Günter Allmaier

List of Publications by Year in descending order

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162 papers 4,503 citations

32 h-index 59 g-index

164 all docs

164 docs citations

164 times ranked 4910 citing authors

#	Article	IF	Citations
1	Sustainable Synthesis of Quinolines and Pyrimidines Catalyzed by Manganese PNP Pincer Complexes. Journal of the American Chemical Society, 2016, 138, 15543-15546.	6.6	300
2	Divergent Coupling of Alcohols and Amines Catalyzed by Isoelectronic Hydride Mn ^I and Fe ^{II} PNP Pincer Complexes. Chemistry - A European Journal, 2016, 22, 12316-12320.	1.7	212
3	Charge-reduced nano electrospray ionization combined with differential mobility analysis of peptides, proteins, glycoproteins, noncovalent protein complexes and viruses. Journal of Mass Spectrometry, 2001, 36, 1038-1052.	0.7	202
4	Transferrin binding and transferrin-mediated cellular uptake of the ruthenium coordination compound KP1019, studied by means of AAS, ESI-MS and CD spectroscopy. Journal of Analytical Atomic Spectrometry, 2004, 19, 46.	1.6	183
5	Manganese-Catalyzed Aminomethylation of Aromatic Compounds with Methanol as a Sustainable C1 Building Block. Journal of the American Chemical Society, 2017, 139, 8812-8815.	6.6	177
6	Co(II) PCP Pincer Complexes as Catalysts for the Alkylation of Aromatic Amines with Primary Alcohols. Organic Letters, 2016, 18, 3462-3465.	2.4	161
7	Efficient Hydrogenation of Ketones and Aldehydes Catalyzed by Well-Defined Iron(II) PNP Pincer Complexes: Evidence for an Insertion Mechanism. Organometallics, 2014, 33, 6905-6914.	1.1	119
8	The renaissance of high-energy CID for structural elucidation of complex lipids: MALDI-TOF/RTOF-MS of alkali cationized triacylglycerols. Journal of the American Society for Mass Spectrometry, 2009, 20, 1037-1047.	1.2	93
9	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. Rapid Communications in Mass Spectrometry. 2004. 18. 1575-1582.	0.7	89
10	Air Stable Iron(II) PNP Pincer Complexes as Efficient Catalysts for the Selective Alkylation of Amines with Alcohols. Advanced Synthesis and Catalysis, 2016, 358, 3824-3831.	2.1	89
11	Comparative method evaluation for size and sizeâ€distribution analysis of gold nanoparticles. Journal of Separation Science, 2013, 36, 2952-2961.	1.3	87
12	Identification and characterization of organic nanoparticles in food. TrAC - Trends in Analytical Chemistry, 2011, 30, 100-112.	5.8	84
13	Determination of Molecular Weight, Particle Size, and Density of High Number Generation PAMAM Dendrimers Using MALDIâ^'TOFâ^'MS and nESâ^'GEMMA. Macromolecules, 2007, 40, 5599-5605.	2.2	81
14	Evaluation of matrixâ€assisted laser desorption/ionization (MALDI) preparation techniques for surface characterization of intact <i>Fusarium</i> spores by MALDI linear timeâ€ofâ€flight mass spectrometry. Rapid Communications in Mass Spectrometry, 2009, 23, 877-884.	0.7	75
15	Biological Variation of the Platelet Proteome in the Elderly Population and Its Implication for Biomarker Research. Molecular and Cellular Proteomics, 2008, 7, 193-203.	2.5	71
16	Thermo-oxidative stability and corrosion properties of ammonium based ionic liquids. Tribology International, 2012, 46, 73-83.	3.0	69
17	Corrosion properties of ammonium based ionic liquids evaluated by SEM-EDX, XPS and ICP-OES. Green Chemistry, 2011, 13, 2869.	4.6	66
18	Nanoscale chemical imaging of individual chemotherapeutic cytarabine-loaded liposomal nanocarriers. Nano Research, 2019, 12, 197-203.	5.8	65

