List of Publications by Year in descending order

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DETE NOVAK

#	Article	IF	CITATIONS
1	Early-onset pulmonary and cutaneous vasculitis driven by constitutively active SRC-family kinase HCK. Journal of Allergy and Clinical Immunology, 2022, 149, 1464-1472.e3.	1.5	10
2	Hydroxyl radical footprinting analysis of a human haptoglobin-hemoglobin complex. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2022, 1870, 140735.	1.1	8
3	In Vitro Evolution Reveals Noncationic Protein–RNA Interaction Mediated by Metal Ions. Molecular Biology and Evolution, 2022, 39, .	3.5	13
4	Utilization of Fast Photochemical Oxidation of Proteins and Both Bottom-up and Top-down Mass Spectrometry for Structural Characterization of a Transcription Factor–dsDNA Complex. Analytical Chemistry, 2022, 94, 3203-3210.	3.2	10
5	Top-Down Detection of Oxidative Protein Footprinting by Collision-Induced Dissociation, Electron-Transfer Dissociation, and Electron-Capture Dissociation. Analytical Chemistry, 2022, 94, 9993-10002.	3.2	8
6	Motif orientation matters: Structural characterization of TEAD1 recognition of genomic DNA. Structure, 2021, 29, 345-356.e8.	1.6	2
7	Impact of parasitic lifestyle and different types of centromere organization on chromosome and genome evolution in the plant genus <i>Cuscuta</i> . New Phytologist, 2021, 229, 2365-2377.	3.5	22
8	Complex sequence organization of heterochromatin in the holocentric plant Cuscuta europaea elucidated by the computational analysis of nanopore reads. Computational and Structural Biotechnology Journal, 2021, 19, 2179-2189.	1.9	9
9	SAP domain forms a flexible part of DNA aperture in Ku70/80. FEBS Journal, 2021, 288, 4382-4393.	2.2	13
10	Early modification of cytochrome c by hydrogen peroxide triggers its fast degradation. International Journal of Biological Macromolecules, 2021, 174, 413-423.	3.6	7
11	LinX: A Software Tool for Uncommon Cross-Linking Chemistry. Journal of Proteome Research, 2021, 20, 2021-2027.	1.8	5
12	Benefits of Ion Mobility Separation and Parallel Accumulation–Serial Fragmentation Technology on timsTOF Pro for the Needs of Fast Photochemical Oxidation of Protein Analysis. ACS Omega, 2021, 6, 10352-10361.	1.6	18
13	Molecular Interactions Driving Intermediate Filament Assembly. Cells, 2021, 10, 2457.	1.8	22
14	Chitinase Chit62J4 Essential for Chitin Processing by Human Microbiome Bacterium Clostridium paraputrificum J4. Molecules, 2021, 26, 5978.	1.7	5
15	Studying Protein–DNA Interactions by Hydrogen/Deuterium Exchange Mass Spectrometry. Methods in Molecular Biology, 2021, 2247, 193-219.	0.4	6
16	Fast Fluoroalkylation of Proteins Uncovers the Structure and Dynamics of Biological Macromolecules. Journal of the American Chemical Society, 2021, 143, 20670-20679.	6.6	13
17	Characterization of repeat arrays in ultraâ€long nanopore reads reveals frequent origin of satellite DNA from retrotransposonâ€derived tandem repeats. Plant Journal, 2020, 101, 484-500.	2.8	76
18	Bacteriocin ASM1 is an O / S â€diglycosylated, plasmidâ€encoded homologue of glycocin F. FEBS Letters, 2020, 594, 1196-1206.	1.3	10

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19	Repeat-sequence turnover shifts fundamentally in species with large genomes. Nature Plants, 2020, 6, 1325-1329.	4.7	87
20	Molecular Mechanism of LEDGF/p75 Dimerization. Structure, 2020, 28, 1288-1299.e7.	1.6	4
