

Xu Li

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

2,472
citations

394421

19
h-index

526287

27
g-index

27
all docs

27
docs citations

27
times ranked

4882
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA damage accumulation in aging brain and its links to Alzheimer's disease progression. <i>Genome Instability & Disease</i> , 2022, 3, 172-178.	1.1	2
2	Identification of an Ultrathin Osteochondral Interface Tissue with Specific Nanostructure at the Human Knee Joint. <i>Nano Letters</i> , 2022, 22, 2309-2319.	9.1	18
3	Protocol for establishing a protein-protein interaction network using tandem affinity purification followed by mass spectrometry in mammalian cells. <i>STAR Protocols</i> , 2022, 3, 101569.	1.2	6
4	A phosphatidic acid-binding lncRNA SNHG9 facilitates LATS1 liquid-liquid phase separation to promote oncogenic YAP signaling. <i>Cell Research</i> , 2021, 31, 1088-1105.	12.0	72
5	Two Novel Pathogenic Variants of TJP2 Gene and the Underlying Molecular Mechanisms in Progressive Familial Intrahepatic Cholestasis Type 4 Patients. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 661599.	3.7	4
6	AXL is a candidate receptor for SARS-CoV-2 that promotes infection of pulmonary and bronchial epithelial cells. <i>Cell Research</i> , 2021, 31, 126-140.	12.0	356
7	Low-density-lipoprotein-receptor-related protein 1 mediates Notch pathway activation. <i>Developmental Cell</i> , 2021, 56, 2902-2919.e8.	7.0	22
8	Elucidation of WW domain ligand binding specificities in the Hippo pathway reveals STXBP4 as YAP inhibitor. <i>EMBO Journal</i> , 2020, 39, e102406.	7.8	23
9	FOXK1 Participates in DNA Damage Response by Controlling 53BP1 Function. <i>Cell Reports</i> , 2020, 32, 108018.	6.4	13
10	MAP4K Interactome Reveals STRN4 as a Key STRIPAK Complex Component in Hippo Pathway Regulation. <i>Cell Reports</i> , 2020, 32, 107860.	6.4	34
11	Tankyrase disrupts metabolic homeostasis and promotes tumorigenesis by inhibiting LKB1-AMPK signalling. <i>Nature Communications</i> , 2019, 10, 4363.	12.8	61
12	SHROOM2 inhibits tumor metastasis through RhoA-ROCK pathway-dependent and -independent mechanisms in nasopharyngeal carcinoma. <i>Cell Death and Disease</i> , 2019, 10, 58.	6.3	40
13	Glioblastoma stem cell-derived exosomes induce M2 macrophages and PD-L1 expression on human monocytes. <i>Oncolmmunology</i> , 2018, 7, e1412909.	4.6	247
14	BAP1 links metabolic regulation of ferroptosis to tumour suppression. <i>Nature Cell Biology</i> , 2018, 20, 1181-1192.	10.3	565
15	Clustered, Regularly Interspaced Short Palindromic Repeats (CRISPR)/Cas9-coupled Affinity Purification/Mass Spectrometry Analysis Revealed a Novel Role of Neurofibromin in mTOR Signaling. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 594-607.	3.8	13
16	Recent progress in mass spectrometry proteomics for biomedical research. <i>Science China Life Sciences</i> , 2017, 60, 1093-1113.	4.9	97
17	Proteomic Analysis of the Human Tankyrase Protein Interaction Network Reveals Its Role in Pexophagy. <i>Cell Reports</i> , 2017, 20, 737-749.	6.4	69
18	A transcriptional coregulator, SPIN1-DOC, attenuates the coactivator activity of Spindlin1. <i>Journal of Biological Chemistry</i> , 2017, 292, 20808-20817.	3.4	28

#	ARTICLE	IF	CITATIONS
19	FOXR2 Interacts with MYC to Promote Its Transcriptional Activities and Tumorigenesis. <i>Cell Reports</i> , 2016, 16, 487-497.	6.4	28
20	Defining the Protein-Protein Interaction Network of the Human Protein Tyrosine Phosphatase Family. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 3030-3044.	3.8	41
21	FOXKs Promote Wnt/ β -Catenin Signaling by Translocating DVL into the Nucleus. <i>Developmental Cell</i> , 2015, 32, 707-718.	7.0	106
22	Proteomic analyses reveal distinct chromatin-associated and soluble transcription factor complexes. <i>Molecular Systems Biology</i> , 2015, 11, 775.	7.2	121
23	Tankyrase Inhibitors Target YAP by Stabilizing Angiomotin Family Proteins. <i>Cell Reports</i> , 2015, 13, 524-532.	6.4	160
24	From pathways to networks: Connecting dots by establishing protein-protein interaction networks in signaling pathways using affinity purification and mass spectrometry. <i>Proteomics</i> , 2015, 15, 188-202.	2.2	20
25	Proteomic Analysis of the Human Cyclin-dependent Kinase Family Reveals a Novel CDK5 Complex Involved in Cell Growth and Migration. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 2986-3000.	3.8	34
26	Defining the Protein-Protein Interaction Network of the Human Hippo Pathway. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 119-131.	3.8	126
27	PTPN14 is required for the density-dependent control of YAP1. <i>Genes and Development</i> , 2012, 26, 1959-1971.	5.9	166