Qingyi Wei

List of Publications by Year
in descending order

592 papers	26,477 citations	7568 77 h-index	15732 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. Seminars in Thoracic and Cardiovascular Surgery, 2022, 34, 737-746.	0.6	3
2	Systems biomarker characteristics of circulating alkaline phosphatase activities for 48 types of human diseases. Current Medical Research and Opinion, 2022, 38, 201-209.	1.9	2
3	Genetic variants in CYP2B6 and HSD17B12 associated with risk of squamous cell carcinoma of the head and neck. International Journal of Cancer, 2022, 151, 553-564.	5.1	7
4	Genetic variants in DDO and PEX5L in peroxisomeâ€related pathways predict nonâ€small cell lung cancer survival. Molecular Carcinogenesis, 2022, 61, 619-628.	2.7	2
5	Deciphering associations between three RNA splicing-related genetic variants and lung cancer risk. Npj Precision Oncology, 2022, 6, .	5.4	1
6	Potentially functional genetic variants of the notch signaling pathway genes predict survival of Chinese patients with esophageal squamous cell carcinoma. Journal of Gene Medicine, 2022, 24, .	2.8	1
7	Circular RNA circCSNK1G3 induces HOXA10 signaling and promotes the growth and metastasis of lung adenocarcinoma cells through hsa-miR-143-3p sponging. Cellular Oncology (Dordrecht), 2021, 44, 297-310.	4.4	21
8	Necessity of preoperative bone scintigraphy for cT1N0 lung cancer: Evidence from retrospective to prospective study. Thoracic Cancer, 2021, 12, 413-414.	1.9	1
9	Potentially functional variants of ERAP1, PSMF1 and NCF2 in the MHC-I-related pathway predict non-small cell lung cancer survival. Cancer Immunology, Immunotherapy, 2021, 70, 2819-2833.	4.2	8
10	Novel functional variants in the Notch pathway and survival of Chinese colorectal cancer. International Journal of Cancer, 2021, 149, 84-96.	5.1	3
11	Association of genetic variants of TMEM135 and PEX5 in the peroxisome pathway with cutaneous melanoma-specific survival. Annals of Translational Medicine, 2021, 9, 396-396.	1.7	3
12	Genetic variants of <scp>CHEK1</scp>, <scp>PRIM2</scp> and <scp>CDK6</scp> in the mitotic phaseâ€related pathway are associated with nonsmall cell lung cancer survival. International Journal of Cancer, 2021, 149, 1302-1312.	5.1	9
13	Potentially functional variants of HBEGF and ITPR3 in GnRH signaling pathway genes predict survival of non-small cell lung cancer patients. Translational Research, 2021, 233, 92-103.	5.0	14
14	Association of genetic variants of FBXO32 and FOXO6 in the FOXO pathway with breast cancer risk. Molecular Carcinogenesis, 2021, 60, 661-670.	2.7	4
15	Genetic Risk for Overall Cancer and the Benefit of Adherence to a Healthy Lifestyle. Cancer Research, 2021, 81, 4618-4627.	0.9	48
16	Electromagnetic navigational bronchoscopyâ€guided dye marking to identify the subsegmental bronchus in thoracoscopic anatomic subsegmentectomy. Thoracic Cancer, 2021, 12, 2819-2821.	1.9	3
17	Genetic variants of SDCCAG8 and MAGI2 in mitosisâ€related pathway genes are independent predictors of cutaneous melanomaâ€specific survival. Cancer Science, 2021, 112, 4355-4364.	3.9	1
18	Genetic Variants of CLPP and M1AP Are Associated With Risk of Non-Small Cell Lung Cancer. Frontiers in Oncology, 2021, 11, 709829.	2.8	1

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19	A pleiotropic ATM variant (rs1800057 C>G) is associated with risk of multiple cancers. Carcinogenesis, 2021, , .	2.8	1
20	Genetic Variants of CLEC4E and BIRC3 in Damage-Associated Molecular Patterns-Related Pathway Genes Predict Non-Small Cell Lung Cancer Survival. Frontiers in Oncology, 2021, 11, 717109.	2.8	6
21	Predictive value of a novel Asian lung cancer screening nomogram based on artificial intelligence and epidemiological characteristics. Thoracic Cancer, 2021, 12, 3130-3140.	1.9	4
