

Jiali Han

List of Publications by Year in descending order

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Version: 2024-02-01

123
papers

6,217
citations

71102

41
h-index

79698

73
g-index

125
all docs

125
docs citations

125
times ranked

10735
citing authors

#	ARTICLE	IF	CITATIONS
1	COXâ€2 inhibitors show no preventive effect in the development of skin cancer. JDDG - Journal of the German Society of Dermatology, 2022, 20, 157-166.	0.8	3
2	Genome-wide association study identifies 48 common genetic variants associated with handedness. Nature Human Behaviour, 2021, 5, 59-70.	12.0	79
3	Looking for Sunshine: Genetic Predisposition to Sun Seeking in 265,000 Individuals of European Ancestry. Journal of Investigative Dermatology, 2021, 141, 779-786.	0.7	5
4	Genome-wide association study in almost 195,000 individuals identifies 50 previously unidentified genetic loci for eye color. Science Advances, 2021, 7, .	10.3	36
5	Circulating markers of cellular immune activation in prediagnostic blood sample and lung cancer risk in the Lung Cancer Cohort Consortium (LC3). International Journal of Cancer, 2020, 146, 2394-2405.	5.1	21
6	Cutaneous nevi and internal cancer risk: Results from two large prospective cohorts of US women. International Journal of Cancer, 2020, 147, 14-20.	5.1	2
7	Recreational and residential sun exposure and risk of endometriosis: a prospective cohort study. Human Reproduction, 2020, 36, 199-210.	0.9	2
8	Pancreatic safety of sodiumâ€glucose cotransporter 2 inhibitors in patients with type 2 diabetes mellitus: A systematic review and metaâ€analysis. Pharmacoepidemiology and Drug Safety, 2020, 29, 161-172.	1.9	21
9	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. Nature Genetics, 2020, 52, 494-504.	21.4	138
10	Genome-wide meta-analysis identifies eight new susceptibility loci for cutaneous squamous cell carcinoma. Nature Communications, 2020, 11, 820.	12.8	30
11	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
12	A Prospective Study of Leukocyte Telomere Length and Risk of Gestational Diabetes in a Multiracial Cohort. Epidemiology, 2019, 30, S10-S16.	2.7	3
13	Association Between Health Maintenance Practices and Skin Cancer Risk as a Possible Source of Detection Bias. JAMA Dermatology, 2019, 155, 353.	4.1	10
14	Circulating high sensitivity C reactive protein concentrations and risk of lung cancer: nested case-control study within Lung Cancer Cohort Consortium. BMJ: British Medical Journal, 2019, 364, k4981.	2.3	36
15	Voriconazole exposure and risk of cutaneous squamous cell carcinoma among lung or hematopoietic cell transplant patients: A systematic review and meta-analysis. Journal of the American Academy of Dermatology, 2019, 80, 500-507.e10.	1.2	30
16	Interaction of body mass index or waistâ€toâ€hip ratio and sun exposure associated with nonmelanoma skin cancer: A prospective study from the Womenâ€™s Health Initiative. Cancer, 2019, 125, 1133-1142.	4.1	6
17	Pioglitazone and bladder cancer risk: a systematic review and metaâ€analysis. Cancer Medicine, 2018, 7, 1070-1080.	2.8	91
18	Melanoma risk prediction using a multilocus genetic risk score in the Women's Health Initiative cohort. Journal of the American Academy of Dermatology, 2018, 79, 36-41.e10.	1.2	22

