

# Zhaoqi Liu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4214820/publications.pdf>

Version: 2024-02-01

18  
papers

1,555  
citations

687363

13  
h-index

940533

16  
g-index

19  
all docs

19  
docs citations

19  
times ranked

3510  
citing authors

#	ARTICLE	IF	CITATIONS
1	SF3B1 mutant-induced missplicing of MAP3K7 causes anemia in myelodysplastic syndromes. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	26
2	The UVSSA complex alleviates MYC-driven transcription stress. Journal of Cell Biology, 2021, 220, .	5.2	6
3	Computing the Role of Alternative Splicing in Cancer. Trends in Cancer, 2021, 7, 347-358.	7.4	19
4	iPLA2 <sup>2</sup> -mediated lipid detoxification controls p53-driven ferroptosis independent of GPX4. Nature Communications, 2021, 12, 3644.	12.8	153
5	Transcriptional control of CBX5 by the RNA-binding proteins RBMX and RBMXL1 maintains chromatin state in myeloid leukemia. Nature Cancer, 2021, 2, 741-757.	13.2	10
6	Mutant SF3B1 promotes AKT- and NF- $\kappa$ B-driven mammary tumorigenesis. Journal of Clinical Investigation, 2021, 131, .	8.2	22
7	A Novel JAK1 Mutant Breast Implant-Associated Anaplastic Large Cell Lymphoma Patient-Derived Xenograft Fostering Pre-Clinical Discoveries. Cancers, 2020, 12, 1603.	3.7	11
8	Pan-cancer analysis identifies mutations in <i>SUGP1</i> that recapitulate mutant SF3B1 splicing dysregulation. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 10305-10312.	7.1	30
9	Genomic Characterization of HIV-Associated Plasmablastic Lymphoma Identifies Pervasive Mutations in the JAK-STAT Pathway. Blood Cancer Discovery, 2020, 1, 112-125.	5.0	40
10	Transcriptional Control of HP1a By the RNA Binding Proteins RbmX/L1 Maintain Chromatin State in Myeloid Leukemia. Blood, 2020, 136, 15-15.	1.4	0
11	Disease-Causing Mutations in SF3B1 Alter Splicing by Disrupting Interaction with SUGP1. Molecular Cell, 2019, 76, 82-95.e7.	9.7	84
12	Secretome analysis of patient-derived GBM tumor spheres identifies midkine as a potent therapeutic target. Experimental and Molecular Medicine, 2019, 51, 1-11.	7.7	28
13	Comprehensive characterisation of compartment-specific long non-coding RNAs associated with pancreatic ductal adenocarcinoma. Gut, 2019, 68, 499-511.	12.1	39
14	Saa3 is a key mediator of the protumorigenic properties of cancer-associated fibroblasts in pancreatic tumors. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1147-E1156.	7.1	128
15	Pharmacogenomic landscape of patient-derived tumor cells informs precision oncology therapy. Nature Genetics, 2018, 50, 1399-1411.	21.4	145
16	Mutations in the RNA Splicing Factor SF3B1 Promote Transformation through MYC Stabilization. Blood, 2018, 132, 882-882.	1.4	0
17	Spatiotemporal genomic architecture informs precision oncology in glioblastoma. Nature Genetics, 2017, 49, 594-599.	21.4	223
18	Clonal evolution of glioblastoma under therapy. Nature Genetics, 2016, 48, 768-776.	21.4	591