

Yue Jiang

List of Publications by Citations

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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	Plasma miRNAs as early biomarkers for detecting hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2015 , 137, 1679-90	7.5	152
66	Genomic Landscape Survey Identifies SRSF1 as a Key Oncodriver in Small Cell Lung Cancer. <i>PLoS Genetics</i> , 2016 , 12, e1005895	6	87
65	Systematic identification of genes with a cancer-testis expression pattern in 19 cancer types. <i>Nature Communications</i> , 2016 , 7, 10499	17.4	80
64	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	35.1	58
63	Associations Between Hepatitis B Virus Infection and Risk of All Cancer Types. <i>JAMA Network Open</i> , 2019 , 2, e195718	10.4	54
62	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
61	Association analysis identifies new risk loci for non-obstructive azoospermia in Chinese men. <i>Nature Communications</i> , 2014 , 5, 3857	17.4	50
60	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
59	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
58	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
57	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
56	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
55	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
54	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	21.7	38
53	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
52	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
51	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

50	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
49	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
48	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
47	Evaluation of genetic variants in autophagy pathway genes as prognostic biomarkers for breast cancer. <i>Gene</i> , 2017 , 627, 549-555	3.8	18
46	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
45	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
44	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
43	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
42	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
41	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
40	Predictive value of prognostic nutritional index in patients with oral squamous cell carcinoma. <i>Oral Diseases</i> , 2020 , 26, 903-911	3.5	10
39	Evaluation of CpG-SNPs in miRNA promoters and risk of breast cancer. <i>Gene</i> , 2018 , 651, 1-8	3.8	10
38	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
37	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
36	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
35	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
34	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
33	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

32	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
31	Early second-trimester serum microRNAs as potential biomarker for nondiabetic macrosomia. <i>BioMed Research International</i> , 2014 , 2014, 394125	3	7
30	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
29	Genetic Variation in the 3' 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
28	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
27	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
26	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
25	Claudin-7 Inhibits Proliferation and Metastasis in Salivary Adenoid Cystic Carcinoma Through Wnt/ECatenin Signaling. <i>Cell Transplantation</i> , 2020 , 29, 963689720943583	4	4
24	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
23	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
22	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
21	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
20	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
19	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
18	Immune landscape and subtypes in primary resectable oral squamous cell carcinoma: prognostic significance and predictive of therapeutic response 2021 , 9,		3
17	The sonic hedgehog signaling pathway is suppressed following PCB exposure during retinal development. <i>Environmental Toxicology</i> , 2019 , 34, 340-347	4.2	3
16	Genetic Variants in the Promoter Region of and the Risk of Breast Cancer. <i>BioMed Research International</i> , 2017 , 2017, 2352874	3	2
15	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

14	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
13	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
12	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
11	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
10	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
9	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
8	Sex-Specific Associations of Testosterone and Genetic Factors With Health Span.. <i>Frontiers in Endocrinology</i> , 2021 , 12, 773464	5.7	1
7	Identification of Recurrent Variants in and across Multiple Cancers in the Chinese Population. <i>BioMed Research International</i> , 2020 , 2020, 6739823	3	1
6	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
5	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
4	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
3	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
2	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	
1	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	