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19	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
20	Impaired functional vitamin B6 status is associated with increased risk of lung cancer. <i>International Journal of Cancer</i> , 2018, 142, 2425-2434.	5.1	12
21	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.	5.1	13
22	Use of antihypertensive drugs and risk of keratinocyte carcinoma: A meta-analysis of observational studies. <i>Pharmacoepidemiology and Drug Safety</i> , 2018, 27, 279-288.	1.9	30
23	Inverse Relationship between Vitiligo-Related Genes and Skin Cancer Risk. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2072-2075.	0.7	20
24	Circulating Folate, Vitamin B6, and Methionine in Relation to Lung Cancer Risk in the Lung Cancer Cohort Consortium (LC3). <i>Journal of the National Cancer Institute</i> , 2018, 110, 57-67.	6.3	40
25	Use of Antihypertensive Drugs and Risk of Malignant Melanoma: A Meta-analysis of Observational Studies. <i>Drug Safety</i> , 2018, 41, 161-169.	3.2	20
26	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
27	Personal history of non-melanoma skin cancer diagnosis and death from melanoma in women. <i>International Journal of Cancer</i> , 2018, 142, 1536-1541.	5.1	3
28	Prospective study of restless legs syndrome and total and cardiovascular mortality among women. <i>Neurology</i> , 2018, 90, e135-e141.	1.1	50
29	Genetic variants in the metzincin metallopeptidase family genes predict melanoma survival. <i>Molecular Carcinogenesis</i> , 2018, 57, 22-31.	2.7	5
30	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. <i>Nature Communications</i> , 2018, 9, 4774.	12.8	87
31	Systematic analyses of a novel lncRNA-associated signature as the prognostic biomarker for Hepatocellular Carcinoma. <i>Cancer Medicine</i> , 2018, 7, 3240-3256.	2.8	35
32	Hierarchical modeling of melanocortin 1 receptor variants with skin cancer risk. <i>Genetic Epidemiology</i> , 2018, 42, 571-586.	1.3	5
33	Meta-analysis of the association between sodium-glucose co-transporter 2 inhibitors and risk of skin cancer among patients with type 2 diabetes. <i>Diabetes, Obesity and Metabolism</i> , 2018, 20, 2919-2924.	4.4	10
34	Genome-wide association study in 176,678 Europeans reveals genetic loci for tanning response to sun exposure. <i>Nature Communications</i> , 2018, 9, 1684.	12.8	80
35	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
36	<i>MC1R</i> variants as melanoma risk factors independent of at-risk phenotypic characteristics: a pooled analysis from the M-SKIP project. <i>Cancer Management and Research</i> , 2018, Volume 10, 1143-1154.	1.9	57

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37	Indoor tanning use among white female students aged 18â€“30. <i>Journal of Dermatological Science</i> , 2017, 85, 253-256.	1.9	2
38	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	7.1	376
39	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.	5.1	11
40	Circulating concentrations of biomarkers and metabolites related to vitamin status, one-carbon and the kynurenine pathways in US, Nordic, Asian, and Australian populations. <i>American Journal of Clinical Nutrition</i> , 2017, 105, 1314-1326.	4.7	22
41	Genetic Variants in WNT2B and BTRC Predict Melanoma Survival. <i>Journal of Investigative Dermatology</i> , 2017, 137, 1749-1756.	0.7	5
42	Association study of genetic variation in <sc>DNA</sc> repair pathway genes and risk of basal cell carcinoma. <i>International Journal of Cancer</i> , 2017, 141, 952-957.	5.1	14
43	Genetic variants in the integrin signaling pathway genes predict cutaneous melanoma survival. <i>International Journal of Cancer</i> , 2017, 140, 1270-1279.	5.1	4
44	Genetic variants in the genes encoding rho GTPases and related regulators predict cutaneous melanomaâ€™specific survival. <i>International Journal of Cancer</i> , 2017, 141, 721-730.	5.1	8
45	A <i>PGC1 ² </i> genetic variant associated with nevus count and melanoma mortality. <i>International Journal of Cancer</i> , 2017, 141, 1066-1067.	5.1	5
46	Cancer risk in the EMPA-REG OUTCOME trial. Reply to Shaikh AMY [letter] and Kohler S, Lee J, George JT et al [letter]. <i>Diabetologia</i> , 2017, 60, 2538-2539.	6.3	2
47	Rare germline variants in known melanoma susceptibility genes in familial melanoma. <i>Human Molecular Genetics</i> , 2017, 26, 4886-4895.	2.9	37
48	SGLT2 inhibitors and risk of cancer in type 2 diabetes: a systematic review and meta-analysis of randomised controlled trials. <i>Diabetologia</i> , 2017, 60, 1862-1872.	6.3	134
49	Phosphodiesterase type 5 inhibitors and risk of melanoma: A meta-analysis. <i>Journal of the American Academy of Dermatology</i> , 2017, 77, 480-488.e9.	1.2	13
50	Height, height-related SNPs, and risk of non-melanoma skin cancer. <i>British Journal of Cancer</i> , 2017, 116, 134-140.	6.4	8
51	Genetic variants of PDGF signaling pathway genes predict cutaneous melanoma survival. <i>Oncotarget</i> , 2017, 8, 74595-74606.	1.8	3
52	Integrated analysis of competing endogenous RNA network revealing lncRNAs as potential prognostic biomarkers in human lung squamous cell carcinoma. <i>Oncotarget</i> , 2017, 8, 65997-66018.	1.8	39
53	Two-stage genome-wide association study identifies a novel susceptibility locus associated with melanoma. <i>Oncotarget</i> , 2017, 8, 17586-17592.	1.8	61
54	Statin use and non-melanoma skin cancer risk: a meta-analysis of randomized controlled trials and observational studies. <i>Oncotarget</i> , 2017, 8, 75411-75417.	1.8	16

