

Liang Xu

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

2,794
citations

218677

26
h-index

265206

42
g-index

44
all docs

44
docs citations

44
times ranked

5672
citing authors

#	ARTICLE	IF	CITATIONS
1	The aberrant upregulation of exon 10-inclusive SREK1 through SRSF10 acts as an oncogenic driver in human hepatocellular carcinoma. <i>Nature Communications</i> , 2022, 13, 1363.	12.8	20
2	MNK1 and MNK2 enforce expression of E2F1, FOXM1, and WEE1 to drive soft tissue sarcoma. <i>Oncogene</i> , 2021, 40, 1851-1867.	5.9	11
3	PON2 subverts metabolic gatekeeper functions in B cells to promote leukemogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	10
4	Topography of transcriptionally active chromatin in glioblastoma. <i>Science Advances</i> , 2021, 7, .	10.3	19
5	Core transcriptional regulatory circuitries in cancer. <i>Oncogene</i> , 2020, 39, 6633-6646.	5.9	41
6	Multiple myeloma: Combination therapy of BET proteolysis targeting chimeric molecule with CDK9 inhibitor. <i>PLoS ONE</i> , 2020, 15, e0232068.	2.5	12
7	TP63, SOX2, and KLF5 Establish a Core Regulatory Circuitry That Controls Epigenetic and Transcription Patterns in Esophageal Squamous Cell Carcinoma Cell Lines. <i>Gastroenterology</i> , 2020, 159, 1311-1327.e19.	1.3	92
8	ETV4 is a theranostic target in clear cell renal cell carcinoma that promotes metastasis by activating the pro-metastatic gene FOSL1 in a PI3K-AKT dependent manner. <i>Cancer Letters</i> , 2020, 482, 74-89.	7.2	19
9	SOX7 regulates MAPK/ERK-BIM mediated apoptosis in cancer cells. <i>Oncogene</i> , 2019, 38, 6196-6210.	5.9	32
10	<p>Candidate tumor suppressor gene IRF6 is involved in human breast cancer pathogenesis via modulating PI3K-regulatory subunit PIK3R2 expression</p>. <i>Cancer Management and Research</i> , 2019, Volume 11, 5557-5572.	1.9	14
11	LNK suppresses interferon signaling in melanoma. <i>Nature Communications</i> , 2019, 10, 2230.	12.8	21
12	Bromodomain and extraterminal proteins foster the core transcriptional regulatory programs and confer vulnerability in liposarcoma. <i>Nature Communications</i> , 2019, 10, 1353.	12.8	39
13	The Upregulation of Trophinin-Associated Protein (TROAP) Predicts a Poor Prognosis in Hepatocellular Carcinoma. <i>Journal of Cancer</i> , 2019, 10, 957-967.	2.5	31
14	Proteolysis targeting chimeric molecules as therapy for multiple myeloma: efficacy, biomarker and drug combinations. <i>Haematologica</i> , 2019, 104, 1209-1220.	3.5	30
15	Along with its favorable prognostic role, CLCA2 inhibits growth and metastasis of nasopharyngeal carcinoma cells via inhibition of FAK/ERK signaling. <i>Journal of Experimental and Clinical Cancer Research</i> , 2018, 37, 34.	8.6	33
16	Global expression profiling and pathway analysis of mouse mammary tumor reveals strain and stage specific dysregulated pathways in breast cancer progression. <i>Cell Cycle</i> , 2018, 17, 963-973.	2.6	6
17	dbCoRC: a database of core transcriptional regulatory circuitries modeled by H3K27ac ChIP-seq signals. <i>Nucleic Acids Research</i> , 2018, 46, D71-D77.	14.5	37
18	Targeting the vulnerability to NAD+ depletion in B-cell acute lymphoblastic leukemia. <i>Leukemia</i> , 2018, 32, 616-625.	7.2	29