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19	Nano ES GEMMA and PDMA, new tools for the analysis of nanobioparticlesâ€"Protein complexes, lipoparticles, and viruses. Journal of the American Society for Mass Spectrometry, 2008, 19, 1062-1068.	1.2	61
20	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. Rapid Communications in Mass Spectrometry, 2005, 19, 728-742.	0.7	52
21	Characterization of cysteinylation of pharmaceutical-grade human serum albumin by electrospray ionization mass spectrometry and low-energy collision-induced dissociation tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2005, 19, 2965-2973.	0.7	51
22	Development of a MALDI two-layer volume sample preparation technique for analysis of colored conidia spores of Fusarium by MALDI linear TOF mass spectrometry. Analytical and Bioanalytical Chemistry, 2009, 395, 1373-1383.	1.9	51
23	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. Journal of Separation Science, 2005, 28, 1764-1778.	1.3	50
24	Synthesis and Reactivity of Four- and Five-Coordinate Low-Spin Cobalt(II) PCP Pincer Complexes and Some Nickel(II) Analogues. Organometallics, 2014, 33, 6132-6140.	1.1	44
25	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. Journal of Mass Spectrometry, 2006, 41, 421-447.	0.7	43
26	Analysis of human plasma lipids and soybean lecithin by means of highâ€performance thinâ€layer chromatography and matrixâ€assisted laser desorption/ionization mass spectrometry. Rapid Communications in Mass Spectrometry, 2009, 23, 2711-2723.	0.7	41
27	Gas-Phase Electrophoretic Molecular Mobility Analysis of Size and Stoichiometry of Complexes of a Common Cold Virus with Antibody and Soluble Receptor Molecules. Analytical Chemistry, 2008, 80, 2261-2264.	3.2	40
28	MALDI Seamless Postsource Decay Fragment Ion Analysis of Sodiated and Lithiated Phospholipids. Analytical Chemistry, 2008, 80, 1664-1678.	3.2	38
29	Analysis of a Common Cold Virus and Its Subviral Particles by Gas-Phase Electrophoretic Mobility Molecular Analysis and Native Mass Spectrometry. Analytical Chemistry, 2015, 87, 8709-8717.	3.2	37
30	Production of reference materials for the detection and size determination of silica nanoparticles in tomato soup. Analytical and Bioanalytical Chemistry, 2014, 406, 3895-907.	1.9	36
31	Characterisation of castor oil by on-line and off-line non-aqueous reverse-phase high-performance liquid chromatography-mass spectrometry (APCI and UV/MALDI). Phytochemical Analysis, 2003, 14, 337-346.	1.2	35
32	A new approach in proteomics of wheat gluten: combining chymotrypsin cleavage and matrix-assisted laser desorption/ionization quadrupole ion trap reflectron tandem mass spectrometry. Rapid Communications in Mass Spectrometry, 2005, 19, 2725-2728.	0.7	35
33	Instrumental Parameters in the MALDI-TOF Mass Spectrometric Analysis of Quaternary Protein Structures. Analytical Chemistry, 2005, 77, 103-110.	3.2	34
34	Three Different Reactions, One Catalyst: A Cu(I) PNP Pincer Complex as Catalyst for C–C and C–N Cross-Couplings. Organic Letters, 2017, 19, 2178-2181.	2.4	34
35	Structural analysis of Bacillus megaterium KM spore peptidoglycan and its dynamics during germination. Microbiology (United Kingdom), 1999, 145, 1033-1041.	0.7	34
36	K4[Fe(CN)6]/Glycerolâ€"A New Liquid Matrix System for Matrix-assisted Laser Desorption/Ionization Mass Spectrometry of Hydrophobic Compounds. Rapid Communications in Mass Spectrometry, 1996, 10, 1278-1282.	0.7	33