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55	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.	3.3	19
56	A genome-wide investigation of food addiction. <i>Obesity</i> , 2016, 24, 1336-1341.	3.0	37
57	Pre-diagnostic leukocyte mitochondrial DNA copy number and skin cancer risk. <i>Carcinogenesis</i> , 2016, 37, 897-903.	2.8	12
58	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
59	Alcohol Intake is Associated with Increased Risk of Squamous Cell Carcinoma of the Skin: Three US Prospective Cohort Studies. <i>Nutrition and Cancer</i> , 2016, 68, 545-553.	2.0	18
60	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.	5.1	21
61	Nonsyndromic cleft lip with or without cleft palate and cancer: Evaluation of a possible common genetic background through the analysis of GWAS data. <i>Genomics Data</i> , 2016, 10, 22-29.	1.3	19
62	Trends in the diagnosis and clinical features of melanoma in situ (MIS) in US men and women: A prospective, observational study. <i>Journal of the American Academy of Dermatology</i> , 2016, 75, 698-705.	1.2	28
63	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
64	Male pattern baldness and risk of incident skin cancer in a cohort of men. <i>International Journal of Cancer</i> , 2016, 139, 2671-2678.	5.1	19
65	Type 2 diabetes mellitus and risk of cutaneous squamous cell carcinoma. <i>Journal of the American Academy of Dermatology</i> , 2016, 75, 831-834.	1.2	3
66	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.	1.2	17
67	Genome-wide association study identifies novel susceptibility loci for cutaneous squamous cell carcinoma. <i>Nature Communications</i> , 2016, 7, 12048.	12.8	117
68	A genome-wide analysis of gene-caffeine consumption interaction on basal cell carcinoma. <i>Carcinogenesis</i> , 2016, 37, bgw107.	2.8	3
69	Tissue-specific Co-expression of Long Non-coding and Coding RNAs Associated with Breast Cancer. <i>Scientific Reports</i> , 2016, 6, 32731.	3.3	35
70	Genome-wide association study identifies 14 novel risk alleles associated with basal cell carcinoma. <i>Nature Communications</i> , 2016, 7, 12510.	12.8	94
71	Association of Melanocortin-1 Receptor Variants with Pigmentary Traits in Humans: A Pooled Analysis from the M-Skip Project. <i>Journal of Investigative Dermatology</i> , 2016, 136, 1914-1917.	0.7	16
72	Response to Letter Regarding Article, "Association of Coffee Consumption With Total and Cause-Specific Mortality in 3 Large Prospective Cohorts". <i>Circulation</i> , 2016, 133, e660.	1.6	1

