

Qingyi Wei

List of Publications by Year in descending order

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Version: 2024-02-01

592
papers

26,477
citations

7561

77
h-index

15716

125
g-index

598
all docs

598
docs citations

598
times ranked

25903
citing authors

#	ARTICLE	IF	CITATIONS
1	Risk of Acute Lung Injury after Esophagectomy. <i>Seminars in Thoracic and Cardiovascular Surgery</i> , 2022, 34, 737-746.	0.4	3
2	Systems biomarker characteristics of circulating alkaline phosphatase activities for 48 types of human diseases. <i>Current Medical Research and Opinion</i> , 2022, 38, 201-209.	0.9	2
3	Genetic variants in <i>CYP2B6</i> and <i>HSD17B12</i> associated with risk of squamous cell carcinoma of the head and neck. <i>International Journal of Cancer</i> , 2022, 151, 553-564.	2.3	7
4	Genetic variants in <i>DDO</i> and <i>PEX5L</i> in peroxisome-related pathways predict non-small cell lung cancer survival. <i>Molecular Carcinogenesis</i> , 2022, 61, 619-628.	1.3	2
5	Deciphering associations between three RNA splicing-related genetic variants and lung cancer risk. <i>Npj Precision Oncology</i> , 2022, 6, .	2.3	1
6	Potentially functional genetic variants of the notch signaling pathway genes predict survival of Chinese patients with esophageal squamous cell carcinoma. <i>Journal of Gene Medicine</i> , 2022, 24, .	1.4	1
7	Circular RNA circCSNK1G3 induces HOXA10 signaling and promotes the growth and metastasis of lung adenocarcinoma cells through hsa-miR-143-3p sponging. <i>Cellular Oncology (Dordrecht)</i> , 2021, 44, 297-310.	2.1	21
8	Necessity of preoperative bone scintigraphy for cT1N0 lung cancer: Evidence from retrospective to prospective study. <i>Thoracic Cancer</i> , 2021, 12, 413-414.	0.8	1
9	Potentially functional variants of ERAP1, PSMF1 and NCF2 in the MHC-I-related pathway predict non-small cell lung cancer survival. <i>Cancer Immunology, Immunotherapy</i> , 2021, 70, 2819-2833.	2.0	8
10	Novel functional variants in the Notch pathway and survival of Chinese colorectal cancer. <i>International Journal of Cancer</i> , 2021, 149, 84-96.	2.3	3
11	Association of genetic variants of TMEM135 and PEX5 in the peroxisome pathway with cutaneous melanoma-specific survival. <i>Annals of Translational Medicine</i> , 2021, 9, 396-396.	0.7	3
12	Genetic variants of <i>CHEK1</i> , <i>PRIM2</i> and <i>CDK6</i> in the mitotic phase-related pathway are associated with nonsmall cell lung cancer survival. <i>International Journal of Cancer</i> , 2021, 149, 1302-1312.	2.3	9
13	Potentially functional variants of HBEGF and ITPR3 in GnRH signaling pathway genes predict survival of non-small cell lung cancer patients. <i>Translational Research</i> , 2021, 233, 92-103.	2.2	14
14	Association of genetic variants of <i>FBXO32</i> and <i>FOXO6</i> in the FOXO pathway with breast cancer risk. <i>Molecular Carcinogenesis</i> , 2021, 60, 661-670.	1.3	4
15	Genetic Risk for Overall Cancer and the Benefit of Adherence to a Healthy Lifestyle. <i>Cancer Research</i> , 2021, 81, 4618-4627.	0.4	48
16	Electromagnetic navigational bronchoscopy-guided dye marking to identify the subsegmental bronchus in thoracoscopic anatomic subsegmentectomy. <i>Thoracic Cancer</i> , 2021, 12, 2819-2821.	0.8	3
17	Genetic variants of SDCCAG8 and MAGI2 in mitosis-related pathway genes are independent predictors of cutaneous melanoma-specific survival. <i>Cancer Science</i> , 2021, 112, 4355-4364.	1.7	1
18	Genetic Variants of CLPP and M1AP Are Associated With Risk of Non-Small Cell Lung Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 709829.	1.3	1

