

GÃ¼nter Allmaier

List of Publications by Year in descending order

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162
papers

4,503
citations

136740

32
h-index

133063

59
g-index

164
all docs

164
docs citations

164
times ranked

4910
citing authors

#	ARTICLE	IF	CITATIONS
1	Calcium ion effect on phospholipid bilayers as cell membrane analogues. <i>Bioelectrochemistry</i> , 2022, 143, 107988.	2.4	11
2	Atmospheric pressure matrix-assisted laser desorption/ionization mass spectrometry of engine oil additive components. <i>Rapid Communications in Mass Spectrometry</i> , 2022, 36, e9271.	0.7	4
3	Online hyphenation of size-exclusion chromatography and gas-phase electrophoresis facilitates the characterization of protein aggregates. <i>Electrophoresis</i> , 2021, 42, 1202-1208.	1.3	1
4	Comparative Analysis of Platelet-Derived Extracellular Vesicles Using Flow Cytometry and Nanoparticle Tracking Analysis. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3839.	1.8	21
5	Adeno-associated Virus Virus-like Particle Characterization via Orthogonal Methods: Nanoelectrospray Differential Mobility Analysis, Asymmetric Flow Field-Flow Fractionation, and Atomic Force Microscopy. <i>ACS Omega</i> , 2021, 6, 16428-16437.	1.6	7
6	Molecular Weight Determination of Adeno-Associate Virus serotype 8 Virus-like Particle either carrying or lacking genome via native nES Gas-phase Electrophoretic Molecular Mobility Analysis (GEMMA) and nESI QTOF Mass Spectrometry. <i>Journal of Mass Spectrometry</i> , 2021, 56, e4786.	0.7	6
7	nES-DMA with Charge-reduction based on Soft X-ray Radiation: Analysis of a Recombinant Monoclonal Antibody. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2021, 1182, 122925.	1.2	1
8	N-terminal VP1 Truncations Favor T = 1 Norovirus-Like Particles. <i>Vaccines</i> , 2021, 9, 8.	2.1	15
9	A possible role of gas-phase electrophoretic mobility molecular analysis (nES GEMMA) in extracellular vesicle research. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 7341-7352.	1.9	2
10	Nano electrospray differential mobility analysis based size-selection of liposomes and very-low density lipoprotein particles for offline hyphenation to MALDI mass spectrometry. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2020, 179, 112998.	1.4	4
11	Bipolar Corona Discharge-Based Charge Equilibration for Nano Electrospray Gas-Phase Electrophoretic Mobility Molecular Analysis of Bio- and Polymer Nanoparticles. <i>Analytical Chemistry</i> , 2020, 92, 8665-8669.	3.2	9
12	Ligand engineering of immobilized nanoclusters on surfaces: ligand exchange reactions with supported Au ₁₁ (PPh ₃) ₇ Br ₃ . <i>Nanoscale</i> , 2020, 12, 12809-12816.	2.8	19
13	Virus-like particle size and molecular weight/mass determination applying gas-phase electrophoresis (native nES GEMMA). <i>Analytical and Bioanalytical Chemistry</i> , 2019, 411, 5951-5962.	1.9	28
14	Native Nano-electrospray Differential Mobility Analyzer (nES GEMMA) Enables Size Selection of Liposomal Nanocarriers Combined with Subsequent Direct Spectroscopic Analysis. <i>Analytical Chemistry</i> , 2019, 91, 3860-3868.	3.2	14
15	Support effect on the reactivity and stability of Au ₂₅ (SR) ₁₈ and Au ₁₄₄ (SR) ₆₀ nanoclusters in liquid phase cyclohexane oxidation. <i>Catalysis Today</i> , 2019, 336, 174-185.	2.2	33
16	Development of an accelerated artificial ageing method for the characterization of degradation products of antioxidants in lubricants by mass spectrometry. <i>European Journal of Mass Spectrometry</i> , 2019, 25, 300-323.	0.5	6
17	In-depth analysis of crocetin ester glycosides from dried/processed stigmas of <i>Crocus sativus</i> L. by HPLC-ESI-MS ² (n = 2, 3). <i>Phytochemical Analysis</i> , 2019, 30, 1.2 346-356.	1.2	6
18	Optimization of MALDI-TOF mass spectrometry imaging for the visualization and comparison of peptide distributions in dry-cured ham muscle fibers. <i>Food Chemistry</i> , 2019, 283, 275-286.	4.2	30