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73	A Genome-Wide Association Study of Cutaneous Squamous Cell Carcinoma among European Descendants. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 714-720.	2.5	34
74	Associations between smoking behavior-related alleles and the risk of melanoma. <i>Oncotarget</i> , 2016, 7, 47366-47375.	1.8	15
75	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
76	Pre-diagnostic leukocyte mitochondrial DNA copy number and risk of lung cancer. <i>Oncotarget</i> , 2016, 7, 27307-27312.	1.8	17
77	Pathway analysis of expression-related SNPs on genome-wide association study of basal cell carcinoma. <i>Oncotarget</i> , 2016, 7, 36885-36895.	1.8	6
78	Explicit Modeling of Ancestry Improves Polygenic Risk Scores and BLUP Prediction. <i>Genetic Epidemiology</i> , 2015, 39, 427-438.	1.3	30
79	Sleep duration and sleep-disordered breathing and the risk of melanoma among US women and men. <i>International Journal of Dermatology</i> , 2015, 54, e492-5.	1.0	10
80	Reply to finasteride and dutasteride may reduce melanoma risk. <i>Cancer</i> , 2015, 121, 3558-3559.	4.1	2
81	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
82	Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. <i>Nature Communications</i> , 2015, 6, 6916.	12.8	154
83	ZBTB7A Suppresses Melanoma Metastasis by Transcriptionally Repressing MCAM. <i>Molecular Cancer Research</i> , 2015, 13, 1206-1217.	3.4	44
84	Total and Cause-Specific Mortality of U.S. Nurses Working Rotating Night Shifts. <i>American Journal of Preventive Medicine</i> , 2015, 48, 241-252.	3.0	139
85	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , 2015, 47, 987-995.	21.4	218
86	Citrus Consumption and Risk of Cutaneous Malignant Melanoma. <i>Journal of Clinical Oncology</i> , 2015, 33, 2500-2508.	1.6	74
87	Plasma miRNAs as early biomarkers for detecting hepatocellular carcinoma. <i>International Journal of Cancer</i> , 2015, 137, 1679-1690.	5.1	188
88	Citrus consumption and risk of basal cell carcinoma and squamous cell carcinoma of the skin. <i>Carcinogenesis</i> , 2015, 36, 1162-1168.	2.8	39
89	Alcohol Intake and Risk of Incident Psoriatic Arthritis in Women. <i>Journal of Rheumatology</i> , 2015, 42, 835-840.	2.0	32
90	Association of Coffee Consumption With Total and Cause-Specific Mortality in 3 Large Prospective Cohorts. <i>Circulation</i> , 2015, 132, 2305-2315.	1.6	175

