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	Genomic Landscapes of Epstein-Barr Virus in Pulmonary Lymphoepithelioma-Like Carcinoma. Journal of Virology, 2022, 96, JVI0169321.	3.4	5
2	Environmental Factors for Epstein-Barr Virus Reactivation in a High-Risk Area of Nasopharyngeal Carcinoma: A Population-Based Study. Open Forum Infectious Diseases, 2022, 9, ofac128.	0.9	8
3	Genomic landscape of Epsteinâ€“Barr virus in familial nasopharyngeal carcinoma. Journal of General Virology, 2022, 103, .	2.9	1
4	Banking of Tumor Tissues: Effect of Preanalytical Variables in the Phase of Pre- and Postacquisition on RNA Integrity. Biopreservation and Biobanking, 2022, , .	1.0	0
5	A polygenic risk score for nasopharyngeal carcinoma shows potential for risk stratification and personalized screening. Nature Communications, 2022, 13, 1966.	12.8	19
6	Comprehensive profiling of 1015 patientsâ€™ exomes reveals genomic-clinical associations in colorectal cancer. Nature Communications, 2022, 13, 2342.	12.8	21
7	The Effects of Alcohol Drinking on Oral Microbiota in the Chinese Population. International Journal of Environmental Research and Public Health, 2022, 19, 5729.	2.6	17
8	Transcriptomeâ€wide association analysis identified candidate susceptibility genes for nasopharyngeal carcinoma. Cancer Communications, 2022, 42, 887-891.	9.2	1
9	Epstein-Barr virus DNA loads in the peripheral blood cells predict the survival of locoregionally-advanced nasopharyngeal carcinoma patients. Cancer Biology and Medicine, 2021, 18, 888-899.	3.0	6
10	ATAD2 interacts with C/EBPÎ² to promote esophageal squamous cell carcinoma metastasis via TGF-Î²1/Smad3 signaling. Journal of Experimental and Clinical Cancer Research, 2021, 40, 109.	8.6	19
11	Occupational exposures and risk of nasopharyngeal carcinoma in a highâ€risk area: A populationâ€based caseâ€control study. Cancer, 2021, 127, 2724-2735.	4.1	10
12	Informatics Management of Tumor Specimens in the Era of Big Data: Challenges and Solutions. Biopreservation and Biobanking, 2021, , .	1.0	3
13	Performance of common genetic variants in risk prediction for colorectal cancer in Chinese: A two-stage and multicenter study. Genomics, 2021, 113, 867-873.	2.9	1
14	Association Between Oral Microbiota and Cigarette Smoking in the Chinese Population. Frontiers in Cellular and Infection Microbiology, 2021, 11, 658203.	3.9	43
15	Residence characteristics and risk of nasopharyngeal carcinoma in southern China: A population-based case-control study. Environment International, 2021, 151, 106455.	10.0	11
16	Association between HLA alleles and Epsteinâ€“Barr virus Ztaâ€IgA serological status in healthy males from southern China. Journal of Gene Medicine, 2021, 23, e3375.	2.8	4
17	Polymorphisms in TYMS for Prediction of Capecitabine-Induced Hand-Foot Syndrome in Chinese Patients with Colorectal Cancer. Cancer Research and Treatment, 2021, 53, 724-732.	3.0	6
18	A comprehensive analysis of genetic diversity of EBV reveals potential high-risk subtypes associated with nasopharyngeal carcinoma in China. Virus Evolution, 2021, 7, veab010.	4.9	13

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19	A fecal-based test for the detection of advanced adenoma and colorectal cancer: a case-control and screening cohort study. BMC Medicine, 2021, 19, 250.	5.5	5
20	Intake of Alcohol and Tea and Risk of Nasopharyngeal Carcinoma: A Population-Based Caseâ€“Control Study in Southern China. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 545-553.	2.5	5
21	Prognostic Value of Oral Epsteinâ€“Barr Virus DNA Load in Locoregionally Advanced Nasopharyngeal Carcinoma. Frontiers in Molecular Biosciences, 2021, 8, 757644.	3.5	2
22	Glycogenes in Oncofetal Chondroitin Sulfate Biosynthesis are Differently Expressed and Correlated With Immune Response in Placenta and Colorectal Cancer. Frontiers in Cell and Developmental Biology, 2021, 9, 763875.	3.7	13
23	AMPKÎ±1 confers survival advantage of colorectal cancer cells under metabolic stress by promoting redox balance through the regulation of glutathione reductase phosphorylation. Oncogene, 2020, 39, 637-650.	5.9	16