22	Genetic variants of , and in the natural killer cell-related pathway are associated with non-small cell lung cancer survival. American Journal of Cancer Research, 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. American Journal of Cancer Research, 2021, 11, 2343-2350.	1.4	2
24	Genetic variants of and in myeloid cell-related pathway genes independently predict cutaneous melanoma-specific survival. American Journal of Cancer Research, 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. Cell Discovery, 2021, 7, 121.	6.7	5
26	Novel genetic variants in KIF16B and NEDD4L in the endosomeâ€related genes are associated with nonsmall cell lung cancer survival. International Journal of Cancer, 2020, 147, 392-403.	5.1	6
27	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. Gut, 2020, 69, 641-651.	12.1	36
28	Common genetic variation and risk of osteosarcoma in a multi-ethnic pediatric and adolescent population. Bone, 2020, 130, 115070.	2.9	22
29	Novel genetic variants in HDAC2 and PPARGC1A of the CREBâ€binding protein pathway predict survival of nonâ€smallâ€cell lung cancer. Molecular Carcinogenesis, 2020, 59, 104-115.	2.7	13
30	Robotic sleeve lobectomy for centrally located nonâ€"small cell lung cancer: A propensity scoreâ€"weighted comparison with thoracoscopic and open surgery. Journal of Thoracic and Cardiovascular Surgery, 2020, 160, 838-846.e2.	0.8	45
31	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. Lancet Oncology, The, 2020, 21, 1378-1386.	10.7	123
32	Evaluated Lymph NodeÂNumber in Population-Based Analyses: Significant Prognostic Factor, but Assess With Caution. Journal of Thoracic Oncology, 2020, 15, e77-e78.	1.1	0
33	Genetic variation associated with childhood and adult stature and risk of MYCNâ€amplified neuroblastoma. Cancer Medicine, 2020, 9, 8216-8225.	2.8	3
34	Genetic variants in TKT and DERA in the nicotinamide adenine dinucleotide phosphate pathway predict melanoma survival. European Journal of Cancer, 2020, 136, 84-94.	2.8	3
35	Novel genetic variants in genes of the Fc gamma receptor-mediated phagocytosis pathway predict non-small cell lung cancer survival. Translational Lung Cancer Research, 2020, 9, 575-586.	2.8	6
36	Associations between genetic variants of KIF5B , FMN1 , and MGAT3 in the cadherin pathway and pancreatic cancer risk. Cancer Medicine, 2020, 9, 9620-9631.	2.8	1

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37	Functional genetic variants of CTNNBIP1 predict platinum treatment response of Chinese epithelial ovarian cancer patients. Journal of Cancer, 2020, 11, 6850-6860.	2.5	6
38	Genetic variants of the peroxisome proliferatorâ€activated receptor (PPAR) signaling pathway genes and risk of pancreatic cancer. Molecular Carcinogenesis, 2020, 59, 930-939.	2.7	11
39	Novel Variants of ELP2 and PIAS1 in the Interferon Gamma Signaling Pathway Are Associated with Nonâ€"Small Cell Lung Cancer Survival. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1679-1688.	2.5	2
40	The Rare Variant rs35356162 in UHRF1BP1 Increases Bladder Cancer Risk in Han Chinese Population. Frontiers in Oncology, 2020, 10, 134.	2.8	16
41	Genetic variants in the human leukocyte antigen region and survival of Chinese patients with non-small cell lung carcinoma. Carcinogenesis, 2020, 41, 1203-1212.	2.8	3
42	Potentially functional genetic variants in PLIN2, SULT2A1 and UGT1A9 genes of the ketone pathway and survival of nonsmall cell lung cancer. International Journal of Cancer, 2020, 147, 1559-1570.	5.1	8
43	Relevance and prognostic ability of Twist, Slug and tumor spread through air spaces in lung adenocarcinoma. Cancer Medicine, 2020, 9, 1986-1998.	2.8	13
44	Novel Genetic Variants of ALG6 and GALNTL4 of the Glycosylation Pathway Predict Cutaneous Melanoma-Specific Survival. Cancers, 2020, 12, 288.	3.7	7
45	Genetic variants in the folate metabolic pathway genes predict cutaneous melanomaâ€specific survival. British Journal of Dermatology, 2020, 183, 719-728.	1.5	4
46	Genetic variants in PDSS1 and SLC16A6 of the ketone body metabolic pathway predict cutaneous melanomaâ€specific survival. Molecular Carcinogenesis, 2020, 59, 640-650.	2.7	9
