Xiahe Huang

List of Publications by Year in descending order

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		304743	254184
52	2,125	22	43
papers	citations	h-index	g-index
F.2	F 2	E2	2406
53	53	53	3406
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Dogs lacking Apolipoprotein E show advanced atherosclerosis leading to apparent clinical complications. Science China Life Sciences, 2022, 65, 1342-1356.	4.9	4
2	The UBP14-CDKB1;1-CDKG2 cascade controls endoreduplication and cell growth in Arabidopsis. Plant Cell, 2022, 34, 1308-1325.	6.6	12
3	An RDHâ€Plin2 axis modulates lipid droplet size by antagonizing Bmm lipase. EMBO Reports, 2022, 23, e52669.	4.5	7
4	Regulation of nitrogen starvation responses by the alarmone (p)ppGpp in rice. Journal of Genetics and Genomics, 2022, 49, 469-480.	3.9	12
5	OsHYPK-mediated protein N-terminal acetylation coordinates plant development and abiotic stress responses in rice. Molecular Plant, 2022, 15, 740-754.	8.3	9
6	Comparative Proteome and Cis-Regulatory Element Analysis Reveals Specific Molecular Pathways Conserved in Dog and Human Brains. Molecular and Cellular Proteomics, 2022, 21, 100261.	3.8	7
7	The plastid-encoded protein Orf2971 is required for protein translocation and chloroplast quality control. Plant Cell, 2022, 34, 3383-3399.	6.6	12
8	Modulation of nitrate-induced phosphate response by the MYB transcription factor RLI1/HINGE1 in the nucleus. Molecular Plant, 2021, 14, 517-529.	8.3	22
9	RNA kinase CLP1/Cbc regulates meiosis initiation in spermatogenesis. Human Molecular Genetics, 2021, 30, 1569-1578.	2.9	7
10	A zinc transporter, transmembrane protein 163, is critical for the biogenesis of platelet dense granules. Blood, 2021, 137, 1804-1817.	1.4	14
11	ERAD-related E2 and E3 enzymes modulate the drought response by regulating the stability of PIP2 aquaporins. Plant Cell, 2021, 33, 2883-2898.	6.6	44
12	A multi-omics investigation of the composition and function of extracellular vesicles along the temporal trajectory of COVID-19. Nature Metabolism, 2021, 3, 909-922.	11.9	132
13	Synergistic interplay of ABA and BR signal in regulating plant growth and adaptation. Nature Plants, 2021, 7, 1108-1118.	9.3	49
14	Plant Phosphopeptides Enrichment by Immobilized Metal Ion Affinity Chromatography. Methods in Molecular Biology, 2021, 2358, 145-157.	0.9	4
15	A Systematic Survey of the Light/Dark-dependent Protein Degradation Events in a Model Cyanobacterium. Molecular and Cellular Proteomics, 2021, 20, 100162.	3.8	2
16	The Quantitative Proteome Atlas of a Model Cyanobacterium. Journal of Genetics and Genomics, 2021, , .	3.9	14
17	NuRD mediates mitochondrial stress–induced longevity via chromatin remodeling in response to acetyl-CoA level. Science Advances, 2020, 6, eabb2529.	10.3	62
18	CAMSAP1 breaks the homeostatic microtubule network to instruct neuronal polarity. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 22193-22203.	7.1	18