24	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study in multiple populations. Lancet Oncology, The, 2020, 21, 306-316.	10.7	49
25	Identification of Novel Loci and New Risk Variant in Known Loci for Colorectal Cancer Risk in East Asians. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 477-486.	2.5	25
26	Targeting Epstein-Barr virus oncoprotein LMP1-mediated high oxidative stress suppresses EBV lytic reactivation and sensitizes tumors to radiation therapy. Theranostics, 2020, 10, 11921-11937.	10.0	19
27	Nasopharyngeal Epsteinâ€Barr virus DNA loads in highâ€risk nasopharyngeal carcinoma families: Familial aggregation and host heritability. Journal of Medical Virology, 2020, 92, 3717-3725.	5.0	4
28	Genome-wide association study identifies genetic susceptibility loci and pathways of radiation-induced acute oral mucositis. Journal of Translational Medicine, 2020, 18, 224.	4.4	29
29	Detection of methylation status of Epsteinâ€Barr virus DNA C promoter in the diagnosis of nasopharyngeal carcinoma. Cancer Science, 2020, 111, 592-600.	3.9	12
30	Wild-type IDH2 contributes to Epsteinâ€“Barr virus-dependent metabolic alterations and tumorigenesis. Molecular Metabolism, 2020, 36, 100966.	6.5	16
31	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	6.3	129
32	Identification of risk loci and a polygenic risk score for lung cancer: a large-scale prospective cohort study in Chinese populations. Lancet Respiratory Medicine,the, 2019, 7, 881-891.	10.7	167
33	Chinese nonmedicinal herbal diet and risk of nasopharyngeal carcinoma: A populationâ€based caseâ€control study. Cancer, 2019, 125, 4462-4470.	4.1	21
34	A circRNA signature predicts postoperative recurrence in stage II/III colon cancer. EMBO Molecular Medicine, 2019, 11, e10168.	6.9	90
35	Association Between Serum Cotinine Level and Serological Markers of Epsteinâ€“Barr Virus in Healthy Subjects in South China Where Nasopharyngeal Carcinoma Is Endemic. Frontiers in Oncology, 2019, 9, 865.	2.8	6
36	Genome sequencing analysis identifies Epsteinâ€“Barr virus subtypes associated with high risk of nasopharyngeal carcinoma. Nature Genetics, 2019, 51, 1131-1136.	21.4	133

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37	Associations between environmental factors and serological Epsteinâ€Barr virus antibodies in patients with nasopharyngeal carcinoma in South China. Cancer Medicine, 2019, 8, 4852-4866.	2.8	15
38	Past and Recent Salted Fish and Preserved Food Intakes Are Weakly Associated with Nasopharyngeal Carcinoma Risk in Adults in Southern China. Journal of Nutrition, 2019, 149, 1596-1605.	2.9	25
39	Discovery of a Pathogenic Variant rs139379666 (p. P2974L) in ATM for Breast Cancer Risk in Chinese Populations. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 1308-1315.	2.5	3
40	Excessive miR-25-3p maturation via N6-methyladenosine stimulated by cigarette smoke promotes pancreatic cancer progression. Nature Communications, 2019, 10, 1858.	12.8	242
41	LIG3 gene polymorphisms and risk of gastric cancer in a Southern Chinese population. Gene, 2019, 705, 90-94.	2.2	6
42	X-chromosome association study reveals genetic susceptibility loci of nasopharyngeal carcinoma. Biology of Sex Differences, 2019, 10, 13.	4.1	12
43	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. Gynecologic Oncology, 2019, 153, 343-355.	1.4	28
44	Body mass index, body shape, and risk of nasopharyngeal carcinoma: A populationâ€based caseâ€“control study in Southern China. Cancer Medicine, 2019, 8, 1835-1844.	2.8	15
45	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. Gastroenterology, 2019, 156, 1455-1466.	1.3	111
46	Genome-wide profiling of Epstein-Barr virus integration by targeted sequencing in Epstein-Barr virus associated malignancies. Theranostics, 2019, 9, 1115-1124.	10.0	56
47	Reproductive history and risk of nasopharyngeal carcinoma: A population-based caseâ€“control study in southern China. Oral Oncology, 2019, 88, 102-108.	1.5	8