47	APOB Genotypes and CDH13 Haplotypes in the Cholesterol-Related Pathway Genes Predict Nonâ€"Small Cell Lung Cancer Survival. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1204-1213.	2.5	7
48	A Genome-Wide Association Study Identifies Two Novel Susceptible Regions for Squamous Cell Carcinoma of the Head and Neck. Cancer Research, 2020, 80, 2451-2460.	0.9	33
49	Novel genetic variants of and involved in immunoregulatory interactions are associated with non-small cell lung cancer survival. American Journal of Cancer Research, 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. American Journal of Cancer Research, 2020, 10, 2160-2173.	1.4	2
51	Associations of novel variants in , and of the ATM pathway genes with pancreatic cancer risk. American Journal of Cancer Research, 2020, 10, 2128-2144.	1.4	2
52	Novel genetic variants of and related lymphangiogenesis signaling pathway predict non-small cell lung cancer survival. American Journal of Cancer Research, 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. American Journal of Cancer Research, 2020, 10, 2582-2595.	1.4	6
54	Novel genetic variants of and of the endosome-related pathway predict cutaneous melanoma-specific survival. American Journal of Cancer Research, 2020, 10, 3382-3394.	1.4	0

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55	Functional variant of MTOR rs2536 and survival of Chinese gastric cancer patients. International Journal of Cancer, 2019, 144, 251-262.	5.1	5
56	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
57	Shortâ€term outcomes of typical versus atypical lung segmentectomy by minimally invasive surgeries. Thoracic Cancer, 2019, 10, 1812-1818.	1.9	11
58	Robotic Bronchial Sleeve Lobectomy for Central Lung Tumors: Technique andÂOutcome. Annals of Thoracic Surgery, 2019, 108, 211-218.	1.3	32
59	A genetic variant within MDM4 3â€²UTR miRNA binding site is associated with HPV16â€positive tumors and survival of oropharyngeal cancer. Molecular Carcinogenesis, 2019, 58, 2276-2285.	2.7	5
60	Genetic variants in glutamine metabolic pathway genes predict cutaneous melanomaâ€specific survival. Molecular Carcinogenesis, 2019, 58, 2091-2103.	2.7	5
61	Impact of visceral pleural invasion on the association of extent of lymphadenectomy and survival in stage I nonâ€small cell lung cancer. Cancer Medicine, 2019, 8, 669-678.	2.8	24
62	Role of Immune Response, Inflammation, and Tumor Immune Responseâ€"Related Cytokines/Chemokines in Melanoma Progression. Journal of Investigative Dermatology, 2019, 139, 2352-2358.e3.	0.7	23
63	Functional genetic variants of RUVBL1 predict overall survival of Chinese patients with epithelial ovarian cancer. Carcinogenesis, 2019, 40, 1209-1219.	2.8	5
64	Three novel genetic variants in NRF2 signaling pathway genes are associated with pancreatic cancer risk. Cancer Science, 2019, 110, 2022-2032.	3.9	14
65	Genetic variants in the liver kinase B1â€AMPâ€activated protein kinase pathway genes and pancreatic cancer risk. Molecular Carcinogenesis, 2019, 58, 1338-1348.	2.7	14
66	Potentially functional genetic variants in the TNF/TNFR signaling pathway genes predict survival of patients with nonâ€small cell lung cancer in the PLCO cancer screening trial. Molecular Carcinogenesis, 2019, 58, 1094-1104.	2.7	9
67	Genetic variants of genes in the NER pathway associated with risk of breast cancer: A largeâ€scale analysis of 14 published GWAS datasets in the DRIVE study. International Journal of Cancer, 2019, 145, 1270-1279.	5.1	13
68	Predictive value of a prognostic model based on pathologic features in lung invasive adenocarcinoma. Lung Cancer, 2019, 131, 14-22.	2.0	18
69	Genome-wide association studies identify susceptibility loci for epithelial ovarian cancer in east Asian women. Gynecologic Oncology, 2019, 153, 343-355.	1.4	28