21	Global analysis of repetitive DNA from unassembled sequence reads using RepeatExplorer2. Nature Protocols, 2020, 15, 3745-3776.	5.5	144
22	Three-Dimensional Printed Target Plates for Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry. Analytical Chemistry, 2020, 92, 12783-12788.	3.2	3
23	Chromosome-scale genome assembly for the duckweed Spirodela intermedia, integrating cytogenetic maps, PacBio and Oxford Nanopore libraries. Scientific Reports, 2020, 10, 19230.	1.6	23
24	Targeting ERK-Hippo Interplay in Cancer Therapy. International Journal of Molecular Sciences, 2020, 21, 3236.	1.8	17
25	Photoinduced damage of AsLOV2 domain is accompanied by increased singlet oxygen production due to flavin dissociation. Scientific Reports, 2020, 10, 4119.	1.6	10
26	Addressing the Molecular Mechanism of Longitudinal Lamin Assembly Using Chimeric Fusions. Cells, 2020, 9, 1633.	1.8	10
27	Influence of cross-linker polarity on selectivity towards lysine side chains. Journal of Proteomics, 2020, 218, 103716.	1.2	5
28	Extraordinary Sequence Diversity and Promiscuity of Centromeric Satellites in the Legume Tribe Fabeae. Molecular Biology and Evolution, 2020, 37, 2341-2356.	3.5	42
29	Differential Genome Size and Repetitive DNA Evolution in Diploid Species of Melampodium sect. Melampodium (Asteraceae). Frontiers in Plant Science, 2020, 11, 362.	1.7	37
30	Binding of eIF3 in complex with eIF5 and eIF1 to the 40S ribosomal subunit is accompanied by dramatic structural changes. Nucleic Acids Research, 2019, 47, 8282-8300.	6.5	20
31	Ergochromes: Heretofore Neglected Side of Ergot Toxicity. Toxins, 2019, 11, 439.	1.5	11
32	Reductantâ€Induced Free Radical Fluoroalkylation of Nitrogen Heterocycles and Innate Aromatic Amino Acid Residues in Peptides and Proteins. Chemistry - A European Journal, 2019, 25, 15779-15785.	1.7	23
33	A reference genome for pea provides insight into legume genome evolution. Nature Genetics, 2019, 51, 1411-1422.	9.4	363
34	MS-Based Approaches Enable the Structural Characterization of Transcription Factor/DNA Response Element Complex. Biomolecules, 2019, 9, 535.	1.8	9
35	The Dark Matter of Large Cereal Genomes: Long Tandem Repeats. International Journal of Molecular Sciences, 2019, 20, 2483.	1.8	19
36	Oligomeric Architecture of Mouse Activating Nkrp1 Receptors on Living Cells. International Journal of Molecular Sciences, 2019, 20, 1884.	1.8	11

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37	Genome invasion by a hypomethylated satellite repeat in Australian crucifer Ballantinia antipoda. Plant Journal, 2019, 99, 1066-1079.	2.8	3
38	First Community-Wide, Comparative Cross-Linking Mass Spectrometry Study. Analytical Chemistry, 2019, 91, 6953-6961.	3.2	100
39	Nondisjunction and unequal spindle organization accompany the drive of <i>Aegilops speltoides</i> B chromosomes. New Phytologist, 2019, 223, 1340-1352.	3.5	26
40	The MEK-ERK-MST1 Axis Potentiates the Activation of the Extrinsic Apoptotic Pathway during GDC-0941 Treatment in Jurkat T Cells. Cells, 2019, 8, 191.	1.8	8
41	Proteases Immobilization for In Situ Time-Limited Proteolysis on MALDI Chips. Catalysts, 2019, 9, 833.	1.6	2
42	The C-type lectin-like receptor Nkrp1b: Structural proteomics reveals features affecting protein conformation and interactions. Journal of Proteomics, 2019, 196, 162-172.	1.2	4
43	Thiopurine intolerance-causing mutations in NUDT15 induce temperature-dependent destabilization of the catalytic site. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2019, 1867, 376-381.	1.1	10
