

# Wei-Hua Jia

## List of Publications by Year in descending order

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140  
papers

7,541  
citations

57758

44  
h-index

60623

81  
g-index

140  
all docs

140  
docs citations

140  
times ranked

10240  
citing authors

#	ARTICLE	IF	CITATIONS
1	A genome-wide association study of nasopharyngeal carcinoma identifies three new susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 599-603.	21.4	374
2	Global trends in incidence and mortality of nasopharyngeal carcinoma. <i>Cancer Letters</i> , 2016, 374, 22-30.	7.2	330
3	Association of the Asp312Asn and Lys751Gln polymorphisms in the XPD gene with the risk of non-Hodgkin's lymphoma: evidence from a meta-analysis. <i>Chinese Journal of Cancer</i> , 2015, 34, 108-14.	4.9	326
4	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2011, 43, 679-684.	21.4	260
5	Excessive miR-25-3p maturation via N6-methyladenosine stimulated by cigarette smoke promotes pancreatic cancer progression. <i>Nature Communications</i> , 2019, 10, 1858.	12.8	242
6	Genome-wide scan for familial nasopharyngeal carcinoma reveals evidence of linkage to chromosome 4. <i>Nature Genetics</i> , 2002, 31, 395-399.	21.4	217
7	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. <i>Nature Genetics</i> , 2014, 46, 533-542.	21.4	212
8	CPT1A-mediated fatty acid oxidation promotes colorectal cancer cell metastasis by inhibiting anoikis. <i>Oncogene</i> , 2018, 37, 6025-6040.	5.9	211
9	Trends in incidence and mortality of nasopharyngeal carcinoma over a 20-year period (1978/1983-2002) in Sihui and Cangwu counties in southern China. <i>BMC Cancer</i> , 2006, 6, 178.	2.6	199
10	Genome-wide association analyses in east Asians identify new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2013, 45, 191-196.	21.4	173
11	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.	10.7	167
12	An Epidemiological and Molecular Study of the Relationship Between Smoking, Risk of Nasopharyngeal Carcinoma, and Epstein-Barr Virus Activation. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1396-1410.	6.3	164
13	Non-viral environmental risk factors for nasopharyngeal carcinoma: A systematic review. <i>Seminars in Cancer Biology</i> , 2012, 22, 117-126.	9.6	151
14	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	12.8	138
15	Genome sequencing analysis identifies Epstein-Barr virus subtypes associated with high risk of nasopharyngeal carcinoma. <i>Nature Genetics</i> , 2019, 51, 1131-1136.	21.4	133
16	Fluctuations of Epstein-Barr Virus Serological Antibodies and Risk for Nasopharyngeal Carcinoma: A Prospective Screening Study with a 20-Year Follow-Up. <i>PLoS ONE</i> , 2011, 6, e19100.	2.5	129
17	Genomic Characterization of Esophageal Squamous Cell Carcinoma Reveals Critical Genes Underlying Tumorigenesis and Poor Prognosis. <i>American Journal of Human Genetics</i> , 2016, 98, 709-727.	6.2	129
18	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	6.3	129