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19	Nanoscale chemical imaging of individual chemotherapeutic cytarabine-loaded liposomal nanocarriers. <i>Nano Research</i> , 2019, 12, 197-203.	5.8	65
20	Optimization of sample preparation for intact cell mass spectrometry (matrix-assisted laser) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 707 Communications in Mass Spectrometry, 2018, 32, 815-823.	0.7	2
21	Size and molecular weight determination of polysaccharides by means of nano electrospray gas-phase electrophoretic mobility molecular analysis (nES GEMMA). <i>Electrophoresis</i> , 2018, 39, 1142-1150.	1.3	12
22	A laser desorption ionization/matrix-assisted laser desorption ionization target system applicable for three distinct types of instruments (LinTOF/curved field RTOF, LinTOF/RTOF and QqRTOF) with different performance characteristics from three vendors. <i>Rapid Communications in Mass Spectrometry</i> , 2018, 32, 649-656.	0.7	3
23	Monolithic anion-exchange chromatography yields rhinovirus of high purity. <i>Journal of Virological Methods</i> , 2018, 251, 15-21.	1.0	12
24	Mass spectrometry-based investigation of measles and mumps virus proteome. <i>Virology Journal</i> , 2018, 15, 160.	1.4	10
25	Three Different Reactions, One Catalyst: A Cu(I) PNP Pincer Complex as Catalyst for C=C and C=N Cross-Couplings. <i>Organic Letters</i> , 2017, 19, 2178-2181.	2.4	34
26	Manganese-Catalyzed Aminomethylation of Aromatic Compounds with Methanol as a Sustainable C1 Building Block. <i>Journal of the American Chemical Society</i> , 2017, 139, 8812-8815.	6.6	177
27	Microchip capillary gel electrophoresis combined with lectin affinity enrichment employing magnetic beads for glycoprotein analysis. <i>Analytical and Bioanalytical Chemistry</i> , 2017, 409, 6625-6634.	1.9	7
28	Elucidation of oxidation and degradation products of oxygen containing fuel components by combined use of a stable isotopic tracer and mass spectrometry. <i>Analytica Chimica Acta</i> , 2017, 993, 47-54.	2.6	6
29	A bio-inspired method for direct measurement of local wall shear rates with micrometer localization using the multimeric protein von Willebrand factor as sensor molecule. <i>Biomicrofluidics</i> , 2017, 11, 044117.	1.2	2
30	Soft X-ray Radiation Applied in the Analysis of Intact Viruses and Antibodies by Means of Nano Electrospray Differential Mobility Analysis. <i>NATO Science for Peace and Security Series A: Chemistry and Biology</i> , 2017, , 149-157.	0.5	3
31	nES GEMMA Analysis of Lectins and Their Interactions with Glycoproteins – Separation, Detection, and Sampling of Noncovalent Biospecific Complexes. <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 77-86.	1.2	8
32	Oxidation Products of Ester-Based Oils with and without Antioxidants Identified by Stable Isotope Labelling and Mass Spectrometry. <i>Applied Sciences (Switzerland)</i> , 2017, 7, 396.	1.3	12
33	Processed stigmas of <i>Crocus sativus</i> L. imaged by MALDI-based MS. <i>Proteomics</i> , 2016, 16, 1726-1730.	1.3	4
34	Quality control of oligonucleotide synthesis by means of matrix-assisted laser desorption/ionization linear time-of-flight mass spectrometry on a nanocoated disposable target. <i>Rapid Communications in Mass Spectrometry</i> , 2016, 30, 665-668.	0.7	1
35	Polymer-based metal nano-coated disposable target for matrix-assisted and matrix-free laser desorption/ionization mass spectrometry. <i>Methods</i> , 2016, 104, 182-193.	1.9	4
36	Identification of <i>Bremia lactucae</i> and <i>Oidium neolycopersici</i> proteins extracted for intact spore MALDI mass spectrometric biotyping. <i>Electrophoresis</i> , 2016, 37, 2940-2952.	1.3	6