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19	<scp>ROCK</scp> â€dependent phosphorylation of <scp>NUP</scp> 62 regulates p63 nuclear transport and squamous cell carcinoma proliferation. EMBO Reports, 2018, 19, 73-88.	4.5	56
20	The developmental transcription factor IRF6 attenuates ABCG2 gene expression and distinctively reverses stemness phenotype in nasopharyngeal carcinoma. Cancer Letters, 2018, 431, 230-243.	7.2	31
21	Functional Genome-wide Screening Identifies Targets and Pathways Sensitizing Pancreatic Cancer Cells to Dasatinib. Journal of Cancer, 2018, 9, 4762-4773.	2.5	25
22	Profiling the B/T cell receptor repertoire of lymphocyte derived cell lines. BMC Cancer, 2018, 18, 940.	2.6	10
23	Co-activation of super-enhancer-driven CCAT1 by TP63 and SOX2 promotes squamous cancer progression. Nature Communications, 2018, 9, 3619.	12.8	179
24	Targetable BET proteins- and E2F1-dependent transcriptional program maintains the malignancy of glioblastoma. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E5086-E5095.	7.1	87
25	A comparison of weekly versus 3-weekly cisplatin during concurrent chemoradiotherapy for locoregionally advanced nasopharyngeal carcinoma using intensity modulated radiation therapy: a matched study. Journal of Cancer, 2018, 9, 92-99.	2.5	17
26	Mutational profiling of acute lymphoblastic leukemia with testicular relapse. Journal of Hematology and Oncology, 2017, 10, 65.	17.0	16
27	BCL6 promotes glioma and serves as a therapeutic target. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 3981-3986.	7.1	58
28	Targeting super-enhancer-associated oncogenes in oesophageal squamous cell carcinoma. Gut, 2017, 66, 1358-1368.	12.1	169
29	ZNF750 is a lineage-specific tumour suppressor in squamous cell carcinoma. Oncogene, 2017, 36, 2243-2254.	5.9	90
30	Mutational Landscape of Pediatric Acute Lymphoblastic Leukemia. Cancer Research, 2017, 77, 390-400.	0.9	77
31	Extracellular serglycin upregulates the CD44 receptor in an autocrine manner to maintain self-renewal in nasopharyngeal carcinoma cells by reciprocally activating the MAPK/Î²-catenin axis. Cell Death and Disease, 2016, 7, e2456-e2456.	6.3	47
32	Tumor vasculogenic mimicry predicts poor prognosis in cancer patients: a meta-analysis. Angiogenesis, 2016, 19, 191-200.	7.2	100
33	CRM1 Inhibition Promotes Cytotoxicity in Ewing Sarcoma Cells by Repressing EWS-FLI1â€Dependent IGF-1 Signaling. Cancer Research, 2016, 76, 2687-2697.	0.9	29
34	EB-virus latent membrane protein 1 potentiates the stemness of nasopharyngeal carcinoma via preferential activation of PI3K/AKT pathway by a positive feedback loop. Oncogene, 2016, 35, 3419-3431.	5.9	52
35	PDZ binding kinase (PBK) is a theranostic target for nasopharyngeal carcinoma: driving tumor growth via ROS signaling and correlating with patient survival. Oncotarget, 2016, 7, 26604-26616.	1.8	23
36	Genomic and Functional Analysis of the E3 Ligase PARK2 in Glioma. Cancer Research, 2015, 75, 1815-1827.	0.9	50

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37	SETDB1 accelerates tumorigenesis by regulating the WNT signalling pathway. <i>Journal of Pathology</i> , 2015, 235, 559-570.	4.5	64
38	An emerging role of PARK2 in cancer. <i>Journal of Molecular Medicine</i> , 2014, 92, 31-42.	3.9	88
39	Genomic and molecular characterization of esophageal squamous cell carcinoma. <i>Nature Genetics</i> , 2014, 46, 467-473.	21.4	523
40	The genomic landscape of nasopharyngeal carcinoma. <i>Nature Genetics</i> , 2014, 46, 866-871.	21.4	317
41	Design of hybrid β -hairpin peptides with enhanced cell specificity and potent anti-inflammatory activity. <i>Biomaterials</i> , 2013, 34, 237-250.	11.4	128
42	Genomic and functional characterizations of phosphodiesterase subtype 4D in human cancers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6109-6114.	7.1	59