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19	Traditional Cantonese diet and nasopharyngeal carcinoma risk: a large-scale case-control study in Guangdong, China. <i>BMC Cancer</i> , 2010, 10, 446.	2.6	118
20	Liquid biopsies to track trastuzumab resistance in metastatic HER2-positive gastric cancer. <i>Gut</i> , 2019, 68, 1152-1161.	12.1	118
21	Establishment of VCA and EBNA1 IgA-based combination by enzyme-linked immunosorbent assay as preferred screening method for nasopharyngeal carcinoma: a two-stage design with a preliminary performance study and a mass screening in southern China. <i>International Journal of Cancer</i> , 2012, 131, 406-416.	5.1	116
22	PIWI-interacting RNA-54265 is oncogenic and a potential therapeutic target in colorectal adenocarcinoma. <i>Theranostics</i> , 2018, 8, 5213-5230.	10.0	115
23	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , 2019, 156, 1455-1466.	1.3	111
24	Two Epstein-Barr Virus-Related Serologic Antibody Tests in Nasopharyngeal Carcinoma Screening: Results From the Initial Phase of a Cluster Randomized Controlled Trial in Southern China. <i>American Journal of Epidemiology</i> , 2013, 177, 242-250.	3.4	108
25	The Pretreatment Albumin to Globulin Ratio Has Predictive Value for Long-Term Mortality in Nasopharyngeal Carcinoma. <i>PLoS ONE</i> , 2014, 9, e94473.	2.5	99
26	Genome-wide association study identifies common variants in SLC39A6 associated with length of survival in esophageal squamous-cell carcinoma. <i>Nature Genetics</i> , 2013, 45, 632-638.	21.4	97
27	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	1.3	97
28	A circRNA signature predicts postoperative recurrence in stage II/III colon cancer. <i>EMBO Molecular Medicine</i> , 2019, 11, e10168.	6.9	90
29	Quantitative Association of Tobacco Smoking With the Risk of Nasopharyngeal Carcinoma: A Comprehensive Meta-Analysis of Studies Conducted Between 1979 and 2011. <i>American Journal of Epidemiology</i> , 2013, 178, 325-338.	3.4	89
30	Association of MTHFR C677T and A1298C polymorphisms with non-Hodgkin lymphoma susceptibility: Evidence from a meta-analysis. <i>Scientific Reports</i> , 2015, 4, 6159.	3.3	83
31	Genome-Wide Identification of a Methylation Gene Panel as a Prognostic Biomarker in Nasopharyngeal Carcinoma. <i>Molecular Cancer Therapeutics</i> , 2015, 14, 2864-2873.	4.1	80
32	Familial risk and clustering of nasopharyngeal carcinoma in Guangdong, China. <i>Cancer</i> , 2004, 101, 363-369.	4.1	79
33	Redox Regulation of Stem-like Cells Through the CD44v-xCT Axis in Colorectal Cancer: Mechanisms and Therapeutic Implications. <i>Theranostics</i> , 2016, 6, 1160-1175.	10.0	75
34	Active and Passive Smoking and Risk of Nasopharyngeal Carcinoma: A Population-Based Case-Control Study in Southern China. <i>American Journal of Epidemiology</i> , 2017, 185, 1272-1280.	3.4	68
35	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.	5.1	66
36	ADAR2 functions as a tumor suppressor via editing IGFBP7 in esophageal squamous cell carcinoma. <i>International Journal of Oncology</i> , 2017, 50, 622-630.	3.3	65