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37	Combining gas-phase electrophoretic mobility molecular analysis (GEMMA), light scattering, field flow fractionation and cryo electron microscopy in a multidimensional approach to characterize liposomal carrier vesicles. <i>International Journal of Pharmaceutics</i> , 2016, 513, 309-318.	2.6	19
38	Nano electrospray gas-phase electrophoretic mobility molecular analysis (nES GEMMA) of liposomes: applicability of the technique for nano vesicle batch control. <i>Analyst, The</i> , 2016, 141, 6042-6050.	1.7	15
39	Sustainable Synthesis of Quinolines and Pyrimidines Catalyzed by Manganese PNP Pincer Complexes. <i>Journal of the American Chemical Society</i> , 2016, 138, 15543-15546.	6.6	300
40	Air Stable Iron(II) PNP Pincer Complexes as Efficient Catalysts for the Selective Alkylation of Amines with Alcohols. <i>Advanced Synthesis and Catalysis</i> , 2016, 358, 3824-3831.	2.1	89
41	Divergent Coupling of Alcohols and Amines Catalyzed by Isoelectronic Hydride Mn ^I and Fe ^{II} PNP Pincer Complexes. <i>Chemistry - A European Journal</i> , 2016, 22, 12316-12320.	1.7	212
42	In vitro RNA release from a human rhinovirus monitored by means of a molecular beacon and chip electrophoresis. <i>Analytical and Bioanalytical Chemistry</i> , 2016, 408, 4209-4217.	1.9	2
43	Co(II) PCP Pincer Complexes as Catalysts for the Alkylation of Aromatic Amines with Primary Alcohols. <i>Organic Letters</i> , 2016, 18, 3462-3465.	2.4	161
44	Identification of mumps virus protein and lipid composition by mass spectrometry. <i>Virology Journal</i> , 2016, 13, 9.	1.4	9
45	Analysis of a Common Cold Virus and Its Subviral Particles by Gas-Phase Electrophoretic Mobility Molecular Analysis and Native Mass Spectrometry. <i>Analytical Chemistry</i> , 2015, 87, 8709-8717.	3.2	37
46	Comprehensive Size-Determination of Whole Virus Vaccine Particles Using Gas-Phase Electrophoretic Mobility Macromolecular Analyzer, Atomic Force Microscopy, and Transmission Electron Microscopy. <i>Analytical Chemistry</i> , 2015, 87, 8657-8664.	3.2	18
47	Collision-induced dissociation of aminophospholipids (PE, MMPE, DMPE, PS): an apparently known fragmentation process revisited. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 5079-5089.	1.9	2
48	Feasibility of the development of reference materials for the detection of Ag nanoparticles in food: neat dispersions and spiked chicken meat. <i>Accreditation and Quality Assurance</i> , 2015, 20, 3-16.	0.4	33
49	Long time storage (archiving) of peptide, protein and tryptic digest samples on disposable nano-coated polymer targets for MALDI MS. <i>EuPA Open Proteomics</i> , 2015, 8, 48-54.	2.5	5
50	A uniform measurement expression for cross method comparison of nanoparticle aggregate size distributions. <i>Analyst, The</i> , 2015, 140, 5257-5267.	1.7	14
51	Challenges of glycoprotein analysis by microchip capillary gel electrophoresis. <i>Electrophoresis</i> , 2015, 36, 1754-1758.	1.3	16
52	Sensitive detection of C-reactive protein in serum by immunoprecipitationâ€“microchip capillary gel electrophoresis. <i>Analytical Biochemistry</i> , 2015, 478, 102-106.	1.1	10
53	Ca ²⁺ concentration-dependent conformational change of FVIII B-domain observed by atomic force microscopy. <i>Analytical and Bioanalytical Chemistry</i> , 2015, 407, 6051-6056.	1.9	11
54	Shear-Dependent Interactions of von Willebrand Factor with Factor VIII and Protease ADAMTS 13 Demonstrated at a Single Molecule Level by Atomic Force Microscopy. <i>Analytical Chemistry</i> , 2015, 87, 10299-10305.	3.2	16