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19	A pleiotropic ATM variant (rs1800057 C>G) is associated with risk of multiple cancers. <i>Carcinogenesis</i> , 2021, , .	1.3	1
20	Genetic Variants of CLEC4E and BIRC3 in Damage-Associated Molecular Patterns-Related Pathway Genes Predict Non-Small Cell Lung Cancer Survival. <i>Frontiers in Oncology</i> , 2021, 11, 717109.	1.3	6
21	Predictive value of a novel Asian lung cancer screening nomogram based on artificial intelligence and epidemiological characteristics. <i>Thoracic Cancer</i> , 2021, 12, 3130-3140.	0.8	4
22	Genetic variants of , and in the natural killer cell-related pathway are associated with non-small cell lung cancer survival. <i>American Journal of Cancer Research</i> , 2021, 11, 2264-2277.	1.4	0
23	Association of pretreatment body mass index with risk of head and neck cancer: a large single-center study. <i>American Journal of Cancer Research</i> , 2021, 11, 2343-2350.	1.4	2
24	Genetic variants of and in myeloid cell-related pathway genes independently predict cutaneous melanoma-specific survival. <i>American Journal of Cancer Research</i> , 2021, 11, 3252-3262.	1.4	0
25	Genome-wide association and functional interrogation identified a variant at 3p26.1 modulating ovarian cancer survival among Chinese women. <i>Cell Discovery</i> , 2021, 7, 121.	3.1	5
26	Novel genetic variants in KIF16B and NEDD4L in the endosomeâ€related genes are associated with nonsmall cell lung cancer survival. <i>International Journal of Cancer</i> , 2020, 147, 392-403.	2.3	6
27	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
28	Common genetic variation and risk of osteosarcoma in a multi-ethnic pediatric and adolescent population. <i>Bone</i> , 2020, 130, 115070.	1.4	22
29	Novel genetic variants in <i>HDAC2</i> and <i>PPARGC1A</i> of the CREBâ€binding protein pathway predict survival of nonâ€smallâ€cell lung cancer. <i>Molecular Carcinogenesis</i> , 2020, 59, 104-115.	1.3	13
30	Robotic sleeve lobectomy for centrally located nonâ€small cell lung cancer: A propensity scoreâ€weighted comparison with thoracoscopic and open surgery. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2020, 160, 838-846.e2.	0.4	45
31	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. <i>Lancet Oncology</i> , The, 2020, 21, 1378-1386.	5.1	123
32	Evaluated Lymph Nodeâ€Number in Population-Based Analyses: Significant Prognostic Factor, but Assess With Caution. <i>Journal of Thoracic Oncology</i> , 2020, 15, e77-e78.	0.5	0
33	Genetic variation associated with childhood and adult stature and risk of <i>MYCN</i> â€amplified neuroblastoma. <i>Cancer Medicine</i> , 2020, 9, 8216-8225.	1.3	3
34	Genetic variants in TKT and DERA in the nicotinamide adenine dinucleotide phosphate pathway predict melanoma survival. <i>European Journal of Cancer</i> , 2020, 136, 84-94.	1.3	3
35	Novel genetic variants in genes of the Fc gamma receptor-mediated phagocytosis pathway predict non-small cell lung cancer survival. <i>Translational Lung Cancer Research</i> , 2020, 9, 575-586.	1.3	6
36	Associations between genetic variants of KIF5B , FMN1 , and MGAT3 in the cadherin pathway and pancreatic cancer risk. <i>Cancer Medicine</i> , 2020, 9, 9620-9631.	1.3	1

