Yehia S Mechref

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

257	10,403	58	88
papers	citations	h-index	g-index
273	11,507 ext. citations	5	6.42
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
257	Mitoquinone Helps Combat the Neurological, Cognitive, and Molecular Consequences of Open Head Traumatic Brain Injury at Chronic Time Point <i>Biomedicines</i> , 2022 , 10,	4.8	3
256	Mitoquinone supplementation alleviates oxidative stress and pathologic outcomes following repetitive mild traumatic brain injury at a chronic time point <i>Experimental Neurology</i> , 2022 , 351, 11398	7 ^{5.7}	2
255	Isomeric separation of permethylated glycans by extra-long reversed-phase liquid chromatography (RPLC)-MS/MS <i>Analyst, The</i> , 2022 ,	5	1
254	Salmonella enterica serovar Typhimurium chitinases modulate the intestinal glycome and promote small intestinal invasion <i>PLoS Pathogens</i> , 2022 , 18, e1010167	7.6	O
253	Glycome Profiling of Cancer Cell Lines Cultivated in Physiological and Commercial Media. <i>Biomolecules</i> , 2022 , 12, 743	5.9	O
252	Changes in the Expression of Renal Brush Border Membrane -Glycome in Model Rats with Chronic Kidney Diseases. <i>Biomolecules</i> , 2021 , 11,	5.9	1
251	Advances in mass spectrometry-based glycoproteomics: An update covering the period 2017-2021. <i>Electrophoresis</i> , 2021 ,	3.6	3
250	Altered -glycomes of Renal Brush-Border Membrane in Model Rats with Chronic Kidney Diseases. <i>Biomolecules</i> , 2021 , 11,	5.9	3
249	Community evaluation of glycoproteomics informatics solutions reveals high-performance search strategies for serum glycopeptide analysis. <i>Nature Methods</i> , 2021 , 18, 1304-1316	21.6	16
248	Analysis of the Neuroproteome Associated With Cell Therapy After Intranigral Grafting in a Mouse Model of Parkinson Disease. <i>Frontiers in Neuroscience</i> , 2021 , 15, 621121	5.1	1
247	Mesoporous Graphitized Carbon Column for Efficient Isomeric Separation of Permethylated Glycans. <i>Analytical Chemistry</i> , 2021 , 93, 5061-5070	7.8	5
246	A General New Method for Calculating the Molecular Nonpolar Surface for Analysis of LC-MS Data. <i>International Journal of Mass Spectrometry</i> , 2021 , 461, 116495-116495	1.9	2
245	-Glycomics of Cerebrospinal Fluid: Method Comparison. <i>Molecules</i> , 2021 , 26,	4.8	2
244	Object classification in analytical chemistry via data-driven discovery of partial differential equations. <i>Computational and Mathematical Methods</i> , 2021 , 3, e1164	0.9	1
243	GlycoHybridSeq: Automated Identification of N-Linked Glycopeptides Using Electron Transfer/High-Energy Collision Dissociation (EThcD). <i>Journal of Proteome Research</i> , 2021 , 20, 3345-3352	5.6	4
242	Browning white adipose tissue using adipose stromal cell-targeted resveratrol-loaded nanoparticles for combating obesity. <i>Journal of Controlled Release</i> , 2021 , 333, 339-351	11.7	7
241	Searching for Novel Candidate Biomarkers of RLS in Blood by Proteomic Analysis. <i>Nature and Science of Sleep</i> , 2021 , 13, 873-883	3.6	4

(2020-2021)