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37	Feasibility of the development of reference materials for the detection of Ag nanoparticles in food: neat dispersions and spiked chicken meat. Accreditation and Quality Assurance, 2015, 20, 3-16.	0.4	33
38	Support effect on the reactivity and stability of Au25(SR)18 and Au144(SR)60 nanoclusters in liquid phase cyclohexane oxidation. Catalysis Today, 2019, 336, 174-185.	2.2	33
39	Characterization of the bga1-encoded glycoside hydrolase family 35â€∫β-galactosidase of Hypocrea jecorina with galacto-β-d-galactanase activity. FEBS Journal, 2007, 274, 1691-1700.	2.2	31
40	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. Analytical and Bioanalytical Chemistry, 2014, 406, 1437-1446.	1.9	30
41	Optimization of MALDI-TOF mass spectrometry imaging for the visualization and comparison of peptide distributions in dry-cured ham muscle fibers. Food Chemistry, 2019, 283, 275-286.	4.2	30
42	Comparison of engine oil degradation observed in laboratory alteration and in the engine by chemometric data evaluation. Tribology International, 2013, 65, 37-47.	3.0	28
43	Virus-like particle size and molecular weight/mass determination applying gas-phase electrophoresis (native nES GEMMA). Analytical and Bioanalytical Chemistry, 2019, 411, 5951-5962.	1.9	28
44	Structural characterization of the cyanelle peptidoglycan of Cyanophora paradoxa by 252Cf plasma desorption mass spectrometry and fast atom bombardment/tandem mass spectrometry. Biological Mass Spectrometry, 1993, 22, 524-536.	0.5	27
45	Investigation of sample preparation and instrumental parameters in the matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of noncovalent peptide/peptide complexes. Rapid Communications in Mass Spectrometry, 2003, 17, 1931-1940.	0.7	27
46	Molecular mass determination of plasma-derived glycoproteins by ultraviolet matrix-assisted laser desorption/ionization time-of-flight mass spectrometry with internal calibration. Journal of Mass Spectrometry, 2002, 37, 1118-1130.	0.7	26
47	Comparison of various nano-differential mobility analysers (nDMAs) applying globular proteins. Journal of Experimental Nanoscience, 2007, 2, 291-301.	1.3	26
48	Analysis and handling of bio-nanoparticles and environmental nanoparticles using electrostatic aerosol mobility. Particuology, 2013, 11, 14-19.	2.0	25
49	Ultrahighâ€performance liquid chromatography/electrospray ionization linear ion trap Orbitrap mass spectrometry of antioxidants (amines and phenols) applied in lubricant engineering. Rapid Communications in Mass Spectrometry, 2014, 28, 63-76.	0.7	25
50	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 variousBacillusspecies. Journal of Mass Spectrometry, 2001, 36, 124-139.	0.7	24
51	MALDI linear TOF mass spectrometry of PEGylated (glyco)proteins. Journal of Mass Spectrometry, 2010, 45, 612-617.	0.7	24
52	Comparison of different tandem mass spectrometric techniques (ESIâ€IT, ESIâ€and IPâ€MALDIâ€QRTOF and) Tj I sativus i> L Rapid Communications in Mass Spectrometry, 2012, 26, 670-678.	ETQq0 0 0 0.7	rgBT /Overl 24
53	MALDI mass spectrometry of biomolecules and synthetic polymers using alkali hexacyanoferrate (II) complexes and glycerol as matrix. International Journal of Mass Spectrometry and Ion Processes, 1997, 169-170, 99-109.	1.9	23
54	Characterization of antithrombin III from human plasma by two-dimensional gel electrophoresis and capillary electrophoretic methods. Electrophoresis, 2003, 24, 4282-4290.	1.3	23

