

# Hongji Dai

## List of Publications by Year in descending order

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Version: 2024-02-01

52  
papers

1,289  
citations

393982

19  
h-index

395343

33  
g-index

62  
all docs

62  
docs citations

62  
times ranked

2728  
citing authors

#	ARTICLE	IF	CITATIONS
1	Probing the formation, structure and free energy relationships of M protein dimers of SARS-CoV-2. Computational and Structural Biotechnology Journal, 2022, 20, 573-582.	1.9	9
2	Editorial: The Role of Sex Dimorphism in Disease Susceptibility and Immune Response. Frontiers in Nutrition, 2022, 9, 849563.	1.6	1
3	Poor oral health and risks of total and site-specific cancers in China: A prospective cohort study of 0.5 million adults. EClinicalMedicine, 2022, 45, 101330.	3.2	7
4	Associations of Nutritional, Environmental, and Metabolic Biomarkers with Diabetes-Related Mortality in U.S. Adults: The Third National Health and Nutrition Examination Surveys between 1988-1994 and 2016. Nutrients, 2022, 14, 2629.	1.7	1
5	Distinct transcriptomic landscapes of cutaneous basal cell carcinomas and squamous cell carcinomas. Genes and Diseases, 2021, 8, 181-192.	1.5	14
6	Detection rate is not higher for women with BBD history in breast cancer screening. Journal of Public Health, 2021, 43, 333-340.	1.0	1
7	Functional Interrogation of Enhancer Connectome Prioritizes Candidate Target Genes at Ovarian Cancer Susceptibility Loci. Frontiers in Genetics, 2021, 12, 646179.	1.1	3
8	Development and evaluation of the screening performance of a low-cost high-risk screening strategy for breast cancer. Cancer Biology and Medicine, 2021, , 1-13.	1.4	1
9	Association Between Genetically Proxied Lipid-Lowering Drug Targets and Renal Cell Carcinoma: A Mendelian Randomization Study. Frontiers in Nutrition, 2021, 8, 755834.	1.6	6
10	Association between thyroid dysfunction and type 2 diabetes: a meta-analysis of prospective observational studies. BMC Medicine, 2021, 19, 257.	2.3	39
11	Identification of an m6A Regulators-Mediated Prognosis Signature For Survival Prediction and Its Relevance to Immune Infiltration in Melanoma. Frontiers in Cell and Developmental Biology, 2021, 9, 718912.	1.8	6
12	Genome-wide association and functional interrogation identified a variant at 3p26.1 modulating ovarian cancer survival among Chinese women. Cell Discovery, 2021, 7, 121.	3.1	5
13	Potentially functional variants in nucleotide excision repair pathway genes predict platinum treatment response of Chinese ovarian cancer patients. Carcinogenesis, 2020, 41, 1229-1237.	1.3	3
14	Functional genetic variants of <i>CTNNBIP1</i> predict platinum treatment response of Chinese epithelial ovarian cancer patients. Journal of Cancer, 2020, 11, 6850-6860.	1.2	6
15	Comparison of spatiotemporal characteristics of the COVID-19 and SARS outbreaks in mainland China. BMC Infectious Diseases, 2020, 20, 805.	1.3	28
16	Tumor markers CA15-3, CA125, CEA and breast cancer survival by molecular subtype: a cohort study. Breast Cancer, 2020, 27, 621-630.	1.3	44
17	Ages at menarche and menopause, and mortality among postmenopausal women. Maturitas, 2019, 130, 50-56.	1.0	22
18	Age at menarche and epithelial ovarian cancer risk: A meta-analysis and Mendelian randomization study. Cancer Medicine, 2019, 8, 4012-4022.	1.3	15