44	Evolutionary history of ergot with a new infrageneric classification (Hypocreales: Clavicipitaceae:) Tj ETQq0 0 0 r	gBT_/Over	lock_10 Tf 50
45	Relapsed acute lymphoblastic leukemia-specific mutations in NT5C2 cluster into hotspots driving intersubunit stimulation. Leukemia, 2018, 32, 1393-1403.	3.3	27
46	Dating the Species Network: Allopolyploidy and Repetitive DNA Evolution in American Daisies (Melampodium sect. Melampodium, Asteraceae). Systematic Biology, 2018, 67, 1010-1024.	2.7	54
47	Impact of Chemical Cross-Linking on Protein Structure and Function. Analytical Chemistry, 2018, 90, 1104-1113.	3.2	44
48	Crystal structure of native βâ€ <i>N</i> â€acetylhexosaminidase isolated from <i>AspergillusÂoryzae</i> sheds light onto its substrate specificity, high stability, and regulation by propeptide. FEBS Journal, 2018, 285, 580-598.	2.2	12
49	Detection and Quantification of Carbohydrate-Deficient Transferrin by MALDI-Compatible Protein Chips Prepared by Ambient Ion Soft Landing. Clinical Chemistry, 2018, 64, 1319-1326.	1.5	14
50	Affinity switching of the LEDGF/p75 IBD interactome is governed by kinase-dependent phosphorylation. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E7053-E7062.	3.3	27
51	Structural basis of the interaction between the putative adhesion-involved and iron-regulated FrpD and FrpC proteins of Neisseria meningitidis. Scientific Reports, 2017, 7, 40408.	1.6	10
52	The conserved tyrosine residue 940 plays a key structural role in membrane interaction of Bordetella adenylate cyclase toxin. Scientific Reports, 2017, 7, 9330.	1.6	18
53	High-level expression and purification of soluble form of human natural killer cell receptor NKR-P1 in HEK293S GnTlâ^' cells. Protein Expression and Purification, 2017, 140, 36-43.	0.6	7
54	Changes in the expression of N- and O-glycopeptides in patients with colorectal cancer and hepatocellular carcinoma quantified by full-MS scan FT-ICR and multiple reaction monitoring. Journal of Proteomics, 2017, 153, 44-52.	1.2	26

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55	The cyanobacterial metabolite nocuolin a is a natural oxadiazine that triggers apoptosis in human cancer cells. PLoS ONE, 2017, 12, e0172850.	1.1	43
56	Reprogramming of leukemic cell metabolism through the naphthoquinonic compound Quambalarine B. Oncotarget, 2017, 8, 103137-103153.	0.8	6
57	A community proposal to integrate proteomics activities in ELIXIR. F1000Research, 2017, 6, 875.	0.8	13
58	Solution structure of the lymphocyte receptor Nkrp1a reveals a distinct conformation of the long loop region as compared to in the crystal structure. Proteins: Structure, Function and Bioinformatics, 2016, 84, 1304-1311.	1.5	7
59	Vitamin B2 as a virulence factor in Pseudogymnoascus destructans skin infection. Scientific Reports, 2016, 6, 33200.	1.6	46
60	Protein Chips Compatible with MALDI Mass Spectrometry Prepared by Ambient Ion Landing. Analytical Chemistry, 2016, 88, 8526-8534.	3.2	14
61	Quambalarine B, a Secondary Metabolite from <i>Quambalaria cyanescens</i> with Potential Anticancer Properties. Journal of Natural Products, 2016, 79, 2304-2314.	1.5	9
62	Shikonin regulates C-MYC and GLUT1 expression through the MST1-YAP1-TEAD1 axis. Experimental Cell Research, 2016, 349, 273-281.	1.2	22