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37	Genome-wide association study identifies new susceptibility loci for epithelial ovarian cancer in Han Chinese women. <i>Nature Communications</i> , 2014, 5, 4682.	12.8	59
38	Familial and large-scale case-control studies identify genes associated with nasopharyngeal carcinoma. <i>Seminars in Cancer Biology</i> , 2012, 22, 96-106.	9.6	58
39	Genome-wide association study identifies three susceptibility loci for laryngeal squamous cell carcinoma in the Chinese population. <i>Nature Genetics</i> , 2014, 46, 1110-1114.	21.4	57
40	Genome-wide profiling of Epstein-Barr virus integration by targeted sequencing in Epstein-Barr virus associated malignancies. <i>Theranostics</i> , 2019, 9, 1115-1124.	10.0	56
41	Quantification of Epstein-Barr virus DNA load in nasopharyngeal brushing samples in the diagnosis of nasopharyngeal carcinoma in southern China. <i>Cancer Science</i> , 2015, 106, 1196-1201.	3.9	54
42	Quantification of familial risk of nasopharyngeal carcinoma in a high-incidence area. <i>Cancer</i> , 2017, 123, 2716-2725.	4.1	54
43	Effect of family history of cancers and environmental factors on risk of nasopharyngeal carcinoma in Guangdong, China. <i>Cancer Epidemiology</i> , 2010, 34, 419-424.	1.9	53
44	High Expression of p300 Has an Unfavorable Impact on Survival in Resectable Esophageal Squamous Cell Carcinoma. <i>Annals of Thoracic Surgery</i> , 2011, 91, 1531-1538.	1.3	49
45	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study in multiple populations. <i>Lancet Oncology</i> , The, 2020, 21, 306-316.	10.7	49
46	TERT Polymorphism rs2736100-C Is Associated with EGFR Mutation-Positive Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 5173-5180.	7.0	47
47	Oral Hygiene and Risk of Nasopharyngeal Carcinoma: A Population-Based Case-Control Study in China. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1201-1207.	2.5	46
48	A GWAS Meta-analysis and Replication Study Identifies a Novel Locus within <i>CLPTM1L/TERT</i> Associated with Nasopharyngeal Carcinoma in Individuals of Chinese Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 188-192.	2.5	45
49	Genome-Wide Association Study of Susceptibility Loci for Radiation-Induced Brain Injury. <i>Journal of the National Cancer Institute</i> , 2019, 111, 620-628.	6.3	45
50	A Case-control and a family-based association study revealing an association between CYP2E1 polymorphisms and nasopharyngeal carcinoma risk in Cantonese. <i>Carcinogenesis</i> , 2009, 30, 2031-2036.	2.8	43
51	Association Between Oral Microbiota and Cigarette Smoking in the Chinese Population. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 658203.	3.9	43
52	Increased RIPK4 expression is associated with progression and poor prognosis in cervical squamous cell carcinoma patients. <i>Scientific Reports</i> , 2015, 5, 11955.	3.3	42
53	An extended genome-wide association study identifies novel susceptibility loci for nasopharyngeal carcinoma. <i>Human Molecular Genetics</i> , 2016, 25, 3626-3634.	2.9	42
54	Comprehensive Pathway-Based Association Study of DNA Repair Gene Variants and the Risk of Nasopharyngeal Carcinoma. <i>Cancer Research</i> , 2011, 71, 3000-3008.	0.9	41

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55	Hepatitis B virus infection is associated with younger median age at diagnosis and death in cancers. <i>International Journal of Cancer</i> , 2017, 141, 152-159.	5.1	38
56	Prognostic efficacy of combining tumor volume with Epstein-Barr virus DNA in patients treated with intensity-modulated radiotherapy for nasopharyngeal carcinoma. <i>Oral Oncology</i> , 2016, 60, 18-24.	1.5	35
57	Epstein-Barr virus <i>mir-bart1-5p</i> detection via nasopharyngeal brush sampling is effective for diagnosing nasopharyngeal carcinoma. <i>Oncotarget</i> , 2016, 7, 4972-4980.	1.8	34
58	Association of BRCA2 N372H polymorphism with cancer susceptibility: A comprehensive review and meta-analysis. <i>Scientific Reports</i> , 2014, 4, 6791.	3.3	33
59	Household inhalants exposure and nasopharyngeal carcinoma risk: a large-scale case-control study in Guangdong, China. <i>BMC Cancer</i> , 2015, 15, 1022.	2.6	32
60	Comparison of Long-Term Survival and Toxicity of Cisplatin Delivered Weekly versus Every Three Weeks Concurrently with Intensity-Modulated Radiotherapy in Nasopharyngeal Carcinoma. <i>PLoS ONE</i> , 2014, 9, e110765.	2.5	31
61	High Expression of LAMP3 Is a Novel Biomarker of Poor Prognosis in Patients with Esophageal Squamous Cell Carcinoma. <i>International Journal of Molecular Sciences</i> , 2015, 16, 17655-17667.	4.1	31
62	The Relationship Between Environmental Factors and the Profile of Epstein-Barr Virus Antibodies in the Lytic and Latent Infection Periods in Healthy Populations from Endemic and Non-Endemic Nasopharyngeal Carcinoma Areas in China. <i>EBioMedicine</i> , 2018, 30, 184-191.	6.1	31
63	Identification of miR-143 as a tumour suppressor in nasopharyngeal carcinoma based on microRNA expression profiling. <i>International Journal of Biochemistry and Cell Biology</i> , 2015, 61, 120-128.	2.8	30
64	Association between genetic variants in the XPG gene and gastric cancer risk in a Southern Chinese population. <i>Aging</i> , 2016, 8, 3311-3320.	3.1	30
65	Genome-wide association study identifies genetic susceptibility loci and pathways of radiation-induced acute oral mucositis. <i>Journal of Translational Medicine</i> , 2020, 18, 224.	4.4	29
66	Development of a population-based cancer case-control study in southern china. <i>Oncotarget</i> , 2017, 8, 87073-87085.	1.8	29
67	A single nucleotide polymorphism in the Epstein-Barr virus genome is strongly associated with a high risk of nasopharyngeal carcinoma. <i>Chinese Journal of Cancer</i> , 2015, 34, 563-72.	4.9	28
68	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. <i>Gynecologic Oncology</i> , 2019, 153, 343-355.	1.4	28
69	<i>XPG</i> Gene Polymorphisms Contribute to Colorectal Cancer Susceptibility: A Two-Stage Case-Control Study. <i>Journal of Cancer</i> , 2016, 7, 1731-1739.	2.5	27
70	Genetic variants in the nucleotide excision repair pathway genes and gastric cancer susceptibility in a southern Chinese population. <i>Cancer Management and Research</i> , 2018, Volume 10, 765-774.	1.9	27
71	Past and Recent Salted Fish and Preserved Food Intakes Are Weakly Associated with Nasopharyngeal Carcinoma Risk in Adults in Southern China. <i>Journal of Nutrition</i> , 2019, 149, 1596-1605.	2.9	25
72	Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 477-486.	2.5	25

