Shuwei Li

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

32	453 citations	10	2 O
papers		h-index	g-index
37 ext. papers	620 ext. citations	8.9 avg, IF	3.49 L-index

#	Paper	IF	Citations
32	LncRNA-422 suppresses the proliferation and growth of colorectal cancer cells by targeting SFPQ <i>Clinical and Translational Medicine</i> , 2022 , 12, e664	5.7	1
31	Association of genetic variants in autophagy-lysosome pathway genes with susceptibility and survival to prostate cancer. <i>Gene</i> , 2022 , 808, 145953	3.8	2
30	Exosomal circLPAR1 functions in colorectal cancer diagnosis and tumorigenesis through suppressing BRD4 via METTL3-eIF3h interaction <i>Molecular Cancer</i> , 2022 , 21, 49	42.1	6
29	Genetic variants in the cholesterol biosynthesis pathway genes and risk of prostate cancer. <i>Gene</i> , 2021 , 774, 145432	3.8	2
28	Genetic variations in the CTLA-4 immune checkpoint pathway are associated with colon cancer risk, prognosis, and immune infiltration via regulation of IQCB1 expression. <i>Archives of Toxicology</i> , 2021 , 95, 2053-2063	5.8	O
27	Evaluation of common genetic variants in vitamin E-related pathway genes and colorectal cancer susceptibility. <i>Archives of Toxicology</i> , 2021 , 95, 2523-2532	5.8	1
26	The biogenesis and biological function of PIWI-interacting RNA in cancer. <i>Journal of Hematology and Oncology</i> , 2021 , 14, 93	22.4	3
25	Genetic variants in Hippo signalling pathway-related genes affect the risk of colorectal cancer. <i>Archives of Toxicology</i> , 2021 , 95, 271-281	5.8	2
24	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	20.1	8
23	Genetic variants in m6A modification genes are associated with colorectal cancer risk. <i>Carcinogenesis</i> , 2020 , 41, 8-17	4.6	30
22	Functional genetic variant of HSD17B12 in the fatty acid biosynthesis pathway predicts the outcome of colorectal cancer. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 14160-14170	5.6	1
21	Remote modulation of lncRNA by risk variant at 16p13 underlying genetic susceptibility to gastric cancer. <i>Science Advances</i> , 2020 , 6, eaay5525	14.3	10
20	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	5.8	2
19	A transcriptomic study for identifying cardia- and non-cardia-specific gastric cancer prognostic factors using genetic algorithm-based methods. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 9457-9465	5.6	2
18	Sex hormones and genetic variants in hormone metabolic pathways associated with the risk of colorectal cancer. <i>Environment International</i> , 2020 , 137, 105543	12.9	7
17	Novel CpG-SNPs in the gastric acid secretion pathway GNAI3 and susceptibility to gastric cancer. <i>Gene</i> , 2020 , 736, 144447	3.8	3
16	Genetic variants in circTUBB interacting with smoking can enhance colorectal cancer risk. <i>Archives of Toxicology</i> , 2020 , 94, 325-333	5.8	2

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15	Association study between genetic variants in retinol metabolism pathway genes and prostate cancer risk. <i>Cancer Medicine</i> , 2020 , 9, 9462-9470	4.8	7
14	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	4.5	
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	6.9	8
12	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	7.5	12
11	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	6.9	9
10	Genetic variants in SLC22A3 contribute to the susceptibility to colorectal cancer. <i>International Journal of Cancer</i> , 2019 , 145, 154-163	7.5	15
9	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	7.6	19
8	Genetic variant in miR-21 binding sites is associated with colorectal cancer risk. <i>Journal of Cellular and Molecular Medicine</i> , 2019 , 23, 2012-2019	5.6	6
7	Evaluation of GWAS-Identified Genetic Variants for Gastric Cancer Survival. <i>EBioMedicine</i> , 2018 , 33, 82-	- 83 .8	5
6	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	42.1	166
5	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018 , 673, 174-180	3.8	9
4	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	5.8	12
3	Circadian clock pathway genes associated with colorectal cancer risk and prognosis. <i>Archives of Toxicology</i> , 2018 , 92, 2681-2689	5.8	24
2	Plasma Mesothelin as a Novel Diagnostic and Prognostic Biomarker in Colorectal Cancer. <i>Journal of Cancer</i> , 2017 , 8, 1355-1361	4.5	10
1	Association of genetic variants in lncRNA H19 with risk of colorectal cancer in a Chinese population. <i>Oncotarget</i> , 2016 , 7, 25470-7	3.3	68