63	Oligomeric interface modulation causes misregulation of purine 5´-nucleotidase in relapsed leukemia. BMC Biology, 2016, 14, 91.	1.7	9
64	Large-scale identification of membrane proteins based on analysis of trypsin-protected transmembrane segments. Journal of Proteomics, 2016, 149, 15-22.	1.2	12
65	1H, 13C and 15N resonance assignments of human DCL-1 (CD302) extracellular domain. Biomolecular NMR Assignments, 2016, 10, 189-192.	0.4	3
66	Planar Functionalized Surfaces for Direct Immunoaffinity Desorption/Ionization Mass Spectrometry. Clinical Chemistry, 2016, 62, 270-278.	1.5	18
67	Pharmacological inhibition of fatty-acid oxidation synergistically enhances the effect of l-asparaginase in childhood ALL cells. Leukemia, 2016, 30, 209-218.	3.3	31
68	Highâ€ŧhroughput workflow for identification of phosphorylated peptides by LCâ€MALDIâ€TOF/TOFâ€MS coupled to <i>in situ</i> enrichment on MALDI plates functionalized by ion landing. Journal of Mass Spectrometry, 2015, 50, 802-811.	0.7	8
69	Nkrp1 Family, from Lectins to Protein Interacting Molecules. Molecules, 2015, 20, 3463-3478.	1.7	6
70	Biologically Active Metabolites Produced by the Basidiomycete Quambalaria cyanescens. PLoS ONE, 2015, 10, e0118913.	1.1	20
71	Mapping protein structural changes by quantitative cross-linking. Methods, 2015, 89, 112-120.	1.9	27
72	Interaction of <i>Bordetella</i> adenylate cyclase toxin with complement receptor 3 involves multivalent glycan binding. FEBS Letters, 2015, 589, 374-379.	1.3	29

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73	Cytotoxic Lipopeptide Muscotoxin A, Isolated from Soil Cyanobacterium <i>Desmonostoc muscorum</i> , Permeabilizes Phospholipid Membranes by Reducing Their Fluidity. Chemical Research in Toxicology, 2015, 28, 216-224.	1.7	16
74	Expression and purification of soluble and stable ectodomain of natural killer cell receptor LLT1 through high-density transfection of suspension adapted HEK293S GnTIâ^' cells. Protein Expression and Purification, 2015, 109, 7-13.	0.6	18
75	Paxillin-dependent regulation of <i>IGF2</i> / <i>H19</i> gene cluster expression. Journal of Cell Science, 2015, 128, 3106-16.	1.2	20
76	Hydnocarpin-Type Flavonolignans: Semisynthesis and Inhibitory Effects onStaphylococcus aureusBiofilm Formation. Journal of Natural Products, 2015, 78, 2095-2103.	1.5	16
77	PSTPIP2, a Protein Associated with Autoinflammatory Disease, Interacts with Inhibitory Enzymes SHIP1 and Csk. Journal of Immunology, 2015, 195, 3416-3426.	0.4	34
78	Lincosamide Synthetase—A Unique Condensation System Combining Elements of Nonribosomal Peptide Synthetase and Mycothiol Metabolism. PLoS ONE, 2015, 10, e0118850.	1.1	27
79	Recombinant Expression, In Vitro Refolding and Characterizing Disulfide Bonds of a Mouse Inhibitory C-Type Lectin-Like Receptor Nkrp1b. Physiological Research, 2015, 64, S85-S93.	0.4	2
80	Target antigens for Hs-14 monoclonal antibody and their various expression in normozoospermic and asthenozoospermic men. Basic and Clinical Andrology, 2015, 25, 11.	0.8	2
81	Target antigens for Hs-14 monoclonal antibody and their various expression in normozoospermic and asthenozoospermic men. Basic and Clinical Andrology, 2015, 25, 11.	0.8	3
82	The Application of an Emerging Technique for Protein–Protein Interaction Interface Mapping: The Combination of Photo-Initiated Cross-Linking Protein Nanoprobes with Mass Spectrometry. International Journal of Molecular Sciences, 2014, 15, 9224-9241.	1.8	8