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73	Association of XPC Gene Polymorphisms with Colorectal Cancer Risk in a Southern Chinese Population: A Case-Control Study and Meta-Analysis. <i>Genes</i> , 2016, 7, 73.	2.4	24
74	Association Between Environmental Factors and Oral Epstein-Barr Virus DNA Loads: A Multicenter Cross-sectional Study in China. <i>Journal of Infectious Diseases</i> , 2019, 219, 400-409.	4.0	22
75	Alcohol and tea consumption in relation to the risk of nasopharyngeal carcinoma in Guangdong, China. <i>Frontiers of Medicine in China</i> , 2010, 4, 448-456.	0.1	21
76	Overexpression of CIP2A is an independent prognostic indicator in nasopharyngeal carcinoma and its depletion suppresses cell proliferation and tumor growth. <i>Molecular Cancer</i> , 2014, 13, 111.	19.2	21
77	High expression of Talin-1 is associated with poor prognosis in patients with nasopharyngeal carcinoma. <i>BMC Cancer</i> , 2015, 15, 332.	2.6	21
78	Chinese nonmedicinal herbal diet and risk of nasopharyngeal carcinoma: A population-based case-control study. <i>Cancer</i> , 2019, 125, 4462-4470.	4.1	21
79	Developing Genetic Epidemiological Models to Predict Risk for Nasopharyngeal Carcinoma in High-Risk Population of China. <i>PLoS ONE</i> , 2013, 8, e56128.	2.5	21
80	Comprehensive profiling of 1015 patients' exomes reveals genomic-clinical associations in colorectal cancer. <i>Nature Communications</i> , 2022, 13, 2342.	12.8	21
81	Antibodies against Epstein-Barr virus gp78 antigen: a novel marker for serological diagnosis of nasopharyngeal carcinoma detected by xMAP technology. <i>Journal of General Virology</i> , 2008, 89, 1152-1158.	2.9	20
82	Medical History, Medication Use, and Risk of Nasopharyngeal Carcinoma. <i>American Journal of Epidemiology</i> , 2018, 187, 2117-2125.	3.4	20
83	Glatiramer acetate reverses cognitive deficits from cranial-irradiated rat by inducing hippocampal neurogenesis. <i>Journal of Neuroimmunology</i> , 2014, 271, 1-7.	2.3	19
84	Targeting Epstein-Barr virus oncoprotein LMP1-mediated high oxidative stress suppresses EBV lytic reactivation and sensitizes tumors to radiation therapy. <i>Theranostics</i> , 2020, 10, 11921-11937.	10.0	19
85	ATAD2 interacts with C/EBP $\beta$ to promote esophageal squamous cell carcinoma metastasis via TGF- $\beta$ 1/Smad3 signaling. <i>Journal of Experimental and Clinical Cancer Research</i> , 2021, 40, 109.	8.6	19
86	A polygenic risk score for nasopharyngeal carcinoma shows potential for risk stratification and personalized screening. <i>Nature Communications</i> , 2022, 13, 1966.	12.8	19
87	Polymorphisms in the XPC gene and gastric cancer susceptibility in a Southern Chinese population. <i>OncoTargets and Therapy</i> , 2016, Volume 9, 5513-5519.	2.0	18
88	The Bidirectional Regulation between MYL5 and HIF-1 $\alpha$ Promotes Cervical Carcinoma Metastasis. <i>Theranostics</i> , 2017, 7, 3768-3780.	10.0	17
89	XPC rs2296147 T>C polymorphism predicted clinical outcome in colorectal cancer. <i>Oncotarget</i> , 2016, 7, 11724-11732.	1.8	17
90	The Effects of Alcohol Drinking on Oral Microbiota in the Chinese Population. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 5729.	2.6	17

