , 2018, 29, 2467-2470.	1.2	18
60	Xylose Migration During Tandem Mass Spectrometry of <i>N</i> -Linked Glycans. <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 729-732.	1.2	19
61	The PeptideAtlas of the Domestic Laying Hen. <i>Journal of Proteome Research</i> , 2017, 16, 1352-1363.	1.8	9
62	Recent advances in glycomics, glycoproteomics and allied topics. <i>Analytical and Bioanalytical Chemistry</i> , 2017, 409, 355-357.	1.9	22
63	Direct screening of enzyme activity using infrared matrix-assisted laser desorption electrospray ionization. <i>Rapid Communications in Mass Spectrometry</i> , 2017, 31, 1868-1874.	0.7	21
64	IR-MALDESI Mass Spectrometry Imaging at 50 Micron Spatial Resolution. <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 2099-2107.	1.2	28
65	Perfluorinated alcohol induced coacervates as extraction media for proteomic analysis. <i>Journal of Chromatography A</i> , 2017, 1523, 293-299.	1.8	12
66	DRILL: An Electrospray Ionization-Mass Spectrometry Interface for Improved Sensitivity via Inertial Droplet Sorting and Electrohydrodynamic Focusing in a Swirling Flow. <i>Analytical Chemistry</i> , 2017, 89, 8981-8987.	3.2	18
67	Direct Analysis of Triterpenes from High-Salt Fermented Cucumbers Using Infrared Matrix-Assisted Laser Desorption Electrospray Ionization (IR-MALDESI). <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 370-375.	1.2	26
68	N-linked glycosite profiling and use of Skyline as a platform for characterization and relative quantification of glycans in differentiating xylem of <i>Populus trichocarpa</i> . <i>Analytical and Bioanalytical Chemistry</i> , 2017, 409, 487-497.	1.9	23
69	Mass spectrometric detection of chlorophyll <i>a</i> and the tetrapyrrole secondary metabolite tolyporphin A in the filamentous cyanobacterium HT-58-2. Approaches to high-throughput screening of intact cyanobacteria. <i>Journal of Porphyrins and Phthalocyanines</i> , 2017, 21, 759-768.	0.4	9
70	Comparative proteomic analysis between nitrogen supplemented and starved conditions in <i>Magnaporthe oryzae</i> . <i>Proteome Science</i> , 2017, 15, 20.	0.7	18
71	Identification of Epigenetic Factor Proteins Expressed in Human Embryonic Stem Cell-Derived Trophoblasts and in Human Placental Trophoblasts. <i>Journal of Proteome Research</i> , 2016, 15, 2433-2444.	1.8	9
72	A cell wall-bound anionic peroxidase, PtrPO21, is involved in lignin polymerization in <i>Populus trichocarpa</i> . <i>Tree Genetics and Genomes</i> , 2016, 12, 1.	0.6	24

#	ARTICLE	IF	CITATIONS
73	Analysis of Antiretrovirals in Single Hair Strands for Evaluation of Drug Adherence with Infrared-Matrix-Assisted Laser Desorption Electrospray Ionization Mass Spectrometry Imaging. <i>Analytical Chemistry</i> , 2016, 88, 1336-1344.	3.2	40
74	Infrared matrix-assisted laser desorption electrospray ionization mass spectrometry imaging analysis of biospecimens. <i>Analyst, The</i> , 2016, 141, 5236-5245.	1.7	50
75	Enhanced Lipidome Coverage in Shotgun Analyses by using Gas-Phase Fractionation. <i>Journal of the American Society for Mass Spectrometry</i> , 2016, 27, 1735-1744.	1.2	17
76	Examining ubiquitinated peptide enrichment efficiency through an epitope labeled protein. <i>Analytical Biochemistry</i> , 2016, 512, 114-119.	1.1	0
77	Whole-body Mass Spectrometry Imaging by Infrared Matrix-assisted Laser Desorption Electrospray Ionization (IR-MALDESI). <i>Journal of Visualized Experiments</i> , 2016, , e53942.	0.2	13
78	A Quantitative Glycomics and Proteomics Combined Purification Strategy. <i>Journal of Visualized Experiments</i> , 2016, , .	0.2	11
79	What if you could only publish 50 papers your entire career?. <i>Analytical and Bioanalytical Chemistry</i> , 2016, 408, 663-664.	1.9	1
80	Composition of Rosenthal Fibers, the Protein Aggregate Hallmark of Alexander Disease. <i>Journal of Proteome Research</i> , 2016, 15, 2265-2282.	1.8	34
81	TransOmic analysis of forebrain sections in Sp2 conditional knockout embryonic mice using IR-MALDESI imaging of lipids and LC-MS/MS label-free proteomics. <i>Analytical and Bioanalytical Chemistry</i> , 2016, 408, 3453-3474.	1.9	14
82	Optimizing Mass Spectrometry Analyses: A Tailored Review on the Utility of Design of Experiments. <i>Journal of the American Society for Mass Spectrometry</i> , 2016, 27, 767-785.	1.2	56
83	MALDESI: Fundamentals, Direct Analysis, and MS Imaging. , 2016, , 169-182.		4
84	Wall modified photonic crystal fibre capillaries as porous layer open tubular columns for in-capillary micro-extraction and capillary chromatography. <i>Analytica Chimica Acta</i> , 2016, 905, 1-7.	2.6	23
85	Polarity switching mass spectrometry imaging of healthy and cancerous hen ovarian tissue sections by infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Analyst, The</i> , 2016, 141, 595-605.	1.7	43
86	Influence of C-Trap Ion Accumulation Time on the Detectability of Analytes in IR-MALDESI MSI. <i>Analytical Chemistry</i> , 2015, 87, 10483-10490.	3.2	17
87	Machine learning reveals sex-specific 17 $\beta$ -estradiol-responsive expression patterns in white perch ( <i>Morone americana</i> ) plasma proteins. <i>Proteomics</i> , 2015, 15, 2678-2690.	1.3	13
88	Mass Spectrometry Imaging Reveals Heterogeneous Efavirenz Distribution within Putative HIV Reservoirs. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 2944-2948.	1.4	67
89	4-Coumaroyl and Caffeoyl Shikimic Acids Inhibit 4-Coumaric Acid:Coenzyme A Ligases and Modulate Metabolic Flux for 3-Hydroxylation in Monolignol Biosynthesis of <i>Populus trichocarpa</i> . <i>Molecular Plant</i> , 2015, 8, 176-187.	3.9	50
90	Analysis of trace fibers by IR-MALDESI imaging coupled with high resolving power MS. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 813-820.	1.9	20