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55	Intact cell mass spectrometry as a progress tracking tool for batch and fed-batch fermentation processes. <i>Analytical Biochemistry</i> , 2015, 470, 25-33.	1.1	4
56	Maldi â€ Massenspektrometrie Und Tandemmassenspektrometrie. , 2015, , 91-94.		0
57	Chip electrophoretic separation of highly homologous ammodytoxin isoforms: Three neurotoxic phospholipases A₂ of <i>Vipera ammodytes ammodytes</i> venom. <i>Electrophoresis</i> , 2014, 35, 2137-2145.	1.3	0
58	Identification of proteins interacting with ammodytoxins in <i>Vipera ammodytes ammodytes</i> venom by immuno-affinity chromatography. <i>Analytical and Bioanalytical Chemistry</i> , 2014, 406, 293-304.	1.9	17
59	Visualization of a protein-protein interaction at a single-molecule level by atomic force microscopy. <i>Analytical and Bioanalytical Chemistry</i> , 2014, 406, 1411-1421.	1.9	12
60	Sizing up large protein complexes by electrospray ionisation-based electrophoretic mobility and native mass spectrometry: morphology selective binding of Fabs to hepatitis B virus capsids. <i>Analytical and Bioanalytical Chemistry</i> , 2014, 406, 1437-1446.	1.9	30
61	Intact cell/intact spore mass spectrometry (IC/ISMS) on polymer-based, nano-coated disposable targets. <i>Molecular and Cellular Probes</i> , 2014, 28, 99-105.	0.9	2
62	Production of reference materials for the detection and size determination of silica nanoparticles in tomato soup. <i>Analytical and Bioanalytical Chemistry</i> , 2014, 406, 3895-907.	1.9	36
63	Ultrahighâ€performance liquid chromatography/electrospray ionization linear ion trap Orbitrap mass spectrometry of antioxidants (amines and phenols) applied in lubricant engineering. <i>Rapid Communications in Mass Spectrometry</i> , 2014, 28, 63-76.	0.7	25
64	Efficient Hydrogenation of Ketones and Aldehydes Catalyzed by Well-Defined Iron(II) PNP Pincer Complexes: Evidence for an Insertion Mechanism. <i>Organometallics</i> , 2014, 33, 6905-6914.	1.1	119
65	Synthesis and Reactivity of Four- and Five-Coordinate Low-Spin Cobalt(II) PCP Pincer Complexes and Some Nickel(II) Analogues. <i>Organometallics</i> , 2014, 33, 6132-6140.	1.1	44
66	Development of a bio-analytical strategy for characterization of vaccine particles combining SEC and nanoES GEMMA. <i>Analyst, The</i> , 2014, 139, 1412-1419.	1.7	14
67	Improved sample preparation for intact cell mass spectrometry (biotyping) of mycelium samples taken from a batch fermentation process of <i>Penicillium chrysogenum</i>. <i>Rapid Communications in Mass Spectrometry</i> , 2014, 28, 957-964.	0.7	4
68	MALDI-TOF Mass Spectrometry Imaging Reveals Molecular Level Changes in Ultrahigh Molecular Weight Polyethylene Joint Implants in Correlation with Lipid Adsorption. <i>Analytical Chemistry</i> , 2014, 86, 9723-9732.	3.2	14
69	Characterization of on-target generated tryptic peptides from <i>Giberella zeae</i> conidia spore proteins by means of matrix-assisted laser desorption/ionization mass spectrometry. <i>Molecular and Cellular Probes</i> , 2014, 28, 91-98.	0.9	4
70	Liquid phase separation of proteins based on electrophoretic effects in an electrospray setup during sample introduction into a gas-phase electrophoretic mobility molecular analyzer (CEâ€GEMMA/CEâ€ESâ€DMA). <i>Analytica Chimica Acta</i> , 2014, 841, 91-98.	2.6	12
71	Analysis of Bio-nanoparticles by Means of Nano ES in Combination with DMA and PDMA: Intact Viruses, Virus-Like-Particles and Vaccine Particles. <i>NATO Science for Peace and Security Series A: Chemistry and Biology</i> , 2014, , 133-147.	0.5	0
72	Comparison of engine oil degradation observed in laboratory alteration and in the engine by chemometric data evaluation. <i>Tribology International</i> , 2013, 65, 37-47.	3.0	28

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73	In-chain neutral hydrocarbon loss from crocin apocarotenoid ester glycosides and the crocetin aglycon (<i>Crocus sativus</i> L.) by ESI-MS (n = 2, 3). <i>Journal of Mass Spectrometry</i> , 2013, 48, 1299-1307.		8
74	Analysis and handling of bio-nanoparticles and environmental nanoparticles using electrostatic aerosol mobility. <i>Particuology</i> , 2013, 11, 14-19.	2.0	25
75	Combining light microscopy, dielectric spectroscopy, MALDI intact cell mass spectrometry, FTIR spectromicroscopy and multivariate data mining for morphological and physiological bioprocess characterization of filamentous organisms. <i>Fungal Genetics and Biology</i> , 2013, 51, 1-11.	0.9	19
76	Characterization of cross-linked gelatin nanoparticles by electrophoretic techniques in the liquid and the gas phase. <i>Electrophoresis</i> , 2013, 34, 3267-3276.	1.3	16
77	Comparative method evaluation for size and size-distribution analysis of gold nanoparticles. <i>Journal of Separation Science</i> , 2013, 36, 2952-2961.	1.3	87
78	Characterization of rhinovirus subviral A particles via capillary electrophoresis, electron microscopy and gas phase electrophoretic mobility molecular analysis: Part II. <i>Electrophoresis</i> , 2013, 34, 1600-1609.	1.3	10
79	Chip electrophoresis of gelatin-based nanoparticles. <i>Electrophoresis</i> , 2013, 34, 2152-2161.	1.3	8
80	Inhibition of extracellular lipase from <i>Streptomyces rimosus</i> with 3,4-dichloroisocoumarin. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2013, 28, 1094-1104.	2.5	3
81	Imaging of a Tribolayer Formed from Ionic Liquids by Laser Desorption/Ionization-Reflection Time-of-Flight Mass Spectrometry. <i>Analytical Chemistry</i> , 2012, 84, 10708-10714.	3.2	13
82	Comparison of different tandem mass spectrometric techniques (ESI-IT, ESI and IP-MALDI-QTOF and) Tj ETQq0 0 0 rgBT /Overl <i>sativus</i> L.. <i>Rapid Communications in Mass Spectrometry</i> , 2012, 26, 670-678.	0.7	24
83	MALDI-based intact spore mass spectrometry of downy and powdery mildews. <i>Journal of Mass Spectrometry</i> , 2012, 47, 978-986.	0.7	21
84	Characterization of rhinovirus subviral A particles via capillary electrophoresis, electron microscopy and gas phase electrophoretic mobility molecular analysis: Part I. <i>Electrophoresis</i> , 2012, 33, 1833-1841.	1.3	23
85	Microchip capillary gel electrophoresis of multiply PEGylated high-molecular-mass glycoproteins. <i>Biotechnology Journal</i> , 2012, 7, 635-641.	1.8	11
86	Thermo-oxidative stability and corrosion properties of ammonium based ionic liquids. <i>Tribology International</i> , 2012, 46, 73-83.	3.0	69
87	Corrosion properties of ammonium based ionic liquids evaluated by SEM-EDX, XPS and ICP-OES. <i>Green Chemistry</i> , 2011, 13, 2869.	4.6	66
88	PEGylated recombinant von Willebrand factor analyzed by means of MALDI-TOF-MS, CGE-on-a-chip and nES-GEMMA. <i>International Journal of Mass Spectrometry</i> , 2011, 305, 157-163.	0.7	6
89	Immunoprecipitation combined with microchip capillary gel electrophoresis: Detection and quantification of β -galactosidase from crude <i>E. coli</i> cell lysate. <i>Biotechnology Journal</i> , 2011, 6, 420-427.	1.8	8
90	Rapid detection of apoptosis in mammalian cells by using intact cell MALDI mass spectrometry. <i>Analyst</i> , 2011, 136, 5181.	1.7	23