70	Potential functional variants in SMC2 and TP53 in the AURORA pathway genes and risk of pancreatic cancer. Carcinogenesis, 2019, 40, 521-528.	2.8	17
71	Lymphocyte telomere length predicts clinical outcomes of HPV-positive oropharyngeal cancer patients after definitive radiotherapy. Carcinogenesis, 2019, 40, 735-741.	2.8	5
72	Genetic variants in ELOVL2 and HSD17B12 predict melanomaâ€specific survival. International Journal of Cancer, 2019, 145, 2619-2628.	5.1	11

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73	Genetic variants in the calcium signaling pathway genes are associated with cutaneous melanoma-specific survival. Carcinogenesis, 2019, 40, 279-288.	2.8	6
74	Genetic variants in RUNX3, AMD1 and MSRA in the methionine metabolic pathway and survival in nonsmall cell lung cancer patients. International Journal of Cancer, 2019, 145, 621-631.	5.1	21
75	Robotic circumferential tracheal resection and reconstruction via a completely portal approach. Thoracic Cancer, 2019, 10, 378-380.	1.9	4
76	Potentially functional genetic variants in the complementâ€related immunity geneâ€set are associated with nonâ€small cell lung cancer survival. International Journal of Cancer, 2019, 144, 1867-1876.	5.1	14
77	TGFÎ²1 Genetic Variants Predict Clinical Outcomes of HPV-Positive Oropharyngeal Cancer Patients after Definitive Radiotherapy. Clinical Cancer Research, 2018, 24, 2225-2233.	7.0	20
78	A TGFâ€Î²1 genetic variant at the miRNA187 binding site significantly modifies risk of HPV16â€associated oropharyngeal cancer. International Journal of Cancer, 2018, 143, 1327-1334.	5.1	7
79	Genetic variants in RORA and DNMT1 associated with cutaneous melanoma survival. International Journal of Cancer, 2018, 142, 2303-2312.	5.1	13
80	Potentially Functional Variants of ATG16L2 Predict Radiation Pneumonitis and Outcomes in Patients with Nonâ€"Small Cell Lung Cancer after Definitive Radiotherapy. Journal of Thoracic Oncology, 2018, 13, 660-675.	1.1	29
81	Inverse Relationship between Vitiligo-Related Genes and Skin Cancer Risk. Journal of Investigative Dermatology, 2018, 138, 2072-2075.	0.7	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. Molecular Carcinogenesis, 2018, 57, 784-793.	2.7	5
83	Novel genetic variants in the P38MAPK pathway gene ZAK and susceptibility to lung cancer. Molecular Carcinogenesis, 2018, 57, 216-224.	2.7	9
84	Mouse double minute 4 variants modify susceptibility to risk of recurrence in patients with squamous cell carcinoma of the oropharynx. Molecular Carcinogenesis, 2018, 57, 361-369.	2.7	6
85	Genetic variants in the plateletâ€derived growth factor subunit B gene associated with pancreatic cancer risk. International Journal of Cancer, 2018, 142, 1322-1331.	5.1	20
86	An ERCC4 regulatory variant predicts gradeâ€3 or â€4 toxicities in patients with advanced nonâ€small cell lung cancer treated by platinumâ€based therapy. International Journal of Cancer, 2018, 142, 1218-1229.	5.1	7
87	Genetic variants in the metzincin metallopeptidase family genes predict melanoma survival. Molecular Carcinogenesis, 2018, 57, 22-31.	2.7	5
88	Predictors of nodal metastasis and prognostic significance of lymph node ratio and total lymph node count in tracheobronchial adenoid cystic carcinoma. Cancer Management and Research, 2018, Volume 10, 5919-5925.	1.9	13
89	Potential clinical application of lncRNAs in non-small cell lung cancer. OncoTargets and Therapy, 2018, Volume 11, 8045-8052.	2.0	79
90	Genetic determinants of childhood and adult height associated with osteosarcoma risk. Cancer, 2018, 124, 3742-3752.	4.1	20

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91	Identification of an eight-gene prognostic signature for lung adenocarcinoma. Cancer Management and Research, 2018, Volume 10, 3383-3392.	1.9	63
92	Identification of SPP1 as a promising biomarker to predict clinical outcome of lung adenocarcinoma individuals. Gene, 2018, 679, 398-404.	2.2	51