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55	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. Rapid Communications in Mass Spectrometry, 2003, 17, 1815-1824.	0.7	23
56	Isolation of esterified fatty acids bound to serum albumin purified from human plasma and characterised by MALDI mass spectrometry. Biologicals, 2007, 35, 43-49.	0.5	23
57	Rapid detection of apoptosis in mammalian cells by using intact cell MALDI mass spectrometry. Analyst, The, 2011, 136, 5181.	1.7	23
58	Characterization of rhinovirus subviral <scp>A</scp> particles via capillary electrophoresis, electron microscopy and gasâ€phase electrophoretic mobility molecular analysis: Part I. Electrophoresis, 2012, 33, 1833-1841.	1.3	23
59	Characterization of moenomycin antibiotic complex by multistage MALDI-IT/RTOF-MS and ESI-IT-MS. Journal of the American Society for Mass Spectrometry, 2006, 17, 1081-1090.	1.2	21
60	MALDIâ€based intact spore mass spectrometry of downy and powdery mildews. Journal of Mass Spectrometry, 2012, 47, 978-986.	0.7	21
61	Comparative Analysis of Platelet-Derived Extracellular Vesicles Using Flow Cytometry and Nanoparticle Tracking Analysis. International Journal of Molecular Sciences, 2021, 22, 3839.	1.8	21
62	Ultraviolet matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of intact hemoglobin complex from whole human blood. Rapid Communications in Mass Spectrometry, 2004, 18, 1932-1938.	0.7	20
63	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. Rapid Communications in Mass Spectrometry, 2006, 20, 3803-3806.	0.7	20
64	GEMMA and MALDI-TOF MS of reactive PEGs for pharmaceutical applications. Journal of Pharmaceutical and Biomedical Analysis, 2010, 52, 432-437.	1.4	20
65	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. Fungal Genetics and Biology, 2013, 51, 1-11.	0.9	19
66	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. International Journal of Pharmaceutics, 2016, 513, 309-318.	2.6	19
67	Ligand engineering of immobilized nanoclusters on surfaces: ligand exchange reactions with supported Au ₁₁ (PPh ₃) ₇ Br ₃ . Nanoscale, 2020, 12, 12809-12816.	2.8	19
68	One-way hydrophobic surface foil for UV matrix-assisted laser desorption/ionization mass spectrometry of peptides. Rapid Communications in Mass Spectrometry, 2002, 16, 899-902.	0.7	18
69	Comprehensive Size-Determination of Whole Virus Vaccine Particles Using Gas-Phase Electrophoretic Mobility Macromolecular Analyzer, Atomic Force Microscopy, and Transmission Electron Microscopy. Analytical Chemistry, 2015, 87, 8657-8664.	3.2	18
70	Time-delayed extraction matrix-assisted laser desorption/ionization time-of-flight mass spectrometry of polyacrylnitrile and other synthetic polymers with the matrixÂ4-hydroxybenzylidene malononitrile. Rapid Communications in Mass Spectrometry, 1998, 12, 1344-1350.	0.7	17
71	Electrospray ionization and atmospheric pressure matrixâ€assisted laser desorption/ionization mass spectrometry of antioxidants applied in lubricants. Rapid Communications in Mass Spectrometry, 2009, 23, 3917-3927.	0.7	17
72	Identification of proteins interacting with ammodytoxins in Vipera ammodytes ammodytes venom by immuno-affinity chromatography. Analytical and Bioanalytical Chemistry, 2014, 406, 293-304.	1.9	17

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73	Characterization of crossâ€linked gelatin nanoparticles by electrophoretic techniques in the liquid and the gas phase. Electrophoresis, 2013, 34, 3267-3276.	1.3	16
74	Challenges of glycoprotein analysis by microchip capillary gel electrophoresis. Electrophoresis, 2015, 36, 1754-1758.	1.3	16
75	Shear-Dependent Interactions of von Willebrand Factor with Factor VIII and Protease ADAMTS 13 Demonstrated at a Single Molecule Level by Atomic Force Microscopy. Analytical Chemistry, 2015, 87, 10299-10305.	3.2	16
76	Mass spectrometric evidence of covalently-bound tetrahydrolipstatin at the catalytic serine of Streptomyces rimosus lipase. Biochimica Et Biophysica Acta - General Subjects, 2007, 1770, 163-170.	1.1	15
77	Mixed volume sample preparation method for intact cell mass spectrometry of Fusarium spores. Journal of Mass Spectrometry, 2009, 44, 1622-1624.	0.7	15
78	A fluorescent derivatization method of proteins for the detection of lowâ€level impurities by microchip capillary gel electrophoresis. Electrophoresis, 2010, 31, 611-617.	1.3	15
79	Nano electrospray gas-phase electrophoretic mobility molecular analysis (nES GEMMA) of liposomes: applicability of the technique for nano vesicle batch control. Analyst, The, 2016, 141, 6042-6050.	1.7	15
80	N-terminal VP1 Truncations Favor T = 1 Norovirus-Like Particles. Vaccines, 2021, 9, 8.	2.1	15
81	Characterization of braun's lipoprotein and determination of its attachment sites to peptidoglycan by 252Cf-PD and MALDI time-of-flight mass spectrometry. Journal of the American Society for Mass Spectrometry, 1995, 6, 892-905.	1.2	14
82	Development of a bio-analytical strategy for characterization of vaccine particles combining SEC and nanoES GEMMA. Analyst, The, 2014, 139, 1412-1419.	1.7	14
83	MALDI-TOF Mass Spectrometry Imaging Reveals Molecular Level Changes in Ultrahigh Molecular Weight Polyethylene Joint Implants in Correlation with Lipid Adsorption. Analytical Chemistry, 2014, 86, 9723-9732.	3.2	14
84	A uniform measurement expression for cross method comparison of nanoparticle aggregate size distributions. Analyst, The, 2015, 140, 5257-5267.	1.7	14
85	Native Nano-electrospray Differential Mobility Analyzer (nES GEMMA) Enables Size Selection of Liposomal Nanocarriers Combined with Subsequent Direct Spectroscopic Analysis. Analytical Chemistry, 2019, 91, 3860-3868.	3.2	14
86	Characterization of covalently inhibited extracellular lipase fromStreptomyces rimosus 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. Journal of Mass Spectrometry, 2004, 39, 1474-1483.	0.7	13
87	The impact of tyrosine kinase 2 (Tyk2) on the proteome of murine macrophages and their response to lipopolysaccharide (LPS). Proteomics, 2008, 8, 3469-3485.	1.3	13
88	Parallel differential mobility analysis for electrostatic characterization and manipulation of nanoparticles and viruses. TrAC - Trends in Analytical Chemistry, 2011, 30, 123-132.	5.8	13
89	Imaging of a Tribolayer Formed from Ionic Liquids by Laser Desorption/Ionization-Reflectron Time-of-Flight Mass Spectrometry. Analytical Chemistry, 2012, 84, 10708-10714.	3.2	13
90	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. Rapid Communications in Mass Spectrometry, 2005, 19, 79-82.	0.7	12

