

# Matthew H Law

## List of Publications by Citations

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**Version:** 2024-04-28

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

83  
papers

2,363  
citations

25  
h-index

47  
g-index

103  
ext. papers

3,196  
ext. citations

9.1  
avg, IF

4.56  
L-index

#	Paper	IF	Citations
83	A novel recurrent mutation in MITF predisposes to familial and sporadic melanoma. <i>Nature</i> , <b>2011</b> , 480, 99-103	50.4	335
82	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases: A Mendelian Randomization Study. <i>JAMA Oncology</i> , <b>2017</b> , 3, 636-651	13.4	236
81	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , <b>2015</b> , 47, 987-995	36.3	162
80	Common variants near ABCA1, AFAP1 and GMDS confer risk of primary open-angle glaucoma. <i>Nature Genetics</i> , <b>2014</b> , 46, 1120-1125	36.3	141
79	The effect on melanoma risk of genes previously associated with telomere length. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106,	9.7	97
78	A variant in FTO shows association with melanoma risk not due to BMI. <i>Nature Genetics</i> , <b>2013</b> , 45, 428-32, 432e1	36.3	95
77	Genome-wide association study of intraocular pressure uncovers new pathways to glaucoma. <i>Nature Genetics</i> , <b>2018</b> , 50, 1067-1071	36.3	86
76	Cardiometabolic effects of genetic upregulation of the interleukin 1 receptor antagonist: a Mendelian randomisation analysis. <i>Lancet Diabetes and Endocrinology</i> , <b>2015</b> , 3, 243-53	18.1	81
75	Multitrait analysis of glaucoma identifies new risk loci and enables polygenic prediction of disease susceptibility and progression. <i>Nature Genetics</i> , <b>2020</b> , 52, 160-166	36.3	78
74	Melanoma genetics: recent findings take us beyond well-traveled pathways. <i>Journal of Investigative Dermatology</i> , <b>2012</b> , 132, 1763-74	4.3	65
73	The role of phospholipases A2 in schizophrenia. <i>Molecular Psychiatry</i> , <b>2006</b> , 11, 547-56	15.1	54
72	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. <i>Nature Communications</i> , <b>2018</b> , 9, 4774	17.4	47
71	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , <b>2020</b> , 52, 494-504	36.3	39
70	ABCB1 (MDR1) polymorphisms and ovarian cancer progression and survival: a comprehensive analysis from the Ovarian Cancer Association Consortium and The Cancer Genome Atlas. <i>Gynecologic Oncology</i> , <b>2013</b> , 131, 8-14	4.9	39
69	Identification of a melanoma susceptibility locus and somatic mutation in TET2. <i>Carcinogenesis</i> , <b>2014</b> , 35, 2097-101	4.6	38
68	Meta-analysis combining new and existing data sets confirms that the TERT-CLPTM1L locus influences melanoma risk. <i>Journal of Investigative Dermatology</i> , <b>2012</b> , 132, 485-7	4.3	38
67	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. <i>Journal of Investigative Dermatology</i> , <b>2018</b> , 138, 2617-2624	4.3	36

66	A common intronic variant of PARP1 confers melanoma risk and mediates melanocyte growth via regulation of MITF. <i>Nature Genetics</i> , <b>2017</b> , 49, 1326-1335	36.3	36
65	A Pilot Randomized Controlled Trial of the Feasibility, Acceptability, and Impact of Giving Information on Personalized Genomic Risk of Melanoma to the Public. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 212-221	4	33
64	Cell-type-specific eQTL of primary melanocytes facilitates identification of melanoma susceptibility genes. <i>Genome Research</i> , <b>2018</b> , 28, 1621-1635	9.7	33
63	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , <b>2020</b> , 11, 3353	17.4	32
62	Vitamin D and overall cancer risk and cancer mortality: a Mendelian randomization study. <i>Human Molecular Genetics</i> , <b>2018</b> , 27, 4315-4322	5.6	32
61	Genome-Wide Association Shows that Pigmentation Genes Play a Role in Skin Aging. <i>Journal of Investigative Dermatology</i> , <b>2017</b> , 137, 1887-1894	4.3	30
60	A study of circulating gliadin antibodies in schizophrenia among a Chinese population. <i>Schizophrenia Bulletin</i> , <b>2012</b> , 38, 514-8	1.3	29
59	Height and overall cancer risk and mortality: evidence from a Mendelian randomisation study on 310,000 UK Biobank participants. <i>British Journal of Cancer</i> , <b>2018</b> , 118, 1262-1267	8.7	27
58	Analysis combining correlated glaucoma traits identifies five new risk loci for open-angle glaucoma. <i>Scientific Reports</i> , <b>2018</b> , 8, 3124	4.9	25
57	Massively parallel reporter assays of melanoma risk variants identify MX2 as a gene promoting melanoma. <i>Nature Communications</i> , <b>2020</b> , 11, 2718	17.4	24
56	Combining common genetic variants and non-genetic risk factors to predict risk of cutaneous melanoma. <i>Human Molecular Genetics</i> , <b>2018</b> , 27, 4145-4156	5.6	21
55	Combined analysis of keratinocyte cancers identifies novel genome-wide loci. <i>Human Molecular Genetics</i> , <b>2019</b> , 28, 3148-3160	5.6	20
54	Genetic association of the AKT1 gene with schizophrenia in a British population. <i>Psychiatric Genetics</i> , <b>2010</b> , 20, 118-22	2.9	20
53	Survival outcomes in patients with multiple primary melanomas. <i>Journal of the European Academy of Dermatology and Venereology</i> , <b>2015</b> , 29, 2120-7	4.6	18
52	The emerging field of polygenic risk scores and perspective for use in clinical care. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, R165-R176	5.6	16
51	Gastroesophageal reflux GWAS identifies risk loci that also associate with subsequent severe esophageal diseases. <i>Nature Communications</i> , <b>2019</b> , 10, 4219	17.4	15
50	Association between coffee consumption and overall risk of being diagnosed with or dying from cancer among >300 000 UK Biobank participants in a large-scale Mendelian randomization study. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 1447-1456	7.8	15
49	Accurate Imputation-Based Screening of Gln368Ter Myocilin Variant in Primary Open-Angle Glaucoma <b>2015</b> , 56, 5087-93		15