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91	A universal product ion nomenclature for $[M\hat{H}]^+$, $[M+H]^+$ and $[M+nNa\hat{(n-1)H}]^+$ ($n=1\hat{3}$) glycerophospholipid precursor ions based on high-energy CID by MALDI-TOF/TOF mass spectrometry. <i>International Journal of Mass Spectrometry</i> , 2011, 301, 90-101.	0.7	9
92	Positive and negative electrospray ionisation travelling wave ion mobility mass spectrometry and low-energy collision-induced dissociation of sialic acid derivatives. <i>Rapid Communications in Mass Spectrometry</i> , 2011, 25, 3235-3244.	0.7	7
93	CID of singly charged antioxidants applied in lubricants by means of a 3D ion trap and a linear ion trap Orbitrap mass spectrometer. <i>Journal of Mass Spectrometry</i> , 2011, 46, 517-528.	0.7	12
94	LDI and ESI MS as well as low energy CID of a self-assembling nanorod-forming fullerene derivative. <i>Journal of Mass Spectrometry</i> , 2011, 46, 1108-1114.	0.7	1
95	Different target surfaces for the analysis of peptides, peptide mixtures and peptide mass fingerprints by AP-MALDI ion trap-mass spectrometry. <i>Journal of Proteomics</i> , 2011, 74, 975-981.	1.2	8
96	Mass spectrometry – One of the pillars of proteomics. <i>Journal of Proteomics</i> , 2011, 74, 915-919.	1.2	4
97	Identification and characterization of organic nanoparticles in food. <i>TrAC - Trends in Analytical Chemistry</i> , 2011, 30, 100-112.	5.8	84
98	Parallel differential mobility analysis for electrostatic characterization and manipulation of nanoparticles and viruses. <i>TrAC - Trends in Analytical Chemistry</i> , 2011, 30, 123-132.	5.8	13
99	Intact Cell/Spore Mass Spectrometry of <i>Fusarium</i> Macro Conidia for Fast Isolate and Species Differentiation. <i>NATO Science for Peace and Security Series A: Chemistry and Biology</i> , 2011, , 47-63.	0.5	4
100	A fluorescent derivatization method of proteins for the detection of low-level impurities by microchip capillary gel electrophoresis. <i>Electrophoresis</i> , 2010, 31, 611-617.	1.3	15
101	Molecular weight determination of high molecular mass (glyco)proteins using CGE-on-chip, planar SDS-PAGE and MALDI-TOF-MS. <i>Electrophoresis</i> , 2010, 31, 3850-3862.	1.3	7
102	Diamond-like carbon coated polymer-based targets in microscope slide format for MALDI mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2010, 45, 566-569.	0.7	7
103	MALDI linear TOF mass spectrometry of PEGylated (glyco)proteins. <i>Journal of Mass Spectrometry</i> , 2010, 45, 612-617.	0.7	24
104	GEMMA and MALDI-TOF MS of reactive PEGs for pharmaceutical applications. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2010, 52, 432-437.	1.4	20
105	A comparison of nano-electrospray gas-phase electrophoretic mobility macromolecular analysis and matrix-assisted laser desorption/ionization linear time-of-flight mass spectrometry for the characterization of the recombinant coagulation glycoprotein von W. <i>Rapid Communications in Mass Spectrometry</i> , 2010, 24, 761-767.	0.7	9
106	Mixed volume sample preparation method for intact cell mass spectrometry of <i>Fusarium</i> spores. <i>Journal of Mass Spectrometry</i> , 2009, 44, 1622-1624.	0.7	15
107	Analysis of antioxidants in insulation cladding of copper wire: a comparison of different mass spectrometric techniques (ESI-IT, MALDI-TOF and TOF-SIMS). <i>Journal of Mass Spectrometry</i> , 2009, 44, 1724-1732.	0.7	10
108	Development of a MALDI two-layer volume sample preparation technique for analysis of colored conidia spores of <i>Fusarium</i> by MALDI linear TOF mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2009, 395, 1373-1383.	1.9	51