93	DNA repair capacity correlates with standardized uptake values from 18 F-fluorodeoxyglucose positron emission tomography/CT in patients with advanced nonâ€“small-cell lung cancer. Chronic Diseases and Translational Medicine, 2018, 4, 109-116.	1.2	1
94	Genetic variants in nucleotide excision repair pathway predict survival of esophageal squamous cell cancer patients receiving platinumâ€based chemotherapy. Molecular Carcinogenesis, 2018, 57, 1553-1565.	2.7	9
95	Variants in Notch signalling pathway genes, <scp>PSEN</scp>1 and <scp>MAML</scp>2, predict overall survival in Chinese patients with epithelial ovarian cancer. Journal of Cellular and Molecular Medicine, 2018, 22, 4975-4984.	3.6	14
96	Genetic variant of*IRAK2*in the tollâ€like receptor signaling pathway and survival of nonâ€small cell lung cancer. International Journal of Cancer, 2018, 143, 2400-2408.	5.1	14
97	Single-nucleotide polymorphisms of stemness genes predicted to regulate RNA splicing, microRNA and oncogenic signaling are associated with prostate cancer survival. Carcinogenesis, 2018, 39, 879-888.	2.8	9
98	Genetic correction improves prediction efficiency of serum tumor biomarkers on digestive cancer risk in the elderly Chinese cohort study. Oncotarget, 2018, 9, 7389-7397.	1.8	7
99	Apoptotic capacity and risk of squamous cell carcinoma of the head and neck. European Journal of Cancer, 2017, 72, 166-176.	2.8	19
100	Modifying effect of mouse double minute-2 promoter variants on risk of recurrence for patients with squamous cell carcinoma of oropharynx. Scientific Reports, 2017, 7, 39765.	3.3	4
101	Pathwayâ€analysis of published genomeâ€wide association studies of lung cancer: A potential role for the CYP4F3 locus. Molecular Carcinogenesis, 2017, 56, 1663-1672.	2.7	13
102	BRCA1 and BRCA2 mutations in ovarian cancer patients from China: ethnicâ€related mutations in BRCA1 associated with an increased risk of ovarian cancer. International Journal of Cancer, 2017, 140, 2051-2059.	5.1	45
103	Reduced mRNA expression of nucleotide excision repair genes in lymphocytes and risk of squamous cell carcinoma of the head and neck. Carcinogenesis, 2017, 38, 504-510.	2.8	6
104	Genetic Variants in WNT2B and BTRC Predict Melanoma Survival. Journal of Investigative Dermatology, 2017, 137, 1749-1756.	0.7	5
105	Associations between RNA splicing regulatory variants of stemnessâ€related genes and racial disparities in susceptibility to prostate cancer. International Journal of Cancer, 2017, 141, 731-743.	5.1	20
106	Genetic variants of PTPN2 are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. Scientific Reports, 2017, 7, 825.	3.3	10
107	Association between Body Mass Index, C-Reactive Protein Levels, and Melanoma Patient Outcomes. Journal of Investigative Dermatology, 2017, 137, 1792-1795.	0.7	40
108	Genetic variants in the integrin signaling pathway genes predict cutaneous melanoma survival. International Journal of Cancer, 2017, 140, 1270-1279.	5.1	4

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109	Genetic variants in the genes encoding rho GTPases and related regulators predict cutaneous melanomaâ€specific survival. International Journal of Cancer, 2017, 141, 721-730.	5.1	8
110	Carbon Quantum Dots Decorated C₃N₄/TiO₂Heterostructure Nanorod Arrays for Enhanced Photoelectrochemical Performance. Journal of the Electrochemical Society, 2017, 164, H515-H520.	2.9	22
111	A PGC1Î² genetic variant associated with nevus count and melanoma mortality. International Journal of Cancer, 2017, 141, 1066-1067.	5.1	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. Carcinogenesis, 2017, 38, 432-438.	2.8	6
113	Towards precision prevention: Technologies for identifying healthy individuals with high risk of disease. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2017, 800-802, 14-28.	1.0	20
114	Functional variants in DCAF4 associated with lung cancer risk in European populations. Carcinogenesis, 2017, 38, 541-551.	2.8	16