#	ARTICLE	IF	CITATIONS
91	Cellular-level mass spectrometry imaging using infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI) by oversampling. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 2265-2271.	1.9	39
92	Quantitative mass spectrometry imaging of emtricitabine in cervical tissue model using infrared matrix-assisted laser desorption electrospray ionization. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 2073-2084.	1.9	66
93	Definitive Screening Design Optimization of Mass Spectrometry Parameters for Sensitive Comparison of Filter and Solid Phase Extraction Purified, INLIGHT Plasma <sup>15</sup> N-Glycans. <i>Analytical Chemistry</i> , 2015, 87, 7305-7312.	3.2	35
94	In-depth LC-MS/MS analysis of the chicken ovarian cancer proteome reveals conserved and novel differentially regulated proteins in humans. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 6851-6863.	1.9	10
95	Phosphorylation is an on/off switch for 5-hydroxyconiferaldehyde O-methyltransferase activity in poplar monolignol biosynthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 8481-8486.	3.3	60
96	Michael Gross: 25 Years of Dedication and Leadership of <i>JASMS</i> (1990-2015). <i>Journal of the American Society for Mass Spectrometry</i> , 2015, 26, 1-4.	1.2	4
97	Influence of Desorption Conditions on Analyte Sensitivity and Internal Energy in Discrete Tissue or Whole Body Imaging by IR-MALDESI. <i>Journal of the American Society for Mass Spectrometry</i> , 2015, 26, 899-910.	1.2	22
98	Phosphoproteome Analysis Links Protein Phosphorylation to Cellular Remodeling and Metabolic Adaptation during <i>Magnaporthe oryzae</i> Appressorium Development. <i>Journal of Proteome Research</i> , 2015, 14, 2408-2424.	1.8	42
99	Activin/Nodal Signaling Switches the Terminal Fate of Human Embryonic Stem Cell-derived Trophoblasts. <i>Journal of Biological Chemistry</i> , 2015, 290, 8834-8848.	1.6	23
100	Global Proteomic Analysis of Functional Compartments in Immature Avian Follicles Using Laser Microdissection Coupled to LC-MS/MS. <i>Journal of Proteome Research</i> , 2015, 14, 3912-3923.	1.8	8
101	Elucidation of Xylem-Specific Transcription Factors and Absolute Quantification of Enzymes Regulating Cellulose Biosynthesis in <i>Populus trichocarpa</i> . <i>Journal of Proteome Research</i> , 2015, 14, 4158-4168.	1.8	14
102	Relative Quantification and Higher-Order Modeling of the Plasma Glycan Cancer Burden Ratio in Ovarian Cancer Case-Control Samples. <i>Journal of Proteome Research</i> , 2015, 14, 4394-4401.	1.8	18
103	Assessing drug and metabolite detection in liver tissue by UV-MALDI and IR-MALDESI mass spectrometry imaging coupled to FT-ICR MS. <i>International Journal of Mass Spectrometry</i> , 2015, 377, 448-455.	0.7	50
104	Mechanisms of Egg Yolk Formation and Implications on Early Life History of White Perch (Morone t. j.)	1.1	38
105	Focus on Advancing High Performance Mass Spectrometry, Honoring Dr. Richard D. Smith, Recipient of the 2013 Award for a Distinguished Contribution in Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2014, 25, 1997-1999.	1.2	0
106	C/EBP $\beta$ regulates CRL4 <sup>Cdt2</sup> -mediated degradation of p21 in response to UVB-induced DNA damage to control the G <sub>1</sub> /S checkpoint. <i>Cell Cycle</i> , 2014, 13, 3602-3610.	1.3	19
107	Evaluating nonpolar surface area and liquid chromatography/mass spectrometry response: an application for site occupancy measurements for enzyme intermediates in polyketide biosynthesis. <i>Rapid Communications in Mass Spectrometry</i> , 2014, 28, 2511-2522.	0.7	4
108	Hierarchical Self-Assembly of Supramolecular Hydrophobic Metallacycles into Ordered Nanostructures. <i>Chemistry - an Asian Journal</i> , 2014, 9, 2928-2936.	1.7	23

#	ARTICLE	IF	CITATIONS
109	Silver dopants for targeted and untargeted direct analysis of unsaturated lipids via infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI). <i>Rapid Communications in Mass Spectrometry</i> , 2014, 28, 2461-2470.	0.7	21
110	Reply to the Comment on: "Utilizing Artificial Neural Networks in MATLAB to Achieve Parts-Per-Billion Mass Measurement Accuracy with a Fourier Transform Ion Cyclotron Resonance Mass Spectrometer" by D. Keith Williams Jr., Alexander L. Kovach, David C. Muddiman, and Kenneth W. Hanck. <i>J. Am. Soc. Mass Spectrom.</i> 20, 1303-1310 (2009). <i>Journal of the American Society for Mass Spectrometry</i> , 2014, 25, 697-697.	1.2	0
111	Are Presentations at ASMS Conferences Publications?. <i>Journal of the American Society for Mass Spectrometry</i> , 2014, 25, 301-302.	1.2	0
112	Accurate Identification of Deamidated Peptides in Global Proteomics Using a Quadrupole Orbitrap Mass Spectrometer. <i>Journal of Proteome Research</i> , 2014, 13, 777-785.	1.8	31
113	4-Coumaroyl and Caffeoyl Shikimic Acids Inhibit 4-Coumaric Acid: Coenzyme A Ligases and Modulate Metabolic Flux for 3-Hydroxylation in Monolignol Biosynthesis of <i>Populus trichocarpa</i> . <i>Molecular Plant</i> , 2014, , .	3.9	0
114	Establishing ion ratio thresholds based on absolute peak area for absolute protein quantification using protein cleavage isotope dilution mass spectrometry. <i>Analyst</i> , The, 2014, 139, 5439-5450.	1.7	9
115	Systems Biology of Lignin Biosynthesis in <i>Populus trichocarpa</i> : Heteromeric 4-Coumaric Acid:Coenzyme A Ligase Protein Complex Formation, Regulation, and Numerical Modeling. <i>Plant Cell</i> , 2014, 26, 876-893.	3.1	75
116	Mass Spectrometry Imaging of Hair Strands Allows for Evaluation of Long Term Antiretroviral Adherence. <i>AIDS Research and Human Retroviruses</i> , 2014, 30, A69-A69.	0.5	2
117	Compartment Proteomics Analysis of White Perch ( <i>Morone americana</i> ) Ovary Using Support Vector Machines. <i>Journal of Proteome Research</i> , 2014, 13, 1515-1526.	1.8	20
118	Complete Proteomic-Based Enzyme Reaction and Inhibition Kinetics Reveal How Monolignol Biosynthetic Enzyme Families Affect Metabolic Flux and Lignin in <i>Populus trichocarpa</i> . <i>Plant Cell</i> , 2014, 26, 894-914.	3.1	136
119	IR-MALDESI Mass Spectrometry Imaging of Biological Tissue Sections Using Ice as a Matrix. <i>Journal of the American Society for Mass Spectrometry</i> , 2014, 25, 319-328.	1.2	119
120	Mapping Antiretroviral Drugs in Tissue by IR-MALDESI MSI Coupled to the Q Exactive and Comparison with LC-MS/MS SRM Assay. <i>Journal of the American Society for Mass Spectrometry</i> , 2014, 25, 2038-2047.	1.2	44
121	Individuality Normalization when Labeling with Isotopic Glycan Hydrazide Tags (INLIGHT): A Novel Glycan-Relative Quantification Strategy. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 1376-1384.	1.2	44
122	Mass Recalibration of FT-ICR Mass Spectrometry Imaging Data Using the Average Frequency Shift of Ambient Ions. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 1137-1145.	1.2	21
123	MSiReader: An Open-Source Interface to View and Analyze High Resolving Power MS Imaging Files on Matlab Platform. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 718-721.	1.2	339
124	Regulation of phenylalanine ammonia-lyase (PAL) gene family in wood forming tissue of <i>Populus trichocarpa</i> . <i>Planta</i> , 2013, 238, 487-497.	1.6	53
125	Multi-peptide nLC-PC-IDMS-SRM-based Assay for the quantification of biomarkers in the chicken ovarian cancer model. <i>Methods</i> , 2013, 61, 323-330.	1.9	13
126	Infrared Matrix-Assisted Laser Desorption Electrospray Ionization (IR-MALDESI) Imaging Source Coupled to a FT-ICR Mass Spectrometer. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 92-100.	1.2	106