83	Systemic AL amyloidosis with unusual cutaneous presentation unmasked by carotenoderma. Amyloid: the International Journal of Experimental and Clinical Investigation: the Official Journal of the International Society of Amyloidosis, 2014, 21, 57-61.	1.4	3
84	Molecular dynamics comparison of E. coli WrbA apoprotein and holoprotein. Journal of Molecular Modeling, 2014, 20, 2400.	0.8	1
85	Role of the EF-hand-like Motif in the 14-3-3 Protein-mediated Activation of Yeast Neutral Trehalase Nth1. Journal of Biological Chemistry, 2014, 289, 13948-13961.	1.6	23
86	SKIP and BIR-1/Survivin have potential to integrate proteome status with gene expression. Journal of Proteomics, 2014, 110, 93-106.	1.2	1
87	Antibacterial properties of lucifensin in Lucilia sericata maggots after septic injury. Asian Pacific Journal of Tropical Biomedicine, 2014, 4, 358-361.	0.5	13
88	L-Asparaginase Causes Metabolic Reprogramming in ALL Cells. Blood, 2014, 124, 922-922.	0.6	1
89	Carbohydrate synthesis and biosynthesis technologies for cracking of the glycan code: Recent advances. Biotechnology Advances, 2013, 31, 17-37.	6.0	14

90 Protein Extraction and Precipitation. , 2013, , 79-90.

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91	Novel Aeruginosinâ€865 from <i>Nostoc</i> sp. as a Potent Antiâ€inflammatory Agent. ChemBioChem, 2013, 14, 2329-2337.	1.3	30
92	Structural basis of the 14-3-3 protein-dependent activation of yeast neutral trehalase Nth1. Biochimica Et Biophysica Acta - General Subjects, 2013, 1830, 4491-4499.	1.1	34
93	Re-evaluation of the involvement of NK cells and C-type lectin-like NK receptors in modulation of immune responses by multivalent GlcNAc-terminated oligosaccharides. Immunology Letters, 2013, 156, 110-117.	1.1	9
94	New insight into the role of a base in the mechanism of imine transfer hydrogenation on a Ru(ii) half-sandwich complex. Dalton Transactions, 2013, 42, 5174.	1.6	27
95	Structural Model of Lymphocyte Receptor NKR-P1C Revealed by Mass Spectrometry and Molecular Modeling. Analytical Chemistry, 2013, 85, 1597-1604.	3.2	19
96	1.2â€Ã resolution crystal structure of <i>Escherichia coli</i> WrbA holoprotein. Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 1748-1757.	2.5	6
97	Lincomycin Biosynthesis Involves a Tyrosine Hydroxylating Heme Protein of an Unusual Enzyme Family. PLoS ONE, 2013, 8, e79974.	1.1	24
98	Mouse Clr-g, a Ligand for NK Cell Activation Receptor NKR-P1F: Crystal Structure and Biophysical Properties. Journal of Immunology, 2012, 189, 4881-4889.	0.4	21
99	Chemical Cross-Linking and H/D Exchange for Fast Refinement of Protein Crystal Structure. Analytical Chemistry, 2012, 84, 867-870.	3.2	30
100	The Cyanobacterial Cyclic Lipopeptides Puwainaphycins F/G Are Inducing Necrosis via Cell Membrane Permeabilization and Subsequent Unusual Actin Relocalization. Chemical Research in Toxicology, 2012, 25, 1203-1211.	1.7	30
101	The combination of hydrogen/deuterium exchange or chemical cross-linking techniques with mass spectrometry: Mapping of human 14-3-3ζ homodimer interface. Journal of Structural Biology, 2012, 179, 10-17.	1.3	19
102	<i>Inâ€situ</i> enrichment of phosphopeptides on MALDI plates modified by ambient ion landing. Journal of Mass Spectrometry, 2012, 47, 1294-1302.	0.7	21
103	Preparation of soluble isotopically labeled NKp30, a human natural cytotoxicity receptor, for structural studies using NMR. Protein Expression and Purification, 2012, 86, 142-150.	0.6	1