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91	Comparing standard and microwave assisted staining protocols for SDS-PAGE of glycoproteins followed by subsequent PMF with MALDI MS. Journal of Proteomics, 2009, 72, 628-639.	1.2	12
92	CID of singly charged antioxidants applied in lubricants by means of a 3D ion trap and a linear ion trap–Orbitrap mass spectrometer. Journal of Mass Spectrometry, 2011, 46, 517-528.	0.7	12
93	Visualization of a protein-protein interaction at a single-molecule level by atomic force microscopy. Analytical and Bioanalytical Chemistry, 2014, 406, 1411-1421.	1.9	12
94	Liquid phase separation of proteins based on electrophoretic effects in an electrospray setup during sample introduction into a gas-phase electrophoretic mobility molecular analyzer (CE–GEMMA/CE–ES–DMA). Analytica Chimica Acta, 2014, 841, 91-98.	2.6	12
95	Oxidation Products of Ester-Based Oils with and without Antioxidants Identified by Stable Isotope Labelling and Mass Spectrometry. Applied Sciences (Switzerland), 2017, 7, 396.	1.3	12
96	Size and molecular weight determination of polysaccharides by means of nano electrospray gasâ€phase electrophoretic mobility molecular analysis (nES GEMMA). Electrophoresis, 2018, 39, 1142-1150.	1.3	12
97	Monolithic anion-exchange chromatography yields rhinovirus of high purity. Journal of Virological Methods, 2018, 251, 15-21.	1.0	12
98	Molecular weight-determination of biosynthetically modified monomeric and oligomeric muropeptides fromEscherichia coliby plasma desorption-mass spectrometry. FEBS Letters, 1993, 316, 181-185.	1.3	11
99	Microchip capillary gel electrophoresis of multiply PEGylated highâ€molecularâ€mass glycoproteins. Biotechnology Journal, 2012, 7, 635-641.	1.8	11
100	Ca2+ concentration-dependent conformational change of FVIII B-domain observed by atomic force microscopy. Analytical and Bioanalytical Chemistry, 2015, 407, 6051-6056.	1.9	11
101	Calcium ion effect on phospholipid bilayers as cell membrane analogues. Bioelectrochemistry, 2022, 143, 107988.	2.4	11
102	Analysis of antioxidants in insulation cladding of copper wire: a comparison of different mass spectrometric techniques (ESI–IT, MALDI–RTOF and RTOF–SIMS). Journal of Mass Spectrometry, 2009, 44, 1724-1732.	0.7	10
103	Improved identification of hordeins by cysteine alkylation with 2â€bromoethylamine, SDSâ€PAGE and subsequent ⟨i⟩inâ€gel⟨/i⟩ tryptic digestion. Journal of Mass Spectrometry, 2009, 44, 1613-1621.	0.7	10
104	Characterization of rhinovirus subviral A particles via capillary electrophoresis, electron microscopy and gas phase electrophoretic mobility molecular analysis: Part II. Electrophoresis, 2013, 34, 1600-1609.	1.3	10
105	Sensitive detection of C-reactive protein in serum by immunoprecipitation–microchip capillary gel electrophoresis. Analytical Biochemistry, 2015, 478, 102-106.	1.1	10
106	Mass spectrometry-based investigation of measles and mumps virus proteome. Virology Journal, 2018, 15, 160.	1.4	10
107	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. Journal of Mass Spectrometry, 2004, 39, 1429-1436.	0.7	9
108	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. Rapid Communications in Mass Spectrometry, 2010, 24, 761-767.	0.7	9