#	ARTICLE	IF	CITATIONS
127	Direct Analysis of Textile Fabrics and Dyes Using Infrared Matrix-Assisted Laser Desorption Electrospray Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , 2013, 85, 831-836.	3.2	46
128	Comparative Proteomic Analysis and IgE Binding Properties of Peanut Seed and Testa (Skin). <i>Journal of Agricultural and Food Chemistry</i> , 2013, 61, 3957-3968.	2.4	23
129	Factorial Experimental Designs Elucidate Significant Variables Affecting Data Acquisition on a Quadrupole Orbitrap Mass Spectrometer. <i>Journal of the American Society for Mass Spectrometry</i> , 2013, 24, 1501-1512.	1.2	27
130	Poly Specific <i>trans</i> -Acyltransferase Machinery Revealed <i>via</i> Engineered Acyl-CoA Synthetases. <i>ACS Chemical Biology</i> , 2013, 8, 200-208.	1.6	60
131	Understanding the Role of Proteolytic Digestion on Discovery and Targeted Proteomic Measurements Using Liquid Chromatography Tandem Mass Spectrometry and Design of Experiments. <i>Journal of Proteome Research</i> , 2013, 12, 5820-5829.	1.8	44
132	Monolignol Pathway 4-Coumaric Acid:Coenzyme A Ligases in <i>Populus. trichocarpa</i> : Novel Specificity, Metabolic Regulation, and Simulation of Coenzyme A Ligation Fluxes Å. <i>Plant Physiology</i> , 2013, 161, 1501-1516.	2.3	54
133	Temporal Analysis of the <i>Magnaporthe Oryzae</i> Proteome During Conidial Germination and Cyclic AMP (cAMP)-mediated Appressorium Formation. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2249-2265.	2.5	39
134	The use of a xylosylated plant glycoprotein as an internal standard accounting for <i>N</i> -linked glycan cleavage and sample preparation variability. <i>Rapid Communications in Mass Spectrometry</i> , 2013, 27, 1354-1358.	0.7	6
135	Targeted Proteomics of the Secretory Pathway Reveals the Secretome of Mouse Embryonic Fibroblasts and Human Embryonic Stem Cells. <i>Molecular and Cellular Proteomics</i> , 2012, 11, 1829-1839.	2.5	31
136	Peptide Production and Decay Rates Affect the Quantitative Accuracy of Protein Cleavage Isotope Dilution Mass Spectrometry (PC-IDMS). <i>Molecular and Cellular Proteomics</i> , 2012, 11, 814-823.	2.5	69
137	Functional redundancy of the two 5-hydroxylases in monolignol biosynthesis of <i>Populus trichocarpa</i> : LC-MS/MS based protein quantification and metabolic flux analysis. <i>Planta</i> , 2012, 236, 795-808.	1.6	19
138	Systematic Comparison of Reverse Phase and Hydrophilic Interaction Liquid Chromatography Platforms for the Analysis of N-Linked Glycans. <i>Analytical Chemistry</i> , 2012, 84, 8198-8206.	3.2	23
139	Evidence for Complex Molecular Architectures for Solvent-Extracted Lignins. <i>ACS Macro Letters</i> , 2012, 1, 568-573.	2.3	33
140	Coordination-Driven Self-Assembly of Charged and Neutral Dendritic Tetrakis(ferrocenyl) Rhomboids. <i>Organometallics</i> , 2012, 31, 7241-7247.	1.1	36
141	Analytical strategies for the global quantification of intact proteins. <i>Amino Acids</i> , 2012, 43, 1109-1117.	1.2	28
142	Comprehensive Quantification of Monolignol-Pathway Enzymes in <i>Populus trichocarpa</i> by Protein Cleavage Isotope Dilution Mass Spectrometry. <i>Journal of Proteome Research</i> , 2012, 11, 3390-3404.	1.8	42
143	In-Depth Analysis of the <i>Magnaporthe oryzae</i> Conidial Proteome. <i>Journal of Proteome Research</i> , 2012, 11, 5827-5835.	1.8	30
144	The subcellular proteome of undifferentiated human embryonic stem cells. <i>Proteomics</i> , 2012, 12, 421-430.	1.3	16

#	ARTICLE	IF	CITATIONS
145	High-resolution mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2012, 403, 1201-1202.	1.9	4
146	Absolute quantification of free glutathione and cysteine in aquatic insects using isotope dilution and selected reaction monitoring. <i>Analytical and Bioanalytical Chemistry</i> , 2012, 402, 357-366.	1.9	8
147	Polyubiquitin Is Required for Growth, Development and Pathogenicity in the Rice Blast Fungus <i>Magnaporthe oryzae</i> . <i>PLoS ONE</i> , 2012, 7, e42868.	1.1	65
148	Stable-Isotope Labeled Hydrophobic Hydrazide Reagents for the Relative Quantification of N-Linked Glycans by Electrospray Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , 2011, 83, 6738-6745.	3.2	90
149	Facile Self-Assembly of Supramolecular Hexakisferrocenyl Triangles via Coordination-Driven Self-Assembly and Their Electrochemical Behavior. <i>Organometallics</i> , 2011, 30, 3637-3642.	1.1	37
150	Performance Characteristics of a New Hybrid Quadrupole Time-of-Flight Tandem Mass Spectrometer (TripleTOF 5600). <i>Analytical Chemistry</i> , 2011, 83, 5442-5446.	3.2	254
151	Increasing proteome coverage with offline RP HPLC coupled to online RP nanoLC-MS. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2011, 879, 610-614.	1.2	24
152	One-year plasma N-linked glycome intra-individual and inter-individual variability in the chicken model of spontaneous ovarian adenocarcinoma. <i>International Journal of Mass Spectrometry</i> , 2011, 305, 79-86.	0.7	8
153	Evaluation of Normalization Methods on GeLC-MS/MS Label-Free Spectral Counting Data to Correct for Variation during Proteomic Workflows. <i>Journal of the American Society for Mass Spectrometry</i> , 2011, 22, 2199-2208.	1.2	77
154	Spectral Accuracy and Sulfur Counting Capabilities of the LTQ-FT-ICR and the LTQ-Orbitrap XL for Small Molecule Analysis. <i>Journal of the American Society for Mass Spectrometry</i> , 2011, 22, 2269-2275.	1.2	45
155	Improving Proteome Coverage on a LTQ-Orbitrap Using Design of Experiments. <i>Journal of the American Society for Mass Spectrometry</i> , 2011, 22, 773-783.	1.2	54
156	Hydrophobic Derivatization of N-linked Glycans for Increased Ion Abundance in Electrospray Ionization Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2011, 22, 1309-1317.	1.2	71
157	Identification of alternative splice variants in <i>Aspergillus flavus</i> through comparison of multiple tandem MS search algorithms. <i>BMC Genomics</i> , 2011, 12, 358.	1.2	12
158	Comparison of stable isotope labeling with amino acids in cell culture and spectral counting for relative quantification of protein expression. <i>Rapid Communications in Mass Spectrometry</i> , 2011, 25, 2524-2532.	0.7	42
159	Global optimization of the infrared matrix-assisted laser desorption electrospray ionization (IR-MALDESI) in Mass Spectrometry, 2011, 25, 3527-3536.	0.7	38
160	Design, modeling, fabrication, and evaluation of the air amplifier for improved detection of biomolecules by electrospray ionization mass spectrometry. <i>International Journal of Mass Spectrometry</i> , 2011, 300, 99-107.	0.7	19
161	Capitalizing on the hydrophobic bias of electrospray ionization through chemical modification in mass spectrometry-based proteomics. <i>Expert Review of Proteomics</i> , 2011, 8, 317-323.	1.3	17
162	John Bennett Fenn (1917-2010). <i>Science</i> , 2011, 331, 160-160.	6.0	1