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109	Evaluation of matrix-assisted laser desorption/ionization (MALDI) preparation techniques for surface characterization of intact <i>Fusarium</i> spores by MALDI linear time-of-flight mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2009, 23, 877-884.	0.7	75
110	Analysis of human plasma lipids and soybean lecithin by means of high-performance thin-layer chromatography and matrix-assisted laser desorption/ionization mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2009, 23, 2711-2723.	0.7	41
111	Electrospray ionization and atmospheric pressure matrix-assisted laser desorption/ionization mass spectrometry of antioxidants applied in lubricants. <i>Rapid Communications in Mass Spectrometry</i> , 2009, 23, 3917-3927.	0.7	17
112	The renaissance of high-energy CID for structural elucidation of complex lipids: MALDI-TOF/RTOF-MS of alkali cationized triacylglycerols. <i>Journal of the American Society for Mass Spectrometry</i> , 2009, 20, 1037-1047.	1.2	93
113	Comparing standard and microwave assisted staining protocols for SDS-PAGE of glycoproteins followed by subsequent PMF with MALDI MS. <i>Journal of Proteomics</i> , 2009, 72, 628-639.	1.2	12
114	Improved identification of hordeins by cysteine alkylation with 2-bromoethylamine, SDS-PAGE and subsequent in-gel tryptic digestion. <i>Journal of Mass Spectrometry</i> , 2009, 44, 1613-1621.	0.7	10
115	The impact of tyrosine kinase 2 (Tyk2) on the proteome of murine macrophages and their response to lipopolysaccharide (LPS). <i>Proteomics</i> , 2008, 8, 3469-3485.	1.3	13
116	Comparing the applicability of CGE-on-chip and SDS-PAGE for fast pre-screening of mouse serum samples prior to proteomics analysis. <i>Electrophoresis</i> , 2008, 29, 4332-4340.	1.3	7
117	Nano ES GEMMA and PDMA, new tools for the analysis of nanobioparticles' Protein complexes, lipoparticles, and viruses. <i>Journal of the American Society for Mass Spectrometry</i> , 2008, 19, 1062-1068.	1.2	61
118	MALDI Seamless Postsource Decay Fragment Ion Analysis of Sodiated and Lithiated Phospholipids. <i>Analytical Chemistry</i> , 2008, 80, 1664-1678.	3.2	38
119	Gas-Phase Electrophoretic Molecular Mobility Analysis of Size and Stoichiometry of Complexes of a Common Cold Virus with Antibody and Soluble Receptor Molecules. <i>Analytical Chemistry</i> , 2008, 80, 2261-2264.	3.2	40
120	Biological Variation of the Platelet Proteome in the Elderly Population and Its Implication for Biomarker Research. <i>Molecular and Cellular Proteomics</i> , 2008, 7, 193-203.	2.5	71
121	Mass spectrometry of proteinous allergens inducing human diseases. , 2008, , 459-485.		0
122	Comparison of various nano-differential mobility analysers (nDMAs) applying globular proteins. <i>Journal of Experimental Nanoscience</i> , 2007, 2, 291-301.	1.3	26
123	Mass spectrometric evidence of covalently-bound tetrahydrolipstatin at the catalytic serine of <i>Streptomyces rimosus</i> lipase. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2007, 1770, 163-170.	1.1	15
124	Determination of Molecular Weight, Particle Size, and Density of High Number Generation PAMAM Dendrimers Using MALDI-TOF-MS and nES-GEMMA. <i>Macromolecules</i> , 2007, 40, 5599-5605.	2.2	81
125	Characterization of the bga1-encoded glycoside hydrolase family 35- β -galactosidase of <i>Hypocrea jecorina</i> with galacto-1 ² -d-galactanase activity. <i>FEBS Journal</i> , 2007, 274, 1691-1700.	2.2	31
126	Isolation of esterified fatty acids bound to serum albumin purified from human plasma and characterised by MALDI mass spectrometry. <i>Biologicals</i> , 2007, 35, 43-49.	0.5	23

