

Yue Jiang

List of Publications by Year in descending order

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Version: 2024-02-01

70
papers

2,144
citations

304368

22
h-index

264894

42
g-index

74
all docs

74
docs citations

74
times ranked

4234
citing authors

#	ARTICLE	IF	CITATIONS
1	Plasma miRNA as early biomarkers for detecting hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2015, 137, 1679-1690.	2.3	188
2	Identification of risk loci and a polygenic risk score for lung cancer: a large-scale prospective cohort study in Chinese populations. <i>Lancet Respiratory Medicine</i> , 2019, 7, 881-891.	5.2	167
3	Genomic Landscape Survey Identifies SRSF1 as a Key Oncodriver in Small Cell Lung Cancer. <i>PLoS Genetics</i> , 2016, 12, e1005895.	1.5	144
4	Systematic identification of genes with a cancer-testis expression pattern in 19 cancer types. <i>Nature Communications</i> , 2016, 7, 10499.	5.8	124
5	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. <i>Lancet Oncology</i> , 2020, 21, 1378-1386.	5.1	123
6	Associations Between Hepatitis B Virus Infection and Risk of All Cancer Types. <i>JAMA Network Open</i> , 2019, 2, e195718.	2.8	114
7	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , 2017, 66, 581-587.	6.1	68
8	Whole-genome sequencing reveals genomic signatures associated with the inflammatory microenvironments in Chinese NSCLC patients. <i>Nature Communications</i> , 2018, 9, 2054.	5.8	68
9	Estimation of heritability for nine common cancers using data from genome-wide association studies in Chinese population. <i>International Journal of Cancer</i> , 2017, 140, 329-336.	2.3	66
10	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. <i>Molecular Cancer</i> , 2021, 20, 67.	7.9	65
11	Association analysis identifies new risk loci for non-obstructive azoospermia in Chinese men. <i>Nature Communications</i> , 2014, 5, 3857.	5.8	64
12	Plasma metabolomics identified novel metabolites associated with risk of type 2 diabetes in two prospective cohorts of Chinese adults. <i>International Journal of Epidemiology</i> , 2016, 45, 1507-1516.	0.9	64
13	A cancer-testis non-coding RNA LIN28B-AS1 activates driver gene LIN28B by interacting with IGF2BP1 in lung adenocarcinoma. <i>Oncogene</i> , 2019, 38, 1611-1624.	2.6	61
14	Identification of Novel T1D Risk Loci and Their Association With Age and Islet Function at Diagnosis in Autoantibody-Positive T1D Individuals: Based on a Two-Stage Genome-Wide Association Study. <i>Diabetes Care</i> , 2019, 42, 1414-1421.	4.3	60
15	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017, 152, 2011-2021.	0.6	58
16	Evaluation of genetic variants in microRNA biosynthesis genes and risk of breast cancer in Chinese women. <i>International Journal of Cancer</i> , 2013, 133, 2216-2224.	2.3	50
17	Genetic variation in a hsa-let-7 binding site in RAD52 is associated with breast cancer susceptibility. <i>Carcinogenesis</i> , 2013, 34, 689-693.	1.3	43
18	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. <i>Gut</i> , 2020, 69, 641-651.	6.1	36