104	Specific Nuclear Localizing Sequence Directs Two Myosin Isoforms to the Cell Nucleus in Calmodulin-Sensitive Manner. PLoS ONE, 2012, 7, e30529.	1.1	44
105	Mapping of interaction between cytochrome P450 2B4 and cytochrome b5: the first evidence of two mutual orientations. Neuroendocrinology Letters, 2012, 33 Suppl 3, 41-7.	0.2	4
106	Poly[ <i>N</i> -(2-hydroxypropyl)methacrylamide]-Based Tissue-Embedding Medium Compatible with MALDI Mass Spectrometry Imaging Experiments. Analytical Chemistry, 2011, 83, 5458-5462.	3.2	48
107	New polyene macrolide family produced by submerged culture of Streptomyces durmitorensis. Journal of Antibiotics, 2011, 64, 717-722.	1.0	17
108	Molecular architecture of mouse activating NKR-P1 receptors. Journal of Structural Biology, 2011, 175, 434-441.	1.3	34

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109	High-level expression of soluble form of mouse natural killer cell receptor NKR-P1C(B6) in Escherichia coli. Protein Expression and Purification, 2011, 77, 178-184.	0.6	19
110	Cysteine <i>S</i> -glycosylation, a new post-translational modification found in glycopeptide bacteriocins. FEBS Letters, 2011, 585, 645-650.	1.3	132
111	Structural Basis for the 14-3-3 Protein-dependent Inhibition of the Regulator of G Protein Signaling 3 (RGS3) Function. Journal of Biological Chemistry, 2011, 286, 43527-43536.	1.6	25
112	Spatial Distribution of Glycerophospholipids in the Ocular Lens. PLoS ONE, 2011, 6, e19441.	1.1	23
113	Dissociation of Nystatin and Amphotericin Analogues: Characterisation of Minor Anti-Fungal Macrolides. European Journal of Mass Spectrometry, 2010, 16, 73-80.	0.5	4
114	Visualizing spatial lipid distribution in porcine lens by MALDI imaging high-resolution mass spectrometry. Journal of Lipid Research, 2010, 51, 2295-2302.	2.0	50
115	A New Type of Membrane Raft-Like Microdomains and Their Possible Involvement in TCR Signaling. Journal of Immunology, 2010, 184, 3689-3696.	0.4	37
116	The Â-galactosidase type A gene aglA from Aspergillus niger encodes a fully functional Â-N-acetylgalactosaminidase. Glycobiology, 2010, 20, 1410-1419.	1.3	9
117	<i>mMass</i> 3: A Cross-Platform Software Environment for Precise Analysis of Mass Spectrometric Data. Analytical Chemistry, 2010, 82, 4648-4651.	3.2	697
118	Characterization of Pseudacyclins Aâ^'E, a Suite of Cyclic Peptides Produced by <i>Pseudallescheria boydii</i> . Journal of Natural Products, 2010, 73, 1027-1032.	1.5	29
119	High-throughput analysis of tetracycline antibiotics and their epimers in liquid hog manure using Ultra Performance Liquid Chromatography with UV detection. Chemosphere, 2010, 78, 353-359.	4.2	46
120	Laser Desorption-Ionization of Lipid Transfers: Tissue Mass Spectrometry Imaging without MALDI Matrix. Analytical Chemistry, 2010, 82, 4994-4997.	3.2	78
121	Effective Removal of Nonionic Detergents in Protein Mass Spectrometry, Hydrogen/Deuterium Exchange, and Proteomics. Analytical Chemistry, 2010, 82, 5107-5116.	3.2	63
122	Utilization of highâ€accuracy FTICRâ€MS data in protein quantitation experiments. Journal of Mass Spectrometry, 2009, 44, 1565-1570.	0.7	4
123	Modified electrophoretic and digestion conditions allow a simplified mass spectrometric evaluation of disulfide bonds. Journal of Mass Spectrometry, 2009, 44, 1571-1578.	0.7	31
124	Hydroxylated anthraquinones produced by Geosmithia species. Folia Microbiologica, 2009, 54, 179-187.	1.1	23
