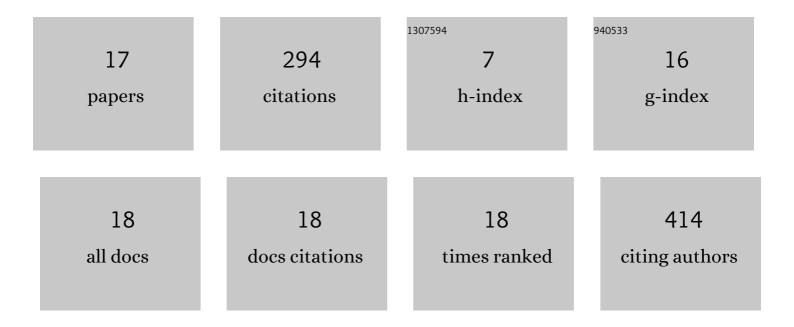
## Wantao Ying

List of Publications by Year in descending order

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Μαντλο Υίνο

#	Article	IF	CITATIONS
1	Bone marrow mesenchymal stem cell-derived exosomal miR-34c-5p ameliorates RIF by inhibiting the core fucosylation of multiple proteins. Molecular Therapy, 2022, 30, 763-781.	8.2	24
2	Strategy for Microscale Extraction and Proteome Profiling of Peripheral Blood Mononuclear Cells. Analytical Chemistry, 2022, 94, 8827-8832.	6.5	0
3	Development of the C12Im-Cl-assisted method for rapid sample preparation in proteomic application. Analytical Methods, 2021, 13, 776-781.	2.7	5
4	Optimised data-independent acquisition strategy recaptures the classification of early-stage hepatocellular carcinoma based on data-dependent acquisition. Journal of Proteomics, 2021, 238, 104152.	2.4	6
5	O-glycoforms of polymeric immunoglobulin A1 in the plasma of patients with IgA nephropathy are associated with pathological phenotypes. Nephrology Dialysis Transplantation, 2021, 37, 33-41.	0.7	5
6	Integrated Strategy for Unbiased Profiling of the Histidine Phosphoproteome. Analytical Chemistry, 2021, 93, 15584-15589.	6.5	4
7	L-cysteine functionalized straticulate C3N4 for the selective enrichment of glycopeptides. Journal of Chromatography A, 2020, 1610, 460545.	3.7	20
8	Site-Specific and Quantitative N-Glycan Heterogeneity Analysis of the Charge Isomers of an Anti-VEGF Recombinant Fusion Protein by High-Resolution Two-Dimensional Gel Electrophoresis and Mass Spectrometry. Analytical Chemistry, 2020, 92, 5695-5700.	6.5	4
9	Relative content detection of oligomannose modification of IgM heavy chain induced by TNP-antigen in an early vertebrate through nanoLC-MS/MS. Talanta, 2020, 219, 121346.	5.5	3
10	Integrated Strategy for Large-Scale Investigation on Protein Core Fucosylation Stoichiometry Based on Glycan-Simplification and Paired-Peaks-Extraction. Analytical Chemistry, 2020, 92, 2896-2901.	6.5	7
11	Identification of hedgehog signaling as a potential oncogenic driver in an aggressive subclass of human hepatocellular carcinoma: A reanalysis of the TCGA cohort. Science China Life Sciences, 2019, 62, 1481-1491.	4.9	5
12	A multi-parallel N-glycopeptide enrichment strategy for high-throughput and in-depth mapping of the N-glycoproteome in metastatic human hepatocellular carcinoma cell lines. Talanta, 2019, 199, 254-261.	5.5	12
13	Phenylboronic acid functionalized C3N4 facultative hydrophilic materials for enhanced enrichment of glycopeptides. Talanta, 2019, 191, 509-518.	5.5	38
14	Systems analysis of singly and multiply O -glycosylated peptides in the human serum glycoproteome via EThcD and HCD mass spectrometry. Journal of Proteomics, 2018, 170, 14-27.	2.4	72
15	Targeted histidine-peptide enrichment improved the accuracy of isobaric-based quantitative proteomics. Analytical Methods, 2016, 8, 5255-5261.	2.7	5
16	In-depth analysis of secretome and N-glycosecretome of human hepatocellular carcinoma metastatic cell lines shed light on metastasis correlated proteins. Oncotarget, 2016, 7, 22031-22049.	1.8	18
17	Strategy Integrating Stepped Fragmentation and Glycan Diagnostic Ion-Based Spectrum Refinement for the Identification of Core Fucosylated Glycoproteome Using Mass Spectrometry. Analytical Chemistry, 2014, 86, 6804-6811.	6.5	59