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19	Combinational therapeutic targeting of BRD4 and CDK7 synergistically induces anticancer effects in head and neck squamous cell carcinoma. <i>Cancer Letters</i> , 2020, 469, 510-523.	3.2	36
20	Hepatitis B virus genotype, mutations, human leukocyte antigen polymorphisms and their interactions in hepatocellular carcinoma: a multi-centre case-control study. <i>Scientific Reports</i> , 2015, 5, 16489.	1.6	35
21	A Screen for Genomic Disorders of Infertility Identifies MAST2 Duplications Associated with Nonobstructive Azoospermia in Humans1. <i>Biology of Reproduction</i> , 2015, 93, 61.	1.2	30
22	Genetic Variants at 1p11.2 and Breast Cancer Risk: A Two-Stage Study in Chinese Women. <i>PLoS ONE</i> , 2011, 6, e21563.	1.1	24
23	Evaluation of genetic variants in autophagy pathway genes as prognostic biomarkers for breast cancer. <i>Gene</i> , 2017, 627, 549-555.	1.0	21
24	Comprehensive characterization of functional eRNAs in lung adenocarcinoma reveals novel regulators and a prognosis-related molecular subtype. <i>Theranostics</i> , 2020, 10, 11264-11277.	4.6	20
25	Genetic Variations in the Flanking Regions of miR-101-2 Are Associated with Increased Risk of Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e86319.	1.1	19
26	Immune landscape and subtypes in primary resectable oral squamous cell carcinoma: prognostic significance and predictive of therapeutic response. , 2021, 9, e002434.		19
27	Association of KCTD10, MVK, and MMAB polymorphisms with dyslipidemia and coronary heart disease in Han Chinese population. <i>Lipids in Health and Disease</i> , 2016, 15, 171.	1.2	18
28	Circulating Tumor Cells Enriched by the Depletion of Leukocytes with Bi-Antibodies in Non-Small Cell Lung Cancer: Potential Clinical Application. <i>PLoS ONE</i> , 2015, 10, e0137076.	1.1	18
29	Predictive value of prognostic nutritional index in patients with oral squamous cell carcinoma. <i>Oral Diseases</i> , 2020, 26, 903-911.	1.5	17
30	Potentially functional polymorphisms in aminoacyl-tRNA synthetases genes are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015, 54, 577-583.	1.3	15
31	Evaluation of CpG-SNPs in miRNA promoters and risk of breast cancer. <i>Gene</i> , 2018, 651, 1-8.	1.0	15
32	Genetic variants at 9p21.3 are associated with risk of esophageal squamous cell carcinoma in a Chinese population. <i>Cancer Science</i> , 2017, 108, 250-255.	1.7	14
33	Implementation of Clinical Diagnostic Criteria and Universal Symptom Survey Contributed to Lower Magnitude and Faster Resolution of the COVID-19 Epidemic in Wuhan. <i>Engineering</i> , 2020, 6, 1141-1146.	3.2	14
34	Epigeneticâ€“smoking interaction reveals histologically heterogeneous effects of TRIM27 DNA methylation on overall survival among earlyâ€“stage NSCLC patients. <i>Molecular Oncology</i> , 2020, 14, 2759-2774.	2.1	13
35	Functional genetic variants in centrosome-related genes CEP72 and YWHAG confer susceptibility to gastric cancer. <i>Archives of Toxicology</i> , 2020, 94, 2861-2872.	1.9	13
36	Associations Between Sleep Quality and Health Span: A Prospective Cohort Study Based on 328,850 UK Biobank Participants. <i>Frontiers in Genetics</i> , 2021, 12, 663449.	1.1	13

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37	Low-frequency germline variants across 6p22.2–6p21.33 are associated with non-obstructive azoospermia in Han Chinese men. <i>Human Molecular Genetics</i> , 2015, 24, 5628-5636.	1.4	12
38	Polymorphisms in CARS are associated with gastric cancer risk: a two-stage case–control study in the Chinese population. <i>Gastric Cancer</i> , 2017, 20, 940-947.	2.7	12
39	Low-frequency nonsynonymous variants in <i>FKBP1</i> and <i>ARPC1B</i> genes are associated with breast cancer risk in Chinese women. <i>Molecular Carcinogenesis</i> , 2017, 56, 774-780.	1.3	12
40	Claudin-7 Inhibits Proliferation and Metastasis in Salivary Adenoid Cystic Carcinoma Through Wnt/ β -Catenin Signaling. <i>Cell Transplantation</i> , 2020, 29, 096368972094358.	1.2	12
41	Potentially functional variants in lncRNAs are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 2048-2057.	1.3	11
42	Integrating of genomic and transcriptomic profiles for the prognostic assessment of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2019, 175, 691-699.	1.1	11
43	Genetic characterisation of sarcomatoid carcinomas reveals multiple novel actionable mutations and identifies <i>KRAS</i> mutation as a biomarker of poor prognosis. <i>Journal of Medical Genetics</i> , 2022, 59, 10-17.	1.5	11
44	Epidemiological, clinical, and 3-dimensional CBCT radiographic characterizations of supernumerary teeth in a non-syndromic adult population: a single-institutional study from 60,104 Chinese subjects. <i>Clinical Oral Investigations</i> , 2020, 24, 4271-4281.	1.4	11
45	Epidemiological, clinical, radiographic characterization of non-syndromic supernumerary teeth in Chinese children and adolescents. <i>Oral Diseases</i> , 2021, 27, 981-992.	1.5	11
46	U-shaped association between telomere length and esophageal squamous cell carcinoma risk: a case-control study in Chinese population. <i>Frontiers of Medicine</i> , 2015, 9, 478-486.	1.5	10
47	Leukocyte telomere length, lipid parameters and gestational diabetes risk: a case-control study in a Chinese population. <i>Scientific Reports</i> , 2019, 9, 8483.	1.6	10
48	Identification of an autophagy-related prognostic signature in head and neck squamous cell carcinoma. <i>Journal of Oral Pathology and Medicine</i> , 2021, 50, 1040-1049.	1.4	10
49	Early Second-Trimester Serum MicroRNAs as Potential Biomarker for Nondiabetic Macrosomia. <i>BioMed Research International</i> , 2014, 2014, 1-6.	0.9	8
50	A cis-eQTL genetic variant of the cancer-testis gene <i>CCDC116</i> is associated with risk of multiple cancers. <i>Human Genetics</i> , 2017, 136, 987-997.	1.8	7
51	Genetic variants affecting telomere length are associated with the prognosis of esophageal squamous cell carcinoma in a Chinese population. <i>Molecular Carcinogenesis</i> , 2017, 56, 1021-1029.	1.3	7
52	Genetic Variants in the Promoter Region of <i>miR-10b</i> and the Risk of Breast Cancer. <i>BioMed Research International</i> , 2017, 2017, 1-7.	0.9	7
53	Genetic Variation in the 3'-Untranslated Region of <i>NBN</i> Gene Is Associated with Gastric Cancer Risk in a Chinese Population. <i>PLoS ONE</i> , 2015, 10, e0139059.	1.1	7
54	Peptidomic analysis of zebrafish embryos exposed to polychlorinated biphenyls and their impact on eye development. <i>Ecotoxicology and Environmental Safety</i> , 2019, 175, 164-172.	2.9	6