#	ARTICLE	IF	CITATIONS
163	Membrane protein complexes catalyze both 4- and 3-hydroxylation of cinnamic acid derivatives in monoglignol biosynthesis. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 21253-21258.	3.3	133
164	N-Linked Global Glycan Profiling by NanoLC Mass Spectrometry. Methods in Molecular Biology, 2011, 790, 87-97.	0.4	1
165	Utilizing Spectral Counting To Quantitatively Characterize Tandem Removal of Abundant Proteins (TRAP) in Human Plasma. Analytical Chemistry, 2010, 82, 10179-10185.	3.2	20
166	The effects of abundant plasma protein depletion on global glycan profiling using NanoLC FT-ICR mass spectrometry. Analytical and Bioanalytical Chemistry, 2010, 396, 1473-1479.	1.9	10
167	Part II: defining and quantifying individual and co-cultured intracellular proteomes of two thermophilic microorganisms by GeLC-MS2 and spectral counting. Analytical and Bioanalytical Chemistry, 2010, 398, 391-404.	1.9	10
168	Part I: characterization of the extracellular proteome of the extreme thermophile Caldicellulosiruptor saccharolyticus by GeLC-MS2. Analytical and Bioanalytical Chemistry, 2010, 398, 377-389.	1.9	16
169	Measuring the intra-individual variability of the plasma proteome in the chicken model of spontaneous ovarian adenocarcinoma. Analytical and Bioanalytical Chemistry, 2010, 398, 737-749.	1.9	25
170	Quantitative top-down proteomics of SILAC labeled human embryonic stem cells. Journal of the American Society for Mass Spectrometry, 2010, 21, 879-889.	1.2	58
171	Self-Assembly of Dendritic Tris(crown ether) Hexagons and Their Complexation with Dibenzylammonium Cations. Journal of Organic Chemistry, 2010, 75, 7373-7380.	1.7	50
172	Direct Comparison of Stable Isotope Labeling by Amino Acids in Cell Culture and Spectral Counting for Quantitative Proteomics. Analytical Chemistry, 2010, 82, 8696-8702.	3.2	86
173	Detection of Alternative Splice Variants at the Proteome Level in <i>Aspergillus flavus</i> . Journal of Proteome Research, 2010, 9, 1209-1217.	1.8	29
174	Interplay of Permanent Charge and Hydrophobicity in the Electrospray Ionization of Glycans. Analytical Chemistry, 2010, 82, 6636-6642.	3.2	53
175	Facile Self-Assembly of Dendritic Multiferrocenyl Hexagons and Their Electrochemistry. Organometallics, 2010, 29, 6137-6140.	1.1	37
176	Increasing the hydrophobicity and electrospray response of glycans through derivatization with novel cationic hydrazides. Chemical Communications, 2010, 46, 237-239.	2.2	52
177	Improving limits of detection for B-type natriuretic peptide using PC-IDMS: An application of the ALiPHAT strategy. Analyst, The, 2010, 135, 36-41.	1.7	20
178	Study of the ionization mechanism in hybrid laser based desorption techniques. Analyst, The, 2010, 135, 880.	1.7	37
179	Intact and top-down characterization of biomolecules and direct analysis using infrared matrix-assisted laser desorption electrospray ionization coupled to FT-ICR mass spectrometry. Journal of the American Society for Mass Spectrometry, 2009, 20, 667-673.	1.2	101
180	Utilizing artificial neural networks in matlab to achieve parts-per-billion mass measurement accuracy with a fourier transform ion cyclotron resonance mass spectrometer. Journal of the American Society for Mass Spectrometry, 2009, 20, 1303-1310.	1.2	15

#	ARTICLE	IF	CITATIONS
181	Characterization of charge separation in the array of Micromachined UltraSonic ElectroSpray (AMUSE) ion source for mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2009, 20, 1684-1687.	1.2	12
182	Atmospheric pressure infrared (10.6 μm) laser desorption electrospray ionization (IR-LDESI) coupled to a LTQ Fourier transform ion cyclotron resonance mass spectrometer. <i>Rapid Communications in Mass Spectrometry</i> , 2009, 23, 1989-1992.	0.7	47
183	Generation of multiply charged peptides and proteins by radio frequency acoustic desorption and ionization for mass spectrometric detection. <i>Journal of the American Society for Mass Spectrometry</i> , 2009, 20, 597-600.	1.2	26
184	Evaluation of the ALiPHAT method for PC-IDMS and correlation of limits-of-detection with nonpolar surface area. <i>Journal of the American Society for Mass Spectrometry</i> , 2009, 20, 2006-2012.	1.2	15
185	Coupling of a vented column with splitless nanoRPLC-ESI-MS for the improved separation and detection of brain natriuretic peptide-32 and its proteolytic peptides. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2009, 877, 948-954.	1.2	48
186	Development of a Robust and High Throughput Method for Profiling N-Linked Glycans Derived from Plasma Glycoproteins by NanoLC-FTICR Mass Spectrometry. <i>Journal of Proteome Research</i> , 2009, 8, 3764-3770.	1.8	42
187	Absolute Quantification of C-Reactive Protein in Human Plasma Derived from Patients with Epithelial Ovarian Cancer Utilizing Protein Cleavage Isotope Dilution Mass Spectrometry. <i>Journal of Proteome Research</i> , 2009, 8, 1085-1090.	1.8	54
188	Development of a nanoLC LTQ Orbitrap Mass Spectrometric Method for Profiling Glycans Derived from Plasma from Healthy, Benign Tumor Control, and Epithelial Ovarian Cancer Patients. <i>Analytical Chemistry</i> , 2009, 81, 1130-1136.	3.2	74
189	Mass Spectrometry-Based Biomarker Discovery: Toward a Global Proteome Index of Individuality. <i>Annual Review of Analytical Chemistry</i> , 2009, 2, 265-277.	2.8	130
190	Tandem Mass Spectrometry of Thiolate-Protected Au Nanoparticles Na <sub>25</sub> (SC <sub>2</sub> H <sub>4</sub> Ph) <sub>18</sub> (S(C <sub>2</sub> H <sub>4</sub> Ph) <sub>4</sub> ) <sub>4</sub> . <i>Journal of the American Chemical Society</i> , 2009, 131, 13844-13851.	1.2	10
191	Synthesis of Six-Component Metallodendrimers via [3 + 3] Coordination-Driven Self-Assembly. <i>Journal of Organic Chemistry</i> , 2009, 74, 3524-3527.	1.7	26
192	Proteomics Characterization of Cell Membrane Blebs in Human Retinal Pigment Epithelium Cells. <i>Molecular and Cellular Proteomics</i> , 2009, 8, 2201-2211.	2.5	38
193	Mass measurement accuracy comparisons between a double-focusing magnetic sector and a time-of-flight mass analyzer. <i>Rapid Communications in Mass Spectrometry</i> , 2008, 22, 1563-1566.	0.7	11
194	Investigations with N-linked protein glycosylations by matrix-assisted laser desorption/ionization Fourier transform ion cyclotron resonance mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2008, 43, 1215-1223.	0.7	13
195	Calibration laws based on multiple linear regression applied to matrix-assisted laser desorption/ionization Fourier transform ion cyclotron resonance mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2008, 43, 1659-1663.	0.7	13
196	Construction of a Versatile High Precision Ambient Ionization Source for Direct Analysis and Imaging. <i>Journal of the American Society for Mass Spectrometry</i> , 2008, 19, 1527-1534.	1.2	42
197	Synthesis, Characterization, and Application of Iodoacetamide Derivatives Utilized for the ALiPHAT Strategy. <i>Journal of the American Chemical Society</i> , 2008, 130, 2122-2123.	6.6	43
198	Human Proteinpedia enables sharing of human protein data. <i>Nature Biotechnology</i> , 2008, 26, 164-167.	9.4	155

