## Haidi Yin

## List of Publications by Year in descending order

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516710 677142 22 684 16 22 citations h-index g-index papers 24 24 24 953 docs citations citing authors all docs times ranked

#	Article	IF	CITATIONS
1	Analysis of Serum Haptoglobin Fucosylation in Hepatocellular Carcinoma and Liver Cirrhosis of Different Etiologies. Journal of Proteome Research, 2014, 13, 2986-2997.	3.7	103
2	Large-Scale Identification of Core-Fucosylated Glycopeptide Sites in Pancreatic Cancer Serum Using Mass Spectrometry. Journal of Proteome Research, 2015, 14, 1968-1978.	3.7	66
3	Atlastin-mediated membrane tethering is critical for cargo mobility and exit from the endoplasmic reticulum. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 14029-14038.	7.1	52
4	Mass-Selected Site-Specific Core-Fucosylation of Ceruloplasmin in Alcohol-Related Hepatocellular Carcinoma. Journal of Proteome Research, 2014, 13, 2887-2896.	3.7	48
5	Analysis of Glycan Variation on Glycoproteins from Serum by the Reverse Lectin-Based ELISA Assay. Journal of Proteome Research, 2014, 13, 2197-2204.	3.7	41
6	ESI–LC–MS Method for Haptoglobin Fucosylation Analysis in Hepatocellular Carcinoma and Liver Cirrhosis. Journal of Proteome Research, 2015, 14, 5388-5395.	3.7	38
7	Mass-Selected Site-Specific Core-Fucosylation of Serum Proteins in Hepatocellular Carcinoma. Journal of Proteome Research, 2015, 14, 4876-4884.	3.7	37
8	Mass Spectrometric N-Glycan Analysis of Haptoglobin from Patient Serum Samples Using a 96-Well Plate Format. Journal of Proteome Research, 2015, 14, 4932-4939.	3.7	30
9	Validation of LRG1 as a Potential Biomarker for Detection of Epithelial Ovarian Cancer by a Blinded Study. PLoS ONE, 2015, 10, e0121112.	2.5	27
10	Extracellular Proteome Analysis of <i>Leptospira interrogans </i> Integrative Biology, 2013, 17, 527-535.	2.0	26
11	Quantitative Analysis of Single Amino Acid Variant Peptides Associated with Pancreatic Cancer in Serum by an Isobaric Labeling Quantitative Method. Journal of Proteome Research, 2014, 13, 6058-6066.	3.7	26
12	An in vitro vesicle formation assay reveals cargo clients and factors that mediate vesicular trafficking. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	25
13	Labelâ€free relative quantification of alphaâ€2â€macroglobulin siteâ€specific coreâ€fucosylation in pancreatic cancer by LCâ€MS/MS. Electrophoresis, 2014, 35, 2108-2115.	2.4	22
14	The analysis of alphaâ€1â€antitrypsin glycosylation with direct LCâ€MS/MS. Electrophoresis, 2018, 39, 2351-2361.	2.4	22
15	Quantitative Analysis of α-1-Antitrypsin Glycosylation Isoforms in HCC Patients Using LC-HCD-PRM-MS. Analytical Chemistry, 2020, 92, 8201-8208.	6.5	21
16	Data storage using peptide sequences. Nature Communications, 2021, 12, 4242.	12.8	20
17	Development of an Integrated Pipeline for Profiling Microbial Proteins from Mouse Fecal Samples by LC–MS/MS. Journal of Proteome Research, 2016, 15, 3635-3642.	3.7	17
18	A distinct giant coat protein complex II vesicle population in Arabidopsis thaliana. Nature Plants, 2021, 7, 1335-1346.	9.3	15

#	Article	IF	CITATION
19	Methods for quantification of glycopeptides by liquid separation and mass spectrometry. Mass Spectrometry Reviews, 2023, 42, 887-917.	5.4	13
20	Impacts of transgenic Bt cotton on the stylet penetration behaviors of <i>Bemisia tabaci</i> biotype B: Evidence from laboratory experiments. Insect Science, 2010, 17, 344-352.	3.0	12
21	Proteomic analysis of Arabidopsis thaliana leaves infested by tobacco whitefly Bemisia tabaci (Gennadius) B biotype. Plant Molecular Biology Reporter, 2012, 30, 379-390.	1.8	12
22	A procedure for the analysis of siteâ€specific and structureâ€specific fucosylation in alphaâ€1â€antitrypsin. Electrophoresis, 2016, 37, 2624-2632.	2.4	10