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109	A universal product ion nomenclature for [Mâ^'H]â^', [M+H]+ and [M+nNaâ^'(nâ^'1)H]+ (n=1–3) glycerophospholipid precursor ions based on high-energy CID by MALDI-TOF/RTOF mass spectrometry. International Journal of Mass Spectrometry, 2011, 301, 90-101.	0.7	9
110	Identification of mumps virus protein and lipid composition by mass spectrometry. Virology Journal, 2016, 13, 9.	1.4	9
111	Bipolar Corona Discharge-Based Charge Equilibration for Nano Electrospray Gas-Phase Electrophoretic Mobility Molecular Analysis of Bio- and Polymer Nanoparticles. Analytical Chemistry, 2020, 92, 8665-8669.	3.2	9
112	Immunoprecipitation combined with microchip capillary gel electrophoresis: Detection and quantification of βâ€galactosidase from crude ⟨i⟩E. coli⟨/i⟩ cell lysate. Biotechnology Journal, 2011, 6, 420-427.	1.8	8
113	Different target surfaces for the analysis of peptides, peptide mixtures and peptide mass fingerprints by AP-MALDI ion trap-mass spectrometry. Journal of Proteomics, 2011, 74, 975-981.	1.2	8
114	Inâ€chain neutral hydrocarbon loss from crocin apocarotenoid ester glycosides and the crocetin aglycon (<i>Crocus sativus</i> L.) by ESIâ€MS ⁿ (n = 2, 3). Journal of Mass Spectrometry, 2048, 1299-1307.	b37	8
115	Chip electrophoresis of gelatinâ€based nanoparticles. Electrophoresis, 2013, 34, 2152-2161.	1.3	8
116	nES GEMMA Analysis of Lectins and Their Interactions with Glycoproteins – Separation, Detection, and Sampling of Noncovalent Biospecific Complexes. Journal of the American Society for Mass Spectrometry, 2017, 28, 77-86.	1.2	8
117	Comparing the applicability of CGEâ€onâ€theâ€chip and SDSâ€PAGE for fast preâ€screening of mouse serum samples prior to proteomics analysis. Electrophoresis, 2008, 29, 4332-4340.	1.3	7
118	Molecular weight determination of high molecular mass (glyco)proteins using CGEâ€onâ€aâ€chip, planar SDSâ€PAGE and MALDIâ€TOFâ€MS. Electrophoresis, 2010, 31, 3850-3862.	1.3	7
119	Diamondâ€like carbon coated polymerâ€based targets in microscope slide format for MALDI mass spectrometry. Journal of Mass Spectrometry, 2010, 45, 566-569.	0.7	7
120	Positive and negative electrospray ionisation travelling wave ion mobility mass spectrometry and lowâ€energy collisionâ€induced dissociation of sialic acid derivatives. Rapid Communications in Mass Spectrometry, 2011, 25, 3235-3244.	0.7	7
121	Microchip capillary gel electrophoresis combined with lectin affinity enrichment employing magnetic beads for glycoprotein analysis. Analytical and Bioanalytical Chemistry, 2017, 409, 6625-6634.	1.9	7
122	Adeno-associated Virus Virus-like Particle Characterization via Orthogonal Methods: Nanoelectrospray Differential Mobility Analysis, Asymmetric Flow Field-Flow Fractionation, and Atomic Force Microscopy. ACS Omega, 2021, 6, 16428-16437.	1.6	7
123	PEGylated recombinant von Willebrand factor analyzed by means of MALDI-TOF-MS, CGE-on-a-chip and nES-GEMMA. International Journal of Mass Spectrometry, 2011, 305, 157-163.	0.7	6
124	Identification of <i>Bremia lactucae</i> and <i>Oidium neolycopersici</i> proteins extracted for intact spore MALDI mass spectrometric biotyping. Electrophoresis, 2016, 37, 2940-2952.	1.3	6
125	Elucidation of oxidation and degradation products of oxygen containing fuel components by combined use of a stable isotopic tracer and mass spectrometry. Analytica Chimica Acta, 2017, 993, 47-54.	2.6	6
126	Development of an accelerated artificial ageing method for the characterization of degradation products of antioxidants in lubricants by mass spectrometry. European Journal of Mass Spectrometry, 2019, 25, 300-323.	0.5	6

