

# Shuwei Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5758830/publications.pdf>

Version: 2024-02-01

35  
papers

773  
citations

623188

14  
h-index

552369

26  
g-index

37  
all docs

37  
docs citations

37  
times ranked

1053  
citing authors

#	ARTICLE	IF	CITATIONS
1	LncRNA MT1JP functions as a ceRNA in regulating FBXW7 through competitively binding to miR-92a-3p in gastric cancer. <i>Molecular Cancer</i> , 2018, 17, 87.	7.9	218
2	Association of genetic variants in lncRNA <i>H19</i> with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016, 7, 25470-25477.	0.8	90
3	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction. <i>Molecular Cancer</i> , 2022, 21, 49.	7.9	72
4	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020, 41, 8-17.	1.3	38
5	Genome-wide long non-coding RNAs identified a panel of novel plasma biomarkers for gastric cancer diagnosis. <i>Gastric Cancer</i> , 2019, 22, 731-741.	2.7	37
6	The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021, 14, 93.	6.9	31
7	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018, 92, 2681-2689.	1.9	30
8	Combinations of single nucleotide polymorphisms identified in genome-wide association studies determine risk for colorectal cancer. <i>International Journal of Cancer</i> , 2019, 145, 2661-2669.	2.3	25
9	Remote modulation of lncRNA <i>GCLET</i> by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. <i>Science Advances</i> , 2020, 6, eaay5525.	4.7	23
10	Systematic evaluation of the effects of genetic variants on PIWI-interacting RNA expression across 33 cancer types. <i>Nucleic Acids Research</i> , 2021, 49, 90-97.	6.5	22
11	Genetic variants in <i>SLC22A3</i> contribute to the susceptibility to colorectal cancer. <i>International Journal of Cancer</i> , 2019, 145, 154-163.	2.3	21
12	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020, 137, 105543.	4.8	16
13	Genetic variants in RPA1 associated with the response to oxaliplatin-based chemotherapy in colorectal cancer. <i>Journal of Gastroenterology</i> , 2019, 54, 939-949.	2.3	15
14	Association study of genetic variants in estrogen metabolic pathway genes and colorectal cancer risk and survival. <i>Archives of Toxicology</i> , 2018, 92, 1991-1999.	1.9	14
15	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017, 8, 1355-1361.	1.2	12
16	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018, 673, 174-180.	1.0	12
17	Association study between genetic variants in retinol metabolism pathway genes and prostate cancer risk. <i>Cancer Medicine</i> , 2020, 9, 9462-9470.	1.3	11
18	A genetic variant located in the miR-532-5p-binding site of TGFBR1 is associated with the colorectal cancer risk. <i>Journal of Gastroenterology</i> , 2019, 54, 141-148.	2.3	9

#	ARTICLE	IF	CITATIONS
19	Genetic variant in miRâ€21 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019, 23, 2012-2019.	1.6	9
20	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018, 33, 82-87.	2.7	7
21	Functional genetic variant of <i>HSD17B12</i> in the fatty acid biosynthesis pathway predicts the outcome of colorectal cancer. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 14160-14170.	1.6	6
22	Association of genetic variants in autophagy-lysosome pathway genes with susceptibility and survival to prostate cancer. <i>Gene</i> , 2022, 808, 145953.	1.0	6
23	LncRNAâ€422 suppresses the proliferation and growth of colorectal cancer cells by targeting SFPQ. <i>Clinical and Translational Medicine</i> , 2022, 12, e664.	1.7	6
24	Novel CpG-SNPs in the gastric acid secretion pathway <i>GNAI3</i> and susceptibility to gastric cancer. <i>Gene</i> , 2020, 736, 144447.	1.0	5
25	Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of <i>IQCB1</i> expression. <i>Archives of Toxicology</i> , 2021, 95, 2053-2063.	1.9	5
26	CoSMed: a user-friendly web server to estimate 5-year survival probability of left-sided and right-sided colorectal cancer patients using molecular data. <i>Bioinformatics</i> , 2021, 38, 278-281.	1.8	5
27	Genetic variants in Ras/Raf/MEK/ERK pathway are associated with gastric cancer risk in Chinese Han population. <i>Archives of Toxicology</i> , 2020, 94, 2683-2690.	1.9	4
28	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021, 95, 2523-2532.	1.9	4
29	High-density lipoprotein, low-density lipoprotein and triglyceride levels and upper gastrointestinal cancers risk: a trans-ancestry Mendelian randomization study. <i>European Journal of Clinical Nutrition</i> , 2022, , .	1.3	4
30	Genetic variants in <i>circTUBB</i> interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020, 94, 325-333.	1.9	3
31	A transcriptomic study for identifying cardiaâ€and nonâ€cardiaâ€specific gastric cancer prognostic factors using genetic algorithmâ€based methods. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 9457-9465.	1.6	3
32	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. <i>Archives of Toxicology</i> , 2021, 95, 271-281.	1.9	3
33	Genetic variants in the cholesterol biosynthesis pathway genes and risk of prostate cancer. <i>Gene</i> , 2021, 774, 145432.	1.0	3
34	Genome-Wide Association Analyses Identify <i>CATSPERE</i> as a Mediator of Colorectal Cancer Susceptibility and Progression. <i>Cancer Research</i> , 2022, 82, 986-997.	0.4	3
35	Genetic variants in the Folic acid Metabolic Pathway Genes predict outcomes of metastatic Colorectal Cancer patients receiving first-line Chemotherapy. <i>Journal of Cancer</i> , 2020, 11, 6507-6515.	1.2	1