125	Automated Ambient Desorptionâ~lonization Platform for Surface Imaging Integrated with a Commercial Fourier Transform Ion Cyclotron Resonance Mass Spectrometer. Analytical Chemistry, 2009, 81, 8479-8487.	3.2	67
126	LC MALDI-TOF MS/MS and LC ESI FTMS analyses of HLA-B27 associated peptides isolated from peripheral blood cells. Immunology Letters, 2008, 116, 79-85.	1.1	7

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127	Liquid chromatography–tandem mass spectrometry characterization of ergocristam degradation products. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2008, 873, 165-172.	1.2	8
128	Abnormal expression and processing of uromodulin in Fabry disease reflects tubular cell storage alteration and is reversible by enzyme replacement therapy. Journal of Inherited Metabolic Disease, 2008, 31, 508-517.	1.7	25
129	Production of (+)-globulol needle crystals on the surface mycelium of Quambalaria cyanescens. Folia Microbiologica, 2008, 53, 15-22.	1.1	10
130	Proteomic analysis of wheat proteins recognized by IgE antibodies of allergic patients. Proteomics, 2008, 8, 1677-1691.	1.3	81
131	Characterization of DESIâ€FTICR mass spectrometry—from ECD to accurate mass tissue analysis. Journal of Mass Spectrometry, 2008, 43, 196-203.	0.7	27
132	Singleâ€ <b>s</b> tep affinity purification of recombinant proteins using a selfâ€excising module from <i>Neisseria meningitidis</i> FrpC. Protein Science, 2008, 17, 1834-1843.	3.1	31
133	Soluble recombinant CD69 receptors optimized to have an exceptional physical and chemical stability display prolonged circulation and remain intact in the blood of mice. FEBS Journal, 2008, 275, 5589-5606.	2.2	26
134	153-P: Analysis of HLA-B27 peptides form healthy individuals and ankylosing spodylitis patients. Human Immunology, 2008, 69, S84.	1.2	0
135	Proteomic analysis uncovers a metabolic phenotype in C. elegans after nhr-40 reduction of function. Biochemical and Biophysical Research Communications, 2008, 374, 49-54.	1.0	5
136	Application of Silicon Nanowires and Indium Tin Oxide Surfaces in Desorption Electrospray Ionization. European Journal of Mass Spectrometry, 2008, 14, 391-399.	0.5	15
137	Intra-Molecular Cross-Linking of Acidic Residues for Protein Structure Studies. European Journal of Mass Spectrometry, 2008, 14, 355-365.	0.5	67
138	Increased expression of secretory actin-binding protein on human spermatozoa is associated with poor semen quality. Human Reproduction, 2007, 22, 1396-1404.	0.4	24
139	Chemical Cross-Linking and Mass Spectrometry as Structure Determination Tools. European Journal of Mass Spectrometry, 2007, 13, 105-113.	0.5	23
140	Monitoring Conformational Changes in Protein Complexes Using Chemical Cross-Linking and Fourier Transform Ion Cyclotron Resonance Mass Spectrometry: The Effect of Calcium Binding on the Calmodulin—Melittin Complex. European Journal of Mass Spectrometry, 2007, 13, 281-290.	0.5	10
141	Synthesis and Structural Study of Organoantimony(III) and Organobismuth(III) Triflates and Cations Containing O,C,O-Pincer Type Ligandsâ€. Organometallics, 2007, 26, 2911-2917.	1.1	53
142	Large Propeptides of Fungal β-N-Acetylhexosaminidases Are Novel Enzyme Regulators That Must Be Intracellularly Processed to Control Activity, Dimerization, and Secretion into the Extracellular Environmentâ€. Biochemistry, 2007, 46, 2719-2734.	1.2	23
