

Haopeng Xiao

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

1,923
citations

331670

21
h-index

414414

32
g-index

34
all docs

34
docs citations

34
times ranked

3244
citing authors

#	ARTICLE	IF	CITATIONS
1	Cysteine 253 of UCP1 regulates energy expenditure and sex-dependent adipose tissue inflammation. <i>Cell Metabolism</i> , 2022, 34, 140-157.e8.	16.2	27
2	Simultaneously Identifying and Distinguishing Glycoproteins with O-GlcNAc and O-GalNAc (the Tn) Tj ETQq0 0 0 rgBTj/Overlock 10 Tf 50	8.5	10
3	AIDA and UCP1 snuggle up to prevent hypothermia. <i>Nature Cell Biology</i> , 2021, 23, 216-218.	10.3	0
4	IRF3 reduces adipose thermogenesis via ISG15-mediated reprogramming of glycolysis. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	43
5	UCP1 governs liver extracellular succinate and inflammatory pathogenesis. <i>Nature Metabolism</i> , 2021, 3, 604-617.	11.9	82
6	Glycogen metabolism links glucose homeostasis to thermogenesis in adipocytes. <i>Nature</i> , 2021, 599, 296-301.	27.8	36
7	pH-Gated Succinate Secretion Regulates Muscle Remodeling in Response to Exercise. <i>Cell</i> , 2020, 183, 62-75.e17.	28.9	129
8	A Quantitative Tissue-Specific Landscape of Protein Redox Regulation during Aging. <i>Cell</i> , 2020, 180, 968-983.e24.	28.9	220
9	Systematic quantification of the dynamics of newly synthesized proteins unveiling their degradation pathways in human cells. <i>Chemical Science</i> , 2020, 11, 3557-3568.	7.4	18
10	Sample multiplexing for targeted pathway proteomics in aging mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 9723-9732.	7.1	73
11	Comprehensive Analysis of Protein Glycation Reveals Its Potential Impacts on Protein Degradation and Gene Expression in Human Cells. <i>Journal of the American Society for Mass Spectrometry</i> , 2019, 30, 2480-2490.	2.8	17
12	Global and site-specific analysis of protein glycosylation in complex biological systems with Mass Spectrometry. <i>Mass Spectrometry Reviews</i> , 2019, 38, 356-379.	5.4	75
13	An enrichment method based on synergistic and reversible covalent interactions for large-scale analysis of glycoproteins. <i>Nature Communications</i> , 2018, 9, 1692.	12.8	127
14	Extracellular vesicles from bone marrow-derived mesenchymal stromal cells support <i>ex vivo</i> survival of human antibody secreting cells. <i>Journal of Extracellular Vesicles</i> , 2018, 7, 1463778.	12.2	27
15	Mass spectrometric analysis of the N-glycoproteome in statin-treated liver cells with two lectin-independent chemical enrichment methods. <i>International Journal of Mass Spectrometry</i> , 2018, 429, 66-75.	1.5	12
16	Factors of the bone marrow microniche that support human plasma cell survival and immunoglobulin secretion. <i>Nature Communications</i> , 2018, 9, 3698.	12.8	95
17	Mass Spectrometry-Based Chemical and Enzymatic Methods for Global Analysis of Protein Glycosylation. <i>Accounts of Chemical Research</i> , 2018, 51, 1796-1806.	15.6	77
18	Global and Site-Specific Analysis Revealing Unexpected and Extensive Protein S-GlcNAcylation in Human Cells. <i>Analytical Chemistry</i> , 2017, 89, 3656-3663.	6.5	21

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19	Specific Identification of Glycoproteins Bearing the Tn Antigen in Human Cells. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 7107-7111.	13.8	48
20	Specific Identification of Glycoproteins Bearing the Tn Antigen in Human Cells. <i>Angewandte Chemie</i> , 2017, 129, 7213-7217.	2.0	2
21	Efficacy, long-term toxicity, and mechanistic studies of gold nanorods photothermal therapy of cancer in xenograft mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E3110-E3118.	7.1	237
22	Global Analysis of Secreted Proteins and Glycoproteins in <i>Saccharomyces cerevisiae</i> . <i>Journal of Proteome Research</i> , 2017, 16, 1039-1049.	3.7	30
23	Evaluation and optimization of reduction and alkylation methods to maximize peptide identification with MS-based proteomics. <i>Molecular BioSystems</i> , 2017, 13, 2574-2582.	2.9	68
24	Simultaneous Quantitation of Glycoprotein Degradation and Synthesis Rates by Integrating Isotope Labeling, Chemical Enrichment, and Multiplexed Proteomics. <i>Analytical Chemistry</i> , 2017, 89, 10361-10367.	6.5	13
25	Evidence for the importance of post-transcriptional regulatory changes in ovarian cancer progression and the contribution of miRNAs. <i>Scientific Reports</i> , 2017, 7, 8171.	3.3	14
26	Targeting cancer cell integrins using gold nanorods in photothermal therapy inhibits migration through affecting cytoskeletal proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E5655-E5663.	7.1	151
27	Quantitative investigation of human cell surface N-glycoprotein dynamics. <i>Chemical Science</i> , 2017, 8, 268-277.	7.4	55
28	Simultaneous Time-Dependent Surface-Enhanced Raman Spectroscopy, Metabolomics, and Proteomics Reveal Cancer Cell Death Mechanisms Associated with Gold Nanorod Photothermal Therapy. <i>Journal of the American Chemical Society</i> , 2016, 138, 15434-15442.	13.7	128
29	Quantification of tunicamycin-induced protein expression and N-glycosylation changes in yeast. <i>Analyst</i> , 2016, 141, 3737-3745.	3.5	30
30	Site-Specific Quantification of Surface N-Glycoproteins in Statin-Treated Liver Cells. <i>Analytical Chemistry</i> , 2016, 88, 3324-3332.	6.5	44
31	A Boronic Acid-Based Enrichment for Site-Specific Identification of the N-glycoproteome Using MS-Based Proteomics. <i>NeuroMethods</i> , 2015, , 31-41.	0.3	3
32	Systematic Investigation of Cellular Response and Pleiotropic Effects in Atorvastatin-Treated Liver Cells by MS-Based Proteomics. <i>Journal of Proteome Research</i> , 2015, 14, 1600-1611.	3.7	9