Xiaolu Zhao

List of Publications by Year in descending order

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567281 642732 1,223 23 15 23 h-index citations g-index papers 23 23 23 2433 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Cholesterylation of Smoothened is a calcium-accelerated autoreaction involving an intramolecular ester intermediate. Cell Research, 2022, 32, 288-301.	12.0	18
2	Enterovirus 71 3C proteolytically processes the histone H3 N-terminal tail during infection. Virologica Sinica, 2022, 37, 314-317.	3.0	1
3	POST1/C12ORF49 regulates the SREBP pathway by promoting site-1 protease maturation. Protein and Cell, 2021, 12, 279-296.	11.0	31
4	Feeding induces cholesterol biosynthesis via the mTORC1–USP20–HMGCR axis. Nature, 2020, 588, 479-484.	27.8	125
5	Identification of Hydroxyproline-Containing Proteins and Hydroxylation of Proline Residues in Rice. Frontiers in Plant Science, 2020, 11, 1207.	3.6	3
6	Determination of RNA Hydroxylmethylation in Mammals by Mass Spectrometry Analysis. Analytical Chemistry, 2019, 91, 10477-10483.	6.5	29
7	Thermostable potassium channel-inhibiting neurotoxins in processed scorpion medicinal material revealed by proteomic analysis: Implications of its pharmaceutical basis in traditional Chinese medicine. Journal of Proteomics, 2019, 206, 103435.	2.4	10
8	P200 family protein IFI204 negatively regulates type I interferon responses by targeting IRF7 in nucleus. PLoS Pathogens, 2019, 15, e1008079.	4.7	23
9	Proteomics Profiling of Host Cell Response via Protein Expression and Phosphorylation upon Dengue Virus Infection. Virologica Sinica, 2019, 34, 549-562.	3.0	23
10	A methylation-phosphorylation switch determines Plk1 kinase activity and function in DNA damage repair. Science Advances, 2019, 5, eaau7566.	10.3	52
11	Isotopic Labeling and Quantitative Proteomics of Acetylation on Histones and Beyond. Methods in Molecular Biology, 2019, 1977, 43-70.	0.9	12
12	Quantitative Proteomics Identified TTC4 as a TBK1 Interactor and a Positive Regulator of SeVâ€Induced Innate Immunity. Proteomics, 2018, 18, 1700403.	2,2	15
13	Assessment of Quantification Precision of Histone Post-Translational Modifications by Using an Ion Trap and down To 50†000 Cells as Starting Material. Journal of Proteome Research, 2018, 17, 234-242.	3.7	10
14	Gcn5-mediated Rph1 acetylation regulates its autophagic degradation under DNA damage stress. Nucleic Acids Research, 2017, 45, 5183-5197.	14.5	27
15	Abnormal levels of histone methylation in the retinas of diabetic rats are reversed by minocycline treatment. Scientific Reports, 2017, 7, 45103.	3.3	18
16	Proteome-wide acetylation dynamics in human cells. Scientific Reports, 2017, 7, 10296.	3.3	42
17	The tumor suppressor PTEN has a critical role in antiviral innate immunity. Nature Immunology, 2016, 17, 241-249.	14.5	138
18	Human Enterovirus Nonstructural Protein 2CATPase Functions as Both an RNA Helicase and ATP-Independent RNA Chaperone. PLoS Pathogens, 2015, 11, e1005067.	4.7	68

XIAOLU ZHAO

#	Article	IF	CITATIONS
19	Comparative Proteomic Analysis of Histone Post-translational Modifications upon Ischemia/Reperfusion-Induced Retinal Injury. Journal of Proteome Research, 2014, 13, 2175-2186.	3.7	16
20	Insulin Increases Phosphorylation of Mitochondrial Proteins in Human Skeletal Muscle <i>in Vivo</i> Journal of Proteome Research, 2014, 13, 2359-2369.	3.7	22
21	Quantitative proteomics reveals olfactory input-dependent alterations in the mouse olfactory bulb proteome. Journal of Proteomics, 2014, 109, 125-142.	2.4	10
22	MicroRNA Directly Enhances Mitochondrial Translation during Muscle Differentiation. Cell, 2014, 158, 607-619.	28.9	385
23	Phosphoproteome Analysis of Functional Mitochondria Isolated from Resting Human Muscle Reveals Extensive Phosphorylation of Inner Membrane Protein Complexes and Enzymes. Molecular and Cellular Proteomics, 2011, 10, M110.000299.	3.8	145