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55	The sonic hedgehog signaling pathway is suppressed following PCB ₁₂₅₄ exposure during retinal development. <i>Environmental Toxicology</i> , 2019, 34, 340-347.	2.1	6
56	Transcriptome-wide association study for persistent hepatitis B virus infection and related hepatocellular carcinoma. <i>Liver International</i> , 2020, 40, 2117-2127.	1.9	6
57	Functional Dissection of CD26 and Its Pharmacological Inhibition by Sitagliptin During Skin Wound Healing. <i>Medical Science Monitor</i> , 2021, 27, e928933.	0.5	5
58	Genetic variants at 10q23.33 are associated with plasma lipid levels in a Chinese population. <i>Journal of Biomedical Research</i> , 2014, 28, 53-8.	0.7	5
59	Weight Change across Adulthood in Relation to Non-Alcoholic Fatty Liver Disease among Non-Obese Individuals. <i>Nutrients</i> , 2022, 14, 2140.	1.7	5
60	Identification of Recurrent Variants in BRCA1 and BRCA2 across Multiple Cancers in the Chinese Population. <i>BioMed Research International</i> , 2020, 2020, 1-11.	0.9	4
61	Sex-Specific Associations of Testosterone and Genetic Factors With Health Span. <i>Frontiers in Endocrinology</i> , 2021, 12, 773464.	1.5	4
62	Pharmacological inhibition of CDK7 by THZ1 impairs tumor growth in p53-mutated HNSCC. <i>Oral Diseases</i> , 2022, 28, 611-620.	1.5	3
63	Investigation of Causal Effect of Type 2 Diabetes Mellitus on Lung Cancer: A Mendelian Randomization Study. <i>Frontiers in Genetics</i> , 2021, 12, 673687.	1.1	3
64	The rs6771157 C/G polymorphism in SCN10A is associated with the risk of atrial fibrillation in a Chinese Han population. <i>Scientific Reports</i> , 2016, 6, 35212.	1.6	2
65	Prognostic role of CD11b+ myeloid-derived suppressor cells in oral squamous cell carcinoma. <i>Archives of Medical Science</i> , 2021, , .	0.4	2
66	Functional characterization and clinical significance of super-enhancers in lung adenocarcinoma. <i>Molecular Carcinogenesis</i> , 2022, 61, 776-786.	1.3	2
67	Genetic Variations in miR-30 Family Member Regulatory Regions Are Associated with Breast Cancer Risk in a Chinese Population. <i>BioMed Research International</i> , 2020, 2020, 1-8.	0.9	1
68	The lab-confirmed interval of COVID-19 clusters and its application in the strength evaluation of prevention and control measures. <i>BMC Infectious Diseases</i> , 2021, 21, 226.	1.3	1
69	Potential functional variants of KIAA genes are associated with breast cancer risk in a case control study. <i>Annals of Translational Medicine</i> , 2021, 9, 549-549.	0.7	1
70	Coding variants in the PCNT and CEP295 genes contribute to breast cancer risk in Chinese women. <i>Pathology Research and Practice</i> , 2021, 225, 153581.	1.0	1