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91	AMPK $\beta$ 1 confers survival advantage of colorectal cancer cells under metabolic stress by promoting redox balance through the regulation of glutathione reductase phosphorylation. <i>Oncogene</i> , 2020, 39, 637-650.	5.9	16
92	Wild-type IDH2 contributes to Epstein-Barr virus-dependent metabolic alterations and tumorigenesis. <i>Molecular Metabolism</i> , 2020, 36, 100966.	6.5	16
93	The association between the polymorphisms of TNF- $\beta$ and non-Hodgkin lymphoma: a meta-analysis. <i>Tumor Biology</i> , 2014, 35, 12509-12517.	1.8	15
94	Cumulative scores based on plasma D-dimer and serum albumin levels predict survival in esophageal squamous cell carcinoma patients treated with transthoracic esophagectomy. <i>Chinese Journal of Cancer</i> , 2016, 35, 11.	4.9	15
95	Fine-mapping of HLA class I and class II genes identified two independent novel variants associated with nasopharyngeal carcinoma susceptibility. <i>Cancer Medicine</i> , 2018, 7, 6308-6316.	2.8	15
96	Associations between environmental factors and serological Epstein-Barr virus antibodies in patients with nasopharyngeal carcinoma in South China. <i>Cancer Medicine</i> , 2019, 8, 4852-4866.	2.8	15
97	Body mass index, body shape, and risk of nasopharyngeal carcinoma: A population-based case-control study in Southern China. <i>Cancer Medicine</i> , 2019, 8, 1835-1844.	2.8	15
98	Genetic association of telomere length with hepatocellular carcinoma risk: A Mendelian randomization analysis. <i>Cancer Epidemiology</i> , 2017, 50, 39-45.	1.9	14
99	Elevated Epstein-Barr virus seroreactivity among unaffected members of families with nasopharyngeal carcinoma. <i>Journal of Medical Virology</i> , 2011, 83, 1792-1798.	5.0	13
100	No association between MTRs1805087 A > G polymorphism and non-Hodgkin lymphoma susceptibility: evidence from 11 486 subjects. <i>Leukemia and Lymphoma</i> , 2015, 56, 763-767.	1.3	13
101	Nasopharyngeal carcinoma risk prediction via salivary detection of host and Epstein-Barr virus genetic variants. <i>Oncotarget</i> , 2017, 8, 95066-95074.	1.8	13
102	Nasopharyngeal brushing: a convenient and feasible sampling method for nucleic acid-based nasopharyngeal carcinoma research. <i>Cancer Communications</i> , 2018, 38, 1-10.	9.2	13
103	A comprehensive analysis of genetic diversity of EBV reveals potential high-risk subtypes associated with nasopharyngeal carcinoma in China. <i>Virus Evolution</i> , 2021, 7, veab010.	4.9	13
104	Glycogenes in Oncofetal Chondroitin Sulfate Biosynthesis are Differently Expressed and Correlated With Immune Response in Placenta and Colorectal Cancer. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 763875.	3.7	13
105	X-chromosome association study reveals genetic susceptibility loci of nasopharyngeal carcinoma. <i>Biology of Sex Differences</i> , 2019, 10, 13.	4.1	12
106	Detection of methylation status of Epstein-Barr virus DNA C promoter in the diagnosis of nasopharyngeal carcinoma. <i>Cancer Science</i> , 2020, 111, 592-600.	3.9	12
107	Residence characteristics and risk of nasopharyngeal carcinoma in southern China: A population-based case-control study. <i>Environment International</i> , 2021, 151, 106455.	10.0	11
108	Occupational exposures and risk of nasopharyngeal carcinoma in a high-risk area: A population-based case-control study. <i>Cancer</i> , 2021, 127, 2724-2735.	4.1	10