#	ARTICLE	IF	CITATIONS
199	A New Family of Multiferrocene Complexes with Enhanced Control of Structure and Stoichiometry via Coordination-Driven Self-Assembly and Their Electrochemistry. <i>Journal of the American Chemical Society</i> , 2008, 130, 839-841.	6.6	160
200	Top-Down Identification and Quantification of Stable Isotope Labeled Proteins from <i>Aspergillus flavus</i> Using Online Nano-Flow Reversed-Phase Liquid Chromatography Coupled to a LTQ-FTICR Mass Spectrometer. <i>Analytical Chemistry</i> , 2008, 80, 4994-5001.	3.2	68
201	Studying O-Linked Protein Glycosylations in Human Plasma. <i>Journal of Proteome Research</i> , 2008, 7, 2562-2568.	1.8	18
202	Coordination-Driven Face-Directed Self-Assembly of Trigonal Prisms. Face-Based Conformational Chirality. <i>Journal of the American Chemical Society</i> , 2008, 130, 7620-7628.	6.6	100
203	Temperature-Dependent Regulation of Proteins in <i>Aspergillus flavus</i> : Whole Organism Stable Isotope Labeling by Amino Acids. <i>Journal of Proteome Research</i> , 2008, 7, 2973-2979.	1.8	36
204	Ambient Aerodynamic Ionization Source for Remote Analyte Sampling and Mass Spectrometric Analysis. <i>Analytical Chemistry</i> , 2008, 80, 5266-5271.	3.2	32
205	Development and Characterization of an Ionization Technique for Analysis of Biological Macromolecules: Liquid Matrix-Assisted Laser Desorption Electrospray Ionization. <i>Analytical Chemistry</i> , 2008, 80, 6773-6778.	3.2	47
206	Coordination-Driven Self-Assembly of Cavity-Cored Multiple Crown Ether Derivatives and Poly[2]pseudorotaxanes. <i>Journal of the American Chemical Society</i> , 2008, 130, 5320-5334.	6.6	113
207	Effect of Plasma Protein Depletion on BNP-32 Recovery. <i>Clinical Chemistry</i> , 2008, 54, 933-934.	1.5	11
208	A Method for Automatically Interpreting Mass Spectra of <sup>18</sup> O-Labeled Isotopic Clusters. <i>Molecular and Cellular Proteomics</i> , 2007, 6, 305-318.	2.5	59
209	Dual Electrospray Ion Source for Electron-Transfer Dissociation on a Hybrid Linear Ion Trap Orbitrap Mass Spectrometer. <i>Analytical Chemistry</i> , 2007, 79, 7916-7919.	3.2	32
210	Parts-Per-Billion Mass Measurement Accuracy Achieved through the Combination of Multiple Linear Regression and Automatic Gain Control in a Fourier Transform Ion Cyclotron Resonance Mass Spectrometer. <i>Analytical Chemistry</i> , 2007, 79, 5058-5063.	3.2	45
211	Elevated Levels of Phosphorylated Fibrinogen- $\alpha$ -Iso-forms and Differential Expression of Other Post-Translationally Modified Proteins in the Plasma of Ovarian Cancer Patients. <i>Proteome Res.</i> 2006, 5, 3318-3325. <i>Journal of Proteome Research</i> , 2007, 6, 1615-1615.	1.8	2
212	Details of the Structure Determination of the Sulfated Steroids PSDS and PADS: A New Components of the Sea Lamprey ( <i>Petromyzon marinus</i> ) Migratory Pheromone. <i>Journal of Organic Chemistry</i> , 2007, 72, 7544-7550.	1.7	41
213	A Highly Efficient Approach to the Self-Assembly of Hexagonal Cavity-Cored Tris[2]pseudorotaxanes from Several Components via Multiple Noncovalent Interactions. <i>Journal of the American Chemical Society</i> , 2007, 129, 14187-14189.	6.6	119
214	Achieving Augmented Limits of Detection for Peptides with Hydrophobic Alkyl Tags. <i>Analytical Chemistry</i> , 2007, 79, 3989-3995.	3.2	43
215	Carbohydrate Analysis by Desorption Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Analytical Chemistry</i> , 2007, 79, 8812-8815.	3.2	38
216	Epithelial Ovarian Cancer: Disease Etiology, Treatment, Detection, and Investigational Gene, Metabolite, and Protein Biomarkers. <i>Journal of Proteome Research</i> , 2007, 6, 2936-2962.	1.8	102