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19	Evaluation of the Potential Risk of Advanced Peak Determination in Distorting Isobaric Labelingâ€Based Singleâ€Shot Proteome Quantitation. Proteomics, 2020, 20, 1900255.	2.2	0
20	A Kinase–Phosphatase–Transcription Factor Module Regulates Adventitious Root Emergence in Arabidopsis Root–Hypocotyl Junctions. Molecular Plant, 2020, 13, 1162-1177.	8.3	13
21	Control of Grain Size and Weight by the GSK2-LARGE1/OML4 Pathway in Rice. Plant Cell, 2020, 32, 1905-1918.	6.6	61
22	CDK4/6 regulate lysosome biogenesis through TFEB/TFE3. Journal of Cell Biology, 2020, 219, .	5 . 2	70
23	Overdosage of Balanced Protein Complexes Reduces Proliferation Rate in Aneuploid Cells. Cell Systems, 2019, 9, 129-142.e5.	6.2	32
24	An unreported NB ―LRR protein SUT 1 is required for the autoimmune response mediated by type one protein phosphatase 4 mutation (topp4â€1) in Arabidopsis. Plant Journal, 2019, 100, 357-373.	5.7	17
25	mTERF5 Acts as a Transcriptional Pausing Factor to Positively Regulate Transcription of Chloroplast psbEFLJ. Molecular Plant, 2019, 12, 1259-1277.	8.3	53
26	Nitration-induced ubiquitination and degradation control quality of ERK1. Biochemical Journal, 2019, 476, 1911-1926.	3.7	9
27	Nitrate–NRT1.1B–SPX4 cascade integrates nitrogen and phosphorus signalling networks in plants. Nature Plants, 2019, 5, 401-413.	9.3	263
28	Extensive protein S-nitrosylation associated with human pancreatic ductal adenocarcinoma pathogenesis. Cell Death and Disease, 2019, 10, 914.	6. 3	31
29	The \hat{I}^25 subunit is essential for intact 26S proteasome assembly to specifically promote plant autotrophic growth under salt stress. New Phytologist, 2019, 221, 1359-1368.	7.3	32
30	BRI1 and BAK1 interact with G proteins and regulate sugar-responsive growth and development in Arabidopsis. Nature Communications, 2018, 9, 1522.	12.8	65
31	Systematic identification of light-regulated cold-responsive proteome in a model cyanobacterium. Journal of Proteomics, 2018, 179, 100-109.	2.4	6
32	Substrate-independent immunomodulatory characteristics of mesenchymal stem cells in three-dimensional culture. PLoS ONE, 2018, 13, e0206811.	2.5	21
33	Phosphorylation of serine/arginineâ€rich splicing factor 1 at tyrosine 19 promotes cell proliferation in pediatric acute lymphoblastic leukemia. Cancer Science, 2018, 109, 3805-3815.	3.9	10
34	cTAGE5/MEA6 plays a critical role in neuronal cellular components trafficking and brain development. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E9449-E9458.	7.1	18
35	Activation of the Oxidative Pentose Phosphate Pathway is Critical for Photomixotrophic Growth of a <i>hik33</i> êDeletion Mutant of <i>Synechocystis</i>	2.2	1
36	Translation repression by maternal RNA binding protein zar1 is essential for early oogenesis in zebrafish. Development (Cambridge), 2017, 144, 128-138.	2.5	45

#	Article	IF	CITATIONS
37	Translating Divergent Environmental Stresses into a Common Proteome Response through Hik33 in a Model Cyanobacterium. Molecular and Cellular Proteomics, 2017, , mcp.M117.068080.	3.8	O
38	A Light Harvesting Complex-Like Protein in Maintenance of Photosynthetic Components in Chlamydomonas. Plant Physiology, 2017, 174, 2419-2433.	4.8	21
39	Translating Divergent Environmental Stresses into a Common Proteome Response through the Histidine Kinase 33 (Hik33) in a Model Cyanobacterium. Molecular and Cellular Proteomics, 2017, 16, 1258-1274.	3.8	26
40	Trophic Mode-Dependent Proteomic Analysis Reveals Functional Significance of Light-Independent Chlorophyll Synthesis in Synechocystis sp. PCC 6803. Molecular Plant, 2017, 10, 73-85.	8.3	22
41	Ablation of SNX6 leads to defects in synaptic function of CA1 pyramidal neurons and spatial memory. ELife, 2017, 6, .	6.0	18
42	SCFSAP controls organ size by targeting PPD proteins for degradation in Arabidopsis thaliana. Nature Communications, 2016, 7, 11192.	12.8	77
43	Quantitative profiling of spreading-coupled protein tyrosine phosphorylation in migratory cells. Scientific Reports, 2016, 6, 31811.	3.3	6
44	Protein kinase C controls lysosome biogenesis independently of mTORC1. Nature Cell Biology, 2016, 18, 1065-1077.	10.3	265
45	Comparative proteome analysis of saccular intracranial aneurysms with iTRAQ quantitative proteomics. Journal of Proteomics, 2016, 130, 120-128.	2.4	19
46	E3 ubiquitin ligase gene <i> <scp>CMPG</scp>1â€"V</i> from <i> Haynaldia villosa</i> L. contributes to powdery mildew resistance in common wheat (<i> Triticum aestivum</i> L.). Plant Journal, 2015, 84, 154-168.	5.7	52
47	Site-Specific Nitrosoproteomic Identification of Endogenously <i>S</i> -Nitrosylated Proteins in Arabidopsis. Plant Physiology, 2015, 167, 1731-1746.	4.8	202
48	Toward the complete proteome of Synechocystis sp. PCC 6803. Photosynthesis Research, 2015, 126, 203-219.	2.9	29
49	Systematically Ranking the Tightness of Membrane Association for Peripheral Membrane Proteins (PMPs) *. Molecular and Cellular Proteomics, 2015, 14, 340-353.	3.8	17
50	Aged monkey brains reveal the role of ubiquitin-conjugating enzyme UBE2N in the synaptosomal accumulation of mutant huntingtin. Human Molecular Genetics, 2015, 24, 1350-1362.	2.9	20
51	Seipin Promotes Adipose Tissue Fat Storage through the ER Ca2+-ATPase SERCA. Cell Metabolism, 2014, 19, 861-871.	16.2	132
52	OsRFPH2-10, a RING-H2 Finger E3 Ubiquitin Ligase, Is Involved in Rice Antiviral Defense in the Early Stages of Rice dwarf virus Infection. Molecular Plant, 2014, 7, 1057-1060.	8.3	33