48	12-hydroxyeicosatetraenoic acid is associated with variability in aspirin-induced platelet inhibition. <i>Journal of Inflammation</i> , <b>2014</b> , 11, 33	6.7	15
47	Effect of increased body mass index on risk of diagnosis or death from cancer. <i>British Journal of Cancer</i> , <b>2019</b> , 120, 565-570	8.7	13
46	The melanoma genomics managing your risk study: A protocol for a randomized controlled trial evaluating the impact of personal genomic risk information on skin cancer prevention behaviors. <i>Contemporary Clinical Trials</i> , <b>2018</b> , 70, 106-116	2.3	13
45	Rare variants analysis of cutaneous malignant melanoma genes in Parkinson's disease. <i>Neurobiology of Aging</i> , <b>2016</b> , 48, 222.e1-222.e7	5.6	12
44	A risk prediction model for the development of subsequent primary melanoma in a population-based cohort. <i>British Journal of Dermatology</i> , <b>2020</b> , 182, 1148-1157	4	12
43	Overlapping genetic architecture between Parkinson disease and melanoma. <i>Acta Neuropathologica</i> , <b>2020</b> , 139, 347-364	14.3	12
42	A comprehensive re-assessment of the association between vitamin D and cancer susceptibility using Mendelian randomization. <i>Nature Communications</i> , <b>2021</b> , 12, 246	17.4	12
41	Polyunsaturated fatty acids and risk of melanoma: A Mendelian randomisation analysis. <i>International Journal of Cancer</i> , <b>2018</b> , 143, 508-514	7.5	11
40	The TGM2 gene is associated with schizophrenia in a British population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2009</b> , 150B, 335-40	3.5	11
39	Chemical cleavage of mismatch (CCM) to locate base mismatches in heteroduplex DNA. <i>Nature Protocols</i> , <b>2006</b> , 1, 2297-304	18.8	11
38	Neural crest-derived tumor neuroblastoma and melanoma share 1p13.2 as susceptibility locus that shows a long-range interaction with the SLC16A1 gene. <i>Carcinogenesis</i> , <b>2020</b> , 41, 284-295	4.6	11
37	Mendelian Randomization Study for Genetically Predicted Polyunsaturated Fatty Acids Levels on Overall Cancer Risk and Mortality. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2019</b> , 28, 1015-1023	4	10
36	Body mass index and height and risk of cutaneous melanoma: Mendelian randomization analyses. <i>International Journal of Epidemiology</i> , <b>2020</b> , 49, 1236-1245	7.8	9
35	Is there a causal relationship between vitamin D and melanoma risk? A Mendelian randomization study. <i>British Journal of Dermatology</i> , <b>2020</b> , 182, 97-103	4	8
34	Vitamin D Pathway Gene Polymorphisms and Keratinocyte Cancers: A Nested Case-Control Study and Meta-Analysis. <i>Anticancer Research</i> , <b>2016</b> , 36, 2145-52	2.3	7
33	PARP1 polymorphisms play opposing roles in melanoma occurrence and survival. <i>International Journal of Cancer</i> , <b>2015</b> , 136, 2488-9	7.5	6
32	Association between functional polymorphisms in genes involved in the MAPK signaling pathways and cutaneous melanoma risk. <i>Carcinogenesis</i> , <b>2013</b> , 34, 885-92	4.6	6
31	Global Biobank Meta-analysis Initiative: powering genetic discovery across human diseases		6