240	MS-based glycomics and glycoproteomics methods enabling isomeric characterization. <i>Mass Spectrometry Reviews</i> , 2021 ,	11	5
239	GlycanGUI: Automated Glycan Annotation and Quantification Using Glucose Unit Index. <i>Frontiers in Chemistry</i> , 2021 , 9, 707382	5	O
238	Glycomics and glycoproteomics: Approaches to address isomeric separation of glycans and glycopeptides. <i>Journal of Separation Science</i> , 2021 , 44, 403-425	3.4	19
237	Analysis of NIST Monoclonal Antibody Reference Material Glycosylation Using the LC-MS/MS-Based Glycoproteomic Approach. <i>Journal of Proteome Research</i> , 2021 , 20, 818-830	5.6	2
236	Determination of Isomeric Glycan Structures by Permethylation and Liquid Chromatography-Mass Spectrometry (LC-MS). <i>Methods in Molecular Biology</i> , 2021 , 2271, 281-301	1.4	
235	Mechanism of how carbamylation reduces albumin binding to FcRn contributing to increased vascular clearance. <i>American Journal of Physiology - Renal Physiology</i> , 2021 , 320, F114-F129	4.3	3
234	Using micro pillar array columns (PAC) for the analysis of permethylated glycans. <i>Analyst, The</i> , 2021 , 146, 4374-4383	5	5
233	Derivatization of Sialylated Glycopeptides (DOSG) Enabling Site-Specific Isomeric Profiling Using LC-MS/MS. <i>Analytical Chemistry</i> , 2021 , 93, 5763-5772	7.8	6
232	Direct Comparison of Glycans and Their Isomers Derived from Spike Glycoprotein 1 of MERS-CoV, SARS-CoV-1, and SARS-CoV-2. <i>Journal of Proteome Research</i> , 2021 , 20, 4357-4365	5.6	4
231	PRM-MS Quantitative Analysis of Isomeric -Glycopeptides Derived from Human Serum Haptoglobin of Patients with Cirrhosis and Hepatocellular Carcinoma. <i>Metabolites</i> , 2021 , 11,	5.6	5
230	Variability in the Glycosylation Patterns of gp120 Proteins from Different Human Immunodeficiency Virus Type 1 Isolates Expressed in Different Host Cells. <i>Journal of Proteome Research</i> , 2021 , 20, 4862-4874	5.6	2
229	Identification and Characterization of Two Structurally Related Dipeptides that Enhance Catalytic Efficiency of Neurolysin. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2021 , 379, 191-202	4.7	3
228	Advances in mass spectrometry-based glycomics-An update covering the period 2017-2021. <i>Electrophoresis</i> , 2021 ,	3.6	1
227	LC-MS/MS in glycomics and glycoproteomics analyses 2021 , 391-441		
226	Vascular Cells Proteome Associated with Bradykinin and Leptin Inflammation and Oxidative Stress Signals. <i>Antioxidants</i> , 2020 , 9,	7.1	2
225	Separation of Permethylated -Glycans, Free Oligosaccharides, and Glycosphingolipid-Glycans Using Porous Graphitized Carbon (PGC) Column. <i>Metabolites</i> , 2020 , 10,	5.6	6
224	Serum Glycomics Profiling of Patients with Primary Restless Legs Syndrome Using LC-MS/MS. Journal of Proteome Research, 2020 , 19, 2933-2941	5.6	7
223	Isomeric Separation of N-Glycopeptides Derived from Glycoproteins by Porous Graphitic Carbon (PGC) LC-MS/MS. <i>Analytical Chemistry</i> , 2020 , 92, 9556-9565	7.8	17