115	Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. Scientific Reports, 2017, 7, 44634.	3.3	10
116	Genetic variants in ERCC1 and XPC predict survival outcome of non-small cell lung cancer patients treated with platinum-based therapy. Scientific Reports, 2017, 7, 10702.	3.3	12
117	Diagnostic accuracy of serum antibodies to human papillomavirus type 16 early antigens in the detection of human papillomavirusâ€"related oropharyngeal cancer. Cancer, 2017, 123, 4886-4894.	4.1	16
118	Assessing the spectrum of germline variation in Fanconi anemia genes among patients with head and neck carcinoma before age 50. Cancer, 2017, 123, 3943-3954.	4.1	37
119	Functional genetic variants of XRCC4 and ERCC1 predict survival of gastric cancer patients treated with chemotherapy by regulating the gene expression. Molecular Carcinogenesis, 2017, 56, 2706-2717.	2.7	4
120	Genetic variants in microRNAâ€binding sites of DNA repair genes as predictors of recurrence in patients with squamous cell carcinoma of the oropharynx. International Journal of Cancer, 2017, 141, 1355-1364.	5.1	9
121	<scp>G</scp>enetic variants of JNK and p38Î± pathways and risk of nonâ€small cell lung cancer in an <scp>E</scp>astern <scp>C</scp>hinese population. International Journal of Cancer, 2017, 140, 807-817.	5.1	8
122	E2F transcription factor 2 variants as predictive biomarkers for recurrence risk in patients with squamous cell carcinoma of the oropharynx. Molecular Carcinogenesis, 2017, 56, 1335-1343.	2.7	13
123	Susceptibility loci of CNOT6 in the general mRNA degradation pathway and lung cancer riskâ€"A reâ€analysis of eight GWASs. Molecular Carcinogenesis, 2017, 56, 1227-1238.	2.7	10
124	Melanoma Expression Genes Identified through Genome-Wide Association Study ofÂBreslow Tumor Thickness. Journal of Investigative Dermatology, 2017, 137, 253-257.	0.7	2
125	A functional variant at the miRNA binding site in E2F1 gene is associated with risk and tumor HPV16 status of oropharynx squamous cell carcinoma. Molecular Carcinogenesis, 2017, 56, 1100-1106.	2.7	12
126	Functional variants in the low-density lipoprotein receptor gene are associated with clear cell renal cell carcinoma susceptibility. Carcinogenesis, 2017, 38, 1241-1248.	2.8	5

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127	Genetic variants of PDGF signaling pathway genes predict cutaneous melanoma survival. Oncotarget, 2017, 8, 74595-74606.	1.8	3
128	THE AUTHORS REPLY. American Journal of Epidemiology, 2017, 186, 625-626.	3.4	0
129	Two-stage genome-wide association study identifies a novel susceptibility locus associated with melanoma. Oncotarget, 2017, 8, 17586-17592.	1.8	61
130	Polymorphisms in nucleotide excision repair genes and risk of primary prostate cancer in Chinese Han populations. Oncotarget, 2017, 8, 24362-24371.	1.8	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.. Journal of Clinical Oncology, 2017, 35, 1583-1583.	1.6	1
133	A functional variant at miRNA-122 binding site in IL-1Î± 3â€² UTR predicts risk of recurrence in patients with oropharyngeal cancer. Oncotarget, 2016, 7, 34472-34479.	1.8	8
134	Genetic variants in the vitamin <scp>D</scp> pathway genes <scp>VDBP</scp>Âand <scp>RXRA</scp> modulate cutaneous melanoma diseaseâ€specific survival. Pigment Cell and Melanoma Research, 2016, 29, 176-185.	3.3	19
135	Genetic variants in <scp>ABCG</scp>1 are associated with survival of nonsmallâ€cell lung cancer patients. International Journal of Cancer, 2016, 138, 2592-2601.	5.1	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. Chest, 2016, 149, A275.	0.8	0
137	Genetic variants in the PIWIâ€piRNA pathway gene DCP1A predict melanoma diseaseâ€specific survival. International Journal of Cancer, 2016, 139, 2730-2737.	5.1	21
138	Effect of human papillomavirus seropositivity andE2F2promoter variants on risk of squamous cell carcinomas of oropharynx and oral cavity. Carcinogenesis, 2016, 37, 1070-1078.	2.8	5