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37	Functional genetic variants of <i>CTNNBIP1</i> predict platinum treatment response of Chinese epithelial ovarian cancer patients. <i>Journal of Cancer</i> , 2020, 11, 6850-6860.	1.2	6
38	Genetic variants of the peroxisome proliferator-activated receptor (PPAR) signaling pathway genes and risk of pancreatic cancer. <i>Molecular Carcinogenesis</i> , 2020, 59, 930-939.	1.3	11
39	Novel Variants of <i>ELP2</i> and <i>PIAS1</i> in the Interferon Gamma Signaling Pathway Are Associated with Non-Small Cell Lung Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1679-1688.	1.1	2
40	The Rare Variant rs35356162 in <i>UHRF1BP1</i> Increases Bladder Cancer Risk in Han Chinese Population. <i>Frontiers in Oncology</i> , 2020, 10, 134.	1.3	16
41	Genetic variants in the human leukocyte antigen region and survival of Chinese patients with non-small cell lung carcinoma. <i>Carcinogenesis</i> , 2020, 41, 1203-1212.	1.3	3
42	Potentially functional genetic variants in <i>PLIN2</i> , <i>SULT2A1</i> and <i>UGT1A9</i> genes of the ketone pathway and survival of nonsmall cell lung cancer. <i>International Journal of Cancer</i> , 2020, 147, 1559-1570.	2.3	8
43	Relevance and prognostic ability of Twist, Slug and tumor spread through air spaces in lung adenocarcinoma. <i>Cancer Medicine</i> , 2020, 9, 1986-1998.	1.3	13
44	Novel Genetic Variants of <i>ALG6</i> and <i>GALNTL4</i> of the Glycosylation Pathway Predict Cutaneous Melanoma-Specific Survival. <i>Cancers</i> , 2020, 12, 288.	1.7	7
45	Genetic variants in the folate metabolic pathway genes predict cutaneous melanoma-specific survival. <i>British Journal of Dermatology</i> , 2020, 183, 719-728.	1.4	4
46	Genetic variants in <i>PDSS1</i> and <i>SLC16A6</i> of the ketone body metabolic pathway predict cutaneous melanoma-specific survival. <i>Molecular Carcinogenesis</i> , 2020, 59, 640-650.	1.3	9
47	<i>APOB</i> Genotypes and <i>CDH13</i> Haplotypes in the Cholesterol-Related Pathway Genes Predict Non-Small Cell Lung Cancer Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1204-1213.	1.1	7
48	A Genome-Wide Association Study Identifies Two Novel Susceptible Regions for Squamous Cell Carcinoma of the Head and Neck. <i>Cancer Research</i> , 2020, 80, 2451-2460.	0.4	33
49	Novel genetic variants of and involved in immunoregulatory interactions are associated with non-small cell lung cancer survival. <i>American Journal of Cancer Research</i> , 2020, 10, 1770-1784.	1.4	2
50	Variants in , and in vitamin D pathway genes are associated with breast cancer risk: a large-scale analysis of 14 GWASs in the DRIVE study. <i>American Journal of Cancer Research</i> , 2020, 10, 2160-2173.	1.4	2
51	Associations of novel variants in , and of the ATM pathway genes with pancreatic cancer risk. <i>American Journal of Cancer Research</i> , 2020, 10, 2128-2144.	1.4	2
52	Novel genetic variants of and related lymphangiogenesis signaling pathway predict non-small cell lung cancer survival. <i>American Journal of Cancer Research</i> , 2020, 10, 2603-2616.	1.4	1
53	Genetic variants of and in the NLRP3 inflammasome pathway are associated with non-small cell lung cancer survival. <i>American Journal of Cancer Research</i> , 2020, 10, 2582-2595.	1.4	6
54	Novel genetic variants of and of the endosome-related pathway predict cutaneous melanoma-specific survival. <i>American Journal of Cancer Research</i> , 2020, 10, 3382-3394.	1.4	0