222	Modulation of proteomic and inflammatory signals by Bradykinin in podocytes. <i>Journal of Advanced Research</i> , 2020 , 24, 409-422	13	2
221	N-Linked Surface Glycan Biosynthesis, Composition, Inhibition, and Function in Cnidarian-Dinoflagellate Symbiosis. <i>Microbial Ecology</i> , 2020 , 80, 223-236	4.4	5
220	Circulating Brain Injury Exosomal Proteins following Moderate-To-Severe Traumatic Brain Injury: Temporal Profile, Outcome Prediction and Therapy Implications. <i>Cells</i> , 2020 , 9,	7.9	26
219	Comparative Membrane -Glycomics of Different Breast Cancer Cell Lines To Understand Breast Cancer Brain Metastasis. <i>Journal of Proteome Research</i> , 2020 , 19, 854-863	5.6	6
218	Effects of the Oncoprotein PAX3-FOXO1 on Modulation of Exosomes Function and Protein Content: Implications on Oxidative Stress Protection and Enhanced Plasticity. <i>Frontiers in Oncology</i> , 2020 , 10, 1784	5.3	2
217	Glucose unit index (GUI) of permethylated glycans for effective identification of glycans and glycan isomers. <i>Analyst, The</i> , 2020 , 145, 6656-6667	5	11
216	Protein Expression Analysis of an In Vitro Murine Model of Prostate Cancer Progression: Towards Identification of High-Potential Therapeutic Targets. <i>Journal of Personalized Medicine</i> , 2020 , 10,	3.6	3
215	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	7.6	49
214	Proteomic Profiling of Rhabdomyosarcoma-Derived Exosomes Yield Insights into Their Functional Role in Paracrine Signaling. <i>Journal of Proteome Research</i> , 2019 , 18, 3567-3579	5.6	5
213	Revealing the Biological Attributes of N-Glycan Isomers in Breast Cancer Brain Metastasis Using Porous Graphitic Carbon (PGC) Liquid Chromatography-Tandem Mass Spectrometry (LC-MS/MS). <i>Journal of Proteome Research</i> , 2019 , 18, 3731-3740	5.6	23
212	N-Glycan Profile of Cerebrospinal Fluids from Alzheimer Disease Patients Using Liquid Chromatography with Mass Spectrometry. <i>Journal of Proteome Research</i> , 2019 , 18, 3770-3779	5.6	22
211	Characterization of glycan isomers using magnetic carbon nanoparticles as a MALDI co-matrix. <i>RSC Advances</i> , 2019 , 9, 20137-20148	3.7	8
210	8-plex LC-MS/MS Analysis of Permethylated -Glycans Achieved by Using Stable Isotopic Iodomethane. <i>Analytical Chemistry</i> , 2019 , 91, 11794-11802	7.8	18
209	Integrated Transcriptomics, Proteomics, and Glycomics Reveals the Association between Up-regulation of Sialylated N-glycans/Integrin and Breast Cancer Brain Metastasis. <i>Scientific Reports</i> , 2019 , 9, 17361	4.9	8
208	Functional pathways associated with human carotid atheroma: a proteomics analysis. <i>Hypertension Research</i> , 2019 , 42, 362-373	4.7	2
207	Implication of the Kallikrein-Kinin system in neurological disorders: Quest for potential biomarkers and mechanisms. <i>Progress in Neurobiology</i> , 2018 , 165-167, 26-50	10.9	41
206	Glycosylation Changes in Brain Cancer. ACS Chemical Neuroscience, 2018, 9, 51-72	5.7	65
205	Enhanced Quantitative LC-MS/MS Analysis of N-linked Glycans Derived from Glycoproteins Using Sodium Deoxycholate Detergent. <i>Journal of Proteome Research</i> , 2018 , 17, 2668-2678	5.6	16

(2017-2018)

204	A carbon nanoparticles-based solid-phase purification method facilitating sensitive MALDI-MS analysis of permethylated N-glycans. <i>Electrophoresis</i> , 2018 , 39, 3087-3095	3.6	9	
203	Carbon Nanoparticles and Graphene Nanosheets as MALDI Matrices in Glycomics: a New Approach to Improve Glycan Profiling in Biological Samples. <i>Journal of the American Society for Mass Spectrometry</i> , 2018 , 29, 1892-1900	3.5	22	
202	Quantitative Glycomics by Mass Spectrometry and Liquid Chromatography Mass Spectrometry 2018 , 1-31			
201	Novel biomarker signatures for idiopathic REM sleep behavior disorder: A proteomic and system biology approach. <i>Neurology</i> , 2018 , 91, e1710-e1715	6.5	15	
200	Clinical application of quantitative glycomics. Expert Review of Proteomics, 2018, 15, 1007-1031	4.2	22	
199	Advances in mass spectrometry-based glycoproteomics. <i>Electrophoresis</i> , 2018 , 39, 3104-3122	3.6	53	
198	Advances in mass spectrometry-based glycomics. <i>Electrophoresis</i> , 2018 , 39, 3063-3081	3.6	47	
197	Magnetic carbon nanocomposites as a MALDI co-matrix enhancing MS-based glycomics. <i>Analytical and Bioanalytical Chemistry</i> , 2018 , 410, 7395-7404	4.4	5	
196	LC-MS/MS glycomics of idiopathic rapid eye movement sleep behavior disorder. <i>Electrophoresis</i> , 2018 , 39, 3096-3103	3.6	10	
195	Glycoprotein Enrichment Analytical Techniques: Advantages and Disadvantages. <i>Methods in Enzymology</i> , 2017 , 585, 397-429	1.7	37	
194	Quantitative Glycomics: A Combined Analytical and Bioinformatics Approach. <i>Methods in Enzymology</i> , 2017 , 585, 431-477	1.7	11	
193	Glycoproteins Enrichment and LC-MS/MS Glycoproteomics in Central Nervous System Applications. <i>Methods in Molecular Biology</i> , 2017 , 1598, 213-227	1.4	9	
192	Isomeric Separation of Permethylated Glycans by Porous Graphitic Carbon (PGC)-LC-MS/MS at High Temperatures. <i>Analytical Chemistry</i> , 2017 , 89, 6590-6597	7.8	70	
191	Comparative membrane proteomics analyses of breast cancer cell lines to understand the molecular mechanism of breast cancer brain metastasis. <i>Electrophoresis</i> , 2017 , 38, 2124-2134	3.6	14	
190	LC-MS/MS isomeric profiling of permethylated N-glycans derived from serum haptoglobin of hepatocellular carcinoma (HCC) and cirrhotic patients. <i>Electrophoresis</i> , 2017 , 38, 2160-2167	3.6	52	
189	Multitargeted Flavonoid Inhibition of the Pathogenic Bacterium Staphylococcus aureus: A Proteomic Characterization. <i>Journal of Proteome Research</i> , 2017 , 16, 2579-2586	5.6	16	
188	Delineation of disease phenotypes associated with esophageal adenocarcinoma by MALDI-IMS-MS analysis of serum N-linked glycans. <i>Analyst, The</i> , 2017 , 142, 1525-1535	5	10	
187	Characterization of isomeric glycan structures by LC-MS/MS. <i>Electrophoresis</i> , 2017 , 38, 2100-2114	3.6	92	