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127	Inâ€depth analysis of crocetin ester glycosides from dried/processed stigmas of <scp><i>Crocus sativus</i></scp> L. by HPLCâ€ESIâ€MS ^{<i>n</i>} (<i>n</i> > (<i>n</i> > 3). Phytochemical Analysis, 2019, 30, 346-356.	1.2	6
128	Molecular Weight Determination of Adenoâ€Associate Virus serotype 8 Virusâ€like Particle either carrying or lacking genome via native nES Gasâ€phase Electrophoretic Molecular Mobility Analysis (GEMMA) and nESI QRTOF Mass Spectrometry. Journal of Mass Spectrometry, 2021, 56, e4786.	0.7	6
129	Electron impact ionization mass spectrometry and tandem mass spectrometry of phenylalkylpyridazines. Organic Mass Spectrometry, 1991, 26, 595-600.	1.3	5
130	Sample preparation for the analysis of glycerophospholipids by matrix-assisted positive and negative ion 252Cf plasma desorption time-of-flight mass spectrometry. European Journal of Mass Spectrometry, 1996, 2, 247.	0.7	5
131	Long time storage (archiving) of peptide, protein and tryptic digest samples on disposable nano-coated polymer targets for MALDI MS. EuPA Open Proteomics, 2015, 8, 48-54.	2.5	5
132	Characterization of Peptidoglycan Trimers after Gel Chromatography and Reversed-phase High-performance Liquid Chromatography by Positive-ion Plasma Desorption Mass Spectrometry. Rapid Communications in Mass Spectrometry, 1996, 10, 1956-1960.	0.7	4
133	Mass spectrometry â€" One of the pillars of proteomics. Journal of Proteomics, 2011, 74, 915-919.	1.2	4
134	Improved sample preparation for intact cell mass spectrometry (biotyping) of mycelium samples taken from a batch fermentation process of <i>Penicillium chrysogenum</i> . Rapid Communications in Mass Spectrometry, 2014, 28, 957-964.	0.7	4
135	Characterization of on-target generated tryptic peptides from Giberella zeae conidia spore proteins by means of matrix-assisted laser desorption/ionization mass spectrometry. Molecular and Cellular Probes, 2014, 28, 91-98.	0.9	4
136	Intact cell mass spectrometry as a progress tracking tool for batch and fed-batch fermentation processes. Analytical Biochemistry, 2015, 470, 25-33.	1.1	4
137	Processed stigmas of <i>Crocus sativus</i> L. imaged by MALDIâ€based MS. Proteomics, 2016, 16, 1726-1730.	1.3	4
138	Polymer-based metal nano-coated disposable target for matrix-assisted and matrix-free laser desorption/ionization mass spectrometry. Methods, 2016, 104, 182-193.	1.9	4
139	Nano electrospray differential mobility analysis based size-selection of liposomes and very-low density lipoprotein particles for offline hyphenation to MALDI mass spectrometry. Journal of Pharmaceutical and Biomedical Analysis, 2020, 179, 112998.	1.4	4
140	Intact Cell/Spore Mass Spectrometry of Fusarium Macro Conidia for Fast Isolate and Species Differentiation. NATO Science for Peace and Security Series A: Chemistry and Biology, 2011, , 47-63.	0.5	4
141	Atmospheric pressure matrixâ€assisted laser desorption/ionization mass spectrometry of engine oil additive components. Rapid Communications in Mass Spectrometry, 2022, 36, e9271.	0.7	4
142	Inhibition of extracellular lipase from Streptomyces rimosus with 3,4-dichloroisocoumarin. Journal of Enzyme Inhibition and Medicinal Chemistry, 2013, 28, 1094-1104.	2.5	3
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