

# Xujun Liang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7994009/publications.pdf>

Version: 2024-02-01

19  
papers

353  
citations

1040056

9  
h-index

839539

18  
g-index

19  
all docs

19  
docs citations

19  
times ranked

464  
citing authors

#	ARTICLE	IF	CITATIONS
1	LRSSL: predict and interpret drug-disease associations based on data integration using sparse subspace learning. <i>Bioinformatics</i> , 2017, 33, 1187-1196.	4.1	106
2	Structural basis for DNA recognition by FOXC2. <i>Nucleic Acids Research</i> , 2019, 47, 3752-3764.	14.5	36
3	Identifying DCN and HSPD1 as Potential Biomarkers in Colon Cancer Using 2D-LC-MS/MS Combined with iTRAQ Technology. <i>Journal of Cancer</i> , 2017, 8, 479-489.	2.5	35
4	Mechanism of forkhead transcription factors binding to a novel palindromic DNA site. <i>Nucleic Acids Research</i> , 2021, 49, 3573-3583.	14.5	28
5	Structural basis of binding of homodimers of the nuclear receptor NR4A2 to selective Nur-responsive DNA elements. <i>Journal of Biological Chemistry</i> , 2019, 294, 19795-19803.	3.4	23
6	lncRNA RP11-624L4.1 Is Associated with Unfavorable Prognosis and Promotes Proliferation via the CDK4/6-Cyclin D1-Rb-E2F1 Pathway in NPC. <i>Molecular Therapy - Nucleic Acids</i> , 2020, 22, 1025-1039.	5.1	20
7	The Significance of Serum S100A9 and TNC Levels as Biomarkers in Colorectal Cancer. <i>Journal of Cancer</i> , 2019, 10, 5315-5323.	2.5	19
8	Quantitative proteomic profiling of tumor-associated vascular endothelial cells in colorectal cancer. <i>Biology Open</i> , 2019, 8, .	1.2	14
9	DNA-binding properties of FOXP3 transcription factor. <i>Acta Biochimica Et Biophysica Sinica</i> , 2017, 49, 792-799.	2.0	12
10	Identification of VWF as a Novel Biomarker in Lung Adenocarcinoma by Comprehensive Analysis. <i>Frontiers in Oncology</i> , 2021, 11, 639600.	2.8	10
11	S-Adenosylmethionine Affects Cell Cycle Pathways and Suppresses Proliferation in Liver Cells. <i>Journal of Cancer</i> , 2019, 10, 4368-4379.	2.5	9
12	Simultaneous Inhibition of EGFR and HER2 via Afatinib Augments the Radiosensitivity of Nasopharyngeal Carcinoma Cells. <i>Journal of Cancer</i> , 2019, 10, 2063-2073.	2.5	8
13	Learning important features from multi-view data to predict drug side effects. <i>Journal of Cheminformatics</i> , 2019, 11, 79.	6.1	8
14	Identification of prognostic genes in the pancreatic adenocarcinoma immune microenvironment by integrated bioinformatics analysis. <i>Cancer Immunology, Immunotherapy</i> , 2022, 71, 1757-1769.	4.2	8
15	Coupling of Cell Surface Biotinylation and SILAC-Based Quantitative Proteomics Identified Myoferlin as a Potential Therapeutic Target for Nasopharyngeal Carcinoma Metastasis. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 621810.	3.7	6
16	Identification of a pyroptosis-related lncRNA risk model for predicting prognosis and immune response in colon adenocarcinoma. <i>World Journal of Surgical Oncology</i> , 2022, 20, 118.	1.9	4
17	Mechanistic Insights into the Preference for Tandem Binding Sites in DNA Recognition by FOXM1. <i>Journal of Molecular Biology</i> , 2022, 434, 167426.	4.2	3
18	A novel machine learning model based on sparse structure learning with adaptive graph regularization for predicting drug side effects. <i>Journal of Biomedical Informatics</i> , 2022, 132, 104131.	4.3	3

#	ARTICLE	IF	CITATIONS
19	Detection and Analysis of RNAs Expression Profile for Methylated Candidate Tumor Suppressor Genes in Nasopharyngeal Carcinoma. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2019, 19, 772-782.	1.7	1