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109	Natural Variations in BRLF1 Promoter Contribute to the Elevated Reactivation Level of Epstein-Barr Virus in Endemic Areas of Nasopharyngeal Carcinoma. <i>EBioMedicine</i> , 2018, 37, 101-109.	6.1	9
110	Decreased oral Epstein-Barr virus DNA loads in patients with nasopharyngeal carcinoma in Southern China: A case-control and a family-based study. <i>Cancer Medicine</i> , 2018, 7, 3453-3464.	2.8	9
111	Weighted Risk Score-Based Multifactor Dimensionality Reduction to Detect Gene-Gene Interactions in Nasopharyngeal Carcinoma. <i>International Journal of Molecular Sciences</i> , 2014, 15, 10724-10737.	4.1	8
112	Pregnancy associated nasopharyngeal carcinoma: A retrospective case-control analysis of maternal survival outcomes. <i>Radiotherapy and Oncology</i> , 2015, 116, 125-130.	0.6	8
113	Association between XRCC3 Thr241Met polymorphism and nasopharyngeal carcinoma risk: evidence from a large-scale case-control study and a meta-analysis. <i>Tumor Biology</i> , 2016, 37, 14825-14830.	1.8	8
114	Reproductive history and risk of nasopharyngeal carcinoma: A population-based case-control study in southern China. <i>Oral Oncology</i> , 2019, 88, 102-108.	1.5	8
115	Environmental Factors for Epstein-Barr Virus Reactivation in a High-Risk Area of Nasopharyngeal Carcinoma: A Population-Based Study. <i>Open Forum Infectious Diseases</i> , 2022, 9, ofac128.	0.9	8
116	A comprehensive genomic characterization of esophageal squamous cell carcinoma: from prognostic analysis to in vivo assay. <i>Chinese Journal of Cancer</i> , 2016, 35, 76.	4.9	7
117	Identification of surrogate endpoints in patients with locoregionally advanced nasopharyngeal carcinoma receiving neoadjuvant chemotherapy plus concurrent chemoradiotherapy versus concurrent chemoradiotherapy alone. <i>BMC Cancer</i> , 2015, 15, 930.	2.6	6
118	CPEB4 interacts with Vimentin and involves in progressive features and poor prognosis of patients with astrocytic tumors. <i>Tumor Biology</i> , 2016, 37, 5075-5087.	1.8	6
119	Association Between Serum Cotinine Level and Serological Markers of Epstein-Barr Virus in Healthy Subjects in South China Where Nasopharyngeal Carcinoma Is Endemic. <i>Frontiers in Oncology</i> , 2019, 9, 865.	2.8	6
120	LIG3 gene polymorphisms and risk of gastric cancer in a Southern Chinese population. <i>Gene</i> , 2019, 705, 90-94.	2.2	6
121	Epstein-Barr virus DNA loads in the peripheral blood cells predict the survival of locoregionally-advanced nasopharyngeal carcinoma patients. <i>Cancer Biology and Medicine</i> , 2021, 18, 888-899.	3.0	6
122	Polymorphisms in TYMS for Prediction of Capecitabine-Induced Hand-Foot Syndrome in Chinese Patients with Colorectal Cancer. <i>Cancer Research and Treatment</i> , 2021, 53, 724-732.	3.0	6
123	A genetic variant in CHRN3-CHRNA6 increases risk of esophageal squamous cell carcinoma in Chinese populations. <i>Carcinogenesis</i> , 2015, 36, 538-542.	2.8	5
124	Potential factors associated with clinical stage of nasopharyngeal carcinoma at diagnosis: a case-control study. <i>Chinese Journal of Cancer</i> , 2017, 36, 71.	4.9	5
125	Implication of comorbidity on the initiation of chemotherapy and survival outcomes in patients with locoregionally advanced nasopharyngeal carcinoma. <i>Oncotarget</i> , 2017, 8, 10594-10601.	1.8	5
126	A fecal-based test for the detection of advanced adenoma and colorectal cancer: a case-control and screening cohort study. <i>BMC Medicine</i> , 2021, 19, 250.	5.5	5



