Haiyan Zheng

List of Publications by Year in descending order

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687363 677142 22 874 13 22 citations h-index g-index papers 26 26 26 1467 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Autophagy maintains tumour growth through circulating arginine. Nature, 2018, 563, 569-573.	27.8	279
2	TMK-based cell-surface auxin signalling activates cell-wall acidification. Nature, 2021, 599, 278-282.	27.8	125
3	Accounting for Protein Subcellular Localization: A Compartmental Map of the Rat Liver Proteome. Molecular and Cellular Proteomics, 2017, 16, 194-212.	3.8	69
4	Role of N-linked glycosylation in the secretion and enzymatic properties of Rhizopus chinensis lipase expressed in Pichia pastoris. Microbial Cell Factories, 2015, 14, 40.	4.0	57
5	CDK8-Cyclin C Mediates Nutritional Regulation of Developmental Transitions through the Ecdysone Receptor in Drosophila. PLoS Biology, 2015, 13, e1002207.	5.6	38
6	O-GlcNAcylation of PERIOD regulates its interaction with CLOCK and timing of circadian transcriptional repression. PLoS Genetics, 2019, 15, e1007953.	3.5	38
7	Proteomic Analysis of Brain and Cerebrospinal Fluid from the Three Major Forms of Neuronal Ceroid Lipofuscinosis Reveals Potential Biomarkers. Journal of Proteome Research, 2017, 16, 3787-3804.	3.7	37
8	Analysis of Brain and Cerebrospinal Fluid from Mouse Models of the Three Major Forms of Neuronal Ceroid Lipofuscinosis Reveals Changes in the Lysosomal Proteome. Molecular and Cellular Proteomics, 2019, 18, 2244-2261.	3.8	29
9	Protein phosphatase 1 catalyzes HBV core protein dephosphorylation and is co-packaged with viral pregenomic RNA into nucleocapsids. PLoS Pathogens, 2020, 16, e1008669.	4.7	26
10	Paraspeckle Protein NONO Promotes TAZ Phase Separation in the Nucleus to Drive the Oncogenic Transcriptional Program. Advanced Science, 2021, 8, e2102653.	11.2	24
11	<scp>NEDD</scp> 4Lâ€mediated Merlin ubiquitination facilitates Hippo pathway activation. EMBO Reports, 2020, 21, e50642.	4.5	18
12	<i>Mycobacterium tuberculosis</i> VapC4 toxin engages small ORFs to initiate an integrated oxidative and copper stress response. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	17
13	Large-scale Proteomic and Phosphoproteomic Analyses of Maize Seedling Leaves During De-etiolation. Genomics, Proteomics and Bioinformatics, 2020, 18, 397-414.	6.9	16
14	Comparative Analysis of Quantitative Mass Spectrometric Methods for Subcellular Proteomics. Journal of Proteome Research, 2020, 19, 1718-1730.	3.7	15
15	CK2 Inhibits TIMELESS Nuclear Export and Modulates CLOCK Transcriptional Activity to Regulate Circadian Rhythms. Current Biology, 2021, 31, 502-514.e7.	3.9	15
16	ATR/ATM-Mediated Phosphorylation of BRCA1 T1394 Promotes Homologous Recombinational Repair and G2â€"M Checkpoint Maintenance. Cancer Research, 2021, 81, 4676-4684.	0.9	14
17	Large-scale Identification and Time-course Quantification of Ubiquitylation Events During Maize Seedling De-etiolation. Genomics, Proteomics and Bioinformatics, 2019, 17, 603-622.	6.9	11
18	Thyroid hormone receptor phosphorylation regulates acute fasting-induced suppression of the hypothalamic–pituitary–thyroid axis. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	11

#	Article	IF	CITATION
19	A comprehensive analysis of the lysine acetylome reveals diverse functions of acetylated proteins during de-etiolation in Zea mays. Journal of Plant Physiology, 2020, 248, 153158.	3.5	9
20	PRMT5 Interacting Partners and Substrates in Oligodendrocyte Lineage Cells. Frontiers in Cellular Neuroscience, 2022, 16, 820226.	3.7	8
21	Protein-aggregating ability of different protoporphyrin-IX nanostructures is dependent on their oxidation and protein-binding capacity. Journal of Biological Chemistry, 2021, 297, 100778.	3.4	6
22	Independent effects of Src kinase and podoplanin on anchorage independent cell growth and migration. Molecular Carcinogenesis, 2022, , .	2.7	1