#	ARTICLE	IF	CITATIONS
217	Coordination-Driven Self-Assembly of Metallodendrimers Possessing Well-Defined and Controllable Cavities as Cores. <i>Journal of the American Chemical Society</i> , 2007, 129, 2120-2129.	6.6	129
218	Sub parts-per-million mass measurement accuracy of intact proteins and product ions achieved using a dual electrospray ionization quadrupole fourier transform ion cyclotron resonance mass spectrometer. <i>Journal of the American Society for Mass Spectrometry</i> , 2007, 18, 1-7.	1.2	18
219	Detection of Attomole Amounts of Analyte by Desorption Electrospray Ionization Mass Spectrometry (DESI-MS) Determined Using Fluorescence Spectroscopy. <i>Journal of the American Society for Mass Spectrometry</i> , 2007, 18, 1093-1096.	1.2	52
220	Remote mass spectrometric sampling of electrospray- and desorption electrospray-generated ions using an air ejector. <i>Journal of the American Society for Mass Spectrometry</i> , 2007, 18, 1844-1847.	1.2	33
221	Probing the mechanisms of an air amplifier using a LTQ-FT-ICR-MS and fluorescence spectroscopy. <i>Journal of the American Society for Mass Spectrometry</i> , 2007, 18, 1909-1913.	1.2	18
222	Effect of matrix crystal structure on ion abundance of carbohydrates by matrix-assisted laser desorption/ionization Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 807-811.	0.7	20
223	Direct characterization of intact polypeptides by matrix-assisted laser desorption electrospray ionization quadrupole Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 1150-1154.	0.7	54
224	Understanding the influence of post-excite radius and axial confinement on quantitative proteomic measurements using Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 1196-1204.	0.7	7
225	Quantitative comparison of a flared and a standard heated metal capillary inlet with a voltage-assisted air amplifier on an electrospray ionization linear ion trap mass spectrometer. <i>Rapid Communications in Mass Spectrometry</i> , 2007, 21, 3207-3212.	0.7	10
226	Differential protein expression in male and female human lumbar cerebrospinal fluid using iTRAQ reagents after abundant protein depletion. <i>Proteomics</i> , 2007, 7, 3726-3734.	1.3	41
227	Chromatin Assembly Factor 1 Interacts with Histone H3 Methylated at Lysine 79 in the Processes of Epigenetic Silencing and DNA Repair. <i>Biochemistry</i> , 2006, 45, 2852-2861.	1.2	64
228	Elevated Levels of Phosphorylated Fibrinogen-Î±-Isoforms and Differential Expression of Other Post-Translationally Modified Proteins in the Plasma of Ovarian Cancer Patients. <i>Journal of Proteome Research</i> , 2006, 5, 3318-3325.	1.8	49
229	"Lewis & Clark" Proteomics. <i>Journal of Proteome Research</i> , 2006, 5, 221-222.	1.8	4
230	Incorporation of 2,6-Di(4-ethyl-dipyridyl)-9-thiabicyclo[3.3.1]nonane into Discrete 2D Supramolecules via Coordination-Driven Self-Assembly. <i>Journal of Organic Chemistry</i> , 2006, 71, 6644-6647.	1.7	24
231	Identification of Subunit-Subunit Interactions in Bacteriophage P22 Procapsids by Chemical Cross-linking and Mass Spectrometry. <i>Journal of Proteome Research</i> , 2006, 5, 370-377.	1.8	49
232	Molecular Architecture via Coordination: Self-Assembly of Nanoscale Hexagonal Metallodendrimers with Designed Building Blocks. <i>Journal of the American Chemical Society</i> , 2006, 128, 10014-10015.	6.6	103
233	Direct high-resolution peptide and protein analysis by desorption electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2006, 20, 3409-3411.	0.7	80
234	Generation and detection of multiply-charged peptides and proteins by matrix-assisted laser desorption electrospray ionization (MALDESI) fourier transform ion cyclotron resonance mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2006, 17, 1712-1716.	1.2	336

#	ARTICLE	IF	CITATIONS
235	Accessible proteomics space and its implications for peak capacity for zero-, one- and two-dimensional separations coupled with FT-ICR and TOF mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2006, 41, 281-288.	0.7	55
236	Leveling response factors in the electrospray ionization process using a heated capillary interface. <i>Journal of the American Society for Mass Spectrometry</i> , 2005, 16, 772-778.	1.2	21
237	Accurate mass precursor ion data and tandem mass spectrometry identify a class I human leukocyte antigen A*0201-presented peptide originating from vaccinia virus. <i>Journal of the American Society for Mass Spectrometry</i> , 2005, 16, 1812-1817.	1.2	24
238	Protein expression profiling of CLL B cells using replicate off-line strong cation exchange chromatography and LC-MS/MS. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2005, 819, 33-39.	1.2	15
239	Derivatives of pentamidine designed to target the Leishmania lipophosphoglycan. <i>Tetrahedron Letters</i> , 2005, 46, 695-698.	0.7	13
240	Effect of post-excitation radius on ion abundance, mass measurement accuracy, and isotopic distributions in Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 915-918.	0.7	14
241	Normalization of relative peptide ratios derived from in-gel digests: applications to protein variant analysis at the peptide level. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 2871-2877.	0.7	4
242	Primary Trabecular Meshwork Cells Incubated in Human Aqueous Humor Differ from Cells Incubated in Serum Supplements. , 2005, 46, 2848.		34
243	DÃ©jÃ Vu All Over Again: Skin Cap Still Contains a High-Potency Glucocorticosteroid. <i>Archives of Dermatology</i> , 2005, 141, 801-3.	1.7	4
244	Quantitative mass spectral evidence for the absence of circulating brain natriuretic peptide (BNP-32) in severe human heart failure. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 17442-17447.	3.3	256
245	Evaluation of Protein Depletion Methods for the Analysis of Total-, Phospho- and Glycoproteins in Lumbar Cerebrospinal Fluid. <i>Journal of Proteome Research</i> , 2005, 4, 837-845.	1.8	58
246	Gentle Protein Ionization Assisted by High-Velocity Gas Flow. <i>Analytical Chemistry</i> , 2005, 77, 6174-6183.	3.2	44
247	Identification of Class II HLA-DRB1*03-Bound Measles Virus Peptides by 2D-Liquid Chromatography Tandem Mass Spectrometry. <i>Journal of Proteome Research</i> , 2005, 4, 2243-2249.	1.8	26
248	Coordination-Driven Self-Assemblies with a Carborane Backbone. <i>Journal of the American Chemical Society</i> , 2005, 127, 12131-12139.	6.6	214
249	Quantitative Protein Expression Analysis Of CLL B Cells from Mutated and Unmutated IgVHSubgroups Using Acid-Cleavable Isotope-Coded Affinity Tag Reagents. <i>Journal of Proteome Research</i> , 2005, 4, 1310-1317.	1.8	15
250	Mapping Sites of Protein Phosphorylation by Mass Spectrometry Utilizing a Chemical-Enzymatic Approach:Ã Characterization of Products from Î±-S1Casein Phosphopeptides. <i>Journal of Proteome Research</i> , 2005, 4, 424-434.	1.8	8
251	Statistical Evaluation of Internal and External Mass Calibration Laws Utilized in Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Analytical Chemistry</i> , 2005, 77, 2406-2414.	3.2	43
252	Reproducibility of retention time using a splitless nanoLC coupled to an ESI-FTICR mass spectrometer. <i>Journal of Biomolecular Techniques</i> , 2005, 16, 414-22.	0.8	1

