

Yue Jiang

List of Publications by Year in Descending Order

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Version: 2024-04-20

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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.

67
papers

1,247
citations

19
h-index

34
g-index

74
ext. papers

1,736
ext. citations

7.4
avg, IF

3.72
L-index

#	Paper	IF	Citations
67	Weight Change across Adulthood in Relation to Non-Alcoholic Fatty Liver Disease among Non-Obese Individuals. <i>Nutrients</i> , 2022 , 14, 2140	6.7	1
66	Sex-Specific Associations of Testosterone and Genetic Factors With Health Span.. <i>Frontiers in Endocrinology</i> , 2021 , 12, 773464	5.7	1
65	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. <i>Molecular Cancer</i> , 2021 , 20, 67	42.1	13
64	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	3.2	
63	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	4.5	4
62	Immune landscape and subtypes in primary resectable oral squamous cell carcinoma: prognostic significance and predictive of therapeutic response 2021 , 9,		3
61	Epidemiological, clinical, radiographic characterization of non-syndromic supernumerary teeth in Chinese children and adolescents. <i>Oral Diseases</i> , 2021 , 27, 981-992	3.5	4
60	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	4	1
59	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	3.3	1
58	Investigation of Causal Effect of Type 2 Diabetes Mellitus on Lung Cancer: A Mendelian Randomization Study. <i>Frontiers in Genetics</i> , 2021 , 12, 673687	4.5	1
57	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	3.4	
56	Functional Dissection of CD26 and Its Pharmacological Inhibition by Sitagliptin During Skin Wound Healing. <i>Medical Science Monitor</i> , 2021 , 27, e928933	3.2	2
55	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	9.7	10
54	Functional genetic variants in centrosome-related genes CEP72 and YWHAG confer susceptibility to gastric cancer. <i>Archives of Toxicology</i> , 2020 , 94, 2861-2872	5.8	3
53	Transcriptome-wide association study for persistent hepatitis B virus infection and related hepatocellular carcinoma. <i>Liver International</i> , 2020 , 40, 2117-2127	7.9	1
52	Predictive value of prognostic nutritional index in patients with oral squamous cell carcinoma. <i>Oral Diseases</i> , 2020 , 26, 903-911	3.5	10
51	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	4.2	1

50	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, 8781348	3	1
49	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	9.9	23
48	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. <i>Lancet Oncology, The</i> , 2020 , 21, 1378-1386	21.7	38
47	Comprehensive characterization of functional eRNAs in lung adenocarcinoma reveals novel regulators and a prognosis-related molecular subtype. <i>Theranostics</i> , 2020 , 10, 11264-11277	12.1	9
46	Claudin-7 Inhibits Proliferation and Metastasis in Salivary Adenoid Cystic Carcinoma Through Wnt/ECatenin Signaling. <i>Cell Transplantation</i> , 2020 , 29, 963689720943583	4	4
45	Genetic characterisation of sarcomatoid carcinomas reveals multiple novel actionable mutations and identifies mutation as a biomarker of poor prognosis. <i>Journal of Medical Genetics</i> , 2020 ,	5.8	2
44	Identification of Recurrent Variants in and across Multiple Cancers in the Chinese Population. <i>BioMed Research International</i> , 2020 , 2020, 6739823	3	1
43	Epigenetic-smoking interaction reveals histologically heterogeneous effects of TRIM27 DNA methylation on overall survival among early-stage NSCLC patients. <i>Molecular Oncology</i> , 2020 , 14, 2759-2774	7.9	7
42	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	19.2	18
41	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	14.6	30
40	Leukocyte telomere length, lipid parameters and gestational diabetes risk: a case-control study in a Chinese population. <i>Scientific Reports</i> , 2019 , 9, 8483	4.9	5
39	Associations Between Hepatitis B Virus Infection and Risk of All Cancer Types. <i>JAMA Network Open</i> , 2019 , 2, e195718	10.4	54
38	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	7	1
37	Integrating of genomic and transcriptomic profiles for the prognostic assessment of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2019 , 175, 691-699	4.4	3
36	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, the</i> , 2019 , 7, 881-891	35.1	58
35	The sonic hedgehog signaling pathway is suppressed following PCB exposure during retinal development. <i>Environmental Toxicology</i> , 2019 , 34, 340-347	4.2	3
34	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	9.2	45
33	Evaluation of CpG-SNPs in miRNA promoters and risk of breast cancer. <i>Gene</i> , 2018 , 651, 1-8	3.8	10