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55	Functional variant of <i>MTOR</i> rs2536 and survival of Chinese gastric cancer patients. <i>International Journal of Cancer</i> , 2019, 144, 251-262.	2.3	5
56	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
57	Short-term outcomes of typical versus atypical lung segmentectomy by minimally invasive surgeries. <i>Thoracic Cancer</i> , 2019, 10, 1812-1818.	0.8	11
58	Robotic Bronchial Sleeve Lobectomy for Central Lung Tumors: Technique and Outcome. <i>Annals of Thoracic Surgery</i> , 2019, 108, 211-218.	0.7	32
59	A genetic variant within <i>MDM4</i> 3'UTR miRNA binding site is associated with HPV16-positive tumors and survival of oropharyngeal cancer. <i>Molecular Carcinogenesis</i> , 2019, 58, 2276-2285.	1.3	5
60	Genetic variants in glutamine metabolic pathway genes predict cutaneous melanoma-specific survival. <i>Molecular Carcinogenesis</i> , 2019, 58, 2091-2103.	1.3	5
61	Impact of visceral pleural invasion on the association of extent of lymphadenectomy and survival in stage I non-small cell lung cancer. <i>Cancer Medicine</i> , 2019, 8, 669-678.	1.3	24
62	Role of Immune Response, Inflammation, and Tumor Immune Response-Related Cytokines/Chemokines in Melanoma Progression. <i>Journal of Investigative Dermatology</i> , 2019, 139, 2352-2358.e3.	0.3	23
63	Functional genetic variants of <i>RUVBL1</i> predict overall survival of Chinese patients with epithelial ovarian cancer. <i>Carcinogenesis</i> , 2019, 40, 1209-1219.	1.3	5
64	Three novel genetic variants in <i>NRF2</i> signaling pathway genes are associated with pancreatic cancer risk. <i>Cancer Science</i> , 2019, 110, 2022-2032.	1.7	14
65	Genetic variants in the liver kinase B1-activated protein kinase pathway genes and pancreatic cancer risk. <i>Molecular Carcinogenesis</i> , 2019, 58, 1338-1348.	1.3	14
66	Potentially functional genetic variants in the <i>TNF/TNFR</i> signaling pathway genes predict survival of patients with non-small cell lung cancer in the PLCO cancer screening trial. <i>Molecular Carcinogenesis</i> , 2019, 58, 1094-1104.	1.3	9
67	Genetic variants of genes in the <i>NER</i> pathway associated with risk of breast cancer: A large-scale analysis of 14 published GWAS datasets in the DRIVE study. <i>International Journal of Cancer</i> , 2019, 145, 1270-1279.	2.3	13
68	Predictive value of a prognostic model based on pathologic features in lung invasive adenocarcinoma. <i>Lung Cancer</i> , 2019, 131, 14-22.	0.9	18
69	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. <i>Gynecologic Oncology</i> , 2019, 153, 343-355.	0.6	28
70	Potential functional variants in <i>SMC2</i> and <i>TP53</i> in the <i>AURORA</i> pathway genes and risk of pancreatic cancer. <i>Carcinogenesis</i> , 2019, 40, 521-528.	1.3	17
71	Lymphocyte telomere length predicts clinical outcomes of HPV-positive oropharyngeal cancer patients after definitive radiotherapy. <i>Carcinogenesis</i> , 2019, 40, 735-741.	1.3	5
72	Genetic variants in <i>ELOVL2</i> and <i>HSD17B12</i> predict melanoma-specific survival. <i>International Journal of Cancer</i> , 2019, 145, 2619-2628.	2.3	11