48	Genome-Wide Association Study of Susceptibility Loci for Radiation-Induced Brain Injury. Journal of the National Cancer Institute, 2019, 111, 620-628.	6.3	45
49	Association Between Environmental Factors and Oral Epstein-Barr Virus DNA Loads: A Multicenter Cross-sectional Study in China. Journal of Infectious Diseases, 2019, 219, 400-409.	4.0	22
50	Liquid biopsies to track trastuzumab resistance in metastatic HER2-positive gastric cancer. Gut, 2019, 68, 1152-1161.	12.1	118
51	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. EBioMedicine, 2018, 30, 184-191.	6.1	31
52	Nasopharyngeal brushing: a convenient and feasible sampling method for nucleic acidâ€based nasopharyngeal carcinoma research. Cancer Communications, 2018, 38, 1-10.	9.2	13
53	Medical History, Medication Use, and Risk of Nasopharyngeal Carcinoma. American Journal of Epidemiology, 2018, 187, 2117-2125.	3.4	20
54	Natural Variations in BRLF1 Promoter Contribute to the Elevated Reactivation Level of Epstein-Barr Virus in Endemic Areas of Nasopharyngeal Carcinoma. EBioMedicine, 2018, 37, 101-109.	6.1	9

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55	Tumor Cell Content and RNA Integrity of Surgical Tissues from Different Types of Tumors and Its Correlation with Ex Vivo and In Vivo Ischemia. Annals of Surgical Oncology, 2018, 25, 3764-3770.	1.5	4
56	Fineâ€mapping of <scp>HLA</scp> class I and class <scp>II</scp> genes identified two independent novel variants associated with nasopharyngeal carcinoma susceptibility. Cancer Medicine, 2018, 7, 6308-6316.	2.8	15
57	PIWI-interacting RNA-54265 is oncogenic and a potential therapeutic target in colorectal adenocarcinoma. Theranostics, 2018, 8, 5213-5230.	10.0	115
58	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. Annals of Surgical Oncology, 2018, 25, 3771-3772.	1.5	2
59	Genetic variants in the nucleotide excision repair pathway genes and gastric cancer susceptibility in a southern Chinese population. Cancer Management and Research, 2018, Volume 10, 765-774.	1.9	27
60	CPT1A-mediated fatty acid oxidation promotes colorectal cancer cell metastasis by inhibiting anoikis. Oncogene, 2018, 37, 6025-6040.	5.9	211
61	Decreased oral Epstein-Barr virus DNA loads in patients with nasopharyngeal carcinoma in Southern China: A case-control and a family-based study. Cancer Medicine, 2018, 7, 3453-3464.	2.8	9
62	Quantification of familial risk of nasopharyngeal carcinoma in a highâ€incidence area. Cancer, 2017, 123, 2716-2725.	4.1	54
63	ADAR2 functions as a tumor suppressor via editing IGFBP7 in esophageal squamous cell carcinoma. International Journal of Oncology, 2017, 50, 622-630.	3.3	65
64	Active and Passive Smoking and Risk of Nasopharyngeal Carcinoma: A Population-Based Case-Control Study in Southern China. American Journal of Epidemiology, 2017, 185, 1272-1280.	3.4	68
65	Hepatitis B virus infection is associated with younger median age at diagnosis and death in cancers. International Journal of Cancer, 2017, 141, 152-159.	5.1	38
66	Genetic association of telomere length with hepatocellular carcinoma risk: A Mendelian randomization analysis. Cancer Epidemiology, 2017, 50, 39-45.	1.9	14
67	Estimation of heritability for nine common cancers using data from genomeâ€wide association studies in Chinese population. International Journal of Cancer, 2017, 140, 329-336.	5.1	66
68	The Bidirectional Regulation between MYL5 and HIF-1Î± Promotes Cervical Carcinoma Metastasis. Theranostics, 2017, 7, 3768-3780.	10.0	17
69	Potential factors associated with clinical stage of nasopharyngeal carcinoma at diagnosis: a caseâ€“control study. Chinese Journal of Cancer, 2017, 36, 71.	4.9	5
70	Nasopharyngeal carcinoma risk prediction via salivary detection of host and Epstein-Barr virus genetic variants. Oncotarget, 2017, 8, 95066-95074.	1.8	13
71	Development of a population-based cancer case-control study in southern china. Oncotarget, 2017, 8, 87073-87085.	1.8	29
72	Implication of comorbidity on the initiation of chemotherapy and survival outcomes in patients with locoregionally advanced nasopharyngeal carcinoma. Oncotarget, 2017, 8, 10594-10601.	1.8	5