139	A comprehensive genomeâ€wide analysis of melanoma Breslow thickness identifies interaction between CDC42 and SCIN genetic variants. International Journal of Cancer, 2016, 139, 2012-2020.	5.1	8
140	Smokeless Tobacco Use and the Risk of Head and Neck Cancer: Pooled Analysis of US Studies in the INHANCE Consortium. American Journal of Epidemiology, 2016, 184, 703-716.	3.4	78
141	The P38Î± rs3804451 Variant Predicts Chemotherapy Response and Survival of Patients with Nonâ€"Small Cell Lung Cancer Treated with Platinum-Based Chemotherapy. Translational Oncology, 2016, 9, 531-539.	3.7	3
142	A Novel Genetic Variant in Long Non-coding RNA Gene NEXN-AS1 is Associated with Risk of Lung Cancer. Scientific Reports, 2016, 6, 34234.	3.3	48
143	Site disparities in apoptotic variants as predictors of risk for second primary malignancy in patients with squamous cell carcinoma of the head and neck. BMC Cancer, 2016, 16, 70.	2.6	2
144	4-Nitroquinoline-1-oxide-induced mutagen sensitivity and risk of cutaneous melanoma. Melanoma Research, 2016, 26, 181-187.	1.2	8

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145	Genetic variant in DNA repair geneGTF2H4is associated with lung cancer risk: a large-scale analysis of six published GWAS datasets in the TRICL consortium. Carcinogenesis, 2016, 37, 888-896.	2.8	15
146	Genetic variants of DNA repair genes predict the survival of patients with esophageal squamous cell cancer receiving platinum-based adjuvant chemotherapy. Journal of Translational Medicine, 2016, 14, 154.	4.4	8
147	Single Nucleotide Polymorphisms in CBLB, aÂRegulator of T-Cell Response, Predict Radiation Pneumonitis and Outcomes After Definitive Radiotherapy for Nonâ€"Small-Cell Lung Cancer. Clinical Lung Cancer, 2016, 17, 253-262.e5.	2.6	16
148	Association of Vitamin D Levels With Outcome in Patients With Melanoma After Adjustment For C-Reactive Protein. Journal of Clinical Oncology, 2016, 34, 1741-1747.	1.6	64
149	Polymorphisms in the <scp>AKT</scp>1 and <scp>AKT</scp>2 genes and oesophageal squamous cell carcinoma risk in an Eastern Chinese population. Journal of Cellular and Molecular Medicine, 2016, 20, 666-677.	3.6	31
150	Squamous cell carcinoma of the oral cavity often overexpresses p16 but is rarely driven by human papillomavirus. Oral Oncology, 2016, 56, 47-53.	1.5	88
151	Genetic variants in miR-196a2 and miR-499 are associated with susceptibility to esophageal squamous cell carcinoma in Chinese Han population. Tumor Biology, 2016, 37, 4777-4784.	1.8	39
152	Polymorphisms of the centrosomal gene (FGFR1OP) and lung cancer risk: a meta-analysis of 14 463 cases and 44 188 controls. Carcinogenesis, 2016, 37, 280-289.	2.8	7
153	Reduced DNA double-strand break repair capacity and risk of squamous cell carcinoma of the head and neckâ€"A case-control study. DNA Repair, 2016, 40, 18-26.	2.8	14
154	A variant at a potentially functional microRNA-binding site in BRIP1 was associated with risk of squamous cell carcinoma of the head and neck. Tumor Biology, 2016, 37, 8057-8066.	1.8	12
155	Genetic variations in the homologous recombination repair pathway genes modify risk of glioma. Journal of Neuro-Oncology, 2016, 126, 11-17.	2.9	10
156	Low frequency of cigarette smoking and the risk of head and neck cancer in the INHANCE consortium pooled analysis. International Journal of Epidemiology, 2016, 45, 835-845.	1.9	40
157	Associations between smoking behavior-related alleles and the risk of melanoma. Oncotarget, 2016, 7, 47366-47375.	1.8	15
158	Genetic variants of genes in the Notch signaling pathway predict overall survival of non-small cell lung cancer patients in the PLCO study. Oncotarget, 2016, 7, 61716-61727.	1.8	17
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160	Pre-miRNA variants as predictors of clinical outcome in patients with squamous cell carcinomas of the nonoropharynx. Oncotarget, 2016, 7, 26444-26453.	1.8	11
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