#	ARTICLE	IF	CITATIONS
253	Discovery of Ovarian Cancer Biomarkers in Serum Using NanoLC Electrospray Ionization TOF and FT-ICR Mass Spectrometry. <i>Disease Markers</i> , 2004, 19, 239-249.	0.6	59
254	Detection of Genetic Variants of Transthyretin by Liquid Chromatography- <sup>19</sup> F-Dual Electrospray Ionization Fourier-Transform Ion-Cyclotron-Resonance Mass Spectrometry. <i>Clinical Chemistry</i> , 2004, 50, 1535-1543.	1.5	47
255	Identification of Transthyretin Variants by Sequential Proteomic and Genomic Analysis. <i>Clinical Chemistry</i> , 2004, 50, 1544-1552.	1.5	61
256	Identification and Characterization of Novel, Naturally Processed Measles Virus Class II HLA-DRB1 Peptides. <i>Journal of Virology</i> , 2004, 78, 42-51.	1.5	31
257	Androgens Negatively Regulate Forkhead Transcription Factor FKHR (FOXO1) through a Proteolytic Mechanism in Prostate Cancer Cells. <i>Journal of Biological Chemistry</i> , 2004, 279, 13866-13877.	1.6	61
258	Informed use of proteolytic inhibitors in biomarker discovery. <i>Rapid Communications in Mass Spectrometry</i> , 2004, 18, 1001-1002.	0.7	5
259	Utility of accurate monoisotopic mass measurements to confidently identify lambda exonuclease generated single-stranded amplicons containing 7-deaza analogs by electrospray ionization FT-ICR mass spectrometry. <i>International Journal of Mass Spectrometry</i> , 2004, 234, 79-87.	0.7	12
260	Determination of the relative energies of activation for the dissociation of aromatic versus aliphatic phosphopeptides by ESI-FTICR-MS and IRMPD. <i>Journal of the American Society for Mass Spectrometry</i> , 2004, 15, 121-127.	1.2	42
261	A method for calculating <sup>16</sup> O/ <sup>18</sup> O peptide ion ratios for the relative quantification of proteomes. <i>Journal of the American Society for Mass Spectrometry</i> , 2004, 15, 437-445.	1.2	83
262	Evaluation of a Cleavable Stable Isotope Labeled Synthetic Peptide for Absolute Protein Quantification Using LC-MS/MS. <i>Journal of Proteome Research</i> , 2004, 3, 658-661.	1.8	83
263	Analytical Performance of a Venturi Device Integrated into an Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometer for Analysis of Nucleic Acids. <i>Analytical Chemistry</i> , 2004, 76, 4118-4122.	3.2	43
264	Analysis of the Low Molecular Weight Fraction of Serum by LC-Dual ESI-FT-ICR Mass Spectrometry: $\hat{A}$ Precision of Retention Time, Mass, and Ion Abundance. <i>Analytical Chemistry</i> , 2004, 76, 5097-5103.	3.2	62
265	Absolute Quantification of the Model Biomarker Prostate-Specific Antigen in Serum by LC-MS/MS Using Protein Cleavage and Isotope Dilution Mass Spectrometry. <i>Journal of Proteome Research</i> , 2004, 3, 644-652.	1.8	245
266	Naturally processed measles virus peptide eluted from class II HLA-DRB1*03 recognized by T lymphocytes from human blood. <i>Virology</i> , 2003, 312, 495-506.	1.1	29
267	<sup>1</sup> H NMR and electrospray mass spectrometry of the mono-ionized		



#	ARTICLE	IF	CITATIONS
271	Enzymatic strategies for the characterization of nucleic acids by electrospray ionization mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2003, 17, 2699-2706.	0.7	16
272	Implications of Hydrophobicity and Free Energy of Solvation for Characterization of Nucleic Acids by Electrospray Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , 2003, 75, 1331-1339.	3.2	74
273	Dual Electrospray Ionization Source for Confident Generation of Accurate Mass Tags Using Liquid Chromatography Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Analytical Chemistry</i> , 2003, 75, 3411-3418.	3.2	62
274	Coordination-Driven Self-Assembly of Supramolecular Cages: Heteroatom-Containing and Complementary Trigonal Prisms. <i>Journal of the American Chemical Society</i> , 2003, 125, 9647-9652.	6.6	79
275	Gas-Phase Ion Unimolecular Dissociation for Rapid Phosphopeptide Mapping by IRMPD in a Penning Ion Trap: An Energetically Favored Process. <i>Journal of the American Chemical Society</i> , 2002, 124, 6546-6547.	6.6	45
276	Dual-micro-ESI source for precise mass determination on a quadrupole time-of-flight mass spectrometer for genomic and proteomic applications. <i>Analytical and Bioanalytical Chemistry</i> , 2002, 373, 538-546.	1.9	24
277	In celebration of the 70th birthday of Dr David M. Hercules. <i>Analytical and Bioanalytical Chemistry</i> , 2002, 373, 517-518.	1.9	0
278	Memoirs, Hercules lineage, students, visiting scientists and staff. <i>Analytical and Bioanalytical Chemistry</i> , 2002, 373, 669-675.	1.9	0
279	Tailoring the gas-phase dissociation and determining the relative energy of activation for dissociation of 7-deaza purine modified oligonucleotides containing a repeating motif. <i>International Journal of Mass Spectrometry</i> , 2002, 219, 139-150.	0.7	26
280	CEPH family 1362 STR database: An online resource for characterization of PCR products using electrospray ionization mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2002, 13, 89-90.	1.2	0
281	Solution composition and thermal denaturation for the production of single-stranded PCR amplicons: Piperidine-induced destabilization of the DNA duplex?. <i>Journal of the American Society for Mass Spectrometry</i> , 2002, 13, 232-240.	1.2	17
282	Evaluation of sample preparation techniques for mass measurements of PCR products using ESI-FT-ICR mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2002, 13, 338-344.	1.2	22
283	Selective, Sensitive, and Rapid Phosphopeptide Identification in Enzymatic Digests Using ESI-FTICR-MS with Infrared Multiphoton Dissociation. <i>Analytical Chemistry</i> , 2001, 73, 3305-3311.	3.2	96
284	High-Mass Accuracy of Product Ions Produced by SORI-CID Using a Dual Electrospray Ionization Source Coupled with FTICR Mass Spectrometry. <i>Analytical Chemistry</i> , 2001, 73, 1247-1251.	3.2	57
285	Genotyping of Simple and Compound Short Tandem Repeat Loci Using Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Analytical Chemistry</i> , 2001, 73, 4514-4521.	3.2	41
286	Detection of double-stranded PCR amplicons at the attomole level electrosprayed from low nanomolar solutions using FT-ICR mass spectrometry. <i>Fresenius' Journal of Analytical Chemistry</i> , 2001, 369, 246-251.	1.5	21
287	Complete sequencing of mono-deprotonated peptide nucleic acids by sustained off-resonance irradiation collision-induced dissociation. <i>Journal of the American Society for Mass Spectrometry</i> , 2001, 12, 805-809.	1.2	9
288	Genotyping short tandem repeats using flow injection and electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2001, 15, 348-350.	0.7	25