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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. Genes, 2016, 7, 73.	2.4	24
74	Polymorphisms in the XPC gene and gastric cancer susceptibility in a Southern Chinese population. OncoTargets and Therapy, 2016, Volume 9, 5513-5519.	2.0	18
75	XPG Gene Polymorphisms Contribute to Colorectal Cancer Susceptibility: A Two-Stage Case-Control Study. Journal of Cancer, 2016, 7, 1731-1739.	2.5	27
76	Redox Regulation of Stem-like Cells Though the CD44v-xCT Axis in Colorectal Cancer: Mechanisms and Therapeutic Implications. Theranostics, 2016, 6, 1160-1175.	10.0	75
77	Oral Hygiene and Risk of Nasopharyngeal Carcinomaâ€"A Population-Based Caseâ€"Control Study in China. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1201-1207.	2.5	46
78	Genomic Characterization of Esophageal Squamous Cell Carcinoma Reveals Critical Genes Underlying Tumorigenesis and Poor Prognosis. American Journal of Human Genetics, 2016, 98, 709-727.	6.2	129
79	A comprehensive genomic characterization of esophageal squamous cell carcinoma: from prognostic analysis to in vivo assay. Chinese Journal of Cancer, 2016, 35, 76.	4.9	7
80	Prognostic efficacy of combining tumor volume with Epstein-Barr virus DNA in patients treated with intensity-modulated radiotherapy for nasopharyngeal carcinoma. Oral Oncology, 2016, 60, 18-24.	1.5	35
81	Association between XRCC3 Thr241Met polymorphism and nasopharyngeal carcinoma risk: evidence from a large-scale case-control study and a meta-analysis. Tumor Biology, 2016, 37, 14825-14830.	1.8	8
82	An extended genome-wide association study identifies novel susceptibility loci for nasopharyngeal carcinoma. Human Molecular Genetics, 2016, 25, 3626-3634.	2.9	42
83	Cumulative scores based on plasma D-dimer and serum albumin levels predict survival in esophageal squamous cell carcinoma patients treated with transthoracic esophagectomy. Chinese Journal of Cancer, 2016, 35, 11.	4.9	15
84	CPEB4 interacts with Vimentin and involves in progressive features and poor prognosis of patients with astrocytic tumors. Tumor Biology, 2016, 37, 5075-5087.	1.8	6
85	Global trends in incidence and mortality of nasopharyngeal carcinoma. Cancer Letters, 2016, 374, 22-30.	7.2	330
86	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. Gastroenterology, 2016, 150, 1633-1645.	1.3	97
87	A GWAS Meta-analysis and Replication Study Identifies a Novel Locus within CLPTM1L/TERT Associated with Nasopharyngeal Carcinoma in Individuals of Chinese Ancestry. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 188-192.	2.5	45
88	Association between genetic variants in the XPG gene and gastric cancer risk in a Southern Chinese population. Aging, 2016, 8, 3311-3320.	3.1	30
89	Epstein-Barr virus mir-bart1-5p detection via nasopharyngeal brush sampling is effective for diagnosing nasopharyngeal carcinoma. Oncotarget, 2016, 7, 4972-4980.	1.8	34
90	XPGrs2296147 T>C polymorphism predicted clinical outcome in colorectal cancer. Oncotarget, 2016, 7, 11724-11732.	1.8	17

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91	A single nucleotide polymorphism in the Epstein-Barr virus genome is strongly associated with a high risk of nasopharyngeal carcinoma. Chinese Journal of Cancer, 2015, 34, 563-72.	4.9	28
92	Identification of surrogate endpoints in patients with locoregionally advanced nasopharyngeal carcinoma receiving neoadjuvant chemotherapy plus concurrent chemoradiotherapy versus concurrent chemoradiotherapy alone. BMC Cancer, 2015, 15, 930.	2.6	6
93	Household inhalants exposure and nasopharyngeal carcinoma risk: a large-scale case-control study in Guangdong, China. BMC Cancer, 2015, 15, 1022.	2.6	32
94	High Expression of LAMP3 Is a Novel Biomarker of Poor Prognosis in Patients with Esophageal Squamous Cell Carcinoma. International Journal of Molecular Sciences, 2015, 16, 17655-17667.	4.1	31