143	Structure of the dimeric N-glycosylated form of fungal β-N-acetylhexosaminidase revealed by computer modeling, vibrational spectroscopy, and biochemical studies. BMC Structural Biology, 2007, 7, 32.	2.3	24
144	Structure and dynamics of dark-state bovine rhodopsin revealed by chemical cross-linking and high-resolution mass spectrometry. Protein Science, 2006, 15, 1303-1317.	3.1	50

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145	Separation of nuclear protein complexes by blue native polyacrylamide gel electrophoresis. Electrophoresis, 2006, 27, 1277-1287.	1.3	28
146	Mass spectrometric analysis of the glycosphingolipid-enriched microdomains of rat natural killer cells. Proteomics, 2005, 5, 113-122.	1.3	31
147	Unambiguous Assignment of Intramolecular Chemical Cross-Links in Modified Mammalian Membrane Proteins by Fourier Transform-Tandem Mass Spectrometry. Analytical Chemistry, 2005, 77, 5101-5106.	3.2	40
148	Heat Shock Protein 90 Recognized as an Iron-Binding Protein Associated with the Plasma Membrane of HeLa Cells. Cellular Physiology and Biochemistry, 2004, 14, 41-46.	1.1	17
149	Lymphocyte activation receptors: new structural paradigms in group V of C-type animal lectins. Biochemical Society Transactions, 2004, 32, 1124-1126.	1.6	10
150	A Top-down method for the determination of residue-specific solvent accessibility in proteins. Journal of Mass Spectrometry, 2004, 39, 322-328.	0.7	61
151	Strategy for selective chemical cross-linking of tyrosine and lysine residues. Journal of the American Society for Mass Spectrometry, 2004, 15, 1604-1611.	1.2	52
152	Mass spectrometry is a powerful tool for identification of proteins associated with lipid rafts of Jurkat T-cell line. Biochemical Society Transactions, 2004, 32, 777-779.	1.6	3
153	Metabolism of Carcinogenic 2-Nitroanisole in Rat, Rabbit, Porcine and Human Hepatic Cytosol. Collection of Czechoslovak Chemical Communications, 2004, 69, 589-602.	1.0	12
154	A Top-Down Approach to Protein Structure Studies Using Chemical Cross-Linking and Fourier Transform Mass Spectrometry. European Journal of Mass Spectrometry, 2003, 9, 623-631.	0.5	46
155	Nuclear DNA helicase II is recruited to IFN-α–activated transcription sites at PML nuclear bodies. Journal of Cell Biology, 2002, 158, 463-473.	2.3	50
156	Non–T Cell Activation Linker (NTAL). Journal of Experimental Medicine, 2002, 196, 1617-1626.	4.2	192
157	Monitoring ofin vitrodeamidation of gliadin peptic fragment by mass spectrometry may reflect one of the molecular mechanisms taking place in celiac disease development. Journal of Mass Spectrometry, 2002, 37, 507-511.	0.7	11
158	A proteolytically truncated form of free CD18, the common chain of leukocyte integrins, as a novel marker of activated myeloid cells. Blood, 2001, 98, 1561-1566.	0.6	23
159	Synthesis of chitooligomer-based glycoconjugates and their binding to the rat natural killer cell activation receptor NKR-P1. Glycoconjugate Journal, 2001, 18, 817-826.	1.4	39
160	Two forms of DNA-dependent RNA polymerase α subunit in streptomycetes. FEMS Microbiology Letters, 2000, 187, 9-14.	0.7	3
161	Human Leukocytes Contain a Large Pool of Free Forms of CD18. Biochemical and Biophysical Research Communications, 2000, 275, 295-299.	1.0	16
162	Two forms of DNA-dependent RNA polymerase α subunit in streptomycetes. FEMS Microbiology Letters, 2000, 187, 9-14.	0.7	2

#	Article	IF	CITATIONS
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