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73	Genetic variants in the calcium signaling pathway genes are associated with cutaneous melanoma-specific survival. <i>Carcinogenesis</i> , 2019, 40, 279-288.	1.3	6
74	Genetic variants in <i>RUNX3</i> , <i>AMD1</i> and <i>MSRA</i> in the methionine metabolic pathway and survival in nonsmall cell lung cancer patients. <i>International Journal of Cancer</i> , 2019, 145, 621-631.	2.3	21
75	Robotic circumferential tracheal resection and reconstruction via a completely portal approach. <i>Thoracic Cancer</i> , 2019, 10, 378-380.	0.8	4
76	Potentially functional genetic variants in the complement-related immunity gene set are associated with non-small cell lung cancer survival. <i>International Journal of Cancer</i> , 2019, 144, 1867-1876.	2.3	14
77	<i>TGFβ1</i> Genetic Variants Predict Clinical Outcomes of HPV-Positive Oropharyngeal Cancer Patients after Definitive Radiotherapy. <i>Clinical Cancer Research</i> , 2018, 24, 2225-2233.	3.2	20
78	A <i>TGFβ1</i> genetic variant at the miRNA187 binding site significantly modifies risk of HPV16-associated oropharyngeal cancer. <i>International Journal of Cancer</i> , 2018, 143, 1327-1334.	2.3	7
79	Genetic variants in <i>RORA</i> and <i>DNMT1</i> associated with cutaneous melanoma survival. <i>International Journal of Cancer</i> , 2018, 142, 2303-2312.	2.3	13
80	Potentially Functional Variants of ATG16L2 Predict Radiation Pneumonitis and Outcomes in Patients with Non-Small Cell Lung Cancer after Definitive Radiotherapy. <i>Journal of Thoracic Oncology</i> , 2018, 13, 660-675.	0.5	29
81	Inverse Relationship between Vitiligo-Related Genes and Skin Cancer Risk. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2072-2075.	0.3	20
82	Associations between expression levels of nucleotide excision repair proteins in lymphoblastoid cells and risk of squamous cell carcinoma of the head and neck. <i>Molecular Carcinogenesis</i> , 2018, 57, 784-793.	1.3	5
83	Novel genetic variants in the P38MAPK pathway gene <i>ZAK</i> and susceptibility to lung cancer. <i>Molecular Carcinogenesis</i> , 2018, 57, 216-224.	1.3	9
84	Mouse double minute 4 variants modify susceptibility to risk of recurrence in patients with squamous cell carcinoma of the oropharynx. <i>Molecular Carcinogenesis</i> , 2018, 57, 361-369.	1.3	6
85	Genetic variants in the platelet-derived growth factor subunit B gene associated with pancreatic cancer risk. <i>International Journal of Cancer</i> , 2018, 142, 1322-1331.	2.3	20
86	An <i>ERCC4</i> regulatory variant predicts grade 3 or 4 toxicities in patients with advanced non-small cell lung cancer treated by platinum-based therapy. <i>International Journal of Cancer</i> , 2018, 142, 1218-1229.	2.3	7
87	Genetic variants in the metzincin metallopeptidase family genes predict melanoma survival. <i>Molecular Carcinogenesis</i> , 2018, 57, 22-31.	1.3	5
88	Predictors of nodal metastasis and prognostic significance of lymph node ratio and total lymph node count in tracheobronchial adenoid cystic carcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 5919-5925.	0.9	13
89	Potential clinical application of lncRNAs in non-small cell lung cancer. <i>OncoTargets and Therapy</i> , 2018, Volume 11, 8045-8052.	1.0	79
90	Genetic determinants of childhood and adult height associated with osteosarcoma risk. <i>Cancer</i> , 2018, 124, 3742-3752.	2.0	20