186	Multivariate classification of disease phenotypes of esophageal adenocarcinoma by pattern recognition analysis of MALDI-TOF mass spectra of serum N-linked glycans. <i>Microchemical Journal</i> , 2017 , 132, 83-88	4.8	3
185	Direct comparison of derivatization strategies for LC-MS/MS analysis of N-glycans. <i>Analyst, The</i> , 2017 , 142, 4446-4455	5	63
184	Proteome profiling in the aorta and kidney of type 1 diabetic rats. PLoS ONE, 2017, 12, e0187752	3.7	11
183	LC-MS/MS analysis of permethylated N-glycans facilitating isomeric characterization. <i>Analytical and Bioanalytical Chemistry</i> , 2017 , 409, 453-466	4.4	61
182	Recent advances in mass spectrometric analysis of glycoproteins. <i>Electrophoresis</i> , 2017 , 38, 162-189	3.6	63
181	Analysis of Permethylated Glycan by Liquid Chromatography (LC) and Mass Spectrometry (MS). <i>Methods in Molecular Biology</i> , 2017 , 1503, 83-96	1.4	22
180	Analysis of Proteins That Rapidly Change Upon Mechanistic/Mammalian Target of Rapamycin Complex 1 (mTORC1) Repression Identifies Parkinson Protein 7 (PARK7) as a Novel Protein Aberrantly Expressed in Tuberous Sclerosis Complex (TSC). <i>Molecular and Cellular Proteomics</i> , 2016 ,	7.6	19
179	15, 426-44 A procedure for the analysis of site-specific and structure-specific fucosylation in alpha-1-antitrypsin. <i>Electrophoresis</i> , 2016 , 37, 2624-2632	3.6	10
178	Iridoid glycoside permethylation enhances chromatographic separation and chemical ionization. <i>Rapid Communications in Mass Spectrometry</i> , 2016 , 30, 2033-42	2.2	4
177	HILIC and ERLIC Enrichment of Glycopeptides Derived from Breast and Brain Cancer Cells. <i>Journal of Proteome Research</i> , 2016 , 15, 3624-3634	5.6	60
176	Multi-omic approaches for characterization of hepatocellular carcinoma. <i>Annual International Conference of the IEEE Engineering in Medicine and Biology Society IEEE Engineering in Medicine and Biology Society Annual International Conference</i> , 2016 , 2016, 3437-3440	0.9	3
175	Parallel data acquisition of in-source fragmented glycopeptides to sequence the glycosylation sites of proteins. <i>Electrophoresis</i> , 2016 , 37, 1420-30	3.6	12
174	Glycosylation and other PTMs alterations in neurodegenerative diseases: Current status and future role in neurotrauma. <i>Electrophoresis</i> , 2016 , 37, 1549-61	3.6	36
173	LC-MS/MS analysis of permethylated free oligosaccharides and N-glycans derived from human, bovine, and goat milk samples. <i>Electrophoresis</i> , 2016 , 37, 1532-48	3.6	61
172	Structure-antioxidant and anti-tumor activity of Teucrium polium phytochemicals. <i>Phytochemistry Letters</i> , 2016 , 15, 81-87	1.9	13
171	Characterization of Pharmaceutical IgG and Biosimilars Using Miniaturized Platforms and LC-MS/MS. <i>Current Pharmaceutical Biotechnology</i> , 2016 , 17, 788-801	2.6	25
170	The Effect of Chronic Methamphetamine Exposure on the Hippocampal and Olfactory Bulb Neuroproteomes of Rats. <i>PLoS ONE</i> , 2016 , 11, e0151034	3.7	10
169	High-temperature LC-MS/MS of permethylated glycans derived from glycoproteins. <i>Electrophoresis</i> , 2016 , 37, 1506-13	3.6	42

