

Fengju Song

List of Publications by Year in descending order

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Version: 2024-02-01

65
papers

2,755
citations

201674

27
h-index

197818

49
g-index

77
all docs

77
docs citations

77
times ranked

5881
citing authors

#	ARTICLE	IF	CITATIONS
1	Effect of estradiol as a continuous variable on breast cancer survival by menopausal status: a cohort study in China. <i>Breast Cancer Research and Treatment</i> , 2022, 194, 103-111.	2.5	1
2	Consumption of flavonoids and risk of hormone-related cancers: a systematic review and meta-analysis of observational studies. <i>Nutrition Journal</i> , 2022, 21, 27.	3.4	14
3	Detection rate is not higher for women with BBD history in breast cancer screening. <i>Journal of Public Health</i> , 2021, 43, 333-340.	1.8	1
4	Utility of Preoperative Inflammatory Markers to Distinguish Epithelial Ovarian Cancer from Benign Ovarian Masses. <i>Journal of Cancer</i> , 2021, 12, 2687-2693.	2.5	8
5	SNPs within microRNA binding sites and the prognosis of breast cancer. <i>Aging</i> , 2021, 13, 7465-7480.	3.1	3
6	Functional Interrogation of Enhancer Connectome Prioritizes Candidate Target Genes at Ovarian Cancer Susceptibility Loci. <i>Frontiers in Genetics</i> , 2021, 12, 646179.	2.3	3
7	Interrogation of gender disparity uncovers androgen receptor as the transcriptional activator for oncogenic miR-125b in gastric cancer. <i>Cell Death and Disease</i> , 2021, 12, 441.	6.3	15
8	Community-based lung cancer screening by low-dose computed tomography in China: First round results and a meta-analysis. <i>European Journal of Radiology</i> , 2021, 144, 109988.	2.6	6
9	Improved diagnosis of thyroid cancer aided with deep learning applied to sonographic text reports: a retrospective, multi-cohort, diagnostic study. <i>Cancer Biology and Medicine</i> , 2021, 19, 733-741.	3.0	4
10	Development and evaluation of the screening performance of a low-cost high-risk screening strategy for breast cancer. <i>Cancer Biology and Medicine</i> , 2021, , 1-13.	3.0	1
11	Identification of an Autophagy-Related Risk Signature Correlates With Immunophenotype and Predicts Immune Checkpoint Blockade Efficacy of Neuroblastoma. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 731380.	3.7	3
12	Parity and risk of developing breast cancer according to tumor subtype: A systematic review and meta-analysis. <i>Cancer Epidemiology</i> , 2021, 75, 102050.	1.9	21
13	Association between low density lipoprotein cholesterol and all-cause mortality: results from the NHANES 1999-2014. <i>Scientific Reports</i> , 2021, 11, 22111.	3.3	15
14	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.	6.7	5
15	Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020, 112, 1223-1232.	2.9	15
16	Performance of ultrasonography screening for breast cancer: a systematic review and meta-analysis. <i>BMC Cancer</i> , 2020, 20, 499.	2.6	38
17	Tumor markers CA15-3, CA125, CEA and breast cancer survival by molecular subtype: a cohort study. <i>Breast Cancer</i> , 2020, 27, 621-630.	2.9	44
18	Identification of a Sixteen-gene Prognostic Biomarker for Lung Adenocarcinoma Using a Machine Learning Method. <i>Journal of Cancer</i> , 2020, 11, 1288-1298.	2.5	37

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19	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020, 52, 494-504.	21.4	138
20	Diagnostic classification of cancers using extreme gradient boosting algorithm and multi-omics data. <i>Computers in Biology and Medicine</i> , 2020, 121, 103761.	7.0	102
21	Analysis of immune-related signatures of lung adenocarcinoma identified two distinct subtypes: implications for immune checkpoint blockade therapy. <i>Aging</i> , 2020, 12, 3312-3339.	3.1	103
22	Ages at menarche and menopause, and mortality among postmenopausal women. <i>Maturitas</i> , 2019, 130, 50-56.	2.4	22
23	Age at menarche and epithelial ovarian cancer risk: A meta-analysis and Mendelian randomization study. <i>Cancer Medicine</i> , 2019, 8, 4012-4022.	2.8	15
24	The new identified biomarkers determine sensitivity to immune check-point blockade therapies in melanoma. <i>Oncolmmunology</i> , 2019, 8, 1608132.	4.6	37
25	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
26	Comparison of breast cancer risk factors among molecular subtypes: A case-only study. <i>Cancer Medicine</i> , 2019, 8, 1882-1892.	2.8	29
27	<p>A Nomogram To Predict The Overall Survival Of Breast Cancer Patients And Guide The Postoperative Adjuvant Chemotherapy In China</p>. <i>Cancer Management and Research</i> , 2019, Volume 11, 10029-10039.	1.9	8
28	Personal history of keratinocyte carcinoma is associated with reduced risk of death from invasive melanoma in men. <i>Journal of the American Academy of Dermatology</i> , 2018, 78, 957-963.	1.2	5
29	Assessment of performance of the Gail model for predicting breast cancer risk: a systematic review and meta-analysis with trial sequential analysis. <i>Breast Cancer Research</i> , 2018, 20, 18.	5.0	66
30	Improved Performance of Adjunctive Ultrasonography After Mammography Screening for Breast Cancer Among Chinese Females. <i>Clinical Breast Cancer</i> , 2018, 18, e353-e361.	2.4	13
31	Histone deacetylase 6 in cancer. <i>Journal of Hematology and Oncology</i> , 2018, 11, 111.	17.0	214
32	SNP rs2071095 in LincRNA H19 is associated with breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2018, 171, 161-171.	2.5	34
33	Association study of genetic variation in <scp>DNA</scp> repair pathway genes and risk of basal cell carcinoma. <i>International Journal of Cancer</i> , 2017, 141, 952-957.	5.1	14
34	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
35	Preliminary effectiveness of breast cancer screening among 1.22 million Chinese females and different cancer patterns between urban and rural women. <i>Scientific Reports</i> , 2016, 6, 39459.	3.3	29
36	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , 2016, 26, ddw414.	2.9	50