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127	Intake of Alcohol and Tea and Risk of Nasopharyngeal Carcinoma: A Population-Based Caseâ€“Control Study in Southern China. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 545-553.	2.5	5
128	Genomic Landscapes of Epstein-Barr Virus in Pulmonary Lymphoepithelioma-Like Carcinoma. <i>Journal of Virology</i> , 2022, 96, JVI0169321.	3.4	5
129	Tumor Cell Content and RNA Integrity of Surgical Tissues from Different Types of Tumors and Its Correlation with Ex Vivo and In Vivo Ischemia. <i>Annals of Surgical Oncology</i> , 2018, 25, 3764-3770.	1.5	4
130	Nasopharyngeal Epsteinâ€“Barr virus DNA loads in highâ€“risk nasopharyngeal carcinoma families: Familial aggregation and host heritability. <i>Journal of Medical Virology</i> , 2020, 92, 3717-3725.	5.0	4
131	Association between HLA alleles and Epsteinâ€“Barr virus Ztaâ€“IgA serological status in healthy males from southern China. <i>Journal of Gene Medicine</i> , 2021, 23, e3375.	2.8	4
132	Discovery of a Pathogenic Variant rs139379666 (p. P2974L) in <i>ATM</i> for Breast Cancer Risk in Chinese Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1308-1315.	2.5	3
133	Informatics Management of Tumor Specimens in the Era of Big Data: Challenges and Solutions. <i>Biopreservation and Biobanking</i> , 2021, , .	1.0	3
134	ASO Author Reflections: Tumor Cell Content and RNA Integrity of Surgical Tissues from Different Types of Tumors and Its Correlation with Ex Vivo and In Vivo Ischemia. <i>Annals of Surgical Oncology</i> , 2018, 25, 3771-3772.	1.5	2
135	Prognostic Value of Oral Epsteinâ€“Barr Virus DNA Load in Locoregionally Advanced Nasopharyngeal Carcinoma. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 757644.	3.5	2
136	Performance of common genetic variants in risk prediction for colorectal cancer in Chinese: A two-stage and multicenter study. <i>Genomics</i> , 2021, 113, 867-873.	2.9	1
137	Genomic landscape of Epsteinâ€“Barr virus in familial nasopharyngeal carcinoma. <i>Journal of General Virology</i> , 2022, 103, .	2.9	1
138	Transcriptomeâ€“wide association analysis identified candidate susceptibility genes for nasopharyngeal carcinoma. <i>Cancer Communications</i> , 2022, 42, 887-891.	9.2	1
139	Association of plgR polymorphisms with nasopharyngeal carcinoma. <i>Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association</i> , Beijing Institute for Cancer Research, 2006, 18, 168-172.	2.2	0
140	Banking of Tumor Tissues: Effect of Preanalytical Variables in the Phase of Pre- and Postacquisition on RNA Integrity. <i>Biopreservation and Biobanking</i> , 2022, , .	1.0	0