(2015-2016)

168	Deciphering glycomics and neuroproteomic alterations in experimental traumatic brain injury: Comparative analysis of aspirin and clopidogrel treatment. <i>Electrophoresis</i> , 2016 , 37, 1562-76	3.6	17	
167	Reliable LC-MS quantitative glycomics using iGlycoMab stable isotope labeled glycans as internal standards. <i>Electrophoresis</i> , 2016 , 37, 1489-97	3.6	19	
166	Quantitative LC-MS/MS Glycomic Analysis of Biological Samples Using AminoxyTMT. <i>Analytical Chemistry</i> , 2016 , 88, 7515-22	7.8	50	
165	LC-MS/MS of permethylated N-glycans derived from model and human blood serum glycoproteins. <i>Electrophoresis</i> , 2016 , 37, 1498-505	3.6	34	
164	Automated Glycan Sequencing from Tandem Mass Spectra of N-Linked Glycopeptides. <i>Analytical Chemistry</i> , 2016 , 88, 5725-32	7.8	27	
163	Preparation and preliminary characterization of recombinant neurolysin for in vivo studies. <i>Journal of Biotechnology</i> , 2016 , 234, 105-115	3.7	15	
162	Quantitation of permethylated N-glycans through multiple-reaction monitoring (MRM) LC-MS/MS. <i>Journal of the American Society for Mass Spectrometry</i> , 2015 , 26, 596-603	3.5	33	
161	Characterization of the Glycosylation Site of Human PSA Prompted by Missense Mutation using LC-MS/MS. <i>Journal of Proteome Research</i> , 2015 , 14, 2872-83	5.6	19	
160	Integrated Transcriptomic and Glycomic Profiling of Glioma Stem Cell Xenografts. <i>Journal of Proteome Research</i> , 2015 , 14, 3932-9	5.6	11	
159	N-linked glycan profiling in neuroblastoma cell lines. <i>Journal of Proteome Research</i> , 2015 , 14, 2074-81	5.6	18	
158	Cytotoxic saponin poliusaposide from Teucrium polium. <i>RSC Advances</i> , 2015 , 5, 27126-27133	3.7	13	
157	Defining glycoprotein cancer biomarkers by MS in conjunction with glycoprotein enrichment. <i>Biomarkers in Medicine</i> , 2015 , 9, 835-44	2.3	30	
156	Identification of Glycopeptides with Multiple Hydroxylysine O-Glycosylation Sites by Tandem Mass Spectrometry. <i>Journal of Proteome Research</i> , 2015 , 14, 5099-108	5.6	8	
155	Teucrium polium phenylethanol and iridoid glycoside characterization and flavonoid inhibition of biofilm-forming Staphylococcus aureus. <i>Journal of Natural Products</i> , 2015 , 78, 2-9	4.9	32	
154	Automated annotation and quantitation of glycans by liquid chromatography/electrospray ionization mass spectrometric analysis using the MultiGlycan-ESI computational tool. <i>Rapid Communications in Mass Spectrometry</i> , 2015 , 29, 135-42	2.2	31	
153	Characterization of the Kallikrein-Kinin System Post Chemical Neuronal Injury: An In Vitro Biochemical and Neuroproteomics Assessment. <i>PLoS ONE</i> , 2015 , 10, e0128601	3.7	5	
152	30th ASMS Asilomar Conference on Advances in Glycomics and Glycoproteomics: Methods and Applications. <i>Journal of the American Society for Mass Spectrometry</i> , 2015 , 26, 1047-50	3.5		
151	LC-MS/MS-based serum proteomics for identification of candidate biomarkers for hepatocellular carcinoma. <i>Proteomics</i> , 2015 , 15, 2369-81	4.8	52	