30	Variants of EVER1 and EVER2 (TMC6 and TMC8) and human papillomavirus status in patients with mucosal squamous cell carcinoma of the head and neck. <i>Cancer Causes and Control</i> , <b>2016</b> , 27, 809-15	2.8	6
29	Association between putative functional variants in the PSMB9 gene and risk of melanoma--re-analysis of published melanoma genome-wide association studies. <i>Pigment Cell and Melanoma Research</i> , <b>2013</b> , 26, 392-401	4.5	4
28	The functional significance of the TGM2 gene in schizophrenia: a correlation of SNPs and circulating IL-2 levels. <i>Journal of Neuroimmunology</i> , <b>2011</b> , 232, 5-7	3.5	4
27	No association between the PPARC gene and schizophrenia in a British population. <i>Prostaglandins Leukotrienes and Essential Fatty Acids</i> , <b>2009</b> , 81, 273-7	2.8	4
26	273 A study of gluten antibody levels in serum among patients with schizophrenia. <i>Schizophrenia Research</i> , <b>2008</b> , 98, 145	3.6	4
25	Genome-wide analyses in 1,987,836 participants identify 39 genetic loci associated with sleep apnoea		4
24	Does polygenic risk influence associations between sun exposure and melanoma? A prospective cohort analysis. <i>British Journal of Dermatology</i> , <b>2020</b> , 183, 303-310	4	4
23	Genomic Risk Score for Melanoma in a Prospective Study of Older Individuals. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 1379-1385	9.7	4
22	Polygenic Risk Scores Allow Risk Stratification for Keratinocyte Cancer in Organ-Transplant Recipients. <i>Journal of Investigative Dermatology</i> , <b>2021</b> , 141, 325-333.e6	4.3	4
21	No association observed between schizophrenia and non-HLA coeliac disease genes: integration with the initial MYO9B association with coeliac disease. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , <b>2011</b> , 156B, 709-19	3.5	3
20	Multiplex melanoma families are enriched for polygenic risk. <i>Human Molecular Genetics</i> , <b>2020</b> , 29, 2976-2985	2.8	3
19	Polyunsaturated Fatty Acid Levels and the Risk of Keratinocyte Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 1591-1598	4	3
18	Risk factors for melanoma by anatomical site: an evaluation of aetiological heterogeneity. <i>British Journal of Dermatology</i> , <b>2021</b> , 184, 1085-1093	4	3
17	Massively parallel reporter assays combined with cell-type specific eQTL informed multiple melanoma loci and identified a pleiotropic function of HIV-1 restriction gene, MX2, in melanoma promotion		2
16	Multitrait genetic association analysis identifies 50 new risk loci for gastro-oesophageal reflux, seven new loci for Barrett's oesophagus and provides insights into clinical heterogeneity in reflux diagnosis. <i>Gut</i> , <b>2021</b> ,	19.2	2
15	Impact of personal genomic risk information on melanoma prevention behaviors and psychological outcomes: a randomized controlled trial. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 2394-2403	8.1	2
14	Cell-type-specific meQTLs extend melanoma GWAS annotation beyond eQTLs and inform melanocyte gene-regulatory mechanisms. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 1631-1646	11	2
13	The Melanoma Genomics Managing Your Risk Study randomised controlled trial: statistical analysis plan. <i>Trials</i> , <b>2020</b> , 21, 594	2.8	1

12	Germline variants are associated with increased primary melanoma tumor thickness at diagnosis. <i>Human Molecular Genetics</i> , <b>2021</b> , 29, 3578-3587	5.6	1
11	Assessment of Polygenic Architecture and Risk Prediction based on Common Variants Across Fourteen Cancers		1
10	Polygenic Risk Scores Stratify Keratinocyte Cancer Risk among Solid Organ Transplant Recipients with Chronic Immunosuppression in a High Ultraviolet Radiation Environment. <i>Journal of Investigative Dermatology</i> , <b>2021</b> , 141, 2866-2875.e2	4.3	1
9	Independent evaluation of melanoma polygenic risk scores in UK and Australian prospective cohorts.. <i>British Journal of Dermatology</i> , <b>2021</b> ,	4	1
8	Is Genetic Risk for Sleep Apnea Causally Linked With Glaucoma Susceptibility? <b>2022</b> , 63, 25		0
7	A UVB-responsive common variant at chromosome band 7p21.1 confers tanning response and melanoma risk via regulation of the aryl hydrocarbon receptor, AHR. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 1611-1630	11	0
6	Assessment of melanoma candidate genes in a meta-analysis of 16534 melanoma cases. <i>Journal of the European Academy of Dermatology and Venereology</i> , <b>2019</b> , 33, e369-e370	4.6	
5	Inherited Contributions to Melanoma Risk <b>2019</b> , 225-248		
4	Genetically determined risk of keratinocyte carcinoma and risk of other cancers. <i>International Journal of Epidemiology</i> , <b>2021</b> , 50, 1316-1324	7.8	
3	A pilot randomised controlled trial examining the feasibility, acceptability and impact of giving information on personalised genomic risk of melanoma to the public, for motivating preventive behaviours.. <i>Journal of Clinical Oncology</i> , <b>2016</b> , 34, 1556-1556	2.2	
2	Inherited Contributions to Melanoma Risk <b>2018</b> , 1-23		
1	Making sense of different measures of skin ageing. <i>British Journal of Dermatology</i> , <b>2020</b> , 182, 1323-1324		