95	High expression of Talin-1 is associated with poor prognosis in patients with nasopharyngeal carcinoma. BMC Cancer, 2015, 15, 332.	2.6	21
96	Identification of miR-143 as a tumour suppressor in nasopharyngeal carcinoma based on microRNA expression profiling. International Journal of Biochemistry and Cell Biology, 2015, 61, 120-128.	2.8	30
97	Pregnancy associated nasopharyngeal carcinoma: A retrospective case-control analysis of maternal survival outcomes. Radiotherapy and Oncology, 2015, 116, 125-130.	0.6	8
98	Increased RIPK4 expression is associated with progression and poor prognosis in cervical squamous cell carcinoma patients. Scientific Reports, 2015, 5, 11955.	3.3	42
99	TERT Polymorphism rs2736100-C Is Associated with EGFR Mutationâ€“Positive Nonâ€“Small Cell Lung Cancer. Clinical Cancer Research, 2015, 21, 5173-5180.	7.0	47
100	A genetic variant in CHRNB3-CHRNA6 increases risk of esophageal squamous cell carcinoma in Chinese populations. Carcinogenesis, 2015, 36, 538-542.	2.8	5
101	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. Nature Communications, 2015, 6, 7138.	12.8	138
102	Association of the Asp312Asn and Lys751Gln polymorphisms in the XPD gene with the risk of non-Hodgkinâ€™s lymphoma: evidence from a meta-analysis. Chinese Journal of Cancer, 2015, 34, 108-14.	4.9	326
103	Quantification of Epsteinâ€“Barr virus <scp>DNA</scp> load in nasopharyngeal brushing samples in the diagnosis of nasopharyngeal carcinoma in southern China. Cancer Science, 2015, 106, 1196-1201.	3.9	54
104	Genome-Wide Identification of a Methylation Gene Panel as a Prognostic Biomarker in Nasopharyngeal Carcinoma. Molecular Cancer Therapeutics, 2015, 14, 2864-2873.	4.1	80
105	No association betweenMTRrs1805087 A > G polymorphism and non-Hodgkin lymphoma susceptibility: evidence from 11 486 subjects. Leukemia and Lymphoma, 2015, 56, 763-767.	1.3	13
106	Association of MTHFR C677T and A1298C polymorphisms with non-Hodgkin lymphoma susceptibility: Evidence from a meta-analysis. Scientific Reports, 2015, 4, 6159.	3.3	83
107	Comparison of Long-Term Survival and Toxicity of Cisplatin Delivered Weekly versus Every Three Weeks Concurrently with Intensity-Modulated Radiotherapy in Nasopharyngeal Carcinoma. PLoS ONE, 2014, 9, e110765.	2.5	31
108	Weighted Risk Score-Based Multifactor Dimensionality Reduction to Detect Gene-Gene Interactions in Nasopharyngeal Carcinoma. International Journal of Molecular Sciences, 2014, 15, 10724-10737.	4.1	8

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109	The association between the polymorphisms of TNF-Î± and non-Hodgkin lymphoma: a meta-analysis. Tumor Biology, 2014, 35, 12509-12517.	1.8	15
110	Large-scale genetic study in East Asians identifies six new loci associated with colorectal cancer risk. Nature Genetics, 2014, 46, 533-542.	21.4	212
111	Genome-wide association study identifies three susceptibility loci for laryngeal squamous cell carcinoma in the Chinese population. Nature Genetics, 2014, 46, 1110-1114.	21.4	57
112	Genome-wide association study identifies new susceptibility loci for epithelial ovarian cancer in Han Chinese women. Nature Communications, 2014, 5, 4682.	12.8	59
113	Overexpression of CIP2A is an independent prognostic indicator in nasopharyngeal carcinoma and its depletion suppresses cell proliferation and tumor growth. Molecular Cancer, 2014, 13, 111.	19.2	21
114	Glatiramer acetate reverses cognitive deficits from cranial-irradiated rat by inducing hippocampal neurogenesis. Journal of Neuroimmunology, 2014, 271, 1-7.	2.3	19
115	Association of BRCA2 N372H polymorphism with cancer susceptibility: A comprehensive review and meta-analysis. Scientific Reports, 2014, 4, 6791.	3.3	33
116	The Pretreatment Albumin to Globulin Ratio Has Predictive Value for Long-Term Mortality in Nasopharyngeal Carcinoma. PLoS ONE, 2014, 9, e94473.	2.5	99
117	Genome-wide association analyses in east Asians identify new susceptibility loci for colorectal cancer. Nature Genetics, 2013, 45, 191-196.	21.4	173