150	Biofilm blocking sesquiterpenes from Teucrium polium. <i>Phytochemistry</i> , 2014 , 103, 107-113	4	31
149	Rapid and sensitive LC-ESI-MS of gangliosides. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2014 , 947-948, 1-7	3.2	30
148	LC-MS profiling of N-Glycans derived from human serum samples for biomarker discovery in hepatocellular carcinoma. <i>Journal of Proteome Research</i> , 2014 , 13, 4859-68	5.6	40
147	LC-MS/MS quantitation of esophagus disease blood serum glycoproteins by enrichment with hydrazide chemistry and lectin affinity chromatography. <i>Journal of Proteome Research</i> , 2014 , 13, 4808-2	2 6 .6	63
146	Computational framework for identification of intact glycopeptides in complex samples. <i>Analytical Chemistry</i> , 2014 , 86, 453-63	7.8	79
145	Label-free glycopeptide quantification for biomarker discovery in human sera. <i>Journal of Proteome Research</i> , 2014 , 13, 4821-32	5.6	34
144	Recent updates on drug abuse analyzed by neuroproteomics studies: Cocaine, Methamphetamine and MDMA. <i>Translational Proteomics</i> , 2014 , 3, 38-52		6
143	Quantitative Glycomics by Mass Spectrometry and Liquid Chromatography-Mass Spectrometry 2014 , 1-22		2
142	Social learning and amygdala disruptions in Nf1 mice are rescued by blocking p21-activated kinase. <i>Nature Neuroscience</i> , 2014 , 17, 1583-90	25.5	82
141	Glycoproteomics: identifying the glycosylation of prostate specific antigen at normal and high isoelectric points by LC-MS/MS. <i>Journal of Proteome Research</i> , 2014 , 13, 5570-80	5.6	38
140	Bioinformatics protocols in glycomics and glycoproteomics. <i>Current Protocols in Protein Science</i> , 2014 , 76, 2.15.1-2.15.7	3.1	9
139	LC-MS/MS identification of the O-glycosylation and hydroxylation of amino acid residues of collagen 日 (II) chain from bovine cartilage. <i>Journal of Proteome Research</i> , 2013 , 12, 3599-609	5.6	31
138	Immobilized metal affinity chromatography and human serum proteomics. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2013 , 934, 26-33	3.2	32
137	Multi-profile Bayesian alignment model for LC-MS data analysis with integration of internal standards. <i>Bioinformatics</i> , 2013 , 29, 2774-80	7.2	15
136	Glycomic profiling of tissue sections by LC-MS. <i>Analytical Chemistry</i> , 2013 , 85, 4074-9	7.8	31
135	Glycomic Profiling Through Capillary Electrophoresis and Microchip Capillary Electrophoresis 2013 , 367	-383	2
134	Leaf Dhurrin Content is a Quantitative Measure of the Level of Pre- and Postflowering Drought Tolerance in Sorghum. <i>Crop Science</i> , 2013 , 53, 1056-1065	2.4	35
133	Comparative glycomic profiling of isotopically permethylated N-glycans by liquid chromatography/electrospray ionization mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2013 , 27, 865-77	2.2	44

(2011-2013)