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127	Molecular weight determination of ultra-high mass compounds on a standard matrix-assisted laser desorption/ionization time-of-flight mass spectrometer: PAMAM dendrimer generation 10 and immunoglobulin M. <i>Rapid Communications in Mass Spectrometry</i> , 2006, 20, 3803-3806.	0.7	20
128	Characterization of moenomycin antibiotic complex by multistage MALDI-IT/RTOF-MS and ESI-IT-MS. <i>Journal of the American Society for Mass Spectrometry</i> , 2006, 17, 1081-1090.	1.2	21
129	Comparison of CID spectra of singly charged polypeptide antibiotic precursor ions obtained by positive-ion vacuum MALDI IT/RTOF and TOF/RTOF, AP-MALDI-IT and ESI-IT mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2006, 41, 421-447.	0.7	43
130	Characterization of N- and O-glycopeptides of recombinant human erythropoietins as potential biomarkers for doping analysis by means of microscale sample purification combined with MALDI-TOF and quadrupole IT/RTOF mass spectrometry. <i>Journal of Separation Science</i> , 2005, 28, 1764-1778.	1.3	50
131	The influence of nonspecific cleavage sites on identification of low molecular mass proteins by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry with seamless post-source decay fragment ion analysis. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 79-82.	0.7	12
132	Characterisation of intact recombinant human erythropoietins applied in doping by means of planar gel electrophoretic techniques and matrix-assisted laser desorption/ionisation linear time-of-flight mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 728-742.	0.7	52
133	A new approach in proteomics of wheat gluten: combining chymotrypsin cleavage and matrix-assisted laser desorption/ionization quadrupole ion trap reflectron tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 2725-2728.	0.7	35
134	Characterization of cysteinylated pharmaceutical-grade human serum albumin by electrospray ionization mass spectrometry and low-energy collision-induced dissociation tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2005, 19, 2965-2973.	0.7	51
135	Instrumental Parameters in the MALDI-TOF Mass Spectrometric Analysis of Quaternary Protein Structures. <i>Analytical Chemistry</i> , 2005, 77, 103-110.	3.2	34
136	Determination of glycopeptide structures by multistage mass spectrometry with low-energy collision-induced dissociation: comparison of electrospray ionization quadrupole ion trap and matrix-assisted laser desorption/ionization quadrupole ion trap reflectron time-of-flight approaches. <i>Rapid Communications in Mass Spectrometry</i> , 2004, 18, 1575-1582.	0.7	89
137	Ultraviolet matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of intact hemoglobin complex from whole human blood. <i>Rapid Communications in Mass Spectrometry</i> , 2004, 18, 1932-1938.	0.7	20
138	Exact molecular mass determination of various forms of native and de-N-glycosylated human plasma-derived antithrombin by means of electrospray ionization ion trap mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2004, 39, 1429-1436.	0.7	9
139	Characterization of covalently inhibited extracellular lipase from <i>Streptomyces rimosus</i> by matrix-assisted laser desorption/ionization time-of-flight and matrix-assisted laser desorption/ionization quadrupole ion trap reflectron time-of-flight mass spectrometry: localization of the active site serine. <i>Journal of Mass Spectrometry</i> , 2004, 39, 1474-1483.	0.7	13
140	Transferrin binding and transferrin-mediated cellular uptake of the ruthenium coordination compound KP1019, studied by means of AAS, ESI-MS and CD spectroscopy. <i>Journal of Analytical Atomic Spectrometry</i> , 2004, 19, 46.	1.6	183
141	Characterization of antithrombin III from human plasma by two-dimensional gel electrophoresis and capillary electrophoretic methods. <i>Electrophoresis</i> , 2003, 24, 4282-4290.	1.3	23
142	Selective solid-phase isolation of methionine-containing peptides and subsequent matrix-assisted laser desorption/ionisation mass spectrometric detection of methionine- and of methionine-sulfoxide-containing peptides. <i>Rapid Communications in Mass Spectrometry</i> , 2003, 17, 1815-1824.	0.7	23
143	Investigation of sample preparation and instrumental parameters in the matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of noncovalent peptide/peptide complexes. <i>Rapid Communications in Mass Spectrometry</i> , 2003, 17, 1931-1940.	0.7	27
144	Characterisation of castor oil by on-line and off-line non-aqueous reverse-phase high-performance liquid chromatography-mass spectrometry (APCI and UV/MALDI). <i>Phytochemical Analysis</i> , 2003, 14, 337-346.	1.2	35