118	Genome-wide association study identifies common variants in SLC39A6 associated with length of survival in esophageal squamous-cell carcinoma. Nature Genetics, 2013, 45, 632-638.	21.4	97
119	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. American Journal of Epidemiology, 2013, 177, 242-250.	3.4	108
120	Quantitative Association of Tobacco Smoking With the Risk of Nasopharyngeal Carcinoma: A Comprehensive Meta-Analysis of Studies Conducted Between 1979 and 2011. American Journal of Epidemiology, 2013, 178, 325-338.	3.4	89
121	Developing Genetic Epidemiological Models to Predict Risk for Nasopharyngeal Carcinoma in High-Risk Population of China. PLoS ONE, 2013, 8, e56128.	2.5	21
122	An Epidemiological and Molecular Study of the Relationship Between Smoking, Risk of Nasopharyngeal Carcinoma, and Epsteinâ€“Barr Virus Activation. Journal of the National Cancer Institute, 2012, 104, 1396-1410.	6.3	164
123	Establishment of VCA and EBNA1 IgAâ€based combination by enzymeâ€linked immunosorbent assay as preferred screening method for nasopharyngeal carcinoma: a twoâ€stage design with a preliminary performance study and a mass screening in southern China. International Journal of Cancer, 2012, 131, 406-416.	5.1	116
124	Non-viral environmental risk factors for nasopharyngeal carcinoma: A systematic review. Seminars in Cancer Biology, 2012, 22, 117-126.	9.6	151
125	Familial and large-scale caseâ€“control studies identify genes associated with nasopharyngeal carcinoma. Seminars in Cancer Biology, 2012, 22, 96-106.	9.6	58
126	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. Nature Genetics, 2011, 43, 679-684.	21.4	260

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127	Fluctuations of Epstein-Barr Virus Serological Antibodies and Risk for Nasopharyngeal Carcinoma: A Prospective Screening Study with a 20-Year Follow-Up. PLoS ONE, 2011, 6, e19100.	2.5	129
128	High Expression of p300 Has an Unfavorable Impact on Survival in Resectable Esophageal Squamous Cell Carcinoma. Annals of Thoracic Surgery, 2011, 91, 1531-1538.	1.3	49
129	Elevated Epsteinâ€“Barr virus seroreactivity among unaffected members of families with nasopharyngeal carcinoma. Journal of Medical Virology, 2011, 83, 1792-1798.	5.0	13
130	Comprehensive Pathway-Based Association Study of DNA Repair Gene Variants and the Risk of Nasopharyngeal Carcinoma. Cancer Research, 2011, 71, 3000-3008.	0.9	41
131	Alcohol and tea consumption in relation to the risk of nasopharyngeal carcinoma in Guangdong, China. Frontiers of Medicine in China, 2010, 4, 448-456.	0.1	21
132	Traditional Cantonese diet and nasopharyngeal carcinoma risk: a large-scale case-control study in Guangdong, China. BMC Cancer, 2010, 10, 446.	2.6	118
133	Effect of family history of cancers and environmental factors on risk of nasopharyngeal carcinoma in Guangdong, China. Cancer Epidemiology, 2010, 34, 419-424.	1.9	53
134	A genome-wide association study of nasopharyngeal carcinoma identifies three new susceptibility loci. Nature Genetics, 2010, 42, 599-603.	21.4	374
135	A Case-control and a family-based association study revealing an association between CYP2E1 polymorphisms and nasopharyngeal carcinoma risk in Cantonese. Carcinogenesis, 2009, 30, 2031-2036.	2.8	43
136	Antibodies against Epsteinâ€“Barr virus gp78 antigen: a novel marker for serological diagnosis of nasopharyngeal carcinoma detected by xMAP technology. Journal of General Virology, 2008, 89, 1152-1158.	2.9	20
137	Association of pIgR polymorphisms with nasopharyngeal carcinoma. Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association, Beijing Institute for Cancer Research, 2006, 18, 168-172.	2.2	0
138	Trends in incidence and mortality of nasopharyngeal carcinoma over a 20â€“25 year period (1978/1983â€“2002) in Sihui and Cangwu counties in southern China. BMC Cancer, 2006, 6, 178.	2.6	199
139	Familial risk and clustering of nasopharyngeal carcinoma in Guangdong, China. Cancer, 2004, 101, 363-369.	4.1	79
140	Genome-wide scan for familial nasopharyngeal carcinoma reveals evidence of linkage to chromosome 4. Nature Genetics, 2002, 31, 395-399.	21.4	217