132	Automated annotation and quantification of glycans using liquid chromatography-mass spectrometry. <i>Bioinformatics</i> , 2013 , 29, 1706-7	7.2	40
131	Quantitative glycomics strategies. <i>Molecular and Cellular Proteomics</i> , 2013 , 12, 874-84	7.6	76
130	Sugar nucleotide quantification using multiple reaction monitoring liquid chromatography/tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2013 , 27, 1794-800	2.2	6
129	GPA: An algorithm for LC/MS based glycan profile annotation 2013 ,		2
128	Interlaboratory study on differential analysis of protein glycosylation by mass spectrometry: the ABRF glycoprotein research multi-institutional study 2012. <i>Molecular and Cellular Proteomics</i> , 2013 , 12, 2935-51	7.6	92
127	Ion mobility-mass spectrometry analysis of serum N-linked glycans from esophageal adenocarcinoma phenotypes. <i>Journal of Proteome Research</i> , 2012 , 11, 6102-10	5.6	40
126	N-linked glycan structures and their expressions change in the blood sera of ovarian cancer patients. <i>Journal of Proteome Research</i> , 2012 , 11, 2282-300	5.6	149
125	Defining putative glycan cancer biomarkers by MS. <i>Bioanalysis</i> , 2012 , 4, 2457-69	2.1	53
124	Delineating diseases by IMS-MS profiling of serum N-linked glycans. <i>Journal of Proteome Research</i> , 2012 , 11, 576-85	5.6	46
123	N-glycan profiling by microchip electrophoresis to differentiate disease states related to esophageal adenocarcinoma. <i>Analytical Chemistry</i> , 2012 , 84, 3621-7	7.8	40
122	Fish consumption, low-level mercury, lipids, and inflammatory markers in children. <i>Environmental Research</i> , 2012 , 112, 204-11	7.9	32
121	Use of CID/ETD mass spectrometry to analyze glycopeptides. <i>Current Protocols in Protein Science</i> , 2012 , Chapter 12, Unit 12.11.1-11	3.1	68
120	Fast and efficient online release of N-glycans from glycoproteins facilitating liquid chromatography-tandem mass spectrometry glycomic profiling. <i>Analytical Chemistry</i> , 2012 , 84, 8790-6	7.8	35
119	Quantification of glycopeptides by multiple reaction monitoring liquid chromatography/tandem mass spectrometry. <i>Rapid Communications in Mass Spectrometry</i> , 2012 , 26, 1941-54	2.2	61
118	Comparing MALDI-MS, RP-LC-MALDI-MS and RP-LC-ESI-MS glycomic profiles of permethylated N-glycans derived from model glycoproteins and human blood serum. <i>Electrophoresis</i> , 2012 , 33, 1768-7	7 ^{3.6}	54
117	Identifying cancer biomarkers by mass spectrometry-based glycomics. <i>Electrophoresis</i> , 2012 , 33, 1755-6	73.6	115
116	Gel-free shotgun proteomic analysis of human milk. <i>Journal of Chromatography A</i> , 2012 , 1227, 219-33	4.5	30
115	Monosaccharide Compositional Analysis of Glycoproteins Using Liquid Chromatography-Mass Spectrometry. <i>Current Proteomics</i> , 2011 , 8, 257-268	0.7	3

114	Glycomic Alterations in the Highly-abundant and Lesser-abundant Blood Serum Protein Fractions for Patients Diagnosed with Hepatocellular Carcinoma. <i>International Journal of Mass Spectrometry</i> , 2011 , 305, 185-198	1.9	15
113	Improving confidence in detection and characterization of protein N-glycosylation sites and microheterogeneity. <i>Rapid Communications in Mass Spectrometry</i> , 2011 , 25, 2007-19	2.2	61
112	Microchip electrophoresis of N-glycans on serpentine separation channels with asymmetrically tapered turns. <i>Electrophoresis</i> , 2011 , 32, 246-53	3.6	30
111	Analysis of glycans derived from glycoconjugates by capillary electrophoresis-mass spectrometry. <i>Electrophoresis</i> , 2011 , 32, 3467-81	3.6	52
110	Enhanced sensitivity of LC-MS analysis of permethylated N-glycans through online purification. <i>Electrophoresis</i> , 2011 , 32, 3516-25	3.6	57
109	Increased protein nitration in mitochondrial diseases: evidence for vessel wall involvement. <i>Molecular and Cellular Proteomics</i> , 2011 , 10, M110.002964	7.6	32
108	Bioinformatic Approaches in Glycomics and Glycoproteomics. <i>Current Proteomics</i> , 2011 , 8, 309-324	0.7	1
107	Monosaccharide Compositional Analysis of Glycoproteins and Glycolipids: Utility in the Diagnosis/Prognosis of Diseases 2011 , 237-267		1
106	High-Sensitivity Analytical Approaches to the Analysis of N-Glycans 2010 , 1-43		2
105	Analysis of site-specific glycosylation of renal and hepatic liglutamyl transpeptidase from normal human tissue. <i>Journal of Biological Chemistry</i> , 2010 , 285, 29511-24	5.4	38
104	Identification of N-glycan serum markers associated with hepatocellular carcinoma from mass spectrometry data. <i>Journal of Proteome Research</i> , 2010 , 9, 104-12	5.6	58
103	Chip-based reversed-phase liquid chromatography-mass spectrometry of permethylated N-linked glycans: a potential methodology for cancer-biomarker discovery. <i>Analytical Chemistry</i> , 2010 , 82, 5095-	1068	139
102	Effects of lead and mercury on the blood proteome of children. <i>Journal of Proteome Research</i> , 2010 , 9, 4443-53	5.6	24
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