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37	MiR-502/SET8 regulatory circuit in pathobiology of breast cancer. <i>Cancer Letters</i> , 2016, 376, 259-267.	7.2	36
38	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , 2016, 139, 2655-2670.	5.1	43
39	A survey of overall life satisfaction and its association with breast diseases in Chinese women. <i>Cancer Medicine</i> , 2016, 5, 111-119.	2.8	16
40	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	9.4	157
41	Roles of low-density lipoprotein receptor-related protein 1 in tumors. <i>Chinese Journal of Cancer</i> , 2016, 35, 6.	4.9	43
42	Associations between smoking behavior-related alleles and the risk of melanoma. <i>Oncotarget</i> , 2016, 7, 47366-47375.	1.8	15
43	Urban-rural disparity of overweight/obesity distribution and its potential trend with breast cancer among Chinese women. <i>Oncotarget</i> , 2016, 7, 56608-56618.	1.8	18
44	Prognostic roles for fibroblast growth factor receptor family members in malignant peripheral nerve sheath tumor. <i>Oncotarget</i> , 2016, 7, 22234-22244.	1.8	14
45	A functional single nucleotide polymorphism of SET8 is prognostic for breast cancer. <i>Oncotarget</i> , 2016, 7, 34277-34287.	1.8	18
46	Comparative genomic analysis reveals bilateral breast cancers are genetically independent. <i>Oncotarget</i> , 2015, 6, 31820-31829.	1.8	20
47	Pleiotropic and Sex-Specific Effects of Cancer GWAS SNPs on Melanoma Risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. <i>PLoS ONE</i> , 2015, 10, e0120491.	2.5	19
48	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , 2015, 47, 987-995.	21.4	218
49	Clonality: A New Marker for Gastric Cancer Survival. <i>Journal of Genetics and Genomics</i> , 2015, 42, 517-519.	3.9	1
50	Ambient particulate matter and lung cancer incidence and mortality: a meta-analysis of prospective studies. <i>European Journal of Public Health</i> , 2015, 25, 324-329.	0.3	74
51	miR-485-5p Binding Site SNP rs8752 in HPGD Gene Is Associated with Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e102093.	2.5	26
52	Replication of Associations between GWAS SNPs and Melanoma Risk in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. <i>Journal of Investigative Dermatology</i> , 2014, 134, 2049-2052.	0.7	21
53	Identification of a melanoma susceptibility locus and somatic mutation in <i>TET2</i> . <i>Carcinogenesis</i> , 2014, 35, 2097-2101.	2.8	41
54	Integrated MicroRNA Network Analyses Identify a Poor-Prognosis Subtype of Gastric Cancer Characterized by the miR-200 Family. <i>Clinical Cancer Research</i> , 2014, 20, 878-889.	7.0	97

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55	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
56	Genome-wide association studies identify several new loci associated with pigmentation traits and skin cancer risk in European Americans. <i>Human Molecular Genetics</i> , 2013, 22, 2948-2959.	2.9	104
57	Risk of a Second Primary Cancer after Non-melanoma Skin Cancer in White Men and Women: A Prospective Cohort Study. <i>PLoS Medicine</i> , 2013, 10, e1001433.	8.4	59
58	Smoking and risk of skin cancer: a prospective analysis and a meta-analysis. <i>International Journal of Epidemiology</i> , 2012, 41, 1694-1705.	1.9	93
59	Exonuclease 1 (EXO1) gene variation and melanoma risk. <i>DNA Repair</i> , 2012, 11, 304-309.	2.8	22
60	Functional SNP in the microRNA-367 binding site in the 3'UTR of the calcium channel ryanodine receptor gene 3 (<i>RYR3</i>) affects breast cancer risk and calcification. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 13653-13658.	7.1	144
61	Genetic variants at the miR-124 binding site on the cytoskeleton-organizing <i>IQGAP1</i> gene confer differential predisposition to breast cancer. <i>International Journal of Oncology</i> , 2011, 38, 1153-61.	3.3	24
62	Definition of a Functional Single Nucleotide Polymorphism in the Cell Migration Inhibitory Gene <i>MIIP</i> That Affects the Risk of Breast Cancer. <i>Cancer Research</i> , 2010, 70, 1024-1032.	0.9	16
63	An miR-502 Binding Site Single-Nucleotide Polymorphism in the 3'-Untranslated Region of the <i>SET8</i> Gene Is Associated with Early Age of Breast Cancer Onset. <i>Clinical Cancer Research</i> , 2009, 15, 6292-6300.	7.0	106
64	A cancer incidence survey in Tianjin: the third largest city in China between 1981 and 2000. <i>Cancer Causes and Control</i> , 2008, 19, 443-450.	1.8	40
65	Analysis of the thyroid carcinoma incidence in Tianjin over a recent twenty-year period. <i>Chinese Journal of Clinical Oncology</i> , 2005, 2, 815-819.	0.0	0