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91	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
92	Psoriasis, psoriatic arthritis and risk of gout in US men and women. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1495-1500.	0.9	67
93	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
94	Identification of a melanoma susceptibility locus and somatic mutation in <i>TET2</i> . <i>Carcinogenesis</i> , 2014, 35, 2097-2101.	2.8	41
95	Association between Cutaneous Nevi and Breast Cancer in the Nurses' Health Study: A Prospective Cohort Study. <i>PLoS Medicine</i> , 2014, 11, e1001659.	8.4	16
96	Sildenafil Use and Increased Risk of Incident Melanoma in US Men. <i>JAMA Internal Medicine</i> , 2014, 174, 964.	5.1	108
97	Hypertension, Antihypertensive Medication Use, and Risk of Psoriasis. <i>JAMA Dermatology</i> , 2014, 150, 957.	4.1	68
98	Long-term Ultraviolet Flux, Other Potential Risk Factors, and Skin Cancer Risk: A Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1080-1089.	2.5	122
99	Severe teenage acne and risk of endometriosis. <i>Human Reproduction</i> , 2014, 29, 2592-2599.	0.9	9
100	PGC-1 Coactivators Regulate MITF and the Tanning Response. <i>Molecular Cell</i> , 2013, 49, 145-157.	9.7	84
101	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
102	Has too much blame been placed on tanning beds for the rise in melanoma diagnosis?. <i>Expert Review of Dermatology</i> , 2013, 8, 135-143.	0.3	1
103	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
104	Joint Effect of Multiple Common SNPs Predicts Melanoma Susceptibility. <i>PLoS ONE</i> , 2013, 8, e85642.	2.5	40
105	Use of Tanning Beds and Incidence of Skin Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 1588-1593.	1.6	183
106	Pre-Diagnostic Plasma 25-Hydroxyvitamin D Levels and Risk of Non-Melanoma Skin Cancer in Women. <i>PLoS ONE</i> , 2012, 7, e35211.	2.5	43
107	A Germline Variant in the Interferon Regulatory Factor 4 Gene as a Novel Skin Cancer Risk Locus. <i>Cancer Research</i> , 2011, 71, 1533-1539.	0.9	45
108	Genome-wide association study identifies novel alleles associated with risk of cutaneous basal cell carcinoma and squamous cell carcinoma. <i>Human Molecular Genetics</i> , 2011, 20, 3718-3724.	2.9	92

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109	Genetic variation in DNA repair pathway genes and premenopausal breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2009, 115, 613-622.	2.5	46
110	A Prospective Study of Telomere Length and the Risk of Skin Cancer. <i>Journal of Investigative Dermatology</i> , 2009, 129, 415-421.	0.7	152
111	Genome-Wide Association Study of Tanning Phenotype in a Population of European Ancestry. <i>Journal of Investigative Dermatology</i> , 2009, 129, 2250-2257.	0.7	122
112	A Genome-Wide Association Study Identifies Novel Alleles Associated with Hair Color and Skin Pigmentation. <i>PLoS Genetics</i> , 2008, 4, e1000074.	3.5	439
113	Polymorphisms in O 6-methylguanine DNA methyltransferase and breast cancer risk. <i>Pharmacogenetics and Genomics</i> , 2006, 16, 469-474.	1.5	20
114	The p53 codon 72 polymorphism, sunburns, and risk of skin cancer in US caucasian women. <i>Molecular Carcinogenesis</i> , 2006, 45, 694-700.	2.7	41
115	Polymorphisms in the MTHFR and VDR genes and skin cancer risk. <i>Carcinogenesis</i> , 2006, 28, 390-397.	2.8	89
116	Risk factors for skin cancers: a nested caseâ€“control study within the Nursesâ€™ Health Study. <i>International Journal of Epidemiology</i> , 2006, 35, 1514-1521.	1.9	182
117	Genetic Variation in XPD, Sun Exposure, and Risk of Skin Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 1539-1544.	2.5	86
118	Genetic Variations in <i>XRCC2</i> and <i>XRCC3</i> Are Not Associated with Endometrial Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 330-331.	2.5	40
119	Polymorphisms in DNA Double-Strand Break Repair Genes and Skin Cancer Risk. <i>Cancer Research</i> , 2004, 64, 3009-3013.	0.9	97
120	Interaction between genetic variations in DNA repair genes and plasma folate on breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 520-4.	2.5	14
121	Polymorphisms in DNA double-strand break repair genes and breast cancer risk in the Nurses' Health Study. <i>Carcinogenesis</i> , 2003, 25, 189-195.	2.8	77
122	A prospective study of XRCC1 haplotypes and their interaction with plasma carotenoids on breast cancer risk. <i>Cancer Research</i> , 2003, 63, 8536-41.	0.9	69
123	No association between a stop codon polymorphism in RAD52 and breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002, 11, 1138-9.	2.5	3