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91	Identification of an eight-gene prognostic signature for lung adenocarcinoma. <i>Cancer Management and Research</i> , 2018, Volume 10, 3383-3392.	0.9	63
92	Identification of SPP1 as a promising biomarker to predict clinical outcome of lung adenocarcinoma individuals. <i>Gene</i> , 2018, 679, 398-404.	1.0	51
93	DNA repair capacity correlates with standardized uptake values from 18 F-fluorodeoxyglucose positron emission tomography/CT in patients with advanced non-small-cell lung cancer. <i>Chronic Diseases and Translational Medicine</i> , 2018, 4, 109-116.	0.9	1
94	Genetic variants in nucleotide excision repair pathway predict survival of esophageal squamous cell cancer patients receiving platinum-based chemotherapy. <i>Molecular Carcinogenesis</i> , 2018, 57, 1553-1565.	1.3	9
95	Variants in Notch signalling pathway genes, <i>PCSEN1</i> and <i>MAML2</i> , predict overall survival in Chinese patients with epithelial ovarian cancer. <i>Journal of Cellular and Molecular Medicine</i> , 2018, 22, 4975-4984.	1.6	14
96	Genetic variant of <i>IRAK2</i> in the toll-like receptor signaling pathway and survival of non-small cell lung cancer. <i>International Journal of Cancer</i> , 2018, 143, 2400-2408.	2.3	14
97	Single-nucleotide polymorphisms of stemness genes predicted to regulate RNA splicing, microRNA and oncogenic signaling are associated with prostate cancer survival. <i>Carcinogenesis</i> , 2018, 39, 879-888.	1.3	9
98	Genetic correction improves prediction efficiency of serum tumor biomarkers on digestive cancer risk in the elderly Chinese cohort study. <i>Oncotarget</i> , 2018, 9, 7389-7397.	0.8	7
99	Apoptotic capacity and risk of squamous cell carcinoma of the head and neck. <i>European Journal of Cancer</i> , 2017, 72, 166-176.	1.3	19
100	Modifying effect of mouse double minute-2 promoter variants on risk of recurrence for patients with squamous cell carcinoma of oropharynx. <i>Scientific Reports</i> , 2017, 7, 39765.	1.6	4
101	Pathway analysis of published genome-wide association studies of lung cancer: A potential role for the <i>CYP4F3</i> locus. <i>Molecular Carcinogenesis</i> , 2017, 56, 1663-1672.	1.3	13
102	<i>BRCA1</i> and <i>BRCA2</i> mutations in ovarian cancer patients from China: ethnic-related mutations in <i>BRCA1</i> associated with an increased risk of ovarian cancer. <i>International Journal of Cancer</i> , 2017, 140, 2051-2059.	2.3	45
103	Reduced mRNA expression of nucleotide excision repair genes in lymphocytes and risk of squamous cell carcinoma of the head and neck. <i>Carcinogenesis</i> , 2017, 38, 504-510.	1.3	6
104	Genetic Variants in <i>WNT2B</i> and <i>BTRC</i> Predict Melanoma Survival. <i>Journal of Investigative Dermatology</i> , 2017, 137, 1749-1756.	0.3	5
105	Associations between RNA splicing regulatory variants of stemness-related genes and racial disparities in susceptibility to prostate cancer. <i>International Journal of Cancer</i> , 2017, 141, 731-743.	2.3	20
106	Genetic variants of <i>PTPN2</i> are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. <i>Scientific Reports</i> , 2017, 7, 825.	1.6	10
107	Association between Body Mass Index, C-Reactive Protein Levels, and Melanoma Patient Outcomes. <i>Journal of Investigative Dermatology</i> , 2017, 137, 1792-1795.	0.3	40
108	Genetic variants in the integrin signaling pathway genes predict cutaneous melanoma survival. <i>International Journal of Cancer</i> , 2017, 140, 1270-1279.	2.3	4

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109	Genetic variants in the genes encoding rho GTPases and related regulators predict cutaneous melanoma-specific survival. <i>International Journal of Cancer</i> , 2017, 141, 721-730.	2.3	8
110	Carbon Quantum Dots Decorated C ₃ N ₄ /TiO ₂ Heterostructure Nanorod Arrays for Enhanced Photoelectrochemical Performance. <i>Journal of the Electrochemical Society</i> , 2017, 164, H515-H520.	1.3	22
111	A <i>PGC1²</i> genetic variant associated with nevus count and melanoma mortality. <i>International Journal of Cancer</i> , 2017, 141, 1066-1067.	2.3	5
112	Association between miRNA-binding site polymorphisms in double-strand break repair genes and risk of recurrence in patients with squamous cell carcinomas of the non-oropharynx. <i>Carcinogenesis</i> , 2017, 38, 432-438.	1.3	6
113	Towards precision prevention: Technologies for identifying healthy individuals with high risk of disease. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2017, 800-802, 14-28.	0.4	20
114	Functional variants in DCAF4 associated with lung cancer risk in European populations. <i>Carcinogenesis</i> , 2017, 38, 541-551.	1.3	16
115	Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. <i>Scientific Reports</i> , 2017, 7, 44634.	1.6	10
116	Genetic variants in ERCC1 and XPC predict survival outcome of non-small cell lung cancer patients treated with platinum-based therapy. <i>Scientific Reports</i> , 2017, 7, 10702.	1.6	12
117	Diagnostic accuracy of serum antibodies to human papillomavirus type 16 early antigens in the detection of human papillomavirus-related oropharyngeal cancer. <i>Cancer</i> , 2017, 123, 4886-4894.	2.0	16
118	Assessing the spectrum of germline variation in Fanconi anemia genes among patients with head and neck carcinoma before age 50. <i>Cancer</i> , 2017, 123, 3943-3954.	2.0	37
119	Functional genetic variants of <i>XRCC4</i> and <i>ERCC1</i> predict survival of gastric cancer patients treated with chemotherapy by regulating the gene expression. <i>Molecular Carcinogenesis</i> , 2017, 56, 2706-2717.	1.3	4
120	Genetic variants in microRNA-binding sites of DNA repair genes as predictors of recurrence in patients with squamous cell carcinoma of the oropharynx. <i>International Journal of Cancer</i> , 2017, 141, 1355-1364.	2.3	9
121	Genetic variants of JNK and p38 pathways and risk of non-small cell lung cancer in an eastern Chinese population. <i>International Journal of Cancer</i> , 2017, 140, 807-817.	2.3	8
122	E2F transcription factor 2 variants as predictive biomarkers for recurrence risk in patients with squamous cell carcinoma of the oropharynx. <i>Molecular Carcinogenesis</i> , 2017, 56, 1335-1343.	1.3	13
123	Susceptibility loci of <i>CNOT6</i> in the general mRNA degradation pathway and lung cancer risk: A re-analysis of eight GWASs. <i>Molecular Carcinogenesis</i> , 2017, 56, 1227-1238.	1.3	10
124	Melanoma Expression Genes Identified through Genome-Wide Association Study of Breslow Tumor Thickness. <i>Journal of Investigative Dermatology</i> , 2017, 137, 253-257.	0.3	2
125	A functional variant at the miRNA binding site in <i>E2F1</i> gene is associated with risk and tumor HPV16 status of oropharynx squamous cell carcinoma. <i>Molecular Carcinogenesis</i> , 2017, 56, 1100-1106.	1.3	12
126	Functional variants in the low-density lipoprotein receptor gene are associated with clear cell renal cell carcinoma susceptibility. <i>Carcinogenesis</i> , 2017, 38, 1241-1248.	1.3	5