#	ARTICLE	IF	CITATIONS
289	Impact of ion cloud densities on the measurement of relative ion abundances in Fourier transform ion cyclotron resonance mass spectrometry: experimental observations of coulombically induced cyclotron radius perturbations and ion cloud dephasing rates. <i>Journal of Mass Spectrometry</i> , 2001, 36, 195-203.	0.7	39
290	Perspectives on the use of electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry for short tandem repeat genotyping in the post-genome era. <i>Journal of Mass Spectrometry</i> , 2001, 36, 589-606.	0.7	39
291	Homogeneous Preparations of $3\text{-}^{32}\text{P}$ -Phosphoglycolate-Terminated Oligodeoxynucleotides from Bleomycin-Treated DNA as Verified by Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Analytical Biochemistry</i> , 2001, 289, 274-280.	1.1	18
292	<title>Genotyping complex short tandem repeats using electrospray ionization Fourier transform ion cyclotron resonance multistage mass spectrometry</title>. , 2000, , .		6
293	An experimental and theoretical study of the gas-phase decomposition of monoprotonated peptide nucleic acids. <i>Journal of the American Society for Mass Spectrometry</i> , 2000, 11, 615-625.	1.2	5
294	A dual electrospray ionization source combined with hexapole accumulation to achieve high mass accuracy of biopolymers in Fourier transform ion cyclotron resonance mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2000, 11, 876-883.	1.2	100
295	Preparation of single-stranded PCR products for electrospray ionization mass spectrometry using the DNA repair enzyme lambda exonuclease. <i>Analyst, The</i> , 2000, 125, 619-626.	1.7	37
296	Quantification of singly charged biomolecules by electrospray ionization fourier transform ion cyclotron resonance mass spectrometry utilizing an internal standard. <i>Rapid Communications in Mass Spectrometry</i> , 1999, 13, 164-171.	0.7	19
297	Characterization of a microdialysis approach to prepare polymerase chain reaction products for electrospray ionization mass spectrometry using on-line ultraviolet absorbance measurements and inductively coupled plasma-atomic emission spectroscopy. <i>Rapid Communications in Mass Spectrometry</i> , 1999, 13, 323-330.	0.7	40
298	Accurate characterization of the tyrosine hydroxylase forensic allele 9.3 through development of electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry. , 1999, 13, 954-962.		40
299	Precise mass measurement of a double-stranded 500 base-pair (309 kDa) polymerase chain reaction product by negative ion electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry. , 1999, 13, 1201-1204.		53
300	Hydrophathic influences on the quantification of equine heart cytochromec using relative ion abundance measurements by electrospray ionization Fourier transform ion cyclotron resonance mass spectrometry. , 1999, 34, 1055-1062.		37
301	Consequences of Nucleic Acid Conformation on the Binding of a Trinuclear Platinum Drugâ€. <i>Biochemistry</i> , 1999, 38, 14731-14737.	1.2	67
302	Hemoglobinase Activity of the Lysine Gingipain Protease (Kgp) of <i>Porphyromonas gingivalis</i> W83. <i>Journal of Bacteriology</i> , 1999, 181, 4905-4913.	1.0	109
303	Nanoelectrospray mass spectrometry using non-metalized, tapered (50 $\hat{\text{a}}$ ' 10 $\hat{\text{a}}$ ... $\hat{\text{a}}$ ¼m) fused-silica capillaries. <i>Rapid Communications in Mass Spectrometry</i> , 1998, 12, 443-448.	0.7	62
304	Comprehensive nomenclature for the fragment ions produced from collisional activation of peptide nucleic acids. <i>Rapid Communications in Mass Spectrometry</i> , 1998, 12, 759-762.	0.7	11
305	Heterogeneity in <i>Bacillus cereus</i> PCR Products Detected by ESI $\hat{\text{a}}$ FTICR Mass Spectrometry. <i>Analytical Chemistry</i> , 1998, 70, 1203-1207.	3.2	31
306	Sequencing and Characterization of Larger Oligonucleotides by Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. <i>Reviews in Analytical Chemistry</i> , 1998, 17, 1-68.	1.5	33

#	ARTICLE	IF	CITATIONS
307	Nanoscale Tectonics: Self-Assembly, Characterization, and Chemistry of a Novel Class of Organoplatinum Square Macrocycles. <i>Journal of the American Chemical Society</i> , 1997, 119, 11611-11619.	6.6	110
308	Transition-Metal-Mediated Rational Design and Self-Assembly of Chiral, Nanoscale Supramolecular Polyhedra with Unique Symmetry. <i>Organometallics</i> , 1997, 16, 3094-3096.	1.1	146
309	Length and Base Composition of PCR-Amplified Nucleic Acids Using Mass Measurements from Electrospray Ionization Mass Spectrometry. <i>Analytical Chemistry</i> , 1997, 69, 1543-1549.	3.2	96
310	Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry. <i>Instrumentation and Applications. Journal of Chemical Education</i> , 1997, 74, 1288.	1.1	39
311	Molecular Architecture via Coordination: Marriage of Crown Ethers and Calixarenes with Molecular Squares, Unique Tetranuclear Metallamacrocycles from Metallacrown Ether and Metallacalixarene Complexes via Self-Assembly. <i>Journal of the American Chemical Society</i> , 1997, 119, 5163-5168.	6.6	127
312	Improving the Microdialysis Procedure for Electrospray Ionization Mass Spectrometry of Biological Samples. , 1997, 32, 425-431.		37
313	Application of sequential paired covariance to liquid chromatography-mass spectrometry data enhancements in both the signal-to-noise ratio and the resolution of analyte peaks in the chromatogram. <i>Journal of Chromatography A</i> , 1997, 771, 1-7.	1.8	16
314	Characterization of PCR Products from Bacilli Using Electrospray Ionization FTICR Mass Spectrometry. <i>Analytical Chemistry</i> , 1996, 68, 3705-3712.	3.2	105
315	Design and Self-Assembly of Nanoscale Organoplatinum Macrocycles. <i>Journal of the American Chemical Society</i> , 1996, 118, 8731-8732.	6.6	96
316	Important Aspects concerning the Quantification of Biomolecules by Time-of-Flight Secondary-Ion Mass Spectrometry. <i>Applied Spectroscopy</i> , 1996, 50, 161-166.	1.2	9
317	Direct quantification of cocaine in urine by time-of-flight mass spectrometry. <i>Fresenius' Journal of Analytical Chemistry</i> , 1996, 354, 103-110.	1.5	19
318	Analysis of Double-stranded Polymerase Chain Reaction Products from the <i>Bacillus cereus</i> Group by Electrospray Ionization Fourier Transform Ion Cyclotron Resonance Mass Spectrometry. , 1996, 10, 29-35.		51
319	A Quantitative Study of in vitro Hepatic Metabolism of Tacrolimus (FK506) Using Secondary Ion and Matrix-assisted Laser Desorption/Ionization Mass Spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 1996, 10, 1215-1218.	0.7	16
320	Enhancement of ion intensity in time-of-flight secondary-ionization mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 1996, 7, 467-472.	1.2	38
321	Charge-state reduction with improved signal intensity of oligonucleotides in electrospray ionization mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 1996, 7, 697-706.	1.2	125
322	Simultaneous quantification of cyclosporin A and its major metabolites by time-of-flight secondary-ion mass spectrometry and matrix-assisted laser desorption/ionization mass spectrometry utilizing data analysis techniques: Comparison with high-performance liquid chromatography. <i>Journal of Mass Spectrometry</i> , 1995, 30, 1469-1479.	0.7	35
323	Application of secondary ion and matrix-assisted laser desorption-ionization time-of-flight mass spectrometry for the quantitative analysis of biological molecules. <i>Mass Spectrometry Reviews</i> , 1995, 14, 383-429.	2.8	53
324	Application of Sequential Paired Covariance to Capillary Electrophoresis Electrospray Ionization Time-of-Flight Mass Spectrometry: Unraveling the Signal from the Noise in the Electropherogram. <i>Analytical Chemistry</i> , 1995, 67, 4371-4375.	3.2	57

#	ARTICLE	IF	CITATIONS
325	Characterization of Polystyrene on Etched Silver Using Ion Scattering and X-ray Photoelectron Spectroscopy: Correlation of Secondary Ion Yield in Time-of-Flight SIMS with Surface Coverage. The Journal of Physical Chemistry, 1994, 98, 11570-11575.	2.9	29
326	Quantitative Measurement of Cyclosporin A in Blood by Time-of-Flight Mass Spectrometry. Analytical Chemistry, 1994, 66, 2362-2368.	3.2	94
327	Normalization Techniques for High-Throughput Screening by Infrared Matrix-Assisted Laser Desorption Electrospray Ionization Mass Spectrometry. Journal of Mass Spectrometry, 0, , .	0.7	4