#	ARTICLE	IF	CITATIONS
145	Molecular mass determination of plasma-derived glycoproteins by ultraviolet matrix-assisted laser desorption/ionization time-of-flight mass spectrometry with internal calibration. <i>Journal of Mass Spectrometry</i> , 2002, 37, 1118-1130.	0.7	26
146	One-way hydrophobic surface foil for UV matrix-assisted laser desorption/ionization mass spectrometry of peptides. <i>Rapid Communications in Mass Spectrometry</i> , 2002, 16, 899-902.	0.7	18
147	Negative and positive ion matrix-assisted laser desorption/ionization time-of-flight mass spectrometry and positive ion nano-electrospray ionization quadrupole ion trap mass spectrometry of peptidoglycan fragments isolated from various <i>Bacillus</i> species. <i>Journal of Mass Spectrometry</i> , 2001, 36, 124-139.	0.7	24
148	Charge-reduced nano electrospray ionization combined with differential mobility analysis of peptides, proteins, glycoproteins, noncovalent protein complexes and viruses. <i>Journal of Mass Spectrometry</i> , 2001, 36, 1038-1052.	0.7	202
149	Structural analysis of <i>Bacillus megaterium</i> KM spore peptidoglycan and its dynamics during germination. <i>Microbiology (United Kingdom)</i> , 1999, 145, 1033-1041.	0.7	34
150	Time-delayed extraction matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of polyacrylonitrile and other synthetic polymers with the matrix 4-hydroxybenzylidene malononitrile. <i>Rapid Communications in Mass Spectrometry</i> , 1998, 12, 1344-1350.	0.7	17
151	MALDI mass spectrometry of biomolecules and synthetic polymers using alkali hexacyanoferrate (II) complexes and glycerol as matrix. <i>International Journal of Mass Spectrometry and Ion Processes</i> , 1997, 169-170, 99-109.	1.9	23
152	Characterization of Calixarenes by Positive- and Negative-ion Californium-252 Plasma Desorption Mass Spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 1997, 11, 427-432.	0.7	1
153	Sample preparation for the analysis of glycerophospholipids by matrix-assisted positive and negative ion ²⁵² Cf plasma desorption time-of-flight mass spectrometry. <i>European Journal of Mass Spectrometry</i> , 1996, 2, 247.	0.7	5
154	K4[Fe(CN)6]/Glycerol A New Liquid Matrix System for Matrix-assisted Laser Desorption/Ionization Mass Spectrometry of Hydrophobic Compounds. <i>Rapid Communications in Mass Spectrometry</i> , 1996, 10, 1278-1282.	0.7	33
155	Characterization of Peptidoglycan Trimers after Gel Chromatography and Reversed-phase High-performance Liquid Chromatography by Positive-ion Plasma Desorption Mass Spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 1996, 10, 1956-1960.	0.7	4
156	Characterization of braunâ€™s lipoprotein and determination of its attachment sites to peptidoglycan by ²⁵² Cf-PD and MALDI time-of-flight mass spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 1995, 6, 892-905.	1.2	14
157	Structural characterization of the cyanelle peptidoglycan of <i>Cyanophora paradoxa</i> by ²⁵² Cf plasma desorption mass spectrometry and fast atom bombardment/tandem mass spectrometry. <i>Biological Mass Spectrometry</i> , 1993, 22, 524-536.	0.5	27
158	Molecular weight-determination of biosynthetically modified monomeric and oligomeric muropeptides from <i>Escherichia coli</i> by plasma desorption-mass spectrometry. <i>FEBS Letters</i> , 1993, 316, 181-185.	1.3	11
159	Electron impact ionization mass spectrometry and tandem mass spectrometry of phenylalkylpyridazines. <i>Organic Mass Spectrometry</i> , 1991, 26, 595-600.	1.3	5
160	Characterization of synthetic N-acetylcysteine conjugates by positive- and negative-ion ²⁵² Cf plasma desorption mass spectrometry. <i>Organic Mass Spectrometry</i> , 1991, 26, 1065-1073.	1.3	2
161	Electron ionization mass spectrometry of phenylpyridazinylmethanols and phenylpyridazinyl ketones. <i>Rapid Communications in Mass Spectrometry</i> , 1991, 5, 421-424.	0.7	1
162	Positive and negative ion ²⁵² Cf plasma desorption mass spectrometry of polar agrochemical metabolites. <i>Biological Mass Spectrometry</i> , 1990, 19, 75-79.	0.5	2