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127	Genetic variants of PDGF signaling pathway genes predict cutaneous melanoma survival. <i>Oncotarget</i> , 2017, 8, 74595-74606.	0.8	3
128	THE AUTHORS REPLY. <i>American Journal of Epidemiology</i> , 2017, 186, 625-626.	1.6	0
129	Two-stage genome-wide association study identifies a novel susceptibility locus associated with melanoma. <i>Oncotarget</i> , 2017, 8, 17586-17592.	0.8	61
130	Polymorphisms in nucleotide excision repair genes and risk of primary prostate cancer in Chinese Han populations. <i>Oncotarget</i> , 2017, 8, 24362-24371.	0.8	21
131	Abstract B58: Single-nucleotide polymorphisms of race-related alternatively spliced genes associate with prostate cancer risk, aggressiveness and/or survival. , 2017, , .		0
132	BRCA1 and BRCA2 mutations in ovarian cancer patients from China: Association of ethnic-specific mutations in BRCA1 with an increased risk of ovarian cancer.. <i>Journal of Clinical Oncology</i> , 2017, 35, 1583-1583.	0.8	1
133	A functional variant at <i>miRNA-122</i> binding site in <i>IL-1β</i> 3' UTR predicts risk of recurrence in patients with oropharyngeal cancer. <i>Oncotarget</i> , 2016, 7, 34472-34479.	0.8	8
134	Genetic variants in the vitamin D pathway genes <i>VDBP</i> and <i>RXRA</i> modulate cutaneous melanoma disease-specific survival. <i>Pigment Cell and Melanoma Research</i> , 2016, 29, 176-185.	1.5	19
135	Genetic variants in <i>ABCG1</i> are associated with survival of nonsmall-cell lung cancer patients. <i>International Journal of Cancer</i> , 2016, 138, 2592-2601.	2.3	41
136	Comparative Effectiveness of 5 Local-Regional Control Strategies for IIIA (N2) Non-small Cell Lung Cancer Using SEER Data: Outcomes After Treatment of 20,468 Patients. <i>Chest</i> , 2016, 149, A275.	0.4	0
137	Genetic variants in the PIWI-miRNA pathway gene <i>DCP1A</i> predict melanoma disease-specific survival. <i>International Journal of Cancer</i> , 2016, 139, 2730-2737.	2.3	21
138	Effect of human papillomavirus seropositivity and <i>E2F2</i> promoter variants on risk of squamous cell carcinomas of oropharynx and oral cavity. <i>Carcinogenesis</i> , 2016, 37, 1070-1078.	1.3	5
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