32	Whole-genome sequencing reveals genomic signatures associated with the inflammatory microenvironments in Chinese NSCLC patients. <i>Nature Communications</i> , 2018 , 9, 2054	17.4	43
31	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	19.2	51
30	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017 , 152, 2011-2021	13.3	32
29	Potentially functional variants in lncRNAs are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2017 , 56, 2048-2057	5	8
28	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	7.6	4
27	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	6.9	9
26	Genetic Variants in the Promoter Region of and the Risk of Breast Cancer. <i>BioMed Research International</i> , 2017 , 2017, 2352874	3	2
25	A cis-eQTL genetic variant of the cancer-testis gene CCDC116 is associated with risk of multiple cancers. <i>Human Genetics</i> , 2017 , 136, 987-997	6.3	4
24	Evaluation of genetic variants in autophagy pathway genes as prognostic biomarkers for breast cancer. <i>Gene</i> , 2017 , 627, 549-555	3.8	18
23	Low-frequency nonsynonymous variants in FKBPL and ARPC1B genes are associated with breast cancer risk in Chinese women. <i>Molecular Carcinogenesis</i> , 2017 , 56, 774-780	5	9
22	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	5	3
21	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	7.5	43
20	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	4.9	1
19	Systematic identification of genes with a cancer-testis expression pattern in 19 cancer types. <i>Nature Communications</i> , 2016 , 7, 10499	17.4	80
18	Genomic Landscape Survey Identifies SRSF1 as a Key Oncodriver in Small Cell Lung Cancer. <i>PLoS Genetics</i> , 2016 , 12, e1005895	6	87
17	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	4.4	11
16	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	7.8	45
15	Plasma miRNAs as early biomarkers for detecting hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2015 , 137, 1679-90	7.5	152

14	A Screen for Genomic Disorders of Infertility Identifies MAST2 Duplications Associated with Nonobstructive Azoospermia in Humans. <i>Biology of Reproduction</i> , 2015 , 93, 61	3.9	18
13	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, 478-86	12	8
12	Potentially functional polymorphisms in aminoacyl-tRNA synthetases genes are associated with breast cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015 , 54, 577-83	5	7
11	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	4.9	31
10	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, 5628-36	5.6	10
9	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	3.7	16
8	Genetic Variation in the 3'UTR Untranslated Region of NBN Gene Is Associated with Gastric Cancer Risk in a Chinese Population. <i>PLoS ONE</i> , 2015 , 10, e0139059	3.7	6
7	Association analysis identifies new risk loci for non-obstructive azoospermia in Chinese men. <i>Nature Communications</i> , 2014 , 5, 3857	17.4	50
6	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	3.7	18
5	Early second-trimester serum microRNAs as potential biomarker for nondiabetic macrosomia. <i>BioMed Research International</i> , 2014 , 2014, 394125	3	7
4	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	1.5	3
3	Evaluation of genetic variants in microRNA biosynthesis genes and risk of breast cancer in Chinese women. <i>International Journal of Cancer</i> , 2013 , 133, 2216-24	7.5	41
2	Genetic variation in a hsa-let-7 binding site in RAD52 is associated with breast cancer susceptibility. <i>Carcinogenesis</i> , 2013 , 34, 689-93	4.6	41
1	Genetic variants at 1p11.2 and breast cancer risk: a two-stage study in Chinese women. <i>PLoS ONE</i> , 2011 , 6, e21563	3.7	21