#	ARTICLE	IF	CITATIONS
19	Functional genetic variants of RUVBL1 predict overall survival of Chinese patients with epithelial ovarian cancer. <i>Carcinogenesis</i> , 2019, 40, 1209-1219.	1.3	5
20	<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.	0.9	8
21	Abstract 1570: Genome-wide association study identifies variants at 3p26.1 linked to survival in ovarian cancer patients among Chinese women. , 2019, , .		0
22	Abstract 3558: Interrogation of onco-microRNA 125b reveals androgen receptor as a key upstream transcriptional factor that is targetable in female gastric cancer. , 2019, , .		0
23	A mutational signature associated with alcohol consumption and prognostically significantly mutated driver genes in esophageal squamous cell carcinoma. <i>Annals of Oncology</i> , 2018, 29, 938-944.	0.6	76
24	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.	2.2	66
25	Improved Performance of Adjunctive Ultrasonography After Mammography Screening for Breast Cancer Among Chinese Females. <i>Clinical Breast Cancer</i> , 2018, 18, e353-e361.	1.1	13
26	Variants in Notch signalling pathway genes, <i><scp>PSEN</scp>1</i> and <i><scp>MAML</scp>2,</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
27	Abstract 3405: A mutational signature associated with alcohol consumption and prognostically mutated driver genes in esophageal squamous cell carcinoma. , 2018, , .		0
28	Association between genetic variation within vitamin D receptorâ€DNA binding sites and risk of basal cell carcinoma. <i>International Journal of Cancer</i> , 2017, 140, 2085-2091.	2.3	11
29	Associations between benign cutaneous nevi and risk of Type 2 diabetes mellitus in men and women: results from two prospective cohort studies. <i>Diabetic Medicine</i> , 2017, 34, 925-933.	1.2	2
30	Two-stage genome-wide association study identifies a novel susceptibility locus associated with melanoma. <i>Oncotarget</i> , 2017, 8, 17586-17592.	0.8	61
31	Predictors for local invasive recurrence of ductal carcinoma in situ of the breast. <i>European Journal of Cancer Prevention</i> , 2016, 25, 19-28.	0.6	34
32	Personal history of psoriasis and risk of incident cancer among women: a population-based cohort study. <i>British Journal of Dermatology</i> , 2016, 174, 1108-1111.	1.4	13
33	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.	1.6	29
34	A survey of overall life satisfaction and its association with breast diseases in Chinese women. <i>Cancer Medicine</i> , 2016, 5, 111-119.	1.3	16
35	Personal history of psoriasis and risk of nonmelanoma skin cancer (NMSC) among women in the United States: A population-based cohort study. <i>Journal of the American Academy of Dermatology</i> , 2016, 75, 731-735.	0.6	17
36	Genome-wide association study identifies novel susceptibility loci for cutaneous squamous cell carcinoma. <i>Nature Communications</i> , 2016, 7, 12048.	5.8	117

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37	Tissue-specific Co-expression of Long Non-coding and Coding RNAs Associated with Breast Cancer. Scientific Reports, 2016, 6, 32731.	1.6	35
38	Genome-wide association study identifies 14 novel risk alleles associated with basal cell carcinoma. Nature Communications, 2016, 7, 12510.	5.8	94
39	069 Genome-wide association study identifies 14 novel risk alleles associated with basal cell carcinoma. Journal of Investigative Dermatology, 2016, 136, S13.	0.3	0
40	Urban-rural disparity of overweight/obesity distribution and its potential trend with breast cancer among Chinese women. Oncotarget, 2016, 7, 56608-56618.	0.8	18
41	A functional single nucleotide polymorphism of SET8 is prognostic for breast cancer. Oncotarget, 2016, 7, 34277-34287.	0.8	18
42	Abstract 4045: Urban-rural disparity of overweight/obesity distribution and its potential association with breast cancer among Chinese females. , 2016, , .		0
43	Abstract 874: Personal history of psoriasis and risk of incident cancer among women: the Nursesâ€™ Health Study. , 2015, , .		0
44	Active and passive smoking with breast cancer risk for Chinese females: a systematic review and meta-analysis. Chinese Journal of Cancer, 2014, 33, 306-316.	4.9	37
45	Distribution of mammographic density and its influential factors among Chinese women. International Journal of Epidemiology, 2014, 43, 1240-1251.	0.9	66
46	A meta-analysis of the association between induced abortion and breast cancer risk among Chinese females. Cancer Causes and Control, 2014, 25, 227-236.	0.8	48
47	OP0008 Insights from the national breast cancer screening programme of 1 million Chinese women. European Journal of Cancer, 2014, 50, e3.	1.3	0
48	Association between Dietary Factors and Breast Cancer Risk among Chinese Females: Systematic Review and Meta-analysis. Asian Pacific Journal of Cancer Prevention, 2014, 15, 1291-1298.	0.5	37
49	Tea Consumption, Alcohol Drinking and Physical Activity Associations with Breast Cancer Risk among Chinese Females: a Systematic Review and Meta-analysis. Asian Pacific Journal of Cancer Prevention, 2013, 14, 7543-7550.	0.5	38
50	The role of polymorphisms in circadian pathway genes in breast tumorigenesis. Breast Cancer Research and Treatment, 2011, 127, 531-540.	1.1	64
51	Abstract A1: An miRâ€‘502 binding site singleâ€‘nucleotide polymorphism in the 3â€‘-untranslated region of the SET8 gene is associated with early age of breast cancer onset. , 2010, , .		1
52	An miR-502â€‘Binding Site Single-Nucleotide Polymorphism in the 3â€‘-Untranslated Region of the SET8 Gene Is Associated with Early Age of Breast Cancer Onset. Clinical Cancer Research, 2009, 15, 6292-6300.	3.2	106